# Full wwPDB X-ray Structure Validation Report ⓘ

Mar 12, 2026 – 07:31 AM UTC

| | | |
|---|---|---|
| PDB ID | : | 4V7Y / pdb_00004v7y |
| Title | : | Structure of the Thermus thermophilus 70S ribosome complexed with azithromycin. |
| Authors | : | Bulkley, D.P.; Innis, C.A.; Blaha, G.; Steitz, T.A. |
| Deposited on | : | 2010-08-18 |
| Resolution | : | 3.00 Å(reported) |

This is a Full wwPDB X-ray Structure Validation Report for a publicly released PDB entry.

We welcome your comments at *validation@mail.wwpdb.org*
A user guide is available at
https://www.wwpdb.org/validation/2017/XrayValidationReportHelp
with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at
http://www.wwpdb.org/validation/2017/FAQs#types.

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The following versions of software and data (see references ⓘ) were used in the production of this report:

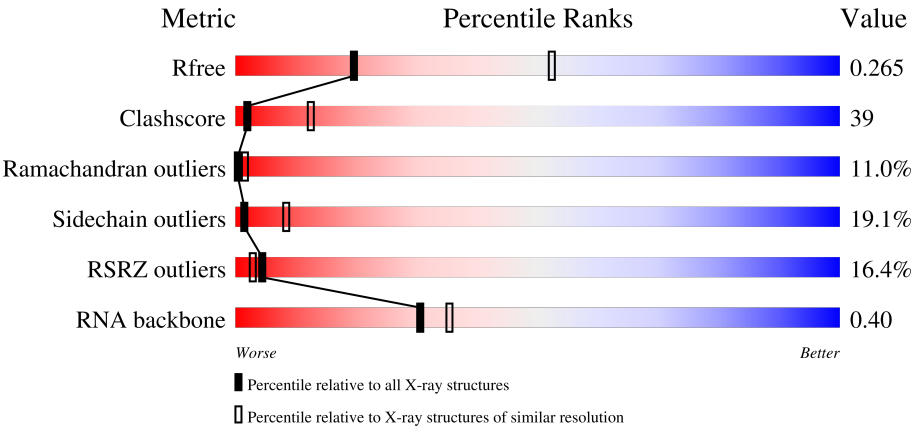
| | | |
|---|---|---|
| MolProbity | : | 4-5-2 with Phenix2.0 |
| Mogul | : | 2022.3.0, CSD as543be (2022) |
| Xtriage (Phenix) | : | 2.0 |
| EDS | : | 3.0 |
| Buster-report | : | wwPDB partial adaption of 1.1.7 (2018) |
| Percentile statistics | : | 20250101.v01 (using entries in the PDB archive January 1st 2025) |
| CCP4 | : | 9.0.010 (Gargrove) |
| Density-Fitness | : | 1.0.12 |
| Ideal geometry (proteins) | : | Engh & Huber (2001) |
| Ideal geometry (DNA, RNA) | : | Parkinson et al. (1996) |
| Validation Pipeline (wwPDB-VP) | : | 2.49 |

# 1 Overall quality at a glance (i)

The following experimental techniques were used to determine the structure:
*X-RAY DIFFRACTION*

The reported resolution of this entry is 3.00 Å.

Percentile scores (ranging between 0-100) for global validation metrics of the entry are shown in the following graphic. The table shows the number of entries on which the scores are based.




| Metric | Whole archive (#Entries) | Similar resolution (#Entries, resolution range(Å)) |
|---|---|---|
| $R_{free}$ | 180053 | 2672 (3.00-3.00) |
| Clashscore | 190562 | 2977 (3.00-3.00) |
| Ramachandran outliers | 187476 | 2877 (3.00-3.00) |
| Sidechain outliers | 187428 | 2880 (3.00-3.00) |
| RSRZ outliers | 180081 | 2671 (3.00-3.00) |
| RNA backbone | 3983 | 1109 (3.20-2.80) |

The table below summarises the geometric issues observed across the polymeric chains and their fit to the electron density. The red, orange, yellow and green segments of the lower bar indicate the fraction of residues that contain outliers for >=3, 2, 1 and 0 types of geometric quality criteria respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions <=5% The upper red bar (where present) indicates the fraction of residues that have poor fit to the electron density. The numeric value is given above the bar.

| Mol | Chain | Length | Quality of chain |
|---|---|---|---|
| 1 | AA | 1522 | 11% 27% 57% 15% · |
| 1 | CA | 1522 | 9% 27% 57% 15% · |
| 2 | AB | 256 | 19% 34% 43% 14% · 8% |
| 2 | CB | 256 | 19% 33% 44% 13% · 8% |

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| Mol | Chain | Length | Quality of chain |
|---|---|---|---|
| 3 | AC | 239 | 13% 38% 39% 8% 13% |
| 3 | CC | 239 | 20% 38% 40% 8% 13% |
| 4 | AD | 209 | 22% 31% 50% 17% |
| 4 | CD | 209 | 22% 32% 49% 17% |
| 5 | AE | 162 | 17% 30% 48% 15% 7% |
| 5 | CE | 162 | 14% 30% 48% 15% 7% |
| 6 | AF | 101 | 14% 29% 56% 15% |
| 6 | CF | 101 | 17% 28% 56% 16% |
| 7 | AG | 156 | 29% 52% 45% |
| 7 | CG | 156 | 25% 51% 46% |
| 8 | AH | 138 | 14% 36% 53% 9% |
| 8 | CH | 138 | 17% 38% 53% 8% |
| 9 | AI | 128 | 35% 37% 52% 10% |
| 9 | CI | 128 | 44% 34% 55% 10% |
| 10 | AJ | 105 | 35% 36% 51% 7% 6% |
| 10 | CJ | 105 | 35% 36% 51% 7% 6% |
| 11 | AK | 129 | 23% 39% 46% 7% 8% |
| 11 | CK | 129 | 19% 38% 44% 9% 8% |
| 12 | AL | 135 | 24% 33% 49% 11% 7% |
| 12 | CL | 135 | 26% 34% 49% 10% 7% |
| 13 | AM | 126 | 32% 42% 42% 6% 9% |
| 13 | CM | 126 | 39% 40% 45% 6% 9% |
| 14 | AN | 61 | 36% 46% 46% 7% |
| 14 | CN | 61 | 34% 46% 46% 7% |
| 15 | AO | 89 | 21% 39% 47% 12% |

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| Mol | Chain | Length | Quality of chain |
|---|---|---|---|
| 15 | CO | 89 | 13% 40% 46% 12% |
| 16 | AP | 88 | 34% 20% 60% 11% 5% |
| 16 | CP | 88 | 31% 24% 56% 14% 5% |
| 17 | AQ | 105 | 18% 48% 39% 9% 5% |
| 17 | CQ | 105 | 15% 49% 41% 5% 5% |
| 18 | AR | 88 | 10% 24% 40% 15% 20% |
| 18 | CR | 88 | 6% 26% 36% 16% 20% |
| 19 | AS | 93 | 32% 37% 38% 11% 15% |
| 19 | CS | 93 | 27% 41% 35% 9% 15% |
| 20 | AT | 106 | 31% 28% 52% 12% 7% |
| 20 | CT | 106 | 25% 23% 58% 12% 7% |
| 21 | AU | 27 | 81% 48% 41% 7% |
| 21 | CU | 27 | 74% 48% 41% 7% |
| 22 | B0 | 85 | 18% 33% 51% 14% |
| 22 | D0 | 85 | 16% 35% 48% 13% |
| 23 | B1 | 98 | 32% 19% 33% 32% 7% 9% |
| 23 | D1 | 98 | 23% 18% 37% 28% 8% 9% |
| 24 | B2 | 72 | 43% 8% 32% 25% 6% 29% |
| 24 | D2 | 72 | 33% 8% 38% 21% 29% |
| 25 | B3 | 60 | 17% 43% 43% 13% |
| 25 | D3 | 60 | 3% 37% 53% 10% |
| 26 | B4 | 71 | 18% 17% 13% 15% 55% |
| 26 | D4 | 71 | 14% 14% 15% 15% 55% |
| 27 | B5 | 60 | 18% 25% 40% 27% 7% |
| 27 | D5 | 60 | 12% 28% 42% 23% 5% |

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| Mol | Chain | Length | Quality of chain |
|---|---|---|---|
| 28 | B6 | 54 | 31% 6% 28% 44% 6% 17% |
| 28 | D6 | 54 | 31% 6% 28% 44% 6% 17% |
| 29 | B7 | 49 | 12% 37% 45% 16% |
| 29 | D7 | 49 | 12% 39% 45% 14% |
| 30 | B8 | 65 | 32% 22% 40% 31% 6% |
| 30 | D8 | 65 | 31% 25% 40% 29% 5% |
| 31 | BA | 2787 | 6% 32% 46% 20% |
| 31 | DA | 2787 | 4% 28% 49% 20% |
| 32 | BB | 122 | 28% 24% 51% 23% |
| 32 | DB | 122 | 20% 22% 52% 23% |
| 33 | BD | 276 | 9% 25% 46% 24% |
| 33 | DD | 276 | 7% 29% 44% 22% |
| 34 | BE | 206 | 12% 25% 47% 23% |
| 34 | DE | 206 | 13% 25% 49% 22% |
| 35 | BF | 210 | 11% 29% 50% 17% |
| 35 | DF | 210 | 10% 30% 47% 19% |
| 36 | BG | 182 | 40% 26% 54% 18% |
| 36 | DG | 182 | 36% 27% 52% 19% |
| 37 | BH | 180 | 39% 30% 36% 17% 6% 11% |
| 37 | DH | 180 | 12% 28% 40% 14% 7% 11% |
| 38 | BI | 148 | 33% 22% 51% 21% |
| 38 | DI | 148 | 28% 20% 54% 20% |
| 39 | BN | 140 | 24% 22% 49% 24% |
| 39 | DN | 140 | 12% 25% 46% 25% |
| 40 | BO | 122 | 2% 29% 54% 16% |

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| Mol | Chain | Length | Quality of chain |
|---|---|---|---|
| 40 | DO | 122 | 7% 34% 48% 16% |
| 41 | BP | 150 | 31% 13% 31% 39% 14% |
| 41 | DP | 150 | 25% 12% 36% 34% 15% |
| 42 | BQ | 141 | 21% 26% 48% 19% |
| 42 | DQ | 141 | 12% 23% 52% 19% |
| 43 | BR | 118 | 11% 25% 53% 19% |
| 43 | DR | 118 | 14% 21% 53% 22% |
| 44 | BS | 112 | 43% 20% 35% 28% 6% 12% |
| 44 | DS | 112 | 28% 19% 36% 28% 6% 12% |
| 45 | BT | 146 | 21% 16% 41% 28% 5% 10% |
| 45 | DT | 146 | 20% 16% 42% 26% 6% 10% |
| 46 | BU | 118 | 19% 32% 45% 21% |
| 46 | DU | 118 | 13% 29% 44% 24% |
| 47 | BV | 101 | 50% 12% 37% 44% 8% |
| 47 | DV | 101 | 21% 13% 37% 44% 7% |
| 48 | BW | 113 | 6% 37% 39% 21% |
| 48 | DW | 113 | 4% 31% 47% 19% |
| 49 | BX | 96 | 30% 15% 41% 31% 10% |
| 49 | DX | 96 | 20% 15% 42% 29% 11% |
| 50 | BY | 110 | 54% 11% 36% 38% 6% 8% |
| 50 | DY | 110 | 38% 9% 38% 38% 6% 8% |
| 51 | BZ | 206 | 26% 23% 44% 17% 14% |
| 51 | DZ | 206 | 19% 24% 44% 17% 14% |

The following table lists non-polymeric compounds, carbohydrate monomers and non-standard residues in protein, DNA, RNA chains that are outliers for geometric or electron-density-fit criteria:

| Mol | Type | Chain | Res | Chirality | Geometry | Clashes | Electron density |
|---|---|---|---|---|---|---|---|
| 52 | MG | AA | 1622 | - | - | - | X |
| 52 | MG | AA | 1646 | - | - | - | X |
| 52 | MG | CA | 1633 | - | - | - | X |
| 52 | MG | CA | 1641 | - | - | - | X |
| 52 | MG | DA | 3125 | - | - | - | X |
| 52 | MG | DA | 3179 | - | - | - | X |
| 52 | MG | DA | 3216 | - | - | - | X |
| 52 | MG | DA | 3280 | - | - | - | X |
| 52 | MG | DA | 3289 | - | - | - | X |
| 52 | MG | DA | 3302 | - | - | - | X |
| 54 | K | DA | 3310 | - | - | - | X |

# 2 Entry composition

There are 55 unique types of molecules in this entry. The entry contains 278000 atoms, of which 0 are hydrogens and 0 are deuteriums.

In the tables below, the ZeroOcc column contains the number of atoms modelled with zero occupancy, the AltConf column contains the number of residues with at least one atom in alternate conformation and the Trace column contains the number of residues modelled with at most 2 atoms.

- Molecule 1 is a RNA chain called 16S rRNA.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 1 | AA | 1504 | Total | C | N | O | P | 0 | 0 | 0 |
| | | | 32329 | 14390 | 5992 | 10444 | 1503 | | | |
| 1 | CA | 1504 | Total | C | N | O | P | 0 | 0 | 0 |
| | | | 32329 | 14390 | 5992 | 10444 | 1503 | | | |

- Molecule 2 is a protein called 30S ribosomal protein S2.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 2 | AB | 235 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1901 | 1213 | 342 | 341 | 5 | | | |
| 2 | CB | 235 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1901 | 1213 | 342 | 341 | 5 | | | |

- Molecule 3 is a protein called 30S ribosomal protein S3.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 3 | AC | 207 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1613 | 1016 | 315 | 281 | 1 | | | |
| 3 | CC | 207 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1613 | 1016 | 315 | 281 | 1 | | | |

- Molecule 4 is a protein called 30S ribosomal protein S4.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 4 | AD | 208 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1703 | 1066 | 339 | 291 | 7 | | | |
| 4 | CD | 208 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1703 | 1066 | 339 | 291 | 7 | | | |

- Molecule 5 is a protein called 30S ribosomal protein S5.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 5 | AE | 151 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1147 | 724 | 218 | 201 | 4 | | | |
| 5 | CE | 151 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1147 | 724 | 218 | 201 | 4 | | | |

- Molecule 6 is a protein called 30S ribosomal protein S6.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 6 | AF | 101 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 843 | 531 | 155 | 154 | 3 | | | |
| 6 | CF | 101 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 843 | 531 | 155 | 154 | 3 | | | |

- Molecule 7 is a protein called 30S ribosomal protein S7.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 7 | AG | 155 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1257 | 781 | 252 | 218 | 6 | | | |
| 7 | CG | 155 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1257 | 781 | 252 | 218 | 6 | | | |

- Molecule 8 is a protein called 30S ribosomal protein S8.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 8 | AH | 138 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1116 | 705 | 215 | 193 | 3 | | | |
| 8 | CH | 138 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1116 | 705 | 215 | 193 | 3 | | | |

- Molecule 9 is a protein called 30S ribosomal protein S9.

| Mol | Chain | Residues | Atoms | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|
| 9 | AI | 127 | Total | C | N | O | 0 | 0 | 0 |
| | | | 1011 | 639 | 198 | 174 | | | |
| 9 | CI | 127 | Total | C | N | O | 0 | 0 | 0 |
| | | | 1011 | 639 | 198 | 174 | | | |

There are 2 discrepancies between the modelled and reference sequences:

| Chain | Residue | Modelled | Actual | Comment | Reference |
|---|---|---|---|---|---|
| AI | 58 | ARG | HIS | conflict | UNP P80374 |
| CI | 58 | ARG | HIS | conflict | UNP P80374 |

- Molecule 10 is a protein called 30S ribosomal protein S10.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 10 | AJ | 99 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 795 | 499 | 157 | 138 | 1 | | | |
| 10 | CJ | 99 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 795 | 499 | 157 | 138 | 1 | | | |

- Molecule 11 is a protein called 30S ribosomal protein S11.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 11 | AK | 119 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 885 | 549 | 168 | 165 | 3 | | | |
| 11 | CK | 119 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 885 | 549 | 168 | 165 | 3 | | | |

- Molecule 12 is a protein called 30S ribosomal protein S12.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 12 | AL | 125 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 971 | 611 | 196 | 163 | 1 | | | |
| 12 | CL | 125 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 971 | 611 | 196 | 163 | 1 | | | |

There are 6 discrepancies between the modelled and reference sequences:

| Chain | Residue | Modelled | Actual | Comment | Reference |
|---|---|---|---|---|---|
| AL | 2 | VAL | - | insertion | UNP Q5SHN3 |
| AL | 3 | ALA | - | insertion | UNP Q5SHN3 |
| AL | 4 | LEU | - | insertion | UNP Q5SHN3 |
| CL | 2 | VAL | - | insertion | UNP Q5SHN3 |
| CL | 3 | ALA | - | insertion | UNP Q5SHN3 |
| CL | 4 | LEU | - | insertion | UNP Q5SHN3 |

- Molecule 13 is a protein called 30S ribosomal protein S13.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 13 | AM | 115 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 921 | 569 | 190 | 160 | 2 | | | |
| 13 | CM | 115 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 921 | 569 | 190 | 160 | 2 | | | |

- Molecule 14 is a protein called 30S ribosomal protein S14.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 14 | AN | 60 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 492 | 312 | 104 | 72 | 4 | | | |
| 14 | CN | 60 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 492 | 312 | 104 | 72 | 4 | | | |

- Molecule 15 is a protein called 30S ribosomal protein S15.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 15 | AO | 88 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 734 | 459 | 147 | 126 | 2 | | | |
| 15 | CO | 88 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 734 | 459 | 147 | 126 | 2 | | | |

- Molecule 16 is a protein called 30S ribosomal protein S16.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 16 | AP | 84 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 701 | 443 | 140 | 117 | 1 | | | |
| 16 | CP | 84 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 701 | 443 | 140 | 117 | 1 | | | |

- Molecule 17 is a protein called 30S ribosomal protein S17.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 17 | AQ | 100 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 824 | 528 | 152 | 142 | 2 | | | |
| 17 | CQ | 100 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 824 | 528 | 152 | 142 | 2 | | | |

- Molecule 18 is a protein called 30S ribosomal protein S18.

| Mol | Chain | Residues | Atoms | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|
| 18 | AR | 70 | Total | C | N | O | 0 | 0 | 0 |
| | | | 574 | 367 | 112 | 95 | | | |
| 18 | CR | 70 | Total | C | N | O | 0 | 0 | 0 |
| | | | 574 | 367 | 112 | 95 | | | |

- Molecule 19 is a protein called 30S ribosomal protein S19.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 19 | AS | 79 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 630 | 403 | 115 | 110 | 2 | | | |

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| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 19 | CS | 79 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 630 | 403 | 115 | 110 | 2 | | | |

- Molecule 20 is a protein called 30S ribosomal protein S20.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 20 | AT | 99 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 763 | 470 | 162 | 129 | 2 | | | |
| 20 | CT | 99 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 763 | 470 | 162 | 129 | 2 | | | |

- Molecule 21 is a protein called 30S ribosomal protein Thx.

| Mol | Chain | Residues | Atoms | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|
| 21 | AU | 25 | Total | C | N | O | 0 | 0 | 1 |
| | | | 209 | 128 | 51 | 30 | | | |
| 21 | CU | 25 | Total | C | N | O | 0 | 0 | 1 |
| | | | 209 | 128 | 51 | 30 | | | |

- Molecule 22 is a protein called 50S ribosomal protein L27.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 22 | B0 | 85 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 650 | 401 | 137 | 111 | 1 | | | |
| 22 | D0 | 85 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 650 | 401 | 137 | 111 | 1 | | | |

- Molecule 23 is a protein called 50S ribosomal protein L28.

| Mol | Chain | Residues | Atoms | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|
| 23 | B1 | 89 | Total | C | N | O | 0 | 0 | 1 |
| | | | 693 | 435 | 140 | 118 | | | |
| 23 | D1 | 89 | Total | C | N | O | 0 | 0 | 1 |
| | | | 693 | 435 | 140 | 118 | | | |

- Molecule 24 is a protein called 50S ribosomal protein L29.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 24 | B2 | 51 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 421 | 263 | 85 | 72 | 1 | | | |
| 24 | D2 | 51 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 421 | 263 | 85 | 72 | 1 | | | |

- Molecule 25 is a protein called 50S ribosomal protein L30.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 25 | B3 | 60 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 468 | 298 | 91 | 78 | 1 | | | |
| 25 | D3 | 60 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 468 | 298 | 91 | 78 | 1 | | | |

- Molecule 26 is a protein called 50S ribosomal protein L31.

| Mol | Chain | Residues | Atoms | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|
| 26 | B4 | 32 | Total | C | N | O | 0 | 0 | 0 |
| | | | 157 | 93 | 32 | 32 | | | |
| 26 | D4 | 32 | Total | C | N | O | 0 | 0 | 0 |
| | | | 157 | 93 | 32 | 32 | | | |

- Molecule 27 is a protein called 50S ribosomal protein L32.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 27 | B5 | 59 | Total | C | N | O | S | 9 | 0 | 0 |
| | | | 459 | 288 | 90 | 76 | 5 | | | |
| 27 | D5 | 59 | Total | C | N | O | S | 9 | 0 | 0 |
| | | | 459 | 288 | 90 | 76 | 5 | | | |

- Molecule 28 is a protein called 50S ribosomal protein L33.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 28 | B6 | 45 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 381 | 235 | 78 | 64 | 4 | | | |
| 28 | D6 | 45 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 381 | 235 | 78 | 64 | 4 | | | |

- Molecule 29 is a protein called 50S ribosomal protein L34.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 29 | B7 | 49 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 419 | 257 | 105 | 55 | 2 | | | |
| 29 | D7 | 49 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 419 | 257 | 105 | 55 | 2 | | | |

- Molecule 30 is a protein called 50S ribosomal protein L35.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 30 | B8 | 64 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 508 | 326 | 102 | 78 | 2 | | | |
| 30 | D8 | 64 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 508 | 326 | 102 | 78 | 2 | | | |

- Molecule 31 is a RNA chain called 23S ribosomal RNA.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 31 | BA | 2725 | Total | C | N | O | P | 0 | 0 | 0 |
| | | | 58698 | 26124 | 10986 | 18864 | 2724 | | | |
| 31 | DA | 2725 | Total | C | N | O | P | 0 | 0 | 0 |
| | | | 58698 | 26124 | 10986 | 18864 | 2724 | | | |

- Molecule 32 is a RNA chain called 5S ribosomal RNA.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 32 | BB | 119 | Total | C | N | O | P | 0 | 0 | 0 |
| | | | 2551 | 1136 | 471 | 826 | 118 | | | |
| 32 | DB | 119 | Total | C | N | O | P | 0 | 0 | 0 |
| | | | 2551 | 1136 | 471 | 826 | 118 | | | |

- Molecule 33 is a protein called 50S ribosomal protein L2.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 33 | BD | 272 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 2105 | 1329 | 417 | 356 | 3 | | | |
| 33 | DD | 272 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 2105 | 1329 | 417 | 356 | 3 | | | |

- Molecule 34 is a protein called 50S ribosomal protein L3.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 34 | BE | 205 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1564 | 988 | 300 | 270 | 6 | | | |
| 34 | DE | 205 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1564 | 988 | 300 | 270 | 6 | | | |

- Molecule 35 is a protein called 50S ribosomal protein L4.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 35 | BF | 208 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1624 | 1035 | 304 | 282 | 3 | | | |

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| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 35 | DF | 208 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1624 | 1035 | 304 | 282 | 3 | | | |

- Molecule 36 is a protein called 50S ribosomal protein L5.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 36 | BG | 181 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1474 | 942 | 268 | 260 | 4 | | | |
| 36 | DG | 181 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1474 | 942 | 268 | 260 | 4 | | | |

- Molecule 37 is a protein called 50S ribosomal protein L6.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 37 | BH | 160 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1223 | 773 | 229 | 220 | 1 | | | |
| 37 | DH | 160 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1223 | 773 | 229 | 220 | 1 | | | |

- Molecule 38 is a protein called 50S ribosomal protein L9.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 38 | BI | 146 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1132 | 723 | 201 | 207 | 1 | | | |
| 38 | DI | 146 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1132 | 723 | 201 | 207 | 1 | | | |

- Molecule 39 is a protein called 50S ribosomal protein L13.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 39 | BN | 139 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1105 | 712 | 207 | 182 | 4 | | | |
| 39 | DN | 139 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1105 | 712 | 207 | 182 | 4 | | | |

- Molecule 40 is a protein called 50S ribosomal protein L14.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 40 | BO | 122 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 933 | 588 | 171 | 170 | 4 | | | |
| 40 | DO | 122 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 933 | 588 | 171 | 170 | 4 | | | |

- Molecule 41 is a protein called 50S ribosomal protein L15.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 41 | BP | 146 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1114 | 692 | 227 | 193 | 2 | | | |
| 41 | DP | 146 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1114 | 692 | 227 | 193 | 2 | | | |

- Molecule 42 is a protein called 50S ribosomal protein L16.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 42 | BQ | 136 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1080 | 688 | 204 | 183 | 5 | | | |
| 42 | DQ | 136 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1080 | 688 | 204 | 183 | 5 | | | |

- Molecule 43 is a protein called 50S ribosomal protein L17.

| Mol | Chain | Residues | Atoms | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|
| 43 | BR | 117 | Total | C | N | O | 0 | 0 | 0 |
| | | | 960 | 599 | 202 | 159 | | | |
| 43 | DR | 117 | Total | C | N | O | 0 | 0 | 0 |
| | | | 960 | 599 | 202 | 159 | | | |

- Molecule 44 is a protein called 50S ribosomal protein L18.

| Mol | Chain | Residues | Atoms | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|
| 44 | BS | 99 | Total | C | N | O | 0 | 0 | 1 |
| | | | 771 | 486 | 155 | 130 | | | |
| 44 | DS | 99 | Total | C | N | O | 0 | 0 | 1 |
| | | | 771 | 486 | 155 | 130 | | | |

- Molecule 45 is a protein called 50S ribosomal protein L19.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 45 | BT | 132 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1100 | 686 | 227 | 186 | 1 | | | |
| 45 | DT | 132 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1100 | 686 | 227 | 186 | 1 | | | |

- Molecule 46 is a protein called 50S ribosomal protein L20.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 46 | BU | 117 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 958 | 604 | 202 | 151 | 1 | | | |
| 46 | DU | 117 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 958 | 604 | 202 | 151 | 1 | | | |

- Molecule 47 is a protein called 50S ribosomal protein L21.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 47 | BV | 101 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 779 | 501 | 142 | 135 | 1 | | | |
| 47 | DV | 101 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 779 | 501 | 142 | 135 | 1 | | | |

- Molecule 48 is a protein called 50S ribosomal protein L22.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 48 | BW | 113 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 896 | 563 | 176 | 155 | 2 | | | |
| 48 | DW | 113 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 896 | 563 | 176 | 155 | 2 | | | |

- Molecule 49 is a protein called 50S ribosomal protein L23.

| Mol | Chain | Residues | Atoms | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|
| 49 | BX | 93 | Total | C | N | O | 0 | 0 | 1 |
| | | | 726 | 471 | 132 | 123 | | | |
| 49 | DX | 93 | Total | C | N | O | 0 | 0 | 1 |
| | | | 726 | 471 | 132 | 123 | | | |

- Molecule 50 is a protein called 50S ribosomal protein L24.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 50 | BY | 101 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 776 | 500 | 149 | 123 | 4 | | | |
| 50 | DY | 101 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 776 | 500 | 149 | 123 | 4 | | | |

- Molecule 51 is a protein called 50S ribosomal protein L25.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 51 | BZ | 177 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1404 | 897 | 253 | 252 | 2 | | | |

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| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|---|---|---|---|---|---|---|---|---|---|---|
| 51 | DZ | 177 | Total | C | N | O | S | 0 | 0 | 1 |
| | | | 1404 | 897 | 253 | 252 | 2 | | | |

- Molecule 52 is MAGNESIUM ION (CCD ID: MG) (formula: Mg).

| Mol | Chain | Residues | Atoms | | ZeroOcc | AltConf |
|---|---|---|---|---|---|---|
| 52 | AA | 51 | Total | Mg | 0 | 0 |
| | | | 51 | 51 | | |
| 52 | B0 | 1 | Total | Mg | 0 | 0 |
| | | | 1 | 1 | | |
| 52 | B1 | 1 | Total | Mg | 0 | 0 |
| | | | 1 | 1 | | |
| 52 | B5 | 2 | Total | Mg | 0 | 0 |
| | | | 2 | 2 | | |
| 52 | B7 | 1 | Total | Mg | 0 | 0 |
| | | | 1 | 1 | | |
| 52 | BA | 349 | Total | Mg | 0 | 0 |
| | | | 349 | 349 | | |
| 52 | BB | 5 | Total | Mg | 0 | 0 |
| | | | 5 | 5 | | |
| 52 | BD | 1 | Total | Mg | 0 | 0 |
| | | | 1 | 1 | | |
| 52 | BE | 1 | Total | Mg | 0 | 0 |
| | | | 1 | 1 | | |
| 52 | BF | 1 | Total | Mg | 0 | 0 |
| | | | 1 | 1 | | |
| 52 | BP | 3 | Total | Mg | 0 | 0 |
| | | | 3 | 3 | | |
| 52 | BQ | 2 | Total | Mg | 0 | 0 |
| | | | 2 | 2 | | |
| 52 | BR | 1 | Total | Mg | 0 | 0 |
| | | | 1 | 1 | | |
| 52 | BU | 1 | Total | Mg | 0 | 0 |
| | | | 1 | 1 | | |
| 52 | BX | 1 | Total | Mg | 0 | 0 |
| | | | 1 | 1 | | |
| 52 | CA | 48 | Total | Mg | 0 | 0 |
| | | | 48 | 48 | | |
| 52 | D0 | 1 | Total | Mg | 0 | 0 |
| | | | 1 | 1 | | |
| 52 | D1 | 1 | Total | Mg | 0 | 0 |
| | | | 1 | 1 | | |

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| Mol | Chain | Residues | Atoms | ZeroOcc | AltConf |
|---|---|---|---|---|---|
| 52 | D5 | 2 | Total Mg<br>2 2 | 0 | 0 |
| 52 | D7 | 1 | Total Mg<br>1 1 | 0 | 0 |
| 52 | DA | 309 | Total Mg<br>309 309 | 0 | 0 |
| 52 | DB | 3 | Total Mg<br>3 3 | 0 | 0 |
| 52 | DD | 1 | Total Mg<br>1 1 | 0 | 0 |
| 52 | DE | 1 | Total Mg<br>1 1 | 0 | 0 |
| 52 | DF | 1 | Total Mg<br>1 1 | 0 | 0 |
| 52 | DP | 1 | Total Mg<br>1 1 | 0 | 0 |
| 52 | DQ | 1 | Total Mg<br>1 1 | 0 | 0 |
| 52 | DR | 1 | Total Mg<br>1 1 | 0 | 0 |
| 52 | DU | 1 | Total Mg<br>1 1 | 0 | 0 |
| 52 | DX | 1 | Total Mg<br>1 1 | 0 | 0 |

- Molecule 53 is ZINC ION (CCD ID: ZN) (formula: Zn).

| Mol | Chain | Residues | Atoms | ZeroOcc | AltConf |
|---|---|---|---|---|---|
| 53 | AD | 1 | Total Zn<br>1 1 | 0 | 0 |
| 53 | AN | 1 | Total Zn<br>1 1 | 0 | 0 |
| 53 | CD | 1 | Total Zn<br>1 1 | 0 | 0 |
| 53 | CN | 1 | Total Zn<br>1 1 | 0 | 0 |

- Molecule 54 is POTASSIUM ION (CCD ID: K) (formula: K).

| Mol | Chain | Residues | Atoms | ZeroOcc | AltConf |
|---|---|---|---|---|---|
| 54 | BA | 1 | Total K<br>1 1 | 0 | 0 |

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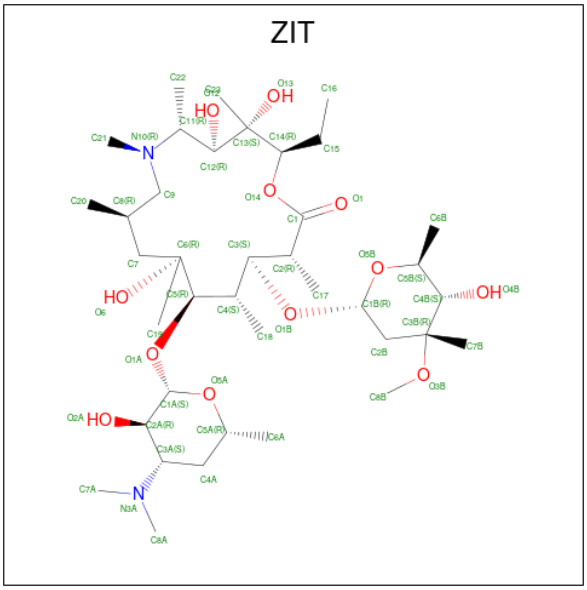
| Mol | Chain | Residues | Atoms | ZeroOcc | AltConf |
|---|---|---|---|---|---|
| 54 | DA | 1 | Total K<br>1 1 | 0 | 0 |

- Molecule 55 is AZITHROMYCIN (CCD ID: ZIT) (formula: $C_{38}H_{72}N_2O_{12}$).



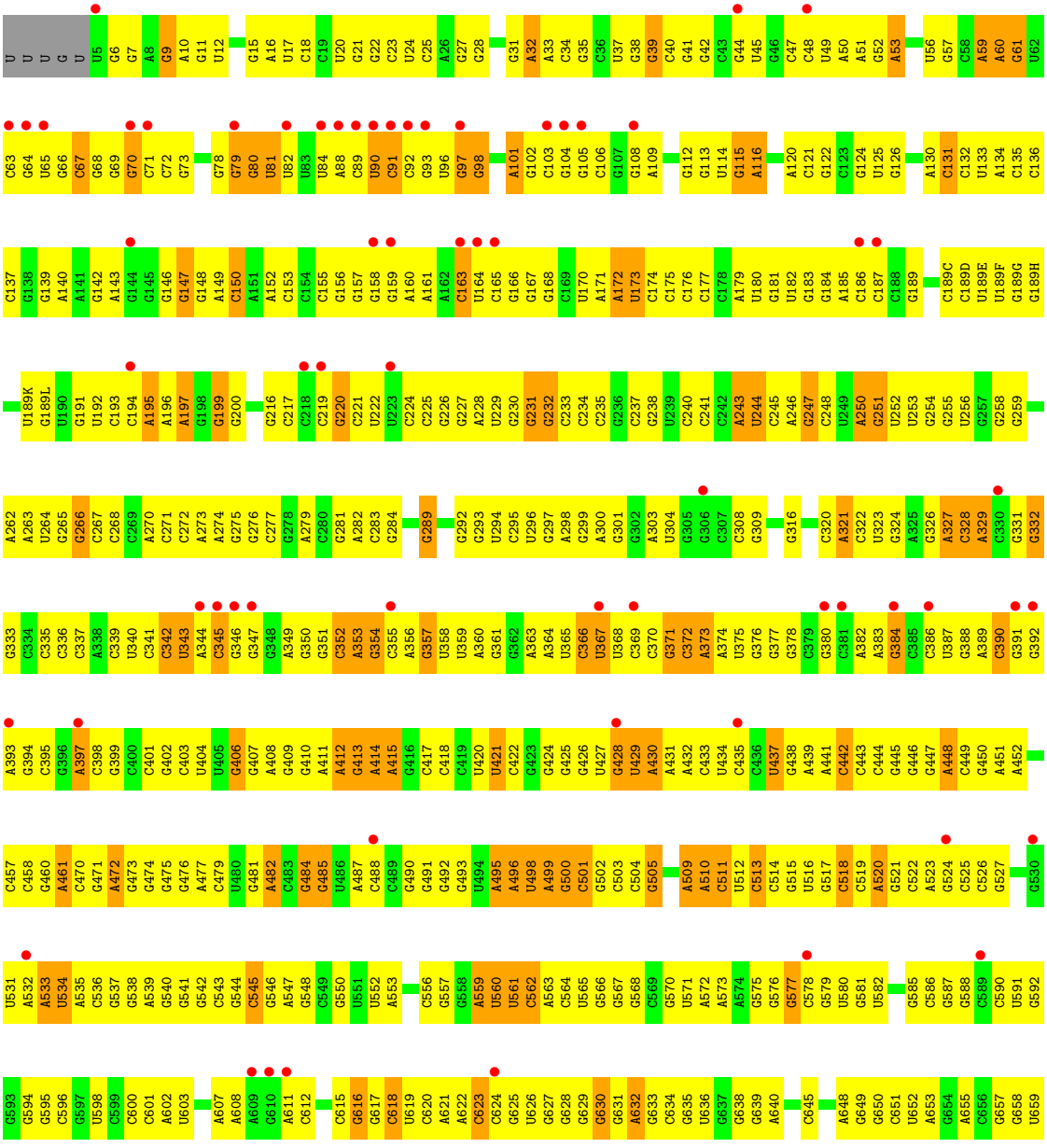

| Mol | Chain | Residues | Atoms | ZeroOcc | AltConf |
|---|---|---|---|---|---|
| 55 | BA | 1 | Total C N O<br>52 38 2 12 | 0 | 0 |
| 55 | DA | 1 | Total C N O<br>52 38 2 12 | 0 | 0 |

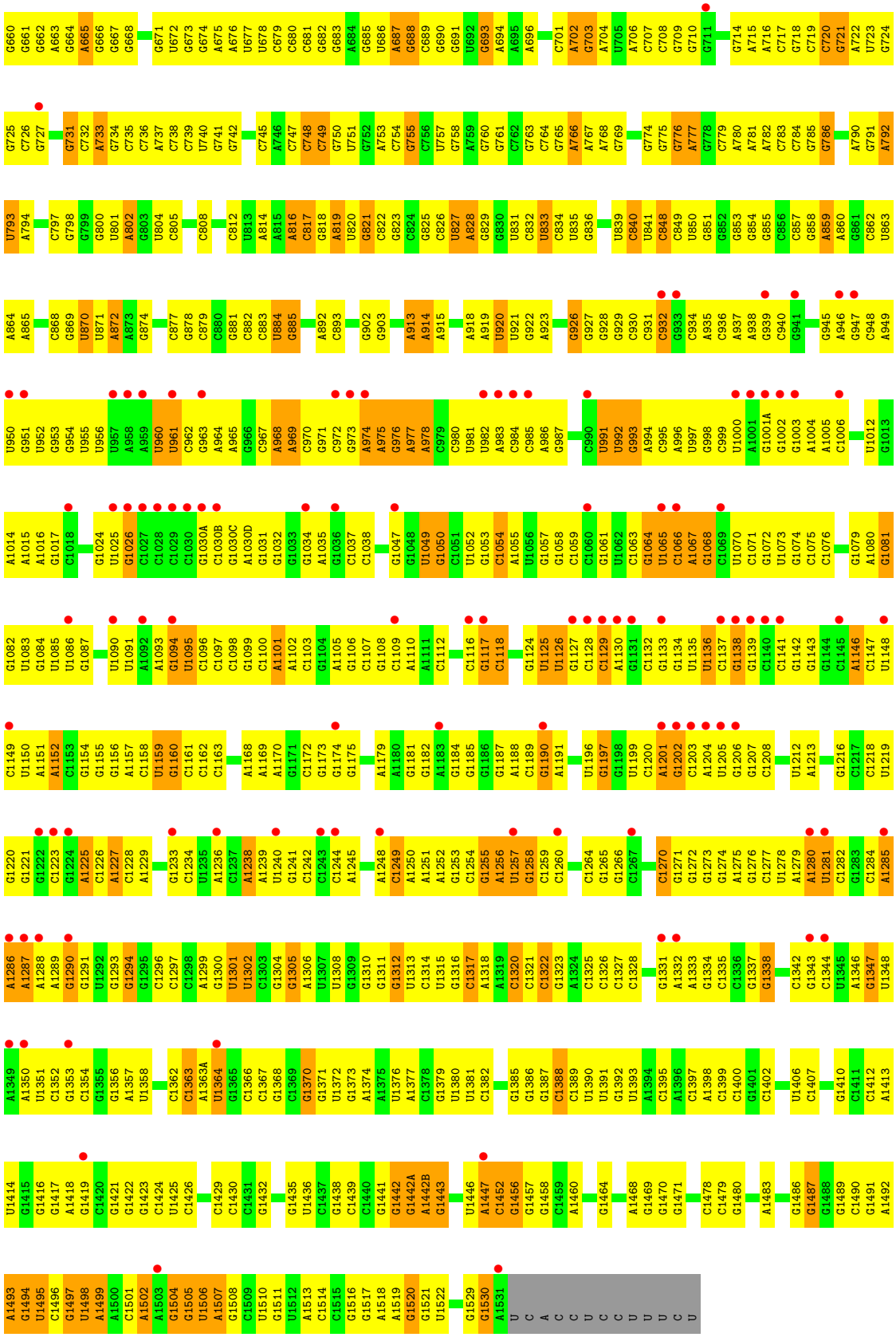
# 3 Residue-property plots (i)

These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and electron density. Residues are color-coded according to the number of geometric quality criteria for which they contain at least one outlier: green = 0, yellow = 1, orange = 2 and red = 3 or more. A red dot above a residue indicates a poor fit to the electron density (RSRZ > 2). Stretches of 2 or more consecutive residues without any outlier are shown as a green connector. Residues present in the sample, but not in the model, are shown in grey.

- Molecule 1: 16S rRNA

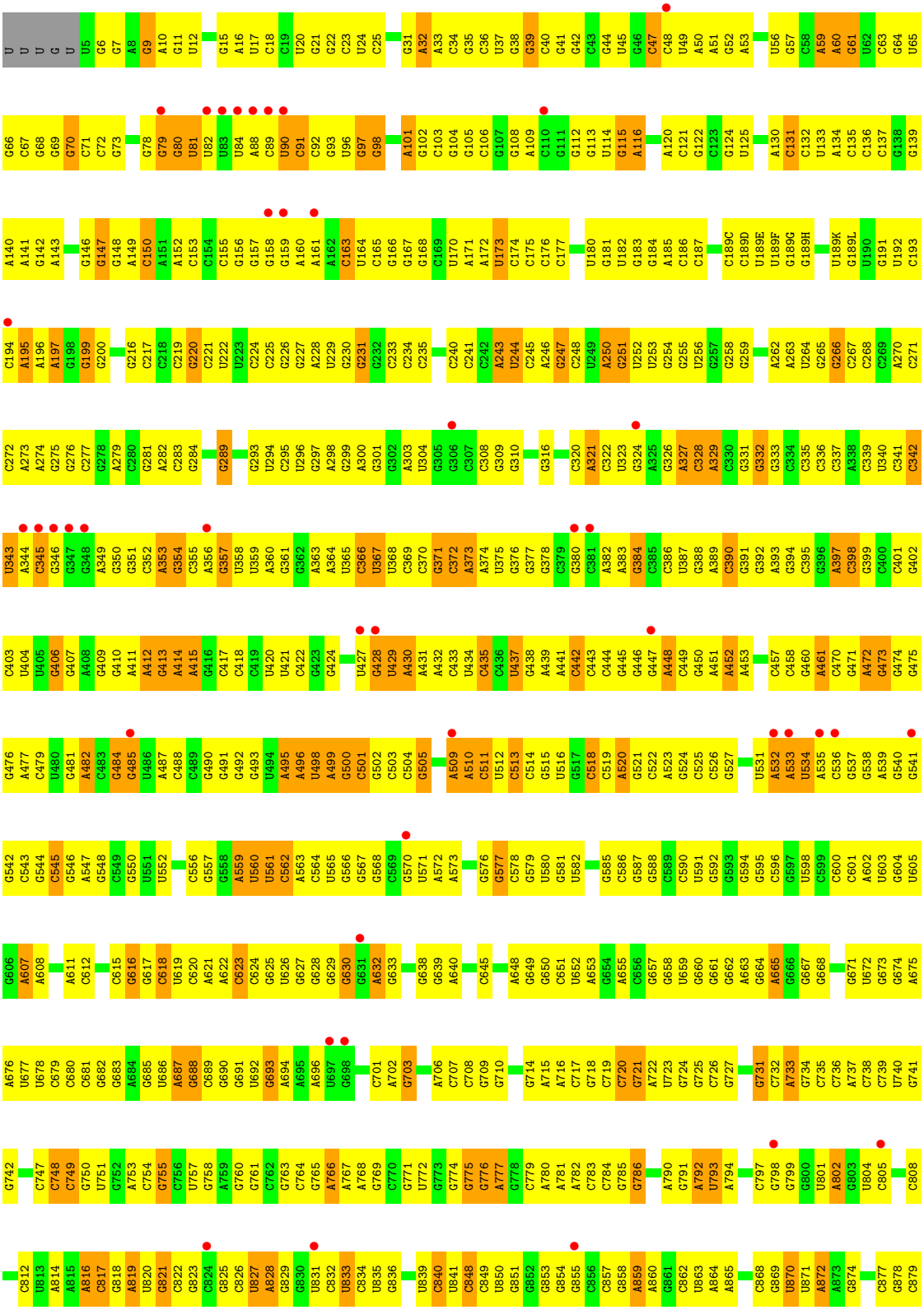
Chain AA: 11% 27% 57% 15%

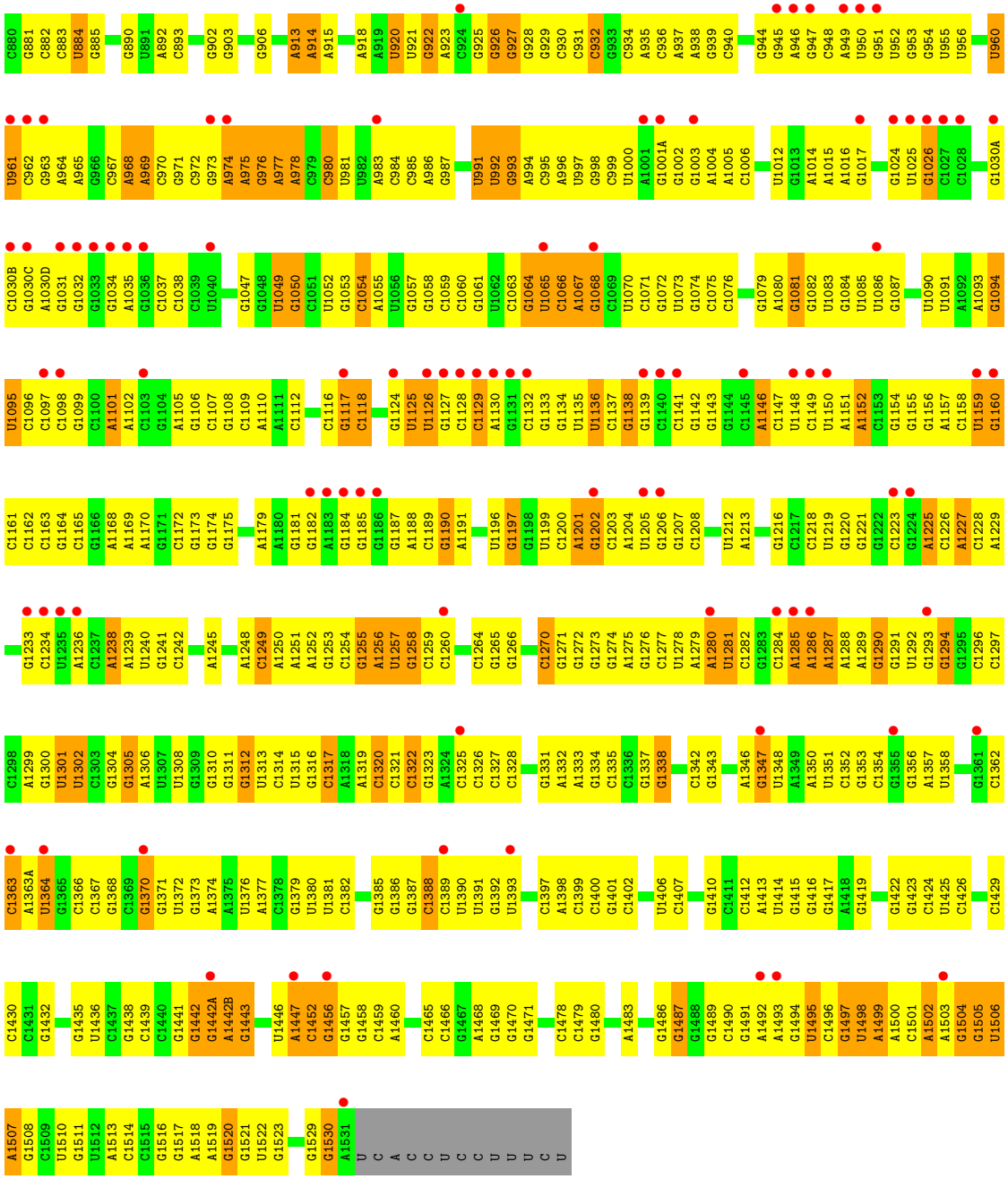




• Molecule 1: 16S rRNA

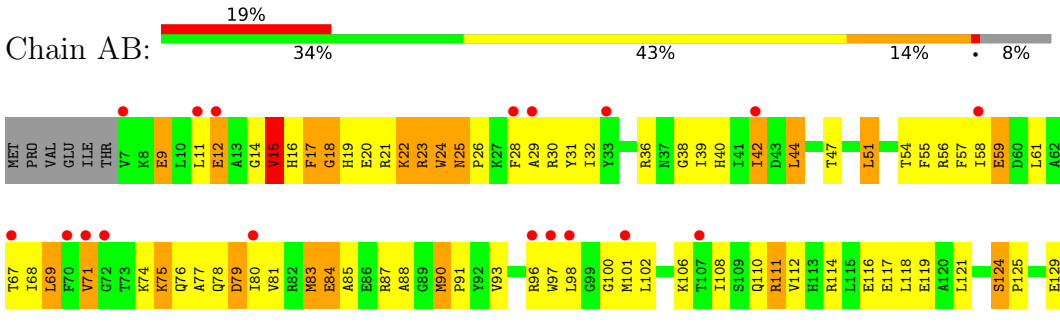
Chain CA: 9% 27% 57% 15%

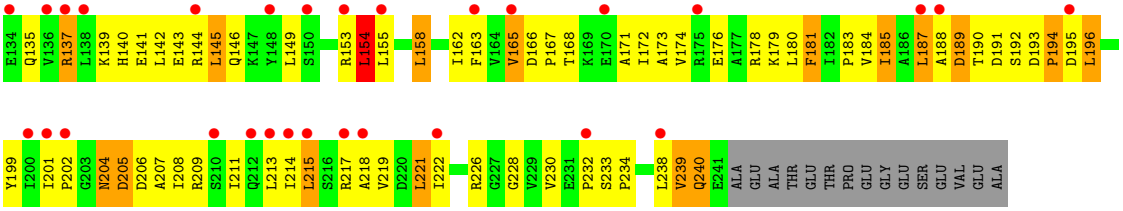




• Molecule 2: 30S ribosomal protein S2

Chain AB:

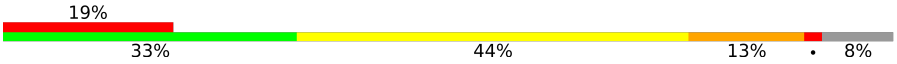


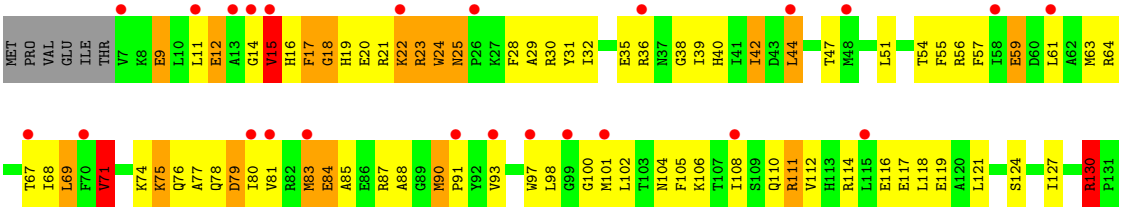


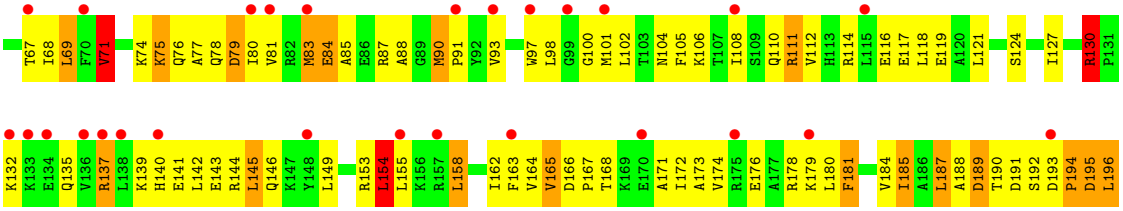


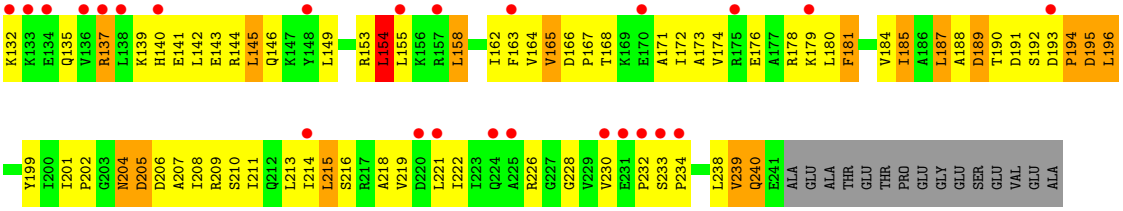
• Molecule 2: 30S ribosomal protein S2

Chain CB:



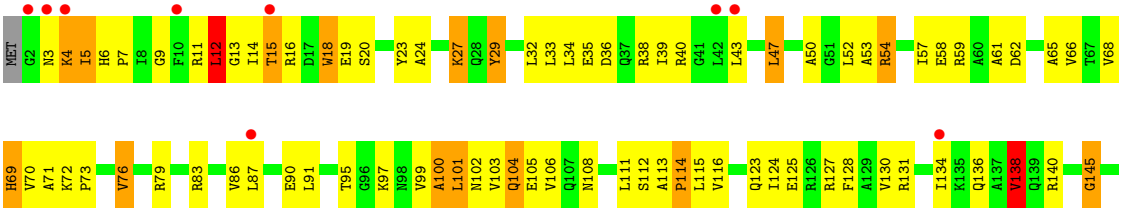


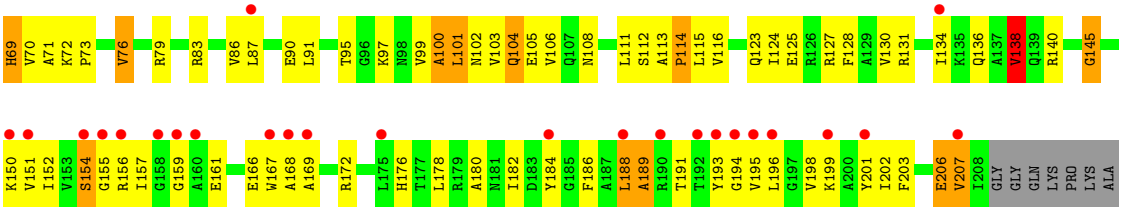


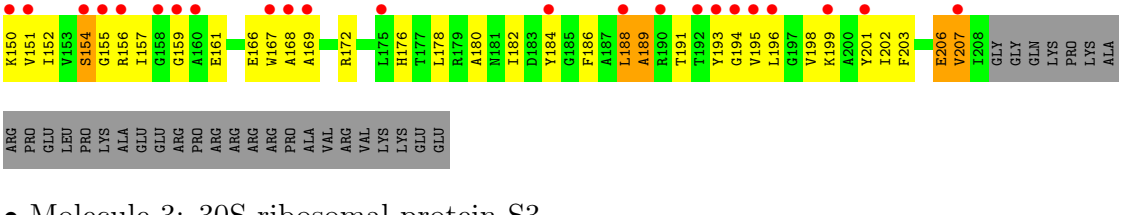


• Molecule 3: 30S ribosomal protein S3

Chain AC:



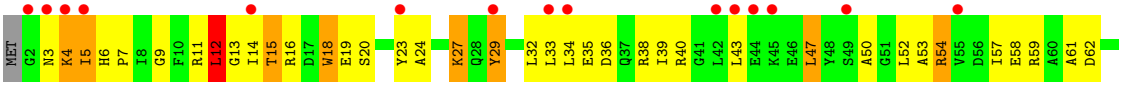


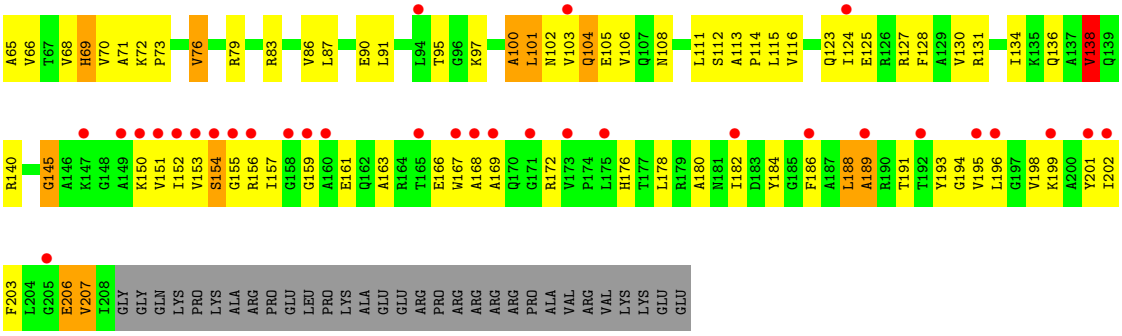




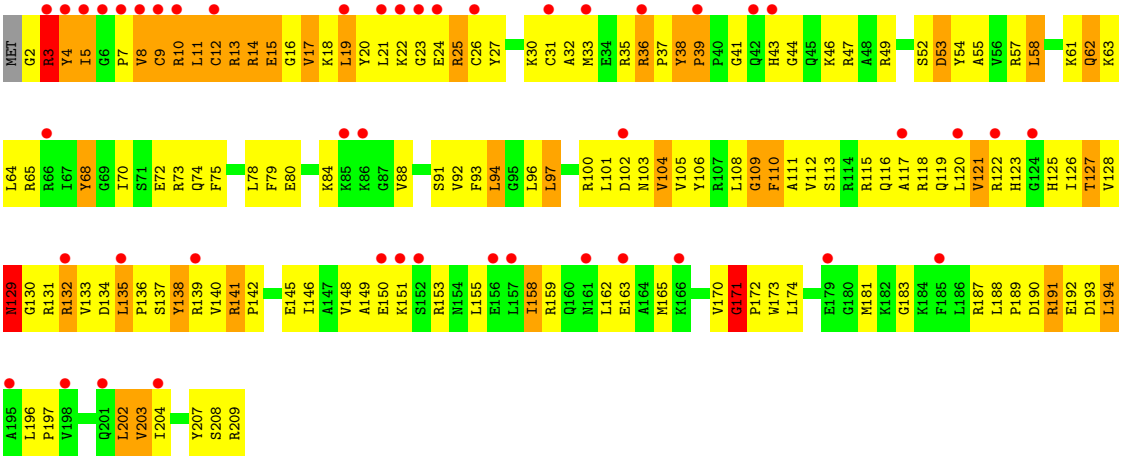
• Molecule 3: 30S ribosomal protein S3

Chain CC:

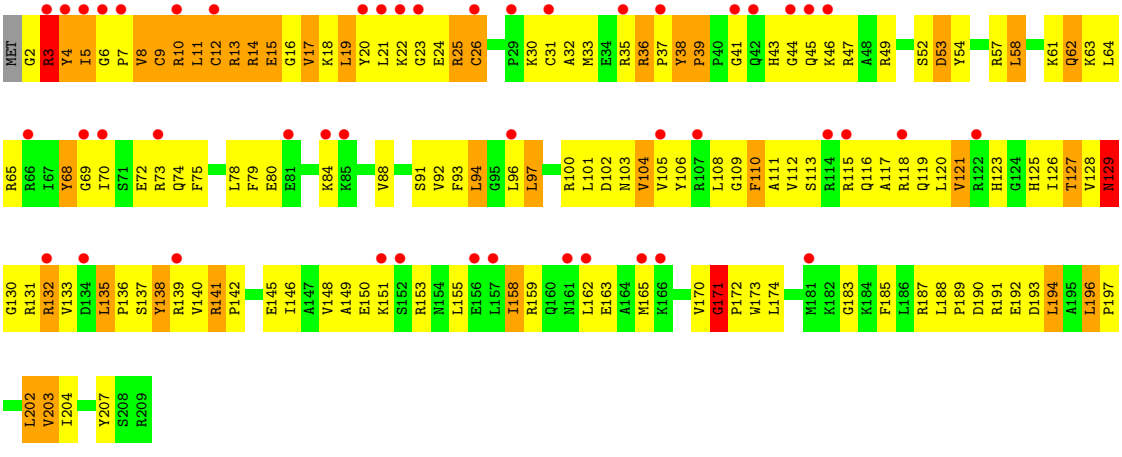




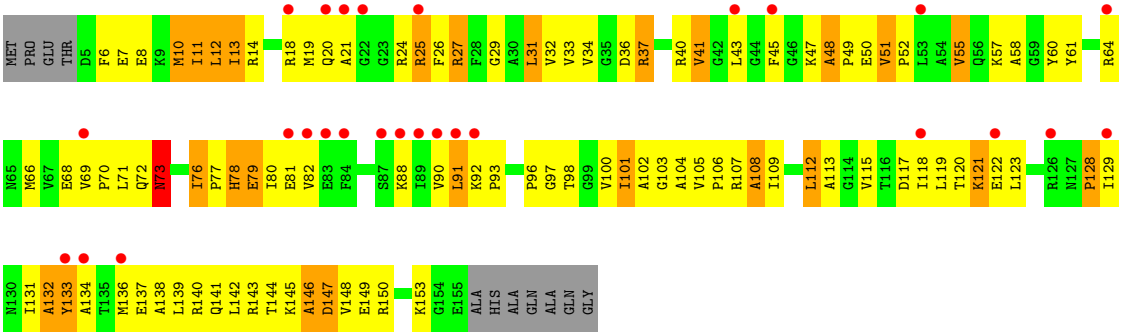

• Molecule 4: 30S ribosomal protein S4




• Molecule 4: 30S ribosomal protein S4

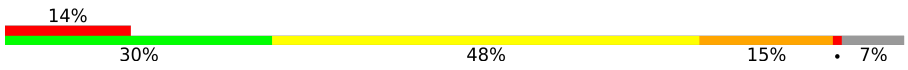


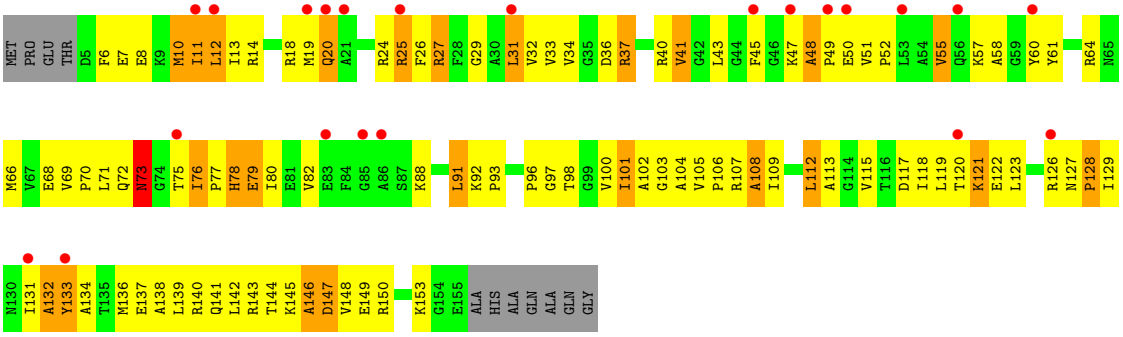

• Molecule 5: 30S ribosomal protein S5



• Molecule 5: 30S ribosomal protein S5

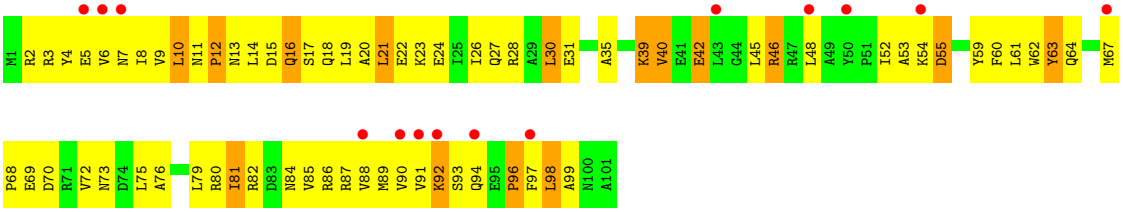
Chain CE:



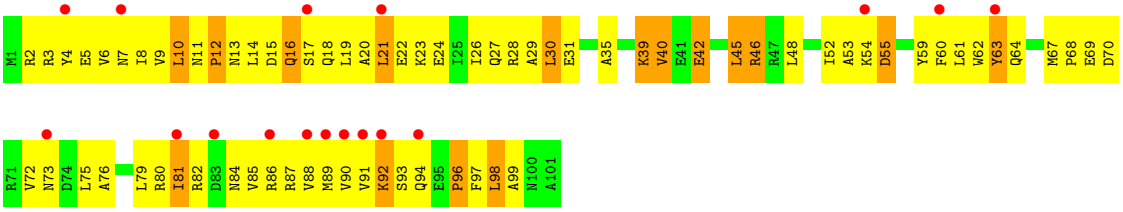
• Molecule 6: 30S ribosomal protein S6

Chain AF:



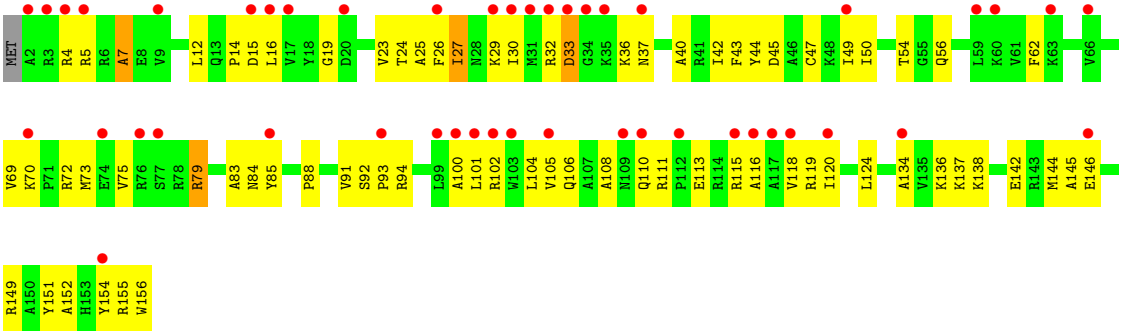
• Molecule 6: 30S ribosomal protein S6

Chain CF:

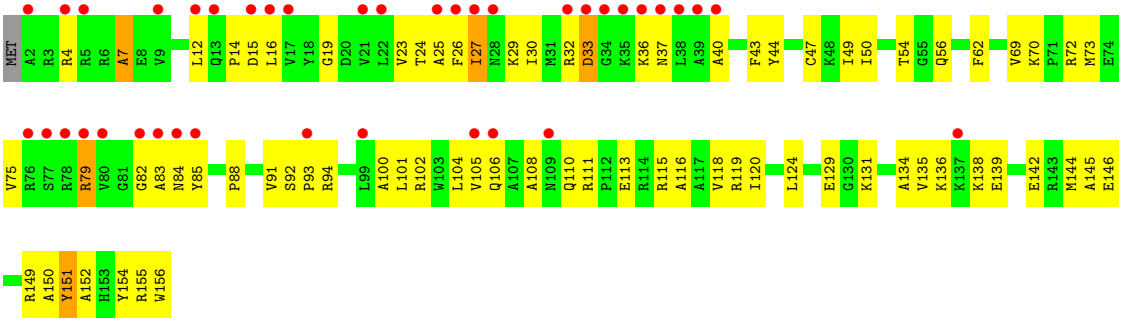


• Molecule 7: 30S ribosomal protein S7

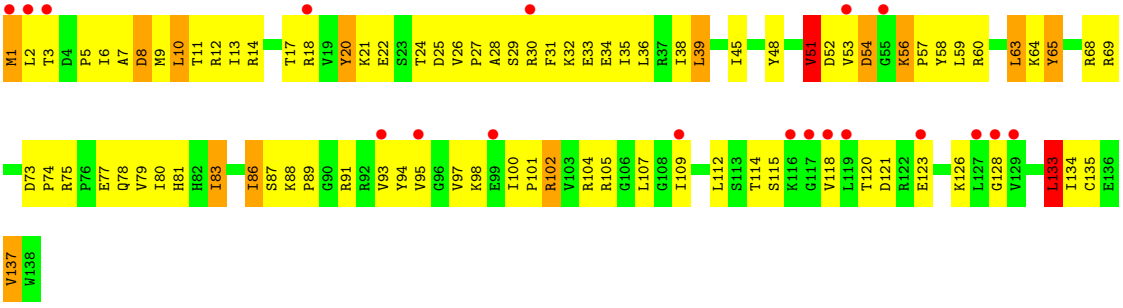
Chain AG:



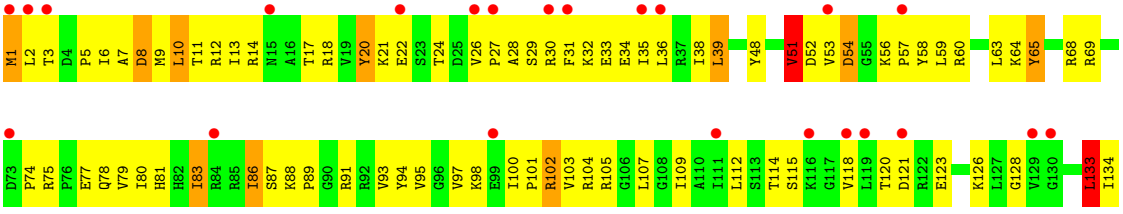

• Molecule 7: 30S ribosomal protein S7



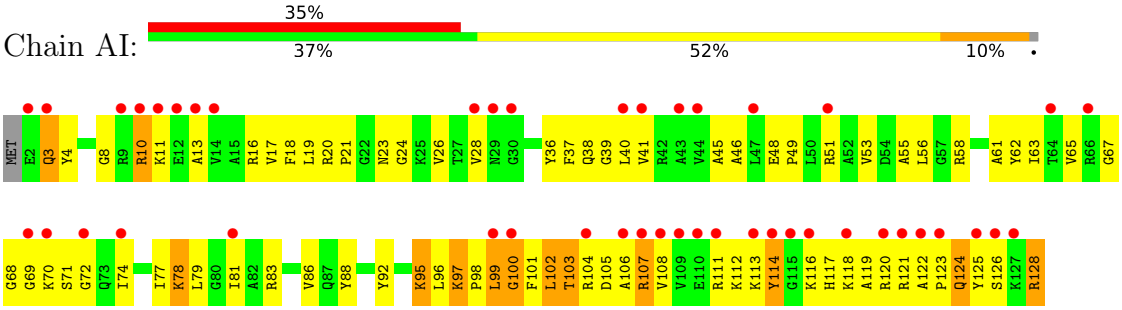

• Molecule 8: 30S ribosomal protein S8



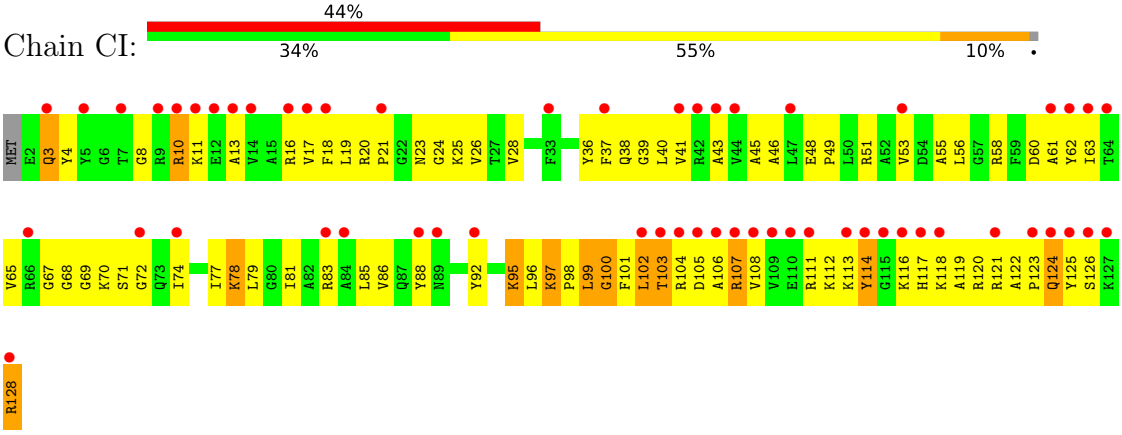

• Molecule 8: 30S ribosomal protein S8



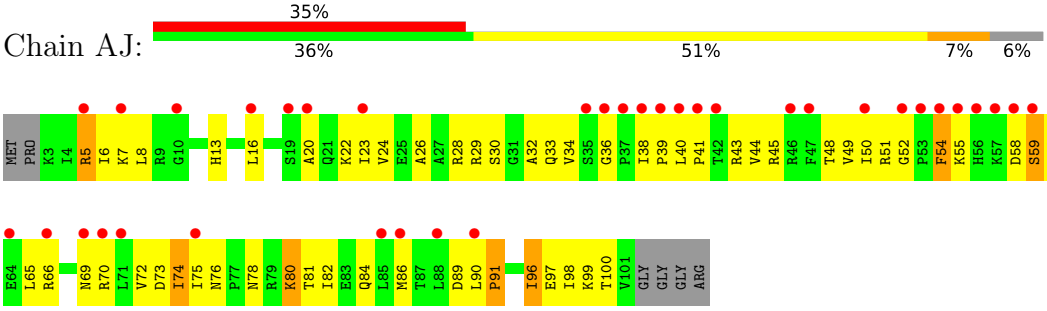
• Molecule 9: 30S ribosomal protein S9



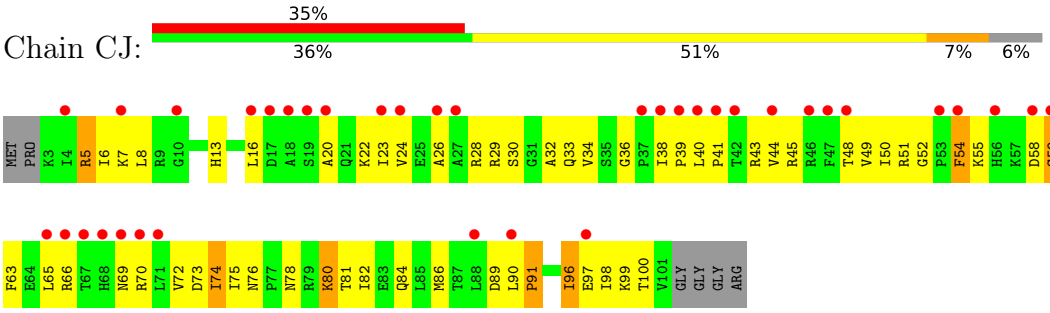

• Molecule 9: 30S ribosomal protein S9




• Molecule 10: 30S ribosomal protein S10

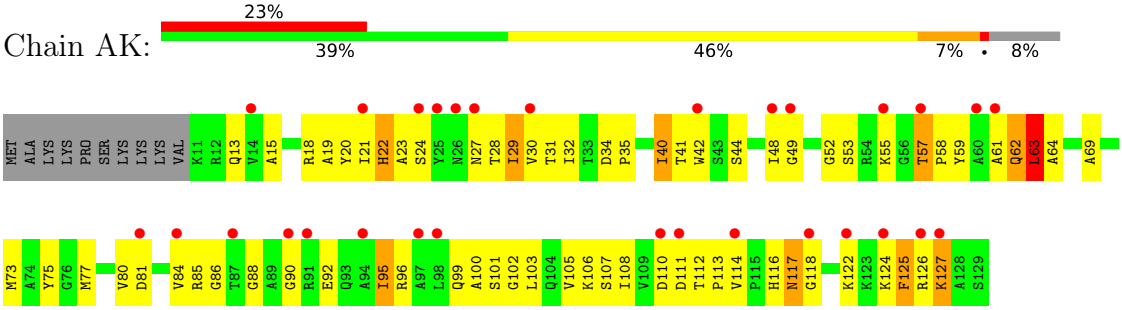



• Molecule 10: 30S ribosomal protein S10



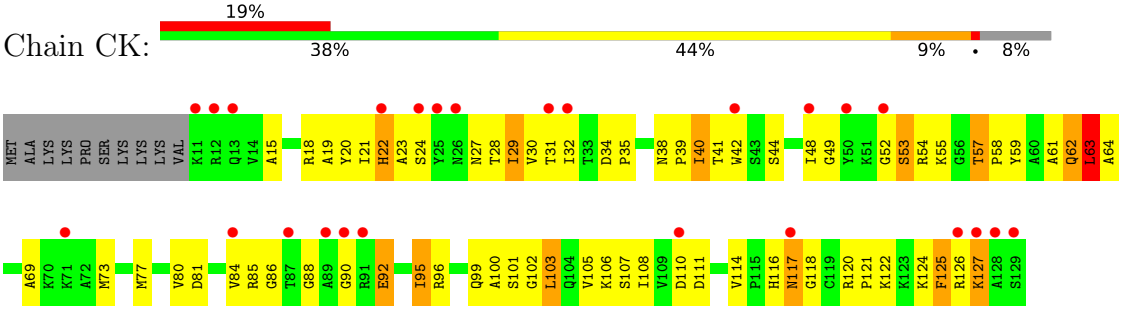
• Molecule 11: 30S ribosomal protein S11

Chain AK:



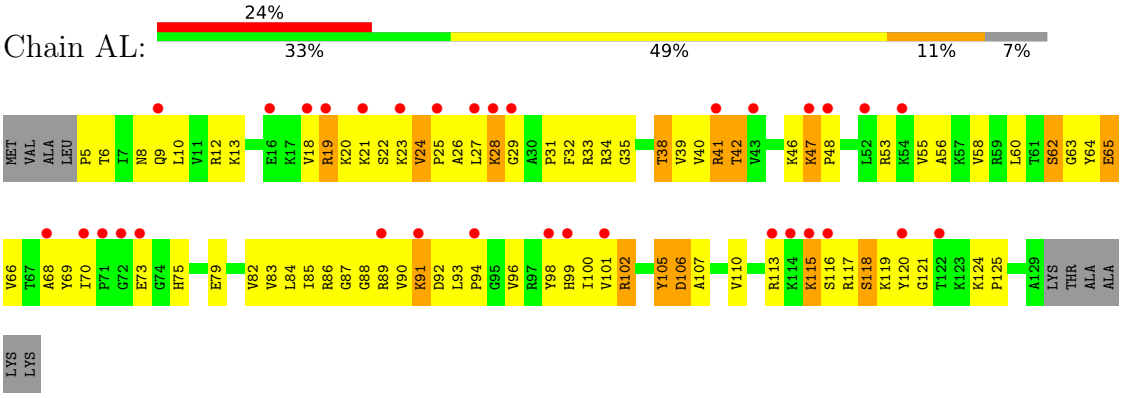

• Molecule 11: 30S ribosomal protein S11

Chain CK:



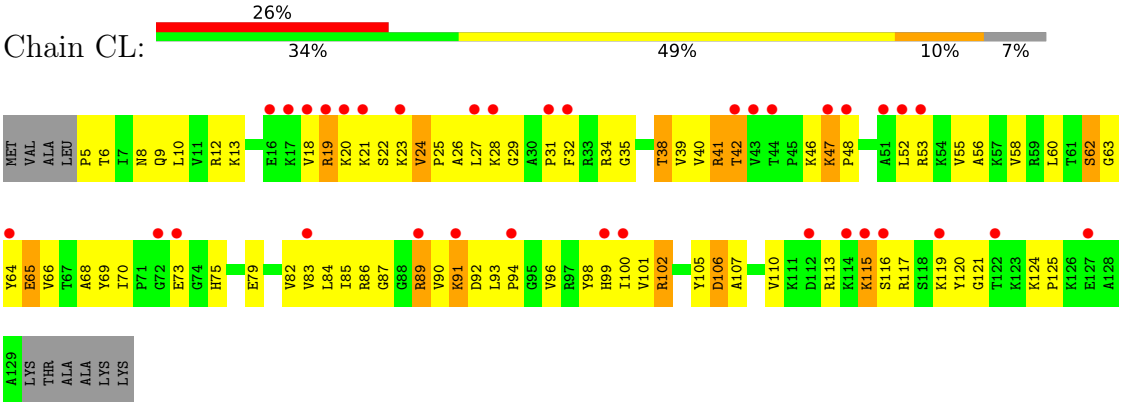

• Molecule 12: 30S ribosomal protein S12

Chain AL:



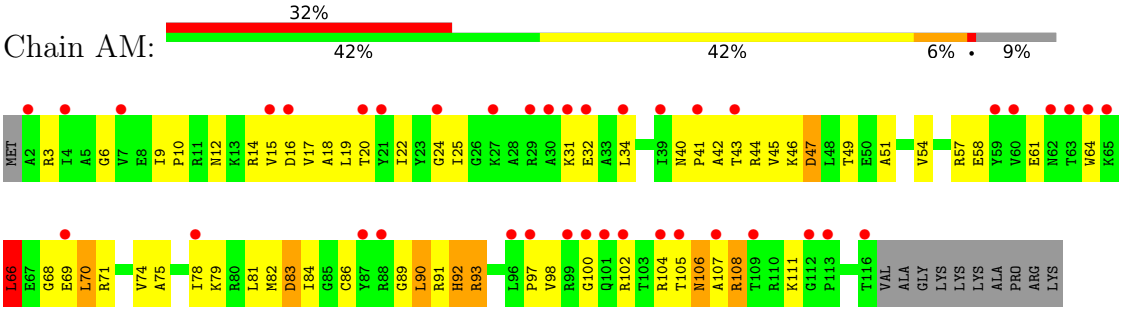

• Molecule 12: 30S ribosomal protein S12

Chain CL:



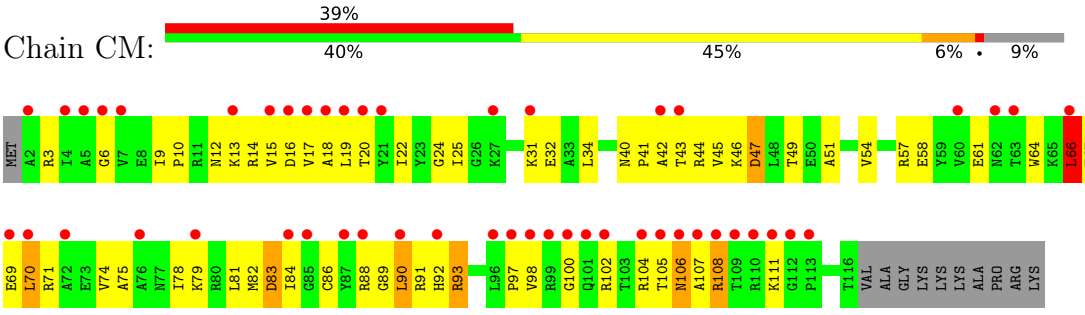

• Molecule 13: 30S ribosomal protein S13

Chain AM:



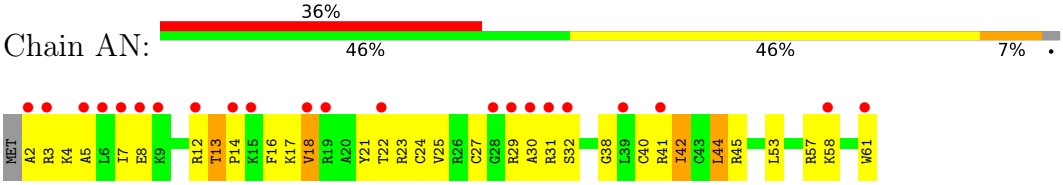

• Molecule 13: 30S ribosomal protein S13

Chain CM:



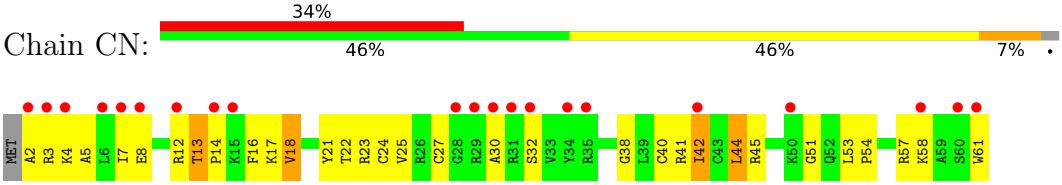

• Molecule 14: 30S ribosomal protein S14

Chain AN:



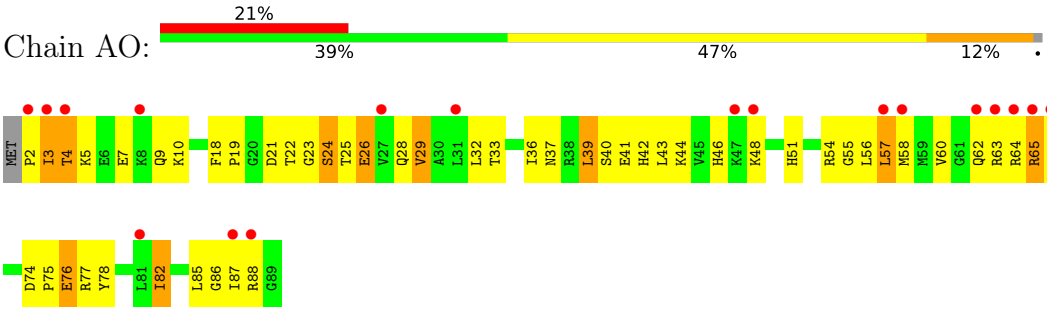

• Molecule 14: 30S ribosomal protein S14

Chain CN:



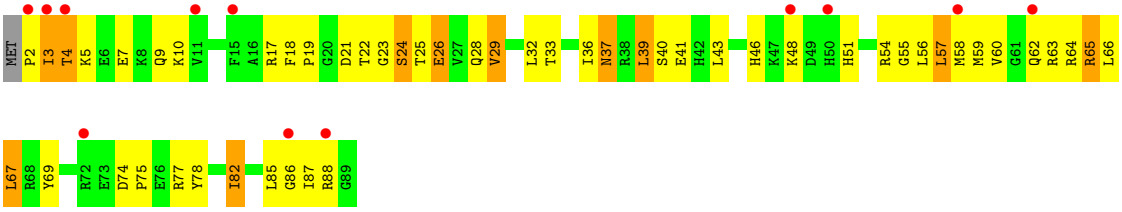

• Molecule 15: 30S ribosomal protein S15

Chain AO:

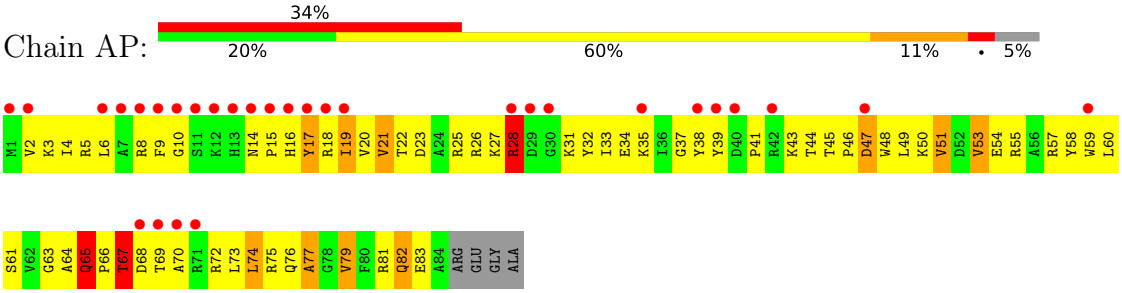



• Molecule 15: 30S ribosomal protein S15

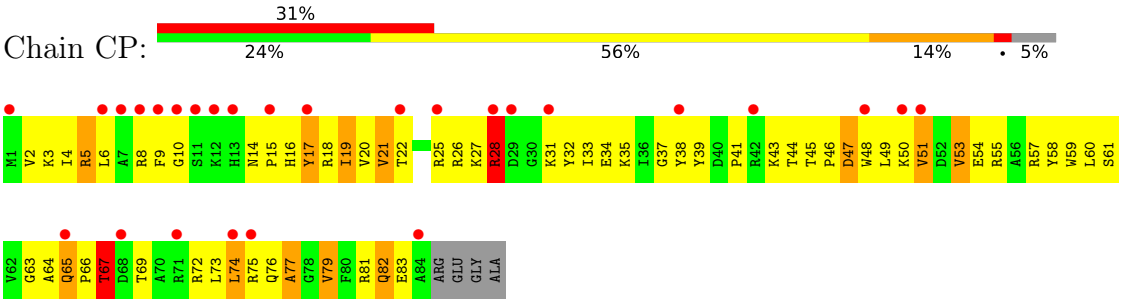
Chain CO:



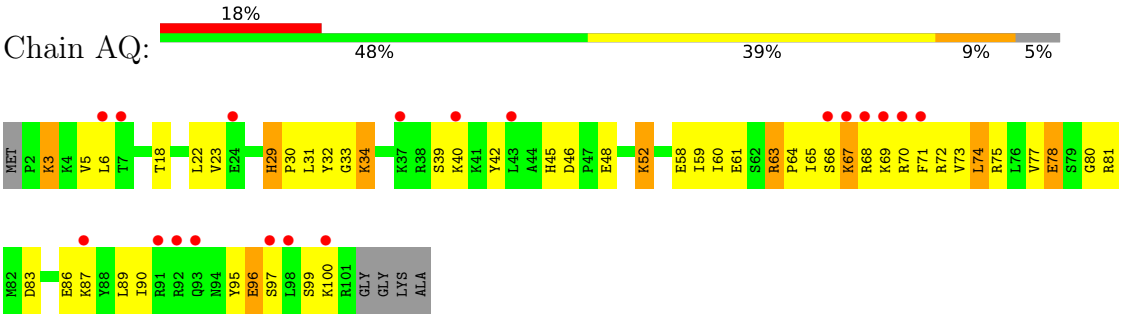

• Molecule 16: 30S ribosomal protein S16



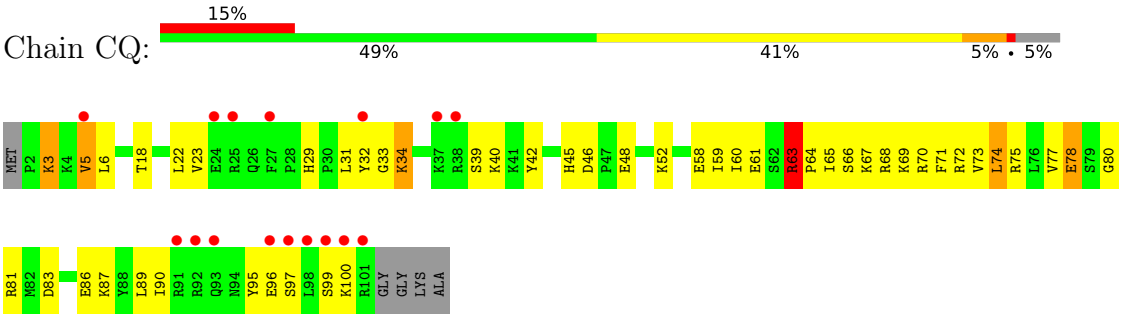

• Molecule 16: 30S ribosomal protein S16




• Molecule 17: 30S ribosomal protein S17

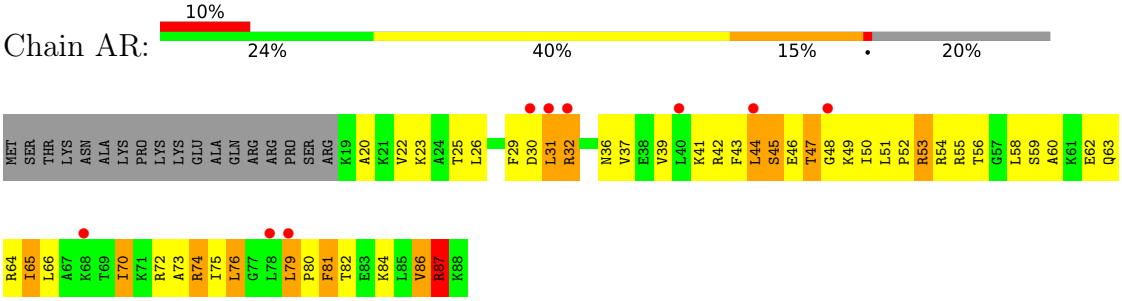



• Molecule 17: 30S ribosomal protein S17



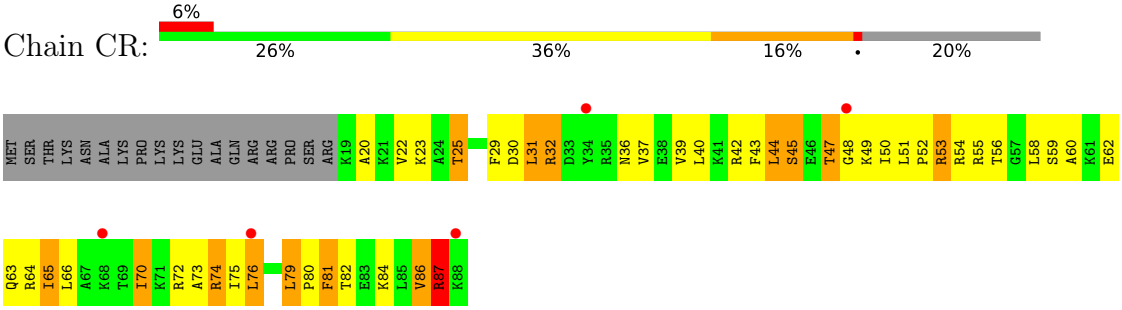
• Molecule 18: 30S ribosomal protein S18

Chain AR:



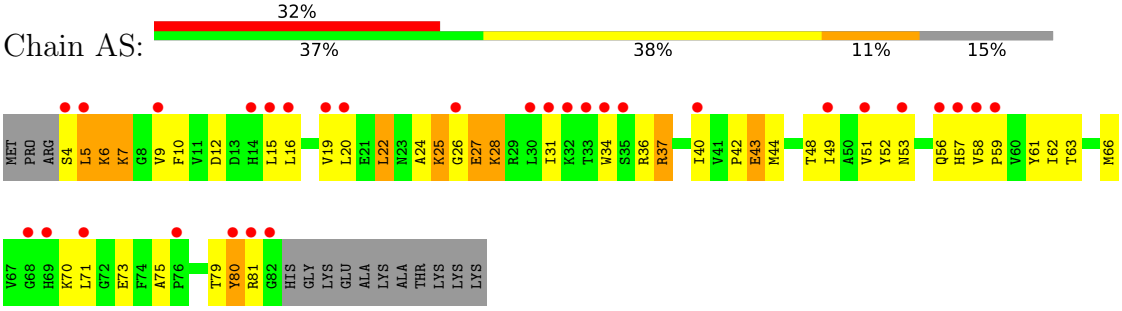

• Molecule 18: 30S ribosomal protein S18

Chain CR:



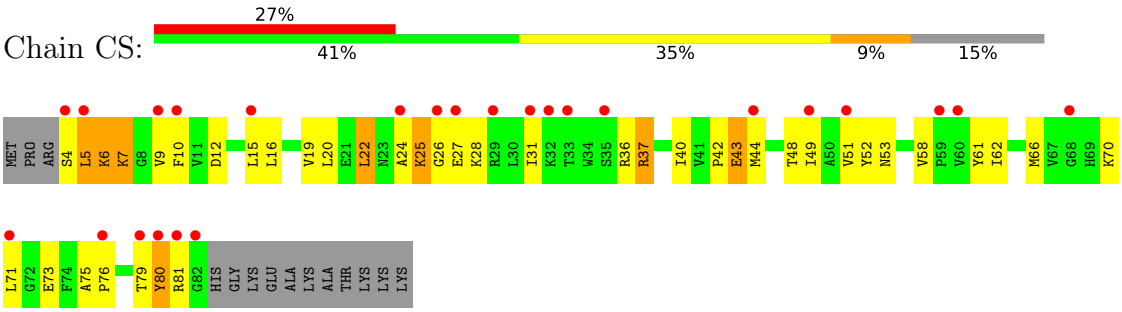

• Molecule 19: 30S ribosomal protein S19

Chain AS:



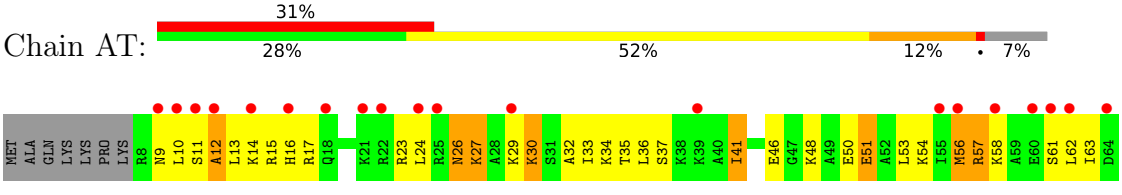

• Molecule 19: 30S ribosomal protein S19

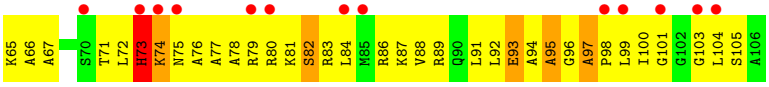
Chain CS:




• Molecule 20: 30S ribosomal protein S20

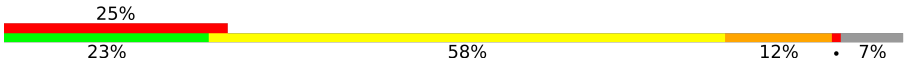
Chain AT:

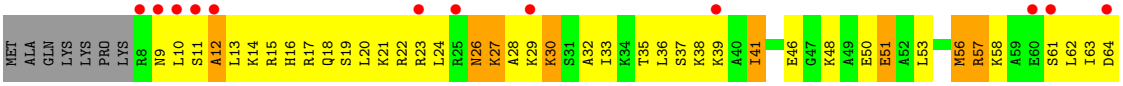


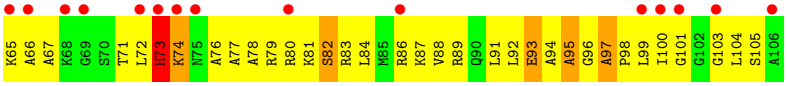



• Molecule 20: 30S ribosomal protein S20

Chain CT:

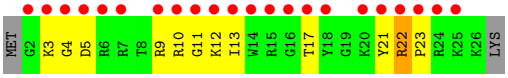


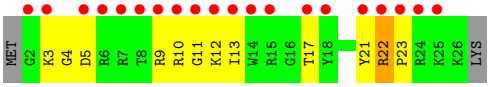

• Molecule 21: 30S ribosomal protein Thx

Chain AU:



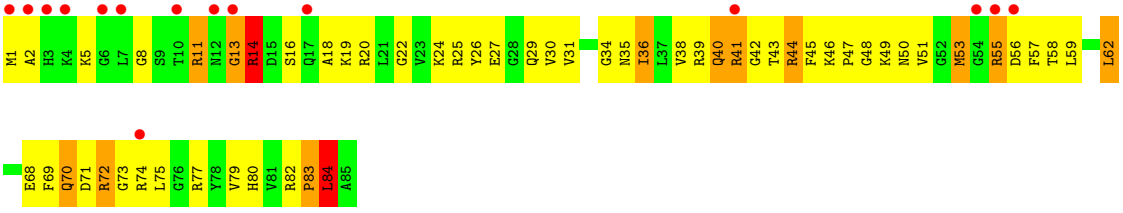

• Molecule 21: 30S ribosomal protein Thx

Chain CU:



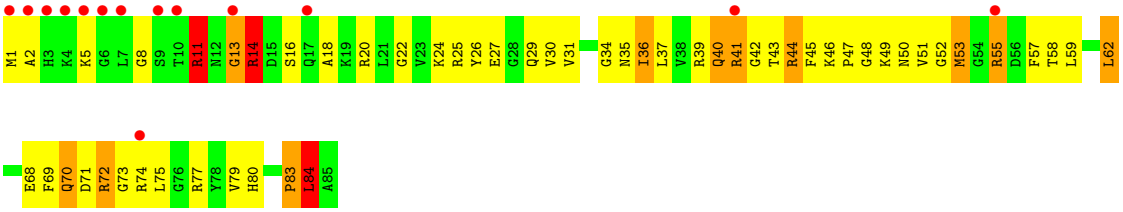

• Molecule 22: 50S ribosomal protein L27

Chain B0:



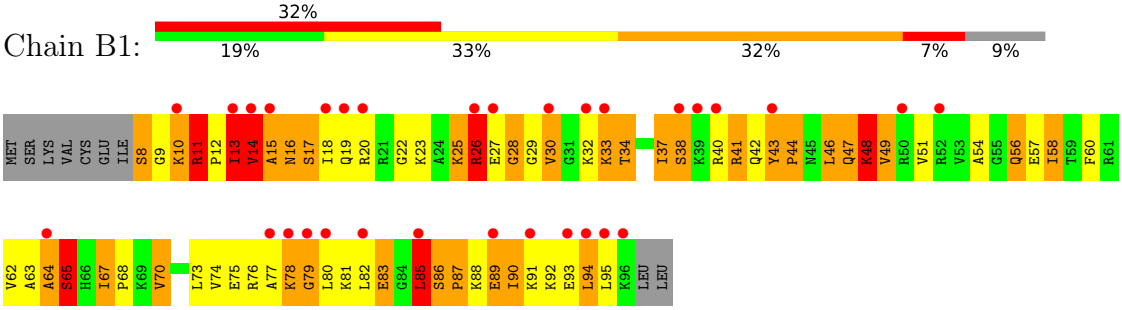

• Molecule 22: 50S ribosomal protein L27

Chain D0:



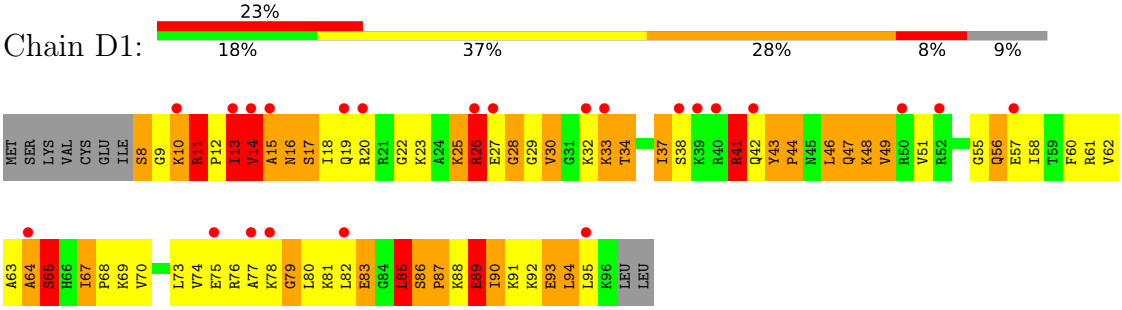
• Molecule 23: 50S ribosomal protein L28

Chain B1:



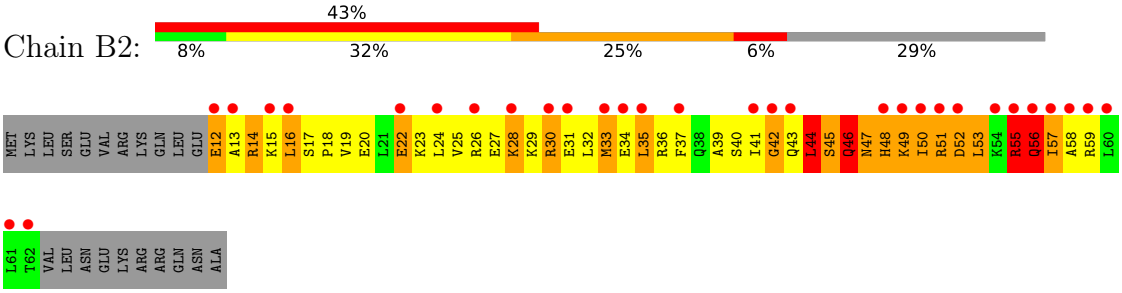

• Molecule 23: 50S ribosomal protein L28

Chain D1:



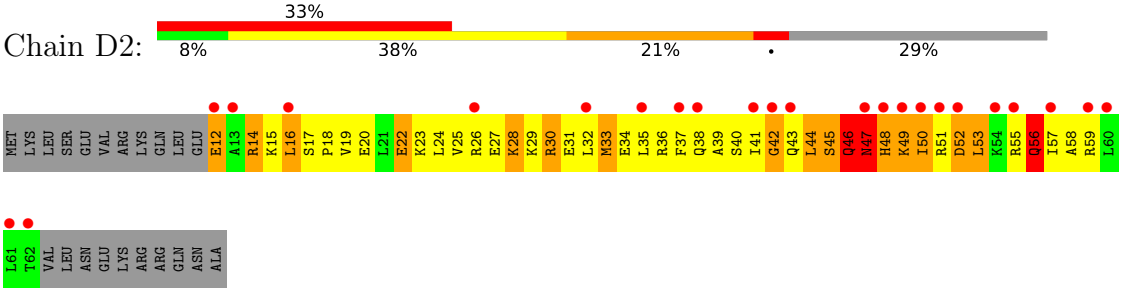

• Molecule 24: 50S ribosomal protein L29

Chain B2:



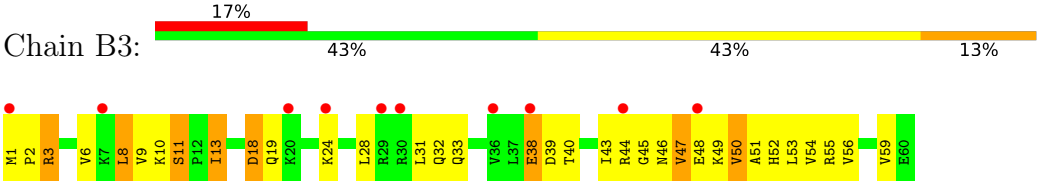

• Molecule 24: 50S ribosomal protein L29

Chain D2:



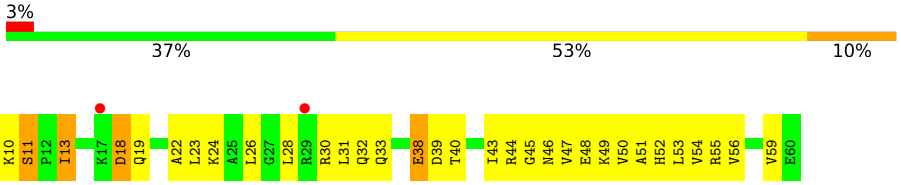

• Molecule 25: 50S ribosomal protein L30

Chain B3:



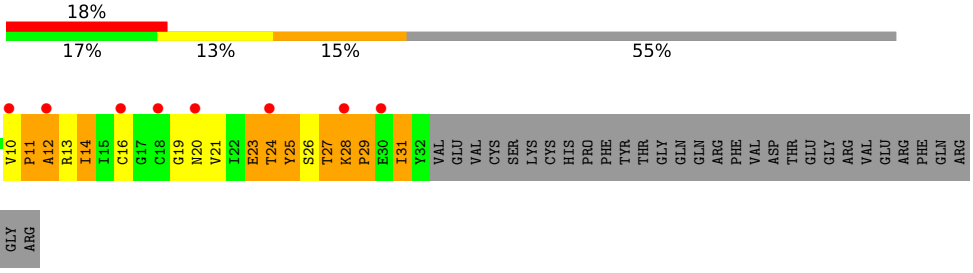
• Molecule 25: 50S ribosomal protein L30

Chain D3:



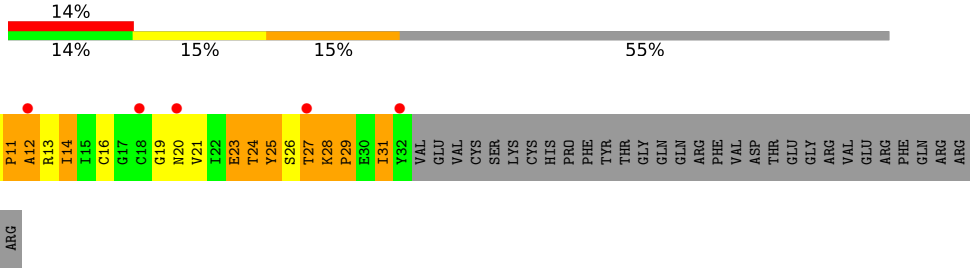

• Molecule 26: 50S ribosomal protein L31

Chain B4:



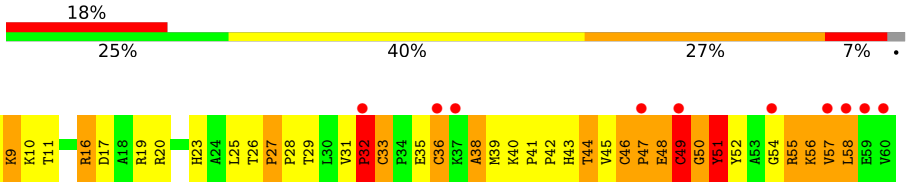

• Molecule 26: 50S ribosomal protein L31

Chain D4:



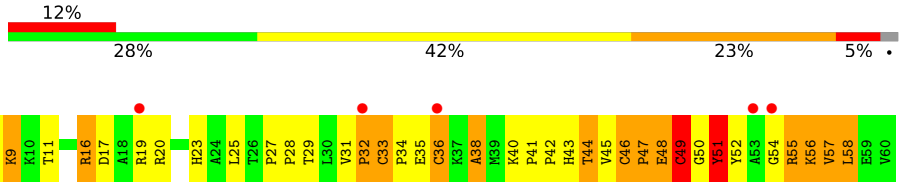

• Molecule 27: 50S ribosomal protein L32

Chain B5:



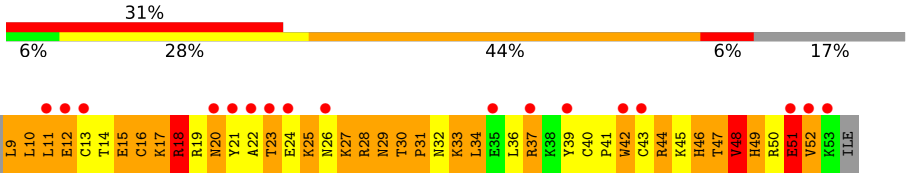

• Molecule 27: 50S ribosomal protein L32

Chain D5:



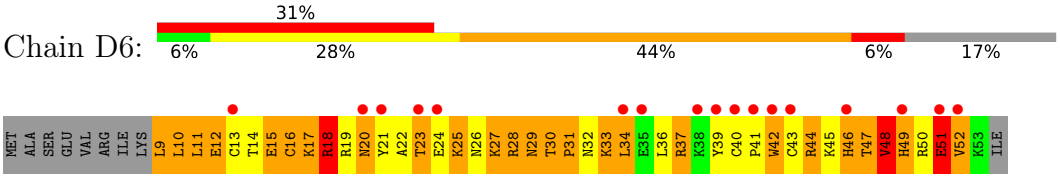

• Molecule 28: 50S ribosomal protein L33

Chain B6:



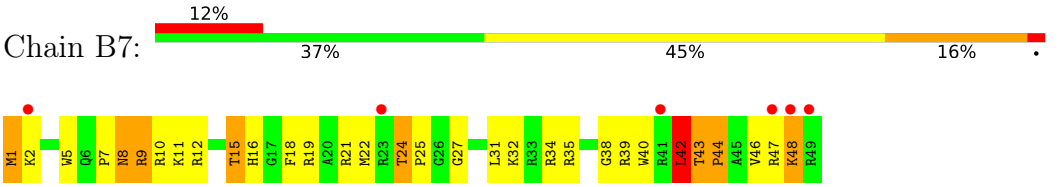

• Molecule 28: 50S ribosomal protein L33

Chain D6:



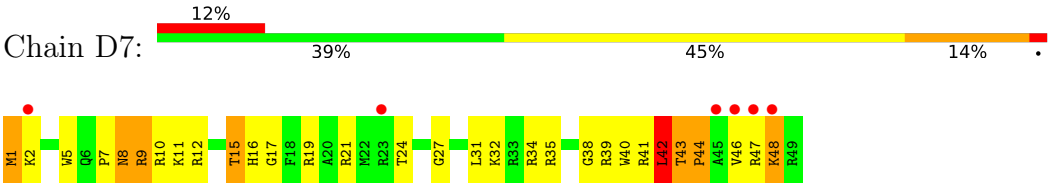

• Molecule 29: 50S ribosomal protein L34

Chain B7:



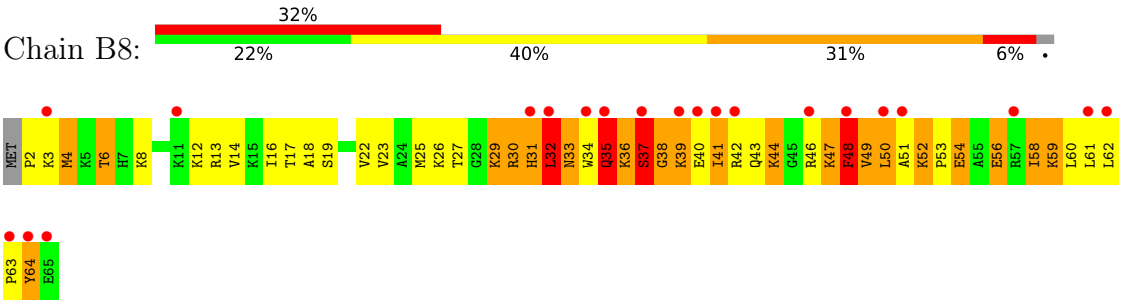

• Molecule 29: 50S ribosomal protein L34

Chain D7:



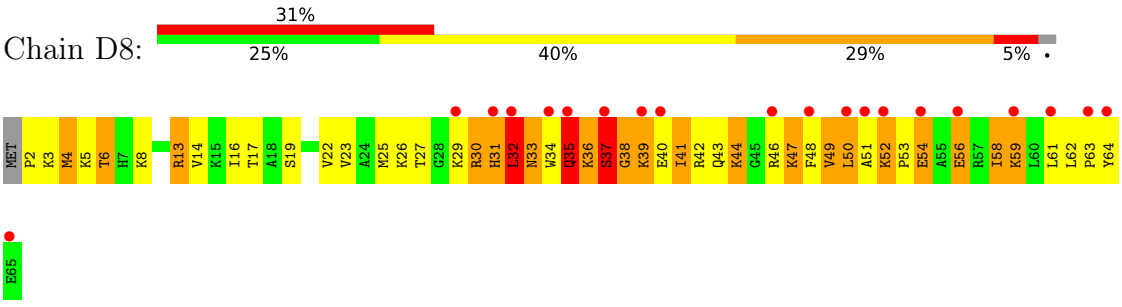

• Molecule 30: 50S ribosomal protein L35

Chain B8:



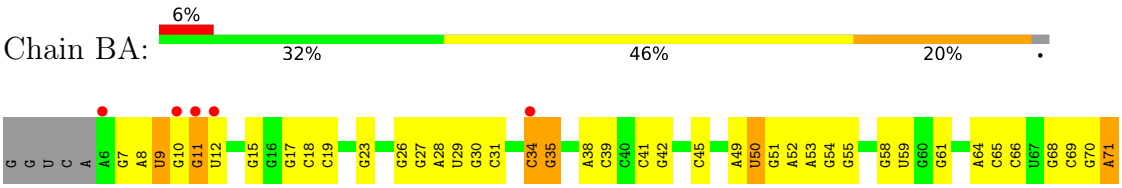

• Molecule 30: 50S ribosomal protein L35

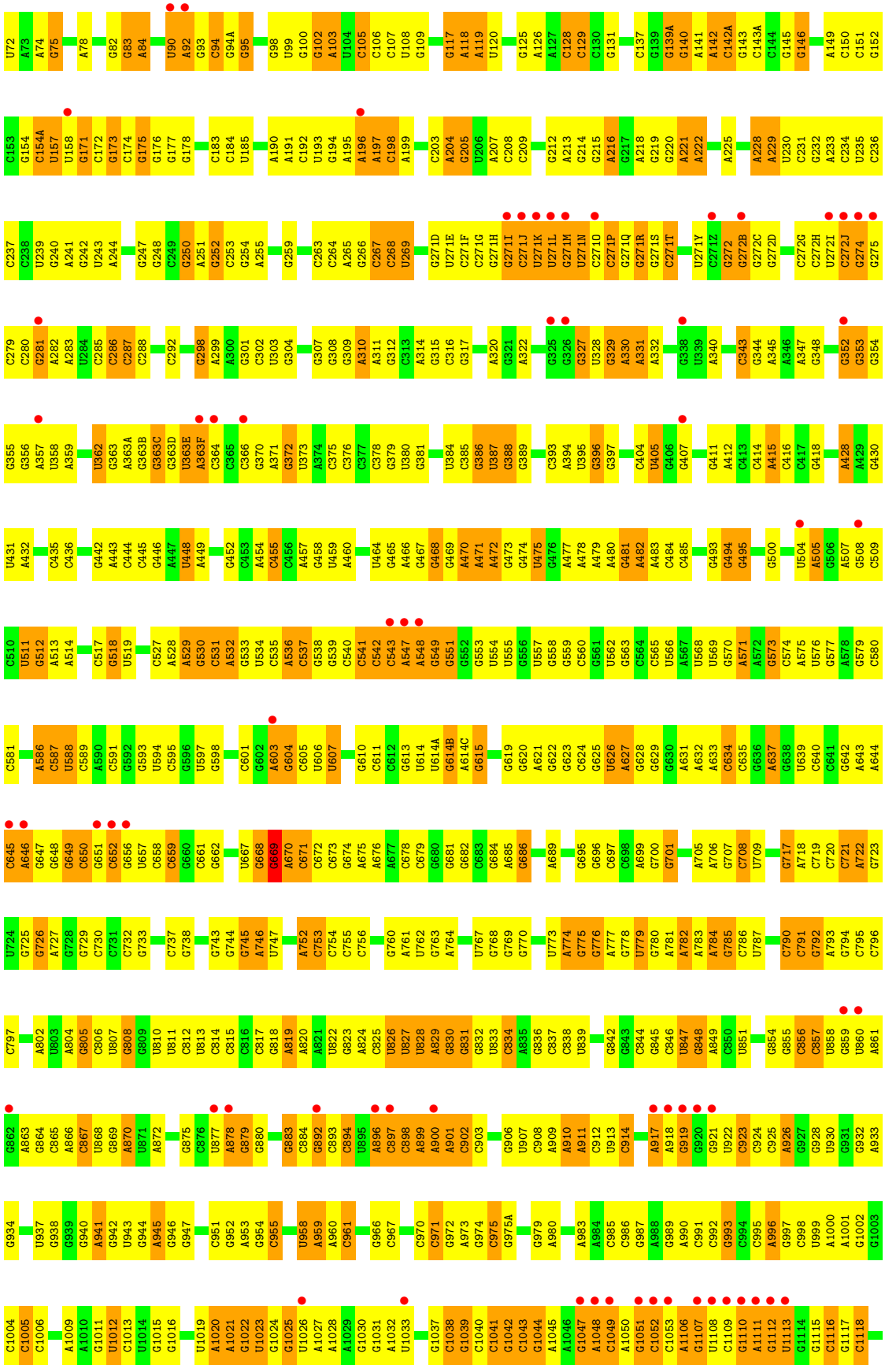
Chain D8:




• Molecule 31: 23S ribosomal RNA

Chain BA:

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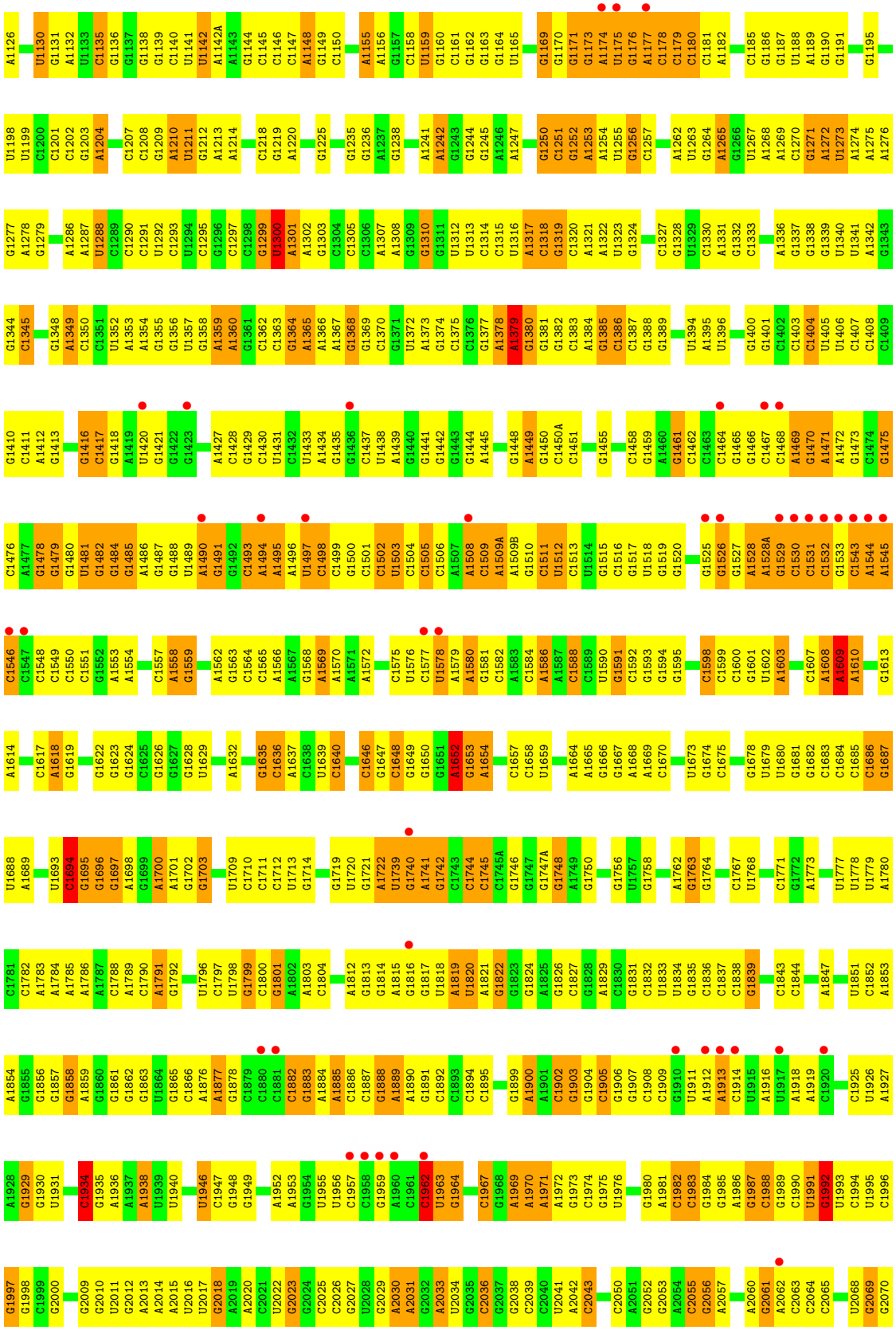
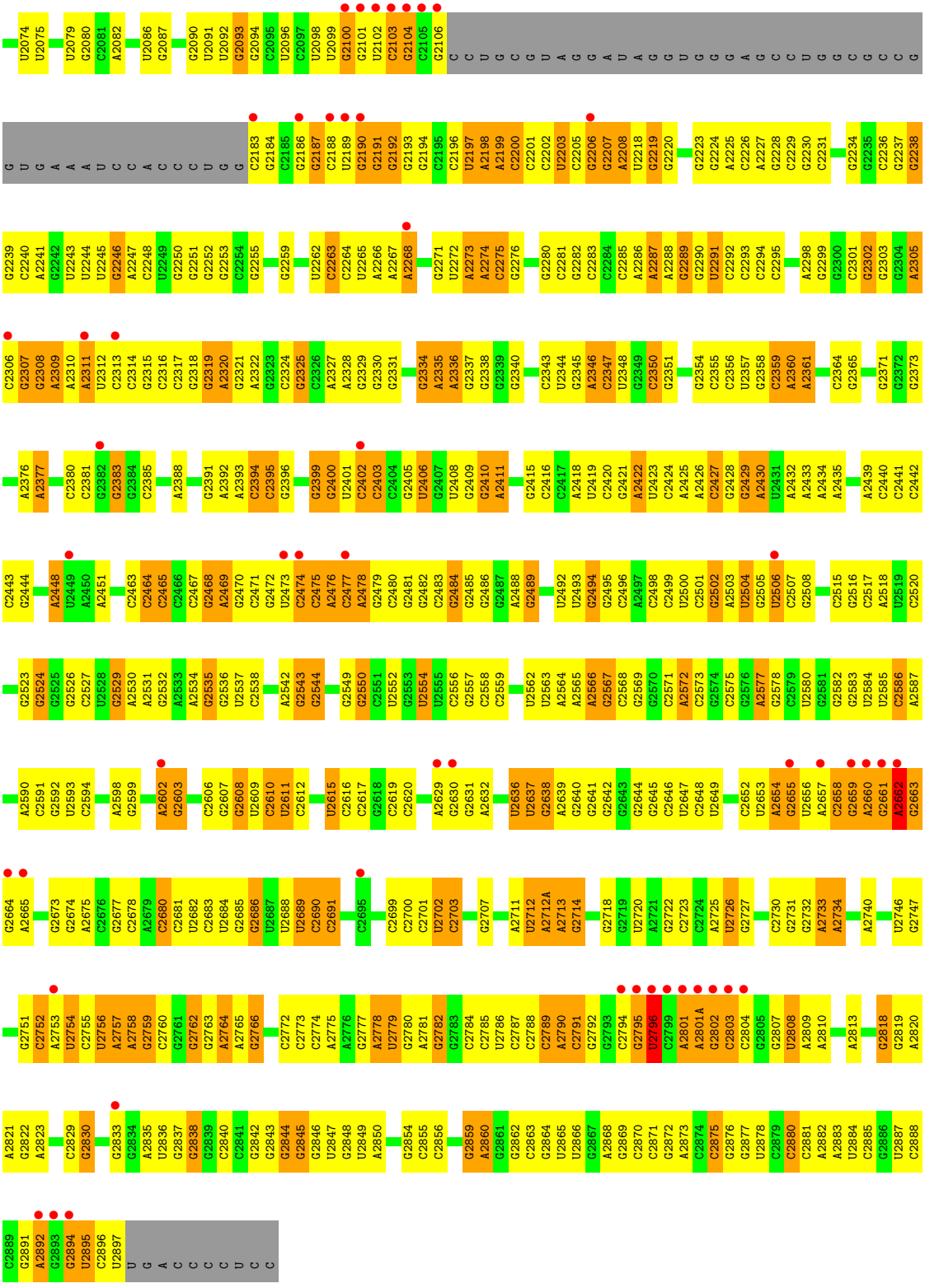
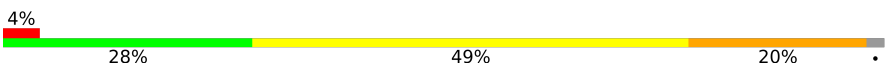
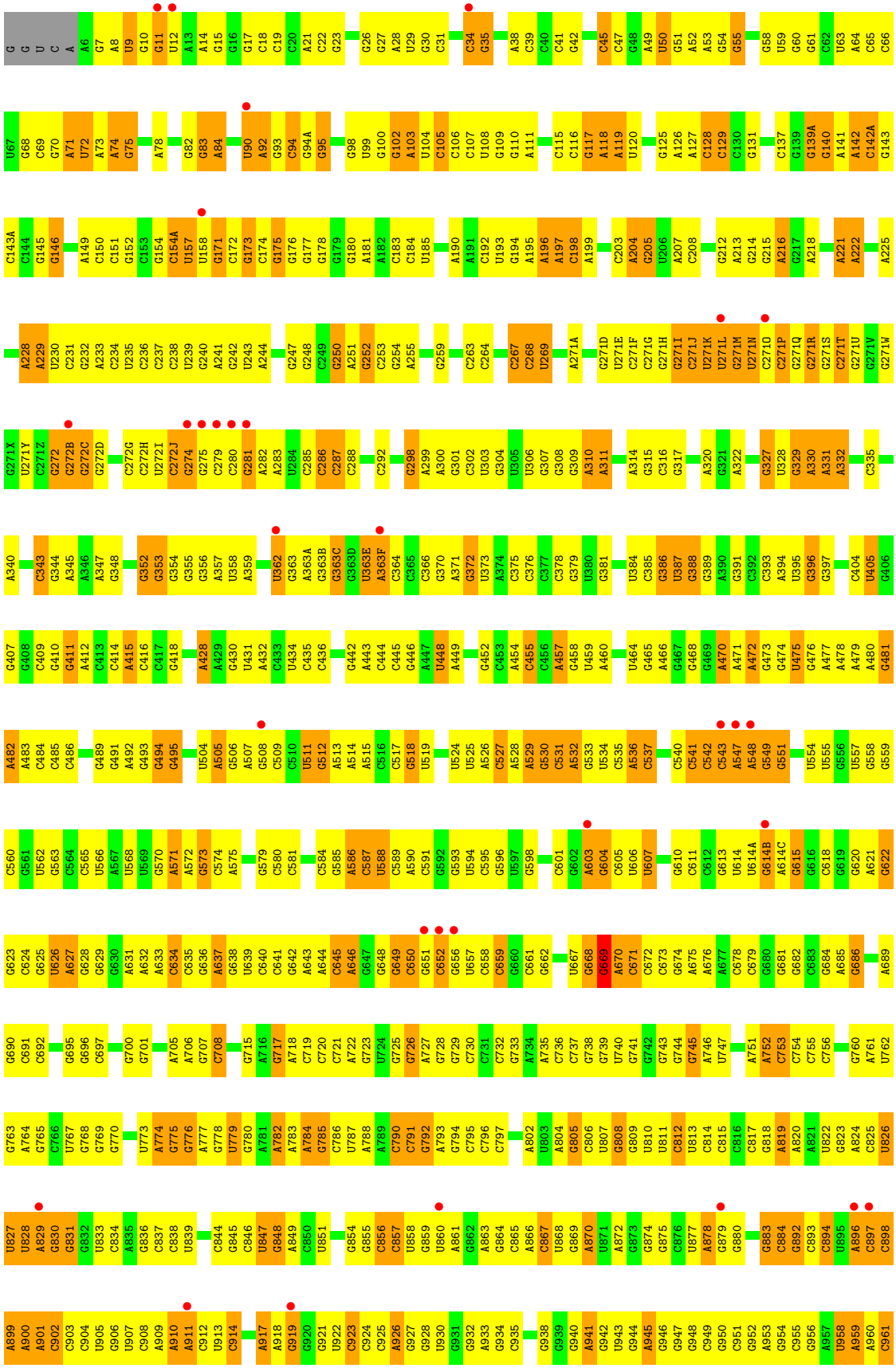


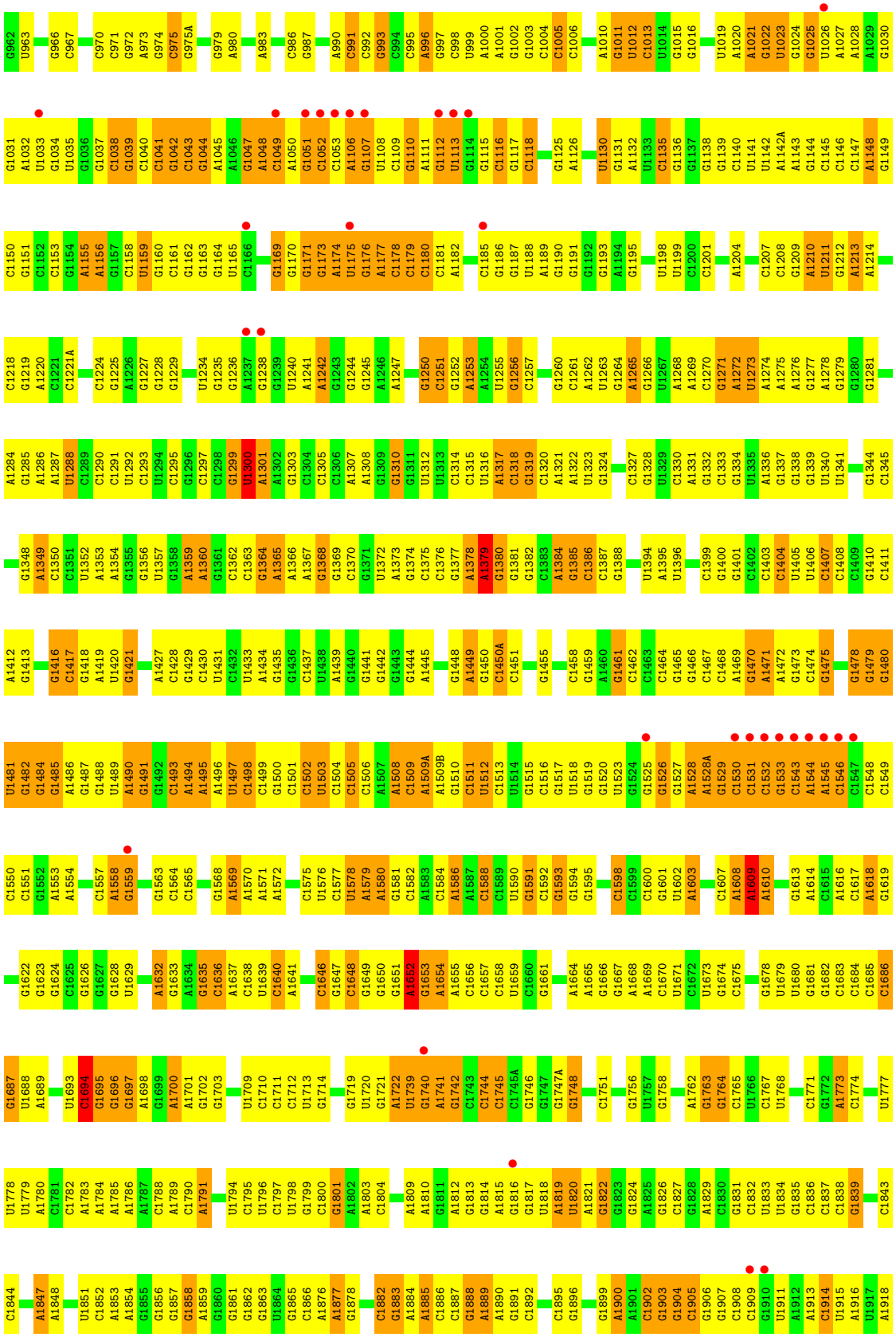


• Molecule 31: 23S ribosomal RNA

Chain DA:



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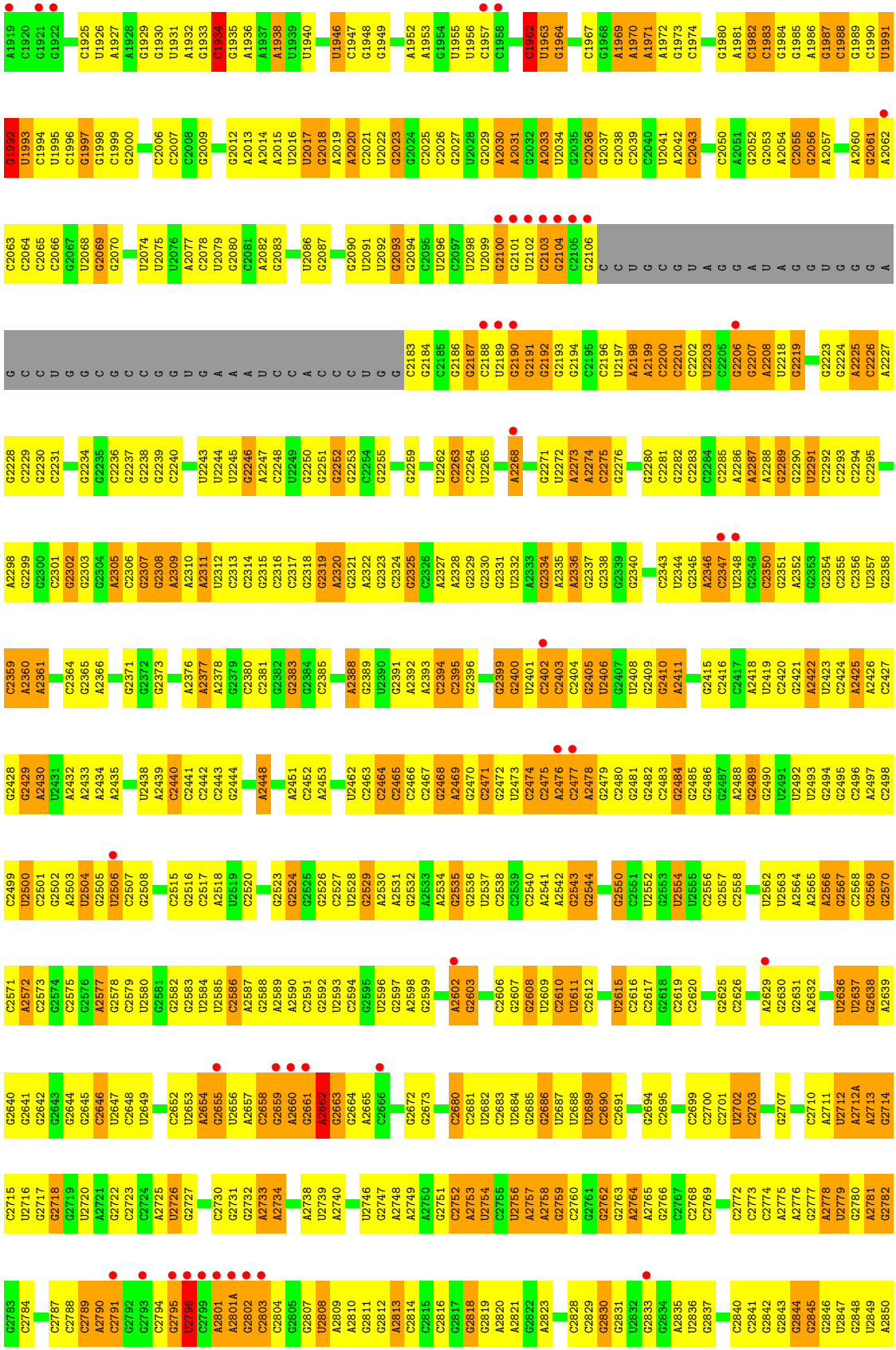

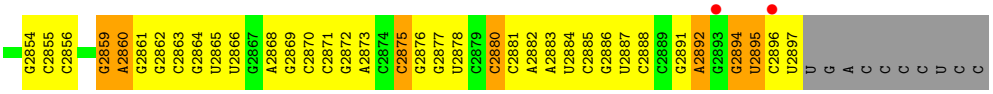


• Molecule 32: 5S ribosomal RNA

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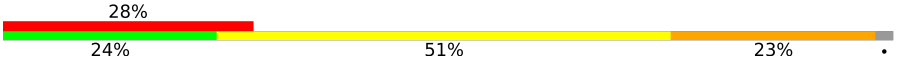


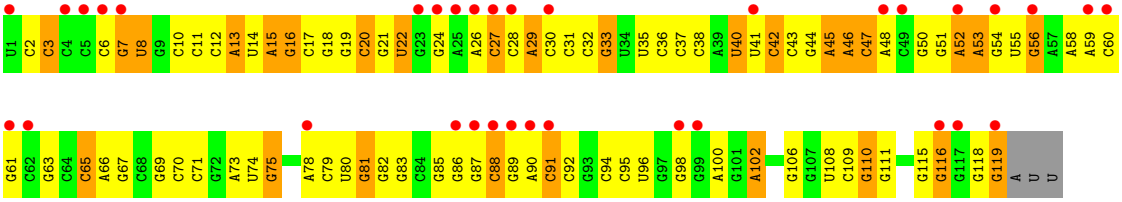


• Molecule 32: 5S ribosomal RNA

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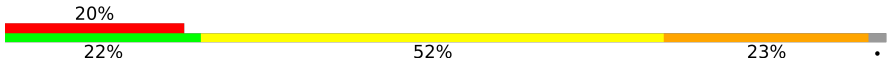
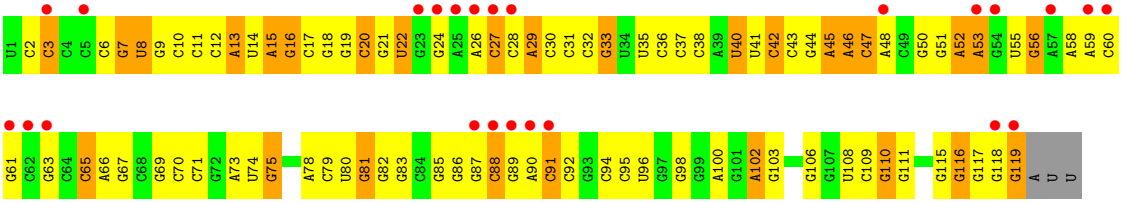
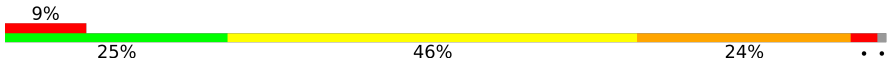
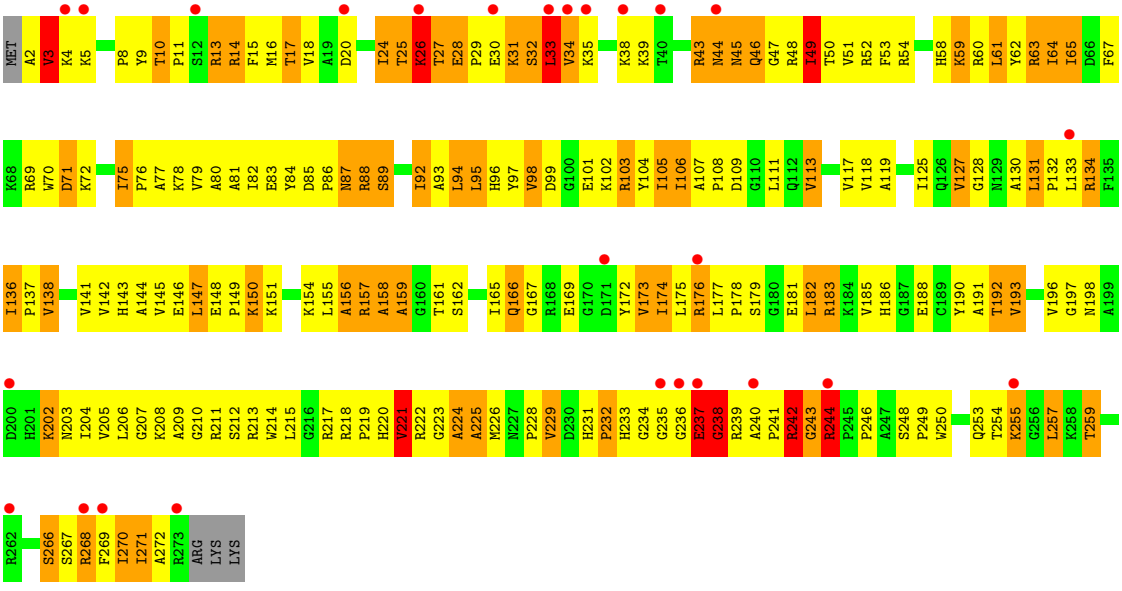



• Molecule 33: 50S ribosomal protein L2

Chain BD:




• Molecule 33: 50S ribosomal protein L2

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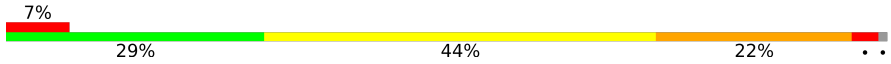

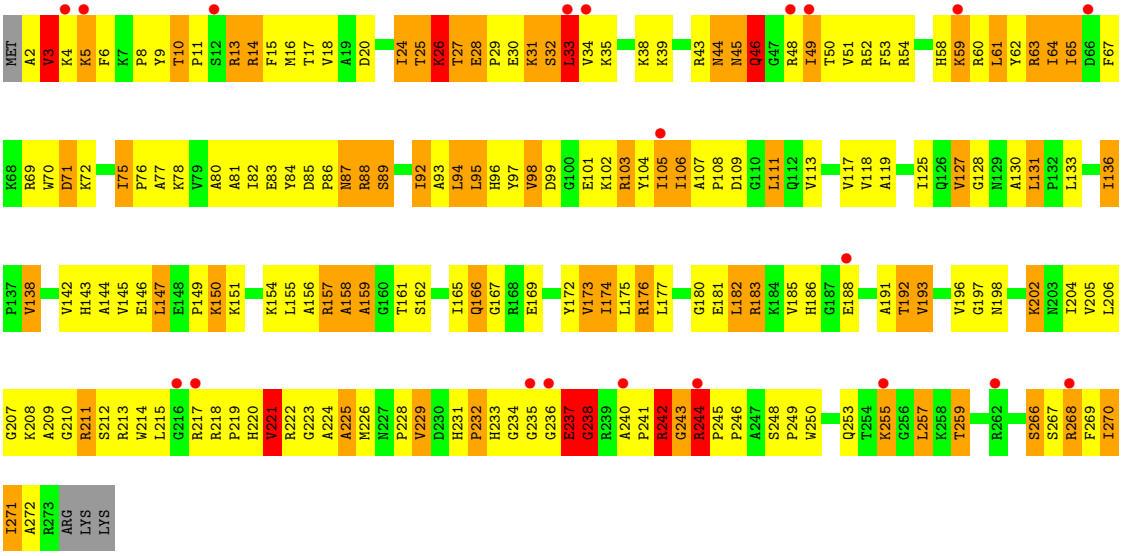

• Molecule 34: 50S ribosomal protein L3

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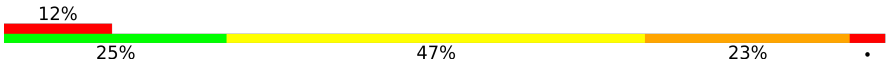


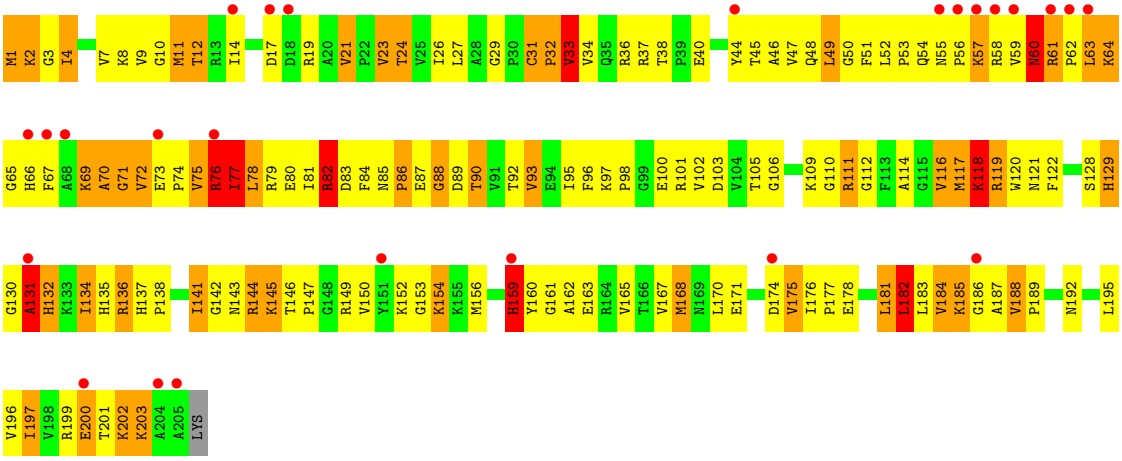

• Molecule 34: 50S ribosomal protein L3

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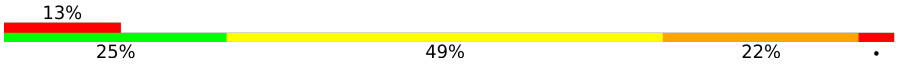
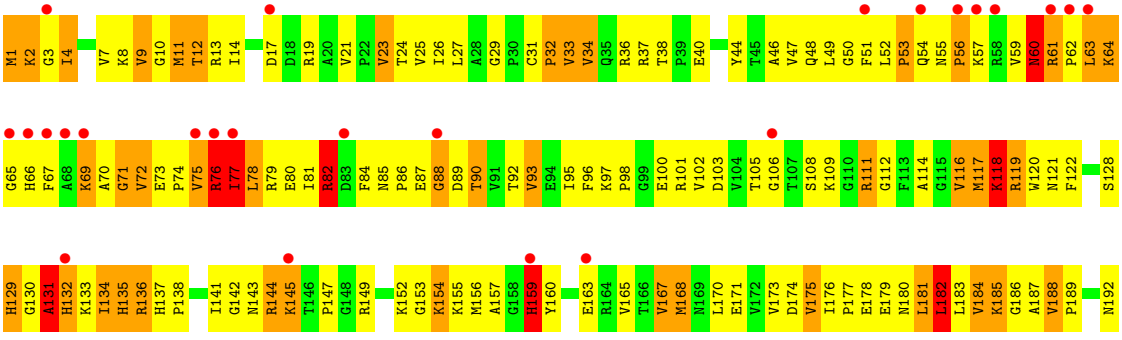
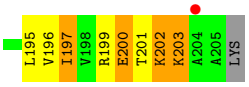
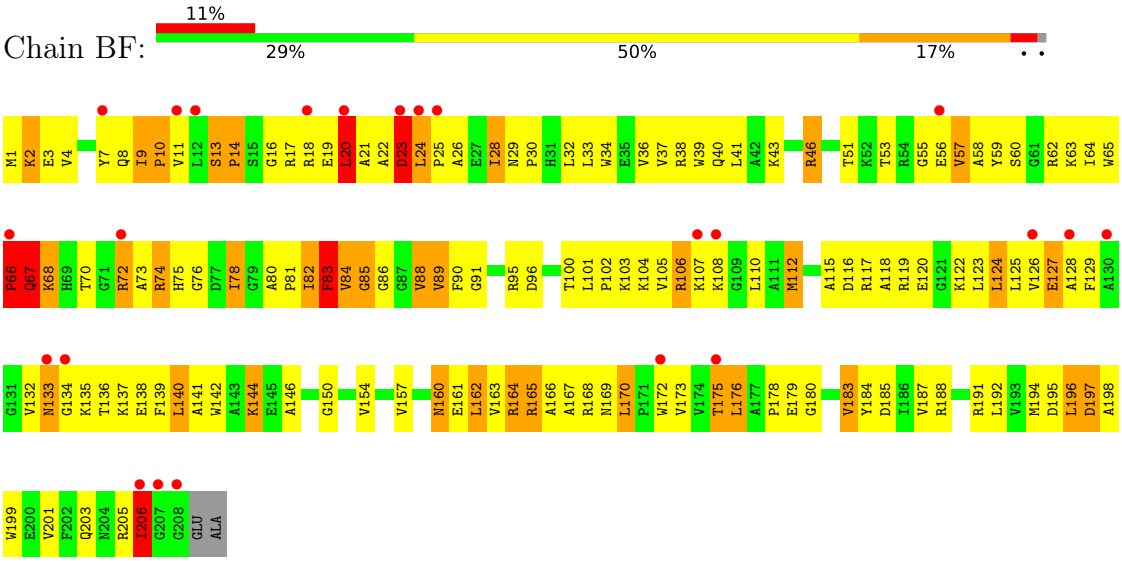




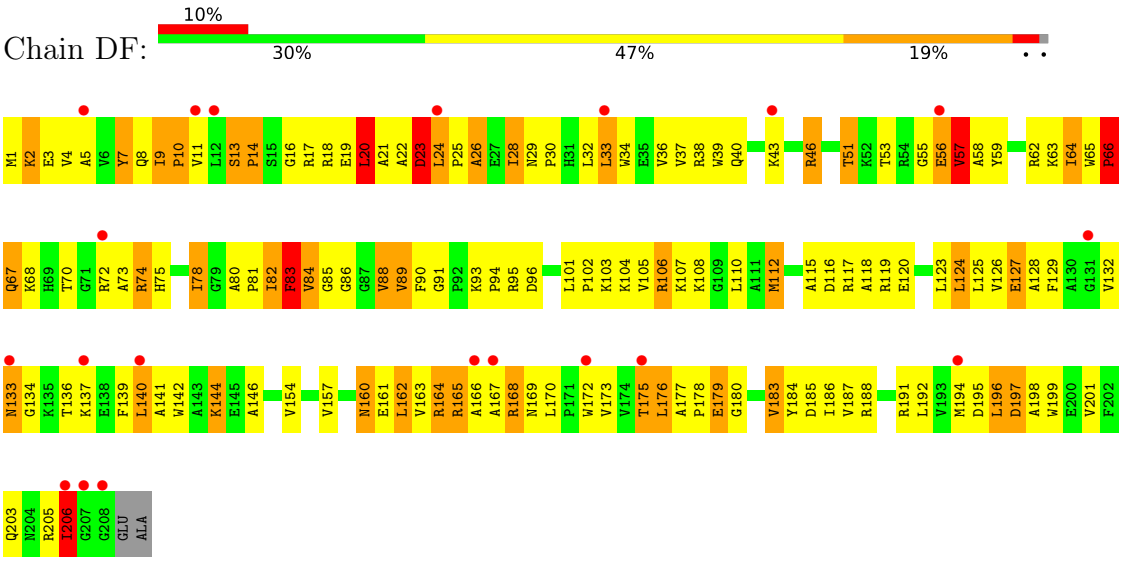

• Molecule 35: 50S ribosomal protein L4

Chain BF:



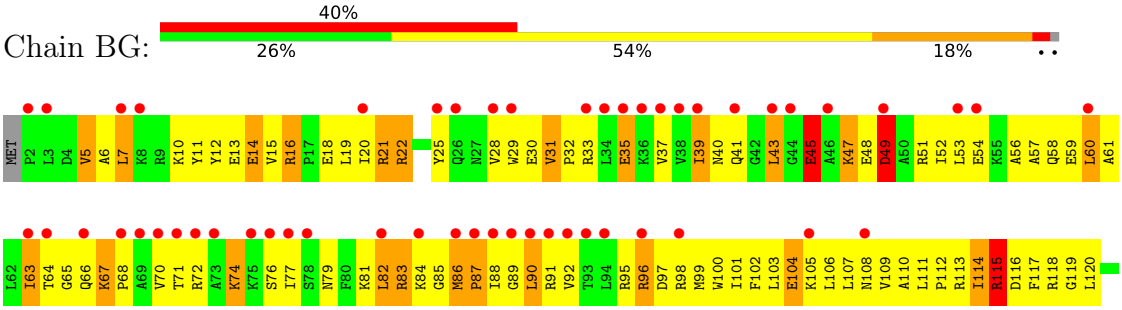

• Molecule 35: 50S ribosomal protein L4

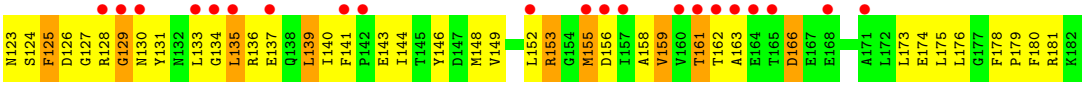
Chain DF:




• Molecule 36: 50S ribosomal protein L5

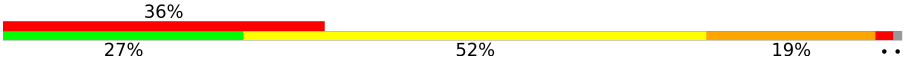
Chain BG:

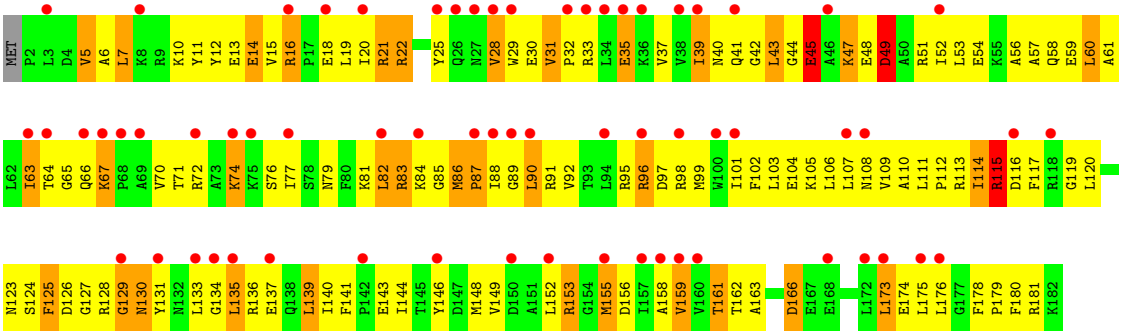




• Molecule 36: 50S ribosomal protein L5

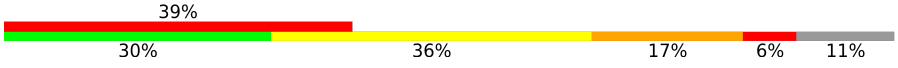
Chain DG:

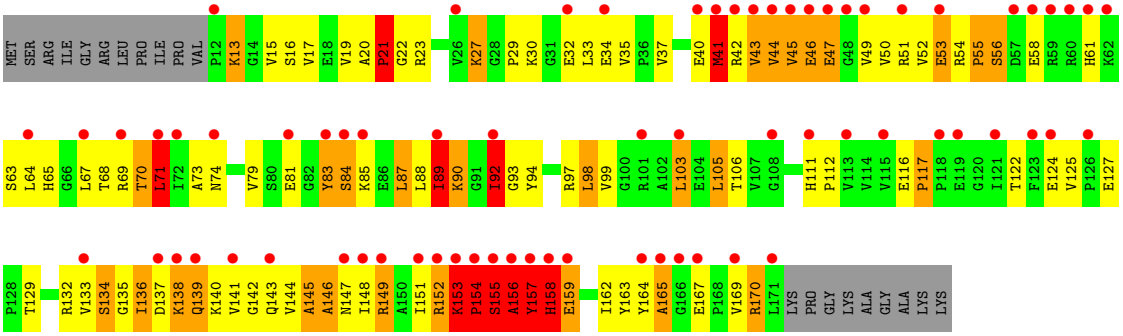


• Molecule 37: 50S ribosomal protein L6

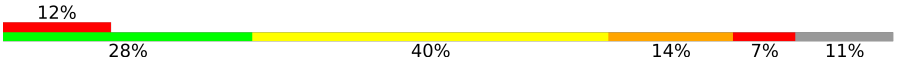
Chain BH:

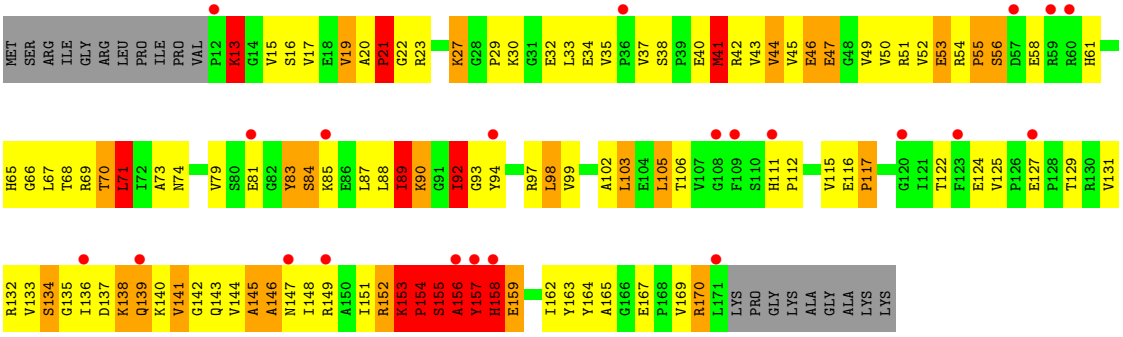


• Molecule 37: 50S ribosomal protein L6

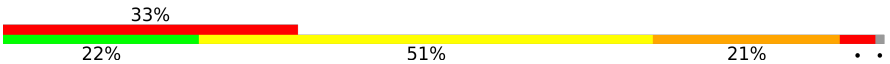
Chain DH:

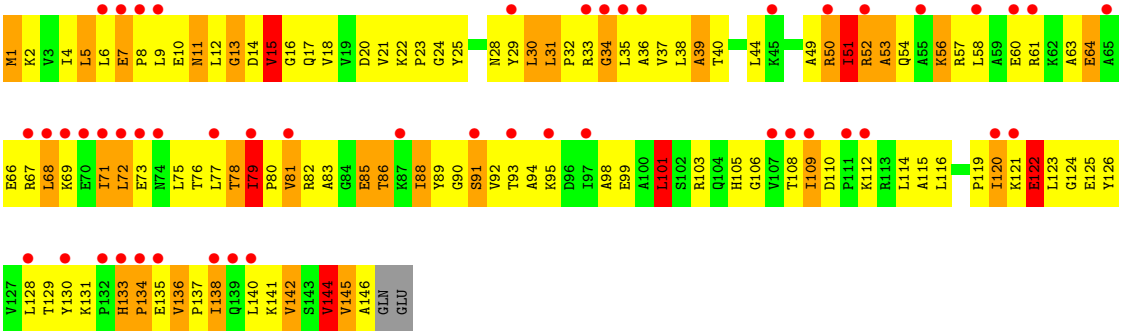


• Molecule 38: 50S ribosomal protein L9

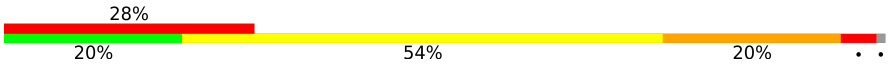
Chain BI:

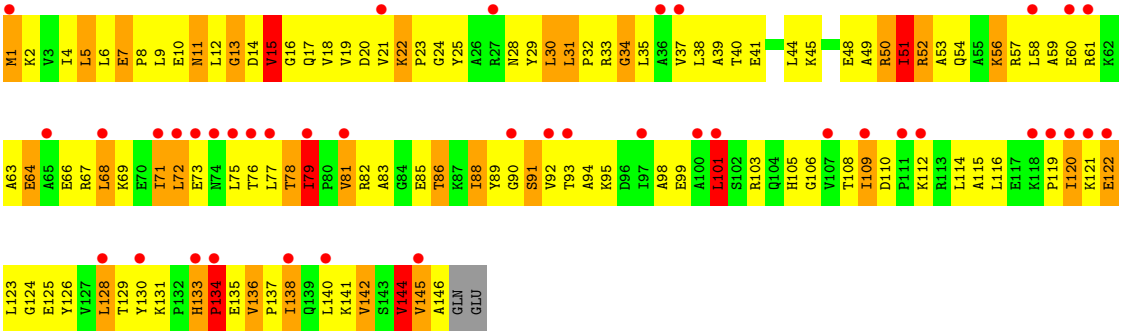




• Molecule 38: 50S ribosomal protein L9

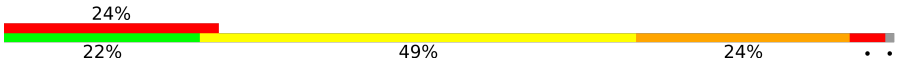
Chain DI:

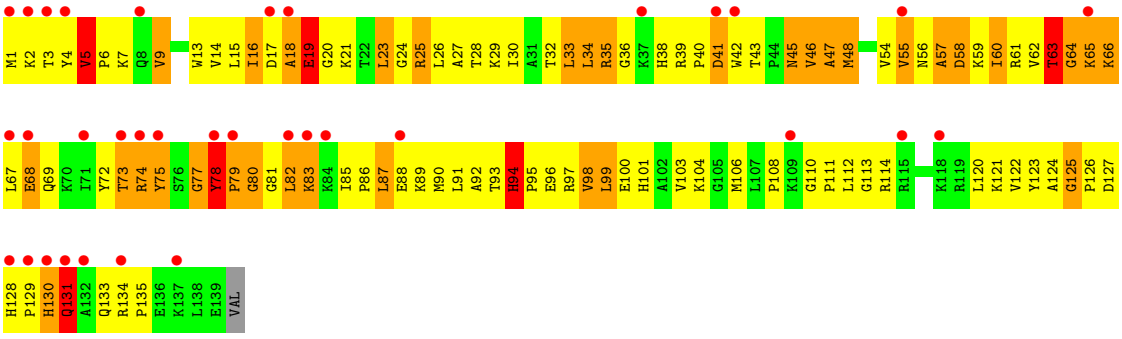


• Molecule 39: 50S ribosomal protein L13

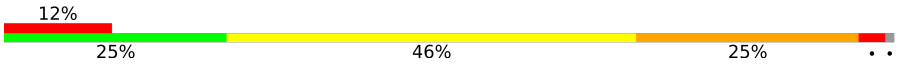
Chain BN:

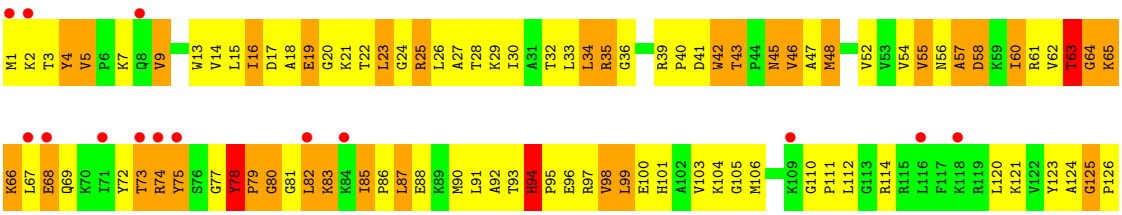


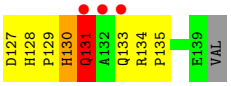



• Molecule 39: 50S ribosomal protein L13

Chain DN:

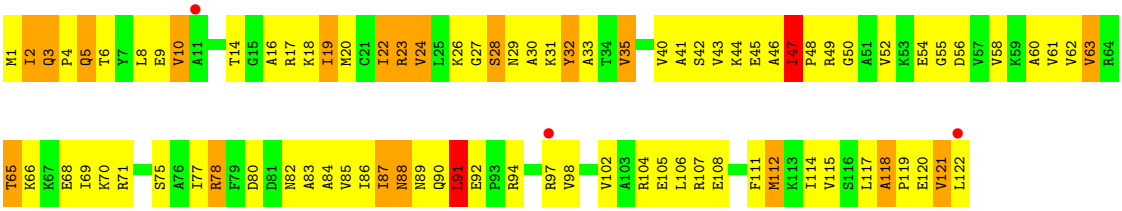




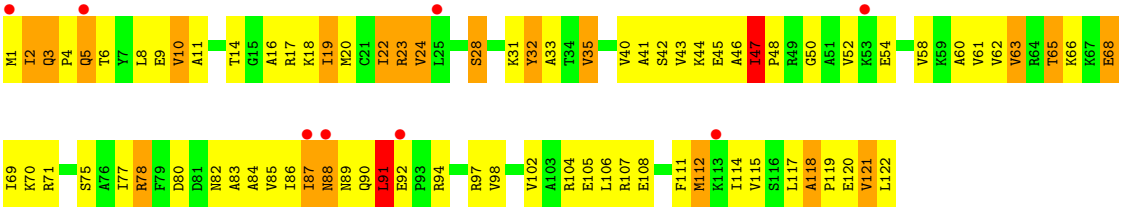

• Molecule 40: 50S ribosomal protein L14

Chain BO:



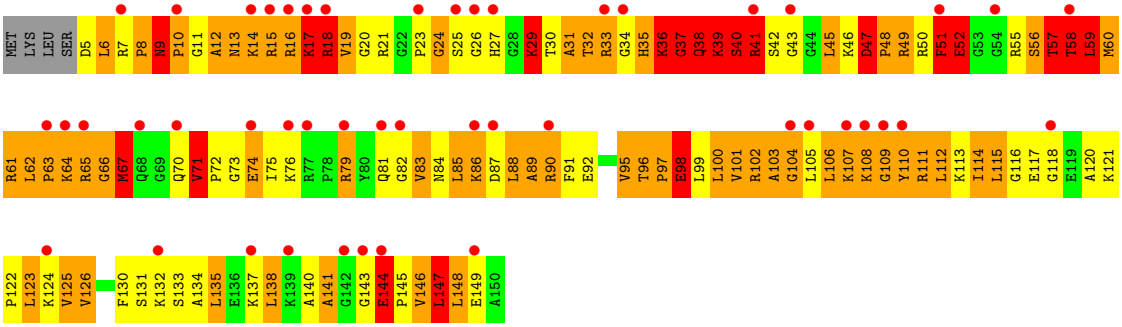
• Molecule 40: 50S ribosomal protein L14

Chain DO:



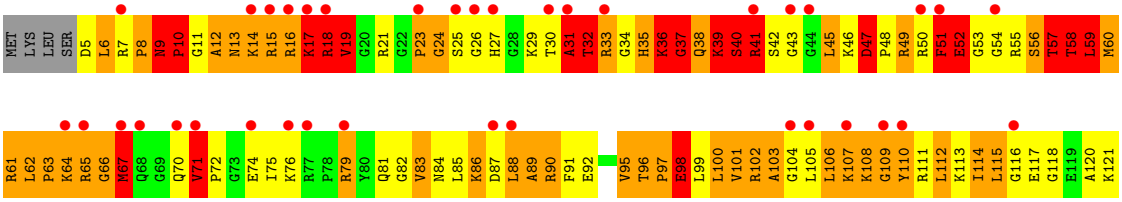
• Molecule 41: 50S ribosomal protein L15

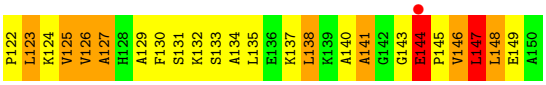
Chain BP:



• Molecule 41: 50S ribosomal protein L15

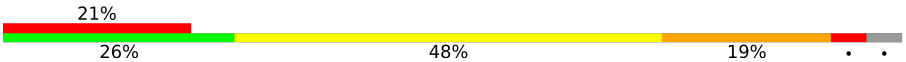
Chain DP:

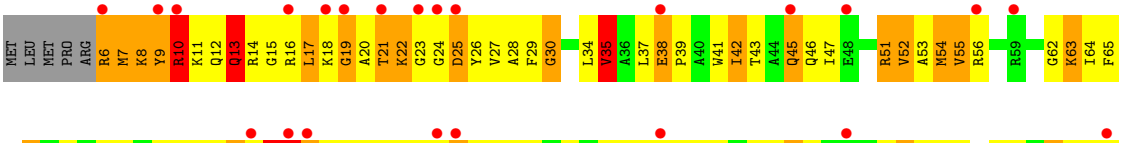


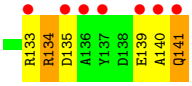



• Molecule 42: 50S ribosomal protein L16

Chain BQ:

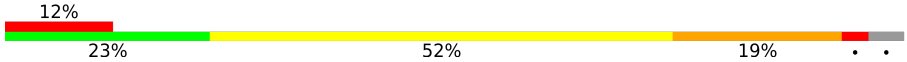


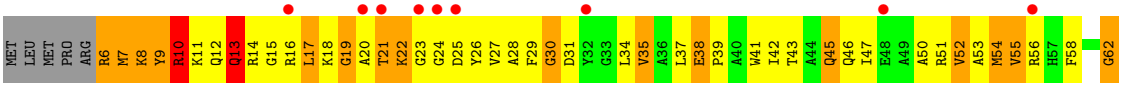



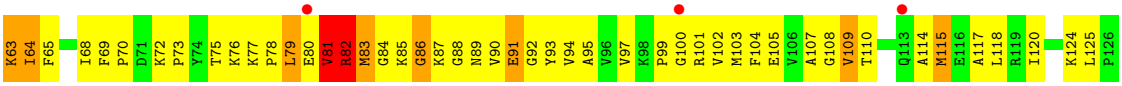


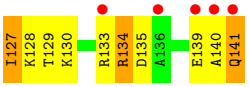

• Molecule 42: 50S ribosomal protein L16

Chain DQ:



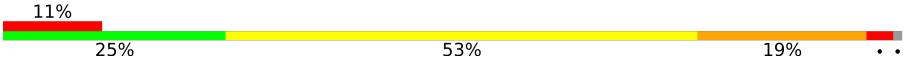



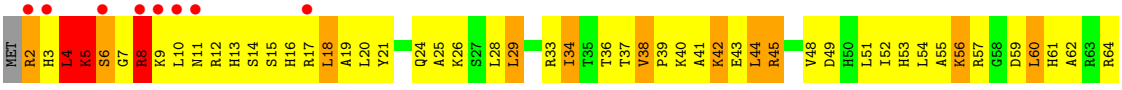



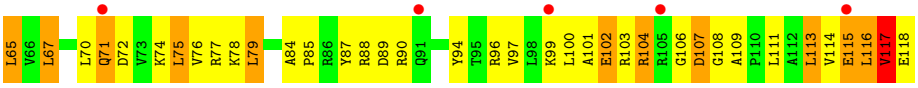




• Molecule 43: 50S ribosomal protein L17

Chain BR:

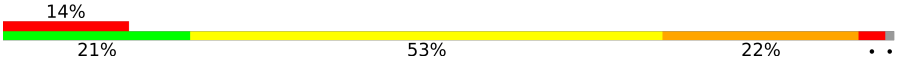


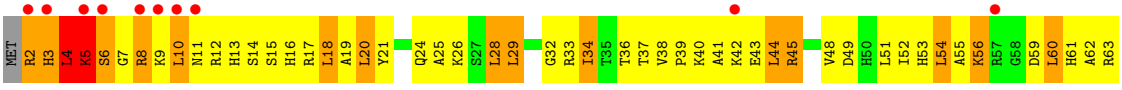



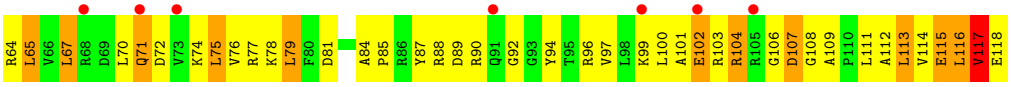




• Molecule 43: 50S ribosomal protein L17

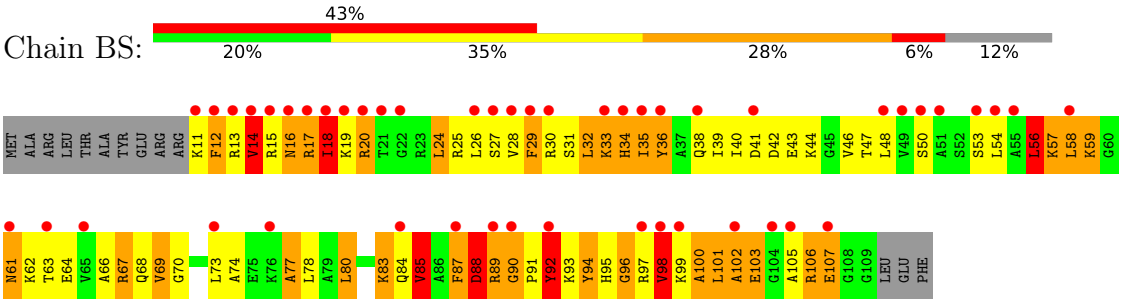
Chain DR:



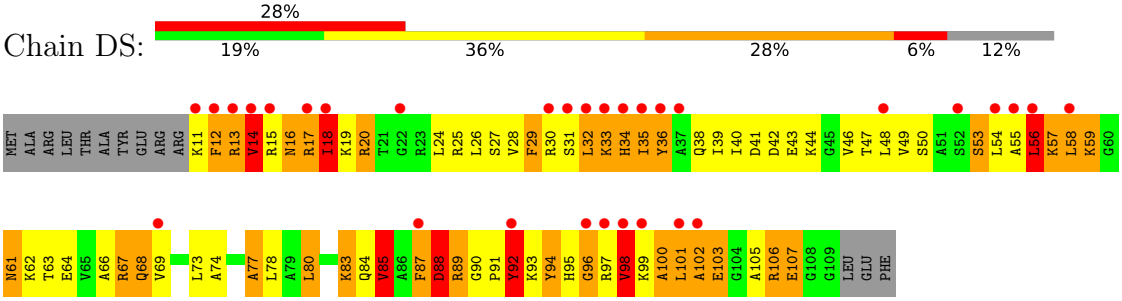





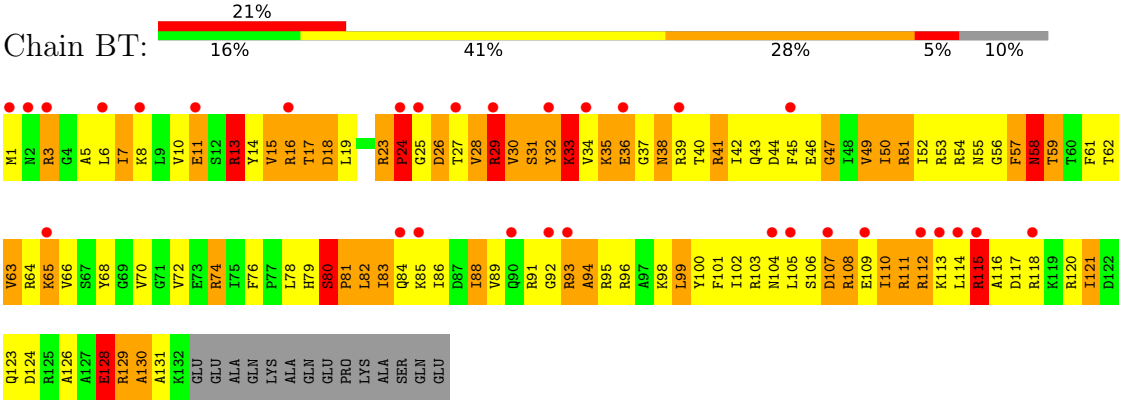

• Molecule 44: 50S ribosomal protein L18



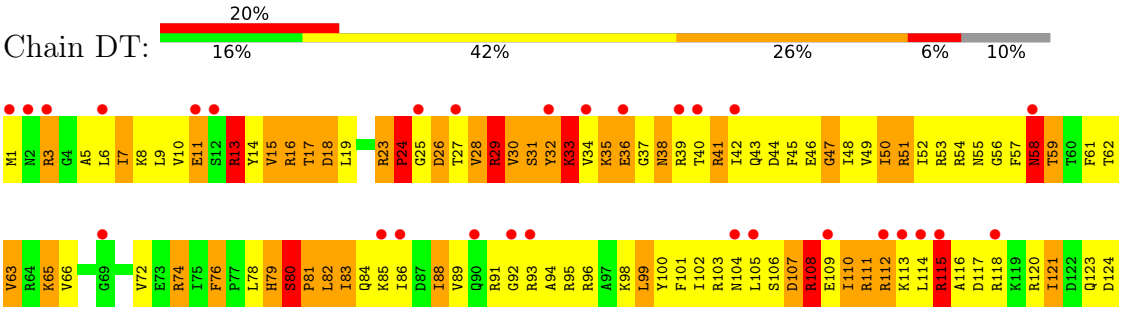

• Molecule 44: 50S ribosomal protein L18

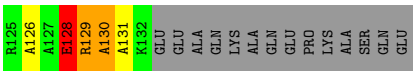



• Molecule 45: 50S ribosomal protein L19



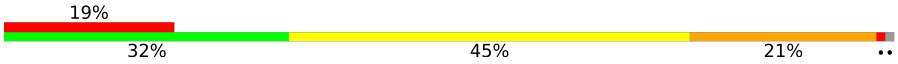

• Molecule 45: 50S ribosomal protein L19

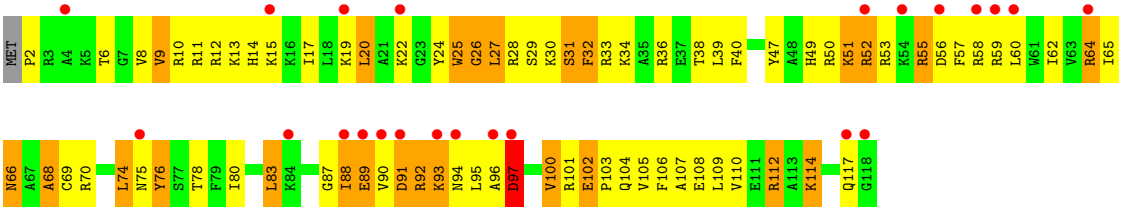





• Molecule 46: 50S ribosomal protein L20

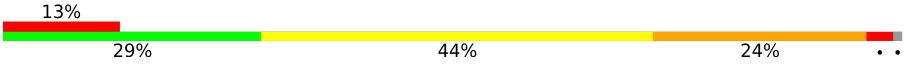
Chain BU:

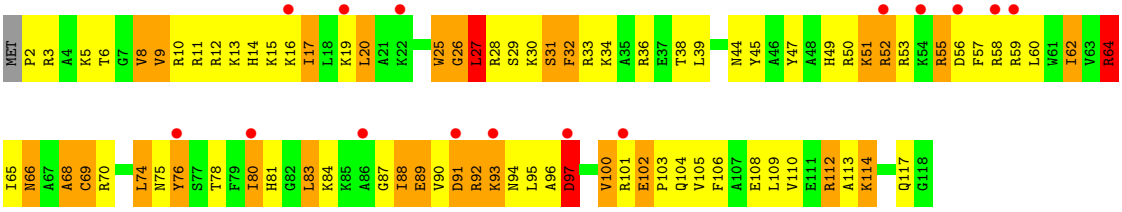



• Molecule 46: 50S ribosomal protein L20

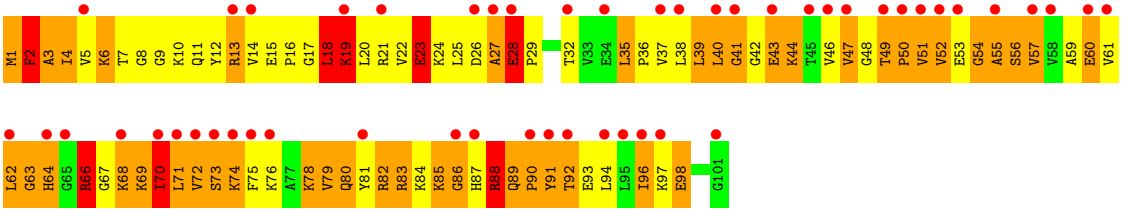
Chain DU:



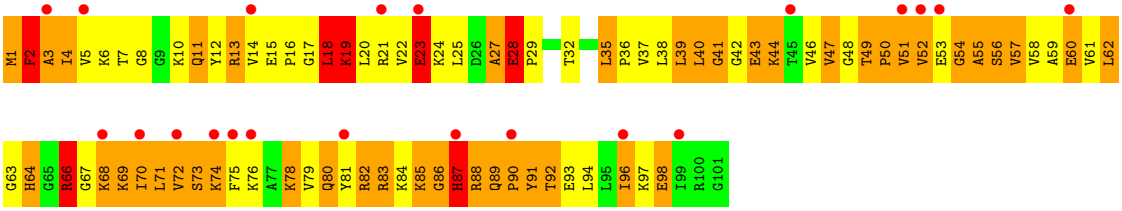

• Molecule 47: 50S ribosomal protein L21

Chain BV:



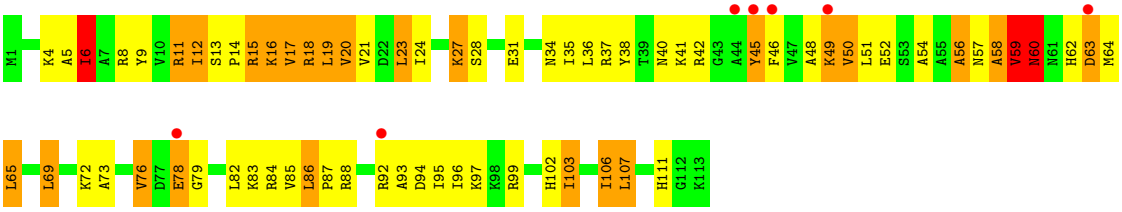

• Molecule 47: 50S ribosomal protein L21

Chain DV:



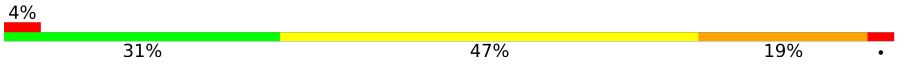

• Molecule 48: 50S ribosomal protein L22

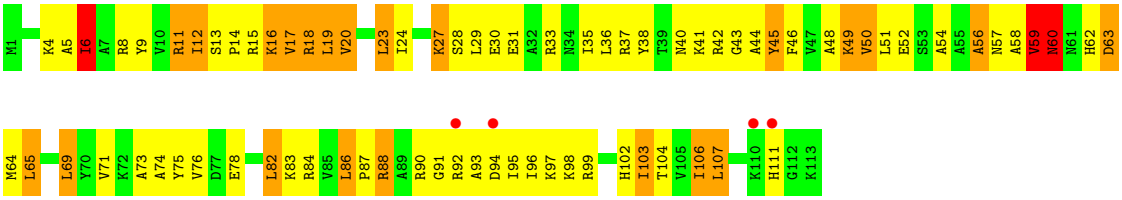
Chain BW:



• Molecule 48: 50S ribosomal protein L22

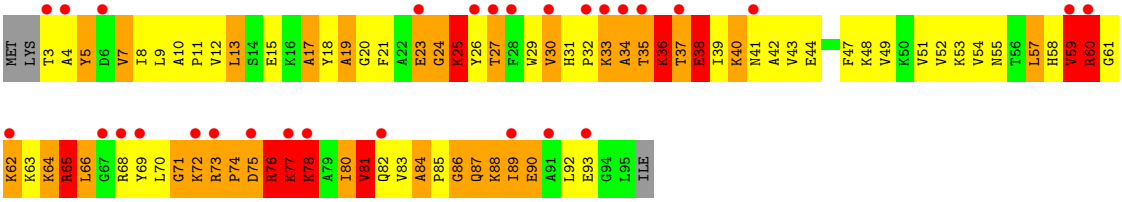
Chain DW:



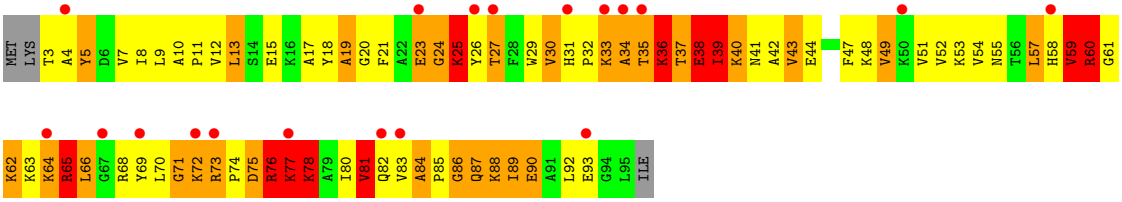
• Molecule 49: 50S ribosomal protein L23

Chain BX:



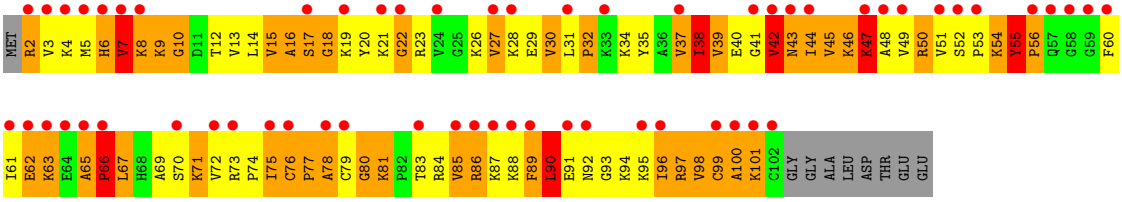
• Molecule 49: 50S ribosomal protein L23

Chain DX:



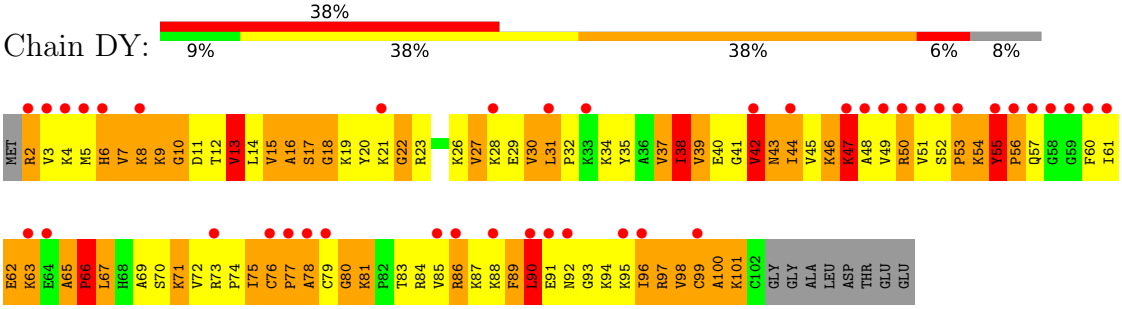
• Molecule 50: 50S ribosomal protein L24

Chain BY:



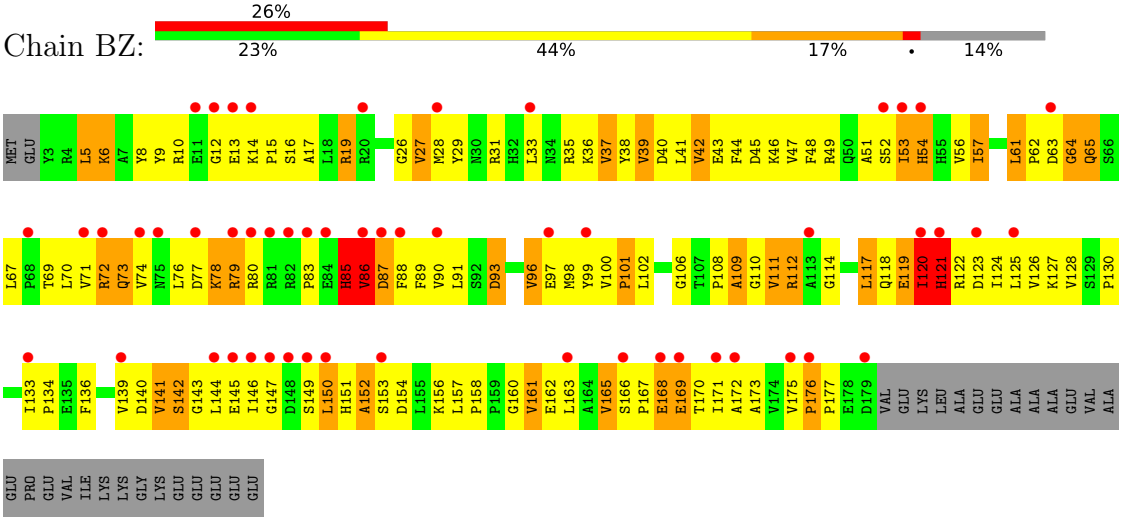
• Molecule 50: 50S ribosomal protein L24

Chain DY:



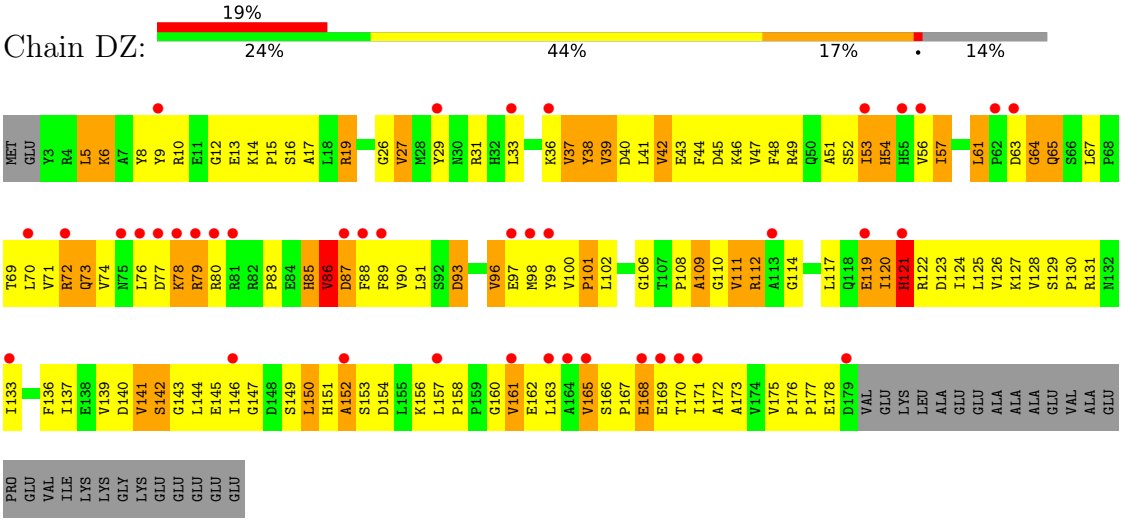

• Molecule 51: 50S ribosomal protein L25

Chain BZ:




• Molecule 51: 50S ribosomal protein L25

Chain DZ:



# 4 Data and refinement statistics (i)

| Property | Value | Source |
|---|---|---|
| Space group | P 21 21 21 | Depositor |
| Cell constants<br>a, b, c, $\alpha$, $\beta$, $\gamma$ | 210.22Å 450.25Å 623.81Å<br>90.00° 90.00° 90.00° | Depositor |
| Resolution (Å) | 49.80 – 3.00<br>49.80 – 3.00 | Depositor<br>EDS |
| % Data completeness<br>(in resolution range) | (Not available) (49.80-3.00)<br>88.7 (49.80-3.00) | Depositor<br>EDS |
| $R_{merge}$ | 0.24 | Depositor |
| $R_{sym}$ | (Not available) | Depositor |
| $< I/\sigma(I) >$ [1] | 1.09 (at 3.01Å) | Xtriage |
| Refinement program | PHENIX | Depositor |
| R, $R_{free}$ | 0.235 , 0.269<br>0.232 , 0.265 | Depositor<br>DCC |
| $R_{free}$ test set | 51892 reflections (4.45%) | wwPDB-VP |
| Wilson B-factor (Å$^2$) | 73.9 | Xtriage |
| Anisotropy | 0.352 | Xtriage |
| Bulk solvent $k_{sol}$(e/Å$^3$), $B_{sol}$(Å$^2$) | 0.28 , 101.0 | EDS |
| L-test for twinning[2] | $< \|L\| > = 0.43$, $< L^2 > = 0.25$ | Xtriage |
| Estimated twinning fraction | No twinning to report. | Xtriage |
| $F_o$,$F_c$ correlation | 0.90 | EDS |
| Total number of atoms | 278000 | wwPDB-VP |
| Average B, all atoms (Å$^2$) | 101.0 | wwPDB-VP |

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.49% of the height of the origin peak. No significant pseudotranslation is detected.*

[1] Intensities estimated from amplitudes.

[2] Theoretical values of $< |L| >$, $< L^2 >$ for acentric reflections are 0.5, 0.333 respectively for untwinned datasets, and 0.375, 0.2 for perfectly twinned datasets.

# 5 Model quality

## 5.1 Standard geometry

Bond lengths and bond angles in the following residue types are not validated in this section: K, MG, ZN, ZIT

The Z score for a bond length (or angle) is the number of standard deviations the observed value is removed from the expected value. A bond length (or angle) with $|Z| > 5$ is considered an outlier worth inspection. RMSZ is the root-mean-square of all Z scores of the bond lengths (or angles).

| Mol | Chain | Bond lengths | | Bond angles | |
|---|---|---|---|---|---|
| | | RMSZ | #$\|Z\|$ >5 | RMSZ | #$\|Z\|$ >5 |
| 1 | AA | 0.29 | 0/36190 | 0.39 | 0/56486 |
| 1 | CA | 0.29 | 0/36190 | 0.39 | 0/56486 |
| 2 | AB | 0.37 | 0/1936 | 0.86 | 4/2611 (0.2%) |
| 2 | CB | 0.37 | 0/1936 | 0.85 | 5/2611 (0.2%) |
| 3 | AC | 0.37 | 0/1637 | 0.84 | 5/2207 (0.2%) |
| 3 | CC | 0.36 | 0/1637 | 0.83 | 5/2207 (0.2%) |
| 4 | AD | 0.42 | 0/1733 | 0.87 | 6/2318 (0.3%) |
| 4 | CD | 0.42 | 0/1733 | 0.88 | 7/2318 (0.3%) |
| 5 | AE | 0.46 | 0/1163 | 0.90 | 1/1566 (0.1%) |
| 5 | CE | 0.43 | 0/1163 | 0.90 | 2/1566 (0.1%) |
| 6 | AF | 0.45 | 0/856 | 0.82 | 0/1154 |
| 6 | CF | 0.44 | 0/856 | 0.81 | 0/1154 |
| 7 | AG | 0.35 | 0/1276 | 0.78 | 0/1709 |
| 7 | CG | 0.36 | 0/1276 | 0.79 | 0/1709 |
| 8 | AH | 0.44 | 0/1136 | 0.88 | 2/1527 (0.1%) |
| 8 | CH | 0.43 | 0/1136 | 0.86 | 2/1527 (0.1%) |
| 9 | AI | 0.35 | 0/1028 | 0.76 | 0/1375 |
| 9 | CI | 0.35 | 0/1028 | 0.74 | 0/1375 |
| 10 | AJ | 0.41 | 0/808 | 0.83 | 0/1087 |
| 10 | CJ | 0.40 | 0/808 | 0.84 | 0/1087 |
| 11 | AK | 0.40 | 0/900 | 0.85 | 3/1213 (0.2%) |
| 11 | CK | 0.41 | 0/900 | 0.86 | 4/1213 (0.3%) |
| 12 | AL | 0.50 | 0/987 | 0.90 | 3/1322 (0.2%) |
| 12 | CL | 0.51 | 0/987 | 0.90 | 0/1322 |
| 13 | AM | 0.36 | 0/928 | 0.83 | 4/1238 (0.3%) |
| 13 | CM | 0.37 | 0/928 | 0.85 | 3/1238 (0.2%) |
| 14 | AN | 0.35 | 0/501 | 0.75 | 0/664 |
| 14 | CN | 0.34 | 0/501 | 0.75 | 0/664 |
| 15 | AO | 0.46 | 0/745 | 0.84 | 0/992 |
| 15 | CO | 0.42 | 0/745 | 0.85 | 0/992 |
| 16 | AP | 0.41 | 0/717 | 0.85 | 2/965 (0.2%) |
| 16 | CP | 0.44 | 0/717 | 0.85 | 3/965 (0.3%) |

| Mol | Chain | Bond lengths | | Bond angles | |
|---|---|---|---|---|---|
| | | RMSZ | #\|Z\| >5 | RMSZ | #\|Z\| >5 |
| 17 | AQ | 0.41 | 0/837 | 0.92 | 3/1119 (0.3%) |
| 17 | CQ | 0.42 | 0/837 | 0.94 | 4/1119 (0.4%) |
| 18 | AR | 0.43 | 0/579 | 0.98 | 4/768 (0.5%) |
| 18 | CR | 0.43 | 0/579 | 0.96 | 5/768 (0.7%) |
| 19 | AS | 0.40 | 0/643 | 0.78 | 0/867 |
| 19 | CS | 0.39 | 0/643 | 0.78 | 0/867 |
| 20 | AT | 0.44 | 0/765 | 0.91 | 1/1007 (0.1%) |
| 20 | CT | 0.44 | 0/765 | 0.90 | 1/1007 (0.1%) |
| 21 | AU | 0.37 | 0/213 | 0.78 | 0/279 |
| 21 | CU | 0.37 | 0/213 | 0.77 | 0/279 |
| 22 | B0 | 0.74 | 0/658 | 1.18 | 3/878 (0.3%) |
| 22 | D0 | 0.61 | 0/658 | 1.17 | 4/878 (0.5%) |
| 23 | B1 | 0.98 | 0/700 | 1.42 | 11/931 (1.2%) |
| 23 | D1 | 0.87 | 1/700 (0.1%) | 1.38 | 11/931 (1.2%) |
| 24 | B2 | 0.94 | 0/423 | 1.20 | 5/560 (0.9%) |
| 24 | D2 | 0.74 | 0/423 | 1.19 | 2/560 (0.4%) |
| 25 | B3 | 0.78 | 0/473 | 1.13 | 3/636 (0.5%) |
| 25 | D3 | 0.59 | 0/473 | 1.12 | 2/636 (0.3%) |
| 26 | B4 | 0.49 | 0/156 | 1.30 | 3/215 (1.4%) |
| 26 | D4 | 0.50 | 0/156 | 1.31 | 3/215 (1.4%) |
| 27 | B5 | 1.03 | 0/473 | 1.87 | 12/639 (1.9%) |
| 27 | D5 | 0.89 | 0/473 | 1.72 | 9/639 (1.4%) |
| 28 | B6 | 0.95 | 0/387 | 1.29 | 2/517 (0.4%) |
| 28 | D6 | 0.81 | 0/387 | 1.26 | 2/517 (0.4%) |
| 29 | B7 | 0.84 | 0/427 | 1.04 | 0/563 |
| 29 | D7 | 0.68 | 0/427 | 1.05 | 0/563 |
| 30 | B8 | 0.97 | 0/516 | 1.51 | 11/681 (1.6%) |
| 30 | D8 | 0.81 | 0/516 | 1.44 | 8/681 (1.2%) |
| 31 | BA | 0.61 | 5/65745 (0.0%) | 0.55 | 13/102639 (0.0%) |
| 31 | DA | 0.48 | 4/65745 (0.0%) | 0.53 | 12/102639 (0.0%) |
| 32 | BB | 0.50 | 0/2853 | 0.50 | 0/4451 |
| 32 | DB | 0.40 | 0/2853 | 0.48 | 0/4451 |
| 33 | BD | 0.80 | 0/2155 | 1.20 | 16/2907 (0.6%) |
| 33 | DD | 0.72 | 1/2155 (0.0%) | 1.19 | 13/2907 (0.4%) |
| 34 | BE | 0.83 | 1/1597 (0.1%) | 1.18 | 9/2155 (0.4%) |
| 34 | DE | 0.73 | 0/1597 | 1.16 | 7/2155 (0.3%) |
| 35 | BF | 0.83 | 0/1659 | 1.14 | 7/2246 (0.3%) |
| 35 | DF | 0.64 | 1/1659 (0.1%) | 1.12 | 10/2246 (0.4%) |
| 36 | BG | 0.43 | 0/1498 | 0.87 | 4/2013 (0.2%) |
| 36 | DG | 0.40 | 0/1498 | 0.86 | 4/2013 (0.2%) |
| 37 | BH | 0.79 | 0/1246 | 1.08 | 7/1684 (0.4%) |
| 37 | DH | 0.54 | 0/1246 | 1.01 | 6/1684 (0.4%) |
| 38 | BI | 0.56 | 0/1147 | 0.96 | 7/1553 (0.5%) |

| Mol | Chain | Bond lengths | | Bond angles | |
|---|---|---|---|---|---|
| | | RMSZ | #\|Z\| >5 | RMSZ | #\|Z\| >5 |
| 38 | DI | 0.51 | 0/1147 | 0.94 | 6/1553 (0.4%) |
| 39 | BN | 0.87 | 0/1132 | 1.26 | 14/1527 (0.9%) |
| 39 | DN | 0.66 | 0/1132 | 1.17 | 9/1527 (0.6%) |
| 40 | BO | 0.71 | 0/943 | 1.09 | 4/1269 (0.3%) |
| 40 | DO | 0.60 | 0/943 | 1.06 | 4/1269 (0.3%) |
| 41 | BP | 0.92 | 1/1131 (0.1%) | 1.42 | 14/1504 (0.9%) |
| 41 | DP | 0.77 | 1/1131 (0.1%) | 1.33 | 15/1504 (1.0%) |
| 42 | BQ | 0.86 | 0/1100 | 1.28 | 7/1470 (0.5%) |
| 42 | DQ | 0.73 | 0/1100 | 1.19 | 7/1470 (0.5%) |
| 43 | BR | 0.91 | 2/974 (0.2%) | 1.18 | 3/1302 (0.2%) |
| 43 | DR | 0.74 | 0/974 | 1.18 | 8/1302 (0.6%) |
| 44 | BS | 0.71 | 0/779 | 1.12 | 3/1038 (0.3%) |
| 44 | DS | 0.56 | 0/779 | 1.07 | 4/1038 (0.4%) |
| 45 | BT | 0.79 | 0/1114 | 1.29 | 16/1488 (1.1%) |
| 45 | DT | 0.66 | 1/1114 (0.1%) | 1.27 | 18/1488 (1.2%) |
| 46 | BU | 0.96 | 0/975 | 1.08 | 3/1297 (0.2%) |
| 46 | DU | 0.70 | 0/975 | 1.04 | 5/1297 (0.4%) |
| 47 | BV | 0.97 | 1/789 (0.1%) | 1.36 | 11/1054 (1.0%) |
| 47 | DV | 0.74 | 0/789 | 1.26 | 5/1054 (0.5%) |
| 48 | BW | 0.88 | 0/907 | 1.17 | 6/1216 (0.5%) |
| 48 | DW | 0.74 | 0/907 | 1.13 | 4/1216 (0.3%) |
| 49 | BX | 0.99 | 0/740 | 1.36 | 13/995 (1.3%) |
| 49 | DX | 0.85 | 1/740 (0.1%) | 1.30 | 11/995 (1.1%) |
| 50 | BY | 0.90 | 1/789 (0.1%) | 1.31 | 12/1053 (1.1%) |
| 50 | DY | 0.76 | 1/789 (0.1%) | 1.28 | 13/1053 (1.2%) |
| 51 | BZ | 0.63 | 0/1436 | 1.01 | 10/1951 (0.5%) |
| 51 | DZ | 0.49 | 0/1436 | 1.00 | 8/1951 (0.4%) |
| All | All | 0.52 | 22/301000 (0.0%) | 0.69 | 518/449812 (0.1%) |

Chiral center outliers are detected by calculating the chiral volume of a chiral center and verifying if the center is modelled as a planar moiety or with the opposite hand.A planarity outlier is detected by checking planarity of atoms in a peptide group, atoms in a mainchain group or atoms of a sidechain that are expected to be planar.

| Mol | Chain | #Chirality outliers | #Planarity outliers |
|---|---|---|---|
| 23 | B1 | 0 | 1 |
| 23 | D1 | 0 | 1 |
| 24 | B2 | 0 | 3 |
| 24 | D2 | 0 | 1 |
| 27 | B5 | 0 | 1 |
| 27 | D5 | 0 | 1 |
| 31 | BA | 21 | 0 |

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| Mol | Chain | #Chirality outliers | #Planarity outliers |
|---|---|---|---|
| 31 | DA | 21 | 0 |
| 33 | BD | 0 | 2 |
| 33 | DD | 0 | 2 |
| 34 | BE | 0 | 2 |
| 34 | DE | 0 | 2 |
| 35 | BF | 0 | 1 |
| 37 | BH | 0 | 2 |
| 37 | DH | 0 | 2 |
| 41 | BP | 0 | 5 |
| 41 | DP | 0 | 4 |
| 42 | BQ | 0 | 1 |
| 42 | DQ | 0 | 1 |
| 43 | BR | 0 | 1 |
| 43 | DR | 0 | 1 |
| 44 | BS | 0 | 1 |
| 44 | DS | 0 | 1 |
| 45 | BT | 0 | 1 |
| 45 | DT | 0 | 1 |
| 47 | BV | 0 | 1 |
| 47 | DV | 0 | 2 |
| 49 | BX | 0 | 3 |
| 49 | DX | 0 | 3 |
| All | All | 42 | 47 |

All (22) bond length outliers are listed below:

| Mol | Chain | Res | Type | Atoms | Z | Observed(Å) | Ideal(Å) |
|---|---|---|---|---|---|---|---|
| 31 | BA | 669 | G | C4'-C3' | -8.47 | 1.40 | 1.53 |
| 31 | DA | 669 | G | C4'-C3' | -6.99 | 1.42 | 1.53 |
| 31 | BA | 1300 | U | C4'-C3' | -6.73 | 1.43 | 1.53 |
| 43 | BR | 38 | VAL | CA-CB | -6.71 | 1.50 | 1.54 |
| 31 | DA | 1300 | U | C4'-C3' | -6.64 | 1.43 | 1.53 |
| 50 | BY | 45 | VAL | CA-CB | 6.21 | 1.65 | 1.55 |
| 31 | BA | 1694 | C | C4'-C3' | -5.96 | 1.43 | 1.52 |
| 34 | BE | 21 | VAL | CA-CB | -5.92 | 1.50 | 1.54 |
| 31 | DA | 1694 | C | C4'-C3' | -5.91 | 1.43 | 1.52 |
| 31 | BA | 652 | C | O3'-P | 5.83 | 1.70 | 1.61 |
| 45 | DT | 79 | HIS | CA-C | 5.73 | 1.59 | 1.52 |
| 50 | DY | 11 | ASP | N-CA | 5.71 | 1.49 | 1.46 |
| 43 | BR | 5 | LYS | CA-C | 5.55 | 1.60 | 1.52 |
| 41 | BP | 39 | LYS | CB-CG | 5.53 | 1.69 | 1.52 |
| 47 | BV | 70 | ILE | CA-CB | 5.44 | 1.61 | 1.54 |

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| Mol | Chain | Res | Type | Atoms | Z | Observed(Å) | Ideal(Å) |
|---|---|---|---|---|---|---|---|
| 49 | DX | 59 | VAL | CA-C | 5.37 | 1.59 | 1.52 |
| 31 | BA | 652 | C | C3'-O3' | 5.21 | 1.50 | 1.43 |
| 31 | DA | 652 | C | O3'-P | 5.21 | 1.69 | 1.61 |
| 35 | DF | 57 | VAL | CA-CB | -5.20 | 1.48 | 1.54 |
| 33 | DD | 46 | GLN | CA-C | 5.15 | 1.59 | 1.53 |
| 41 | DP | 31 | ALA | CA-CB | 5.15 | 1.62 | 1.53 |
| 23 | D1 | 14 | VAL | CA-C | 5.06 | 1.59 | 1.52 |

All (518) bond angle outliers are listed below:

| Mol | Chain | Res | Type | Atoms | Z | Observed($^o$) | Ideal($^o$) |
|---|---|---|---|---|---|---|---|
| 27 | B5 | 46 | CYS | CA-C-N | 16.33 | 140.25 | 119.84 |
| 27 | B5 | 46 | CYS | C-N-CA | 16.33 | 140.25 | 119.84 |
| 27 | D5 | 46 | CYS | CA-C-N | 14.91 | 138.48 | 119.84 |
| 27 | D5 | 46 | CYS | C-N-CA | 14.91 | 138.48 | 119.84 |
| 41 | BP | 59 | LEU | N-CA-C | -13.90 | 95.64 | 112.59 |
| 41 | DP | 59 | LEU | N-CA-C | -12.39 | 97.48 | 112.59 |
| 45 | BT | 29 | ARG | N-CA-C | 11.05 | 123.30 | 108.38 |
| 22 | B0 | 13 | GLY | N-CA-C | -11.04 | 100.19 | 112.33 |
| 45 | DT | 29 | ARG | N-CA-C | 10.96 | 122.71 | 108.24 |
| 42 | DQ | 81 | VAL | N-CA-C | 10.72 | 120.08 | 111.62 |
| 47 | BV | 49 | THR | CA-C-N | 10.51 | 132.98 | 119.84 |
| 47 | BV | 49 | THR | C-N-CA | 10.51 | 132.98 | 119.84 |
| 27 | D5 | 4 | HIS | CA-C-N | -10.35 | 109.12 | 119.78 |
| 27 | D5 | 4 | HIS | C-N-CA | -10.35 | 109.12 | 119.78 |
| 42 | BQ | 10 | ARG | N-CA-C | 10.20 | 124.58 | 109.07 |
| 49 | BX | 87 | GLN | N-CA-C | 10.07 | 124.85 | 109.23 |
| 42 | BQ | 81 | VAL | N-CA-C | 9.94 | 119.47 | 111.62 |
| 49 | DX | 87 | GLN | N-CA-C | 9.91 | 124.59 | 109.23 |
| 31 | BA | 1379 | A | N9-C1'-C2' | 9.81 | 126.72 | 112.00 |
| 31 | DA | 1379 | A | N9-C1'-C2' | 9.72 | 126.58 | 112.00 |
| 47 | DV | 49 | THR | CA-C-N | 9.67 | 131.93 | 119.84 |
| 47 | DV | 49 | THR | C-N-CA | 9.67 | 131.93 | 119.84 |
| 27 | B5 | 4 | HIS | CA-C-N | -9.53 | 110.22 | 119.85 |
| 27 | B5 | 4 | HIS | C-N-CA | -9.53 | 110.22 | 119.85 |
| 33 | BD | 268 | ARG | N-CA-C | -9.52 | 99.26 | 113.72 |
| 42 | DQ | 10 | ARG | N-CA-C | 9.40 | 123.36 | 109.07 |
| 22 | D0 | 13 | GLY | N-CA-C | -9.37 | 102.02 | 112.33 |
| 45 | BT | 23 | ARG | CA-C-N | 9.30 | 131.47 | 119.84 |
| 45 | BT | 23 | ARG | C-N-CA | 9.30 | 131.47 | 119.84 |
| 51 | DZ | 54 | HIS | N-CA-C | -9.27 | 104.05 | 114.62 |
| 38 | BI | 54 | GLN | N-CA-C | -9.16 | 101.66 | 112.92 |

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| Mol | Chain | Res | Type | Atoms | Z | Observed($^o$) | Ideal($^o$) |
|---|---|---|---|---|---|---|---|
| 45 | DT | 23 | ARG | CA-C-N | 9.08 | 131.19 | 119.84 |
| 45 | DT | 23 | ARG | C-N-CA | 9.08 | 131.19 | 119.84 |
| 39 | BN | 35 | ARG | N-CA-C | -9.05 | 98.04 | 110.35 |
| 41 | BP | 41 | ARG | N-CA-C | -8.95 | 92.32 | 107.80 |
| 30 | B8 | 19 | SER | N-CA-C | -8.93 | 102.39 | 113.38 |
| 30 | B8 | 33 | ASN | N-CA-C | -8.89 | 95.50 | 113.29 |
| 27 | B5 | 31 | VAL | CA-C-N | 8.86 | 130.91 | 119.84 |
| 27 | B5 | 31 | VAL | C-N-CA | 8.86 | 130.91 | 119.84 |
| 50 | DY | 65 | ALA | CA-C-N | 8.85 | 130.90 | 119.84 |
| 50 | DY | 65 | ALA | C-N-CA | 8.85 | 130.90 | 119.84 |
| 39 | DN | 35 | ARG | N-CA-C | -8.76 | 98.15 | 110.50 |
| 51 | BZ | 54 | HIS | N-CA-C | -8.76 | 104.64 | 114.62 |
| 38 | DI | 54 | GLN | N-CA-C | -8.68 | 102.25 | 112.92 |
| 28 | D6 | 48 | VAL | N-CA-C | 8.65 | 121.61 | 112.96 |
| 40 | BO | 118 | ALA | CA-C-N | 8.62 | 128.61 | 119.05 |
| 40 | BO | 118 | ALA | C-N-CA | 8.62 | 128.61 | 119.05 |
| 41 | BP | 66 | GLY | N-CA-C | -8.60 | 100.79 | 112.57 |
| 49 | BX | 75 | ASP | N-CA-C | -8.55 | 99.01 | 110.55 |
| 49 | BX | 5 | TYR | N-CA-C | -8.44 | 102.00 | 113.30 |
| 33 | DD | 268 | ARG | N-CA-C | -8.40 | 100.95 | 113.72 |
| 24 | D2 | 53 | LEU | N-CA-C | -8.37 | 98.44 | 110.24 |
| 43 | BR | 6 | SER | N-CA-C | 8.31 | 121.53 | 109.14 |
| 27 | D5 | 31 | VAL | CA-C-N | 8.31 | 130.23 | 119.84 |
| 27 | D5 | 31 | VAL | C-N-CA | 8.31 | 130.23 | 119.84 |
| 43 | BR | 8 | ARG | N-CA-C | 8.15 | 128.16 | 110.80 |
| 26 | B4 | 28 | LYS | CA-C-N | 8.15 | 130.03 | 119.84 |
| 26 | B4 | 28 | LYS | C-N-CA | 8.15 | 130.03 | 119.84 |
| 40 | DO | 118 | ALA | CA-C-N | 8.14 | 128.60 | 119.32 |
| 40 | DO | 118 | ALA | C-N-CA | 8.14 | 128.60 | 119.32 |
| 26 | D4 | 28 | LYS | CA-C-N | 8.10 | 129.96 | 119.84 |
| 26 | D4 | 28 | LYS | C-N-CA | 8.10 | 129.96 | 119.84 |
| 45 | DT | 76 | PHE | CA-C-N | 8.10 | 127.82 | 119.56 |
| 45 | DT | 76 | PHE | C-N-CA | 8.10 | 127.82 | 119.56 |
| 30 | D8 | 33 | ASN | N-CA-C | -8.05 | 97.19 | 113.29 |
| 35 | DF | 68 | LYS | N-CA-C | 8.00 | 124.83 | 114.75 |
| 18 | AR | 81 | PHE | N-CA-C | -7.97 | 103.35 | 113.23 |
| 33 | DD | 237 | GLU | N-CA-C | 7.96 | 122.26 | 110.48 |
| 43 | DR | 6 | SER | N-CA-C | 7.93 | 120.78 | 108.96 |
| 41 | BP | 37 | GLY | N-CA-C | 7.86 | 131.81 | 113.18 |
| 43 | DR | 8 | ARG | N-CA-C | 7.84 | 127.50 | 110.80 |
| 50 | BY | 65 | ALA | CA-C-N | 7.81 | 129.61 | 119.84 |
| 50 | BY | 65 | ALA | C-N-CA | 7.81 | 129.61 | 119.84 |

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| Mol | Chain | Res | Type | Atoms | Z | Observed($^o$) | Ideal($^o$) |
|---|---|---|---|---|---|---|---|
| 33 | BD | 243 | GLY | N-CA-C | -7.78 | 104.03 | 115.63 |
| 24 | B2 | 53 | LEU | N-CA-C | -7.78 | 99.27 | 110.24 |
| 30 | D8 | 49 | VAL | N-CA-C | -7.77 | 96.89 | 108.85 |
| 34 | DE | 132 | HIS | CA-C-N | -7.75 | 109.93 | 122.26 |
| 34 | DE | 132 | HIS | C-N-CA | -7.75 | 109.93 | 122.26 |
| 36 | DG | 125 | PHE | N-CA-C | -7.75 | 101.95 | 111.40 |
| 3 | AC | 188 | LEU | N-CA-C | 7.74 | 115.13 | 108.78 |
| 46 | BU | 97 | ASP | N-CA-C | -7.71 | 103.67 | 113.23 |
| 33 | BD | 237 | GLU | N-CA-C | 7.68 | 122.53 | 110.32 |
| 49 | DX | 5 | TYR | N-CA-C | -7.67 | 103.02 | 113.30 |
| 4 | CD | 171 | GLY | CA-C-N | 7.66 | 127.37 | 119.56 |
| 4 | CD | 171 | GLY | C-N-CA | 7.66 | 127.37 | 119.56 |
| 34 | BE | 132 | HIS | CA-C-N | -7.62 | 110.15 | 122.26 |
| 34 | BE | 132 | HIS | C-N-CA | -7.62 | 110.15 | 122.26 |
| 30 | D8 | 19 | SER | N-CA-C | -7.61 | 104.02 | 113.38 |
| 50 | BY | 46 | LYS | N-CA-C | 7.55 | 126.89 | 110.80 |
| 33 | DD | 180 | GLY | N-CA-C | -7.55 | 104.66 | 114.85 |
| 37 | BH | 67 | LEU | N-CA-C | -7.45 | 103.10 | 111.07 |
| 3 | CC | 188 | LEU | N-CA-C | 7.44 | 114.88 | 108.78 |
| 33 | DD | 238 | GLY | N-CA-C | -7.41 | 95.61 | 113.18 |
| 41 | DP | 41 | ARG | N-CA-C | -7.39 | 95.01 | 107.80 |
| 4 | AD | 171 | GLY | CA-C-N | 7.37 | 127.07 | 119.56 |
| 4 | AD | 171 | GLY | C-N-CA | 7.37 | 127.07 | 119.56 |
| 33 | BD | 43 | ARG | N-CA-C | 7.35 | 120.35 | 111.82 |
| 18 | CR | 81 | PHE | N-CA-C | -7.34 | 104.12 | 113.23 |
| 50 | DY | 63 | LYS | N-CA-C | 7.33 | 117.92 | 108.24 |
| 23 | B1 | 29 | GLY | N-CA-C | -7.28 | 105.68 | 115.36 |
| 27 | B5 | 51 | TYR | CA-CB-CG | 7.25 | 126.96 | 113.90 |
| 41 | DP | 66 | GLY | N-CA-C | -7.25 | 101.53 | 112.41 |
| 50 | BY | 6 | HIS | N-CA-C | 7.23 | 117.42 | 107.73 |
| 33 | DD | 221 | VAL | N-CA-C | 7.21 | 118.50 | 108.12 |
| 33 | BD | 221 | VAL | N-CA-C | 7.20 | 118.48 | 108.12 |
| 45 | BT | 14 | TYR | N-CA-C | -7.18 | 103.82 | 112.59 |
| 49 | DX | 62 | LYS | N-CA-C | 7.18 | 120.14 | 109.59 |
| 49 | BX | 62 | LYS | N-CA-C | 7.16 | 120.11 | 109.59 |
| 35 | BF | 64 | ILE | N-CA-C | 7.15 | 118.77 | 110.62 |
| 49 | BX | 59 | VAL | N-CA-C | 7.14 | 124.20 | 109.34 |
| 38 | DI | 79 | ILE | N-CA-C | 7.13 | 115.67 | 107.89 |
| 41 | BP | 86 | LYS | N-CA-C | -7.13 | 103.25 | 112.23 |
| 33 | BD | 238 | GLY | N-CA-C | -7.12 | 96.30 | 113.18 |
| 42 | BQ | 72 | LYS | CA-C-N | 7.12 | 126.88 | 119.76 |
| 42 | BQ | 72 | LYS | C-N-CA | 7.12 | 126.88 | 119.76 |

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| Mol | Chain | Res | Type | Atoms | Z | Observed($^o$) | Ideal($^o$) |
|---|---|---|---|---|---|---|---|
| 45 | DT | 14 | TYR | N-CA-C | -7.12 | 103.91 | 112.59 |
| 33 | BD | 13 | ARG | N-CA-C | -7.11 | 103.07 | 114.09 |
| 45 | BT | 30 | VAL | CB-CA-C | -7.10 | 102.37 | 110.96 |
| 50 | DY | 46 | LYS | N-CA-C | 7.10 | 125.92 | 110.80 |
| 31 | BA | 1379 | A | O4'-C1'-N9 | 7.09 | 119.14 | 108.50 |
| 35 | BF | 68 | LYS | N-CA-C | 7.05 | 123.63 | 114.75 |
| 42 | BQ | 38 | GLU | CA-C-N | 7.03 | 126.79 | 119.76 |
| 42 | BQ | 38 | GLU | C-N-CA | 7.03 | 126.79 | 119.76 |
| 36 | BG | 125 | PHE | N-CA-C | -7.00 | 102.86 | 111.40 |
| 24 | D2 | 48 | HIS | N-CA-C | -7.00 | 105.16 | 113.21 |
| 50 | DY | 43 | ASN | N-CA-C | 7.00 | 118.84 | 110.44 |
| 35 | DF | 64 | ILE | N-CA-C | 6.93 | 118.52 | 110.62 |
| 50 | BY | 43 | ASN | N-CA-C | 6.93 | 118.75 | 110.44 |
| 49 | DX | 59 | VAL | N-CA-C | 6.89 | 123.68 | 109.34 |
| 44 | BS | 16 | ASN | N-CA-C | -6.88 | 103.83 | 111.82 |
| 45 | BT | 128 | GLU | N-CA-C | -6.88 | 104.01 | 112.88 |
| 33 | DD | 243 | GLY | N-CA-C | -6.88 | 105.60 | 115.64 |
| 41 | BP | 60 | MET | N-CA-C | -6.88 | 103.57 | 112.23 |
| 44 | DS | 16 | ASN | N-CA-C | -6.87 | 103.85 | 111.82 |
| 31 | DA | 1379 | A | O4'-C1'-N9 | 6.86 | 118.79 | 108.50 |
| 35 | DF | 83 | PHE | N-CA-C | 6.83 | 124.68 | 113.89 |
| 31 | DA | 1379 | A | O4'-C1'-C2' | 6.83 | 114.43 | 107.60 |
| 23 | B1 | 11 | ARG | CA-C-N | -6.82 | 111.78 | 119.28 |
| 23 | B1 | 11 | ARG | C-N-CA | -6.82 | 111.78 | 119.28 |
| 45 | BT | 80 | SER | N-CA-C | 6.81 | 124.87 | 109.81 |
| 27 | D5 | 51 | TYR | CA-CB-CG | 6.79 | 126.12 | 113.90 |
| 47 | DV | 85 | LYS | N-CA-C | -6.79 | 104.15 | 113.18 |
| 30 | B8 | 29 | LYS | N-CA-C | -6.79 | 105.00 | 113.28 |
| 45 | BT | 76 | PHE | CA-C-N | 6.77 | 126.47 | 119.56 |
| 45 | BT | 76 | PHE | C-N-CA | 6.77 | 126.47 | 119.56 |
| 37 | DH | 67 | LEU | N-CA-C | -6.74 | 103.02 | 111.11 |
| 12 | AL | 118 | SER | N-CA-C | -6.72 | 102.82 | 111.02 |
| 51 | DZ | 109 | ALA | N-CA-C | -6.71 | 100.78 | 110.24 |
| 46 | DU | 69 | CYS | N-CA-C | -6.71 | 103.78 | 112.23 |
| 41 | DP | 86 | LYS | N-CA-C | -6.68 | 103.81 | 112.23 |
| 39 | DN | 131 | GLN | N-CA-C | 6.67 | 120.04 | 110.10 |
| 33 | DD | 13 | ARG | N-CA-C | -6.66 | 103.77 | 114.09 |
| 38 | BI | 79 | ILE | N-CA-C | 6.64 | 115.52 | 107.61 |
| 22 | B0 | 22 | GLY | N-CA-C | 6.64 | 118.82 | 112.04 |
| 30 | B8 | 48 | PHE | N-CA-C | -6.64 | 101.32 | 110.35 |
| 5 | CE | 48 | ALA | CA-C-N | 6.63 | 126.32 | 119.82 |
| 5 | CE | 48 | ALA | C-N-CA | 6.63 | 126.32 | 119.82 |

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| Mol | Chain | Res | Type | Atoms | Z | Observed($^o$) | Ideal($^o$) |
|---|---|---|---|---|---|---|---|
| 37 | BH | 167 | GLU | CA-C-N | 6.62 | 128.12 | 119.84 |
| 37 | BH | 167 | GLU | C-N-CA | 6.62 | 128.12 | 119.84 |
| 39 | BN | 110 | GLY | CA-C-N | 6.61 | 126.06 | 118.85 |
| 39 | BN | 110 | GLY | C-N-CA | 6.61 | 126.06 | 118.85 |
| 3 | AC | 188 | LEU | CA-C-O | 6.60 | 121.77 | 117.94 |
| 35 | BF | 83 | PHE | N-CA-C | 6.60 | 124.32 | 113.89 |
| 46 | BU | 69 | CYS | N-CA-C | -6.60 | 103.92 | 112.23 |
| 26 | B4 | 26 | SER | N-CA-C | -6.59 | 105.63 | 113.21 |
| 30 | D8 | 33 | ASN | N-CA-CB | -6.59 | 103.47 | 114.27 |
| 34 | BE | 168 | MET | N-CA-C | 6.58 | 120.25 | 109.72 |
| 47 | BV | 66 | ARG | N-CA-C | 6.54 | 116.69 | 108.45 |
| 39 | BN | 131 | GLN | N-CA-C | 6.52 | 119.82 | 110.10 |
| 23 | D1 | 29 | GLY | N-CA-C | -6.52 | 106.69 | 115.36 |
| 50 | BY | 63 | LYS | N-CA-C | 6.50 | 117.71 | 108.74 |
| 34 | DE | 159 | HIS | CB-CA-C | 6.50 | 119.51 | 110.16 |
| 31 | DA | 1544 | A | N9-C1'-C2' | 6.49 | 121.73 | 112.00 |
| 51 | BZ | 109 | ALA | N-CA-C | -6.47 | 101.12 | 110.24 |
| 31 | BA | 1379 | A | O4'-C1'-C2' | 6.46 | 114.06 | 107.60 |
| 41 | BP | 20 | GLY | N-CA-C | -6.45 | 103.60 | 112.25 |
| 46 | DU | 19 | LYS | N-CA-C | -6.45 | 104.25 | 111.28 |
| 46 | DU | 97 | ASP | N-CA-C | -6.44 | 105.24 | 113.23 |
| 31 | BA | 1544 | A | N9-C1'-C2' | 6.41 | 121.62 | 112.00 |
| 50 | DY | 6 | HIS | N-CA-C | 6.40 | 116.31 | 107.73 |
| 41 | DP | 37 | GLY | N-CA-C | 6.39 | 128.33 | 113.18 |
| 51 | BZ | 86 | VAL | CB-CA-C | -6.39 | 100.66 | 110.95 |
| 27 | B5 | 27 | PRO | O-C-N | 6.38 | 124.15 | 121.15 |
| 23 | D1 | 16 | ASN | N-CA-C | 6.37 | 119.57 | 109.50 |
| 48 | DW | 12 | ILE | N-CA-C | -6.37 | 98.33 | 108.90 |
| 49 | BX | 78 | LYS | N-CA-C | 6.36 | 119.61 | 109.24 |
| 3 | AC | 29 | TYR | N-CA-C | -6.36 | 106.11 | 114.31 |
| 47 | DV | 66 | ARG | N-CA-C | 6.35 | 116.45 | 108.45 |
| 33 | BD | 158 | ALA | N-CA-C | 6.34 | 117.86 | 111.07 |
| 30 | B8 | 33 | ASN | N-CA-CB | -6.34 | 103.87 | 114.27 |
| 23 | B1 | 16 | ASN | N-CA-C | 6.33 | 119.50 | 109.50 |
| 33 | DD | 136 | ILE | CA-C-N | 6.32 | 127.75 | 119.84 |
| 33 | DD | 136 | ILE | C-N-CA | 6.32 | 127.75 | 119.84 |
| 30 | B8 | 50 | LEU | N-CA-C | 6.30 | 119.18 | 108.90 |
| 23 | B1 | 58 | ILE | N-CA-C | 6.29 | 116.92 | 108.11 |
| 30 | D8 | 50 | LEU | N-CA-C | 6.28 | 119.13 | 108.90 |
| 23 | B1 | 15 | ALA | N-CA-C | 6.24 | 119.15 | 109.60 |
| 45 | DT | 80 | SER | N-CA-C | 6.24 | 123.61 | 109.81 |
| 44 | DS | 56 | LEU | N-CA-C | -6.24 | 106.31 | 114.04 |

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| Mol | Chain | Res | Type | Atoms | Z | Observed($^o$) | Ideal($^o$) |
|---|---|---|---|---|---|---|---|
| 28 | B6 | 48 | VAL | N-CA-C | 6.23 | 122.31 | 109.34 |
| 38 | BI | 53 | ALA | N-CA-C | 6.23 | 117.91 | 110.44 |
| 51 | BZ | 136 | PHE | N-CA-C | 6.22 | 116.06 | 107.73 |
| 3 | CC | 188 | LEU | CA-C-O | 6.21 | 121.55 | 117.94 |
| 26 | D4 | 26 | SER | N-CA-C | -6.21 | 106.07 | 113.21 |
| 37 | BH | 155 | SER | N-CA-C | 6.19 | 123.98 | 110.80 |
| 45 | DT | 128 | GLU | N-CA-C | -6.19 | 104.89 | 112.88 |
| 41 | BP | 38 | GLN | N-CA-C | 6.17 | 123.94 | 110.80 |
| 3 | CC | 29 | TYR | N-CA-C | -6.17 | 106.35 | 114.31 |
| 45 | BT | 13 | ARG | N-CA-C | -6.17 | 103.13 | 110.41 |
| 49 | DX | 39 | ILE | N-CA-C | -6.16 | 105.72 | 112.80 |
| 31 | DA | 1934 | C | C4'-C3'-C2' | 6.15 | 108.75 | 102.60 |
| 47 | BV | 85 | LYS | N-CA-C | -6.14 | 104.86 | 112.72 |
| 39 | DN | 110 | GLY | CA-C-N | 6.14 | 125.55 | 118.85 |
| 39 | DN | 110 | GLY | C-N-CA | 6.14 | 125.55 | 118.85 |
| 34 | BE | 11 | MET | N-CA-C | 6.13 | 119.19 | 109.81 |
| 2 | AB | 23 | ARG | N-CA-C | -6.09 | 104.49 | 112.41 |
| 45 | DT | 130 | ALA | N-CA-C | -6.09 | 104.44 | 112.72 |
| 13 | CM | 66 | LEU | N-CA-C | 6.08 | 113.77 | 108.78 |
| 33 | BD | 107 | ALA | CA-C-N | 6.08 | 126.05 | 120.03 |
| 33 | BD | 107 | ALA | C-N-CA | 6.08 | 126.05 | 120.03 |
| 41 | BP | 58 | THR | N-CA-C | -6.08 | 97.85 | 110.80 |
| 39 | BN | 25 | ARG | N-CA-C | -6.08 | 104.77 | 111.82 |
| 45 | BT | 57 | PHE | N-CA-C | -6.07 | 100.33 | 109.79 |
| 22 | D0 | 22 | GLY | N-CA-C | 6.07 | 118.23 | 112.04 |
| 49 | BX | 7 | VAL | N-CA-C | 6.05 | 116.23 | 110.42 |
| 23 | D1 | 11 | ARG | CA-C-N | -6.05 | 112.33 | 119.05 |
| 23 | D1 | 11 | ARG | C-N-CA | -6.05 | 112.33 | 119.05 |
| 20 | AT | 12 | ALA | N-CA-C | -6.05 | 105.90 | 113.28 |
| 38 | BI | 50 | ARG | N-CA-C | -6.05 | 104.02 | 111.40 |
| 44 | DS | 40 | ILE | N-CA-C | 6.05 | 116.64 | 108.17 |
| 31 | BA | 1934 | C | C4'-C3'-C2' | 6.04 | 108.64 | 102.60 |
| 23 | D1 | 58 | ILE | N-CA-C | 6.04 | 116.57 | 108.11 |
| 41 | DP | 54 | GLY | N-CA-C | -6.04 | 103.53 | 114.46 |
| 23 | B1 | 17 | SER | N-CA-C | 6.04 | 118.64 | 109.95 |
| 48 | DW | 69 | LEU | N-CA-C | 6.03 | 119.37 | 109.72 |
| 50 | DY | 5 | MET | N-CA-C | 6.03 | 119.27 | 110.17 |
| 47 | BV | 88 | ARG | N-CA-C | -6.02 | 99.88 | 109.76 |
| 27 | B5 | 38 | ALA | N-CA-C | 6.02 | 117.86 | 109.31 |
| 51 | BZ | 110 | GLY | N-CA-C | -6.00 | 98.96 | 113.18 |
| 38 | DI | 50 | ARG | N-CA-C | -5.99 | 104.67 | 111.14 |
| 49 | BX | 60 | ARG | N-CA-C | 5.99 | 123.55 | 110.80 |

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| Mol | Chain | Res | Type | Atoms | Z | Observed($^o$) | Ideal($^o$) |
|---|---|---|---|---|---|---|---|
| 4 | CD | 39 | PRO | CA-C-N | 5.98 | 127.31 | 119.84 |
| 4 | CD | 39 | PRO | C-N-CA | 5.98 | 127.31 | 119.84 |
| 45 | DT | 30 | VAL | CB-CA-C | -5.95 | 103.76 | 110.96 |
| 41 | DP | 19 | VAL | N-CA-C | 5.95 | 112.57 | 106.21 |
| 51 | BZ | 152 | ALA | CB-CA-C | -5.94 | 109.71 | 116.54 |
| 31 | BA | 2662 | A | N9-C1'-C2' | 5.94 | 120.90 | 112.00 |
| 45 | DT | 30 | VAL | N-CA-C | 5.92 | 117.29 | 108.23 |
| 35 | DF | 4 | VAL | N-CA-C | 5.91 | 115.12 | 106.55 |
| 40 | BO | 47 | ILE | CA-C-N | 5.90 | 127.22 | 119.84 |
| 40 | BO | 47 | ILE | C-N-CA | 5.90 | 127.22 | 119.84 |
| 2 | AB | 25 | ASN | CA-C-N | 5.90 | 127.21 | 119.84 |
| 2 | AB | 25 | ASN | C-N-CA | 5.90 | 127.21 | 119.84 |
| 51 | DZ | 152 | ALA | CB-CA-C | -5.90 | 109.76 | 116.54 |
| 45 | BT | 47 | GLY | N-CA-C | 5.89 | 121.93 | 110.73 |
| 48 | BW | 12 | ILE | N-CA-C | -5.89 | 99.11 | 108.90 |
| 34 | BE | 187 | ALA | N-CA-C | 5.88 | 119.21 | 107.62 |
| 50 | DY | 11 | ASP | N-CA-C | 5.88 | 116.98 | 108.34 |
| 4 | CD | 38 | TYR | CA-C-N | 5.87 | 126.43 | 120.38 |
| 4 | CD | 38 | TYR | C-N-CA | 5.87 | 126.43 | 120.38 |
| 30 | B8 | 49 | VAL | N-CA-C | -5.86 | 99.85 | 108.99 |
| 47 | BV | 63 | GLY | N-CA-C | -5.86 | 103.63 | 112.41 |
| 49 | DX | 60 | ARG | N-CA-C | 5.86 | 123.27 | 110.80 |
| 4 | AD | 39 | PRO | CA-C-N | 5.85 | 127.15 | 119.84 |
| 4 | AD | 39 | PRO | C-N-CA | 5.85 | 127.15 | 119.84 |
| 51 | DZ | 136 | PHE | N-CA-C | 5.84 | 115.56 | 107.73 |
| 50 | BY | 7 | VAL | N-CA-C | 5.84 | 121.48 | 109.34 |
| 41 | BP | 97 | PRO | N-CA-C | -5.84 | 102.03 | 111.14 |
| 3 | AC | 188 | LEU | CB-CA-C | -5.83 | 108.67 | 117.07 |
| 50 | BY | 8 | LYS | N-CA-C | 5.83 | 118.08 | 109.69 |
| 30 | D8 | 35 | GLN | N-CA-C | 5.80 | 123.16 | 110.80 |
| 37 | DH | 155 | SER | N-CA-C | 5.80 | 123.16 | 110.80 |
| 31 | DA | 472 | A | C4'-C3'-C2' | 5.79 | 108.39 | 102.60 |
| 24 | B2 | 48 | HIS | N-CA-C | -5.79 | 106.45 | 112.93 |
| 28 | D6 | 16 | CYS | N-CA-C | 5.78 | 118.22 | 107.99 |
| 22 | D0 | 11 | ARG | N-CA-C | -5.76 | 106.29 | 113.38 |
| 35 | BF | 4 | VAL | N-CA-C | 5.76 | 114.90 | 106.55 |
| 31 | DA | 1652 | A | C4'-C3'-C2' | 5.76 | 108.36 | 102.60 |
| 51 | DZ | 110 | GLY | N-CA-C | -5.76 | 99.53 | 113.18 |
| 33 | BD | 136 | ILE | CA-C-N | 5.75 | 127.03 | 119.84 |
| 33 | BD | 136 | ILE | C-N-CA | 5.75 | 127.03 | 119.84 |
| 40 | DO | 47 | ILE | CA-C-N | 5.75 | 127.03 | 119.84 |
| 40 | DO | 47 | ILE | C-N-CA | 5.75 | 127.03 | 119.84 |

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| Mol | Chain | Res | Type | Atoms | Z | Observed($^o$) | Ideal($^o$) |
|---|---|---|---|---|---|---|---|
| 2 | AB | 124 | SER | N-CA-C | 5.74 | 113.39 | 108.22 |
| 27 | D5 | 46 | CYS | N-CA-C | 5.73 | 122.48 | 109.81 |
| 51 | DZ | 86 | VAL | CB-CA-C | -5.72 | 101.74 | 110.95 |
| 31 | DA | 1992 | G | C2'-C3'-O3' | 5.72 | 118.08 | 109.50 |
| 35 | DF | 23 | ASP | N-CA-C | -5.72 | 105.97 | 113.17 |
| 4 | AD | 38 | TYR | CA-C-N | 5.71 | 126.26 | 120.38 |
| 4 | AD | 38 | TYR | C-N-CA | 5.71 | 126.26 | 120.38 |
| 31 | DA | 1962 | C | N1-C1'-C2' | 5.70 | 122.55 | 114.00 |
| 33 | DD | 14 | ARG | N-CA-C | -5.68 | 106.25 | 112.72 |
| 23 | D1 | 55 | GLY | N-CA-C | -5.68 | 99.72 | 113.18 |
| 50 | DY | 12 | THR | N-CA-C | 5.68 | 117.62 | 107.80 |
| 31 | BA | 472 | A | C4'-C3'-C2' | 5.67 | 108.28 | 102.60 |
| 34 | BE | 159 | HIS | CB-CA-C | 5.66 | 121.68 | 110.42 |
| 38 | DI | 79 | ILE | CA-C-N | 5.65 | 126.91 | 119.84 |
| 38 | DI | 79 | ILE | C-N-CA | 5.65 | 126.91 | 119.84 |
| 44 | BS | 56 | LEU | N-CA-C | -5.65 | 107.04 | 114.04 |
| 13 | AM | 66 | LEU | N-CA-C | 5.64 | 113.41 | 108.78 |
| 18 | AR | 86 | VAL | N-CA-C | 5.64 | 115.69 | 106.72 |
| 41 | DP | 71 | VAL | CA-C-N | -5.64 | 114.07 | 120.89 |
| 41 | DP | 71 | VAL | C-N-CA | -5.64 | 114.07 | 120.89 |
| 24 | B2 | 51 | ARG | N-CA-C | -5.63 | 98.80 | 110.80 |
| 13 | CM | 66 | LEU | CA-C-O | 5.63 | 121.21 | 117.94 |
| 22 | B0 | 84 | LEU | N-CA-C | 5.62 | 115.97 | 108.38 |
| 25 | B3 | 11 | SER | CA-C-N | 5.62 | 126.11 | 120.04 |
| 25 | B3 | 11 | SER | C-N-CA | 5.62 | 126.11 | 120.04 |
| 28 | B6 | 16 | CYS | N-CA-C | 5.61 | 117.93 | 107.99 |
| 23 | D1 | 43 | TYR | CA-C-N | 5.61 | 126.86 | 119.84 |
| 23 | D1 | 43 | TYR | C-N-CA | 5.61 | 126.86 | 119.84 |
| 31 | DA | 2662 | A | N9-C1'-C2' | 5.61 | 120.41 | 112.00 |
| 43 | DR | 6 | SER | CA-C-N | 5.61 | 126.86 | 120.53 |
| 43 | DR | 6 | SER | C-N-CA | 5.61 | 126.86 | 120.53 |
| 38 | BI | 66 | GLU | N-CA-C | -5.60 | 105.79 | 113.30 |
| 34 | DE | 187 | ALA | N-CA-C | 5.60 | 118.65 | 107.62 |
| 34 | DE | 168 | MET | N-CA-C | 5.59 | 118.67 | 109.72 |
| 41 | DP | 38 | GLN | N-CA-C | 5.57 | 122.66 | 110.80 |
| 42 | DQ | 64 | ILE | N-CA-C | 5.56 | 115.87 | 107.75 |
| 11 | CK | 57 | THR | CA-C-N | 5.56 | 125.22 | 119.05 |
| 11 | CK | 57 | THR | C-N-CA | 5.56 | 125.22 | 119.05 |
| 23 | D1 | 17 | SER | N-CA-C | 5.55 | 117.94 | 109.95 |
| 49 | DX | 61 | GLY | N-CA-C | 5.55 | 118.59 | 110.38 |
| 23 | D1 | 41 | ARG | N-CA-C | 5.55 | 118.28 | 109.96 |
| 13 | AM | 66 | LEU | CA-C-O | 5.54 | 121.16 | 117.94 |

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| Mol | Chain | Res | Type | Atoms | Z | Observed($^o$) | Ideal($^o$) |
|---|---|---|---|---|---|---|---|
| 48 | DW | 60 | ASN | N-CA-C | 5.54 | 117.76 | 111.11 |
| 39 | DN | 5 | VAL | N-CA-C | 5.54 | 120.85 | 108.88 |
| 48 | BW | 60 | ASN | N-CA-C | 5.54 | 117.75 | 111.11 |
| 43 | DR | 112 | ALA | N-CA-C | 5.54 | 117.52 | 108.55 |
| 36 | DG | 45 | GLU | N-CA-C | 5.53 | 116.87 | 108.07 |
| 44 | BS | 40 | ILE | N-CA-C | 5.53 | 115.91 | 108.17 |
| 31 | BA | 1652 | A | C4'-C3'-C2' | 5.52 | 108.12 | 102.60 |
| 23 | B1 | 70 | VAL | CB-CA-C | -5.52 | 104.82 | 111.88 |
| 35 | BF | 23 | ASP | N-CA-C | -5.52 | 106.22 | 113.17 |
| 48 | BW | 85 | VAL | N-CA-C | 5.52 | 116.20 | 109.30 |
| 31 | BA | 1992 | G | C2'-C3'-O3' | 5.51 | 117.77 | 109.50 |
| 22 | D0 | 84 | LEU | N-CA-C | 5.51 | 115.83 | 108.38 |
| 36 | BG | 45 | GLU | N-CA-C | 5.51 | 116.83 | 108.07 |
| 38 | BI | 79 | ILE | CA-C-N | 5.51 | 126.73 | 119.84 |
| 38 | BI | 79 | ILE | C-N-CA | 5.51 | 126.73 | 119.84 |
| 2 | CB | 25 | ASN | CA-C-N | 5.51 | 126.72 | 119.84 |
| 2 | CB | 25 | ASN | C-N-CA | 5.51 | 126.72 | 119.84 |
| 42 | DQ | 72 | LYS | CA-C-N | 5.50 | 125.26 | 119.76 |
| 42 | DQ | 72 | LYS | C-N-CA | 5.50 | 125.26 | 119.76 |
| 17 | AQ | 67 | LYS | N-CA-C | -5.50 | 103.27 | 110.53 |
| 8 | CH | 56 | LYS | CA-C-N | 5.50 | 125.77 | 119.83 |
| 8 | CH | 56 | LYS | C-N-CA | 5.50 | 125.77 | 119.83 |
| 5 | AE | 48 | ALA | N-CA-C | 5.50 | 113.08 | 108.13 |
| 31 | BA | 1962 | C | N1-C1'-C2' | 5.49 | 122.24 | 114.00 |
| 39 | BN | 5 | VAL | N-CA-C | 5.49 | 120.74 | 108.88 |
| 49 | DX | 78 | LYS | N-CA-C | 5.49 | 118.19 | 109.24 |
| 50 | DY | 8 | LYS | N-CA-C | 5.49 | 117.59 | 109.69 |
| 20 | CT | 12 | ALA | N-CA-C | -5.49 | 106.36 | 113.16 |
| 41 | DP | 52 | GLU | N-CA-C | 5.49 | 122.49 | 110.80 |
| 12 | AL | 88 | GLY | N-CA-C | -5.49 | 100.18 | 113.18 |
| 39 | BN | 122 | VAL | CB-CA-C | -5.48 | 103.39 | 110.84 |
| 33 | DD | 158 | ALA | N-CA-C | 5.48 | 116.94 | 111.07 |
| 16 | AP | 21 | VAL | N-CA-C | 5.48 | 116.00 | 107.28 |
| 34 | DE | 11 | MET | N-CA-C | 5.47 | 118.19 | 109.81 |
| 48 | BW | 69 | LEU | N-CA-C | 5.47 | 118.47 | 109.72 |
| 49 | DX | 75 | ASP | N-CA-C | -5.47 | 103.68 | 110.41 |
| 35 | DF | 68 | LYS | CA-C-N | -5.47 | 112.80 | 123.13 |
| 35 | DF | 68 | LYS | C-N-CA | -5.47 | 112.80 | 123.13 |
| 18 | AR | 32 | ARG | N-CA-C | -5.46 | 106.47 | 114.39 |
| 48 | BW | 6 | ILE | CB-CA-C | -5.46 | 102.33 | 111.29 |
| 3 | AC | 138 | VAL | N-CA-C | -5.46 | 107.12 | 111.81 |
| 3 | CC | 138 | VAL | N-CA-C | -5.46 | 107.12 | 111.81 |

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| Mol | Chain | Res | Type | Atoms | Z | Observed($^o$) | Ideal($^o$) |
|---|---|---|---|---|---|---|---|
| 18 | CR | 44 | LEU | N-CA-C | 5.46 | 117.86 | 110.06 |
| 49 | BX | 80 | ILE | CB-CA-C | -5.45 | 104.94 | 112.24 |
| 34 | BE | 141 | ILE | CB-CA-C | -5.44 | 104.94 | 110.93 |
| 2 | CB | 124 | SER | N-CA-C | 5.44 | 113.12 | 108.22 |
| 33 | BD | 49 | ILE | N-CA-CB | 5.43 | 116.99 | 110.31 |
| 34 | BE | 31 | CYS | CA-C-N | 5.43 | 126.62 | 119.84 |
| 34 | BE | 31 | CYS | C-N-CA | 5.43 | 126.62 | 119.84 |
| 45 | DT | 80 | SER | CA-C-N | -5.43 | 113.06 | 119.84 |
| 45 | DT | 80 | SER | C-N-CA | -5.43 | 113.06 | 119.84 |
| 8 | AH | 56 | LYS | CA-C-N | 5.42 | 125.69 | 119.83 |
| 8 | AH | 56 | LYS | C-N-CA | 5.42 | 125.69 | 119.83 |
| 45 | BT | 130 | ALA | N-CA-C | -5.42 | 104.00 | 111.81 |
| 16 | CP | 77 | ALA | N-CA-C | -5.42 | 106.58 | 114.12 |
| 2 | CB | 23 | ARG | N-CA-C | -5.42 | 105.36 | 112.41 |
| 17 | CQ | 29 | HIS | CA-C-N | 5.41 | 126.61 | 119.84 |
| 17 | CQ | 29 | HIS | C-N-CA | 5.41 | 126.61 | 119.84 |
| 49 | BX | 61 | GLY | N-CA-C | 5.41 | 118.38 | 110.38 |
| 50 | DY | 55 | TYR | CA-C-N | 5.40 | 126.59 | 119.84 |
| 50 | DY | 55 | TYR | C-N-CA | 5.40 | 126.59 | 119.84 |
| 11 | CK | 53 | SER | N-CA-C | -5.40 | 106.53 | 113.23 |
| 23 | D1 | 15 | ALA | N-CA-C | 5.40 | 117.86 | 109.60 |
| 34 | DE | 135 | HIS | N-CA-C | 5.40 | 116.97 | 111.14 |
| 36 | BG | 114 | ILE | CB-CA-C | -5.40 | 106.53 | 112.68 |
| 16 | CP | 21 | VAL | N-CA-C | 5.38 | 115.84 | 107.28 |
| 33 | DD | 107 | ALA | CA-C-N | 5.38 | 125.36 | 120.03 |
| 33 | DD | 107 | ALA | C-N-CA | 5.38 | 125.36 | 120.03 |
| 44 | DS | 68 | GLN | N-CA-C | -5.38 | 105.52 | 111.71 |
| 18 | CR | 32 | ARG | N-CA-C | -5.38 | 106.59 | 114.39 |
| 45 | BT | 30 | VAL | N-CA-C | 5.38 | 116.46 | 108.23 |
| 41 | BP | 126 | VAL | CB-CA-C | -5.37 | 105.68 | 111.59 |
| 50 | BY | 12 | THR | N-CA-C | 5.37 | 117.09 | 107.80 |
| 45 | DT | 47 | GLY | N-CA-C | 5.37 | 120.93 | 110.73 |
| 39 | DN | 94 | HIS | CA-C-N | 5.36 | 126.21 | 120.47 |
| 39 | DN | 94 | HIS | C-N-CA | 5.36 | 126.21 | 120.47 |
| 18 | AR | 44 | LEU | N-CA-C | 5.35 | 117.71 | 110.06 |
| 30 | B8 | 37 | SER | N-CA-C | 5.34 | 122.18 | 110.80 |
| 3 | CC | 188 | LEU | CB-CA-C | -5.34 | 109.39 | 117.07 |
| 37 | DH | 167 | GLU | CA-C-N | 5.33 | 126.50 | 119.84 |
| 37 | DH | 167 | GLU | C-N-CA | 5.33 | 126.50 | 119.84 |
| 11 | AK | 40 | ILE | N-CA-C | -5.33 | 107.23 | 111.81 |
| 39 | BN | 41 | ASP | N-CA-C | -5.33 | 106.62 | 113.23 |
| 31 | BA | 2796 | U | N1-C1'-C2' | 5.32 | 119.98 | 112.00 |

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| Mol | Chain | Res | Type | Atoms | Z | Observed($^o$) | Ideal($^o$) |
|---|---|---|---|---|---|---|---|
| 33 | BD | 14 | ARG | N-CA-C | -5.32 | 106.66 | 112.72 |
| 37 | DH | 158 | HIS | N-CA-C | 5.32 | 122.12 | 110.80 |
| 45 | BT | 80 | SER | CA-C-N | -5.31 | 113.20 | 119.84 |
| 45 | BT | 80 | SER | C-N-CA | -5.31 | 113.20 | 119.84 |
| 49 | BX | 64 | LYS | N-CA-C | 5.31 | 118.09 | 109.06 |
| 39 | DN | 5 | VAL | CA-C-N | 5.31 | 127.42 | 120.25 |
| 39 | DN | 5 | VAL | C-N-CA | 5.31 | 127.42 | 120.25 |
| 41 | DP | 97 | PRO | N-CA-C | -5.31 | 102.86 | 111.14 |
| 30 | B8 | 35 | GLN | N-CA-C | 5.30 | 122.09 | 110.80 |
| 13 | AM | 66 | LEU | CB-CA-C | -5.29 | 109.45 | 117.07 |
| 35 | BF | 100 | THR | N-CA-C | 5.28 | 117.31 | 108.23 |
| 13 | CM | 66 | LEU | CB-CA-C | -5.28 | 109.47 | 117.07 |
| 31 | BA | 1609 | A | C3'-C2'-C1' | 5.28 | 106.78 | 101.50 |
| 51 | BZ | 176 | PRO | CA-C-N | 5.27 | 126.43 | 119.84 |
| 51 | BZ | 176 | PRO | C-N-CA | 5.27 | 126.43 | 119.84 |
| 16 | AP | 77 | ALA | N-CA-C | -5.27 | 106.79 | 114.12 |
| 36 | BG | 114 | ILE | N-CA-C | 5.27 | 114.20 | 106.55 |
| 38 | DI | 66 | GLU | N-CA-C | -5.27 | 106.23 | 113.30 |
| 51 | BZ | 61 | LEU | CA-C-N | 5.27 | 126.43 | 119.84 |
| 51 | BZ | 61 | LEU | C-N-CA | 5.27 | 126.43 | 119.84 |
| 36 | DG | 114 | ILE | N-CA-C | 5.27 | 114.19 | 106.55 |
| 33 | BD | 229 | VAL | CB-CA-C | -5.26 | 104.37 | 112.05 |
| 18 | CR | 86 | VAL | N-CA-C | 5.25 | 115.07 | 106.72 |
| 17 | AQ | 29 | HIS | CA-C-N | 5.24 | 126.39 | 119.84 |
| 17 | AQ | 29 | HIS | C-N-CA | 5.24 | 126.39 | 119.84 |
| 16 | CP | 5 | ARG | N-CA-C | 5.24 | 116.77 | 108.96 |
| 30 | B8 | 49 | VAL | CA-C-N | -5.23 | 115.61 | 122.99 |
| 30 | B8 | 49 | VAL | C-N-CA | -5.23 | 115.61 | 122.99 |
| 25 | B3 | 50 | VAL | CB-CA-C | -5.23 | 106.17 | 112.19 |
| 27 | D5 | 38 | ALA | N-CA-C | 5.23 | 116.74 | 109.31 |
| 35 | DF | 177 | ALA | CA-C-N | 5.23 | 125.28 | 119.32 |
| 35 | DF | 177 | ALA | C-N-CA | 5.23 | 125.28 | 119.32 |
| 47 | BV | 49 | THR | N-CA-C | 5.23 | 118.55 | 108.45 |
| 23 | B1 | 43 | TYR | CA-C-N | 5.23 | 126.38 | 119.84 |
| 23 | B1 | 43 | TYR | C-N-CA | 5.23 | 126.38 | 119.84 |
| 46 | BU | 19 | LYS | N-CA-C | -5.23 | 105.58 | 111.28 |
| 36 | DG | 114 | ILE | CB-CA-C | -5.22 | 106.72 | 112.68 |
| 11 | AK | 57 | THR | CA-C-N | 5.22 | 124.85 | 119.05 |
| 11 | AK | 57 | THR | C-N-CA | 5.22 | 124.85 | 119.05 |
| 47 | BV | 43 | GLU | N-CA-C | 5.22 | 121.92 | 110.80 |
| 31 | DA | 1609 | A | C3'-C2'-C1' | 5.22 | 106.72 | 101.50 |
| 41 | DP | 53 | GLY | N-CA-C | -5.22 | 100.81 | 113.18 |

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| Mol | Chain | Res | Type | Atoms | Z | Observed($^o$) | Ideal($^o$) |
|---|---|---|---|---|---|---|---|
| 41 | DP | 58 | THR | N-CA-C | -5.22 | 99.69 | 110.80 |
| 31 | DA | 2796 | U | N1-C1'-C2' | 5.21 | 119.82 | 112.00 |
| 49 | DX | 64 | LYS | N-CA-C | 5.21 | 117.92 | 109.06 |
| 50 | BY | 5 | MET | N-CA-C | 5.21 | 118.04 | 110.17 |
| 12 | AL | 105 | TYR | CB-CA-C | -5.20 | 109.03 | 115.89 |
| 11 | CK | 40 | ILE | N-CA-C | -5.20 | 107.34 | 111.81 |
| 47 | BV | 28 | GLU | CA-C-N | -5.19 | 114.64 | 120.14 |
| 47 | BV | 28 | GLU | C-N-CA | -5.19 | 114.64 | 120.14 |
| 42 | DQ | 38 | GLU | CA-C-N | 5.17 | 124.93 | 119.76 |
| 42 | DQ | 38 | GLU | C-N-CA | 5.17 | 124.93 | 119.76 |
| 43 | DR | 10 | LEU | CA-C-N | -5.17 | 113.02 | 122.12 |
| 43 | DR | 10 | LEU | C-N-CA | -5.17 | 113.02 | 122.12 |
| 49 | DX | 43 | VAL | N-CA-C | -5.16 | 105.36 | 110.62 |
| 47 | BV | 9 | GLY | N-CA-C | -5.15 | 106.00 | 114.76 |
| 41 | DP | 116 | GLY | N-CA-C | 5.15 | 125.38 | 113.18 |
| 2 | CB | 71 | VAL | N-CA-C | 5.14 | 116.06 | 108.45 |
| 39 | BN | 18 | ALA | N-CA-C | -5.14 | 106.21 | 113.20 |
| 45 | DT | 13 | ARG | N-CA-C | -5.13 | 104.36 | 110.41 |
| 24 | B2 | 44 | LEU | N-CA-C | 5.12 | 120.00 | 113.18 |
| 46 | DU | 8 | VAL | N-CA-C | 5.12 | 115.40 | 110.74 |
| 41 | BP | 71 | VAL | N-CA-C | 5.11 | 119.92 | 108.88 |
| 30 | D8 | 49 | VAL | CA-C-N | -5.11 | 115.78 | 122.99 |
| 30 | D8 | 49 | VAL | C-N-CA | -5.11 | 115.78 | 122.99 |
| 43 | DR | 3 | HIS | N-CA-C | 5.11 | 117.95 | 107.37 |
| 41 | BP | 116 | GLY | N-CA-C | 5.11 | 125.28 | 113.18 |
| 49 | BX | 17 | ALA | N-CA-C | -5.10 | 105.64 | 111.14 |
| 48 | BW | 111 | HIS | N-CA-C | 5.09 | 115.01 | 108.34 |
| 37 | BH | 43 | VAL | N-CA-C | 5.09 | 115.10 | 107.51 |
| 45 | DT | 48 | ILE | CB-CA-C | -5.09 | 104.24 | 111.21 |
| 39 | BN | 94 | HIS | CA-C-N | 5.09 | 125.92 | 120.47 |
| 39 | BN | 94 | HIS | C-N-CA | 5.09 | 125.92 | 120.47 |
| 27 | B5 | 46 | CYS | N-CA-C | 5.09 | 122.43 | 107.37 |
| 39 | BN | 5 | VAL | CA-C-N | 5.08 | 127.11 | 120.25 |
| 39 | BN | 5 | VAL | C-N-CA | 5.08 | 127.11 | 120.25 |
| 13 | AM | 92 | HIS | N-CA-C | -5.08 | 106.59 | 112.89 |
| 27 | B5 | 6 | VAL | CA-C-N | 5.08 | 125.08 | 119.90 |
| 27 | B5 | 6 | VAL | C-N-CA | 5.08 | 125.08 | 119.90 |
| 43 | BR | 57 | ARG | N-CA-C | -5.08 | 103.44 | 110.35 |
| 45 | DT | 108 | ARG | N-CA-C | -5.08 | 107.21 | 113.41 |
| 41 | BP | 29 | LYS | CD-CE-NZ | 5.07 | 128.13 | 111.90 |
| 42 | BQ | 35 | VAL | N-CA-C | 5.07 | 116.27 | 108.71 |
| 18 | CR | 25 | THR | N-CA-C | -5.07 | 107.14 | 113.38 |

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| Mol | Chain | Res | Type | Atoms | Z | Observed($^o$) | Ideal($^o$) |
|---|---|---|---|---|---|---|---|
| 50 | DY | 13 | VAL | N-CA-C | 5.07 | 116.66 | 108.85 |
| 39 | BN | 47 | ALA | N-CA-C | 5.07 | 118.12 | 111.28 |
| 48 | DW | 111 | HIS | N-CA-C | 5.07 | 114.98 | 108.34 |
| 47 | DV | 43 | GLU | N-CA-C | 5.06 | 121.58 | 110.80 |
| 37 | BH | 45 | VAL | CB-CA-C | -5.06 | 104.23 | 111.31 |
| 33 | BD | 224 | ALA | N-CA-C | -5.05 | 102.43 | 109.96 |
| 35 | DF | 26 | ALA | CB-CA-C | -5.05 | 109.16 | 116.54 |
| 17 | CQ | 63 | ARG | CA-C-N | 5.05 | 126.16 | 119.84 |
| 17 | CQ | 63 | ARG | C-N-CA | 5.05 | 126.16 | 119.84 |
| 45 | DT | 79 | HIS | N-CA-C | 5.05 | 117.56 | 108.48 |
| 49 | BX | 7 | VAL | CB-CA-C | -5.04 | 105.51 | 111.97 |
| 37 | DH | 19 | VAL | N-CA-C | 5.04 | 115.23 | 108.17 |
| 35 | BF | 67 | GLN | CB-CA-C | -5.04 | 102.80 | 110.06 |
| 46 | DU | 27 | LEU | N-CA-C | -5.03 | 106.97 | 113.01 |
| 50 | BY | 55 | TYR | CA-C-N | 5.03 | 126.13 | 119.84 |
| 50 | BY | 55 | TYR | C-N-CA | 5.03 | 126.13 | 119.84 |
| 23 | B1 | 78 | LYS | N-CA-C | -5.03 | 107.17 | 113.15 |
| 25 | D3 | 11 | SER | CA-C-N | 5.03 | 125.47 | 120.04 |
| 25 | D3 | 11 | SER | C-N-CA | 5.03 | 125.47 | 120.04 |
| 31 | BA | 1204 | A | C3'-C2'-C1' | -5.02 | 96.48 | 101.50 |
| 51 | DZ | 61 | LEU | CA-C-N | 5.02 | 126.11 | 119.84 |
| 51 | DZ | 61 | LEU | C-N-CA | 5.02 | 126.11 | 119.84 |
| 24 | B2 | 55 | ARG | N-CA-C | -5.01 | 105.56 | 113.02 |
| 37 | BH | 158 | HIS | N-CA-C | 5.01 | 121.46 | 110.80 |
| 4 | CD | 12 | CYS | N-CA-C | -5.00 | 100.15 | 110.80 |

All (42) chirality outliers are listed below:

| Mol | Chain | Res | Type | Atom |
|---|---|---|---|---|
| 31 | BA | 100 | G | C1' |
| 31 | BA | 472 | A | C3' |
| 31 | BA | 669 | G | C1',C3',C4' |
| 31 | BA | 945 | A | C1' |
| 31 | BA | 1300 | U | C1',C3',C4' |
| 31 | BA | 1379 | A | C1' |
| 31 | BA | 1544 | A | C1' |
| 31 | BA | 1609 | A | C2' |
| 31 | BA | 1652 | A | C3' |
| 31 | BA | 1694 | C | C3',C4' |
| 31 | BA | 1934 | C | C3' |
| 31 | BA | 1962 | C | C1',C4' |
| 31 | BA | 2286 | A | C1' |

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| Mol | Chain | Res | Type | Atom |
|---|---|---|---|---|
| 31 | BA | 2662 | A | C1’ |
| 31 | BA | 2796 | U | C1’ |
| 31 | DA | 100 | G | C1’ |
| 31 | DA | 472 | A | C3’ |
| 31 | DA | 669 | G | C1’,C3’,C4’ |
| 31 | DA | 945 | A | C1’ |
| 31 | DA | 1300 | U | C1’,C3’,C4’ |
| 31 | DA | 1379 | A | C1’ |
| 31 | DA | 1544 | A | C1’ |
| 31 | DA | 1609 | A | C2’ |
| 31 | DA | 1652 | A | C3’ |
| 31 | DA | 1694 | C | C3’,C4’ |
| 31 | DA | 1934 | C | C3’ |
| 31 | DA | 1962 | C | C1’,C4’ |
| 31 | DA | 2286 | A | C1’ |
| 31 | DA | 2662 | A | C1’ |
| 31 | DA | 2796 | U | C1’ |

All (47) planarity outliers are listed below:

| Mol | Chain | Res | Type | Group |
|---|---|---|---|---|
| 23 | B1 | 30 | VAL | Peptide |
| 24 | B2 | 55 | ARG | Peptide |
| 24 | B2 | 56 | GLN | Peptide |
| 24 | B2 | 57 | ILE | Peptide |
| 27 | B5 | 51 | TYR | Peptide |
| 33 | BD | 237 | GLU | Peptide |
| 33 | BD | 244 | ARG | Peptide |
| 34 | BE | 131 | ALA | Peptide |
| 34 | BE | 76 | ARG | Peptide |
| 35 | BF | 85 | GLY | Peptide |
| 37 | BH | 154 | PRO | Peptide |
| 37 | BH | 156 | ALA | Peptide |
| 41 | BP | 37 | GLY | Peptide |
| 41 | BP | 39 | LYS | Peptide |
| 41 | BP | 41 | ARG | Peptide |
| 41 | BP | 51 | PHE | Peptide |
| 41 | BP | 57 | THR | Peptide |
| 42 | BQ | 10 | ARG | Peptide |
| 43 | BR | 7 | GLY | Peptide |
| 44 | BS | 88 | ASP | Peptide |
| 45 | BT | 29 | ARG | Peptide |

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| Mol | Chain | Res | Type | Group |
|---|---|---|---|---|
| 47 | BV | 81 | TYR | Peptide |
| 49 | BX | 38 | GLU | Peptide |
| 49 | BX | 76 | ARG | Peptide |
| 49 | BX | 77 | LYS | Peptide |
| 23 | D1 | 30 | VAL | Peptide |
| 24 | D2 | 56 | GLN | Peptide |
| 27 | D5 | 51 | TYR | Peptide |
| 33 | DD | 237 | GLU | Peptide |
| 33 | DD | 244 | ARG | Peptide |
| 34 | DE | 131 | ALA | Peptide |
| 34 | DE | 76 | ARG | Peptide |
| 37 | DH | 154 | PRO | Peptide |
| 37 | DH | 156 | ALA | Peptide |
| 41 | DP | 37 | GLY | Peptide |
| 41 | DP | 41 | ARG | Peptide |
| 41 | DP | 51 | PHE | Peptide |
| 41 | DP | 57 | THR | Peptide |
| 42 | DQ | 10 | ARG | Peptide |
| 43 | DR | 7 | GLY | Peptide |
| 44 | DS | 88 | ASP | Peptide |
| 45 | DT | 29 | ARG | Peptide |
| 47 | DV | 81 | TYR | Peptide |
| 47 | DV | 87 | HIS | Peptide |
| 49 | DX | 38 | GLU | Peptide |
| 49 | DX | 76 | ARG | Peptide |
| 49 | DX | 77 | LYS | Peptide |

## 5.2 Too-close contacts (i)

In the following table, the Non-H and H(model) columns list the number of non-hydrogen atoms and hydrogen atoms in the chain respectively. The H(added) column lists the number of hydrogen atoms added and optimized by MolProbity. The Clashes column lists the number of clashes within the asymmetric unit, whereas Symm-Clashes lists symmetry-related clashes.

| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|---|---|---|---|---|---|---|
| 1 | AA | 32329 | 0 | 16318 | 1411 | 0 |
| 1 | CA | 32329 | 0 | 16318 | 1383 | 0 |
| 2 | AB | 1901 | 0 | 1951 | 177 | 0 |
| 2 | CB | 1901 | 0 | 1951 | 175 | 0 |
| 3 | AC | 1613 | 0 | 1677 | 119 | 0 |
| 3 | CC | 1613 | 0 | 1677 | 120 | 0 |
| 4 | AD | 1703 | 0 | 1763 | 175 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|---|---|---|---|---|---|---|
| 4 | CD | 1703 | 0 | 1763 | 175 | 0 |
| 5 | AE | 1147 | 0 | 1207 | 113 | 0 |
| 5 | CE | 1147 | 0 | 1207 | 119 | 0 |
| 6 | AF | 843 | 0 | 857 | 83 | 0 |
| 6 | CF | 843 | 0 | 857 | 90 | 0 |
| 7 | AG | 1257 | 0 | 1296 | 60 | 0 |
| 7 | CG | 1257 | 0 | 1296 | 63 | 0 |
| 8 | AH | 1116 | 0 | 1177 | 89 | 0 |
| 8 | CH | 1116 | 0 | 1177 | 89 | 0 |
| 9 | AI | 1011 | 0 | 1042 | 87 | 0 |
| 9 | CI | 1011 | 0 | 1042 | 88 | 0 |
| 10 | AJ | 795 | 0 | 840 | 85 | 0 |
| 10 | CJ | 795 | 0 | 840 | 87 | 0 |
| 11 | AK | 885 | 0 | 904 | 67 | 0 |
| 11 | CK | 885 | 0 | 904 | 73 | 0 |
| 12 | AL | 971 | 0 | 1057 | 111 | 0 |
| 12 | CL | 971 | 0 | 1057 | 110 | 0 |
| 13 | AM | 921 | 0 | 976 | 64 | 0 |
| 13 | CM | 921 | 0 | 976 | 68 | 0 |
| 14 | AN | 492 | 0 | 530 | 35 | 0 |
| 14 | CN | 492 | 0 | 529 | 35 | 0 |
| 15 | AO | 734 | 0 | 771 | 61 | 0 |
| 15 | CO | 734 | 0 | 771 | 60 | 0 |
| 16 | AP | 701 | 0 | 720 | 95 | 0 |
| 16 | CP | 701 | 0 | 720 | 97 | 0 |
| 17 | AQ | 824 | 0 | 891 | 48 | 0 |
| 17 | CQ | 824 | 0 | 891 | 50 | 0 |
| 18 | AR | 574 | 0 | 644 | 63 | 0 |
| 18 | CR | 574 | 0 | 644 | 64 | 0 |
| 19 | AS | 630 | 0 | 652 | 42 | 0 |
| 19 | CS | 630 | 0 | 652 | 36 | 0 |
| 20 | AT | 763 | 0 | 861 | 81 | 0 |
| 20 | CT | 763 | 0 | 861 | 84 | 0 |
| 21 | AU | 209 | 0 | 221 | 12 | 0 |
| 21 | CU | 209 | 0 | 221 | 12 | 0 |
| 22 | B0 | 650 | 0 | 654 | 71 | 0 |
| 22 | D0 | 650 | 0 | 654 | 67 | 0 |
| 23 | B1 | 693 | 0 | 764 | 150 | 0 |
| 23 | D1 | 693 | 0 | 764 | 147 | 0 |
| 24 | B2 | 421 | 0 | 461 | 124 | 1 |
| 24 | D2 | 421 | 0 | 461 | 128 | 0 |
| 25 | B3 | 468 | 0 | 523 | 38 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|---|---|---|---|---|---|---|
| 25 | D3 | 468 | 0 | 523 | 54 | 0 |
| 26 | B4 | 157 | 0 | 69 | 15 | 0 |
| 26 | D4 | 157 | 0 | 69 | 15 | 0 |
| 27 | B5 | 459 | 0 | 478 | 88 | 0 |
| 27 | D5 | 459 | 0 | 480 | 88 | 0 |
| 28 | B6 | 381 | 0 | 390 | 97 | 0 |
| 28 | D6 | 381 | 0 | 390 | 93 | 0 |
| 29 | B7 | 419 | 0 | 467 | 40 | 0 |
| 29 | D7 | 419 | 0 | 467 | 41 | 0 |
| 30 | B8 | 508 | 0 | 576 | 158 | 0 |
| 30 | D8 | 508 | 0 | 576 | 147 | 0 |
| 31 | BA | 58698 | 0 | 29590 | 2399 | 0 |
| 31 | DA | 58698 | 0 | 29591 | 2594 | 1 |
| 32 | BB | 2551 | 0 | 1295 | 154 | 0 |
| 32 | DB | 2551 | 0 | 1295 | 171 | 0 |
| 33 | BD | 2105 | 0 | 2182 | 348 | 0 |
| 33 | DD | 2105 | 0 | 2182 | 343 | 0 |
| 34 | BE | 1564 | 0 | 1629 | 219 | 0 |
| 34 | DE | 1564 | 0 | 1629 | 221 | 0 |
| 35 | BF | 1624 | 0 | 1677 | 183 | 0 |
| 35 | DF | 1624 | 0 | 1677 | 186 | 0 |
| 36 | BG | 1474 | 0 | 1534 | 158 | 0 |
| 36 | DG | 1474 | 0 | 1534 | 158 | 0 |
| 37 | BH | 1223 | 0 | 1282 | 144 | 0 |
| 37 | DH | 1223 | 0 | 1282 | 133 | 0 |
| 38 | BI | 1132 | 0 | 1218 | 145 | 0 |
| 38 | DI | 1132 | 0 | 1218 | 163 | 0 |
| 39 | BN | 1105 | 0 | 1180 | 187 | 0 |
| 39 | DN | 1105 | 0 | 1180 | 189 | 0 |
| 40 | BO | 933 | 0 | 996 | 89 | 0 |
| 40 | DO | 933 | 0 | 996 | 81 | 0 |
| 41 | BP | 1114 | 0 | 1187 | 277 | 0 |
| 41 | DP | 1114 | 0 | 1187 | 266 | 0 |
| 42 | BQ | 1080 | 0 | 1127 | 167 | 0 |
| 42 | DQ | 1080 | 0 | 1127 | 167 | 0 |
| 43 | BR | 960 | 0 | 1021 | 123 | 0 |
| 43 | DR | 960 | 0 | 1021 | 123 | 0 |
| 44 | BS | 771 | 0 | 832 | 152 | 0 |
| 44 | DS | 771 | 0 | 832 | 153 | 0 |
| 45 | BT | 1100 | 0 | 1164 | 184 | 0 |
| 45 | DT | 1100 | 0 | 1164 | 174 | 0 |
| 46 | BU | 958 | 0 | 1015 | 150 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|---|---|---|---|---|---|---|
| 46 | DU | 958 | 0 | 1015 | 161 | 0 |
| 47 | BV | 779 | 0 | 851 | 221 | 0 |
| 47 | DV | 779 | 0 | 851 | 226 | 0 |
| 48 | BW | 896 | 0 | 953 | 78 | 0 |
| 48 | DW | 896 | 0 | 953 | 86 | 0 |
| 49 | BX | 726 | 0 | 778 | 172 | 0 |
| 49 | DX | 726 | 0 | 778 | 174 | 0 |
| 50 | BY | 776 | 0 | 870 | 184 | 0 |
| 50 | DY | 776 | 0 | 870 | 191 | 0 |
| 51 | BZ | 1404 | 0 | 1432 | 148 | 0 |
| 51 | DZ | 1404 | 0 | 1432 | 149 | 0 |
| 52 | AA | 51 | 0 | 0 | 0 | 0 |
| 52 | B0 | 1 | 0 | 0 | 0 | 0 |
| 52 | B1 | 1 | 0 | 0 | 0 | 0 |
| 52 | B5 | 2 | 0 | 0 | 0 | 0 |
| 52 | B7 | 1 | 0 | 0 | 0 | 0 |
| 52 | BA | 349 | 0 | 0 | 0 | 0 |
| 52 | BB | 5 | 0 | 0 | 0 | 0 |
| 52 | BD | 1 | 0 | 0 | 0 | 0 |
| 52 | BE | 1 | 0 | 0 | 0 | 0 |
| 52 | BF | 1 | 0 | 0 | 0 | 0 |
| 52 | BP | 3 | 0 | 0 | 0 | 0 |
| 52 | BQ | 2 | 0 | 0 | 0 | 0 |
| 52 | BR | 1 | 0 | 0 | 0 | 0 |
| 52 | BU | 1 | 0 | 0 | 0 | 0 |
| 52 | BX | 1 | 0 | 0 | 0 | 0 |
| 52 | CA | 48 | 0 | 0 | 0 | 0 |
| 52 | D0 | 1 | 0 | 0 | 0 | 0 |
| 52 | D1 | 1 | 0 | 0 | 0 | 0 |
| 52 | D5 | 2 | 0 | 0 | 0 | 0 |
| 52 | D7 | 1 | 0 | 0 | 0 | 0 |
| 52 | DA | 309 | 0 | 0 | 0 | 0 |
| 52 | DB | 3 | 0 | 0 | 0 | 0 |
| 52 | DD | 1 | 0 | 0 | 0 | 0 |
| 52 | DE | 1 | 0 | 0 | 0 | 0 |
| 52 | DF | 1 | 0 | 0 | 0 | 0 |
| 52 | DP | 1 | 0 | 0 | 0 | 0 |
| 52 | DQ | 1 | 0 | 0 | 0 | 0 |
| 52 | DR | 1 | 0 | 0 | 0 | 0 |
| 52 | DU | 1 | 0 | 0 | 0 | 0 |
| 52 | DX | 1 | 0 | 0 | 0 | 0 |
| 53 | AD | 1 | 0 | 0 | 0 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|---|---|---|---|---|---|---|
| 53 | AN | 1 | 0 | 0 | 0 | 0 |
| 53 | CD | 1 | 0 | 0 | 0 | 0 |
| 53 | CN | 1 | 0 | 0 | 0 | 0 |
| 54 | BA | 1 | 0 | 0 | 0 | 0 |
| 54 | DA | 1 | 0 | 0 | 0 | 0 |
| 55 | BA | 52 | 0 | 72 | 3 | 0 |
| 55 | DA | 52 | 0 | 72 | 3 | 0 |
| All | All | 278000 | 0 | 189246 | 17928 | 1 |

The all-atom clashscore is defined as the number of clashes found per 1000 atoms (including hydrogen atoms). The all-atom clashscore for this structure is 39.

All (17928) close contacts within the same asymmetric unit are listed below, sorted by their clash magnitude.

| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 42:BQ:81:VAL:O | 42:BQ:82:ARG:HG2 | 1.13 | 1.31 |
| 42:DQ:81:VAL:O | 42:DQ:82:ARG:HG2 | 1.25 | 1.27 |
| 41:BP:59:LEU:HA | 41:BP:61:ARG:NH1 | 1.49 | 1.25 |
| 41:DP:59:LEU:HA | 41:DP:61:ARG:NH1 | 1.55 | 1.20 |
| 31:DA:2206:G:N2 | 31:DA:2207:G:H5' | 1.58 | 1.19 |
| 31:BA:2206:G:N2 | 31:BA:2207:G:H5' | 1.58 | 1.18 |
| 33:BD:35:LYS:HD2 | 33:BD:104:TYR:CD1 | 1.80 | 1.17 |
| 33:DD:35:LYS:NZ | 33:DD:104:TYR:HB2 | 1.58 | 1.17 |
| 2:AB:111:ARG:HH11 | 2:AB:111:ARG:HG2 | 1.08 | 1.17 |
| 47:BV:72:VAL:HA | 47:BV:88:ARG:HH22 | 1.07 | 1.17 |
| 24:D2:49:LYS:HD2 | 24:D2:53:LEU:HD12 | 1.24 | 1.16 |
| 31:DA:2334:G:H21 | 44:DS:18:ILE:HD11 | 1.08 | 1.16 |
| 31:DA:2701:C:H3' | 31:DA:2702:U:C5' | 1.75 | 1.16 |
| 1:AA:59:A:H5" | 1:AA:60:A:H5" | 1.20 | 1.16 |
| 31:DA:2394:C:OP1 | 41:DP:63:PRO:HD2 | 1.43 | 1.16 |
| 31:BA:870:A:H5" | 42:BQ:7:MET:HB2 | 1.24 | 1.16 |
| 39:BN:47:ALA:HB2 | 39:BN:112:LEU:HD11 | 1.29 | 1.15 |
| 24:B2:49:LYS:HD2 | 24:B2:53:LEU:HD12 | 1.17 | 1.15 |
| 47:DV:19:LYS:HG3 | 47:DV:20:LEU:H | 1.08 | 1.14 |
| 33:DD:35:LYS:HD2 | 33:DD:104:TYR:CD1 | 1.82 | 1.14 |
| 51:DZ:151:HIS:HB3 | 51:DZ:170:THR:HA | 1.15 | 1.14 |
| 2:CB:111:ARG:HH11 | 2:CB:111:ARG:HG2 | 1.09 | 1.14 |
| 46:DU:64:ARG:CZ | 46:DU:64:ARG:HA | 1.78 | 1.14 |
| 44:BS:85:VAL:HG23 | 44:BS:106:ARG:HB2 | 1.30 | 1.13 |
| 47:BV:19:LYS:HG3 | 47:BV:20:LEU:H | 1.05 | 1.13 |
| 50:BY:95:LYS:HD3 | 50:BY:100:ALA:HB1 | 1.17 | 1.13 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:870:A:H5" | 42:DQ:7:MET:HB2 | 1.18 | 1.13 |
| 31:BA:2701:C:H3' | 31:BA:2702:U:C5' | 1.76 | 1.13 |
| 33:BD:35:LYS:NZ | 33:BD:104:TYR:HB2 | 1.62 | 1.13 |
| 31:DA:1821:A:H2' | 31:DA:1822:G:H5" | 1.26 | 1.12 |
| 46:BU:64:ARG:CZ | 46:BU:64:ARG:HA | 1.79 | 1.12 |
| 50:DY:28:LYS:HE2 | 50:DY:30:VAL:HG22 | 1.29 | 1.12 |
| 47:BV:43:GLU:H | 47:BV:48:GLY:HA2 | 1.04 | 1.12 |
| 31:BA:2334:G:H21 | 44:BS:18:ILE:HD11 | 1.06 | 1.12 |
| 47:DV:43:GLU:H | 47:DV:48:GLY:HA2 | 1.07 | 1.12 |
| 28:D6:10:LEU:HD12 | 30:D8:35:GLN:HE22 | 1.14 | 1.11 |
| 31:BA:2394:C:OP1 | 41:BP:63:PRO:HD2 | 1.48 | 1.11 |
| 47:BV:13:ARG:HG3 | 47:BV:13:ARG:HH11 | 1.14 | 1.11 |
| 7:CG:113:GLU:HB2 | 7:CG:119:ARG:HG2 | 1.33 | 1.11 |
| 33:DD:255:LYS:H | 33:DD:255:LYS:HE3 | 1.14 | 1.10 |
| 47:DV:72:VAL:HA | 47:DV:88:ARG:HH22 | 1.12 | 1.10 |
| 33:BD:255:LYS:H | 33:BD:255:LYS:HE3 | 1.13 | 1.10 |
| 41:BP:140:ALA:HB1 | 25:D3:38:GLU:HG2 | 1.19 | 1.10 |
| 33:DD:35:LYS:HD3 | 33:DD:63:ARG:HB3 | 1.30 | 1.10 |
| 44:DS:89:ARG:O | 44:DS:92:TYR:HB3 | 1.50 | 1.10 |
| 50:BY:28:LYS:HE2 | 50:BY:30:VAL:HG22 | 1.31 | 1.10 |
| 47:DV:13:ARG:HH11 | 47:DV:13:ARG:HG3 | 1.13 | 1.09 |
| 30:B8:32:LEU:HG | 30:B8:34:TRP:HE3 | 1.14 | 1.09 |
| 51:BZ:151:HIS:HB3 | 51:BZ:170:THR:HA | 1.15 | 1.09 |
| 44:DS:28:VAL:HB | 44:DS:89:ARG:HB2 | 1.15 | 1.09 |
| 50:DY:95:LYS:HD3 | 50:DY:100:ALA:HB1 | 1.30 | 1.09 |
| 35:DF:101:LEU:HD12 | 35:DF:102:PRO:HD2 | 1.21 | 1.09 |
| 7:AG:113:GLU:HB2 | 7:AG:119:ARG:HG2 | 1.32 | 1.09 |
| 16:AP:28:ARG:HH11 | 16:AP:28:ARG:HG2 | 1.12 | 1.09 |
| 45:BT:50:ILE:HD11 | 45:BT:102:ILE:HD11 | 1.34 | 1.09 |
| 28:B6:10:LEU:HD12 | 30:B8:35:GLN:HE22 | 1.13 | 1.09 |
| 40:BO:23:ARG:HH11 | 40:BO:23:ARG:HG2 | 1.15 | 1.09 |
| 1:CA:59:A:H5" | 1:CA:60:A:H5" | 1.28 | 1.09 |
| 2:CB:185:ILE:HG22 | 2:CB:199:TYR:HB2 | 1.32 | 1.09 |
| 44:DS:34:HIS:HB3 | 44:DS:53:SER:HB3 | 1.34 | 1.08 |
| 2:AB:185:ILE:HG22 | 2:AB:199:TYR:HB2 | 1.30 | 1.08 |
| 33:BD:183:ARG:HH11 | 33:BD:183:ARG:HG2 | 1.18 | 1.08 |
| 44:BS:28:VAL:HB | 44:BS:89:ARG:HB2 | 1.13 | 1.08 |
| 41:BP:24:GLY:HA3 | 41:BP:33:ARG:HH21 | 1.06 | 1.08 |
| 41:DP:24:GLY:HA3 | 41:DP:33:ARG:HH21 | 1.09 | 1.08 |
| 23:B1:17:SER:O | 23:B1:44:PRO:HD2 | 1.53 | 1.07 |
| 28:B6:15:GLU:CG | 28:B6:18:ARG:HE | 1.65 | 1.07 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 33:BD:35:LYS:HD3 | 33:BD:63:ARG:HB3 | 1.34 | 1.07 |
| 31:DA:1899:G:H22 | 31:DA:1902:C:N4 | 1.50 | 1.07 |
| 33:DD:44:ASN:HB3 | 33:DD:49:ILE:HA | 1.33 | 1.07 |
| 33:DD:183:ARG:HH11 | 33:DD:183:ARG:HG2 | 1.17 | 1.07 |
| 31:BA:2206:G:H21 | 31:BA:2207:G:C5' | 1.67 | 1.07 |
| 42:BQ:9:TYR:O | 42:BQ:9:TYR:HD2 | 1.37 | 1.07 |
| 51:DZ:53:ILE:HG22 | 51:DZ:71:VAL:HB | 1.37 | 1.07 |
| 31:BA:1821:A:H2' | 31:BA:1822:G:H5" | 1.34 | 1.07 |
| 46:BU:64:ARG:HA | 46:BU:64:ARG:NH2 | 1.70 | 1.07 |
| 39:DN:40:PRO:HA | 46:DU:64:ARG:HH22 | 1.11 | 1.07 |
| 49:DX:77:LYS:HG2 | 49:DX:78:LYS:HG3 | 1.37 | 1.07 |
| 35:BF:101:LEU:HD12 | 35:BF:102:PRO:HD2 | 1.33 | 1.07 |
| 47:BV:85:LYS:O | 47:BV:87:HIS:N | 1.88 | 1.06 |
| 23:B1:41:ARG:HG3 | 23:B1:41:ARG:HH11 | 0.92 | 1.06 |
| 31:DA:511:U:H3' | 31:DA:512:G:H5" | 1.31 | 1.06 |
| 31:DA:571:A:H5' | 31:DA:2030:A:H62 | 0.94 | 1.06 |
| 50:BY:45:VAL:HG13 | 50:BY:62:GLU:HB2 | 1.33 | 1.06 |
| 4:CD:128:VAL:HG12 | 4:CD:129:ASN:ND2 | 1.69 | 1.06 |
| 31:DA:1986:A:H2' | 31:DA:1987:G:H5" | 1.38 | 1.06 |
| 44:DS:85:VAL:HG23 | 44:DS:106:ARG:HB2 | 1.36 | 1.06 |
| 47:DV:47:VAL:HG13 | 47:DV:48:GLY:H | 1.20 | 1.06 |
| 31:DA:2206:G:H21 | 31:DA:2207:G:C5' | 1.67 | 1.06 |
| 50:DY:45:VAL:HG13 | 50:DY:62:GLU:HB2 | 1.38 | 1.06 |
| 31:BA:571:A:H5' | 31:BA:2030:A:H62 | 0.93 | 1.06 |
| 49:BX:77:LYS:HG2 | 49:BX:78:LYS:HG3 | 1.33 | 1.06 |
| 39:DN:47:ALA:HB2 | 39:DN:112:LEU:HD11 | 1.35 | 1.05 |
| 32:BB:87:G:H3' | 32:BB:88:C:H5" | 1.35 | 1.05 |
| 46:DU:64:ARG:HA | 46:DU:64:ARG:NH2 | 1.68 | 1.05 |
| 31:BA:2562:U:H1' | 40:BO:23:ARG:HH12 | 1.16 | 1.05 |
| 33:DD:27:THR:HG21 | 33:DD:83:GLU:HG2 | 1.35 | 1.05 |
| 39:DN:1:MET:HB3 | 47:DV:20:LEU:HD22 | 1.38 | 1.05 |
| 41:BP:120:ALA:O | 25:D3:1:MET:HA | 1.54 | 1.05 |
| 49:BX:25:LYS:HG3 | 49:BX:26:TYR:H | 1.15 | 1.05 |
| 16:CP:28:ARG:HH11 | 16:CP:28:ARG:HG2 | 1.18 | 1.05 |
| 44:BS:89:ARG:O | 44:BS:92:TYR:HB3 | 1.56 | 1.04 |
| 44:BS:34:HIS:HB3 | 44:BS:53:SER:HB3 | 1.32 | 1.04 |
| 51:BZ:53:ILE:HG22 | 51:BZ:71:VAL:HB | 1.33 | 1.04 |
| 31:DA:197:A:H8 | 31:DA:197:A:H5' | 1.18 | 1.04 |
| 31:DA:2012:G:H4' | 48:DW:96:ILE:HD11 | 1.35 | 1.04 |
| 31:DA:2415:G:H4' | 41:DP:67:MET:H | 1.21 | 1.04 |
| 49:DX:25:LYS:HG3 | 49:DX:26:TYR:H | 1.15 | 1.04 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 39:BN:40:PRO:HA | 46:BU:64:ARG:HH22 | 1.15 | 1.04 |
| 41:BP:29:LYS:HD2 | 41:BP:29:LYS:H | 1.19 | 1.04 |
| 49:BX:63:LYS:HD2 | 49:BX:70:LEU:HD13 | 1.37 | 1.04 |
| 10:AJ:34:VAL:HG22 | 10:AJ:74:ILE:HG22 | 1.40 | 1.04 |
| 31:BA:1696:G:H2' | 31:BA:1697:G:H5" | 1.40 | 1.04 |
| 31:BA:1899:G:H22 | 31:BA:1902:C:N4 | 1.55 | 1.04 |
| 31:DA:571:A:H5' | 31:DA:2030:A:N6 | 1.73 | 1.04 |
| 31:DA:1484:G:N2 | 31:DA:1505:C:H41 | 1.55 | 1.04 |
| 34:DE:36:ARG:HH21 | 34:DE:88:GLY:HA2 | 1.17 | 1.04 |
| 31:BA:571:A:H5' | 31:BA:2030:A:N6 | 1.73 | 1.03 |
| 31:BA:2415:G:H4' | 41:BP:67:MET:H | 1.18 | 1.03 |
| 33:BD:44:ASN:HB3 | 33:BD:49:ILE:HA | 1.35 | 1.03 |
| 47:BV:75:PHE:HE1 | 47:BV:89:GLN:HB3 | 1.21 | 1.03 |
| 31:DA:171:G:H2' | 31:DA:172:C:O4' | 1.57 | 1.03 |
| 31:BA:2305:A:H5" | 36:BG:134:GLY:HA3 | 1.40 | 1.03 |
| 50:DY:10:GLY:HA2 | 50:DY:27:VAL:HG13 | 1.04 | 1.03 |
| 31:BA:2317:C:H2' | 31:BA:2318:G:H5' | 1.35 | 1.03 |
| 39:BN:1:MET:HB3 | 47:BV:20:LEU:HD22 | 1.40 | 1.03 |
| 47:BV:82:ARG:HH11 | 47:BV:82:ARG:CG | 1.70 | 1.03 |
| 31:DA:662:G:OP1 | 41:DP:18:ARG:HD2 | 1.58 | 1.03 |
| 31:DA:1747(A):G:H2' | 31:DA:1748:G:H5' | 1.40 | 1.03 |
| 50:BY:96:ILE:HD12 | 50:BY:99:CYS:SG | 1.99 | 1.03 |
| 25:D3:8:LEU:HD13 | 25:D3:31:LEU:HD23 | 1.36 | 1.03 |
| 32:DB:87:G:H3' | 32:DB:88:C:H5" | 1.36 | 1.03 |
| 20:AT:50:GLU:HB3 | 20:AT:100:ILE:HG12 | 1.38 | 1.03 |
| 29:B7:8:ASN:C | 29:B7:8:ASN:HD22 | 1.66 | 1.03 |
| 31:BA:2317:C:C2' | 31:BA:2318:G:H5' | 1.89 | 1.03 |
| 33:BD:35:LYS:HD2 | 33:BD:104:TYR:CE1 | 1.94 | 1.03 |
| 45:BT:23:ARG:HB2 | 45:BT:24:PRO:HD2 | 1.36 | 1.03 |
| 47:BV:19:LYS:HG3 | 47:BV:20:LEU:N | 1.67 | 1.03 |
| 13:CM:3:ARG:HH22 | 36:DG:139:LEU:HD13 | 1.24 | 1.03 |
| 32:DB:15:A:H5' | 32:DB:16:G:C8 | 1.93 | 1.03 |
| 40:DO:23:ARG:HH11 | 40:DO:23:ARG:HG2 | 1.23 | 1.03 |
| 31:BA:171:G:H2' | 31:BA:172:C:O4' | 1.56 | 1.02 |
| 51:BZ:151:HIS:CB | 51:BZ:170:THR:HA | 1.88 | 1.02 |
| 31:DA:1798:U:H5' | 33:DD:259:THR:HG22 | 1.35 | 1.02 |
| 1:CA:1090:U:H2' | 1:CA:1091:U:H6 | 1.22 | 1.02 |
| 31:BA:2012:G:H4' | 48:BW:96:ILE:HD11 | 1.38 | 1.02 |
| 41:BP:16:ARG:HD3 | 41:BP:18:ARG:H | 1.23 | 1.02 |
| 20:CT:50:GLU:HB3 | 20:CT:100:ILE:HG12 | 1.40 | 1.02 |
| 41:DP:29:LYS:HD2 | 41:DP:29:LYS:H | 1.21 | 1.02 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 42:DQ:9:TYR:O | 42:DQ:9:TYR:HD2 | 1.42 | 1.02 |
| 45:DT:23:ARG:HB2 | 45:DT:24:PRO:HD2 | 1.37 | 1.02 |
| 12:AL:47:LYS:HB3 | 12:AL:48:PRO:HD3 | 1.41 | 1.02 |
| 31:BA:1529:G:H21 | 31:BA:1530:C:H5" | 1.21 | 1.02 |
| 47:BV:18:LEU:HD22 | 47:BV:19:LYS:HA | 1.42 | 1.02 |
| 30:D8:32:LEU:HG | 30:D8:34:TRP:HE3 | 1.20 | 1.02 |
| 31:DA:1899:G:N2 | 31:DA:1902:C:H41 | 1.57 | 1.02 |
| 47:DV:1:MET:HE3 | 47:DV:44:LYS:HB2 | 1.41 | 1.02 |
| 50:DY:96:ILE:HD12 | 50:DY:99:CYS:SG | 1.99 | 1.02 |
| 27:B5:16:ARG:HH11 | 27:B5:16:ARG:HG2 | 1.24 | 1.02 |
| 31:BA:102:G:O2' | 31:BA:103:A:OP2 | 1.75 | 1.02 |
| 33:BD:49:ILE:HD11 | 33:BD:52:ARG:HA | 1.38 | 1.02 |
| 42:BQ:81:VAL:O | 42:BQ:82:ARG:CG | 2.08 | 1.02 |
| 12:CL:47:LYS:HB3 | 12:CL:48:PRO:HD3 | 1.41 | 1.02 |
| 22:D0:13:GLY:O | 22:D0:14:ARG:HB2 | 1.56 | 1.02 |
| 23:D1:17:SER:O | 23:D1:44:PRO:HD2 | 1.57 | 1.02 |
| 51:DZ:151:HIS:CB | 51:DZ:170:THR:HA | 1.89 | 1.02 |
| 23:B1:87:PRO:HD2 | 23:B1:88:LYS:H | 1.24 | 1.01 |
| 31:BA:511:U:H3' | 31:BA:512:G:H5" | 1.34 | 1.01 |
| 31:BA:1484:G:N2 | 31:BA:1505:C:H41 | 1.58 | 1.01 |
| 31:DA:2317:C:H2' | 31:DA:2318:G:H5' | 1.36 | 1.01 |
| 31:DA:2562:U:H1' | 40:DO:23:ARG:HH12 | 1.21 | 1.01 |
| 31:DA:2701:C:C3' | 31:DA:2702:U:H5" | 1.91 | 1.01 |
| 4:AD:128:VAL:HG12 | 4:AD:129:ASN:ND2 | 1.75 | 1.01 |
| 37:BH:158:HIS:NE2 | 37:BH:170:ARG:HA | 1.76 | 1.01 |
| 10:CJ:34:VAL:HG22 | 10:CJ:74:ILE:HG22 | 1.40 | 1.01 |
| 31:DA:2206:G:H21 | 31:DA:2207:G:H5' | 0.85 | 1.01 |
| 31:DA:2305:A:H5" | 36:DG:134:GLY:HA3 | 1.42 | 1.01 |
| 38:DI:53:ALA:CB | 38:DI:56:LYS:HG3 | 1.91 | 1.01 |
| 1:AA:1090:U:H2' | 1:AA:1091:U:H6 | 1.22 | 1.01 |
| 24:B2:44:LEU:O | 24:B2:44:LEU:HD12 | 1.60 | 1.01 |
| 39:BN:42:TRP:HB3 | 46:BU:64:ARG:NH1 | 1.74 | 1.01 |
| 23:D1:47:GLN:HG2 | 31:DA:2230:G:H1' | 1.42 | 1.01 |
| 23:D1:87:PRO:HD2 | 23:D1:88:LYS:H | 1.24 | 1.01 |
| 31:DA:197:A:H5' | 31:DA:197:A:C8 | 1.95 | 1.01 |
| 39:DN:42:TRP:HB3 | 46:DU:64:ARG:NH1 | 1.74 | 1.01 |
| 45:DT:50:ILE:HD11 | 45:DT:102:ILE:HD11 | 1.41 | 1.01 |
| 31:BA:1658:C:OP1 | 34:BE:132:HIS:CE1 | 2.13 | 1.01 |
| 36:BG:43:LEU:HD12 | 36:BG:153:ARG:HD2 | 1.43 | 1.01 |
| 31:DA:1529:G:H21 | 31:DA:1530:C:H5" | 1.25 | 1.01 |
| 33:DD:35:LYS:HD2 | 33:DD:104:TYR:CE1 | 1.96 | 1.01 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 40:DO:115:VAL:HG13 | 40:DO:121:VAL:HG21 | 1.40 | 1.01 |
| 31:BA:1986:A:C2' | 31:BA:1987:G:H5" | 1.89 | 1.00 |
| 31:BA:2701:C:C3' | 31:BA:2702:U:H5" | 1.91 | 1.00 |
| 23:D1:41:ARG:HH11 | 23:D1:41:ARG:HG3 | 0.88 | 1.00 |
| 33:DD:35:LYS:HD3 | 33:DD:63:ARG:CB | 1.92 | 1.00 |
| 44:DS:89:ARG:HA | 44:DS:89:ARG:HE | 1.26 | 1.00 |
| 49:DX:63:LYS:HD2 | 49:DX:70:LEU:HD13 | 1.40 | 1.00 |
| 32:BB:15:A:H5' | 32:BB:16:G:C8 | 1.97 | 1.00 |
| 32:BB:20:C:H2' | 32:BB:21:G:H5" | 1.44 | 1.00 |
| 34:DE:132:HIS:CD2 | 34:DE:135:HIS:NE2 | 2.30 | 1.00 |
| 47:BV:75:PHE:CE1 | 47:BV:89:GLN:HB3 | 1.96 | 1.00 |
| 31:DA:102:G:O2' | 31:DA:103:A:OP2 | 1.78 | 1.00 |
| 31:DA:1986:A:C2' | 31:DA:1987:G:H5" | 1.90 | 1.00 |
| 28:D6:15:GLU:CG | 28:D6:18:ARG:HE | 1.75 | 1.00 |
| 31:DA:1019:U:H3 | 31:DA:1142(A):A:H62 | 1.04 | 1.00 |
| 31:DA:2317:C:C2' | 31:DA:2318:G:H5' | 1.90 | 1.00 |
| 31:BA:2206:G:H21 | 31:BA:2207:G:H5' | 0.85 | 1.00 |
| 43:DR:4:LEU:O | 43:DR:4:LEU:HD12 | 1.62 | 1.00 |
| 34:BE:73:GLU:HG3 | 34:BE:74:PRO:HD2 | 1.44 | 1.00 |
| 33:DD:49:ILE:HD11 | 33:DD:52:ARG:HA | 1.40 | 1.00 |
| 4:AD:128:VAL:HG12 | 4:AD:129:ASN:HD22 | 1.24 | 0.99 |
| 22:B0:13:GLY:O | 22:B0:14:ARG:HB2 | 1.54 | 0.99 |
| 24:B2:32:LEU:HD21 | 31:BA:61:G:O2' | 1.60 | 0.99 |
| 34:BE:36:ARG:HH21 | 34:BE:88:GLY:HA2 | 1.25 | 0.99 |
| 31:BA:1986:A:H2' | 31:BA:1987:G:H5" | 1.42 | 0.99 |
| 3:CC:47:LEU:HD21 | 3:CC:68:VAL:HG11 | 1.44 | 0.99 |
| 31:BA:1697:G:H5' | 31:BA:1697:G:H8 | 1.26 | 0.99 |
| 50:BY:76:CYS:SG | 50:BY:77:PRO:HD2 | 2.00 | 0.99 |
| 31:BA:197:A:H8 | 31:BA:197:A:H5' | 1.22 | 0.99 |
| 34:BE:132:HIS:CD2 | 34:BE:135:HIS:NE2 | 2.31 | 0.99 |
| 23:D1:41:ARG:HG3 | 23:D1:41:ARG:NH1 | 1.62 | 0.99 |
| 23:B1:19:GLN:HE21 | 31:BA:379:G:H21 | 1.08 | 0.99 |
| 24:B2:56:GLN:H | 24:B2:56:GLN:NE2 | 1.59 | 0.99 |
| 50:BY:10:GLY:HA2 | 50:BY:27:VAL:HG13 | 0.99 | 0.99 |
| 31:DA:1697:G:H8 | 31:DA:1697:G:H5' | 1.21 | 0.99 |
| 30:B8:32:LEU:O | 30:B8:33:ASN:HB3 | 1.61 | 0.99 |
| 44:BS:89:ARG:HA | 44:BS:89:ARG:HE | 1.26 | 0.99 |
| 41:DP:16:ARG:HD3 | 41:DP:18:ARG:H | 1.22 | 0.99 |
| 43:BR:11:ASN:OD1 | 43:BR:12:ARG:N | 1.95 | 0.98 |
| 31:BA:102:G:H5" | 31:BA:102:G:H8 | 1.24 | 0.98 |
| 47:DV:82:ARG:CG | 47:DV:82:ARG:HH11 | 1.77 | 0.98 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 47:BV:19:LYS:CG | 47:BV:20:LEU:H | 1.75 | 0.98 |
| 31:DA:996:A:H4' | 46:DU:92:ARG:NE | 1.77 | 0.98 |
| 1:CA:1256:A:H61 | 1:CA:1278:U:H1' | 1.26 | 0.98 |
| 25:B3:8:LEU:HD13 | 25:B3:31:LEU:HD23 | 1.44 | 0.98 |
| 31:DA:2656:U:H3 | 31:DA:2665:A:H2 | 1.08 | 0.98 |
| 50:BY:10:GLY:CA | 50:BY:27:VAL:HG13 | 1.94 | 0.98 |
| 47:DV:18:LEU:HD22 | 47:DV:19:LYS:HA | 1.46 | 0.98 |
| 1:AA:664:G:H22 | 1:AA:741:G:H1 | 1.04 | 0.98 |
| 37:DH:158:HIS:NE2 | 37:DH:170:ARG:HA | 1.78 | 0.98 |
| 39:BN:45:ASN:HD22 | 39:BN:45:ASN:H | 1.08 | 0.98 |
| 31:DA:2829:C:H2' | 31:DA:2830:G:H5" | 1.45 | 0.98 |
| 50:DY:28:LYS:HB2 | 50:DY:37:VAL:HB | 1.46 | 0.98 |
| 27:D5:2:ALA:HA | 31:DA:2015:A:H1' | 1.45 | 0.97 |
| 33:DD:39:LYS:HB2 | 33:DD:62:TYR:HB2 | 1.45 | 0.97 |
| 16:AP:4:ILE:HG13 | 16:AP:21:VAL:HG12 | 1.46 | 0.97 |
| 31:BA:1902:C:O2' | 33:BD:244:ARG:HB2 | 1.63 | 0.97 |
| 31:BA:71:A:H2 | 49:BX:31:HIS:HE1 | 0.99 | 0.97 |
| 31:BA:996:A:H4' | 46:BU:92:ARG:NE | 1.78 | 0.97 |
| 41:BP:143:GLY:C | 41:BP:145:PRO:HD3 | 1.89 | 0.97 |
| 1:CA:664:G:H22 | 1:CA:741:G:H1 | 1.06 | 0.97 |
| 24:D2:32:LEU:HD21 | 31:DA:61:G:O2' | 1.64 | 0.97 |
| 34:DE:73:GLU:HG3 | 34:DE:74:PRO:HD2 | 1.45 | 0.97 |
| 31:BA:1798:U:H5' | 33:BD:259:THR:HG22 | 1.41 | 0.97 |
| 36:DG:43:LEU:HD12 | 36:DG:153:ARG:HD2 | 1.44 | 0.97 |
| 4:CD:128:VAL:HG12 | 4:CD:129:ASN:HD22 | 1.19 | 0.97 |
| 33:BD:35:LYS:HD3 | 33:BD:63:ARG:CB | 1.93 | 0.97 |
| 45:DT:30:VAL:HG21 | 45:DT:83:ILE:HG13 | 1.46 | 0.97 |
| 30:B8:62:LEU:HD13 | 31:BA:242:G:H5" | 1.45 | 0.97 |
| 33:BD:39:LYS:HB2 | 33:BD:62:TYR:HB2 | 1.45 | 0.97 |
| 43:BR:4:LEU:HD12 | 43:BR:4:LEU:O | 1.64 | 0.97 |
| 47:BV:43:GLU:N | 47:BV:48:GLY:HA2 | 1.80 | 0.97 |
| 23:B1:13:ILE:HG12 | 23:B1:14:VAL:N | 1.80 | 0.97 |
| 30:D8:32:LEU:O | 30:D8:33:ASN:HB3 | 1.64 | 0.97 |
| 31:DA:102:G:H5" | 31:DA:102:G:H8 | 1.30 | 0.97 |
| 31:DA:145:G:H2' | 31:DA:146:G:H5" | 1.47 | 0.97 |
| 23:B1:85:LEU:HB3 | 23:B1:87:PRO:HD3 | 1.47 | 0.97 |
| 31:DA:1658:C:OP1 | 34:DE:132:HIS:CE1 | 2.18 | 0.97 |
| 1:AA:1256:A:H61 | 1:AA:1278:U:H1' | 1.27 | 0.96 |
| 23:D1:85:LEU:HB3 | 23:D1:87:PRO:HD3 | 1.45 | 0.96 |
| 28:B6:15:GLU:HG2 | 28:B6:18:ARG:HE | 1.29 | 0.96 |
| 23:B1:41:ARG:HG3 | 23:B1:41:ARG:NH1 | 1.66 | 0.96 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 34:DE:61:ARG:H | 34:DE:62:PRO:HD2 | 1.29 | 0.96 |
| 31:BA:659:C:H5' | 31:BA:659:C:H6 | 1.25 | 0.96 |
| 33:BD:25:THR:HG21 | 33:BD:81:ALA:HB1 | 1.47 | 0.96 |
| 28:B6:9:LEU:HD22 | 28:B6:10:LEU:N | 1.80 | 0.96 |
| 34:BE:61:ARG:H | 34:BE:62:PRO:HD2 | 1.30 | 0.96 |
| 49:DX:60:ARG:HB2 | 49:DX:72:LYS:O | 1.65 | 0.96 |
| 24:B2:14:ARG:O | 24:B2:18:PRO:HD3 | 1.63 | 0.96 |
| 30:B8:32:LEU:HG | 30:B8:34:TRP:CE3 | 2.01 | 0.96 |
| 47:BV:47:VAL:HG13 | 47:BV:48:GLY:H | 1.27 | 0.96 |
| 47:DV:79:VAL:O | 47:DV:80:GLN:HB3 | 1.66 | 0.96 |
| 32:DB:20:C:H2' | 32:DB:21:G:H5" | 1.44 | 0.96 |
| 3:AC:47:LEU:HD21 | 3:AC:68:VAL:HG11 | 1.45 | 0.95 |
| 42:BQ:22:LYS:HE2 | 42:BQ:22:LYS:HA | 1.46 | 0.95 |
| 47:BV:79:VAL:O | 47:BV:80:GLN:HB3 | 1.63 | 0.95 |
| 27:D5:16:ARG:HH11 | 27:D5:16:ARG:HG2 | 1.26 | 0.95 |
| 1:AA:1502:A:H2 | 1:AA:1505:G:H1 | 0.98 | 0.95 |
| 42:BQ:23:GLY:O | 42:BQ:100:GLY:HA3 | 1.64 | 0.95 |
| 47:BV:1:MET:HE3 | 47:BV:44:LYS:HB2 | 1.44 | 0.95 |
| 29:D7:8:ASN:HD22 | 29:D7:8:ASN:C | 1.74 | 0.95 |
| 32:DB:7:G:H4' | 44:DS:29:PHE:CD1 | 2.01 | 0.95 |
| 44:DS:61:ASN:HD22 | 44:DS:62:LYS:H | 1.01 | 0.95 |
| 28:B6:15:GLU:CD | 28:B6:18:ARG:HE | 1.75 | 0.95 |
| 31:BA:2068:U:H3 | 31:BA:2430:A:H2 | 0.98 | 0.95 |
| 8:CH:1:MET:HE2 | 8:CH:1:MET:H3 | 1.28 | 0.95 |
| 3:AC:20:SER:HB2 | 3:AC:40:ARG:HH22 | 1.30 | 0.95 |
| 38:DI:52:ARG:O | 38:DI:53:ALA:HB3 | 1.66 | 0.95 |
| 23:D1:13:ILE:HG12 | 23:D1:14:VAL:N | 1.76 | 0.95 |
| 23:D1:16:ASN:HB3 | 23:D1:46:LEU:HG | 1.49 | 0.95 |
| 31:DA:1696:G:H2' | 31:DA:1697:G:H5" | 1.46 | 0.95 |
| 31:BA:1899:G:N2 | 31:BA:1902:C:H41 | 1.64 | 0.95 |
| 50:BY:28:LYS:HB2 | 50:BY:37:VAL:HB | 1.45 | 0.95 |
| 31:DA:1790:C:H5" | 31:DA:1791:A:OP1 | 1.67 | 0.95 |
| 50:DY:76:CYS:SG | 50:DY:77:PRO:HD2 | 2.06 | 0.95 |
| 1:AA:475:G:H2' | 1:AA:476:G:H8 | 1.30 | 0.95 |
| 38:BI:9:LEU:H | 38:BI:13:GLY:HA3 | 1.29 | 0.95 |
| 31:DA:746:A:C2 | 55:DA:3311:ZIT:H161 | 2.00 | 0.95 |
| 47:DV:43:GLU:N | 47:DV:48:GLY:HA2 | 1.82 | 0.95 |
| 33:BD:27:THR:HG21 | 33:BD:83:GLU:HG2 | 1.46 | 0.95 |
| 31:BA:746:A:C2 | 55:BA:3351:ZIT:H161 | 2.02 | 0.95 |
| 31:DA:154:G:H1 | 31:DA:172:C:H42 | 1.14 | 0.95 |
| 27:B5:4:HIS:HB3 | 27:B5:5:PRO:HD3 | 1.49 | 0.94 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:197:A:H5' | 31:BA:197:A:C8 | 1.99 | 0.94 |
| 41:BP:41:ARG:NH2 | 41:BP:41:ARG:HA | 1.82 | 0.94 |
| 29:D7:8:ASN:ND2 | 29:D7:11:LYS:H | 1.64 | 0.94 |
| 31:DA:659:C:H5' | 31:DA:659:C:H6 | 1.32 | 0.94 |
| 49:BX:82:GLN:O | 49:BX:85:PRO:HD2 | 1.67 | 0.94 |
| 1:CA:15:G:H4' | 5:CE:24:ARG:HH12 | 1.29 | 0.94 |
| 16:CP:4:ILE:HG13 | 16:CP:21:VAL:HG12 | 1.48 | 0.94 |
| 47:DV:19:LYS:CG | 47:DV:20:LEU:H | 1.79 | 0.94 |
| 18:AR:79:LEU:HD23 | 18:AR:80:PRO:HD2 | 1.50 | 0.94 |
| 31:BA:1169:G:H1 | 31:BA:1180:C:H42 | 0.96 | 0.94 |
| 30:B8:52:LYS:H | 30:B8:53:PRO:HD2 | 1.32 | 0.94 |
| 50:BY:10:GLY:HA2 | 50:BY:27:VAL:CG1 | 1.94 | 0.94 |
| 3:CC:20:SER:HB2 | 3:CC:40:ARG:HH22 | 1.31 | 0.94 |
| 47:DV:75:PHE:HE1 | 47:DV:89:GLN:HB3 | 1.30 | 0.94 |
| 1:AA:737:A:H2' | 1:AA:738:C:C6 | 2.02 | 0.94 |
| 23:B1:16:ASN:HB3 | 23:B1:46:LEU:HG | 1.49 | 0.94 |
| 18:CR:79:LEU:HD23 | 18:CR:80:PRO:HD2 | 1.48 | 0.94 |
| 31:BA:145:G:H2' | 31:BA:146:G:H5" | 1.47 | 0.94 |
| 33:BD:25:THR:O | 33:BD:27:THR:HB | 1.68 | 0.94 |
| 34:DE:36:ARG:NH2 | 34:DE:88:GLY:HA2 | 1.83 | 0.94 |
| 2:AB:158:LEU:H | 2:AB:158:LEU:HD12 | 1.31 | 0.94 |
| 31:BA:1019:U:H3 | 31:BA:1142(A):A:H62 | 1.09 | 0.94 |
| 31:DA:1024:G:H3' | 31:DA:1025:G:H5" | 1.49 | 0.94 |
| 31:BA:662:G:OP1 | 41:BP:18:ARG:HD2 | 1.68 | 0.94 |
| 44:BS:92:TYR:HD1 | 44:BS:93:LYS:H | 1.16 | 0.94 |
| 47:BV:15:GLU:HB3 | 47:BV:16:PRO:HD2 | 1.47 | 0.94 |
| 28:D6:15:GLU:CD | 28:D6:18:ARG:HE | 1.75 | 0.94 |
| 44:DS:28:VAL:HB | 44:DS:89:ARG:CB | 1.98 | 0.94 |
| 46:DU:83:LEU:HB3 | 46:DU:88:ILE:HD11 | 1.48 | 0.94 |
| 47:DV:19:LYS:HG3 | 47:DV:20:LEU:N | 1.72 | 0.94 |
| 44:BS:28:VAL:HB | 44:BS:89:ARG:CB | 1.97 | 0.94 |
| 29:B7:8:ASN:ND2 | 29:B7:11:LYS:H | 1.66 | 0.94 |
| 31:BA:1747(A):G:H2' | 31:BA:1748:G:H5' | 1.46 | 0.94 |
| 44:BS:14:VAL:HG12 | 44:BS:15:ARG:H | 1.32 | 0.94 |
| 45:BT:100:TYR:HB3 | 45:BT:103:ARG:HE | 1.33 | 0.94 |
| 1:CA:737:A:H2' | 1:CA:738:C:C6 | 2.03 | 0.94 |
| 2:CB:158:LEU:H | 2:CB:158:LEU:HD12 | 1.31 | 0.94 |
| 31:DA:71:A:H2 | 49:DX:31:HIS:HE1 | 1.00 | 0.94 |
| 45:DT:100:TYR:HB3 | 45:DT:103:ARG:HE | 1.33 | 0.94 |
| 24:B2:49:LYS:CD | 24:B2:53:LEU:HD12 | 1.99 | 0.93 |
| 12:CL:8:ASN:HD22 | 17:CQ:34:LYS:HE2 | 1.33 | 0.93 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 30:D8:62:LEU:HD13 | 31:DA:242:G:H5" | 1.49 | 0.93 |
| 31:DA:875:G:H4' | 51:DZ:170:THR:HG21 | 1.50 | 0.93 |
| 31:DA:2334:G:N2 | 44:DS:18:ILE:HD11 | 1.84 | 0.93 |
| 38:DI:9:LEU:H | 38:DI:13:GLY:HA3 | 1.28 | 0.93 |
| 35:BF:123:LEU:HD12 | 35:BF:124:LEU:H | 1.33 | 0.93 |
| 47:BV:22:VAL:O | 47:BV:23:GLU:HB2 | 1.67 | 0.93 |
| 31:DA:1484:G:H21 | 31:DA:1505:C:N4 | 1.64 | 0.93 |
| 33:DD:25:THR:HG21 | 33:DD:81:ALA:HB1 | 1.50 | 0.93 |
| 46:BU:92:ARG:HB2 | 47:BV:11:GLN:NE2 | 1.83 | 0.93 |
| 45:BT:30:VAL:HG21 | 45:BT:83:ILE:HG13 | 1.50 | 0.93 |
| 50:BY:39:VAL:HG12 | 50:BY:40:GLU:H | 1.31 | 0.93 |
| 23:D1:19:GLN:HE21 | 31:DA:379:G:H21 | 1.14 | 0.93 |
| 41:DP:143:GLY:C | 41:DP:145:PRO:HD3 | 1.91 | 0.93 |
| 31:DA:71:A:H2 | 49:DX:31:HIS:CE1 | 1.86 | 0.93 |
| 47:DV:15:GLU:HB3 | 47:DV:16:PRO:HD2 | 1.49 | 0.93 |
| 31:BA:2334:G:N2 | 44:BS:18:ILE:HD11 | 1.83 | 0.93 |
| 31:DA:1771:C:HO2' | 31:DA:1786:A:H8 | 0.94 | 0.93 |
| 47:DV:75:PHE:CE1 | 47:DV:89:GLN:HB3 | 2.02 | 0.93 |
| 30:B8:31:HIS:CG | 31:BA:2419:U:O4 | 2.22 | 0.93 |
| 31:BA:2681:C:H5 | 31:BA:2725:A:H62 | 0.94 | 0.93 |
| 31:DA:528:A:N1 | 31:DA:2042:A:H2' | 1.82 | 0.93 |
| 31:DA:2415:G:O3' | 41:DP:66:GLY:HA3 | 1.69 | 0.93 |
| 1:AA:15:G:H4' | 5:AE:24:ARG:HH12 | 1.31 | 0.93 |
| 31:BA:102:G:H5" | 31:BA:102:G:C8 | 2.03 | 0.93 |
| 31:BA:1484:G:H21 | 31:BA:1505:C:N4 | 1.65 | 0.93 |
| 39:DN:45:ASN:H | 39:DN:45:ASN:HD22 | 1.13 | 0.93 |
| 1:CA:475:G:H2' | 1:CA:476:G:H8 | 1.33 | 0.93 |
| 23:D1:85:LEU:C | 23:D1:87:PRO:HD3 | 1.94 | 0.93 |
| 34:DE:93:VAL:H | 34:DE:95:ILE:CD1 | 1.82 | 0.93 |
| 23:B1:85:LEU:C | 23:B1:87:PRO:HD3 | 1.94 | 0.93 |
| 1:CA:1442(A):G:H3' | 1:CA:1442(B):A:H5" | 1.50 | 0.93 |
| 31:DA:1278:A:OP1 | 43:DR:36:THR:HG22 | 1.67 | 0.93 |
| 15:AO:54:ARG:HG2 | 15:AO:58:MET:HE2 | 1.50 | 0.92 |
| 31:DA:1039:G:H1 | 31:DA:1116:C:H42 | 1.17 | 0.92 |
| 31:DA:2068:U:H3 | 31:DA:2430:A:H2 | 1.04 | 0.92 |
| 50:DY:8:LYS:HZ1 | 50:DY:74:PRO:HD3 | 1.34 | 0.92 |
| 1:AA:954:G:H21 | 1:AA:1227:A:H62 | 1.16 | 0.92 |
| 8:AH:1:MET:HE2 | 8:AH:1:MET:H3 | 1.34 | 0.92 |
| 40:BO:115:VAL:HG13 | 40:BO:121:VAL:HG21 | 1.49 | 0.92 |
| 15:CO:56:LEU:O | 15:CO:60:VAL:HG23 | 1.68 | 0.92 |
| 28:D6:12:GLU:HA | 28:D6:23:THR:HA | 1.49 | 0.92 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 33:DD:35:LYS:HZ3 | 33:DD:104:TYR:HB2 | 1.23 | 0.92 |
| 1:CA:1090:U:H2' | 1:CA:1091:U:C6 | 2.04 | 0.92 |
| 16:CP:45:THR:HG22 | 16:CP:47:ASP:H | 1.34 | 0.92 |
| 51:DZ:39:VAL:HG21 | 51:DZ:44:PHE:HB2 | 1.51 | 0.92 |
| 1:AA:445:G:H2' | 1:AA:446:G:H8 | 1.33 | 0.92 |
| 31:BA:2701:C:H3' | 31:BA:2702:U:H5" | 0.95 | 0.92 |
| 51:BZ:69:THR:HG22 | 51:BZ:90:VAL:HA | 1.51 | 0.92 |
| 28:D6:9:LEU:HD22 | 28:D6:10:LEU:N | 1.84 | 0.92 |
| 33:DD:25:THR:O | 33:DD:27:THR:HB | 1.69 | 0.92 |
| 42:DQ:23:GLY:O | 42:DQ:100:GLY:HA3 | 1.69 | 0.92 |
| 47:DV:22:VAL:O | 47:DV:23:GLU:HB2 | 1.70 | 0.92 |
| 15:AO:56:LEU:O | 15:AO:60:VAL:HG23 | 1.68 | 0.92 |
| 28:B6:28:ARG:HA | 28:B6:32:ASN:HD22 | 1.33 | 0.92 |
| 31:BA:2415:G:O3' | 41:BP:66:GLY:HA3 | 1.69 | 0.92 |
| 43:DR:24:GLN:HE22 | 43:DR:36:THR:HG21 | 1.34 | 0.92 |
| 11:AK:22:HIS:HB3 | 11:AK:29:ILE:HG23 | 1.50 | 0.92 |
| 37:BH:137:ASP:O | 37:BH:138:LYS:HB2 | 1.70 | 0.92 |
| 11:CK:22:HIS:HB3 | 11:CK:29:ILE:HG23 | 1.49 | 0.92 |
| 1:AA:376:G:H4' | 16:AP:5:ARG:HH11 | 1.30 | 0.92 |
| 1:AA:1442(A):G:H3' | 1:AA:1442(B):A:H5" | 1.51 | 0.92 |
| 48:BW:9:TYR:H | 48:BW:102:HIS:HD2 | 1.13 | 0.92 |
| 29:D7:8:ASN:HD21 | 29:D7:11:LYS:H | 0.98 | 0.92 |
| 49:DX:60:ARG:HE | 49:DX:74:PRO:HG2 | 1.34 | 0.92 |
| 24:B2:23:LYS:HB2 | 49:BX:5:TYR:HE1 | 1.32 | 0.92 |
| 31:BA:71:A:H2 | 49:BX:31:HIS:CE1 | 1.87 | 0.92 |
| 32:BB:7:G:H4' | 44:BS:29:PHE:CD1 | 2.05 | 0.92 |
| 41:BP:17:LYS:O | 41:BP:19:VAL:N | 2.02 | 0.92 |
| 30:B8:30:ARG:HH21 | 41:BP:62:LEU:HB2 | 1.33 | 0.92 |
| 31:BA:875:G:H4' | 51:BZ:170:THR:HG21 | 1.50 | 0.92 |
| 41:BP:51:PHE:O | 41:BP:52:GLU:HB2 | 1.68 | 0.92 |
| 1:CA:1502:A:H2 | 1:CA:1505:G:H1 | 0.95 | 0.92 |
| 41:DP:47:ASP:HB3 | 41:DP:48:PRO:C | 1.95 | 0.92 |
| 29:B7:8:ASN:HD21 | 29:B7:11:LYS:H | 1.03 | 0.92 |
| 37:DH:44:VAL:HG12 | 37:DH:45:VAL:H | 1.35 | 0.92 |
| 38:BI:133:HIS:HB2 | 38:BI:134:PRO:HD2 | 1.51 | 0.91 |
| 35:BF:67:GLN:O | 35:BF:67:GLN:HG3 | 1.68 | 0.91 |
| 24:D2:14:ARG:O | 24:D2:18:PRO:HD3 | 1.69 | 0.91 |
| 50:DY:10:GLY:CA | 50:DY:27:VAL:HG13 | 1.99 | 0.91 |
| 31:BA:669:G:H8 | 31:BA:669:G:HO2' | 1.14 | 0.91 |
| 31:BA:1039:G:H1 | 31:BA:1116:C:H42 | 1.15 | 0.91 |
| 31:BA:2829:C:H2' | 31:BA:2830:G:H5" | 1.48 | 0.91 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2808:U:H5' | 31:DA:2891:G:O6 | 1.69 | 0.91 |
| 32:DB:50:G:OP1 | 44:DS:63:THR:HG23 | 1.70 | 0.91 |
| 38:DI:133:HIS:HB2 | 38:DI:134:PRO:HD2 | 1.51 | 0.91 |
| 31:DA:659:C:H6 | 31:DA:659:C:C5' | 1.83 | 0.91 |
| 35:BF:46:ARG:HH11 | 35:BF:46:ARG:HG2 | 1.31 | 0.91 |
| 44:BS:61:ASN:HD22 | 44:BS:62:LYS:H | 0.97 | 0.91 |
| 30:D8:30:ARG:HH21 | 41:DP:62:LEU:HB2 | 1.35 | 0.91 |
| 35:DF:67:GLN:HG3 | 35:DF:67:GLN:O | 1.71 | 0.91 |
| 51:DZ:69:THR:HG22 | 51:DZ:90:VAL:HA | 1.51 | 0.91 |
| 1:CA:376:G:H4' | 16:CP:5:ARG:HH11 | 1.33 | 0.91 |
| 24:D2:44:LEU:HD12 | 24:D2:44:LEU:O | 1.69 | 0.91 |
| 28:D6:12:GLU:HB3 | 28:D6:23:THR:HG22 | 1.50 | 0.91 |
| 35:DF:18:ARG:HG2 | 35:DF:19:GLU:H | 1.36 | 0.91 |
| 18:AR:50:ILE:HD11 | 18:AR:70:ILE:HG21 | 1.53 | 0.91 |
| 22:B0:53:MET:HE3 | 22:B0:57:PHE:HD1 | 1.36 | 0.91 |
| 31:BA:2656:U:H3 | 31:BA:2665:A:H2 | 1.10 | 0.91 |
| 43:BR:8:ARG:HE | 43:BR:8:ARG:HA | 1.34 | 0.91 |
| 30:D8:32:LEU:HG | 30:D8:34:TRP:CE3 | 2.06 | 0.91 |
| 31:DA:658:C:H2' | 31:DA:659:C:H5" | 1.51 | 0.91 |
| 31:DA:1403:C:H5" | 31:DA:1471:A:H1' | 1.52 | 0.91 |
| 31:DA:1497:U:H5' | 31:DA:1498:C:H5 | 1.33 | 0.91 |
| 27:B5:2:ALA:HA | 31:BA:2015:A:H1' | 1.51 | 0.91 |
| 51:BZ:53:ILE:CG2 | 51:BZ:71:VAL:HB | 2.00 | 0.91 |
| 1:CA:735:C:H2' | 1:CA:736:C:H6 | 1.36 | 0.91 |
| 8:AH:86:ILE:HG21 | 8:AH:133:LEU:HD13 | 1.53 | 0.91 |
| 30:B8:52:LYS:N | 30:B8:53:PRO:HD2 | 1.86 | 0.91 |
| 31:BA:658:C:H2' | 31:BA:659:C:H5" | 1.51 | 0.91 |
| 30:D8:31:HIS:CG | 31:DA:2419:U:O4 | 2.25 | 0.91 |
| 1:AA:1090:U:H2' | 1:AA:1091:U:C6 | 2.05 | 0.90 |
| 31:BA:1590:U:H2' | 31:BA:1591:G:H5" | 1.53 | 0.90 |
| 48:DW:9:TYR:H | 48:DW:102:HIS:HD2 | 1.13 | 0.90 |
| 33:BD:175:LEU:HD12 | 33:BD:185:VAL:HG21 | 1.53 | 0.90 |
| 42:BQ:140:ALA:HB3 | 51:BZ:53:ILE:HG13 | 1.54 | 0.90 |
| 31:DA:1378:A:H4' | 31:DA:1379:A:OP1 | 1.71 | 0.90 |
| 31:DA:2701:C:H3' | 31:DA:2702:U:H5" | 0.94 | 0.90 |
| 43:DR:8:ARG:HA | 43:DR:8:ARG:HE | 1.35 | 0.90 |
| 37:BH:43:VAL:HG21 | 37:BH:52:VAL:HG13 | 1.53 | 0.90 |
| 27:D5:4:HIS:HB3 | 27:D5:5:PRO:HD3 | 1.52 | 0.90 |
| 35:DF:46:ARG:HH11 | 35:DF:46:ARG:HG2 | 1.34 | 0.90 |
| 31:BA:1678:G:N2 | 31:BA:1989:G:H22 | 1.70 | 0.90 |
| 24:D2:56:GLN:H | 24:D2:56:GLN:NE2 | 1.68 | 0.90 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:796:C:H2' | 31:DA:797:C:C6 | 2.06 | 0.90 |
| 31:DA:2681:C:H5 | 31:DA:2725:A:H62 | 0.91 | 0.90 |
| 35:DF:24:LEU:HB3 | 35:DF:25:PRO:HD2 | 1.54 | 0.90 |
| 39:DN:3:THR:HG22 | 39:DN:4:TYR:H | 1.36 | 0.90 |
| 47:DV:62:LEU:HD22 | 47:DV:98:GLU:HB2 | 1.53 | 0.90 |
| 31:BA:2787:C:H1' | 34:BE:61:ARG:HB2 | 1.52 | 0.90 |
| 46:BU:88:ILE:C | 46:BU:90:VAL:H | 1.80 | 0.90 |
| 31:DA:2472:G:N1 | 31:DA:2477:C:OP1 | 2.04 | 0.90 |
| 31:DA:2577:A:H5" | 31:DA:2578:G:H5' | 1.54 | 0.90 |
| 34:DE:93:VAL:H | 34:DE:95:ILE:HD13 | 1.32 | 0.90 |
| 47:DV:85:LYS:O | 47:DV:87:HIS:N | 2.05 | 0.90 |
| 31:BA:2415:G:H4' | 41:BP:67:MET:N | 1.87 | 0.90 |
| 33:BD:35:LYS:HZ3 | 33:BD:104:TYR:HB2 | 1.29 | 0.90 |
| 33:BD:35:LYS:HG2 | 33:BD:64:ILE:N | 1.87 | 0.90 |
| 24:D2:46:GLN:HE21 | 24:D2:47:ASN:N | 1.69 | 0.90 |
| 31:DA:2415:G:H4' | 41:DP:67:MET:N | 1.87 | 0.90 |
| 38:DI:53:ALA:HB2 | 38:DI:56:LYS:HG3 | 1.53 | 0.90 |
| 30:B8:32:LEU:HB3 | 30:B8:34:TRP:H | 1.35 | 0.90 |
| 31:BA:154:G:H1 | 31:BA:172:C:H42 | 1.18 | 0.90 |
| 31:BA:286:C:H42 | 31:BA:355:G:H1 | 1.20 | 0.90 |
| 31:BA:659:C:H6 | 31:BA:659:C:C5' | 1.83 | 0.90 |
| 31:BA:1378:A:H4' | 31:BA:1379:A:OP1 | 1.69 | 0.90 |
| 37:BH:85:LYS:NZ | 37:BH:145:ALA:HA | 1.86 | 0.90 |
| 8:CH:86:ILE:HG21 | 8:CH:133:LEU:HD13 | 1.54 | 0.90 |
| 31:DA:2223:G:H2' | 31:DA:2224:G:H5' | 1.51 | 0.90 |
| 42:DQ:140:ALA:HB3 | 51:DZ:53:ILE:HG13 | 1.54 | 0.90 |
| 2:AB:101:MET:HA | 2:AB:108:ILE:HG13 | 1.54 | 0.90 |
| 24:B2:56:GLN:H | 24:B2:56:GLN:HE21 | 1.15 | 0.90 |
| 41:BP:58:THR:O | 41:BP:61:ARG:CZ | 2.20 | 0.90 |
| 23:D1:41:ARG:HH11 | 23:D1:41:ARG:CG | 1.80 | 0.90 |
| 31:DA:676:A:H8 | 31:DA:2069:G:N2 | 1.69 | 0.90 |
| 31:DA:2787:C:H1' | 34:DE:61:ARG:HB2 | 1.52 | 0.90 |
| 41:DP:17:LYS:O | 41:DP:19:VAL:N | 2.05 | 0.90 |
| 23:B1:89:GLU:CD | 23:B1:89:GLU:H | 1.76 | 0.90 |
| 24:B2:46:GLN:HE21 | 24:B2:47:ASN:N | 1.68 | 0.90 |
| 31:BA:993:G:H5' | 47:BV:75:PHE:CE2 | 2.07 | 0.90 |
| 49:BX:60:ARG:HE | 49:BX:74:PRO:HG2 | 1.35 | 0.90 |
| 35:DF:123:LEU:HD12 | 35:DF:124:LEU:H | 1.34 | 0.90 |
| 47:DV:19:LYS:HG2 | 47:DV:96:ILE:HG22 | 1.52 | 0.90 |
| 46:DU:88:ILE:C | 46:DU:90:VAL:H | 1.79 | 0.90 |
| 31:BA:587:C:C5 | 41:BP:33:ARG:HG2 | 2.08 | 0.89 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1497:U:H5' | 31:BA:1498:C:H5 | 1.36 | 0.89 |
| 23:D1:86:SER:N | 23:D1:87:PRO:HD3 | 1.87 | 0.89 |
| 28:B6:10:LEU:HD12 | 30:B8:35:GLN:NE2 | 1.86 | 0.89 |
| 34:BE:116:VAL:O | 34:BE:117:MET:HB3 | 1.72 | 0.89 |
| 33:DD:235:GLY:O | 33:DD:237:GLU:HG2 | 1.72 | 0.89 |
| 40:DO:19:ILE:HG22 | 40:DO:43:VAL:HA | 1.54 | 0.89 |
| 44:DS:92:TYR:HD1 | 44:DS:93:LYS:H | 1.18 | 0.89 |
| 4:AD:189:PRO:HB2 | 4:AD:194:LEU:HD21 | 1.54 | 0.89 |
| 28:B6:12:GLU:HA | 28:B6:23:THR:HA | 1.52 | 0.89 |
| 31:BA:1744:C:H2' | 31:BA:1745:C:H5' | 1.54 | 0.89 |
| 39:BN:3:THR:HG22 | 39:BN:4:TYR:H | 1.37 | 0.89 |
| 1:CA:17:U:H2' | 1:CA:18:C:C6 | 2.07 | 0.89 |
| 15:CO:54:ARG:HG2 | 15:CO:58:MET:HE2 | 1.52 | 0.89 |
| 44:DS:92:TYR:CD1 | 44:DS:93:LYS:N | 2.41 | 0.89 |
| 23:B1:47:GLN:HG2 | 31:BA:2230:G:H1' | 1.51 | 0.89 |
| 41:BP:141:ALA:HB3 | 25:D3:1:MET:SD | 2.13 | 0.89 |
| 46:BU:92:ARG:HD2 | 47:BV:11:GLN:CG | 2.02 | 0.89 |
| 23:B1:10:LYS:HB2 | 23:B1:14:VAL:H | 1.35 | 0.89 |
| 31:BA:2223:G:H2' | 31:BA:2224:G:H5' | 1.54 | 0.89 |
| 35:BF:18:ARG:HG2 | 35:BF:19:GLU:H | 1.35 | 0.89 |
| 23:D1:89:GLU:CD | 23:D1:89:GLU:H | 1.74 | 0.89 |
| 31:DA:993:G:H5' | 47:DV:75:PHE:CE2 | 2.07 | 0.89 |
| 33:DD:87:ASN:N | 33:DD:87:ASN:HD22 | 1.69 | 0.89 |
| 49:DX:82:GLN:O | 49:DX:85:PRO:HD2 | 1.72 | 0.89 |
| 12:AL:8:ASN:HD22 | 17:AQ:34:LYS:HE2 | 1.36 | 0.89 |
| 31:BA:2808:U:H5' | 31:BA:2891:G:O6 | 1.71 | 0.89 |
| 24:D2:16:LEU:H | 24:D2:18:PRO:HD2 | 1.38 | 0.89 |
| 39:DN:3:THR:C | 39:DN:4:TYR:CD1 | 2.50 | 0.89 |
| 28:D6:10:LEU:HD12 | 30:D8:35:GLN:NE2 | 1.86 | 0.89 |
| 28:D6:46:HIS:HB2 | 28:D6:47:THR:N | 1.87 | 0.89 |
| 27:B5:46:CYS:SG | 27:B5:47:PRO:HD2 | 2.13 | 0.89 |
| 30:B8:52:LYS:H | 30:B8:53:PRO:CD | 1.84 | 0.89 |
| 31:BA:658:C:C2' | 31:BA:659:C:H5" | 2.02 | 0.89 |
| 46:BU:92:ARG:HD2 | 47:BV:11:GLN:HG3 | 1.55 | 0.89 |
| 1:CA:250:A:H4' | 1:CA:251:G:O5' | 1.73 | 0.89 |
| 34:DE:152:LYS:HD3 | 39:DN:78:TYR:HB2 | 1.55 | 0.89 |
| 24:B2:45:SER:O | 24:B2:47:ASN:ND2 | 2.05 | 0.89 |
| 49:BX:60:ARG:HB2 | 49:BX:72:LYS:O | 1.71 | 0.89 |
| 31:DA:1378:A:O2' | 31:DA:1379:A:H5" | 1.72 | 0.89 |
| 36:DG:41:GLN:HG2 | 36:DG:155:MET:HB3 | 1.55 | 0.89 |
| 32:DB:15:A:H5' | 32:DB:16:G:H8 | 1.38 | 0.89 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 34:DE:92:THR:H | 34:DE:95:ILE:HD11 | 1.38 | 0.89 |
| 51:DZ:53:ILE:CG2 | 51:DZ:71:VAL:HB | 2.02 | 0.89 |
| 51:DZ:165:VAL:HG12 | 51:DZ:166:SER:H | 1.37 | 0.89 |
| 16:AP:45:THR:HG22 | 16:AP:47:ASP:H | 1.36 | 0.88 |
| 31:BA:1024:G:H3' | 31:BA:1025:G:H5" | 1.52 | 0.88 |
| 31:BA:1790:C:H5" | 31:BA:1791:A:OP1 | 1.73 | 0.88 |
| 48:BW:9:TYR:H | 48:BW:102:HIS:CD2 | 1.91 | 0.88 |
| 50:BY:47:LYS:NZ | 50:BY:47:LYS:HB3 | 1.88 | 0.88 |
| 5:CE:51:VAL:HB | 5:CE:52:PRO:HD3 | 1.56 | 0.88 |
| 31:DA:197:A:H8 | 31:DA:197:A:C5' | 1.84 | 0.88 |
| 47:DV:24:LYS:HB2 | 47:DV:92:THR:OG1 | 1.72 | 0.88 |
| 28:B6:12:GLU:HB3 | 28:B6:23:THR:HG22 | 1.53 | 0.88 |
| 37:BH:44:VAL:HG12 | 37:BH:45:VAL:H | 1.38 | 0.88 |
| 4:CD:189:PRO:HB2 | 4:CD:194:LEU:HD21 | 1.55 | 0.88 |
| 33:DD:63:ARG:HG3 | 33:DD:63:ARG:HH11 | 1.38 | 0.88 |
| 24:B2:16:LEU:H | 24:B2:18:PRO:HD2 | 1.36 | 0.88 |
| 1:CA:954:G:H21 | 1:CA:1227:A:H62 | 1.17 | 0.88 |
| 1:CA:1442(A):G:H3' | 1:CA:1442(B):A:C5' | 2.02 | 0.88 |
| 31:DA:102:G:H5" | 31:DA:102:G:C8 | 2.08 | 0.88 |
| 35:DF:2:LYS:HG3 | 35:DF:25:PRO:HB2 | 1.54 | 0.88 |
| 33:BD:35:LYS:CD | 33:BD:104:TYR:CD1 | 2.56 | 0.88 |
| 27:D5:55:ARG:HD2 | 27:D5:56:LYS:H | 1.38 | 0.88 |
| 35:DF:89:VAL:HG12 | 35:DF:90:PHE:N | 1.88 | 0.88 |
| 46:DU:92:ARG:HB2 | 47:DV:11:GLN:NE2 | 1.89 | 0.88 |
| 1:AA:685:G:O2' | 1:AA:686:U:H5' | 1.74 | 0.88 |
| 23:B1:87:PRO:CD | 23:B1:88:LYS:H | 1.86 | 0.88 |
| 31:BA:1109:C:H5 | 31:BA:1110:G:C4 | 1.92 | 0.88 |
| 34:BE:3:GLY:HA3 | 34:BE:81:ILE:HG13 | 1.56 | 0.88 |
| 47:BV:2:PHE:HE1 | 47:BV:13:ARG:CZ | 1.86 | 0.88 |
| 47:BV:62:LEU:HD22 | 47:BV:98:GLU:HB2 | 1.55 | 0.88 |
| 50:BY:45:VAL:HG11 | 50:BY:60:PHE:O | 1.74 | 0.88 |
| 24:D2:49:LYS:CD | 24:D2:53:LEU:HD12 | 2.04 | 0.88 |
| 24:D2:56:GLN:H | 24:D2:56:GLN:HE21 | 1.17 | 0.88 |
| 32:BB:20:C:C2' | 32:BB:21:G:H5" | 2.03 | 0.88 |
| 41:BP:24:GLY:HA3 | 41:BP:33:ARG:NH2 | 1.87 | 0.88 |
| 1:CA:445:G:H2' | 1:CA:446:G:H8 | 1.36 | 0.88 |
| 33:DD:35:LYS:HZ1 | 33:DD:104:TYR:HB2 | 1.36 | 0.88 |
| 40:DO:66:LYS:H | 40:DO:82:ASN:ND2 | 1.71 | 0.88 |
| 31:BA:2307:G:H21 | 31:BA:2308:G:H5' | 1.37 | 0.88 |
| 31:DA:2469:A:H2 | 31:DA:2481:G:H21 | 1.22 | 0.88 |
| 36:DG:85:GLY:O | 36:DG:87:PRO:HD2 | 1.74 | 0.88 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 50:DY:10:GLY:HA2 | 50:DY:27:VAL:CG1 | 1.98 | 0.88 |
| 31:BA:676:A:H8 | 31:BA:2069:G:N2 | 1.70 | 0.88 |
| 41:BP:47:ASP:HB3 | 41:BP:48:PRO:C | 1.98 | 0.88 |
| 49:BX:25:LYS:CG | 49:BX:26:TYR:H | 1.87 | 0.88 |
| 22:D0:53:MET:HE3 | 22:D0:57:PHE:HD1 | 1.38 | 0.88 |
| 31:DA:1590:U:H2' | 31:DA:1591:G:H5" | 1.55 | 0.88 |
| 1:AA:1065:U:H1' | 1:AA:1066:C:OP2 | 1.74 | 0.88 |
| 51:BZ:39:VAL:HG21 | 51:BZ:44:PHE:HB2 | 1.55 | 0.88 |
| 39:DN:123:TYR:OH | 39:DN:130:HIS:HD2 | 1.57 | 0.88 |
| 46:DU:92:ARG:HD2 | 47:DV:11:GLN:HG3 | 1.54 | 0.88 |
| 1:AA:17:U:H2' | 1:AA:18:C:C6 | 2.08 | 0.88 |
| 31:BA:1378:A:O2' | 31:BA:1379:A:H5" | 1.74 | 0.88 |
| 31:BA:2068:U:N3 | 31:BA:2430:A:H2 | 1.71 | 0.88 |
| 38:BI:53:ALA:HB2 | 38:BI:56:LYS:HG3 | 1.56 | 0.88 |
| 43:BR:24:GLN:HE22 | 43:BR:36:THR:HG21 | 1.35 | 0.88 |
| 1:CA:1065:U:H1' | 1:CA:1066:C:OP2 | 1.74 | 0.88 |
| 23:D1:87:PRO:CD | 23:D1:88:LYS:H | 1.86 | 0.88 |
| 37:DH:137:ASP:O | 37:DH:138:LYS:HB2 | 1.73 | 0.88 |
| 42:DQ:22:LYS:HE2 | 42:DQ:22:LYS:HA | 1.56 | 0.88 |
| 28:B6:9:LEU:HD22 | 28:B6:10:LEU:H | 1.38 | 0.87 |
| 31:BA:2250:G:C5 | 42:BQ:82:ARG:HD2 | 2.09 | 0.87 |
| 32:BB:87:G:C3' | 32:BB:88:C:H5" | 2.03 | 0.87 |
| 45:BT:29:ARG:HG2 | 45:BT:86:ILE:H | 1.39 | 0.87 |
| 31:DA:993:G:H5' | 47:DV:75:PHE:CZ | 2.09 | 0.87 |
| 31:DA:1109:C:H5 | 31:DA:1110:G:C4 | 1.92 | 0.87 |
| 31:DA:1821:A:C2' | 31:DA:1822:G:H5" | 2.04 | 0.87 |
| 33:DD:186:HIS:HD2 | 33:DD:188:GLU:H | 1.22 | 0.87 |
| 1:AA:1442(A):G:H3' | 1:AA:1442(B):A:C5' | 2.02 | 0.87 |
| 31:BA:1021:A:H3' | 31:BA:1021:A:H8 | 1.39 | 0.87 |
| 49:BX:25:LYS:HG3 | 49:BX:26:TYR:N | 1.89 | 0.87 |
| 47:DV:19:LYS:HZ3 | 47:DV:20:LEU:HB2 | 1.37 | 0.87 |
| 35:BF:24:LEU:HB3 | 35:BF:25:PRO:HD2 | 1.53 | 0.87 |
| 45:BT:3:ARG:HB2 | 45:BT:6:LEU:HB3 | 1.55 | 0.87 |
| 49:BX:36:LYS:O | 49:BX:36:LYS:HD2 | 1.74 | 0.87 |
| 31:DA:1038:C:H42 | 31:DA:1117:G:H1 | 1.19 | 0.87 |
| 39:DN:40:PRO:HA | 46:DU:64:ARG:NH2 | 1.87 | 0.87 |
| 31:DA:1430:C:H2' | 31:DA:1431:U:C6 | 2.10 | 0.87 |
| 33:DD:76:PRO:HG2 | 33:DD:98:VAL:HG21 | 1.57 | 0.87 |
| 33:DD:175:LEU:HD12 | 33:DD:185:VAL:HG21 | 1.53 | 0.87 |
| 39:BN:131:GLN:CD | 39:BN:134:ARG:HB3 | 1.99 | 0.87 |
| 44:BS:14:VAL:CG1 | 44:BS:15:ARG:N | 2.37 | 0.87 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 49:BX:37:THR:HG23 | 49:BX:54:VAL:HB | 1.55 | 0.87 |
| 50:BY:45:VAL:HG11 | 50:BY:60:PHE:C | 1.99 | 0.87 |
| 18:CR:50:ILE:HD11 | 18:CR:70:ILE:HG21 | 1.56 | 0.87 |
| 43:DR:10:LEU:HB3 | 43:DR:17:ARG:NE | 1.89 | 0.87 |
| 27:B5:8:LYS:O | 27:B5:9:LYS:HG2 | 1.74 | 0.87 |
| 31:BA:2810:A:H2' | 34:BE:61:ARG:NH2 | 1.88 | 0.87 |
| 35:BF:89:VAL:HG12 | 35:BF:90:PHE:N | 1.89 | 0.87 |
| 41:BP:58:THR:O | 41:BP:61:ARG:NE | 2.07 | 0.87 |
| 41:BP:59:LEU:HA | 41:BP:61:ARG:HH11 | 1.33 | 0.87 |
| 30:D8:52:LYS:H | 30:D8:53:PRO:HD2 | 1.40 | 0.87 |
| 31:DA:1405:U:H2' | 31:DA:1406:U:C6 | 2.10 | 0.87 |
| 31:DA:2681:C:H5 | 31:DA:2725:A:N6 | 1.73 | 0.87 |
| 44:BS:92:TYR:CD1 | 44:BS:93:LYS:N | 2.43 | 0.87 |
| 1:CA:685:G:O2' | 1:CA:686:U:H5' | 1.73 | 0.87 |
| 31:DA:2810:A:H2' | 34:DE:61:ARG:NH2 | 1.90 | 0.87 |
| 49:DX:60:ARG:NE | 49:DX:74:PRO:HG2 | 1.90 | 0.87 |
| 50:DY:27:VAL:HG12 | 50:DY:29:GLU:H | 1.38 | 0.87 |
| 31:BA:1341:U:C2 | 49:BX:77:LYS:HE2 | 2.09 | 0.87 |
| 1:CA:1502:A:H2 | 1:CA:1505:G:N1 | 1.70 | 0.87 |
| 32:DB:87:G:C3' | 32:DB:88:C:H5" | 2.04 | 0.87 |
| 39:DN:42:TRP:HB3 | 46:DU:64:ARG:HH11 | 1.38 | 0.87 |
| 50:DY:28:LYS:O | 50:DY:38:ILE:HB | 1.75 | 0.87 |
| 35:BF:2:LYS:HG3 | 35:BF:25:PRO:HB2 | 1.56 | 0.87 |
| 36:BG:82:LEU:HB3 | 36:BG:87:PRO:HG3 | 1.56 | 0.87 |
| 46:BU:83:LEU:HB3 | 46:BU:88:ILE:HD11 | 1.55 | 0.87 |
| 48:BW:64:MET:O | 48:BW:65:LEU:HB2 | 1.73 | 0.87 |
| 32:DB:20:C:C2' | 32:DB:21:G:H5" | 2.04 | 0.87 |
| 1:AA:626:U:H2' | 1:AA:627:G:H8 | 1.38 | 0.86 |
| 30:B8:35:GLN:HE21 | 30:B8:36:LYS:HZ3 | 1.18 | 0.86 |
| 31:BA:1696:G:C2' | 31:BA:1697:G:H5" | 2.04 | 0.86 |
| 47:BV:72:VAL:HA | 47:BV:88:ARG:NH2 | 1.90 | 0.86 |
| 47:BV:72:VAL:CA | 47:BV:88:ARG:HH22 | 1.86 | 0.86 |
| 30:D8:52:LYS:N | 30:D8:53:PRO:HD2 | 1.90 | 0.86 |
| 31:DA:2308:G:O6 | 31:DA:2310:A:H2' | 1.73 | 0.86 |
| 32:DB:7:G:H3' | 32:DB:8:U:H5" | 1.57 | 0.86 |
| 49:DX:36:LYS:O | 49:DX:36:LYS:HD2 | 1.73 | 0.86 |
| 31:BA:1934:C:H6 | 31:BA:1934:C:H5" | 1.38 | 0.86 |
| 31:DA:1169:G:H1 | 31:DA:1180:C:N4 | 1.73 | 0.86 |
| 36:DG:124:SER:HB2 | 36:DG:131:TYR:CE1 | 2.09 | 0.86 |
| 45:DT:3:ARG:HB2 | 45:DT:6:LEU:HB3 | 1.56 | 0.86 |
| 47:DV:82:ARG:HH11 | 47:DV:82:ARG:HG3 | 1.39 | 0.86 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 49:BX:60:ARG:NE | 49:BX:74:PRO:HG2 | 1.89 | 0.86 |
| 33:DD:35:LYS:HG2 | 33:DD:64:ILE:N | 1.89 | 0.86 |
| 37:DH:43:VAL:HG21 | 37:DH:52:VAL:HG13 | 1.56 | 0.86 |
| 1:AA:250:A:H4' | 1:AA:251:G:O5' | 1.71 | 0.86 |
| 33:BD:65:ILE:HD11 | 33:BD:67:PHE:CE1 | 2.09 | 0.86 |
| 33:BD:235:GLY:O | 33:BD:237:GLU:HG2 | 1.76 | 0.86 |
| 40:BO:19:ILE:HG22 | 40:BO:43:VAL:HA | 1.54 | 0.86 |
| 24:D2:16:LEU:N | 24:D2:18:PRO:HD2 | 1.90 | 0.86 |
| 34:DE:3:GLY:HA3 | 34:DE:81:ILE:HG13 | 1.55 | 0.86 |
| 35:DF:178:PRO:HB2 | 35:DF:201:VAL:HG11 | 1.56 | 0.86 |
| 1:AA:1502:A:H2 | 1:AA:1505:G:N1 | 1.73 | 0.86 |
| 28:B6:10:LEU:CD1 | 30:B8:35:GLN:HE22 | 1.88 | 0.86 |
| 36:BG:127:GLY:HA2 | 36:BG:166:ASP:HB3 | 1.56 | 0.86 |
| 40:BO:66:LYS:H | 40:BO:82:ASN:ND2 | 1.73 | 0.86 |
| 47:BV:19:LYS:HZ1 | 47:BV:20:LEU:HB2 | 1.41 | 0.86 |
| 23:D1:10:LYS:HB2 | 23:D1:14:VAL:H | 1.39 | 0.86 |
| 31:DA:1678:G:N2 | 31:DA:1989:G:H22 | 1.73 | 0.86 |
| 31:DA:1902:C:O2' | 33:DD:244:ARG:HB2 | 1.76 | 0.86 |
| 46:DU:75:ASN:HB2 | 46:DU:78:THR:OG1 | 1.73 | 0.86 |
| 24:B2:16:LEU:N | 24:B2:18:PRO:HD2 | 1.91 | 0.86 |
| 33:BD:166:GLN:HE21 | 33:BD:166:GLN:HA | 1.40 | 0.86 |
| 34:BE:93:VAL:H | 34:BE:95:ILE:HD13 | 1.41 | 0.86 |
| 31:BA:528:A:N1 | 31:BA:2042:A:H2' | 1.89 | 0.86 |
| 17:CQ:86:GLU:O | 17:CQ:90:ILE:HG12 | 1.76 | 0.86 |
| 28:D6:10:LEU:CD1 | 30:D8:35:GLN:HE22 | 1.87 | 0.86 |
| 45:DT:29:ARG:HG2 | 45:DT:86:ILE:H | 1.39 | 0.86 |
| 31:BA:2472:G:N1 | 31:BA:2477:C:OP1 | 2.09 | 0.86 |
| 32:BB:7:G:C3' | 32:BB:8:U:H5" | 2.06 | 0.86 |
| 50:BY:27:VAL:HG12 | 50:BY:29:GLU:H | 1.40 | 0.86 |
| 1:CA:102:G:H2' | 1:CA:103:C:H6 | 1.41 | 0.86 |
| 33:DD:93:ALA:HB3 | 33:DD:105:ILE:HG23 | 1.58 | 0.86 |
| 43:DR:11:ASN:OD1 | 43:DR:12:ARG:N | 2.08 | 0.86 |
| 47:DV:70:ILE:HG13 | 47:DV:90:PRO:CB | 2.06 | 0.86 |
| 24:B2:49:LYS:HD2 | 24:B2:53:LEU:CD1 | 2.05 | 0.86 |
| 36:BG:124:SER:HB2 | 36:BG:131:TYR:CE1 | 2.10 | 0.86 |
| 41:BP:62:LEU:N | 41:BP:62:LEU:HD22 | 1.90 | 0.86 |
| 13:CM:3:ARG:NH2 | 36:DG:139:LEU:HD13 | 1.91 | 0.86 |
| 33:DD:166:GLN:HA | 33:DD:166:GLN:HE21 | 1.41 | 0.86 |
| 34:DE:38:THR:HG22 | 34:DE:40:GLU:H | 1.40 | 0.86 |
| 39:DN:40:PRO:HB3 | 46:DU:68:ALA:HB2 | 1.56 | 0.86 |
| 43:DR:33:ARG:HG3 | 43:DR:115:GLU:HG3 | 1.58 | 0.86 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 2:AB:111:ARG:HG2 | 2:AB:111:ARG:NH1 | 1.84 | 0.86 |
| 31:BA:1430:C:H2' | 31:BA:1431:U:C6 | 2.11 | 0.86 |
| 31:BA:1779:U:H5 | 31:BA:1784:A:N7 | 1.74 | 0.86 |
| 34:BE:92:THR:H | 34:BE:95:ILE:HD11 | 1.41 | 0.86 |
| 51:BZ:165:VAL:HG12 | 51:BZ:166:SER:H | 1.39 | 0.86 |
| 2:CB:101:MET:HA | 2:CB:108:ILE:HG13 | 1.55 | 0.86 |
| 2:CB:111:ARG:HG2 | 2:CB:111:ARG:NH1 | 1.86 | 0.86 |
| 23:D1:92:LYS:C | 23:D1:94:LEU:H | 1.80 | 0.86 |
| 27:D5:57:VAL:HG23 | 27:D5:58:LEU:H | 1.40 | 0.86 |
| 31:DA:1051:G:H5' | 31:DA:2752:C:O2' | 1.76 | 0.86 |
| 1:AA:1169:A:H2' | 1:AA:1170:A:C8 | 2.10 | 0.85 |
| 46:BU:8:VAL:HG11 | 46:BU:12:ARG:CZ | 2.04 | 0.85 |
| 1:CA:673:G:H2' | 1:CA:674:G:C8 | 2.11 | 0.85 |
| 31:DA:1744:C:H2' | 31:DA:1745:C:H5' | 1.55 | 0.85 |
| 23:B1:92:LYS:C | 23:B1:94:LEU:H | 1.82 | 0.85 |
| 24:B2:26:ARG:CZ | 24:B2:29:LYS:HE2 | 2.05 | 0.85 |
| 29:B7:9:ARG:NH1 | 31:BA:1310:G:OP2 | 2.08 | 0.85 |
| 31:BA:1286:A:O2' | 31:BA:1288:U:OP2 | 1.94 | 0.85 |
| 32:BB:50:G:OP1 | 44:BS:63:THR:HG23 | 1.75 | 0.85 |
| 1:CA:1169:A:H2' | 1:CA:1170:A:C8 | 2.11 | 0.85 |
| 31:DA:1779:U:H5 | 31:DA:1784:A:N7 | 1.73 | 0.85 |
| 1:AA:735:C:H2' | 1:AA:736:C:H6 | 1.39 | 0.85 |
| 33:DD:35:LYS:CD | 33:DD:104:TYR:CD1 | 2.58 | 0.85 |
| 49:DX:25:LYS:HG3 | 49:DX:26:TYR:N | 1.91 | 0.85 |
| 1:AA:59:A:H5" | 1:AA:60:A:C5' | 2.05 | 0.85 |
| 8:AH:102:ARG:H | 8:AH:102:ARG:HE | 1.24 | 0.85 |
| 23:B1:34:THR:HG21 | 31:BA:388:G:P | 2.16 | 0.85 |
| 31:BA:993:G:H5' | 47:BV:75:PHE:CZ | 2.11 | 0.85 |
| 39:BN:3:THR:C | 39:BN:4:TYR:CD1 | 2.54 | 0.85 |
| 31:DA:286:C:H42 | 31:DA:355:G:H1 | 1.21 | 0.85 |
| 46:DU:8:VAL:HG11 | 46:DU:12:ARG:CZ | 2.06 | 0.85 |
| 33:BD:35:LYS:HZ1 | 33:BD:104:TYR:HB2 | 1.36 | 0.85 |
| 34:BE:93:VAL:H | 34:BE:95:ILE:CD1 | 1.88 | 0.85 |
| 46:BU:75:ASN:HB2 | 46:BU:78:THR:OG1 | 1.76 | 0.85 |
| 39:DN:131:GLN:CD | 39:DN:134:ARG:HB3 | 2.02 | 0.85 |
| 41:DP:24:GLY:HA3 | 41:DP:33:ARG:NH2 | 1.90 | 0.85 |
| 16:AP:22:THR:HG22 | 16:AP:32:TYR:HA | 1.58 | 0.85 |
| 31:BA:1403:C:H5" | 31:BA:1471:A:H1' | 1.56 | 0.85 |
| 31:BA:2334:G:H21 | 44:BS:18:ILE:CD1 | 1.89 | 0.85 |
| 33:BD:255:LYS:HE3 | 33:BD:255:LYS:N | 1.92 | 0.85 |
| 8:CH:102:ARG:HE | 8:CH:102:ARG:H | 1.25 | 0.85 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2523:G:H2' | 31:DA:2524:G:H5' | 1.57 | 0.85 |
| 32:DB:7:G:C3' | 32:DB:8:U:H5" | 2.06 | 0.85 |
| 36:DG:82:LEU:HB3 | 36:DG:87:PRO:HG3 | 1.58 | 0.85 |
| 31:BA:1598:C:H5' | 49:BX:37:THR:HB | 1.59 | 0.85 |
| 31:BA:2681:C:H5 | 31:BA:2725:A:N6 | 1.75 | 0.85 |
| 33:BD:93:ALA:HB3 | 33:BD:105:ILE:HG23 | 1.59 | 0.85 |
| 28:D6:28:ARG:HA | 28:D6:32:ASN:HD22 | 1.39 | 0.85 |
| 31:BA:2308:G:O6 | 31:BA:2310:A:H2' | 1.76 | 0.85 |
| 31:BA:2463:C:H2' | 31:BA:2464:C:H5' | 1.59 | 0.85 |
| 36:BG:13:GLU:O | 36:BG:14:GLU:HB2 | 1.75 | 0.85 |
| 39:BN:42:TRP:HB3 | 46:BU:64:ARG:HH11 | 1.38 | 0.85 |
| 47:BV:24:LYS:HB2 | 47:BV:92:THR:OG1 | 1.75 | 0.85 |
| 1:CA:1070:U:H2' | 1:CA:1071:C:H6 | 1.42 | 0.85 |
| 41:DP:41:ARG:NH2 | 41:DP:41:ARG:HA | 1.91 | 0.85 |
| 44:DS:29:PHE:N | 44:DS:89:ARG:HD2 | 1.92 | 0.85 |
| 49:DX:37:THR:HG23 | 49:DX:54:VAL:HB | 1.56 | 0.85 |
| 1:AA:491:G:H2' | 1:AA:492:G:H8 | 1.42 | 0.85 |
| 19:AS:40:ILE:HD13 | 19:AS:62:ILE:HD11 | 1.59 | 0.85 |
| 50:BY:81:LYS:HG2 | 50:BY:96:ILE:HG22 | 1.59 | 0.85 |
| 31:DA:1169:G:H1 | 31:DA:1180:C:H42 | 0.90 | 0.85 |
| 48:DW:59:VAL:HG12 | 48:DW:60:ASN:N | 1.89 | 0.85 |
| 28:B6:46:HIS:HB2 | 28:B6:47:THR:N | 1.91 | 0.85 |
| 30:B8:59:LYS:HB2 | 30:B8:59:LYS:NZ | 1.92 | 0.85 |
| 39:BN:40:PRO:HA | 46:BU:64:ARG:NH2 | 1.92 | 0.85 |
| 42:BQ:9:TYR:O | 42:BQ:9:TYR:CD2 | 2.28 | 0.85 |
| 16:CP:20:VAL:HG21 | 16:CP:32:TYR:CG | 2.12 | 0.85 |
| 31:DA:1598:C:H5' | 49:DX:37:THR:HB | 1.57 | 0.85 |
| 41:DP:58:THR:O | 41:DP:61:ARG:CZ | 2.24 | 0.85 |
| 31:BA:2469:A:H2 | 31:BA:2481:G:H21 | 1.21 | 0.84 |
| 36:BG:41:GLN:HG2 | 36:BG:155:MET:HB3 | 1.57 | 0.84 |
| 24:D2:23:LYS:HB2 | 49:DX:5:TYR:HE1 | 1.42 | 0.84 |
| 31:DA:587:C:H4' | 31:DA:588:U:OP2 | 1.76 | 0.84 |
| 1:AA:1070:U:H2' | 1:AA:1071:C:H6 | 1.42 | 0.84 |
| 31:BA:2310:A:O2' | 31:BA:2311:A:H5" | 1.76 | 0.84 |
| 31:DA:2068:U:N3 | 31:DA:2430:A:H2 | 1.74 | 0.84 |
| 48:DW:9:TYR:H | 48:DW:102:HIS:CD2 | 1.93 | 0.84 |
| 50:DY:96:ILE:HG13 | 50:DY:99:CYS:C | 2.02 | 0.84 |
| 4:AD:57:ARG:HH22 | 5:AE:107:ARG:HD3 | 1.42 | 0.84 |
| 5:AE:51:VAL:HB | 5:AE:52:PRO:HD3 | 1.60 | 0.84 |
| 10:AJ:40:LEU:HB2 | 10:AJ:41:PRO:HD2 | 1.59 | 0.84 |
| 23:B1:86:SER:N | 23:B1:87:PRO:HD3 | 1.91 | 0.84 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 27:B5:57:VAL:HG23 | 27:B5:58:LEU:H | 1.39 | 0.84 |
| 50:BY:96:ILE:HG13 | 50:BY:99:CYS:C | 2.02 | 0.84 |
| 49:DX:60:ARG:HG2 | 49:DX:74:PRO:HD3 | 1.57 | 0.84 |
| 1:AA:1505:G:H4' | 1:AA:1506:U:H5" | 1.58 | 0.84 |
| 30:B8:32:LEU:CB | 30:B8:34:TRP:H | 1.89 | 0.84 |
| 31:BA:197:A:H8 | 31:BA:197:A:C5' | 1.89 | 0.84 |
| 34:BE:36:ARG:NH2 | 34:BE:88:GLY:HA2 | 1.91 | 0.84 |
| 44:BS:14:VAL:HG12 | 44:BS:15:ARG:N | 1.92 | 0.84 |
| 47:BV:15:GLU:HB3 | 47:BV:16:PRO:CD | 2.07 | 0.84 |
| 49:BX:60:ARG:HG2 | 49:BX:74:PRO:HD3 | 1.59 | 0.84 |
| 50:BY:75:ILE:HD11 | 50:BY:79:CYS:HA | 1.57 | 0.84 |
| 31:DA:493:G:H2' | 31:DA:494:G:H5" | 1.59 | 0.84 |
| 31:DA:1697:G:H5' | 31:DA:1697:G:C8 | 2.11 | 0.84 |
| 31:DA:2463:C:H2' | 31:DA:2464:C:H5' | 1.60 | 0.84 |
| 27:B5:55:ARG:HD2 | 27:B5:56:LYS:H | 1.41 | 0.84 |
| 41:BP:71:VAL:HG12 | 41:BP:72:PRO:HD3 | 1.59 | 0.84 |
| 47:BV:71:LEU:HD13 | 47:BV:72:VAL:HG23 | 1.59 | 0.84 |
| 1:AA:445:G:H2' | 1:AA:446:G:C8 | 2.12 | 0.84 |
| 31:BA:71:A:H8 | 31:BA:71:A:H5' | 1.42 | 0.84 |
| 34:BE:199:ARG:O | 34:BE:200:GLU:HB3 | 1.74 | 0.84 |
| 47:DV:2:PHE:HE1 | 47:DV:13:ARG:CZ | 1.89 | 0.84 |
| 33:BD:186:HIS:HD2 | 33:BD:188:GLU:H | 1.25 | 0.84 |
| 10:CJ:40:LEU:HB2 | 10:CJ:41:PRO:HD2 | 1.57 | 0.84 |
| 31:DA:1341:U:C2 | 49:DX:77:LYS:HE2 | 2.13 | 0.84 |
| 42:DQ:30:GLY:HA2 | 42:DQ:107:ALA:HB2 | 1.59 | 0.84 |
| 46:DU:92:ARG:HD2 | 47:DV:11:GLN:CG | 2.08 | 0.84 |
| 16:AP:22:THR:CG2 | 16:AP:32:TYR:HA | 2.08 | 0.84 |
| 35:BF:126:VAL:HG21 | 35:BF:129:PHE:CZ | 2.13 | 0.84 |
| 47:BV:83:ARG:O | 47:BV:84:LYS:HG3 | 1.77 | 0.84 |
| 31:DA:2327:A:H2' | 31:DA:2328:A:C8 | 2.13 | 0.84 |
| 41:DP:105:LEU:O | 41:DP:106:LEU:HB2 | 1.77 | 0.84 |
| 31:DA:527:C:OP2 | 31:DA:2779:U:H5 | 1.60 | 0.84 |
| 31:DA:1696:G:C2' | 31:DA:1697:G:H5" | 2.06 | 0.84 |
| 36:DG:13:GLU:O | 36:DG:14:GLU:HB2 | 1.77 | 0.84 |
| 48:DW:64:MET:O | 48:DW:65:LEU:HB2 | 1.77 | 0.84 |
| 1:AA:627:G:H2' | 1:AA:628:G:H8 | 1.40 | 0.84 |
| 5:AE:10:MET:HB2 | 5:AE:32:VAL:HG22 | 1.60 | 0.84 |
| 10:AJ:78:ASN:HB2 | 10:AJ:81:THR:HG23 | 1.60 | 0.84 |
| 48:BW:59:VAL:HG12 | 48:BW:60:ASN:N | 1.93 | 0.84 |
| 1:CA:1442:G:O2' | 1:CA:1442(A):G:H5" | 1.77 | 0.84 |
| 2:CB:172:ILE:HD12 | 2:CB:172:ILE:H | 1.43 | 0.84 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 23:D1:19:GLN:HG3 | 23:D1:44:PRO:HG3 | 1.59 | 0.84 |
| 28:D6:10:LEU:H | 28:D6:10:LEU:HD22 | 1.42 | 0.84 |
| 31:DA:1639:U:H2' | 31:DA:1640:C:H5" | 1.60 | 0.84 |
| 31:DA:2733:A:H2' | 31:DA:2734:A:H5" | 1.59 | 0.84 |
| 28:B6:27:LYS:HD3 | 31:BA:2285:C:OP2 | 1.78 | 0.83 |
| 33:BD:63:ARG:HH11 | 33:BD:63:ARG:HG3 | 1.40 | 0.83 |
| 45:BT:91:ARG:HB2 | 45:BT:116:ALA:HA | 1.60 | 0.83 |
| 31:DA:571:A:C5' | 31:DA:2030:A:H62 | 1.87 | 0.83 |
| 47:DV:15:GLU:HB3 | 47:DV:16:PRO:CD | 2.08 | 0.83 |
| 16:AP:20:VAL:HG21 | 16:AP:32:TYR:CG | 2.13 | 0.83 |
| 31:BA:2523:G:H2' | 31:BA:2524:G:H5' | 1.59 | 0.83 |
| 32:BB:15:A:H5' | 32:BB:16:G:H8 | 1.43 | 0.83 |
| 37:BH:89:ILE:HD11 | 37:BH:129:THR:HB | 1.61 | 0.83 |
| 49:BX:33:LYS:C | 49:BX:35:THR:H | 1.83 | 0.83 |
| 2:CB:91:PRO:HG3 | 2:CB:154:LEU:HB2 | 1.60 | 0.83 |
| 36:DG:127:GLY:HA2 | 36:DG:166:ASP:HB3 | 1.57 | 0.83 |
| 17:AQ:86:GLU:O | 17:AQ:90:ILE:HG12 | 1.78 | 0.83 |
| 43:BR:5:LYS:N | 43:BR:5:LYS:HD2 | 1.92 | 0.83 |
| 28:D6:15:GLU:HG2 | 28:D6:18:ARG:HE | 1.43 | 0.83 |
| 30:D8:59:LYS:HB2 | 30:D8:59:LYS:NZ | 1.92 | 0.83 |
| 34:DE:52:LEU:O | 34:DE:74:PRO:HA | 1.79 | 0.83 |
| 31:BA:543:C:C6 | 31:BA:547:A:N7 | 2.46 | 0.83 |
| 31:BA:1146:C:H2' | 31:BA:1147:C:H5' | 1.58 | 0.83 |
| 31:BA:1170:G:H1 | 31:BA:1179:C:H42 | 1.25 | 0.83 |
| 31:BA:2392:A:H2 | 31:BA:2424:C:H42 | 1.25 | 0.83 |
| 34:BE:197:ILE:HD11 | 34:BE:199:ARG:NH2 | 1.92 | 0.83 |
| 47:BV:69:LYS:HB2 | 47:BV:93:GLU:OE2 | 1.78 | 0.83 |
| 33:DD:58:HIS:HD2 | 33:DD:59:LYS:N | 1.76 | 0.83 |
| 37:DH:149:ARG:HA | 37:DH:162:ILE:HG13 | 1.60 | 0.83 |
| 49:DX:25:LYS:CG | 49:DX:26:TYR:H | 1.89 | 0.83 |
| 50:DY:47:LYS:NZ | 50:DY:47:LYS:HB3 | 1.94 | 0.83 |
| 50:DY:81:LYS:HG2 | 50:DY:96:ILE:HG22 | 1.60 | 0.83 |
| 31:DA:71:A:C2 | 49:DX:31:HIS:HE1 | 1.93 | 0.83 |
| 31:DA:1188:U:C2' | 31:DA:1189:A:H5' | 2.09 | 0.83 |
| 31:DA:1934:C:H5" | 31:DA:1934:C:H6 | 1.39 | 0.83 |
| 31:DA:2307:G:H21 | 31:DA:2308:G:H5' | 1.42 | 0.83 |
| 33:BD:210:GLY:O | 33:BD:211:ARG:HB3 | 1.78 | 0.83 |
| 35:BF:101:LEU:HD12 | 35:BF:102:PRO:CD | 2.08 | 0.83 |
| 31:DA:286:C:H2' | 31:DA:287:C:H5' | 1.59 | 0.83 |
| 50:DY:39:VAL:HG12 | 50:DY:40:GLU:H | 1.41 | 0.83 |
| 31:BA:1494:A:H4' | 31:BA:1495:A:OP1 | 1.78 | 0.83 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:1505:G:H4' | 1:CA:1506:U:H5" | 1.58 | 0.83 |
| 13:CM:66:LEU:H | 13:CM:66:LEU:HD12 | 1.44 | 0.83 |
| 33:DD:255:LYS:HE3 | 33:DD:255:LYS:N | 1.93 | 0.83 |
| 1:AA:664:G:N2 | 1:AA:741:G:H1 | 1.77 | 0.83 |
| 31:BA:493:G:H2' | 31:BA:494:G:H5" | 1.60 | 0.83 |
| 35:BF:178:PRO:HB2 | 35:BF:201:VAL:HG11 | 1.60 | 0.83 |
| 1:CA:522:C:H41 | 12:CL:53:ARG:HH22 | 1.24 | 0.83 |
| 31:DA:271(F):C:H2' | 31:DA:271(G):C:H6 | 1.43 | 0.83 |
| 33:DD:35:LYS:CD | 33:DD:63:ARG:HB3 | 2.06 | 0.83 |
| 35:DF:101:LEU:HD12 | 35:DF:102:PRO:CD | 2.07 | 0.83 |
| 39:DN:3:THR:HA | 39:DN:4:TYR:CE1 | 2.12 | 0.83 |
| 31:BA:286:C:H2' | 31:BA:287:C:H5' | 1.60 | 0.83 |
| 31:BA:529:A:H62 | 31:BA:2041:U:H3 | 1.27 | 0.83 |
| 31:BA:1771:C:HO2' | 31:BA:1786:A:H8 | 0.83 | 0.83 |
| 12:CL:102:ARG:CG | 12:CL:102:ARG:HH11 | 1.92 | 0.83 |
| 27:D5:8:LYS:O | 27:D5:9:LYS:HG2 | 1.77 | 0.83 |
| 33:DD:108:PRO:HB3 | 33:DD:143:HIS:HE1 | 1.43 | 0.83 |
| 41:DP:38:GLN:HG3 | 41:DP:39:LYS:H | 1.44 | 0.83 |
| 27:B5:35:GLU:HB2 | 27:B5:49:CYS:HB3 | 1.60 | 0.83 |
| 1:CA:1003:G:H2' | 1:CA:1004:A:H4' | 1.61 | 0.83 |
| 31:DA:2106:G:H1' | 31:DA:2184:G:N2 | 1.94 | 0.83 |
| 33:DD:210:GLY:O | 33:DD:211:ARG:HB3 | 1.76 | 0.83 |
| 36:DG:47:LYS:HD3 | 36:DG:81:LYS:HD2 | 1.59 | 0.83 |
| 37:DH:41:MET:HE2 | 37:DH:55:PRO:HD3 | 1.61 | 0.83 |
| 39:DN:3:THR:C | 39:DN:4:TYR:CG | 2.57 | 0.83 |
| 1:AA:200:G:H1 | 1:AA:217:C:H42 | 1.26 | 0.82 |
| 12:AL:38:THR:HG23 | 12:AL:39:VAL:HG23 | 1.59 | 0.82 |
| 31:BA:271(F):C:H2' | 31:BA:271(G):C:H6 | 1.44 | 0.82 |
| 31:BA:1603:A:H5' | 31:BA:1603:A:H8 | 1.43 | 0.82 |
| 31:BA:2733:A:H2' | 31:BA:2734:A:H5" | 1.60 | 0.82 |
| 33:BD:35:LYS:CD | 33:BD:63:ARG:HB3 | 2.09 | 0.82 |
| 36:BG:47:LYS:HD3 | 36:BG:81:LYS:HD2 | 1.60 | 0.82 |
| 45:BT:30:VAL:HG23 | 45:BT:30:VAL:O | 1.77 | 0.82 |
| 49:BX:82:GLN:HB3 | 49:BX:85:PRO:HG2 | 1.61 | 0.82 |
| 4:CD:57:ARG:HH22 | 5:CE:107:ARG:HD3 | 1.39 | 0.82 |
| 31:DA:819:A:C4 | 31:DA:1189:A:C2 | 2.66 | 0.82 |
| 31:DA:1021:A:H3' | 31:DA:1021:A:H8 | 1.44 | 0.82 |
| 37:DH:85:LYS:NZ | 37:DH:145:ALA:HA | 1.94 | 0.82 |
| 32:BB:7:G:H3' | 32:BB:8:U:H5" | 1.59 | 0.82 |
| 39:BN:3:THR:HA | 39:BN:4:TYR:CE1 | 2.13 | 0.82 |
| 50:BY:45:VAL:HG13 | 50:BY:62:GLU:CB | 2.08 | 0.82 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:626:U:H2' | 1:CA:627:G:H8 | 1.45 | 0.82 |
| 18:CR:59:SER:HB3 | 18:CR:62:GLU:HG3 | 1.59 | 0.82 |
| 23:D1:34:THR:HG21 | 31:DA:388:G:P | 2.18 | 0.82 |
| 30:D8:52:LYS:H | 30:D8:53:PRO:CD | 1.91 | 0.82 |
| 31:DA:997:G:OP1 | 46:DU:93:LYS:HD3 | 1.79 | 0.82 |
| 1:AA:673:G:H2' | 1:AA:674:G:C8 | 2.13 | 0.82 |
| 1:AA:1281:U:H4' | 1:AA:1282:C:OP2 | 1.79 | 0.82 |
| 13:AM:66:LEU:H | 13:AM:66:LEU:HD12 | 1.45 | 0.82 |
| 36:BG:85:GLY:O | 36:BG:87:PRO:HD2 | 1.79 | 0.82 |
| 37:BH:149:ARG:HA | 37:BH:162:ILE:HG13 | 1.61 | 0.82 |
| 47:BV:83:ARG:CG | 47:BV:83:ARG:HH11 | 1.92 | 0.82 |
| 5:CE:10:MET:HB2 | 5:CE:32:VAL:HG22 | 1.60 | 0.82 |
| 31:DA:543:C:C6 | 31:DA:547:A:N7 | 2.47 | 0.82 |
| 34:DE:197:ILE:HD11 | 34:DE:199:ARG:NH2 | 1.93 | 0.82 |
| 1:AA:522:C:H41 | 12:AL:53:ARG:HH22 | 1.26 | 0.82 |
| 12:AL:102:ARG:HH11 | 12:AL:102:ARG:CG | 1.91 | 0.82 |
| 31:BA:796:C:H2' | 31:BA:797:C:C6 | 2.14 | 0.82 |
| 33:BD:108:PRO:HB3 | 33:BD:143:HIS:CE1 | 2.15 | 0.82 |
| 34:BE:152:LYS:HD3 | 39:BN:78:TYR:HB2 | 1.61 | 0.82 |
| 1:CA:627:G:H2' | 1:CA:628:G:H8 | 1.41 | 0.82 |
| 2:CB:84:GLU:HB3 | 2:CB:219:VAL:HG21 | 1.60 | 0.82 |
| 31:BA:1292:U:H2' | 31:BA:1293:C:C6 | 2.15 | 0.82 |
| 32:BB:88:C:H2' | 32:BB:89:G:C8 | 2.14 | 0.82 |
| 38:BI:133:HIS:HB2 | 38:BI:134:PRO:CD | 2.08 | 0.82 |
| 41:BP:124:LYS:HG2 | 41:BP:143:GLY:HA3 | 1.60 | 0.82 |
| 43:BR:33:ARG:HG3 | 43:BR:115:GLU:HG3 | 1.61 | 0.82 |
| 44:BS:29:PHE:N | 44:BS:89:ARG:HD2 | 1.94 | 0.82 |
| 44:BS:34:HIS:HB3 | 44:BS:53:SER:CB | 2.09 | 0.82 |
| 50:BY:8:LYS:HZ1 | 50:BY:74:PRO:HD3 | 1.44 | 0.82 |
| 50:BY:39:VAL:HG12 | 50:BY:40:GLU:N | 1.91 | 0.82 |
| 4:CD:62:GLN:HE21 | 4:CD:62:GLN:HA | 1.44 | 0.82 |
| 38:DI:52:ARG:HA | 38:DI:52:ARG:HE | 1.44 | 0.82 |
| 43:DR:5:LYS:N | 43:DR:5:LYS:HD2 | 1.94 | 0.82 |
| 49:DX:60:ARG:HG2 | 49:DX:74:PRO:CD | 2.09 | 0.82 |
| 2:AB:91:PRO:HG3 | 2:AB:154:LEU:HB2 | 1.61 | 0.82 |
| 31:BA:1278:A:OP1 | 43:BR:36:THR:HG22 | 1.80 | 0.82 |
| 33:BD:35:LYS:NZ | 33:BD:65:ILE:HA | 1.95 | 0.82 |
| 38:BI:53:ALA:CB | 38:BI:56:LYS:HG3 | 2.10 | 0.82 |
| 42:BQ:81:VAL:C | 42:BQ:82:ARG:HG2 | 2.02 | 0.82 |
| 47:BV:82:ARG:HH11 | 47:BV:82:ARG:HG3 | 1.43 | 0.82 |
| 50:BY:95:LYS:CD | 50:BY:100:ALA:HB1 | 2.06 | 0.82 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 30:D8:22:VAL:HB | 30:D8:53:PRO:HB3 | 1.61 | 0.82 |
| 32:DB:88:C:H2' | 32:DB:89:G:C8 | 2.14 | 0.82 |
| 50:DY:45:VAL:HG11 | 50:DY:60:PHE:C | 2.04 | 0.82 |
| 30:B8:32:LEU:HD23 | 30:B8:35:GLN:CA | 2.09 | 0.82 |
| 33:BD:108:PRO:HB3 | 33:BD:143:HIS:HE1 | 1.44 | 0.82 |
| 34:BE:52:LEU:O | 34:BE:74:PRO:HA | 1.79 | 0.82 |
| 44:BS:89:ARG:HA | 44:BS:89:ARG:NE | 1.94 | 0.82 |
| 19:CS:40:ILE:HD13 | 19:CS:62:ILE:HD11 | 1.61 | 0.82 |
| 50:DY:45:VAL:HG11 | 50:DY:60:PHE:O | 1.79 | 0.82 |
| 31:BA:2327:A:H2' | 31:BA:2328:A:C8 | 2.15 | 0.82 |
| 31:BA:2463:C:C2' | 31:BA:2464:C:H5' | 2.09 | 0.82 |
| 51:BZ:166:SER:HB2 | 51:BZ:168:GLU:N | 1.95 | 0.82 |
| 31:DA:911:A:H2' | 42:DQ:9:TYR:OH | 1.80 | 0.82 |
| 31:DA:1434:A:O2' | 31:DA:1435:G:H5' | 1.79 | 0.82 |
| 31:DA:1494:A:H4' | 31:DA:1495:A:OP1 | 1.79 | 0.82 |
| 31:DA:2334:G:H21 | 44:DS:18:ILE:CD1 | 1.88 | 0.82 |
| 35:DF:126:VAL:HG21 | 35:DF:129:PHE:CZ | 2.15 | 0.82 |
| 37:DH:85:LYS:HD2 | 37:DH:141:VAL:HG13 | 1.62 | 0.82 |
| 41:DP:17:LYS:C | 41:DP:19:VAL:H | 1.86 | 0.82 |
| 50:DY:75:ILE:HD11 | 50:DY:79:CYS:HA | 1.62 | 0.82 |
| 1:AA:353:A:H5' | 1:AA:353:A:H8 | 1.44 | 0.82 |
| 31:BA:1188:U:C2' | 31:BA:1189:A:H5' | 2.10 | 0.82 |
| 49:BX:60:ARG:HG2 | 49:BX:74:PRO:CD | 2.10 | 0.82 |
| 1:CA:353:A:H8 | 1:CA:353:A:H5' | 1.44 | 0.82 |
| 8:CH:20:TYR:HD1 | 8:CH:65:TYR:CD2 | 1.97 | 0.82 |
| 24:D2:55:ARG:HH22 | 49:DX:3:THR:HG22 | 1.45 | 0.82 |
| 31:DA:2571:C:H5' | 31:DA:2572:A:H5" | 1.62 | 0.82 |
| 44:DS:34:HIS:HB3 | 44:DS:53:SER:CB | 2.10 | 0.82 |
| 46:DU:112:ARG:HH11 | 46:DU:112:ARG:CG | 1.93 | 0.82 |
| 46:BU:112:ARG:HH11 | 46:BU:112:ARG:CG | 1.92 | 0.82 |
| 47:BV:82:ARG:HH11 | 47:BV:82:ARG:HG2 | 1.43 | 0.82 |
| 12:CL:38:THR:HG23 | 12:CL:39:VAL:HG23 | 1.62 | 0.82 |
| 28:D6:27:LYS:HD3 | 31:DA:2285:C:OP2 | 1.80 | 0.82 |
| 31:DA:1146:C:H2' | 31:DA:1147:C:H5' | 1.60 | 0.82 |
| 36:DG:64:THR:HG23 | 36:DG:65:GLY:H | 1.45 | 0.82 |
| 38:DI:133:HIS:HB2 | 38:DI:134:PRO:CD | 2.09 | 0.82 |
| 44:DS:89:ARG:HA | 44:DS:89:ARG:NE | 1.94 | 0.82 |
| 18:AR:59:SER:HB3 | 18:AR:62:GLU:HG3 | 1.61 | 0.81 |
| 31:BA:2571:C:H5' | 31:BA:2572:A:H5" | 1.62 | 0.81 |
| 32:BB:86:G:H2' | 32:BB:87:G:C8 | 2.15 | 0.81 |
| 35:BF:123:LEU:HD12 | 35:BF:124:LEU:N | 1.95 | 0.81 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 36:BG:64:THR:HG23 | 36:BG:65:GLY:H | 1.44 | 0.81 |
| 45:BT:30:VAL:HG12 | 45:BT:44:ASP:OD2 | 1.80 | 0.81 |
| 47:BV:70:ILE:HG13 | 47:BV:90:PRO:CB | 2.08 | 0.81 |
| 31:DA:1286:A:O2' | 31:DA:1288:U:OP2 | 1.97 | 0.81 |
| 31:DA:2273:A:O2' | 31:DA:2274:A:H5' | 1.80 | 0.81 |
| 34:DE:95:ILE:H | 34:DE:95:ILE:HD12 | 1.44 | 0.81 |
| 35:DF:103:LYS:HA | 35:DF:106:ARG:HG3 | 1.63 | 0.81 |
| 27:B5:50:GLY:HA3 | 27:B5:56:LYS:HG2 | 1.62 | 0.81 |
| 31:BA:2845:G:O2' | 31:BA:2846:G:H5' | 1.81 | 0.81 |
| 43:BR:10:LEU:HB3 | 43:BR:17:ARG:NE | 1.95 | 0.81 |
| 16:CP:22:THR:CG2 | 16:CP:32:TYR:HA | 2.10 | 0.81 |
| 31:DA:93:G:O2' | 50:DY:51:VAL:HG21 | 1.80 | 0.81 |
| 33:DD:183:ARG:HH11 | 33:DD:183:ARG:CG | 1.91 | 0.81 |
| 45:DT:91:ARG:HB2 | 45:DT:116:ALA:HA | 1.61 | 0.81 |
| 31:BA:71:A:C2 | 49:BX:31:HIS:HE1 | 1.92 | 0.81 |
| 31:BA:139(A):G:N2 | 49:BX:44:GLU:OE1 | 2.12 | 0.81 |
| 31:BA:1179:C:O2 | 31:BA:1179:C:H2' | 1.80 | 0.81 |
| 33:BD:25:THR:O | 33:BD:25:THR:HG23 | 1.81 | 0.81 |
| 39:BN:18:ALA:HB1 | 39:BN:21:LYS:HG3 | 1.62 | 0.81 |
| 41:BP:17:LYS:C | 41:BP:19:VAL:H | 1.87 | 0.81 |
| 47:BV:13:ARG:HG3 | 47:BV:13:ARG:NH1 | 1.85 | 0.81 |
| 49:BX:77:LYS:CG | 49:BX:78:LYS:HG3 | 2.11 | 0.81 |
| 24:D2:26:ARG:CZ | 24:D2:29:LYS:HE2 | 2.10 | 0.81 |
| 1:CA:445:G:H2' | 1:CA:446:G:C8 | 2.15 | 0.81 |
| 20:CT:89:ARG:NH2 | 20:CT:104:LEU:HD21 | 1.95 | 0.81 |
| 31:DA:2250:G:C5 | 42:DQ:82:ARG:HD2 | 2.15 | 0.81 |
| 38:DI:75:LEU:HD12 | 38:DI:76:THR:H | 1.46 | 0.81 |
| 47:DV:13:ARG:HG3 | 47:DV:13:ARG:NH1 | 1.84 | 0.81 |
| 31:BA:1405:U:H2' | 31:BA:1406:U:C6 | 2.14 | 0.81 |
| 42:BQ:30:GLY:HA2 | 42:BQ:107:ALA:HB2 | 1.60 | 0.81 |
| 1:CA:403:C:O2' | 1:CA:404:U:H5' | 1.81 | 0.81 |
| 25:D3:6:VAL:HG13 | 25:D3:54:VAL:HG11 | 1.60 | 0.81 |
| 38:DI:53:ALA:HB1 | 38:DI:57:ARG:H | 1.43 | 0.81 |
| 1:AA:102:G:H2' | 1:AA:103:C:H6 | 1.42 | 0.81 |
| 20:AT:89:ARG:NH2 | 20:AT:104:LEU:HD21 | 1.95 | 0.81 |
| 23:B1:19:GLN:HG3 | 23:B1:44:PRO:HG3 | 1.63 | 0.81 |
| 33:BD:76:PRO:HG2 | 33:BD:98:VAL:HG21 | 1.62 | 0.81 |
| 43:BR:11:ASN:CG | 43:BR:12:ARG:H | 1.89 | 0.81 |
| 13:CM:81:LEU:HB3 | 13:CM:89:GLY:HA2 | 1.62 | 0.81 |
| 31:DA:587:C:C5 | 41:DP:33:ARG:HG2 | 2.16 | 0.81 |
| 31:DA:658:C:C2' | 31:DA:659:C:H5" | 2.08 | 0.81 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:870:A:C5' | 42:DQ:7:MET:HB2 | 2.07 | 0.81 |
| 31:DA:2562:U:H1' | 40:DO:23:ARG:NH1 | 1.96 | 0.81 |
| 33:DD:108:PRO:HB3 | 33:DD:143:HIS:CE1 | 2.15 | 0.81 |
| 42:DQ:35:VAL:HG13 | 42:DQ:130:LYS:HB3 | 1.60 | 0.81 |
| 49:DX:36:LYS:NZ | 49:DX:38:GLU:O | 2.13 | 0.81 |
| 31:BA:2036:C:H5' | 31:BA:2036:C:H6 | 1.45 | 0.81 |
| 41:BP:38:GLN:HG3 | 41:BP:39:LYS:H | 1.45 | 0.81 |
| 50:BY:35:TYR:CE2 | 50:BY:69:ALA:HB3 | 2.15 | 0.81 |
| 31:DA:71:A:OP2 | 31:DA:71:A:H3' | 1.81 | 0.81 |
| 31:DA:1019:U:H3 | 31:DA:1142(A):A:N6 | 1.79 | 0.81 |
| 33:BD:183:ARG:HH11 | 33:BD:183:ARG:CG | 1.94 | 0.81 |
| 38:BI:75:LEU:HD12 | 38:BI:76:THR:H | 1.46 | 0.81 |
| 44:BS:61:ASN:ND2 | 44:BS:62:LYS:H | 1.76 | 0.81 |
| 50:BY:97:ARG:HG3 | 50:BY:97:ARG:O | 1.79 | 0.81 |
| 1:CA:1435:G:H2' | 1:CA:1436:U:C6 | 2.16 | 0.81 |
| 31:DA:2463:C:C2' | 31:DA:2464:C:H5' | 2.11 | 0.81 |
| 32:DB:29:A:P | 44:DS:32:LEU:HD12 | 2.21 | 0.81 |
| 47:DV:72:VAL:HA | 47:DV:88:ARG:NH2 | 1.95 | 0.81 |
| 1:AA:1442:G:O2' | 1:AA:1442(A):G:H5" | 1.81 | 0.81 |
| 30:B8:32:LEU:C | 30:B8:34:TRP:H | 1.89 | 0.81 |
| 31:BA:1762:A:H8 | 31:BA:1762:A:O5' | 1.64 | 0.81 |
| 39:BN:73:THR:O | 39:BN:75:TYR:N | 2.12 | 0.81 |
| 1:CA:365:U:H5" | 1:CA:366:C:OP1 | 1.81 | 0.81 |
| 1:CA:444:C:H2' | 1:CA:445:G:H8 | 1.46 | 0.81 |
| 2:CB:67:THR:HG22 | 2:CB:90:MET:HE1 | 1.62 | 0.81 |
| 31:DA:142:A:H5' | 31:DA:142(A):C:OP2 | 1.81 | 0.81 |
| 1:AA:17:U:H2' | 1:AA:18:C:H6 | 1.44 | 0.81 |
| 31:BA:1039:G:H1 | 31:BA:1116:C:N4 | 1.79 | 0.81 |
| 31:BA:2273:A:O2' | 31:BA:2274:A:H5' | 1.78 | 0.81 |
| 16:CP:72:ARG:HH21 | 16:CP:73:LEU:HD21 | 1.46 | 0.81 |
| 31:DA:1179:C:H2' | 31:DA:1179:C:O2 | 1.80 | 0.81 |
| 35:DF:123:LEU:HD12 | 35:DF:124:LEU:N | 1.94 | 0.81 |
| 49:DX:82:GLN:HB3 | 49:DX:85:PRO:HG2 | 1.62 | 0.81 |
| 1:AA:1432:G:OP1 | 45:BT:107:ASP:HB2 | 1.80 | 0.80 |
| 4:AD:119:GLN:HG2 | 4:AD:123:HIS:HD2 | 1.46 | 0.80 |
| 24:B2:46:GLN:HE21 | 24:B2:46:GLN:C | 1.89 | 0.80 |
| 39:BN:3:THR:C | 39:BN:4:TYR:CG | 2.59 | 0.80 |
| 50:BY:8:LYS:CE | 50:BY:72:VAL:HG23 | 2.11 | 0.80 |
| 1:CA:877:C:H5" | 8:CH:88:LYS:HD3 | 1.61 | 0.80 |
| 1:CA:1238:A:H62 | 1:CA:1299:A:N6 | 1.78 | 0.80 |
| 24:D2:46:GLN:HE21 | 24:D2:46:GLN:C | 1.88 | 0.80 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 27:D5:46:CYS:SG | 27:D5:47:PRO:HD2 | 2.21 | 0.80 |
| 30:D8:32:LEU:CB | 30:D8:35:GLN:H | 1.94 | 0.80 |
| 31:DA:796:C:H2' | 31:DA:797:C:H6 | 1.44 | 0.80 |
| 31:DA:966:G:H2' | 31:DA:967:C:H6 | 1.46 | 0.80 |
| 31:DA:2310:A:O2' | 31:DA:2311:A:H5" | 1.80 | 0.80 |
| 50:DY:8:LYS:CE | 50:DY:72:VAL:HG23 | 2.10 | 0.80 |
| 1:AA:1003:G:H2' | 1:AA:1004:A:H4' | 1.62 | 0.80 |
| 4:AD:119:GLN:HG2 | 4:AD:123:HIS:CD2 | 2.17 | 0.80 |
| 13:AM:91:ARG:HB2 | 13:AM:98:VAL:HG21 | 1.63 | 0.80 |
| 31:BA:1696:G:H2' | 31:BA:1697:G:C5' | 2.11 | 0.80 |
| 31:BA:2577:A:H5" | 31:BA:2578:G:H5' | 1.61 | 0.80 |
| 9:CI:4:TYR:HB2 | 9:CI:19:LEU:HB2 | 1.63 | 0.80 |
| 24:D2:46:GLN:NE2 | 24:D2:47:ASN:N | 2.29 | 0.80 |
| 39:DN:40:PRO:CA | 46:DU:64:ARG:HH22 | 1.94 | 0.80 |
| 39:DN:65:LYS:HA | 39:DN:65:LYS:CE | 2.12 | 0.80 |
| 41:DP:59:LEU:HA | 41:DP:61:ARG:HH11 | 1.43 | 0.80 |
| 1:AA:877:C:H5" | 8:AH:88:LYS:HD3 | 1.63 | 0.80 |
| 41:BP:121:LYS:HD3 | 25:D3:2:PRO:HD3 | 1.63 | 0.80 |
| 3:CC:52:LEU:HD23 | 3:CC:52:LEU:H | 1.45 | 0.80 |
| 19:CS:36:ARG:HH12 | 19:CS:75:ALA:HB3 | 1.46 | 0.80 |
| 22:D0:41:ARG:H | 22:D0:41:ARG:HD2 | 1.44 | 0.80 |
| 23:D1:19:GLN:OE1 | 23:D1:44:PRO:HB3 | 1.81 | 0.80 |
| 35:DF:20:LEU:HD13 | 35:DF:203:GLN:OE1 | 1.81 | 0.80 |
| 44:DS:24:LEU:HB3 | 44:DS:85:VAL:HG12 | 1.63 | 0.80 |
| 1:AA:975:A:H4' | 1:AA:976:G:H5" | 1.64 | 0.80 |
| 1:AA:1238:A:H62 | 1:AA:1299:A:N6 | 1.79 | 0.80 |
| 12:AL:124:LYS:HD2 | 12:AL:125:PRO:HD2 | 1.64 | 0.80 |
| 31:BA:1494:A:C4' | 31:BA:1495:A:OP1 | 2.29 | 0.80 |
| 31:BA:1798:U:H5' | 33:BD:259:THR:CG2 | 2.11 | 0.80 |
| 31:BA:2106:G:H1' | 31:BA:2184:G:N2 | 1.95 | 0.80 |
| 40:BO:104:ARG:CZ | 45:BT:33:LYS:HD2 | 2.12 | 0.80 |
| 43:BR:116:LEU:O | 43:BR:117:VAL:HB | 1.79 | 0.80 |
| 47:BV:64:HIS:HD2 | 47:BV:94:LEU:HD21 | 1.47 | 0.80 |
| 1:CA:491:G:H2' | 1:CA:492:G:H8 | 1.47 | 0.80 |
| 37:DH:89:ILE:HD11 | 37:DH:129:THR:HB | 1.63 | 0.80 |
| 41:DP:101:VAL:HG12 | 41:DP:106:LEU:HB3 | 1.63 | 0.80 |
| 33:BD:35:LYS:HD3 | 33:BD:63:ARG:CA | 2.11 | 0.80 |
| 33:BD:58:HIS:HD2 | 33:BD:59:LYS:N | 1.79 | 0.80 |
| 31:DA:1497:U:H5' | 31:DA:1498:C:C5 | 2.16 | 0.80 |
| 42:DQ:9:TYR:O | 42:DQ:9:TYR:CD2 | 2.31 | 0.80 |
| 28:B6:15:GLU:HB3 | 28:B6:18:ARG:HG2 | 1.62 | 0.80 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 30:B8:22:VAL:HB | 30:B8:53:PRO:HB3 | 1.61 | 0.80 |
| 31:BA:27:G:N2 | 31:BA:512:G:H1' | 1.97 | 0.80 |
| 31:BA:93:G:O2' | 50:BY:51:VAL:HG21 | 1.82 | 0.80 |
| 31:BA:2660:A:H5' | 31:BA:2661:G:N2 | 1.97 | 0.80 |
| 33:BD:25:THR:HG22 | 33:BD:82:ILE:O | 1.80 | 0.80 |
| 45:BT:83:ILE:HG13 | 45:BT:84:GLN:H | 1.46 | 0.80 |
| 1:CA:532:A:H61 | 3:CC:193:TYR:HB3 | 1.46 | 0.80 |
| 28:D6:13:CYS:HB3 | 28:D6:49:HIS:HB3 | 1.61 | 0.80 |
| 31:DA:27:G:N2 | 31:DA:512:G:H1' | 1.97 | 0.80 |
| 31:DA:1292:U:H2' | 31:DA:1293:C:C6 | 2.16 | 0.80 |
| 48:DW:18:ARG:HH11 | 48:DW:18:ARG:HG2 | 1.45 | 0.80 |
| 2:AB:172:ILE:H | 2:AB:172:ILE:HD12 | 1.45 | 0.80 |
| 25:B3:54:VAL:HG12 | 25:B3:55:ARG:N | 1.97 | 0.80 |
| 28:B6:10:LEU:H | 28:B6:10:LEU:HD22 | 1.47 | 0.80 |
| 31:BA:1051:G:H5' | 31:BA:2752:C:O2' | 1.81 | 0.80 |
| 44:BS:101:LEU:HD13 | 44:BS:102:ALA:O | 1.82 | 0.80 |
| 5:CE:72:GLN:O | 5:CE:73:ASN:HB2 | 1.82 | 0.80 |
| 31:DA:84:A:N6 | 31:DA:102:G:H1' | 1.97 | 0.80 |
| 31:DA:1798:U:H5' | 33:DD:259:THR:CG2 | 2.09 | 0.80 |
| 15:AO:55:GLY:HA2 | 15:AO:58:MET:HE3 | 1.63 | 0.80 |
| 19:AS:36:ARG:HH12 | 19:AS:75:ALA:HB3 | 1.46 | 0.80 |
| 31:BA:1021:A:H3' | 31:BA:1021:A:C8 | 2.17 | 0.80 |
| 33:BD:87:ASN:N | 33:BD:87:ASN:HD22 | 1.80 | 0.80 |
| 46:BU:90:VAL:HG12 | 46:BU:91:ASP:H | 1.46 | 0.80 |
| 31:DA:2395:C:H5' | 31:DA:2395:C:H6 | 1.44 | 0.80 |
| 34:DE:76:ARG:HG3 | 34:DE:195:LEU:HD12 | 1.63 | 0.80 |
| 18:AR:58:LEU:HB3 | 18:AR:62:GLU:HB2 | 1.64 | 0.80 |
| 23:B1:41:ARG:HH11 | 23:B1:41:ARG:CG | 1.82 | 0.80 |
| 30:B8:30:ARG:O | 30:B8:31:HIS:C | 2.23 | 0.80 |
| 30:B8:32:LEU:CB | 30:B8:35:GLN:H | 1.94 | 0.80 |
| 31:BA:83:G:N2 | 31:BA:102:G:O2' | 2.12 | 0.80 |
| 31:BA:143:G:H1' | 49:BX:38:GLU:HG3 | 1.63 | 0.80 |
| 31:BA:911:A:H2' | 42:BQ:9:TYR:OH | 1.82 | 0.80 |
| 1:CA:1281:U:H4' | 1:CA:1282:C:OP2 | 1.80 | 0.80 |
| 19:CS:42:PRO:O | 19:CS:43:GLU:HB3 | 1.81 | 0.80 |
| 24:D2:49:LYS:HD2 | 24:D2:53:LEU:CD1 | 2.11 | 0.80 |
| 40:DO:23:ARG:HG2 | 40:DO:23:ARG:NH1 | 1.97 | 0.80 |
| 49:DX:33:LYS:C | 49:DX:35:THR:H | 1.89 | 0.80 |
| 1:AA:1169:A:H2' | 1:AA:1170:A:H8 | 1.43 | 0.80 |
| 1:AA:1199:U:H4' | 10:AJ:54:PHE:CZ | 2.16 | 0.80 |
| 2:AB:154:LEU:HD23 | 2:AB:154:LEU:H | 1.47 | 0.80 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 6:AF:24:GLU:HG3 | 6:AF:28:ARG:NH1 | 1.97 | 0.80 |
| 31:BA:2092:U:H4' | 31:BA:2093:G:O5' | 1.82 | 0.80 |
| 39:BN:83:LYS:HE3 | 39:BN:85:ILE:HD11 | 1.62 | 0.80 |
| 45:BT:29:ARG:CB | 45:BT:85:LYS:HA | 2.12 | 0.80 |
| 16:CP:22:THR:HG22 | 16:CP:32:TYR:HA | 1.63 | 0.80 |
| 31:DA:1300:U:O2' | 31:DA:1626:G:C2 | 2.34 | 0.80 |
| 16:AP:28:ARG:HH11 | 16:AP:28:ARG:CG | 1.91 | 0.79 |
| 1:CA:737:A:H2' | 1:CA:738:C:H6 | 1.46 | 0.79 |
| 1:CA:975:A:H4' | 1:CA:976:G:H5" | 1.64 | 0.79 |
| 5:CE:50:GLU:HG3 | 5:CE:52:PRO:HD2 | 1.65 | 0.79 |
| 18:CR:58:LEU:HB3 | 18:CR:62:GLU:HB2 | 1.64 | 0.79 |
| 27:D5:35:GLU:HB2 | 27:D5:49:CYS:HB3 | 1.64 | 0.79 |
| 23:B1:19:GLN:NE2 | 31:BA:379:G:H21 | 1.79 | 0.79 |
| 31:BA:1639:U:H2' | 31:BA:1640:C:H5" | 1.64 | 0.79 |
| 41:BP:41:ARG:HA | 41:BP:41:ARG:HH21 | 1.44 | 0.79 |
| 1:CA:1446:U:H4' | 1:CA:1447:A:N7 | 1.98 | 0.79 |
| 4:CD:148:VAL:HG12 | 4:CD:149:ALA:H | 1.45 | 0.79 |
| 15:CO:55:GLY:HA2 | 15:CO:58:MET:HE3 | 1.64 | 0.79 |
| 30:D8:32:LEU:CB | 30:D8:34:TRP:H | 1.96 | 0.79 |
| 31:DA:1833:U:H2' | 31:DA:1834:U:H6 | 1.47 | 0.79 |
| 1:AA:444:C:H2' | 1:AA:445:G:H8 | 1.45 | 0.79 |
| 1:AA:707:C:O2' | 1:AA:708:C:H5' | 1.82 | 0.79 |
| 1:AA:1201:A:H1' | 1:AA:1202:G:OP2 | 1.83 | 0.79 |
| 1:AA:1291:G:H4' | 9:AI:38:GLN:O | 1.82 | 0.79 |
| 31:BA:1012:U:O4 | 39:BN:28:THR:HG21 | 1.82 | 0.79 |
| 33:BD:70:TRP:HZ3 | 33:BD:146:GLU:OE2 | 1.66 | 0.79 |
| 38:BI:144:VAL:HG12 | 38:BI:145:VAL:H | 1.47 | 0.79 |
| 50:BY:28:LYS:O | 50:BY:38:ILE:HB | 1.83 | 0.79 |
| 31:DA:142:A:C8 | 31:DA:1408:C:H1' | 2.17 | 0.79 |
| 31:DA:1406:U:H2' | 31:DA:1407:C:C6 | 2.17 | 0.79 |
| 31:DA:2660:A:H3' | 31:DA:2660:A:N3 | 1.97 | 0.79 |
| 32:DB:47:C:O2' | 44:DS:93:LYS:HG2 | 1.82 | 0.79 |
| 45:DT:83:ILE:HG13 | 45:DT:84:GLN:H | 1.44 | 0.79 |
| 9:AI:96:LEU:HG | 9:AI:102:LEU:HB2 | 1.64 | 0.79 |
| 31:BA:1529:G:N2 | 31:BA:1530:C:H5" | 1.97 | 0.79 |
| 32:BB:20:C:H2' | 32:BB:21:G:C5' | 2.12 | 0.79 |
| 34:BE:59:VAL:HG22 | 34:BE:63:LEU:HA | 1.64 | 0.79 |
| 39:BN:65:LYS:HA | 39:BN:65:LYS:CE | 2.11 | 0.79 |
| 4:CD:119:GLN:HG2 | 4:CD:123:HIS:HD2 | 1.47 | 0.79 |
| 10:CJ:78:ASN:HB2 | 10:CJ:81:THR:HG23 | 1.62 | 0.79 |
| 11:CK:21:ILE:HB | 11:CK:84:VAL:HA | 1.64 | 0.79 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:71:A:H5' | 31:DA:71:A:H8 | 1.48 | 0.79 |
| 47:DV:5:VAL:CG2 | 47:DV:36:PRO:HB2 | 2.11 | 0.79 |
| 1:AA:1321:C:C5' | 1:AA:1322:C:H5" | 2.12 | 0.79 |
| 4:AD:148:VAL:HG12 | 4:AD:149:ALA:H | 1.46 | 0.79 |
| 25:B3:6:VAL:HG13 | 25:B3:54:VAL:HG11 | 1.63 | 0.79 |
| 31:BA:1038:C:H42 | 31:BA:1117:G:H1 | 1.29 | 0.79 |
| 31:BA:2562:U:H1' | 40:BO:23:ARG:NH1 | 1.96 | 0.79 |
| 45:BT:23:ARG:O | 45:BT:25:GLY:N | 2.16 | 0.79 |
| 48:BW:18:ARG:HG2 | 48:BW:18:ARG:HH11 | 1.48 | 0.79 |
| 9:CI:96:LEU:HG | 9:CI:102:LEU:HB2 | 1.62 | 0.79 |
| 23:D1:8:SER:N | 23:D1:46:LEU:HD11 | 1.96 | 0.79 |
| 31:DA:1170:G:H1 | 31:DA:1179:C:H42 | 1.25 | 0.79 |
| 44:DS:61:ASN:ND2 | 44:DS:62:LYS:H | 1.80 | 0.79 |
| 1:AA:1446:U:H4' | 1:AA:1447:A:N7 | 1.98 | 0.79 |
| 31:BA:1495:A:N3 | 31:BA:1496:A:C2 | 2.51 | 0.79 |
| 31:BA:2712:U:H5" | 31:BA:2712:U:O2 | 1.83 | 0.79 |
| 39:BN:123:TYR:OH | 39:BN:130:HIS:HD2 | 1.64 | 0.79 |
| 5:CE:101:ILE:HD11 | 5:CE:119:LEU:HA | 1.64 | 0.79 |
| 30:D8:32:LEU:HB3 | 30:D8:34:TRP:H | 1.47 | 0.79 |
| 31:DA:870:A:OP1 | 42:DQ:7:MET:HE2 | 1.82 | 0.79 |
| 33:DD:35:LYS:HD3 | 33:DD:63:ARG:CA | 2.12 | 0.79 |
| 33:DD:65:ILE:HD11 | 33:DD:67:PHE:CE1 | 2.18 | 0.79 |
| 36:DG:60:LEU:O | 36:DG:64:THR:HG22 | 1.83 | 0.79 |
| 41:DP:58:THR:O | 41:DP:61:ARG:NE | 2.16 | 0.79 |
| 45:DT:23:ARG:O | 45:DT:25:GLY:N | 2.16 | 0.79 |
| 47:DV:62:LEU:CD2 | 47:DV:98:GLU:HB2 | 2.13 | 0.79 |
| 51:DZ:149:SER:HB2 | 51:DZ:172:ALA:O | 1.83 | 0.79 |
| 1:AA:1226:C:C4 | 13:AM:104:ARG:HB2 | 2.17 | 0.79 |
| 31:BA:1697:G:H5' | 31:BA:1697:G:C8 | 2.16 | 0.79 |
| 36:BG:131:TYR:HB3 | 36:BG:159:VAL:HG13 | 1.64 | 0.79 |
| 42:BQ:8:LYS:HD2 | 42:BQ:9:TYR:H | 1.47 | 0.79 |
| 51:BZ:151:HIS:HB3 | 51:BZ:170:THR:CA | 2.08 | 0.79 |
| 13:CM:68:GLY:O | 13:CM:69:GLU:HB2 | 1.83 | 0.79 |
| 38:DI:2:LYS:HB2 | 38:DI:39:ALA:HB3 | 1.64 | 0.79 |
| 8:AH:10:LEU:HD23 | 8:AH:10:LEU:N | 1.97 | 0.79 |
| 23:B1:19:GLN:OE1 | 23:B1:44:PRO:HB3 | 1.83 | 0.79 |
| 28:B6:22:ALA:HB2 | 28:B6:39:TYR:CE2 | 2.17 | 0.79 |
| 31:BA:659:C:H5' | 31:BA:659:C:C6 | 2.15 | 0.79 |
| 34:BE:76:ARG:HG3 | 34:BE:195:LEU:HD12 | 1.64 | 0.79 |
| 1:CA:1201:A:H1' | 1:CA:1202:G:OP2 | 1.82 | 0.79 |
| 2:CB:88:ALA:HB2 | 2:CB:219:VAL:HG13 | 1.64 | 0.79 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:676:A:H8 | 31:DA:2069:G:H21 | 0.87 | 0.79 |
| 31:DA:1012:U:O4 | 39:DN:28:THR:HG21 | 1.83 | 0.79 |
| 31:DA:2393:A:H5' | 41:DP:62:LEU:HB3 | 1.62 | 0.79 |
| 33:DD:25:THR:HG22 | 33:DD:82:ILE:O | 1.83 | 0.79 |
| 34:DE:59:VAL:HG22 | 34:DE:63:LEU:HA | 1.64 | 0.79 |
| 34:DE:116:VAL:O | 34:DE:117:MET:HB3 | 1.80 | 0.79 |
| 46:DU:88:ILE:O | 46:DU:90:VAL:N | 2.14 | 0.79 |
| 9:AI:4:TYR:HB2 | 9:AI:19:LEU:HB2 | 1.64 | 0.79 |
| 16:AP:72:ARG:HH21 | 16:AP:73:LEU:HD21 | 1.47 | 0.79 |
| 28:B6:46:HIS:CB | 28:B6:47:THR:N | 2.46 | 0.79 |
| 1:CA:80:G:H1 | 1:CA:89:C:N4 | 1.81 | 0.79 |
| 1:CA:707:C:O2' | 1:CA:708:C:H5' | 1.83 | 0.79 |
| 31:DA:1039:G:H1 | 31:DA:1116:C:N4 | 1.80 | 0.79 |
| 35:DF:32:LEU:HD11 | 35:DF:105:VAL:HG13 | 1.64 | 0.79 |
| 46:DU:65:ILE:HG12 | 46:DU:96:ALA:CB | 2.13 | 0.79 |
| 2:AB:88:ALA:HB2 | 2:AB:219:VAL:HG13 | 1.65 | 0.79 |
| 31:BA:1169:G:H1 | 31:BA:1180:C:N4 | 1.78 | 0.79 |
| 31:BA:2712:U:H1' | 31:BA:2712(A):A:C8 | 2.17 | 0.79 |
| 50:BY:47:LYS:HB3 | 50:BY:47:LYS:HZ3 | 1.46 | 0.79 |
| 1:CA:17:U:H2' | 1:CA:18:C:H6 | 1.45 | 0.79 |
| 1:CA:1226:C:C4 | 13:CM:104:ARG:HB2 | 2.17 | 0.79 |
| 13:CM:91:ARG:HB2 | 13:CM:98:VAL:HG21 | 1.64 | 0.79 |
| 31:DA:154:G:H1 | 31:DA:172:C:N4 | 1.80 | 0.79 |
| 31:DA:2476:A:C2 | 31:DA:2477:C:H5" | 2.18 | 0.79 |
| 50:DY:38:ILE:HG22 | 50:DY:39:VAL:N | 1.97 | 0.79 |
| 1:AA:737:A:H2' | 1:AA:738:C:H6 | 1.46 | 0.78 |
| 1:AA:1435:G:H2' | 1:AA:1436:U:C6 | 2.17 | 0.78 |
| 3:AC:52:LEU:H | 3:AC:52:LEU:HD23 | 1.46 | 0.78 |
| 8:AH:20:TYR:HD1 | 8:AH:65:TYR:CD2 | 2.00 | 0.78 |
| 27:B5:46:CYS:SG | 27:B5:47:PRO:CD | 2.71 | 0.78 |
| 28:B6:15:GLU:CD | 28:B6:18:ARG:NE | 2.40 | 0.78 |
| 31:BA:71:A:H3' | 31:BA:71:A:OP2 | 1.83 | 0.78 |
| 31:BA:747:U:O2 | 31:BA:2014:A:H1' | 1.82 | 0.78 |
| 41:BP:84:ASN:OD1 | 41:BP:117:GLU:HB2 | 1.82 | 0.78 |
| 51:BZ:149:SER:HB2 | 51:BZ:172:ALA:O | 1.83 | 0.78 |
| 28:D6:46:HIS:CB | 28:D6:47:THR:N | 2.46 | 0.78 |
| 31:DA:747:U:O2 | 31:DA:2014:A:H1' | 1.81 | 0.78 |
| 31:DA:1494:A:C4' | 31:DA:1495:A:OP1 | 2.31 | 0.78 |
| 33:DD:25:THR:O | 33:DD:25:THR:HG23 | 1.83 | 0.78 |
| 33:DD:186:HIS:CD2 | 33:DD:188:GLU:H | 2.01 | 0.78 |
| 42:DQ:81:VAL:O | 42:DQ:82:ARG:CG | 2.21 | 0.78 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:1250:A:H61 | 1:AA:1354:C:H1' | 1.48 | 0.78 |
| 2:AB:67:THR:HG22 | 2:AB:90:MET:HE1 | 1.65 | 0.78 |
| 38:BI:77:LEU:HD21 | 38:BI:101:LEU:HD22 | 1.63 | 0.78 |
| 47:BV:90:PRO:HG2 | 47:BV:91:TYR:H | 1.47 | 0.78 |
| 1:CA:1321:C:C5' | 1:CA:1322:C:H5" | 2.13 | 0.78 |
| 20:CT:56:MET:HG3 | 20:CT:88:VAL:HG21 | 1.63 | 0.78 |
| 31:DA:833:U:H2' | 31:DA:834:C:H6 | 1.48 | 0.78 |
| 31:DA:1430:C:H2' | 31:DA:1431:U:H6 | 1.46 | 0.78 |
| 1:AA:191:G:C4 | 20:AT:105:SER:HB3 | 2.19 | 0.78 |
| 1:AA:1152:A:H5" | 10:AJ:13:HIS:CD2 | 2.17 | 0.78 |
| 20:AT:50:GLU:CB | 20:AT:100:ILE:HG12 | 2.13 | 0.78 |
| 20:AT:56:MET:HG3 | 20:AT:88:VAL:HG21 | 1.63 | 0.78 |
| 31:BA:84:A:N6 | 31:BA:102:G:H1' | 1.97 | 0.78 |
| 31:BA:1771:C:O2' | 31:BA:1786:A:H8 | 1.65 | 0.78 |
| 31:BA:2475:C:H42 | 31:BA:2529:G:H22 | 1.28 | 0.78 |
| 34:BE:51:PHE:O | 34:BE:74:PRO:HB3 | 1.83 | 0.78 |
| 44:BS:24:LEU:HB3 | 44:BS:85:VAL:HG12 | 1.65 | 0.78 |
| 33:DD:27:THR:O | 33:DD:29:PRO:HD2 | 1.83 | 0.78 |
| 37:DH:85:LYS:HZ3 | 37:DH:145:ALA:HA | 1.45 | 0.78 |
| 47:DV:72:VAL:CA | 47:DV:88:ARG:HH22 | 1.92 | 0.78 |
| 50:DY:35:TYR:CE2 | 50:DY:69:ALA:HB3 | 2.18 | 0.78 |
| 1:AA:80:G:H1 | 1:AA:89:C:N4 | 1.81 | 0.78 |
| 5:AE:101:ILE:HD11 | 5:AE:119:LEU:HA | 1.66 | 0.78 |
| 8:AH:102:ARG:HE | 8:AH:102:ARG:N | 1.81 | 0.78 |
| 13:AM:68:GLY:O | 13:AM:69:GLU:HB2 | 1.84 | 0.78 |
| 19:AS:42:PRO:O | 19:AS:43:GLU:HB3 | 1.82 | 0.78 |
| 24:B2:14:ARG:NH1 | 24:B2:57:ILE:H | 1.80 | 0.78 |
| 27:B5:50:GLY:O | 27:B5:51:TYR:HD1 | 1.67 | 0.78 |
| 31:BA:102:G:HO2' | 31:BA:103:A:P | 2.07 | 0.78 |
| 48:BW:4:LYS:HB2 | 48:BW:106:ILE:HG22 | 1.65 | 0.78 |
| 1:CA:1289:A:H3' | 1:CA:1290:G:H8 | 1.49 | 0.78 |
| 28:D6:9:LEU:HD22 | 28:D6:10:LEU:H | 1.44 | 0.78 |
| 29:D7:9:ARG:NH1 | 31:DA:1310:G:OP2 | 2.17 | 0.78 |
| 31:DA:2712:U:H1' | 31:DA:2712(A):A:C8 | 2.18 | 0.78 |
| 40:DO:104:ARG:CZ | 45:DT:33:LYS:HD2 | 2.14 | 0.78 |
| 42:DQ:81:VAL:C | 42:DQ:82:ARG:HG2 | 2.08 | 0.78 |
| 9:AI:40:LEU:HD11 | 9:AI:70:LYS:HG2 | 1.66 | 0.78 |
| 31:BA:833:U:H2' | 31:BA:834:C:H6 | 1.48 | 0.78 |
| 37:BH:85:LYS:HD2 | 37:BH:141:VAL:HG13 | 1.64 | 0.78 |
| 1:CA:1169:A:H2' | 1:CA:1170:A:H8 | 1.45 | 0.78 |
| 9:CI:40:LEU:HD11 | 9:CI:70:LYS:HG2 | 1.66 | 0.78 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 30:D8:59:LYS:CD | 41:DP:50:ARG:HB3 | 2.13 | 0.78 |
| 32:DB:20:C:H2' | 32:DB:21:G:C5' | 2.13 | 0.78 |
| 44:DS:14:VAL:HG12 | 44:DS:15:ARG:H | 1.47 | 0.78 |
| 50:DY:45:VAL:HG13 | 50:DY:62:GLU:CB | 2.12 | 0.78 |
| 5:AE:50:GLU:HG3 | 5:AE:52:PRO:HD2 | 1.66 | 0.78 |
| 26:B4:12:ALA:O | 36:BG:101:ILE:HD11 | 1.83 | 0.78 |
| 30:B8:32:LEU:HD23 | 30:B8:35:GLN:N | 1.98 | 0.78 |
| 31:BA:1887:C:H2' | 31:BA:1888:G:H5' | 1.63 | 0.78 |
| 1:CA:1442(B):A:N1 | 45:DT:118:ARG:NH2 | 2.32 | 0.78 |
| 2:CB:154:LEU:HD23 | 2:CB:154:LEU:H | 1.49 | 0.78 |
| 31:DA:1681:G:H8 | 31:DA:1681:G:OP2 | 1.67 | 0.78 |
| 31:DA:1876:A:H2' | 31:DA:1877:A:C8 | 2.19 | 0.78 |
| 47:DV:64:HIS:HD2 | 47:DV:94:LEU:HD21 | 1.48 | 0.78 |
| 50:DY:39:VAL:HG12 | 50:DY:40:GLU:N | 1.97 | 0.78 |
| 1:AA:859:A:H2' | 1:AA:860:A:O4' | 1.84 | 0.78 |
| 4:AD:62:GLN:HA | 4:AD:62:GLN:HE21 | 1.49 | 0.78 |
| 31:BA:2756:U:H4' | 31:BA:2757:A:OP1 | 1.84 | 0.78 |
| 1:CA:1291:G:H4' | 9:CI:38:GLN:O | 1.83 | 0.78 |
| 8:CH:102:ARG:HE | 8:CH:102:ARG:N | 1.80 | 0.78 |
| 31:DA:90:U:H1' | 31:DA:92:A:H5" | 1.65 | 0.78 |
| 31:DA:1021:A:H62 | 31:DA:1141:U:H3 | 1.31 | 0.78 |
| 41:DP:124:LYS:HG2 | 41:DP:143:GLY:HA3 | 1.64 | 0.78 |
| 44:DS:101:LEU:HD13 | 44:DS:102:ALA:O | 1.83 | 0.78 |
| 5:AE:72:GLN:O | 5:AE:73:ASN:HB2 | 1.82 | 0.78 |
| 1:CA:200:G:H1 | 1:CA:217:C:H42 | 1.28 | 0.78 |
| 1:CA:1412:C:H2' | 1:CA:1413:A:H8 | 1.47 | 0.78 |
| 2:CB:163:PHE:HA | 2:CB:185:ILE:HG12 | 1.64 | 0.78 |
| 31:DA:2092:U:H4' | 31:DA:2093:G:O5' | 1.81 | 0.78 |
| 31:DA:2475:C:H42 | 31:DA:2529:G:H22 | 1.29 | 0.78 |
| 39:DN:9:VAL:HG11 | 39:DN:39:ARG:HH22 | 1.48 | 0.78 |
| 42:DQ:20:ALA:HB2 | 42:DQ:99:PRO:HD2 | 1.66 | 0.78 |
| 23:B1:62:VAL:HG22 | 23:B1:63:ALA:H | 1.49 | 0.78 |
| 31:BA:527:C:OP2 | 31:BA:2779:U:H5 | 1.66 | 0.78 |
| 32:BB:116:G:C5' | 32:BB:116:G:H8 | 1.97 | 0.78 |
| 31:DA:1884:A:C2' | 31:DA:1885:A:H5' | 2.14 | 0.78 |
| 2:AB:84:GLU:HB3 | 2:AB:219:VAL:HG21 | 1.65 | 0.78 |
| 16:AP:74:LEU:O | 16:AP:79:VAL:HG23 | 1.83 | 0.78 |
| 24:B2:55:ARG:HH22 | 49:BX:3:THR:HG22 | 1.48 | 0.78 |
| 31:BA:1022:G:O2' | 31:BA:1023:U:OP2 | 2.02 | 0.78 |
| 39:BN:9:VAL:HG11 | 39:BN:39:ARG:HH22 | 1.49 | 0.78 |
| 47:BV:19:LYS:HG2 | 47:BV:96:ILE:HG22 | 1.65 | 0.78 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:826:C:H2' | 1:CA:827:U:C6 | 2.19 | 0.78 |
| 1:CA:1199:U:H4' | 10:CJ:54:PHE:CZ | 2.18 | 0.78 |
| 2:CB:97:TRP:CH2 | 2:CB:173:ALA:HA | 2.19 | 0.78 |
| 4:CD:119:GLN:HG2 | 4:CD:123:HIS:CD2 | 2.18 | 0.78 |
| 24:D2:45:SER:O | 24:D2:47:ASN:ND2 | 2.17 | 0.78 |
| 31:DA:2030:A:H4' | 31:DA:2031:A:OP1 | 1.83 | 0.78 |
| 47:DV:70:ILE:HG13 | 47:DV:90:PRO:HB3 | 1.66 | 0.78 |
| 48:DW:4:LYS:HB2 | 48:DW:106:ILE:HG22 | 1.64 | 0.78 |
| 23:B1:10:LYS:HB2 | 23:B1:14:VAL:N | 1.99 | 0.77 |
| 49:BX:36:LYS:NZ | 49:BX:38:GLU:O | 2.13 | 0.77 |
| 16:CP:28:ARG:HH11 | 16:CP:28:ARG:CG | 1.97 | 0.77 |
| 33:DD:183:ARG:HG2 | 33:DD:183:ARG:NH1 | 1.98 | 0.77 |
| 1:AA:437:U:OP1 | 4:AD:155:LEU:HD22 | 1.85 | 0.77 |
| 31:BA:90:U:H1' | 31:BA:92:A:H5" | 1.65 | 0.77 |
| 31:DA:1887:C:H2' | 31:DA:1888:G:H5' | 1.66 | 0.77 |
| 31:DA:2223:G:C2' | 31:DA:2224:G:H5' | 2.14 | 0.77 |
| 39:DN:42:TRP:HA | 39:DN:48:MET:HE1 | 1.64 | 0.77 |
| 41:BP:95:VAL:HG22 | 41:BP:125:VAL:HB | 1.65 | 0.77 |
| 43:BR:53:HIS:HD2 | 43:BR:94:TYR:OH | 1.67 | 0.77 |
| 50:BY:35:TYR:CD2 | 50:BY:69:ALA:HB3 | 2.20 | 0.77 |
| 1:CA:266:G:H5" | 1:CA:268:C:H41 | 1.48 | 0.77 |
| 28:D6:10:LEU:H | 28:D6:10:LEU:CD2 | 1.96 | 0.77 |
| 31:DA:674:G:O2' | 35:DF:74:ARG:HG3 | 1.84 | 0.77 |
| 31:DA:2712(A):A:H5" | 31:DA:2713:A:OP2 | 1.83 | 0.77 |
| 31:DA:2790:A:H2' | 31:DA:2791:C:H5" | 1.66 | 0.77 |
| 32:DB:8:U:H5' | 32:DB:8:U:H6 | 1.48 | 0.77 |
| 33:DD:35:LYS:NZ | 33:DD:65:ILE:HA | 1.99 | 0.77 |
| 39:DN:128:HIS:CD2 | 39:DN:131:GLN:HB2 | 2.19 | 0.77 |
| 1:AA:403:C:O2' | 1:AA:404:U:H5' | 1.82 | 0.77 |
| 2:AB:163:PHE:HA | 2:AB:185:ILE:HG12 | 1.65 | 0.77 |
| 30:B8:35:GLN:NE2 | 30:B8:36:LYS:HZ3 | 1.81 | 0.77 |
| 31:BA:1300:U:O2' | 31:BA:1626:G:C2 | 2.38 | 0.77 |
| 31:BA:2395:C:H5' | 31:BA:2395:C:H6 | 1.50 | 0.77 |
| 41:BP:17:LYS:HG3 | 41:BP:19:VAL:HG23 | 1.66 | 0.77 |
| 1:CA:664:G:N2 | 1:CA:741:G:H1 | 1.82 | 0.77 |
| 26:D4:12:ALA:O | 36:DG:101:ILE:HD11 | 1.85 | 0.77 |
| 31:DA:1614:A:H61 | 48:DW:88:ARG:H | 1.33 | 0.77 |
| 41:DP:84:ASN:OD1 | 41:DP:117:GLU:HB2 | 1.85 | 0.77 |
| 1:AA:1289:A:H3' | 1:AA:1290:G:H8 | 1.50 | 0.77 |
| 13:AM:81:LEU:HB3 | 13:AM:89:GLY:HA2 | 1.64 | 0.77 |
| 23:B1:46:LEU:HD12 | 23:B1:46:LEU:N | 1.99 | 0.77 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1497:U:H5' | 31:BA:1498:C:C5 | 2.18 | 0.77 |
| 41:BP:140:ALA:CB | 25:D3:38:GLU:HG2 | 2.10 | 0.77 |
| 30:D8:32:LEU:C | 30:D8:34:TRP:H | 1.88 | 0.77 |
| 31:DA:1502:C:H2' | 31:DA:1502:C:O2 | 1.84 | 0.77 |
| 31:DA:2660:A:H5' | 31:DA:2661:G:N2 | 2.00 | 0.77 |
| 39:DN:18:ALA:HB1 | 39:DN:21:LYS:HG3 | 1.64 | 0.77 |
| 42:DQ:140:ALA:HB1 | 51:DZ:99:TYR:HB2 | 1.67 | 0.77 |
| 50:DY:35:TYR:CD2 | 50:DY:69:ALA:HB3 | 2.20 | 0.77 |
| 1:AA:192:U:H2' | 1:AA:193:C:H6 | 1.48 | 0.77 |
| 1:AA:365:U:H5" | 1:AA:366:C:OP1 | 1.85 | 0.77 |
| 32:BB:8:U:H5' | 32:BB:8:U:H6 | 1.48 | 0.77 |
| 33:BD:27:THR:O | 33:BD:29:PRO:HD2 | 1.83 | 0.77 |
| 40:BO:23:ARG:HG2 | 40:BO:23:ARG:NH1 | 1.93 | 0.77 |
| 1:CA:472:A:H4' | 16:CP:82:GLN:HE22 | 1.50 | 0.77 |
| 6:CF:62:TRP:CH2 | 6:CF:64:GLN:HB2 | 2.19 | 0.77 |
| 31:DA:966:G:H2' | 31:DA:967:C:C6 | 2.19 | 0.77 |
| 41:DP:62:LEU:H | 41:DP:62:LEU:HD13 | 1.47 | 0.77 |
| 1:AA:626:U:H2' | 1:AA:627:G:C8 | 2.20 | 0.77 |
| 31:BA:819:A:C4 | 31:BA:1189:A:C2 | 2.72 | 0.77 |
| 31:BA:1884:A:C2' | 31:BA:1885:A:H5' | 2.15 | 0.77 |
| 31:BA:2660:A:H3' | 31:BA:2660:A:N3 | 1.99 | 0.77 |
| 33:BD:27:THR:CG2 | 33:BD:28:GLU:N | 2.48 | 0.77 |
| 39:BN:128:HIS:CD2 | 39:BN:131:GLN:HB2 | 2.19 | 0.77 |
| 50:BY:8:LYS:HE2 | 50:BY:72:VAL:HG23 | 1.67 | 0.77 |
| 1:CA:437:U:OP1 | 4:CD:155:LEU:HD22 | 1.84 | 0.77 |
| 1:CA:878:G:H5' | 8:CH:89:PRO:HG2 | 1.67 | 0.77 |
| 28:D6:20:ASN:CG | 28:D6:21:TYR:H | 1.92 | 0.77 |
| 31:DA:2392:A:H2 | 31:DA:2424:C:H42 | 1.29 | 0.77 |
| 34:DE:152:LYS:HG2 | 39:DN:78:TYR:CD2 | 2.20 | 0.77 |
| 45:DT:29:ARG:CB | 45:DT:85:LYS:HA | 2.14 | 0.77 |
| 1:AA:1070:U:H2' | 1:AA:1071:C:C6 | 2.20 | 0.77 |
| 11:AK:21:ILE:HB | 11:AK:84:VAL:HA | 1.65 | 0.77 |
| 31:BA:1019:U:H3 | 31:BA:1142(A):A:N6 | 1.83 | 0.77 |
| 31:BA:1108:U:H2' | 31:BA:1109:C:H5' | 1.67 | 0.77 |
| 37:BH:41:MET:HE2 | 37:BH:55:PRO:HD3 | 1.66 | 0.77 |
| 39:BN:78:TYR:HD1 | 39:BN:79:PRO:CD | 1.98 | 0.77 |
| 43:BR:4:LEU:O | 43:BR:4:LEU:CD1 | 2.32 | 0.77 |
| 45:BT:106:SER:O | 45:BT:107:ASP:CG | 2.27 | 0.77 |
| 31:DA:669:G:HO2' | 31:DA:669:G:H8 | 1.31 | 0.77 |
| 1:AA:1412:C:H2' | 1:AA:1413:A:H8 | 1.49 | 0.77 |
| 24:B2:52:ASP:O | 24:B2:56:GLN:NE2 | 2.18 | 0.77 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 28:B6:13:CYS:HB3 | 28:B6:49:HIS:HB3 | 1.65 | 0.77 |
| 31:BA:1021:A:H62 | 31:BA:1141:U:H3 | 1.32 | 0.77 |
| 31:BA:1614:A:H61 | 48:BW:88:ARG:H | 1.33 | 0.77 |
| 33:BD:127:VAL:HA | 33:BD:193:VAL:HG22 | 1.67 | 0.77 |
| 35:BF:66:PRO:O | 35:BF:67:GLN:HB3 | 1.84 | 0.77 |
| 41:BP:105:LEU:O | 41:BP:106:LEU:HB2 | 1.85 | 0.77 |
| 8:CH:10:LEU:N | 8:CH:10:LEU:HD23 | 1.99 | 0.77 |
| 31:DA:1527:G:H5" | 31:DA:1528:A:OP1 | 1.85 | 0.77 |
| 38:DI:77:LEU:HD21 | 38:DI:101:LEU:HD22 | 1.67 | 0.77 |
| 1:AA:826:C:H2' | 1:AA:827:U:C6 | 2.20 | 0.77 |
| 3:AC:180:ALA:HB1 | 3:AC:182:ILE:HG13 | 1.65 | 0.77 |
| 4:AD:61:LYS:HB2 | 4:AD:203:VAL:HG22 | 1.66 | 0.77 |
| 6:AF:62:TRP:CH2 | 6:AF:64:GLN:HB2 | 2.19 | 0.77 |
| 30:B8:59:LYS:HB2 | 30:B8:59:LYS:HZ1 | 1.50 | 0.77 |
| 31:BA:102:G:H8 | 31:BA:102:G:C5' | 1.96 | 0.77 |
| 31:BA:587:C:H4' | 31:BA:588:U:OP2 | 1.84 | 0.77 |
| 39:BN:45:ASN:H | 39:BN:45:ASN:ND2 | 1.81 | 0.77 |
| 41:BP:62:LEU:HD22 | 41:BP:62:LEU:H | 1.48 | 0.77 |
| 49:BX:24:GLY:HA3 | 49:BX:80:ILE:HG13 | 1.66 | 0.77 |
| 1:CA:191:G:C4 | 20:CT:105:SER:HB3 | 2.19 | 0.77 |
| 20:CT:50:GLU:CB | 20:CT:100:ILE:HG12 | 2.15 | 0.77 |
| 31:DA:1899:G:H21 | 31:DA:1902:C:H5 | 1.29 | 0.77 |
| 35:DF:53:THR:HG22 | 35:DF:56:GLU:OE1 | 1.85 | 0.77 |
| 2:AB:87:ARG:HE | 2:AB:233:SER:HB3 | 1.48 | 0.76 |
| 2:AB:97:TRP:CH2 | 2:AB:173:ALA:HA | 2.19 | 0.76 |
| 45:BT:50:ILE:HD11 | 45:BT:102:ILE:CD1 | 2.15 | 0.76 |
| 45:BT:54:ARG:HA | 45:BT:59:THR:HB | 1.67 | 0.76 |
| 23:D1:10:LYS:HB2 | 23:D1:14:VAL:N | 2.00 | 0.76 |
| 23:D1:47:GLN:HB2 | 31:DA:397:G:H5" | 1.67 | 0.76 |
| 45:DT:30:VAL:HG21 | 45:DT:83:ILE:CG1 | 2.16 | 0.76 |
| 1:AA:166:G:H2' | 1:AA:167:G:H8 | 1.49 | 0.76 |
| 10:AJ:49:VAL:HG13 | 14:AN:41:ARG:HB2 | 1.65 | 0.76 |
| 23:B1:87:PRO:HD2 | 23:B1:88:LYS:N | 1.99 | 0.76 |
| 33:BD:2:ALA:O | 33:BD:3:VAL:HB | 1.83 | 0.76 |
| 37:BH:70:THR:HG22 | 37:BH:74:ASN:HD21 | 1.51 | 0.76 |
| 42:BQ:27:VAL:HG13 | 42:BQ:105:GLU:OE1 | 1.85 | 0.76 |
| 1:CA:166:G:H2' | 1:CA:167:G:H8 | 1.49 | 0.76 |
| 16:CP:28:ARG:HG2 | 16:CP:28:ARG:NH1 | 1.95 | 0.76 |
| 24:D2:52:ASP:O | 24:D2:56:GLN:NE2 | 2.18 | 0.76 |
| 31:DA:1696:G:H2' | 31:DA:1697:G:C5' | 2.15 | 0.76 |
| 35:DF:132:VAL:HG22 | 35:DF:133:ASN:H | 1.50 | 0.76 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 38:DI:6:LEU:HD12 | 38:DI:34:GLY:O | 1.84 | 0.76 |
| 39:DN:65:LYS:O | 39:DN:69:GLN:HB2 | 1.85 | 0.76 |
| 1:AA:475:G:H2' | 1:AA:476:G:C8 | 2.18 | 0.76 |
| 18:AR:31:LEU:HD23 | 18:AR:31:LEU:H | 1.50 | 0.76 |
| 31:BA:2790:A:H2' | 31:BA:2791:C:H5" | 1.67 | 0.76 |
| 42:BQ:140:ALA:HB1 | 51:BZ:99:TYR:HB2 | 1.67 | 0.76 |
| 1:CA:1152:A:H5" | 10:CJ:13:HIS:CD2 | 2.20 | 0.76 |
| 24:D2:14:ARG:NH1 | 24:D2:57:ILE:H | 1.83 | 0.76 |
| 31:DA:542:C:N3 | 31:DA:543:C:N4 | 2.33 | 0.76 |
| 31:DA:1108:U:H2' | 31:DA:1109:C:H5' | 1.68 | 0.76 |
| 32:DB:28:C:C2 | 32:DB:29:A:C8 | 2.73 | 0.76 |
| 39:DN:78:TYR:HD1 | 39:DN:79:PRO:CD | 1.98 | 0.76 |
| 51:DZ:166:SER:HB2 | 51:DZ:168:GLU:N | 1.99 | 0.76 |
| 1:AA:247:G:OP2 | 17:AQ:100:LYS:HG2 | 1.86 | 0.76 |
| 31:BA:61:G:H1 | 31:BA:94:C:H42 | 1.33 | 0.76 |
| 31:BA:2476:A:C2 | 31:BA:2477:C:H5" | 2.19 | 0.76 |
| 34:BE:24:THR:HG23 | 34:BE:184:VAL:HG23 | 1.67 | 0.76 |
| 34:BE:38:THR:HG22 | 34:BE:40:GLU:H | 1.50 | 0.76 |
| 35:BF:53:THR:HG22 | 35:BF:56:GLU:OE1 | 1.85 | 0.76 |
| 37:BH:88:LEU:O | 37:BH:89:ILE:HG23 | 1.85 | 0.76 |
| 39:BN:42:TRP:HA | 39:BN:48:MET:HE1 | 1.65 | 0.76 |
| 42:BQ:35:VAL:HG13 | 42:BQ:130:LYS:HB3 | 1.66 | 0.76 |
| 44:BS:83:LYS:HE3 | 44:BS:105:ALA:HB2 | 1.66 | 0.76 |
| 51:BZ:109:ALA:HB1 | 51:BZ:145:GLU:OE2 | 1.85 | 0.76 |
| 1:CA:1250:A:H61 | 1:CA:1354:C:H1' | 1.50 | 0.76 |
| 6:CF:30:LEU:HB2 | 6:CF:35:ALA:HB3 | 1.67 | 0.76 |
| 31:DA:1529:G:N2 | 31:DA:1530:C:H5" | 1.99 | 0.76 |
| 32:DB:86:G:H2' | 32:DB:87:G:C8 | 2.19 | 0.76 |
| 42:DQ:81:VAL:O | 42:DQ:82:ARG:NH1 | 2.17 | 0.76 |
| 46:DU:93:LYS:HD3 | 46:DU:93:LYS:H | 1.50 | 0.76 |
| 23:B1:12:PRO:HD2 | 23:B1:62:VAL:HG23 | 1.68 | 0.76 |
| 34:BE:136:ARG:HH11 | 34:BE:136:ARG:HG2 | 1.49 | 0.76 |
| 39:BN:40:PRO:HB3 | 46:BU:68:ALA:HB2 | 1.67 | 0.76 |
| 1:CA:341:C:O2' | 1:CA:342:C:H5' | 1.85 | 0.76 |
| 31:DA:1179:C:H3' | 31:DA:1180:C:H5" | 1.67 | 0.76 |
| 31:DA:2036:C:H5' | 31:DA:2036:C:H6 | 1.51 | 0.76 |
| 34:DE:14:ILE:HG12 | 34:DE:21:VAL:CG2 | 2.15 | 0.76 |
| 39:DN:17:ASP:OD2 | 39:DN:19:GLU:HB3 | 1.86 | 0.76 |
| 39:DN:126:PRO:O | 39:DN:127:ASP:HB2 | 1.84 | 0.76 |
| 41:DP:71:VAL:HG12 | 41:DP:72:PRO:HD3 | 1.66 | 0.76 |
| 44:DS:14:VAL:CG1 | 44:DS:15:ARG:N | 2.48 | 0.76 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 51:DZ:73:GLN:HG2 | 51:DZ:87:ASP:OD1 | 1.84 | 0.76 |
| 1:AA:266:G:H5" | 1:AA:268:C:H41 | 1.51 | 0.76 |
| 24:B2:30:ARG:HD2 | 24:B2:30:ARG:H | 1.49 | 0.76 |
| 31:BA:674:G:O2' | 35:BF:74:ARG:HG3 | 1.85 | 0.76 |
| 32:BB:32:C:C2 | 32:BB:51:G:N2 | 2.53 | 0.76 |
| 33:BD:35:LYS:HZ1 | 33:BD:65:ILE:HA | 1.48 | 0.76 |
| 33:BD:75:ILE:HG21 | 33:BD:99:ASP:HB2 | 1.68 | 0.76 |
| 47:BV:5:VAL:CG2 | 47:BV:36:PRO:HB2 | 2.14 | 0.76 |
| 2:CB:87:ARG:HE | 2:CB:233:SER:HB3 | 1.48 | 0.76 |
| 25:D3:54:VAL:HG12 | 25:D3:55:ARG:N | 2.01 | 0.76 |
| 31:DA:529:A:H62 | 31:DA:2041:U:H3 | 1.33 | 0.76 |
| 31:DA:659:C:H5' | 31:DA:659:C:C6 | 2.20 | 0.76 |
| 31:DA:2523:G:H2' | 31:DA:2524:G:C5' | 2.14 | 0.76 |
| 47:DV:64:HIS:CG | 47:DV:64:HIS:O | 2.38 | 0.76 |
| 47:DV:66:ARG:HG2 | 47:DV:94:LEU:HG | 1.68 | 0.76 |
| 38:BI:8:PRO:HA | 38:BI:13:GLY:O | 1.86 | 0.76 |
| 47:BV:13:ARG:HH11 | 47:BV:13:ARG:CG | 1.97 | 0.76 |
| 30:D8:30:ARG:O | 30:D8:31:HIS:C | 2.28 | 0.76 |
| 32:DB:74:U:H2' | 32:DB:75:G:H5" | 1.65 | 0.76 |
| 35:DF:3:GLU:O | 35:DF:19:GLU:HA | 1.86 | 0.76 |
| 24:B2:46:GLN:NE2 | 24:B2:47:ASN:N | 2.33 | 0.76 |
| 31:BA:71:A:H5' | 31:BA:71:A:C8 | 2.20 | 0.76 |
| 31:BA:2660:A:H5' | 31:BA:2661:G:H21 | 1.51 | 0.76 |
| 35:BF:103:LYS:HA | 35:BF:106:ARG:HG3 | 1.66 | 0.76 |
| 47:BV:69:LYS:HB2 | 47:BV:93:GLU:CD | 2.11 | 0.76 |
| 4:CD:61:LYS:HB2 | 4:CD:203:VAL:HG22 | 1.68 | 0.76 |
| 28:D6:15:GLU:CD | 28:D6:18:ARG:NE | 2.43 | 0.76 |
| 31:DA:1173:G:H3' | 31:DA:1174:A:C5' | 2.15 | 0.76 |
| 34:DE:14:ILE:HG12 | 34:DE:21:VAL:HG23 | 1.67 | 0.76 |
| 35:DF:194:MET:HE3 | 35:DF:199:TRP:HD1 | 1.50 | 0.76 |
| 37:DH:144:VAL:O | 37:DH:148:ILE:HG12 | 1.84 | 0.76 |
| 50:DY:28:LYS:HD2 | 50:DY:37:VAL:HG12 | 1.68 | 0.76 |
| 31:BA:1173:G:H3' | 31:BA:1174:A:C5' | 2.15 | 0.76 |
| 32:BB:74:U:H2' | 32:BB:75:G:H5" | 1.68 | 0.76 |
| 50:BY:28:LYS:HD2 | 50:BY:37:VAL:HG12 | 1.67 | 0.76 |
| 1:CA:327:A:H3' | 1:CA:328:C:H5" | 1.68 | 0.76 |
| 5:CE:26:PHE:O | 5:CE:27:ARG:HB2 | 1.85 | 0.76 |
| 6:CF:24:GLU:HG3 | 6:CF:28:ARG:NH1 | 2.00 | 0.76 |
| 13:CM:49:THR:HG22 | 13:CM:51:ALA:H | 1.50 | 0.76 |
| 31:DA:141:A:C8 | 31:DA:1408:C:O2' | 2.37 | 0.76 |
| 31:DA:542:C:C4 | 31:DA:543:C:N4 | 2.54 | 0.76 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2790:A:H2' | 31:DA:2791:C:C5' | 2.15 | 0.76 |
| 31:DA:2875:C:H4' | 45:DT:5:ALA:HB2 | 1.66 | 0.76 |
| 38:DI:8:PRO:HA | 38:DI:13:GLY:O | 1.84 | 0.76 |
| 38:DI:53:ALA:HB1 | 38:DI:57:ARG:N | 1.99 | 0.76 |
| 40:DO:35:VAL:HA | 40:DO:62:VAL:HG12 | 1.68 | 0.76 |
| 45:DT:88:ILE:HG22 | 45:DT:89:VAL:HG23 | 1.67 | 0.76 |
| 2:AB:180:LEU:O | 2:AB:181:PHE:HB2 | 1.85 | 0.76 |
| 27:B5:16:ARG:NH1 | 27:B5:17:ASP:OD1 | 2.19 | 0.76 |
| 30:B8:32:LEU:HB3 | 30:B8:34:TRP:N | 2.00 | 0.76 |
| 31:BA:142:A:C8 | 31:BA:1408:C:H1' | 2.21 | 0.76 |
| 31:BA:752:A:H4' | 31:BA:753:C:O5' | 1.84 | 0.76 |
| 31:BA:814:C:H41 | 41:BP:27:HIS:CD2 | 2.03 | 0.76 |
| 31:BA:870:A:C5' | 42:BQ:7:MET:HB2 | 2.11 | 0.76 |
| 31:BA:1798:U:C5' | 33:BD:259:THR:HG22 | 2.16 | 0.76 |
| 31:BA:2475:C:H6 | 31:BA:2475:C:H5" | 1.51 | 0.76 |
| 35:BF:3:GLU:O | 35:BF:19:GLU:HA | 1.85 | 0.76 |
| 36:BG:60:LEU:O | 36:BG:64:THR:HG22 | 1.86 | 0.76 |
| 1:CA:475:G:H2' | 1:CA:476:G:C8 | 2.20 | 0.76 |
| 27:D5:50:GLY:HA3 | 27:D5:56:LYS:HG2 | 1.68 | 0.76 |
| 27:D5:54:GLY:O | 27:D5:56:LYS:NZ | 2.19 | 0.76 |
| 31:DA:631:A:OP1 | 41:DP:64:LYS:HE3 | 1.85 | 0.76 |
| 31:DA:1670:C:O2 | 34:DE:129:HIS:HE1 | 1.69 | 0.76 |
| 32:DB:44:G:H5" | 32:DB:45:A:OP1 | 1.86 | 0.76 |
| 33:DD:70:TRP:HZ3 | 33:DD:146:GLU:OE2 | 1.69 | 0.76 |
| 44:DS:24:LEU:HB3 | 44:DS:85:VAL:CG1 | 2.14 | 0.76 |
| 47:DV:51:VAL:HG12 | 47:DV:52:VAL:H | 1.50 | 0.76 |
| 47:DV:83:ARG:HH11 | 47:DV:83:ARG:CG | 1.99 | 0.76 |
| 51:DZ:109:ALA:HB1 | 51:DZ:145:GLU:OE2 | 1.85 | 0.76 |
| 1:AA:377:G:OP1 | 16:AP:3:LYS:HD2 | 1.86 | 0.75 |
| 1:AA:1117:G:H4' | 9:AI:104:ARG:NH2 | 2.01 | 0.75 |
| 28:B6:33:LYS:O | 28:B6:34:LEU:HB2 | 1.86 | 0.75 |
| 1:CA:1070:U:H2' | 1:CA:1071:C:C6 | 2.22 | 0.75 |
| 27:D5:16:ARG:HH11 | 27:D5:16:ARG:CG | 2.00 | 0.75 |
| 31:DA:598:G:H5' | 41:DP:15:ARG:HD2 | 1.67 | 0.75 |
| 31:DA:752:A:H4' | 31:DA:753:C:O5' | 1.85 | 0.75 |
| 31:DA:2654:A:OP1 | 31:DA:2654:A:H8 | 1.69 | 0.75 |
| 45:DT:54:ARG:HA | 45:DT:59:THR:HB | 1.66 | 0.75 |
| 49:DX:65:ARG:HA | 49:DX:65:ARG:NE | 2.02 | 0.75 |
| 1:AA:148:G:O2' | 1:AA:149:A:H5' | 1.87 | 0.75 |
| 31:BA:184:C:H2' | 31:BA:185:U:H6 | 1.49 | 0.75 |
| 31:BA:542:C:C4 | 31:BA:543:C:N4 | 2.54 | 0.75 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 38:BI:91:SER:HB3 | 38:BI:121:LYS:HB2 | 1.68 | 0.75 |
| 42:BQ:83:MET:HG2 | 42:BQ:83:MET:O | 1.86 | 0.75 |
| 1:CA:859:A:H2' | 1:CA:860:A:O4' | 1.86 | 0.75 |
| 31:DA:1021:A:H3' | 31:DA:1021:A:C8 | 2.20 | 0.75 |
| 36:DG:131:TYR:HB3 | 36:DG:159:VAL:HG13 | 1.68 | 0.75 |
| 39:DN:73:THR:O | 39:DN:75:TYR:N | 2.17 | 0.75 |
| 39:DN:83:LYS:HE3 | 39:DN:85:ILE:HD11 | 1.68 | 0.75 |
| 46:DU:92:ARG:HD3 | 46:DU:94:ASN:HB3 | 1.68 | 0.75 |
| 1:AA:499:A:H4' | 1:AA:500:G:OP1 | 1.86 | 0.75 |
| 31:BA:2685:G:H2' | 31:BA:2686:G:H5" | 1.67 | 0.75 |
| 33:BD:186:HIS:CD2 | 33:BD:188:GLU:H | 2.03 | 0.75 |
| 38:BI:2:LYS:HB2 | 38:BI:39:ALA:HB3 | 1.66 | 0.75 |
| 1:CA:1030(A):G:H1' | 1:CA:1031:G:H22 | 1.51 | 0.75 |
| 1:CA:1285:A:H1' | 1:CA:1286:A:OP2 | 1.86 | 0.75 |
| 3:CC:180:ALA:HB1 | 3:CC:182:ILE:HG13 | 1.66 | 0.75 |
| 31:DA:1798:U:C5' | 33:DD:259:THR:HG22 | 2.13 | 0.75 |
| 32:DB:32:C:C2 | 32:DB:51:G:N2 | 2.53 | 0.75 |
| 27:B5:54:GLY:O | 27:B5:56:LYS:NZ | 2.18 | 0.75 |
| 31:BA:943:U:OP2 | 41:BP:38:GLN:OE1 | 2.04 | 0.75 |
| 31:BA:1430:C:H2' | 31:BA:1431:U:H6 | 1.49 | 0.75 |
| 31:BA:2223:G:C2' | 31:BA:2224:G:H5' | 2.16 | 0.75 |
| 35:BF:46:ARG:HH11 | 35:BF:46:ARG:CG | 2.00 | 0.75 |
| 31:DA:2206:G:C2 | 31:DA:2207:G:H5' | 2.21 | 0.75 |
| 32:DB:116:G:C5' | 32:DB:116:G:H8 | 1.99 | 0.75 |
| 33:DD:2:ALA:O | 33:DD:3:VAL:HB | 1.83 | 0.75 |
| 34:DE:199:ARG:O | 34:DE:200:GLU:HB3 | 1.83 | 0.75 |
| 47:DV:69:LYS:HB2 | 47:DV:93:GLU:OE2 | 1.86 | 0.75 |
| 28:B6:14:THR:O | 28:B6:49:HIS:HA | 1.85 | 0.75 |
| 31:BA:997:G:OP1 | 46:BU:93:LYS:HD3 | 1.86 | 0.75 |
| 47:BV:19:LYS:HB3 | 47:BV:96:ILE:O | 1.87 | 0.75 |
| 49:BX:55:ASN:HB2 | 49:BX:78:LYS:HD2 | 1.68 | 0.75 |
| 1:CA:1117:G:H4' | 9:CI:104:ARG:NH2 | 2.00 | 0.75 |
| 10:CJ:26:ALA:HB1 | 10:CJ:29:ARG:HH21 | 1.51 | 0.75 |
| 12:CL:60:LEU:HD21 | 12:CL:66:VAL:HG22 | 1.68 | 0.75 |
| 30:D8:32:LEU:HB3 | 30:D8:34:TRP:HB3 | 1.68 | 0.75 |
| 33:DD:35:LYS:HZ1 | 33:DD:65:ILE:HA | 1.51 | 0.75 |
| 34:DE:132:HIS:CG | 34:DE:135:HIS:NE2 | 2.54 | 0.75 |
| 37:DH:85:LYS:HD3 | 37:DH:133:VAL:HB | 1.69 | 0.75 |
| 38:DI:61:ARG:O | 38:DI:133:HIS:HE1 | 1.69 | 0.75 |
| 38:DI:144:VAL:HG12 | 38:DI:145:VAL:H | 1.49 | 0.75 |
| 50:DY:8:LYS:NZ | 50:DY:74:PRO:HD3 | 2.01 | 0.75 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:38:G:C2 | 1:AA:397:A:C2 | 2.74 | 0.75 |
| 31:BA:1527:G:H5" | 31:BA:1528:A:OP1 | 1.86 | 0.75 |
| 34:BE:95:ILE:HD12 | 34:BE:95:ILE:H | 1.49 | 0.75 |
| 38:BI:138:ILE:H | 38:BI:138:ILE:HD12 | 1.52 | 0.75 |
| 42:BQ:81:VAL:O | 42:BQ:82:ARG:NH1 | 2.18 | 0.75 |
| 45:BT:13:ARG:NH2 | 45:BT:15:VAL:HG13 | 2.01 | 0.75 |
| 23:D1:23:LYS:HB2 | 23:D1:37:ILE:HG22 | 1.68 | 0.75 |
| 31:DA:1495:A:N3 | 31:DA:1496:A:C2 | 2.55 | 0.75 |
| 31:DA:1747(A):G:C2' | 31:DA:1748:G:H5' | 2.15 | 0.75 |
| 32:DB:82:G:C2' | 32:DB:83:G:H5' | 2.16 | 0.75 |
| 43:DR:116:LEU:O | 43:DR:117:VAL:HB | 1.86 | 0.75 |
| 45:DT:38:ASN:C | 45:DT:38:ASN:HD22 | 1.94 | 0.75 |
| 47:DV:83:ARG:O | 47:DV:84:LYS:HG3 | 1.86 | 0.75 |
| 1:AA:114:U:H2' | 1:AA:115:G:C8 | 2.21 | 0.75 |
| 10:AJ:63:PHE:HZ | 14:AN:45:ARG:HG3 | 1.50 | 0.75 |
| 23:B1:73:LEU:HD21 | 23:B1:94:LEU:HA | 1.67 | 0.75 |
| 31:BA:154:G:H1 | 31:BA:172:C:N4 | 1.83 | 0.75 |
| 31:BA:1821:A:C2' | 31:BA:1822:G:H5" | 2.14 | 0.75 |
| 31:BA:2206:G:C2 | 31:BA:2207:G:H5' | 2.21 | 0.75 |
| 1:CA:114:U:H2' | 1:CA:115:G:C8 | 2.21 | 0.75 |
| 2:CB:137:ARG:HH11 | 2:CB:137:ARG:HA | 1.51 | 0.75 |
| 23:D1:73:LEU:HD21 | 23:D1:94:LEU:HA | 1.68 | 0.75 |
| 38:DI:53:ALA:HB1 | 38:DI:56:LYS:HG3 | 1.69 | 0.75 |
| 41:DP:62:LEU:N | 41:DP:62:LEU:HD22 | 2.02 | 0.75 |
| 45:DT:30:VAL:HG12 | 45:DT:44:ASP:OD2 | 1.85 | 0.75 |
| 1:AA:390:C:O3' | 16:AP:28:ARG:NH2 | 2.20 | 0.75 |
| 1:AA:472:A:H4' | 16:AP:82:GLN:HE22 | 1.50 | 0.75 |
| 1:AA:735:C:O2' | 1:AA:736:C:H5' | 1.87 | 0.75 |
| 1:AA:1279:A:H5" | 1:AA:1280:A:OP1 | 1.86 | 0.75 |
| 28:B6:15:GLU:CG | 28:B6:18:ARG:NE | 2.47 | 0.75 |
| 30:B8:32:LEU:HD23 | 30:B8:35:GLN:H | 1.51 | 0.75 |
| 31:BA:1899:G:H21 | 31:BA:1902:C:H5 | 1.32 | 0.75 |
| 31:BA:2657:A:H5' | 31:BA:2658:C:OP2 | 1.86 | 0.75 |
| 33:BD:35:LYS:HZ3 | 33:BD:104:TYR:HD1 | 1.35 | 0.75 |
| 35:BF:32:LEU:HD11 | 35:BF:105:VAL:HG13 | 1.68 | 0.75 |
| 39:BN:17:ASP:OD2 | 39:BN:19:GLU:HB3 | 1.87 | 0.75 |
| 41:BP:107:LYS:C | 41:BP:109:GLY:H | 1.94 | 0.75 |
| 1:CA:735:C:H2' | 1:CA:736:C:C6 | 2.20 | 0.75 |
| 16:CP:43:LYS:HA | 16:CP:48:TRP:HB3 | 1.69 | 0.75 |
| 31:DA:143:G:H1' | 49:DX:38:GLU:HG3 | 1.67 | 0.75 |
| 47:DV:71:LEU:HD13 | 47:DV:72:VAL:HG23 | 1.68 | 0.75 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 12:AL:60:LEU:HD21 | 12:AL:66:VAL:HG22 | 1.69 | 0.75 |
| 13:AM:49:THR:HG22 | 13:AM:51:ALA:H | 1.52 | 0.75 |
| 22:B0:41:ARG:HD2 | 22:B0:41:ARG:H | 1.50 | 0.75 |
| 35:BF:63:LYS:CE | 35:BF:67:GLN:HB2 | 2.16 | 0.75 |
| 41:BP:16:ARG:CD | 41:BP:18:ARG:HB2 | 2.16 | 0.75 |
| 42:BQ:20:ALA:HB2 | 42:BQ:99:PRO:HD2 | 1.68 | 0.75 |
| 42:BQ:43:THR:OG1 | 42:BQ:46:GLN:HG3 | 1.86 | 0.75 |
| 47:BV:64:HIS:O | 47:BV:64:HIS:CG | 2.37 | 0.75 |
| 8:CH:10:LEU:HD13 | 8:CH:83:ILE:HD11 | 1.68 | 0.75 |
| 29:D7:8:ASN:C | 29:D7:8:ASN:ND2 | 2.41 | 0.75 |
| 31:DA:314:A:O2' | 31:DA:315:G:H5' | 1.86 | 0.75 |
| 31:DA:806:C:OP2 | 41:DP:39:LYS:HD2 | 1.86 | 0.75 |
| 31:DA:1493:C:H2' | 31:DA:1493:C:O2 | 1.86 | 0.75 |
| 32:DB:30:C:OP2 | 44:DS:32:LEU:HD11 | 1.87 | 0.75 |
| 37:DH:98:LEU:HD22 | 37:DH:125:VAL:HG23 | 1.69 | 0.75 |
| 38:DI:133:HIS:CB | 38:DI:134:PRO:HD2 | 2.17 | 0.75 |
| 49:DX:77:LYS:CG | 49:DX:78:LYS:HG3 | 2.14 | 0.75 |
| 31:BA:1833:U:H2' | 31:BA:1834:U:H6 | 1.52 | 0.74 |
| 45:BT:38:ASN:HD22 | 45:BT:38:ASN:C | 1.94 | 0.74 |
| 1:CA:38:G:C2 | 1:CA:397:A:C2 | 2.74 | 0.74 |
| 1:CA:59:A:H5" | 1:CA:60:A:C5' | 2.13 | 0.74 |
| 1:CA:775:G:C2' | 1:CA:776:G:H5' | 2.17 | 0.74 |
| 1:CA:1279:A:H5" | 1:CA:1280:A:OP1 | 1.86 | 0.74 |
| 12:CL:25:PRO:O | 12:CL:27:LEU:HD22 | 1.87 | 0.74 |
| 23:D1:12:PRO:HD2 | 23:D1:62:VAL:HG23 | 1.68 | 0.74 |
| 31:DA:61:G:H1 | 31:DA:94:C:H42 | 1.34 | 0.74 |
| 31:DA:102:G:H8 | 31:DA:102:G:C5' | 1.99 | 0.74 |
| 31:DA:145:G:C2' | 31:DA:146:G:H5" | 2.17 | 0.74 |
| 31:DA:212:G:O2' | 31:DA:213:A:H5' | 1.86 | 0.74 |
| 33:DD:127:VAL:HA | 33:DD:193:VAL:HG22 | 1.69 | 0.74 |
| 34:DE:51:PHE:O | 34:DE:74:PRO:HB3 | 1.87 | 0.74 |
| 35:DF:63:LYS:CE | 35:DF:67:GLN:HB2 | 2.16 | 0.74 |
| 1:AA:1096:C:O2' | 1:AA:1097:C:H5' | 1.88 | 0.74 |
| 31:BA:141:A:C8 | 31:BA:1408:C:O2' | 2.39 | 0.74 |
| 31:BA:2655:G:H2' | 31:BA:2655:G:N3 | 2.02 | 0.74 |
| 31:BA:2789:C:OP1 | 31:BA:2789:C:H4' | 1.87 | 0.74 |
| 37:BH:124:GLU:HB2 | 37:BH:132:ARG:HG2 | 1.69 | 0.74 |
| 42:BQ:38:GLU:OE2 | 42:BQ:127:ILE:HG22 | 1.87 | 0.74 |
| 1:CA:247:G:OP2 | 17:CQ:100:LYS:HG2 | 1.87 | 0.74 |
| 1:CA:499:A:H4' | 1:CA:500:G:OP1 | 1.87 | 0.74 |
| 1:CA:1079:G:H2' | 1:CA:1080:A:C8 | 2.23 | 0.74 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 20:CT:50:GLU:HB3 | 20:CT:100:ILE:CG1 | 2.17 | 0.74 |
| 31:DA:1603:A:H8 | 31:DA:1603:A:H5' | 1.51 | 0.74 |
| 31:DA:2756:U:H4' | 31:DA:2757:A:OP1 | 1.87 | 0.74 |
| 39:DN:47:ALA:HB2 | 39:DN:112:LEU:CD1 | 2.16 | 0.74 |
| 47:DV:47:VAL:HG13 | 47:DV:48:GLY:N | 2.01 | 0.74 |
| 50:DY:20:TYR:CD2 | 50:DY:41:GLY:HA2 | 2.22 | 0.74 |
| 1:AA:627:G:H2' | 1:AA:628:G:C8 | 2.22 | 0.74 |
| 1:AA:878:G:H5' | 8:AH:89:PRO:HG2 | 1.69 | 0.74 |
| 4:AD:31:CYS:C | 4:AD:33:MET:H | 1.93 | 0.74 |
| 7:AG:15:ASP:HB3 | 7:AG:19:GLY:H | 1.53 | 0.74 |
| 31:BA:1502:C:H2' | 31:BA:1502:C:O2 | 1.86 | 0.74 |
| 31:BA:2393:A:H5' | 41:BP:62:LEU:HB3 | 1.67 | 0.74 |
| 50:BY:38:ILE:HG22 | 50:BY:39:VAL:N | 2.00 | 0.74 |
| 1:CA:159:G:O2' | 1:CA:160:A:C8 | 2.39 | 0.74 |
| 31:DA:370:G:H4' | 31:DA:371:A:OP2 | 1.86 | 0.74 |
| 31:DA:2818:G:O2' | 31:DA:2819:G:H5' | 1.87 | 0.74 |
| 33:DD:35:LYS:HZ3 | 33:DD:104:TYR:CB | 1.99 | 0.74 |
| 43:DR:10:LEU:HD22 | 43:DR:17:ARG:HD2 | 1.69 | 0.74 |
| 47:DV:25:LEU:H | 47:DV:94:LEU:CD1 | 1.99 | 0.74 |
| 6:AF:30:LEU:HB2 | 6:AF:35:ALA:HB3 | 1.67 | 0.74 |
| 20:AT:50:GLU:HB3 | 20:AT:100:ILE:CG1 | 2.17 | 0.74 |
| 31:BA:2818:G:O2' | 31:BA:2819:G:H5' | 1.87 | 0.74 |
| 31:BA:2875:C:H4' | 45:BT:5:ALA:HB2 | 1.69 | 0.74 |
| 49:BX:63:LYS:HZ2 | 49:BX:70:LEU:HD21 | 1.53 | 0.74 |
| 49:BX:65:ARG:NE | 49:BX:65:ARG:HA | 2.02 | 0.74 |
| 2:CB:162:ILE:HD11 | 2:CB:184:VAL:HG22 | 1.69 | 0.74 |
| 4:CD:31:CYS:C | 4:CD:33:MET:H | 1.93 | 0.74 |
| 38:DI:6:LEU:O | 38:DI:15:VAL:HB | 1.87 | 0.74 |
| 43:DR:4:LEU:O | 43:DR:4:LEU:CD1 | 2.35 | 0.74 |
| 46:DU:90:VAL:HG12 | 46:DU:91:ASP:H | 1.52 | 0.74 |
| 1:AA:598:U:H4' | 8:AH:94:TYR:CD2 | 2.22 | 0.74 |
| 37:BH:98:LEU:HD22 | 37:BH:125:VAL:HG23 | 1.68 | 0.74 |
| 41:BP:30:THR:HG22 | 41:BP:31:ALA:N | 2.01 | 0.74 |
| 45:BT:13:ARG:HH21 | 45:BT:15:VAL:HG13 | 1.51 | 0.74 |
| 45:BT:32:TYR:HB3 | 45:BT:81:PRO:O | 1.86 | 0.74 |
| 47:BV:62:LEU:CD2 | 47:BV:98:GLU:HB2 | 2.17 | 0.74 |
| 22:D0:20:ARG:NH1 | 31:DA:2357:U:OP1 | 2.21 | 0.74 |
| 31:DA:84:A:H61 | 31:DA:102:G:H1' | 1.50 | 0.74 |
| 31:DA:2475:C:H5' | 31:DA:2476:A:OP2 | 1.87 | 0.74 |
| 49:DX:65:ARG:CZ | 49:DX:66:LEU:H | 2.00 | 0.74 |
| 31:BA:598:G:H5' | 41:BP:15:ARG:HD2 | 1.69 | 0.74 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 33:BD:96:HIS:CE1 | 33:BD:102:LYS:HE2 | 2.22 | 0.74 |
| 35:BF:194:MET:HE3 | 35:BF:199:TRP:HD1 | 1.50 | 0.74 |
| 37:BH:16:SER:HB3 | 37:BH:27:LYS:HG3 | 1.70 | 0.74 |
| 41:BP:101:VAL:HG12 | 41:BP:106:LEU:HB3 | 1.69 | 0.74 |
| 46:BU:93:LYS:HD3 | 46:BU:93:LYS:H | 1.53 | 0.74 |
| 1:CA:192:U:H2' | 1:CA:193:C:H6 | 1.52 | 0.74 |
| 10:CJ:63:PHE:HZ | 14:CN:45:ARG:HG3 | 1.52 | 0.74 |
| 34:DE:132:HIS:CD2 | 34:DE:135:HIS:CE1 | 2.75 | 0.74 |
| 41:DP:51:PHE:HB3 | 41:DP:52:GLU:OE2 | 1.88 | 0.74 |
| 31:BA:142:A:H5' | 31:BA:142(A):C:OP2 | 1.87 | 0.74 |
| 31:BA:811:U:O2 | 31:BA:1250:G:H3' | 1.88 | 0.74 |
| 31:BA:2790:A:H2' | 31:BA:2791:C:C5' | 2.16 | 0.74 |
| 32:BB:47:C:O2' | 44:BS:93:LYS:HG2 | 1.87 | 0.74 |
| 2:CB:180:LEU:O | 2:CB:181:PHE:HB2 | 1.86 | 0.74 |
| 24:D2:30:ARG:H | 24:D2:30:ARG:HD2 | 1.51 | 0.74 |
| 31:DA:2498:C:O2' | 31:DA:2499:C:H5' | 1.88 | 0.74 |
| 49:DX:60:ARG:HE | 49:DX:74:PRO:CG | 2.00 | 0.74 |
| 50:DY:95:LYS:CD | 50:DY:100:ALA:HB1 | 2.14 | 0.74 |
| 1:AA:1285:A:H1' | 1:AA:1286:A:OP2 | 1.88 | 0.74 |
| 28:B6:10:LEU:H | 28:B6:10:LEU:CD2 | 2.00 | 0.74 |
| 30:B8:6:THR:HB | 30:B8:63:PRO:HG3 | 1.69 | 0.74 |
| 35:BF:124:LEU:HD12 | 35:BF:125:LEU:N | 2.02 | 0.74 |
| 10:CJ:49:VAL:HG13 | 14:CN:41:ARG:HB2 | 1.70 | 0.74 |
| 28:D6:33:LYS:O | 28:D6:34:LEU:HB2 | 1.87 | 0.74 |
| 30:D8:59:LYS:HB2 | 30:D8:59:LYS:HZ2 | 1.50 | 0.74 |
| 31:DA:2683:C:OP1 | 45:DT:53:ARG:NH2 | 2.21 | 0.74 |
| 37:DH:70:THR:HG22 | 37:DH:74:ASN:HD21 | 1.52 | 0.74 |
| 1:AA:1030(A):G:H1' | 1:AA:1031:G:H22 | 1.52 | 0.74 |
| 30:B8:35:GLN:NE2 | 30:B8:36:LYS:NZ | 2.35 | 0.74 |
| 31:BA:84:A:H61 | 31:BA:102:G:H1' | 1.49 | 0.74 |
| 31:BA:1484:G:H21 | 31:BA:1505:C:H41 | 0.81 | 0.74 |
| 31:BA:1528(A):A:C5 | 31:BA:1529:G:H8 | 2.06 | 0.74 |
| 32:BB:15:A:H1' | 32:BB:110:G:C8 | 2.23 | 0.74 |
| 34:BE:132:HIS:CG | 34:BE:135:HIS:NE2 | 2.56 | 0.74 |
| 37:BH:85:LYS:HZ3 | 37:BH:145:ALA:HA | 1.50 | 0.74 |
| 41:BP:91:PHE:HE2 | 41:BP:95:VAL:HG12 | 1.53 | 0.74 |
| 1:CA:377:G:OP1 | 16:CP:3:LYS:HD2 | 1.88 | 0.74 |
| 1:CA:1412:C:H2' | 1:CA:1413:A:C8 | 2.23 | 0.74 |
| 15:CO:33:THR:HG21 | 15:CO:85:LEU:HD22 | 1.70 | 0.74 |
| 23:D1:46:LEU:HD12 | 23:D1:46:LEU:N | 2.03 | 0.74 |
| 31:DA:814:C:H5" | 47:DV:86:GLY:HA3 | 1.68 | 0.74 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1396:U:O2 | 31:DA:1396:U:H2' | 1.86 | 0.74 |
| 31:DA:1762:A:H8 | 31:DA:1762:A:O5' | 1.70 | 0.74 |
| 49:DX:55:ASN:HB2 | 49:DX:78:LYS:HD2 | 1.68 | 0.74 |
| 1:AA:341:C:O2' | 1:AA:342:C:H5' | 1.88 | 0.74 |
| 1:AA:392:G:H2' | 1:AA:393:A:H8 | 1.52 | 0.74 |
| 15:AO:78:TYR:O | 15:AO:82:ILE:HG22 | 1.88 | 0.74 |
| 22:B0:53:MET:HE3 | 22:B0:57:PHE:CD1 | 2.22 | 0.74 |
| 24:B2:56:GLN:NE2 | 24:B2:56:GLN:N | 2.34 | 0.74 |
| 31:BA:271(L):U:H4' | 31:BA:271(M):G:C5 | 2.23 | 0.74 |
| 33:BD:16:MET:HB3 | 33:BD:207:GLY:HA3 | 1.69 | 0.74 |
| 47:BV:69:LYS:HG3 | 47:BV:70:ILE:N | 2.02 | 0.74 |
| 30:D8:6:THR:HB | 30:D8:63:PRO:HG3 | 1.69 | 0.74 |
| 40:DO:98:VAL:HG12 | 40:DO:117:LEU:HB3 | 1.69 | 0.74 |
| 41:DP:24:GLY:CA | 41:DP:33:ARG:HH21 | 1.94 | 0.74 |
| 31:BA:827:U:O2' | 31:BA:2068:U:C2 | 2.37 | 0.73 |
| 31:BA:1210:A:H4' | 31:BA:1211:U:OP2 | 1.87 | 0.73 |
| 35:BF:63:LYS:HE2 | 35:BF:67:GLN:HB2 | 1.69 | 0.73 |
| 41:BP:62:LEU:H | 41:BP:62:LEU:HD13 | 1.53 | 0.73 |
| 45:BT:29:ARG:HB3 | 45:BT:85:LYS:HA | 1.70 | 0.73 |
| 46:BU:88:ILE:O | 46:BU:90:VAL:N | 2.16 | 0.73 |
| 29:D7:8:ASN:HD21 | 29:D7:11:LYS:N | 1.81 | 0.73 |
| 29:D7:47:ARG:C | 29:D7:48:LYS:HD3 | 2.13 | 0.73 |
| 30:D8:32:LEU:HD23 | 30:D8:35:GLN:CA | 2.18 | 0.73 |
| 41:DP:17:LYS:HG3 | 41:DP:19:VAL:HG23 | 1.70 | 0.73 |
| 1:AA:328:C:O2 | 1:AA:328:C:H2' | 1.87 | 0.73 |
| 1:AA:333:G:H4' | 20:AT:16:HIS:CE1 | 2.24 | 0.73 |
| 2:AB:137:ARG:HA | 2:AB:137:ARG:HH11 | 1.53 | 0.73 |
| 12:AL:47:LYS:HB3 | 12:AL:48:PRO:CD | 2.17 | 0.73 |
| 12:AL:47:LYS:CB | 12:AL:48:PRO:HD3 | 2.18 | 0.73 |
| 13:AM:10:PRO:O | 13:AM:45:VAL:HG11 | 1.88 | 0.73 |
| 16:AP:43:LYS:HA | 16:AP:48:TRP:HB3 | 1.70 | 0.73 |
| 23:B1:26:ARG:CB | 23:B1:34:THR:HA | 2.17 | 0.73 |
| 31:BA:1956:U:H2' | 31:BA:1957:C:H5' | 1.70 | 0.73 |
| 35:BF:198:ALA:O | 35:BF:201:VAL:HG12 | 1.89 | 0.73 |
| 45:BT:28:VAL:HG21 | 45:BT:46:GLU:HG3 | 1.69 | 0.73 |
| 46:BU:83:LEU:HG | 46:BU:88:ILE:HG12 | 1.69 | 0.73 |
| 46:BU:88:ILE:C | 46:BU:90:VAL:N | 2.45 | 0.73 |
| 28:D6:22:ALA:HB2 | 28:D6:39:TYR:CE2 | 2.22 | 0.73 |
| 31:DA:1218:C:H2' | 31:DA:1219:G:H5' | 1.69 | 0.73 |
| 31:DA:1528(A):A:C5 | 31:DA:1529:G:H8 | 2.06 | 0.73 |
| 31:DA:1956:U:H2' | 31:DA:1957:C:H5' | 1.70 | 0.73 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 45:DT:65:LYS:HE3 | 45:DT:66:VAL:H | 1.54 | 0.73 |
| 30:B8:32:LEU:O | 30:B8:33:ASN:CB | 2.36 | 0.73 |
| 31:BA:1448:G:H1' | 31:BA:1528:A:H62 | 1.53 | 0.73 |
| 31:BA:2723:C:H5" | 43:BR:2:ARG:CD | 2.18 | 0.73 |
| 39:BN:47:ALA:HB2 | 39:BN:112:LEU:CD1 | 2.12 | 0.73 |
| 50:BY:7:VAL:HB | 50:BY:8:LYS:NZ | 2.02 | 0.73 |
| 31:DA:139(A):G:N2 | 49:DX:44:GLU:OE1 | 2.20 | 0.73 |
| 31:DA:310:A:OP1 | 50:DY:17:SER:O | 2.07 | 0.73 |
| 31:DA:1256:G:H5' | 31:DA:1257:C:OP2 | 1.88 | 0.73 |
| 31:DA:2335:A:O2' | 31:DA:2336:A:H5" | 1.87 | 0.73 |
| 51:DZ:151:HIS:HB3 | 51:DZ:170:THR:CA | 2.08 | 0.73 |
| 1:AA:60:A:P | 1:AA:60:A:H8 | 2.11 | 0.73 |
| 6:AF:26:ILE:HG22 | 6:AF:30:LEU:HD21 | 1.69 | 0.73 |
| 30:B8:46:ARG:NH2 | 41:BP:65:ARG:HH22 | 1.84 | 0.73 |
| 31:BA:1109:C:H5 | 31:BA:1110:G:C5 | 2.06 | 0.73 |
| 31:BA:1179:C:H3' | 31:BA:1180:C:H5" | 1.68 | 0.73 |
| 31:BA:1747(A):G:H2' | 31:BA:1748:G:C5' | 2.16 | 0.73 |
| 31:BA:1971:A:H1' | 33:BD:240:ALA:O | 1.87 | 0.73 |
| 31:BA:2030:A:H4' | 31:BA:2031:A:OP1 | 1.87 | 0.73 |
| 33:BD:255:LYS:H | 33:BD:255:LYS:CE | 1.98 | 0.73 |
| 42:BQ:104:PHE:HE1 | 42:BQ:125:LEU:HD11 | 1.52 | 0.73 |
| 47:BV:52:VAL:O | 47:BV:53:GLU:HB3 | 1.89 | 0.73 |
| 47:BV:66:ARG:HG2 | 47:BV:94:LEU:HG | 1.70 | 0.73 |
| 1:CA:826:C:H2' | 1:CA:827:U:H6 | 1.53 | 0.73 |
| 12:CL:47:LYS:CB | 12:CL:48:PRO:HD3 | 2.18 | 0.73 |
| 16:CP:74:LEU:O | 16:CP:79:VAL:HG23 | 1.88 | 0.73 |
| 23:D1:19:GLN:NE2 | 31:DA:379:G:H21 | 1.86 | 0.73 |
| 23:D1:85:LEU:HB3 | 23:D1:87:PRO:CD | 2.18 | 0.73 |
| 41:DP:108:LYS:C | 41:DP:110:TYR:H | 1.93 | 0.73 |
| 43:DR:4:LEU:O | 43:DR:6:SER:N | 2.21 | 0.73 |
| 45:DT:25:GLY:O | 45:DT:26:ASP:HB2 | 1.89 | 0.73 |
| 27:B5:51:TYR:CD2 | 27:B5:52:TYR:CZ | 2.76 | 0.73 |
| 28:B6:20:ASN:CG | 28:B6:21:TYR:H | 1.96 | 0.73 |
| 30:B8:34:TRP:O | 30:B8:35:GLN:HB2 | 1.88 | 0.73 |
| 31:BA:2758:A:H2' | 31:BA:2759:G:H5" | 1.71 | 0.73 |
| 38:BI:83:ALA:HA | 38:BI:89:TYR:CD1 | 2.24 | 0.73 |
| 50:BY:45:VAL:HG11 | 50:BY:62:GLU:H | 1.53 | 0.73 |
| 1:CA:627:G:H2' | 1:CA:628:G:C8 | 2.23 | 0.73 |
| 7:CG:15:ASP:HB3 | 7:CG:19:GLY:H | 1.52 | 0.73 |
| 22:D0:74:ARG:HH22 | 32:DB:13:A:H8 | 1.36 | 0.73 |
| 23:D1:87:PRO:HD2 | 23:D1:88:LYS:N | 2.01 | 0.73 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 30:D8:46:ARG:NH2 | 41:DP:65:ARG:HH22 | 1.87 | 0.73 |
| 31:DA:1980:G:O2' | 31:DA:1982:C:OP2 | 2.04 | 0.73 |
| 41:DP:95:VAL:HG22 | 41:DP:125:VAL:HB | 1.71 | 0.73 |
| 42:DQ:42:ILE:HD13 | 42:DQ:97:VAL:HG21 | 1.70 | 0.73 |
| 42:DQ:104:PHE:HE1 | 42:DQ:125:LEU:HD11 | 1.54 | 0.73 |
| 45:DT:106:SER:O | 45:DT:107:ASP:CG | 2.31 | 0.73 |
| 50:DY:96:ILE:CD1 | 50:DY:99:CYS:SG | 2.77 | 0.73 |
| 1:AA:159:G:O2' | 1:AA:160:A:C8 | 2.41 | 0.73 |
| 1:AA:1442(A):G:H8 | 45:BT:118:ARG:HH11 | 1.36 | 0.73 |
| 31:BA:1188:U:H2' | 31:BA:1189:A:H5' | 1.69 | 0.73 |
| 31:BA:2292:C:O2' | 31:BA:2293:C:H5' | 1.89 | 0.73 |
| 31:BA:2598:A:P | 33:BD:236:GLY:HA3 | 2.28 | 0.73 |
| 35:BF:203:GLN:HA | 35:BF:206:ILE:O | 1.88 | 0.73 |
| 1:CA:224:C:H2' | 1:CA:225:C:C6 | 2.24 | 0.73 |
| 1:CA:1228:C:H2' | 1:CA:1229:A:H8 | 1.53 | 0.73 |
| 17:CQ:18:THR:HG23 | 17:CQ:69:LYS:HE3 | 1.67 | 0.73 |
| 31:DA:954:G:H5" | 42:DQ:13:GLN:HG2 | 1.69 | 0.73 |
| 37:DH:16:SER:HB3 | 37:DH:27:LYS:HG3 | 1.69 | 0.73 |
| 42:DQ:27:VAL:HG13 | 42:DQ:105:GLU:OE1 | 1.89 | 0.73 |
| 31:BA:322:A:H5' | 31:BA:340:A:H1' | 1.70 | 0.73 |
| 37:BH:99:VAL:HG12 | 37:BH:99:VAL:O | 1.87 | 0.73 |
| 1:CA:922:G:H4' | 5:CE:20:GLN:HA | 1.69 | 0.73 |
| 6:CF:67:MET:HB2 | 6:CF:68:PRO:HD2 | 1.70 | 0.73 |
| 23:D1:26:ARG:CB | 23:D1:34:THR:HA | 2.17 | 0.73 |
| 31:DA:811:U:O2 | 31:DA:1250:G:H3' | 1.88 | 0.73 |
| 31:DA:2292:C:O2' | 31:DA:2293:C:H5' | 1.89 | 0.73 |
| 38:DI:91:SER:HB3 | 38:DI:121:LYS:HB2 | 1.69 | 0.73 |
| 38:DI:101:LEU:O | 38:DI:101:LEU:HD12 | 1.89 | 0.73 |
| 44:DS:83:LYS:HE3 | 44:DS:105:ALA:HB2 | 1.69 | 0.73 |
| 50:DY:95:LYS:HD3 | 50:DY:100:ALA:CB | 2.15 | 0.73 |
| 1:AA:224:C:H2' | 1:AA:225:C:C6 | 2.24 | 0.73 |
| 5:AE:26:PHE:O | 5:AE:27:ARG:HB2 | 1.86 | 0.73 |
| 8:AH:1:MET:HE2 | 8:AH:1:MET:N | 2.03 | 0.73 |
| 31:BA:2682:U:C6 | 34:BE:11:MET:HE2 | 2.24 | 0.73 |
| 38:BI:133:HIS:CB | 38:BI:134:PRO:HD2 | 2.18 | 0.73 |
| 1:CA:819:A:H4' | 1:CA:820:U:OP2 | 1.88 | 0.73 |
| 1:CA:1096:C:O2' | 1:CA:1097:C:H5' | 1.88 | 0.73 |
| 3:CC:182:ILE:HG12 | 3:CC:203:PHE:HD1 | 1.52 | 0.73 |
| 18:CR:53:ARG:HH21 | 18:CR:60:ALA:N | 1.87 | 0.73 |
| 27:D5:4:HIS:HB3 | 27:D5:5:PRO:CD | 2.18 | 0.73 |
| 29:D7:16:HIS:HB2 | 29:D7:44:PRO:HG2 | 1.71 | 0.73 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:83:G:H1 | 31:DA:102:G:H2' | 1.51 | 0.73 |
| 31:DA:184:C:H2' | 31:DA:185:U:H6 | 1.52 | 0.73 |
| 31:DA:1899:G:N2 | 31:DA:1902:C:C5 | 2.57 | 0.73 |
| 31:DA:2789:C:H4' | 31:DA:2789:C:OP1 | 1.86 | 0.73 |
| 33:DD:75:ILE:HG21 | 33:DD:99:ASP:HB2 | 1.71 | 0.73 |
| 35:DF:46:ARG:HH11 | 35:DF:46:ARG:CG | 2.02 | 0.73 |
| 42:DQ:83:MET:HG2 | 42:DQ:83:MET:O | 1.89 | 0.73 |
| 50:DY:7:VAL:HB | 50:DY:8:LYS:HZ2 | 1.54 | 0.73 |
| 1:AA:243:A:H4' | 1:AA:244:U:O5' | 1.88 | 0.73 |
| 1:AA:327:A:H3' | 1:AA:328:C:H5" | 1.69 | 0.73 |
| 1:AA:1079:G:H2' | 1:AA:1080:A:C8 | 2.23 | 0.73 |
| 18:AR:53:ARG:HH21 | 18:AR:60:ALA:N | 1.86 | 0.73 |
| 25:B3:52:HIS:CD2 | 32:BB:83:G:H5" | 2.24 | 0.73 |
| 30:B8:59:LYS:CD | 41:BP:50:ARG:HB3 | 2.19 | 0.73 |
| 31:BA:481:G:O2' | 31:BA:482:A:OP2 | 2.06 | 0.73 |
| 31:BA:1495:A:H2' | 31:BA:1496:A:N3 | 2.03 | 0.73 |
| 31:BA:2565:A:H5" | 31:BA:2566:A:OP2 | 1.87 | 0.73 |
| 39:BN:40:PRO:CA | 46:BU:64:ARG:HH22 | 1.96 | 0.73 |
| 45:BT:28:VAL:CG2 | 45:BT:46:GLU:HG3 | 2.19 | 0.73 |
| 47:BV:51:VAL:HG12 | 47:BV:52:VAL:H | 1.52 | 0.73 |
| 47:BV:70:ILE:HG13 | 47:BV:90:PRO:HB3 | 1.70 | 0.73 |
| 50:BY:45:VAL:CG1 | 50:BY:62:GLU:H | 2.01 | 0.73 |
| 1:CA:148:G:O2' | 1:CA:149:A:H5' | 1.88 | 0.73 |
| 1:CA:243:A:H4' | 1:CA:244:U:O5' | 1.88 | 0.73 |
| 1:CA:600:C:H2' | 1:CA:601:C:C6 | 2.24 | 0.73 |
| 1:CA:1226:C:H42 | 13:CM:104:ARG:HD2 | 1.54 | 0.73 |
| 4:CD:138:TYR:C | 4:CD:138:TYR:HD2 | 1.96 | 0.73 |
| 18:CR:31:LEU:HD23 | 18:CR:31:LEU:H | 1.52 | 0.73 |
| 31:DA:288:C:H42 | 31:DA:353:G:H1 | 1.36 | 0.73 |
| 31:DA:1188:U:H2' | 31:DA:1189:A:H5' | 1.71 | 0.73 |
| 31:DA:1946:U:H2' | 31:DA:1947:C:H6 | 1.53 | 0.73 |
| 34:DE:24:THR:HG23 | 34:DE:184:VAL:HG23 | 1.70 | 0.73 |
| 47:DV:62:LEU:HA | 47:DV:98:GLU:HA | 1.69 | 0.73 |
| 1:AA:735:C:H2' | 1:AA:736:C:C6 | 2.23 | 0.73 |
| 3:AC:50:ALA:HB1 | 3:AC:70:VAL:HG11 | 1.71 | 0.73 |
| 8:AH:20:TYR:HA | 8:AH:65:TYR:CE2 | 2.23 | 0.73 |
| 16:AP:22:THR:HA | 16:AP:33:ILE:HG12 | 1.71 | 0.73 |
| 16:AP:28:ARG:HG2 | 16:AP:28:ARG:NH1 | 1.90 | 0.73 |
| 24:B2:45:SER:O | 24:B2:48:HIS:N | 2.20 | 0.73 |
| 45:BT:25:GLY:O | 45:BT:26:ASP:HB2 | 1.86 | 0.73 |
| 49:BX:60:ARG:HE | 49:BX:74:PRO:CG | 2.01 | 0.73 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 49:BX:82:GLN:C | 49:BX:85:PRO:HD2 | 2.13 | 0.73 |
| 1:CA:333:G:H4' | 20:CT:16:HIS:CE1 | 2.24 | 0.73 |
| 2:CB:11:LEU:HB3 | 2:CB:213:LEU:HD11 | 1.71 | 0.73 |
| 11:CK:111:ASP:HA | 18:CR:84:LYS:HE2 | 1.71 | 0.73 |
| 16:CP:22:THR:HA | 16:CP:33:ILE:HG12 | 1.70 | 0.73 |
| 27:D5:40:LYS:HE2 | 27:D5:46:CYS:HB3 | 1.71 | 0.73 |
| 43:DR:11:ASN:CG | 43:DR:12:ARG:H | 1.95 | 0.73 |
| 22:B0:72:ARG:HB2 | 22:B0:75:LEU:HB2 | 1.70 | 0.72 |
| 23:B1:67:ILE:N | 23:B1:68:PRO:HD2 | 2.02 | 0.72 |
| 24:B2:37:PHE:HE2 | 24:B2:40:SER:HA | 1.54 | 0.72 |
| 31:BA:142:A:H8 | 31:BA:1595:G:H21 | 1.37 | 0.72 |
| 31:BA:1657:C:O2' | 31:BA:1658:C:H5' | 1.89 | 0.72 |
| 31:BA:2029:G:H2' | 31:BA:2031:A:OP2 | 1.89 | 0.72 |
| 32:BB:29:A:P | 44:BS:32:LEU:HD12 | 2.29 | 0.72 |
| 49:BX:33:LYS:O | 49:BX:35:THR:N | 2.22 | 0.72 |
| 23:D1:37:ILE:HG21 | 31:DA:2080:G:OP1 | 1.88 | 0.72 |
| 23:D1:67:ILE:N | 23:D1:68:PRO:HD2 | 2.04 | 0.72 |
| 23:D1:85:LEU:CB | 23:D1:87:PRO:HD3 | 2.19 | 0.72 |
| 24:D2:37:PHE:HE2 | 24:D2:40:SER:HA | 1.53 | 0.72 |
| 31:DA:1495:A:H2' | 31:DA:1496:A:N3 | 2.04 | 0.72 |
| 33:DD:35:LYS:HZ3 | 33:DD:104:TYR:HD1 | 1.36 | 0.72 |
| 37:DH:124:GLU:HB2 | 37:DH:132:ARG:HG2 | 1.68 | 0.72 |
| 47:DV:18:LEU:HD22 | 47:DV:19:LYS:CA | 2.19 | 0.72 |
| 1:AA:1456:G:H2' | 1:AA:1457:G:O4' | 1.88 | 0.72 |
| 31:BA:7:G:H2' | 31:BA:8:A:O4' | 1.89 | 0.72 |
| 31:BA:542:C:N3 | 31:BA:543:C:N4 | 2.36 | 0.72 |
| 31:BA:2653:U:H3' | 31:BA:2654:A:H5" | 1.69 | 0.72 |
| 36:BG:67:LYS:HD2 | 36:BG:67:LYS:H | 1.54 | 0.72 |
| 42:BQ:75:THR:HA | 42:BQ:88:GLY:HA2 | 1.70 | 0.72 |
| 47:BV:69:LYS:CB | 47:BV:93:GLU:CD | 2.62 | 0.72 |
| 16:CP:4:ILE:HG13 | 16:CP:21:VAL:CG1 | 2.19 | 0.72 |
| 31:DA:833:U:H2' | 31:DA:834:C:C6 | 2.23 | 0.72 |
| 31:DA:1336:A:H2' | 31:DA:1337:G:C8 | 2.24 | 0.72 |
| 31:DA:1434:A:H61 | 31:DA:1558:A:N6 | 1.86 | 0.72 |
| 31:DA:2657:A:H5' | 31:DA:2658:C:OP2 | 1.89 | 0.72 |
| 33:DD:30:GLU:HG3 | 33:DD:63:ARG:CZ | 2.19 | 0.72 |
| 45:DT:56:GLY:O | 45:DT:59:THR:CG2 | 2.37 | 0.72 |
| 1:AA:600:C:H2' | 1:AA:601:C:C6 | 2.25 | 0.72 |
| 32:BB:82:G:C2' | 32:BB:83:G:H5' | 2.19 | 0.72 |
| 37:BH:144:VAL:O | 37:BH:148:ILE:HG12 | 1.88 | 0.72 |
| 12:CL:47:LYS:HB3 | 12:CL:48:PRO:CD | 2.17 | 0.72 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 13:CM:10:PRO:O | 13:CM:45:VAL:HG11 | 1.89 | 0.72 |
| 23:D1:49:VAL:HG11 | 31:DA:2091:U:O2' | 1.88 | 0.72 |
| 31:DA:2777:G:H5" | 31:DA:2778:A:H5' | 1.71 | 0.72 |
| 31:DA:2787:C:C1' | 34:DE:61:ARG:HB2 | 2.19 | 0.72 |
| 33:DD:58:HIS:CD2 | 33:DD:59:LYS:N | 2.57 | 0.72 |
| 34:DE:111:ARG:NH1 | 43:DR:2:ARG:HH21 | 1.87 | 0.72 |
| 45:DT:109:GLU:HA | 45:DT:112:ARG:HG3 | 1.71 | 0.72 |
| 50:DY:8:LYS:HE2 | 50:DY:72:VAL:HG23 | 1.68 | 0.72 |
| 1:AA:192:U:H2' | 1:AA:193:C:C6 | 2.24 | 0.72 |
| 1:AA:1441:G:H5" | 1:AA:1442:G:O5' | 1.90 | 0.72 |
| 3:AC:182:ILE:HG12 | 3:AC:203:PHE:HD1 | 1.54 | 0.72 |
| 10:AJ:26:ALA:HB1 | 10:AJ:29:ARG:HH21 | 1.53 | 0.72 |
| 31:BA:1777:U:O2' | 31:BA:1778:U:H5' | 1.89 | 0.72 |
| 34:BE:61:ARG:H | 34:BE:62:PRO:CD | 2.02 | 0.72 |
| 35:BF:20:LEU:HD13 | 35:BF:203:GLN:OE1 | 1.90 | 0.72 |
| 35:BF:132:VAL:HG22 | 35:BF:133:ASN:H | 1.54 | 0.72 |
| 45:BT:91:ARG:CB | 45:BT:116:ALA:HA | 2.20 | 0.72 |
| 51:BZ:27:VAL:HG23 | 51:BZ:36:LYS:HA | 1.70 | 0.72 |
| 4:CD:57:ARG:NH2 | 5:CE:107:ARG:HD3 | 2.04 | 0.72 |
| 6:CF:89:MET:HG2 | 6:CF:91:VAL:HG23 | 1.71 | 0.72 |
| 31:DA:196:A:O4' | 41:DP:46:LYS:HE2 | 1.89 | 0.72 |
| 31:DA:1406:U:H2' | 31:DA:1407:C:H6 | 1.54 | 0.72 |
| 35:DF:8:GLN:HB3 | 35:DF:126:VAL:HA | 1.71 | 0.72 |
| 39:DN:78:TYR:H | 39:DN:79:PRO:HD3 | 1.55 | 0.72 |
| 39:DN:82:LEU:H | 39:DN:82:LEU:HD12 | 1.55 | 0.72 |
| 41:DP:107:LYS:C | 41:DP:109:GLY:H | 1.97 | 0.72 |
| 30:B8:58:ILE:O | 30:B8:61:LEU:HG | 1.90 | 0.72 |
| 31:BA:1876:A:H2' | 31:BA:1877:A:C8 | 2.24 | 0.72 |
| 31:BA:1899:G:H22 | 31:BA:1902:C:H41 | 0.78 | 0.72 |
| 31:BA:2523:G:H2' | 31:BA:2524:G:C5' | 2.19 | 0.72 |
| 33:BD:25:THR:HG21 | 33:BD:81:ALA:CB | 2.19 | 0.72 |
| 41:BP:10:PRO:CD | 41:BP:11:GLY:N | 2.51 | 0.72 |
| 44:BS:24:LEU:HB3 | 44:BS:85:VAL:CG1 | 2.20 | 0.72 |
| 50:BY:95:LYS:HD3 | 50:BY:100:ALA:CB | 2.08 | 0.72 |
| 1:CA:673:G:H2' | 1:CA:674:G:H8 | 1.55 | 0.72 |
| 24:D2:45:SER:O | 24:D2:48:HIS:N | 2.21 | 0.72 |
| 31:DA:672:C:H2' | 31:DA:673:C:H6 | 1.54 | 0.72 |
| 45:DT:28:VAL:CG2 | 45:DT:46:GLU:HG3 | 2.20 | 0.72 |
| 45:DT:82:LEU:HD12 | 45:DT:82:LEU:N | 2.04 | 0.72 |
| 45:DT:109:GLU:HB3 | 45:DT:113:LYS:HE3 | 1.71 | 0.72 |
| 1:AA:1442(A):G:C8 | 45:BT:118:ARG:HD2 | 2.25 | 0.72 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 17:AQ:74:LEU:HD12 | 17:AQ:75:ARG:HG2 | 1.71 | 0.72 |
| 31:BA:212:G:O2' | 31:BA:213:A:H5' | 1.88 | 0.72 |
| 31:BA:1019:U:HO2' | 31:BA:1021:A:H2 | 1.36 | 0.72 |
| 31:BA:2475:C:H5' | 31:BA:2476:A:OP2 | 1.88 | 0.72 |
| 39:BN:57:ALA:C | 39:BN:58:ASP:O | 2.31 | 0.72 |
| 39:BN:65:LYS:O | 39:BN:69:GLN:HB2 | 1.89 | 0.72 |
| 3:CC:70:VAL:HG12 | 3:CC:72:LYS:H | 1.54 | 0.72 |
| 4:CD:23:GLY:HA3 | 4:CD:112:VAL:HG22 | 1.71 | 0.72 |
| 31:DA:1777:U:O2' | 31:DA:1778:U:H5' | 1.89 | 0.72 |
| 34:DE:136:ARG:HH11 | 34:DE:136:ARG:HG2 | 1.51 | 0.72 |
| 40:DO:32:TYR:N | 40:DO:32:TYR:HD1 | 1.88 | 0.72 |
| 44:DS:17:ARG:O | 44:DS:18:ILE:HB | 1.88 | 0.72 |
| 1:AA:1203:C:OP1 | 14:AN:3:ARG:HD3 | 1.89 | 0.72 |
| 2:AB:162:ILE:HD11 | 2:AB:184:VAL:HG22 | 1.69 | 0.72 |
| 28:B6:18:ARG:HG3 | 28:B6:19:ARG:H | 1.54 | 0.72 |
| 31:BA:1713:U:O2' | 31:BA:1714:G:H5' | 1.89 | 0.72 |
| 33:BD:183:ARG:HG2 | 33:BD:183:ARG:NH1 | 1.97 | 0.72 |
| 40:BO:35:VAL:HA | 40:BO:62:VAL:HG12 | 1.70 | 0.72 |
| 45:BT:35:LYS:O | 45:BT:37:GLY:N | 2.23 | 0.72 |
| 45:BT:65:LYS:HE3 | 45:BT:66:VAL:H | 1.52 | 0.72 |
| 50:BY:20:TYR:CD2 | 50:BY:41:GLY:HA2 | 2.24 | 0.72 |
| 1:CA:522:C:H41 | 12:CL:53:ARG:NH2 | 1.87 | 0.72 |
| 7:CG:113:GLU:HB2 | 7:CG:119:ARG:CG | 2.18 | 0.72 |
| 27:D5:40:LYS:CE | 27:D5:46:CYS:HB3 | 2.20 | 0.72 |
| 30:D8:32:LEU:C | 30:D8:34:TRP:N | 2.46 | 0.72 |
| 31:DA:7:G:H2' | 31:DA:8:A:O4' | 1.89 | 0.72 |
| 31:DA:2029:G:H2' | 31:DA:2031:A:OP2 | 1.90 | 0.72 |
| 31:DA:2287:A:N6 | 31:DA:2344:U:H3 | 1.87 | 0.72 |
| 33:DD:16:MET:HB3 | 33:DD:207:GLY:HA3 | 1.72 | 0.72 |
| 46:DU:88:ILE:C | 46:DU:90:VAL:N | 2.45 | 0.72 |
| 3:AC:70:VAL:HG12 | 3:AC:72:LYS:H | 1.52 | 0.72 |
| 10:AJ:6:ILE:HG13 | 10:AJ:72:VAL:O | 1.90 | 0.72 |
| 24:B2:23:LYS:HB2 | 49:BX:5:TYR:CE1 | 2.19 | 0.72 |
| 37:BH:85:LYS:HD3 | 37:BH:133:VAL:HB | 1.72 | 0.72 |
| 41:BP:64:LYS:O | 41:BP:66:GLY:N | 2.22 | 0.72 |
| 1:CA:390:C:O3' | 16:CP:28:ARG:NH2 | 2.23 | 0.72 |
| 8:CH:20:TYR:HA | 8:CH:65:TYR:CE2 | 2.25 | 0.72 |
| 22:D0:72:ARG:HB2 | 22:D0:75:LEU:HB2 | 1.69 | 0.72 |
| 30:D8:34:TRP:O | 30:D8:35:GLN:HB2 | 1.90 | 0.72 |
| 31:DA:329:G:H22 | 50:DY:19:LYS:NZ | 1.88 | 0.72 |
| 31:DA:910:A:H62 | 42:DQ:12:GLN:HA | 1.52 | 0.72 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2287:A:H62 | 31:DA:2344:U:H3 | 1.38 | 0.72 |
| 38:DI:61:ARG:O | 38:DI:133:HIS:CE1 | 2.43 | 0.72 |
| 38:DI:83:ALA:HA | 38:DI:89:TYR:CD1 | 2.24 | 0.72 |
| 38:DI:133:HIS:ND1 | 38:DI:134:PRO:HD2 | 2.05 | 0.72 |
| 42:DQ:75:THR:HA | 42:DQ:88:GLY:HA2 | 1.71 | 0.72 |
| 45:DT:27:THR:O | 45:DT:28:VAL:HG23 | 1.90 | 0.72 |
| 11:AK:24:SER:HB3 | 11:AK:27:ASN:O | 1.90 | 0.72 |
| 11:AK:34:ASP:HB3 | 11:AK:40:ILE:HD11 | 1.72 | 0.72 |
| 25:B3:52:HIS:HD2 | 32:BB:83:G:H4' | 1.53 | 0.72 |
| 31:BA:2359:C:H2' | 31:BA:2360:A:H5' | 1.71 | 0.72 |
| 32:BB:30:C:OP2 | 44:BS:32:LEU:HD11 | 1.90 | 0.72 |
| 34:BE:132:HIS:CD2 | 34:BE:135:HIS:CE1 | 2.77 | 0.72 |
| 37:BH:89:ILE:O | 37:BH:90:LYS:HG2 | 1.89 | 0.72 |
| 40:BO:32:TYR:N | 40:BO:32:TYR:HD1 | 1.88 | 0.72 |
| 41:BP:48:PRO:O | 41:BP:50:ARG:N | 2.23 | 0.72 |
| 42:BQ:82:ARG:O | 42:BQ:83:MET:HB3 | 1.88 | 0.72 |
| 45:BT:109:GLU:HB3 | 45:BT:113:LYS:HE3 | 1.72 | 0.72 |
| 50:BY:44:ILE:H | 50:BY:44:ILE:HD12 | 1.54 | 0.72 |
| 50:BY:45:VAL:CG1 | 50:BY:62:GLU:HB2 | 2.17 | 0.72 |
| 5:CE:100:VAL:HG13 | 5:CE:118:ILE:HG22 | 1.72 | 0.72 |
| 31:DA:1528(A):A:N7 | 31:DA:1529:G:C8 | 2.57 | 0.72 |
| 31:DA:2012:G:H4' | 48:DW:96:ILE:CD1 | 2.16 | 0.72 |
| 34:DE:93:VAL:N | 34:DE:95:ILE:HD13 | 2.04 | 0.72 |
| 45:DT:50:ILE:HD11 | 45:DT:102:ILE:CD1 | 2.17 | 0.72 |
| 46:DU:25:TRP:CD1 | 46:DU:26:GLY:N | 2.58 | 0.72 |
| 1:AA:1063:C:H3' | 1:AA:1064:G:H2' | 1.70 | 0.72 |
| 4:AD:23:GLY:HA3 | 4:AD:112:VAL:HG22 | 1.70 | 0.72 |
| 31:BA:370:G:H4' | 31:BA:371:A:OP2 | 1.87 | 0.72 |
| 31:BA:833:U:H2' | 31:BA:834:C:C6 | 2.25 | 0.72 |
| 35:BF:89:VAL:HG12 | 35:BF:90:PHE:H | 1.54 | 0.72 |
| 45:BT:56:GLY:O | 45:BT:59:THR:CG2 | 2.38 | 0.72 |
| 1:CA:976:G:H5' | 1:CA:1358:U:O2' | 1.90 | 0.72 |
| 1:CA:1228:C:H2' | 1:CA:1229:A:C8 | 2.25 | 0.72 |
| 10:CJ:34:VAL:CG2 | 10:CJ:74:ILE:HG22 | 2.18 | 0.72 |
| 12:CL:124:LYS:HD2 | 12:CL:125:PRO:HD2 | 1.71 | 0.72 |
| 31:DA:2661:G:N7 | 31:DA:2662:A:C2 | 2.57 | 0.72 |
| 35:DF:65:TRP:O | 35:DF:67:GLN:N | 2.23 | 0.72 |
| 37:DH:40:GLU:O | 37:DH:41:MET:HB2 | 1.90 | 0.72 |
| 39:DN:55:VAL:HG12 | 39:DN:125:GLY:HA3 | 1.72 | 0.72 |
| 47:DV:83:ARG:HH11 | 47:DV:83:ARG:HG3 | 1.53 | 0.72 |
| 1:AA:639:G:O2' | 1:AA:640:A:H5' | 1.90 | 0.71 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 27:B5:6:VAL:HG13 | 27:B5:7:PRO:HD2 | 1.72 | 0.71 |
| 31:BA:314:A:O2' | 31:BA:315:G:H5' | 1.90 | 0.71 |
| 31:BA:943:U:OP2 | 41:BP:38:GLN:CD | 2.33 | 0.71 |
| 31:BA:1278:A:O2' | 31:BA:1279:G:H5' | 1.90 | 0.71 |
| 31:BA:1434:A:O2' | 31:BA:1435:G:H5' | 1.90 | 0.71 |
| 31:BA:1658:C:OP1 | 34:BE:132:HIS:ND1 | 2.23 | 0.71 |
| 45:BT:88:ILE:HG22 | 45:BT:89:VAL:HG23 | 1.72 | 0.71 |
| 46:BU:104:GLN:HB2 | 47:BV:43:GLU:OE1 | 1.90 | 0.71 |
| 46:BU:112:ARG:HH11 | 46:BU:112:ARG:HG2 | 1.54 | 0.71 |
| 1:CA:1063:C:H3' | 1:CA:1064:G:H2' | 1.70 | 0.71 |
| 15:CO:87:ILE:HG22 | 15:CO:88:ARG:N | 2.04 | 0.71 |
| 30:D8:35:GLN:HA | 31:DA:2420:C:P | 2.30 | 0.71 |
| 31:DA:322:A:H5' | 31:DA:340:A:H1' | 1.72 | 0.71 |
| 31:DA:1747(A):G:H2' | 31:DA:1748:G:C5' | 2.18 | 0.71 |
| 31:DA:2203:U:H1' | 33:DD:151:LYS:HE2 | 1.72 | 0.71 |
| 31:DA:2488:A:H2' | 31:DA:2489:G:O4' | 1.89 | 0.71 |
| 31:DA:2685:G:H2' | 31:DA:2686:G:H5" | 1.72 | 0.71 |
| 33:DD:223:GLY:HA3 | 33:DD:231:HIS:CE1 | 2.25 | 0.71 |
| 42:DQ:8:LYS:HD2 | 42:DQ:9:TYR:H | 1.55 | 0.71 |
| 47:DV:71:LEU:HD13 | 47:DV:72:VAL:H | 1.54 | 0.71 |
| 1:AA:819:A:H4' | 1:AA:820:U:OP2 | 1.89 | 0.71 |
| 12:AL:41:ARG:CG | 12:AL:42:THR:H | 2.03 | 0.71 |
| 17:AQ:18:THR:HG23 | 17:AQ:69:LYS:HE3 | 1.70 | 0.71 |
| 31:BA:2307:G:H21 | 31:BA:2308:G:C5' | 2.02 | 0.71 |
| 31:BA:2506:U:H5' | 31:BA:2506:U:C6 | 2.24 | 0.71 |
| 37:BH:155:SER:O | 37:BH:157:TYR:N | 2.23 | 0.71 |
| 39:BN:18:ALA:HB1 | 39:BN:21:LYS:CG | 2.20 | 0.71 |
| 41:BP:108:LYS:C | 41:BP:110:TYR:H | 1.96 | 0.71 |
| 1:CA:1203:C:OP1 | 14:CN:3:ARG:HD3 | 1.90 | 0.71 |
| 31:DA:779:U:OP1 | 33:DD:49:ILE:HG22 | 1.90 | 0.71 |
| 31:DA:1210:A:H5' | 31:DA:1212:G:H5' | 1.73 | 0.71 |
| 31:DA:2845:G:O2' | 31:DA:2846:G:H5' | 1.89 | 0.71 |
| 32:DB:75:G:H5' | 32:DB:75:G:H8 | 1.56 | 0.71 |
| 35:DF:164:ARG:HH11 | 35:DF:164:ARG:HG2 | 1.54 | 0.71 |
| 41:DP:64:LYS:O | 41:DP:66:GLY:N | 2.23 | 0.71 |
| 50:DY:75:ILE:HD11 | 50:DY:79:CYS:C | 2.15 | 0.71 |
| 23:B1:85:LEU:CB | 23:B1:87:PRO:HD3 | 2.20 | 0.71 |
| 24:B2:25:VAL:HG13 | 24:B2:26:ARG:CD | 2.20 | 0.71 |
| 24:B2:31:GLU:HG2 | 24:B2:37:PHE:HD1 | 1.55 | 0.71 |
| 28:B6:20:ASN:O | 28:B6:21:TYR:CG | 2.42 | 0.71 |
| 33:BD:58:HIS:CD2 | 33:BD:59:LYS:N | 2.59 | 0.71 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 37:BH:156:ALA:C | 37:BH:158:HIS:N | 2.49 | 0.71 |
| 42:BQ:141:GLN:NE2 | 51:BZ:72:ARG:HG2 | 2.05 | 0.71 |
| 44:BS:61:ASN:HD22 | 44:BS:62:LYS:N | 1.81 | 0.71 |
| 4:CD:138:TYR:C | 4:CD:138:TYR:CD2 | 2.69 | 0.71 |
| 10:CJ:6:ILE:HG13 | 10:CJ:72:VAL:O | 1.90 | 0.71 |
| 22:D0:8:GLY:HA3 | 31:DA:2255:G:H21 | 1.54 | 0.71 |
| 24:D2:15:LYS:O | 24:D2:16:LEU:CB | 2.39 | 0.71 |
| 31:DA:271(L):U:H4' | 31:DA:271(M):G:C5 | 2.25 | 0.71 |
| 31:DA:614(C):A:H4' | 31:DA:615:G:OP1 | 1.90 | 0.71 |
| 31:DA:1484:G:H21 | 31:DA:1505:C:H41 | 0.80 | 0.71 |
| 41:DP:120:ALA:HB1 | 41:DP:138:LEU:HB3 | 1.72 | 0.71 |
| 1:AA:826:C:H2' | 1:AA:827:U:H6 | 1.53 | 0.71 |
| 1:AA:976:G:H5' | 1:AA:1358:U:O2' | 1.89 | 0.71 |
| 1:AA:1256:A:N6 | 1:AA:1278:U:H1' | 2.03 | 0.71 |
| 1:AA:1412:C:H2' | 1:AA:1413:A:C8 | 2.25 | 0.71 |
| 4:AD:108:LEU:HD11 | 4:AD:174:LEU:HD22 | 1.72 | 0.71 |
| 5:AE:10:MET:CB | 5:AE:32:VAL:HG22 | 2.20 | 0.71 |
| 13:AM:25:ILE:HD11 | 13:AM:66:LEU:HD23 | 1.70 | 0.71 |
| 31:BA:1181:C:O2' | 31:BA:1182:A:H5' | 1.90 | 0.71 |
| 31:BA:2787:C:H1' | 34:BE:61:ARG:HD3 | 1.71 | 0.71 |
| 33:BD:175:LEU:HD12 | 33:BD:185:VAL:CG2 | 2.20 | 0.71 |
| 38:BI:133:HIS:ND1 | 38:BI:134:PRO:HD2 | 2.04 | 0.71 |
| 39:BN:126:PRO:O | 39:BN:127:ASP:HB2 | 1.90 | 0.71 |
| 41:BP:30:THR:HG22 | 41:BP:31:ALA:H | 1.54 | 0.71 |
| 41:BP:47:ASP:HB3 | 41:BP:48:PRO:O | 1.90 | 0.71 |
| 42:BQ:89:ASN:O | 42:BQ:91:GLU:N | 2.23 | 0.71 |
| 1:CA:1446:U:O2' | 1:CA:1447:A:C8 | 2.43 | 0.71 |
| 28:D6:14:THR:O | 28:D6:49:HIS:HA | 1.89 | 0.71 |
| 30:D8:59:LYS:HD3 | 41:DP:50:ARG:HB3 | 1.72 | 0.71 |
| 31:DA:142:A:H8 | 31:DA:1595:G:H21 | 1.36 | 0.71 |
| 31:DA:669:G:H4' | 31:DA:670:A:OP2 | 1.88 | 0.71 |
| 31:DA:684:G:C2 | 31:DA:774:A:C2 | 2.78 | 0.71 |
| 31:DA:1448:G:H1' | 31:DA:1528:A:H62 | 1.52 | 0.71 |
| 38:DI:51:ILE:O | 38:DI:53:ALA:N | 2.23 | 0.71 |
| 45:DT:28:VAL:HG21 | 45:DT:46:GLU:HG3 | 1.71 | 0.71 |
| 45:DT:128:GLU:C | 45:DT:130:ALA:H | 1.99 | 0.71 |
| 4:AD:138:TYR:HD2 | 4:AD:138:TYR:C | 1.97 | 0.71 |
| 12:AL:25:PRO:O | 12:AL:27:LEU:HD22 | 1.90 | 0.71 |
| 31:BA:1681:G:H8 | 31:BA:1681:G:OP2 | 1.73 | 0.71 |
| 35:BF:164:ARG:HH11 | 35:BF:164:ARG:HG2 | 1.54 | 0.71 |
| 1:CA:163:C:H2' | 1:CA:164:U:C6 | 2.26 | 0.71 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 10:CJ:75:ILE:HG13 | 10:CJ:76:ASN:N | 2.06 | 0.71 |
| 31:DA:631:A:O2' | 41:DP:67:MET:HB3 | 1.91 | 0.71 |
| 31:DA:1410:G:H2' | 31:DA:1411:C:H6 | 1.55 | 0.71 |
| 31:DA:2506:U:H5' | 31:DA:2506:U:C6 | 2.25 | 0.71 |
| 31:DA:2758:A:H2' | 31:DA:2759:G:H5" | 1.71 | 0.71 |
| 33:DD:96:HIS:CE1 | 33:DD:102:LYS:HE2 | 2.25 | 0.71 |
| 41:DP:97:PRO:O | 41:DP:98:GLU:HB3 | 1.89 | 0.71 |
| 1:AA:377:G:O2' | 1:AA:378:G:H5' | 1.89 | 0.71 |
| 1:AA:542:G:H2' | 1:AA:543:C:H6 | 1.56 | 0.71 |
| 1:AA:775:G:C2' | 1:AA:776:G:H5' | 2.20 | 0.71 |
| 9:AI:114:TYR:HD2 | 9:AI:114:TYR:H | 1.39 | 0.71 |
| 23:B1:8:SER:N | 23:B1:46:LEU:HD11 | 2.05 | 0.71 |
| 24:B2:32:LEU:HD21 | 31:BA:61:G:HO2' | 1.55 | 0.71 |
| 33:BD:27:THR:HG23 | 33:BD:28:GLU:N | 2.05 | 0.71 |
| 44:BS:28:VAL:HG12 | 44:BS:29:PHE:N | 2.05 | 0.71 |
| 47:BV:70:ILE:HG13 | 47:BV:90:PRO:HB2 | 1.71 | 0.71 |
| 51:BZ:73:GLN:HG2 | 51:BZ:87:ASP:OD1 | 1.90 | 0.71 |
| 51:BZ:165:VAL:HG12 | 51:BZ:166:SER:N | 2.05 | 0.71 |
| 1:CA:1072:G:H2' | 1:CA:1073:U:C6 | 2.25 | 0.71 |
| 31:DA:1019:U:HO2' | 31:DA:1021:A:H2 | 1.36 | 0.71 |
| 31:DA:1109:C:H5 | 31:DA:1110:G:C5 | 2.07 | 0.71 |
| 31:DA:1317:A:H2' | 31:DA:1318:C:H6 | 1.56 | 0.71 |
| 31:DA:1504:C:O2' | 31:DA:1505:C:C5' | 2.38 | 0.71 |
| 31:DA:2580:U:H5" | 34:DE:131:ALA:H | 1.55 | 0.71 |
| 42:DQ:89:ASN:O | 42:DQ:91:GLU:N | 2.24 | 0.71 |
| 42:DQ:140:ALA:HA | 51:DZ:99:TYR:CD2 | 2.26 | 0.71 |
| 47:DV:1:MET:HE2 | 47:DV:1:MET:HA | 1.72 | 0.71 |
| 47:DV:19:LYS:HB3 | 47:DV:96:ILE:O | 1.91 | 0.71 |
| 1:AA:503:C:OP2 | 12:AL:116:SER:HB3 | 1.90 | 0.71 |
| 1:AA:659:U:C2' | 1:AA:660:G:H5' | 2.21 | 0.71 |
| 1:AA:1057:G:H5" | 3:AC:154:SER:HB2 | 1.71 | 0.71 |
| 16:AP:4:ILE:HG13 | 16:AP:21:VAL:CG1 | 2.18 | 0.71 |
| 25:B3:52:HIS:CD2 | 32:BB:83:G:H4' | 2.25 | 0.71 |
| 31:BA:1614:A:N6 | 48:BW:88:ARG:H | 1.88 | 0.71 |
| 39:BN:19:GLU:HG3 | 39:BN:20:GLY:H | 1.55 | 0.71 |
| 39:BN:78:TYR:H | 39:BN:79:PRO:HD3 | 1.56 | 0.71 |
| 42:BQ:6:ARG:O | 42:BQ:7:MET:HG2 | 1.90 | 0.71 |
| 42:BQ:140:ALA:HA | 51:BZ:99:TYR:CD2 | 2.25 | 0.71 |
| 1:CA:1172:C:H2' | 1:CA:1173:G:C8 | 2.26 | 0.71 |
| 13:CM:25:ILE:HD11 | 13:CM:66:LEU:HD23 | 1.71 | 0.71 |
| 31:DA:626:U:N3 | 41:DP:105:LEU:HG | 2.06 | 0.71 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1614:A:N6 | 48:DW:88:ARG:H | 1.87 | 0.71 |
| 47:DV:70:ILE:HG13 | 47:DV:90:PRO:HB2 | 1.71 | 0.71 |
| 51:DZ:27:VAL:HG23 | 51:DZ:36:LYS:HA | 1.71 | 0.71 |
| 1:AA:658:G:H2' | 1:AA:659:U:H6 | 1.55 | 0.71 |
| 22:B0:8:GLY:HA3 | 31:BA:2255:G:H21 | 1.55 | 0.71 |
| 28:B6:15:GLU:HG2 | 28:B6:18:ARG:NE | 2.06 | 0.71 |
| 31:BA:953:A:C2' | 31:BA:954:G:H5' | 2.20 | 0.71 |
| 31:BA:1528(A):A:N7 | 31:BA:1529:G:C8 | 2.58 | 0.71 |
| 31:BA:2359:C:C2' | 31:BA:2360:A:H5' | 2.20 | 0.71 |
| 46:BU:74:LEU:HD12 | 46:BU:74:LEU:N | 2.06 | 0.71 |
| 47:BV:1:MET:HE2 | 47:BV:1:MET:HA | 1.73 | 0.71 |
| 1:CA:428:G:H4' | 1:CA:429:U:O5' | 1.90 | 0.71 |
| 3:CC:50:ALA:HB1 | 3:CC:70:VAL:HG11 | 1.73 | 0.71 |
| 6:CF:46:ARG:HH12 | 18:CR:37:VAL:HG21 | 1.54 | 0.71 |
| 30:D8:52:LYS:N | 30:D8:53:PRO:CD | 2.52 | 0.71 |
| 31:DA:329:G:H4' | 31:DA:330:A:OP2 | 1.91 | 0.71 |
| 31:DA:848:G:H2' | 31:DA:849:A:C8 | 2.25 | 0.71 |
| 33:DD:87:ASN:N | 33:DD:87:ASN:ND2 | 2.34 | 0.71 |
| 47:DV:90:PRO:HG2 | 47:DV:91:TYR:H | 1.55 | 0.71 |
| 1:AA:491:G:H2' | 1:AA:492:G:C8 | 2.25 | 0.71 |
| 1:AA:1446:U:O2' | 1:AA:1447:A:C8 | 2.44 | 0.71 |
| 23:B1:85:LEU:HB3 | 23:B1:87:PRO:CD | 2.20 | 0.71 |
| 31:BA:184:C:H2' | 31:BA:185:U:C6 | 2.26 | 0.71 |
| 31:BA:1173:G:H3' | 31:BA:1174:A:H5' | 1.73 | 0.71 |
| 31:BA:1493:C:H5 | 31:BA:2206:G:O2' | 1.74 | 0.71 |
| 34:BE:14:ILE:HG12 | 34:BE:21:VAL:CG2 | 2.21 | 0.71 |
| 35:BF:65:TRP:O | 35:BF:67:GLN:N | 2.24 | 0.71 |
| 36:BG:33:ARG:H | 36:BG:162:THR:HB | 1.56 | 0.71 |
| 38:BI:1:MET:HG3 | 38:BI:23:PRO:HA | 1.73 | 0.71 |
| 39:BN:55:VAL:HG12 | 39:BN:125:GLY:HA3 | 1.73 | 0.71 |
| 47:BV:2:PHE:CE1 | 47:BV:13:ARG:CZ | 2.71 | 0.71 |
| 1:CA:1057:G:H5" | 3:CC:154:SER:HB2 | 1.71 | 0.71 |
| 17:CQ:74:LEU:HD12 | 17:CQ:75:ARG:HG2 | 1.72 | 0.71 |
| 31:DA:484:C:H2' | 31:DA:485:C:C6 | 2.26 | 0.71 |
| 31:DA:827:U:O2' | 31:DA:2068:U:C2 | 2.40 | 0.71 |
| 31:DA:1779:U:C5 | 31:DA:1784:A:N7 | 2.58 | 0.71 |
| 1:AA:741:G:H2' | 1:AA:742:G:O4' | 1.91 | 0.71 |
| 1:AA:1226:C:H42 | 13:AM:104:ARG:HD2 | 1.56 | 0.71 |
| 10:AJ:34:VAL:CG2 | 10:AJ:74:ILE:HG22 | 2.18 | 0.71 |
| 30:B8:32:LEU:C | 30:B8:34:TRP:N | 2.44 | 0.71 |
| 31:BA:1042:G:H5" | 31:BA:1043:C:OP2 | 1.91 | 0.71 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1887:C:C2' | 31:BA:1888:G:H5' | 2.21 | 0.71 |
| 34:BE:152:LYS:HG2 | 39:BN:78:TYR:CD2 | 2.26 | 0.71 |
| 47:BV:18:LEU:HD22 | 47:BV:19:LYS:CA | 2.19 | 0.71 |
| 1:CA:1441:G:H5" | 1:CA:1442:G:O5' | 1.90 | 0.71 |
| 1:CA:1456:G:H2' | 1:CA:1457:G:O4' | 1.91 | 0.71 |
| 27:D5:46:CYS:SG | 27:D5:47:PRO:CD | 2.78 | 0.71 |
| 28:D6:18:ARG:HG3 | 28:D6:19:ARG:H | 1.55 | 0.71 |
| 31:DA:511:U:H5" | 31:DA:512:G:OP2 | 1.90 | 0.71 |
| 31:DA:1359:A:H2' | 31:DA:1360:A:H5' | 1.73 | 0.71 |
| 31:DA:1504:C:O2' | 31:DA:1505:C:H5' | 1.91 | 0.71 |
| 31:DA:2475:C:H6 | 31:DA:2475:C:H5" | 1.56 | 0.71 |
| 33:DD:228:PRO:HD3 | 33:DD:235:GLY:HA3 | 1.71 | 0.71 |
| 34:DE:61:ARG:H | 34:DE:62:PRO:CD | 2.02 | 0.71 |
| 41:DP:57:THR:HB | 41:DP:59:LEU:N | 2.06 | 0.71 |
| 11:AK:111:ASP:HA | 18:AR:84:LYS:HE2 | 1.73 | 0.70 |
| 22:B0:74:ARG:HH22 | 32:BB:13:A:H8 | 1.38 | 0.70 |
| 31:BA:588:U:O4 | 31:BA:670:A:H1' | 1.91 | 0.70 |
| 31:BA:2464:C:O2' | 31:BA:2465:C:P | 2.49 | 0.70 |
| 31:BA:2542:A:H5' | 31:BA:2543:G:OP1 | 1.90 | 0.70 |
| 37:BH:40:GLU:O | 37:BH:41:MET:HB2 | 1.90 | 0.70 |
| 38:BI:101:LEU:HD12 | 38:BI:101:LEU:O | 1.89 | 0.70 |
| 41:BP:36:LYS:O | 41:BP:38:GLN:HG2 | 1.90 | 0.70 |
| 41:BP:95:VAL:HA | 41:BP:99:LEU:HD23 | 1.73 | 0.70 |
| 42:BQ:8:LYS:CD | 42:BQ:9:TYR:H | 2.04 | 0.70 |
| 44:BS:28:VAL:CB | 44:BS:89:ARG:HB2 | 2.07 | 0.70 |
| 46:BU:90:VAL:O | 46:BU:92:ARG:N | 2.22 | 0.70 |
| 50:BY:75:ILE:HD11 | 50:BY:79:CYS:CA | 2.20 | 0.70 |
| 1:CA:328:C:H2' | 1:CA:328:C:O2 | 1.90 | 0.70 |
| 35:DF:198:ALA:O | 35:DF:201:VAL:HG12 | 1.91 | 0.70 |
| 35:DF:203:GLN:HA | 35:DF:206:ILE:O | 1.91 | 0.70 |
| 37:DH:88:LEU:O | 37:DH:89:ILE:HG23 | 1.90 | 0.70 |
| 38:DI:88:ILE:HG22 | 38:DI:89:TYR:H | 1.54 | 0.70 |
| 39:DN:45:ASN:H | 39:DN:45:ASN:ND2 | 1.86 | 0.70 |
| 41:DP:29:LYS:H | 41:DP:29:LYS:CD | 1.97 | 0.70 |
| 50:DY:7:VAL:HB | 50:DY:8:LYS:NZ | 2.06 | 0.70 |
| 31:BA:806:C:OP2 | 41:BP:39:LYS:HD2 | 1.91 | 0.70 |
| 31:BA:996:A:H4' | 46:BU:92:ARG:HE | 1.56 | 0.70 |
| 31:BA:2476:A:C2 | 31:BA:2477:C:C6 | 2.79 | 0.70 |
| 31:BA:2787:C:C1' | 34:BE:61:ARG:HB2 | 2.20 | 0.70 |
| 38:BI:81:VAL:HG13 | 38:BI:88:ILE:HG23 | 1.73 | 0.70 |
| 41:BP:120:ALA:HB1 | 41:BP:138:LEU:HB3 | 1.72 | 0.70 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 42:BQ:42:ILE:HD13 | 42:BQ:97:VAL:HG21 | 1.72 | 0.70 |
| 22:D0:53:MET:HE3 | 22:D0:57:PHE:CD1 | 2.24 | 0.70 |
| 27:D5:55:ARG:CD | 27:D5:56:LYS:H | 2.03 | 0.70 |
| 31:DA:71:A:H5' | 31:DA:71:A:C8 | 2.26 | 0.70 |
| 31:DA:1042:G:H5" | 31:DA:1043:C:OP2 | 1.92 | 0.70 |
| 31:DA:1209:G:H21 | 31:DA:1210:A:H62 | 1.39 | 0.70 |
| 33:DD:94:LEU:HB2 | 33:DD:104:TYR:CE2 | 2.26 | 0.70 |
| 38:DI:71:ILE:O | 38:DI:75:LEU:HB3 | 1.91 | 0.70 |
| 40:DO:2:ILE:HD12 | 40:DO:6:THR:HG21 | 1.71 | 0.70 |
| 46:DU:95:LEU:CD1 | 47:DV:11:GLN:HG3 | 2.21 | 0.70 |
| 47:DV:13:ARG:HH11 | 47:DV:13:ARG:CG | 1.97 | 0.70 |
| 49:DX:24:GLY:HA3 | 49:DX:80:ILE:CG1 | 2.21 | 0.70 |
| 30:B8:32:LEU:HB3 | 30:B8:34:TRP:HB3 | 1.74 | 0.70 |
| 31:BA:1218:C:H2' | 31:BA:1219:G:H5' | 1.72 | 0.70 |
| 31:BA:1300:U:H3' | 31:BA:1301:A:C5' | 2.21 | 0.70 |
| 38:BI:6:LEU:HD12 | 38:BI:34:GLY:O | 1.91 | 0.70 |
| 38:BI:71:ILE:CG1 | 38:BI:72:LEU:HD22 | 2.21 | 0.70 |
| 47:BV:62:LEU:HA | 47:BV:98:GLU:HA | 1.72 | 0.70 |
| 47:BV:66:ARG:CD | 47:BV:67:GLY:N | 2.54 | 0.70 |
| 1:CA:579:G:H2' | 1:CA:580:U:H6 | 1.57 | 0.70 |
| 1:CA:741:G:H2' | 1:CA:742:G:O4' | 1.91 | 0.70 |
| 25:D3:52:HIS:HD2 | 32:DB:83:G:H4' | 1.54 | 0.70 |
| 31:DA:267:C:H2' | 31:DA:268:C:H6 | 1.56 | 0.70 |
| 31:DA:1509:C:H4' | 31:DA:1509:C:OP1 | 1.91 | 0.70 |
| 33:DD:175:LEU:HD12 | 33:DD:185:VAL:CG2 | 2.21 | 0.70 |
| 35:DF:128:ALA:O | 35:DF:142:TRP:NE1 | 2.25 | 0.70 |
| 41:DP:41:ARG:HA | 41:DP:41:ARG:HH21 | 1.53 | 0.70 |
| 44:DS:33:LYS:HB3 | 44:DS:34:HIS:HD2 | 1.56 | 0.70 |
| 29:B7:8:ASN:C | 29:B7:8:ASN:ND2 | 2.36 | 0.70 |
| 34:BE:203:LYS:O | 34:BE:203:LYS:HD2 | 1.92 | 0.70 |
| 38:BI:1:MET:HB2 | 38:BI:21:VAL:O | 1.90 | 0.70 |
| 39:BN:131:GLN:HG2 | 39:BN:133:GLN:O | 1.92 | 0.70 |
| 41:BP:48:PRO:O | 41:BP:49:ARG:C | 2.32 | 0.70 |
| 46:BU:65:ILE:HG12 | 46:BU:96:ALA:CB | 2.21 | 0.70 |
| 3:CC:9:GLY:HA2 | 3:CC:12:LEU:HD23 | 1.74 | 0.70 |
| 9:CI:114:TYR:HD2 | 9:CI:114:TYR:H | 1.36 | 0.70 |
| 25:D3:52:HIS:CD2 | 32:DB:83:G:H5" | 2.26 | 0.70 |
| 31:DA:676:A:H2 | 31:DA:802:A:H61 | 1.39 | 0.70 |
| 41:DP:47:ASP:HB3 | 41:DP:48:PRO:O | 1.92 | 0.70 |
| 45:DT:35:LYS:O | 45:DT:37:GLY:N | 2.25 | 0.70 |
| 1:AA:52:G:C2' | 1:AA:53:A:H5' | 2.20 | 0.70 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 6:AF:46:ARG:HH12 | 18:AR:37:VAL:HG21 | 1.57 | 0.70 |
| 7:AG:73:MET:HA | 7:AG:91:VAL:HG23 | 1.73 | 0.70 |
| 15:AO:82:ILE:HG12 | 15:AO:87:ILE:HB | 1.74 | 0.70 |
| 24:B2:14:ARG:CZ | 24:B2:57:ILE:HG22 | 2.21 | 0.70 |
| 31:BA:329:G:OP2 | 50:BY:71:LYS:HE3 | 1.91 | 0.70 |
| 31:BA:2488:A:H2' | 31:BA:2489:G:O4' | 1.91 | 0.70 |
| 37:BH:70:THR:HG22 | 37:BH:74:ASN:ND2 | 2.06 | 0.70 |
| 38:BI:71:ILE:O | 38:BI:75:LEU:HB3 | 1.90 | 0.70 |
| 41:BP:131:SER:C | 41:BP:133:SER:H | 1.99 | 0.70 |
| 41:BP:146:VAL:HG22 | 41:BP:147:LEU:N | 2.07 | 0.70 |
| 44:BS:95:HIS:CG | 44:BS:96:GLY:H | 2.08 | 0.70 |
| 31:DA:579:G:H2' | 31:DA:580:C:C6 | 2.27 | 0.70 |
| 33:DD:255:LYS:H | 33:DD:255:LYS:CE | 2.00 | 0.70 |
| 41:DP:16:ARG:CD | 41:DP:18:ARG:HB2 | 2.22 | 0.70 |
| 48:DW:54:ALA:HB1 | 48:DW:107:LEU:HD22 | 1.71 | 0.70 |
| 51:DZ:165:VAL:HG12 | 51:DZ:166:SER:N | 2.06 | 0.70 |
| 1:AA:1172:C:H2' | 1:AA:1173:G:C8 | 2.26 | 0.70 |
| 2:AB:204:ASN:ND2 | 2:AB:206:ASP:H | 1.90 | 0.70 |
| 6:AF:67:MET:HB2 | 6:AF:68:PRO:HD2 | 1.73 | 0.70 |
| 8:AH:5:PRO:HB2 | 8:AH:32:LYS:HE2 | 1.74 | 0.70 |
| 31:BA:631:A:O2' | 41:BP:67:MET:HB3 | 1.91 | 0.70 |
| 31:BA:1509:C:OP1 | 31:BA:1509:C:H4' | 1.91 | 0.70 |
| 36:BG:76:SER:HB2 | 36:BG:83:ARG:HB3 | 1.73 | 0.70 |
| 39:BN:82:LEU:HD12 | 39:BN:82:LEU:H | 1.54 | 0.70 |
| 41:BP:144:GLU:N | 41:BP:145:PRO:HD3 | 2.06 | 0.70 |
| 46:BU:64:ARG:CZ | 46:BU:64:ARG:CA | 2.66 | 0.70 |
| 47:BV:82:ARG:O | 47:BV:82:ARG:HD3 | 1.92 | 0.70 |
| 1:CA:377:G:O2' | 1:CA:378:G:H5' | 1.91 | 0.70 |
| 4:CD:141:ARG:HB3 | 4:CD:142:PRO:CD | 2.22 | 0.70 |
| 25:D3:6:VAL:HG13 | 25:D3:54:VAL:CG1 | 2.21 | 0.70 |
| 27:D5:57:VAL:HB | 27:D5:58:LEU:HD12 | 1.74 | 0.70 |
| 29:D7:40:TRP:CD2 | 31:DA:459:U:H5" | 2.27 | 0.70 |
| 30:D8:25:MET:HG3 | 41:DP:64:LYS:CB | 2.21 | 0.70 |
| 31:DA:1713:U:O2' | 31:DA:1714:G:H5' | 1.91 | 0.70 |
| 34:DE:36:ARG:NH2 | 34:DE:88:GLY:CA | 2.54 | 0.70 |
| 35:DF:175:THR:O | 35:DF:176:LEU:HB2 | 1.91 | 0.70 |
| 46:DU:90:VAL:HG13 | 47:DV:39:LEU:HG | 1.74 | 0.70 |
| 31:BA:780:G:H21 | 31:BA:783:A:H62 | 1.40 | 0.70 |
| 31:BA:2287:A:N6 | 31:BA:2344:U:H3 | 1.89 | 0.70 |
| 31:BA:2287:A:H62 | 31:BA:2344:U:H3 | 1.40 | 0.70 |
| 35:BF:67:GLN:O | 35:BF:67:GLN:CG | 2.38 | 0.70 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:559:A:H4' | 1:CA:560:U:C5' | 2.22 | 0.70 |
| 1:CA:659:U:C2' | 1:CA:660:G:H5' | 2.22 | 0.70 |
| 24:D2:32:LEU:HD21 | 31:DA:61:G:HO2' | 1.57 | 0.70 |
| 31:DA:1658:C:OP1 | 34:DE:132:HIS:ND1 | 2.24 | 0.70 |
| 32:DB:15:A:H1' | 32:DB:110:G:C8 | 2.26 | 0.70 |
| 39:DN:78:TYR:H | 39:DN:79:PRO:CD | 2.05 | 0.70 |
| 45:DT:32:TYR:HB3 | 45:DT:81:PRO:O | 1.91 | 0.70 |
| 46:DU:112:ARG:HH11 | 46:DU:112:ARG:HG2 | 1.55 | 0.70 |
| 1:AA:1423:G:H5' | 40:BO:49:ARG:NH2 | 2.05 | 0.70 |
| 24:B2:26:ARG:HA | 24:B2:29:LYS:HE3 | 1.74 | 0.70 |
| 31:BA:2683:C:OP1 | 45:BT:53:ARG:NH2 | 2.24 | 0.70 |
| 33:BD:35:LYS:HE3 | 33:BD:64:ILE:C | 2.17 | 0.70 |
| 33:BD:70:TRP:CH2 | 33:BD:150:LYS:HA | 2.27 | 0.70 |
| 33:BD:166:GLN:HE21 | 33:BD:166:GLN:CA | 2.05 | 0.70 |
| 41:BP:10:PRO:HD2 | 41:BP:11:GLY:H | 1.56 | 0.70 |
| 25:D3:52:HIS:CD2 | 32:DB:83:G:H4' | 2.26 | 0.70 |
| 1:AA:163:C:H2' | 1:AA:164:U:C6 | 2.25 | 0.70 |
| 3:AC:9:GLY:HA2 | 3:AC:12:LEU:HD23 | 1.74 | 0.70 |
| 4:AD:57:ARG:NH2 | 5:AE:107:ARG:HD3 | 2.07 | 0.70 |
| 11:AK:69:ALA:HB1 | 11:AK:103:LEU:HD23 | 1.74 | 0.70 |
| 31:BA:494:G:O2' | 31:BA:495:G:H5' | 1.90 | 0.70 |
| 31:BA:779:U:OP1 | 33:BD:49:ILE:HG22 | 1.91 | 0.70 |
| 31:BA:1503:U:C4 | 31:BA:1504:C:N4 | 2.60 | 0.70 |
| 31:BA:1504:C:O2' | 31:BA:1505:C:C5' | 2.40 | 0.70 |
| 31:BA:2223:G:H2' | 31:BA:2224:G:C5' | 2.22 | 0.70 |
| 31:BA:2360:A:O2' | 31:BA:2361:A:P | 2.49 | 0.70 |
| 34:BE:102:VAL:HA | 34:BE:200:GLU:HA | 1.73 | 0.70 |
| 39:BN:18:ALA:O | 39:BN:21:LYS:HB2 | 1.92 | 0.70 |
| 43:BR:5:LYS:HD2 | 43:BR:5:LYS:H | 1.54 | 0.70 |
| 45:BT:58:ASN:C | 45:BT:58:ASN:HD22 | 2.00 | 0.70 |
| 47:BV:83:ARG:HH11 | 47:BV:83:ARG:HG3 | 1.57 | 0.70 |
| 50:BY:17:SER:HA | 50:BY:71:LYS:HD2 | 1.71 | 0.70 |
| 1:CA:735:C:O2' | 1:CA:736:C:H5' | 1.91 | 0.70 |
| 1:CA:1274:G:N2 | 1:CA:1275:A:H62 | 1.88 | 0.70 |
| 5:CE:80:ILE:HD11 | 5:CE:138:ALA:HB1 | 1.73 | 0.70 |
| 31:DA:1116:C:H2' | 31:DA:1117:G:H5' | 1.73 | 0.70 |
| 31:DA:1418:G:OP1 | 31:DA:1588:C:O2' | 2.08 | 0.70 |
| 33:DD:25:THR:HG21 | 33:DD:81:ALA:CB | 2.21 | 0.70 |
| 36:DG:18:GLU:O | 36:DG:22:ARG:HB2 | 1.91 | 0.70 |
| 38:DI:1:MET:HG3 | 38:DI:23:PRO:HA | 1.73 | 0.70 |
| 44:DS:89:ARG:O | 44:DS:92:TYR:CB | 2.35 | 0.70 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 45:DT:42:ILE:HD13 | 45:DT:83:ILE:HD11 | 1.73 | 0.70 |
| 50:DY:75:ILE:HD11 | 50:DY:79:CYS:CA | 2.22 | 0.70 |
| 30:B8:6:THR:HG22 | 30:B8:63:PRO:HD3 | 1.73 | 0.70 |
| 30:B8:58:ILE:HG22 | 41:BP:49:ARG:HD2 | 1.72 | 0.70 |
| 31:BA:1410:G:H2' | 31:BA:1411:C:H6 | 1.57 | 0.70 |
| 50:BY:28:LYS:HA | 50:BY:39:VAL:H | 1.55 | 0.70 |
| 34:DE:117:MET:O | 34:DE:118:LYS:HB2 | 1.91 | 0.70 |
| 37:DH:155:SER:O | 37:DH:157:TYR:N | 2.25 | 0.70 |
| 38:DI:71:ILE:CG1 | 38:DI:72:LEU:HD22 | 2.21 | 0.70 |
| 49:DX:33:LYS:C | 49:DX:35:THR:N | 2.46 | 0.70 |
| 1:AA:1274:G:N2 | 1:AA:1275:A:H62 | 1.90 | 0.69 |
| 31:BA:1116:C:H2' | 31:BA:1117:G:H5' | 1.73 | 0.69 |
| 40:BO:32:TYR:N | 40:BO:32:TYR:CD1 | 2.58 | 0.69 |
| 1:CA:503:C:OP2 | 12:CL:116:SER:HB3 | 1.91 | 0.69 |
| 8:CH:86:ILE:HG22 | 8:CH:87:SER:N | 2.06 | 0.69 |
| 23:D1:73:LEU:HD13 | 23:D1:90:ILE:HG22 | 1.73 | 0.69 |
| 31:DA:83:G:N2 | 31:DA:102:G:O2' | 2.22 | 0.69 |
| 31:DA:587:C:C4' | 31:DA:588:U:OP2 | 2.40 | 0.69 |
| 36:DG:161:THR:HG22 | 36:DG:163:ALA:H | 1.57 | 0.69 |
| 38:DI:51:ILE:O | 38:DI:51:ILE:HG22 | 1.91 | 0.69 |
| 49:DX:82:GLN:C | 49:DX:85:PRO:HD2 | 2.15 | 0.69 |
| 2:AB:61:LEU:CD2 | 2:AB:68:ILE:HD11 | 2.22 | 0.69 |
| 23:B1:10:LYS:CD | 23:B1:14:VAL:HA | 2.22 | 0.69 |
| 31:BA:676:A:H2 | 31:BA:802:A:H61 | 1.40 | 0.69 |
| 31:BA:2307:G:N2 | 31:BA:2308:G:H5' | 2.07 | 0.69 |
| 33:BD:145:VAL:HG12 | 33:BD:146:GLU:O | 1.92 | 0.69 |
| 35:BF:7:TYR:HD1 | 35:BF:8:GLN:H | 1.41 | 0.69 |
| 37:BH:30:LYS:NZ | 37:BH:81:GLU:HA | 2.07 | 0.69 |
| 43:BR:5:LYS:H | 43:BR:5:LYS:CD | 2.05 | 0.69 |
| 29:D7:8:ASN:ND2 | 29:D7:11:LYS:N | 2.39 | 0.69 |
| 31:DA:481:G:O2' | 31:DA:482:A:OP2 | 2.08 | 0.69 |
| 45:DT:29:ARG:HB3 | 45:DT:85:LYS:HA | 1.72 | 0.69 |
| 50:DY:17:SER:HA | 50:DY:71:LYS:HD2 | 1.73 | 0.69 |
| 1:AA:671:G:H2' | 1:AA:672:U:H6 | 1.57 | 0.69 |
| 5:AE:68:GLU:O | 5:AE:70:PRO:HD3 | 1.92 | 0.69 |
| 6:AF:69:GLU:O | 6:AF:72:VAL:HG12 | 1.92 | 0.69 |
| 27:B5:55:ARG:CD | 27:B5:56:LYS:H | 2.05 | 0.69 |
| 27:B5:57:VAL:HB | 27:B5:58:LEU:HD12 | 1.73 | 0.69 |
| 31:BA:825:C:H2' | 31:BA:826:U:O5' | 1.92 | 0.69 |
| 31:BA:1488:G:C6 | 31:BA:1489:U:N3 | 2.58 | 0.69 |
| 35:BF:128:ALA:O | 35:BF:142:TRP:NE1 | 2.24 | 0.69 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 38:BI:61:ARG:O | 38:BI:133:HIS:HE1 | 1.73 | 0.69 |
| 1:CA:626:U:H2' | 1:CA:627:G:C8 | 2.27 | 0.69 |
| 1:CA:775:G:O2' | 1:CA:776:G:H5' | 1.92 | 0.69 |
| 5:CE:10:MET:CB | 5:CE:32:VAL:HG22 | 2.23 | 0.69 |
| 30:D8:31:HIS:CD2 | 31:DA:2419:U:O4 | 2.45 | 0.69 |
| 31:DA:1109:C:C5 | 31:DA:1110:G:C4 | 2.80 | 0.69 |
| 31:DA:1688:U:H1' | 31:DA:1701:A:C6 | 2.27 | 0.69 |
| 33:DD:228:PRO:CD | 33:DD:235:GLY:HA3 | 2.21 | 0.69 |
| 36:DG:76:SER:HB2 | 36:DG:83:ARG:HB3 | 1.73 | 0.69 |
| 1:AA:1228:C:H2' | 1:AA:1229:A:H8 | 1.56 | 0.69 |
| 23:B1:47:GLN:HB2 | 31:BA:397:G:H5" | 1.73 | 0.69 |
| 31:BA:1744:C:C2' | 31:BA:1745:C:H5' | 2.21 | 0.69 |
| 31:BA:1899:G:N2 | 31:BA:1902:C:C5 | 2.61 | 0.69 |
| 33:BD:35:LYS:HZ3 | 33:BD:104:TYR:CB | 2.04 | 0.69 |
| 35:BF:9:ILE:HG12 | 35:BF:14:PRO:C | 2.17 | 0.69 |
| 37:BH:153:LYS:HD3 | 37:BH:153:LYS:H | 1.57 | 0.69 |
| 30:D8:6:THR:HG21 | 31:DA:243:U:OP1 | 1.92 | 0.69 |
| 31:DA:814:C:H41 | 41:DP:27:HIS:CD2 | 2.10 | 0.69 |
| 31:DA:854:G:H2' | 31:DA:855:G:H8 | 1.58 | 0.69 |
| 31:DA:1744:C:C2' | 31:DA:1745:C:H5' | 2.22 | 0.69 |
| 31:DA:2359:C:C2' | 31:DA:2360:A:H5' | 2.21 | 0.69 |
| 31:DA:2565:A:H5" | 31:DA:2566:A:OP2 | 1.92 | 0.69 |
| 33:DD:27:THR:CG2 | 33:DD:28:GLU:N | 2.54 | 0.69 |
| 42:DQ:89:ASN:O | 42:DQ:92:GLY:N | 2.23 | 0.69 |
| 47:DV:47:VAL:CG1 | 47:DV:48:GLY:H | 2.03 | 0.69 |
| 4:AD:91:SER:HA | 4:AD:94:LEU:HD12 | 1.74 | 0.69 |
| 23:B1:26:ARG:HG2 | 23:B1:34:THR:OG1 | 1.92 | 0.69 |
| 28:B6:16:CYS:O | 28:B6:18:ARG:NH2 | 2.25 | 0.69 |
| 31:BA:684:G:C2 | 31:BA:774:A:C2 | 2.81 | 0.69 |
| 31:BA:1493:C:C5 | 31:BA:2206:G:O2' | 2.44 | 0.69 |
| 31:BA:2335:A:O2' | 31:BA:2336:A:H5" | 1.92 | 0.69 |
| 33:BD:77:ALA:HB2 | 33:BD:97:TYR:CD2 | 2.27 | 0.69 |
| 33:BD:83:GLU:HB2 | 33:BD:92:ILE:CD1 | 2.23 | 0.69 |
| 35:BF:8:GLN:HB3 | 35:BF:126:VAL:HA | 1.73 | 0.69 |
| 36:BG:18:GLU:O | 36:BG:22:ARG:HB2 | 1.93 | 0.69 |
| 5:CE:68:GLU:O | 5:CE:70:PRO:HD3 | 1.93 | 0.69 |
| 8:CH:5:PRO:HB2 | 8:CH:32:LYS:HE2 | 1.73 | 0.69 |
| 8:CH:58:TYR:O | 8:CH:59:LEU:HD23 | 1.92 | 0.69 |
| 24:D2:31:GLU:HG2 | 24:D2:37:PHE:HD1 | 1.58 | 0.69 |
| 31:DA:1300:U:H3' | 31:DA:1301:A:C5' | 2.22 | 0.69 |
| 31:DA:2469:A:O2' | 42:DQ:56:ARG:HG2 | 1.92 | 0.69 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2787:C:H1' | 34:DE:61:ARG:HD3 | 1.73 | 0.69 |
| 33:DD:35:LYS:HE3 | 33:DD:64:ILE:C | 2.18 | 0.69 |
| 41:DP:36:LYS:O | 41:DP:38:GLN:HG2 | 1.93 | 0.69 |
| 41:DP:51:PHE:O | 41:DP:52:GLU:HB2 | 1.93 | 0.69 |
| 41:DP:65:ARG:HB2 | 41:DP:65:ARG:HH11 | 1.57 | 0.69 |
| 41:DP:146:VAL:HG22 | 41:DP:147:LEU:N | 2.08 | 0.69 |
| 46:DU:64:ARG:CZ | 46:DU:64:ARG:CA | 2.66 | 0.69 |
| 47:DV:62:LEU:HD12 | 47:DV:96:ILE:HD13 | 1.74 | 0.69 |
| 1:AA:428:G:H4' | 1:AA:429:U:O5' | 1.93 | 0.69 |
| 24:B2:51:ARG:O | 24:B2:52:ASP:CB | 2.40 | 0.69 |
| 31:BA:2498:C:O2' | 31:BA:2499:C:H5' | 1.92 | 0.69 |
| 36:BG:111:LEU:HA | 36:BG:114:ILE:HG12 | 1.75 | 0.69 |
| 41:BP:65:ARG:HH11 | 41:BP:65:ARG:HB2 | 1.58 | 0.69 |
| 1:CA:503:C:H2' | 1:CA:504:C:H6 | 1.57 | 0.69 |
| 1:CA:542:G:H2' | 1:CA:543:C:H6 | 1.56 | 0.69 |
| 1:CA:930:C:O2' | 1:CA:931:C:H5' | 1.93 | 0.69 |
| 23:D1:34:THR:CG2 | 31:DA:388:G:OP1 | 2.40 | 0.69 |
| 29:D7:5:TRP:CD1 | 29:D7:7:PRO:HG3 | 2.27 | 0.69 |
| 31:DA:875:G:C4' | 51:DZ:170:THR:HG21 | 2.23 | 0.69 |
| 31:DA:1146:C:C2' | 31:DA:1147:C:H5' | 2.22 | 0.69 |
| 31:DA:1173:G:H3' | 31:DA:1174:A:H5' | 1.73 | 0.69 |
| 31:DA:1448:G:H1' | 31:DA:1528:A:N6 | 2.07 | 0.69 |
| 37:DH:89:ILE:O | 37:DH:90:LYS:HG2 | 1.92 | 0.69 |
| 39:DN:42:TRP:CA | 39:DN:48:MET:HE1 | 2.21 | 0.69 |
| 42:DQ:78:PRO:O | 42:DQ:79:LEU:HB2 | 1.93 | 0.69 |
| 50:DY:28:LYS:HA | 50:DY:39:VAL:H | 1.57 | 0.69 |
| 1:AA:930:C:O2' | 1:AA:931:C:H5' | 1.93 | 0.69 |
| 1:AA:1337:G:H5" | 1:AA:1338:G:OP1 | 1.92 | 0.69 |
| 31:BA:267:C:H2' | 31:BA:268:C:H6 | 1.57 | 0.69 |
| 31:BA:896:A:C2 | 31:BA:898:C:H5" | 2.28 | 0.69 |
| 31:BA:1256:G:H5' | 31:BA:1257:C:OP2 | 1.92 | 0.69 |
| 39:BN:27:ALA:HB3 | 39:BN:106:MET:CE | 2.23 | 0.69 |
| 39:BN:42:TRP:CA | 39:BN:48:MET:HE1 | 2.23 | 0.69 |
| 41:BP:124:LYS:HG2 | 41:BP:143:GLY:CA | 2.22 | 0.69 |
| 44:BS:17:ARG:O | 44:BS:18:ILE:HB | 1.90 | 0.69 |
| 45:BT:128:GLU:C | 45:BT:130:ALA:H | 2.00 | 0.69 |
| 49:BX:24:GLY:HA3 | 49:BX:80:ILE:CG1 | 2.23 | 0.69 |
| 50:BY:8:LYS:NZ | 50:BY:74:PRO:HD3 | 2.07 | 0.69 |
| 1:CA:392:G:H2' | 1:CA:393:A:H8 | 1.57 | 0.69 |
| 1:CA:598:U:H4' | 8:CH:94:TYR:CD2 | 2.26 | 0.69 |
| 1:CA:601:C:H2' | 1:CA:602:A:H8 | 1.58 | 0.69 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:1256:A:N6 | 1:CA:1278:U:H1' | 2.03 | 0.69 |
| 2:CB:61:LEU:CD2 | 2:CB:68:ILE:HD11 | 2.22 | 0.69 |
| 30:D8:58:ILE:O | 30:D8:61:LEU:HG | 1.93 | 0.69 |
| 31:DA:588:U:O4 | 31:DA:670:A:H1' | 1.92 | 0.69 |
| 31:DA:2660:A:H5' | 31:DA:2661:G:H21 | 1.56 | 0.69 |
| 33:DD:25:THR:O | 33:DD:27:THR:N | 2.26 | 0.69 |
| 33:DD:70:TRP:CH2 | 33:DD:150:LYS:HA | 2.28 | 0.69 |
| 33:DD:130:ALA:C | 33:DD:131:LEU:HD12 | 2.18 | 0.69 |
| 38:DI:38:LEU:HD12 | 38:DI:38:LEU:H | 1.58 | 0.69 |
| 39:DN:123:TYR:OH | 39:DN:130:HIS:CD2 | 2.44 | 0.69 |
| 41:DP:144:GLU:N | 41:DP:145:PRO:HD3 | 2.07 | 0.69 |
| 47:DV:69:LYS:HG3 | 47:DV:70:ILE:N | 2.08 | 0.69 |
| 2:AB:112:VAL:HG22 | 2:AB:149:LEU:HD13 | 1.72 | 0.69 |
| 3:AC:58:GLU:H | 3:AC:65:ALA:HB3 | 1.57 | 0.69 |
| 15:AO:64:ARG:HH11 | 15:AO:64:ARG:HG3 | 1.58 | 0.69 |
| 18:AR:43:PHE:C | 18:AR:44:LEU:HD12 | 2.18 | 0.69 |
| 29:B7:47:ARG:C | 29:B7:48:LYS:HD3 | 2.18 | 0.69 |
| 31:BA:145:G:C2' | 31:BA:146:G:H5" | 2.20 | 0.69 |
| 31:BA:614(C):A:H4' | 31:BA:615:G:OP1 | 1.92 | 0.69 |
| 31:BA:675:A:C4 | 31:BA:804:A:C2 | 2.81 | 0.69 |
| 31:BA:966:G:H2' | 31:BA:967:C:H6 | 1.58 | 0.69 |
| 31:BA:1108:U:C2' | 31:BA:1109:C:H5' | 2.23 | 0.69 |
| 31:BA:1403:C:C5' | 31:BA:1471:A:H1' | 2.23 | 0.69 |
| 31:BA:1822:G:H8 | 31:BA:1822:G:C5' | 2.06 | 0.69 |
| 33:BD:8:PRO:HB3 | 33:BD:14:ARG:HB3 | 1.75 | 0.69 |
| 33:BD:211:ARG:O | 33:BD:215:LEU:HG | 1.93 | 0.69 |
| 41:BP:97:PRO:O | 41:BP:98:GLU:HB3 | 1.91 | 0.69 |
| 43:BR:33:ARG:HD3 | 43:BR:115:GLU:OE2 | 1.93 | 0.69 |
| 43:BR:67:LEU:HD13 | 43:BR:76:VAL:HG21 | 1.75 | 0.69 |
| 44:BS:33:LYS:HB3 | 44:BS:34:HIS:HD2 | 1.56 | 0.69 |
| 45:BT:28:VAL:HG22 | 45:BT:46:GLU:HA | 1.75 | 0.69 |
| 45:BT:100:TYR:HD2 | 45:BT:103:ARG:HH21 | 1.38 | 0.69 |
| 45:BT:109:GLU:HA | 45:BT:112:ARG:HG3 | 1.74 | 0.69 |
| 50:BY:7:VAL:HB | 50:BY:8:LYS:HD2 | 1.74 | 0.69 |
| 50:BY:81:LYS:CG | 50:BY:96:ILE:HG22 | 2.23 | 0.69 |
| 50:BY:96:ILE:CD1 | 50:BY:99:CYS:SG | 2.79 | 0.69 |
| 1:CA:559:A:H5" | 1:CA:560:U:H3' | 1.74 | 0.69 |
| 1:CA:671:G:H2' | 1:CA:672:U:H6 | 1.58 | 0.69 |
| 1:CA:1442:G:C8 | 1:CA:1442(B):A:C2 | 2.81 | 0.69 |
| 1:CA:1504:G:OP1 | 1:CA:1507:A:H4' | 1.92 | 0.69 |
| 11:CK:24:SER:HB3 | 11:CK:27:ASN:O | 1.92 | 0.69 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 11:CK:34:ASP:HB3 | 11:CK:40:ILE:HD11 | 1.75 | 0.69 |
| 15:CO:78:TYR:O | 15:CO:82:ILE:HG22 | 1.93 | 0.69 |
| 16:CP:53:VAL:HG12 | 16:CP:79:VAL:HG13 | 1.74 | 0.69 |
| 23:D1:10:LYS:CD | 23:D1:14:VAL:HA | 2.23 | 0.69 |
| 23:D1:92:LYS:C | 23:D1:94:LEU:N | 2.47 | 0.69 |
| 28:D6:15:GLU:HB3 | 28:D6:18:ARG:HG2 | 1.72 | 0.69 |
| 31:DA:951:C:O2' | 31:DA:952:G:H5' | 1.92 | 0.69 |
| 31:DA:1022:G:O2' | 31:DA:1023:U:OP2 | 2.06 | 0.69 |
| 31:DA:1181:C:O2' | 31:DA:1182:A:H5' | 1.91 | 0.69 |
| 33:DD:83:GLU:HB2 | 33:DD:92:ILE:CD1 | 2.23 | 0.69 |
| 34:DE:24:THR:HG21 | 34:DE:188:VAL:HG13 | 1.73 | 0.69 |
| 34:DE:117:MET:SD | 34:DE:136:ARG:HB3 | 2.32 | 0.69 |
| 35:DF:9:ILE:HG12 | 35:DF:14:PRO:C | 2.17 | 0.69 |
| 36:DG:33:ARG:H | 36:DG:162:THR:HB | 1.57 | 0.69 |
| 37:DH:156:ALA:C | 37:DH:158:HIS:N | 2.51 | 0.69 |
| 38:DI:94:ALA:HB1 | 38:DI:114:LEU:HD12 | 1.75 | 0.69 |
| 38:DI:138:ILE:H | 38:DI:138:ILE:HD12 | 1.58 | 0.69 |
| 43:DR:5:LYS:HD2 | 43:DR:5:LYS:H | 1.57 | 0.69 |
| 44:DS:14:VAL:HG12 | 44:DS:15:ARG:N | 2.07 | 0.69 |
| 47:DV:69:LYS:HB2 | 47:DV:93:GLU:CD | 2.18 | 0.69 |
| 49:DX:24:GLY:HA3 | 49:DX:80:ILE:HG13 | 1.75 | 0.69 |
| 49:DX:59:VAL:HG23 | 49:DX:74:PRO:HD2 | 1.74 | 0.69 |
| 2:AB:11:LEU:HB3 | 2:AB:213:LEU:HD11 | 1.75 | 0.69 |
| 2:AB:61:LEU:HA | 2:AB:64:ARG:HG2 | 1.75 | 0.69 |
| 18:AR:79:LEU:HD23 | 18:AR:80:PRO:CD | 2.23 | 0.69 |
| 23:B1:34:THR:CG2 | 31:BA:388:G:OP1 | 2.41 | 0.69 |
| 31:BA:141:A:H8 | 31:BA:1408:C:O2' | 1.73 | 0.69 |
| 31:BA:2476:A:C6 | 31:BA:2477:C:C5 | 2.80 | 0.69 |
| 33:BD:63:ARG:HG3 | 33:BD:63:ARG:NH1 | 2.07 | 0.69 |
| 47:BV:73:SER:OG | 47:BV:74:LYS:N | 2.22 | 0.69 |
| 49:BX:33:LYS:C | 49:BX:35:THR:N | 2.42 | 0.69 |
| 1:CA:1226:C:N4 | 13:CM:104:ARG:HD2 | 2.08 | 0.69 |
| 8:CH:1:MET:HE2 | 8:CH:1:MET:N | 2.04 | 0.69 |
| 9:CI:17:VAL:HG11 | 9:CI:81:ILE:HD13 | 1.75 | 0.69 |
| 23:D1:26:ARG:HG2 | 23:D1:34:THR:OG1 | 1.93 | 0.69 |
| 31:DA:272:G:H4' | 31:DA:272(B):G:OP1 | 1.92 | 0.69 |
| 31:DA:518:G:H4' | 48:DW:18:ARG:NH1 | 2.07 | 0.69 |
| 31:DA:2262:U:O2' | 31:DA:2263:C:H5' | 1.91 | 0.69 |
| 31:DA:2598:A:P | 33:DD:236:GLY:HA3 | 2.33 | 0.69 |
| 37:DH:70:THR:HG22 | 37:DH:74:ASN:ND2 | 2.07 | 0.69 |
| 37:DH:153:LYS:HD3 | 37:DH:153:LYS:H | 1.56 | 0.69 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 43:DR:24:GLN:NE2 | 43:DR:36:THR:HG21 | 2.06 | 0.69 |
| 1:AA:1073:U:H2' | 1:AA:1074:G:H8 | 1.58 | 0.69 |
| 1:AA:1228:C:H2' | 1:AA:1229:A:C8 | 2.28 | 0.69 |
| 23:B1:49:VAL:HG11 | 31:BA:2091:U:O2' | 1.92 | 0.69 |
| 23:B1:92:LYS:C | 23:B1:94:LEU:N | 2.50 | 0.69 |
| 31:BA:543:C:C5 | 31:BA:547:A:N7 | 2.60 | 0.69 |
| 31:BA:814:C:H5" | 47:BV:86:GLY:HA3 | 1.73 | 0.69 |
| 37:BH:137:ASP:O | 37:BH:138:LYS:CB | 2.40 | 0.69 |
| 38:BI:108:THR:C | 38:BI:109:ILE:HG13 | 2.17 | 0.69 |
| 38:BI:123:LEU:HD23 | 38:BI:142:VAL:HB | 1.75 | 0.69 |
| 2:CB:204:ASN:ND2 | 2:CB:206:ASP:H | 1.91 | 0.69 |
| 3:CC:58:GLU:H | 3:CC:65:ALA:HB3 | 1.58 | 0.69 |
| 7:CG:116:ALA:O | 7:CG:120:ILE:HG12 | 1.93 | 0.69 |
| 27:D5:16:ARG:NH1 | 27:D5:17:ASP:OD1 | 2.26 | 0.69 |
| 27:D5:50:GLY:O | 27:D5:51:TYR:HD1 | 1.75 | 0.69 |
| 31:DA:1887:C:C2' | 31:DA:1888:G:H5' | 2.23 | 0.69 |
| 31:DA:2286:A:H5" | 31:DA:2287:A:O4' | 1.93 | 0.69 |
| 37:DH:43:VAL:HG11 | 37:DH:53:GLU:H | 1.58 | 0.69 |
| 40:DO:32:TYR:N | 40:DO:32:TYR:CD1 | 2.59 | 0.69 |
| 41:DP:48:PRO:O | 41:DP:49:ARG:C | 2.35 | 0.69 |
| 24:B2:26:ARG:NH1 | 24:B2:29:LYS:HE2 | 2.07 | 0.68 |
| 29:B7:16:HIS:CB | 29:B7:44:PRO:HG2 | 2.22 | 0.68 |
| 31:BA:83:G:H1 | 31:BA:102:G:H2' | 1.58 | 0.68 |
| 31:BA:1590:U:C2' | 31:BA:1591:G:H5" | 2.22 | 0.68 |
| 31:BA:1701:A:H2' | 31:BA:1702:G:H5' | 1.75 | 0.68 |
| 31:BA:2654:A:OP1 | 31:BA:2654:A:H8 | 1.75 | 0.68 |
| 37:BH:43:VAL:HG11 | 37:BH:53:GLU:H | 1.56 | 0.68 |
| 38:BI:71:ILE:HG12 | 38:BI:72:LEU:HD22 | 1.75 | 0.68 |
| 39:BN:78:TYR:H | 39:BN:79:PRO:CD | 2.06 | 0.68 |
| 42:BQ:8:LYS:HD2 | 42:BQ:9:TYR:N | 2.07 | 0.68 |
| 43:BR:10:LEU:HD22 | 43:BR:17:ARG:HD2 | 1.74 | 0.68 |
| 45:BT:82:LEU:HD12 | 45:BT:82:LEU:N | 2.08 | 0.68 |
| 49:BX:63:LYS:HZ2 | 49:BX:70:LEU:CD2 | 2.06 | 0.68 |
| 3:CC:104:GLN:CD | 3:CC:105:GLU:H | 2.00 | 0.68 |
| 31:DA:535:C:C2' | 31:DA:536:A:H5' | 2.23 | 0.68 |
| 31:DA:1694:C:O2' | 31:DA:1695:G:C4 | 2.46 | 0.68 |
| 31:DA:1784:A:H4' | 31:DA:1785:A:C5' | 2.23 | 0.68 |
| 31:DA:2359:C:H2' | 31:DA:2360:A:H5' | 1.74 | 0.68 |
| 31:DA:2360:A:O2' | 31:DA:2361:A:P | 2.51 | 0.68 |
| 33:DD:142:VAL:HG23 | 33:DD:192:THR:C | 2.18 | 0.68 |
| 38:DI:71:ILE:HG12 | 38:DI:72:LEU:HD22 | 1.75 | 0.68 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 38:DI:108:THR:C | 38:DI:109:ILE:HG13 | 2.18 | 0.68 |
| 41:DP:91:PHE:HE2 | 41:DP:95:VAL:HG12 | 1.58 | 0.68 |
| 43:DR:3:HIS:O | 43:DR:4:LEU:HB3 | 1.91 | 0.68 |
| 45:DT:30:VAL:O | 45:DT:30:VAL:HG23 | 1.91 | 0.68 |
| 50:DY:97:ARG:O | 50:DY:97:ARG:HG3 | 1.91 | 0.68 |
| 4:AD:12:CYS:HA | 4:AD:19:LEU:HD12 | 1.74 | 0.68 |
| 7:AG:32:ARG:O | 7:AG:33:ASP:HB2 | 1.92 | 0.68 |
| 31:BA:9:U:C4 | 31:BA:2629:A:N6 | 2.62 | 0.68 |
| 31:BA:953:A:O2' | 31:BA:954:G:H5' | 1.93 | 0.68 |
| 31:BA:2208:A:H1' | 31:BA:2219:G:C5 | 2.28 | 0.68 |
| 35:BF:16:GLY:O | 35:BF:17:ARG:HG3 | 1.93 | 0.68 |
| 35:BF:22:ALA:HA | 35:BF:26:ALA:HB2 | 1.75 | 0.68 |
| 37:BH:158:HIS:NE2 | 37:BH:169:VAL:O | 2.26 | 0.68 |
| 40:BO:98:VAL:HG12 | 40:BO:117:LEU:HB3 | 1.75 | 0.68 |
| 45:BT:30:VAL:HG21 | 45:BT:83:ILE:CG1 | 2.22 | 0.68 |
| 47:BV:90:PRO:CG | 47:BV:91:TYR:H | 2.06 | 0.68 |
| 1:CA:783:C:O2' | 1:CA:784:C:H5' | 1.94 | 0.68 |
| 31:DA:1503:U:C4 | 31:DA:1504:C:N4 | 2.61 | 0.68 |
| 31:DA:2223:G:H2' | 31:DA:2224:G:C5' | 2.22 | 0.68 |
| 39:DN:58:ASP:OD1 | 39:DN:124:ALA:HB1 | 1.93 | 0.68 |
| 44:DS:28:VAL:CB | 44:DS:89:ARG:HB2 | 2.08 | 0.68 |
| 48:DW:88:ARG:HB3 | 48:DW:92:ARG:HB3 | 1.75 | 0.68 |
| 1:AA:1190:G:H3' | 3:AC:3:ASN:HD21 | 1.58 | 0.68 |
| 4:AD:141:ARG:HB3 | 4:AD:142:PRO:CD | 2.23 | 0.68 |
| 8:AH:86:ILE:HG22 | 8:AH:87:SER:N | 2.07 | 0.68 |
| 31:BA:329:G:H22 | 50:BY:19:LYS:NZ | 1.91 | 0.68 |
| 31:BA:721:C:H2' | 31:BA:721:C:O2 | 1.93 | 0.68 |
| 31:BA:1639:U:C2' | 31:BA:1640:C:H5" | 2.23 | 0.68 |
| 35:BF:3:GLU:HB2 | 35:BF:20:LEU:H | 1.58 | 0.68 |
| 50:BY:2:ARG:N | 50:BY:4:LYS:HG2 | 2.09 | 0.68 |
| 1:CA:658:G:H2' | 1:CA:659:U:H6 | 1.57 | 0.68 |
| 24:D2:48:HIS:NE2 | 31:DA:75:G:H4' | 2.08 | 0.68 |
| 31:DA:94:C:H5' | 31:DA:94(A):G:OP2 | 1.93 | 0.68 |
| 31:DA:588:U:H2' | 31:DA:589:C:C6 | 2.29 | 0.68 |
| 31:DA:2801(A):A:H4' | 31:DA:2802:G:H5' | 1.74 | 0.68 |
| 36:DG:148:MET:HA | 36:DG:148:MET:HE3 | 1.76 | 0.68 |
| 41:DP:30:THR:HG22 | 41:DP:31:ALA:N | 2.08 | 0.68 |
| 41:DP:47:ASP:HB3 | 41:DP:48:PRO:CA | 2.22 | 0.68 |
| 45:DT:58:ASN:HD22 | 45:DT:58:ASN:C | 2.02 | 0.68 |
| 45:DT:91:ARG:CB | 45:DT:116:ALA:HA | 2.22 | 0.68 |
| 50:DY:2:ARG:C | 50:DY:4:LYS:H | 2.02 | 0.68 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:1442(A):G:C3' | 1:AA:1442(B):A:H5" | 2.24 | 0.68 |
| 4:AD:172:PRO:HB2 | 4:AD:187:ARG:HH22 | 1.59 | 0.68 |
| 5:AE:78:HIS:HE1 | 5:AE:142:LEU:HA | 1.57 | 0.68 |
| 16:AP:82:GLN:N | 16:AP:82:GLN:HE21 | 1.91 | 0.68 |
| 23:B1:37:ILE:HG21 | 31:BA:2080:G:OP1 | 1.92 | 0.68 |
| 31:BA:1047:G:N2 | 31:BA:1111:A:H62 | 1.92 | 0.68 |
| 34:BE:117:MET:O | 34:BE:118:LYS:HB2 | 1.92 | 0.68 |
| 42:BQ:29:PHE:O | 42:BQ:30:GLY:O | 2.11 | 0.68 |
| 50:BY:7:VAL:HB | 50:BY:8:LYS:HZ2 | 1.58 | 0.68 |
| 1:CA:1073:U:H2' | 1:CA:1074:G:H8 | 1.57 | 0.68 |
| 6:CF:26:ILE:HG22 | 6:CF:30:LEU:HD21 | 1.75 | 0.68 |
| 31:DA:184:C:H2' | 31:DA:185:U:C6 | 2.28 | 0.68 |
| 31:DA:1022:G:N2 | 31:DA:1142(A):A:C2 | 2.61 | 0.68 |
| 31:DA:2315:G:H2' | 31:DA:2316:C:C6 | 2.28 | 0.68 |
| 44:DS:95:HIS:CG | 44:DS:96:GLY:H | 2.11 | 0.68 |
| 45:DT:32:TYR:CG | 45:DT:81:PRO:HB2 | 2.28 | 0.68 |
| 5:AE:100:VAL:HG13 | 5:AE:118:ILE:HG22 | 1.74 | 0.68 |
| 28:B6:18:ARG:CG | 28:B6:19:ARG:H | 2.07 | 0.68 |
| 30:B8:31:HIS:CD2 | 31:BA:2419:U:O4 | 2.47 | 0.68 |
| 30:B8:32:LEU:HD23 | 30:B8:35:GLN:HA | 1.74 | 0.68 |
| 31:BA:1493:C:O2 | 31:BA:1493:C:H2' | 1.92 | 0.68 |
| 31:BA:2777:G:H5" | 31:BA:2778:A:H5' | 1.75 | 0.68 |
| 33:BD:25:THR:O | 33:BD:27:THR:N | 2.27 | 0.68 |
| 33:BD:94:LEU:HB2 | 33:BD:104:TYR:CE2 | 2.27 | 0.68 |
| 36:BG:161:THR:HG22 | 36:BG:163:ALA:H | 1.59 | 0.68 |
| 1:CA:192:U:H2' | 1:CA:193:C:C6 | 2.28 | 0.68 |
| 1:CA:1074:G:C4 | 1:CA:1102:A:C2 | 2.82 | 0.68 |
| 5:CE:78:HIS:HE1 | 5:CE:142:LEU:HA | 1.57 | 0.68 |
| 7:CG:73:MET:HA | 7:CG:91:VAL:HG23 | 1.73 | 0.68 |
| 23:D1:41:ARG:NH1 | 23:D1:41:ARG:CG | 2.47 | 0.68 |
| 24:D2:23:LYS:HB2 | 49:DX:5:TYR:CE1 | 2.28 | 0.68 |
| 31:DA:721:C:H2' | 31:DA:721:C:O2 | 1.93 | 0.68 |
| 31:DA:896:A:C2 | 31:DA:898:C:H5" | 2.29 | 0.68 |
| 31:DA:999:U:O2' | 31:DA:1000:A:H5' | 1.93 | 0.68 |
| 31:DA:1403:C:C5' | 31:DA:1471:A:H1' | 2.21 | 0.68 |
| 31:DA:1493:C:C5 | 31:DA:2206:G:O2' | 2.47 | 0.68 |
| 31:DA:2236:C:C2' | 31:DA:2237:G:H5' | 2.24 | 0.68 |
| 31:DA:2746:U:H4' | 37:DH:138:LYS:HB3 | 1.75 | 0.68 |
| 35:DF:160:ASN:HD22 | 35:DF:160:ASN:C | 2.02 | 0.68 |
| 39:DN:18:ALA:HB1 | 39:DN:21:LYS:CG | 2.22 | 0.68 |
| 45:DT:13:ARG:NH2 | 45:DT:15:VAL:HG13 | 2.08 | 0.68 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 47:DV:2:PHE:CE1 | 47:DV:13:ARG:CZ | 2.75 | 0.68 |
| 13:AM:10:PRO:HB2 | 13:AM:18:ALA:HB1 | 1.74 | 0.68 |
| 15:AO:87:ILE:HG22 | 15:AO:88:ARG:N | 2.08 | 0.68 |
| 30:B8:32:LEU:HB3 | 30:B8:35:GLN:H | 1.57 | 0.68 |
| 31:BA:1406:U:H2' | 31:BA:1407:C:C6 | 2.29 | 0.68 |
| 31:BA:1784:A:H4' | 31:BA:1785:A:C5' | 2.24 | 0.68 |
| 31:BA:2801(A):A:H4' | 31:BA:2802:G:H5' | 1.76 | 0.68 |
| 33:BD:27:THR:CG2 | 33:BD:28:GLU:H | 2.07 | 0.68 |
| 33:BD:130:ALA:C | 33:BD:131:LEU:HD12 | 2.18 | 0.68 |
| 46:BU:25:TRP:CD1 | 46:BU:26:GLY:N | 2.62 | 0.68 |
| 47:BV:36:PRO:HG2 | 47:BV:62:LEU:HD11 | 1.75 | 0.68 |
| 1:CA:60:A:H8 | 1:CA:60:A:P | 2.17 | 0.68 |
| 13:CM:10:PRO:HB2 | 13:CM:18:ALA:HB1 | 1.75 | 0.68 |
| 35:DF:16:GLY:O | 35:DF:17:ARG:HG3 | 1.93 | 0.68 |
| 39:DN:56:ASN:C | 39:DN:57:ALA:O | 2.35 | 0.68 |
| 6:AF:89:MET:HG2 | 6:AF:91:VAL:HG23 | 1.74 | 0.68 |
| 31:BA:286:C:C2' | 31:BA:287:C:H5' | 2.24 | 0.68 |
| 31:BA:288:C:H42 | 31:BA:353:G:H1 | 1.40 | 0.68 |
| 31:BA:1339:G:N2 | 31:BA:1603:A:H1' | 2.08 | 0.68 |
| 31:BA:1396:U:H2' | 31:BA:1396:U:O2 | 1.94 | 0.68 |
| 41:BP:51:PHE:HB3 | 41:BP:52:GLU:OE2 | 1.94 | 0.68 |
| 1:CA:828:A:H2' | 1:CA:829:G:O4' | 1.94 | 0.68 |
| 1:CA:1513:A:H2' | 1:CA:1514:C:C6 | 2.28 | 0.68 |
| 24:D2:51:ARG:O | 24:D2:52:ASP:CB | 2.41 | 0.68 |
| 31:DA:1590:U:C2' | 31:DA:1591:G:H5" | 2.24 | 0.68 |
| 31:DA:1833:U:H2' | 31:DA:1834:U:C6 | 2.27 | 0.68 |
| 46:DU:83:LEU:HG | 46:DU:88:ILE:HG12 | 1.76 | 0.68 |
| 47:DV:69:LYS:CB | 47:DV:93:GLU:CD | 2.67 | 0.68 |
| 50:DY:45:VAL:CG1 | 50:DY:62:GLU:HB2 | 2.19 | 0.68 |
| 1:AA:503:C:H2' | 1:AA:504:C:H6 | 1.58 | 0.68 |
| 29:B7:8:ASN:HD21 | 29:B7:11:LYS:N | 1.85 | 0.68 |
| 31:BA:910:A:H62 | 42:BQ:12:GLN:HA | 1.59 | 0.68 |
| 31:BA:1670:C:O2 | 34:BE:129:HIS:HE1 | 1.77 | 0.68 |
| 31:BA:1694:C:O2' | 31:BA:1695:G:C4 | 2.46 | 0.68 |
| 39:BN:56:ASN:C | 39:BN:57:ALA:O | 2.34 | 0.68 |
| 41:BP:24:GLY:CA | 41:BP:33:ARG:HH21 | 1.96 | 0.68 |
| 46:BU:51:LYS:O | 46:BU:52:ARG:C | 2.36 | 0.68 |
| 1:CA:639:G:O2' | 1:CA:640:A:H5' | 1.93 | 0.68 |
| 1:CA:1190:G:H3' | 3:CC:3:ASN:HD21 | 1.59 | 0.68 |
| 4:CD:12:CYS:HA | 4:CD:19:LEU:HD12 | 1.76 | 0.68 |
| 30:D8:25:MET:HB2 | 41:DP:62:LEU:HD23 | 1.76 | 0.68 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2567:G:H2' | 31:DA:2568:C:C6 | 2.29 | 0.68 |
| 33:DD:166:GLN:HE21 | 33:DD:166:GLN:CA | 2.07 | 0.68 |
| 37:DH:99:VAL:HG12 | 37:DH:99:VAL:O | 1.94 | 0.68 |
| 39:DN:4:TYR:CD1 | 39:DN:4:TYR:N | 2.61 | 0.68 |
| 42:DQ:82:ARG:O | 42:DQ:83:MET:HB3 | 1.92 | 0.68 |
| 1:AA:1074:G:C4 | 1:AA:1102:A:C2 | 2.82 | 0.68 |
| 15:AO:3:ILE:HG12 | 15:AO:3:ILE:O | 1.93 | 0.68 |
| 22:B0:20:ARG:NH1 | 31:BA:2357:U:OP1 | 2.27 | 0.68 |
| 31:BA:1021:A:C8 | 31:BA:1021:A:C3' | 2.77 | 0.68 |
| 31:BA:1332:G:C8 | 31:BA:1332:G:H5' | 2.28 | 0.68 |
| 31:BA:1504:C:O2' | 31:BA:1505:C:H5' | 1.93 | 0.68 |
| 31:BA:2286:A:H5" | 31:BA:2287:A:O4' | 1.94 | 0.68 |
| 37:BH:158:HIS:HE2 | 37:BH:170:ARG:HA | 1.58 | 0.68 |
| 1:CA:52:G:C2' | 1:CA:53:A:H5' | 2.24 | 0.68 |
| 1:CA:450:G:OP1 | 16:CP:43:LYS:NZ | 2.27 | 0.68 |
| 8:CH:97:VAL:O | 8:CH:100:ILE:HG13 | 1.94 | 0.68 |
| 11:CK:20:TYR:C | 11:CK:21:ILE:HD12 | 2.19 | 0.68 |
| 13:CM:92:HIS:CE1 | 13:CM:98:VAL:HG23 | 2.29 | 0.68 |
| 24:D2:56:GLN:NE2 | 24:D2:56:GLN:N | 2.40 | 0.68 |
| 31:DA:329:G:H1 | 50:DY:19:LYS:HD2 | 1.59 | 0.68 |
| 39:DN:27:ALA:HB3 | 39:DN:106:MET:CE | 2.24 | 0.68 |
| 47:DV:82:ARG:HH11 | 47:DV:82:ARG:HG2 | 1.57 | 0.68 |
| 15:AO:33:THR:HG21 | 15:AO:85:LEU:HD22 | 1.76 | 0.68 |
| 31:BA:2208:A:H1' | 31:BA:2219:G:C4 | 2.29 | 0.68 |
| 33:BD:35:LYS:CD | 33:BD:104:TYR:HD1 | 2.07 | 0.68 |
| 41:BP:146:VAL:HG13 | 41:BP:147:LEU:H | 1.58 | 0.68 |
| 46:BU:95:LEU:CD1 | 47:BV:11:GLN:HG3 | 2.24 | 0.68 |
| 1:CA:450:G:H5" | 16:CP:41:PRO:O | 1.94 | 0.68 |
| 2:CB:111:ARG:HH11 | 2:CB:111:ARG:CG | 1.99 | 0.68 |
| 6:CF:21:LEU:O | 6:CF:24:GLU:HB3 | 1.94 | 0.68 |
| 31:DA:286:C:C2' | 31:DA:287:C:H5' | 2.23 | 0.68 |
| 31:DA:518:G:H2' | 31:DA:519:U:C6 | 2.29 | 0.68 |
| 31:DA:996:A:H4' | 46:DU:92:ARG:HE | 1.57 | 0.68 |
| 44:DS:28:VAL:HG12 | 44:DS:29:PHE:N | 2.07 | 0.68 |
| 48:DW:73:ALA:HB3 | 48:DW:106:ILE:HD11 | 1.76 | 0.68 |
| 51:DZ:71:VAL:HG22 | 51:DZ:88:PHE:HE2 | 1.57 | 0.68 |
| 3:AC:104:GLN:CD | 3:AC:105:GLU:H | 2.01 | 0.67 |
| 3:AC:127:ARG:N | 3:AC:127:ARG:HD2 | 2.09 | 0.67 |
| 4:AD:138:TYR:C | 4:AD:138:TYR:CD2 | 2.70 | 0.67 |
| 10:AJ:75:ILE:HG13 | 10:AJ:76:ASN:N | 2.08 | 0.67 |
| 11:AK:29:ILE:HB | 11:AK:44:SER:HB3 | 1.75 | 0.67 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 18:AR:56:THR:OG1 | 18:AR:58:LEU:HD13 | 1.94 | 0.67 |
| 22:B0:43:THR:HG22 | 31:BA:2331:G:O3' | 1.94 | 0.67 |
| 28:B6:51:GLU:O | 28:B6:52:VAL:HB | 1.94 | 0.67 |
| 31:BA:94:C:H5' | 31:BA:94(A):G:OP2 | 1.95 | 0.67 |
| 31:BA:1673:U:O4 | 34:BE:129:HIS:HD2 | 1.76 | 0.67 |
| 31:BA:1836:C:O2' | 31:BA:1837:C:H5' | 1.95 | 0.67 |
| 33:BD:71:ASP:HB3 | 33:BD:103:ARG:NH2 | 2.08 | 0.67 |
| 39:BN:2:LYS:HE2 | 46:BU:95:LEU:HD21 | 1.75 | 0.67 |
| 39:BN:78:TYR:N | 39:BN:79:PRO:HD3 | 2.09 | 0.67 |
| 46:BU:49:HIS:HA | 46:BU:52:ARG:HB2 | 1.75 | 0.67 |
| 1:CA:15:G:H4' | 5:CE:24:ARG:NH1 | 2.08 | 0.67 |
| 4:CD:18:LYS:HE3 | 4:CD:31:CYS:SG | 2.33 | 0.67 |
| 12:CL:87:GLY:HA2 | 12:CL:98:TYR:HA | 1.75 | 0.67 |
| 15:CO:82:ILE:HG12 | 15:CO:87:ILE:HB | 1.76 | 0.67 |
| 23:D1:62:VAL:HG22 | 23:D1:63:ALA:H | 1.59 | 0.67 |
| 31:DA:141:A:H8 | 31:DA:1408:C:O2' | 1.76 | 0.67 |
| 42:DQ:30:GLY:CA | 42:DQ:107:ALA:HB2 | 2.24 | 0.67 |
| 49:DX:89:ILE:HA | 49:DX:92:LEU:HD12 | 1.74 | 0.67 |
| 1:AA:1513:A:H2' | 1:AA:1514:C:C6 | 2.29 | 0.67 |
| 2:AB:165:VAL:HG23 | 2:AB:166:ASP:H | 1.59 | 0.67 |
| 23:B1:30:VAL:O | 23:B1:30:VAL:HG12 | 1.92 | 0.67 |
| 24:B2:15:LYS:O | 24:B2:16:LEU:CB | 2.42 | 0.67 |
| 31:BA:2810:A:H2' | 34:BE:61:ARG:HH21 | 1.58 | 0.67 |
| 33:BD:158:ALA:O | 33:BD:159:ALA:HB2 | 1.94 | 0.67 |
| 34:BE:1:MET:HB2 | 34:BE:84:PHE:HB2 | 1.76 | 0.67 |
| 38:BI:6:LEU:O | 38:BI:15:VAL:HB | 1.94 | 0.67 |
| 39:BN:2:LYS:HE2 | 46:BU:95:LEU:CD2 | 2.25 | 0.67 |
| 42:BQ:17:LEU:HD23 | 42:BQ:17:LEU:N | 2.09 | 0.67 |
| 42:BQ:22:LYS:HA | 42:BQ:22:LYS:CE | 2.24 | 0.67 |
| 47:BV:71:LEU:HD13 | 47:BV:72:VAL:H | 1.58 | 0.67 |
| 1:CA:102:G:H2' | 1:CA:103:C:C6 | 2.27 | 0.67 |
| 1:CA:892:A:H2' | 1:CA:893:C:C6 | 2.30 | 0.67 |
| 1:CA:1442(A):G:C3' | 1:CA:1442(B):A:H5" | 2.24 | 0.67 |
| 27:D5:51:TYR:CD2 | 27:D5:52:TYR:CZ | 2.82 | 0.67 |
| 31:DA:1925:C:O2' | 31:DA:1926:U:H5' | 1.94 | 0.67 |
| 34:DE:1:MET:HB2 | 34:DE:84:PHE:HB2 | 1.76 | 0.67 |
| 42:DQ:38:GLU:OE2 | 42:DQ:127:ILE:HG22 | 1.94 | 0.67 |
| 1:AA:66:G:H4' | 1:AA:173:U:C5 | 2.30 | 0.67 |
| 1:AA:522:C:H41 | 12:AL:53:ARG:NH2 | 1.92 | 0.67 |
| 1:AA:1226:C:N4 | 13:AM:104:ARG:HD2 | 2.10 | 0.67 |
| 3:AC:20:SER:HB2 | 3:AC:40:ARG:NH2 | 2.06 | 0.67 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:272:G:H4' | 31:BA:272(B):G:OP1 | 1.92 | 0.67 |
| 31:BA:579:G:H2' | 31:BA:580:C:C6 | 2.29 | 0.67 |
| 31:BA:631:A:OP1 | 41:BP:64:LYS:HE3 | 1.94 | 0.67 |
| 31:BA:2661:G:N7 | 31:BA:2662:A:C2 | 2.61 | 0.67 |
| 36:BG:25:TYR:CZ | 36:BG:32:PRO:HD3 | 2.30 | 0.67 |
| 37:BH:153:LYS:HD3 | 37:BH:153:LYS:N | 2.09 | 0.67 |
| 38:BI:61:ARG:O | 38:BI:133:HIS:CE1 | 2.48 | 0.67 |
| 2:CB:187:LEU:HD13 | 2:CB:187:LEU:O | 1.95 | 0.67 |
| 24:D2:14:ARG:CZ | 24:D2:57:ILE:HG22 | 2.24 | 0.67 |
| 33:DD:63:ARG:HG3 | 33:DD:63:ARG:NH1 | 2.03 | 0.67 |
| 36:DG:111:LEU:HA | 36:DG:114:ILE:HG12 | 1.75 | 0.67 |
| 39:DN:18:ALA:O | 39:DN:21:LYS:HB2 | 1.94 | 0.67 |
| 41:DP:146:VAL:HG13 | 41:DP:147:LEU:H | 1.57 | 0.67 |
| 46:DU:104:GLN:HB2 | 47:DV:43:GLU:OE1 | 1.94 | 0.67 |
| 50:DY:60:PHE:HA | 50:DY:62:GLU:OE2 | 1.94 | 0.67 |
| 27:B5:40:LYS:CE | 27:B5:46:CYS:HB3 | 2.24 | 0.67 |
| 27:B5:48:GLU:O | 27:B5:50:GLY:N | 2.27 | 0.67 |
| 31:BA:1181:C:C2' | 31:BA:1182:A:H5' | 2.23 | 0.67 |
| 31:BA:1225:G:OP1 | 47:BV:88:ARG:HB3 | 1.95 | 0.67 |
| 33:BD:30:GLU:HG3 | 33:BD:63:ARG:CZ | 2.24 | 0.67 |
| 39:BN:78:TYR:CE1 | 39:BN:79:PRO:HB3 | 2.29 | 0.67 |
| 50:BY:46:LYS:HB2 | 50:BY:47:LYS:HE2 | 1.76 | 0.67 |
| 51:BZ:101:PRO:O | 51:BZ:102:LEU:HD23 | 1.94 | 0.67 |
| 7:CG:32:ARG:O | 7:CG:33:ASP:HB2 | 1.94 | 0.67 |
| 8:CH:77:GLU:HG3 | 8:CH:78:GLN:H | 1.60 | 0.67 |
| 16:CP:45:THR:HG23 | 16:CP:46:PRO:HD2 | 1.75 | 0.67 |
| 23:D1:46:LEU:HD12 | 23:D1:46:LEU:H | 1.59 | 0.67 |
| 31:DA:2055:C:H5' | 31:DA:2056:G:O5' | 1.93 | 0.67 |
| 31:DA:2653:U:H3' | 31:DA:2654:A:H5" | 1.75 | 0.67 |
| 33:DD:17:THR:HG23 | 33:DD:205:VAL:HB | 1.76 | 0.67 |
| 33:DD:43:ARG:HH11 | 33:DD:44:ASN:CG | 2.02 | 0.67 |
| 37:DH:153:LYS:HD3 | 37:DH:153:LYS:N | 2.09 | 0.67 |
| 42:DQ:8:LYS:CG | 42:DQ:9:TYR:H | 2.07 | 0.67 |
| 43:DR:33:ARG:HD3 | 43:DR:115:GLU:OE2 | 1.94 | 0.67 |
| 1:AA:327:A:C4 | 1:AA:329:A:C8 | 2.83 | 0.67 |
| 11:AK:29:ILE:HB | 11:AK:44:SER:CB | 2.25 | 0.67 |
| 12:AL:41:ARG:HG2 | 12:AL:42:THR:H | 1.60 | 0.67 |
| 29:B7:8:ASN:ND2 | 29:B7:11:LYS:N | 2.42 | 0.67 |
| 30:B8:59:LYS:NZ | 30:B8:59:LYS:CB | 2.57 | 0.67 |
| 35:BF:160:ASN:C | 35:BF:160:ASN:HD22 | 2.02 | 0.67 |
| 36:BG:47:LYS:HG3 | 36:BG:82:LEU:HD11 | 1.76 | 0.67 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 43:BR:116:LEU:O | 43:BR:117:VAL:CB | 2.43 | 0.67 |
| 45:BT:42:ILE:O | 45:BT:42:ILE:HG13 | 1.95 | 0.67 |
| 48:BW:86:LEU:C | 48:BW:86:LEU:HD12 | 2.19 | 0.67 |
| 1:CA:509:A:H2' | 1:CA:510:A:C8 | 2.28 | 0.67 |
| 2:CB:112:VAL:HG22 | 2:CB:149:LEU:HD13 | 1.75 | 0.67 |
| 31:DA:543:C:C5 | 31:DA:547:A:N7 | 2.63 | 0.67 |
| 31:DA:2682:U:C6 | 34:DE:11:MET:HE2 | 2.30 | 0.67 |
| 32:DB:7:G:H4' | 44:DS:29:PHE:HD1 | 1.56 | 0.67 |
| 34:DE:95:ILE:HD12 | 34:DE:95:ILE:N | 2.10 | 0.67 |
| 41:DP:131:SER:C | 41:DP:133:SER:H | 2.01 | 0.67 |
| 47:DV:25:LEU:HB2 | 47:DV:94:LEU:HD13 | 1.75 | 0.67 |
| 50:DY:44:ILE:HD12 | 50:DY:44:ILE:H | 1.58 | 0.67 |
| 4:AD:13:ARG:O | 4:AD:15:GLU:N | 2.28 | 0.67 |
| 11:AK:34:ASP:HB2 | 11:AK:35:PRO:HD2 | 1.77 | 0.67 |
| 31:BA:737:C:H2' | 31:BA:738:G:O5' | 1.95 | 0.67 |
| 31:BA:2315:G:H2' | 31:BA:2316:C:C6 | 2.30 | 0.67 |
| 32:BB:44:G:H5" | 32:BB:45:A:OP1 | 1.95 | 0.67 |
| 38:BI:94:ALA:HB1 | 38:BI:114:LEU:HD12 | 1.77 | 0.67 |
| 47:BV:47:VAL:HG13 | 47:BV:48:GLY:N | 2.07 | 0.67 |
| 1:CA:135:C:H2' | 1:CA:136:C:H5' | 1.75 | 0.67 |
| 2:CB:114:ARG:HD2 | 2:CB:141:GLU:OE1 | 1.94 | 0.67 |
| 3:CC:100:ALA:O | 3:CC:101:LEU:HB2 | 1.95 | 0.67 |
| 25:D3:43:ILE:O | 25:D3:47:VAL:HG23 | 1.95 | 0.67 |
| 31:DA:861:A:C2 | 31:DA:917:A:C4 | 2.83 | 0.67 |
| 31:DA:953:A:O2' | 31:DA:954:G:H5' | 1.93 | 0.67 |
| 31:DA:2875:C:O2' | 45:DT:5:ALA:HB3 | 1.95 | 0.67 |
| 39:DN:65:LYS:HA | 39:DN:65:LYS:HE2 | 1.77 | 0.67 |
| 41:DP:10:PRO:CD | 41:DP:11:GLY:N | 2.56 | 0.67 |
| 45:DT:28:VAL:O | 45:DT:29:ARG:HD3 | 1.95 | 0.67 |
| 46:DU:14:HIS:HD2 | 46:DU:32:PHE:CB | 2.08 | 0.67 |
| 2:AB:211:ILE:O | 2:AB:215:LEU:HB2 | 1.95 | 0.67 |
| 31:BA:548:A:O2' | 31:BA:549:G:OP1 | 2.12 | 0.67 |
| 31:BA:1448:G:H1' | 31:BA:1528:A:N6 | 2.10 | 0.67 |
| 31:BA:2203:U:H1' | 33:BD:151:LYS:HE2 | 1.75 | 0.67 |
| 31:BA:2746:U:H4' | 37:BH:138:LYS:HB3 | 1.75 | 0.67 |
| 41:BP:17:LYS:O | 41:BP:17:LYS:HG2 | 1.94 | 0.67 |
| 43:BR:44:LEU:HD22 | 43:BR:48:VAL:HG23 | 1.76 | 0.67 |
| 50:BY:28:LYS:CD | 50:BY:37:VAL:HG12 | 2.23 | 0.67 |
| 6:CF:17:SER:O | 6:CF:21:LEU:HD22 | 1.95 | 0.67 |
| 10:CJ:63:PHE:HB3 | 14:CN:57:ARG:O | 1.94 | 0.67 |
| 15:CO:62:GLN:HA | 15:CO:65:ARG:NH1 | 2.09 | 0.67 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 16:CP:82:GLN:N | 16:CP:82:GLN:HE21 | 1.91 | 0.67 |
| 31:DA:737:C:H2' | 31:DA:738:G:O5' | 1.94 | 0.67 |
| 35:DF:63:LYS:HE2 | 35:DF:67:GLN:CB | 2.25 | 0.67 |
| 36:DG:67:LYS:HD2 | 36:DG:67:LYS:H | 1.59 | 0.67 |
| 37:DH:30:LYS:NZ | 37:DH:81:GLU:HA | 2.09 | 0.67 |
| 50:DY:37:VAL:HG13 | 50:DY:69:ALA:HA | 1.76 | 0.67 |
| 1:AA:102:G:H2' | 1:AA:103:C:C6 | 2.28 | 0.67 |
| 6:AF:21:LEU:O | 6:AF:24:GLU:HB3 | 1.94 | 0.67 |
| 12:AL:83:VAL:HG11 | 12:AL:100:ILE:HG12 | 1.77 | 0.67 |
| 31:BA:825:C:C2' | 31:BA:826:U:O5' | 2.42 | 0.67 |
| 33:BD:223:GLY:HA3 | 33:BD:231:HIS:CE1 | 2.30 | 0.67 |
| 36:BG:7:LEU:HB2 | 36:BG:104:GLU:OE2 | 1.95 | 0.67 |
| 37:BH:158:HIS:NE2 | 37:BH:170:ARG:CA | 2.56 | 0.67 |
| 2:CB:29:ALA:O | 2:CB:32:ILE:HG22 | 1.93 | 0.67 |
| 23:D1:37:ILE:HD11 | 31:DA:2079:U:H4' | 1.76 | 0.67 |
| 30:D8:32:LEU:O | 30:D8:33:ASN:CB | 2.40 | 0.67 |
| 31:DA:854:G:H2' | 31:DA:855:G:C8 | 2.30 | 0.67 |
| 31:DA:2307:G:H21 | 31:DA:2308:G:C5' | 2.07 | 0.67 |
| 32:DB:7:G:H4' | 44:DS:29:PHE:CE1 | 2.29 | 0.67 |
| 32:DB:74:U:C2' | 32:DB:75:G:H5" | 2.25 | 0.67 |
| 33:DD:3:VAL:H | 33:DD:20:ASP:HB2 | 1.58 | 0.67 |
| 35:DF:24:LEU:HB3 | 35:DF:25:PRO:CD | 2.24 | 0.67 |
| 50:DY:7:VAL:HB | 50:DY:8:LYS:HD2 | 1.76 | 0.67 |
| 1:AA:601:C:H2' | 1:AA:602:A:H8 | 1.57 | 0.67 |
| 1:AA:688:G:H2' | 1:AA:689:C:H6 | 1.60 | 0.67 |
| 8:AH:58:TYR:O | 8:AH:59:LEU:HD23 | 1.95 | 0.67 |
| 9:AI:17:VAL:HG11 | 9:AI:81:ILE:HD13 | 1.75 | 0.67 |
| 12:AL:87:GLY:HA2 | 12:AL:98:TYR:HA | 1.75 | 0.67 |
| 15:AO:62:GLN:HA | 15:AO:65:ARG:NH1 | 2.09 | 0.67 |
| 31:BA:1429:G:H2' | 31:BA:1430:C:C6 | 2.30 | 0.67 |
| 31:BA:1987:G:H5' | 31:BA:1987:G:H8 | 1.60 | 0.67 |
| 32:BB:7:G:H4' | 44:BS:29:PHE:HD1 | 1.58 | 0.67 |
| 34:BE:23:VAL:HA | 34:BE:186:GLY:H | 1.60 | 0.67 |
| 38:BI:88:ILE:HG22 | 38:BI:89:TYR:H | 1.59 | 0.67 |
| 41:BP:57:THR:HB | 41:BP:59:LEU:N | 2.09 | 0.67 |
| 44:BS:89:ARG:O | 44:BS:92:TYR:CB | 2.39 | 0.67 |
| 1:CA:523:A:H61 | 12:CL:53:ARG:HH12 | 1.43 | 0.67 |
| 1:CA:748:C:H4' | 1:CA:749:C:O5' | 1.94 | 0.67 |
| 2:CB:44:LEU:H | 2:CB:44:LEU:HD12 | 1.59 | 0.67 |
| 3:CC:104:GLN:NE2 | 3:CC:105:GLU:H | 1.93 | 0.67 |
| 31:DA:197:A:C8 | 31:DA:197:A:C5' | 2.68 | 0.67 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:384:U:H2' | 31:DA:385:C:H6 | 1.60 | 0.67 |
| 31:DA:662:G:P | 41:DP:18:ARG:HD2 | 2.35 | 0.67 |
| 31:DA:1278:A:O2' | 31:DA:1279:G:H5' | 1.95 | 0.67 |
| 31:DA:1971:A:H1' | 33:DD:240:ALA:O | 1.95 | 0.67 |
| 35:DF:80:ALA:O | 35:DF:83:PHE:HB2 | 1.94 | 0.67 |
| 39:DN:78:TYR:N | 39:DN:79:PRO:HD3 | 2.09 | 0.67 |
| 41:DP:108:LYS:C | 41:DP:110:TYR:N | 2.53 | 0.67 |
| 42:DQ:43:THR:OG1 | 42:DQ:46:GLN:HG3 | 1.94 | 0.67 |
| 44:DS:87:PHE:O | 44:DS:88:ASP:HB2 | 1.94 | 0.67 |
| 45:DT:100:TYR:HD2 | 45:DT:103:ARG:HH21 | 1.43 | 0.67 |
| 47:DV:52:VAL:O | 47:DV:53:GLU:HB3 | 1.94 | 0.67 |
| 1:AA:622:A:C8 | 1:AA:623:C:C6 | 2.83 | 0.67 |
| 2:AB:67:THR:HG21 | 2:AB:155:LEU:HG | 1.77 | 0.67 |
| 30:B8:52:LYS:N | 30:B8:53:PRO:CD | 2.50 | 0.67 |
| 31:BA:286:C:N4 | 31:BA:355:G:H1 | 1.91 | 0.67 |
| 31:BA:535:C:C2' | 31:BA:536:A:H5' | 2.25 | 0.67 |
| 31:BA:1146:C:C2' | 31:BA:1147:C:H5' | 2.25 | 0.67 |
| 31:BA:1418:G:OP1 | 31:BA:1588:C:O2' | 2.13 | 0.67 |
| 31:BA:1719:G:C2' | 31:BA:1720:U:H5' | 2.25 | 0.67 |
| 33:BD:17:THR:HG23 | 33:BD:205:VAL:HB | 1.75 | 0.67 |
| 34:BE:60:ASN:ND2 | 34:BE:60:ASN:N | 2.42 | 0.67 |
| 34:BE:76:ARG:CG | 34:BE:195:LEU:HD12 | 2.24 | 0.67 |
| 41:BP:47:ASP:HB3 | 41:BP:48:PRO:CA | 2.24 | 0.67 |
| 42:BQ:78:PRO:O | 42:BQ:79:LEU:HB2 | 1.94 | 0.67 |
| 47:BV:25:LEU:H | 47:BV:94:LEU:CD1 | 2.07 | 0.67 |
| 47:BV:62:LEU:HD12 | 47:BV:96:ILE:HD13 | 1.76 | 0.67 |
| 3:CC:127:ARG:HD2 | 3:CC:127:ARG:N | 2.10 | 0.67 |
| 15:CO:64:ARG:HG3 | 15:CO:64:ARG:HH11 | 1.59 | 0.67 |
| 28:D6:20:ASN:O | 28:D6:21:TYR:CG | 2.48 | 0.67 |
| 31:DA:1108:U:C2' | 31:DA:1109:C:H5' | 2.24 | 0.67 |
| 31:DA:1161:C:H1' | 47:DV:8:GLY:O | 1.95 | 0.67 |
| 31:DA:1783:A:C2 | 31:DA:2587:A:C5 | 2.83 | 0.67 |
| 31:DA:2265:U:H4' | 42:DQ:13:GLN:HE22 | 1.60 | 0.67 |
| 31:DA:2542:A:H5' | 31:DA:2543:G:OP1 | 1.95 | 0.67 |
| 39:DN:2:LYS:HE2 | 46:DU:95:LEU:HD21 | 1.77 | 0.67 |
| 49:DX:33:LYS:O | 49:DX:35:THR:N | 2.27 | 0.67 |
| 1:AA:590:C:H2' | 1:AA:591:U:C6 | 2.30 | 0.66 |
| 1:AA:748:C:H4' | 1:AA:749:C:O5' | 1.95 | 0.66 |
| 1:AA:828:A:H2' | 1:AA:829:G:O4' | 1.94 | 0.66 |
| 1:AA:1072:G:H2' | 1:AA:1073:U:C6 | 2.29 | 0.66 |
| 1:AA:1442:G:C8 | 1:AA:1442(B):A:C2 | 2.83 | 0.66 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 5:AE:33:VAL:HG11 | 5:AE:109:ILE:HA | 1.77 | 0.66 |
| 30:B8:47:LYS:HD2 | 30:B8:48:PHE:O | 1.95 | 0.66 |
| 33:BD:228:PRO:CD | 33:BD:235:GLY:HA3 | 2.24 | 0.66 |
| 50:BY:10:GLY:O | 50:BY:27:VAL:HG22 | 1.96 | 0.66 |
| 1:CA:102:G:C4 | 1:CA:103:C:C5 | 2.83 | 0.66 |
| 1:CA:1076:C:C2 | 1:CA:1082:G:N2 | 2.62 | 0.66 |
| 1:CA:1342:C:H1' | 9:CI:124:GLN:HE22 | 1.59 | 0.66 |
| 3:CC:43:LEU:O | 3:CC:47:LEU:HB3 | 1.95 | 0.66 |
| 24:D2:26:ARG:HA | 24:D2:29:LYS:HE3 | 1.76 | 0.66 |
| 31:DA:271(E):U:H3 | 31:DA:271(S):G:H1 | 1.41 | 0.66 |
| 31:DA:1181:C:C2' | 31:DA:1182:A:H5' | 2.25 | 0.66 |
| 39:DN:91:LEU:HA | 39:DN:95:PRO:HB3 | 1.77 | 0.66 |
| 39:DN:131:GLN:HG2 | 39:DN:133:GLN:O | 1.94 | 0.66 |
| 41:DP:95:VAL:HA | 41:DP:99:LEU:HD23 | 1.77 | 0.66 |
| 27:B5:16:ARG:HH11 | 27:B5:16:ARG:CG | 2.04 | 0.66 |
| 31:BA:203:C:H3' | 31:BA:204:A:H5" | 1.77 | 0.66 |
| 31:BA:796:C:H2' | 31:BA:797:C:H6 | 1.56 | 0.66 |
| 31:BA:1109:C:C5 | 31:BA:1110:G:C4 | 2.79 | 0.66 |
| 31:BA:2286:A:C8 | 31:BA:2286:A:O2' | 2.48 | 0.66 |
| 31:BA:2723:C:H5" | 43:BR:2:ARG:HD3 | 1.76 | 0.66 |
| 33:BD:65:ILE:HD11 | 33:BD:67:PHE:CD1 | 2.29 | 0.66 |
| 41:BP:16:ARG:HG3 | 41:BP:17:LYS:N | 2.09 | 0.66 |
| 2:CB:77:ALA:HB2 | 2:CB:211:ILE:HD13 | 1.76 | 0.66 |
| 4:CD:91:SER:HA | 4:CD:94:LEU:HD12 | 1.77 | 0.66 |
| 12:CL:83:VAL:HG11 | 12:CL:100:ILE:HG12 | 1.77 | 0.66 |
| 18:CR:79:LEU:HD23 | 18:CR:80:PRO:CD | 2.22 | 0.66 |
| 30:D8:32:LEU:HB3 | 30:D8:35:GLN:H | 1.59 | 0.66 |
| 31:DA:244:A:C2 | 31:DA:255:A:C4 | 2.83 | 0.66 |
| 31:DA:1882:C:H2' | 31:DA:1882:C:O2 | 1.94 | 0.66 |
| 32:DB:8:U:H6 | 32:DB:8:U:C5' | 2.07 | 0.66 |
| 35:DF:67:GLN:O | 35:DF:67:GLN:CG | 2.43 | 0.66 |
| 36:DG:47:LYS:HE2 | 36:DG:81:LYS:HB2 | 1.78 | 0.66 |
| 39:DN:74:ARG:NH2 | 39:DN:101:HIS:HB3 | 2.09 | 0.66 |
| 46:DU:92:ARG:NH1 | 46:DU:94:ASN:HD22 | 1.94 | 0.66 |
| 1:AA:102:G:C4 | 1:AA:103:C:C5 | 2.84 | 0.66 |
| 1:AA:509:A:H2' | 1:AA:510:A:C8 | 2.30 | 0.66 |
| 1:AA:1076:C:C2 | 1:AA:1082:G:N2 | 2.64 | 0.66 |
| 2:AB:44:LEU:H | 2:AB:44:LEU:HD12 | 1.58 | 0.66 |
| 7:AG:113:GLU:HB2 | 7:AG:119:ARG:CG | 2.17 | 0.66 |
| 23:B1:25:LYS:C | 23:B1:26:ARG:HG3 | 2.19 | 0.66 |
| 31:BA:661:C:H4' | 41:BP:16:ARG:NH1 | 2.11 | 0.66 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1747(A):G:C2' | 31:BA:1748:G:H5' | 2.22 | 0.66 |
| 33:BD:35:LYS:HA | 33:BD:64:ILE:HG22 | 1.77 | 0.66 |
| 39:BN:74:ARG:NH2 | 39:BN:101:HIS:HB3 | 2.09 | 0.66 |
| 51:BZ:5:LEU:HD22 | 51:BZ:6:LYS:N | 2.11 | 0.66 |
| 2:CB:20:GLU:HG3 | 2:CB:191:ASP:HB2 | 1.77 | 0.66 |
| 2:CB:61:LEU:HA | 2:CB:64:ARG:HG2 | 1.75 | 0.66 |
| 2:CB:211:ILE:O | 2:CB:215:LEU:HB2 | 1.95 | 0.66 |
| 4:CD:158:ILE:HG23 | 4:CD:162:LEU:HD12 | 1.78 | 0.66 |
| 4:CD:172:PRO:HB2 | 4:CD:187:ARG:HH22 | 1.59 | 0.66 |
| 10:CJ:30:SER:OG | 10:CJ:81:THR:HG22 | 1.95 | 0.66 |
| 24:D2:25:VAL:HG13 | 24:D2:26:ARG:CD | 2.24 | 0.66 |
| 31:DA:2655:G:N3 | 31:DA:2655:G:H2' | 2.08 | 0.66 |
| 31:DA:2661:G:C8 | 31:DA:2662:A:C2 | 2.83 | 0.66 |
| 38:DI:1:MET:HB2 | 38:DI:21:VAL:O | 1.95 | 0.66 |
| 38:DI:52:ARG:O | 38:DI:53:ALA:CB | 2.37 | 0.66 |
| 1:AA:224:C:H2' | 1:AA:225:C:H6 | 1.61 | 0.66 |
| 30:B8:6:THR:CG2 | 30:B8:63:PRO:HD3 | 2.25 | 0.66 |
| 31:BA:1464:C:HO2' | 31:BA:1528:A:H8 | 1.40 | 0.66 |
| 38:BI:28:ASN:C | 38:BI:32:PRO:HG2 | 2.19 | 0.66 |
| 1:CA:502:G:C2 | 1:CA:503:C:O2 | 2.48 | 0.66 |
| 31:DA:1899:G:N2 | 31:DA:1902:C:H5 | 1.93 | 0.66 |
| 46:DU:12:ARG:O | 46:DU:15:LYS:HG2 | 1.95 | 0.66 |
| 47:DV:61:VAL:C | 47:DV:62:LEU:HD23 | 2.21 | 0.66 |
| 49:DX:41:ASN:HA | 49:DX:44:GLU:HG2 | 1.77 | 0.66 |
| 16:AP:48:TRP:CD1 | 16:AP:48:TRP:H | 2.13 | 0.66 |
| 20:AT:48:LYS:HB3 | 20:AT:51:GLU:HG3 | 1.77 | 0.66 |
| 28:B6:15:GLU:CD | 28:B6:18:ARG:HD2 | 2.21 | 0.66 |
| 30:B8:59:LYS:HD3 | 41:BP:50:ARG:HB3 | 1.75 | 0.66 |
| 31:BA:1719:G:H2' | 31:BA:1720:U:H5' | 1.76 | 0.66 |
| 31:BA:2272:U:H5" | 31:BA:2273:A:OP1 | 1.94 | 0.66 |
| 34:BE:33:VAL:HG12 | 34:BE:90:THR:H | 1.58 | 0.66 |
| 35:BF:83:PHE:O | 35:BF:85:GLY:N | 2.29 | 0.66 |
| 38:BI:38:LEU:H | 38:BI:38:LEU:HD12 | 1.60 | 0.66 |
| 43:BR:4:LEU:O | 43:BR:6:SER:N | 2.27 | 0.66 |
| 1:CA:1342:C:H4' | 9:CI:125:TYR:HB3 | 1.77 | 0.66 |
| 2:CB:233:SER:HB2 | 2:CB:234:PRO:HD2 | 1.76 | 0.66 |
| 12:CL:41:ARG:CG | 12:CL:42:THR:H | 2.08 | 0.66 |
| 16:CP:48:TRP:H | 16:CP:48:TRP:CD1 | 2.12 | 0.66 |
| 31:DA:1210:A:H5' | 31:DA:1212:G:C5' | 2.25 | 0.66 |
| 31:DA:2873:A:C2 | 43:DR:6:SER:HB2 | 2.30 | 0.66 |
| 35:DF:7:TYR:HD1 | 35:DF:8:GLN:H | 1.43 | 0.66 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 35:DF:132:VAL:HG22 | 35:DF:133:ASN:N | 2.11 | 0.66 |
| 36:DG:111:LEU:HA | 36:DG:114:ILE:CG1 | 2.26 | 0.66 |
| 39:DN:30:ILE:O | 39:DN:34:LEU:HD22 | 1.95 | 0.66 |
| 1:AA:559:A:H5" | 1:AA:560:U:H3' | 1.76 | 0.66 |
| 7:AG:116:ALA:O | 7:AG:120:ILE:HG12 | 1.95 | 0.66 |
| 31:BA:445:C:O2' | 31:BA:446:G:H5' | 1.95 | 0.66 |
| 31:BA:830:G:H4' | 31:BA:831:G:OP2 | 1.94 | 0.66 |
| 31:BA:1688:U:H1' | 31:BA:1701:A:C6 | 2.30 | 0.66 |
| 32:BB:28:C:C2 | 32:BB:29:A:C8 | 2.83 | 0.66 |
| 33:BD:35:LYS:NZ | 33:BD:64:ILE:O | 2.25 | 0.66 |
| 37:BH:85:LYS:HE2 | 37:BH:145:ALA:HB2 | 1.78 | 0.66 |
| 44:BS:42:ASP:C | 44:BS:44:LYS:H | 2.04 | 0.66 |
| 46:BU:90:VAL:HG13 | 47:BV:39:LEU:HG | 1.78 | 0.66 |
| 51:BZ:128:VAL:HG23 | 51:BZ:160:GLY:O | 1.95 | 0.66 |
| 1:CA:64:G:H4' | 1:CA:65:U:H5" | 1.78 | 0.66 |
| 1:CA:155:C:H2' | 1:CA:156:G:H8 | 1.60 | 0.66 |
| 4:CD:108:LEU:HD11 | 4:CD:174:LEU:HD22 | 1.77 | 0.66 |
| 9:CI:114:TYR:HD2 | 9:CI:114:TYR:N | 1.94 | 0.66 |
| 27:D5:55:ARG:C | 27:D5:56:LYS:HG3 | 2.20 | 0.66 |
| 31:DA:1164:G:H2' | 31:DA:1165:U:C6 | 2.30 | 0.66 |
| 33:DD:71:ASP:HB3 | 33:DD:103:ARG:NH2 | 2.10 | 0.66 |
| 33:DD:145:VAL:HG12 | 33:DD:146:GLU:O | 1.96 | 0.66 |
| 34:DE:76:ARG:CG | 34:DE:195:LEU:HD12 | 2.25 | 0.66 |
| 41:DP:115:LEU:HA | 41:DP:134:ALA:HB2 | 1.78 | 0.66 |
| 46:DU:49:HIS:HA | 46:DU:52:ARG:HB2 | 1.77 | 0.66 |
| 48:DW:4:LYS:CB | 48:DW:106:ILE:HG22 | 2.26 | 0.66 |
| 31:BA:271(E):U:H3 | 31:BA:271(S):G:H1 | 1.42 | 0.66 |
| 31:BA:571:A:C5' | 31:BA:2030:A:H62 | 1.88 | 0.66 |
| 31:BA:806:C:OP2 | 41:BP:39:LYS:CD | 2.43 | 0.66 |
| 31:BA:870:A:OP1 | 42:BQ:7:MET:HE2 | 1.95 | 0.66 |
| 32:BB:116:G:C5' | 32:BB:116:G:C8 | 2.79 | 0.66 |
| 40:BO:90:GLN:O | 40:BO:91:LEU:HB2 | 1.94 | 0.66 |
| 41:BP:62:LEU:H | 41:BP:62:LEU:CD2 | 2.07 | 0.66 |
| 46:BU:88:ILE:O | 46:BU:88:ILE:CD1 | 2.44 | 0.66 |
| 1:CA:622:A:C8 | 1:CA:623:C:C6 | 2.84 | 0.66 |
| 1:CA:1342:C:H1' | 9:CI:124:GLN:NE2 | 2.09 | 0.66 |
| 4:CD:128:VAL:O | 4:CD:130:GLY:N | 2.28 | 0.66 |
| 31:DA:128:C:H6 | 31:DA:128:C:H5" | 1.61 | 0.66 |
| 31:DA:1429:G:H2' | 31:DA:1430:C:C6 | 2.30 | 0.66 |
| 33:DD:44:ASN:CB | 33:DD:49:ILE:HA | 2.21 | 0.66 |
| 33:DD:77:ALA:HB2 | 33:DD:97:TYR:CD2 | 2.31 | 0.66 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 36:DG:25:TYR:CZ | 36:DG:32:PRO:HD3 | 2.31 | 0.66 |
| 41:DP:56:SER:O | 41:DP:58:THR:N | 2.28 | 0.66 |
| 49:DX:35:THR:HB | 49:DX:75:ASP:OD2 | 1.96 | 0.66 |
| 1:AA:359:U:H2' | 1:AA:360:A:H8 | 1.61 | 0.66 |
| 2:AB:29:ALA:O | 2:AB:32:ILE:HG22 | 1.95 | 0.66 |
| 6:AF:17:SER:O | 6:AF:21:LEU:HD22 | 1.96 | 0.66 |
| 18:AR:62:GLU:HA | 18:AR:65:ILE:HD11 | 1.77 | 0.66 |
| 23:B1:37:ILE:HD11 | 31:BA:2079:U:H4' | 1.76 | 0.66 |
| 23:B1:62:VAL:HG22 | 23:B1:63:ALA:N | 2.10 | 0.66 |
| 24:B2:25:VAL:HG13 | 24:B2:26:ARG:HD2 | 1.77 | 0.66 |
| 31:BA:848:G:H2' | 31:BA:849:A:C8 | 2.30 | 0.66 |
| 31:BA:2394:C:H2' | 31:BA:2395:C:H5' | 1.77 | 0.66 |
| 35:BF:24:LEU:HB3 | 35:BF:25:PRO:CD | 2.24 | 0.66 |
| 41:BP:10:PRO:CD | 41:BP:11:GLY:H | 2.07 | 0.66 |
| 43:BR:3:HIS:O | 43:BR:4:LEU:HB3 | 1.96 | 0.66 |
| 1:CA:1337:G:H5" | 1:CA:1338:G:OP1 | 1.94 | 0.66 |
| 5:CE:33:VAL:HG11 | 5:CE:109:ILE:HA | 1.78 | 0.66 |
| 31:DA:286:C:N4 | 31:DA:355:G:H1 | 1.92 | 0.66 |
| 31:DA:1210:A:H4' | 31:DA:1211:U:OP2 | 1.95 | 0.66 |
| 34:DE:167:VAL:HG13 | 34:DE:170:LEU:HD11 | 1.76 | 0.66 |
| 35:DF:89:VAL:HG12 | 35:DF:90:PHE:H | 1.57 | 0.66 |
| 36:DG:47:LYS:HG3 | 36:DG:82:LEU:HD11 | 1.76 | 0.66 |
| 38:DI:110:ASP:C | 38:DI:112:LYS:H | 2.03 | 0.66 |
| 50:DY:81:LYS:CG | 50:DY:96:ILE:HG22 | 2.26 | 0.66 |
| 2:AB:28:PHE:HD1 | 2:AB:190:THR:HG22 | 1.61 | 0.66 |
| 2:AB:188:ALA:HB1 | 2:AB:192:SER:HB2 | 1.77 | 0.66 |
| 16:AP:53:VAL:HG12 | 16:AP:79:VAL:HG13 | 1.77 | 0.66 |
| 31:BA:511:U:C3' | 31:BA:512:G:H5" | 2.19 | 0.66 |
| 43:BR:24:GLN:NE2 | 43:BR:36:THR:HG21 | 2.08 | 0.66 |
| 45:BT:28:VAL:O | 45:BT:29:ARG:HD3 | 1.96 | 0.66 |
| 1:CA:359:U:H2' | 1:CA:360:A:H8 | 1.61 | 0.66 |
| 5:CE:57:LYS:O | 5:CE:61:TYR:HD2 | 1.78 | 0.66 |
| 11:CK:34:ASP:HB2 | 11:CK:35:PRO:HD2 | 1.78 | 0.66 |
| 15:CO:3:ILE:O | 15:CO:3:ILE:HG12 | 1.96 | 0.66 |
| 25:D3:11:SER:OG | 25:D3:13:ILE:HG12 | 1.95 | 0.66 |
| 31:DA:626:U:H3 | 41:DP:105:LEU:HG | 1.60 | 0.66 |
| 31:DA:780:G:H21 | 31:DA:783:A:H62 | 1.43 | 0.66 |
| 31:DA:1493:C:H5 | 31:DA:2206:G:O2' | 1.77 | 0.66 |
| 31:DA:2402:C:H2' | 31:DA:2403:C:H5' | 1.78 | 0.66 |
| 36:DG:137:GLU:HA | 36:DG:152:LEU:HD22 | 1.78 | 0.66 |
| 45:DT:13:ARG:HH21 | 45:DT:15:VAL:HG13 | 1.61 | 0.66 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:155:C:H2' | 1:AA:156:G:H8 | 1.61 | 0.66 |
| 1:AA:1342:C:H1' | 9:AI:124:GLN:HE22 | 1.61 | 0.66 |
| 8:AH:10:LEU:HD13 | 8:AH:83:ILE:HD11 | 1.78 | 0.66 |
| 10:AJ:5:ARG:HG3 | 10:AJ:73:ASP:OD1 | 1.96 | 0.66 |
| 35:BF:22:ALA:CA | 35:BF:26:ALA:HB2 | 2.26 | 0.66 |
| 45:BT:32:TYR:CG | 45:BT:81:PRO:HB2 | 2.31 | 0.66 |
| 2:CB:188:ALA:HB1 | 2:CB:192:SER:HB2 | 1.77 | 0.66 |
| 23:D1:16:ASN:HB3 | 23:D1:46:LEU:CG | 2.26 | 0.66 |
| 27:D5:2:ALA:CA | 31:DA:2015:A:H1' | 2.23 | 0.66 |
| 42:DQ:8:LYS:CD | 42:DQ:9:TYR:H | 2.08 | 0.66 |
| 47:DV:62:LEU:HB3 | 47:DV:98:GLU:CB | 2.26 | 0.66 |
| 1:AA:103:C:OP2 | 20:AT:14:LYS:HD3 | 1.96 | 0.65 |
| 1:AA:622:A:C8 | 1:AA:623:C:C5 | 2.83 | 0.65 |
| 1:AA:659:U:H2' | 1:AA:660:G:H5' | 1.78 | 0.65 |
| 1:AA:667:G:H4' | 15:AO:51:HIS:CE1 | 2.31 | 0.65 |
| 3:AC:100:ALA:O | 3:AC:101:LEU:HB2 | 1.94 | 0.65 |
| 13:AM:92:HIS:CE1 | 13:AM:98:VAL:HG23 | 2.31 | 0.65 |
| 20:AT:97:ALA:O | 20:AT:99:LEU:N | 2.29 | 0.65 |
| 28:B6:15:GLU:CD | 28:B6:18:ARG:CD | 2.69 | 0.65 |
| 31:BA:404:C:H4' | 31:BA:405:U:H5' | 1.78 | 0.65 |
| 31:BA:518:G:H4' | 48:BW:18:ARG:NH1 | 2.12 | 0.65 |
| 31:BA:2469:A:O2' | 42:BQ:56:ARG:HG2 | 1.95 | 0.65 |
| 45:BT:13:ARG:HH21 | 45:BT:15:VAL:CG1 | 2.09 | 0.65 |
| 1:CA:983:A:H2 | 1:CA:984:C:C5 | 2.14 | 0.65 |
| 9:CI:55:ALA:HB1 | 9:CI:58:ARG:HD2 | 1.77 | 0.65 |
| 31:DA:649:G:H2' | 31:DA:650:C:C6 | 2.31 | 0.65 |
| 31:DA:924:C:H2' | 31:DA:925:C:C6 | 2.31 | 0.65 |
| 31:DA:1405:U:H2' | 31:DA:1406:U:H6 | 1.56 | 0.65 |
| 31:DA:2286:A:C8 | 31:DA:2286:A:O2' | 2.47 | 0.65 |
| 34:DE:117:MET:O | 34:DE:117:MET:HG2 | 1.96 | 0.65 |
| 35:DF:83:PHE:O | 35:DF:85:GLY:N | 2.30 | 0.65 |
| 37:DH:35:VAL:O | 37:DH:37:VAL:HG23 | 1.96 | 0.65 |
| 37:DH:158:HIS:HE2 | 37:DH:170:ARG:HA | 1.59 | 0.65 |
| 49:DX:60:ARG:NH2 | 49:DX:74:PRO:HG2 | 2.11 | 0.65 |
| 1:AA:135:C:H2' | 1:AA:136:C:H5' | 1.77 | 0.65 |
| 1:AA:775:G:O2' | 1:AA:776:G:H5' | 1.96 | 0.65 |
| 30:B8:25:MET:HG3 | 41:BP:64:LYS:CB | 2.26 | 0.65 |
| 31:BA:1171:G:H8 | 31:BA:1171:G:OP2 | 1.79 | 0.65 |
| 31:BA:2619:C:O2' | 31:BA:2620:C:H5' | 1.96 | 0.65 |
| 31:BA:2722:G:O2' | 43:BR:5:LYS:HB2 | 1.96 | 0.65 |
| 31:BA:2759:G:H5' | 31:BA:2759:G:C8 | 2.31 | 0.65 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 32:BB:8:U:H6 | 32:BB:8:U:C5' | 2.08 | 0.65 |
| 33:BD:267:SER:O | 33:BD:268:ARG:HB2 | 1.97 | 0.65 |
| 38:BI:75:LEU:HD12 | 38:BI:76:THR:N | 2.10 | 0.65 |
| 41:BP:29:LYS:H | 41:BP:29:LYS:CD | 1.93 | 0.65 |
| 45:BT:29:ARG:HG3 | 45:BT:30:VAL:HG13 | 1.77 | 0.65 |
| 46:BU:92:ARG:HD3 | 46:BU:94:ASN:HB3 | 1.77 | 0.65 |
| 1:CA:66:G:H4' | 1:CA:173:U:C5 | 2.31 | 0.65 |
| 1:CA:353:A:H5' | 1:CA:353:A:C8 | 2.31 | 0.65 |
| 1:CA:437:U:O2' | 1:CA:438:G:H5' | 1.96 | 0.65 |
| 1:CA:559:A:H4' | 1:CA:560:U:H5" | 1.78 | 0.65 |
| 1:CA:590:C:H2' | 1:CA:591:U:C6 | 2.31 | 0.65 |
| 11:CK:29:ILE:HB | 11:CK:44:SER:CB | 2.25 | 0.65 |
| 24:D2:41:ILE:HG13 | 31:DA:95:G:N3 | 2.11 | 0.65 |
| 28:D6:13:CYS:O | 28:D6:21:TYR:HA | 1.95 | 0.65 |
| 28:D6:16:CYS:SG | 28:D6:48:VAL:HG23 | 2.35 | 0.65 |
| 31:DA:49:A:C3' | 31:DA:50:U:H5' | 2.25 | 0.65 |
| 31:DA:128:C:H2' | 31:DA:129:C:O4' | 1.95 | 0.65 |
| 31:DA:527:C:N4 | 31:DA:2779:U:OP2 | 2.29 | 0.65 |
| 31:DA:672:C:H2' | 31:DA:673:C:C6 | 2.30 | 0.65 |
| 31:DA:1210:A:H5" | 31:DA:1211:U:H3' | 1.79 | 0.65 |
| 31:DA:1464:C:HO2' | 31:DA:1528:A:H8 | 1.40 | 0.65 |
| 37:DH:106:THR:HG22 | 37:DH:112:PRO:HB3 | 1.78 | 0.65 |
| 39:DN:78:TYR:CE1 | 39:DN:79:PRO:HB3 | 2.31 | 0.65 |
| 50:DY:2:ARG:N | 50:DY:4:LYS:HG2 | 2.11 | 0.65 |
| 1:AA:881:G:P | 12:AL:12:ARG:HH22 | 2.18 | 0.65 |
| 1:AA:1504:G:OP1 | 1:AA:1507:A:H4' | 1.95 | 0.65 |
| 16:AP:4:ILE:HB | 16:AP:66:PRO:HB3 | 1.78 | 0.65 |
| 23:B1:73:LEU:HD13 | 23:B1:90:ILE:HG22 | 1.78 | 0.65 |
| 31:BA:708:C:O2 | 31:BA:708:C:H2' | 1.96 | 0.65 |
| 31:BA:828:U:O2' | 31:BA:829:A:H5' | 1.96 | 0.65 |
| 31:BA:1164:G:H2' | 31:BA:1165:U:C6 | 2.31 | 0.65 |
| 31:BA:1882:C:O2 | 31:BA:1882:C:H2' | 1.96 | 0.65 |
| 38:BI:10:GLU:O | 38:BI:12:LEU:HD23 | 1.95 | 0.65 |
| 38:BI:25:TYR:CD1 | 38:BI:30:LEU:HD11 | 2.31 | 0.65 |
| 46:BU:88:ILE:O | 46:BU:88:ILE:HD13 | 1.96 | 0.65 |
| 4:CD:148:VAL:HG12 | 4:CD:149:ALA:N | 2.12 | 0.65 |
| 5:CE:51:VAL:O | 5:CE:55:VAL:HG23 | 1.96 | 0.65 |
| 11:CK:29:ILE:HB | 11:CK:44:SER:HB3 | 1.77 | 0.65 |
| 23:D1:25:LYS:C | 23:D1:26:ARG:HG3 | 2.21 | 0.65 |
| 27:D5:6:VAL:HG13 | 27:D5:7:PRO:HD2 | 1.77 | 0.65 |
| 30:D8:59:LYS:NZ | 30:D8:59:LYS:CB | 2.59 | 0.65 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:953:A:C2' | 31:DA:954:G:H5' | 2.27 | 0.65 |
| 31:DA:2307:G:N2 | 31:DA:2308:G:H5' | 2.11 | 0.65 |
| 31:DA:2476:A:C6 | 31:DA:2477:C:C5 | 2.84 | 0.65 |
| 34:DE:95:ILE:CD1 | 34:DE:95:ILE:H | 2.10 | 0.65 |
| 34:DE:163:GLU:O | 34:DE:165:VAL:HG23 | 1.95 | 0.65 |
| 35:DF:3:GLU:HB2 | 35:DF:20:LEU:H | 1.62 | 0.65 |
| 35:DF:124:LEU:HD12 | 35:DF:125:LEU:N | 2.11 | 0.65 |
| 1:AA:559:A:H4' | 1:AA:560:U:C5' | 2.26 | 0.65 |
| 5:AE:57:LYS:O | 5:AE:61:TYR:HD2 | 1.79 | 0.65 |
| 22:B0:39:ARG:HH21 | 31:BA:2355:C:H1' | 1.61 | 0.65 |
| 27:B5:40:LYS:HE2 | 27:B5:46:CYS:HB3 | 1.78 | 0.65 |
| 31:BA:2632:A:N3 | 34:BE:61:ARG:NH1 | 2.44 | 0.65 |
| 36:BG:111:LEU:HA | 36:BG:114:ILE:CG1 | 2.25 | 0.65 |
| 36:BG:148:MET:HA | 36:BG:148:MET:HE3 | 1.77 | 0.65 |
| 51:BZ:71:VAL:HG22 | 51:BZ:88:PHE:HE2 | 1.62 | 0.65 |
| 51:BZ:143:GLY:C | 51:BZ:144:LEU:HD22 | 2.21 | 0.65 |
| 1:CA:750:G:N3 | 15:CO:23:GLY:HA3 | 2.11 | 0.65 |
| 1:CA:881:G:P | 12:CL:12:ARG:HH22 | 2.19 | 0.65 |
| 4:CD:149:ALA:O | 4:CD:153:ARG:HG3 | 1.95 | 0.65 |
| 10:CJ:5:ARG:HG3 | 10:CJ:73:ASP:OD1 | 1.96 | 0.65 |
| 30:D8:23:VAL:CG1 | 30:D8:46:ARG:HB3 | 2.26 | 0.65 |
| 31:DA:659:C:C5' | 31:DA:659:C:C6 | 2.74 | 0.65 |
| 38:DI:123:LEU:HD23 | 38:DI:142:VAL:HB | 1.77 | 0.65 |
| 42:DQ:6:ARG:O | 42:DQ:7:MET:HG2 | 1.97 | 0.65 |
| 43:DR:33:ARG:CG | 43:DR:115:GLU:HG3 | 2.26 | 0.65 |
| 45:DT:28:VAL:HG22 | 45:DT:46:GLU:HA | 1.78 | 0.65 |
| 48:DW:86:LEU:C | 48:DW:86:LEU:HD12 | 2.20 | 0.65 |
| 8:AH:77:GLU:HG3 | 8:AH:78:GLN:H | 1.60 | 0.65 |
| 31:BA:117:G:H5" | 31:BA:118:A:OP2 | 1.96 | 0.65 |
| 31:BA:484:C:H2' | 31:BA:485:C:C6 | 2.32 | 0.65 |
| 31:BA:954:G:H5" | 42:BQ:13:GLN:HG2 | 1.79 | 0.65 |
| 31:BA:1568:G:P | 33:BD:63:ARG:HH22 | 2.19 | 0.65 |
| 34:BE:111:ARG:NH1 | 43:BR:2:ARG:HH21 | 1.93 | 0.65 |
| 37:BH:46:GLU:O | 37:BH:47:GLU:HB2 | 1.96 | 0.65 |
| 41:BP:13:ASN:N | 41:BP:13:ASN:HD22 | 1.95 | 0.65 |
| 47:BV:89:GLN:HE21 | 47:BV:91:TYR:HB2 | 1.61 | 0.65 |
| 49:BX:59:VAL:HG23 | 49:BX:74:PRO:HD2 | 1.76 | 0.65 |
| 50:BY:75:ILE:HD11 | 50:BY:79:CYS:C | 2.21 | 0.65 |
| 51:BZ:111:VAL:HG13 | 51:BZ:112:ARG:N | 2.11 | 0.65 |
| 29:D7:16:HIS:CB | 29:D7:44:PRO:HG2 | 2.26 | 0.65 |
| 30:D8:25:MET:HG3 | 41:DP:64:LYS:HB3 | 1.78 | 0.65 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:943:U:OP2 | 41:DP:38:GLN:CD | 2.40 | 0.65 |
| 31:DA:1719:G:H2' | 31:DA:1720:U:H5' | 1.78 | 0.65 |
| 32:DB:37:C:C5 | 32:DB:38:C:C5 | 2.84 | 0.65 |
| 33:DD:32:SER:C | 33:DD:33:LEU:HG | 2.21 | 0.65 |
| 33:DD:35:LYS:HG2 | 33:DD:64:ILE:H | 1.61 | 0.65 |
| 36:DG:7:LEU:HB2 | 36:DG:104:GLU:OE2 | 1.97 | 0.65 |
| 37:DH:68:THR:O | 37:DH:69:ARG:C | 2.39 | 0.65 |
| 14:AN:23:ARG:NH1 | 14:AN:30:ALA:HB2 | 2.12 | 0.65 |
| 24:B2:14:ARG:NH1 | 24:B2:57:ILE:N | 2.43 | 0.65 |
| 31:BA:2275:C:O2 | 42:BQ:83:MET:HG3 | 1.96 | 0.65 |
| 33:BD:3:VAL:H | 33:BD:20:ASP:HB2 | 1.61 | 0.65 |
| 33:BD:58:HIS:HD2 | 33:BD:59:LYS:H | 1.44 | 0.65 |
| 39:BN:65:LYS:HA | 39:BN:65:LYS:HE2 | 1.79 | 0.65 |
| 41:BP:112:LEU:HD22 | 41:BP:113:LYS:N | 2.12 | 0.65 |
| 1:CA:184:G:H2' | 1:CA:185:A:H8 | 1.61 | 0.65 |
| 1:CA:973:G:H3' | 1:CA:974:A:H5" | 1.79 | 0.65 |
| 31:DA:59:U:OP1 | 49:DX:72:LYS:HE2 | 1.96 | 0.65 |
| 31:DA:943:U:OP2 | 41:DP:38:GLN:OE1 | 2.14 | 0.65 |
| 34:DE:102:VAL:HA | 34:DE:200:GLU:HA | 1.77 | 0.65 |
| 37:DH:137:ASP:O | 37:DH:138:LYS:CB | 2.44 | 0.65 |
| 38:DI:75:LEU:HD12 | 38:DI:76:THR:N | 2.11 | 0.65 |
| 39:DN:47:ALA:CB | 39:DN:112:LEU:HD11 | 2.22 | 0.65 |
| 47:DV:1:MET:CE | 47:DV:44:LYS:HB2 | 2.23 | 0.65 |
| 50:DY:28:LYS:CD | 50:DY:37:VAL:HG12 | 2.26 | 0.65 |
| 1:AA:1342:C:H1' | 9:AI:124:GLN:NE2 | 2.12 | 0.65 |
| 4:AD:18:LYS:HE3 | 4:AD:31:CYS:SG | 2.36 | 0.65 |
| 27:B5:33:CYS:SG | 27:B5:49:CYS:CB | 2.85 | 0.65 |
| 27:B5:55:ARG:C | 27:B5:56:LYS:HG3 | 2.20 | 0.65 |
| 31:BA:228:A:H2' | 31:BA:230:U:O4' | 1.96 | 0.65 |
| 31:BA:669:G:H4' | 31:BA:670:A:OP2 | 1.94 | 0.65 |
| 33:BD:35:LYS:HG2 | 33:BD:64:ILE:H | 1.61 | 0.65 |
| 41:BP:108:LYS:C | 41:BP:110:TYR:N | 2.55 | 0.65 |
| 43:BR:17:ARG:O | 43:BR:20:LEU:HB3 | 1.96 | 0.65 |
| 51:BZ:10:ARG:HH21 | 51:BZ:26:GLY:H | 1.44 | 0.65 |
| 1:CA:254:G:OP1 | 17:CQ:67:LYS:O | 2.15 | 0.65 |
| 4:CD:62:GLN:HA | 4:CD:62:GLN:NE2 | 2.12 | 0.65 |
| 23:D1:26:ARG:HB3 | 23:D1:34:THR:HA | 1.77 | 0.65 |
| 28:D6:20:ASN:CG | 28:D6:21:TYR:N | 2.54 | 0.65 |
| 31:DA:308:G:O2' | 50:DY:19:LYS:HE3 | 1.96 | 0.65 |
| 31:DA:661:C:H4' | 41:DP:16:ARG:NH1 | 2.11 | 0.65 |
| 32:DB:21:G:O6 | 32:DB:63:G:C5 | 2.50 | 0.65 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 49:DX:60:ARG:CZ | 49:DX:74:PRO:HG2 | 2.27 | 0.65 |
| 1:AA:827:U:H5” | 1:AA:828:A:OP2 | 1.97 | 0.65 |
| 1:AA:939:G:H5” | 7:AG:102:ARG:HH12 | 1.62 | 0.65 |
| 1:AA:1096:C:H2’ | 1:AA:1097:C:H6 | 1.61 | 0.65 |
| 2:AB:22:LYS:NZ | 2:AB:40:HIS:HE1 | 1.94 | 0.65 |
| 24:B2:33:MET:HG2 | 49:BX:11:PRO:HD3 | 1.78 | 0.65 |
| 31:BA:197:A:C8 | 31:BA:197:A:C5’ | 2.72 | 0.65 |
| 31:BA:1833:U:H2’ | 31:BA:1834:U:C6 | 2.30 | 0.65 |
| 31:BA:1946:U:H2’ | 31:BA:1947:C:H6 | 1.59 | 0.65 |
| 31:BA:2317:C:H2’ | 31:BA:2318:G:C5’ | 2.20 | 0.65 |
| 34:BE:134:ILE:HG12 | 34:BE:134:ILE:O | 1.96 | 0.65 |
| 45:BT:78:LEU:O | 45:BT:78:LEU:HD23 | 1.97 | 0.65 |
| 50:BY:2:ARG:C | 50:BY:4:LYS:H | 2.02 | 0.65 |
| 1:CA:409:G:H2’ | 1:CA:410:G:H5’ | 1.78 | 0.65 |
| 1:CA:963:G:H21 | 10:CJ:55:LYS:CE | 2.09 | 0.65 |
| 1:CA:1106:G:H5” | 3:CC:172:ARG:HG2 | 1.79 | 0.65 |
| 4:CD:13:ARG:O | 4:CD:15:GLU:N | 2.30 | 0.65 |
| 28:D6:51:GLU:O | 28:D6:52:VAL:HB | 1.95 | 0.65 |
| 31:DA:92:A:H2’ | 31:DA:93:G:O4’ | 1.97 | 0.65 |
| 31:DA:830:G:H4’ | 31:DA:831:G:OP2 | 1.97 | 0.65 |
| 31:DA:979:G:H3’ | 31:DA:980:A:H5” | 1.79 | 0.65 |
| 31:DA:1047:G:N2 | 31:DA:1111:A:H62 | 1.95 | 0.65 |
| 31:DA:1163:G:H5’ | 47:DV:92:THR:HG21 | 1.78 | 0.65 |
| 31:DA:1639:U:C2’ | 31:DA:1640:C:H5” | 2.25 | 0.65 |
| 32:DB:116:G:C5’ | 32:DB:116:G:C8 | 2.80 | 0.65 |
| 35:DF:183:VAL:O | 35:DF:187:VAL:HG23 | 1.96 | 0.65 |
| 48:DW:82:LEU:HG | 48:DW:84:ARG:HH21 | 1.61 | 0.65 |
| 1:AA:543:C:C2 | 1:AA:544:G:C8 | 2.85 | 0.65 |
| 1:AA:1497:G:H2’ | 1:AA:1498:U:H5’ | 1.77 | 0.65 |
| 31:BA:626:U:N3 | 41:BP:105:LEU:HG | 2.11 | 0.65 |
| 33:BD:228:PRO:HD3 | 33:BD:235:GLY:HA3 | 1.76 | 0.65 |
| 35:BF:205:ARG:O | 35:BF:206:ILE:HG13 | 1.97 | 0.65 |
| 49:BX:78:LYS:O | 49:BX:78:LYS:HD3 | 1.96 | 0.65 |
| 51:BZ:29:TYR:HE2 | 51:BZ:87:ASP:HB2 | 1.61 | 0.65 |
| 2:CB:178:ARG:NH2 | 8:CH:74:PRO:HG3 | 2.11 | 0.65 |
| 4:CD:126:ILE:HG22 | 4:CD:127:THR:H | 1.60 | 0.65 |
| 12:CL:55:VAL:HG12 | 12:CL:56:ALA:H | 1.62 | 0.65 |
| 27:D5:32:PRO:O | 27:D5:33:CYS:HB3 | 1.95 | 0.65 |
| 30:D8:58:ILE:HG22 | 41:DP:49:ARG:HD2 | 1.79 | 0.65 |
| 31:DA:102:G:O2’ | 31:DA:103:A:P | 2.55 | 0.65 |
| 31:DA:806:C:OP2 | 41:DP:39:LYS:CD | 2.45 | 0.65 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 35:DF:66:PRO:O | 35:DF:67:GLN:HB3 | 1.97 | 0.65 |
| 39:DN:2:LYS:HE2 | 46:DU:95:LEU:CD2 | 2.27 | 0.65 |
| 39:DN:128:HIS:NE2 | 39:DN:131:GLN:HB2 | 2.12 | 0.65 |
| 44:DS:17:ARG:C | 44:DS:19:LYS:H | 2.04 | 0.65 |
| 47:DV:5:VAL:HG23 | 47:DV:36:PRO:HB2 | 1.79 | 0.65 |
| 1:AA:1162:C:H2' | 1:AA:1163:C:C6 | 2.32 | 0.65 |
| 2:AB:114:ARG:HD2 | 2:AB:141:GLU:OE1 | 1.96 | 0.65 |
| 23:B1:73:LEU:HD11 | 23:B1:94:LEU:HG | 1.79 | 0.65 |
| 31:BA:92:A:H2' | 31:BA:93:G:O4' | 1.97 | 0.65 |
| 31:BA:1497:U:H3 | 31:BA:1578:U:P | 2.20 | 0.65 |
| 31:BA:1987:G:H5' | 31:BA:1987:G:C8 | 2.32 | 0.65 |
| 42:BQ:16:ARG:HH11 | 42:BQ:16:ARG:HB2 | 1.62 | 0.65 |
| 45:BT:31:SER:C | 45:BT:32:TYR:CD2 | 2.75 | 0.65 |
| 47:BV:23:GLU:O | 47:BV:24:LYS:C | 2.40 | 0.65 |
| 1:CA:963:G:H21 | 10:CJ:55:LYS:HE2 | 1.61 | 0.65 |
| 1:CA:963:G:N3 | 10:CJ:55:LYS:NZ | 2.45 | 0.65 |
| 1:CA:1117:G:H4' | 9:CI:104:ARG:CZ | 2.27 | 0.65 |
| 2:CB:165:VAL:HG23 | 2:CB:166:ASP:H | 1.62 | 0.65 |
| 8:CH:77:GLU:HG3 | 8:CH:78:GLN:N | 2.10 | 0.65 |
| 9:CI:97:LYS:HB3 | 9:CI:98:PRO:HD3 | 1.79 | 0.65 |
| 12:CL:8:ASN:ND2 | 17:CQ:34:LYS:HE2 | 2.08 | 0.65 |
| 22:D0:43:THR:HG22 | 31:DA:2331:G:O3' | 1.96 | 0.65 |
| 22:D0:74:ARG:NH2 | 32:DB:13:A:OP2 | 2.30 | 0.65 |
| 24:D2:14:ARG:NH1 | 24:D2:57:ILE:N | 2.44 | 0.65 |
| 30:D8:6:THR:HG22 | 30:D8:63:PRO:HD3 | 1.78 | 0.65 |
| 31:DA:825:C:H2' | 31:DA:826:U:O5' | 1.98 | 0.65 |
| 31:DA:2236:C:H2' | 31:DA:2237:G:H5' | 1.76 | 0.65 |
| 31:DA:2394:C:H2' | 31:DA:2395:C:H5' | 1.79 | 0.65 |
| 42:DQ:54:MET:HG3 | 42:DQ:117:ALA:HB1 | 1.79 | 0.65 |
| 1:AA:450:G:H5" | 16:AP:41:PRO:O | 1.97 | 0.64 |
| 1:AA:820:U:H4' | 1:AA:821:G:OP2 | 1.97 | 0.64 |
| 1:AA:946:A:H2' | 1:AA:947:G:H8 | 1.62 | 0.64 |
| 1:AA:1321:C:H5' | 1:AA:1322:C:H5" | 1.78 | 0.64 |
| 1:AA:1442(A):G:H8 | 45:BT:118:ARG:NH1 | 1.95 | 0.64 |
| 2:AB:187:LEU:HD13 | 2:AB:187:LEU:O | 1.95 | 0.64 |
| 5:AE:80:ILE:HD11 | 5:AE:138:ALA:HB1 | 1.79 | 0.64 |
| 13:AM:46:LYS:HG3 | 13:AM:47:ASP:H | 1.61 | 0.64 |
| 31:BA:84:A:H5' | 50:BY:9:LYS:HD2 | 1.79 | 0.64 |
| 31:BA:2870:C:C2' | 31:BA:2871:C:H5' | 2.27 | 0.64 |
| 33:BD:35:LYS:CG | 33:BD:64:ILE:N | 2.58 | 0.64 |
| 39:BN:129:PRO:O | 39:BN:130:HIS:HB2 | 1.96 | 0.64 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 40:BO:63:VAL:HG13 | 40:BO:84:ALA:HA | 1.80 | 0.64 |
| 41:BP:107:LYS:O | 41:BP:109:GLY:N | 2.29 | 0.64 |
| 49:BX:65:ARG:O | 49:BX:66:LEU:HB2 | 1.97 | 0.64 |
| 1:CA:775:G:H2' | 1:CA:776:G:H5' | 1.79 | 0.64 |
| 7:CG:26:PHE:O | 7:CG:30:ILE:HG12 | 1.96 | 0.64 |
| 9:CI:103:THR:HG22 | 9:CI:105:ASP:H | 1.63 | 0.64 |
| 31:DA:151:C:O2' | 31:DA:152:G:H5' | 1.97 | 0.64 |
| 33:DD:27:THR:HG23 | 33:DD:28:GLU:N | 2.13 | 0.64 |
| 41:DP:16:ARG:CZ | 41:DP:16:ARG:HB2 | 2.27 | 0.64 |
| 45:DT:29:ARG:CG | 45:DT:84:GLN:HG3 | 2.27 | 0.64 |
| 50:DY:45:VAL:HG11 | 50:DY:62:GLU:H | 1.62 | 0.64 |
| 1:AA:450:G:OP1 | 16:AP:43:LYS:NZ | 2.30 | 0.64 |
| 2:AB:116:GLU:HA | 2:AB:119:GLU:CB | 2.28 | 0.64 |
| 4:AD:158:ILE:HG23 | 4:AD:162:LEU:HD12 | 1.80 | 0.64 |
| 23:B1:10:LYS:O | 23:B1:13:ILE:HG23 | 1.96 | 0.64 |
| 31:BA:588:U:H2' | 31:BA:589:C:C6 | 2.33 | 0.64 |
| 31:BA:1336:A:H2' | 31:BA:1337:G:C8 | 2.32 | 0.64 |
| 31:BA:1784:A:H4' | 31:BA:1785:A:O5' | 1.97 | 0.64 |
| 32:BB:21:G:O2' | 32:BB:22:U:P | 2.54 | 0.64 |
| 37:BH:156:ALA:C | 37:BH:158:HIS:H | 2.05 | 0.64 |
| 39:BN:63:THR:O | 39:BN:64:GLY:O | 2.15 | 0.64 |
| 39:BN:78:TYR:CD1 | 39:BN:79:PRO:HB3 | 2.32 | 0.64 |
| 1:CA:327:A:C4 | 1:CA:329:A:C8 | 2.85 | 0.64 |
| 1:CA:622:A:C8 | 1:CA:623:C:C5 | 2.86 | 0.64 |
| 3:CC:134:ILE:HG23 | 3:CC:151:VAL:HB | 1.79 | 0.64 |
| 4:CD:20:TYR:CD2 | 4:CD:26:CYS:HB3 | 2.33 | 0.64 |
| 18:CR:62:GLU:HA | 18:CR:65:ILE:HD11 | 1.79 | 0.64 |
| 20:CT:48:LYS:HB3 | 20:CT:51:GLU:HG3 | 1.79 | 0.64 |
| 31:DA:176:G:O2' | 31:DA:177:G:H5' | 1.97 | 0.64 |
| 31:DA:2052:G:C8 | 34:DE:141:ILE:HD11 | 2.32 | 0.64 |
| 31:DA:2712:U:O2 | 31:DA:2712:U:H5" | 1.97 | 0.64 |
| 31:DA:2723:C:H5" | 43:DR:2:ARG:CD | 2.27 | 0.64 |
| 31:DA:2733:A:C2' | 31:DA:2734:A:H5" | 2.27 | 0.64 |
| 35:DF:63:LYS:HE2 | 35:DF:67:GLN:HB2 | 1.77 | 0.64 |
| 40:DO:63:VAL:HG13 | 40:DO:84:ALA:HA | 1.79 | 0.64 |
| 40:DO:90:GLN:O | 40:DO:91:LEU:HB2 | 1.96 | 0.64 |
| 41:DP:112:LEU:HD22 | 41:DP:113:LYS:N | 2.13 | 0.64 |
| 8:AH:97:VAL:O | 8:AH:100:ILE:HG13 | 1.96 | 0.64 |
| 9:AI:61:ALA:HB1 | 9:AI:63:ILE:HD11 | 1.78 | 0.64 |
| 12:AL:55:VAL:HG12 | 12:AL:56:ALA:H | 1.62 | 0.64 |
| 15:AO:3:ILE:HD13 | 15:AO:3:ILE:H | 1.63 | 0.64 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 28:B6:15:GLU:OE1 | 28:B6:43:CYS:HB2 | 1.98 | 0.64 |
| 31:BA:518:G:H2' | 31:BA:519:U:C6 | 2.32 | 0.64 |
| 31:BA:587:C:C4' | 31:BA:588:U:OP2 | 2.45 | 0.64 |
| 33:BD:43:ARG:HH11 | 33:BD:44:ASN:CG | 2.04 | 0.64 |
| 35:BF:18:ARG:CG | 35:BF:19:GLU:H | 2.10 | 0.64 |
| 38:BI:63:ALA:HB3 | 38:BI:64:GLU:OE1 | 1.97 | 0.64 |
| 42:BQ:54:MET:HG3 | 42:BQ:117:ALA:HB1 | 1.79 | 0.64 |
| 46:BU:12:ARG:O | 46:BU:15:LYS:HG2 | 1.98 | 0.64 |
| 47:BV:36:PRO:CG | 47:BV:62:LEU:HD11 | 2.27 | 0.64 |
| 49:BX:60:ARG:NH2 | 49:BX:74:PRO:HG2 | 2.13 | 0.64 |
| 1:CA:159:G:H2' | 1:CA:161:A:OP2 | 1.97 | 0.64 |
| 1:CA:1109:C:H2' | 1:CA:1110:A:O4' | 1.97 | 0.64 |
| 15:CO:87:ILE:HG22 | 15:CO:88:ARG:H | 1.60 | 0.64 |
| 23:D1:19:GLN:CD | 23:D1:44:PRO:HB3 | 2.21 | 0.64 |
| 28:D6:44:ARG:O | 28:D6:45:LYS:HG2 | 1.98 | 0.64 |
| 31:DA:1784:A:H4' | 31:DA:1785:A:O5' | 1.97 | 0.64 |
| 35:DF:65:TRP:HZ3 | 35:DF:73:ALA:O | 1.79 | 0.64 |
| 41:DP:17:LYS:O | 41:DP:17:LYS:HG2 | 1.96 | 0.64 |
| 42:DQ:34:LEU:HD11 | 42:DQ:129:THR:HB | 1.80 | 0.64 |
| 42:DQ:141:GLN:NE2 | 51:DZ:72:ARG:HG2 | 2.12 | 0.64 |
| 43:DR:67:LEU:HD13 | 43:DR:76:VAL:HG21 | 1.79 | 0.64 |
| 44:DS:16:ASN:O | 44:DS:19:LYS:HB3 | 1.97 | 0.64 |
| 49:DX:73:ARG:O | 49:DX:74:PRO:C | 2.41 | 0.64 |
| 50:DY:96:ILE:HB | 50:DY:99:CYS:HB3 | 1.79 | 0.64 |
| 51:DZ:128:VAL:HG23 | 51:DZ:160:GLY:O | 1.97 | 0.64 |
| 1:AA:15:G:H4' | 5:AE:24:ARG:NH1 | 2.10 | 0.64 |
| 1:AA:105:G:H2' | 1:AA:106:C:C6 | 2.32 | 0.64 |
| 1:AA:437:U:C5 | 1:AA:438:G:N7 | 2.65 | 0.64 |
| 1:AA:1342:C:H4' | 9:AI:125:TYR:HB3 | 1.78 | 0.64 |
| 4:AD:149:ALA:O | 4:AD:153:ARG:HG3 | 1.96 | 0.64 |
| 25:B3:54:VAL:CG1 | 25:B3:55:ARG:N | 2.59 | 0.64 |
| 30:B8:35:GLN:HA | 31:BA:2420:C:P | 2.38 | 0.64 |
| 31:BA:528:A:H5" | 31:BA:528:A:C8 | 2.32 | 0.64 |
| 31:BA:2712(A):A:H5" | 31:BA:2713:A:OP2 | 1.96 | 0.64 |
| 36:BG:47:LYS:HE2 | 36:BG:81:LYS:HB2 | 1.78 | 0.64 |
| 41:BP:71:VAL:CG1 | 41:BP:72:PRO:HD3 | 2.27 | 0.64 |
| 41:BP:86:LYS:HB3 | 41:BP:118:GLY:HA3 | 1.79 | 0.64 |
| 43:BR:87:TYR:O | 43:BR:89:ASP:N | 2.27 | 0.64 |
| 49:BX:35:THR:HB | 49:BX:75:ASP:OD2 | 1.97 | 0.64 |
| 49:BX:60:ARG:CZ | 49:BX:74:PRO:HG2 | 2.26 | 0.64 |
| 1:CA:376:G:OP2 | 16:CP:67:THR:HG21 | 1.98 | 0.64 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:437:U:H5" | 4:CD:155:LEU:HD13 | 1.79 | 0.64 |
| 2:CB:22:LYS:NZ | 2:CB:40:HIS:HE1 | 1.94 | 0.64 |
| 31:DA:588:U:OP2 | 31:DA:588:U:C6 | 2.50 | 0.64 |
| 31:DA:675:A:C4 | 31:DA:804:A:C2 | 2.85 | 0.64 |
| 31:DA:2280:G:C2' | 31:DA:2281:C:H5' | 2.28 | 0.64 |
| 38:DI:52:ARG:HA | 38:DI:52:ARG:NE | 2.11 | 0.64 |
| 39:DN:9:VAL:HG11 | 39:DN:39:ARG:NH2 | 2.12 | 0.64 |
| 39:DN:40:PRO:CA | 46:DU:64:ARG:NH2 | 2.58 | 0.64 |
| 45:DT:29:ARG:HG3 | 45:DT:84:GLN:HG3 | 1.79 | 0.64 |
| 46:DU:65:ILE:HG12 | 46:DU:96:ALA:HB1 | 1.79 | 0.64 |
| 1:AA:254:G:OP1 | 17:AQ:67:LYS:O | 2.15 | 0.64 |
| 1:AA:409:G:H2' | 1:AA:410:G:H5' | 1.79 | 0.64 |
| 1:AA:502:G:C2 | 1:AA:503:C:O2 | 2.50 | 0.64 |
| 1:AA:992:U:H1' | 1:AA:993:G:OP2 | 1.97 | 0.64 |
| 1:AA:1172:C:H2' | 1:AA:1173:G:H8 | 1.62 | 0.64 |
| 1:AA:1316:G:H2' | 1:AA:1317:C:H5" | 1.80 | 0.64 |
| 17:AQ:22:LEU:HD12 | 17:AQ:23:VAL:N | 2.13 | 0.64 |
| 24:B2:48:HIS:NE2 | 31:BA:75:G:H4' | 2.13 | 0.64 |
| 29:B7:16:HIS:HB2 | 29:B7:44:PRO:HG2 | 1.79 | 0.64 |
| 31:BA:196:A:O4' | 41:BP:46:LYS:HE2 | 1.96 | 0.64 |
| 31:BA:875:G:C4' | 51:BZ:170:THR:HG21 | 2.24 | 0.64 |
| 31:BA:2250:G:C4 | 42:BQ:82:ARG:HD2 | 2.32 | 0.64 |
| 37:BH:158:HIS:HE2 | 37:BH:170:ARG:CA | 2.11 | 0.64 |
| 41:BP:121:LYS:CD | 25:D3:2:PRO:HD3 | 2.28 | 0.64 |
| 41:BP:131:SER:O | 41:BP:133:SER:N | 2.30 | 0.64 |
| 45:BT:29:ARG:CG | 45:BT:84:GLN:HG3 | 2.28 | 0.64 |
| 1:CA:688:G:H2' | 1:CA:689:C:H6 | 1.62 | 0.64 |
| 2:CB:28:PHE:HD1 | 2:CB:190:THR:HG22 | 1.63 | 0.64 |
| 13:CM:97:PRO:C | 13:CM:98:VAL:HA | 2.23 | 0.64 |
| 17:CQ:22:LEU:HD12 | 17:CQ:23:VAL:N | 2.13 | 0.64 |
| 28:D6:10:LEU:N | 28:D6:10:LEU:HD22 | 2.10 | 0.64 |
| 31:DA:1225:G:OP1 | 47:DV:88:ARG:HB3 | 1.98 | 0.64 |
| 31:DA:1719:G:C2' | 31:DA:1720:U:H5' | 2.28 | 0.64 |
| 31:DA:2571:C:C5' | 31:DA:2572:A:H5" | 2.28 | 0.64 |
| 31:DA:2632:A:HO2' | 31:DA:2811:G:HO2' | 1.44 | 0.64 |
| 31:DA:2753:A:O2' | 31:DA:2754:U:H5' | 1.97 | 0.64 |
| 37:DH:84:SER:O | 37:DH:85:LYS:HB3 | 1.97 | 0.64 |
| 38:DI:63:ALA:HB3 | 38:DI:64:GLU:OE1 | 1.97 | 0.64 |
| 45:DT:29:ARG:HG3 | 45:DT:30:VAL:HG13 | 1.77 | 0.64 |
| 50:DY:45:VAL:CG1 | 50:DY:62:GLU:H | 2.10 | 0.64 |
| 1:AA:983:A:H2 | 1:AA:984:C:C5 | 2.16 | 0.64 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:1109:C:H2' | 1:AA:1110:A:O4' | 1.96 | 0.64 |
| 2:AB:20:GLU:HG3 | 2:AB:191:ASP:HB2 | 1.79 | 0.64 |
| 2:AB:22:LYS:HZ3 | 2:AB:22:LYS:HA | 1.62 | 0.64 |
| 2:AB:77:ALA:HB2 | 2:AB:211:ILE:HD13 | 1.79 | 0.64 |
| 2:AB:194:PRO:O | 2:AB:196:LEU:N | 2.30 | 0.64 |
| 9:AI:103:THR:HG22 | 9:AI:105:ASP:H | 1.62 | 0.64 |
| 16:AP:45:THR:HG23 | 16:AP:46:PRO:HD2 | 1.79 | 0.64 |
| 23:B1:26:ARG:HB3 | 23:B1:34:THR:HA | 1.79 | 0.64 |
| 29:B7:43:THR:HG23 | 29:B7:44:PRO:HD2 | 1.79 | 0.64 |
| 31:BA:102:G:O2' | 31:BA:103:A:P | 2.54 | 0.64 |
| 31:BA:966:G:H2' | 31:BA:967:C:C6 | 2.32 | 0.64 |
| 32:BB:74:U:C2' | 32:BB:75:G:H5" | 2.28 | 0.64 |
| 38:BI:109:ILE:HG22 | 38:BI:130:TYR:OH | 1.97 | 0.64 |
| 41:BP:17:LYS:O | 41:BP:17:LYS:CG | 2.44 | 0.64 |
| 45:BT:42:ILE:HD13 | 45:BT:83:ILE:HD11 | 1.80 | 0.64 |
| 46:BU:95:LEU:C | 46:BU:97:ASP:H | 2.06 | 0.64 |
| 47:BV:28:GLU:HG3 | 47:BV:29:PRO:HD3 | 1.77 | 0.64 |
| 50:BY:27:VAL:C | 50:BY:29:GLU:OE1 | 2.41 | 0.64 |
| 1:CA:250:A:H1' | 1:CA:251:G:OP2 | 1.97 | 0.64 |
| 1:CA:579:G:C4 | 1:CA:580:U:C5 | 2.85 | 0.64 |
| 1:CA:939:G:H5" | 7:CG:102:ARG:HH12 | 1.63 | 0.64 |
| 3:CC:73:PRO:O | 3:CC:76:VAL:HG22 | 1.97 | 0.64 |
| 9:CI:61:ALA:HB1 | 9:CI:63:ILE:HD11 | 1.78 | 0.64 |
| 12:CL:102:ARG:HH11 | 12:CL:102:ARG:HG2 | 1.61 | 0.64 |
| 13:CM:46:LYS:HG3 | 13:CM:47:ASP:H | 1.62 | 0.64 |
| 30:D8:32:LEU:HD23 | 30:D8:35:GLN:HA | 1.79 | 0.64 |
| 30:D8:47:LYS:HD2 | 30:D8:48:PHE:O | 1.96 | 0.64 |
| 31:DA:2199:A:H2' | 31:DA:2199:A:N3 | 2.12 | 0.64 |
| 31:DA:2759:G:C8 | 31:DA:2759:G:H5' | 2.33 | 0.64 |
| 31:DA:2807:G:H3' | 31:DA:2808:U:H5" | 1.79 | 0.64 |
| 31:DA:2876:G:H4' | 45:DT:3:ARG:HD3 | 1.79 | 0.64 |
| 33:DD:8:PRO:HB3 | 33:DD:14:ARG:HB3 | 1.78 | 0.64 |
| 34:DE:33:VAL:HG12 | 34:DE:90:THR:H | 1.62 | 0.64 |
| 40:DO:60:ALA:HB2 | 40:DO:86:ILE:HA | 1.78 | 0.64 |
| 42:DQ:8:LYS:HD2 | 42:DQ:9:TYR:N | 2.13 | 0.64 |
| 42:DQ:17:LEU:N | 42:DQ:17:LEU:HD23 | 2.12 | 0.64 |
| 45:DT:107:ASP:H | 45:DT:110:ILE:HG13 | 1.63 | 0.64 |
| 3:AC:104:GLN:NE2 | 3:AC:105:GLU:H | 1.95 | 0.64 |
| 4:AD:128:VAL:O | 4:AD:130:GLY:N | 2.30 | 0.64 |
| 27:B5:32:PRO:O | 27:B5:33:CYS:HB3 | 1.97 | 0.64 |
| 31:BA:1478:G:C2' | 31:BA:1479:G:H5' | 2.28 | 0.64 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2314:C:O2' | 31:BA:2315:G:H5' | 1.98 | 0.64 |
| 33:BD:267:SER:C | 33:BD:269:PHE:H | 2.03 | 0.64 |
| 42:BQ:8:LYS:CG | 42:BQ:9:TYR:H | 2.11 | 0.64 |
| 45:BT:78:LEU:O | 45:BT:79:HIS:ND1 | 2.31 | 0.64 |
| 48:BW:88:ARG:HB3 | 48:BW:92:ARG:HB3 | 1.78 | 0.64 |
| 1:CA:386:C:O2' | 1:CA:387:U:H5' | 1.97 | 0.64 |
| 6:CF:63:TYR:N | 6:CF:63:TYR:HD2 | 1.95 | 0.64 |
| 7:CG:40:ALA:O | 7:CG:44:TYR:CD1 | 2.51 | 0.64 |
| 10:CJ:39:PRO:HB3 | 10:CJ:70:ARG:HH12 | 1.62 | 0.64 |
| 18:CR:43:PHE:C | 18:CR:44:LEU:HD12 | 2.23 | 0.64 |
| 31:DA:996:A:C4' | 46:DU:92:ARG:NE | 2.58 | 0.64 |
| 34:DE:203:LYS:O | 34:DE:203:LYS:HD2 | 1.97 | 0.64 |
| 50:DY:47:LYS:HB3 | 50:DY:47:LYS:HZ3 | 1.61 | 0.64 |
| 10:AJ:30:SER:OG | 10:AJ:81:THR:HG22 | 1.96 | 0.64 |
| 28:B6:16:CYS:SG | 28:B6:48:VAL:HG23 | 2.37 | 0.64 |
| 28:B6:20:ASN:CG | 28:B6:21:TYR:N | 2.56 | 0.64 |
| 28:B6:29:ASN:O | 28:B6:30:THR:C | 2.41 | 0.64 |
| 28:B6:36:LEU:HD13 | 28:B6:50:ARG:CZ | 2.28 | 0.64 |
| 31:BA:1040:C:O2' | 31:BA:1041:C:P | 2.56 | 0.64 |
| 31:BA:1779:U:C5 | 31:BA:1784:A:N7 | 2.61 | 0.64 |
| 31:BA:2012:G:H4' | 48:BW:96:ILE:CD1 | 2.20 | 0.64 |
| 34:BE:154:LYS:HA | 34:BE:154:LYS:HE3 | 1.80 | 0.64 |
| 36:BG:137:GLU:HA | 36:BG:152:LEU:HD22 | 1.79 | 0.64 |
| 37:BH:68:THR:O | 37:BH:69:ARG:C | 2.41 | 0.64 |
| 42:BQ:54:MET:HB3 | 42:BQ:64:ILE:HD11 | 1.79 | 0.64 |
| 45:BT:65:LYS:CE | 45:BT:66:VAL:H | 2.10 | 0.64 |
| 5:CE:101:ILE:HD13 | 5:CE:118:ILE:O | 1.97 | 0.64 |
| 12:CL:55:VAL:HG12 | 12:CL:56:ALA:N | 2.12 | 0.64 |
| 24:D2:33:MET:HG2 | 49:DX:11:PRO:HD3 | 1.78 | 0.64 |
| 30:D8:35:GLN:HA | 31:DA:2420:C:OP2 | 1.98 | 0.64 |
| 31:DA:9:U:C4 | 31:DA:2629:A:N6 | 2.66 | 0.64 |
| 31:DA:329:G:OP2 | 50:DY:71:LYS:HE3 | 1.98 | 0.64 |
| 31:DA:918:A:H5" | 32:DB:98:G:O2' | 1.98 | 0.64 |
| 31:DA:1987:G:C8 | 31:DA:1987:G:H5' | 2.31 | 0.64 |
| 31:DA:2263:C:O2' | 31:DA:2264:C:H5' | 1.97 | 0.64 |
| 34:DE:38:THR:HG22 | 34:DE:40:GLU:N | 2.12 | 0.64 |
| 50:DY:46:LYS:HB2 | 50:DY:47:LYS:HE2 | 1.78 | 0.64 |
| 1:AA:1392:G:O2' | 1:AA:1393:U:H5' | 1.97 | 0.64 |
| 4:AD:8:VAL:HB | 4:AD:21:LEU:HD12 | 1.79 | 0.64 |
| 30:B8:39:LYS:CE | 30:B8:39:LYS:O | 2.46 | 0.64 |
| 31:BA:2759:G:C5' | 31:BA:2759:G:H8 | 2.11 | 0.64 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2807:G:H3' | 31:BA:2808:U:H5" | 1.79 | 0.64 |
| 33:BD:266:SER:O | 33:BD:267:SER:CB | 2.45 | 0.64 |
| 34:BE:111:ARG:HG3 | 43:BR:2:ARG:HG3 | 1.80 | 0.64 |
| 37:BH:30:LYS:HZ1 | 37:BH:81:GLU:HA | 1.63 | 0.64 |
| 37:BH:35:VAL:O | 37:BH:37:VAL:HG23 | 1.97 | 0.64 |
| 40:BO:2:ILE:HD12 | 40:BO:6:THR:HG21 | 1.80 | 0.64 |
| 45:BT:27:THR:O | 45:BT:28:VAL:HG23 | 1.97 | 0.64 |
| 49:BX:70:LEU:O | 49:BX:71:GLY:O | 2.14 | 0.64 |
| 50:BY:76:CYS:SG | 50:BY:77:PRO:CD | 2.82 | 0.64 |
| 51:BZ:101:PRO:HA | 51:BZ:123:ASP:HB3 | 1.79 | 0.64 |
| 4:CD:8:VAL:HB | 4:CD:21:LEU:HD12 | 1.80 | 0.64 |
| 6:CF:82:ARG:HA | 6:CF:82:ARG:HH11 | 1.63 | 0.64 |
| 30:D8:32:LEU:HB3 | 30:D8:34:TRP:N | 2.12 | 0.64 |
| 31:DA:1963:U:H4' | 31:DA:1964:G:OP1 | 1.98 | 0.64 |
| 40:DO:66:LYS:H | 40:DO:82:ASN:HD22 | 1.46 | 0.64 |
| 1:AA:1225:A:H2' | 1:AA:1225:A:N3 | 2.13 | 0.64 |
| 6:AF:18:GLN:HA | 6:AF:21:LEU:CD2 | 2.28 | 0.64 |
| 17:AQ:59:ILE:CG2 | 17:AQ:71:PHE:HB3 | 2.28 | 0.64 |
| 30:B8:4:MET:SD | 30:B8:61:LEU:HD12 | 2.38 | 0.64 |
| 31:BA:1747(A):G:C2' | 31:BA:1748:G:C5' | 2.76 | 0.64 |
| 31:BA:1986:A:O2' | 31:BA:1987:G:H5" | 1.98 | 0.64 |
| 31:BA:2661:G:C8 | 31:BA:2662:A:C2 | 2.85 | 0.64 |
| 34:BE:116:VAL:CG1 | 34:BE:122:PHE:CD2 | 2.81 | 0.64 |
| 35:BF:63:LYS:HE2 | 35:BF:67:GLN:CB | 2.27 | 0.64 |
| 36:BG:71:THR:HG22 | 36:BG:72:ARG:N | 2.13 | 0.64 |
| 41:BP:56:SER:O | 41:BP:58:THR:N | 2.31 | 0.64 |
| 44:BS:85:VAL:CG2 | 44:BS:106:ARG:HB2 | 2.19 | 0.64 |
| 45:BT:102:ILE:HB | 45:BT:110:ILE:HD13 | 1.81 | 0.64 |
| 50:BY:95:LYS:HE2 | 50:BY:101:LYS:HA | 1.79 | 0.64 |
| 1:CA:103:C:OP2 | 20:CT:14:LYS:HD3 | 1.97 | 0.64 |
| 15:CO:39:LEU:HD13 | 15:CO:56:LEU:HB2 | 1.80 | 0.64 |
| 25:D3:54:VAL:CG1 | 25:D3:55:ARG:N | 2.60 | 0.64 |
| 34:DE:23:VAL:HA | 34:DE:186:GLY:H | 1.63 | 0.64 |
| 41:DP:17:LYS:C | 41:DP:19:VAL:N | 2.53 | 0.64 |
| 48:DW:59:VAL:CG1 | 48:DW:60:ASN:N | 2.58 | 0.64 |
| 51:DZ:39:VAL:CG2 | 51:DZ:44:PHE:HB2 | 2.27 | 0.64 |
| 1:AA:952:U:H2' | 1:AA:953:G:H8 | 1.63 | 0.63 |
| 31:BA:49:A:C3' | 31:BA:50:U:H5' | 2.28 | 0.63 |
| 31:BA:1799:G:N7 | 33:BD:179:SER:OG | 2.23 | 0.63 |
| 31:BA:2199:A:H2' | 31:BA:2199:A:N3 | 2.13 | 0.63 |
| 31:BA:2571:C:C5' | 31:BA:2572:A:H5" | 2.28 | 0.63 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 33:BD:106:ILE:O | 33:BD:106:ILE:HD13 | 1.97 | 0.63 |
| 34:BE:46:ALA:HA | 34:BE:82:ARG:O | 1.97 | 0.63 |
| 41:BP:113:LYS:HE3 | 41:BP:130:PHE:O | 1.99 | 0.63 |
| 45:BT:107:ASP:H | 45:BT:110:ILE:HG13 | 1.63 | 0.63 |
| 47:BV:1:MET:CE | 47:BV:44:LYS:HB2 | 2.26 | 0.63 |
| 49:BX:65:ARG:CZ | 49:BX:66:LEU:H | 2.10 | 0.63 |
| 50:BY:60:PHE:HA | 50:BY:62:GLU:OE2 | 1.97 | 0.63 |
| 24:D2:26:ARG:NH1 | 24:D2:29:LYS:HE2 | 2.12 | 0.63 |
| 30:D8:59:LYS:HD2 | 41:DP:50:ARG:HB3 | 1.78 | 0.63 |
| 31:DA:404:C:H4' | 31:DA:405:U:H5' | 1.78 | 0.63 |
| 31:DA:588:U:OP2 | 31:DA:588:U:H6 | 1.80 | 0.63 |
| 31:DA:825:C:C2' | 31:DA:826:U:O5' | 2.46 | 0.63 |
| 31:DA:1171:G:C8 | 31:DA:1173:G:H1' | 2.33 | 0.63 |
| 31:DA:1209:G:H21 | 31:DA:1210:A:N6 | 1.95 | 0.63 |
| 31:DA:2632:A:N3 | 34:DE:61:ARG:NH1 | 2.46 | 0.63 |
| 33:DD:211:ARG:O | 33:DD:215:LEU:HG | 1.98 | 0.63 |
| 34:DE:137:HIS:HB3 | 34:DE:138:PRO:CD | 2.28 | 0.63 |
| 38:DI:28:ASN:C | 38:DI:32:PRO:HG2 | 2.22 | 0.63 |
| 41:DP:13:ASN:N | 41:DP:13:ASN:HD22 | 1.96 | 0.63 |
| 41:DP:107:LYS:O | 41:DP:109:GLY:N | 2.30 | 0.63 |
| 42:DQ:16:ARG:HB2 | 42:DQ:16:ARG:HH11 | 1.61 | 0.63 |
| 44:DS:66:ALA:O | 44:DS:67:ARG:CB | 2.45 | 0.63 |
| 47:DV:51:VAL:HG12 | 47:DV:52:VAL:N | 2.12 | 0.63 |
| 48:DW:13:SER:HB3 | 48:DW:16:LYS:HD3 | 1.78 | 0.63 |
| 48:DW:18:ARG:HH11 | 48:DW:18:ARG:CG | 2.12 | 0.63 |
| 3:AC:34:LEU:O | 3:AC:38:ARG:HG2 | 1.98 | 0.63 |
| 3:AC:43:LEU:O | 3:AC:47:LEU:HB3 | 1.97 | 0.63 |
| 4:AD:12:CYS:HA | 4:AD:19:LEU:CD1 | 2.28 | 0.63 |
| 9:AI:55:ALA:HB1 | 9:AI:58:ARG:HD2 | 1.79 | 0.63 |
| 23:B1:65:SER:H | 23:B1:67:ILE:CD1 | 2.12 | 0.63 |
| 31:BA:814:C:C5 | 41:BP:27:HIS:CE1 | 2.87 | 0.63 |
| 31:BA:1171:G:C8 | 31:BA:1173:G:H1' | 2.33 | 0.63 |
| 31:BA:1359:A:H2' | 31:BA:1360:A:H5' | 1.79 | 0.63 |
| 31:BA:2236:C:C2' | 31:BA:2237:G:H5' | 2.28 | 0.63 |
| 34:BE:24:THR:HG21 | 34:BE:188:VAL:HG13 | 1.79 | 0.63 |
| 41:BP:114:ILE:HG13 | 41:BP:115:LEU:N | 2.13 | 0.63 |
| 49:BX:83:VAL:O | 49:BX:84:ALA:HB3 | 1.96 | 0.63 |
| 1:CA:1096:C:H2' | 1:CA:1097:C:H6 | 1.63 | 0.63 |
| 1:CA:1316:G:H2' | 1:CA:1317:C:H5" | 1.79 | 0.63 |
| 3:CC:6:HIS:NE2 | 3:CC:184:TYR:HE2 | 1.96 | 0.63 |
| 5:CE:76:ILE:HG12 | 5:CE:77:PRO:HD2 | 1.79 | 0.63 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 16:CP:6:LEU:HD23 | 16:CP:17:TYR:CG | 2.33 | 0.63 |
| 31:DA:228:A:H2' | 31:DA:230:U:O4' | 1.98 | 0.63 |
| 31:DA:247:G:H4' | 31:DA:386:G:C5 | 2.33 | 0.63 |
| 31:DA:1204:A:N1 | 31:DA:1241:A:C2 | 2.66 | 0.63 |
| 31:DA:1526:G:C6 | 31:DA:1527:G:C2 | 2.87 | 0.63 |
| 31:DA:2464:C:O2' | 31:DA:2465:C:P | 2.56 | 0.63 |
| 33:DD:52:ARG:NH1 | 33:DD:53:PHE:HE2 | 1.96 | 0.63 |
| 33:DD:169:GLU:HG3 | 33:DD:174:ILE:HD11 | 1.78 | 0.63 |
| 43:DR:34:ILE:HG22 | 43:DR:114:VAL:HB | 1.80 | 0.63 |
| 45:DT:42:ILE:O | 45:DT:42:ILE:HG13 | 1.96 | 0.63 |
| 45:DT:65:LYS:CE | 45:DT:66:VAL:H | 2.11 | 0.63 |
| 47:DV:62:LEU:HD22 | 47:DV:98:GLU:CB | 2.28 | 0.63 |
| 51:DZ:19:ARG:HH11 | 51:DZ:19:ARG:HG2 | 1.62 | 0.63 |
| 12:AL:55:VAL:HG12 | 12:AL:56:ALA:N | 2.14 | 0.63 |
| 23:B1:17:SER:O | 23:B1:44:PRO:CD | 2.40 | 0.63 |
| 31:BA:659:C:C5' | 31:BA:659:C:C6 | 2.74 | 0.63 |
| 34:BE:14:ILE:HG12 | 34:BE:21:VAL:HG23 | 1.80 | 0.63 |
| 37:BH:155:SER:O | 37:BH:155:SER:OG | 2.15 | 0.63 |
| 38:BI:14:ASP:HB2 | 38:BI:17:GLN:OE1 | 1.98 | 0.63 |
| 38:BI:50:ARG:C | 38:BI:52:ARG:H | 2.06 | 0.63 |
| 39:BN:9:VAL:HG11 | 39:BN:39:ARG:NH2 | 2.14 | 0.63 |
| 39:BN:123:TYR:OH | 39:BN:130:HIS:CD2 | 2.51 | 0.63 |
| 41:BP:10:PRO:HD2 | 41:BP:11:GLY:N | 2.13 | 0.63 |
| 43:BR:5:LYS:N | 43:BR:5:LYS:CD | 2.62 | 0.63 |
| 44:BS:66:ALA:O | 44:BS:67:ARG:CB | 2.46 | 0.63 |
| 3:CC:20:SER:HB2 | 3:CC:40:ARG:NH2 | 2.08 | 0.63 |
| 23:D1:26:ARG:HB2 | 23:D1:34:THR:HA | 1.80 | 0.63 |
| 28:D6:15:GLU:CG | 28:D6:18:ARG:NE | 2.57 | 0.63 |
| 31:DA:298:G:H5" | 31:DA:299:A:OP1 | 1.97 | 0.63 |
| 31:DA:548:A:O2' | 31:DA:549:G:OP1 | 2.15 | 0.63 |
| 31:DA:1381:G:H2' | 31:DA:1382:G:H5' | 1.80 | 0.63 |
| 31:DA:1822:G:C5' | 31:DA:1822:G:H8 | 2.12 | 0.63 |
| 31:DA:2870:C:C2' | 31:DA:2871:C:H5' | 2.28 | 0.63 |
| 32:DB:66:A:C5 | 32:DB:109:C:C5 | 2.87 | 0.63 |
| 32:DB:75:G:H8 | 32:DB:75:G:C5' | 2.11 | 0.63 |
| 34:DE:10:GLY:HA3 | 45:DT:8:LYS:HE3 | 1.79 | 0.63 |
| 37:DH:17:VAL:HG21 | 37:DH:50:VAL:HG21 | 1.80 | 0.63 |
| 41:DP:105:LEU:HD12 | 41:DP:105:LEU:N | 2.13 | 0.63 |
| 42:DQ:78:PRO:O | 42:DQ:79:LEU:CB | 2.46 | 0.63 |
| 44:DS:61:ASN:HD22 | 44:DS:62:LYS:N | 1.86 | 0.63 |
| 51:DZ:101:PRO:HA | 51:DZ:123:ASP:HB3 | 1.79 | 0.63 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:392:G:H2' | 1:AA:393:A:C8 | 2.33 | 0.63 |
| 1:AA:437:U:O2' | 1:AA:438:G:H5' | 1.98 | 0.63 |
| 1:AA:1133:G:N3 | 1:AA:1142:G:N2 | 2.46 | 0.63 |
| 2:AB:19:HIS:O | 2:AB:39:ILE:HG23 | 1.98 | 0.63 |
| 6:AF:63:TYR:N | 6:AF:63:TYR:HD2 | 1.97 | 0.63 |
| 23:B1:16:ASN:HB3 | 23:B1:46:LEU:CG | 2.25 | 0.63 |
| 27:B5:50:GLY:O | 27:B5:51:TYR:CD1 | 2.50 | 0.63 |
| 30:B8:23:VAL:CG1 | 30:B8:46:ARG:HB3 | 2.29 | 0.63 |
| 31:BA:151:C:O2' | 31:BA:152:G:H5' | 1.99 | 0.63 |
| 31:BA:528:A:H5" | 31:BA:528:A:H8 | 1.64 | 0.63 |
| 31:BA:1603:A:H5' | 31:BA:1603:A:C8 | 2.29 | 0.63 |
| 33:BD:32:SER:C | 33:BD:33:LEU:HG | 2.23 | 0.63 |
| 40:BO:2:ILE:HG13 | 40:BO:8:LEU:HD11 | 1.79 | 0.63 |
| 1:CA:632:A:C8 | 1:CA:633:G:C8 | 2.85 | 0.63 |
| 1:CA:1160:G:H2' | 1:CA:1160:G:N3 | 2.13 | 0.63 |
| 16:CP:55:ARG:HA | 16:CP:55:ARG:HE | 1.62 | 0.63 |
| 20:CT:97:ALA:O | 20:CT:99:LEU:N | 2.31 | 0.63 |
| 24:D2:37:PHE:CE2 | 24:D2:40:SER:HA | 2.33 | 0.63 |
| 29:D7:43:THR:HG23 | 29:D7:44:PRO:HD2 | 1.79 | 0.63 |
| 31:DA:49:A:C4' | 31:DA:50:U:H5' | 2.28 | 0.63 |
| 31:DA:322:A:H5' | 31:DA:340:A:C1' | 2.28 | 0.63 |
| 31:DA:330:A:H2 | 31:DA:1210:A:HO2' | 1.46 | 0.63 |
| 31:DA:1654:A:H1' | 31:DA:2823:A:H5' | 1.80 | 0.63 |
| 33:DD:166:GLN:HA | 33:DD:166:GLN:NE2 | 2.12 | 0.63 |
| 36:DG:124:SER:HB2 | 36:DG:131:TYR:HE1 | 1.63 | 0.63 |
| 38:DI:14:ASP:HB2 | 38:DI:17:GLN:OE1 | 1.98 | 0.63 |
| 41:DP:131:SER:O | 41:DP:133:SER:N | 2.31 | 0.63 |
| 47:DV:25:LEU:H | 47:DV:94:LEU:HD13 | 1.63 | 0.63 |
| 49:DX:76:ARG:O | 49:DX:77:LYS:HB2 | 1.96 | 0.63 |
| 51:DZ:5:LEU:HD22 | 51:DZ:6:LYS:N | 2.14 | 0.63 |
| 1:AA:9:G:H5' | 5:AE:122:GLU:OE2 | 1.98 | 0.63 |
| 1:AA:357:G:O2' | 1:AA:358:U:H5' | 1.98 | 0.63 |
| 1:AA:952:U:H4' | 1:AA:964:A:H61 | 1.64 | 0.63 |
| 1:AA:1190:G:OP1 | 3:AC:4:LYS:HA | 1.99 | 0.63 |
| 2:AB:21:ARG:HB3 | 2:AB:39:ILE:HA | 1.81 | 0.63 |
| 3:AC:36:ASP:OD2 | 3:AC:57:ILE:HG21 | 1.98 | 0.63 |
| 18:AR:59:SER:H | 18:AR:62:GLU:CD | 2.06 | 0.63 |
| 23:B1:26:ARG:HB2 | 23:B1:34:THR:HA | 1.79 | 0.63 |
| 31:BA:626:U:H5" | 31:BA:627:A:H5' | 1.80 | 0.63 |
| 1:CA:779:C:O2' | 1:CA:780:A:H5' | 1.98 | 0.63 |
| 1:CA:1162:C:H2' | 1:CA:1163:C:C6 | 2.34 | 0.63 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 6:CF:63:TYR:N | 6:CF:63:TYR:CD2 | 2.66 | 0.63 |
| 28:D6:15:GLU:OE1 | 28:D6:43:CYS:HB2 | 1.98 | 0.63 |
| 31:DA:1902:C:H1' | 33:DD:244:ARG:HD3 | 1.81 | 0.63 |
| 31:DA:2208:A:H1' | 31:DA:2219:G:C5 | 2.33 | 0.63 |
| 32:DB:21:G:HO2' | 32:DB:22:U:H6 | 1.43 | 0.63 |
| 32:DB:29:A:C2 | 32:DB:30:C:C2 | 2.86 | 0.63 |
| 44:DS:42:ASP:C | 44:DS:44:LYS:H | 2.06 | 0.63 |
| 44:DS:67:ARG:H | 44:DS:69:VAL:HG12 | 1.63 | 0.63 |
| 50:DY:79:CYS:O | 50:DY:80:GLY:C | 2.41 | 0.63 |
| 51:DZ:111:VAL:HG13 | 51:DZ:112:ARG:N | 2.13 | 0.63 |
| 1:AA:673:G:H2' | 1:AA:674:G:H8 | 1.58 | 0.63 |
| 1:AA:877:C:H5" | 8:AH:88:LYS:CD | 2.28 | 0.63 |
| 3:AC:73:PRO:O | 3:AC:76:VAL:HG22 | 1.99 | 0.63 |
| 7:AG:26:PHE:O | 7:AG:30:ILE:HG12 | 1.98 | 0.63 |
| 31:BA:1925:C:O2' | 31:BA:1926:U:H5' | 1.99 | 0.63 |
| 31:BA:2236:C:H2' | 31:BA:2237:G:H5' | 1.80 | 0.63 |
| 31:BA:2580:U:H5" | 34:BE:131:ALA:H | 1.64 | 0.63 |
| 33:BD:218:ARG:HB3 | 33:BD:219:PRO:HD2 | 1.80 | 0.63 |
| 36:BG:76:SER:CB | 36:BG:83:ARG:HB3 | 2.29 | 0.63 |
| 37:BH:32:GLU:O | 37:BH:33:LEU:HD23 | 1.99 | 0.63 |
| 38:BI:64:GLU:HA | 38:BI:67:ARG:HB2 | 1.81 | 0.63 |
| 38:BI:110:ASP:C | 38:BI:112:LYS:H | 2.04 | 0.63 |
| 39:BN:40:PRO:C | 46:BU:64:ARG:NH2 | 2.57 | 0.63 |
| 41:BP:85:LEU:HD23 | 41:BP:85:LEU:H | 1.64 | 0.63 |
| 50:BY:96:ILE:HB | 50:BY:99:CYS:HB3 | 1.80 | 0.63 |
| 1:CA:579:G:H2' | 1:CA:580:U:C6 | 2.33 | 0.63 |
| 2:CB:67:THR:HG21 | 2:CB:155:LEU:HG | 1.79 | 0.63 |
| 2:CB:194:PRO:O | 2:CB:196:LEU:N | 2.31 | 0.63 |
| 6:CF:69:GLU:CD | 6:CF:69:GLU:H | 2.06 | 0.63 |
| 18:CR:59:SER:H | 18:CR:62:GLU:CD | 2.07 | 0.63 |
| 31:DA:669:G:O2' | 31:DA:669:G:C8 | 2.49 | 0.63 |
| 31:DA:2016:U:H2' | 31:DA:2017:U:C6 | 2.33 | 0.63 |
| 31:DA:2250:G:C6 | 42:DQ:82:ARG:HD2 | 2.32 | 0.63 |
| 31:DA:2632:A:H1' | 34:DE:61:ARG:NH1 | 2.13 | 0.63 |
| 33:DD:65:ILE:HD11 | 33:DD:67:PHE:CD1 | 2.33 | 0.63 |
| 33:DD:106:ILE:HD13 | 33:DD:106:ILE:O | 1.97 | 0.63 |
| 36:DG:18:GLU:HG2 | 36:DG:175:LEU:HD21 | 1.79 | 0.63 |
| 38:DI:64:GLU:HA | 38:DI:67:ARG:HB2 | 1.81 | 0.63 |
| 42:DQ:37:LEU:HB2 | 42:DQ:128:LYS:O | 1.98 | 0.63 |
| 1:AA:376:G:H2' | 1:AA:377:G:H8 | 1.63 | 0.63 |
| 1:AA:579:G:H2' | 1:AA:580:U:H6 | 1.64 | 0.63 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:674:G:H2' | 1:AA:675:A:H8 | 1.64 | 0.63 |
| 23:B1:19:GLN:CD | 23:B1:44:PRO:HB3 | 2.24 | 0.63 |
| 27:B5:57:VAL:HG23 | 27:B5:58:LEU:N | 2.11 | 0.63 |
| 31:BA:84:A:C5' | 50:BY:9:LYS:HD2 | 2.29 | 0.63 |
| 31:BA:573:G:N1 | 31:BA:2030:A:H3' | 2.14 | 0.63 |
| 31:BA:1636:C:H2' | 31:BA:1637:A:H8 | 1.64 | 0.63 |
| 31:BA:2713:A:H3' | 31:BA:2714:G:H5' | 1.81 | 0.63 |
| 36:BG:7:LEU:CD2 | 36:BG:176:LEU:HD22 | 2.29 | 0.63 |
| 37:BH:106:THR:HG22 | 37:BH:112:PRO:HB3 | 1.78 | 0.63 |
| 37:BH:138:LYS:C | 37:BH:140:LYS:N | 2.55 | 0.63 |
| 37:BH:138:LYS:O | 37:BH:139:GLN:C | 2.41 | 0.63 |
| 39:BN:128:HIS:NE2 | 39:BN:131:GLN:HB2 | 2.14 | 0.63 |
| 47:BV:79:VAL:O | 47:BV:80:GLN:CB | 2.42 | 0.63 |
| 1:CA:992:U:H1' | 1:CA:993:G:OP2 | 1.98 | 0.63 |
| 1:CA:1497:G:H2' | 1:CA:1498:U:H5' | 1.80 | 0.63 |
| 17:CQ:59:ILE:CG2 | 17:CQ:71:PHE:HB3 | 2.28 | 0.63 |
| 29:D7:5:TRP:NE1 | 29:D7:7:PRO:HG3 | 2.13 | 0.63 |
| 31:DA:107:C:H2' | 31:DA:108:U:H6 | 1.62 | 0.63 |
| 31:DA:528:A:H8 | 31:DA:528:A:H5" | 1.64 | 0.63 |
| 31:DA:867:C:C5 | 31:DA:868:U:C5 | 2.86 | 0.63 |
| 31:DA:1162:G:H1' | 47:DV:91:TYR:OH | 1.98 | 0.63 |
| 31:DA:1558:A:H1' | 31:DA:1559:G:OP2 | 1.98 | 0.63 |
| 31:DA:1796:U:H2' | 31:DA:1797:C:C6 | 2.34 | 0.63 |
| 31:DA:1839:G:C8 | 31:DA:1927:A:H1' | 2.33 | 0.63 |
| 31:DA:2402:C:C2' | 31:DA:2403:C:H5' | 2.27 | 0.63 |
| 31:DA:2483:C:O2 | 31:DA:2483:C:H2' | 1.99 | 0.63 |
| 31:DA:2829:C:C2' | 31:DA:2830:G:H5" | 2.27 | 0.63 |
| 33:DD:35:LYS:HA | 33:DD:64:ILE:HG22 | 1.79 | 0.63 |
| 44:DS:92:TYR:HD1 | 44:DS:93:LYS:N | 1.86 | 0.63 |
| 47:DV:18:LEU:HD13 | 47:DV:98:GLU:OE1 | 1.97 | 0.63 |
| 51:DZ:150:LEU:N | 51:DZ:150:LEU:HD13 | 2.13 | 0.63 |
| 1:AA:159:G:H2' | 1:AA:161:A:OP2 | 1.99 | 0.63 |
| 1:AA:192:U:O4' | 20:AT:103:GLY:HA2 | 1.98 | 0.63 |
| 1:AA:946:A:H2' | 1:AA:947:G:C8 | 2.33 | 0.63 |
| 1:AA:949:A:H1' | 1:AA:1364:U:H3 | 1.64 | 0.63 |
| 10:AJ:39:PRO:HB3 | 10:AJ:70:ARG:HH12 | 1.63 | 0.63 |
| 23:B1:64:ALA:HA | 23:B1:67:ILE:CG1 | 2.29 | 0.63 |
| 31:BA:774:A:H2 | 31:BA:787:U:HO2' | 1.46 | 0.63 |
| 33:BD:169:GLU:HG3 | 33:BD:174:ILE:HD11 | 1.81 | 0.63 |
| 34:BE:60:ASN:N | 34:BE:60:ASN:HD22 | 1.97 | 0.63 |
| 35:BF:22:ALA:HB1 | 35:BF:26:ALA:HB1 | 1.79 | 0.63 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:659:U:H2' | 1:CA:660:G:H5' | 1.80 | 0.63 |
| 1:CA:1072:G:C5 | 1:CA:1073:U:C4 | 2.87 | 0.63 |
| 1:CA:1190:G:OP1 | 3:CC:4:LYS:HA | 1.98 | 0.63 |
| 30:D8:53:PRO:HA | 30:D8:56:GLU:HB2 | 1.80 | 0.63 |
| 31:DA:445:C:O2' | 31:DA:446:G:H5' | 1.99 | 0.63 |
| 31:DA:573:G:N1 | 31:DA:2030:A:H3' | 2.14 | 0.63 |
| 31:DA:942:G:H5' | 41:DP:35:HIS:HB2 | 1.81 | 0.63 |
| 31:DA:1747(A):G:C2' | 31:DA:1748:G:C5' | 2.76 | 0.63 |
| 31:DA:1783:A:C2 | 31:DA:2587:A:C4 | 2.86 | 0.63 |
| 31:DA:2865:U:C4 | 31:DA:2866:U:C4 | 2.87 | 0.63 |
| 33:DD:35:LYS:NZ | 33:DD:64:ILE:O | 2.25 | 0.63 |
| 33:DD:267:SER:C | 33:DD:269:PHE:H | 2.07 | 0.63 |
| 49:DX:78:LYS:HD3 | 49:DX:78:LYS:O | 1.99 | 0.63 |
| 51:DZ:106:GLY:HA3 | 51:DZ:142:SER:HB3 | 1.80 | 0.63 |
| 1:AA:973:G:H3' | 1:AA:974:A:H5" | 1.81 | 0.63 |
| 3:AC:6:HIS:NE2 | 3:AC:184:TYR:HE2 | 1.97 | 0.63 |
| 8:AH:77:GLU:HG3 | 8:AH:78:GLN:N | 2.13 | 0.63 |
| 9:AI:97:LYS:HB3 | 9:AI:98:PRO:HD3 | 1.80 | 0.63 |
| 12:AL:102:ARG:HH11 | 12:AL:102:ARG:HG3 | 1.62 | 0.63 |
| 16:AP:55:ARG:HA | 16:AP:55:ARG:HE | 1.64 | 0.63 |
| 19:AS:19:VAL:HG12 | 19:AS:44:MET:HE2 | 1.80 | 0.63 |
| 30:B8:25:MET:HB2 | 41:BP:62:LEU:HD23 | 1.81 | 0.63 |
| 31:BA:795:C:H2' | 31:BA:796:C:C6 | 2.34 | 0.63 |
| 33:BD:9:TYR:O | 33:BD:10:THR:HG22 | 1.99 | 0.63 |
| 34:BE:1:MET:CB | 34:BE:84:PHE:HB2 | 2.29 | 0.63 |
| 42:BQ:78:PRO:O | 42:BQ:79:LEU:CB | 2.47 | 0.63 |
| 48:BW:73:ALA:HB3 | 48:BW:106:ILE:HD11 | 1.78 | 0.63 |
| 48:BW:95:ILE:O | 48:BW:95:ILE:HG13 | 1.99 | 0.63 |
| 1:CA:108:G:O6 | 20:CT:15:ARG:HD2 | 1.99 | 0.63 |
| 2:CB:97:TRP:HZ2 | 2:CB:102:LEU:HD13 | 1.62 | 0.63 |
| 7:CG:37:ASN:HD21 | 9:CI:40:LEU:HD22 | 1.64 | 0.63 |
| 16:CP:20:VAL:HG23 | 16:CP:35:LYS:HA | 1.79 | 0.63 |
| 31:DA:1022:G:C6 | 31:DA:1140:C:C4 | 2.86 | 0.63 |
| 31:DA:1106:A:C2' | 31:DA:1107:G:O5' | 2.47 | 0.63 |
| 31:DA:1484:G:C2' | 31:DA:1485:G:O5' | 2.47 | 0.63 |
| 31:DA:1836:C:O2' | 31:DA:1837:C:H5' | 1.98 | 0.63 |
| 31:DA:2208:A:H1' | 31:DA:2219:G:C4 | 2.34 | 0.63 |
| 34:DE:61:ARG:C | 34:DE:63:LEU:H | 2.07 | 0.63 |
| 34:DE:154:LYS:HE3 | 34:DE:154:LYS:HA | 1.79 | 0.63 |
| 40:DO:35:VAL:HG13 | 40:DO:65:THR:HG22 | 1.81 | 0.63 |
| 41:DP:48:PRO:O | 41:DP:50:ARG:N | 2.31 | 0.63 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 47:DV:28:GLU:HG3 | 47:DV:29:PRO:HD3 | 1.80 | 0.63 |
| 47:DV:89:GLN:HE21 | 47:DV:91:TYR:HB2 | 1.64 | 0.63 |
| 1:AA:353:A:H5' | 1:AA:353:A:C8 | 2.29 | 0.62 |
| 1:AA:561:U:O2' | 1:AA:562:C:P | 2.57 | 0.62 |
| 1:AA:963:G:H21 | 10:AJ:55:LYS:CE | 2.11 | 0.62 |
| 2:AB:14:GLY:O | 2:AB:15:VAL:HG13 | 1.99 | 0.62 |
| 12:AL:8:ASN:ND2 | 17:AQ:34:LYS:HE2 | 2.11 | 0.62 |
| 16:AP:76:GLN:O | 16:AP:76:GLN:HG2 | 1.99 | 0.62 |
| 31:BA:244:A:C2 | 31:BA:255:A:C4 | 2.87 | 0.62 |
| 31:BA:384:U:H2' | 31:BA:385:C:H6 | 1.63 | 0.62 |
| 31:BA:979:G:H3' | 31:BA:980:A:H5" | 1.81 | 0.62 |
| 31:BA:1405:U:H2' | 31:BA:1406:U:H6 | 1.61 | 0.62 |
| 31:BA:2402:C:H2' | 31:BA:2403:C:H5' | 1.81 | 0.62 |
| 33:BD:158:ALA:O | 33:BD:159:ALA:CB | 2.47 | 0.62 |
| 36:BG:124:SER:HB2 | 36:BG:131:TYR:HE1 | 1.61 | 0.62 |
| 37:BH:153:LYS:HB2 | 37:BH:154:PRO:HD3 | 1.81 | 0.62 |
| 3:CC:36:ASP:OD2 | 3:CC:57:ILE:HG21 | 1.99 | 0.62 |
| 30:D8:25:MET:HB2 | 41:DP:62:LEU:CD2 | 2.28 | 0.62 |
| 31:DA:543:C:H42 | 31:DA:551:G:H1 | 1.47 | 0.62 |
| 31:DA:1021:A:C8 | 31:DA:1021:A:C3' | 2.81 | 0.62 |
| 31:DA:1484:G:N2 | 31:DA:1505:C:N4 | 2.34 | 0.62 |
| 31:DA:1652:A:O2' | 31:DA:1653:G:H5' | 1.98 | 0.62 |
| 31:DA:1884:A:H2' | 31:DA:1885:A:H5' | 1.81 | 0.62 |
| 31:DA:1973:G:H2' | 31:DA:1974:C:C6 | 2.33 | 0.62 |
| 31:DA:2475:C:C5' | 31:DA:2476:A:OP2 | 2.47 | 0.62 |
| 31:DA:2656:U:N3 | 31:DA:2665:A:H2 | 1.90 | 0.62 |
| 31:DA:2759:G:H8 | 31:DA:2759:G:C5' | 2.11 | 0.62 |
| 32:DB:2:C:H2' | 32:DB:3:C:H6 | 1.63 | 0.62 |
| 35:DF:101:LEU:CD1 | 35:DF:102:PRO:HD2 | 2.14 | 0.62 |
| 45:DT:31:SER:C | 45:DT:32:TYR:CD2 | 2.77 | 0.62 |
| 46:DU:28:ARG:HG2 | 46:DU:38:THR:OG1 | 1.99 | 0.62 |
| 1:AA:52:G:O2' | 1:AA:53:A:H5' | 1.98 | 0.62 |
| 1:AA:437:U:H5" | 4:AD:155:LEU:HD13 | 1.81 | 0.62 |
| 1:AA:1072:G:C5 | 1:AA:1073:U:C4 | 2.87 | 0.62 |
| 3:AC:111:LEU:HD21 | 3:AC:145:GLY:O | 1.99 | 0.62 |
| 24:B2:41:ILE:O | 24:B2:42:GLY:C | 2.42 | 0.62 |
| 30:B8:61:LEU:N | 30:B8:63:PRO:HD2 | 2.14 | 0.62 |
| 31:BA:128:C:H2' | 31:BA:129:C:O4' | 1.99 | 0.62 |
| 31:BA:547:A:O2' | 31:BA:548:A:OP2 | 2.16 | 0.62 |
| 31:BA:1209:G:H21 | 31:BA:1210:A:H62 | 1.47 | 0.62 |
| 31:BA:1884:A:H2' | 31:BA:1885:A:H5' | 1.81 | 0.62 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1956:U:C2' | 31:BA:1957:C:H5' | 2.29 | 0.62 |
| 31:BA:2464:C:O2' | 31:BA:2465:C:OP2 | 2.17 | 0.62 |
| 40:BO:66:LYS:H | 40:BO:82:ASN:HD22 | 1.46 | 0.62 |
| 42:BQ:82:ARG:HG2 | 42:BQ:82:ARG:HH11 | 1.64 | 0.62 |
| 50:BY:29:GLU:OE1 | 50:BY:29:GLU:N | 2.31 | 0.62 |
| 50:BY:37:VAL:HG13 | 50:BY:69:ALA:HA | 1.81 | 0.62 |
| 1:CA:877:C:H5" | 8:CH:88:LYS:CD | 2.28 | 0.62 |
| 1:CA:1321:C:H5' | 1:CA:1322:C:H5" | 1.80 | 0.62 |
| 2:CB:22:LYS:HZ3 | 2:CB:22:LYS:HA | 1.63 | 0.62 |
| 12:CL:91:LYS:O | 12:CL:91:LYS:HG3 | 1.98 | 0.62 |
| 13:CM:34:LEU:HD13 | 13:CM:41:PRO:HG3 | 1.81 | 0.62 |
| 19:CS:19:VAL:HG12 | 19:CS:44:MET:HE2 | 1.81 | 0.62 |
| 24:D2:55:ARG:HH22 | 49:DX:3:THR:CG2 | 2.10 | 0.62 |
| 27:D5:51:TYR:HB3 | 27:D5:52:TYR:CD2 | 2.34 | 0.62 |
| 28:D6:18:ARG:CG | 28:D6:19:ARG:H | 2.09 | 0.62 |
| 31:DA:528:A:C2 | 31:DA:2042:A:H2' | 2.34 | 0.62 |
| 31:DA:1218:C:C2' | 31:DA:1219:G:H5' | 2.28 | 0.62 |
| 31:DA:2199:A:H1' | 38:DI:28:ASN:ND2 | 2.13 | 0.62 |
| 31:DA:2469:A:H2 | 31:DA:2481:G:N2 | 1.95 | 0.62 |
| 38:DI:133:HIS:CB | 38:DI:134:PRO:CD | 2.77 | 0.62 |
| 39:DN:19:GLU:HG3 | 39:DN:20:GLY:H | 1.65 | 0.62 |
| 41:DP:86:LYS:HB3 | 41:DP:118:GLY:HA3 | 1.79 | 0.62 |
| 47:DV:80:GLN:OE1 | 47:DV:80:GLN:O | 2.17 | 0.62 |
| 51:DZ:74:VAL:HG22 | 51:DZ:86:VAL:HG12 | 1.81 | 0.62 |
| 1:AA:963:G:H21 | 10:AJ:55:LYS:HE2 | 1.64 | 0.62 |
| 1:AA:1097:C:H1' | 1:AA:1170:A:H1' | 1.81 | 0.62 |
| 13:AM:97:PRO:C | 13:AM:98:VAL:HA | 2.23 | 0.62 |
| 24:B2:55:ARG:HH22 | 49:BX:3:THR:CG2 | 2.12 | 0.62 |
| 28:B6:42:TRP:CZ2 | 31:BA:642:G:O3' | 2.52 | 0.62 |
| 28:B6:44:ARG:O | 28:B6:45:LYS:HG2 | 1.98 | 0.62 |
| 31:BA:588:U:OP2 | 31:BA:588:U:H6 | 1.82 | 0.62 |
| 31:BA:1839:G:C8 | 31:BA:1927:A:H1' | 2.35 | 0.62 |
| 31:BA:2517:C:C2 | 31:BA:2542:A:N6 | 2.67 | 0.62 |
| 33:BD:27:THR:HG22 | 33:BD:28:GLU:H | 1.63 | 0.62 |
| 34:BE:116:VAL:HG13 | 34:BE:122:PHE:CG | 2.34 | 0.62 |
| 35:BF:22:ALA:HA | 35:BF:26:ALA:CB | 2.28 | 0.62 |
| 35:BF:205:ARG:C | 35:BF:206:ILE:HG13 | 2.25 | 0.62 |
| 39:BN:4:TYR:CD1 | 39:BN:4:TYR:N | 2.66 | 0.62 |
| 41:BP:65:ARG:HB2 | 41:BP:65:ARG:NH1 | 2.14 | 0.62 |
| 41:BP:115:LEU:HA | 41:BP:134:ALA:HB2 | 1.81 | 0.62 |
| 50:BY:88:LYS:NZ | 50:BY:93:GLY:HA3 | 2.13 | 0.62 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:322:C:H5 | 1:CA:328:C:H5 | 1.47 | 0.62 |
| 1:CA:437:U:C5 | 1:CA:438:G:N7 | 2.67 | 0.62 |
| 16:CP:4:ILE:HB | 16:CP:66:PRO:HB3 | 1.79 | 0.62 |
| 27:D5:4:HIS:CB | 27:D5:5:PRO:HD3 | 2.28 | 0.62 |
| 31:DA:1171:G:H8 | 31:DA:1171:G:OP2 | 1.82 | 0.62 |
| 31:DA:1987:G:H5' | 31:DA:1987:G:H8 | 1.63 | 0.62 |
| 31:DA:2030:A:H5" | 31:DA:2030:A:H8 | 1.64 | 0.62 |
| 31:DA:2713:A:H3' | 31:DA:2714:G:H5' | 1.81 | 0.62 |
| 38:DI:51:ILE:O | 38:DI:51:ILE:CG2 | 2.47 | 0.62 |
| 42:DQ:54:MET:HB3 | 42:DQ:64:ILE:HD11 | 1.81 | 0.62 |
| 51:DZ:29:TYR:HE2 | 51:DZ:87:ASP:HB2 | 1.64 | 0.62 |
| 1:AA:64:G:H4' | 1:AA:65:U:H5" | 1.81 | 0.62 |
| 31:BA:272(G):C:H42 | 31:BA:363(C):G:H1 | 1.46 | 0.62 |
| 31:BA:1210:A:H5" | 31:BA:1211:U:H3' | 1.82 | 0.62 |
| 31:BA:1526:G:C6 | 31:BA:1527:G:C2 | 2.87 | 0.62 |
| 39:BN:16:ILE:O | 39:BN:54:VAL:HA | 2.00 | 0.62 |
| 45:BT:28:VAL:HG22 | 45:BT:46:GLU:CA | 2.30 | 0.62 |
| 45:BT:38:ASN:HD22 | 45:BT:40:THR:H | 1.46 | 0.62 |
| 1:CA:78:G:H1 | 1:CA:91:C:H42 | 1.47 | 0.62 |
| 1:CA:538:G:OP2 | 12:CL:115:LYS:HG3 | 2.00 | 0.62 |
| 1:CA:833:U:H2' | 1:CA:834:C:H6 | 1.64 | 0.62 |
| 1:CA:952:U:H4' | 1:CA:964:A:H61 | 1.65 | 0.62 |
| 1:CA:999:C:H2' | 1:CA:1000:U:C6 | 2.34 | 0.62 |
| 18:CR:56:THR:OG1 | 18:CR:58:LEU:HD13 | 1.98 | 0.62 |
| 23:D1:30:VAL:HG12 | 23:D1:30:VAL:O | 1.99 | 0.62 |
| 23:D1:65:SER:H | 23:D1:67:ILE:CD1 | 2.12 | 0.62 |
| 31:DA:2275:C:O2 | 42:DQ:83:MET:HG3 | 2.00 | 0.62 |
| 33:DD:266:SER:O | 33:DD:267:SER:CB | 2.46 | 0.62 |
| 42:DQ:20:ALA:HB2 | 42:DQ:99:PRO:CD | 2.29 | 0.62 |
| 51:DZ:101:PRO:O | 51:DZ:102:LEU:HD23 | 2.00 | 0.62 |
| 1:AA:523:A:H61 | 12:AL:53:ARG:HH12 | 1.47 | 0.62 |
| 1:AA:709:G:H2' | 1:AA:710:G:H8 | 1.65 | 0.62 |
| 1:AA:750:G:N3 | 15:AO:23:GLY:HA3 | 2.15 | 0.62 |
| 2:AB:97:TRP:HZ2 | 2:AB:102:LEU:HD13 | 1.64 | 0.62 |
| 5:AE:57:LYS:HB3 | 5:AE:61:TYR:HE2 | 1.64 | 0.62 |
| 24:B2:41:ILE:HG13 | 31:BA:95:G:N3 | 2.15 | 0.62 |
| 31:BA:626:U:H3 | 41:BP:105:LEU:HG | 1.65 | 0.62 |
| 31:BA:1449:A:HO2' | 31:BA:1530:C:H5 | 1.45 | 0.62 |
| 31:BA:1481:U:H5' | 31:BA:1482:G:OP2 | 1.98 | 0.62 |
| 31:BA:2656:U:N3 | 31:BA:2665:A:H2 | 1.92 | 0.62 |
| 32:BB:7:G:H4' | 44:BS:29:PHE:CE1 | 2.35 | 0.62 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 34:BE:147:PRO:HB2 | 34:BE:149:ARG:HG2 | 1.80 | 0.62 |
| 35:BF:141:ALA:O | 35:BF:144:LYS:HB3 | 1.99 | 0.62 |
| 47:BV:25:LEU:H | 47:BV:94:LEU:HD13 | 1.62 | 0.62 |
| 1:CA:503:C:H2' | 1:CA:504:C:C6 | 2.34 | 0.62 |
| 1:CA:983:A:H2 | 1:CA:984:C:C6 | 2.17 | 0.62 |
| 1:CA:1172:C:H2' | 1:CA:1173:G:H8 | 1.61 | 0.62 |
| 1:CA:1392:G:O2' | 1:CA:1393:U:H5' | 1.99 | 0.62 |
| 11:CK:69:ALA:HB1 | 11:CK:103:LEU:HD23 | 1.79 | 0.62 |
| 26:D4:11:PRO:C | 26:D4:13:ARG:H | 2.08 | 0.62 |
| 28:D6:42:TRP:CZ2 | 31:DA:642:G:O3' | 2.53 | 0.62 |
| 30:D8:32:LEU:HD23 | 30:D8:35:GLN:N | 2.14 | 0.62 |
| 31:DA:2406:U:O4 | 41:DP:70:GLN:HB3 | 1.99 | 0.62 |
| 31:DA:2532:G:O2' | 31:DA:2657:A:N6 | 2.33 | 0.62 |
| 31:DA:2859:G:H3' | 31:DA:2859:G:C8 | 2.35 | 0.62 |
| 33:DD:144:ALA:HB3 | 33:DD:192:THR:HG23 | 1.82 | 0.62 |
| 36:DG:71:THR:HG22 | 36:DG:72:ARG:N | 2.14 | 0.62 |
| 46:DU:90:VAL:O | 46:DU:92:ARG:N | 2.31 | 0.62 |
| 49:DX:30:VAL:HG23 | 49:DX:76:ARG:HA | 1.80 | 0.62 |
| 1:AA:78:G:H22 | 1:AA:91:C:H42 | 1.47 | 0.62 |
| 1:AA:184:G:H2' | 1:AA:185:A:H8 | 1.63 | 0.62 |
| 2:AB:233:SER:HB2 | 2:AB:234:PRO:HD2 | 1.79 | 0.62 |
| 3:AC:134:ILE:HG23 | 3:AC:151:VAL:HB | 1.81 | 0.62 |
| 9:AI:11:LYS:HG2 | 9:AI:11:LYS:O | 1.99 | 0.62 |
| 9:AI:114:TYR:N | 9:AI:114:TYR:CD2 | 2.67 | 0.62 |
| 9:AI:114:TYR:HD1 | 10:AJ:60:ARG:HG2 | 1.64 | 0.62 |
| 11:AK:20:TYR:C | 11:AK:21:ILE:HD12 | 2.24 | 0.62 |
| 15:AO:87:ILE:HG22 | 15:AO:88:ARG:H | 1.64 | 0.62 |
| 16:AP:20:VAL:HG23 | 16:AP:35:LYS:HA | 1.79 | 0.62 |
| 23:B1:64:ALA:HA | 23:B1:67:ILE:HG13 | 1.81 | 0.62 |
| 24:B2:33:MET:HG2 | 49:BX:11:PRO:CD | 2.29 | 0.62 |
| 29:B7:40:TRP:CD2 | 31:BA:459:U:H5" | 2.34 | 0.62 |
| 31:BA:2199:A:H1' | 38:BI:28:ASN:ND2 | 2.14 | 0.62 |
| 31:BA:2875:C:O2' | 45:BT:5:ALA:HB3 | 2.00 | 0.62 |
| 33:BD:80:ALA:HB3 | 33:BD:94:LEU:HD13 | 1.80 | 0.62 |
| 34:BE:36:ARG:NH2 | 34:BE:88:GLY:CA | 2.60 | 0.62 |
| 35:BF:154:VAL:HB | 35:BF:173:VAL:HG22 | 1.81 | 0.62 |
| 37:BH:111:HIS:ND1 | 37:BH:112:PRO:HD2 | 2.14 | 0.62 |
| 1:CA:78:G:H22 | 1:CA:91:C:H42 | 1.47 | 0.62 |
| 6:CF:10:LEU:HD12 | 6:CF:10:LEU:N | 2.13 | 0.62 |
| 24:D2:26:ARG:HG2 | 49:DX:5:TYR:O | 2.00 | 0.62 |
| 30:D8:2:PRO:N | 31:DA:591:C:H1' | 2.15 | 0.62 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:966:G:C4 | 31:DA:967:C:C5 | 2.87 | 0.62 |
| 31:DA:2272:U:H5" | 31:DA:2273:A:OP1 | 1.99 | 0.62 |
| 31:DA:2476:A:C2 | 31:DA:2477:C:C6 | 2.88 | 0.62 |
| 34:DE:63:LEU:O | 34:DE:64:LYS:C | 2.42 | 0.62 |
| 34:DE:111:ARG:HG3 | 43:DR:2:ARG:HG3 | 1.82 | 0.62 |
| 44:DS:74:ALA:HB1 | 44:DS:103:GLU:CB | 2.29 | 0.62 |
| 1:AA:262:A:H2' | 1:AA:263:A:C8 | 2.35 | 0.62 |
| 1:AA:559:A:H4' | 1:AA:560:U:H5" | 1.82 | 0.62 |
| 1:AA:590:C:H2' | 1:AA:591:U:H6 | 1.64 | 0.62 |
| 23:B1:23:LYS:HB2 | 23:B1:37:ILE:HG22 | 1.81 | 0.62 |
| 31:BA:322:A:H5' | 31:BA:340:A:C1' | 2.30 | 0.62 |
| 31:BA:1210:A:H5' | 31:BA:1212:G:H5' | 1.81 | 0.62 |
| 31:BA:2632:A:H1' | 34:BE:61:ARG:NH1 | 2.15 | 0.62 |
| 39:BN:68:GLU:HA | 39:BN:86:PRO:HB3 | 1.81 | 0.62 |
| 41:BP:140:ALA:HB1 | 25:D3:38:GLU:CG | 2.13 | 0.62 |
| 42:BQ:141:GLN:HA | 51:BZ:53:ILE:HB | 1.81 | 0.62 |
| 44:BS:74:ALA:HB1 | 44:BS:103:GLU:CB | 2.29 | 0.62 |
| 49:BX:89:ILE:HA | 49:BX:92:LEU:HD12 | 1.79 | 0.62 |
| 50:BY:16:ALA:HA | 50:BY:21:LYS:HD2 | 1.81 | 0.62 |
| 1:CA:491:G:H2' | 1:CA:492:G:C8 | 2.30 | 0.62 |
| 6:CF:18:GLN:HA | 6:CF:21:LEU:CD2 | 2.29 | 0.62 |
| 14:CN:23:ARG:NH1 | 14:CN:30:ALA:HB2 | 2.15 | 0.62 |
| 31:DA:511:U:C3' | 31:DA:512:G:H5" | 2.17 | 0.62 |
| 31:DA:1502:C:H5' | 31:DA:1503:U:OP2 | 2.00 | 0.62 |
| 31:DA:1657:C:O2' | 31:DA:1658:C:H5' | 1.99 | 0.62 |
| 33:DD:183:ARG:CG | 33:DD:183:ARG:NH1 | 2.56 | 0.62 |
| 50:DY:52:SER:C | 50:DY:54:LYS:H | 2.08 | 0.62 |
| 1:AA:81:U:H2' | 1:AA:82:U:C5 | 2.34 | 0.62 |
| 1:AA:748:C:H1' | 1:AA:749:C:OP2 | 1.99 | 0.62 |
| 1:AA:1117:G:H4' | 9:AI:104:ARG:CZ | 2.29 | 0.62 |
| 1:AA:1391:U:H2' | 1:AA:1392:G:C8 | 2.35 | 0.62 |
| 4:AD:148:VAL:HG12 | 4:AD:149:ALA:N | 2.13 | 0.62 |
| 5:AE:76:ILE:HG12 | 5:AE:77:PRO:HD2 | 1.80 | 0.62 |
| 24:B2:26:ARG:HG2 | 49:BX:5:TYR:O | 1.99 | 0.62 |
| 30:B8:29:LYS:O | 30:B8:32:LEU:N | 2.32 | 0.62 |
| 31:BA:128:C:H3' | 31:BA:128:C:C6 | 2.34 | 0.62 |
| 31:BA:271(H):G:O2' | 31:BA:271(I):G:H5" | 2.00 | 0.62 |
| 31:BA:1963:U:H4' | 31:BA:1964:G:OP1 | 1.98 | 0.62 |
| 36:BG:139:LEU:HA | 36:BG:144:ILE:HG23 | 1.81 | 0.62 |
| 37:BH:85:LYS:NZ | 37:BH:145:ALA:CA | 2.62 | 0.62 |
| 41:BP:88:LEU:C | 41:BP:90:ARG:N | 2.56 | 0.62 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 45:BT:102:ILE:HB | 45:BT:110:ILE:CD1 | 2.29 | 0.62 |
| 1:CA:543:C:C2 | 1:CA:544:G:C8 | 2.88 | 0.62 |
| 1:CA:625:G:H2' | 1:CA:626:U:H6 | 1.64 | 0.62 |
| 1:CA:667:G:H4' | 15:CO:51:HIS:CE1 | 2.35 | 0.62 |
| 1:CA:719:C:H5 | 1:CA:720:C:C4 | 2.17 | 0.62 |
| 1:CA:1238:A:H62 | 1:CA:1299:A:H62 | 1.46 | 0.62 |
| 9:CI:3:GLN:HB3 | 9:CI:20:ARG:NH1 | 2.15 | 0.62 |
| 23:D1:86:SER:N | 23:D1:87:PRO:CD | 2.62 | 0.62 |
| 28:D6:19:ARG:HG3 | 28:D6:20:ASN:H | 1.63 | 0.62 |
| 31:DA:912:C:C2 | 31:DA:913:U:C5 | 2.88 | 0.62 |
| 31:DA:966:G:C6 | 31:DA:967:C:N4 | 2.68 | 0.62 |
| 31:DA:1185:C:H5" | 31:DA:1186:G:OP1 | 2.00 | 0.62 |
| 31:DA:1434:A:C2' | 31:DA:1435:G:H5' | 2.29 | 0.62 |
| 31:DA:1481:U:H5' | 31:DA:1482:G:OP2 | 1.98 | 0.62 |
| 31:DA:1488:G:C6 | 31:DA:1489:U:N3 | 2.62 | 0.62 |
| 32:DB:87:G:H3' | 32:DB:88:C:C5' | 2.22 | 0.62 |
| 33:DD:35:LYS:CD | 33:DD:104:TYR:HD1 | 2.11 | 0.62 |
| 33:DD:158:ALA:O | 33:DD:159:ALA:HB2 | 1.99 | 0.62 |
| 36:DG:139:LEU:HA | 36:DG:144:ILE:HG23 | 1.81 | 0.62 |
| 39:DN:57:ALA:C | 39:DN:58:ASP:O | 2.37 | 0.62 |
| 41:DP:114:ILE:HG13 | 41:DP:115:LEU:N | 2.15 | 0.62 |
| 41:DP:124:LYS:HG2 | 41:DP:143:GLY:CA | 2.29 | 0.62 |
| 49:DX:54:VAL:C | 49:DX:55:ASN:HD22 | 2.07 | 0.62 |
| 50:DY:37:VAL:HG11 | 50:DY:72:VAL:HG21 | 1.82 | 0.62 |
| 1:AA:131:C:H2' | 1:AA:132:C:C6 | 2.34 | 0.62 |
| 1:AA:193:C:H2' | 1:AA:194:C:H6 | 1.63 | 0.62 |
| 1:AA:999:C:H2' | 1:AA:1000:U:C6 | 2.35 | 0.62 |
| 1:AA:1128:C:H5' | 9:AI:16:ARG:NH2 | 2.15 | 0.62 |
| 2:AB:185:ILE:HG22 | 2:AB:199:TYR:CB | 2.19 | 0.62 |
| 2:AB:218:ALA:O | 2:AB:222:ILE:HG13 | 1.99 | 0.62 |
| 6:AF:62:TRP:O | 6:AF:62:TRP:HE3 | 1.82 | 0.62 |
| 12:AL:91:LYS:O | 12:AL:91:LYS:HG3 | 2.00 | 0.62 |
| 24:B2:37:PHE:CE2 | 24:B2:40:SER:HA | 2.34 | 0.62 |
| 27:B5:2:ALA:N | 31:BA:747:U:N3 | 2.48 | 0.62 |
| 27:B5:4:HIS:CB | 27:B5:5:PRO:HD3 | 2.25 | 0.62 |
| 31:BA:620:G:H4' | 31:BA:621:A:C5' | 2.29 | 0.62 |
| 31:BA:1218:C:C2' | 31:BA:1219:G:H5' | 2.29 | 0.62 |
| 39:BN:58:ASP:OD1 | 39:BN:124:ALA:HB1 | 2.00 | 0.62 |
| 39:BN:68:GLU:HG3 | 39:BN:88:GLU:OE1 | 2.00 | 0.62 |
| 42:BQ:34:LEU:HD11 | 42:BQ:129:THR:HB | 1.80 | 0.62 |
| 44:BS:87:PHE:O | 44:BS:88:ASP:HB2 | 1.98 | 0.62 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 47:BV:61:VAL:C | 47:BV:62:LEU:HD23 | 2.24 | 0.62 |
| 48:BW:24:ILE:HD12 | 48:BW:24:ILE:O | 2.00 | 0.62 |
| 49:BX:76:ARG:O | 49:BX:77:LYS:HB2 | 2.00 | 0.62 |
| 1:CA:224:C:H2' | 1:CA:225:C:H6 | 1.61 | 0.62 |
| 1:CA:355:C:H5' | 1:CA:389:A:OP2 | 2.00 | 0.62 |
| 1:CA:1388:C:H2' | 1:CA:1389:C:H6 | 1.65 | 0.62 |
| 5:CE:101:ILE:HG12 | 5:CE:101:ILE:O | 1.98 | 0.62 |
| 22:D0:39:ARG:HH21 | 31:DA:2355:C:H1' | 1.65 | 0.62 |
| 24:D2:33:MET:HG2 | 49:DX:11:PRO:CD | 2.30 | 0.62 |
| 27:D5:57:VAL:HG23 | 27:D5:58:LEU:N | 2.15 | 0.62 |
| 30:D8:6:THR:CG2 | 30:D8:63:PRO:HD3 | 2.29 | 0.62 |
| 31:DA:314:A:C2' | 31:DA:315:G:H5' | 2.30 | 0.62 |
| 31:DA:1288:U:C2 | 31:DA:1327:C:O2 | 2.53 | 0.62 |
| 31:DA:1332:G:C8 | 31:DA:1332:G:H5' | 2.35 | 0.62 |
| 31:DA:1899:G:H22 | 31:DA:1902:C:H41 | 0.74 | 0.62 |
| 34:DE:46:ALA:HA | 34:DE:82:ARG:O | 1.99 | 0.62 |
| 34:DE:116:VAL:HG13 | 34:DE:122:PHE:CG | 2.35 | 0.62 |
| 37:DH:158:HIS:NE2 | 37:DH:170:ARG:CA | 2.59 | 0.62 |
| 38:DI:2:LYS:HB2 | 38:DI:39:ALA:CB | 2.30 | 0.62 |
| 44:DS:85:VAL:CG2 | 44:DS:106:ARG:HB2 | 2.24 | 0.62 |
| 50:DY:46:LYS:O | 50:DY:47:LYS:HB3 | 1.99 | 0.62 |
| 51:DZ:71:VAL:HG22 | 51:DZ:88:PHE:CE2 | 2.34 | 0.62 |
| 7:AG:40:ALA:O | 7:AG:44:TYR:CD1 | 2.52 | 0.62 |
| 24:B2:46:GLN:C | 24:B2:48:HIS:H | 2.07 | 0.62 |
| 26:B4:5:ILE:C | 36:BG:67:LYS:HG2 | 2.25 | 0.62 |
| 31:BA:908:C:O2' | 31:BA:909:A:H5' | 2.00 | 0.62 |
| 31:BA:2317:C:O2 | 31:BA:2318:G:O4' | 2.18 | 0.62 |
| 37:BH:40:GLU:C | 37:BH:41:MET:HE3 | 2.25 | 0.62 |
| 38:BI:2:LYS:HB2 | 38:BI:39:ALA:CB | 2.29 | 0.62 |
| 39:BN:62:VAL:HG22 | 39:BN:66:LYS:HG3 | 1.82 | 0.62 |
| 45:BT:17:THR:O | 45:BT:18:ASP:HB3 | 1.99 | 0.62 |
| 45:BT:61:PHE:CZ | 45:BT:85:LYS:HE2 | 2.35 | 0.62 |
| 50:BY:8:LYS:HZ1 | 50:BY:73:ARG:HA | 1.63 | 0.62 |
| 1:CA:946:A:H2' | 1:CA:947:G:H8 | 1.64 | 0.62 |
| 1:CA:1067:A:H1' | 1:CA:1068:G:C8 | 2.35 | 0.62 |
| 26:D4:19:GLY:C | 26:D4:21:VAL:H | 2.07 | 0.62 |
| 31:DA:154:G:H2' | 31:DA:154(A):C:O2 | 1.99 | 0.62 |
| 31:DA:531:C:H4' | 31:DA:532:A:H5" | 1.82 | 0.62 |
| 31:DA:1021:A:N6 | 31:DA:1141:U:H3 | 1.98 | 0.62 |
| 31:DA:2052:G:O4' | 34:DE:142:GLY:HA3 | 1.99 | 0.62 |
| 31:DA:2524:G:H5' | 31:DA:2524:G:H8 | 1.65 | 0.62 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 33:DD:35:LYS:CG | 33:DD:64:ILE:N | 2.62 | 0.62 |
| 37:DH:30:LYS:HZ1 | 37:DH:81:GLU:HA | 1.62 | 0.62 |
| 38:DI:81:VAL:HG13 | 38:DI:88:ILE:HG23 | 1.80 | 0.62 |
| 39:DN:129:PRO:O | 39:DN:130:HIS:HB2 | 1.99 | 0.62 |
| 40:DO:31:LYS:HB3 | 40:DO:32:TYR:CD1 | 2.34 | 0.62 |
| 42:DQ:63:LYS:HB2 | 42:DQ:63:LYS:NZ | 2.14 | 0.62 |
| 43:DR:8:ARG:HA | 43:DR:8:ARG:NE | 2.12 | 0.62 |
| 45:DT:38:ASN:C | 45:DT:38:ASN:ND2 | 2.56 | 0.62 |
| 51:DZ:150:LEU:HD13 | 51:DZ:150:LEU:H | 1.62 | 0.62 |
| 9:AI:111:ARG:O | 9:AI:113:LYS:HD2 | 1.99 | 0.61 |
| 10:AJ:32:ALA:CB | 10:AJ:76:ASN:HB3 | 2.29 | 0.61 |
| 11:AK:34:ASP:HB2 | 11:AK:35:PRO:CD | 2.29 | 0.61 |
| 12:AL:102:ARG:HH11 | 12:AL:102:ARG:HG2 | 1.64 | 0.61 |
| 31:BA:69:C:H2' | 31:BA:69:C:O2 | 1.99 | 0.61 |
| 31:BA:2712:U:O2' | 31:BA:2712(A):A:OP2 | 2.16 | 0.61 |
| 34:BE:38:THR:HG22 | 34:BE:40:GLU:N | 2.15 | 0.61 |
| 37:BH:20:ALA:HB1 | 37:BH:21:PRO:HD2 | 1.82 | 0.61 |
| 37:BH:158:HIS:CD2 | 37:BH:170:ARG:O | 2.52 | 0.61 |
| 42:BQ:141:GLN:CD | 51:BZ:72:ARG:HG2 | 2.25 | 0.61 |
| 44:BS:33:LYS:HB3 | 44:BS:34:HIS:CD2 | 2.35 | 0.61 |
| 46:BU:8:VAL:CG1 | 46:BU:12:ARG:HG3 | 2.30 | 0.61 |
| 49:BX:73:ARG:O | 49:BX:74:PRO:C | 2.43 | 0.61 |
| 1:CA:590:C:H2' | 1:CA:591:U:H6 | 1.63 | 0.61 |
| 1:CA:748:C:H1' | 1:CA:749:C:OP2 | 1.99 | 0.61 |
| 1:CA:946:A:H2' | 1:CA:947:G:C8 | 2.35 | 0.61 |
| 1:CA:952:U:H2' | 1:CA:953:G:H8 | 1.65 | 0.61 |
| 1:CA:1225:A:H2' | 1:CA:1225:A:N3 | 2.14 | 0.61 |
| 3:CC:34:LEU:O | 3:CC:38:ARG:HG2 | 2.00 | 0.61 |
| 6:CF:69:GLU:O | 6:CF:72:VAL:HG12 | 2.00 | 0.61 |
| 23:D1:85:LEU:C | 23:D1:87:PRO:CD | 2.70 | 0.61 |
| 28:D6:19:ARG:NH2 | 31:DA:2401:U:OP1 | 2.33 | 0.61 |
| 28:D6:19:ARG:O | 28:D6:20:ASN:O | 2.18 | 0.61 |
| 31:DA:83:G:H22 | 31:DA:102:G:HO2' | 1.48 | 0.61 |
| 31:DA:1515:G:H2' | 31:DA:1516:C:C6 | 2.35 | 0.61 |
| 35:DF:141:ALA:O | 35:DF:144:LYS:HB3 | 2.00 | 0.61 |
| 37:DH:158:HIS:NE2 | 37:DH:169:VAL:O | 2.31 | 0.61 |
| 38:DI:109:ILE:HG22 | 38:DI:130:TYR:OH | 2.00 | 0.61 |
| 41:DP:122:PRO:HA | 41:DP:141:ALA:O | 2.00 | 0.61 |
| 45:DT:78:LEU:O | 45:DT:78:LEU:HD23 | 1.99 | 0.61 |
| 48:DW:54:ALA:CB | 48:DW:107:LEU:HD22 | 2.30 | 0.61 |
| 1:AA:250:A:H1' | 1:AA:251:G:OP2 | 2.00 | 0.61 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:356:A:H2' | 1:AA:357:G:H8 | 1.64 | 0.61 |
| 1:AA:625:G:H2' | 1:AA:626:U:H6 | 1.65 | 0.61 |
| 1:AA:767:A:H2' | 1:AA:768:A:O4' | 2.00 | 0.61 |
| 12:AL:87:GLY:H | 12:AL:99:HIS:H | 1.48 | 0.61 |
| 20:AT:58:LYS:HE3 | 20:AT:62:LEU:HD11 | 1.82 | 0.61 |
| 25:B3:39:ASP:OD1 | 25:B3:44:ARG:HG3 | 2.01 | 0.61 |
| 27:B5:57:VAL:O | 27:B5:58:LEU:HG | 2.00 | 0.61 |
| 31:BA:1484:G:C2' | 31:BA:1485:G:O5' | 2.47 | 0.61 |
| 31:BA:2402:C:C2' | 31:BA:2403:C:H5' | 2.30 | 0.61 |
| 33:BD:71:ASP:HB3 | 33:BD:103:ARG:HH22 | 1.64 | 0.61 |
| 34:BE:27:LEU:HD22 | 45:BT:1:MET:CE | 2.30 | 0.61 |
| 34:BE:59:VAL:C | 34:BE:60:ASN:ND2 | 2.57 | 0.61 |
| 34:BE:63:LEU:O | 34:BE:64:LYS:C | 2.42 | 0.61 |
| 50:BY:79:CYS:O | 50:BY:80:GLY:C | 2.42 | 0.61 |
| 51:BZ:166:SER:HB2 | 51:BZ:168:GLU:H | 1.65 | 0.61 |
| 1:CA:509:A:H4' | 1:CA:510:A:OP1 | 2.00 | 0.61 |
| 2:CB:178:ARG:NH2 | 8:CH:68:ARG:HH22 | 1.98 | 0.61 |
| 3:CC:111:LEU:HD21 | 3:CC:145:GLY:O | 2.00 | 0.61 |
| 12:CL:87:GLY:H | 12:CL:99:HIS:H | 1.48 | 0.61 |
| 23:D1:64:ALA:HA | 23:D1:67:ILE:HG13 | 1.82 | 0.61 |
| 27:D5:48:GLU:O | 27:D5:50:GLY:N | 2.34 | 0.61 |
| 30:D8:14:VAL:CG1 | 30:D8:22:VAL:HG13 | 2.30 | 0.61 |
| 31:DA:1179:C:C3' | 31:DA:1180:C:H5" | 2.29 | 0.61 |
| 31:DA:1502:C:O2 | 31:DA:1502:C:C2' | 2.49 | 0.61 |
| 31:DA:2657:A:H2 | 31:DA:2664:G:N2 | 1.97 | 0.61 |
| 33:DD:176:ARG:HH11 | 33:DD:176:ARG:HG2 | 1.64 | 0.61 |
| 34:DE:1:MET:CB | 34:DE:84:PHE:HB2 | 2.30 | 0.61 |
| 35:DF:139:PHE:HB2 | 35:DF:166:ALA:HB1 | 1.81 | 0.61 |
| 41:DP:13:ASN:HD22 | 41:DP:13:ASN:H | 1.49 | 0.61 |
| 41:DP:101:VAL:HB | 41:DP:107:LYS:HA | 1.82 | 0.61 |
| 51:DZ:61:LEU:HB2 | 51:DZ:65:GLN:HB2 | 1.82 | 0.61 |
| 4:AD:126:ILE:HG22 | 4:AD:127:THR:H | 1.63 | 0.61 |
| 6:AF:63:TYR:N | 6:AF:63:TYR:CD2 | 2.67 | 0.61 |
| 22:B0:74:ARG:NH2 | 32:BB:13:A:OP2 | 2.34 | 0.61 |
| 31:BA:542:C:C2' | 31:BA:543:C:OP1 | 2.49 | 0.61 |
| 31:BA:807:U:H2' | 31:BA:808:G:O5' | 2.00 | 0.61 |
| 31:BA:861:A:C2 | 31:BA:917:A:C4 | 2.88 | 0.61 |
| 31:BA:2610:C:H4' | 31:BA:2611:U:OP2 | 1.99 | 0.61 |
| 31:BA:2859:G:H3' | 31:BA:2859:G:C8 | 2.35 | 0.61 |
| 39:BN:67:LEU:C | 39:BN:69:GLN:H | 2.08 | 0.61 |
| 39:BN:91:LEU:HA | 39:BN:95:PRO:HB3 | 1.81 | 0.61 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 47:BV:18:LEU:HD13 | 47:BV:98:GLU:OE1 | 2.00 | 0.61 |
| 49:BX:29:TRP:CH2 | 49:BX:76:ARG:NH1 | 2.68 | 0.61 |
| 50:BY:14:LEU:HG | 50:BY:15:VAL:N | 2.15 | 0.61 |
| 50:BY:61:ILE:HG22 | 50:BY:61:ILE:O | 2.01 | 0.61 |
| 51:BZ:67:LEU:H | 51:BZ:67:LEU:HD12 | 1.65 | 0.61 |
| 1:CA:9:G:H5' | 5:CE:122:GLU:OE2 | 1.99 | 0.61 |
| 1:CA:81:U:H2' | 1:CA:82:U:C5 | 2.35 | 0.61 |
| 1:CA:674:G:H2' | 1:CA:675:A:H8 | 1.64 | 0.61 |
| 1:CA:801:U:H2' | 1:CA:802:A:H8 | 1.65 | 0.61 |
| 1:CA:1128:C:H5' | 9:CI:16:ARG:NH2 | 2.15 | 0.61 |
| 2:CB:19:HIS:O | 2:CB:39:ILE:HG23 | 1.99 | 0.61 |
| 16:CP:9:PHE:CE2 | 16:CP:18:ARG:NE | 2.68 | 0.61 |
| 16:CP:39:TYR:HA | 16:CP:48:TRP:O | 2.00 | 0.61 |
| 28:D6:15:GLU:CD | 28:D6:41:PRO:HG3 | 2.25 | 0.61 |
| 28:D6:29:ASN:O | 28:D6:30:THR:C | 2.43 | 0.61 |
| 31:DA:117:G:H5" | 31:DA:118:A:OP2 | 2.00 | 0.61 |
| 31:DA:1478:G:C2' | 31:DA:1479:G:H5' | 2.31 | 0.61 |
| 32:DB:21:G:O2' | 32:DB:22:U:P | 2.58 | 0.61 |
| 32:DB:74:U:C3' | 32:DB:75:G:H5" | 2.31 | 0.61 |
| 37:DH:46:GLU:O | 37:DH:47:GLU:HB2 | 2.00 | 0.61 |
| 37:DH:153:LYS:HB2 | 37:DH:154:PRO:HD3 | 1.83 | 0.61 |
| 37:DH:158:HIS:HE2 | 37:DH:170:ARG:CA | 2.13 | 0.61 |
| 40:DO:77:ILE:HG13 | 45:DT:74:ARG:HG2 | 1.82 | 0.61 |
| 41:DP:16:ARG:HG3 | 41:DP:17:LYS:N | 2.13 | 0.61 |
| 41:DP:65:ARG:HB2 | 41:DP:65:ARG:NH1 | 2.14 | 0.61 |
| 49:DX:63:LYS:HZ1 | 49:DX:70:LEU:HD21 | 1.66 | 0.61 |
| 49:DX:83:VAL:O | 49:DX:84:ALA:HB3 | 2.00 | 0.61 |
| 1:AA:359:U:H2' | 1:AA:360:A:C8 | 2.36 | 0.61 |
| 1:AA:1278:U:O4 | 10:AJ:99:LYS:HE3 | 2.00 | 0.61 |
| 2:AB:21:ARG:CB | 2:AB:39:ILE:HA | 2.30 | 0.61 |
| 2:AB:178:ARG:NH2 | 8:AH:74:PRO:HG3 | 2.15 | 0.61 |
| 4:AD:133:VAL:HG13 | 4:AD:135:LEU:HD22 | 1.83 | 0.61 |
| 5:AE:96:PRO:HA | 5:AE:117:ASP:OD2 | 2.01 | 0.61 |
| 7:AG:37:ASN:HD21 | 9:AI:40:LEU:HD22 | 1.64 | 0.61 |
| 17:AQ:59:ILE:HG22 | 17:AQ:71:PHE:HD1 | 1.65 | 0.61 |
| 30:B8:32:LEU:CD2 | 30:B8:35:GLN:H | 2.12 | 0.61 |
| 31:BA:588:U:OP2 | 31:BA:588:U:C6 | 2.53 | 0.61 |
| 31:BA:1051:G:C2 | 31:BA:1052:C:N4 | 2.68 | 0.61 |
| 31:BA:1963:U:H2' | 31:BA:1963:U:O2 | 2.01 | 0.61 |
| 31:BA:2196:C:O2' | 31:BA:2197:U:H5' | 2.00 | 0.61 |
| 39:BN:131:GLN:NE2 | 39:BN:134:ARG:HA | 2.13 | 0.61 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 40:BO:31:LYS:HB3 | 40:BO:32:TYR:CD1 | 2.35 | 0.61 |
| 42:BQ:20:ALA:HB2 | 42:BQ:99:PRO:CD | 2.31 | 0.61 |
| 42:BQ:43:THR:HB | 42:BQ:45:GLN:HG2 | 1.82 | 0.61 |
| 43:BR:33:ARG:CG | 43:BR:115:GLU:HG3 | 2.27 | 0.61 |
| 45:BT:78:LEU:C | 45:BT:79:HIS:ND1 | 2.58 | 0.61 |
| 1:CA:105:G:H2' | 1:CA:106:C:C6 | 2.35 | 0.61 |
| 1:CA:709:G:H2' | 1:CA:710:G:H8 | 1.65 | 0.61 |
| 1:CA:820:U:H4' | 1:CA:821:G:OP2 | 1.99 | 0.61 |
| 10:CJ:32:ALA:CB | 10:CJ:76:ASN:HB3 | 2.30 | 0.61 |
| 12:CL:6:THR:HG23 | 12:CL:9:GLN:HE21 | 1.65 | 0.61 |
| 12:CL:27:LEU:HD11 | 12:CL:64:TYR:CE1 | 2.35 | 0.61 |
| 18:CR:50:ILE:HD12 | 18:CR:70:ILE:HD12 | 1.82 | 0.61 |
| 24:D2:57:ILE:HG13 | 24:D2:58:ALA:C | 2.25 | 0.61 |
| 31:DA:1040:C:O2' | 31:DA:1041:C:P | 2.58 | 0.61 |
| 31:DA:1568:G:P | 33:DD:63:ARG:HH22 | 2.24 | 0.61 |
| 31:DA:1786:A:C2 | 31:DA:2606:C:H1' | 2.35 | 0.61 |
| 31:DA:2849:U:O4 | 45:DT:23:ARG:NH2 | 2.32 | 0.61 |
| 37:DH:85:LYS:HD2 | 37:DH:141:VAL:CG1 | 2.29 | 0.61 |
| 50:DY:95:LYS:HE2 | 50:DY:101:LYS:HA | 1.82 | 0.61 |
| 1:AA:503:C:H2' | 1:AA:504:C:C6 | 2.35 | 0.61 |
| 1:AA:556:C:O2' | 1:AA:557:G:H5' | 2.00 | 0.61 |
| 1:AA:579:G:C4 | 1:AA:580:U:C5 | 2.88 | 0.61 |
| 6:AF:82:ARG:HH11 | 6:AF:82:ARG:HA | 1.66 | 0.61 |
| 10:AJ:63:PHE:CZ | 14:AN:45:ARG:HG3 | 2.34 | 0.61 |
| 13:AM:82:MET:HB2 | 13:AM:93:ARG:NH1 | 2.16 | 0.61 |
| 28:B6:13:CYS:O | 28:B6:21:TYR:HA | 1.99 | 0.61 |
| 31:BA:673:C:H5" | 35:BF:81:PRO:HD2 | 1.81 | 0.61 |
| 31:BA:1030:G:OP2 | 42:BQ:128:LYS:HE2 | 2.00 | 0.61 |
| 31:BA:1179:C:C3' | 31:BA:1180:C:H5" | 2.29 | 0.61 |
| 31:BA:1299:G:H5" | 31:BA:1300:U:OP1 | 2.00 | 0.61 |
| 31:BA:1778:U:H2' | 31:BA:1784:A:N6 | 2.15 | 0.61 |
| 31:BA:2016:U:H2' | 31:BA:2017:U:C6 | 2.35 | 0.61 |
| 33:BD:43:ARG:HD2 | 33:BD:44:ASN:OD1 | 2.01 | 0.61 |
| 33:BD:267:SER:C | 33:BD:269:PHE:N | 2.49 | 0.61 |
| 37:BH:85:LYS:HD2 | 37:BH:141:VAL:CG1 | 2.29 | 0.61 |
| 39:BN:39:ARG:HE | 39:BN:41:ASP:HB2 | 1.65 | 0.61 |
| 50:BY:17:SER:OG | 50:BY:18:GLY:N | 2.33 | 0.61 |
| 1:CA:1321:C:C3' | 1:CA:1322:C:H5" | 2.30 | 0.61 |
| 9:CI:114:TYR:HD1 | 10:CJ:60:ARG:HG2 | 1.64 | 0.61 |
| 23:D1:64:ALA:HA | 23:D1:67:ILE:CG1 | 2.30 | 0.61 |
| 31:DA:795:C:H2' | 31:DA:796:C:C6 | 2.35 | 0.61 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:993:G:H1' | 47:DV:91:TYR:CD1 | 2.36 | 0.61 |
| 31:DA:1019:U:O2' | 31:DA:1021:A:H2 | 1.83 | 0.61 |
| 31:DA:1434:A:H61 | 31:DA:1558:A:H62 | 1.49 | 0.61 |
| 38:DI:9:LEU:HB2 | 38:DI:12:LEU:O | 2.00 | 0.61 |
| 39:DN:13:TRP:CZ3 | 39:DN:130:HIS:CE1 | 2.88 | 0.61 |
| 41:DP:85:LEU:HD23 | 41:DP:85:LEU:H | 1.65 | 0.61 |
| 41:DP:120:ALA:CB | 41:DP:138:LEU:HB3 | 2.31 | 0.61 |
| 47:DV:36:PRO:HG2 | 47:DV:62:LEU:HD11 | 1.80 | 0.61 |
| 51:DZ:5:LEU:HG | 51:DZ:47:VAL:HG21 | 1.81 | 0.61 |
| 1:AA:322:C:H5 | 1:AA:328:C:H5 | 1.49 | 0.61 |
| 1:AA:355:C:H5' | 1:AA:389:A:OP2 | 2.01 | 0.61 |
| 1:AA:833:U:H2' | 1:AA:834:C:H6 | 1.64 | 0.61 |
| 1:AA:892:A:H2' | 1:AA:893:C:C6 | 2.35 | 0.61 |
| 2:AB:24:TRP:CG | 2:AB:25:ASN:H | 2.19 | 0.61 |
| 2:AB:158:LEU:H | 2:AB:158:LEU:CD1 | 2.08 | 0.61 |
| 5:AE:71:LEU:O | 5:AE:72:GLN:HG3 | 2.00 | 0.61 |
| 6:AF:69:GLU:CD | 6:AF:69:GLU:H | 2.08 | 0.61 |
| 9:AI:3:GLN:HB3 | 9:AI:20:ARG:NH1 | 2.16 | 0.61 |
| 13:AM:34:LEU:HD13 | 13:AM:41:PRO:HG3 | 1.82 | 0.61 |
| 28:B6:19:ARG:HG3 | 28:B6:20:ASN:H | 1.64 | 0.61 |
| 31:BA:854:G:H2' | 31:BA:855:G:H8 | 1.65 | 0.61 |
| 31:BA:1047:G:H21 | 31:BA:1111:A:H62 | 1.47 | 0.61 |
| 31:BA:2733:A:C2' | 31:BA:2734:A:H5" | 2.30 | 0.61 |
| 31:BA:2865:U:C4 | 31:BA:2866:U:C4 | 2.89 | 0.61 |
| 47:BV:25:LEU:HB2 | 47:BV:94:LEU:HD13 | 1.82 | 0.61 |
| 49:BX:24:GLY:O | 49:BX:25:LYS:O | 2.18 | 0.61 |
| 49:BX:54:VAL:C | 49:BX:55:ASN:HD22 | 2.08 | 0.61 |
| 51:BZ:39:VAL:CG2 | 51:BZ:44:PHE:HB2 | 2.30 | 0.61 |
| 1:CA:687:A:N3 | 1:CA:688:G:H1' | 2.16 | 0.61 |
| 1:CA:1278:U:O4 | 10:CJ:99:LYS:HE3 | 1.99 | 0.61 |
| 4:CD:108:LEU:HD23 | 4:CD:183:GLY:HA3 | 1.81 | 0.61 |
| 5:CE:147:ASP:HA | 5:CE:150:ARG:HH11 | 1.65 | 0.61 |
| 11:CK:31:THR:HA | 11:CK:42:TRP:HA | 1.82 | 0.61 |
| 29:D7:34:ARG:HB2 | 29:D7:42:LEU:HD22 | 1.81 | 0.61 |
| 31:DA:212:G:C2' | 31:DA:213:A:H5' | 2.30 | 0.61 |
| 31:DA:309:G:O3' | 50:DY:18:GLY:HA3 | 2.00 | 0.61 |
| 31:DA:626:U:C2 | 41:DP:105:LEU:HG | 2.36 | 0.61 |
| 31:DA:1545:A:H2' | 31:DA:1546:C:H5' | 1.83 | 0.61 |
| 31:DA:2836:U:H2' | 31:DA:2837:G:C8 | 2.36 | 0.61 |
| 47:DV:25:LEU:H | 47:DV:94:LEU:HD12 | 1.64 | 0.61 |
| 50:DY:8:LYS:HZ1 | 50:DY:73:ARG:HA | 1.64 | 0.61 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 51:DZ:143:GLY:C | 51:DZ:144:LEU:HD22 | 2.25 | 0.61 |
| 1:AA:1158:C:N3 | 1:AA:1181:G:N2 | 2.49 | 0.61 |
| 1:AA:1162:C:H2' | 1:AA:1163:C:H6 | 1.65 | 0.61 |
| 1:AA:1256:A:H61 | 1:AA:1278:U:C1' | 2.09 | 0.61 |
| 10:AJ:63:PHE:HB3 | 14:AN:57:ARG:O | 1.99 | 0.61 |
| 20:AT:97:ALA:O | 20:AT:99:LEU:HG | 2.00 | 0.61 |
| 24:B2:57:ILE:HG13 | 24:B2:58:ALA:C | 2.26 | 0.61 |
| 31:BA:795:C:H2' | 31:BA:796:C:H6 | 1.64 | 0.61 |
| 31:BA:854:G:H2' | 31:BA:855:G:C8 | 2.36 | 0.61 |
| 31:BA:1265:A:OP1 | 31:BA:1265:A:H8 | 1.84 | 0.61 |
| 31:BA:1592:C:O2' | 31:BA:1593:G:H5' | 2.00 | 0.61 |
| 31:BA:1902:C:H1' | 33:BD:244:ARG:HD3 | 1.82 | 0.61 |
| 31:BA:2830:G:H8 | 31:BA:2830:G:C5' | 2.13 | 0.61 |
| 32:BB:71:C:H2' | 32:BB:71:C:O2 | 2.00 | 0.61 |
| 37:BH:138:LYS:O | 37:BH:140:LYS:N | 2.34 | 0.61 |
| 46:BU:14:HIS:HD2 | 46:BU:32:PHE:CB | 2.13 | 0.61 |
| 48:BW:54:ALA:HB1 | 48:BW:107:LEU:HD22 | 1.82 | 0.61 |
| 49:BX:36:LYS:O | 49:BX:38:GLU:N | 2.32 | 0.61 |
| 50:BY:8:LYS:HD3 | 50:BY:28:LYS:HZ2 | 1.66 | 0.61 |
| 1:CA:623:C:C4 | 1:CA:624:C:C5 | 2.89 | 0.61 |
| 1:CA:1133:G:N3 | 1:CA:1142:G:N2 | 2.48 | 0.61 |
| 11:CK:34:ASP:HB2 | 11:CK:35:PRO:CD | 2.31 | 0.61 |
| 23:D1:37:ILE:HD12 | 31:DA:2079:U:O2' | 2.00 | 0.61 |
| 30:D8:29:LYS:O | 30:D8:32:LEU:N | 2.33 | 0.61 |
| 31:DA:547:A:O2' | 31:DA:548:A:OP2 | 2.18 | 0.61 |
| 31:DA:1348:G:H2' | 31:DA:1349:A:H5' | 1.82 | 0.61 |
| 35:DF:205:ARG:O | 35:DF:206:ILE:HG13 | 2.01 | 0.61 |
| 41:DP:79:ARG:HH21 | 41:DP:109:GLY:HA2 | 1.66 | 0.61 |
| 41:DP:113:LYS:HE3 | 41:DP:130:PHE:O | 2.01 | 0.61 |
| 50:DY:27:VAL:HB | 50:DY:29:GLU:OE1 | 2.00 | 0.61 |
| 51:DZ:64:GLY:O | 51:DZ:65:GLN:O | 2.19 | 0.61 |
| 1:AA:22:G:H2' | 1:AA:23:C:C6 | 2.35 | 0.61 |
| 1:AA:386:C:O2' | 1:AA:387:U:H5' | 2.00 | 0.61 |
| 1:AA:562:C:H4' | 1:AA:563:A:O5' | 2.01 | 0.61 |
| 1:AA:1348:U:H4' | 9:AI:120:ARG:HD2 | 1.82 | 0.61 |
| 1:AA:1521:G:H2' | 1:AA:1522:U:C6 | 2.36 | 0.61 |
| 5:AE:43:LEU:HD21 | 5:AE:132:ALA:HB1 | 1.83 | 0.61 |
| 24:B2:23:LYS:HA | 49:BX:5:TYR:CD1 | 2.35 | 0.61 |
| 31:BA:511:U:H5" | 31:BA:512:G:OP2 | 2.00 | 0.61 |
| 31:BA:1162:G:H1' | 47:BV:91:TYR:OH | 2.01 | 0.61 |
| 31:BA:1786:A:C2 | 31:BA:2606:C:H1' | 2.36 | 0.61 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2870:C:H2' | 31:BA:2871:C:H5' | 1.83 | 0.61 |
| 34:BE:117:MET:SD | 34:BE:136:ARG:HB3 | 2.40 | 0.61 |
| 35:BF:124:LEU:HD12 | 35:BF:124:LEU:C | 2.24 | 0.61 |
| 41:BP:13:ASN:HD22 | 41:BP:13:ASN:H | 1.48 | 0.61 |
| 41:BP:16:ARG:CZ | 41:BP:16:ARG:HB2 | 2.31 | 0.61 |
| 44:BS:16:ASN:ND2 | 44:BS:92:TYR:CZ | 2.68 | 0.61 |
| 46:BU:92:ARG:HD2 | 47:BV:11:GLN:HG2 | 1.81 | 0.61 |
| 47:BV:82:ARG:HG3 | 47:BV:82:ARG:NH1 | 2.11 | 0.61 |
| 14:CN:40:CYS:SG | 14:CN:42:ILE:HG12 | 2.41 | 0.61 |
| 30:D8:35:GLN:CB | 31:DA:2420:C:OP1 | 2.49 | 0.61 |
| 31:DA:795:C:H2' | 31:DA:796:C:H6 | 1.65 | 0.61 |
| 31:DA:807:U:H2' | 31:DA:808:G:O5' | 2.00 | 0.61 |
| 31:DA:2314:C:O2' | 31:DA:2315:G:H5' | 2.00 | 0.61 |
| 31:DA:2523:G:C2' | 31:DA:2524:G:C5' | 2.78 | 0.61 |
| 37:DH:44:VAL:HG12 | 37:DH:45:VAL:N | 2.12 | 0.61 |
| 39:DN:32:THR:O | 39:DN:35:ARG:O | 2.19 | 0.61 |
| 50:DY:37:VAL:HG22 | 50:DY:67:LEU:O | 2.01 | 0.61 |
| 1:AA:149:A:HO2' | 1:AA:150:C:H6 | 1.44 | 0.61 |
| 1:AA:344:A:O2' | 1:AA:346:G:N7 | 2.24 | 0.61 |
| 1:AA:538:G:OP2 | 12:AL:115:LYS:HG3 | 2.00 | 0.61 |
| 1:AA:719:C:H5 | 1:AA:720:C:C4 | 2.19 | 0.61 |
| 1:AA:775:G:H2' | 1:AA:776:G:H5' | 1.82 | 0.61 |
| 1:AA:1321:C:C3' | 1:AA:1322:C:H5" | 2.31 | 0.61 |
| 1:AA:1388:C:H2' | 1:AA:1389:C:H6 | 1.63 | 0.61 |
| 2:AB:24:TRP:CG | 2:AB:25:ASN:N | 2.68 | 0.61 |
| 2:AB:71:VAL:HG13 | 2:AB:93:VAL:HB | 1.83 | 0.61 |
| 5:AE:145:LYS:O | 5:AE:149:GLU:HG2 | 2.01 | 0.61 |
| 10:AJ:6:ILE:HG22 | 10:AJ:98:ILE:HG13 | 1.82 | 0.61 |
| 12:AL:90:VAL:O | 12:AL:92:ASP:N | 2.27 | 0.61 |
| 23:B1:34:THR:CG2 | 31:BA:388:G:P | 2.86 | 0.61 |
| 25:B3:19:GLN:NE2 | 25:B3:52:HIS:HE1 | 1.99 | 0.61 |
| 27:B5:52:TYR:CG | 27:B5:52:TYR:O | 2.53 | 0.61 |
| 31:BA:649:G:H2' | 31:BA:650:C:C6 | 2.35 | 0.61 |
| 31:BA:1636:C:H2' | 31:BA:1637:A:C8 | 2.35 | 0.61 |
| 33:BD:94:LEU:C | 33:BD:94:LEU:HD22 | 2.26 | 0.61 |
| 34:BE:182:LEU:HD12 | 34:BE:183:LEU:N | 2.16 | 0.61 |
| 42:BQ:9:TYR:HD2 | 42:BQ:9:TYR:C | 2.08 | 0.61 |
| 1:CA:336:C:O2' | 1:CA:337:C:H5' | 2.01 | 0.61 |
| 1:CA:827:U:H5" | 1:CA:828:A:OP2 | 2.01 | 0.61 |
| 1:CA:1097:C:H1' | 1:CA:1170:A:H1' | 1.81 | 0.61 |
| 2:CB:24:TRP:CG | 2:CB:25:ASN:H | 2.18 | 0.61 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 5:CE:43:LEU:HD21 | 5:CE:132:ALA:HB1 | 1.83 | 0.61 |
| 9:CI:48:GLU:N | 9:CI:49:PRO:HD2 | 2.16 | 0.61 |
| 31:DA:753:C:O5' | 31:DA:753:C:H6 | 1.84 | 0.61 |
| 31:DA:1803:A:O2' | 33:DD:259:THR:HG21 | 2.01 | 0.61 |
| 33:DD:76:PRO:HG2 | 33:DD:98:VAL:CG2 | 2.29 | 0.61 |
| 33:DD:80:ALA:HB3 | 33:DD:94:LEU:HD13 | 1.82 | 0.61 |
| 33:DD:218:ARG:HB3 | 33:DD:219:PRO:HD2 | 1.83 | 0.61 |
| 39:DN:67:LEU:C | 39:DN:69:GLN:H | 2.08 | 0.61 |
| 40:DO:112:MET:HE2 | 40:DO:112:MET:HA | 1.83 | 0.61 |
| 41:DP:62:LEU:H | 41:DP:62:LEU:HD22 | 1.65 | 0.61 |
| 46:DU:76:TYR:CZ | 46:DU:80:ILE:HG13 | 2.36 | 0.61 |
| 48:DW:95:ILE:O | 48:DW:95:ILE:HG13 | 2.00 | 0.61 |
| 49:DX:60:ARG:HH21 | 49:DX:74:PRO:HG2 | 1.65 | 0.61 |
| 1:AA:1106:G:H5" | 3:AC:172:ARG:HG2 | 1.81 | 0.61 |
| 1:AA:1238:A:H62 | 1:AA:1299:A:H62 | 1.48 | 0.61 |
| 8:AH:53:VAL:O | 8:AH:54:ASP:HB2 | 2.01 | 0.61 |
| 9:AI:114:TYR:HD2 | 9:AI:114:TYR:N | 1.96 | 0.61 |
| 23:B1:65:SER:N | 23:B1:67:ILE:CD1 | 2.63 | 0.61 |
| 31:BA:494:G:H5" | 31:BA:494:G:H8 | 1.66 | 0.61 |
| 31:BA:1484:G:N2 | 31:BA:1505:C:N4 | 2.37 | 0.61 |
| 31:BA:2876:G:H4' | 45:BT:3:ARG:HD3 | 1.83 | 0.61 |
| 33:BD:8:PRO:HB3 | 33:BD:14:ARG:CB | 2.31 | 0.61 |
| 34:BE:95:ILE:HD12 | 34:BE:95:ILE:N | 2.14 | 0.61 |
| 47:BV:62:LEU:HB3 | 47:BV:98:GLU:CB | 2.30 | 0.61 |
| 47:BV:83:ARG:CG | 47:BV:83:ARG:NH1 | 2.58 | 0.61 |
| 1:CA:253:U:H2' | 1:CA:254:G:H8 | 1.66 | 0.61 |
| 1:CA:674:G:H21 | 11:CK:116:HIS:HB2 | 1.64 | 0.61 |
| 10:CJ:6:ILE:HD11 | 10:CJ:72:VAL:HB | 1.83 | 0.61 |
| 12:CL:21:LYS:N | 12:CL:21:LYS:HD2 | 2.16 | 0.61 |
| 31:DA:1763:G:H4' | 31:DA:1763:G:OP1 | 2.01 | 0.61 |
| 31:DA:1956:U:C2' | 31:DA:1957:C:H5' | 2.30 | 0.61 |
| 37:DH:41:MET:O | 37:DH:43:VAL:HG13 | 2.00 | 0.61 |
| 39:DN:15:LEU:HD13 | 39:DN:16:ILE:N | 2.15 | 0.61 |
| 49:DX:29:TRP:CH2 | 49:DX:76:ARG:NH1 | 2.69 | 0.61 |
| 1:AA:1372:U:OP1 | 9:AI:72:GLY:N | 2.34 | 0.60 |
| 31:BA:107:C:H2' | 31:BA:108:U:H6 | 1.66 | 0.60 |
| 31:BA:1019:U:O2' | 31:BA:1021:A:H2 | 1.83 | 0.60 |
| 31:BA:1191:G:OP1 | 41:BP:35:HIS:ND1 | 2.34 | 0.60 |
| 32:BB:81:G:H5" | 32:BB:82:G:OP2 | 2.00 | 0.60 |
| 35:BF:80:ALA:O | 35:BF:83:PHE:HB2 | 2.01 | 0.60 |
| 36:BG:133:LEU:HD12 | 36:BG:133:LEU:C | 2.26 | 0.60 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 37:BH:156:ALA:O | 37:BH:157:TYR:C | 2.44 | 0.60 |
| 42:BQ:20:ALA:O | 42:BQ:22:LYS:N | 2.34 | 0.60 |
| 50:BY:44:ILE:HD12 | 50:BY:44:ILE:N | 2.15 | 0.60 |
| 51:BZ:74:VAL:HG22 | 51:BZ:86:VAL:HG12 | 1.83 | 0.60 |
| 1:CA:52:G:O2' | 1:CA:53:A:H5' | 2.00 | 0.60 |
| 1:CA:192:U:O4' | 20:CT:103:GLY:HA2 | 2.01 | 0.60 |
| 2:CB:71:VAL:HG13 | 2:CB:93:VAL:HB | 1.82 | 0.60 |
| 6:CF:19:LEU:HD11 | 6:CF:59:TYR:CZ | 2.35 | 0.60 |
| 31:DA:620:G:H4' | 31:DA:621:A:C5' | 2.30 | 0.60 |
| 31:DA:2476:A:N3 | 31:DA:2477:C:H5" | 2.16 | 0.60 |
| 31:DA:2762:G:H5" | 31:DA:2762:G:H8 | 1.66 | 0.60 |
| 33:DD:43:ARG:HD2 | 33:DD:44:ASN:OD1 | 2.01 | 0.60 |
| 36:DG:7:LEU:CD2 | 36:DG:176:LEU:HD22 | 2.31 | 0.60 |
| 36:DG:76:SER:CB | 36:DG:83:ARG:HB3 | 2.31 | 0.60 |
| 38:DI:50:ARG:C | 38:DI:52:ARG:H | 2.09 | 0.60 |
| 38:DI:79:ILE:HG22 | 38:DI:81:VAL:HG23 | 1.81 | 0.60 |
| 45:DT:129:ARG:NH1 | 45:DT:131:ALA:HB3 | 2.16 | 0.60 |
| 47:DV:79:VAL:O | 47:DV:80:GLN:CB | 2.47 | 0.60 |
| 1:AA:922:G:H4' | 5:AE:20:GLN:HA | 1.83 | 0.60 |
| 1:AA:1067:A:H1' | 1:AA:1068:G:C8 | 2.35 | 0.60 |
| 1:AA:1160:G:N3 | 1:AA:1160:G:H2' | 2.14 | 0.60 |
| 1:AA:1372:U:H5" | 9:AI:71:SER:HB3 | 1.84 | 0.60 |
| 1:AA:1379:G:C6 | 1:AA:1380:U:O4 | 2.54 | 0.60 |
| 9:AI:26:VAL:HG22 | 9:AI:61:ALA:HB3 | 1.81 | 0.60 |
| 12:AL:18:VAL:HG23 | 12:AL:19:ARG:H | 1.65 | 0.60 |
| 27:B5:27:PRO:HB3 | 48:BW:23:LEU:HD11 | 1.83 | 0.60 |
| 27:B5:50:GLY:HA3 | 27:B5:56:LYS:CG | 2.30 | 0.60 |
| 30:B8:25:MET:HG3 | 41:BP:64:LYS:HB3 | 1.83 | 0.60 |
| 30:B8:32:LEU:HB3 | 30:B8:34:TRP:CA | 2.30 | 0.60 |
| 31:BA:329:G:H1 | 50:BY:19:LYS:HD2 | 1.65 | 0.60 |
| 31:BA:1163:G:H5' | 47:BV:92:THR:HG21 | 1.84 | 0.60 |
| 45:BT:29:ARG:HG3 | 45:BT:84:GLN:HG3 | 1.82 | 0.60 |
| 2:CB:187:LEU:HD23 | 2:CB:201:ILE:HG22 | 1.83 | 0.60 |
| 2:CB:228:GLY:O | 2:CB:230:VAL:HG13 | 2.00 | 0.60 |
| 13:CM:82:MET:HB2 | 13:CM:93:ARG:NH1 | 2.16 | 0.60 |
| 31:DA:1549:C:O2' | 31:DA:1550:C:H5' | 2.01 | 0.60 |
| 37:DH:20:ALA:HB1 | 37:DH:21:PRO:HD2 | 1.83 | 0.60 |
| 37:DH:158:HIS:CD2 | 37:DH:170:ARG:O | 2.54 | 0.60 |
| 47:DV:73:SER:OG | 47:DV:74:LYS:N | 2.34 | 0.60 |
| 1:AA:1285:A:H4' | 1:AA:1286:A:O5' | 2.02 | 0.60 |
| 1:AA:1311:G:N2 | 1:AA:1327:C:C2 | 2.69 | 0.60 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 3:AC:29:TYR:O | 3:AC:33:LEU:HB2 | 2.01 | 0.60 |
| 6:AF:19:LEU:HD11 | 6:AF:59:TYR:CZ | 2.36 | 0.60 |
| 21:AU:9:ARG:O | 21:AU:13:ILE:HG13 | 2.01 | 0.60 |
| 25:B3:6:VAL:HG13 | 25:B3:54:VAL:CG1 | 2.31 | 0.60 |
| 31:BA:534:U:O2' | 46:BU:49:HIS:CD2 | 2.55 | 0.60 |
| 31:BA:2524:G:H5' | 31:BA:2524:G:H8 | 1.66 | 0.60 |
| 31:BA:2712:U:O2' | 31:BA:2712(A):A:P | 2.59 | 0.60 |
| 32:BB:82:G:H2' | 32:BB:83:G:H5' | 1.82 | 0.60 |
| 33:BD:142:VAL:HG23 | 33:BD:192:THR:C | 2.26 | 0.60 |
| 35:BF:9:ILE:HG23 | 35:BF:13:SER:O | 2.01 | 0.60 |
| 37:BH:41:MET:O | 37:BH:43:VAL:HG13 | 1.99 | 0.60 |
| 38:BI:53:ALA:HB2 | 38:BI:56:LYS:CG | 2.31 | 0.60 |
| 41:BP:122:PRO:HA | 41:BP:141:ALA:O | 2.01 | 0.60 |
| 47:BV:90:PRO:HG2 | 47:BV:91:TYR:N | 2.17 | 0.60 |
| 50:BY:46:LYS:O | 50:BY:47:LYS:HB3 | 2.01 | 0.60 |
| 1:CA:189(K):U:H2' | 1:CA:189(L):G:C8 | 2.37 | 0.60 |
| 1:CA:509:A:O2' | 1:CA:510:A:P | 2.60 | 0.60 |
| 4:CD:22:LYS:O | 4:CD:113:SER:HB3 | 2.01 | 0.60 |
| 20:CT:73:HIS:O | 20:CT:76:ALA:HB3 | 2.01 | 0.60 |
| 23:D1:63:ALA:O | 23:D1:64:ALA:C | 2.45 | 0.60 |
| 25:D3:19:GLN:HE22 | 25:D3:52:HIS:CE1 | 2.19 | 0.60 |
| 27:D5:27:PRO:HB3 | 48:DW:23:LEU:HD11 | 1.82 | 0.60 |
| 31:DA:588:U:H2' | 31:DA:589:C:H6 | 1.66 | 0.60 |
| 31:DA:1592:C:O2' | 31:DA:1593:G:H5' | 2.01 | 0.60 |
| 31:DA:2657:A:C2 | 31:DA:2664:G:N2 | 2.68 | 0.60 |
| 33:DD:35:LYS:NZ | 33:DD:104:TYR:CB | 2.50 | 0.60 |
| 34:DE:147:PRO:HB2 | 34:DE:149:ARG:HG2 | 1.83 | 0.60 |
| 35:DF:9:ILE:HG23 | 35:DF:13:SER:O | 2.01 | 0.60 |
| 35:DF:194:MET:HE3 | 35:DF:199:TRP:CD1 | 2.34 | 0.60 |
| 36:DG:16:ARG:O | 36:DG:20:ILE:HG13 | 2.01 | 0.60 |
| 37:DH:92:ILE:O | 37:DH:94:TYR:N | 2.33 | 0.60 |
| 42:DQ:20:ALA:O | 42:DQ:22:LYS:N | 2.34 | 0.60 |
| 49:DX:24:GLY:O | 49:DX:25:LYS:O | 2.18 | 0.60 |
| 49:DX:35:THR:O | 49:DX:36:LYS:O | 2.19 | 0.60 |
| 1:AA:41:G:H2' | 1:AA:42:G:C8 | 2.36 | 0.60 |
| 1:AA:299:G:C6 | 1:AA:300:A:C6 | 2.88 | 0.60 |
| 1:AA:1279:A:H2 | 10:AJ:43:ARG:HH12 | 1.47 | 0.60 |
| 20:AT:56:MET:HG2 | 20:AT:84:LEU:CD1 | 2.32 | 0.60 |
| 23:B1:19:GLN:HE21 | 31:BA:379:G:N2 | 1.89 | 0.60 |
| 23:B1:65:SER:N | 23:B1:67:ILE:HD11 | 2.15 | 0.60 |
| 26:B4:25:TYR:C | 26:B4:27:THR:H | 2.08 | 0.60 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 27:B5:51:TYR:CD2 | 27:B5:52:TYR:CE2 | 2.89 | 0.60 |
| 31:BA:128:C:H6 | 31:BA:128:C:H5" | 1.66 | 0.60 |
| 31:BA:2753:A:O2' | 31:BA:2754:U:H5' | 2.02 | 0.60 |
| 35:BF:175:THR:O | 35:BF:176:LEU:HB2 | 2.01 | 0.60 |
| 37:BH:92:ILE:O | 37:BH:94:TYR:N | 2.35 | 0.60 |
| 45:BT:38:ASN:ND2 | 45:BT:40:THR:H | 1.98 | 0.60 |
| 49:BX:38:GLU:N | 49:BX:38:GLU:OE1 | 2.34 | 0.60 |
| 1:CA:1064:G:H1' | 1:CA:1065:U:OP2 | 2.01 | 0.60 |
| 1:CA:1392:G:N2 | 1:CA:1502:A:C8 | 2.68 | 0.60 |
| 2:CB:14:GLY:O | 2:CB:15:VAL:HG13 | 2.01 | 0.60 |
| 24:D2:41:ILE:O | 24:D2:42:GLY:C | 2.44 | 0.60 |
| 31:DA:176:G:C2' | 31:DA:177:G:H5' | 2.31 | 0.60 |
| 31:DA:760:G:H2' | 31:DA:761:A:O4' | 2.02 | 0.60 |
| 31:DA:1991:U:H2' | 31:DA:1992:G:H5" | 1.83 | 0.60 |
| 31:DA:2476:A:C5 | 31:DA:2477:C:C5 | 2.89 | 0.60 |
| 34:DE:60:ASN:N | 34:DE:60:ASN:ND2 | 2.49 | 0.60 |
| 37:DH:85:LYS:HE2 | 37:DH:145:ALA:HB2 | 1.84 | 0.60 |
| 39:DN:1:MET:HB3 | 47:DV:20:LEU:CD2 | 2.25 | 0.60 |
| 42:DQ:9:TYR:HD2 | 42:DQ:9:TYR:C | 2.07 | 0.60 |
| 1:AA:22:G:H2' | 1:AA:23:C:H6 | 1.67 | 0.60 |
| 1:AA:189(K):U:H2' | 1:AA:189(L):G:C8 | 2.37 | 0.60 |
| 2:AB:116:GLU:HA | 2:AB:119:GLU:HB2 | 1.82 | 0.60 |
| 3:AC:155:GLY:O | 3:AC:156:ARG:HB2 | 2.01 | 0.60 |
| 22:B0:77:ARG:NH2 | 31:BA:857:C:H5' | 2.16 | 0.60 |
| 25:B3:11:SER:OG | 25:B3:13:ILE:HG12 | 2.01 | 0.60 |
| 28:B6:10:LEU:N | 28:B6:10:LEU:HD22 | 2.15 | 0.60 |
| 30:B8:39:LYS:O | 30:B8:39:LYS:HE2 | 2.01 | 0.60 |
| 31:BA:17:G:H4' | 46:BU:25:TRP:CH2 | 2.35 | 0.60 |
| 31:BA:308:G:O2' | 50:BY:19:LYS:HE3 | 2.00 | 0.60 |
| 31:BA:1528(A):A:C5 | 31:BA:1529:G:C8 | 2.90 | 0.60 |
| 31:BA:2476:A:C5 | 31:BA:2477:C:C5 | 2.89 | 0.60 |
| 31:BA:2712:U:O2 | 31:BA:2712:U:C5' | 2.49 | 0.60 |
| 34:BE:70:ALA:O | 34:BE:71:GLY:C | 2.45 | 0.60 |
| 35:BF:194:MET:HE3 | 35:BF:199:TRP:CD1 | 2.34 | 0.60 |
| 40:BO:60:ALA:HB2 | 40:BO:86:ILE:HA | 1.82 | 0.60 |
| 45:BT:91:ARG:HA | 45:BT:117:ASP:H | 1.67 | 0.60 |
| 1:CA:1216:G:H5" | 14:CN:5:ALA:CB | 2.32 | 0.60 |
| 5:CE:51:VAL:HB | 5:CE:52:PRO:CD | 2.29 | 0.60 |
| 8:CH:9:MET:HG2 | 8:CH:10:LEU:HD23 | 1.83 | 0.60 |
| 31:DA:823:G:O2' | 31:DA:824:A:H5' | 2.02 | 0.60 |
| 31:DA:2577:A:H5" | 31:DA:2578:G:C5' | 2.31 | 0.60 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 33:DD:267:SER:C | 33:DD:269:PHE:N | 2.55 | 0.60 |
| 39:DN:3:THR:HA | 39:DN:4:TYR:CD1 | 2.36 | 0.60 |
| 44:DS:56:LEU:O | 44:DS:57:LYS:HB2 | 2.01 | 0.60 |
| 46:DU:27:LEU:N | 46:DU:27:LEU:HD23 | 2.17 | 0.60 |
| 47:DV:23:GLU:O | 47:DV:24:LYS:C | 2.45 | 0.60 |
| 47:DV:51:VAL:CG1 | 47:DV:52:VAL:H | 2.13 | 0.60 |
| 2:AB:22:LYS:NZ | 2:AB:22:LYS:HA | 2.17 | 0.60 |
| 2:AB:178:ARG:NH2 | 8:AH:68:ARG:HH22 | 2.00 | 0.60 |
| 2:AB:219:VAL:HA | 2:AB:222:ILE:HD12 | 1.83 | 0.60 |
| 16:AP:39:TYR:HA | 16:AP:48:TRP:O | 2.02 | 0.60 |
| 18:AR:50:ILE:CD1 | 18:AR:70:ILE:HG21 | 2.28 | 0.60 |
| 25:B3:43:ILE:O | 25:B3:47:VAL:HG23 | 2.02 | 0.60 |
| 27:B5:4:HIS:HB3 | 27:B5:5:PRO:CD | 2.16 | 0.60 |
| 29:B7:5:TRP:CD1 | 29:B7:7:PRO:HG3 | 2.37 | 0.60 |
| 31:BA:672:C:H2' | 31:BA:673:C:H6 | 1.66 | 0.60 |
| 31:BA:1434:A:H61 | 31:BA:1558:A:N6 | 1.98 | 0.60 |
| 31:BA:1528:A:H2' | 31:BA:1528:A:N3 | 2.17 | 0.60 |
| 31:BA:2580:U:C5' | 34:BE:131:ALA:H | 2.15 | 0.60 |
| 38:BI:79:ILE:HG22 | 38:BI:81:VAL:HG23 | 1.83 | 0.60 |
| 39:BN:27:ALA:HB3 | 39:BN:106:MET:HE2 | 1.84 | 0.60 |
| 40:BO:14:THR:HG21 | 40:BO:86:ILE:HD13 | 1.83 | 0.60 |
| 41:BP:101:VAL:HB | 41:BP:107:LYS:HA | 1.83 | 0.60 |
| 43:BR:71:GLN:CA | 43:BR:71:GLN:HE21 | 2.14 | 0.60 |
| 44:BS:34:HIS:CD2 | 44:BS:34:HIS:N | 2.67 | 0.60 |
| 47:BV:51:VAL:HG12 | 47:BV:52:VAL:N | 2.15 | 0.60 |
| 1:CA:505:G:C6 | 1:CA:535:A:C2 | 2.89 | 0.60 |
| 1:CA:767:A:H2' | 1:CA:768:A:O4' | 2.01 | 0.60 |
| 1:CA:1279:A:H2 | 10:CJ:43:ARG:HH12 | 1.49 | 0.60 |
| 1:CA:1399:C:C2 | 1:CA:1502:A:N6 | 2.69 | 0.60 |
| 5:CE:101:ILE:CD1 | 5:CE:119:LEU:HA | 2.32 | 0.60 |
| 27:D5:40:LYS:HZ3 | 27:D5:46:CYS:HB3 | 1.66 | 0.60 |
| 27:D5:52:TYR:O | 27:D5:52:TYR:CG | 2.53 | 0.60 |
| 28:D6:36:LEU:HD13 | 28:D6:50:ARG:CZ | 2.31 | 0.60 |
| 31:DA:267:C:H2' | 31:DA:268:C:C6 | 2.37 | 0.60 |
| 31:DA:518:G:H2' | 31:DA:519:U:H6 | 1.66 | 0.60 |
| 31:DA:901:A:H5' | 31:DA:902:C:OP2 | 2.00 | 0.60 |
| 31:DA:1464:C:O2' | 31:DA:1528:A:C8 | 2.55 | 0.60 |
| 31:DA:2317:C:H2' | 31:DA:2318:G:C5' | 2.22 | 0.60 |
| 31:DA:2830:G:C5' | 31:DA:2830:G:H8 | 2.14 | 0.60 |
| 39:DN:46:VAL:HG13 | 39:DN:48:MET:HG3 | 1.84 | 0.60 |
| 40:DO:20:MET:HE3 | 40:DO:44:LYS:HE3 | 1.82 | 0.60 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 41:DP:17:LYS:O | 41:DP:17:LYS:CG | 2.50 | 0.60 |
| 42:DQ:141:GLN:HA | 51:DZ:53:ILE:HB | 1.83 | 0.60 |
| 43:DR:56:LYS:HE2 | 43:DR:94:TYR:CE2 | 2.36 | 0.60 |
| 47:DV:90:PRO:CG | 47:DV:91:TYR:H | 2.13 | 0.60 |
| 50:DY:61:ILE:HG22 | 50:DY:61:ILE:O | 2.02 | 0.60 |
| 1:AA:1090:U:C2 | 1:AA:1091:U:C5 | 2.90 | 0.60 |
| 4:AD:108:LEU:HD23 | 4:AD:183:GLY:HA3 | 1.82 | 0.60 |
| 10:AJ:51:ARG:HE | 10:AJ:61:GLU:HB2 | 1.67 | 0.60 |
| 16:AP:6:LEU:HD23 | 16:AP:17:TYR:CG | 2.36 | 0.60 |
| 19:AS:24:ALA:O | 19:AS:25:LYS:HB2 | 2.01 | 0.60 |
| 26:B4:19:GLY:C | 26:B4:21:VAL:H | 2.08 | 0.60 |
| 31:BA:543:C:H42 | 31:BA:551:G:H1 | 1.50 | 0.60 |
| 31:BA:1024:G:C3' | 31:BA:1025:G:H5" | 2.30 | 0.60 |
| 31:BA:2055:C:H5' | 31:BA:2056:G:O5' | 2.01 | 0.60 |
| 31:BA:2476:A:N3 | 31:BA:2477:C:H5" | 2.17 | 0.60 |
| 33:BD:166:GLN:HA | 33:BD:166:GLN:NE2 | 2.13 | 0.60 |
| 34:BE:93:VAL:N | 34:BE:95:ILE:HD13 | 2.13 | 0.60 |
| 34:BE:95:ILE:CD1 | 34:BE:95:ILE:H | 2.15 | 0.60 |
| 39:BN:47:ALA:CB | 39:BN:112:LEU:HD11 | 2.19 | 0.60 |
| 39:BN:90:MET:O | 39:BN:93:THR:O | 2.20 | 0.60 |
| 41:BP:50:ARG:HG2 | 41:BP:50:ARG:NH2 | 2.15 | 0.60 |
| 44:BS:74:ALA:O | 44:BS:77:ALA:HB3 | 2.01 | 0.60 |
| 1:CA:1285:A:H4' | 1:CA:1286:A:O5' | 2.02 | 0.60 |
| 1:CA:1416:G:H2' | 1:CA:1417:G:O4' | 2.01 | 0.60 |
| 2:CB:17:PHE:CD1 | 2:CB:44:LEU:HD11 | 2.37 | 0.60 |
| 2:CB:116:GLU:HA | 2:CB:119:GLU:CB | 2.30 | 0.60 |
| 5:CE:145:LYS:O | 5:CE:149:GLU:HG2 | 2.02 | 0.60 |
| 22:D0:74:ARG:HG2 | 32:DB:12:C:O2' | 2.02 | 0.60 |
| 25:D3:6:VAL:HG22 | 25:D3:56:VAL:HG22 | 1.83 | 0.60 |
| 27:D5:16:ARG:NH2 | 31:DA:517:C:OP1 | 2.34 | 0.60 |
| 31:DA:271(H):G:O2' | 31:DA:271(I):G:H5" | 2.01 | 0.60 |
| 31:DA:632:A:O2' | 31:DA:633:A:H5' | 2.01 | 0.60 |
| 31:DA:1025:G:C4 | 31:DA:1135:C:H1' | 2.37 | 0.60 |
| 31:DA:1047:G:H21 | 31:DA:1111:A:H62 | 1.49 | 0.60 |
| 31:DA:1359:A:C8 | 31:DA:1372:U:O4 | 2.54 | 0.60 |
| 31:DA:1478:G:HO2' | 31:DA:1558:A:H2 | 1.49 | 0.60 |
| 31:DA:1832:C:N4 | 31:DA:1833:U:C4 | 2.70 | 0.60 |
| 31:DA:2394:C:OP1 | 41:DP:63:PRO:CD | 2.35 | 0.60 |
| 31:DA:2722:G:O2' | 43:DR:5:LYS:HB2 | 2.02 | 0.60 |
| 33:DD:223:GLY:HA3 | 33:DD:231:HIS:ND1 | 2.17 | 0.60 |
| 37:DH:156:ALA:C | 37:DH:158:HIS:H | 2.09 | 0.60 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 38:DI:81:VAL:HG11 | 38:DI:123:LEU:HD21 | 1.83 | 0.60 |
| 42:DQ:29:PHE:O | 42:DQ:30:GLY:O | 2.19 | 0.60 |
| 43:DR:53:HIS:HD2 | 43:DR:94:TYR:OH | 1.84 | 0.60 |
| 49:DX:36:LYS:O | 49:DX:38:GLU:N | 2.35 | 0.60 |
| 1:AA:783:C:O2' | 1:AA:784:C:H5' | 2.01 | 0.60 |
| 1:AA:1325:C:H4' | 21:AU:17:THR:HG21 | 1.82 | 0.60 |
| 5:AE:147:ASP:HA | 5:AE:150:ARG:HH11 | 1.67 | 0.60 |
| 7:AG:79:ARG:NE | 7:AG:84:ASN:HD21 | 2.00 | 0.60 |
| 14:AN:40:CYS:SG | 14:AN:42:ILE:HG12 | 2.41 | 0.60 |
| 28:B6:25:LYS:O | 31:BA:2286:A:H2 | 1.84 | 0.60 |
| 31:BA:901:A:H5' | 31:BA:902:C:OP2 | 2.01 | 0.60 |
| 31:BA:1986:A:H2' | 31:BA:1987:G:C5' | 2.23 | 0.60 |
| 31:BA:2262:U:O2' | 31:BA:2263:C:H5' | 2.01 | 0.60 |
| 31:BA:2759:G:C8 | 31:BA:2759:G:C5' | 2.85 | 0.60 |
| 34:BE:111:ARG:HA | 43:BR:2:ARG:HG3 | 1.82 | 0.60 |
| 47:BV:5:VAL:HG23 | 47:BV:36:PRO:HB2 | 1.84 | 0.60 |
| 1:CA:1311:G:N2 | 1:CA:1327:C:C2 | 2.70 | 0.60 |
| 2:CB:21:ARG:CB | 2:CB:39:ILE:HA | 2.32 | 0.60 |
| 23:D1:46:LEU:H | 23:D1:46:LEU:CD1 | 2.08 | 0.60 |
| 31:DA:1701:A:H2' | 31:DA:1702:G:H5' | 1.84 | 0.60 |
| 31:DA:2759:G:C8 | 31:DA:2759:G:C5' | 2.85 | 0.60 |
| 33:DD:58:HIS:HD2 | 33:DD:59:LYS:H | 1.49 | 0.60 |
| 35:DF:18:ARG:CG | 35:DF:19:GLU:H | 2.10 | 0.60 |
| 42:DQ:43:THR:HB | 42:DQ:45:GLN:HG2 | 1.83 | 0.60 |
| 45:DT:23:ARG:HB2 | 45:DT:24:PRO:CD | 2.23 | 0.60 |
| 47:DV:21:ARG:HG2 | 47:DV:93:GLU:OE1 | 2.02 | 0.60 |
| 1:AA:142:G:C2 | 1:AA:143:A:C8 | 2.90 | 0.60 |
| 1:AA:983:A:H2 | 1:AA:984:C:C6 | 2.20 | 0.60 |
| 6:AF:67:MET:HE1 | 6:AF:75:LEU:HD12 | 1.84 | 0.60 |
| 14:AN:4:LYS:O | 14:AN:7:ILE:HG12 | 2.01 | 0.60 |
| 28:B6:20:ASN:O | 28:B6:21:TYR:CD1 | 2.55 | 0.60 |
| 30:B8:39:LYS:HE2 | 30:B8:39:LYS:C | 2.27 | 0.60 |
| 31:BA:59:U:OP1 | 49:BX:72:LYS:HE2 | 2.02 | 0.60 |
| 31:BA:154:G:H2' | 31:BA:154(A):C:O2 | 2.01 | 0.60 |
| 31:BA:1981:A:H5" | 31:BA:1982:C:OP2 | 2.01 | 0.60 |
| 31:BA:2291:U:H5" | 31:BA:2380:C:O2 | 2.02 | 0.60 |
| 32:BB:29:A:C2 | 32:BB:30:C:C2 | 2.89 | 0.60 |
| 33:BD:176:ARG:O | 33:BD:176:ARG:HG3 | 2.01 | 0.60 |
| 38:BI:4:ILE:HD11 | 38:BI:44:LEU:HD23 | 1.84 | 0.60 |
| 39:BN:15:LEU:HD13 | 39:BN:16:ILE:N | 2.17 | 0.60 |
| 43:BR:72:ASP:HB3 | 43:BR:75:LEU:HB2 | 1.83 | 0.60 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 43:BR:101:ALA:O | 43:BR:102:GLU:HB2 | 2.02 | 0.60 |
| 45:BT:65:LYS:HG3 | 45:BT:66:VAL:N | 2.17 | 0.60 |
| 1:CA:147:G:H2' | 1:CA:148:G:H5' | 1.84 | 0.60 |
| 2:CB:21:ARG:HB3 | 2:CB:39:ILE:HA | 1.83 | 0.60 |
| 3:CC:150:LYS:HB3 | 3:CC:201:TYR:HB2 | 1.83 | 0.60 |
| 31:DA:143:G:H2' | 31:DA:143(A):C:H6 | 1.67 | 0.60 |
| 31:DA:570:G:H2' | 31:DA:2030:A:C6 | 2.37 | 0.60 |
| 31:DA:1484:G:H2' | 31:DA:1485:G:O5' | 2.02 | 0.60 |
| 31:DA:2660:A:H5' | 31:DA:2661:G:C2 | 2.36 | 0.60 |
| 31:DA:2688:U:H5 | 31:DA:2720:U:OP2 | 1.84 | 0.60 |
| 43:DR:44:LEU:HD22 | 43:DR:48:VAL:HG23 | 1.82 | 0.60 |
| 47:DV:64:HIS:CD2 | 47:DV:94:LEU:HD21 | 2.36 | 0.60 |
| 51:DZ:151:HIS:HA | 51:DZ:171:ILE:HG12 | 1.84 | 0.60 |
| 1:AA:233:C:H2' | 1:AA:234:C:H6 | 1.67 | 0.60 |
| 1:AA:370:C:H2' | 1:AA:371:G:O4' | 2.01 | 0.60 |
| 5:AE:18:ARG:HH21 | 5:AE:25:ARG:HG2 | 1.65 | 0.60 |
| 6:AF:3:ARG:HB3 | 6:AF:93:SER:HB2 | 1.84 | 0.60 |
| 9:AI:48:GLU:N | 9:AI:49:PRO:HD2 | 2.16 | 0.60 |
| 30:B8:53:PRO:HA | 30:B8:56:GLU:HB2 | 1.83 | 0.60 |
| 31:BA:49:A:C4' | 31:BA:50:U:H5' | 2.31 | 0.60 |
| 31:BA:176:G:C2' | 31:BA:177:G:H5' | 2.30 | 0.60 |
| 31:BA:547:A:C8 | 31:BA:549:G:O6 | 2.55 | 0.60 |
| 31:BA:1115:G:H2' | 31:BA:1116:C:H5" | 1.84 | 0.60 |
| 31:BA:1245:G:OP1 | 41:BP:16:ARG:HG2 | 2.02 | 0.60 |
| 31:BA:1983:C:O2' | 31:BA:1984:G:H5' | 2.01 | 0.60 |
| 31:BA:2532:G:O2' | 31:BA:2657:A:N6 | 2.34 | 0.60 |
| 31:BA:2810:A:C2' | 34:BE:61:ARG:NH2 | 2.64 | 0.60 |
| 32:BB:74:U:C3' | 32:BB:75:G:H5" | 2.32 | 0.60 |
| 32:BB:87:G:H3' | 32:BB:88:C:C5' | 2.21 | 0.60 |
| 33:BD:144:ALA:HB3 | 33:BD:192:THR:HG23 | 1.84 | 0.60 |
| 33:BD:176:ARG:HH11 | 33:BD:176:ARG:HG2 | 1.67 | 0.60 |
| 34:BE:167:VAL:HG13 | 34:BE:170:LEU:HD11 | 1.84 | 0.60 |
| 35:BF:132:VAL:HG22 | 35:BF:133:ASN:N | 2.17 | 0.60 |
| 36:BG:108:ASN:O | 36:BG:112:PRO:HG2 | 2.02 | 0.60 |
| 45:BT:38:ASN:C | 45:BT:38:ASN:ND2 | 2.56 | 0.60 |
| 48:BW:59:VAL:CG1 | 48:BW:60:ASN:N | 2.63 | 0.60 |
| 49:BX:35:THR:O | 49:BX:36:LYS:O | 2.20 | 0.60 |
| 1:CA:359:U:H2' | 1:CA:360:A:C8 | 2.36 | 0.60 |
| 1:CA:562:C:H4' | 1:CA:563:A:O5' | 2.01 | 0.60 |
| 1:CA:673:G:O3' | 6:CF:87:ARG:NH2 | 2.35 | 0.60 |
| 1:CA:681:C:N3 | 1:CA:710:G:C2 | 2.70 | 0.60 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:949:A:H1' | 1:CA:1364:U:H3 | 1.66 | 0.60 |
| 1:CA:1189:C:O3' | 3:CC:5:ILE:HD12 | 2.02 | 0.60 |
| 1:CA:1521:G:H2' | 1:CA:1522:U:C6 | 2.37 | 0.60 |
| 4:CD:12:CYS:HA | 4:CD:19:LEU:CD1 | 2.31 | 0.60 |
| 5:CE:18:ARG:HH21 | 5:CE:25:ARG:HG2 | 1.67 | 0.60 |
| 24:D2:34:GLU:O | 24:D2:36:ARG:N | 2.35 | 0.60 |
| 31:DA:234:C:H2' | 31:DA:235:U:C6 | 2.37 | 0.60 |
| 31:DA:1336:A:H2' | 31:DA:1337:G:H8 | 1.66 | 0.60 |
| 31:DA:2360:A:HO2' | 31:DA:2361:A:C5' | 2.14 | 0.60 |
| 31:DA:2713:A:H3' | 31:DA:2714:G:C5' | 2.32 | 0.60 |
| 41:DP:83:VAL:CG1 | 41:DP:112:LEU:HD21 | 2.31 | 0.60 |
| 41:DP:140:ALA:O | 41:DP:141:ALA:HB2 | 2.01 | 0.60 |
| 45:DT:109:GLU:HA | 45:DT:112:ARG:CG | 2.31 | 0.60 |
| 49:DX:72:LYS:C | 49:DX:74:PRO:HD3 | 2.27 | 0.60 |
| 1:AA:724:G:C2 | 1:AA:725:G:C8 | 2.89 | 0.59 |
| 2:AB:17:PHE:CD1 | 2:AB:44:LEU:HD11 | 2.37 | 0.59 |
| 3:AC:6:HIS:HD2 | 3:AC:7:PRO:HD2 | 1.66 | 0.59 |
| 3:AC:130:VAL:O | 3:AC:134:ILE:HG12 | 2.02 | 0.59 |
| 16:AP:9:PHE:CE2 | 16:AP:18:ARG:NE | 2.70 | 0.59 |
| 29:B7:11:LYS:HE2 | 31:BA:686:G:H5" | 1.84 | 0.59 |
| 31:BA:632:A:O2' | 31:BA:633:A:H5' | 2.01 | 0.59 |
| 31:BA:662:G:P | 41:BP:18:ARG:HD2 | 2.41 | 0.59 |
| 31:BA:823:G:O2' | 31:BA:824:A:H5' | 2.02 | 0.59 |
| 31:BA:1317:A:H2' | 31:BA:1318:C:H6 | 1.65 | 0.59 |
| 31:BA:2186:G:C3' | 31:BA:2187:G:H5" | 2.32 | 0.59 |
| 31:BA:2801(A):A:HO2' | 31:BA:2803:C:H5 | 1.48 | 0.59 |
| 33:BD:95:LEU:H | 33:BD:95:LEU:HD12 | 1.66 | 0.59 |
| 34:BE:181:LEU:HD21 | 45:BT:7:ILE:HG23 | 1.84 | 0.59 |
| 45:BT:30:VAL:HG22 | 45:BT:84:GLN:O | 2.01 | 0.59 |
| 51:BZ:64:GLY:O | 51:BZ:65:GLN:O | 2.20 | 0.59 |
| 1:CA:524:G:H2' | 1:CA:525:C:C6 | 2.37 | 0.59 |
| 1:CA:1391:U:H2' | 1:CA:1392:G:C8 | 2.37 | 0.59 |
| 1:CA:1442:G:O2' | 1:CA:1442(A):G:C5' | 2.48 | 0.59 |
| 3:CC:6:HIS:HD2 | 3:CC:7:PRO:HD2 | 1.66 | 0.59 |
| 26:D4:5:ILE:C | 36:DG:67:LYS:HG2 | 2.27 | 0.59 |
| 27:D5:51:TYR:HB3 | 27:D5:52:TYR:O | 2.02 | 0.59 |
| 31:DA:863:A:O2' | 31:DA:864:G:H5' | 2.01 | 0.59 |
| 31:DA:1501:C:H2' | 31:DA:1502:C:H6 | 1.67 | 0.59 |
| 31:DA:1839:G:N7 | 31:DA:1927:A:H1' | 2.16 | 0.59 |
| 31:DA:2376:A:OP1 | 31:DA:2376:A:H8 | 1.85 | 0.59 |
| 31:DA:2580:U:C5' | 34:DE:131:ALA:H | 2.15 | 0.59 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2723:C:H5” | 43:DR:2:ARG:HD3 | 1.85 | 0.59 |
| 31:DA:2870:C:H2’ | 31:DA:2871:C:H5’ | 1.84 | 0.59 |
| 32:DB:66:A:C6 | 32:DB:109:C:C5 | 2.89 | 0.59 |
| 32:DB:81:G:H5” | 32:DB:82:G:OP2 | 2.02 | 0.59 |
| 38:DI:10:GLU:O | 38:DI:12:LEU:HD23 | 2.01 | 0.59 |
| 39:DN:63:THR:O | 39:DN:64:GLY:O | 2.20 | 0.59 |
| 42:DQ:76:LYS:H | 42:DQ:88:GLY:HA2 | 1.67 | 0.59 |
| 47:DV:36:PRO:CG | 47:DV:62:LEU:HD11 | 2.32 | 0.59 |
| 50:DY:100:ALA:O | 50:DY:101:LYS:HB2 | 2.02 | 0.59 |
| 1:AA:78:G:H1 | 1:AA:91:C:H42 | 1.48 | 0.59 |
| 1:AA:376:G:H4’ | 16:AP:5:ARG:NH1 | 2.10 | 0.59 |
| 1:AA:672:U:O2 | 1:AA:672:U:H2’ | 2.00 | 0.59 |
| 1:AA:1159:U:H4’ | 1:AA:1160:G:OP1 | 2.01 | 0.59 |
| 27:B5:2:ALA:N | 31:BA:747:U:C4 | 2.70 | 0.59 |
| 31:BA:1502:C:O2 | 31:BA:1502:C:C2’ | 2.49 | 0.59 |
| 31:BA:2657:A:H2 | 31:BA:2664:G:N2 | 2.01 | 0.59 |
| 33:BD:25:THR:HG23 | 33:BD:27:THR:HB | 1.83 | 0.59 |
| 39:BN:3:THR:HA | 39:BN:4:TYR:CD1 | 2.37 | 0.59 |
| 47:BV:54:GLY:O | 47:BV:56:SER:N | 2.31 | 0.59 |
| 50:BY:44:ILE:H | 50:BY:44:ILE:CD1 | 2.12 | 0.59 |
| 51:BZ:151:HIS:HA | 51:BZ:171:ILE:HG12 | 1.83 | 0.59 |
| 1:CA:386:C:C2’ | 1:CA:387:U:H5’ | 2.32 | 0.59 |
| 1:CA:1049:U:H4’ | 1:CA:1050:G:O5’ | 2.02 | 0.59 |
| 1:CA:1399:C:H4’ | 1:CA:1400:C:H5” | 1.85 | 0.59 |
| 6:CF:3:ARG:HB3 | 6:CF:93:SER:HB2 | 1.84 | 0.59 |
| 8:CH:31:PHE:O | 8:CH:35:ILE:HG13 | 2.02 | 0.59 |
| 31:DA:1331:A:H2’ | 31:DA:1333:C:H5 | 1.67 | 0.59 |
| 31:DA:1686:C:C2’ | 31:DA:1687:G:H5’ | 2.32 | 0.59 |
| 35:DF:129:PHE:CD2 | 35:DF:163:VAL:HG21 | 2.36 | 0.59 |
| 39:DN:39:ARG:HE | 39:DN:41:ASP:HB2 | 1.67 | 0.59 |
| 39:DN:96:GLU:CD | 39:DN:96:GLU:H | 2.10 | 0.59 |
| 49:DX:40:LYS:HG3 | 49:DX:51:VAL:HB | 1.83 | 0.59 |
| 1:AA:376:G:OP2 | 16:AP:67:THR:HG21 | 2.02 | 0.59 |
| 1:AA:1452:C:H5’ | 1:AA:1456:G:C4 | 2.37 | 0.59 |
| 3:AC:113:ALA:C | 3:AC:115:LEU:H | 2.08 | 0.59 |
| 4:AD:31:CYS:SG | 4:AD:31:CYS:O | 2.59 | 0.59 |
| 10:AJ:6:ILE:HD11 | 10:AJ:72:VAL:HB | 1.83 | 0.59 |
| 15:AO:39:LEU:HD13 | 15:AO:56:LEU:HB2 | 1.82 | 0.59 |
| 20:AT:13:LEU:H | 20:AT:13:LEU:HD12 | 1.67 | 0.59 |
| 20:AT:73:HIS:O | 20:AT:76:ALA:HB3 | 2.02 | 0.59 |
| 31:BA:960:A:H5” | 31:BA:961:C:OP2 | 2.01 | 0.59 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1332:G:N2 | 31:BA:1609:A:O2' | 2.35 | 0.59 |
| 31:BA:1348:G:H2' | 31:BA:1349:A:H5' | 1.83 | 0.59 |
| 32:BB:21:G:O6 | 32:BB:63:G:C5 | 2.55 | 0.59 |
| 35:BF:22:ALA:C | 35:BF:26:ALA:HB2 | 2.26 | 0.59 |
| 39:BN:75:TYR:N | 39:BN:75:TYR:CD1 | 2.70 | 0.59 |
| 46:BU:59:ARG:O | 46:BU:60:LEU:C | 2.46 | 0.59 |
| 1:CA:341:C:C2' | 1:CA:342:C:H5' | 2.32 | 0.59 |
| 1:CA:1159:U:H4' | 1:CA:1160:G:OP1 | 2.02 | 0.59 |
| 1:CA:1372:U:H5" | 9:CI:71:SER:HB3 | 1.83 | 0.59 |
| 1:CA:1372:U:OP1 | 9:CI:72:GLY:N | 2.35 | 0.59 |
| 1:CA:1521:G:H2' | 1:CA:1522:U:H6 | 1.67 | 0.59 |
| 8:CH:53:VAL:O | 8:CH:54:ASP:HB2 | 2.02 | 0.59 |
| 9:CI:111:ARG:O | 9:CI:113:LYS:HD2 | 2.02 | 0.59 |
| 22:D0:24:LYS:O | 22:D0:25:ARG:HD2 | 2.02 | 0.59 |
| 23:D1:34:THR:CG2 | 31:DA:388:G:P | 2.87 | 0.59 |
| 23:D1:65:SER:N | 23:D1:67:ILE:CD1 | 2.66 | 0.59 |
| 30:D8:14:VAL:HG13 | 30:D8:22:VAL:HG13 | 1.83 | 0.59 |
| 31:DA:1803:A:H4' | 33:DD:259:THR:CG2 | 2.33 | 0.59 |
| 31:DA:2415:G:C4' | 41:DP:67:MET:H | 2.08 | 0.59 |
| 33:DD:172:TYR:CD1 | 33:DD:186:HIS:HA | 2.37 | 0.59 |
| 38:DI:4:ILE:HD11 | 38:DI:44:LEU:HD23 | 1.85 | 0.59 |
| 44:DS:34:HIS:N | 44:DS:34:HIS:CD2 | 2.71 | 0.59 |
| 45:DT:38:ASN:ND2 | 45:DT:40:THR:H | 2.00 | 0.59 |
| 1:AA:922:G:C6 | 1:AA:923:A:C6 | 2.90 | 0.59 |
| 2:AB:228:GLY:O | 2:AB:230:VAL:HG13 | 2.02 | 0.59 |
| 3:AC:6:HIS:CD2 | 3:AC:7:PRO:HD2 | 2.38 | 0.59 |
| 5:AE:12:LEU:O | 5:AE:13:ILE:HD12 | 2.02 | 0.59 |
| 24:B2:34:GLU:O | 24:B2:36:ARG:N | 2.36 | 0.59 |
| 31:BA:143:G:C1' | 49:BX:38:GLU:HG3 | 2.31 | 0.59 |
| 31:BA:624:C:C2' | 31:BA:625:G:H5' | 2.31 | 0.59 |
| 31:BA:966:G:C6 | 31:BA:967:C:N4 | 2.70 | 0.59 |
| 31:BA:1502:C:H5' | 31:BA:1503:U:OP2 | 2.02 | 0.59 |
| 31:BA:1654:A:H1' | 31:BA:2823:A:H5' | 1.82 | 0.59 |
| 41:BP:50:ARG:HG2 | 41:BP:50:ARG:HH21 | 1.67 | 0.59 |
| 46:BU:91:ASP:OD2 | 46:BU:96:ALA:HB2 | 2.01 | 0.59 |
| 51:BZ:149:SER:HB3 | 51:BZ:173:ALA:HA | 1.84 | 0.59 |
| 51:BZ:150:LEU:N | 51:BZ:150:LEU:HD13 | 2.18 | 0.59 |
| 1:CA:56:U:H2' | 1:CA:57:G:C8 | 2.38 | 0.59 |
| 1:CA:299:G:C6 | 1:CA:300:A:C6 | 2.91 | 0.59 |
| 1:CA:392:G:H2' | 1:CA:393:A:C8 | 2.37 | 0.59 |
| 1:CA:774:G:H2' | 1:CA:775:G:H5' | 1.85 | 0.59 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:1313:U:P | 19:CS:6:LYS:HG3 | 2.42 | 0.59 |
| 6:CF:4:TYR:HA | 6:CF:91:VAL:O | 2.03 | 0.59 |
| 6:CF:61:LEU:HB3 | 6:CF:63:TYR:HE2 | 1.67 | 0.59 |
| 7:CG:75:VAL:HG12 | 7:CG:88:PRO:HB3 | 1.83 | 0.59 |
| 9:CI:83:ARG:O | 9:CI:86:VAL:HG12 | 2.02 | 0.59 |
| 17:CQ:59:ILE:HG22 | 17:CQ:71:PHE:HD1 | 1.68 | 0.59 |
| 31:DA:1022:G:N2 | 31:DA:1142(A):A:H2 | 2.00 | 0.59 |
| 31:DA:1051:G:C2 | 31:DA:1052:C:N4 | 2.70 | 0.59 |
| 31:DA:1052:C:H6 | 31:DA:1052:C:H3' | 1.68 | 0.59 |
| 31:DA:1191:G:OP1 | 41:DP:35:HIS:ND1 | 2.35 | 0.59 |
| 31:DA:1509(B):A:H2' | 31:DA:1510:G:H8 | 1.67 | 0.59 |
| 31:DA:2023:G:H5' | 31:DA:2617:C:H4' | 1.84 | 0.59 |
| 31:DA:2201:C:O2' | 31:DA:2202:C:H5' | 2.02 | 0.59 |
| 31:DA:2637:U:H5' | 31:DA:2637:U:H6 | 1.66 | 0.59 |
| 34:DE:116:VAL:CG1 | 34:DE:122:PHE:CD2 | 2.85 | 0.59 |
| 34:DE:182:LEU:HD12 | 34:DE:183:LEU:N | 2.16 | 0.59 |
| 38:DI:67:ARG:O | 38:DI:68:LEU:HB2 | 2.02 | 0.59 |
| 39:DN:39:ARG:HE | 39:DN:41:ASP:CG | 2.10 | 0.59 |
| 42:DQ:42:ILE:HD13 | 42:DQ:97:VAL:CG2 | 2.32 | 0.59 |
| 45:DT:29:ARG:HG2 | 45:DT:86:ILE:N | 2.16 | 0.59 |
| 45:DT:38:ASN:HD22 | 45:DT:40:THR:H | 1.48 | 0.59 |
| 45:DT:102:ILE:HB | 45:DT:110:ILE:HD13 | 1.83 | 0.59 |
| 46:DU:100:VAL:O | 46:DU:103:PRO:HD3 | 2.02 | 0.59 |
| 48:DW:62:HIS:O | 48:DW:64:MET:HG3 | 2.02 | 0.59 |
| 50:DY:37:VAL:HG11 | 50:DY:72:VAL:CG2 | 2.32 | 0.59 |
| 51:DZ:149:SER:HB3 | 51:DZ:173:ALA:HA | 1.84 | 0.59 |
| 1:AA:1064:G:H1' | 1:AA:1065:U:OP2 | 2.02 | 0.59 |
| 1:AA:1346:A:N1 | 1:AA:1374:A:H5" | 2.18 | 0.59 |
| 2:AB:101:MET:HG2 | 2:AB:108:ILE:HG21 | 1.84 | 0.59 |
| 10:AJ:32:ALA:HB2 | 10:AJ:76:ASN:HB3 | 1.85 | 0.59 |
| 22:B0:53:MET:HB2 | 22:B0:59:LEU:HD23 | 1.85 | 0.59 |
| 23:B1:46:LEU:CD1 | 23:B1:46:LEU:H | 2.07 | 0.59 |
| 23:B1:63:ALA:O | 23:B1:64:ALA:C | 2.45 | 0.59 |
| 26:B4:11:PRO:C | 26:B4:13:ARG:H | 2.09 | 0.59 |
| 31:BA:999:U:O2' | 31:BA:1000:A:H5' | 2.02 | 0.59 |
| 31:BA:1434:A:C2' | 31:BA:1435:G:H5' | 2.32 | 0.59 |
| 31:BA:1484:G:H2' | 31:BA:1485:G:O5' | 2.02 | 0.59 |
| 31:BA:2475:C:C5' | 31:BA:2476:A:OP2 | 2.49 | 0.59 |
| 33:BD:8:PRO:CB | 33:BD:14:ARG:HB3 | 2.32 | 0.59 |
| 33:BD:35:LYS:HG2 | 33:BD:64:ILE:CA | 2.32 | 0.59 |
| 34:BE:109:LYS:HD3 | 43:BR:2:ARG:NH1 | 2.17 | 0.59 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 41:BP:121:LYS:HG3 | 41:BP:122:PRO:HD2 | 1.83 | 0.59 |
| 1:CA:193:C:H2' | 1:CA:194:C:H6 | 1.66 | 0.59 |
| 12:CL:90:VAL:O | 12:CL:92:ASP:N | 2.32 | 0.59 |
| 18:CR:50:ILE:CD1 | 18:CR:70:ILE:HG21 | 2.29 | 0.59 |
| 22:D0:26:TYR:CE2 | 31:DA:857:C:H1' | 2.36 | 0.59 |
| 24:D2:25:VAL:HG13 | 24:D2:26:ARG:HD2 | 1.83 | 0.59 |
| 25:D3:47:VAL:HG11 | 25:D3:56:VAL:HG21 | 1.84 | 0.59 |
| 31:DA:527:C:OP2 | 31:DA:2779:U:C5 | 2.50 | 0.59 |
| 31:DA:528:A:H5" | 31:DA:528:A:C8 | 2.36 | 0.59 |
| 31:DA:1022:G:C5 | 31:DA:1140:C:N4 | 2.70 | 0.59 |
| 31:DA:1899:G:N2 | 31:DA:1902:C:N4 | 2.31 | 0.59 |
| 31:DA:2335:A:C8 | 31:DA:2337:G:C5 | 2.90 | 0.59 |
| 31:DA:2631:G:N2 | 34:DE:61:ARG:HH12 | 1.99 | 0.59 |
| 31:DA:2652:C:C2' | 31:DA:2653:U:H5' | 2.32 | 0.59 |
| 33:DD:25:THR:HG23 | 33:DD:27:THR:HB | 1.85 | 0.59 |
| 34:DE:87:GLU:O | 34:DE:89:ASP:N | 2.35 | 0.59 |
| 39:DN:3:THR:CA | 39:DN:4:TYR:CD1 | 2.85 | 0.59 |
| 39:DN:23:LEU:CD1 | 39:DN:98:VAL:HG12 | 2.32 | 0.59 |
| 39:DN:78:TYR:HD1 | 39:DN:79:PRO:N | 1.99 | 0.59 |
| 42:DQ:140:ALA:CB | 51:DZ:53:ILE:HG13 | 2.31 | 0.59 |
| 44:DS:34:HIS:CE1 | 44:DS:54:LEU:HB2 | 2.37 | 0.59 |
| 45:DT:80:SER:HB3 | 45:DT:81:PRO:HD3 | 1.84 | 0.59 |
| 50:DY:88:LYS:NZ | 50:DY:93:GLY:HA3 | 2.17 | 0.59 |
| 1:AA:341:C:C2' | 1:AA:342:C:H5' | 2.33 | 0.59 |
| 1:AA:524:G:H2' | 1:AA:525:C:C6 | 2.38 | 0.59 |
| 1:AA:616:G:C2 | 1:AA:617:G:C8 | 2.91 | 0.59 |
| 1:AA:623:C:C4 | 1:AA:624:C:C5 | 2.90 | 0.59 |
| 5:AE:51:VAL:HB | 5:AE:52:PRO:CD | 2.32 | 0.59 |
| 23:B1:85:LEU:C | 23:B1:87:PRO:CD | 2.71 | 0.59 |
| 25:B3:19:GLN:NE2 | 25:B3:52:HIS:CE1 | 2.71 | 0.59 |
| 28:B6:19:ARG:NH2 | 31:BA:2401:U:OP1 | 2.35 | 0.59 |
| 31:BA:542:C:H2' | 31:BA:543:C:OP1 | 2.03 | 0.59 |
| 31:BA:2196:C:C2' | 31:BA:2197:U:H5' | 2.32 | 0.59 |
| 34:BE:61:ARG:C | 34:BE:63:LEU:H | 2.09 | 0.59 |
| 41:BP:120:ALA:CB | 41:BP:138:LEU:HB3 | 2.32 | 0.59 |
| 42:BQ:89:ASN:O | 42:BQ:92:GLY:N | 2.27 | 0.59 |
| 44:BS:34:HIS:CE1 | 44:BS:54:LEU:HB2 | 2.38 | 0.59 |
| 44:BS:67:ARG:H | 44:BS:69:VAL:HG12 | 1.68 | 0.59 |
| 47:BV:5:VAL:HG21 | 47:BV:36:PRO:HB2 | 1.85 | 0.59 |
| 49:BX:64:LYS:O | 49:BX:65:ARG:HB2 | 2.01 | 0.59 |
| 50:BY:52:SER:C | 50:BY:54:LYS:H | 2.10 | 0.59 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 51:BZ:106:GLY:HA3 | 51:BZ:142:SER:HB3 | 1.85 | 0.59 |
| 51:BZ:120:ILE:HG22 | 51:BZ:120:ILE:O | 2.02 | 0.59 |
| 1:CA:114:U:H2' | 1:CA:115:G:H8 | 1.68 | 0.59 |
| 3:CC:105:GLU:HG2 | 3:CC:106:VAL:H | 1.67 | 0.59 |
| 5:CE:80:ILE:CD1 | 5:CE:138:ALA:HB1 | 2.32 | 0.59 |
| 9:CI:26:VAL:HG22 | 9:CI:61:ALA:HB3 | 1.83 | 0.59 |
| 16:CP:76:GLN:HG2 | 16:CP:76:GLN:O | 2.02 | 0.59 |
| 31:DA:269:U:H2' | 31:DA:269:U:O2 | 2.01 | 0.59 |
| 31:DA:445:C:OP1 | 46:DU:2:PRO:HA | 2.03 | 0.59 |
| 31:DA:1188:U:O2' | 31:DA:1189:A:H5' | 2.02 | 0.59 |
| 31:DA:1339:G:N2 | 31:DA:1603:A:H1' | 2.18 | 0.59 |
| 31:DA:2655:G:N3 | 31:DA:2664:G:O6 | 2.36 | 0.59 |
| 32:DB:82:G:H2' | 32:DB:83:G:H5' | 1.84 | 0.59 |
| 33:DD:85:ASP:HB2 | 33:DD:92:ILE:HG12 | 1.85 | 0.59 |
| 38:DI:133:HIS:CG | 38:DI:134:PRO:HD2 | 2.38 | 0.59 |
| 44:DS:29:PHE:HD2 | 44:DS:30:ARG:N | 2.01 | 0.59 |
| 46:DU:51:LYS:O | 46:DU:52:ARG:C | 2.46 | 0.59 |
| 50:DY:75:ILE:HD11 | 50:DY:80:GLY:N | 2.18 | 0.59 |
| 2:AB:187:LEU:HD23 | 2:AB:201:ILE:HG22 | 1.85 | 0.59 |
| 27:B5:57:VAL:C | 27:B5:58:LEU:HG | 2.27 | 0.59 |
| 30:B8:50:LEU:HD12 | 30:B8:51:ALA:N | 2.18 | 0.59 |
| 31:BA:320:A:OP2 | 35:BF:137:LYS:HE3 | 2.02 | 0.59 |
| 31:BA:535:C:O2' | 31:BA:536:A:H5' | 2.02 | 0.59 |
| 31:BA:753:C:O5' | 31:BA:753:C:H6 | 1.85 | 0.59 |
| 31:BA:1106:A:C2' | 31:BA:1107:G:O5' | 2.51 | 0.59 |
| 31:BA:2567:G:H2' | 31:BA:2568:C:C6 | 2.37 | 0.59 |
| 31:BA:2652:C:C2' | 31:BA:2653:U:H5' | 2.32 | 0.59 |
| 31:BA:2836:U:H2' | 31:BA:2837:G:C8 | 2.37 | 0.59 |
| 41:BP:30:THR:CG2 | 41:BP:31:ALA:H | 2.15 | 0.59 |
| 50:BY:37:VAL:HG11 | 50:BY:72:VAL:HG21 | 1.84 | 0.59 |
| 51:BZ:71:VAL:HG22 | 51:BZ:88:PHE:CE2 | 2.36 | 0.59 |
| 1:CA:7:G:N2 | 5:CE:121:LYS:HG2 | 2.18 | 0.59 |
| 1:CA:142:G:C2 | 1:CA:143:A:C8 | 2.91 | 0.59 |
| 1:CA:375:U:O3' | 16:CP:6:LEU:HB2 | 2.02 | 0.59 |
| 1:CA:509:A:O2' | 1:CA:510:A:O5' | 2.21 | 0.59 |
| 1:CA:921:U:O2 | 5:CE:19:MET:HB2 | 2.02 | 0.59 |
| 1:CA:922:G:O2' | 1:CA:1398:A:N1 | 2.31 | 0.59 |
| 1:CA:1325:C:H4' | 21:CU:17:THR:HG21 | 1.84 | 0.59 |
| 1:CA:1348:U:H4' | 9:CI:120:ARG:HD2 | 1.82 | 0.59 |
| 2:CB:219:VAL:HA | 2:CB:222:ILE:HD12 | 1.85 | 0.59 |
| 3:CC:6:HIS:CD2 | 3:CC:7:PRO:HD2 | 2.38 | 0.59 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 12:CL:18:VAL:HG23 | 12:CL:19:ARG:H | 1.67 | 0.59 |
| 30:D8:4:MET:SD | 30:D8:61:LEU:HD12 | 2.43 | 0.59 |
| 30:D8:61:LEU:C | 30:D8:63:PRO:HD2 | 2.28 | 0.59 |
| 31:DA:84:A:N1 | 31:DA:98:G:O2' | 2.34 | 0.59 |
| 31:DA:102:G:C2' | 31:DA:103:A:OP2 | 2.51 | 0.59 |
| 31:DA:128:C:C6 | 31:DA:128:C:H3' | 2.38 | 0.59 |
| 31:DA:535:C:O2' | 31:DA:536:A:H5' | 2.03 | 0.59 |
| 31:DA:2186:G:C3' | 31:DA:2187:G:H5" | 2.32 | 0.59 |
| 32:DB:28:C:H2' | 32:DB:29:A:H8 | 1.66 | 0.59 |
| 33:DD:35:LYS:HB3 | 33:DD:63:ARG:HA | 1.85 | 0.59 |
| 39:DN:111:PRO:HA | 39:DN:114:ARG:NH1 | 2.17 | 0.59 |
| 46:DU:8:VAL:CG1 | 46:DU:12:ARG:HG3 | 2.33 | 0.59 |
| 46:DU:106:PHE:O | 46:DU:110:VAL:HG23 | 2.03 | 0.59 |
| 5:AE:101:ILE:HG12 | 5:AE:101:ILE:O | 2.01 | 0.59 |
| 8:AH:9:MET:HG2 | 8:AH:10:LEU:HD23 | 1.83 | 0.59 |
| 30:B8:14:VAL:HG13 | 30:B8:22:VAL:HG13 | 1.85 | 0.59 |
| 31:BA:924:C:H2' | 31:BA:925:C:C6 | 2.38 | 0.59 |
| 31:BA:2197:U:H1' | 31:BA:2198:A:C8 | 2.38 | 0.59 |
| 31:BA:2660:A:H5' | 31:BA:2661:G:C2 | 2.37 | 0.59 |
| 31:BA:2862:G:H2' | 31:BA:2863:C:H6 | 1.67 | 0.59 |
| 32:BB:75:G:H8 | 32:BB:75:G:H5' | 1.67 | 0.59 |
| 35:BF:22:ALA:O | 35:BF:26:ALA:HB2 | 2.03 | 0.59 |
| 38:BI:56:LYS:HZ2 | 38:BI:57:ARG:N | 2.01 | 0.59 |
| 47:BV:80:GLN:O | 47:BV:80:GLN:OE1 | 2.21 | 0.59 |
| 50:BY:100:ALA:O | 50:BY:101:LYS:HB2 | 2.02 | 0.59 |
| 1:CA:137:C:H42 | 1:CA:226:G:H1 | 1.50 | 0.59 |
| 1:CA:1442(B):A:OP1 | 1:CA:1442(B):A:H4' | 2.02 | 0.59 |
| 2:CB:116:GLU:HA | 2:CB:119:GLU:HB2 | 1.84 | 0.59 |
| 3:CC:155:GLY:O | 3:CC:156:ARG:HB2 | 2.02 | 0.59 |
| 15:CO:3:ILE:HD13 | 15:CO:3:ILE:H | 1.68 | 0.59 |
| 20:CT:61:SER:O | 20:CT:65:LYS:HG3 | 2.02 | 0.59 |
| 20:CT:63:ILE:HD13 | 20:CT:80:ARG:HB2 | 1.84 | 0.59 |
| 31:DA:1963:U:O2 | 31:DA:1963:U:H2' | 2.01 | 0.59 |
| 31:DA:2392:A:H2 | 31:DA:2424:C:N4 | 1.99 | 0.59 |
| 31:DA:2476:A:H2 | 31:DA:2477:C:H5" | 1.66 | 0.59 |
| 31:DA:2702:U:HO2' | 31:DA:2703:C:H5 | 1.48 | 0.59 |
| 31:DA:2734:A:C5' | 31:DA:2734:A:H8 | 2.15 | 0.59 |
| 33:DD:71:ASP:HB3 | 33:DD:103:ARG:HH22 | 1.66 | 0.59 |
| 35:DF:22:ALA:O | 35:DF:26:ALA:HB2 | 2.03 | 0.59 |
| 38:DI:25:TYR:CD1 | 38:DI:30:LEU:HD11 | 2.37 | 0.59 |
| 40:DO:23:ARG:HD2 | 40:DO:24:VAL:H | 1.67 | 0.59 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 44:DS:33:LYS:HB3 | 44:DS:34:HIS:CD2 | 2.37 | 0.59 |
| 1:AA:336:C:O2' | 1:AA:337:C:H5' | 2.03 | 0.59 |
| 23:B1:86:SER:HA | 23:B1:89:GLU:OE2 | 2.01 | 0.59 |
| 27:B5:51:TYR:HB3 | 27:B5:52:TYR:O | 2.02 | 0.59 |
| 31:BA:942:G:H5' | 41:BP:35:HIS:HB2 | 1.85 | 0.59 |
| 31:BA:1934:C:H5" | 31:BA:1934:C:C6 | 2.29 | 0.59 |
| 31:BA:2307:G:N2 | 31:BA:2308:G:C5' | 2.65 | 0.59 |
| 33:BD:44:ASN:CB | 33:BD:49:ILE:HA | 2.23 | 0.59 |
| 34:BE:75:VAL:C | 34:BE:77:ILE:N | 2.61 | 0.59 |
| 34:BE:137:HIS:HB3 | 34:BE:138:PRO:CD | 2.33 | 0.59 |
| 1:CA:774:G:C2' | 1:CA:775:G:H5' | 2.32 | 0.59 |
| 1:CA:853:G:H2' | 1:CA:854:G:H8 | 1.68 | 0.59 |
| 2:CB:168:THR:HG21 | 2:CB:192:SER:HA | 1.84 | 0.59 |
| 3:CC:113:ALA:C | 3:CC:115:LEU:H | 2.11 | 0.59 |
| 6:CF:62:TRP:HE3 | 6:CF:62:TRP:O | 1.85 | 0.59 |
| 20:CT:56:MET:HG2 | 20:CT:84:LEU:CD1 | 2.33 | 0.59 |
| 21:CU:9:ARG:O | 21:CU:13:ILE:HG13 | 2.03 | 0.59 |
| 24:D2:15:LYS:O | 24:D2:16:LEU:HB3 | 2.02 | 0.59 |
| 31:DA:580:C:H2' | 31:DA:581:C:H6 | 1.67 | 0.59 |
| 31:DA:1464:C:O2' | 31:DA:1528:A:H8 | 1.84 | 0.59 |
| 31:DA:2688:U:C5 | 31:DA:2720:U:OP2 | 2.56 | 0.59 |
| 32:DB:7:G:O5' | 44:DS:29:PHE:CE1 | 2.56 | 0.59 |
| 37:DH:41:MET:CE | 37:DH:55:PRO:HD3 | 2.32 | 0.59 |
| 1:AA:253:U:H2' | 1:AA:254:G:H8 | 1.67 | 0.59 |
| 1:AA:502:G:C6 | 1:AA:503:C:N3 | 2.71 | 0.59 |
| 2:AB:204:ASN:HD21 | 2:AB:207:ALA:H | 1.50 | 0.59 |
| 5:AE:101:ILE:CD1 | 5:AE:119:LEU:HA | 2.33 | 0.59 |
| 10:AJ:34:VAL:HG13 | 10:AJ:73:ASP:O | 2.02 | 0.59 |
| 12:AL:27:LEU:HD11 | 12:AL:64:TYR:CE1 | 2.37 | 0.59 |
| 17:AQ:67:LYS:HA | 17:AQ:70:ARG:HH12 | 1.68 | 0.59 |
| 30:B8:6:THR:HG21 | 31:BA:243:U:OP1 | 2.02 | 0.59 |
| 31:BA:310:A:OP1 | 50:BY:17:SER:O | 2.20 | 0.59 |
| 31:BA:568:U:H5' | 31:BA:945:A:C2 | 2.36 | 0.59 |
| 31:BA:1381:G:H2' | 31:BA:1382:G:H5' | 1.85 | 0.59 |
| 33:BD:69:ARG:NH2 | 33:BD:128:GLY:O | 2.33 | 0.59 |
| 34:BE:21:VAL:HG23 | 34:BE:21:VAL:O | 2.01 | 0.59 |
| 40:BO:20:MET:HE3 | 40:BO:44:LYS:HE3 | 1.85 | 0.59 |
| 44:BS:88:ASP:CG | 44:BS:89:ARG:N | 2.61 | 0.59 |
| 46:BU:106:PHE:O | 46:BU:110:VAL:HG23 | 2.03 | 0.59 |
| 51:BZ:145:GLU:HG3 | 51:BZ:146:ILE:H | 1.68 | 0.59 |
| 1:CA:147:G:C2' | 1:CA:148:G:H5' | 2.32 | 0.59 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 5:CE:57:LYS:HB3 | 5:CE:61:TYR:HE2 | 1.67 | 0.59 |
| 23:D1:10:LYS:O | 23:D1:13:ILE:HG23 | 2.03 | 0.59 |
| 28:D6:16:CYS:O | 28:D6:18:ARG:NH2 | 2.36 | 0.59 |
| 31:DA:8:A:H2' | 31:DA:9:U:C5 | 2.38 | 0.59 |
| 31:DA:234:C:H2' | 31:DA:235:U:H6 | 1.68 | 0.59 |
| 31:DA:1636:C:H2' | 31:DA:1637:A:C8 | 2.37 | 0.59 |
| 31:DA:1636:C:H2' | 31:DA:1637:A:H8 | 1.67 | 0.59 |
| 31:DA:1722:A:O2' | 31:DA:1739:U:H5" | 2.02 | 0.59 |
| 33:DD:27:THR:CG2 | 33:DD:28:GLU:H | 2.16 | 0.59 |
| 34:DE:171:GLU:HB2 | 34:DE:185:LYS:HG2 | 1.85 | 0.59 |
| 37:DH:111:HIS:ND1 | 37:DH:112:PRO:HD2 | 2.17 | 0.59 |
| 43:DR:87:TYR:O | 43:DR:89:ASP:N | 2.30 | 0.59 |
| 46:DU:47:TYR:HA | 46:DU:50:ARG:NH2 | 2.17 | 0.59 |
| 1:AA:687:A:N3 | 1:AA:688:G:H1' | 2.17 | 0.58 |
| 1:AA:1216:G:H5" | 14:AN:5:ALA:CB | 2.33 | 0.58 |
| 1:AA:1301:U:H3' | 1:AA:1302:U:H5" | 1.84 | 0.58 |
| 1:AA:1313:U:P | 19:AS:6:LYS:HG3 | 2.43 | 0.58 |
| 2:AB:185:ILE:CG2 | 2:AB:199:TYR:HB2 | 2.20 | 0.58 |
| 5:AE:18:ARG:NH2 | 5:AE:25:ARG:HG2 | 2.18 | 0.58 |
| 24:B2:14:ARG:O | 24:B2:18:PRO:CD | 2.46 | 0.58 |
| 31:BA:212:G:C2' | 31:BA:213:A:H5' | 2.32 | 0.58 |
| 31:BA:1021:A:N6 | 31:BA:1141:U:H3 | 2.00 | 0.58 |
| 31:BA:1763:G:OP1 | 31:BA:1763:G:H4' | 2.02 | 0.58 |
| 31:BA:1771:C:H1' | 31:BA:1786:A:C8 | 2.38 | 0.58 |
| 31:BA:2030:A:H5" | 31:BA:2030:A:H8 | 1.67 | 0.58 |
| 31:BA:2631:G:N3 | 31:BA:2810:A:H2 | 2.02 | 0.58 |
| 34:BE:33:VAL:CG1 | 34:BE:90:THR:H | 2.16 | 0.58 |
| 38:BI:9:LEU:HB2 | 38:BI:12:LEU:O | 2.03 | 0.58 |
| 38:BI:67:ARG:O | 38:BI:68:LEU:HB2 | 2.03 | 0.58 |
| 41:BP:61:ARG:H | 41:BP:61:ARG:HD2 | 1.68 | 0.58 |
| 42:BQ:18:LYS:O | 42:BQ:19:GLY:C | 2.46 | 0.58 |
| 45:BT:109:GLU:HA | 45:BT:112:ARG:CG | 2.33 | 0.58 |
| 51:BZ:150:LEU:HD23 | 51:BZ:171:ILE:HD11 | 1.85 | 0.58 |
| 1:CA:376:G:H2' | 1:CA:377:G:H8 | 1.68 | 0.58 |
| 1:CA:542:G:P | 4:CD:10:ARG:HH21 | 2.24 | 0.58 |
| 1:CA:672:U:H2' | 1:CA:672:U:O2 | 2.02 | 0.58 |
| 3:CC:29:TYR:O | 3:CC:33:LEU:HB2 | 2.02 | 0.58 |
| 7:CG:79:ARG:NE | 7:CG:84:ASN:HD21 | 2.01 | 0.58 |
| 10:CJ:32:ALA:HB2 | 10:CJ:76:ASN:HB3 | 1.85 | 0.58 |
| 25:D3:39:ASP:OD1 | 25:D3:44:ARG:HG3 | 2.02 | 0.58 |
| 26:D4:25:TYR:C | 26:D4:27:THR:H | 2.11 | 0.58 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 30:D8:35:GLN:HE21 | 30:D8:36:LYS:HG3 | 1.66 | 0.58 |
| 31:DA:541:C:H2' | 31:DA:542:C:C6 | 2.38 | 0.58 |
| 31:DA:1024:G:C3' | 31:DA:1025:G:H5" | 2.29 | 0.58 |
| 32:DB:46:A:C6 | 32:DB:47:C:C4 | 2.91 | 0.58 |
| 38:DI:78:THR:HA | 38:DI:141:LYS:O | 2.03 | 0.58 |
| 39:DN:68:GLU:HA | 39:DN:86:PRO:HB3 | 1.83 | 0.58 |
| 40:DO:65:THR:HG23 | 40:DO:69:ILE:HD11 | 1.83 | 0.58 |
| 46:DU:76:TYR:C | 46:DU:76:TYR:CD2 | 2.81 | 0.58 |
| 1:AA:632:A:C8 | 1:AA:633:G:C8 | 2.91 | 0.58 |
| 4:AD:14:ARG:HA | 4:AD:39:PRO:HB3 | 1.84 | 0.58 |
| 12:AL:119:LYS:HB2 | 12:AL:120:TYR:HD1 | 1.69 | 0.58 |
| 17:AQ:22:LEU:HD12 | 17:AQ:23:VAL:H | 1.67 | 0.58 |
| 22:B0:1:MET:HA | 31:BA:2451:A:H4' | 1.85 | 0.58 |
| 24:B2:49:LYS:O | 24:B2:50:ILE:C | 2.46 | 0.58 |
| 31:BA:314:A:C2' | 31:BA:315:G:H5' | 2.34 | 0.58 |
| 31:BA:1022:G:N2 | 31:BA:1142(A):A:C2 | 2.72 | 0.58 |
| 31:BA:1568:G:H21 | 33:BD:58:HIS:HE1 | 1.52 | 0.58 |
| 31:BA:2334:G:H4' | 31:BA:2335:A:OP2 | 2.04 | 0.58 |
| 31:BA:2476:A:H2 | 31:BA:2477:C:H5" | 1.66 | 0.58 |
| 31:BA:2577:A:H5" | 31:BA:2578:G:C5' | 2.31 | 0.58 |
| 31:BA:2636:U:O2' | 31:BA:2637:U:H5" | 2.03 | 0.58 |
| 34:BE:75:VAL:O | 34:BE:77:ILE:N | 2.36 | 0.58 |
| 37:BH:84:SER:O | 37:BH:85:LYS:HB3 | 2.02 | 0.58 |
| 39:BN:42:TRP:HB3 | 46:BU:64:ARG:CZ | 2.32 | 0.58 |
| 43:BR:8:ARG:HA | 43:BR:8:ARG:NE | 2.12 | 0.58 |
| 44:BS:56:LEU:C | 44:BS:56:LEU:HD23 | 2.27 | 0.58 |
| 49:BX:41:ASN:HA | 49:BX:44:GLU:HG2 | 1.85 | 0.58 |
| 50:BY:8:LYS:HD2 | 50:BY:8:LYS:N | 2.19 | 0.58 |
| 1:CA:16:A:O2' | 1:CA:17:U:H5' | 2.03 | 0.58 |
| 1:CA:262:A:H2' | 1:CA:263:A:C8 | 2.37 | 0.58 |
| 1:CA:1366:C:OP1 | 9:CI:117:HIS:CE1 | 2.56 | 0.58 |
| 1:CA:1392:G:C2' | 1:CA:1393:U:H5' | 2.33 | 0.58 |
| 23:D1:37:ILE:HG21 | 31:DA:2080:G:P | 2.43 | 0.58 |
| 31:DA:61:G:H1 | 31:DA:94:C:N4 | 2.01 | 0.58 |
| 31:DA:727:A:C2 | 33:DD:9:TYR:CD2 | 2.91 | 0.58 |
| 31:DA:2228:G:C5 | 31:DA:2229:C:C4 | 2.91 | 0.58 |
| 31:DA:2281:C:O2' | 31:DA:2282:G:H5' | 2.03 | 0.58 |
| 31:DA:2327:A:H2' | 31:DA:2328:A:H8 | 1.66 | 0.58 |
| 31:DA:2443:C:O2' | 31:DA:2444:G:H5' | 2.02 | 0.58 |
| 32:DB:15:A:H1' | 32:DB:110:G:N9 | 2.18 | 0.58 |
| 46:DU:74:LEU:HD12 | 46:DU:74:LEU:N | 2.18 | 0.58 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 46:DU:91:ASP:OD2 | 46:DU:96:ALA:HB2 | 2.03 | 0.58 |
| 50:DY:29:GLU:OE1 | 50:DY:29:GLU:N | 2.36 | 0.58 |
| 1:AA:80:G:H1 | 1:AA:89:C:H41 | 1.51 | 0.58 |
| 1:AA:299:G:C6 | 1:AA:300:A:N1 | 2.71 | 0.58 |
| 1:AA:477:A:O2' | 1:AA:479:C:H5' | 2.02 | 0.58 |
| 1:AA:542:G:P | 4:AD:10:ARG:HH21 | 2.26 | 0.58 |
| 1:AA:601:C:H2' | 1:AA:602:A:C8 | 2.38 | 0.58 |
| 1:AA:626:U:C2 | 1:AA:627:G:C8 | 2.91 | 0.58 |
| 1:AA:1423:G:C5' | 40:BO:49:ARG:NH2 | 2.67 | 0.58 |
| 27:B5:51:TYR:HB3 | 27:B5:52:TYR:CD2 | 2.38 | 0.58 |
| 29:B7:15:THR:HG22 | 29:B7:16:HIS:N | 2.17 | 0.58 |
| 30:B8:2:PRO:N | 31:BA:591:C:H1' | 2.19 | 0.58 |
| 31:BA:267:C:H2' | 31:BA:268:C:C6 | 2.38 | 0.58 |
| 31:BA:535:C:O3' | 46:BU:53:ARG:NH1 | 2.36 | 0.58 |
| 31:BA:760:G:H2' | 31:BA:761:A:O4' | 2.02 | 0.58 |
| 31:BA:992:C:C2' | 31:BA:993:G:O5' | 2.51 | 0.58 |
| 31:BA:1161:C:H1' | 47:BV:8:GLY:O | 2.04 | 0.58 |
| 31:BA:1501:C:H2' | 31:BA:1502:C:H6 | 1.68 | 0.58 |
| 31:BA:1558:A:H1' | 31:BA:1559:G:OP2 | 2.03 | 0.58 |
| 31:BA:2636:U:O2' | 31:BA:2637:U:C5' | 2.51 | 0.58 |
| 34:BE:103:ASP:OD2 | 34:BE:168:MET:HE2 | 2.02 | 0.58 |
| 37:BH:17:VAL:HG21 | 37:BH:50:VAL:HG21 | 1.85 | 0.58 |
| 41:BP:140:ALA:O | 41:BP:141:ALA:HB2 | 2.03 | 0.58 |
| 42:BQ:30:GLY:CA | 42:BQ:107:ALA:HB2 | 2.31 | 0.58 |
| 43:BR:34:ILE:HG22 | 43:BR:114:VAL:HB | 1.84 | 0.58 |
| 43:BR:103:ARG:HD2 | 43:BR:108:GLY:O | 2.03 | 0.58 |
| 44:BS:42:ASP:O | 44:BS:43:GLU:HB2 | 2.02 | 0.58 |
| 44:BS:95:HIS:ND1 | 44:BS:96:GLY:N | 2.52 | 0.58 |
| 49:BX:72:LYS:C | 49:BX:74:PRO:HD3 | 2.28 | 0.58 |
| 1:CA:370:C:H2' | 1:CA:371:G:O4' | 2.02 | 0.58 |
| 1:CA:577:G:C8 | 1:CA:816:A:C6 | 2.91 | 0.58 |
| 6:CF:53:ALA:O | 6:CF:54:LYS:HB2 | 2.02 | 0.58 |
| 10:CJ:34:VAL:HG13 | 10:CJ:73:ASP:O | 2.03 | 0.58 |
| 30:D8:62:LEU:CD1 | 31:DA:242:G:H5" | 2.29 | 0.58 |
| 31:DA:836:G:C5 | 31:DA:837:C:C4 | 2.91 | 0.58 |
| 31:DA:1030:G:OP2 | 42:DQ:128:LYS:HE2 | 2.03 | 0.58 |
| 31:DA:1257:C:H4' | 35:DF:83:PHE:CE2 | 2.38 | 0.58 |
| 31:DA:1671:U:HO2' | 31:DA:1673:U:H5 | 1.50 | 0.58 |
| 31:DA:1686:C:H2' | 31:DA:1687:G:H5' | 1.84 | 0.58 |
| 33:DD:102:LYS:C | 33:DD:103:ARG:HG2 | 2.29 | 0.58 |
| 38:DI:110:ASP:C | 38:DI:112:LYS:N | 2.60 | 0.58 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 39:DN:16:ILE:O | 39:DN:54:VAL:HA | 2.02 | 0.58 |
| 39:DN:68:GLU:HG3 | 39:DN:88:GLU:OE1 | 2.02 | 0.58 |
| 46:DU:25:TRP:CD1 | 46:DU:26:GLY:H | 2.20 | 0.58 |
| 1:AA:60:A:P | 1:AA:60:A:C8 | 2.95 | 0.58 |
| 1:AA:539:A:OP2 | 12:AL:115:LYS:HE3 | 2.03 | 0.58 |
| 1:AA:1012:U:O5' | 1:AA:1012:U:H6 | 1.87 | 0.58 |
| 1:AA:1080:A:H5' | 5:AE:14:ARG:NH2 | 2.19 | 0.58 |
| 1:AA:1218:C:H2' | 1:AA:1219:U:C6 | 2.37 | 0.58 |
| 1:AA:1399:C:C2 | 1:AA:1502:A:N6 | 2.71 | 0.58 |
| 12:AL:6:THR:HG23 | 12:AL:9:GLN:HE21 | 1.69 | 0.58 |
| 23:B1:37:ILE:HG21 | 31:BA:2080:G:P | 2.43 | 0.58 |
| 30:B8:61:LEU:HD22 | 31:BA:593:G:O3' | 2.04 | 0.58 |
| 31:BA:61:G:H1 | 31:BA:94:C:N4 | 2.00 | 0.58 |
| 31:BA:329:G:H4' | 31:BA:330:A:OP2 | 2.02 | 0.58 |
| 31:BA:737:C:C2' | 31:BA:738:G:O5' | 2.52 | 0.58 |
| 31:BA:1264:G:H3' | 31:BA:1265:A:H5" | 1.85 | 0.58 |
| 31:BA:1832:C:N4 | 31:BA:1833:U:C4 | 2.71 | 0.58 |
| 31:BA:2884:U:C6 | 31:BA:2885:C:C6 | 2.91 | 0.58 |
| 33:BD:231:HIS:CD2 | 33:BD:249:PRO:HA | 2.38 | 0.58 |
| 39:BN:40:PRO:CA | 46:BU:64:ARG:NH2 | 2.60 | 0.58 |
| 47:BV:80:GLN:OE1 | 47:BV:80:GLN:C | 2.46 | 0.58 |
| 47:BV:85:LYS:C | 47:BV:87:HIS:N | 2.62 | 0.58 |
| 1:CA:724:G:C2 | 1:CA:725:G:C8 | 2.92 | 0.58 |
| 1:CA:779:C:C2' | 1:CA:780:A:H5' | 2.34 | 0.58 |
| 1:CA:1065:U:C1' | 1:CA:1066:C:OP2 | 2.50 | 0.58 |
| 2:CB:24:TRP:CG | 2:CB:25:ASN:N | 2.69 | 0.58 |
| 5:CE:12:LEU:O | 5:CE:13:ILE:HD12 | 2.02 | 0.58 |
| 12:CL:6:THR:OG1 | 12:CL:9:GLN:HG3 | 2.03 | 0.58 |
| 31:DA:717:G:H2' | 31:DA:718:A:O4' | 2.04 | 0.58 |
| 31:DA:819:A:N3 | 31:DA:1189:A:C2 | 2.71 | 0.58 |
| 31:DA:1528(A):A:C5 | 31:DA:1529:G:C8 | 2.90 | 0.58 |
| 31:DA:2862:G:H2' | 31:DA:2863:C:H6 | 1.69 | 0.58 |
| 32:DB:43:C:H4' | 36:DG:66:GLN:NE2 | 2.18 | 0.58 |
| 33:DD:143:HIS:HD2 | 33:DD:144:ALA:HB2 | 1.69 | 0.58 |
| 41:DP:30:THR:HG22 | 41:DP:31:ALA:H | 1.68 | 0.58 |
| 51:DZ:145:GLU:HG3 | 51:DZ:146:ILE:H | 1.67 | 0.58 |
| 1:AA:339:C:OP2 | 40:BO:97:ARG:NH1 | 2.36 | 0.58 |
| 1:AA:687:A:H1' | 1:AA:688:G:OP2 | 2.04 | 0.58 |
| 3:AC:23:TYR:CG | 3:AC:24:ALA:N | 2.72 | 0.58 |
| 6:AF:5:GLU:HG3 | 6:AF:93:SER:OG | 2.02 | 0.58 |
| 13:AM:66:LEU:HD12 | 13:AM:66:LEU:N | 2.18 | 0.58 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 23:B1:37:ILE:HD12 | 31:BA:2079:U:O2' | 2.03 | 0.58 |
| 28:B6:19:ARG:O | 28:B6:20:ASN:O | 2.21 | 0.58 |
| 31:BA:528:A:C2 | 31:BA:2042:A:H2' | 2.39 | 0.58 |
| 31:BA:626:U:C2 | 41:BP:105:LEU:HG | 2.39 | 0.58 |
| 31:BA:1050:A:C2 | 31:BA:2751:G:C4 | 2.90 | 0.58 |
| 31:BA:1204:A:N1 | 31:BA:1241:A:C2 | 2.71 | 0.58 |
| 31:BA:1786:A:H2 | 31:BA:2606:C:H1' | 1.67 | 0.58 |
| 31:BA:1839:G:N7 | 31:BA:1927:A:H1' | 2.18 | 0.58 |
| 31:BA:2280:G:C2' | 31:BA:2281:C:H5' | 2.34 | 0.58 |
| 31:BA:2873:A:C2 | 43:BR:6:SER:HB2 | 2.39 | 0.58 |
| 37:BH:146:ALA:O | 37:BH:147:ASN:C | 2.46 | 0.58 |
| 38:BI:81:VAL:HG11 | 38:BI:123:LEU:HD21 | 1.85 | 0.58 |
| 40:BO:65:THR:HG23 | 40:BO:69:ILE:HD11 | 1.85 | 0.58 |
| 41:BP:16:ARG:HD3 | 41:BP:18:ARG:HB2 | 1.84 | 0.58 |
| 43:BR:53:HIS:CD2 | 43:BR:94:TYR:OH | 2.54 | 0.58 |
| 44:BS:28:VAL:C | 44:BS:89:ARG:HD2 | 2.28 | 0.58 |
| 49:BX:60:ARG:HH21 | 49:BX:74:PRO:HG2 | 1.68 | 0.58 |
| 49:BX:85:PRO:O | 49:BX:87:GLN:N | 2.37 | 0.58 |
| 1:CA:233:C:H2' | 1:CA:234:C:H6 | 1.69 | 0.58 |
| 1:CA:502:G:C6 | 1:CA:503:C:N3 | 2.71 | 0.58 |
| 1:CA:1301:U:H3' | 1:CA:1302:U:H5" | 1.85 | 0.58 |
| 3:CC:86:VAL:O | 3:CC:90:GLU:HG2 | 2.03 | 0.58 |
| 5:CE:55:VAL:O | 5:CE:58:ALA:HB3 | 2.04 | 0.58 |
| 29:D7:11:LYS:HE2 | 31:DA:686:G:H5" | 1.86 | 0.58 |
| 30:D8:61:LEU:N | 30:D8:63:PRO:HD2 | 2.19 | 0.58 |
| 31:DA:511:U:H3' | 31:DA:512:G:C5' | 2.20 | 0.58 |
| 31:DA:535:C:O3' | 46:DU:53:ARG:NH1 | 2.37 | 0.58 |
| 31:DA:1265:A:OP1 | 31:DA:1265:A:H8 | 1.87 | 0.58 |
| 31:DA:1268:A:C2 | 31:DA:2013:A:C4 | 2.91 | 0.58 |
| 31:DA:1386:C:H2' | 31:DA:1387:C:C6 | 2.39 | 0.58 |
| 39:DN:30:ILE:HG22 | 39:DN:34:LEU:CD2 | 2.33 | 0.58 |
| 50:DY:27:VAL:C | 50:DY:29:GLU:OE1 | 2.46 | 0.58 |
| 1:AA:892:A:C6 | 1:AA:893:C:C4 | 2.92 | 0.58 |
| 1:AA:977:A:H2' | 1:AA:978:A:H5' | 1.86 | 0.58 |
| 1:AA:1065:U:C1' | 1:AA:1066:C:OP2 | 2.50 | 0.58 |
| 3:AC:113:ALA:HB3 | 3:AC:114:PRO:HD3 | 1.85 | 0.58 |
| 4:AD:20:TYR:CD2 | 4:AD:26:CYS:HB3 | 2.38 | 0.58 |
| 8:AH:104:ARG:O | 8:AH:105:ARG:HB2 | 2.04 | 0.58 |
| 12:AL:21:LYS:N | 12:AL:21:LYS:HD2 | 2.18 | 0.58 |
| 12:AL:70:ILE:HG12 | 12:AL:100:ILE:HD12 | 1.86 | 0.58 |
| 22:B0:74:ARG:HG2 | 32:BB:12:C:O2' | 2.04 | 0.58 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 27:B5:40:LYS:HZ2 | 27:B5:46:CYS:H | 1.52 | 0.58 |
| 31:BA:1509(A):A:H2' | 31:BA:1509(B):A:H8 | 1.68 | 0.58 |
| 31:BA:1882:C:H5' | 31:BA:1883:G:OP2 | 2.03 | 0.58 |
| 31:BA:1925:C:C2' | 31:BA:1926:U:H5' | 2.34 | 0.58 |
| 31:BA:2713:A:H3' | 31:BA:2714:G:C5' | 2.33 | 0.58 |
| 33:BD:186:HIS:HD2 | 33:BD:188:GLU:N | 2.00 | 0.58 |
| 41:BP:24:GLY:HA2 | 41:BP:33:ARG:HE | 1.69 | 0.58 |
| 43:BR:24:GLN:HE22 | 43:BR:36:THR:CG2 | 2.12 | 0.58 |
| 1:CA:1003:G:H2' | 1:CA:1004:A:C4' | 2.32 | 0.58 |
| 1:CA:1216:G:OP1 | 14:CN:2:ALA:HA | 2.04 | 0.58 |
| 4:CD:100:ARG:NH2 | 4:CD:118:ARG:HH22 | 2.01 | 0.58 |
| 7:CG:23:VAL:O | 7:CG:27:ILE:HD12 | 2.04 | 0.58 |
| 13:CM:24:GLY:C | 13:CM:25:ILE:HD12 | 2.28 | 0.58 |
| 27:D5:57:VAL:C | 27:D5:58:LEU:HG | 2.28 | 0.58 |
| 31:DA:542:C:C2' | 31:DA:543:C:OP1 | 2.50 | 0.58 |
| 31:DA:1784:A:H4' | 31:DA:1785:A:H5" | 1.84 | 0.58 |
| 33:DD:25:THR:O | 33:DD:26:LYS:C | 2.46 | 0.58 |
| 34:DE:111:ARG:CZ | 43:DR:2:ARG:HH21 | 2.17 | 0.58 |
| 45:DT:10:VAL:O | 45:DT:13:ARG:HG2 | 2.02 | 0.58 |
| 46:DU:65:ILE:HG12 | 46:DU:96:ALA:HB3 | 1.85 | 0.58 |
| 49:DX:18:TYR:O | 49:DX:20:GLY:N | 2.37 | 0.58 |
| 51:DZ:152:ALA:HB2 | 51:DZ:168:GLU:HA | 1.84 | 0.58 |
| 1:AA:375:U:O3' | 16:AP:6:LEU:HB2 | 2.03 | 0.58 |
| 1:AA:414:A:H2' | 1:AA:415:A:H8 | 1.68 | 0.58 |
| 12:AL:22:SER:O | 12:AL:24:VAL:N | 2.37 | 0.58 |
| 13:AM:15:VAL:O | 13:AM:19:LEU:HD23 | 2.04 | 0.58 |
| 20:AT:89:ARG:HH21 | 20:AT:104:LEU:HD21 | 1.66 | 0.58 |
| 30:B8:31:HIS:O | 30:B8:32:LEU:C | 2.46 | 0.58 |
| 31:BA:176:G:O2' | 31:BA:177:G:H5' | 2.04 | 0.58 |
| 31:BA:570:G:H2' | 31:BA:2030:A:C6 | 2.38 | 0.58 |
| 31:BA:866:A:C6 | 31:BA:914:C:C6 | 2.92 | 0.58 |
| 31:BA:1047:G:N2 | 31:BA:1111:A:N6 | 2.52 | 0.58 |
| 31:BA:1528:A:O2' | 31:BA:1528(A):A:O5' | 2.22 | 0.58 |
| 32:BB:15:A:H1' | 32:BB:110:G:N9 | 2.18 | 0.58 |
| 33:BD:87:ASN:N | 33:BD:87:ASN:ND2 | 2.44 | 0.58 |
| 34:BE:136:ARG:HH11 | 34:BE:136:ARG:CG | 2.17 | 0.58 |
| 36:BG:16:ARG:HE | 36:BG:31:VAL:HG11 | 1.68 | 0.58 |
| 40:BO:4:PRO:O | 40:BO:5:GLN:HB2 | 2.02 | 0.58 |
| 40:BO:23:ARG:HH11 | 40:BO:23:ARG:CG | 1.99 | 0.58 |
| 44:BS:29:PHE:H | 44:BS:89:ARG:HD2 | 1.67 | 0.58 |
| 45:BT:56:GLY:O | 45:BT:59:THR:HG22 | 2.01 | 0.58 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:155:C:H2' | 1:CA:156:G:C8 | 2.38 | 0.58 |
| 1:CA:661:G:C2 | 1:CA:662:G:C8 | 2.92 | 0.58 |
| 1:CA:1227:A:OP2 | 13:CM:111:LYS:HE2 | 2.04 | 0.58 |
| 5:CE:96:PRO:HA | 5:CE:117:ASP:OD2 | 2.02 | 0.58 |
| 31:DA:954:G:C5 | 31:DA:955:C:C5 | 2.92 | 0.58 |
| 31:DA:1279:G:H5' | 43:DR:34:ILE:HD11 | 1.85 | 0.58 |
| 31:DA:1603:A:H5' | 31:DA:1603:A:C8 | 2.36 | 0.58 |
| 31:DA:1697:G:H8 | 31:DA:1697:G:C5' | 2.07 | 0.58 |
| 31:DA:1818:U:H2' | 33:DD:157:ARG:HG3 | 1.84 | 0.58 |
| 31:DA:2022:U:O2' | 31:DA:2617:C:H5' | 2.03 | 0.58 |
| 33:DD:24:ILE:CG2 | 33:DD:24:ILE:O | 2.52 | 0.58 |
| 33:DD:158:ALA:O | 33:DD:159:ALA:CB | 2.50 | 0.58 |
| 33:DD:267:SER:O | 33:DD:268:ARG:HB2 | 2.02 | 0.58 |
| 36:DG:108:ASN:O | 36:DG:112:PRO:HG2 | 2.04 | 0.58 |
| 37:DH:29:PRO:HD2 | 37:DH:79:VAL:O | 2.04 | 0.58 |
| 39:DN:42:TRP:HB3 | 46:DU:64:ARG:CZ | 2.32 | 0.58 |
| 43:DR:72:ASP:HB3 | 43:DR:75:LEU:HB2 | 1.85 | 0.58 |
| 1:AA:109:A:H2' | 1:AA:326:G:N2 | 2.18 | 0.58 |
| 1:AA:509:A:H4' | 1:AA:510:A:OP1 | 2.02 | 0.58 |
| 1:AA:617:G:N1 | 1:AA:618:C:C4 | 2.72 | 0.58 |
| 1:AA:779:C:O2' | 1:AA:780:A:H5' | 2.03 | 0.58 |
| 1:AA:1003:G:H2' | 1:AA:1004:A:C4' | 2.32 | 0.58 |
| 1:AA:1490:C:O2' | 1:AA:1491:G:H5' | 2.04 | 0.58 |
| 3:AC:34:LEU:O | 3:AC:34:LEU:HD23 | 2.04 | 0.58 |
| 4:AD:22:LYS:O | 4:AD:113:SER:HB3 | 2.02 | 0.58 |
| 7:AG:32:ARG:O | 7:AG:33:ASP:CB | 2.52 | 0.58 |
| 9:AI:83:ARG:O | 9:AI:86:VAL:HG12 | 2.04 | 0.58 |
| 10:AJ:70:ARG:HG3 | 10:AJ:70:ARG:HH11 | 1.68 | 0.58 |
| 18:AR:50:ILE:HD12 | 18:AR:70:ILE:HD12 | 1.84 | 0.58 |
| 23:B1:41:ARG:NH2 | 31:BA:205:G:O6 | 2.35 | 0.58 |
| 24:B2:25:VAL:HG13 | 24:B2:26:ARG:HD3 | 1.85 | 0.58 |
| 30:B8:35:GLN:HE21 | 30:B8:36:LYS:HG3 | 1.68 | 0.58 |
| 31:BA:784:A:H5' | 31:BA:785:G:OP1 | 2.04 | 0.58 |
| 31:BA:966:G:C4 | 31:BA:967:C:C5 | 2.92 | 0.58 |
| 31:BA:1315:C:H42 | 31:BA:1337:G:H1 | 1.51 | 0.58 |
| 31:BA:1784:A:H4' | 31:BA:1785:A:H5" | 1.84 | 0.58 |
| 31:BA:2191:G:H2' | 31:BA:2192:G:O5' | 2.03 | 0.58 |
| 39:BN:77:GLY:O | 39:BN:78:TYR:HB3 | 2.04 | 0.58 |
| 42:BQ:75:THR:HG22 | 42:BQ:88:GLY:HA3 | 1.85 | 0.58 |
| 47:BV:19:LYS:CE | 47:BV:20:LEU:H | 2.16 | 0.58 |
| 47:BV:25:LEU:C | 47:BV:27:ALA:H | 2.10 | 0.58 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 51:BZ:61:LEU:HB2 | 51:BZ:65:GLN:HB2 | 1.86 | 0.58 |
| 51:BZ:119:GLU:C | 51:BZ:121:HIS:H | 2.12 | 0.58 |
| 1:CA:22:G:H2' | 1:CA:23:C:C6 | 2.38 | 0.58 |
| 1:CA:417:C:O2' | 1:CA:418:C:H5' | 2.02 | 0.58 |
| 1:CA:484:G:H4' | 1:CA:485:G:OP1 | 2.04 | 0.58 |
| 1:CA:1346:A:N1 | 1:CA:1374:A:H5" | 2.18 | 0.58 |
| 2:CB:185:ILE:HG22 | 2:CB:199:TYR:CB | 2.21 | 0.58 |
| 3:CC:6:HIS:HE2 | 3:CC:184:TYR:HE2 | 1.51 | 0.58 |
| 5:CE:10:MET:HG3 | 5:CE:13:ILE:HD13 | 1.85 | 0.58 |
| 16:CP:64:ALA:O | 16:CP:65:GLN:C | 2.47 | 0.58 |
| 27:D5:57:VAL:O | 27:D5:58:LEU:HG | 2.03 | 0.58 |
| 30:D8:35:GLN:CG | 31:DA:2420:C:OP1 | 2.52 | 0.58 |
| 31:DA:84:A:C5' | 50:DY:9:LYS:HD2 | 2.34 | 0.58 |
| 31:DA:626:U:H5" | 31:DA:627:A:H5' | 1.83 | 0.58 |
| 31:DA:1158:C:O2' | 31:DA:1159:U:H5' | 2.04 | 0.58 |
| 31:DA:1204:A:C2 | 31:DA:1241:A:N1 | 2.72 | 0.58 |
| 31:DA:1582:C:O2' | 31:DA:1586:A:C8 | 2.56 | 0.58 |
| 31:DA:1689:A:H62 | 31:DA:1698:A:H2 | 1.52 | 0.58 |
| 31:DA:1882:C:H5' | 31:DA:1883:G:OP2 | 2.04 | 0.58 |
| 31:DA:2196:C:O2' | 31:DA:2197:U:H5' | 2.03 | 0.58 |
| 31:DA:2631:G:N3 | 31:DA:2810:A:H2 | 2.02 | 0.58 |
| 31:DA:2810:A:H2' | 34:DE:61:ARG:HH21 | 1.66 | 0.58 |
| 33:DD:173:VAL:HG23 | 33:DD:174:ILE:N | 2.16 | 0.58 |
| 34:DE:47:VAL:HG12 | 34:DE:49:LEU:HD22 | 1.85 | 0.58 |
| 36:DG:174:GLU:HG3 | 36:DG:180:PHE:HD1 | 1.68 | 0.58 |
| 41:DP:16:ARG:HD3 | 41:DP:18:ARG:N | 2.07 | 0.58 |
| 41:DP:16:ARG:HH11 | 41:DP:16:ARG:C | 2.11 | 0.58 |
| 42:DQ:20:ALA:C | 42:DQ:22:LYS:H | 2.12 | 0.58 |
| 1:AA:354:G:C6 | 1:AA:355:C:N4 | 2.72 | 0.58 |
| 3:AC:86:VAL:O | 3:AC:90:GLU:HG2 | 2.04 | 0.58 |
| 4:AD:100:ARG:NH2 | 4:AD:118:ARG:HH22 | 2.02 | 0.58 |
| 8:AH:10:LEU:N | 8:AH:10:LEU:CD2 | 2.67 | 0.58 |
| 8:AH:31:PHE:O | 8:AH:35:ILE:HG13 | 2.03 | 0.58 |
| 25:B3:19:GLN:HE22 | 25:B3:52:HIS:CE1 | 2.20 | 0.58 |
| 27:B5:2:ALA:CA | 31:BA:2015:A:H1' | 2.29 | 0.58 |
| 30:B8:14:VAL:CG1 | 30:B8:22:VAL:HG13 | 2.33 | 0.58 |
| 31:BA:642:G:H21 | 31:BA:646:A:H2 | 1.50 | 0.58 |
| 31:BA:866:A:C6 | 31:BA:914:C:C5 | 2.92 | 0.58 |
| 31:BA:1767:C:O2' | 31:BA:1768:U:H5' | 2.04 | 0.58 |
| 31:BA:2023:G:H5' | 31:BA:2617:C:H4' | 1.85 | 0.58 |
| 32:BB:37:C:C5 | 32:BB:38:C:C5 | 2.92 | 0.58 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 35:BF:160:ASN:HD22 | 35:BF:162:LEU:H | 1.51 | 0.58 |
| 38:BI:133:HIS:CG | 38:BI:134:PRO:HD2 | 2.38 | 0.58 |
| 42:BQ:20:ALA:C | 42:BQ:22:LYS:H | 2.11 | 0.58 |
| 47:BV:62:LEU:HB3 | 47:BV:98:GLU:HB2 | 1.86 | 0.58 |
| 49:BX:33:LYS:HA | 49:BX:35:THR:HG22 | 1.85 | 0.58 |
| 1:CA:1016:A:H2' | 1:CA:1017:G:O4' | 2.04 | 0.58 |
| 1:CA:1090:U:C2 | 1:CA:1091:U:C5 | 2.92 | 0.58 |
| 1:CA:1306:A:H1' | 1:CA:1332:A:C2 | 2.39 | 0.58 |
| 4:CD:30:LYS:HA | 4:CD:35:ARG:HD2 | 1.86 | 0.58 |
| 13:CM:25:ILE:CD1 | 13:CM:66:LEU:HD23 | 2.33 | 0.58 |
| 19:CS:24:ALA:O | 19:CS:25:LYS:HB2 | 2.02 | 0.58 |
| 20:CT:32:ALA:O | 20:CT:36:LEU:HD23 | 2.04 | 0.58 |
| 20:CT:97:ALA:O | 20:CT:99:LEU:HG | 2.04 | 0.58 |
| 22:D0:43:THR:O | 22:D0:45:PHE:N | 2.37 | 0.58 |
| 23:D1:65:SER:N | 23:D1:67:ILE:HD11 | 2.18 | 0.58 |
| 31:DA:810:U:O2' | 41:DP:33:ARG:CZ | 2.51 | 0.58 |
| 31:DA:1316:U:O2' | 31:DA:1317:A:H5' | 2.03 | 0.58 |
| 31:DA:2206:G:H3' | 31:DA:2206:G:N3 | 2.19 | 0.58 |
| 35:DF:170:LEU:HD23 | 35:DF:172:TRP:NE1 | 2.19 | 0.58 |
| 39:DN:3:THR:HG22 | 39:DN:4:TYR:N | 2.13 | 0.58 |
| 44:DS:29:PHE:N | 44:DS:89:ARG:CD | 2.65 | 0.58 |
| 44:DS:29:PHE:H | 44:DS:89:ARG:CD | 2.17 | 0.58 |
| 49:DX:38:GLU:N | 49:DX:38:GLU:OE1 | 2.37 | 0.58 |
| 49:DX:65:ARG:O | 49:DX:66:LEU:HB2 | 2.03 | 0.58 |
| 51:DZ:166:SER:HB2 | 51:DZ:168:GLU:H | 1.69 | 0.58 |
| 1:AA:417:C:O2' | 1:AA:418:C:H5' | 2.03 | 0.58 |
| 1:AA:539:A:H2' | 1:AA:540:G:C8 | 2.39 | 0.58 |
| 1:AA:579:G:H2' | 1:AA:580:U:C6 | 2.39 | 0.58 |
| 1:AA:1049:U:H4' | 1:AA:1050:G:O5' | 2.04 | 0.58 |
| 1:AA:1306:A:H1' | 1:AA:1332:A:C2 | 2.39 | 0.58 |
| 1:AA:1366:C:OP1 | 9:AI:117:HIS:CE1 | 2.56 | 0.58 |
| 18:AR:66:LEU:O | 18:AR:70:ILE:HG13 | 2.04 | 0.58 |
| 22:B0:72:ARG:O | 22:B0:73:GLY:C | 2.45 | 0.58 |
| 30:B8:32:LEU:HB3 | 30:B8:35:GLN:N | 2.19 | 0.58 |
| 30:B8:35:GLN:CB | 31:BA:2420:C:OP1 | 2.52 | 0.58 |
| 31:BA:1406:U:H2' | 31:BA:1407:C:H6 | 1.66 | 0.58 |
| 31:BA:1495:A:C4 | 31:BA:1496:A:C2 | 2.92 | 0.58 |
| 31:BA:1742:G:H3' | 31:BA:1742:G:H8 | 1.69 | 0.58 |
| 31:BA:1991:U:H2' | 31:BA:1992:G:H5" | 1.86 | 0.58 |
| 31:BA:2469:A:H2 | 31:BA:2481:G:N2 | 1.96 | 0.58 |
| 34:BE:59:VAL:CG2 | 34:BE:63:LEU:HA | 2.34 | 0.58 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 49:BX:34:ALA:O | 49:BX:36:LYS:HE3 | 2.04 | 0.58 |
| 1:CA:411:A:C4 | 1:CA:413:G:O4' | 2.57 | 0.58 |
| 1:CA:539:A:OP2 | 12:CL:115:LYS:HE3 | 2.04 | 0.58 |
| 1:CA:706:A:N7 | 1:CA:707:C:H5 | 2.01 | 0.58 |
| 1:CA:1162:C:H2' | 1:CA:1163:C:H6 | 1.69 | 0.58 |
| 2:CB:71:VAL:CG1 | 2:CB:93:VAL:HB | 2.34 | 0.58 |
| 6:CF:7:ASN:O | 6:CF:88:VAL:HA | 2.04 | 0.58 |
| 10:CJ:90:LEU:N | 10:CJ:91:PRO:HD3 | 2.19 | 0.58 |
| 14:CN:4:LYS:O | 14:CN:7:ILE:HG12 | 2.04 | 0.58 |
| 29:D7:8:ASN:ND2 | 29:D7:11:LYS:HB3 | 2.19 | 0.58 |
| 31:DA:547:A:C8 | 31:DA:549:G:O6 | 2.57 | 0.58 |
| 31:DA:2280:G:H2' | 31:DA:2281:C:H5' | 1.85 | 0.58 |
| 31:DA:2652:C:O2' | 31:DA:2653:U:H5' | 2.03 | 0.58 |
| 31:DA:2850:A:OP2 | 31:DA:2866:U:C5 | 2.56 | 0.58 |
| 32:DB:46:A:C5 | 32:DB:47:C:C5 | 2.92 | 0.58 |
| 36:DG:133:LEU:C | 36:DG:133:LEU:HD12 | 2.29 | 0.58 |
| 39:DN:40:PRO:C | 46:DU:64:ARG:NH2 | 2.62 | 0.58 |
| 41:DP:105:LEU:O | 41:DP:106:LEU:CB | 2.52 | 0.58 |
| 44:DS:28:VAL:C | 44:DS:89:ARG:HD2 | 2.28 | 0.58 |
| 44:DS:29:PHE:CD2 | 44:DS:29:PHE:C | 2.82 | 0.58 |
| 46:DU:83:LEU:CB | 46:DU:88:ILE:HD11 | 2.28 | 0.58 |
| 49:DX:55:ASN:HB2 | 49:DX:78:LYS:CD | 2.33 | 0.58 |
| 1:AA:147:G:C2' | 1:AA:148:G:H5' | 2.34 | 0.57 |
| 1:AA:240:C:H2' | 1:AA:241:C:H6 | 1.69 | 0.57 |
| 1:AA:560:U:H4' | 1:AA:561:U:O5' | 2.04 | 0.57 |
| 1:AA:1016:A:H2' | 1:AA:1017:G:O4' | 2.03 | 0.57 |
| 2:AB:67:THR:C | 2:AB:68:ILE:HD12 | 2.29 | 0.57 |
| 4:AD:30:LYS:HA | 4:AD:35:ARG:HD2 | 1.86 | 0.57 |
| 18:AR:62:GLU:O | 18:AR:65:ILE:HD12 | 2.03 | 0.57 |
| 31:BA:174:C:H3' | 31:BA:175:G:H5" | 1.85 | 0.57 |
| 31:BA:1288:U:C2 | 31:BA:1327:C:O2 | 2.57 | 0.57 |
| 31:BA:1509(B):A:H2' | 31:BA:1510:G:H8 | 1.69 | 0.57 |
| 31:BA:1843:C:O2' | 31:BA:1844:C:H5' | 2.04 | 0.57 |
| 31:BA:2392:A:H2 | 31:BA:2424:C:N4 | 1.97 | 0.57 |
| 33:BD:25:THR:O | 33:BD:25:THR:CG2 | 2.51 | 0.57 |
| 41:BP:105:LEU:O | 41:BP:106:LEU:CB | 2.52 | 0.57 |
| 42:BQ:54:MET:HB3 | 42:BQ:64:ILE:CD1 | 2.34 | 0.57 |
| 44:BS:38:GLN:HB3 | 44:BS:47:THR:HG21 | 1.85 | 0.57 |
| 45:BT:80:SER:HB3 | 45:BT:81:PRO:HD3 | 1.85 | 0.57 |
| 1:CA:601:C:H2' | 1:CA:602:A:C8 | 2.38 | 0.57 |
| 12:CL:102:ARG:HH11 | 12:CL:102:ARG:HG3 | 1.66 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 30:D8:32:LEU:HB3 | 30:D8:34:TRP:CB | 2.34 | 0.57 |
| 31:DA:330:A:H2 | 31:DA:1210:A:H2' | 1.69 | 0.57 |
| 31:DA:528:A:C2 | 31:DA:2043:C:H5' | 2.39 | 0.57 |
| 31:DA:1582:C:O2' | 31:DA:1586:A:H8 | 1.85 | 0.57 |
| 31:DA:1684:C:O2' | 31:DA:1685:C:H5' | 2.04 | 0.57 |
| 31:DA:2506:U:H4' | 31:DA:2507:C:OP1 | 2.04 | 0.57 |
| 31:DA:2615:U:H2' | 31:DA:2616:C:H6 | 1.69 | 0.57 |
| 35:DF:22:ALA:HA | 35:DF:26:ALA:HB2 | 1.86 | 0.57 |
| 39:DN:78:TYR:CD1 | 39:DN:79:PRO:HB3 | 2.39 | 0.57 |
| 39:DN:131:GLN:NE2 | 39:DN:134:ARG:HA | 2.18 | 0.57 |
| 41:DP:10:PRO:CD | 41:DP:11:GLY:H | 2.17 | 0.57 |
| 42:DQ:9:TYR:CD2 | 42:DQ:9:TYR:C | 2.81 | 0.57 |
| 43:DR:101:ALA:O | 43:DR:102:GLU:HB2 | 2.04 | 0.57 |
| 45:DT:91:ARG:HA | 45:DT:117:ASP:H | 1.69 | 0.57 |
| 50:DY:8:LYS:HD3 | 50:DY:28:LYS:HZ2 | 1.68 | 0.57 |
| 50:DY:28:LYS:CE | 50:DY:30:VAL:HG22 | 2.20 | 0.57 |
| 1:AA:411:A:C4 | 1:AA:413:G:O4' | 2.57 | 0.57 |
| 1:AA:458:C:H2' | 1:AA:460:G:H8 | 1.69 | 0.57 |
| 1:AA:1109:C:P | 3:AC:176:HIS:HD1 | 2.27 | 0.57 |
| 1:AA:1452:C:O4' | 1:AA:1456:G:C2 | 2.57 | 0.57 |
| 10:AJ:48:THR:HG23 | 10:AJ:62:HIS:HB3 | 1.85 | 0.57 |
| 28:B6:18:ARG:CG | 28:B6:19:ARG:N | 2.65 | 0.57 |
| 30:B8:22:VAL:HB | 30:B8:53:PRO:CB | 2.33 | 0.57 |
| 31:BA:527:C:N4 | 31:BA:2779:U:OP2 | 2.36 | 0.57 |
| 31:BA:2631:G:N2 | 34:BE:61:ARG:HH12 | 2.02 | 0.57 |
| 31:BA:2762:G:H8 | 31:BA:2762:G:H5" | 1.68 | 0.57 |
| 31:BA:2829:C:C2' | 31:BA:2830:G:H5" | 2.29 | 0.57 |
| 33:BD:35:LYS:HD3 | 33:BD:63:ARG:C | 2.28 | 0.57 |
| 33:BD:35:LYS:CE | 33:BD:64:ILE:C | 2.77 | 0.57 |
| 35:BF:101:LEU:CD1 | 35:BF:102:PRO:HD2 | 2.23 | 0.57 |
| 41:BP:16:ARG:HD3 | 41:BP:18:ARG:N | 2.07 | 0.57 |
| 41:BP:79:ARG:HH21 | 41:BP:109:GLY:HA2 | 1.69 | 0.57 |
| 42:BQ:42:ILE:HD13 | 42:BQ:97:VAL:CG2 | 2.34 | 0.57 |
| 49:BX:40:LYS:HG3 | 49:BX:51:VAL:HB | 1.85 | 0.57 |
| 50:BY:27:VAL:HB | 50:BY:29:GLU:OE1 | 2.04 | 0.57 |
| 1:CA:611:A:H61 | 1:CA:629:G:H1 | 1.50 | 0.57 |
| 1:CA:640:A:N3 | 8:CH:115:SER:HB2 | 2.19 | 0.57 |
| 1:CA:1442(B):A:C2 | 45:DT:118:ARG:CZ | 2.86 | 0.57 |
| 6:CF:27:GLN:O | 6:CF:31:GLU:HB2 | 2.03 | 0.57 |
| 6:CF:61:LEU:HD23 | 6:CF:63:TYR:OH | 2.04 | 0.57 |
| 25:D3:8:LEU:HD13 | 25:D3:31:LEU:CD2 | 2.24 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:17:G:H2' | 31:DA:18:C:C6 | 2.39 | 0.57 |
| 31:DA:1292:U:H2' | 31:DA:1293:C:H6 | 1.64 | 0.57 |
| 31:DA:1812:A:O2' | 33:DD:45:ASN:HB2 | 2.04 | 0.57 |
| 31:DA:1843:C:O2' | 31:DA:1844:C:H5' | 2.04 | 0.57 |
| 31:DA:2101:G:C6 | 31:DA:2102:U:C5 | 2.92 | 0.57 |
| 33:DD:30:GLU:CD | 33:DD:63:ARG:NE | 2.62 | 0.57 |
| 33:DD:186:HIS:CD2 | 33:DD:188:GLU:HB2 | 2.39 | 0.57 |
| 35:DF:3:GLU:HA | 35:DF:24:LEU:HB3 | 1.87 | 0.57 |
| 37:DH:56:SER:HB2 | 37:DH:58:GLU:HG3 | 1.87 | 0.57 |
| 40:DO:31:LYS:C | 40:DO:32:TYR:CD1 | 2.82 | 0.57 |
| 44:DS:54:LEU:HA | 44:DS:57:LYS:O | 2.03 | 0.57 |
| 45:DT:61:PHE:CZ | 45:DT:85:LYS:HE2 | 2.39 | 0.57 |
| 48:DW:6:ILE:HA | 48:DW:103:ILE:O | 2.04 | 0.57 |
| 1:AA:1189:C:O3' | 3:AC:5:ILE:HD12 | 2.03 | 0.57 |
| 1:AA:1442(A):G:C8 | 45:BT:118:ARG:NH1 | 2.71 | 0.57 |
| 1:AA:1493:A:H2' | 31:BA:1913:A:N1 | 2.19 | 0.57 |
| 5:AE:34:VAL:O | 5:AE:41:VAL:HA | 2.04 | 0.57 |
| 30:B8:35:GLN:CG | 31:BA:2420:C:OP1 | 2.51 | 0.57 |
| 31:BA:1292:U:H2' | 31:BA:1293:C:H6 | 1.64 | 0.57 |
| 31:BA:1722:A:C6 | 31:BA:1741:A:N1 | 2.72 | 0.57 |
| 31:BA:1801:G:OP2 | 33:BD:154:LYS:HE3 | 2.04 | 0.57 |
| 34:BE:89:ASP:O | 34:BE:90:THR:HB | 2.04 | 0.57 |
| 35:BF:3:GLU:HA | 35:BF:24:LEU:HB3 | 1.86 | 0.57 |
| 39:BN:27:ALA:CB | 39:BN:106:MET:HE2 | 2.34 | 0.57 |
| 44:BS:29:PHE:N | 44:BS:89:ARG:CD | 2.67 | 0.57 |
| 1:CA:1256:A:H61 | 1:CA:1278:U:C1' | 2.09 | 0.57 |
| 1:CA:1379:G:C6 | 1:CA:1380:U:O4 | 2.58 | 0.57 |
| 2:CB:163:PHE:HA | 2:CB:185:ILE:O | 2.05 | 0.57 |
| 5:CE:71:LEU:HD23 | 5:CE:71:LEU:N | 2.19 | 0.57 |
| 24:D2:23:LYS:HA | 49:DX:5:TYR:CD1 | 2.39 | 0.57 |
| 24:D2:49:LYS:O | 24:D2:50:ILE:C | 2.47 | 0.57 |
| 30:D8:32:LEU:H | 30:D8:32:LEU:HD13 | 1.70 | 0.57 |
| 31:DA:272(G):C:H42 | 31:DA:363(C):G:H1 | 1.52 | 0.57 |
| 31:DA:1106:A:H2' | 31:DA:1107:G:O5' | 2.04 | 0.57 |
| 31:DA:1204:A:N1 | 31:DA:1241:A:H2 | 2.03 | 0.57 |
| 31:DA:2317:C:O2 | 31:DA:2318:G:O4' | 2.21 | 0.57 |
| 31:DA:2636:U:O2' | 31:DA:2637:U:H5" | 2.04 | 0.57 |
| 31:DA:2801(A):A:HO2' | 31:DA:2803:C:H5 | 1.46 | 0.57 |
| 34:DE:111:ARG:HA | 43:DR:2:ARG:HG3 | 1.85 | 0.57 |
| 37:DH:84:SER:O | 37:DH:133:VAL:O | 2.23 | 0.57 |
| 41:DP:121:LYS:HG3 | 41:DP:122:PRO:HD2 | 1.86 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 42:DQ:141:GLN:HG3 | 51:DZ:72:ARG:HH11 | 1.70 | 0.57 |
| 44:DS:67:ARG:H | 44:DS:69:VAL:CG1 | 2.18 | 0.57 |
| 49:DX:65:ARG:CZ | 49:DX:66:LEU:N | 2.65 | 0.57 |
| 50:DY:17:SER:OG | 50:DY:18:GLY:N | 2.34 | 0.57 |
| 1:AA:509:A:O2' | 1:AA:510:A:O5' | 2.23 | 0.57 |
| 5:AE:144:THR:O | 5:AE:148:VAL:HG23 | 2.04 | 0.57 |
| 14:AN:25:VAL:HG23 | 14:AN:38:GLY:O | 2.04 | 0.57 |
| 20:AT:61:SER:O | 20:AT:65:LYS:HG3 | 2.04 | 0.57 |
| 28:B6:51:GLU:O | 28:B6:52:VAL:CB | 2.51 | 0.57 |
| 31:BA:234:C:H2' | 31:BA:235:U:H6 | 1.69 | 0.57 |
| 31:BA:1803:A:H4' | 33:BD:259:THR:CG2 | 2.33 | 0.57 |
| 32:BB:6:C:C2 | 32:BB:116:G:N2 | 2.73 | 0.57 |
| 33:BD:149:PRO:O | 33:BD:150:LYS:HB2 | 2.04 | 0.57 |
| 34:BE:171:GLU:HB2 | 34:BE:185:LYS:HG2 | 1.85 | 0.57 |
| 38:BI:12:LEU:O | 38:BI:12:LEU:HG | 2.04 | 0.57 |
| 38:BI:38:LEU:HB2 | 38:BI:40:THR:CG2 | 2.34 | 0.57 |
| 38:BI:83:ALA:HA | 38:BI:89:TYR:HD1 | 1.69 | 0.57 |
| 41:BP:35:HIS:O | 41:BP:36:LYS:HB2 | 2.04 | 0.57 |
| 42:BQ:76:LYS:H | 42:BQ:88:GLY:HA2 | 1.70 | 0.57 |
| 45:BT:13:ARG:HA | 45:BT:13:ARG:NE | 2.18 | 0.57 |
| 45:BT:129:ARG:NH1 | 45:BT:131:ALA:HB3 | 2.18 | 0.57 |
| 46:BU:92:ARG:NH1 | 46:BU:94:ASN:HD22 | 2.02 | 0.57 |
| 50:BY:28:LYS:CE | 50:BY:37:VAL:HG12 | 2.35 | 0.57 |
| 1:CA:477:A:O2' | 1:CA:479:C:H5' | 2.04 | 0.57 |
| 1:CA:951:G:H1' | 1:CA:970:C:O2' | 2.03 | 0.57 |
| 1:CA:1058:G:H2' | 1:CA:1059:C:O4' | 2.04 | 0.57 |
| 2:CB:101:MET:HG2 | 2:CB:108:ILE:HG21 | 1.87 | 0.57 |
| 13:CM:86:CYS:HB2 | 19:CS:73:GLU:HB3 | 1.86 | 0.57 |
| 22:D0:43:THR:HG22 | 31:DA:2331:G:O2' | 2.04 | 0.57 |
| 24:D2:30:ARG:HH11 | 24:D2:30:ARG:HG3 | 1.69 | 0.57 |
| 30:D8:32:LEU:HD23 | 30:D8:35:GLN:H | 1.69 | 0.57 |
| 30:D8:50:LEU:HD12 | 30:D8:51:ALA:N | 2.19 | 0.57 |
| 31:DA:84:A:H5' | 50:DY:9:LYS:HD2 | 1.85 | 0.57 |
| 31:DA:142:A:H1' | 31:DA:1408:C:O4' | 2.03 | 0.57 |
| 31:DA:534:U:O2' | 46:DU:49:HIS:CD2 | 2.57 | 0.57 |
| 31:DA:1635:G:H8 | 31:DA:1635:G:H5' | 1.70 | 0.57 |
| 31:DA:1722:A:C6 | 31:DA:1741:A:N1 | 2.73 | 0.57 |
| 31:DA:2593:U:H2' | 31:DA:2594:C:H6 | 1.70 | 0.57 |
| 33:DD:30:GLU:CD | 33:DD:63:ARG:HE | 2.13 | 0.57 |
| 35:DF:74:ARG:HG2 | 35:DF:74:ARG:O | 2.03 | 0.57 |
| 41:DP:87:ASP:O | 41:DP:90:ARG:HB2 | 2.04 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 51:DZ:120:ILE:HG22 | 51:DZ:120:ILE:O | 2.04 | 0.57 |
| 1:AA:619:U:H2' | 4:AD:135:LEU:HD21 | 1.86 | 0.57 |
| 1:AA:681:C:N3 | 1:AA:710:G:C2 | 2.72 | 0.57 |
| 1:AA:822:C:O2' | 1:AA:823:G:H5' | 2.04 | 0.57 |
| 5:AE:101:ILE:HD13 | 5:AE:118:ILE:O | 2.03 | 0.57 |
| 6:AF:4:TYR:HA | 6:AF:91:VAL:O | 2.05 | 0.57 |
| 10:AJ:90:LEU:N | 10:AJ:91:PRO:HD3 | 2.19 | 0.57 |
| 15:AO:67:LEU:CD2 | 15:AO:78:TYR:HE1 | 2.18 | 0.57 |
| 24:B2:29:LYS:O | 24:B2:32:LEU:N | 2.37 | 0.57 |
| 26:B4:14:ILE:HA | 36:BG:5:VAL:HG13 | 1.87 | 0.57 |
| 27:B5:40:LYS:NZ | 27:B5:46:CYS:HB3 | 2.20 | 0.57 |
| 27:B5:55:ARG:O | 27:B5:56:LYS:HG3 | 2.04 | 0.57 |
| 31:BA:470:A:C2 | 31:BA:471:A:C4 | 2.92 | 0.57 |
| 31:BA:676:A:H8 | 31:BA:2069:G:H21 | 0.85 | 0.57 |
| 31:BA:1300:U:H3' | 31:BA:1301:A:H5" | 1.85 | 0.57 |
| 31:BA:1531:C:H3' | 31:BA:1532:C:C5' | 2.34 | 0.57 |
| 31:BA:1741:A:N7 | 31:BA:1742:G:C2 | 2.73 | 0.57 |
| 31:BA:2099:U:H2' | 31:BA:2099:U:O2 | 2.04 | 0.57 |
| 34:BE:70:ALA:O | 34:BE:72:VAL:N | 2.37 | 0.57 |
| 39:BN:111:PRO:HA | 39:BN:114:ARG:NH1 | 2.19 | 0.57 |
| 40:BO:18:LYS:HB2 | 40:BO:45:GLU:HG2 | 1.86 | 0.57 |
| 41:BP:67:MET:HA | 41:BP:67:MET:HE3 | 1.86 | 0.57 |
| 41:BP:131:SER:C | 41:BP:133:SER:N | 2.60 | 0.57 |
| 43:BR:2:ARG:N | 43:BR:2:ARG:HD2 | 2.20 | 0.57 |
| 44:BS:67:ARG:H | 44:BS:69:VAL:CG1 | 2.17 | 0.57 |
| 49:BX:85:PRO:O | 49:BX:86:GLY:C | 2.46 | 0.57 |
| 1:CA:539:A:H2' | 1:CA:540:G:C8 | 2.39 | 0.57 |
| 1:CA:556:C:O2' | 1:CA:557:G:H5' | 2.04 | 0.57 |
| 1:CA:617:G:C6 | 1:CA:618:C:C5 | 2.93 | 0.57 |
| 5:CE:71:LEU:O | 5:CE:72:GLN:HG3 | 2.04 | 0.57 |
| 13:CM:83:ASP:CG | 13:CM:84:ILE:H | 2.13 | 0.57 |
| 22:D0:48:GLY:HA3 | 22:D0:80:HIS:ND1 | 2.20 | 0.57 |
| 25:D3:8:LEU:CD1 | 25:D3:31:LEU:HA | 2.34 | 0.57 |
| 25:D3:19:GLN:NE2 | 25:D3:52:HIS:HE1 | 2.02 | 0.57 |
| 27:D5:2:ALA:HA | 31:DA:2015:A:C1' | 2.28 | 0.57 |
| 27:D5:2:ALA:N | 31:DA:747:U:C4 | 2.73 | 0.57 |
| 28:D6:20:ASN:O | 28:D6:21:TYR:CD1 | 2.57 | 0.57 |
| 31:DA:743:G:C2' | 31:DA:744:G:H5' | 2.34 | 0.57 |
| 31:DA:1487:G:C2 | 31:DA:1488:G:C4 | 2.93 | 0.57 |
| 31:DA:1509(A):A:H2' | 31:DA:1509(B):A:H8 | 1.69 | 0.57 |
| 31:DA:1786:A:H2 | 31:DA:2606:C:H1' | 1.68 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1827:C:OP2 | 33:DD:222:ARG:HD2 | 2.04 | 0.57 |
| 31:DA:2275:C:O2' | 42:DQ:83:MET:HA | 2.05 | 0.57 |
| 35:DF:132:VAL:CG2 | 35:DF:133:ASN:H | 2.17 | 0.57 |
| 41:DP:10:PRO:HD2 | 41:DP:11:GLY:N | 2.19 | 0.57 |
| 41:DP:108:LYS:O | 41:DP:110:TYR:N | 2.37 | 0.57 |
| 44:DS:56:LEU:HD23 | 44:DS:56:LEU:C | 2.29 | 0.57 |
| 46:DU:95:LEU:C | 46:DU:97:ASP:H | 2.12 | 0.57 |
| 1:AA:484:G:H4' | 1:AA:485:G:OP1 | 2.04 | 0.57 |
| 1:AA:617:G:C6 | 1:AA:618:C:C5 | 2.92 | 0.57 |
| 3:AC:27:LYS:HA | 3:AC:27:LYS:NZ | 2.19 | 0.57 |
| 6:AF:27:GLN:O | 6:AF:31:GLU:HB2 | 2.04 | 0.57 |
| 19:AS:6:LYS:HD3 | 19:AS:7:LYS:HE3 | 1.86 | 0.57 |
| 24:B2:41:ILE:O | 24:B2:43:GLN:N | 2.37 | 0.57 |
| 29:B7:10:ARG:NH2 | 31:BA:1378:A:OP1 | 2.37 | 0.57 |
| 30:B8:25:MET:HB2 | 41:BP:62:LEU:CD2 | 2.35 | 0.57 |
| 31:BA:229:A:H3' | 31:BA:230:U:H5' | 1.86 | 0.57 |
| 31:BA:389:G:N1 | 41:BP:70:GLN:HG3 | 2.19 | 0.57 |
| 31:BA:995:C:O2 | 46:BU:57:PHE:CD2 | 2.58 | 0.57 |
| 31:BA:1169:G:C8 | 31:BA:1169:G:H3' | 2.40 | 0.57 |
| 31:BA:1225:G:OP1 | 47:BV:88:ARG:HD2 | 2.04 | 0.57 |
| 31:BA:1818:U:H2' | 33:BD:157:ARG:HG3 | 1.87 | 0.57 |
| 31:BA:1973:G:H2' | 31:BA:1974:C:C6 | 2.39 | 0.57 |
| 31:BA:2406:U:O4 | 41:BP:70:GLN:HB3 | 2.05 | 0.57 |
| 31:BA:2897:U:O2 | 31:BA:2897:U:H2' | 2.04 | 0.57 |
| 32:BB:21:G:O2' | 32:BB:22:U:O5' | 2.22 | 0.57 |
| 32:BB:31:C:H4' | 36:BG:29:TRP:CH2 | 2.39 | 0.57 |
| 33:BD:102:LYS:C | 33:BD:103:ARG:HG2 | 2.29 | 0.57 |
| 33:BD:222:ARG:O | 33:BD:225:ALA:HB3 | 2.04 | 0.57 |
| 41:BP:8:PRO:O | 41:BP:9:ASN:C | 2.47 | 0.57 |
| 41:BP:30:THR:CG2 | 41:BP:31:ALA:N | 2.67 | 0.57 |
| 42:BQ:141:GLN:HE21 | 51:BZ:71:VAL:C | 2.12 | 0.57 |
| 50:BY:45:VAL:CG1 | 50:BY:62:GLU:OE2 | 2.52 | 0.57 |
| 51:BZ:152:ALA:HB2 | 51:BZ:168:GLU:HA | 1.87 | 0.57 |
| 1:CA:59:A:H2' | 1:CA:59:A:N3 | 2.17 | 0.57 |
| 1:CA:109:A:H2' | 1:CA:326:G:N2 | 2.20 | 0.57 |
| 1:CA:1218:C:H2' | 1:CA:1219:U:C6 | 2.39 | 0.57 |
| 2:CB:204:ASN:HD21 | 2:CB:207:ALA:H | 1.51 | 0.57 |
| 4:CD:14:ARG:HA | 4:CD:39:PRO:HB3 | 1.87 | 0.57 |
| 4:CD:146:ILE:HD12 | 4:CD:146:ILE:N | 2.20 | 0.57 |
| 5:CE:88:LYS:HB3 | 5:CE:123:LEU:HB2 | 1.85 | 0.57 |
| 10:CJ:70:ARG:HH11 | 10:CJ:70:ARG:HG3 | 1.68 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 14:CN:25:VAL:HG23 | 14:CN:38:GLY:O | 2.05 | 0.57 |
| 18:CR:62:GLU:O | 18:CR:65:ILE:HD12 | 2.03 | 0.57 |
| 23:D1:73:LEU:HD11 | 23:D1:94:LEU:HG | 1.85 | 0.57 |
| 30:D8:39:LYS:C | 30:D8:39:LYS:HE2 | 2.30 | 0.57 |
| 31:DA:558:G:OP1 | 39:DN:111:PRO:HD2 | 2.04 | 0.57 |
| 31:DA:580:C:H2' | 31:DA:581:C:C6 | 2.38 | 0.57 |
| 31:DA:673:C:H5" | 35:DF:81:PRO:HD2 | 1.86 | 0.57 |
| 31:DA:1528:A:N3 | 31:DA:1528:A:H2' | 2.18 | 0.57 |
| 31:DA:1709:U:O2' | 31:DA:2859:G:H1' | 2.04 | 0.57 |
| 31:DA:1741:A:N7 | 31:DA:1742:G:C2 | 2.73 | 0.57 |
| 31:DA:2314:C:O2 | 31:DA:2315:G:C8 | 2.58 | 0.57 |
| 31:DA:2334:G:H4' | 31:DA:2335:A:OP2 | 2.02 | 0.57 |
| 32:DB:31:C:H4' | 36:DG:29:TRP:CH2 | 2.40 | 0.57 |
| 33:DD:35:LYS:CE | 33:DD:65:ILE:HG22 | 2.34 | 0.57 |
| 34:DE:134:ILE:HG12 | 34:DE:134:ILE:O | 2.04 | 0.57 |
| 37:DH:40:GLU:C | 37:DH:41:MET:HE3 | 2.30 | 0.57 |
| 41:DP:83:VAL:HG12 | 41:DP:112:LEU:HD21 | 1.86 | 0.57 |
| 45:DT:13:ARG:HH21 | 45:DT:15:VAL:CG1 | 2.17 | 0.57 |
| 45:DT:102:ILE:HB | 45:DT:110:ILE:CD1 | 2.34 | 0.57 |
| 49:DX:18:TYR:O | 49:DX:19:ALA:C | 2.47 | 0.57 |
| 50:DY:45:VAL:CG1 | 50:DY:62:GLU:OE2 | 2.53 | 0.57 |
| 1:AA:640:A:N3 | 8:AH:115:SER:HB2 | 2.19 | 0.57 |
| 1:AA:706:A:N7 | 1:AA:707:C:H5 | 2.02 | 0.57 |
| 1:AA:779:C:C2' | 1:AA:780:A:H5' | 2.35 | 0.57 |
| 1:AA:1058:G:H2' | 1:AA:1059:C:O4' | 2.04 | 0.57 |
| 2:AB:168:THR:HG21 | 2:AB:192:SER:HA | 1.85 | 0.57 |
| 7:AG:75:VAL:HG12 | 7:AG:88:PRO:HB3 | 1.85 | 0.57 |
| 13:AM:24:GLY:C | 13:AM:25:ILE:HD12 | 2.29 | 0.57 |
| 13:AM:25:ILE:CD1 | 13:AM:66:LEU:HD23 | 2.33 | 0.57 |
| 13:AM:84:ILE:HG12 | 19:AS:66:MET:HE1 | 1.87 | 0.57 |
| 15:AO:67:LEU:HD22 | 15:AO:78:TYR:HE1 | 1.70 | 0.57 |
| 30:B8:62:LEU:O | 30:B8:64:TYR:N | 2.37 | 0.57 |
| 31:BA:531:C:H4' | 31:BA:532:A:H5" | 1.86 | 0.57 |
| 31:BA:645:C:H3' | 31:BA:645:C:O2 | 2.03 | 0.57 |
| 31:BA:1006:C:C2 | 31:BA:1138:G:N2 | 2.73 | 0.57 |
| 31:BA:1247:A:OP1 | 35:BF:95:ARG:NH2 | 2.34 | 0.57 |
| 31:BA:1635:G:H5' | 31:BA:1635:G:H8 | 1.70 | 0.57 |
| 31:BA:1980:G:O2' | 31:BA:1982:C:OP2 | 2.12 | 0.57 |
| 31:BA:2206:G:H3' | 31:BA:2206:G:N3 | 2.19 | 0.57 |
| 34:BE:87:GLU:O | 34:BE:89:ASP:N | 2.37 | 0.57 |
| 36:BG:16:ARG:O | 36:BG:20:ILE:HG13 | 2.04 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 40:BO:23:ARG:HD2 | 40:BO:24:VAL:H | 1.68 | 0.57 |
| 42:BQ:63:LYS:HB2 | 42:BQ:63:LYS:NZ | 2.19 | 0.57 |
| 42:BQ:75:THR:HA | 42:BQ:89:ASN:H | 1.70 | 0.57 |
| 49:BX:55:ASN:HB2 | 49:BX:78:LYS:CD | 2.34 | 0.57 |
| 1:CA:822:C:O2' | 1:CA:823:G:H5' | 2.04 | 0.57 |
| 1:CA:977:A:H2' | 1:CA:978:A:H5' | 1.85 | 0.57 |
| 1:CA:1109:C:P | 3:CC:176:HIS:HD1 | 2.27 | 0.57 |
| 10:CJ:6:ILE:HG22 | 10:CJ:98:ILE:HG13 | 1.84 | 0.57 |
| 27:D5:50:GLY:O | 27:D5:51:TYR:CD1 | 2.57 | 0.57 |
| 31:DA:672:C:O2' | 31:DA:673:C:H5' | 2.05 | 0.57 |
| 31:DA:1225:G:OP1 | 47:DV:88:ARG:HD2 | 2.05 | 0.57 |
| 31:DA:1264:G:H3' | 31:DA:1265:A:H5" | 1.86 | 0.57 |
| 31:DA:1981:A:H5" | 31:DA:1982:C:OP2 | 2.05 | 0.57 |
| 31:DA:2523:G:C2' | 31:DA:2524:G:H5" | 2.35 | 0.57 |
| 31:DA:2658:C:H5' | 31:DA:2659:G:OP2 | 2.04 | 0.57 |
| 32:DB:10:C:C4 | 32:DB:11:C:C5 | 2.92 | 0.57 |
| 33:DD:125:ILE:HG22 | 33:DD:125:ILE:O | 2.04 | 0.57 |
| 39:DN:58:ASP:C | 39:DN:60:ILE:H | 2.12 | 0.57 |
| 40:DO:9:GLU:O | 40:DO:83:ALA:HA | 2.05 | 0.57 |
| 46:DU:88:ILE:O | 46:DU:88:ILE:CD1 | 2.53 | 0.57 |
| 47:DV:53:GLU:O | 47:DV:55:ALA:N | 2.38 | 0.57 |
| 48:DW:56:ALA:O | 48:DW:57:ASN:C | 2.45 | 0.57 |
| 49:DX:70:LEU:O | 49:DX:71:GLY:O | 2.22 | 0.57 |
| 1:AA:155:C:H2' | 1:AA:156:G:C8 | 2.39 | 0.57 |
| 1:AA:951:G:H1' | 1:AA:970:C:O2' | 2.05 | 0.57 |
| 1:AA:1347:G:N2 | 1:AA:1373:G:H2' | 2.20 | 0.57 |
| 5:AE:50:GLU:OE2 | 5:AE:51:VAL:HG23 | 2.05 | 0.57 |
| 30:B8:32:LEU:CG | 30:B8:35:GLN:H | 2.18 | 0.57 |
| 31:BA:547:A:H8 | 31:BA:549:G:C6 | 2.23 | 0.57 |
| 31:BA:902:C:O2' | 31:BA:903:C:H5' | 2.05 | 0.57 |
| 31:BA:951:C:O2' | 31:BA:952:G:H5' | 2.04 | 0.57 |
| 31:BA:1112:G:H1' | 31:BA:1113:U:OP1 | 2.05 | 0.57 |
| 31:BA:1316:U:H2' | 31:BA:1317:A:C8 | 2.40 | 0.57 |
| 37:BH:44:VAL:HG12 | 37:BH:45:VAL:N | 2.16 | 0.57 |
| 38:BI:78:THR:HA | 38:BI:141:LYS:O | 2.05 | 0.57 |
| 39:BN:96:GLU:CD | 39:BN:96:GLU:H | 2.13 | 0.57 |
| 42:BQ:140:ALA:CB | 51:BZ:53:ILE:HG13 | 2.30 | 0.57 |
| 44:BS:29:PHE:HD2 | 44:BS:30:ARG:N | 2.03 | 0.57 |
| 47:BV:43:GLU:O | 47:BV:44:LYS:O | 2.23 | 0.57 |
| 49:BX:30:VAL:HG23 | 49:BX:76:ARG:HA | 1.87 | 0.57 |
| 50:BY:37:VAL:HG23 | 50:BY:38:ILE:N | 2.19 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:458:C:H2' | 1:CA:460:G:H8 | 1.70 | 0.57 |
| 1:CA:1004:A:H2' | 1:CA:1038:C:O2 | 2.04 | 0.57 |
| 1:CA:1347:G:N2 | 1:CA:1373:G:H2' | 2.20 | 0.57 |
| 9:CI:114:TYR:N | 9:CI:114:TYR:CD2 | 2.66 | 0.57 |
| 11:CK:57:THR:HG22 | 11:CK:59:TYR:H | 1.70 | 0.57 |
| 13:CM:84:ILE:HG12 | 19:CS:66:MET:HE1 | 1.86 | 0.57 |
| 16:CP:26:ARG:HD2 | 16:CP:31:LYS:O | 2.05 | 0.57 |
| 17:CQ:40:LYS:HD2 | 17:CQ:42:TYR:CZ | 2.39 | 0.57 |
| 22:D0:42:GLY:HA2 | 22:D0:57:PHE:CD2 | 2.39 | 0.57 |
| 30:D8:43:GLN:O | 30:D8:44:LYS:HD2 | 2.03 | 0.57 |
| 31:DA:195:A:H4' | 31:DA:251:A:O2' | 2.03 | 0.57 |
| 31:DA:2033:A:H4' | 31:DA:2034:U:OP1 | 2.05 | 0.57 |
| 31:DA:2197:U:H1' | 31:DA:2198:A:C8 | 2.39 | 0.57 |
| 31:DA:2529:G:H5" | 31:DA:2530:A:H5" | 1.86 | 0.57 |
| 31:DA:2712:U:O2 | 31:DA:2712:U:C5' | 2.52 | 0.57 |
| 34:DE:75:VAL:C | 34:DE:77:ILE:N | 2.63 | 0.57 |
| 39:DN:3:THR:CA | 39:DN:4:TYR:CE1 | 2.86 | 0.57 |
| 45:DT:17:THR:O | 45:DT:18:ASP:HB3 | 2.04 | 0.57 |
| 45:DT:28:VAL:HG22 | 45:DT:46:GLU:CA | 2.34 | 0.57 |
| 48:DW:15:ARG:HA | 48:DW:18:ARG:HD2 | 1.87 | 0.57 |
| 1:AA:147:G:H2' | 1:AA:148:G:H5' | 1.86 | 0.57 |
| 1:AA:963:G:N3 | 10:AJ:55:LYS:NZ | 2.46 | 0.57 |
| 1:AA:1190:G:H3' | 3:AC:3:ASN:ND2 | 2.20 | 0.57 |
| 4:AD:118:ARG:O | 4:AD:121:VAL:HG23 | 2.05 | 0.57 |
| 5:AE:10:MET:HG3 | 5:AE:13:ILE:HD13 | 1.87 | 0.57 |
| 6:AF:10:LEU:HD12 | 6:AF:10:LEU:N | 2.20 | 0.57 |
| 6:AF:18:GLN:HA | 6:AF:21:LEU:HD22 | 1.86 | 0.57 |
| 7:AG:138:LYS:HE2 | 7:AG:142:GLU:OE2 | 2.05 | 0.57 |
| 10:AJ:89:ASP:C | 10:AJ:91:PRO:HD3 | 2.30 | 0.57 |
| 13:AM:83:ASP:CG | 13:AM:84:ILE:H | 2.12 | 0.57 |
| 28:B6:24:GLU:HA | 28:B6:24:GLU:OE1 | 2.05 | 0.57 |
| 31:BA:1210:A:H5' | 31:BA:1212:G:C5' | 2.34 | 0.57 |
| 31:BA:2658:C:H5' | 31:BA:2659:G:OP2 | 2.05 | 0.57 |
| 32:BB:66:A:H61 | 32:BB:108:U:H2' | 1.69 | 0.57 |
| 33:BD:267:SER:O | 33:BD:268:ARG:CB | 2.53 | 0.57 |
| 34:BE:111:ARG:CZ | 43:BR:2:ARG:HH21 | 2.17 | 0.57 |
| 34:BE:134:ILE:O | 34:BE:134:ILE:CG1 | 2.53 | 0.57 |
| 44:BS:56:LEU:O | 44:BS:57:LYS:HB2 | 2.03 | 0.57 |
| 45:BT:30:VAL:O | 45:BT:30:VAL:CG2 | 2.51 | 0.57 |
| 46:BU:8:VAL:HG13 | 46:BU:12:ARG:HG3 | 1.86 | 0.57 |
| 49:BX:82:GLN:CD | 49:BX:83:VAL:HG22 | 2.29 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:131:C:H2' | 1:CA:132:C:C6 | 2.40 | 0.57 |
| 1:CA:577:G:C4 | 1:CA:816:A:C2 | 2.92 | 0.57 |
| 1:CA:1158:C:N3 | 1:CA:1181:G:N2 | 2.52 | 0.57 |
| 5:CE:18:ARG:NH2 | 5:CE:25:ARG:HG2 | 2.20 | 0.57 |
| 7:CG:115:ARG:HB2 | 7:CG:118:VAL:HG22 | 1.86 | 0.57 |
| 12:CL:41:ARG:HG2 | 12:CL:42:THR:H | 1.70 | 0.57 |
| 13:CM:15:VAL:O | 13:CM:19:LEU:HD23 | 2.05 | 0.57 |
| 17:CQ:22:LEU:HD12 | 17:CQ:23:VAL:H | 1.69 | 0.57 |
| 20:CT:58:LYS:HE3 | 20:CT:62:LEU:HD11 | 1.87 | 0.57 |
| 22:D0:74:ARG:NH2 | 32:DB:13:A:H8 | 2.02 | 0.57 |
| 23:D1:41:ARG:NH2 | 31:DA:205:G:O6 | 2.38 | 0.57 |
| 30:D8:35:GLN:NE2 | 30:D8:36:LYS:NZ | 2.52 | 0.57 |
| 30:D8:35:GLN:HE21 | 30:D8:36:LYS:HZ3 | 1.53 | 0.57 |
| 31:DA:69:C:H2' | 31:DA:69:C:O2 | 2.05 | 0.57 |
| 31:DA:672:C:C2 | 31:DA:673:C:C5 | 2.92 | 0.57 |
| 31:DA:814:C:C5 | 41:DP:27:HIS:CE1 | 2.93 | 0.57 |
| 31:DA:1270:C:H5" | 31:DA:1271:G:O5' | 2.04 | 0.57 |
| 31:DA:1449:A:HO2' | 31:DA:1530:C:H5 | 1.51 | 0.57 |
| 31:DA:1778:U:H2' | 31:DA:1784:A:N6 | 2.20 | 0.57 |
| 31:DA:2611:U:H5' | 31:DA:2611:U:H6 | 1.70 | 0.57 |
| 33:DD:95:LEU:H | 33:DD:95:LEU:HD12 | 1.70 | 0.57 |
| 36:DG:16:ARG:HE | 36:DG:31:VAL:HG11 | 1.70 | 0.57 |
| 39:DN:57:ALA:O | 39:DN:58:ASP:CG | 2.48 | 0.57 |
| 42:DQ:8:LYS:CG | 42:DQ:9:TYR:N | 2.68 | 0.57 |
| 43:DR:104:ARG:HD2 | 43:DR:111:LEU:HD11 | 1.86 | 0.57 |
| 50:DY:37:VAL:HG21 | 50:DY:67:LEU:HD23 | 1.87 | 0.57 |
| 1:AA:986:A:H2' | 1:AA:987:G:O4' | 2.05 | 0.57 |
| 1:AA:1071:C:O2' | 1:AA:1072:G:H5' | 2.05 | 0.57 |
| 1:AA:1281:U:H5' | 1:AA:1282:C:H5 | 1.69 | 0.57 |
| 28:B6:19:ARG:CG | 28:B6:20:ASN:H | 2.18 | 0.57 |
| 31:BA:912:C:C2 | 31:BA:913:U:C5 | 2.92 | 0.57 |
| 31:BA:1652:A:O2' | 31:BA:1653:G:H5' | 2.05 | 0.57 |
| 31:BA:1722:A:O2' | 31:BA:1739:U:H5" | 2.04 | 0.57 |
| 31:BA:2312:U:OP1 | 36:BG:74:LYS:HG3 | 2.04 | 0.57 |
| 31:BA:2475:C:N4 | 31:BA:2529:G:H22 | 2.01 | 0.57 |
| 31:BA:2734:A:H8 | 31:BA:2734:A:C5' | 2.18 | 0.57 |
| 34:BE:116:VAL:HG13 | 34:BE:122:PHE:CD2 | 2.40 | 0.57 |
| 35:BF:46:ARG:CG | 35:BF:46:ARG:NH1 | 2.65 | 0.57 |
| 39:BN:57:ALA:O | 39:BN:58:ASP:CG | 2.48 | 0.57 |
| 41:BP:64:LYS:O | 41:BP:65:ARG:C | 2.48 | 0.57 |
| 44:BS:92:TYR:HD1 | 44:BS:93:LYS:N | 1.86 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 46:BU:95:LEU:HD22 | 47:BV:4:ILE:HD13 | 1.86 | 0.57 |
| 50:BY:19:LYS:HB3 | 50:BY:20:TYR:CD1 | 2.40 | 0.57 |
| 1:CA:457:C:H2' | 1:CA:458:C:C6 | 2.40 | 0.57 |
| 1:CA:674:G:N2 | 11:CK:116:HIS:HB2 | 2.19 | 0.57 |
| 2:CB:178:ARG:HH21 | 8:CH:68:ARG:HH22 | 1.52 | 0.57 |
| 2:CB:218:ALA:O | 2:CB:222:ILE:HG13 | 2.04 | 0.57 |
| 15:CO:87:ILE:CG2 | 15:CO:88:ARG:H | 2.18 | 0.57 |
| 27:D5:33:CYS:SG | 27:D5:49:CYS:CB | 2.93 | 0.57 |
| 30:D8:25:MET:HG3 | 41:DP:64:LYS:HB2 | 1.86 | 0.57 |
| 31:DA:174:C:H3' | 31:DA:175:G:H5" | 1.87 | 0.57 |
| 31:DA:737:C:C2' | 31:DA:738:G:O5' | 2.52 | 0.57 |
| 31:DA:2050:C:H1' | 34:DE:156:MET:CE | 2.35 | 0.57 |
| 35:DF:3:GLU:O | 35:DF:24:LEU:HG | 2.05 | 0.57 |
| 41:DP:131:SER:C | 41:DP:133:SER:N | 2.61 | 0.57 |
| 42:DQ:75:THR:HG22 | 42:DQ:88:GLY:HA3 | 1.85 | 0.57 |
| 42:DQ:134:ARG:HH21 | 51:DZ:122:ARG:CZ | 2.18 | 0.57 |
| 47:DV:5:VAL:HG21 | 47:DV:36:PRO:HB2 | 1.84 | 0.57 |
| 50:DY:76:CYS:SG | 50:DY:77:PRO:CD | 2.86 | 0.57 |
| 1:AA:59:A:H2' | 1:AA:59:A:N3 | 2.19 | 0.56 |
| 2:AB:106:LYS:O | 2:AB:110:GLN:HG3 | 2.05 | 0.56 |
| 3:AC:95:THR:HG22 | 3:AC:97:LYS:H | 1.70 | 0.56 |
| 31:BA:867:C:C5 | 31:BA:868:U:C5 | 2.93 | 0.56 |
| 31:BA:1508:A:C2' | 31:BA:1509:C:OP1 | 2.54 | 0.56 |
| 31:BA:1657:C:H2' | 31:BA:1658:C:C6 | 2.40 | 0.56 |
| 31:BA:2473:U:N3 | 31:BA:2474:C:C6 | 2.73 | 0.56 |
| 32:BB:10:C:C4 | 32:BB:11:C:C5 | 2.93 | 0.56 |
| 48:BW:82:LEU:HG | 48:BW:84:ARG:HH21 | 1.69 | 0.56 |
| 1:CA:986:A:H2' | 1:CA:987:G:O4' | 2.05 | 0.56 |
| 1:CA:1228:C:P | 13:CM:108:ARG:HH22 | 2.28 | 0.56 |
| 3:CC:32:LEU:HD22 | 3:CC:59:ARG:NH1 | 2.20 | 0.56 |
| 3:CC:130:VAL:O | 3:CC:134:ILE:HG12 | 2.05 | 0.56 |
| 12:CL:70:ILE:HG12 | 12:CL:100:ILE:HD12 | 1.85 | 0.56 |
| 22:D0:1:MET:HA | 31:DA:2451:A:H4' | 1.87 | 0.56 |
| 25:D3:19:GLN:NE2 | 25:D3:52:HIS:CE1 | 2.73 | 0.56 |
| 31:DA:218:A:C2 | 31:DA:235:U:H4' | 2.38 | 0.56 |
| 31:DA:641:C:O2' | 31:DA:2350:C:OP1 | 2.18 | 0.56 |
| 31:DA:645:C:H3' | 31:DA:645:C:O2 | 2.05 | 0.56 |
| 31:DA:919:G:H5' | 32:DB:81:G:H1' | 1.87 | 0.56 |
| 31:DA:2734:A:H8 | 31:DA:2734:A:H5' | 1.69 | 0.56 |
| 40:DO:18:LYS:HB2 | 40:DO:45:GLU:HG2 | 1.87 | 0.56 |
| 44:DS:89:ARG:HE | 44:DS:89:ARG:CA | 2.10 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 45:DT:13:ARG:HA | 45:DT:13:ARG:NE | 2.20 | 0.56 |
| 47:DV:19:LYS:HG2 | 47:DV:96:ILE:CG2 | 2.31 | 0.56 |
| 49:DX:82:GLN:CD | 49:DX:83:VAL:HG22 | 2.30 | 0.56 |
| 1:AA:343:U:O2' | 1:AA:346:G:O6 | 2.23 | 0.56 |
| 1:AA:386:C:C2' | 1:AA:387:U:H5' | 2.35 | 0.56 |
| 4:AD:111:ALA:HB2 | 4:AD:120:LEU:HD12 | 1.87 | 0.56 |
| 5:AE:51:VAL:O | 5:AE:55:VAL:HG23 | 2.05 | 0.56 |
| 6:AF:61:LEU:HB3 | 6:AF:63:TYR:HE2 | 1.69 | 0.56 |
| 10:AJ:61:GLU:OE1 | 14:AN:58:LYS:HE2 | 2.05 | 0.56 |
| 11:AK:106:LYS:HG3 | 11:AK:106:LYS:O | 2.04 | 0.56 |
| 13:AM:86:CYS:HB2 | 19:AS:73:GLU:HB3 | 1.86 | 0.56 |
| 23:B1:46:LEU:HD12 | 23:B1:46:LEU:H | 1.64 | 0.56 |
| 27:B5:16:ARG:NH2 | 31:BA:517:C:OP1 | 2.37 | 0.56 |
| 29:B7:16:HIS:HB3 | 29:B7:44:PRO:HG2 | 1.86 | 0.56 |
| 31:BA:128:C:C6 | 31:BA:128:C:C3' | 2.88 | 0.56 |
| 31:BA:269:U:O2 | 31:BA:269:U:H2' | 2.03 | 0.56 |
| 31:BA:727:A:C2 | 33:BD:9:TYR:CD2 | 2.93 | 0.56 |
| 31:BA:1022:G:C6 | 31:BA:1140:C:C4 | 2.93 | 0.56 |
| 31:BA:1170:G:H1 | 31:BA:1179:C:N4 | 2.00 | 0.56 |
| 31:BA:1372:U:H2' | 31:BA:1373:A:O4' | 2.05 | 0.56 |
| 31:BA:1812:A:O2' | 33:BD:45:ASN:HB2 | 2.05 | 0.56 |
| 31:BA:2052:G:C8 | 34:BE:141:ILE:HD11 | 2.40 | 0.56 |
| 31:BA:2652:C:O2' | 31:BA:2653:U:H5' | 2.05 | 0.56 |
| 33:BD:35:LYS:NZ | 33:BD:104:TYR:CD1 | 2.73 | 0.56 |
| 34:BE:1:MET:O | 34:BE:2:LYS:C | 2.48 | 0.56 |
| 39:BN:39:ARG:HE | 39:BN:41:ASP:CG | 2.12 | 0.56 |
| 39:BN:72:TYR:O | 39:BN:73:THR:C | 2.48 | 0.56 |
| 43:BR:71:GLN:HE21 | 43:BR:71:GLN:HA | 1.71 | 0.56 |
| 46:BU:112:ARG:CG | 46:BU:112:ARG:NH1 | 2.59 | 0.56 |
| 48:BW:9:TYR:N | 48:BW:102:HIS:HD2 | 1.95 | 0.56 |
| 50:BY:37:VAL:HG22 | 50:BY:67:LEU:O | 2.04 | 0.56 |
| 51:BZ:5:LEU:HG | 51:BZ:47:VAL:HG21 | 1.86 | 0.56 |
| 1:CA:22:G:H2' | 1:CA:23:C:H6 | 1.71 | 0.56 |
| 1:CA:240:C:H2' | 1:CA:241:C:H6 | 1.70 | 0.56 |
| 1:CA:626:U:C2 | 1:CA:627:G:C8 | 2.94 | 0.56 |
| 1:CA:707:C:OP1 | 11:CK:85:ARG:NH1 | 2.38 | 0.56 |
| 1:CA:927:G:OP2 | 1:CA:1503:A:C4 | 2.58 | 0.56 |
| 3:CC:27:LYS:NZ | 3:CC:27:LYS:HA | 2.20 | 0.56 |
| 6:CF:18:GLN:HA | 6:CF:21:LEU:HD22 | 1.87 | 0.56 |
| 6:CF:42:GLU:OE1 | 6:CF:59:TYR:HE2 | 1.88 | 0.56 |
| 23:D1:86:SER:HA | 23:D1:89:GLU:OE2 | 2.04 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 27:D5:50:GLY:HA3 | 27:D5:56:LYS:CG | 2.34 | 0.56 |
| 27:D5:51:TYR:CD2 | 27:D5:52:TYR:CE2 | 2.93 | 0.56 |
| 31:DA:389:G:N1 | 41:DP:70:GLN:HG3 | 2.20 | 0.56 |
| 31:DA:542:C:H2' | 31:DA:543:C:OP1 | 2.05 | 0.56 |
| 31:DA:615:G:OP1 | 35:DF:40:GLN:NE2 | 2.38 | 0.56 |
| 31:DA:908:C:O2' | 31:DA:909:A:H5' | 2.05 | 0.56 |
| 31:DA:960:A:H61 | 42:DQ:82:ARG:HH21 | 1.52 | 0.56 |
| 31:DA:1131:G:OP2 | 31:DA:2515:C:H4' | 2.04 | 0.56 |
| 31:DA:2418:A:H2' | 31:DA:2419:U:C6 | 2.40 | 0.56 |
| 37:DH:32:GLU:O | 37:DH:33:LEU:HD23 | 2.05 | 0.56 |
| 41:DP:88:LEU:C | 41:DP:90:ARG:N | 2.60 | 0.56 |
| 47:DV:83:ARG:CG | 47:DV:83:ARG:NH1 | 2.62 | 0.56 |
| 49:DX:58:HIS:O | 49:DX:59:VAL:HG13 | 2.05 | 0.56 |
| 1:AA:509:A:O2' | 1:AA:510:A:P | 2.64 | 0.56 |
| 1:AA:629:G:H2' | 1:AA:630:G:O4' | 2.05 | 0.56 |
| 3:AC:150:LYS:HB3 | 3:AC:201:TYR:HB2 | 1.86 | 0.56 |
| 6:AF:62:TRP:O | 6:AF:62:TRP:CE3 | 2.57 | 0.56 |
| 8:AH:20:TYR:HD1 | 8:AH:65:TYR:HD2 | 1.52 | 0.56 |
| 20:AT:99:LEU:C | 20:AT:100:ILE:HD12 | 2.30 | 0.56 |
| 28:B6:15:GLU:CD | 28:B6:41:PRO:HG3 | 2.30 | 0.56 |
| 29:B7:19:ARG:HH11 | 29:B7:19:ARG:HG2 | 1.70 | 0.56 |
| 31:BA:234:C:H2' | 31:BA:235:U:C6 | 2.40 | 0.56 |
| 31:BA:1209:G:H21 | 31:BA:1210:A:N6 | 2.03 | 0.56 |
| 31:BA:2101:G:C6 | 31:BA:2102:U:C5 | 2.93 | 0.56 |
| 31:BA:2639:A:C2' | 31:BA:2640:G:H5' | 2.36 | 0.56 |
| 31:BA:2880:C:O2' | 43:BR:90:ARG:HD3 | 2.04 | 0.56 |
| 32:BB:116:G:H8 | 32:BB:116:G:H5" | 1.68 | 0.56 |
| 33:BD:35:LYS:CE | 33:BD:65:ILE:HG22 | 2.35 | 0.56 |
| 33:BD:183:ARG:CG | 33:BD:183:ARG:NH1 | 2.57 | 0.56 |
| 33:BD:186:HIS:CD2 | 33:BD:188:GLU:HB2 | 2.39 | 0.56 |
| 35:BF:65:TRP:CZ3 | 35:BF:75:HIS:HD2 | 2.22 | 0.56 |
| 35:BF:74:ARG:HG2 | 35:BF:74:ARG:O | 2.05 | 0.56 |
| 37:BH:70:THR:O | 37:BH:71:LEU:C | 2.48 | 0.56 |
| 38:BI:29:TYR:C | 38:BI:32:PRO:HD2 | 2.30 | 0.56 |
| 40:BO:112:MET:HE2 | 40:BO:112:MET:HA | 1.86 | 0.56 |
| 41:BP:141:ALA:HB3 | 25:D3:1:MET:CE | 2.35 | 0.56 |
| 44:BS:12:PHE:CD1 | 44:BS:12:PHE:O | 2.58 | 0.56 |
| 44:BS:26:LEU:O | 44:BS:88:ASP:HB3 | 2.06 | 0.56 |
| 44:BS:29:PHE:H | 44:BS:89:ARG:CD | 2.19 | 0.56 |
| 45:BT:104:ASN:O | 45:BT:105:LEU:HG | 2.05 | 0.56 |
| 46:BU:27:LEU:HB3 | 46:BU:31:SER:HB3 | 1.86 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 47:BV:53:GLU:O | 47:BV:55:ALA:N | 2.38 | 0.56 |
| 47:BV:69:LYS:HB3 | 47:BV:93:GLU:CD | 2.29 | 0.56 |
| 48:BW:64:MET:HE2 | 48:BW:69:LEU:HD23 | 1.87 | 0.56 |
| 1:CA:60:A:H4' | 1:CA:61:G:O5' | 2.05 | 0.56 |
| 1:CA:457:C:H2' | 1:CA:458:C:H6 | 1.69 | 0.56 |
| 1:CA:594:G:H1 | 1:CA:645:C:H42 | 1.53 | 0.56 |
| 1:CA:687:A:H1' | 1:CA:688:G:OP2 | 2.05 | 0.56 |
| 1:CA:706:A:C8 | 1:CA:707:C:H5 | 2.23 | 0.56 |
| 1:CA:1452:C:H5' | 1:CA:1456:G:C4 | 2.40 | 0.56 |
| 2:CB:22:LYS:NZ | 2:CB:22:LYS:HA | 2.20 | 0.56 |
| 2:CB:106:LYS:O | 2:CB:110:GLN:HG3 | 2.05 | 0.56 |
| 2:CB:201:ILE:HG21 | 2:CB:214:ILE:HG21 | 1.87 | 0.56 |
| 5:CE:107:ARG:O | 5:CE:108:ALA:C | 2.48 | 0.56 |
| 7:CG:32:ARG:O | 7:CG:33:ASP:CB | 2.54 | 0.56 |
| 15:CO:37:ASN:N | 15:CO:37:ASN:HD22 | 2.04 | 0.56 |
| 16:CP:48:TRP:H | 16:CP:48:TRP:HD1 | 1.52 | 0.56 |
| 20:CT:56:MET:HG2 | 20:CT:84:LEU:HD11 | 1.87 | 0.56 |
| 28:D6:18:ARG:CG | 28:D6:19:ARG:N | 2.66 | 0.56 |
| 29:D7:10:ARG:NH2 | 31:DA:1378:A:OP1 | 2.38 | 0.56 |
| 30:D8:31:HIS:O | 30:D8:33:ASN:N | 2.38 | 0.56 |
| 31:DA:192:C:H2' | 31:DA:193:U:H5' | 1.87 | 0.56 |
| 31:DA:848:G:N9 | 31:DA:933:A:H8 | 2.03 | 0.56 |
| 31:DA:1049:C:H1' | 31:DA:1113:U:O2' | 2.04 | 0.56 |
| 31:DA:1198:U:H2' | 31:DA:1199:U:C6 | 2.40 | 0.56 |
| 31:DA:1352:U:O2' | 31:DA:1353:A:H5' | 2.05 | 0.56 |
| 31:DA:1372:U:H2' | 31:DA:1373:A:O4' | 2.05 | 0.56 |
| 31:DA:1374:G:C5 | 31:DA:1375:C:C4 | 2.93 | 0.56 |
| 31:DA:2099:U:O2 | 31:DA:2099:U:H2' | 2.05 | 0.56 |
| 31:DA:2199:A:OP2 | 31:DA:2200:C:H5 | 1.88 | 0.56 |
| 31:DA:2395:C:H6 | 31:DA:2395:C:C5' | 2.18 | 0.56 |
| 31:DA:2619:C:O2' | 31:DA:2620:C:H5' | 2.05 | 0.56 |
| 31:DA:2884:U:C6 | 31:DA:2885:C:C6 | 2.93 | 0.56 |
| 34:DE:14:ILE:CG1 | 34:DE:21:VAL:CG2 | 2.83 | 0.56 |
| 34:DE:76:ARG:O | 34:DE:77:ILE:HG22 | 2.05 | 0.56 |
| 34:DE:109:LYS:HD3 | 43:DR:2:ARG:NH1 | 2.20 | 0.56 |
| 34:DE:112:GLY:O | 34:DE:159:HIS:HA | 2.04 | 0.56 |
| 39:DN:62:VAL:HG22 | 39:DN:66:LYS:HG3 | 1.87 | 0.56 |
| 43:DR:5:LYS:H | 43:DR:5:LYS:CD | 2.17 | 0.56 |
| 43:DR:24:GLN:HE22 | 43:DR:36:THR:CG2 | 2.09 | 0.56 |
| 44:DS:58:LEU:HD21 | 44:DS:68:GLN:HB2 | 1.88 | 0.56 |
| 47:DV:82:ARG:HG3 | 47:DV:82:ARG:NH1 | 2.12 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 48:DW:28:SER:OG | 48:DW:31:GLU:HB2 | 2.05 | 0.56 |
| 49:DX:85:PRO:O | 49:DX:86:GLY:C | 2.47 | 0.56 |
| 50:DY:37:VAL:O | 50:DY:38:ILE:HG12 | 2.04 | 0.56 |
| 1:AA:32:A:H2' | 1:AA:33:A:C8 | 2.40 | 0.56 |
| 1:AA:227:G:O2' | 1:AA:228:A:H5' | 2.06 | 0.56 |
| 1:AA:1423:G:H5' | 40:BO:49:ARG:HH22 | 1.70 | 0.56 |
| 1:AA:1495:U:O2' | 31:BA:1919:A:N1 | 2.38 | 0.56 |
| 8:AH:48:TYR:HA | 8:AH:60:ARG:O | 2.05 | 0.56 |
| 16:AP:48:TRP:H | 16:AP:48:TRP:HD1 | 1.53 | 0.56 |
| 30:B8:38:GLY:C | 30:B8:40:GLU:H | 2.14 | 0.56 |
| 31:BA:541:C:H2' | 31:BA:542:C:C6 | 2.40 | 0.56 |
| 31:BA:2061:G:C2 | 31:BA:2063:C:C4 | 2.94 | 0.56 |
| 31:BA:2689:U:H5" | 31:BA:2690:C:H5' | 1.87 | 0.56 |
| 31:BA:2759:G:H5' | 31:BA:2759:G:H8 | 1.70 | 0.56 |
| 31:BA:2773:C:O2' | 31:BA:2774:C:H5' | 2.05 | 0.56 |
| 31:BA:2894:G:N3 | 31:BA:2894:G:H2' | 2.20 | 0.56 |
| 33:BD:16:MET:HE3 | 33:BD:206:LEU:O | 2.05 | 0.56 |
| 36:BG:18:GLU:HG2 | 36:BG:175:LEU:HD21 | 1.88 | 0.56 |
| 36:BG:174:GLU:HG3 | 36:BG:180:PHE:HD1 | 1.71 | 0.56 |
| 39:BN:39:ARG:HE | 39:BN:41:ASP:CB | 2.18 | 0.56 |
| 42:BQ:141:GLN:HG3 | 51:BZ:72:ARG:HH11 | 1.70 | 0.56 |
| 1:CA:561:U:O2' | 1:CA:562:C:P | 2.63 | 0.56 |
| 1:CA:1012:U:O5' | 1:CA:1012:U:H6 | 1.87 | 0.56 |
| 3:CC:23:TYR:CG | 3:CC:24:ALA:N | 2.74 | 0.56 |
| 3:CC:34:LEU:O | 3:CC:34:LEU:HD23 | 2.05 | 0.56 |
| 5:CE:50:GLU:OE2 | 5:CE:51:VAL:HG23 | 2.05 | 0.56 |
| 18:CR:72:ARG:O | 18:CR:76:LEU:HD23 | 2.05 | 0.56 |
| 23:D1:13:ILE:O | 23:D1:14:VAL:HB | 2.05 | 0.56 |
| 28:D6:15:GLU:CD | 28:D6:18:ARG:CD | 2.78 | 0.56 |
| 31:DA:185:U:H4' | 31:DA:218:A:H4' | 1.87 | 0.56 |
| 31:DA:203:C:H3' | 31:DA:204:A:H5" | 1.87 | 0.56 |
| 31:DA:1654:A:OP2 | 43:DR:3:HIS:HB2 | 2.06 | 0.56 |
| 31:DA:1679:U:H2' | 31:DA:1680:U:H5' | 1.87 | 0.56 |
| 31:DA:2661:G:C8 | 31:DA:2662:A:N3 | 2.74 | 0.56 |
| 31:DA:2773:C:O2' | 31:DA:2774:C:H5' | 2.05 | 0.56 |
| 32:DB:71:C:O2 | 32:DB:71:C:H2' | 2.05 | 0.56 |
| 33:DD:97:TYR:HB2 | 33:DD:101:GLU:O | 2.05 | 0.56 |
| 36:DG:103:LEU:HD23 | 36:DG:106:LEU:HD23 | 1.88 | 0.56 |
| 37:DH:163:TYR:N | 37:DH:163:TYR:CD1 | 2.74 | 0.56 |
| 41:DP:50:ARG:HG2 | 41:DP:50:ARG:HH21 | 1.70 | 0.56 |
| 44:DS:16:ASN:ND2 | 44:DS:92:TYR:CZ | 2.73 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 49:DX:12:VAL:HG13 | 49:DX:17:ALA:HB2 | 1.88 | 0.56 |
| 1:AA:1004:A:H2' | 1:AA:1038:C:O2 | 2.05 | 0.56 |
| 24:B2:55:ARG:H | 24:B2:56:GLN:NE2 | 2.03 | 0.56 |
| 31:BA:218:A:C2 | 31:BA:235:U:H4' | 2.41 | 0.56 |
| 31:BA:746:A:C2 | 55:BA:3351:ZIT:C16 | 2.84 | 0.56 |
| 31:BA:1185:C:H5" | 31:BA:1186:G:OP1 | 2.05 | 0.56 |
| 31:BA:1404:C:O2 | 31:BA:1404:C:H2' | 2.05 | 0.56 |
| 31:BA:1813:G:H2' | 31:BA:1814:G:H5' | 1.88 | 0.56 |
| 32:BB:2:C:H2' | 32:BB:3:C:H6 | 1.69 | 0.56 |
| 34:BE:66:HIS:O | 34:BE:66:HIS:CG | 2.59 | 0.56 |
| 34:BE:195:LEU:C | 34:BE:195:LEU:HD23 | 2.30 | 0.56 |
| 36:BG:115:ARG:NH1 | 36:BG:136:ARG:HG3 | 2.20 | 0.56 |
| 40:BO:31:LYS:HB3 | 40:BO:32:TYR:CE1 | 2.41 | 0.56 |
| 44:BS:54:LEU:HA | 44:BS:57:LYS:O | 2.05 | 0.56 |
| 46:BU:27:LEU:HD23 | 46:BU:27:LEU:N | 2.21 | 0.56 |
| 1:CA:650:G:O2' | 1:CA:651:C:H5' | 2.05 | 0.56 |
| 6:CF:23:LYS:O | 6:CF:27:GLN:HG2 | 2.06 | 0.56 |
| 10:CJ:51:ARG:HE | 10:CJ:61:GLU:HB2 | 1.70 | 0.56 |
| 16:CP:39:TYR:HB2 | 16:CP:49:LEU:HD12 | 1.88 | 0.56 |
| 20:CT:24:LEU:C | 20:CT:24:LEU:HD13 | 2.31 | 0.56 |
| 22:D0:41:ARG:HB2 | 31:DA:2330:G:O2' | 2.05 | 0.56 |
| 26:D4:14:ILE:HA | 36:DG:5:VAL:HG13 | 1.86 | 0.56 |
| 31:DA:68:G:H2' | 31:DA:69:C:C6 | 2.40 | 0.56 |
| 31:DA:528:A:H2 | 31:DA:2043:C:H5' | 1.69 | 0.56 |
| 31:DA:614:U:O2 | 31:DA:614:U:O5' | 2.24 | 0.56 |
| 31:DA:1040:C:O2' | 31:DA:1041:C:OP2 | 2.23 | 0.56 |
| 31:DA:1531:C:H3' | 31:DA:1532:C:C5' | 2.35 | 0.56 |
| 31:DA:1742:G:H3' | 31:DA:1742:G:H8 | 1.70 | 0.56 |
| 31:DA:1925:C:C2' | 31:DA:1926:U:H5' | 2.36 | 0.56 |
| 33:DD:86:PRO:HG2 | 33:DD:87:ASN:HD21 | 1.70 | 0.56 |
| 36:DG:47:LYS:HD3 | 36:DG:81:LYS:CD | 2.34 | 0.56 |
| 38:DI:29:TYR:C | 38:DI:32:PRO:HD2 | 2.31 | 0.56 |
| 38:DI:56:LYS:HZ2 | 38:DI:57:ARG:N | 2.03 | 0.56 |
| 38:DI:82:ARG:O | 38:DI:89:TYR:HB2 | 2.04 | 0.56 |
| 41:DP:50:ARG:HG2 | 41:DP:50:ARG:NH2 | 2.19 | 0.56 |
| 44:DS:29:PHE:H | 44:DS:89:ARG:HD2 | 1.65 | 0.56 |
| 45:DT:51:ARG:HG3 | 45:DT:98:LYS:HG3 | 1.87 | 0.56 |
| 45:DT:101:PHE:HE2 | 45:DT:113:LYS:HD2 | 1.70 | 0.56 |
| 1:AA:134:A:H61 | 16:AP:25:ARG:HH12 | 1.53 | 0.56 |
| 1:AA:611:A:H61 | 1:AA:629:G:H1 | 1.52 | 0.56 |
| 1:AA:1274:G:H22 | 1:AA:1275:A:H62 | 1.54 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 2:AB:28:PHE:CD1 | 2:AB:190:THR:HG22 | 2.41 | 0.56 |
| 24:B2:31:GLU:HG2 | 24:B2:37:PHE:CD1 | 2.40 | 0.56 |
| 30:B8:29:LYS:O | 30:B8:30:ARG:C | 2.48 | 0.56 |
| 30:B8:35:GLN:OE1 | 31:BA:2421:G:OP2 | 2.24 | 0.56 |
| 31:BA:8:A:H2' | 31:BA:9:U:C5 | 2.41 | 0.56 |
| 31:BA:534:U:O2' | 46:BU:49:HIS:HD2 | 1.88 | 0.56 |
| 31:BA:557:U:O2 | 39:BN:45:ASN:HB2 | 2.05 | 0.56 |
| 31:BA:814:C:C5' | 47:BV:86:GLY:HA3 | 2.36 | 0.56 |
| 31:BA:2478:A:C2' | 31:BA:2479:G:H5' | 2.35 | 0.56 |
| 32:BB:116:G:C8 | 32:BB:116:G:H5' | 2.41 | 0.56 |
| 33:BD:35:LYS:HZ2 | 33:BD:64:ILE:C | 2.09 | 0.56 |
| 33:BD:85:ASP:HB2 | 33:BD:92:ILE:HG12 | 1.88 | 0.56 |
| 47:BV:66:ARG:HD2 | 47:BV:67:GLY:N | 2.20 | 0.56 |
| 50:BY:2:ARG:C | 50:BY:4:LYS:N | 2.63 | 0.56 |
| 51:BZ:29:TYR:CE2 | 51:BZ:87:ASP:HB2 | 2.39 | 0.56 |
| 51:BZ:150:LEU:HD13 | 51:BZ:150:LEU:H | 1.70 | 0.56 |
| 1:CA:542:G:H2' | 1:CA:543:C:C6 | 2.40 | 0.56 |
| 1:CA:1446:U:HO2' | 1:CA:1447:A:H8 | 1.44 | 0.56 |
| 3:CC:113:ALA:HB3 | 3:CC:114:PRO:HD3 | 1.88 | 0.56 |
| 6:CF:5:GLU:HG2 | 6:CF:62:TRP:CZ2 | 2.40 | 0.56 |
| 9:CI:4:TYR:HA | 9:CI:88:TYR:CE1 | 2.41 | 0.56 |
| 22:D0:11:ARG:O | 22:D0:14:ARG:NH2 | 2.39 | 0.56 |
| 24:D2:46:GLN:C | 24:D2:48:HIS:H | 2.12 | 0.56 |
| 31:DA:28:A:C5 | 31:DA:29:U:C5 | 2.93 | 0.56 |
| 31:DA:173:G:C6 | 31:DA:174:C:C4 | 2.94 | 0.56 |
| 31:DA:471:A:H8 | 31:DA:471:A:O5' | 1.88 | 0.56 |
| 31:DA:494:G:H5" | 31:DA:494:G:H8 | 1.69 | 0.56 |
| 31:DA:557:U:O2 | 39:DN:45:ASN:HB2 | 2.05 | 0.56 |
| 31:DA:708:C:H2' | 31:DA:708:C:O2 | 2.05 | 0.56 |
| 31:DA:1000:A:C6 | 31:DA:1001:A:C6 | 2.93 | 0.56 |
| 31:DA:1115:G:H2' | 31:DA:1116:C:H5" | 1.88 | 0.56 |
| 31:DA:1648:C:H2' | 31:DA:1649:G:O5' | 2.05 | 0.56 |
| 31:DA:1695:G:N2 | 31:DA:1696:G:C8 | 2.74 | 0.56 |
| 31:DA:1973:G:H2' | 31:DA:1974:C:H6 | 1.70 | 0.56 |
| 31:DA:2788:C:O2' | 31:DA:2809:A:N3 | 2.36 | 0.56 |
| 33:DD:8:PRO:CB | 33:DD:14:ARG:HB3 | 2.35 | 0.56 |
| 34:DE:103:ASP:OD2 | 34:DE:168:MET:HE2 | 2.05 | 0.56 |
| 34:DE:195:LEU:HD23 | 34:DE:195:LEU:C | 2.30 | 0.56 |
| 35:DF:205:ARG:C | 35:DF:206:ILE:HG13 | 2.30 | 0.56 |
| 39:DN:13:TRP:CH2 | 39:DN:130:HIS:HE1 | 2.23 | 0.56 |
| 39:DN:68:GLU:HA | 39:DN:86:PRO:CB | 2.36 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 41:DP:10:PRO:HD2 | 41:DP:11:GLY:H | 1.70 | 0.56 |
| 42:DQ:54:MET:HB3 | 42:DQ:64:ILE:CD1 | 2.36 | 0.56 |
| 43:DR:10:LEU:HB3 | 43:DR:17:ARG:CD | 2.35 | 0.56 |
| 44:DS:28:VAL:HG11 | 44:DS:97:ARG:NH2 | 2.21 | 0.56 |
| 46:DU:64:ARG:NH2 | 46:DU:64:ARG:CA | 2.58 | 0.56 |
| 47:DV:62:LEU:HB3 | 47:DV:98:GLU:HB2 | 1.87 | 0.56 |
| 47:DV:66:ARG:CD | 47:DV:67:GLY:N | 2.69 | 0.56 |
| 50:DY:14:LEU:HG | 50:DY:15:VAL:N | 2.21 | 0.56 |
| 50:DY:42:VAL:HB | 50:DY:65:ALA:HB3 | 1.87 | 0.56 |
| 50:DY:77:PRO:O | 50:DY:78:ALA:HB2 | 2.06 | 0.56 |
| 1:AA:84:U:H5 | 1:AA:88:A:C8 | 2.23 | 0.56 |
| 1:AA:191:G:H1' | 20:AT:105:SER:HA | 1.87 | 0.56 |
| 1:AA:525:C:H2' | 1:AA:526:C:C6 | 2.40 | 0.56 |
| 1:AA:707:C:OP1 | 11:AK:85:ARG:NH1 | 2.39 | 0.56 |
| 1:AA:949:A:H1' | 1:AA:1364:U:N3 | 2.21 | 0.56 |
| 16:AP:18:ARG:HD3 | 16:AP:35:LYS:HD2 | 1.88 | 0.56 |
| 20:AT:56:MET:HG2 | 20:AT:84:LEU:HD11 | 1.88 | 0.56 |
| 31:BA:142:A:H1' | 31:BA:1408:C:O4' | 2.06 | 0.56 |
| 31:BA:1324:G:C4 | 31:BA:1328:G:O6 | 2.59 | 0.56 |
| 31:BA:2593:U:H2' | 31:BA:2594:C:H6 | 1.70 | 0.56 |
| 39:BN:3:THR:CA | 39:BN:4:TYR:CE1 | 2.88 | 0.56 |
| 41:BP:55:ARG:O | 41:BP:56:SER:OG | 2.19 | 0.56 |
| 42:BQ:8:LYS:CG | 42:BQ:9:TYR:N | 2.69 | 0.56 |
| 48:BW:13:SER:HB3 | 48:BW:16:LYS:HD3 | 1.88 | 0.56 |
| 48:BW:64:MET:O | 48:BW:65:LEU:CB | 2.46 | 0.56 |
| 51:BZ:149:SER:CB | 51:BZ:173:ALA:HA | 2.36 | 0.56 |
| 1:CA:521:G:H4' | 12:CL:73:GLU:HG2 | 1.88 | 0.56 |
| 1:CA:1362:C:O2' | 1:CA:1363:C:H5" | 2.06 | 0.56 |
| 2:CB:213:LEU:O | 2:CB:213:LEU:HD23 | 2.05 | 0.56 |
| 6:CF:67:MET:HE1 | 6:CF:75:LEU:HD12 | 1.88 | 0.56 |
| 12:CL:21:LYS:HD2 | 12:CL:21:LYS:H | 1.69 | 0.56 |
| 13:CM:66:LEU:HD12 | 13:CM:66:LEU:N | 2.18 | 0.56 |
| 24:D2:47:ASN:HA | 24:D2:50:ILE:O | 2.06 | 0.56 |
| 27:D5:36:CYS:SG | 27:D5:49:CYS:CB | 2.93 | 0.56 |
| 28:D6:51:GLU:O | 28:D6:52:VAL:CB | 2.54 | 0.56 |
| 31:DA:751:A:H5' | 48:DW:90:ARG:HA | 1.87 | 0.56 |
| 31:DA:794:G:H2' | 31:DA:795:C:C6 | 2.41 | 0.56 |
| 31:DA:1496:A:C8 | 31:DA:1577:C:O2' | 2.59 | 0.56 |
| 31:DA:1508:A:C2' | 31:DA:1509:C:OP1 | 2.53 | 0.56 |
| 31:DA:1987:G:H8 | 31:DA:1987:G:C5' | 2.19 | 0.56 |
| 31:DA:2061:G:N2 | 31:DA:2063:C:C2 | 2.73 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 32:DB:21:G:O2' | 32:DB:22:U:O5' | 2.24 | 0.56 |
| 34:DE:70:ALA:O | 34:DE:71:GLY:C | 2.48 | 0.56 |
| 41:DP:79:ARG:HH21 | 41:DP:109:GLY:CA | 2.19 | 0.56 |
| 44:DS:36:TYR:N | 44:DS:36:TYR:CD1 | 2.74 | 0.56 |
| 47:DV:25:LEU:C | 47:DV:27:ALA:H | 2.13 | 0.56 |
| 51:DZ:44:PHE:C | 51:DZ:44:PHE:CD1 | 2.83 | 0.56 |
| 1:AA:628:G:O2' | 1:AA:629:G:H5' | 2.05 | 0.56 |
| 2:AB:114:ARG:HD3 | 2:AB:114:ARG:O | 2.06 | 0.56 |
| 3:AC:6:HIS:HE2 | 3:AC:184:TYR:HE2 | 1.52 | 0.56 |
| 3:AC:53:ALA:O | 3:AC:54:ARG:HB2 | 2.05 | 0.56 |
| 24:B2:14:ARG:NH1 | 24:B2:57:ILE:HG22 | 2.21 | 0.56 |
| 24:B2:14:ARG:CZ | 24:B2:57:ILE:CG2 | 2.83 | 0.56 |
| 30:B8:61:LEU:C | 30:B8:63:PRO:HD2 | 2.30 | 0.56 |
| 31:BA:661:C:H4' | 41:BP:18:ARG:HG2 | 1.86 | 0.56 |
| 31:BA:836:G:C5 | 31:BA:837:C:C4 | 2.94 | 0.56 |
| 31:BA:892:G:C5 | 31:BA:893:C:C5 | 2.94 | 0.56 |
| 31:BA:1686:C:C2' | 31:BA:1687:G:H5' | 2.35 | 0.56 |
| 31:BA:1686:C:H2' | 31:BA:1687:G:H5' | 1.88 | 0.56 |
| 31:BA:2849:U:O4 | 45:BT:23:ARG:NH2 | 2.39 | 0.56 |
| 32:BB:7:G:C2' | 32:BB:8:U:H5" | 2.36 | 0.56 |
| 32:BB:15:A:O2' | 32:BB:110:G:C8 | 2.59 | 0.56 |
| 32:BB:86:G:H8 | 32:BB:86:G:O5' | 1.88 | 0.56 |
| 33:BD:35:LYS:HE3 | 33:BD:63:ARG:C | 2.30 | 0.56 |
| 33:BD:86:PRO:HG2 | 33:BD:87:ASN:ND2 | 2.21 | 0.56 |
| 34:BE:10:GLY:HA3 | 45:BT:8:LYS:HE3 | 1.86 | 0.56 |
| 35:BF:139:PHE:HB2 | 35:BF:166:ALA:HB1 | 1.86 | 0.56 |
| 37:BH:85:LYS:HZ1 | 37:BH:145:ALA:HA | 1.67 | 0.56 |
| 39:BN:3:THR:CA | 39:BN:4:TYR:CD1 | 2.88 | 0.56 |
| 39:BN:57:ALA:O | 39:BN:58:ASP:O | 2.24 | 0.56 |
| 41:BP:16:ARG:C | 41:BP:16:ARG:HH11 | 2.13 | 0.56 |
| 44:BS:61:ASN:ND2 | 44:BS:64:GLU:OE2 | 2.39 | 0.56 |
| 49:BX:83:VAL:O | 49:BX:84:ALA:CB | 2.53 | 0.56 |
| 1:CA:963:G:N2 | 10:CJ:55:LYS:HE2 | 2.21 | 0.56 |
| 4:CD:109:GLY:O | 4:CD:111:ALA:N | 2.39 | 0.56 |
| 10:CJ:40:LEU:HD21 | 10:CJ:69:ASN:HB3 | 1.86 | 0.56 |
| 11:CK:99:GLN:O | 11:CK:101:SER:N | 2.38 | 0.56 |
| 13:CM:46:LYS:HG3 | 13:CM:47:ASP:N | 2.21 | 0.56 |
| 20:CT:99:LEU:C | 20:CT:100:ILE:HD12 | 2.30 | 0.56 |
| 23:D1:60:PHE:HZ | 23:D1:90:ILE:HG21 | 1.71 | 0.56 |
| 23:D1:62:VAL:HG22 | 23:D1:63:ALA:N | 2.20 | 0.56 |
| 30:D8:31:HIS:O | 30:D8:32:LEU:C | 2.49 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:303:U:H2' | 31:DA:304:G:H8 | 1.70 | 0.56 |
| 31:DA:858:U:O2 | 31:DA:2268:A:H2' | 2.05 | 0.56 |
| 31:DA:1210:A:C5' | 31:DA:1211:U:H3' | 2.35 | 0.56 |
| 31:DA:1317:A:H2' | 31:DA:1318:C:C6 | 2.39 | 0.56 |
| 31:DA:1822:G:H8 | 31:DA:1822:G:H5' | 1.69 | 0.56 |
| 34:DE:197:ILE:HD11 | 34:DE:199:ARG:CZ | 2.35 | 0.56 |
| 38:DI:38:LEU:HB2 | 38:DI:40:THR:HG23 | 1.87 | 0.56 |
| 42:DQ:139:GLU:HG2 | 42:DQ:139:GLU:O | 2.06 | 0.56 |
| 46:DU:25:TRP:HD1 | 46:DU:26:GLY:N | 2.03 | 0.56 |
| 1:AA:355:C:C2' | 1:AA:356:A:H5' | 2.36 | 0.56 |
| 1:AA:674:G:H21 | 11:AK:116:HIS:HB2 | 1.71 | 0.56 |
| 1:AA:1168:A:C6 | 1:AA:1169:A:C6 | 2.94 | 0.56 |
| 1:AA:1216:G:OP1 | 14:AN:2:ALA:HA | 2.06 | 0.56 |
| 3:AC:105:GLU:HG2 | 3:AC:106:VAL:H | 1.69 | 0.56 |
| 3:AC:188:LEU:O | 3:AC:189:ALA:HB2 | 2.06 | 0.56 |
| 4:AD:49:ARG:HA | 4:AD:49:ARG:HE | 1.70 | 0.56 |
| 23:B1:10:LYS:O | 23:B1:13:ILE:CG2 | 2.54 | 0.56 |
| 23:B1:86:SER:N | 23:B1:87:PRO:CD | 2.65 | 0.56 |
| 31:BA:71:A:C2 | 49:BX:31:HIS:CE1 | 2.79 | 0.56 |
| 31:BA:352:G:O2' | 31:BA:353:G:O5' | 2.23 | 0.56 |
| 31:BA:511:U:H3' | 31:BA:512:G:C5' | 2.23 | 0.56 |
| 31:BA:620:G:H4' | 31:BA:621:A:H5" | 1.88 | 0.56 |
| 31:BA:858:U:O2 | 31:BA:2268:A:H2' | 2.06 | 0.56 |
| 31:BA:911:A:C2' | 42:BQ:9:TYR:OH | 2.53 | 0.56 |
| 31:BA:952:G:C6 | 31:BA:953:A:N7 | 2.73 | 0.56 |
| 31:BA:996:A:C4' | 46:BU:92:ARG:NE | 2.60 | 0.56 |
| 31:BA:1359:A:C8 | 31:BA:1372:U:O4 | 2.59 | 0.56 |
| 31:BA:1472:A:C2' | 31:BA:1473:G:H5' | 2.36 | 0.56 |
| 31:BA:1494:A:H2' | 31:BA:1494:A:N3 | 2.21 | 0.56 |
| 31:BA:1648:C:H2' | 31:BA:1649:G:O5' | 2.06 | 0.56 |
| 31:BA:2312:U:O3' | 36:BG:71:THR:HG21 | 2.05 | 0.56 |
| 33:BD:35:LYS:NZ | 33:BD:104:TYR:HD1 | 2.04 | 0.56 |
| 38:BI:110:ASP:C | 38:BI:112:LYS:N | 2.60 | 0.56 |
| 45:BT:83:ILE:HG13 | 45:BT:84:GLN:N | 2.20 | 0.56 |
| 47:BV:62:LEU:HD22 | 47:BV:98:GLU:CB | 2.32 | 0.56 |
| 47:BV:64:HIS:CD2 | 47:BV:94:LEU:HD21 | 2.35 | 0.56 |
| 50:BY:89:PHE:O | 50:BY:90:LEU:HB3 | 2.06 | 0.56 |
| 1:CA:495:A:H4' | 1:CA:496:A:OP1 | 2.06 | 0.56 |
| 4:CD:33:MET:CE | 4:CD:37:PRO:HA | 2.36 | 0.56 |
| 5:CE:43:LEU:HB2 | 5:CE:136:MET:SD | 2.46 | 0.56 |
| 5:CE:112:LEU:HD23 | 5:CE:112:LEU:N | 2.21 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 10:CJ:48:THR:HG23 | 10:CJ:62:HIS:HB3 | 1.87 | 0.56 |
| 17:CQ:67:LYS:HA | 17:CQ:70:ARG:HH12 | 1.71 | 0.56 |
| 23:D1:27:GLU:OE2 | 23:D1:32:LYS:HB2 | 2.06 | 0.56 |
| 31:DA:38:A:H2' | 31:DA:39:C:C6 | 2.40 | 0.56 |
| 31:DA:195:A:H61 | 31:DA:198:C:H3' | 1.71 | 0.56 |
| 31:DA:746:A:C2 | 55:DA:3311:ZIT:C16 | 2.82 | 0.56 |
| 31:DA:774:A:H2 | 31:DA:787:U:HO2' | 1.49 | 0.56 |
| 31:DA:1386:C:H2' | 31:DA:1387:C:H6 | 1.69 | 0.56 |
| 31:DA:1495:A:C4 | 31:DA:1496:A:C2 | 2.94 | 0.56 |
| 31:DA:2191:G:H2' | 31:DA:2192:G:O5' | 2.06 | 0.56 |
| 31:DA:2243:U:O2' | 31:DA:2244:U:H5' | 2.06 | 0.56 |
| 31:DA:2312:U:O3' | 36:DG:71:THR:HG21 | 2.06 | 0.56 |
| 31:DA:2343:C:HO2' | 31:DA:2373:G:HO2' | 1.49 | 0.56 |
| 32:DB:8:U:H5' | 32:DB:8:U:C6 | 2.37 | 0.56 |
| 35:DF:154:VAL:HB | 35:DF:173:VAL:HG22 | 1.88 | 0.56 |
| 37:DH:138:LYS:O | 37:DH:139:GLN:C | 2.49 | 0.56 |
| 38:DI:83:ALA:HA | 38:DI:89:TYR:HD1 | 1.68 | 0.56 |
| 41:DP:24:GLY:HA2 | 41:DP:33:ARG:HE | 1.71 | 0.56 |
| 47:DV:18:LEU:HD22 | 47:DV:19:LYS:N | 2.21 | 0.56 |
| 47:DV:19:LYS:CE | 47:DV:20:LEU:H | 2.18 | 0.56 |
| 49:DX:64:LYS:O | 49:DX:65:ARG:HB2 | 2.06 | 0.56 |
| 1:AA:457:C:H2' | 1:AA:458:C:C6 | 2.41 | 0.56 |
| 1:AA:629:G:C4 | 1:AA:630:G:C8 | 2.94 | 0.56 |
| 1:AA:658:G:C6 | 1:AA:749:C:N4 | 2.74 | 0.56 |
| 1:AA:665:A:H1' | 1:AA:733:A:O4' | 2.06 | 0.56 |
| 1:AA:995:C:H1' | 14:AN:8:GLU:OE2 | 2.06 | 0.56 |
| 1:AA:1116:C:H3' | 1:AA:1117:G:H5" | 1.88 | 0.56 |
| 1:AA:1438:G:H2' | 1:AA:1439:C:C6 | 2.41 | 0.56 |
| 6:AF:61:LEU:HD23 | 6:AF:63:TYR:OH | 2.06 | 0.56 |
| 7:AG:23:VAL:O | 7:AG:27:ILE:HD12 | 2.05 | 0.56 |
| 8:AH:112:LEU:HB2 | 8:AH:133:LEU:HA | 1.88 | 0.56 |
| 10:AJ:32:ALA:HB1 | 10:AJ:75:ILE:HD11 | 1.88 | 0.56 |
| 16:AP:53:VAL:O | 16:AP:57:ARG:HG2 | 2.06 | 0.56 |
| 28:B6:40:CYS:SG | 28:B6:45:LYS:NZ | 2.64 | 0.56 |
| 31:BA:930:U:O4' | 31:BA:930:U:O2 | 2.20 | 0.56 |
| 31:BA:1110:G:OP1 | 31:BA:1110:G:H4' | 2.06 | 0.56 |
| 31:BA:1487:G:C2 | 31:BA:1488:G:C4 | 2.94 | 0.56 |
| 31:BA:2529:G:H5" | 31:BA:2530:A:H5" | 1.87 | 0.56 |
| 31:BA:2615:U:H2' | 31:BA:2616:C:H6 | 1.70 | 0.56 |
| 32:BB:7:G:O5' | 44:BS:29:PHE:CE1 | 2.59 | 0.56 |
| 33:BD:45:ASN:CG | 33:BD:46:GLN:N | 2.60 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 39:BN:28:THR:N | 39:BN:106:MET:HE3 | 2.21 | 0.56 |
| 40:BO:31:LYS:C | 40:BO:32:TYR:CD1 | 2.84 | 0.56 |
| 43:BR:56:LYS:HE2 | 43:BR:94:TYR:CE2 | 2.41 | 0.56 |
| 50:BY:37:VAL:HG11 | 50:BY:72:VAL:CG2 | 2.36 | 0.56 |
| 1:CA:41:G:H2' | 1:CA:42:G:C8 | 2.40 | 0.56 |
| 1:CA:134:A:H61 | 16:CP:25:ARG:HH12 | 1.54 | 0.56 |
| 1:CA:1410:G:C4 | 1:CA:1491:G:N2 | 2.74 | 0.56 |
| 2:CB:55:PHE:HE1 | 2:CB:218:ALA:HA | 1.71 | 0.56 |
| 2:CB:172:ILE:H | 2:CB:172:ILE:CD1 | 2.17 | 0.56 |
| 5:CE:100:VAL:HG13 | 5:CE:118:ILE:CG2 | 2.35 | 0.56 |
| 6:CF:98:LEU:H | 6:CF:98:LEU:HD12 | 1.70 | 0.56 |
| 9:CI:11:LYS:HG2 | 9:CI:11:LYS:O | 2.05 | 0.56 |
| 23:D1:11:ARG:CB | 23:D1:12:PRO:CD | 2.84 | 0.56 |
| 30:D8:39:LYS:CE | 30:D8:39:LYS:O | 2.53 | 0.56 |
| 31:DA:229:A:H3' | 31:DA:230:U:H5' | 1.87 | 0.56 |
| 31:DA:303:U:H2' | 31:DA:304:G:C8 | 2.41 | 0.56 |
| 31:DA:534:U:O2' | 46:DU:49:HIS:HD2 | 1.89 | 0.56 |
| 31:DA:848:G:N3 | 31:DA:933:A:H1' | 2.20 | 0.56 |
| 31:DA:911:A:C2' | 42:DQ:9:TYR:OH | 2.53 | 0.56 |
| 31:DA:1245:G:OP1 | 41:DP:16:ARG:HG2 | 2.04 | 0.56 |
| 31:DA:1580:A:OP2 | 31:DA:1580:A:H8 | 1.89 | 0.56 |
| 31:DA:1598:C:O2 | 31:DA:1598:C:H2' | 2.06 | 0.56 |
| 31:DA:2394:C:P | 41:DP:63:PRO:HD2 | 2.44 | 0.56 |
| 35:DF:22:ALA:CA | 35:DF:26:ALA:HB2 | 2.36 | 0.56 |
| 35:DF:160:ASN:HD22 | 35:DF:162:LEU:H | 1.53 | 0.56 |
| 38:DI:38:LEU:HB2 | 38:DI:40:THR:CG2 | 2.36 | 0.56 |
| 38:DI:88:ILE:HG22 | 38:DI:89:TYR:N | 2.21 | 0.56 |
| 41:DP:97:PRO:O | 41:DP:98:GLU:CB | 2.52 | 0.56 |
| 44:DS:99:LYS:O | 44:DS:101:LEU:HB3 | 2.06 | 0.56 |
| 51:DZ:54:HIS:HB3 | 51:DZ:101:PRO:HD3 | 1.88 | 0.56 |
| 51:DZ:77:ASP:O | 51:DZ:79:ARG:N | 2.39 | 0.56 |
| 1:AA:457:C:H2' | 1:AA:458:C:H6 | 1.71 | 0.55 |
| 1:AA:658:G:H1' | 15:AO:22:THR:HB | 1.89 | 0.55 |
| 4:AD:119:GLN:CG | 4:AD:123:HIS:CD2 | 2.87 | 0.55 |
| 6:AF:89:MET:SD | 18:AR:76:LEU:HD21 | 2.46 | 0.55 |
| 9:AI:4:TYR:HA | 9:AI:88:TYR:CE1 | 2.41 | 0.55 |
| 23:B1:13:ILE:O | 23:B1:14:VAL:HB | 2.06 | 0.55 |
| 31:BA:195:A:H4' | 31:BA:251:A:O2' | 2.05 | 0.55 |
| 31:BA:236:C:H2' | 31:BA:237:C:C6 | 2.41 | 0.55 |
| 31:BA:631:A:OP1 | 41:BP:64:LYS:CE | 2.54 | 0.55 |
| 31:BA:790:C:O2' | 31:BA:791:C:H5' | 2.07 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2265:U:H4' | 42:BQ:13:GLN:HE22 | 1.70 | 0.55 |
| 31:BA:2523:G:C2' | 31:BA:2524:G:C5' | 2.83 | 0.55 |
| 31:BA:2723:C:H5" | 43:BR:2:ARG:HD2 | 1.87 | 0.55 |
| 35:BF:1:MET:O | 35:BF:2:LYS:C | 2.48 | 0.55 |
| 38:BI:49:ALA:HA | 38:BI:52:ARG:HB2 | 1.88 | 0.55 |
| 39:BN:13:TRP:CZ3 | 39:BN:130:HIS:CE1 | 2.94 | 0.55 |
| 48:BW:4:LYS:HE3 | 48:BW:6:ILE:HD11 | 1.88 | 0.55 |
| 51:BZ:19:ARG:HH11 | 51:BZ:19:ARG:HG2 | 1.70 | 0.55 |
| 51:BZ:29:TYR:HE2 | 51:BZ:87:ASP:CB | 2.18 | 0.55 |
| 1:CA:308:C:H2' | 1:CA:309:G:H8 | 1.72 | 0.55 |
| 1:CA:376:G:P | 16:CP:67:THR:HG21 | 2.46 | 0.55 |
| 1:CA:792:A:H4' | 1:CA:793:U:O5' | 2.06 | 0.55 |
| 1:CA:1049:U:H4' | 1:CA:1050:G:C5' | 2.36 | 0.55 |
| 1:CA:1116:C:H3' | 1:CA:1117:G:H5" | 1.88 | 0.55 |
| 10:CJ:63:PHE:CZ | 14:CN:45:ARG:HG3 | 2.36 | 0.55 |
| 13:CM:71:ARG:O | 13:CM:71:ARG:HG3 | 2.06 | 0.55 |
| 15:CO:54:ARG:O | 15:CO:57:LEU:HB2 | 2.05 | 0.55 |
| 31:DA:143:G:H2' | 31:DA:143(A):C:C6 | 2.41 | 0.55 |
| 31:DA:951:C:C2' | 31:DA:952:G:H5' | 2.36 | 0.55 |
| 31:DA:2203:U:C1' | 33:DD:151:LYS:HE2 | 2.35 | 0.55 |
| 31:DA:2317:C:H2' | 31:DA:2317:C:O2 | 2.06 | 0.55 |
| 31:DA:2752:C:H2' | 31:DA:2752:C:O2 | 2.05 | 0.55 |
| 31:DA:2810:A:C2' | 34:DE:61:ARG:NH2 | 2.67 | 0.55 |
| 32:DB:2:C:H2' | 32:DB:3:C:C6 | 2.41 | 0.55 |
| 34:DE:27:LEU:HD22 | 45:DT:1:MET:CE | 2.35 | 0.55 |
| 34:DE:136:ARG:HG2 | 34:DE:136:ARG:NH1 | 2.21 | 0.55 |
| 40:DO:71:ARG:HE | 40:DO:105:GLU:CD | 2.14 | 0.55 |
| 46:DU:112:ARG:CG | 46:DU:112:ARG:NH1 | 2.59 | 0.55 |
| 48:DW:88:ARG:NH1 | 48:DW:94:ASP:OD1 | 2.37 | 0.55 |
| 1:AA:356:A:C2' | 1:AA:357:G:O5' | 2.54 | 0.55 |
| 1:AA:505:G:C6 | 1:AA:535:A:C2 | 2.94 | 0.55 |
| 1:AA:556:C:C2' | 1:AA:557:G:H5' | 2.36 | 0.55 |
| 1:AA:774:G:C2' | 1:AA:775:G:H5' | 2.36 | 0.55 |
| 1:AA:1228:C:P | 13:AM:108:ARG:HH22 | 2.28 | 0.55 |
| 2:AB:55:PHE:HE1 | 2:AB:218:ALA:HA | 1.71 | 0.55 |
| 3:AC:113:ALA:C | 3:AC:115:LEU:N | 2.65 | 0.55 |
| 16:AP:8:ARG:O | 16:AP:9:PHE:CG | 2.59 | 0.55 |
| 25:B3:8:LEU:CD1 | 25:B3:31:LEU:HA | 2.36 | 0.55 |
| 25:B3:45:GLY:HA3 | 31:BA:851:U:O2' | 2.06 | 0.55 |
| 30:B8:59:LYS:HD2 | 41:BP:50:ARG:HB3 | 1.86 | 0.55 |
| 31:BA:146:G:H8 | 31:BA:146:G:C5' | 2.18 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:615:G:OP1 | 35:BF:40:GLN:NE2 | 2.39 | 0.55 |
| 31:BA:1052:C:H3' | 31:BA:1052:C:H6 | 1.70 | 0.55 |
| 31:BA:1339:G:H21 | 31:BA:1603:A:H1' | 1.71 | 0.55 |
| 31:BA:2657:A:C2 | 31:BA:2664:G:N2 | 2.72 | 0.55 |
| 33:BD:89:SER:HB2 | 33:BD:159:ALA:HB2 | 1.86 | 0.55 |
| 34:BE:116:VAL:HG11 | 34:BE:122:PHE:CD2 | 2.42 | 0.55 |
| 41:BP:102:ARG:O | 41:BP:103:ALA:HB2 | 2.06 | 0.55 |
| 47:BV:51:VAL:CG1 | 47:BV:52:VAL:H | 2.17 | 0.55 |
| 1:CA:60:A:P | 1:CA:60:A:C8 | 3.00 | 0.55 |
| 1:CA:541:G:H2' | 1:CA:542:G:H8 | 1.70 | 0.55 |
| 1:CA:658:G:H1' | 15:CO:22:THR:HB | 1.89 | 0.55 |
| 1:CA:665:A:H1' | 1:CA:733:A:O4' | 2.07 | 0.55 |
| 1:CA:1157:A:C1' | 1:CA:1181:G:H21 | 2.19 | 0.55 |
| 1:CA:1259:C:H42 | 1:CA:1276:G:H1 | 1.52 | 0.55 |
| 3:CC:182:ILE:HG12 | 3:CC:203:PHE:CD1 | 2.38 | 0.55 |
| 4:CD:88:VAL:HG13 | 5:CE:97:GLY:HA3 | 1.88 | 0.55 |
| 31:DA:2299:G:N1 | 31:DA:2318:G:C8 | 2.74 | 0.55 |
| 34:DE:77:ILE:HG23 | 34:DE:78:LEU:N | 2.21 | 0.55 |
| 37:DH:156:ALA:O | 37:DH:157:TYR:C | 2.48 | 0.55 |
| 41:DP:8:PRO:O | 41:DP:9:ASN:C | 2.49 | 0.55 |
| 41:DP:140:ALA:O | 41:DP:141:ALA:CB | 2.53 | 0.55 |
| 46:DU:29:SER:OG | 46:DU:30:LYS:HE3 | 2.06 | 0.55 |
| 48:DW:17:VAL:HG21 | 48:DW:103:ILE:HD13 | 1.89 | 0.55 |
| 1:AA:375:U:H5" | 16:AP:6:LEU:HD22 | 1.88 | 0.55 |
| 1:AA:662:G:H2' | 1:AA:663:A:C8 | 2.41 | 0.55 |
| 1:AA:1095:U:H2' | 1:AA:1096:C:C6 | 2.41 | 0.55 |
| 1:AA:1130:A:H1' | 1:AA:1146:A:C2 | 2.41 | 0.55 |
| 3:AC:189:ALA:HB3 | 3:AC:196:LEU:HB2 | 1.88 | 0.55 |
| 4:AD:112:VAL:HG12 | 4:AD:116:GLN:OE1 | 2.05 | 0.55 |
| 5:AE:43:LEU:HB2 | 5:AE:136:MET:SD | 2.46 | 0.55 |
| 7:AG:23:VAL:HG13 | 7:AG:43:PHE:CZ | 2.41 | 0.55 |
| 11:AK:31:THR:HA | 11:AK:42:TRP:HA | 1.86 | 0.55 |
| 11:AK:99:GLN:O | 11:AK:101:SER:N | 2.39 | 0.55 |
| 24:B2:30:ARG:HG3 | 24:B2:30:ARG:HH11 | 1.71 | 0.55 |
| 28:B6:27:LYS:HE3 | 31:BA:2285:C:H5 | 1.71 | 0.55 |
| 30:B8:32:LEU:HB3 | 30:B8:34:TRP:CB | 2.35 | 0.55 |
| 31:BA:394:A:C6 | 31:BA:395:U:C4 | 2.94 | 0.55 |
| 31:BA:2637:U:H5' | 31:BA:2637:U:H6 | 1.70 | 0.55 |
| 31:BA:2639:A:H2' | 31:BA:2640:G:H5' | 1.87 | 0.55 |
| 31:BA:2752:C:H2' | 31:BA:2752:C:O2 | 2.06 | 0.55 |
| 35:BF:7:TYR:CD1 | 35:BF:8:GLN:N | 2.74 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 37:BH:84:SER:O | 37:BH:133:VAL:O | 2.24 | 0.55 |
| 39:BN:17:ASP:C | 39:BN:19:GLU:H | 2.14 | 0.55 |
| 48:BW:62:HIS:O | 48:BW:64:MET:HG3 | 2.05 | 0.55 |
| 1:CA:411:A:C6 | 1:CA:429:U:C4 | 2.94 | 0.55 |
| 1:CA:629:G:C4 | 1:CA:630:G:C8 | 2.94 | 0.55 |
| 3:CC:116:VAL:HG21 | 3:CC:202:ILE:HD11 | 1.88 | 0.55 |
| 6:CF:2:ARG:HB2 | 6:CF:4:TYR:CE2 | 2.41 | 0.55 |
| 7:CG:138:LYS:HE2 | 7:CG:142:GLU:OE2 | 2.07 | 0.55 |
| 8:CH:87:SER:HA | 8:CH:93:VAL:HG23 | 1.89 | 0.55 |
| 8:CH:104:ARG:O | 8:CH:105:ARG:HB2 | 2.06 | 0.55 |
| 15:CO:56:LEU:HD21 | 31:DA:715:G:C2 | 2.40 | 0.55 |
| 16:CP:39:TYR:HB2 | 16:CP:49:LEU:CD1 | 2.36 | 0.55 |
| 25:D3:1:MET:HB3 | 25:D3:2:PRO:HD2 | 1.88 | 0.55 |
| 31:DA:642:G:H21 | 31:DA:646:A:H2 | 1.51 | 0.55 |
| 31:DA:838:C:C4 | 31:DA:839:U:C5 | 2.94 | 0.55 |
| 31:DA:1472:A:C2' | 31:DA:1473:G:H5' | 2.37 | 0.55 |
| 31:DA:1493:C:O2 | 31:DA:1493:C:C2' | 2.51 | 0.55 |
| 31:DA:1856:G:H2' | 31:DA:1857:G:H5' | 1.88 | 0.55 |
| 31:DA:1986:A:H2' | 31:DA:1987:G:C5' | 2.24 | 0.55 |
| 31:DA:2274:A:C5 | 31:DA:2276:G:C8 | 2.94 | 0.55 |
| 32:DB:86:G:H8 | 32:DB:86:G:O5' | 1.88 | 0.55 |
| 39:DN:39:ARG:HE | 39:DN:41:ASP:CB | 2.19 | 0.55 |
| 40:DO:4:PRO:O | 40:DO:5:GLN:HB2 | 2.06 | 0.55 |
| 50:DY:28:LYS:C | 50:DY:38:ILE:HB | 2.30 | 0.55 |
| 1:AA:56:U:H2' | 1:AA:57:G:C8 | 2.42 | 0.55 |
| 1:AA:760:G:H2' | 1:AA:761:G:H5' | 1.88 | 0.55 |
| 1:AA:955:U:H1' | 1:AA:1227:A:N6 | 2.22 | 0.55 |
| 1:AA:1416:G:H2' | 1:AA:1417:G:O4' | 2.07 | 0.55 |
| 1:AA:1442(B):A:H4' | 1:AA:1442(B):A:OP1 | 2.06 | 0.55 |
| 2:AB:213:LEU:O | 2:AB:213:LEU:HD23 | 2.05 | 0.55 |
| 4:AD:33:MET:CE | 4:AD:37:PRO:HA | 2.37 | 0.55 |
| 17:AQ:33:GLY:O | 17:AQ:34:LYS:C | 2.49 | 0.55 |
| 24:B2:30:ARG:CZ | 24:B2:30:ARG:HB3 | 2.36 | 0.55 |
| 28:B6:15:GLU:OE2 | 28:B6:18:ARG:NE | 2.40 | 0.55 |
| 31:BA:573:G:O2' | 31:BA:574:C:H3' | 2.06 | 0.55 |
| 31:BA:993:G:H1' | 47:BV:91:TYR:CD1 | 2.41 | 0.55 |
| 31:BA:1025:G:C4 | 31:BA:1135:C:H1' | 2.42 | 0.55 |
| 31:BA:1386:C:H2' | 31:BA:1387:C:C6 | 2.41 | 0.55 |
| 31:BA:1654:A:OP2 | 43:BR:3:HIS:HB2 | 2.06 | 0.55 |
| 31:BA:2360:A:O2' | 31:BA:2361:A:OP2 | 2.24 | 0.55 |
| 35:BF:24:LEU:O | 35:BF:25:PRO:C | 2.47 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 40:BO:77:ILE:HG13 | 45:BT:74:ARG:HG2 | 1.87 | 0.55 |
| 1:CA:1452:C:O4' | 1:CA:1456:G:C2 | 2.59 | 0.55 |
| 6:CF:79:LEU:HB2 | 6:CF:88:VAL:HG21 | 1.88 | 0.55 |
| 10:CJ:61:GLU:OE1 | 14:CN:58:LYS:HE2 | 2.06 | 0.55 |
| 11:CK:80:VAL:O | 11:CK:106:LYS:HB3 | 2.07 | 0.55 |
| 13:CM:91:ARG:HD3 | 19:CS:81:ARG:HH21 | 1.71 | 0.55 |
| 18:CR:63:GLN:HA | 18:CR:63:GLN:OE1 | 2.07 | 0.55 |
| 27:D5:40:LYS:NZ | 27:D5:46:CYS:HB3 | 2.21 | 0.55 |
| 28:D6:13:CYS:HA | 28:D6:50:ARG:O | 2.05 | 0.55 |
| 31:DA:477:A:O2' | 31:DA:478:A:H5' | 2.07 | 0.55 |
| 31:DA:493:G:C2' | 31:DA:494:G:H5" | 2.32 | 0.55 |
| 31:DA:494:G:O2' | 31:DA:495:G:H5' | 2.06 | 0.55 |
| 31:DA:1410:G:H2' | 31:DA:1411:C:C6 | 2.40 | 0.55 |
| 31:DA:2298:A:H2' | 31:DA:2299:G:O4' | 2.07 | 0.55 |
| 31:DA:2582:G:C2 | 31:DA:2583:G:C8 | 2.95 | 0.55 |
| 33:DD:18:VAL:HG23 | 33:DD:211:ARG:HH12 | 1.71 | 0.55 |
| 33:DD:86:PRO:HG2 | 33:DD:87:ASN:ND2 | 2.21 | 0.55 |
| 34:DE:9:VAL:HG13 | 34:DE:25:VAL:O | 2.05 | 0.55 |
| 46:DU:87:GLY:O | 47:DV:52:VAL:HA | 2.07 | 0.55 |
| 48:DW:45:TYR:O | 48:DW:46:PHE:C | 2.49 | 0.55 |
| 49:DX:24:GLY:HA3 | 49:DX:80:ILE:HG12 | 1.88 | 0.55 |
| 49:DX:85:PRO:O | 49:DX:87:GLN:N | 2.40 | 0.55 |
| 51:DZ:149:SER:CB | 51:DZ:173:ALA:HA | 2.37 | 0.55 |
| 1:AA:495:A:H4' | 1:AA:496:A:OP1 | 2.06 | 0.55 |
| 1:AA:542:G:H2' | 1:AA:543:C:C6 | 2.41 | 0.55 |
| 1:AA:665:A:H2' | 1:AA:732:C:O2 | 2.07 | 0.55 |
| 1:AA:731:G:OP1 | 1:AA:766:A:H1' | 2.07 | 0.55 |
| 1:AA:836:G:C6 | 1:AA:851:G:C6 | 2.94 | 0.55 |
| 1:AA:853:G:H2' | 1:AA:854:G:H8 | 1.70 | 0.55 |
| 1:AA:1315:U:H2' | 1:AA:1316:G:O4' | 2.07 | 0.55 |
| 1:AA:1346:A:H5" | 9:AI:120:ARG:HH12 | 1.72 | 0.55 |
| 2:AB:201:ILE:HG21 | 2:AB:214:ILE:HG21 | 1.89 | 0.55 |
| 3:AC:14:ILE:O | 3:AC:15:THR:HB | 2.07 | 0.55 |
| 12:AL:102:ARG:HG3 | 12:AL:102:ARG:NH1 | 2.21 | 0.55 |
| 20:AT:10:LEU:O | 20:AT:12:ALA:N | 2.31 | 0.55 |
| 20:AT:63:ILE:HD13 | 20:AT:80:ARG:HB2 | 1.88 | 0.55 |
| 28:B6:42:TRP:HA | 28:B6:42:TRP:HE3 | 1.71 | 0.55 |
| 31:BA:536:A:H2' | 31:BA:537:C:C6 | 2.40 | 0.55 |
| 31:BA:717:G:H2' | 31:BA:718:A:O4' | 2.07 | 0.55 |
| 31:BA:1028:A:N3 | 31:BA:2486:G:O2' | 2.31 | 0.55 |
| 31:BA:1709:U:O2' | 31:BA:2859:G:H1' | 2.06 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1797:C:O2' | 33:BD:259:THR:HB | 2.06 | 0.55 |
| 31:BA:2682:U:C5 | 34:BE:11:MET:HE2 | 2.40 | 0.55 |
| 31:BA:2688:U:H5 | 31:BA:2720:U:OP2 | 1.90 | 0.55 |
| 33:BD:118:VAL:HG22 | 33:BD:119:ALA:N | 2.22 | 0.55 |
| 45:BT:10:VAL:O | 45:BT:13:ARG:HG2 | 2.06 | 0.55 |
| 46:BU:100:VAL:O | 46:BU:103:PRO:HD3 | 2.06 | 0.55 |
| 46:BU:101:ARG:C | 46:BU:102:GLU:HG2 | 2.32 | 0.55 |
| 47:BV:90:PRO:CG | 47:BV:91:TYR:N | 2.68 | 0.55 |
| 49:BX:72:LYS:HB2 | 49:BX:74:PRO:CD | 2.36 | 0.55 |
| 50:BY:77:PRO:O | 50:BY:78:ALA:HB2 | 2.05 | 0.55 |
| 1:CA:375:U:H5" | 16:CP:6:LEU:HD22 | 1.88 | 0.55 |
| 1:CA:532:A:H61 | 3:CC:193:TYR:CB | 2.17 | 0.55 |
| 1:CA:791:G:C6 | 1:CA:792:A:N7 | 2.74 | 0.55 |
| 1:CA:1130:A:H1' | 1:CA:1146:A:C2 | 2.42 | 0.55 |
| 3:CC:150:LYS:HG3 | 3:CC:169:ALA:HB2 | 1.88 | 0.55 |
| 4:CD:119:GLN:CG | 4:CD:123:HIS:CD2 | 2.89 | 0.55 |
| 7:CG:27:ILE:HD11 | 7:CG:43:PHE:CD2 | 2.42 | 0.55 |
| 10:CJ:89:ASP:C | 10:CJ:91:PRO:HD3 | 2.31 | 0.55 |
| 11:CK:102:GLY:C | 11:CK:103:LEU:HD22 | 2.32 | 0.55 |
| 28:D6:16:CYS:O | 28:D6:17:LYS:HB2 | 2.05 | 0.55 |
| 31:DA:547:A:H8 | 31:DA:549:G:C6 | 2.25 | 0.55 |
| 31:DA:607:U:H3 | 31:DA:621:A:H2 | 1.45 | 0.55 |
| 31:DA:1037:G:H1 | 31:DA:1118:C:H42 | 1.54 | 0.55 |
| 31:DA:1141:U:H4' | 31:DA:1142(A):A:O4' | 2.06 | 0.55 |
| 31:DA:1678:G:N2 | 31:DA:1989:G:N2 | 2.52 | 0.55 |
| 31:DA:2070:G:C2 | 31:DA:2442:C:C2 | 2.94 | 0.55 |
| 31:DA:2534:A:C2 | 31:DA:2535:G:H1' | 2.41 | 0.55 |
| 42:DQ:23:GLY:HA3 | 42:DQ:99:PRO:C | 2.31 | 0.55 |
| 42:DQ:108:GLY:C | 42:DQ:109:VAL:HG22 | 2.31 | 0.55 |
| 46:DU:117:GLN:OE1 | 46:DU:117:GLN:HA | 2.06 | 0.55 |
| 50:DY:38:ILE:HG22 | 50:DY:39:VAL:H | 1.72 | 0.55 |
| 1:AA:836:G:C6 | 1:AA:851:G:C5 | 2.94 | 0.55 |
| 2:AB:71:VAL:CG1 | 2:AB:93:VAL:HB | 2.37 | 0.55 |
| 10:AJ:40:LEU:HD21 | 10:AJ:69:ASN:HB3 | 1.88 | 0.55 |
| 13:AM:46:LYS:HG3 | 13:AM:47:ASP:N | 2.21 | 0.55 |
| 20:AT:104:LEU:C | 20:AT:104:LEU:HD23 | 2.32 | 0.55 |
| 31:BA:906:G:H2' | 31:BA:907:U:O5' | 2.07 | 0.55 |
| 31:BA:1037:G:H1 | 31:BA:1118:C:H42 | 1.55 | 0.55 |
| 31:BA:1649:G:C6 | 31:BA:2009:G:C6 | 2.95 | 0.55 |
| 31:BA:1783:A:C2 | 31:BA:2587:A:C5 | 2.94 | 0.55 |
| 31:BA:2681:C:O2 | 31:BA:2681:C:H2' | 2.06 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 35:BF:83:PHE:C | 35:BF:84:VAL:HG23 | 2.32 | 0.55 |
| 37:BH:148:ILE:O | 37:BH:151:ILE:HG12 | 2.05 | 0.55 |
| 39:BN:3:THR:HG22 | 39:BN:4:TYR:N | 2.14 | 0.55 |
| 41:BP:62:LEU:N | 41:BP:62:LEU:HD13 | 2.21 | 0.55 |
| 42:BQ:9:TYR:CD2 | 42:BQ:9:TYR:C | 2.82 | 0.55 |
| 44:BS:58:LEU:HD21 | 44:BS:68:GLN:HB2 | 1.89 | 0.55 |
| 46:BU:10:ARG:O | 46:BU:11:ARG:C | 2.47 | 0.55 |
| 46:BU:87:GLY:O | 47:BV:52:VAL:HA | 2.07 | 0.55 |
| 50:BY:28:LYS:HD2 | 50:BY:37:VAL:CG1 | 2.36 | 0.55 |
| 1:CA:96:U:O2' | 1:CA:97:G:P | 2.64 | 0.55 |
| 1:CA:629:G:H2' | 1:CA:630:G:O4' | 2.05 | 0.55 |
| 1:CA:922:G:N3 | 1:CA:1398:A:H2 | 2.04 | 0.55 |
| 1:CA:1118:C:H1' | 1:CA:1179:A:C4 | 2.42 | 0.55 |
| 4:CD:133:VAL:HG13 | 4:CD:135:LEU:HD22 | 1.87 | 0.55 |
| 6:CF:15:ASP:C | 6:CF:17:SER:H | 2.15 | 0.55 |
| 28:D6:36:LEU:C | 28:D6:37:ARG:HD2 | 2.32 | 0.55 |
| 31:DA:542:C:O5' | 31:DA:542:C:H6 | 1.89 | 0.55 |
| 31:DA:1027:A:N6 | 31:DA:1126:A:C4 | 2.75 | 0.55 |
| 31:DA:2250:G:C4 | 42:DQ:82:ARG:HD2 | 2.41 | 0.55 |
| 31:DA:2307:G:N2 | 31:DA:2308:G:C5' | 2.69 | 0.55 |
| 31:DA:2830:G:H5" | 31:DA:2830:G:H8 | 1.72 | 0.55 |
| 31:DA:2897:U:H2' | 31:DA:2897:U:O2 | 2.05 | 0.55 |
| 32:DB:116:G:H8 | 32:DB:116:G:H5" | 1.70 | 0.55 |
| 33:DD:35:LYS:CE | 33:DD:64:ILE:C | 2.79 | 0.55 |
| 45:DT:121:ILE:O | 45:DT:124:ASP:HB2 | 2.07 | 0.55 |
| 47:DV:69:LYS:HB3 | 47:DV:93:GLU:CD | 2.31 | 0.55 |
| 50:DY:15:VAL:HG12 | 50:DY:16:ALA:H | 1.72 | 0.55 |
| 1:AA:79:G:C4' | 1:AA:80:G:OP1 | 2.55 | 0.55 |
| 1:AA:165:C:H2' | 1:AA:166:G:C8 | 2.41 | 0.55 |
| 1:AA:228:A:H2' | 1:AA:229:U:O4' | 2.06 | 0.55 |
| 1:AA:1157:A:C1' | 1:AA:1181:G:H21 | 2.19 | 0.55 |
| 2:AB:20:GLU:O | 2:AB:40:HIS:HB2 | 2.07 | 0.55 |
| 3:AC:32:LEU:HD22 | 3:AC:59:ARG:NH1 | 2.22 | 0.55 |
| 3:AC:150:LYS:HG3 | 3:AC:169:ALA:HB2 | 1.89 | 0.55 |
| 9:AI:53:VAL:C | 9:AI:92:TYR:HE2 | 2.15 | 0.55 |
| 11:AK:125:PHE:HD1 | 11:AK:125:PHE:H | 1.55 | 0.55 |
| 18:AR:45:SER:H | 18:AR:51:LEU:HD11 | 1.72 | 0.55 |
| 31:BA:357:A:N3 | 31:BA:358:U:O2 | 2.39 | 0.55 |
| 31:BA:1543:C:C6 | 31:BA:1543:C:OP2 | 2.59 | 0.55 |
| 31:BA:1701:A:C2' | 31:BA:1702:G:H5' | 2.37 | 0.55 |
| 31:BA:1994:C:O2' | 31:BA:1995:U:H5' | 2.06 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 33:BD:70:TRP:CZ3 | 33:BD:146:GLU:OE2 | 2.55 | 0.55 |
| 36:BG:107:LEU:HD11 | 36:BG:178:PHE:CE1 | 2.41 | 0.55 |
| 37:BH:29:PRO:HD2 | 37:BH:79:VAL:O | 2.07 | 0.55 |
| 41:BP:66:GLY:O | 41:BP:67:MET:O | 2.25 | 0.55 |
| 41:BP:140:ALA:O | 25:D3:1:MET:SD | 2.65 | 0.55 |
| 42:BQ:23:GLY:O | 42:BQ:100:GLY:CA | 2.47 | 0.55 |
| 42:BQ:139:GLU:HG2 | 42:BQ:139:GLU:O | 2.07 | 0.55 |
| 45:BT:45:PHE:HE2 | 45:BT:63:VAL:HG22 | 1.72 | 0.55 |
| 1:CA:115:G:H4' | 1:CA:116:A:O5' | 2.06 | 0.55 |
| 1:CA:949:A:H1' | 1:CA:1364:U:N3 | 2.22 | 0.55 |
| 2:CB:204:ASN:HD22 | 2:CB:205:ASP:N | 2.05 | 0.55 |
| 3:CC:52:LEU:H | 3:CC:52:LEU:CD2 | 2.17 | 0.55 |
| 3:CC:53:ALA:O | 3:CC:54:ARG:HB2 | 2.05 | 0.55 |
| 3:CC:69:HIS:N | 3:CC:69:HIS:CD2 | 2.73 | 0.55 |
| 3:CC:71:ALA:HA | 3:CC:106:VAL:HB | 1.89 | 0.55 |
| 4:CD:49:ARG:HA | 4:CD:49:ARG:HE | 1.71 | 0.55 |
| 4:CD:118:ARG:O | 4:CD:121:VAL:HG23 | 2.05 | 0.55 |
| 21:CU:5:ASP:O | 21:CU:11:GLY:HA3 | 2.07 | 0.55 |
| 27:D5:2:ALA:N | 31:DA:747:U:N3 | 2.53 | 0.55 |
| 31:DA:280:C:H2' | 31:DA:281:G:O5' | 2.06 | 0.55 |
| 31:DA:352:G:O2' | 31:DA:353:G:H3' | 2.07 | 0.55 |
| 31:DA:357:A:N3 | 31:DA:358:U:O2 | 2.40 | 0.55 |
| 31:DA:1107:G:H2' | 31:DA:1108:U:O4' | 2.07 | 0.55 |
| 31:DA:1991:U:C2' | 31:DA:1992:G:H5" | 2.37 | 0.55 |
| 31:DA:2247:A:H2' | 31:DA:2248:C:H6 | 1.71 | 0.55 |
| 31:DA:2358:G:H1 | 41:DP:55:ARG:HH22 | 1.55 | 0.55 |
| 31:DA:2402:C:C3' | 31:DA:2403:C:H5' | 2.36 | 0.55 |
| 31:DA:2801(A):A:C4' | 31:DA:2802:G:H5' | 2.36 | 0.55 |
| 33:DD:231:HIS:CD2 | 33:DD:249:PRO:HA | 2.42 | 0.55 |
| 39:DN:95:PRO:HD2 | 39:DN:96:GLU:OE2 | 2.07 | 0.55 |
| 48:DW:20:VAL:O | 48:DW:23:LEU:HB2 | 2.06 | 0.55 |
| 50:DY:37:VAL:HG23 | 50:DY:38:ILE:N | 2.20 | 0.55 |
| 51:DZ:52:SER:OG | 51:DZ:53:ILE:N | 2.38 | 0.55 |
| 1:AA:491:G:C4 | 1:AA:492:G:C8 | 2.95 | 0.55 |
| 1:AA:1132:C:H2' | 1:AA:1133:G:O4' | 2.07 | 0.55 |
| 4:AD:62:GLN:HA | 4:AD:62:GLN:NE2 | 2.16 | 0.55 |
| 4:AD:146:ILE:HD12 | 4:AD:146:ILE:N | 2.22 | 0.55 |
| 27:B5:4:HIS:HD2 | 31:BA:2056:G:H1 | 1.54 | 0.55 |
| 28:B6:42:TRP:HA | 28:B6:42:TRP:CE3 | 2.41 | 0.55 |
| 31:BA:352:G:O2' | 31:BA:353:G:H3' | 2.06 | 0.55 |
| 31:BA:1496:A:C8 | 31:BA:1577:C:O2' | 2.59 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2262:U:C2' | 31:BA:2263:C:H5' | 2.36 | 0.55 |
| 31:BA:2483:C:H2' | 31:BA:2483:C:O2 | 2.04 | 0.55 |
| 34:BE:163:GLU:O | 34:BE:165:VAL:HG23 | 2.06 | 0.55 |
| 37:BH:56:SER:HB2 | 37:BH:58:GLU:HG3 | 1.88 | 0.55 |
| 41:BP:62:LEU:N | 41:BP:62:LEU:CD2 | 2.59 | 0.55 |
| 41:BP:105:LEU:HD12 | 41:BP:105:LEU:N | 2.22 | 0.55 |
| 41:BP:107:LYS:C | 41:BP:109:GLY:N | 2.63 | 0.55 |
| 45:BT:99:LEU:CD1 | 45:BT:99:LEU:O | 2.55 | 0.55 |
| 45:BT:124:ASP:C | 45:BT:126:ALA:H | 2.14 | 0.55 |
| 1:CA:414:A:H2' | 1:CA:415:A:H8 | 1.71 | 0.55 |
| 2:CB:28:PHE:CD1 | 2:CB:190:THR:HG22 | 2.42 | 0.55 |
| 2:CB:67:THR:C | 2:CB:68:ILE:HD12 | 2.32 | 0.55 |
| 5:CE:34:VAL:O | 5:CE:41:VAL:HA | 2.07 | 0.55 |
| 5:CE:101:ILE:HD11 | 5:CE:119:LEU:HD23 | 1.89 | 0.55 |
| 8:CH:64:LYS:HG2 | 8:CH:79:VAL:HG21 | 1.87 | 0.55 |
| 11:CK:106:LYS:O | 11:CK:106:LYS:HG3 | 2.06 | 0.55 |
| 12:CL:25:PRO:C | 12:CL:27:LEU:H | 2.14 | 0.55 |
| 15:CO:87:ILE:CG2 | 15:CO:88:ARG:N | 2.69 | 0.55 |
| 29:D7:1:MET:O | 29:D7:2:LYS:C | 2.49 | 0.55 |
| 30:D8:6:THR:CG2 | 31:DA:243:U:OP1 | 2.54 | 0.55 |
| 31:DA:71:A:C2 | 49:DX:31:HIS:CE1 | 2.78 | 0.55 |
| 31:DA:309:G:H4' | 50:DY:18:GLY:HA3 | 1.88 | 0.55 |
| 31:DA:902:C:O2' | 31:DA:903:C:H5' | 2.06 | 0.55 |
| 31:DA:1316:U:H2' | 31:DA:1317:A:C8 | 2.41 | 0.55 |
| 31:DA:1578:U:H2' | 31:DA:1578:U:O2 | 2.05 | 0.55 |
| 31:DA:2840:C:H6 | 31:DA:2840:C:O5' | 1.90 | 0.55 |
| 32:DB:116:G:C8 | 32:DB:116:G:H5' | 2.42 | 0.55 |
| 33:DD:76:PRO:O | 33:DD:98:VAL:HG23 | 2.07 | 0.55 |
| 33:DD:176:ARG:HG2 | 33:DD:176:ARG:NH1 | 2.22 | 0.55 |
| 35:DF:22:ALA:HB1 | 35:DF:26:ALA:HB1 | 1.89 | 0.55 |
| 35:DF:88:VAL:HG11 | 35:DF:91:GLY:HA3 | 1.89 | 0.55 |
| 35:DF:124:LEU:HD12 | 35:DF:124:LEU:C | 2.31 | 0.55 |
| 42:DQ:22:LYS:HA | 42:DQ:22:LYS:CE | 2.35 | 0.55 |
| 43:DR:29:LEU:HB3 | 43:DR:75:LEU:HD11 | 1.89 | 0.55 |
| 47:DV:80:GLN:OE1 | 47:DV:80:GLN:C | 2.50 | 0.55 |
| 51:DZ:119:GLU:C | 51:DZ:121:HIS:H | 2.14 | 0.55 |
| 1:AA:1015:A:N6 | 1:AA:1016:A:C6 | 2.74 | 0.55 |
| 1:AA:1049:U:H4' | 1:AA:1050:G:C5' | 2.37 | 0.55 |
| 1:AA:1259:C:H42 | 1:AA:1276:G:H1 | 1.54 | 0.55 |
| 1:AA:1321:C:H5" | 1:AA:1322:C:H2' | 1.88 | 0.55 |
| 2:AB:44:LEU:HA | 2:AB:47:THR:HB | 1.89 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 2:AB:178:ARG:HH21 | 8:AH:68:ARG:HH22 | 1.54 | 0.55 |
| 6:AF:15:ASP:C | 6:AF:17:SER:H | 2.13 | 0.55 |
| 8:AH:6:ILE:H | 8:AH:6:ILE:HD12 | 1.71 | 0.55 |
| 15:AO:3:ILE:HD13 | 15:AO:3:ILE:N | 2.21 | 0.55 |
| 15:AO:65:ARG:HB2 | 15:AO:65:ARG:HH11 | 1.72 | 0.55 |
| 16:AP:64:ALA:O | 16:AP:65:GLN:C | 2.49 | 0.55 |
| 24:B2:40:SER:O | 24:B2:44:LEU:HB3 | 2.06 | 0.55 |
| 27:B5:47:PRO:C | 27:B5:48:GLU:OE2 | 2.50 | 0.55 |
| 31:BA:910:A:N7 | 42:BQ:13:GLN:HB2 | 2.21 | 0.55 |
| 31:BA:1515:G:H2' | 31:BA:1516:C:C6 | 2.41 | 0.55 |
| 31:BA:1580:A:H8 | 31:BA:1580:A:OP2 | 1.89 | 0.55 |
| 31:BA:1582:C:O2' | 31:BA:1586:A:C8 | 2.59 | 0.55 |
| 33:BD:76:PRO:HG2 | 33:BD:98:VAL:CG2 | 2.35 | 0.55 |
| 33:BD:133:LEU:HD23 | 33:BD:136:ILE:HD12 | 1.88 | 0.55 |
| 33:BD:223:GLY:HA3 | 33:BD:231:HIS:ND1 | 2.22 | 0.55 |
| 33:BD:224:ALA:O | 33:BD:225:ALA:CB | 2.54 | 0.55 |
| 35:BF:65:TRP:HZ3 | 35:BF:73:ALA:O | 1.90 | 0.55 |
| 36:BG:110:ALA:HA | 36:BG:140:ILE:O | 2.07 | 0.55 |
| 38:BI:95:LYS:O | 38:BI:99:GLU:HB2 | 2.07 | 0.55 |
| 39:BN:27:ALA:CB | 39:BN:106:MET:CE | 2.84 | 0.55 |
| 39:BN:67:LEU:C | 39:BN:69:GLN:N | 2.64 | 0.55 |
| 41:BP:85:LEU:HB3 | 41:BP:114:ILE:HD12 | 1.89 | 0.55 |
| 42:BQ:23:GLY:HA3 | 42:BQ:99:PRO:C | 2.31 | 0.55 |
| 44:BS:36:TYR:CD1 | 44:BS:36:TYR:N | 2.74 | 0.55 |
| 44:BS:98:VAL:HG13 | 44:BS:100:ALA:H | 1.71 | 0.55 |
| 1:CA:80:G:H1 | 1:CA:89:C:H41 | 1.50 | 0.55 |
| 1:CA:163:C:H2' | 1:CA:164:U:H6 | 1.71 | 0.55 |
| 1:CA:355:C:C2' | 1:CA:356:A:H5' | 2.37 | 0.55 |
| 1:CA:955:U:H1' | 1:CA:1227:A:N6 | 2.22 | 0.55 |
| 1:CA:995:C:H1' | 14:CN:8:GLU:OE2 | 2.07 | 0.55 |
| 2:CB:44:LEU:HA | 2:CB:47:THR:HB | 1.89 | 0.55 |
| 2:CB:158:LEU:H | 2:CB:158:LEU:CD1 | 2.08 | 0.55 |
| 3:CC:35:GLU:O | 3:CC:39:ILE:HG13 | 2.07 | 0.55 |
| 4:CD:112:VAL:HG12 | 4:CD:116:GLN:OE1 | 2.07 | 0.55 |
| 6:CF:72:VAL:HG13 | 6:CF:73:ASN:N | 2.22 | 0.55 |
| 9:CI:53:VAL:HB | 9:CI:92:TYR:CE2 | 2.42 | 0.55 |
| 18:CR:66:LEU:O | 18:CR:70:ILE:HG13 | 2.07 | 0.55 |
| 31:DA:795:C:O2' | 31:DA:796:C:H5' | 2.06 | 0.55 |
| 31:DA:1028:A:N3 | 31:DA:2486:G:O2' | 2.31 | 0.55 |
| 31:DA:1112:G:H1' | 31:DA:1113:U:OP1 | 2.07 | 0.55 |
| 31:DA:1518:U:H2' | 31:DA:1519:G:O4' | 2.06 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1786:A:H1' | 31:DA:1938:A:N6 | 2.22 | 0.55 |
| 31:DA:2360:A:O2' | 31:DA:2361:A:O5' | 2.25 | 0.55 |
| 31:DA:2507:C:C2 | 31:DA:2508:G:C8 | 2.95 | 0.55 |
| 31:DA:2781:A:H5" | 31:DA:2782:G:H5' | 1.89 | 0.55 |
| 31:DA:2810:A:N6 | 31:DA:2891:G:O2' | 2.40 | 0.55 |
| 33:DD:25:THR:O | 33:DD:25:THR:CG2 | 2.54 | 0.55 |
| 36:DG:64:THR:HG23 | 36:DG:65:GLY:N | 2.19 | 0.55 |
| 44:DS:96:GLY:O | 44:DS:98:VAL:HB | 2.07 | 0.55 |
| 46:DU:88:ILE:O | 46:DU:88:ILE:HD13 | 2.07 | 0.55 |
| 49:DX:82:GLN:NE2 | 49:DX:83:VAL:HG22 | 2.22 | 0.55 |
| 51:DZ:156:LYS:O | 51:DZ:158:PRO:HD3 | 2.07 | 0.55 |
| 1:AA:948:C:C5 | 13:AM:106:ASN:ND2 | 2.75 | 0.55 |
| 2:AB:208:ILE:HA | 2:AB:211:ILE:HD12 | 1.88 | 0.55 |
| 3:AC:53:ALA:HB2 | 3:AC:115:LEU:HD21 | 1.89 | 0.55 |
| 4:AD:79:PHE:CD1 | 4:AD:207:TYR:HD1 | 2.25 | 0.55 |
| 5:AE:10:MET:HB2 | 5:AE:32:VAL:CG2 | 2.35 | 0.55 |
| 15:AO:64:ARG:HG3 | 15:AO:64:ARG:NH1 | 2.22 | 0.55 |
| 22:B0:24:LYS:O | 22:B0:25:ARG:HD2 | 2.06 | 0.55 |
| 23:B1:10:LYS:HD3 | 23:B1:14:VAL:HA | 1.89 | 0.55 |
| 24:B2:51:ARG:O | 24:B2:52:ASP:HB3 | 2.07 | 0.55 |
| 28:B6:30:THR:O | 28:B6:31:PRO:C | 2.50 | 0.55 |
| 31:BA:780:G:C2 | 31:BA:782:A:C2 | 2.95 | 0.55 |
| 31:BA:1518:U:H2' | 31:BA:1519:G:O4' | 2.07 | 0.55 |
| 31:BA:2186:G:H2' | 31:BA:2187:G:H5" | 1.88 | 0.55 |
| 31:BA:2191:G:C2' | 31:BA:2192:G:O5' | 2.56 | 0.55 |
| 31:BA:2228:G:C5 | 31:BA:2229:C:C4 | 2.95 | 0.55 |
| 31:BA:2250:G:C6 | 42:BQ:82:ARG:HD2 | 2.41 | 0.55 |
| 31:BA:2343:C:HO2' | 31:BA:2373:G:HO2' | 1.52 | 0.55 |
| 32:BB:43:C:H4' | 36:BG:66:GLN:NE2 | 2.21 | 0.55 |
| 44:BS:91:PRO:O | 44:BS:93:LYS:N | 2.40 | 0.55 |
| 45:BT:23:ARG:HB2 | 45:BT:24:PRO:CD | 2.23 | 0.55 |
| 46:BU:65:ILE:HG12 | 46:BU:96:ALA:HB3 | 1.90 | 0.55 |
| 1:CA:191:G:N3 | 20:CT:103:GLY:O | 2.40 | 0.55 |
| 1:CA:376:G:H4' | 16:CP:5:ARG:NH1 | 2.13 | 0.55 |
| 1:CA:616:G:C2 | 1:CA:617:G:C8 | 2.94 | 0.55 |
| 1:CA:617:G:N1 | 1:CA:618:C:C4 | 2.75 | 0.55 |
| 1:CA:665:A:H2' | 1:CA:732:C:O2 | 2.07 | 0.55 |
| 1:CA:1236:A:O2' | 1:CA:1304:G:H4' | 2.07 | 0.55 |
| 3:CC:39:ILE:HG21 | 3:CC:57:ILE:HD11 | 1.88 | 0.55 |
| 3:CC:112:SER:O | 3:CC:116:VAL:HG23 | 2.06 | 0.55 |
| 6:CF:62:TRP:O | 6:CF:62:TRP:CE3 | 2.60 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 14:CN:24:CYS:HB3 | 14:CN:27:CYS:O | 2.07 | 0.55 |
| 18:CR:59:SER:HB3 | 18:CR:62:GLU:CG | 2.36 | 0.55 |
| 19:CS:6:LYS:HD3 | 19:CS:7:LYS:HE3 | 1.88 | 0.55 |
| 20:CT:104:LEU:HD23 | 20:CT:104:LEU:C | 2.32 | 0.55 |
| 24:D2:25:VAL:HG13 | 24:D2:26:ARG:HD3 | 1.87 | 0.55 |
| 27:D5:42:PRO:HB2 | 27:D5:43:HIS:HD2 | 1.71 | 0.55 |
| 30:D8:22:VAL:HB | 30:D8:53:PRO:CB | 2.34 | 0.55 |
| 31:DA:143:G:C1' | 49:DX:38:GLU:HG3 | 2.35 | 0.55 |
| 31:DA:620:G:H4' | 31:DA:621:A:H5" | 1.89 | 0.55 |
| 31:DA:997:G:O2' | 31:DA:998:C:H5' | 2.07 | 0.55 |
| 31:DA:1349:A:H2' | 31:DA:1350:C:OP1 | 2.06 | 0.55 |
| 31:DA:1396:U:O2 | 31:DA:1396:U:C2' | 2.55 | 0.55 |
| 31:DA:1946:U:H2' | 31:DA:1947:C:C6 | 2.39 | 0.55 |
| 31:DA:2093:G:O5' | 38:DI:24:GLY:HA3 | 2.07 | 0.55 |
| 35:DF:1:MET:O | 35:DF:2:LYS:C | 2.49 | 0.55 |
| 37:DH:138:LYS:C | 37:DH:140:LYS:N | 2.60 | 0.55 |
| 40:DO:31:LYS:C | 40:DO:32:TYR:HD1 | 2.15 | 0.55 |
| 44:DS:17:ARG:C | 44:DS:19:LYS:N | 2.65 | 0.55 |
| 47:DV:43:GLU:O | 47:DV:44:LYS:O | 2.25 | 0.55 |
| 49:DX:33:LYS:HA | 49:DX:35:THR:HG22 | 1.89 | 0.55 |
| 1:AA:63:C:O2' | 1:AA:380:G:H4' | 2.07 | 0.54 |
| 1:AA:444:C:H2' | 1:AA:445:G:C8 | 2.35 | 0.54 |
| 1:AA:1423:G:C5' | 40:BO:49:ARG:HH21 | 2.21 | 0.54 |
| 6:AF:42:GLU:OE1 | 6:AF:59:TYR:HE2 | 1.90 | 0.54 |
| 6:AF:79:LEU:HB2 | 6:AF:88:VAL:HG21 | 1.89 | 0.54 |
| 9:AI:46:ALA:HB2 | 9:AI:74:ILE:HG23 | 1.89 | 0.54 |
| 23:B1:12:PRO:HD2 | 23:B1:62:VAL:CG2 | 2.35 | 0.54 |
| 30:B8:25:MET:HG3 | 41:BP:64:LYS:HB2 | 1.89 | 0.54 |
| 30:B8:39:LYS:O | 30:B8:39:LYS:HG2 | 2.06 | 0.54 |
| 31:BA:667:U:H2' | 31:BA:668:G:H5' | 1.89 | 0.54 |
| 31:BA:1410:G:H2' | 31:BA:1411:C:C6 | 2.39 | 0.54 |
| 31:BA:2655:G:N3 | 31:BA:2664:G:O6 | 2.40 | 0.54 |
| 39:BN:65:LYS:O | 39:BN:69:GLN:CB | 2.55 | 0.54 |
| 41:BP:17:LYS:C | 41:BP:19:VAL:N | 2.53 | 0.54 |
| 44:BS:26:LEU:HD13 | 44:BS:87:PHE:HD1 | 1.71 | 0.54 |
| 51:BZ:77:ASP:O | 51:BZ:79:ARG:N | 2.41 | 0.54 |
| 1:CA:628:G:O2' | 1:CA:629:G:H5' | 2.06 | 0.54 |
| 1:CA:658:G:C4 | 1:CA:659:U:C5 | 2.95 | 0.54 |
| 1:CA:836:G:C6 | 1:CA:851:G:C5 | 2.94 | 0.54 |
| 1:CA:1438:G:H2' | 1:CA:1439:C:C6 | 2.43 | 0.54 |
| 2:CB:83:MET:HE2 | 2:CB:83:MET:HA | 1.89 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 9:CI:114:TYR:HE1 | 10:CJ:60:ARG:O | 1.89 | 0.54 |
| 16:CP:53:VAL:O | 16:CP:57:ARG:HG2 | 2.07 | 0.54 |
| 20:CT:74:LYS:C | 20:CT:76:ALA:H | 2.15 | 0.54 |
| 28:D6:15:GLU:HG2 | 28:D6:18:ARG:NE | 2.19 | 0.54 |
| 31:DA:768:G:O2' | 31:DA:1379:A:N6 | 2.40 | 0.54 |
| 31:DA:1362:C:C2' | 31:DA:1363:C:H5' | 2.37 | 0.54 |
| 31:DA:1694:C:O2' | 31:DA:1695:G:C5 | 2.60 | 0.54 |
| 34:DE:143:ASN:HB2 | 34:DE:147:PRO:HD2 | 1.89 | 0.54 |
| 37:DH:146:ALA:O | 37:DH:147:ASN:C | 2.49 | 0.54 |
| 38:DI:10:GLU:C | 38:DI:12:LEU:H | 2.15 | 0.54 |
| 47:DV:15:GLU:O | 47:DV:98:GLU:OE2 | 2.24 | 0.54 |
| 1:AA:501:C:O2' | 1:AA:502:G:H5' | 2.07 | 0.54 |
| 1:AA:651:C:O2' | 1:AA:652:U:H5' | 2.06 | 0.54 |
| 1:AA:722:A:H2' | 1:AA:724:G:C8 | 2.41 | 0.54 |
| 1:AA:1026:G:N3 | 1:AA:1026:G:H2' | 2.21 | 0.54 |
| 1:AA:1105:A:H2' | 1:AA:1106:G:H8 | 1.73 | 0.54 |
| 1:AA:1399:C:H4' | 1:AA:1400:C:H5" | 1.90 | 0.54 |
| 3:AC:150:LYS:HE3 | 3:AC:167:TRP:HE1 | 1.72 | 0.54 |
| 5:AE:40:ARG:HH21 | 5:AE:66:MET:HE3 | 1.72 | 0.54 |
| 6:AF:8:ILE:HG22 | 6:AF:10:LEU:CD1 | 2.37 | 0.54 |
| 7:AG:115:ARG:HB2 | 7:AG:118:VAL:HG22 | 1.88 | 0.54 |
| 18:AR:63:GLN:HA | 18:AR:63:GLN:OE1 | 2.08 | 0.54 |
| 31:BA:102:G:C8 | 31:BA:102:G:C5' | 2.79 | 0.54 |
| 31:BA:173:G:C6 | 31:BA:174:C:C4 | 2.95 | 0.54 |
| 31:BA:280:C:H2' | 31:BA:281:G:O5' | 2.07 | 0.54 |
| 31:BA:643:A:C2 | 31:BA:644:A:C4 | 2.95 | 0.54 |
| 31:BA:910:A:C8 | 42:BQ:13:GLN:HB2 | 2.42 | 0.54 |
| 31:BA:2320:A:H2' | 31:BA:2320:A:N3 | 2.21 | 0.54 |
| 31:BA:2443:C:O2' | 31:BA:2444:G:H5' | 2.07 | 0.54 |
| 31:BA:2496:C:OP1 | 42:BQ:81:VAL:HG12 | 2.08 | 0.54 |
| 32:BB:102:A:H8 | 32:BB:102:A:OP2 | 1.90 | 0.54 |
| 33:BD:24:ILE:HD11 | 33:BD:84:TYR:N | 2.21 | 0.54 |
| 34:BE:77:ILE:HG23 | 34:BE:78:LEU:N | 2.20 | 0.54 |
| 41:BP:89:ALA:O | 41:BP:90:ARG:C | 2.50 | 0.54 |
| 48:BW:18:ARG:HH11 | 48:BW:18:ARG:CG | 2.14 | 0.54 |
| 1:CA:299:G:C6 | 1:CA:300:A:N1 | 2.75 | 0.54 |
| 1:CA:892:A:C6 | 1:CA:893:C:C4 | 2.95 | 0.54 |
| 1:CA:1242:C:P | 21:CU:10:ARG:HH22 | 2.30 | 0.54 |
| 9:CI:46:ALA:HB2 | 9:CI:74:ILE:HG23 | 1.89 | 0.54 |
| 23:D1:10:LYS:HD3 | 23:D1:14:VAL:HA | 1.89 | 0.54 |
| 24:D2:41:ILE:O | 24:D2:43:GLN:N | 2.40 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:241:A:O4' | 31:DA:243:U:C6 | 2.60 | 0.54 |
| 31:DA:610:G:H2' | 31:DA:611:C:C6 | 2.42 | 0.54 |
| 31:DA:806:C:P | 41:DP:39:LYS:HG3 | 2.47 | 0.54 |
| 31:DA:1116:C:C2' | 31:DA:1117:G:H5' | 2.37 | 0.54 |
| 31:DA:2320:A:N3 | 31:DA:2320:A:H2' | 2.22 | 0.54 |
| 31:DA:2475:C:N4 | 31:DA:2529:G:H22 | 2.01 | 0.54 |
| 31:DA:2484:G:C2 | 31:DA:2485:G:C8 | 2.95 | 0.54 |
| 33:DD:27:THR:HG22 | 33:DD:28:GLU:H | 1.72 | 0.54 |
| 35:DF:24:LEU:O | 35:DF:25:PRO:C | 2.47 | 0.54 |
| 39:DN:67:LEU:C | 39:DN:69:GLN:N | 2.65 | 0.54 |
| 39:DN:77:GLY:O | 39:DN:78:TYR:HB3 | 2.05 | 0.54 |
| 40:DO:35:VAL:HG13 | 40:DO:65:THR:CG2 | 2.37 | 0.54 |
| 40:DO:50:GLY:C | 40:DO:52:VAL:H | 2.16 | 0.54 |
| 45:DT:30:VAL:HG22 | 45:DT:84:GLN:O | 2.08 | 0.54 |
| 46:DU:8:VAL:HG13 | 46:DU:12:ARG:HG3 | 1.90 | 0.54 |
| 49:DX:65:ARG:NH1 | 49:DX:66:LEU:N | 2.54 | 0.54 |
| 50:DY:44:ILE:HD12 | 50:DY:44:ILE:N | 2.21 | 0.54 |
| 51:DZ:8:TYR:O | 51:DZ:37:VAL:HG12 | 2.08 | 0.54 |
| 1:AA:881:G:OP2 | 12:AL:12:ARG:NH2 | 2.38 | 0.54 |
| 5:AE:100:VAL:HG13 | 5:AE:118:ILE:CG2 | 2.37 | 0.54 |
| 8:AH:87:SER:HA | 8:AH:93:VAL:HG23 | 1.89 | 0.54 |
| 13:AM:68:GLY:O | 13:AM:69:GLU:CB | 2.54 | 0.54 |
| 31:BA:330:A:H2 | 31:BA:1210:A:H2' | 1.73 | 0.54 |
| 31:BA:477:A:O2' | 31:BA:478:A:H5' | 2.06 | 0.54 |
| 31:BA:614(A):U:H4' | 31:BA:614(B):G:H5" | 1.90 | 0.54 |
| 31:BA:754:C:O4' | 31:BA:1618:A:H2 | 1.90 | 0.54 |
| 31:BA:1210:A:C5' | 31:BA:1211:U:H3' | 2.37 | 0.54 |
| 31:BA:1210:A:C4' | 31:BA:1211:U:OP2 | 2.55 | 0.54 |
| 31:BA:2061:G:N2 | 31:BA:2063:C:C2 | 2.75 | 0.54 |
| 31:BA:2801(A):A:C4' | 31:BA:2802:G:H5' | 2.37 | 0.54 |
| 33:BD:270:ILE:C | 33:BD:271:ILE:HG13 | 2.32 | 0.54 |
| 35:BF:3:GLU:O | 35:BF:24:LEU:HG | 2.07 | 0.54 |
| 35:BF:195:ASP:CG | 35:BF:197:ASP:HB3 | 2.31 | 0.54 |
| 36:BG:6:ALA:HB3 | 36:BG:104:GLU:OE1 | 2.07 | 0.54 |
| 42:BQ:108:GLY:C | 42:BQ:109:VAL:HG22 | 2.32 | 0.54 |
| 1:CA:84:U:H5 | 1:CA:88:A:C8 | 2.24 | 0.54 |
| 3:CC:95:THR:HG22 | 3:CC:97:LYS:H | 1.71 | 0.54 |
| 4:CD:33:MET:HA | 4:CD:33:MET:HE3 | 1.89 | 0.54 |
| 8:CH:10:LEU:N | 8:CH:10:LEU:CD2 | 2.70 | 0.54 |
| 10:CJ:32:ALA:HB1 | 10:CJ:75:ILE:HD11 | 1.90 | 0.54 |
| 18:CR:74:ARG:HB3 | 18:CR:81:PHE:CE1 | 2.42 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 23:D1:19:GLN:HE21 | 31:DA:379:G:N2 | 1.94 | 0.54 |
| 24:D2:30:ARG:CZ | 24:D2:30:ARG:HB3 | 2.36 | 0.54 |
| 31:DA:197:A:C8 | 31:DA:197:A:C4' | 2.90 | 0.54 |
| 31:DA:661:C:H4' | 41:DP:18:ARG:HG2 | 1.90 | 0.54 |
| 31:DA:814:C:C5' | 47:DV:86:GLY:HA3 | 2.34 | 0.54 |
| 31:DA:1528:A:O2' | 31:DA:1528(A):A:O5' | 2.24 | 0.54 |
| 31:DA:1885:A:H2' | 31:DA:1886:C:O4' | 2.08 | 0.54 |
| 31:DA:2016:U:H2' | 31:DA:2017:U:H6 | 1.70 | 0.54 |
| 31:DA:2610:C:H4' | 31:DA:2611:U:OP2 | 2.05 | 0.54 |
| 33:DD:39:LYS:HB2 | 33:DD:62:TYR:CB | 2.31 | 0.54 |
| 37:DH:148:ILE:O | 37:DH:151:ILE:HG12 | 2.07 | 0.54 |
| 49:DX:34:ALA:O | 49:DX:36:LYS:HE3 | 2.07 | 0.54 |
| 51:DZ:29:TYR:HE2 | 51:DZ:87:ASP:CB | 2.21 | 0.54 |
| 51:DZ:141:VAL:HG23 | 51:DZ:144:LEU:HD23 | 1.88 | 0.54 |
| 1:AA:272:C:H2' | 1:AA:273:A:H8 | 1.72 | 0.54 |
| 1:AA:1362:C:O2' | 1:AA:1363:C:H5" | 2.06 | 0.54 |
| 1:AA:1410:G:C4 | 1:AA:1491:G:N2 | 2.76 | 0.54 |
| 2:AB:121:LEU:O | 2:AB:121:LEU:HD23 | 2.07 | 0.54 |
| 3:AC:157:ILE:CD1 | 3:AC:166:GLU:HB2 | 2.38 | 0.54 |
| 6:AF:99:ALA:HB1 | 18:AR:23:LYS:HZ2 | 1.73 | 0.54 |
| 12:AL:25:PRO:C | 12:AL:27:LEU:H | 2.16 | 0.54 |
| 27:B5:36:CYS:C | 27:B5:38:ALA:H | 2.15 | 0.54 |
| 27:B5:52:TYR:O | 27:B5:52:TYR:CD1 | 2.61 | 0.54 |
| 29:B7:38:GLY:O | 29:B7:39:ARG:C | 2.50 | 0.54 |
| 30:B8:2:PRO:N | 31:BA:591:C:O2 | 2.40 | 0.54 |
| 31:BA:157:U:H4' | 31:BA:171:G:N2 | 2.23 | 0.54 |
| 31:BA:542:C:N4 | 31:BA:543:C:N4 | 2.55 | 0.54 |
| 31:BA:1593:G:H2' | 31:BA:1594:G:C8 | 2.42 | 0.54 |
| 31:BA:1796:U:H2' | 31:BA:1797:C:C6 | 2.42 | 0.54 |
| 31:BA:2292:C:C2' | 31:BA:2293:C:H5' | 2.37 | 0.54 |
| 31:BA:2402:C:C3' | 31:BA:2403:C:H5' | 2.37 | 0.54 |
| 31:BA:2416:C:O5' | 31:BA:2416:C:H6 | 1.90 | 0.54 |
| 31:BA:2661:G:O2' | 31:BA:2662:A:O5' | 2.20 | 0.54 |
| 32:BB:46:A:C5 | 32:BB:47:C:C5 | 2.95 | 0.54 |
| 33:BD:25:THR:O | 33:BD:26:LYS:C | 2.50 | 0.54 |
| 38:BI:51:ILE:HG22 | 38:BI:51:ILE:O | 2.06 | 0.54 |
| 39:BN:68:GLU:HA | 39:BN:86:PRO:CB | 2.37 | 0.54 |
| 41:BP:35:HIS:O | 41:BP:36:LYS:CB | 2.55 | 0.54 |
| 41:BP:108:LYS:O | 41:BP:110:TYR:N | 2.41 | 0.54 |
| 49:BX:58:HIS:O | 49:BX:59:VAL:HG13 | 2.07 | 0.54 |
| 51:BZ:98:MET:HE3 | 51:BZ:99:TYR:O | 2.07 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:651:C:O2' | 1:CA:652:U:H5' | 2.07 | 0.54 |
| 1:CA:722:A:H2' | 1:CA:724:G:C8 | 2.42 | 0.54 |
| 1:CA:836:G:C6 | 1:CA:851:G:C6 | 2.95 | 0.54 |
| 1:CA:1072:G:C6 | 1:CA:1073:U:C4 | 2.95 | 0.54 |
| 8:CH:48:TYR:HA | 8:CH:60:ARG:O | 2.07 | 0.54 |
| 9:CI:53:VAL:C | 9:CI:92:TYR:HE2 | 2.15 | 0.54 |
| 12:CL:60:LEU:HD21 | 12:CL:66:VAL:CG2 | 2.36 | 0.54 |
| 16:CP:34:GLU:OE2 | 16:CP:55:ARG:HD3 | 2.07 | 0.54 |
| 18:CR:73:ALA:HB1 | 18:CR:79:LEU:HD12 | 1.89 | 0.54 |
| 23:D1:19:GLN:CG | 23:D1:44:PRO:HG3 | 2.33 | 0.54 |
| 24:D2:41:ILE:HG21 | 31:DA:95:G:H21 | 1.71 | 0.54 |
| 31:DA:52:A:C2' | 31:DA:53:A:H5' | 2.36 | 0.54 |
| 31:DA:128:C:C6 | 31:DA:128:C:C3' | 2.91 | 0.54 |
| 31:DA:221:A:H4' | 31:DA:222:A:O5' | 2.07 | 0.54 |
| 31:DA:790:C:O2' | 31:DA:791:C:H5' | 2.08 | 0.54 |
| 31:DA:814:C:O2' | 31:DA:815:C:H5' | 2.07 | 0.54 |
| 31:DA:866:A:C6 | 31:DA:914:C:C6 | 2.96 | 0.54 |
| 31:DA:1300:U:H3' | 31:DA:1301:A:H5" | 1.88 | 0.54 |
| 31:DA:1497:U:H3 | 31:DA:1578:U:P | 2.30 | 0.54 |
| 31:DA:1545:A:H2' | 31:DA:1546:C:C5' | 2.37 | 0.54 |
| 31:DA:2193:G:H2' | 31:DA:2194:G:O4' | 2.08 | 0.54 |
| 31:DA:2302:G:C6 | 31:DA:2315:G:C6 | 2.96 | 0.54 |
| 31:DA:2850:A:OP2 | 31:DA:2866:U:H5 | 1.88 | 0.54 |
| 39:DN:65:LYS:O | 39:DN:69:GLN:CB | 2.53 | 0.54 |
| 40:DO:112:MET:HA | 40:DO:112:MET:CE | 2.38 | 0.54 |
| 41:DP:29:LYS:HD2 | 41:DP:29:LYS:N | 2.05 | 0.54 |
| 48:DW:16:LYS:O | 48:DW:19:LEU:HB2 | 2.08 | 0.54 |
| 1:AA:1240:U:P | 7:AG:116:ALA:HB2 | 2.48 | 0.54 |
| 1:AA:1242:C:P | 21:AU:10:ARG:HH22 | 2.29 | 0.54 |
| 3:AC:112:SER:O | 3:AC:116:VAL:HG23 | 2.07 | 0.54 |
| 11:AK:23:ALA:HB3 | 11:AK:86:GLY:O | 2.08 | 0.54 |
| 11:AK:27:ASN:OD1 | 11:AK:28:THR:N | 2.41 | 0.54 |
| 13:AM:54:VAL:HG22 | 13:AM:57:ARG:HH21 | 1.72 | 0.54 |
| 15:AO:4:THR:OG1 | 15:AO:7:GLU:HB2 | 2.07 | 0.54 |
| 31:BA:157:U:H5' | 31:BA:171:G:H22 | 1.72 | 0.54 |
| 31:BA:493:G:C2' | 31:BA:494:G:H5" | 2.35 | 0.54 |
| 31:BA:1040:C:O2' | 31:BA:1041:C:OP2 | 2.20 | 0.54 |
| 31:BA:1803:A:C8 | 31:BA:1804:C:C5 | 2.95 | 0.54 |
| 31:BA:1899:G:N2 | 31:BA:1902:C:N4 | 2.37 | 0.54 |
| 31:BA:2314:C:O2 | 31:BA:2315:G:C8 | 2.61 | 0.54 |
| 33:BD:35:LYS:HB3 | 33:BD:63:ARG:HA | 1.88 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 39:BN:23:LEU:CD1 | 39:BN:98:VAL:HG12 | 2.37 | 0.54 |
| 39:BN:23:LEU:HD13 | 39:BN:98:VAL:HG12 | 1.89 | 0.54 |
| 39:BN:65:LYS:O | 39:BN:69:GLN:HG3 | 2.07 | 0.54 |
| 40:BO:2:ILE:N | 40:BO:2:ILE:HD13 | 2.22 | 0.54 |
| 1:CA:79:G:C4' | 1:CA:80:G:OP1 | 2.55 | 0.54 |
| 1:CA:228:A:H2' | 1:CA:229:U:O4' | 2.08 | 0.54 |
| 1:CA:579:G:C5 | 1:CA:580:U:C5 | 2.95 | 0.54 |
| 1:CA:826:C:H4' | 8:CH:12:ARG:HD3 | 1.89 | 0.54 |
| 1:CA:1126:U:H2' | 1:CA:1127:G:O4' | 2.08 | 0.54 |
| 1:CA:1168:A:C6 | 1:CA:1169:A:C6 | 2.95 | 0.54 |
| 6:CF:5:GLU:HG3 | 6:CF:93:SER:OG | 2.07 | 0.54 |
| 7:CG:151:TYR:OH | 11:CK:54:ARG:HD3 | 2.08 | 0.54 |
| 15:CO:4:THR:OG1 | 15:CO:7:GLU:HB2 | 2.07 | 0.54 |
| 22:D0:53:MET:HB2 | 22:D0:59:LEU:HD23 | 1.89 | 0.54 |
| 31:DA:157:U:H5' | 31:DA:171:G:H22 | 1.73 | 0.54 |
| 31:DA:606:U:H4' | 31:DA:658:C:H4' | 1.89 | 0.54 |
| 31:DA:631:A:OP1 | 41:DP:64:LYS:CE | 2.53 | 0.54 |
| 31:DA:1050:A:C2 | 31:DA:2751:G:C4 | 2.95 | 0.54 |
| 31:DA:2593:U:H2' | 31:DA:2594:C:C6 | 2.43 | 0.54 |
| 31:DA:2859:G:H4' | 31:DA:2860:A:OP1 | 2.08 | 0.54 |
| 32:DB:40:U:H1' | 32:DB:45:A:H61 | 1.72 | 0.54 |
| 33:DD:8:PRO:HB3 | 33:DD:14:ARG:CB | 2.37 | 0.54 |
| 33:DD:83:GLU:HB2 | 33:DD:92:ILE:HD11 | 1.89 | 0.54 |
| 34:DE:181:LEU:HD21 | 45:DT:7:ILE:HG23 | 1.89 | 0.54 |
| 35:DF:10:PRO:HG2 | 35:DF:13:SER:OG | 2.08 | 0.54 |
| 38:DI:95:LYS:O | 38:DI:99:GLU:HB2 | 2.08 | 0.54 |
| 42:DQ:75:THR:HA | 42:DQ:89:ASN:H | 1.72 | 0.54 |
| 42:DQ:141:GLN:HE21 | 51:DZ:71:VAL:C | 2.16 | 0.54 |
| 46:DU:92:ARG:O | 46:DU:93:LYS:C | 2.49 | 0.54 |
| 48:DW:92:ARG:HH11 | 48:DW:92:ARG:HG2 | 1.72 | 0.54 |
| 49:DX:59:VAL:HG22 | 49:DX:74:PRO:O | 2.08 | 0.54 |
| 1:AA:577:G:C8 | 1:AA:816:A:C6 | 2.96 | 0.54 |
| 1:AA:706:A:C8 | 1:AA:707:C:H5 | 2.26 | 0.54 |
| 1:AA:738:C:H2' | 1:AA:739:C:H6 | 1.72 | 0.54 |
| 2:AB:80:ILE:HD13 | 2:AB:208:ILE:HG23 | 1.89 | 0.54 |
| 3:AC:69:HIS:CD2 | 3:AC:69:HIS:N | 2.75 | 0.54 |
| 4:AD:104:VAL:O | 4:AD:108:LEU:HD13 | 2.08 | 0.54 |
| 6:AF:3:ARG:HH11 | 6:AF:3:ARG:HG3 | 1.72 | 0.54 |
| 7:AG:27:ILE:HD11 | 7:AG:43:PHE:CD2 | 2.42 | 0.54 |
| 31:BA:435:C:C5 | 31:BA:436:C:C5 | 2.95 | 0.54 |
| 31:BA:443:A:H1' | 31:BA:1201:C:O4' | 2.07 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:743:G:C2' | 31:BA:744:G:H5' | 2.38 | 0.54 |
| 31:BA:1582:C:O2' | 31:BA:1586:A:H8 | 1.91 | 0.54 |
| 31:BA:1635:G:H5' | 31:BA:1635:G:C8 | 2.42 | 0.54 |
| 31:BA:1822:G:C5' | 31:BA:1822:G:C8 | 2.88 | 0.54 |
| 34:BE:114:ALA:HB3 | 34:BE:160:TYR:HB3 | 1.89 | 0.54 |
| 35:BF:170:LEU:HD23 | 35:BF:172:TRP:NE1 | 2.22 | 0.54 |
| 36:BG:111:LEU:O | 36:BG:114:ILE:HG12 | 2.08 | 0.54 |
| 37:BH:144:VAL:HA | 37:BH:147:ASN:OD1 | 2.08 | 0.54 |
| 39:BN:19:GLU:HG3 | 39:BN:20:GLY:N | 2.22 | 0.54 |
| 41:BP:97:PRO:O | 41:BP:98:GLU:CB | 2.54 | 0.54 |
| 43:BR:41:ALA:O | 43:BR:42:LYS:C | 2.51 | 0.54 |
| 47:BV:54:GLY:C | 47:BV:56:SER:H | 2.13 | 0.54 |
| 51:BZ:153:SER:O | 51:BZ:154:ASP:OD2 | 2.25 | 0.54 |
| 1:CA:11:G:C5 | 1:CA:12:U:C5 | 2.95 | 0.54 |
| 1:CA:191:G:H1' | 20:CT:105:SER:HA | 1.90 | 0.54 |
| 1:CA:266:G:H5" | 1:CA:268:C:N4 | 2.19 | 0.54 |
| 1:CA:356:A:C2' | 1:CA:357:G:O5' | 2.55 | 0.54 |
| 1:CA:760:G:H2' | 1:CA:761:G:H5' | 1.89 | 0.54 |
| 1:CA:1026:G:H2' | 1:CA:1026:G:N3 | 2.23 | 0.54 |
| 1:CA:1071:C:O2' | 1:CA:1072:G:H5' | 2.08 | 0.54 |
| 2:CB:114:ARG:O | 2:CB:114:ARG:HD3 | 2.08 | 0.54 |
| 3:CC:182:ILE:HG12 | 3:CC:203:PHE:HA | 1.90 | 0.54 |
| 5:CE:40:ARG:HH21 | 5:CE:66:MET:HE3 | 1.71 | 0.54 |
| 13:CM:54:VAL:HG22 | 13:CM:57:ARG:HH21 | 1.73 | 0.54 |
| 20:CT:13:LEU:HD12 | 20:CT:13:LEU:H | 1.73 | 0.54 |
| 20:CT:26:ASN:HB2 | 20:CT:71:THR:HG23 | 1.89 | 0.54 |
| 20:CT:89:ARG:HH21 | 20:CT:104:LEU:HD21 | 1.68 | 0.54 |
| 22:D0:77:ARG:NH2 | 31:DA:857:C:H5' | 2.22 | 0.54 |
| 23:D1:89:GLU:CD | 23:D1:89:GLU:N | 2.53 | 0.54 |
| 31:DA:443:A:H1' | 31:DA:1201:C:O4' | 2.08 | 0.54 |
| 31:DA:828:U:O2' | 31:DA:829:A:H5' | 2.08 | 0.54 |
| 31:DA:933:A:H2' | 31:DA:934:G:H5' | 1.90 | 0.54 |
| 31:DA:993:G:N3 | 47:DV:91:TYR:HE1 | 2.05 | 0.54 |
| 31:DA:1404:C:H2' | 31:DA:1404:C:O2 | 2.06 | 0.54 |
| 31:DA:1412:A:H2' | 31:DA:1413:G:C8 | 2.43 | 0.54 |
| 31:DA:2312:U:OP1 | 36:DG:74:LYS:HG3 | 2.07 | 0.54 |
| 31:DA:2517:C:C6 | 31:DA:2542:A:N1 | 2.76 | 0.54 |
| 31:DA:2536:G:C6 | 31:DA:2537:U:C4 | 2.95 | 0.54 |
| 33:DD:35:LYS:HZ2 | 33:DD:64:ILE:C | 2.15 | 0.54 |
| 34:DE:116:VAL:HG13 | 34:DE:122:PHE:CD2 | 2.42 | 0.54 |
| 41:DP:57:THR:HB | 41:DP:59:LEU:H | 1.71 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 48:DW:64:MET:HE2 | 48:DW:69:LEU:HD23 | 1.89 | 0.54 |
| 50:DY:89:PHE:O | 50:DY:90:LEU:HB3 | 2.08 | 0.54 |
| 51:DZ:8:TYR:N | 51:DZ:8:TYR:CD1 | 2.75 | 0.54 |
| 1:AA:113:G:H2' | 1:AA:114:U:C6 | 2.42 | 0.54 |
| 1:AA:521:G:H4' | 12:AL:73:GLU:HG2 | 1.90 | 0.54 |
| 1:AA:650:G:O2' | 1:AA:651:C:H5' | 2.07 | 0.54 |
| 1:AA:826:C:H4' | 8:AH:12:ARG:HD3 | 1.90 | 0.54 |
| 1:AA:946:A:C2 | 1:AA:1236:A:C2 | 2.96 | 0.54 |
| 1:AA:1442:G:O2' | 1:AA:1442(A):G:C5' | 2.52 | 0.54 |
| 3:AC:156:ARG:NH2 | 3:AC:161:GLU:HA | 2.22 | 0.54 |
| 5:AE:12:LEU:C | 5:AE:12:LEU:HD22 | 2.33 | 0.54 |
| 15:AO:18:PHE:CE1 | 15:AO:21:ASP:HB2 | 2.43 | 0.54 |
| 24:B2:12:GLU:CD | 24:B2:12:GLU:C | 2.76 | 0.54 |
| 24:B2:32:LEU:HD12 | 24:B2:32:LEU:O | 2.08 | 0.54 |
| 31:BA:782:A:H5' | 31:BA:783:A:C2 | 2.43 | 0.54 |
| 31:BA:947:G:N2 | 31:BA:971:C:C2 | 2.76 | 0.54 |
| 31:BA:1146:C:H2' | 31:BA:1147:C:C5' | 2.35 | 0.54 |
| 31:BA:1278:A:O2' | 43:BR:34:ILE:HD11 | 2.07 | 0.54 |
| 31:BA:1489:U:H2' | 31:BA:1490:A:OP2 | 2.08 | 0.54 |
| 31:BA:1545:A:H2' | 31:BA:1546:C:H5' | 1.88 | 0.54 |
| 31:BA:2190:G:H2' | 31:BA:2191:G:H5' | 1.90 | 0.54 |
| 31:BA:2299:G:N1 | 31:BA:2318:G:C8 | 2.76 | 0.54 |
| 31:BA:2317:C:H2' | 31:BA:2317:C:O2 | 2.06 | 0.54 |
| 31:BA:2870:C:H5" | 43:BR:65:LEU:HD21 | 1.90 | 0.54 |
| 33:BD:89:SER:HB2 | 33:BD:159:ALA:CB | 2.38 | 0.54 |
| 33:BD:97:TYR:HB2 | 33:BD:101:GLU:O | 2.07 | 0.54 |
| 35:BF:89:VAL:CG1 | 35:BF:90:PHE:N | 2.63 | 0.54 |
| 35:BF:178:PRO:HG2 | 35:BF:179:GLU:OE1 | 2.07 | 0.54 |
| 41:BP:91:PHE:HE2 | 41:BP:95:VAL:CG1 | 2.21 | 0.54 |
| 42:BQ:134:ARG:HH21 | 51:BZ:122:ARG:CZ | 2.20 | 0.54 |
| 49:BX:82:GLN:CD | 49:BX:83:VAL:H | 2.16 | 0.54 |
| 1:CA:343:U:O2' | 1:CA:346:G:O6 | 2.26 | 0.54 |
| 1:CA:1132:C:H2' | 1:CA:1133:G:O4' | 2.08 | 0.54 |
| 1:CA:1315:U:H2' | 1:CA:1316:G:O4' | 2.08 | 0.54 |
| 2:CB:20:GLU:O | 2:CB:40:HIS:HB2 | 2.07 | 0.54 |
| 3:CC:73:PRO:HA | 3:CC:76:VAL:HG13 | 1.89 | 0.54 |
| 4:CD:80:GLU:O | 4:CD:84:LYS:HG2 | 2.08 | 0.54 |
| 8:CH:36:LEU:HD23 | 8:CH:39:LEU:HD23 | 1.89 | 0.54 |
| 8:CH:112:LEU:HB2 | 8:CH:133:LEU:HA | 1.89 | 0.54 |
| 30:D8:62:LEU:O | 30:D8:64:TYR:N | 2.40 | 0.54 |
| 31:DA:17:G:H4' | 46:DU:25:TRP:CH2 | 2.42 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:236:C:H2' | 31:DA:237:C:C6 | 2.43 | 0.54 |
| 31:DA:1169:G:H3' | 31:DA:1169:G:C8 | 2.43 | 0.54 |
| 31:DA:2186:G:H2' | 31:DA:2187:G:H5" | 1.89 | 0.54 |
| 31:DA:2661:G:O2' | 31:DA:2662:A:O5' | 2.19 | 0.54 |
| 31:DA:2894:G:H2' | 31:DA:2894:G:N3 | 2.21 | 0.54 |
| 33:DD:145:VAL:HG11 | 33:DD:175:LEU:HD11 | 1.90 | 0.54 |
| 35:DF:22:ALA:C | 35:DF:26:ALA:HB2 | 2.33 | 0.54 |
| 39:DN:27:ALA:HB3 | 39:DN:106:MET:HE2 | 1.89 | 0.54 |
| 39:DN:90:MET:O | 39:DN:93:THR:O | 2.26 | 0.54 |
| 41:DP:67:MET:HA | 41:DP:67:MET:HE3 | 1.90 | 0.54 |
| 42:DQ:18:LYS:O | 42:DQ:19:GLY:C | 2.50 | 0.54 |
| 1:AA:60:A:H4' | 1:AA:61:G:O5' | 2.07 | 0.54 |
| 1:AA:411:A:C6 | 1:AA:429:U:C4 | 2.95 | 0.54 |
| 1:AA:661:G:C2 | 1:AA:662:G:C8 | 2.96 | 0.54 |
| 1:AA:892:A:C5 | 1:AA:893:C:C4 | 2.96 | 0.54 |
| 2:AB:9:GLU:CD | 2:AB:9:GLU:H | 2.15 | 0.54 |
| 10:AJ:82:ILE:O | 10:AJ:82:ILE:HG22 | 2.08 | 0.54 |
| 12:AL:21:LYS:HD2 | 12:AL:21:LYS:H | 1.71 | 0.54 |
| 15:AO:78:TYR:OH | 15:AO:88:ARG:HD2 | 2.08 | 0.54 |
| 17:AQ:65:ILE:HD12 | 17:AQ:65:ILE:N | 2.23 | 0.54 |
| 30:B8:62:LEU:CD1 | 31:BA:242:G:H5" | 2.28 | 0.54 |
| 31:BA:1268:A:C2 | 31:BA:2013:A:C4 | 2.96 | 0.54 |
| 31:BA:1349:A:H2' | 31:BA:1350:C:OP1 | 2.08 | 0.54 |
| 31:BA:1386:C:H2' | 31:BA:1387:C:H6 | 1.73 | 0.54 |
| 31:BA:1684:C:O2' | 31:BA:1685:C:H5' | 2.08 | 0.54 |
| 31:BA:2376:A:OP1 | 31:BA:2376:A:H8 | 1.91 | 0.54 |
| 31:BA:2593:U:H2' | 31:BA:2594:C:C6 | 2.43 | 0.54 |
| 35:BF:84:VAL:O | 35:BF:85:GLY:C | 2.49 | 0.54 |
| 39:BN:32:THR:O | 39:BN:35:ARG:O | 2.24 | 0.54 |
| 44:BS:27:SER:HB3 | 44:BS:89:ARG:NH1 | 2.22 | 0.54 |
| 1:CA:357:G:O2' | 1:CA:358:U:H5' | 2.07 | 0.54 |
| 1:CA:738:C:H2' | 1:CA:739:C:H6 | 1.72 | 0.54 |
| 1:CA:932:C:H4' | 7:CG:4:ARG:NH2 | 2.23 | 0.54 |
| 1:CA:1392:G:N2 | 1:CA:1502:A:H8 | 2.06 | 0.54 |
| 1:CA:1470:G:C2' | 1:CA:1471:G:H5' | 2.38 | 0.54 |
| 6:CF:55:ASP:HB2 | 6:CF:86:ARG:HH12 | 1.73 | 0.54 |
| 12:CL:102:ARG:HG3 | 12:CL:102:ARG:NH1 | 2.23 | 0.54 |
| 24:D2:14:ARG:O | 24:D2:18:PRO:CD | 2.50 | 0.54 |
| 25:D3:8:LEU:HD13 | 25:D3:31:LEU:HA | 1.90 | 0.54 |
| 27:D5:52:TYR:O | 27:D5:52:TYR:CD1 | 2.61 | 0.54 |
| 30:D8:38:GLY:C | 30:D8:40:GLU:H | 2.14 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 30:D8:39:LYS:HE2 | 30:D8:39:LYS:O | 2.08 | 0.54 |
| 31:DA:536:A:H2' | 31:DA:537:C:C6 | 2.43 | 0.54 |
| 31:DA:670:A:H4' | 31:DA:671:C:OP1 | 2.07 | 0.54 |
| 31:DA:986:C:C2' | 31:DA:987:G:H5' | 2.37 | 0.54 |
| 31:DA:1170:G:H1 | 31:DA:1179:C:N4 | 2.00 | 0.54 |
| 31:DA:1319:G:C6 | 31:DA:1320:C:N4 | 2.76 | 0.54 |
| 31:DA:1669:A:H5" | 31:DA:2550:G:OP1 | 2.08 | 0.54 |
| 31:DA:1801:G:OP2 | 33:DD:154:LYS:HE3 | 2.08 | 0.54 |
| 31:DA:1902:C:H2' | 31:DA:1903:G:O5' | 2.08 | 0.54 |
| 31:DA:2472:G:C2 | 31:DA:2477:C:OP1 | 2.61 | 0.54 |
| 31:DA:2758:A:C2' | 31:DA:2759:G:H5" | 2.38 | 0.54 |
| 33:DD:70:TRP:CD1 | 33:DD:70:TRP:C | 2.86 | 0.54 |
| 35:DF:46:ARG:CG | 35:DF:46:ARG:NH1 | 2.67 | 0.54 |
| 41:DP:16:ARG:HD3 | 41:DP:18:ARG:HB2 | 1.89 | 0.54 |
| 43:DR:116:LEU:O | 43:DR:117:VAL:CB | 2.53 | 0.54 |
| 44:DS:26:LEU:O | 44:DS:88:ASP:HB3 | 2.06 | 0.54 |
| 49:DX:63:LYS:HZ1 | 49:DX:70:LEU:CD2 | 2.20 | 0.54 |
| 50:DY:10:GLY:O | 50:DY:27:VAL:HG22 | 2.08 | 0.54 |
| 51:DZ:19:ARG:HG2 | 51:DZ:19:ARG:NH1 | 2.23 | 0.54 |
| 1:AA:472:A:C4' | 16:AP:82:GLN:HE22 | 2.20 | 0.54 |
| 1:AA:564:C:C2' | 1:AA:565:U:H5' | 2.38 | 0.54 |
| 1:AA:683:G:C2 | 1:AA:708:C:N3 | 2.76 | 0.54 |
| 1:AA:801:U:H2' | 1:AA:802:A:H8 | 1.73 | 0.54 |
| 1:AA:1084:G:OP1 | 1:AA:1086:U:C4 | 2.61 | 0.54 |
| 1:AA:1169:A:C2' | 1:AA:1170:A:H8 | 2.18 | 0.54 |
| 2:AB:98:LEU:HB2 | 2:AB:101:MET:HE3 | 1.90 | 0.54 |
| 9:AI:114:TYR:HE1 | 10:AJ:60:ARG:O | 1.91 | 0.54 |
| 12:AL:27:LEU:O | 12:AL:29:GLY:N | 2.41 | 0.54 |
| 12:AL:119:LYS:HB2 | 12:AL:120:TYR:CD1 | 2.43 | 0.54 |
| 13:AM:10:PRO:CB | 13:AM:18:ALA:HB1 | 2.36 | 0.54 |
| 16:AP:18:ARG:O | 16:AP:19:ILE:C | 2.50 | 0.54 |
| 20:AT:26:ASN:HB2 | 20:AT:71:THR:HG23 | 1.89 | 0.54 |
| 22:B0:53:MET:HB2 | 22:B0:59:LEU:CD2 | 2.36 | 0.54 |
| 23:B1:41:ARG:NH1 | 23:B1:41:ARG:CG | 2.50 | 0.54 |
| 31:BA:272(J):C:C2' | 31:BA:274:G:OP1 | 2.56 | 0.54 |
| 31:BA:669:G:C8 | 31:BA:669:G:O2' | 2.50 | 0.54 |
| 31:BA:954:G:C5 | 31:BA:955:C:C5 | 2.95 | 0.54 |
| 31:BA:1225:G:OP1 | 47:BV:88:ARG:CB | 2.55 | 0.54 |
| 31:BA:2033:A:H4' | 31:BA:2034:U:OP1 | 2.07 | 0.54 |
| 32:BB:15:A:C2' | 32:BB:16:G:OP1 | 2.56 | 0.54 |
| 32:BB:75:G:H8 | 32:BB:75:G:C5' | 2.21 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 33:BD:35:LYS:CE | 33:BD:104:TYR:CD1 | 2.89 | 0.54 |
| 33:BD:48:ARG:NH1 | 33:BD:48:ARG:HG3 | 2.22 | 0.54 |
| 35:BF:10:PRO:HG2 | 35:BF:13:SER:OG | 2.08 | 0.54 |
| 37:BH:141:VAL:HG12 | 37:BH:142:GLY:N | 2.23 | 0.54 |
| 38:BI:5:LEU:O | 38:BI:6:LEU:HD23 | 2.08 | 0.54 |
| 40:BO:35:VAL:HG13 | 40:BO:65:THR:HG22 | 1.89 | 0.54 |
| 41:BP:57:THR:HB | 41:BP:59:LEU:H | 1.71 | 0.54 |
| 41:BP:140:ALA:O | 41:BP:141:ALA:CB | 2.56 | 0.54 |
| 44:BS:17:ARG:C | 44:BS:19:LYS:H | 2.16 | 0.54 |
| 45:BT:29:ARG:HG2 | 45:BT:86:ILE:N | 2.17 | 0.54 |
| 46:BU:92:ARG:CB | 47:BV:11:GLN:NE2 | 2.65 | 0.54 |
| 48:BW:97:LYS:HE2 | 48:BW:99:ARG:NH2 | 2.23 | 0.54 |
| 1:CA:84:U:C5 | 1:CA:88:A:C8 | 2.96 | 0.54 |
| 1:CA:804:U:H5” | 1:CA:805:C:OP2 | 2.07 | 0.54 |
| 1:CA:1015:A:N6 | 1:CA:1016:A:C6 | 2.76 | 0.54 |
| 1:CA:1216:G:H5” | 14:CN:5:ALA:HB2 | 1.89 | 0.54 |
| 1:CA:1334:G:OP2 | 1:CA:1334:G:H8 | 1.91 | 0.54 |
| 2:CB:189:ASP:OD1 | 2:CB:189:ASP:N | 2.41 | 0.54 |
| 3:CC:6:HIS:NE2 | 3:CC:184:TYR:CE2 | 2.76 | 0.54 |
| 27:D5:4:HIS:HD2 | 31:DA:2056:G:H1 | 1.55 | 0.54 |
| 28:D6:40:CYS:SG | 28:D6:45:LYS:CD | 2.96 | 0.54 |
| 31:DA:1006:C:C2 | 31:DA:1138:G:N2 | 2.75 | 0.54 |
| 31:DA:1198:U:H2’ | 31:DA:1199:U:H6 | 1.73 | 0.54 |
| 31:DA:1213:A:O2’ | 31:DA:1214:A:H5’ | 2.08 | 0.54 |
| 31:DA:1278:A:O2’ | 43:DR:34:ILE:HD11 | 2.07 | 0.54 |
| 31:DA:1290:C:H2’ | 31:DA:1291:C:H6 | 1.72 | 0.54 |
| 31:DA:1767:C:O2’ | 31:DA:1768:U:H5’ | 2.08 | 0.54 |
| 31:DA:2061:G:C2 | 31:DA:2063:C:C4 | 2.96 | 0.54 |
| 31:DA:2287:A:C4 | 31:DA:2289:G:C8 | 2.95 | 0.54 |
| 31:DA:2808:U:C2’ | 31:DA:2809:A:H5’ | 2.38 | 0.54 |
| 32:DB:13:A:H2’ | 32:DB:70:C:O2’ | 2.06 | 0.54 |
| 33:DD:224:ALA:O | 33:DD:225:ALA:CB | 2.55 | 0.54 |
| 40:DO:31:LYS:HB3 | 40:DO:32:TYR:CE1 | 2.42 | 0.54 |
| 44:DS:24:LEU:O | 44:DS:85:VAL:HG12 | 2.08 | 0.54 |
| 45:DT:78:LEU:C | 45:DT:79:HIS:ND1 | 2.66 | 0.54 |
| 48:DW:97:LYS:HE2 | 48:DW:99:ARG:NH2 | 2.23 | 0.54 |
| 1:AA:96:U:O2’ | 1:AA:97:G:P | 2.66 | 0.54 |
| 1:AA:1227:A:OP2 | 13:AM:111:LYS:HE2 | 2.08 | 0.54 |
| 1:AA:1334:G:H8 | 1:AA:1334:G:OP2 | 1.92 | 0.54 |
| 10:AJ:96:ILE:N | 10:AJ:96:ILE:HD13 | 2.23 | 0.54 |
| 11:AK:21:ILE:HG13 | 11:AK:30:VAL:HG12 | 1.88 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 18:AR:52:PRO:O | 18:AR:56:THR:HG23 | 2.08 | 0.54 |
| 22:B0:77:ARG:HH22 | 31:BA:857:C:H5' | 1.73 | 0.54 |
| 31:BA:554:U:O2' | 31:BA:555:U:H5' | 2.08 | 0.54 |
| 31:BA:1494:A:N3 | 31:BA:1494:A:C2' | 2.71 | 0.54 |
| 31:BA:2202:C:O2 | 33:BD:151:LYS:NZ | 2.34 | 0.54 |
| 31:BA:2358:G:H1 | 41:BP:55:ARG:HH22 | 1.56 | 0.54 |
| 44:BS:96:GLY:O | 44:BS:98:VAL:HB | 2.07 | 0.54 |
| 46:BU:65:ILE:HG12 | 46:BU:96:ALA:HB1 | 1.90 | 0.54 |
| 49:BX:26:TYR:OH | 49:BX:89:ILE:HG21 | 2.07 | 0.54 |
| 50:BY:28:LYS:HB2 | 50:BY:37:VAL:CB | 2.30 | 0.54 |
| 1:CA:189(K):U:H2' | 1:CA:189(L):G:H8 | 1.71 | 0.54 |
| 1:CA:719:C:C5 | 1:CA:720:C:C4 | 2.95 | 0.54 |
| 1:CA:754:C:H3' | 1:CA:754:C:O2 | 2.08 | 0.54 |
| 7:CG:79:ARG:HE | 7:CG:84:ASN:HD21 | 1.56 | 0.54 |
| 10:CJ:96:ILE:N | 10:CJ:96:ILE:HD13 | 2.23 | 0.54 |
| 15:CO:25:THR:O | 15:CO:26:GLU:C | 2.51 | 0.54 |
| 15:CO:78:TYR:OH | 15:CO:88:ARG:HD2 | 2.07 | 0.54 |
| 23:D1:12:PRO:HD2 | 23:D1:62:VAL:CG2 | 2.36 | 0.54 |
| 23:D1:17:SER:O | 23:D1:44:PRO:CD | 2.45 | 0.54 |
| 24:D2:47:ASN:C | 24:D2:49:LYS:H | 2.14 | 0.54 |
| 28:D6:32:ASN:O | 28:D6:33:LYS:HB2 | 2.08 | 0.54 |
| 29:D7:39:ARG:NH2 | 31:DA:468:G:N7 | 2.50 | 0.54 |
| 30:D8:46:ARG:HH22 | 41:DP:65:ARG:HH22 | 1.56 | 0.54 |
| 31:DA:263:C:H2' | 31:DA:264:C:O4' | 2.07 | 0.54 |
| 31:DA:1489:U:H2' | 31:DA:1490:A:OP2 | 2.07 | 0.54 |
| 31:DA:2292:C:C2' | 31:DA:2293:C:H5' | 2.37 | 0.54 |
| 31:DA:2639:A:C2' | 31:DA:2640:G:H5' | 2.38 | 0.54 |
| 32:DB:7:G:C2' | 32:DB:8:U:H5" | 2.38 | 0.54 |
| 33:DD:4:LYS:NZ | 33:DD:20:ASP:HA | 2.23 | 0.54 |
| 33:DD:35:LYS:HG2 | 33:DD:64:ILE:CA | 2.37 | 0.54 |
| 33:DD:69:ARG:NH2 | 33:DD:128:GLY:O | 2.33 | 0.54 |
| 35:DF:195:ASP:CG | 35:DF:197:ASP:HB3 | 2.33 | 0.54 |
| 36:DG:13:GLU:O | 36:DG:14:GLU:CB | 2.54 | 0.54 |
| 36:DG:110:ALA:HA | 36:DG:140:ILE:O | 2.08 | 0.54 |
| 39:DN:131:GLN:CD | 39:DN:134:ARG:CB | 2.79 | 0.54 |
| 40:DO:14:THR:HG21 | 40:DO:86:ILE:HD13 | 1.90 | 0.54 |
| 50:DY:19:LYS:HB3 | 50:DY:20:TYR:CD1 | 2.43 | 0.54 |
| 51:DZ:39:VAL:HG23 | 51:DZ:40:ASP:N | 2.23 | 0.54 |
| 1:AA:108:G:O6 | 20:AT:15:ARG:HD2 | 2.08 | 0.53 |
| 1:AA:115:G:H4' | 1:AA:116:A:O5' | 2.09 | 0.53 |
| 1:AA:355:C:C2 | 1:AA:356:A:C8 | 2.97 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:996:A:H2' | 1:AA:997:U:O4' | 2.08 | 0.53 |
| 1:AA:1201:A:C1' | 1:AA:1202:G:OP2 | 2.56 | 0.53 |
| 1:AA:1356:G:H2' | 1:AA:1357:A:C8 | 2.43 | 0.53 |
| 1:AA:1468:A:H2' | 1:AA:1469:G:O4' | 2.08 | 0.53 |
| 6:AF:98:LEU:H | 6:AF:98:LEU:HD12 | 1.73 | 0.53 |
| 10:AJ:40:LEU:HD23 | 10:AJ:40:LEU:H | 1.73 | 0.53 |
| 22:B0:11:ARG:O | 22:B0:14:ARG:NH2 | 2.41 | 0.53 |
| 31:BA:675:A:C5 | 31:BA:804:A:C2 | 2.97 | 0.53 |
| 31:BA:993:G:N2 | 47:BV:91:TYR:OH | 2.42 | 0.53 |
| 31:BA:1464:C:O2' | 31:BA:1528:A:H8 | 1.91 | 0.53 |
| 31:BA:2415:G:C4' | 41:BP:67:MET:H | 2.06 | 0.53 |
| 31:BA:2473:U:C4 | 31:BA:2474:C:C5 | 2.96 | 0.53 |
| 33:BD:70:TRP:C | 33:BD:70:TRP:CD1 | 2.86 | 0.53 |
| 35:BF:183:VAL:O | 35:BF:187:VAL:HG23 | 2.08 | 0.53 |
| 41:BP:24:GLY:CA | 41:BP:33:ARG:HE | 2.21 | 0.53 |
| 47:BV:15:GLU:O | 47:BV:98:GLU:OE2 | 2.26 | 0.53 |
| 48:BW:5:ALA:HB3 | 48:BW:50:VAL:HG23 | 1.90 | 0.53 |
| 50:BY:28:LYS:HE3 | 50:BY:37:VAL:HG12 | 1.90 | 0.53 |
| 51:BZ:156:LYS:O | 51:BZ:158:PRO:HD3 | 2.07 | 0.53 |
| 1:CA:180:U:H2' | 1:CA:181:G:H5' | 1.90 | 0.53 |
| 1:CA:227:G:O2' | 1:CA:228:A:H5' | 2.08 | 0.53 |
| 1:CA:411:A:H2' | 1:CA:412:A:H4' | 1.90 | 0.53 |
| 1:CA:498:U:O2 | 1:CA:498:U:H2' | 2.08 | 0.53 |
| 1:CA:1281:U:H5' | 1:CA:1282:C:H5 | 1.72 | 0.53 |
| 1:CA:1346:A:H5" | 9:CI:120:ARG:HH12 | 1.73 | 0.53 |
| 10:CJ:39:PRO:HB3 | 10:CJ:70:ARG:NH1 | 2.22 | 0.53 |
| 12:CL:32:PHE:HB3 | 12:CL:84:LEU:HD21 | 1.90 | 0.53 |
| 15:CO:67:LEU:HD22 | 15:CO:78:TYR:HE1 | 1.73 | 0.53 |
| 17:CQ:77:VAL:O | 17:CQ:78:GLU:HB3 | 2.07 | 0.53 |
| 19:CS:44:MET:HA | 19:CS:44:MET:HE3 | 1.89 | 0.53 |
| 27:D5:32:PRO:O | 27:D5:33:CYS:CB | 2.55 | 0.53 |
| 31:DA:415:A:H2' | 31:DA:416:C:H6 | 1.73 | 0.53 |
| 31:DA:995:C:N3 | 39:DN:4:TYR:CE1 | 2.77 | 0.53 |
| 31:DA:2639:A:H2' | 31:DA:2640:G:H5' | 1.90 | 0.53 |
| 33:DD:186:HIS:HD2 | 33:DD:188:GLU:N | 1.99 | 0.53 |
| 34:DE:60:ASN:OD1 | 34:DE:62:PRO:HD2 | 2.07 | 0.53 |
| 36:DG:81:LYS:O | 36:DG:82:LEU:O | 2.25 | 0.53 |
| 39:DN:56:ASN:H | 39:DN:125:GLY:HA3 | 1.73 | 0.53 |
| 41:DP:98:GLU:HG3 | 41:DP:99:LEU:N | 2.22 | 0.53 |
| 44:DS:98:VAL:HG13 | 44:DS:100:ALA:H | 1.73 | 0.53 |
| 48:DW:5:ALA:HB3 | 48:DW:50:VAL:HG23 | 1.89 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 50:DY:81:LYS:HD3 | 50:DY:97:ARG:O | 2.08 | 0.53 |
| 1:AA:175:C:H2' | 1:AA:176:C:H6 | 1.72 | 0.53 |
| 1:AA:774:G:H2' | 1:AA:775:G:H5' | 1.91 | 0.53 |
| 1:AA:1184:G:H2' | 1:AA:1185:G:H8 | 1.73 | 0.53 |
| 1:AA:1392:G:C2' | 1:AA:1393:U:H5' | 2.38 | 0.53 |
| 4:AD:88:VAL:HG13 | 5:AE:97:GLY:HA3 | 1.90 | 0.53 |
| 11:AK:81:ASP:OD2 | 11:AK:106:LYS:HG2 | 2.08 | 0.53 |
| 25:B3:18:ASP:HB2 | 25:B3:49:LYS:HE3 | 1.91 | 0.53 |
| 26:B4:29:PRO:C | 26:B4:31:ILE:H | 2.16 | 0.53 |
| 31:BA:287:C:C2 | 31:BA:288:C:C6 | 2.96 | 0.53 |
| 31:BA:1169:G:N2 | 31:BA:1181:C:C2 | 2.76 | 0.53 |
| 31:BA:2478:A:H2' | 31:BA:2479:G:H5' | 1.90 | 0.53 |
| 35:BF:65:TRP:CH2 | 35:BF:75:HIS:HD2 | 2.26 | 0.53 |
| 38:BI:109:ILE:CG2 | 38:BI:130:TYR:OH | 2.56 | 0.53 |
| 38:BI:136:VAL:HG22 | 38:BI:136:VAL:O | 2.08 | 0.53 |
| 41:BP:87:ASP:O | 41:BP:90:ARG:HB2 | 2.08 | 0.53 |
| 43:BR:71:GLN:HA | 43:BR:71:GLN:NE2 | 2.23 | 0.53 |
| 44:BS:29:PHE:C | 44:BS:29:PHE:CD2 | 2.85 | 0.53 |
| 48:BW:17:VAL:O | 48:BW:18:ARG:C | 2.50 | 0.53 |
| 48:BW:28:SER:OG | 48:BW:31:GLU:HB2 | 2.08 | 0.53 |
| 48:BW:92:ARG:HH11 | 48:BW:92:ARG:HG2 | 1.74 | 0.53 |
| 1:CA:113:G:H2' | 1:CA:114:U:C6 | 2.43 | 0.53 |
| 1:CA:316:G:OP2 | 1:CA:351:G:O2' | 2.25 | 0.53 |
| 1:CA:619:U:H2' | 4:CD:135:LEU:HD21 | 1.90 | 0.53 |
| 1:CA:662:G:H2' | 1:CA:663:A:C8 | 2.44 | 0.53 |
| 3:CC:53:ALA:HB2 | 3:CC:115:LEU:HD21 | 1.90 | 0.53 |
| 6:CF:20:ALA:O | 6:CF:23:LYS:HB2 | 2.08 | 0.53 |
| 11:CK:125:PHE:N | 11:CK:125:PHE:CD1 | 2.76 | 0.53 |
| 12:CL:86:ARG:HB2 | 12:CL:101:VAL:CG2 | 2.37 | 0.53 |
| 16:CP:18:ARG:O | 16:CP:19:ILE:C | 2.52 | 0.53 |
| 24:D2:32:LEU:O | 24:D2:33:MET:C | 2.51 | 0.53 |
| 29:D7:15:THR:HG22 | 29:D7:16:HIS:N | 2.22 | 0.53 |
| 30:D8:32:LEU:HB3 | 30:D8:34:TRP:CA | 2.38 | 0.53 |
| 31:DA:214:G:H1' | 31:DA:216:A:O2' | 2.09 | 0.53 |
| 31:DA:528:A:C2 | 31:DA:2043:C:C5' | 2.91 | 0.53 |
| 31:DA:542:C:N4 | 31:DA:543:C:N4 | 2.56 | 0.53 |
| 31:DA:719:C:H2' | 31:DA:720:C:C6 | 2.43 | 0.53 |
| 31:DA:860:U:C5 | 31:DA:917:A:N7 | 2.77 | 0.53 |
| 31:DA:866:A:C6 | 31:DA:914:C:C5 | 2.96 | 0.53 |
| 31:DA:948:G:O2' | 31:DA:949:C:H5' | 2.09 | 0.53 |
| 31:DA:952:G:C6 | 31:DA:953:A:N7 | 2.76 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1324:G:C4 | 31:DA:1328:G:O6 | 2.61 | 0.53 |
| 31:DA:1543:C:OP2 | 31:DA:1543:C:C6 | 2.61 | 0.53 |
| 31:DA:2859:G:O2' | 31:DA:2860:A:P | 2.67 | 0.53 |
| 31:DA:2875:C:C4' | 45:DT:5:ALA:HB2 | 2.38 | 0.53 |
| 33:DD:35:LYS:HD3 | 33:DD:63:ARG:C | 2.33 | 0.53 |
| 33:DD:71:ASP:OD2 | 33:DD:103:ARG:NH2 | 2.41 | 0.53 |
| 34:DE:75:VAL:O | 34:DE:77:ILE:N | 2.42 | 0.53 |
| 39:DN:128:HIS:CD2 | 39:DN:128:HIS:O | 2.60 | 0.53 |
| 41:DP:107:LYS:C | 41:DP:109:GLY:N | 2.66 | 0.53 |
| 42:DQ:23:GLY:HA3 | 42:DQ:99:PRO:O | 2.09 | 0.53 |
| 44:DS:42:ASP:O | 44:DS:43:GLU:HB2 | 2.08 | 0.53 |
| 44:DS:66:ALA:O | 44:DS:67:ARG:HB3 | 2.08 | 0.53 |
| 46:DU:59:ARG:O | 46:DU:60:LEU:C | 2.51 | 0.53 |
| 50:DY:97:ARG:O | 50:DY:98:VAL:C | 2.50 | 0.53 |
| 1:AA:191:G:N3 | 20:AT:103:GLY:O | 2.41 | 0.53 |
| 3:AC:182:ILE:HG12 | 3:AC:203:PHE:CD1 | 2.40 | 0.53 |
| 5:AE:112:LEU:N | 5:AE:112:LEU:HD23 | 2.23 | 0.53 |
| 6:AF:2:ARG:HB2 | 6:AF:4:TYR:CE2 | 2.42 | 0.53 |
| 10:AJ:8:LEU:HG | 10:AJ:96:ILE:HG22 | 1.89 | 0.53 |
| 18:AR:73:ALA:HB1 | 18:AR:79:LEU:HD12 | 1.90 | 0.53 |
| 23:B1:60:PHE:HZ | 23:B1:90:ILE:HG21 | 1.73 | 0.53 |
| 24:B2:26:ARG:HG2 | 49:BX:5:TYR:HB3 | 1.90 | 0.53 |
| 27:B5:42:PRO:HB2 | 27:B5:43:HIS:HD2 | 1.73 | 0.53 |
| 31:BA:271(P):C:O2' | 31:BA:271(Q):G:H5' | 2.09 | 0.53 |
| 31:BA:518:G:H2' | 31:BA:519:U:H6 | 1.74 | 0.53 |
| 31:BA:1669:A:H5" | 31:BA:2550:G:OP1 | 2.08 | 0.53 |
| 33:BD:177:LEU:HD23 | 33:BD:177:LEU:N | 2.24 | 0.53 |
| 34:BE:136:ARG:HG2 | 34:BE:136:ARG:NH1 | 2.22 | 0.53 |
| 34:BE:188:VAL:HB | 34:BE:189:PRO:HD2 | 1.89 | 0.53 |
| 37:BH:83:TYR:HA | 37:BH:135:GLY:O | 2.08 | 0.53 |
| 41:BP:98:GLU:HG3 | 41:BP:99:LEU:N | 2.23 | 0.53 |
| 1:CA:272:C:H2' | 1:CA:273:A:H8 | 1.73 | 0.53 |
| 1:CA:734:G:C2 | 1:CA:735:C:C2 | 2.96 | 0.53 |
| 1:CA:1135:U:H4' | 1:CA:1136:U:H5 | 1.74 | 0.53 |
| 1:CA:1356:G:H2' | 1:CA:1357:A:C8 | 2.44 | 0.53 |
| 2:CB:121:LEU:O | 2:CB:121:LEU:HD23 | 2.07 | 0.53 |
| 3:CC:189:ALA:HB3 | 3:CC:196:LEU:HB2 | 1.90 | 0.53 |
| 6:CF:8:ILE:HG22 | 6:CF:10:LEU:CD1 | 2.38 | 0.53 |
| 12:CL:102:ARG:CG | 12:CL:102:ARG:NH1 | 2.59 | 0.53 |
| 15:CO:5:LYS:O | 15:CO:9:GLN:HG2 | 2.08 | 0.53 |
| 15:CO:65:ARG:HB2 | 15:CO:65:ARG:HH11 | 1.72 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 23:D1:16:ASN:CB | 23:D1:46:LEU:HG | 2.32 | 0.53 |
| 24:D2:26:ARG:HG2 | 49:DX:5:TYR:HB3 | 1.90 | 0.53 |
| 25:D3:44:ARG:O | 25:D3:48:GLU:HG2 | 2.08 | 0.53 |
| 31:DA:639:U:H2' | 31:DA:640:C:C6 | 2.43 | 0.53 |
| 31:DA:719:C:H2' | 31:DA:720:C:H6 | 1.73 | 0.53 |
| 31:DA:893:C:H2' | 31:DA:894:C:O5' | 2.09 | 0.53 |
| 31:DA:1110:G:OP1 | 31:DA:1110:G:H4' | 2.07 | 0.53 |
| 31:DA:1208:C:H2' | 31:DA:1209:G:H5' | 1.90 | 0.53 |
| 31:DA:2523:G:C2' | 31:DA:2524:G:H5' | 2.34 | 0.53 |
| 31:DA:2636:U:O2' | 31:DA:2637:U:C5' | 2.56 | 0.53 |
| 32:DB:37:C:H2' | 32:DB:38:C:H5' | 1.90 | 0.53 |
| 34:DE:66:HIS:O | 34:DE:66:HIS:CG | 2.62 | 0.53 |
| 37:DH:153:LYS:HG2 | 37:DH:154:PRO:N | 2.23 | 0.53 |
| 41:DP:30:THR:O | 41:DP:33:ARG:N | 2.39 | 0.53 |
| 41:DP:102:ARG:O | 41:DP:103:ALA:HB2 | 2.08 | 0.53 |
| 42:DQ:8:LYS:HG3 | 42:DQ:9:TYR:H | 1.73 | 0.53 |
| 46:DU:10:ARG:HG2 | 46:DU:14:HIS:CE1 | 2.43 | 0.53 |
| 47:DV:82:ARG:CG | 47:DV:82:ARG:NH1 | 2.48 | 0.53 |
| 51:DZ:10:ARG:HH21 | 51:DZ:26:GLY:H | 1.57 | 0.53 |
| 1:AA:66:G:C2 | 1:AA:67:C:C6 | 2.96 | 0.53 |
| 1:AA:84:U:C5 | 1:AA:88:A:C8 | 2.95 | 0.53 |
| 1:AA:137:C:H42 | 1:AA:226:G:H1 | 1.56 | 0.53 |
| 1:AA:189(C):C:H2' | 1:AA:189(D):C:O4' | 2.08 | 0.53 |
| 1:AA:356:A:H2' | 1:AA:357:G:C8 | 2.43 | 0.53 |
| 1:AA:409:G:C2' | 1:AA:410:G:H5' | 2.39 | 0.53 |
| 1:AA:411:A:H2' | 1:AA:412:A:H4' | 1.89 | 0.53 |
| 1:AA:854:G:H3' | 1:AA:871:U:O4 | 2.09 | 0.53 |
| 1:AA:872:A:C4 | 1:AA:874:G:N7 | 2.77 | 0.53 |
| 1:AA:921:U:O2 | 5:AE:19:MET:HB2 | 2.08 | 0.53 |
| 1:AA:953:G:H5' | 1:AA:965:A:H61 | 1.73 | 0.53 |
| 1:AA:1072:G:C6 | 1:AA:1073:U:C4 | 2.96 | 0.53 |
| 1:AA:1150:U:O4 | 1:AA:1151:A:N6 | 2.41 | 0.53 |
| 2:AB:102:LEU:HD12 | 2:AB:102:LEU:N | 2.23 | 0.53 |
| 6:AF:5:GLU:HG2 | 6:AF:62:TRP:CZ2 | 2.44 | 0.53 |
| 13:AM:91:ARG:HD3 | 19:AS:81:ARG:HH21 | 1.72 | 0.53 |
| 30:B8:31:HIS:O | 30:B8:33:ASN:N | 2.42 | 0.53 |
| 31:BA:65:C:H2' | 31:BA:66:C:H6 | 1.73 | 0.53 |
| 31:BA:614:U:O2 | 31:BA:614:U:O5' | 2.26 | 0.53 |
| 31:BA:863:A:O2' | 31:BA:864:G:H5' | 2.08 | 0.53 |
| 31:BA:911:A:N9 | 42:BQ:9:TYR:OH | 2.39 | 0.53 |
| 31:BA:1049:C:H2' | 31:BA:1049:C:O2 | 2.09 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2036:C:H5' | 31:BA:2036:C:C6 | 2.34 | 0.53 |
| 31:BA:2308:G:C2 | 31:BA:2309:A:C6 | 2.96 | 0.53 |
| 31:BA:2584:U:H2' | 31:BA:2585:U:H2' | 1.89 | 0.53 |
| 31:BA:2702:U:HO2' | 31:BA:2703:C:H5 | 1.51 | 0.53 |
| 31:BA:2810:A:N6 | 31:BA:2891:G:O2' | 2.42 | 0.53 |
| 34:BE:117:MET:O | 34:BE:117:MET:HG2 | 2.08 | 0.53 |
| 42:BQ:18:LYS:O | 42:BQ:19:GLY:O | 2.25 | 0.53 |
| 42:BQ:75:THR:CA | 42:BQ:88:GLY:HA2 | 2.39 | 0.53 |
| 48:BW:92:ARG:O | 48:BW:93:ALA:HB3 | 2.09 | 0.53 |
| 50:BY:14:LEU:HD11 | 50:BY:22:GLY:HA2 | 1.89 | 0.53 |
| 50:BY:88:LYS:O | 50:BY:89:PHE:HB2 | 2.08 | 0.53 |
| 1:CA:173:U:C2 | 1:CA:197:A:N1 | 2.76 | 0.53 |
| 1:CA:189(E):U:O2' | 1:CA:189(F):U:H5' | 2.08 | 0.53 |
| 1:CA:356:A:H2' | 1:CA:357:G:H8 | 1.72 | 0.53 |
| 1:CA:564:C:C2' | 1:CA:565:U:H5' | 2.38 | 0.53 |
| 1:CA:625:G:H2' | 1:CA:626:U:C6 | 2.43 | 0.53 |
| 1:CA:685:G:N2 | 1:CA:686:U:C4 | 2.76 | 0.53 |
| 1:CA:1003:G:C2 | 1:CA:1004:A:H1' | 2.43 | 0.53 |
| 1:CA:1091:U:O2 | 1:CA:1093:A:C8 | 2.62 | 0.53 |
| 1:CA:1277:C:O2' | 1:CA:1279:A:H1' | 2.08 | 0.53 |
| 1:CA:1478:C:O2' | 1:CA:1479:C:H5' | 2.09 | 0.53 |
| 2:CB:98:LEU:HB2 | 2:CB:101:MET:HE3 | 1.89 | 0.53 |
| 4:CD:2:GLY:O | 4:CD:3:ARG:C | 2.51 | 0.53 |
| 4:CD:3:ARG:O | 4:CD:5:ILE:HG13 | 2.09 | 0.53 |
| 18:CR:53:ARG:C | 18:CR:55:ARG:N | 2.65 | 0.53 |
| 22:D0:53:MET:HB2 | 22:D0:59:LEU:CD2 | 2.38 | 0.53 |
| 28:D6:30:THR:O | 28:D6:31:PRO:C | 2.51 | 0.53 |
| 31:DA:194:G:H2' | 31:DA:195:A:O4' | 2.09 | 0.53 |
| 31:DA:819:A:OP2 | 31:DA:1187:G:N2 | 2.24 | 0.53 |
| 31:DA:2751:G:H3' | 31:DA:2752:C:H6 | 1.72 | 0.53 |
| 35:DF:65:TRP:CZ3 | 35:DF:75:HIS:HD2 | 2.27 | 0.53 |
| 36:DG:111:LEU:O | 36:DG:114:ILE:HG12 | 2.08 | 0.53 |
| 37:DH:92:ILE:C | 37:DH:94:TYR:H | 2.16 | 0.53 |
| 49:DX:36:LYS:NZ | 49:DX:38:GLU:C | 2.66 | 0.53 |
| 1:AA:373:A:O2' | 1:AA:374:A:H5' | 2.07 | 0.53 |
| 1:AA:541:G:H2' | 1:AA:542:G:H8 | 1.73 | 0.53 |
| 4:AD:33:MET:HE3 | 4:AD:33:MET:HA | 1.90 | 0.53 |
| 10:AJ:58:ASP:O | 10:AJ:59:SER:C | 2.52 | 0.53 |
| 23:B1:27:GLU:OE2 | 23:B1:32:LYS:HB2 | 2.09 | 0.53 |
| 23:B1:67:ILE:H | 23:B1:67:ILE:HD12 | 1.73 | 0.53 |
| 24:B2:28:LYS:HD3 | 24:B2:43:GLN:HB3 | 1.90 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 30:B8:39:LYS:HE3 | 30:B8:42:ARG:NH1 | 2.24 | 0.53 |
| 31:BA:92:A:O2' | 31:BA:93:G:H5' | 2.09 | 0.53 |
| 31:BA:272(J):C:H42 | 31:BA:363(A):A:N6 | 2.06 | 0.53 |
| 31:BA:542:C:O5' | 31:BA:542:C:H6 | 1.91 | 0.53 |
| 31:BA:634:C:H2' | 31:BA:635:C:C6 | 2.43 | 0.53 |
| 31:BA:2193:G:H2' | 31:BA:2194:G:O4' | 2.08 | 0.53 |
| 33:BD:4:LYS:NZ | 33:BD:20:ASP:HA | 2.22 | 0.53 |
| 34:BE:197:ILE:HD11 | 34:BE:199:ARG:CZ | 2.39 | 0.53 |
| 34:BE:201:THR:HG22 | 34:BE:202:LYS:N | 2.24 | 0.53 |
| 36:BG:131:TYR:HB3 | 36:BG:159:VAL:CG1 | 2.37 | 0.53 |
| 37:BH:44:VAL:O | 37:BH:46:GLU:OE2 | 2.26 | 0.53 |
| 38:BI:10:GLU:C | 38:BI:12:LEU:H | 2.16 | 0.53 |
| 38:BI:50:ARG:C | 38:BI:52:ARG:N | 2.66 | 0.53 |
| 38:BI:130:TYR:O | 38:BI:131:LYS:HG3 | 2.08 | 0.53 |
| 48:BW:54:ALA:CB | 48:BW:107:LEU:HD22 | 2.38 | 0.53 |
| 49:BX:72:LYS:HB2 | 49:BX:74:PRO:HD3 | 1.90 | 0.53 |
| 50:BY:28:LYS:CE | 50:BY:30:VAL:HG22 | 2.22 | 0.53 |
| 51:BZ:67:LEU:HD12 | 51:BZ:67:LEU:N | 2.23 | 0.53 |
| 1:CA:501:C:O2' | 1:CA:502:G:H5' | 2.09 | 0.53 |
| 1:CA:556:C:C2' | 1:CA:557:G:H5' | 2.39 | 0.53 |
| 1:CA:953:G:H5' | 1:CA:965:A:H61 | 1.73 | 0.53 |
| 1:CA:1470:G:O2' | 1:CA:1471:G:H5' | 2.07 | 0.53 |
| 4:CD:79:PHE:CD1 | 4:CD:207:TYR:HD1 | 2.26 | 0.53 |
| 4:CD:111:ALA:HB2 | 4:CD:120:LEU:HD12 | 1.91 | 0.53 |
| 7:CG:75:VAL:HG21 | 7:CG:144:MET:HB3 | 1.90 | 0.53 |
| 10:CJ:8:LEU:HG | 10:CJ:96:ILE:HG22 | 1.89 | 0.53 |
| 11:CK:125:PHE:H | 11:CK:125:PHE:HD1 | 1.56 | 0.53 |
| 15:CO:64:ARG:HG3 | 15:CO:64:ARG:NH1 | 2.24 | 0.53 |
| 29:D7:35:ARG:HG3 | 29:D7:42:LEU:HD11 | 1.91 | 0.53 |
| 30:D8:27:THR:HA | 41:DP:62:LEU:CD1 | 2.39 | 0.53 |
| 31:DA:470:A:OP1 | 35:DF:59:TYR:HE2 | 1.91 | 0.53 |
| 31:DA:906:G:H2' | 31:DA:907:U:O5' | 2.08 | 0.53 |
| 31:DA:2291:U:H5" | 31:DA:2380:C:O2 | 2.08 | 0.53 |
| 31:DA:2400:G:C5 | 31:DA:2401:U:C5 | 2.97 | 0.53 |
| 31:DA:2517:C:C2 | 31:DA:2542:A:N6 | 2.76 | 0.53 |
| 33:DD:53:PHE:CD1 | 33:DD:220:HIS:HA | 2.43 | 0.53 |
| 34:DE:89:ASP:O | 34:DE:90:THR:HB | 2.07 | 0.53 |
| 45:DT:56:GLY:O | 45:DT:59:THR:HG22 | 2.08 | 0.53 |
| 46:DU:55:ARG:HA | 46:DU:58:ARG:HD2 | 1.91 | 0.53 |
| 49:DX:35:THR:O | 49:DX:36:LYS:C | 2.51 | 0.53 |
| 1:AA:189(K):U:H2' | 1:AA:189(L):G:H8 | 1.72 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:323:U:OP1 | 20:AT:26:ASN:ND2 | 2.41 | 0.53 |
| 1:AA:1255:G:H5' | 1:AA:1256:A:OP1 | 2.08 | 0.53 |
| 1:AA:1317:C:OP1 | 14:AN:17:LYS:HG2 | 2.08 | 0.53 |
| 1:AA:1521:G:H2' | 1:AA:1522:U:H6 | 1.70 | 0.53 |
| 3:AC:39:ILE:HG21 | 3:AC:57:ILE:HD11 | 1.90 | 0.53 |
| 4:AD:9:CYS:HA | 4:AD:12:CYS:HB2 | 1.91 | 0.53 |
| 7:AG:92:SER:OG | 7:AG:93:PRO:HD2 | 2.09 | 0.53 |
| 11:AK:69:ALA:HB1 | 11:AK:103:LEU:CD2 | 2.39 | 0.53 |
| 16:AP:39:TYR:HB2 | 16:AP:49:LEU:HD12 | 1.90 | 0.53 |
| 16:AP:39:TYR:HB2 | 16:AP:49:LEU:CD1 | 2.39 | 0.53 |
| 23:B1:10:LYS:HD2 | 23:B1:14:VAL:HA | 1.90 | 0.53 |
| 23:B1:65:SER:H | 23:B1:67:ILE:HD12 | 1.73 | 0.53 |
| 31:BA:358:U:H5 | 31:BA:359:A:N7 | 2.06 | 0.53 |
| 31:BA:644:A:O2' | 31:BA:645:C:O2 | 2.27 | 0.53 |
| 31:BA:672:C:H2' | 31:BA:673:C:C6 | 2.43 | 0.53 |
| 31:BA:807:U:C2' | 31:BA:808:G:O5' | 2.57 | 0.53 |
| 31:BA:1276:A:O2' | 43:BR:16:HIS:HE1 | 1.92 | 0.53 |
| 31:BA:1528:A:O2' | 31:BA:1528(A):A:C8 | 2.58 | 0.53 |
| 31:BA:1549:C:O2' | 31:BA:1550:C:H5' | 2.09 | 0.53 |
| 31:BA:1899:G:O2' | 31:BA:1900:A:H5" | 2.09 | 0.53 |
| 31:BA:2661:G:C8 | 31:BA:2662:A:N3 | 2.77 | 0.53 |
| 35:BF:2:LYS:O | 35:BF:25:PRO:HG2 | 2.09 | 0.53 |
| 35:BF:129:PHE:HE1 | 35:BF:142:TRP:CH2 | 2.26 | 0.53 |
| 36:BG:60:LEU:C | 36:BG:60:LEU:HD13 | 2.33 | 0.53 |
| 36:BG:119:GLY:H | 36:BG:181:ARG:HH21 | 1.56 | 0.53 |
| 38:BI:108:THR:O | 38:BI:109:ILE:HG13 | 2.08 | 0.53 |
| 39:BN:9:VAL:CG1 | 39:BN:39:ARG:HH22 | 2.21 | 0.53 |
| 39:BN:131:GLN:CD | 39:BN:134:ARG:CB | 2.76 | 0.53 |
| 44:BS:16:ASN:O | 44:BS:19:LYS:HB3 | 2.09 | 0.53 |
| 45:BT:120:ARG:HA | 45:BT:123:GLN:HG2 | 1.91 | 0.53 |
| 46:BU:90:VAL:HG12 | 46:BU:91:ASP:N | 2.19 | 0.53 |
| 47:BV:21:ARG:HG2 | 47:BV:93:GLU:OE1 | 2.08 | 0.53 |
| 47:BV:56:SER:O | 47:BV:57:VAL:HB | 2.08 | 0.53 |
| 49:BX:65:ARG:CZ | 49:BX:66:LEU:N | 2.72 | 0.53 |
| 1:CA:115:G:C2 | 1:CA:289:G:N7 | 2.76 | 0.53 |
| 1:CA:189(C):C:H2' | 1:CA:189(D):C:O4' | 2.08 | 0.53 |
| 1:CA:250:A:C4' | 1:CA:251:G:O5' | 2.52 | 0.53 |
| 1:CA:491:G:C4 | 1:CA:492:G:C8 | 2.97 | 0.53 |
| 1:CA:523:A:H61 | 12:CL:53:ARG:NH1 | 2.06 | 0.53 |
| 1:CA:1190:G:H3' | 3:CC:3:ASN:ND2 | 2.21 | 0.53 |
| 1:CA:1256:A:O3' | 1:CA:1257:U:H4' | 2.08 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:1317:C:OP1 | 14:CN:17:LYS:HG2 | 2.07 | 0.53 |
| 7:CG:23:VAL:HG13 | 7:CG:43:PHE:CZ | 2.44 | 0.53 |
| 13:CM:10:PRO:CB | 13:CM:18:ALA:HB1 | 2.37 | 0.53 |
| 24:D2:14:ARG:CZ | 24:D2:57:ILE:CG2 | 2.87 | 0.53 |
| 25:D3:45:GLY:HA3 | 31:DA:851:U:O2' | 2.09 | 0.53 |
| 28:D6:15:GLU:CD | 28:D6:18:ARG:HD2 | 2.33 | 0.53 |
| 29:D7:5:TRP:CZ3 | 31:DA:464:U:H4' | 2.43 | 0.53 |
| 31:DA:157:U:H4' | 31:DA:171:G:N2 | 2.22 | 0.53 |
| 31:DA:271(T):C:O2 | 31:DA:271(T):C:H2' | 2.09 | 0.53 |
| 31:DA:568:U:H5' | 31:DA:945:A:C2 | 2.44 | 0.53 |
| 31:DA:601:C:O2 | 31:DA:605:C:H4' | 2.09 | 0.53 |
| 31:DA:960:A:H5" | 31:DA:961:C:OP2 | 2.09 | 0.53 |
| 31:DA:1673:U:O4 | 34:DE:129:HIS:HD2 | 1.91 | 0.53 |
| 31:DA:1836:C:C2' | 31:DA:1837:C:H5' | 2.39 | 0.53 |
| 31:DA:1934:C:H5" | 31:DA:1934:C:C6 | 2.31 | 0.53 |
| 31:DA:2584:U:H2' | 31:DA:2585:U:H2' | 1.89 | 0.53 |
| 34:DE:59:VAL:C | 34:DE:60:ASN:ND2 | 2.67 | 0.53 |
| 35:DF:7:TYR:CD1 | 35:DF:8:GLN:N | 2.76 | 0.53 |
| 38:DI:99:GLU:HG3 | 38:DI:103:ARG:CZ | 2.39 | 0.53 |
| 42:DQ:35:VAL:CG1 | 42:DQ:130:LYS:HB3 | 2.35 | 0.53 |
| 43:DR:9:LYS:O | 43:DR:10:LEU:HD23 | 2.09 | 0.53 |
| 46:DU:25:TRP:HD1 | 46:DU:26:GLY:H | 1.55 | 0.53 |
| 47:DV:40:LEU:HD12 | 47:DV:40:LEU:O | 2.09 | 0.53 |
| 49:DX:26:TYR:OH | 49:DX:89:ILE:HG21 | 2.09 | 0.53 |
| 50:DY:28:LYS:CE | 50:DY:37:VAL:HG12 | 2.39 | 0.53 |
| 1:AA:316:G:OP2 | 1:AA:351:G:O2' | 2.27 | 0.53 |
| 1:AA:840:C:H4' | 1:AA:848:C:O2 | 2.09 | 0.53 |
| 1:AA:1494:G:N2 | 31:BA:1912:A:N3 | 2.57 | 0.53 |
| 2:AB:189:ASP:N | 2:AB:189:ASP:OD1 | 2.42 | 0.53 |
| 3:AC:52:LEU:H | 3:AC:52:LEU:CD2 | 2.18 | 0.53 |
| 5:AE:55:VAL:O | 5:AE:58:ALA:HB3 | 2.07 | 0.53 |
| 7:AG:79:ARG:HE | 7:AG:84:ASN:HD21 | 1.55 | 0.53 |
| 11:AK:125:PHE:N | 11:AK:125:PHE:CD1 | 2.76 | 0.53 |
| 17:AQ:77:VAL:O | 17:AQ:78:GLU:HB3 | 2.08 | 0.53 |
| 20:AT:89:ARG:HD2 | 20:AT:104:LEU:HD11 | 1.91 | 0.53 |
| 23:B1:89:GLU:O | 23:B1:93:GLU:N | 2.40 | 0.53 |
| 28:B6:32:ASN:O | 28:B6:33:LYS:HB2 | 2.08 | 0.53 |
| 28:B6:32:ASN:OD1 | 28:B6:33:LYS:N | 2.42 | 0.53 |
| 31:BA:725:G:C6 | 31:BA:726:G:N1 | 2.77 | 0.53 |
| 31:BA:1116:C:C2' | 31:BA:1117:G:H5' | 2.38 | 0.53 |
| 31:BA:1270:C:H5" | 31:BA:1271:G:O5' | 2.07 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1569:A:O2' | 33:BD:38:LYS:HE2 | 2.09 | 0.53 |
| 31:BA:2052:G:O4' | 34:BE:142:GLY:HA3 | 2.09 | 0.53 |
| 31:BA:2247:A:H2' | 31:BA:2248:C:H6 | 1.74 | 0.53 |
| 31:BA:2472:G:C2 | 31:BA:2477:C:OP1 | 2.61 | 0.53 |
| 31:BA:2859:G:H4' | 31:BA:2860:A:OP1 | 2.09 | 0.53 |
| 31:BA:2884:U:C5 | 31:BA:2885:C:C6 | 2.96 | 0.53 |
| 33:BD:125:ILE:HG22 | 33:BD:125:ILE:O | 2.06 | 0.53 |
| 36:BG:13:GLU:O | 36:BG:14:GLU:CB | 2.52 | 0.53 |
| 36:BG:47:LYS:HD3 | 36:BG:81:LYS:CD | 2.35 | 0.53 |
| 37:BH:47:GLU:C | 37:BH:49:VAL:H | 2.15 | 0.53 |
| 39:BN:82:LEU:HD12 | 39:BN:82:LEU:N | 2.24 | 0.53 |
| 40:BO:61:VAL:O | 40:BO:63:VAL:HG12 | 2.08 | 0.53 |
| 41:BP:79:ARG:HH21 | 41:BP:109:GLY:CA | 2.21 | 0.53 |
| 41:BP:91:PHE:CE2 | 41:BP:95:VAL:HG12 | 2.39 | 0.53 |
| 41:BP:140:ALA:C | 25:D3:1:MET:SD | 2.91 | 0.53 |
| 43:BR:13:HIS:O | 43:BR:14:SER:C | 2.52 | 0.53 |
| 44:BS:28:VAL:HG11 | 44:BS:97:ARG:NH2 | 2.24 | 0.53 |
| 46:BU:92:ARG:O | 46:BU:93:LYS:C | 2.52 | 0.53 |
| 51:BZ:141:VAL:HG23 | 51:BZ:144:LEU:HD23 | 1.89 | 0.53 |
| 1:CA:165:C:H2' | 1:CA:166:G:C8 | 2.43 | 0.53 |
| 1:CA:373:A:O2' | 1:CA:374:A:H5' | 2.09 | 0.53 |
| 1:CA:764:C:H2' | 1:CA:765:G:H8 | 1.74 | 0.53 |
| 1:CA:963:G:H21 | 10:CJ:55:LYS:HD3 | 1.73 | 0.53 |
| 1:CA:1128:C:O2' | 1:CA:1130:A:C8 | 2.54 | 0.53 |
| 5:CE:12:LEU:C | 5:CE:12:LEU:HD22 | 2.33 | 0.53 |
| 12:CL:27:LEU:O | 12:CL:29:GLY:N | 2.41 | 0.53 |
| 13:CM:68:GLY:O | 13:CM:69:GLU:CB | 2.54 | 0.53 |
| 26:D4:29:PRO:C | 26:D4:31:ILE:H | 2.15 | 0.53 |
| 31:DA:65:C:H2' | 31:DA:66:C:H6 | 1.74 | 0.53 |
| 31:DA:272(J):C:C2' | 31:DA:274:G:OP1 | 2.57 | 0.53 |
| 31:DA:454:A:H4' | 31:DA:455:C:OP2 | 2.08 | 0.53 |
| 31:DA:848:G:C4 | 31:DA:933:A:C8 | 2.97 | 0.53 |
| 31:DA:856:C:C6 | 31:DA:856:C:H5" | 2.43 | 0.53 |
| 31:DA:910:A:C8 | 42:DQ:13:GLN:HB2 | 2.43 | 0.53 |
| 31:DA:1146:C:H2' | 31:DA:1147:C:C5' | 2.33 | 0.53 |
| 31:DA:1210:A:C5' | 31:DA:1212:G:H5' | 2.37 | 0.53 |
| 33:DD:50:THR:O | 33:DD:51:VAL:HG23 | 2.07 | 0.53 |
| 33:DD:65:ILE:HD13 | 33:DD:65:ILE:O | 2.09 | 0.53 |
| 33:DD:177:LEU:N | 33:DD:177:LEU:HD23 | 2.23 | 0.53 |
| 33:DD:211:ARG:HA | 33:DD:214:TRP:CD2 | 2.44 | 0.53 |
| 40:DO:120:GLU:OE2 | 40:DO:122:LEU:HD21 | 2.08 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 41:DP:40:SER:O | 41:DP:41:ARG:HD2 | 2.08 | 0.53 |
| 43:DR:5:LYS:N | 43:DR:5:LYS:CD | 2.69 | 0.53 |
| 44:DS:27:SER:HB3 | 44:DS:89:ARG:NH1 | 2.24 | 0.53 |
| 44:DS:29:PHE:H | 44:DS:89:ARG:HG2 | 1.74 | 0.53 |
| 1:AA:357:G:C2 | 1:AA:358:U:C5 | 2.97 | 0.53 |
| 1:AA:808:C:P | 15:AO:48:LYS:HE3 | 2.49 | 0.53 |
| 1:AA:1084:G:C5 | 1:AA:1085:U:C4 | 2.97 | 0.53 |
| 1:AA:1091:U:O2 | 1:AA:1093:A:C8 | 2.62 | 0.53 |
| 1:AA:1129:C:H4' | 1:AA:1130:A:H5' | 1.90 | 0.53 |
| 5:AE:102:ALA:HB1 | 5:AE:106:PRO:HG2 | 1.90 | 0.53 |
| 21:AU:5:ASP:O | 21:AU:11:GLY:HA3 | 2.08 | 0.53 |
| 24:B2:47:ASN:HA | 24:B2:50:ILE:O | 2.09 | 0.53 |
| 25:B3:44:ARG:O | 25:B3:48:GLU:HG2 | 2.08 | 0.53 |
| 30:B8:32:LEU:HD13 | 30:B8:32:LEU:H | 1.72 | 0.53 |
| 31:BA:2298:A:H2' | 31:BA:2299:G:O4' | 2.09 | 0.53 |
| 31:BA:2314:C:C2' | 31:BA:2315:G:H5' | 2.39 | 0.53 |
| 31:BA:2408:U:H2' | 31:BA:2409:G:C8 | 2.44 | 0.53 |
| 32:BB:66:A:C5 | 32:BB:109:C:C5 | 2.97 | 0.53 |
| 33:BD:18:VAL:HG23 | 33:BD:211:ARG:HH12 | 1.73 | 0.53 |
| 33:BD:143:HIS:HD2 | 33:BD:144:ALA:HB2 | 1.74 | 0.53 |
| 39:BN:46:VAL:HG13 | 39:BN:48:MET:HG3 | 1.90 | 0.53 |
| 41:BP:66:GLY:O | 41:BP:67:MET:C | 2.51 | 0.53 |
| 42:BQ:20:ALA:CB | 42:BQ:99:PRO:HG2 | 2.39 | 0.53 |
| 45:BT:101:PHE:HE2 | 45:BT:113:LYS:HD2 | 1.73 | 0.53 |
| 49:BX:35:THR:O | 49:BX:36:LYS:C | 2.52 | 0.53 |
| 49:BX:65:ARG:O | 49:BX:66:LEU:CB | 2.57 | 0.53 |
| 50:BY:75:ILE:O | 50:BY:76:CYS:HB2 | 2.08 | 0.53 |
| 1:CA:175:C:H2' | 1:CA:176:C:H6 | 1.73 | 0.53 |
| 1:CA:473:G:H5' | 16:CP:81:ARG:HG3 | 1.90 | 0.53 |
| 1:CA:1157:A:H4' | 1:CA:1158:C:O5' | 2.09 | 0.53 |
| 1:CA:1255:G:H5' | 1:CA:1256:A:OP1 | 2.08 | 0.53 |
| 6:CF:89:MET:SD | 18:CR:76:LEU:HD21 | 2.49 | 0.53 |
| 8:CH:109:ILE:HG22 | 8:CH:137:VAL:HB | 1.90 | 0.53 |
| 11:CK:23:ALA:HB3 | 11:CK:86:GLY:O | 2.09 | 0.53 |
| 12:CL:40:VAL:HG12 | 12:CL:40:VAL:O | 2.07 | 0.53 |
| 13:CM:16:ASP:HB3 | 13:CM:41:PRO:HB3 | 1.91 | 0.53 |
| 29:D7:48:LYS:HD3 | 29:D7:48:LYS:N | 2.23 | 0.53 |
| 31:DA:107:C:H2' | 31:DA:108:U:C6 | 2.44 | 0.53 |
| 31:DA:330:A:H2 | 31:DA:1210:A:O2' | 1.91 | 0.53 |
| 31:DA:620:G:H5' | 31:DA:620:G:N3 | 2.24 | 0.53 |
| 31:DA:797:C:OP2 | 35:DF:62:ARG:HG3 | 2.09 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1381:G:C2' | 31:DA:1382:G:H5' | 2.38 | 0.53 |
| 31:DA:1722:A:C2 | 31:DA:1740:G:H2' | 2.43 | 0.53 |
| 31:DA:2245:U:H5' | 31:DA:2246:G:H5' | 1.91 | 0.53 |
| 37:DH:47:GLU:C | 37:DH:49:VAL:H | 2.17 | 0.53 |
| 39:DN:42:TRP:CB | 46:DU:64:ARG:NH1 | 2.62 | 0.53 |
| 40:DO:60:ALA:CB | 40:DO:86:ILE:HA | 2.39 | 0.53 |
| 41:DP:45:LEU:HD22 | 41:DP:46:LYS:H | 1.73 | 0.53 |
| 44:DS:26:LEU:HD13 | 44:DS:87:PHE:HD1 | 1.74 | 0.53 |
| 50:DY:28:LYS:HD2 | 50:DY:37:VAL:CG1 | 2.39 | 0.53 |
| 1:AA:308:C:H2' | 1:AA:309:G:H8 | 1.74 | 0.53 |
| 1:AA:625:G:H2' | 1:AA:626:U:C6 | 2.44 | 0.53 |
| 1:AA:1125:U:O4 | 10:AJ:5:ARG:HD2 | 2.09 | 0.53 |
| 1:AA:1350:A:OP1 | 9:AI:121:ARG:HG3 | 2.09 | 0.53 |
| 1:AA:1470:G:C2' | 1:AA:1471:G:H5' | 2.38 | 0.53 |
| 3:AC:35:GLU:O | 3:AC:39:ILE:HG13 | 2.08 | 0.53 |
| 9:AI:18:PHE:HB3 | 9:AI:20:ARG:NH1 | 2.24 | 0.53 |
| 9:AI:53:VAL:HB | 9:AI:92:TYR:CE2 | 2.44 | 0.53 |
| 12:AL:27:LEU:C | 12:AL:29:GLY:N | 2.64 | 0.53 |
| 23:B1:89:GLU:CD | 23:B1:89:GLU:N | 2.54 | 0.53 |
| 24:B2:47:ASN:N | 24:B2:47:ASN:HD22 | 2.07 | 0.53 |
| 27:B5:42:PRO:CB | 27:B5:43:HIS:HD2 | 2.22 | 0.53 |
| 30:B8:35:GLN:HA | 31:BA:2420:C:OP2 | 2.09 | 0.53 |
| 30:B8:53:PRO:O | 30:B8:54:GLU:C | 2.52 | 0.53 |
| 31:BA:53:A:H2' | 31:BA:54:G:O4' | 2.09 | 0.53 |
| 31:BA:197:A:C8 | 31:BA:197:A:C4' | 2.92 | 0.53 |
| 31:BA:1107:G:H2' | 31:BA:1108:U:O4' | 2.08 | 0.53 |
| 31:BA:2302:G:C6 | 31:BA:2315:G:C6 | 2.96 | 0.53 |
| 31:BA:2758:A:C2' | 31:BA:2759:G:H5" | 2.37 | 0.53 |
| 34:BE:78:LEU:N | 34:BE:78:LEU:HD23 | 2.24 | 0.53 |
| 40:BO:43:VAL:HG12 | 40:BO:54:GLU:HA | 1.91 | 0.53 |
| 40:BO:112:MET:HA | 40:BO:112:MET:CE | 2.39 | 0.53 |
| 44:BS:66:ALA:O | 44:BS:67:ARG:HB3 | 2.09 | 0.53 |
| 44:BS:89:ARG:HE | 44:BS:89:ARG:CA | 2.10 | 0.53 |
| 44:BS:99:LYS:O | 44:BS:101:LEU:HB3 | 2.09 | 0.53 |
| 45:BT:33:LYS:NZ | 45:BT:33:LYS:HA | 2.24 | 0.53 |
| 47:BV:49:THR:HG23 | 47:BV:50:PRO:HD2 | 1.90 | 0.53 |
| 51:BZ:44:PHE:CD1 | 51:BZ:44:PHE:C | 2.87 | 0.53 |
| 51:BZ:54:HIS:HB3 | 51:BZ:101:PRO:HD3 | 1.91 | 0.53 |
| 1:CA:560:U:H4' | 1:CA:561:U:O5' | 2.08 | 0.53 |
| 1:CA:1150:U:O4 | 1:CA:1151:A:N6 | 2.42 | 0.53 |
| 3:CC:14:ILE:O | 3:CC:15:THR:HB | 2.08 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 6:CF:99:ALA:HB1 | 18:CR:23:LYS:HZ2 | 1.74 | 0.53 |
| 30:D8:43:GLN:C | 30:D8:44:LYS:HD2 | 2.34 | 0.53 |
| 31:DA:49:A:H3' | 31:DA:50:U:H5' | 1.91 | 0.53 |
| 31:DA:330:A:H2 | 31:DA:1210:A:C2' | 2.22 | 0.53 |
| 31:DA:460:A:C2 | 31:DA:470:A:C5 | 2.97 | 0.53 |
| 31:DA:607:U:OP1 | 35:DF:102:PRO:HA | 2.09 | 0.53 |
| 31:DA:623:G:H2' | 31:DA:624:C:C6 | 2.44 | 0.53 |
| 31:DA:1027:A:C6 | 31:DA:1126:A:C4 | 2.97 | 0.53 |
| 31:DA:1047:G:N2 | 31:DA:1111:A:N6 | 2.56 | 0.53 |
| 31:DA:1332:G:N2 | 31:DA:1609:A:O2' | 2.42 | 0.53 |
| 31:DA:1531:C:H5' | 31:DA:1532:C:OP2 | 2.09 | 0.53 |
| 31:DA:2262:U:C2' | 31:DA:2263:C:H5' | 2.39 | 0.53 |
| 32:DB:28:C:H2' | 32:DB:29:A:C8 | 2.44 | 0.53 |
| 33:DD:211:ARG:HA | 33:DD:214:TRP:CG | 2.44 | 0.53 |
| 34:DE:69:LYS:O | 34:DE:70:ALA:C | 2.51 | 0.53 |
| 34:DE:152:LYS:HE2 | 39:DN:78:TYR:HD2 | 1.74 | 0.53 |
| 35:DF:22:ALA:HA | 35:DF:26:ALA:CB | 2.37 | 0.53 |
| 35:DF:57:VAL:HG13 | 35:DF:58:ALA:N | 2.24 | 0.53 |
| 38:DI:68:LEU:HA | 38:DI:71:ILE:HG23 | 1.91 | 0.53 |
| 50:DY:15:VAL:O | 50:DY:16:ALA:HB2 | 2.08 | 0.53 |
| 1:AA:1202:G:C2' | 1:AA:1203:C:H5' | 2.39 | 0.53 |
| 2:AB:21:ARG:O | 2:AB:21:ARG:HG3 | 2.07 | 0.53 |
| 2:AB:172:ILE:H | 2:AB:172:ILE:CD1 | 2.19 | 0.53 |
| 4:AD:30:LYS:C | 4:AD:32:ALA:H | 2.17 | 0.53 |
| 5:AE:80:ILE:CD1 | 5:AE:138:ALA:HB1 | 2.37 | 0.53 |
| 5:AE:120:THR:O | 5:AE:121:LYS:CB | 2.57 | 0.53 |
| 6:AF:96:PRO:HB3 | 18:AR:30:ASP:CG | 2.34 | 0.53 |
| 10:AJ:24:VAL:O | 10:AJ:28:ARG:HG3 | 2.09 | 0.53 |
| 13:AM:16:ASP:HB3 | 13:AM:41:PRO:HB3 | 1.90 | 0.53 |
| 20:AT:56:MET:CG | 20:AT:88:VAL:HG21 | 2.35 | 0.53 |
| 22:B0:26:TYR:CE2 | 31:BA:857:C:H1' | 2.44 | 0.53 |
| 22:B0:50:ASN:HD22 | 22:B0:83:PRO:HD3 | 1.74 | 0.53 |
| 31:BA:309:G:H4' | 50:BY:18:GLY:HA3 | 1.90 | 0.53 |
| 31:BA:1719:G:H2' | 31:BA:1720:U:C5' | 2.39 | 0.53 |
| 31:BA:2280:G:H2' | 31:BA:2281:C:H5' | 1.89 | 0.53 |
| 31:BA:2830:G:C5' | 31:BA:2830:G:C8 | 2.91 | 0.53 |
| 32:BB:40:U:H1' | 32:BB:45:A:H61 | 1.74 | 0.53 |
| 33:BD:27:THR:O | 33:BD:28:GLU:HB2 | 2.09 | 0.53 |
| 33:BD:35:LYS:CD | 33:BD:63:ARG:C | 2.82 | 0.53 |
| 34:BE:182:LEU:HD12 | 34:BE:182:LEU:C | 2.34 | 0.53 |
| 35:BF:199:TRP:CZ2 | 35:BF:203:GLN:NE2 | 2.77 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 37:BH:85:LYS:CE | 37:BH:145:ALA:N | 2.72 | 0.53 |
| 37:BH:127:GLU:HA | 37:BH:127:GLU:OE1 | 2.09 | 0.53 |
| 37:BH:153:LYS:HB2 | 37:BH:154:PRO:CD | 2.39 | 0.53 |
| 41:BP:17:LYS:O | 41:BP:19:VAL:HG23 | 2.09 | 0.53 |
| 46:BU:29:SER:OG | 46:BU:30:LYS:HE3 | 2.09 | 0.53 |
| 46:BU:51:LYS:O | 46:BU:53:ARG:N | 2.42 | 0.53 |
| 49:BX:37:THR:C | 49:BX:38:GLU:OE1 | 2.52 | 0.53 |
| 51:BZ:8:TYR:N | 51:BZ:8:TYR:CD1 | 2.76 | 0.53 |
| 1:CA:832:C:N4 | 1:CA:855:G:C6 | 2.77 | 0.53 |
| 1:CA:1129:C:H4' | 1:CA:1130:A:H5' | 1.92 | 0.53 |
| 6:CF:3:ARG:HH11 | 6:CF:3:ARG:HG3 | 1.74 | 0.53 |
| 27:D5:41:PRO:HG2 | 27:D5:44:THR:OG1 | 2.09 | 0.53 |
| 30:D8:26:LYS:HB2 | 30:D8:44:LYS:HG3 | 1.91 | 0.53 |
| 31:DA:806:C:OP2 | 41:DP:39:LYS:CG | 2.57 | 0.53 |
| 31:DA:807:U:C2' | 31:DA:808:G:O5' | 2.57 | 0.53 |
| 31:DA:860:U:C5 | 31:DA:2268:A:C8 | 2.96 | 0.53 |
| 31:DA:1268:A:H2' | 31:DA:1269:A:O4' | 2.08 | 0.53 |
| 31:DA:1315:C:H42 | 31:DA:1337:G:H1 | 1.56 | 0.53 |
| 31:DA:1515:G:H2' | 31:DA:1516:C:H6 | 1.74 | 0.53 |
| 31:DA:1593:G:H2' | 31:DA:1594:G:C8 | 2.44 | 0.53 |
| 31:DA:1783:A:N1 | 31:DA:2587:A:C4 | 2.77 | 0.53 |
| 31:DA:2199:A:C8 | 31:DA:2200:C:C5 | 2.97 | 0.53 |
| 31:DA:2360:A:O2' | 31:DA:2361:A:OP2 | 2.28 | 0.53 |
| 31:DA:2731:G:C6 | 31:DA:2732:G:O6 | 2.62 | 0.53 |
| 32:DB:13:A:N1 | 32:DB:69:G:O2' | 2.38 | 0.53 |
| 33:DD:35:LYS:CE | 33:DD:104:TYR:CD1 | 2.92 | 0.53 |
| 33:DD:84:TYR:HE2 | 33:DD:86:PRO:HB3 | 1.74 | 0.53 |
| 33:DD:270:ILE:C | 33:DD:271:ILE:HG13 | 2.32 | 0.53 |
| 36:DG:60:LEU:HD13 | 36:DG:60:LEU:C | 2.34 | 0.53 |
| 39:DN:13:TRP:O | 39:DN:135:PRO:HG2 | 2.09 | 0.53 |
| 41:DP:16:ARG:O | 41:DP:18:ARG:N | 2.42 | 0.53 |
| 44:DS:38:GLN:HB3 | 44:DS:47:THR:HG21 | 1.90 | 0.53 |
| 45:DT:124:ASP:C | 45:DT:126:ALA:H | 2.16 | 0.53 |
| 47:DV:25:LEU:HD12 | 47:DV:94:LEU:HD22 | 1.91 | 0.53 |
| 1:AA:33:A:H2' | 1:AA:34:C:C6 | 2.44 | 0.52 |
| 1:AA:200:G:H1 | 1:AA:217:C:N4 | 2.03 | 0.52 |
| 1:AA:818:G:HO2' | 1:AA:820:U:H6 | 1.54 | 0.52 |
| 1:AA:862:C:O2' | 1:AA:863:U:H5' | 2.09 | 0.52 |
| 1:AA:932:C:H4' | 7:AG:4:ARG:NH2 | 2.23 | 0.52 |
| 1:AA:963:G:N2 | 10:AJ:55:LYS:HE2 | 2.24 | 0.52 |
| 1:AA:1236:A:O2' | 1:AA:1304:G:H4' | 2.09 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:1240:U:OP2 | 7:AG:116:ALA:HB2 | 2.09 | 0.52 |
| 3:AC:71:ALA:HA | 3:AC:106:VAL:HB | 1.89 | 0.52 |
| 5:AE:101:ILE:HD11 | 5:AE:119:LEU:HD23 | 1.90 | 0.52 |
| 8:AH:64:LYS:HG2 | 8:AH:79:VAL:HG21 | 1.90 | 0.52 |
| 23:B1:25:LYS:O | 23:B1:26:ARG:HB3 | 2.10 | 0.52 |
| 31:BA:795:C:O2' | 31:BA:796:C:H5' | 2.09 | 0.52 |
| 31:BA:828:U:H4' | 31:BA:831:G:N1 | 2.24 | 0.52 |
| 31:BA:893:C:H2' | 31:BA:894:C:O5' | 2.09 | 0.52 |
| 31:BA:1049:C:H1' | 31:BA:1113:U:O2' | 2.08 | 0.52 |
| 31:BA:1648:C:C2' | 31:BA:1649:G:O5' | 2.57 | 0.52 |
| 31:BA:2327:A:H2' | 31:BA:2328:A:H8 | 1.70 | 0.52 |
| 34:BE:134:ILE:HD13 | 34:BE:134:ILE:H | 1.74 | 0.52 |
| 35:BF:53:THR:C | 35:BF:55:GLY:N | 2.66 | 0.52 |
| 38:BI:38:LEU:HB2 | 38:BI:40:THR:HG23 | 1.89 | 0.52 |
| 39:BN:75:TYR:CD2 | 39:BN:83:LYS:NZ | 2.68 | 0.52 |
| 40:BO:71:ARG:HE | 40:BO:105:GLU:CD | 2.15 | 0.52 |
| 47:BV:40:LEU:O | 47:BV:40:LEU:HD12 | 2.09 | 0.52 |
| 47:BV:52:VAL:O | 47:BV:53:GLU:CB | 2.57 | 0.52 |
| 50:BY:76:CYS:O | 50:BY:99:CYS:SG | 2.67 | 0.52 |
| 51:BZ:29:TYR:HA | 51:BZ:33:LEU:O | 2.08 | 0.52 |
| 1:CA:1321:C:H5" | 1:CA:1322:C:H2' | 1.89 | 0.52 |
| 4:CD:9:CYS:HA | 4:CD:12:CYS:HB2 | 1.91 | 0.52 |
| 10:CJ:26:ALA:HB1 | 10:CJ:29:ARG:NH2 | 2.22 | 0.52 |
| 11:CK:126:ARG:O | 11:CK:127:LYS:C | 2.52 | 0.52 |
| 16:CP:21:VAL:HG22 | 16:CP:34:GLU:O | 2.09 | 0.52 |
| 23:D1:13:ILE:HG12 | 23:D1:14:VAL:CA | 2.37 | 0.52 |
| 23:D1:76:ARG:HB3 | 23:D1:78:LYS:HE3 | 1.91 | 0.52 |
| 25:D3:45:GLY:O | 25:D3:48:GLU:HB2 | 2.09 | 0.52 |
| 29:D7:34:ARG:NH1 | 31:DA:466:A:OP1 | 2.41 | 0.52 |
| 31:DA:1362:C:O2' | 31:DA:1363:C:H5' | 2.09 | 0.52 |
| 31:DA:1684:C:C2' | 31:DA:1685:C:H5' | 2.40 | 0.52 |
| 31:DA:1857:G:C6 | 31:DA:1858:G:N1 | 2.77 | 0.52 |
| 31:DA:2531:A:H2 | 31:DA:2658:C:O2 | 1.92 | 0.52 |
| 31:DA:2854:G:H2' | 31:DA:2855:C:C6 | 2.44 | 0.52 |
| 31:DA:2869:G:H2' | 31:DA:2870:C:O4' | 2.10 | 0.52 |
| 34:DE:3:GLY:O | 34:DE:4:ILE:HB | 2.09 | 0.52 |
| 39:DN:17:ASP:C | 39:DN:19:GLU:H | 2.17 | 0.52 |
| 40:DO:24:VAL:CG2 | 40:DO:33:ALA:HB2 | 2.39 | 0.52 |
| 42:DQ:141:GLN:CD | 51:DZ:72:ARG:HG2 | 2.35 | 0.52 |
| 42:DQ:141:GLN:HE22 | 51:DZ:89:PHE:HB3 | 1.74 | 0.52 |
| 43:DR:71:GLN:HE21 | 43:DR:71:GLN:CA | 2.21 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 51:DZ:29:TYR:CE2 | 51:DZ:87:ASP:HB2 | 2.43 | 0.52 |
| 51:DZ:98:MET:HE3 | 51:DZ:99:TYR:O | 2.09 | 0.52 |
| 1:AA:174:C:C5 | 1:AA:175:C:H5 | 2.27 | 0.52 |
| 1:AA:266:G:H5" | 1:AA:268:C:N4 | 2.21 | 0.52 |
| 1:AA:579:G:C5 | 1:AA:580:U:C5 | 2.97 | 0.52 |
| 1:AA:792:A:H4' | 1:AA:793:U:O5' | 2.08 | 0.52 |
| 1:AA:827:U:N3 | 1:AA:870:U:C4 | 2.77 | 0.52 |
| 1:AA:1126:U:H2' | 1:AA:1127:G:O4' | 2.09 | 0.52 |
| 1:AA:1470:G:O2' | 1:AA:1471:G:H5' | 2.09 | 0.52 |
| 11:AK:57:THR:HG22 | 11:AK:59:TYR:H | 1.75 | 0.52 |
| 12:AL:32:PHE:HB3 | 12:AL:84:LEU:HD21 | 1.91 | 0.52 |
| 12:AL:60:LEU:HD21 | 12:AL:66:VAL:CG2 | 2.38 | 0.52 |
| 15:AO:5:LYS:O | 15:AO:9:GLN:HG2 | 2.08 | 0.52 |
| 24:B2:15:LYS:O | 24:B2:16:LEU:HB3 | 2.08 | 0.52 |
| 25:B3:1:MET:HB3 | 25:B3:2:PRO:HD2 | 1.91 | 0.52 |
| 25:B3:45:GLY:O | 25:B3:48:GLU:HB2 | 2.09 | 0.52 |
| 31:BA:90:U:O2 | 31:BA:90:U:O4' | 2.25 | 0.52 |
| 31:BA:241:A:O4' | 31:BA:243:U:C6 | 2.62 | 0.52 |
| 31:BA:559:G:H22 | 46:BU:49:HIS:CD2 | 2.27 | 0.52 |
| 31:BA:701:G:N2 | 31:BA:732:C:C2 | 2.77 | 0.52 |
| 31:BA:1131:G:HO2' | 31:BA:1132:A:H8 | 1.57 | 0.52 |
| 31:BA:1331:A:H2' | 31:BA:1333:C:H5 | 1.73 | 0.52 |
| 31:BA:1531:C:H5' | 31:BA:1532:C:OP2 | 2.09 | 0.52 |
| 31:BA:1742:G:H3' | 31:BA:1742:G:C8 | 2.43 | 0.52 |
| 31:BA:1836:C:C2' | 31:BA:1837:C:H5' | 2.39 | 0.52 |
| 31:BA:2287:A:C4 | 31:BA:2289:G:C8 | 2.97 | 0.52 |
| 33:BD:172:TYR:CD1 | 33:BD:186:HIS:HA | 2.44 | 0.52 |
| 35:BF:22:ALA:CA | 35:BF:26:ALA:CB | 2.87 | 0.52 |
| 35:BF:132:VAL:CG2 | 35:BF:133:ASN:H | 2.22 | 0.52 |
| 35:BF:185:ASP:OD1 | 35:BF:188:ARG:NH1 | 2.39 | 0.52 |
| 36:BG:71:THR:HB | 36:BG:89:GLY:HA3 | 1.91 | 0.52 |
| 40:BO:10:VAL:HG13 | 40:BO:17:ARG:O | 2.09 | 0.52 |
| 42:BQ:141:GLN:HE22 | 51:BZ:89:PHE:HB3 | 1.72 | 0.52 |
| 46:BU:117:GLN:HA | 46:BU:117:GLN:OE1 | 2.08 | 0.52 |
| 2:CB:187:LEU:HD11 | 2:CB:204:ASN:O | 2.09 | 0.52 |
| 5:CE:68:GLU:O | 5:CE:68:GLU:HG3 | 2.09 | 0.52 |
| 6:CF:30:LEU:HD23 | 6:CF:30:LEU:H | 1.75 | 0.52 |
| 8:CH:6:ILE:H | 8:CH:6:ILE:HD12 | 1.73 | 0.52 |
| 9:CI:18:PHE:HB2 | 9:CI:62:TYR:O | 2.09 | 0.52 |
| 11:CK:105:VAL:HG23 | 11:CK:105:VAL:O | 2.08 | 0.52 |
| 12:CL:27:LEU:C | 12:CL:29:GLY:N | 2.63 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 13:CM:81:LEU:HB3 | 13:CM:89:GLY:CA | 2.38 | 0.52 |
| 31:DA:384:U:H2' | 31:DA:385:C:C6 | 2.42 | 0.52 |
| 31:DA:848:G:C4 | 31:DA:933:A:H8 | 2.27 | 0.52 |
| 31:DA:1049:C:O2 | 31:DA:1049:C:H2' | 2.10 | 0.52 |
| 31:DA:1563:G:C5 | 31:DA:1564:C:C5 | 2.97 | 0.52 |
| 31:DA:1657:C:H5" | 34:DE:133:LYS:O | 2.09 | 0.52 |
| 31:DA:1803:A:C8 | 31:DA:1804:C:C5 | 2.97 | 0.52 |
| 32:DB:94:C:H2' | 32:DB:95:C:H6 | 1.73 | 0.52 |
| 33:DD:62:TYR:CE1 | 33:DD:64:ILE:HA | 2.44 | 0.52 |
| 34:DE:70:ALA:O | 34:DE:72:VAL:N | 2.43 | 0.52 |
| 37:DH:43:VAL:HG11 | 37:DH:53:GLU:O | 2.09 | 0.52 |
| 39:DN:97:ARG:O | 39:DN:100:GLU:N | 2.42 | 0.52 |
| 41:DP:62:LEU:H | 41:DP:62:LEU:CD2 | 2.21 | 0.52 |
| 50:DY:22:GLY:O | 50:DY:23:ARG:HG2 | 2.10 | 0.52 |
| 1:AA:1207:G:H2' | 1:AA:1208:C:C6 | 2.44 | 0.52 |
| 2:AB:187:LEU:HD11 | 2:AB:204:ASN:O | 2.09 | 0.52 |
| 15:AO:87:ILE:CG2 | 15:AO:88:ARG:H | 2.21 | 0.52 |
| 17:AQ:87:LYS:HE2 | 17:AQ:87:LYS:HA | 1.91 | 0.52 |
| 31:BA:610:G:H2' | 31:BA:611:C:C6 | 2.44 | 0.52 |
| 31:BA:1044:G:N2 | 31:BA:1112:G:O6 | 2.42 | 0.52 |
| 31:BA:1593:G:H2' | 31:BA:1594:G:H8 | 1.75 | 0.52 |
| 31:BA:2642:G:N2 | 31:BA:2773:C:C2 | 2.78 | 0.52 |
| 31:BA:2830:G:H5" | 31:BA:2830:G:H8 | 1.73 | 0.52 |
| 31:BA:2854:G:H2' | 31:BA:2855:C:C6 | 2.43 | 0.52 |
| 33:BD:30:GLU:CD | 33:BD:63:ARG:NE | 2.68 | 0.52 |
| 33:BD:131:LEU:HD12 | 33:BD:131:LEU:N | 2.24 | 0.52 |
| 46:BU:25:TRP:HD1 | 46:BU:26:GLY:N | 2.08 | 0.52 |
| 49:BX:65:ARG:HD3 | 49:BX:65:ARG:C | 2.34 | 0.52 |
| 51:BZ:102:LEU:HG | 51:BZ:123:ASP:HA | 1.91 | 0.52 |
| 1:CA:170:U:O2' | 1:CA:171:A:H5' | 2.09 | 0.52 |
| 1:CA:340:U:O2' | 1:CA:341:C:H5' | 2.09 | 0.52 |
| 2:CB:208:ILE:HA | 2:CB:211:ILE:HD12 | 1.91 | 0.52 |
| 16:CP:59:TRP:O | 16:CP:64:ALA:HB3 | 2.09 | 0.52 |
| 20:CT:67:ALA:HB2 | 20:CT:77:ALA:HB2 | 1.91 | 0.52 |
| 27:D5:56:LYS:O | 27:D5:57:VAL:C | 2.53 | 0.52 |
| 31:DA:435:C:C5 | 31:DA:436:C:C5 | 2.97 | 0.52 |
| 31:DA:1657:C:H2' | 31:DA:1658:C:C6 | 2.43 | 0.52 |
| 31:DA:2432:A:H2' | 31:DA:2433:A:C8 | 2.44 | 0.52 |
| 31:DA:2536:G:C5 | 31:DA:2537:U:C5 | 2.97 | 0.52 |
| 31:DA:2859:G:C8 | 31:DA:2859:G:C3' | 2.92 | 0.52 |
| 32:DB:15:A:HO2' | 32:DB:110:G:H8 | 1.48 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 32:DB:66:A:H61 | 32:DB:108:U:H2' | 1.74 | 0.52 |
| 36:DG:106:LEU:HD12 | 36:DG:110:ALA:HB3 | 1.89 | 0.52 |
| 38:DI:9:LEU:N | 38:DI:13:GLY:HA3 | 2.12 | 0.52 |
| 38:DI:129:THR:HG22 | 38:DI:130:TYR:O | 2.08 | 0.52 |
| 39:DN:55:VAL:HG12 | 39:DN:126:PRO:HA | 1.92 | 0.52 |
| 46:DU:92:ARG:O | 46:DU:95:LEU:N | 2.42 | 0.52 |
| 47:DV:4:ILE:O | 47:DV:39:LEU:CB | 2.57 | 0.52 |
| 47:DV:19:LYS:HE2 | 47:DV:20:LEU:HD12 | 1.91 | 0.52 |
| 47:DV:49:THR:HG23 | 47:DV:50:PRO:HD2 | 1.90 | 0.52 |
| 50:DY:16:ALA:HA | 50:DY:21:LYS:HD2 | 1.89 | 0.52 |
| 50:DY:52:SER:O | 50:DY:54:LYS:N | 2.42 | 0.52 |
| 1:AA:189(E):U:O2' | 1:AA:189(F):U:H5' | 2.09 | 0.52 |
| 6:AF:23:LYS:O | 6:AF:27:GLN:HG2 | 2.10 | 0.52 |
| 6:AF:96:PRO:HB3 | 18:AR:30:ASP:OD2 | 2.10 | 0.52 |
| 7:AG:62:PHE:HA | 7:AG:124:LEU:HD22 | 1.92 | 0.52 |
| 12:AL:6:THR:OG1 | 12:AL:9:GLN:HG3 | 2.09 | 0.52 |
| 23:B1:85:LEU:CA | 23:B1:87:PRO:HD3 | 2.40 | 0.52 |
| 23:B1:87:PRO:O | 23:B1:91:LYS:N | 2.34 | 0.52 |
| 29:B7:1:MET:O | 29:B7:2:LYS:C | 2.52 | 0.52 |
| 31:BA:195:A:H61 | 31:BA:198:C:H3' | 1.75 | 0.52 |
| 31:BA:247:G:H4' | 31:BA:386:G:C5 | 2.44 | 0.52 |
| 31:BA:819:A:N3 | 31:BA:1189:A:C2 | 2.78 | 0.52 |
| 31:BA:1000:A:C6 | 31:BA:1001:A:C6 | 2.98 | 0.52 |
| 31:BA:1106:A:H2' | 31:BA:1107:G:O5' | 2.08 | 0.52 |
| 31:BA:1488:G:C5 | 31:BA:1489:U:N3 | 2.78 | 0.52 |
| 31:BA:1784:A:C4' | 31:BA:1785:A:H5" | 2.39 | 0.52 |
| 31:BA:2308:G:H2' | 31:BA:2309:A:C8 | 2.45 | 0.52 |
| 32:BB:78:A:C2 | 32:BB:100:A:C4 | 2.98 | 0.52 |
| 33:BD:35:LYS:NZ | 33:BD:104:TYR:CB | 2.54 | 0.52 |
| 34:BE:47:VAL:HG12 | 34:BE:49:LEU:HD22 | 1.90 | 0.52 |
| 38:BI:57:ARG:O | 38:BI:60:GLU:HB2 | 2.09 | 0.52 |
| 41:BP:88:LEU:O | 41:BP:90:ARG:N | 2.43 | 0.52 |
| 42:BQ:16:ARG:HB2 | 42:BQ:16:ARG:NH1 | 2.25 | 0.52 |
| 45:BT:99:LEU:O | 45:BT:99:LEU:HD13 | 2.10 | 0.52 |
| 45:BT:100:TYR:HD2 | 45:BT:103:ARG:NH2 | 2.07 | 0.52 |
| 47:BV:2:PHE:HE1 | 47:BV:13:ARG:NH2 | 2.06 | 0.52 |
| 48:BW:56:ALA:O | 48:BW:57:ASN:C | 2.52 | 0.52 |
| 49:BX:25:LYS:HG3 | 49:BX:26:TYR:CD1 | 2.44 | 0.52 |
| 50:BY:15:VAL:O | 50:BY:16:ALA:HB2 | 2.08 | 0.52 |
| 1:CA:159:G:HO2' | 1:CA:160:A:H8 | 1.49 | 0.52 |
| 1:CA:525:C:H2' | 1:CA:526:C:C6 | 2.44 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:718:G:H5' | 11:CK:117:ASN:HB2 | 1.90 | 0.52 |
| 1:CA:808:C:P | 15:CO:48:LYS:HE3 | 2.50 | 0.52 |
| 1:CA:996:A:H2' | 1:CA:997:U:O4' | 2.10 | 0.52 |
| 1:CA:1287:A:C2 | 1:CA:1353:G:H1' | 2.44 | 0.52 |
| 1:CA:1501:C:H5" | 1:CA:1502:A:OP2 | 2.09 | 0.52 |
| 2:CB:102:LEU:HD12 | 2:CB:102:LEU:N | 2.24 | 0.52 |
| 3:CC:11:ARG:O | 3:CC:13:GLY:N | 2.43 | 0.52 |
| 9:CI:112:LYS:HE3 | 9:CI:116:LYS:O | 2.10 | 0.52 |
| 15:CO:18:PHE:CE1 | 15:CO:21:ASP:HB2 | 2.45 | 0.52 |
| 18:CR:45:SER:H | 18:CR:51:LEU:HD11 | 1.74 | 0.52 |
| 20:CT:56:MET:CG | 20:CT:88:VAL:HG21 | 2.35 | 0.52 |
| 22:D0:72:ARG:O | 22:D0:75:LEU:HB2 | 2.10 | 0.52 |
| 30:D8:27:THR:HA | 41:DP:62:LEU:HD11 | 1.91 | 0.52 |
| 30:D8:32:LEU:HB3 | 30:D8:35:GLN:N | 2.24 | 0.52 |
| 31:DA:107:C:C2 | 31:DA:108:U:C5 | 2.96 | 0.52 |
| 31:DA:892:G:C5 | 31:DA:893:C:C5 | 2.98 | 0.52 |
| 31:DA:1563:G:H2' | 31:DA:1564:C:H6 | 1.73 | 0.52 |
| 31:DA:2801:A:H4' | 31:DA:2801(A):A:H5' | 1.91 | 0.52 |
| 32:DB:27:C:C4 | 32:DB:28:C:C4 | 2.98 | 0.52 |
| 33:DD:35:LYS:HE3 | 33:DD:63:ARG:C | 2.34 | 0.52 |
| 36:DG:107:LEU:HD11 | 36:DG:178:PHE:CE1 | 2.44 | 0.52 |
| 36:DG:120:LEU:HB2 | 36:DG:179:PRO:O | 2.09 | 0.52 |
| 41:DP:62:LEU:HD13 | 41:DP:62:LEU:N | 2.20 | 0.52 |
| 42:DQ:7:MET:O | 42:DQ:10:ARG:NE | 2.39 | 0.52 |
| 46:DU:14:HIS:CD2 | 46:DU:32:PHE:CB | 2.89 | 0.52 |
| 1:AA:349:A:O2' | 1:AA:350:G:H5' | 2.08 | 0.52 |
| 1:AA:718:G:H5' | 11:AK:117:ASN:HB2 | 1.90 | 0.52 |
| 1:AA:1067:A:H1' | 1:AA:1068:G:H8 | 1.75 | 0.52 |
| 1:AA:1125:U:H2' | 1:AA:1126:U:OP2 | 2.10 | 0.52 |
| 2:AB:83:MET:HE2 | 2:AB:83:MET:HA | 1.90 | 0.52 |
| 3:AC:73:PRO:HA | 3:AC:76:VAL:HG13 | 1.92 | 0.52 |
| 6:AF:20:ALA:O | 6:AF:23:LYS:HB2 | 2.09 | 0.52 |
| 9:AI:112:LYS:HE3 | 9:AI:116:LYS:O | 2.10 | 0.52 |
| 10:AJ:32:ALA:HB1 | 10:AJ:75:ILE:CG1 | 2.40 | 0.52 |
| 12:AL:47:LYS:CB | 12:AL:48:PRO:CD | 2.84 | 0.52 |
| 16:AP:14:ASN:N | 16:AP:15:PRO:HD3 | 2.23 | 0.52 |
| 22:B0:1:MET:CB | 31:BA:2602:A:H62 | 2.23 | 0.52 |
| 23:B1:34:THR:HG21 | 31:BA:387:U:O3' | 2.08 | 0.52 |
| 28:B6:36:LEU:C | 28:B6:37:ARG:HD2 | 2.34 | 0.52 |
| 31:BA:17:G:H2' | 31:BA:18:C:C6 | 2.44 | 0.52 |
| 31:BA:271(Q):G:O2' | 31:BA:271(R):G:P | 2.68 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:271(T):C:H2' | 31:BA:271(T):C:O2 | 2.09 | 0.52 |
| 31:BA:768:G:O2' | 31:BA:1379:A:N6 | 2.42 | 0.52 |
| 31:BA:993:G:C5' | 47:BV:75:PHE:CE2 | 2.89 | 0.52 |
| 31:BA:995:C:N3 | 39:BN:4:TYR:CE1 | 2.77 | 0.52 |
| 31:BA:1488:G:C2 | 31:BA:1489:U:O2 | 2.62 | 0.52 |
| 31:BA:1722:A:C2 | 31:BA:1740:G:H2' | 2.44 | 0.52 |
| 33:BD:44:ASN:HB3 | 33:BD:49:ILE:CA | 2.25 | 0.52 |
| 36:BG:25:TYR:CE2 | 36:BG:32:PRO:HD3 | 2.44 | 0.52 |
| 39:BN:42:TRP:CB | 46:BU:64:ARG:NH1 | 2.62 | 0.52 |
| 39:BN:78:TYR:HD1 | 39:BN:79:PRO:HD3 | 1.73 | 0.52 |
| 40:BO:107:ARG:NH1 | 40:BO:112:MET:HE1 | 2.24 | 0.52 |
| 46:BU:14:HIS:CD2 | 46:BU:32:PHE:CB | 2.92 | 0.52 |
| 1:CA:175:C:C2 | 1:CA:176:C:C5 | 2.98 | 0.52 |
| 1:CA:441:A:H3' | 1:CA:442:C:C6 | 2.44 | 0.52 |
| 1:CA:963:G:H21 | 10:CJ:55:LYS:CD | 2.23 | 0.52 |
| 1:CA:1321:C:H3' | 1:CA:1322:C:H5" | 1.92 | 0.52 |
| 3:CC:188:LEU:O | 3:CC:189:ALA:HB2 | 2.09 | 0.52 |
| 5:CE:120:THR:O | 5:CE:121:LYS:CB | 2.57 | 0.52 |
| 8:CH:20:TYR:HD1 | 8:CH:65:TYR:HD2 | 1.50 | 0.52 |
| 16:CP:4:ILE:HD12 | 16:CP:4:ILE:N | 2.25 | 0.52 |
| 22:D0:1:MET:CB | 31:DA:2602:A:H62 | 2.23 | 0.52 |
| 22:D0:71:ASP:C | 22:D0:72:ARG:HG2 | 2.34 | 0.52 |
| 27:D5:36:CYS:HG | 27:D5:49:CYS:CB | 2.20 | 0.52 |
| 31:DA:53:A:H2' | 31:DA:54:G:O4' | 2.09 | 0.52 |
| 31:DA:287:C:C2 | 31:DA:288:C:C6 | 2.98 | 0.52 |
| 31:DA:309:G:N3 | 31:DA:329:G:O2' | 2.42 | 0.52 |
| 31:DA:481:G:C4 | 31:DA:507:A:C2 | 2.98 | 0.52 |
| 31:DA:570:G:H2' | 31:DA:2030:A:N6 | 2.25 | 0.52 |
| 31:DA:573:G:O2' | 31:DA:574:C:H3' | 2.10 | 0.52 |
| 31:DA:727:A:H2 | 33:DD:9:TYR:CD2 | 2.27 | 0.52 |
| 31:DA:754:C:H2' | 31:DA:755:C:H6 | 1.74 | 0.52 |
| 31:DA:838:C:H2' | 31:DA:839:U:H6 | 1.74 | 0.52 |
| 31:DA:1721:G:C2 | 31:DA:1739:U:OP2 | 2.63 | 0.52 |
| 31:DA:1742:G:H3' | 31:DA:1742:G:C8 | 2.44 | 0.52 |
| 31:DA:2287:A:C4 | 31:DA:2289:G:N7 | 2.78 | 0.52 |
| 31:DA:2557:G:H2' | 31:DA:2558:C:C6 | 2.45 | 0.52 |
| 31:DA:2713:A:C3' | 31:DA:2714:G:C5' | 2.87 | 0.52 |
| 35:DF:65:TRP:CH2 | 35:DF:75:HIS:HD2 | 2.27 | 0.52 |
| 37:DH:153:LYS:HB2 | 37:DH:154:PRO:CD | 2.39 | 0.52 |
| 38:DI:5:LEU:O | 38:DI:6:LEU:HD23 | 2.09 | 0.52 |
| 38:DI:136:VAL:O | 38:DI:136:VAL:HG22 | 2.10 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 43:DR:10:LEU:HB3 | 43:DR:17:ARG:CZ | 2.37 | 0.52 |
| 1:AA:152:A:N6 | 1:AA:170:U:C2 | 2.77 | 0.52 |
| 1:AA:914:A:O2' | 1:AA:915:A:H5' | 2.08 | 0.52 |
| 1:AA:1362:C:C2' | 1:AA:1363:C:H5" | 2.39 | 0.52 |
| 4:AD:17:VAL:HG11 | 4:AD:197:PRO:HB2 | 1.92 | 0.52 |
| 12:AL:41:ARG:HG2 | 12:AL:42:THR:N | 2.23 | 0.52 |
| 13:AM:71:ARG:O | 13:AM:71:ARG:HG3 | 2.09 | 0.52 |
| 27:B5:56:LYS:O | 27:B5:57:VAL:C | 2.51 | 0.52 |
| 28:B6:9:LEU:C | 28:B6:9:LEU:HD13 | 2.35 | 0.52 |
| 29:B7:5:TRP:NE1 | 29:B7:7:PRO:HG3 | 2.24 | 0.52 |
| 31:BA:102:G:C2' | 31:BA:103:A:OP2 | 2.58 | 0.52 |
| 31:BA:143:G:H2' | 31:BA:143(A):C:H6 | 1.75 | 0.52 |
| 31:BA:271(K):U:H3' | 31:BA:271(L):U:H5' | 1.92 | 0.52 |
| 31:BA:415:A:H2' | 31:BA:416:C:H6 | 1.74 | 0.52 |
| 31:BA:573:G:C6 | 31:BA:2030:A:H3' | 2.45 | 0.52 |
| 31:BA:1469:A:H2' | 31:BA:1470:G:O5' | 2.10 | 0.52 |
| 31:BA:1592:C:C2' | 31:BA:1593:G:H5' | 2.40 | 0.52 |
| 31:BA:1987:G:H8 | 31:BA:1987:G:C5' | 2.23 | 0.52 |
| 31:BA:2534:A:C2 | 31:BA:2535:G:H1' | 2.45 | 0.52 |
| 31:BA:2632:A:H1' | 34:BE:61:ARG:HH12 | 1.73 | 0.52 |
| 31:BA:2644:G:O2' | 31:BA:2645:G:H5' | 2.10 | 0.52 |
| 31:BA:2713:A:C3' | 31:BA:2714:G:C5' | 2.87 | 0.52 |
| 33:BD:39:LYS:HB2 | 33:BD:62:TYR:CB | 2.29 | 0.52 |
| 33:BD:48:ARG:HG3 | 33:BD:48:ARG:HH11 | 1.73 | 0.52 |
| 36:BG:81:LYS:O | 36:BG:82:LEU:O | 2.27 | 0.52 |
| 48:BW:15:ARG:HA | 48:BW:18:ARG:HD2 | 1.91 | 0.52 |
| 1:CA:49:U:C2 | 1:CA:361:G:N2 | 2.78 | 0.52 |
| 1:CA:130:A:H1' | 1:CA:263:A:O2' | 2.10 | 0.52 |
| 1:CA:402:G:C6 | 1:CA:403:C:C4 | 2.98 | 0.52 |
| 1:CA:447:G:C6 | 1:CA:485:G:H1' | 2.45 | 0.52 |
| 1:CA:1162:C:C2 | 1:CA:1175:G:N2 | 2.78 | 0.52 |
| 1:CA:1530:G:OP1 | 1:CA:1530:G:H4' | 2.10 | 0.52 |
| 3:CC:18:TRP:CD1 | 14:CN:53:LEU:O | 2.63 | 0.52 |
| 4:CD:30:LYS:C | 4:CD:32:ALA:H | 2.17 | 0.52 |
| 5:CE:6:PHE:CE1 | 5:CE:66:MET:HE1 | 2.44 | 0.52 |
| 6:CF:12:PRO:O | 6:CF:14:LEU:N | 2.41 | 0.52 |
| 6:CF:96:PRO:HB3 | 18:CR:30:ASP:CG | 2.34 | 0.52 |
| 10:CJ:45:ARG:HB2 | 10:CJ:65:LEU:HB3 | 1.91 | 0.52 |
| 10:CJ:58:ASP:O | 10:CJ:59:SER:C | 2.52 | 0.52 |
| 11:CK:81:ASP:OD2 | 11:CK:106:LYS:HG2 | 2.09 | 0.52 |
| 12:CL:119:LYS:HB2 | 12:CL:120:TYR:HD1 | 1.74 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 20:CT:80:ARG:O | 20:CT:84:LEU:HB2 | 2.09 | 0.52 |
| 22:D0:43:THR:H | 31:DA:2331:G:H4' | 1.74 | 0.52 |
| 27:D5:27:PRO:HB3 | 48:DW:23:LEU:CD1 | 2.39 | 0.52 |
| 31:DA:142:A:H8 | 31:DA:1408:C:H1' | 1.69 | 0.52 |
| 31:DA:358:U:H5 | 31:DA:359:A:N7 | 2.08 | 0.52 |
| 31:DA:430:G:H5" | 31:DA:431:U:OP2 | 2.10 | 0.52 |
| 31:DA:1158:C:C2' | 31:DA:1159:U:H5' | 2.40 | 0.52 |
| 31:DA:1225:G:OP1 | 47:DV:88:ARG:CB | 2.57 | 0.52 |
| 31:DA:2287:A:C5 | 31:DA:2289:G:C5 | 2.98 | 0.52 |
| 31:DA:2632:A:C2 | 34:DE:61:ARG:HD2 | 2.44 | 0.52 |
| 32:DB:65:C:N4 | 32:DB:109:C:C2 | 2.78 | 0.52 |
| 32:DB:102:A:H8 | 32:DB:102:A:OP2 | 1.92 | 0.52 |
| 33:DD:176:ARG:O | 33:DD:176:ARG:HG3 | 2.08 | 0.52 |
| 35:DF:199:TRP:CZ2 | 35:DF:203:GLN:NE2 | 2.78 | 0.52 |
| 37:DH:85:LYS:NZ | 37:DH:145:ALA:CA | 2.71 | 0.52 |
| 38:DI:94:ALA:HB2 | 38:DI:116:LEU:HD23 | 1.92 | 0.52 |
| 39:DN:128:HIS:O | 39:DN:128:HIS:CG | 2.62 | 0.52 |
| 41:DP:70:GLN:HG3 | 41:DP:71:VAL:H | 1.75 | 0.52 |
| 45:DT:28:VAL:O | 45:DT:29:ARG:CD | 2.58 | 0.52 |
| 45:DT:104:ASN:O | 45:DT:105:LEU:HG | 2.08 | 0.52 |
| 49:DX:72:LYS:HB2 | 49:DX:74:PRO:CD | 2.39 | 0.52 |
| 51:DZ:121:HIS:CE1 | 51:DZ:169:GLU:HG2 | 2.44 | 0.52 |
| 1:AA:1052:U:H2' | 1:AA:1055:A:OP1 | 2.10 | 0.52 |
| 1:AA:1321:C:H5' | 1:AA:1322:C:C5' | 2.39 | 0.52 |
| 2:AB:204:ASN:HD22 | 2:AB:205:ASP:N | 2.07 | 0.52 |
| 4:AD:109:GLY:O | 4:AD:111:ALA:N | 2.42 | 0.52 |
| 8:AH:86:ILE:HB | 8:AH:133:LEU:HD22 | 1.92 | 0.52 |
| 12:AL:83:VAL:CG1 | 12:AL:100:ILE:HG23 | 2.39 | 0.52 |
| 20:AT:93:GLU:HG2 | 20:AT:93:GLU:O | 2.09 | 0.52 |
| 22:B0:74:ARG:NH2 | 32:BB:13:A:H8 | 2.06 | 0.52 |
| 23:B1:19:GLN:CG | 23:B1:44:PRO:HG3 | 2.37 | 0.52 |
| 23:B1:76:ARG:HB3 | 23:B1:78:LYS:HE3 | 1.90 | 0.52 |
| 24:B2:30:ARG:H | 24:B2:30:ARG:CD | 2.10 | 0.52 |
| 25:B3:8:LEU:HD13 | 25:B3:31:LEU:CD2 | 2.29 | 0.52 |
| 28:B6:10:LEU:N | 28:B6:10:LEU:CD2 | 2.69 | 0.52 |
| 31:BA:83:G:H22 | 31:BA:102:G:HO2' | 1.50 | 0.52 |
| 31:BA:775:G:C4 | 31:BA:794:G:C8 | 2.97 | 0.52 |
| 31:BA:1038:C:OP2 | 31:BA:1038:C:H6 | 1.92 | 0.52 |
| 31:BA:2000:G:OP2 | 43:BR:3:HIS:CE1 | 2.63 | 0.52 |
| 31:BA:2662:A:H4' | 31:BA:2663:G:O4' | 2.10 | 0.52 |
| 31:BA:2688:U:C5 | 31:BA:2720:U:OP2 | 2.63 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2689:U:C4' | 31:BA:2690:C:H5' | 2.39 | 0.52 |
| 35:BF:53:THR:HG23 | 35:BF:55:GLY:N | 2.24 | 0.52 |
| 41:BP:38:GLN:CG | 41:BP:39:LYS:H | 2.18 | 0.52 |
| 41:BP:85:LEU:H | 41:BP:85:LEU:CD2 | 2.22 | 0.52 |
| 43:BR:41:ALA:O | 43:BR:44:LEU:N | 2.43 | 0.52 |
| 43:BR:84:ALA:HB3 | 43:BR:85:PRO:HD3 | 1.92 | 0.52 |
| 45:BT:128:GLU:OE1 | 45:BT:129:ARG:N | 2.42 | 0.52 |
| 49:BX:59:VAL:HG22 | 49:BX:74:PRO:O | 2.09 | 0.52 |
| 51:BZ:121:HIS:CE1 | 51:BZ:169:GLU:HG2 | 2.44 | 0.52 |
| 1:CA:366:C:O2' | 1:CA:367:U:H5" | 2.10 | 0.52 |
| 1:CA:685:G:C2 | 1:CA:686:U:C4 | 2.98 | 0.52 |
| 1:CA:892:A:C5 | 1:CA:893:C:C4 | 2.97 | 0.52 |
| 1:CA:913:A:H4' | 1:CA:914:A:O5' | 2.10 | 0.52 |
| 1:CA:948:C:C5 | 13:CM:106:ASN:ND2 | 2.77 | 0.52 |
| 1:CA:1084:G:OP1 | 1:CA:1086:U:C4 | 2.63 | 0.52 |
| 1:CA:1274:G:H22 | 1:CA:1275:A:H62 | 1.52 | 0.52 |
| 1:CA:1347:G:C8 | 9:CI:107:ARG:HB3 | 2.45 | 0.52 |
| 3:CC:157:ILE:CD1 | 3:CC:166:GLU:HB2 | 2.39 | 0.52 |
| 5:CE:101:ILE:HA | 5:CE:107:ARG:NH2 | 2.25 | 0.52 |
| 8:CH:109:ILE:CG2 | 8:CH:137:VAL:HB | 2.40 | 0.52 |
| 9:CI:18:PHE:HB3 | 9:CI:20:ARG:NH1 | 2.25 | 0.52 |
| 11:CK:110:ASP:O | 18:CR:84:LYS:HD2 | 2.10 | 0.52 |
| 17:CQ:33:GLY:O | 17:CQ:34:LYS:C | 2.53 | 0.52 |
| 20:CT:89:ARG:HD2 | 20:CT:104:LEU:HD11 | 1.92 | 0.52 |
| 23:D1:10:LYS:HD2 | 23:D1:14:VAL:HA | 1.92 | 0.52 |
| 25:D3:18:ASP:HB2 | 25:D3:49:LYS:HE3 | 1.92 | 0.52 |
| 30:D8:4:MET:O | 30:D8:62:LEU:CD1 | 2.58 | 0.52 |
| 31:DA:320:A:OP2 | 35:DF:137:LYS:HE3 | 2.09 | 0.52 |
| 31:DA:330:A:HO2' | 31:DA:331:A:H8 | 1.58 | 0.52 |
| 31:DA:754:C:H2' | 31:DA:755:C:C6 | 2.45 | 0.52 |
| 31:DA:784:A:H5' | 31:DA:785:G:OP1 | 2.08 | 0.52 |
| 31:DA:878:A:N6 | 31:DA:900:A:C8 | 2.77 | 0.52 |
| 31:DA:1108:U:H2' | 31:DA:1109:C:C5' | 2.39 | 0.52 |
| 31:DA:1224:C:O2' | 47:DV:87:HIS:N | 2.43 | 0.52 |
| 31:DA:2312:U:H4' | 36:DG:71:THR:HG21 | 1.91 | 0.52 |
| 31:DA:2515:C:O2' | 31:DA:2516:G:H5' | 2.09 | 0.52 |
| 31:DA:2646:C:H2' | 31:DA:2647:U:O4' | 2.10 | 0.52 |
| 31:DA:2884:U:C5 | 31:DA:2885:C:C6 | 2.98 | 0.52 |
| 33:DD:24:ILE:HD11 | 33:DD:84:TYR:N | 2.25 | 0.52 |
| 37:DH:144:VAL:O | 37:DH:144:VAL:HG12 | 2.10 | 0.52 |
| 39:DN:23:LEU:HD13 | 39:DN:98:VAL:HG12 | 1.91 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 44:DS:26:LEU:HA | 44:DS:39:ILE:HD13 | 1.91 | 0.52 |
| 46:DU:31:SER:O | 46:DU:33:ARG:N | 2.42 | 0.52 |
| 1:AA:114:U:H2' | 1:AA:115:G:H8 | 1.68 | 0.52 |
| 1:AA:132:C:O2' | 1:AA:133:U:H5' | 2.09 | 0.52 |
| 1:AA:577:G:C4 | 1:AA:816:A:C2 | 2.97 | 0.52 |
| 1:AA:658:G:C4 | 1:AA:659:U:C5 | 2.98 | 0.52 |
| 1:AA:706:A:H5" | 11:AK:22:HIS:CD2 | 2.45 | 0.52 |
| 1:AA:949:A:C2 | 1:AA:1233:G:N3 | 2.78 | 0.52 |
| 4:AD:117:ALA:O | 4:AD:121:VAL:HG22 | 2.09 | 0.52 |
| 5:AE:88:LYS:HB3 | 5:AE:123:LEU:HB2 | 1.92 | 0.52 |
| 6:AF:72:VAL:HG13 | 6:AF:73:ASN:N | 2.24 | 0.52 |
| 6:AF:99:ALA:HB3 | 18:AR:29:PHE:CE2 | 2.45 | 0.52 |
| 16:AP:26:ARG:HD2 | 16:AP:31:LYS:O | 2.10 | 0.52 |
| 20:AT:14:LYS:HB2 | 20:AT:17:ARG:NH2 | 2.25 | 0.52 |
| 25:B3:28:LEU:HA | 25:B3:33:GLN:OE1 | 2.09 | 0.52 |
| 31:BA:639:U:H2' | 31:BA:640:C:C6 | 2.44 | 0.52 |
| 31:BA:672:C:O2' | 31:BA:673:C:H5' | 2.10 | 0.52 |
| 31:BA:986:C:C2' | 31:BA:987:G:H5' | 2.40 | 0.52 |
| 31:BA:1720:U:H2' | 31:BA:1721:G:O4' | 2.10 | 0.52 |
| 31:BA:1803:A:O2' | 33:BD:259:THR:HG21 | 2.10 | 0.52 |
| 31:BA:1819:A:H5" | 33:BD:158:ALA:CB | 2.40 | 0.52 |
| 31:BA:2850:A:OP2 | 31:BA:2866:U:C5 | 2.63 | 0.52 |
| 32:BB:46:A:C6 | 32:BB:47:C:C4 | 2.98 | 0.52 |
| 33:BD:106:ILE:HD13 | 33:BD:106:ILE:C | 2.34 | 0.52 |
| 36:BG:52:ILE:HG22 | 36:BG:54:GLU:HG3 | 1.91 | 0.52 |
| 39:BN:28:THR:N | 39:BN:106:MET:CE | 2.73 | 0.52 |
| 42:BQ:21:THR:O | 51:BZ:78:LYS:HE2 | 2.10 | 0.52 |
| 44:BS:26:LEU:HA | 44:BS:39:ILE:HD13 | 1.91 | 0.52 |
| 45:BT:51:ARG:HG3 | 45:BT:98:LYS:HG3 | 1.91 | 0.52 |
| 47:BV:28:GLU:CG | 47:BV:29:PRO:HD3 | 2.40 | 0.52 |
| 49:BX:89:ILE:CG2 | 49:BX:89:ILE:O | 2.56 | 0.52 |
| 1:CA:323:U:OP1 | 20:CT:26:ASN:ND2 | 2.43 | 0.52 |
| 1:CA:814:A:N7 | 1:CA:816:A:C4 | 2.78 | 0.52 |
| 1:CA:1274:G:N2 | 1:CA:1275:A:N6 | 2.57 | 0.52 |
| 1:CA:1446:U:O2' | 1:CA:1447:A:H8 | 1.89 | 0.52 |
| 6:CF:8:ILE:HG22 | 6:CF:10:LEU:HD11 | 1.90 | 0.52 |
| 15:CO:3:ILE:HD13 | 15:CO:3:ILE:N | 2.24 | 0.52 |
| 16:CP:8:ARG:O | 16:CP:9:PHE:CG | 2.62 | 0.52 |
| 16:CP:39:TYR:C | 16:CP:39:TYR:CD1 | 2.86 | 0.52 |
| 20:CT:10:LEU:O | 20:CT:12:ALA:N | 2.35 | 0.52 |
| 24:D2:48:HIS:O | 24:D2:49:LYS:HD3 | 2.09 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 24:D2:51:ARG:O | 24:D2:52:ASP:HB3 | 2.09 | 0.52 |
| 24:D2:55:ARG:H | 24:D2:56:GLN:NE2 | 2.08 | 0.52 |
| 30:D8:29:LYS:O | 30:D8:30:ARG:C | 2.53 | 0.52 |
| 31:DA:643:A:C2 | 31:DA:644:A:C4 | 2.97 | 0.52 |
| 31:DA:828:U:H4' | 31:DA:831:G:N1 | 2.25 | 0.52 |
| 31:DA:995:C:O2 | 46:DU:57:PHE:CD2 | 2.63 | 0.52 |
| 31:DA:1441:G:H2' | 31:DA:1442:G:H8 | 1.73 | 0.52 |
| 31:DA:1784:A:C4' | 31:DA:1785:A:H5" | 2.40 | 0.52 |
| 31:DA:1986:A:O2' | 31:DA:1987:G:H5" | 2.09 | 0.52 |
| 31:DA:1992:G:O2' | 31:DA:1993:U:OP2 | 2.22 | 0.52 |
| 31:DA:2191:G:C2' | 31:DA:2192:G:O5' | 2.58 | 0.52 |
| 31:DA:2196:C:C2' | 31:DA:2197:U:H5' | 2.40 | 0.52 |
| 31:DA:2873:A:N3 | 43:DR:6:SER:HB2 | 2.25 | 0.52 |
| 38:DI:57:ARG:O | 38:DI:60:GLU:HB2 | 2.10 | 0.52 |
| 39:DN:78:TYR:HD1 | 39:DN:79:PRO:HD3 | 1.74 | 0.52 |
| 44:DS:74:ALA:O | 44:DS:77:ALA:HB3 | 2.08 | 0.52 |
| 47:DV:19:LYS:CG | 47:DV:96:ILE:HG22 | 2.33 | 0.52 |
| 48:DW:82:LEU:HG | 48:DW:84:ARG:NH2 | 2.25 | 0.52 |
| 50:DY:88:LYS:O | 50:DY:89:PHE:HB2 | 2.10 | 0.52 |
| 1:AA:175:C:C2 | 1:AA:176:C:C5 | 2.96 | 0.52 |
| 1:AA:1003:G:C2 | 1:AA:1004:A:H1' | 2.45 | 0.52 |
| 1:AA:1118:C:H1' | 1:AA:1179:A:C4 | 2.44 | 0.52 |
| 1:AA:1157:A:H4' | 1:AA:1158:C:O5' | 2.10 | 0.52 |
| 5:AE:71:LEU:N | 5:AE:71:LEU:HD23 | 2.24 | 0.52 |
| 8:AH:26:VAL:HG22 | 8:AH:27:PRO:O | 2.09 | 0.52 |
| 15:AO:66:LEU:O | 15:AO:69:TYR:HB3 | 2.09 | 0.52 |
| 23:B1:34:THR:CG2 | 31:BA:387:U:O3' | 2.57 | 0.52 |
| 23:B1:87:PRO:HB2 | 23:B1:91:LYS:HZ2 | 1.75 | 0.52 |
| 30:B8:43:GLN:O | 30:B8:44:LYS:HD2 | 2.09 | 0.52 |
| 31:BA:9:U:C4 | 31:BA:2629:A:C6 | 2.98 | 0.52 |
| 31:BA:767:U:O2' | 31:BA:768:G:H5' | 2.09 | 0.52 |
| 31:BA:814:C:C5 | 41:BP:27:HIS:NE2 | 2.78 | 0.52 |
| 31:BA:1250:G:H4' | 31:BA:1251:C:OP2 | 2.10 | 0.52 |
| 31:BA:2312:U:H4' | 36:BG:71:THR:HG21 | 1.92 | 0.52 |
| 35:BF:51:THR:OG1 | 35:BF:91:GLY:HA3 | 2.10 | 0.52 |
| 35:BF:53:THR:HG23 | 35:BF:55:GLY:H | 1.74 | 0.52 |
| 37:BH:163:TYR:N | 37:BH:163:TYR:CD1 | 2.78 | 0.52 |
| 39:BN:75:TYR:N | 39:BN:75:TYR:HD1 | 2.07 | 0.52 |
| 1:CA:32:A:H2' | 1:CA:33:A:C8 | 2.45 | 0.52 |
| 1:CA:194:C:H2' | 1:CA:195:A:H5" | 1.91 | 0.52 |
| 1:CA:472:A:C4' | 16:CP:82:GLN:HE22 | 2.21 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:683:G:C2 | 1:CA:708:C:N3 | 2.78 | 0.52 |
| 1:CA:939:G:H5" | 7:CG:102:ARG:NH1 | 2.25 | 0.52 |
| 1:CA:1184:G:H2' | 1:CA:1185:G:H8 | 1.74 | 0.52 |
| 2:CB:80:ILE:HD13 | 2:CB:208:ILE:HG23 | 1.92 | 0.52 |
| 8:CH:29:SER:HB3 | 8:CH:32:LYS:HD2 | 1.92 | 0.52 |
| 9:CI:55:ALA:CB | 9:CI:58:ARG:HD2 | 2.39 | 0.52 |
| 11:CK:21:ILE:HG13 | 11:CK:30:VAL:HG12 | 1.90 | 0.52 |
| 15:CO:67:LEU:CD2 | 15:CO:78:TYR:HE1 | 2.23 | 0.52 |
| 20:CT:93:GLU:HG2 | 20:CT:93:GLU:O | 2.09 | 0.52 |
| 22:D0:50:ASN:HD22 | 22:D0:83:PRO:HD3 | 1.75 | 0.52 |
| 23:D1:87:PRO:HB2 | 23:D1:91:LYS:NZ | 2.25 | 0.52 |
| 31:DA:452:G:C2 | 31:DA:458:G:C5 | 2.98 | 0.52 |
| 31:DA:869:G:C4 | 31:DA:870:A:C8 | 2.97 | 0.52 |
| 31:DA:992:C:C2' | 31:DA:993:G:O5' | 2.58 | 0.52 |
| 31:DA:1130:U:O2 | 31:DA:2025:C:H5" | 2.10 | 0.52 |
| 31:DA:1331:A:H2' | 31:DA:1333:C:C5 | 2.45 | 0.52 |
| 31:DA:2564:A:C2 | 31:DA:2647:U:H4' | 2.44 | 0.52 |
| 31:DA:2642:G:N2 | 31:DA:2773:C:C2 | 2.78 | 0.52 |
| 33:DD:43:ARG:HH11 | 33:DD:44:ASN:ND2 | 2.07 | 0.52 |
| 36:DG:29:TRP:C | 36:DG:31:VAL:N | 2.66 | 0.52 |
| 36:DG:76:SER:HB3 | 36:DG:84:LYS:H | 1.74 | 0.52 |
| 38:DI:12:LEU:O | 38:DI:12:LEU:HG | 2.08 | 0.52 |
| 39:DN:72:TYR:O | 39:DN:73:THR:C | 2.52 | 0.52 |
| 42:DQ:23:GLY:O | 42:DQ:99:PRO:O | 2.28 | 0.52 |
| 46:DU:27:LEU:HB3 | 46:DU:31:SER:HB3 | 1.92 | 0.52 |
| 49:DX:88:LYS:O | 49:DX:89:ILE:HB | 2.09 | 0.52 |
| 1:AA:832:C:O2' | 1:AA:833:U:P | 2.67 | 0.52 |
| 1:AA:1135:U:H4' | 1:AA:1136:U:H5 | 1.74 | 0.52 |
| 3:AC:18:TRP:CD1 | 14:AN:53:LEU:O | 2.63 | 0.52 |
| 4:AD:52:SER:O | 4:AD:54:TYR:N | 2.42 | 0.52 |
| 4:AD:79:PHE:CD1 | 4:AD:207:TYR:CD1 | 2.98 | 0.52 |
| 4:AD:131:ARG:HD3 | 4:AD:131:ARG:H | 1.75 | 0.52 |
| 5:AE:107:ARG:O | 5:AE:108:ALA:C | 2.53 | 0.52 |
| 8:AH:6:ILE:HD12 | 8:AH:6:ILE:N | 2.25 | 0.52 |
| 12:AL:20:LYS:HD3 | 12:AL:20:LYS:H | 1.75 | 0.52 |
| 15:AO:87:ILE:CG2 | 15:AO:88:ARG:N | 2.72 | 0.52 |
| 19:AS:44:MET:HA | 19:AS:44:MET:HE3 | 1.92 | 0.52 |
| 20:AT:29:LYS:O | 20:AT:33:ILE:HG12 | 2.09 | 0.52 |
| 20:AT:80:ARG:O | 20:AT:84:LEU:HB2 | 2.08 | 0.52 |
| 22:B0:70:GLN:O | 22:B0:77:ARG:HA | 2.10 | 0.52 |
| 23:B1:25:LYS:O | 23:B1:26:ARG:CB | 2.57 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 27:B5:2:ALA:HA | 31:BA:2015:A:C1' | 2.34 | 0.52 |
| 27:B5:40:LYS:HZ3 | 27:B5:46:CYS:HB3 | 1.75 | 0.52 |
| 30:B8:30:ARG:O | 30:B8:31:HIS:O | 2.28 | 0.52 |
| 31:BA:513:A:C2 | 31:BA:514:A:C5 | 2.98 | 0.52 |
| 31:BA:528:A:H2 | 31:BA:2043:C:H5' | 1.74 | 0.52 |
| 31:BA:860:U:C5 | 31:BA:917:A:N7 | 2.78 | 0.52 |
| 31:BA:918:A:H5" | 32:BB:98:G:O2' | 2.09 | 0.52 |
| 31:BA:1108:U:H2' | 31:BA:1109:C:C5' | 2.38 | 0.52 |
| 31:BA:1822:G:H8 | 31:BA:1822:G:H5' | 1.74 | 0.52 |
| 31:BA:2199:A:OP2 | 31:BA:2200:C:H5 | 1.93 | 0.52 |
| 31:BA:2850:A:OP2 | 31:BA:2866:U:H5 | 1.93 | 0.52 |
| 31:BA:2869:G:H2' | 31:BA:2870:C:O4' | 2.10 | 0.52 |
| 33:BD:76:PRO:O | 33:BD:98:VAL:HG23 | 2.09 | 0.52 |
| 36:BG:64:THR:HG23 | 36:BG:65:GLY:N | 2.20 | 0.52 |
| 40:BO:9:GLU:O | 40:BO:83:ALA:HA | 2.10 | 0.52 |
| 40:BO:50:GLY:C | 40:BO:52:VAL:H | 2.16 | 0.52 |
| 40:BO:60:ALA:CB | 40:BO:86:ILE:HA | 2.39 | 0.52 |
| 44:BS:24:LEU:O | 44:BS:85:VAL:HG12 | 2.10 | 0.52 |
| 50:BY:42:VAL:HB | 50:BY:65:ALA:HB3 | 1.92 | 0.52 |
| 1:CA:166:G:C4 | 1:CA:167:G:C8 | 2.98 | 0.52 |
| 1:CA:987:G:N2 | 1:CA:1219:U:C2 | 2.78 | 0.52 |
| 1:CA:1125:U:H2' | 1:CA:1126:U:OP2 | 2.10 | 0.52 |
| 1:CA:1201:A:C1' | 1:CA:1202:G:OP2 | 2.55 | 0.52 |
| 1:CA:1362:C:C2' | 1:CA:1363:C:H5" | 2.39 | 0.52 |
| 3:CC:113:ALA:C | 3:CC:115:LEU:N | 2.68 | 0.52 |
| 3:CC:156:ARG:NH2 | 3:CC:161:GLU:HA | 2.24 | 0.52 |
| 7:CG:15:ASP:O | 7:CG:19:GLY:HA2 | 2.10 | 0.52 |
| 23:D1:65:SER:H | 23:D1:67:ILE:HD12 | 1.74 | 0.52 |
| 31:DA:271(H):G:O6 | 31:DA:271(Q):G:C6 | 2.63 | 0.52 |
| 31:DA:393:C:C4 | 31:DA:394:A:N7 | 2.78 | 0.52 |
| 31:DA:910:A:N7 | 42:DQ:13:GLN:HB2 | 2.24 | 0.52 |
| 31:DA:923:C:H2' | 31:DA:924:C:H6 | 1.75 | 0.52 |
| 31:DA:1048:A:OP2 | 31:DA:1110:G:N1 | 2.43 | 0.52 |
| 31:DA:1833:U:O2' | 31:DA:1969:A:N1 | 2.40 | 0.52 |
| 31:DA:2052:G:C2 | 31:DA:2053:G:C8 | 2.98 | 0.52 |
| 31:DA:2475:C:H5" | 31:DA:2475:C:C6 | 2.42 | 0.52 |
| 31:DA:2526:G:C5 | 31:DA:2527:C:C5 | 2.97 | 0.52 |
| 31:DA:2602:A:H4' | 31:DA:2603:G:O5' | 2.10 | 0.52 |
| 31:DA:2632:A:H1' | 34:DE:61:ARG:HH12 | 1.74 | 0.52 |
| 32:DB:81:G:O6 | 32:DB:96:U:O2 | 2.28 | 0.52 |
| 33:DD:142:VAL:HG23 | 33:DD:192:THR:O | 2.09 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 34:DE:201:THR:HG22 | 34:DE:202:LYS:N | 2.25 | 0.52 |
| 48:DW:92:ARG:HG2 | 48:DW:92:ARG:NH1 | 2.25 | 0.52 |
| 49:DX:82:GLN:CD | 49:DX:83:VAL:H | 2.18 | 0.52 |
| 1:AA:63:C:H42 | 1:AA:104:G:H1 | 1.58 | 0.51 |
| 1:AA:373:A:H2' | 1:AA:374:A:H8 | 1.76 | 0.51 |
| 1:AA:397:A:N7 | 1:AA:548:G:C8 | 2.78 | 0.51 |
| 1:AA:914:A:C2' | 1:AA:915:A:H5' | 2.40 | 0.51 |
| 1:AA:1154:G:H2' | 1:AA:1155:G:H8 | 1.75 | 0.51 |
| 3:AC:116:VAL:HG21 | 3:AC:202:ILE:HD11 | 1.90 | 0.51 |
| 9:AI:18:PHE:HB2 | 9:AI:62:TYR:O | 2.09 | 0.51 |
| 25:B3:52:HIS:CD2 | 32:BB:83:G:C5' | 2.92 | 0.51 |
| 26:B4:23:GLU:O | 26:B4:24:THR:CB | 2.59 | 0.51 |
| 30:B8:27:THR:HA | 41:BP:62:LEU:CD1 | 2.40 | 0.51 |
| 31:BA:603:A:O2' | 31:BA:604:G:OP2 | 2.27 | 0.51 |
| 31:BA:670:A:H4' | 31:BA:671:C:OP1 | 2.10 | 0.51 |
| 31:BA:1204:A:N1 | 31:BA:1241:A:H2 | 2.08 | 0.51 |
| 31:BA:1341:U:N3 | 49:BX:77:LYS:HE2 | 2.24 | 0.51 |
| 31:BA:1366:A:H2' | 31:BA:1367:A:H5' | 1.93 | 0.51 |
| 31:BA:1378:A:O2' | 31:BA:1379:A:C5' | 2.52 | 0.51 |
| 31:BA:1451:C:N3 | 31:BA:1459:G:O6 | 2.43 | 0.51 |
| 31:BA:2273:A:C2' | 31:BA:2274:A:H5' | 2.40 | 0.51 |
| 31:BA:2484:G:C2 | 31:BA:2485:G:C8 | 2.98 | 0.51 |
| 31:BA:2599:G:OP2 | 33:BD:236:GLY:N | 2.43 | 0.51 |
| 31:BA:2840:C:O5' | 31:BA:2840:C:H6 | 1.93 | 0.51 |
| 33:BD:86:PRO:HG2 | 33:BD:87:ASN:HD21 | 1.75 | 0.51 |
| 33:BD:182:LEU:O | 33:BD:271:ILE:HD12 | 2.10 | 0.51 |
| 34:BE:51:PHE:CD1 | 34:BE:52:LEU:N | 2.78 | 0.51 |
| 36:BG:43:LEU:HD23 | 36:BG:88:ILE:HG22 | 1.92 | 0.51 |
| 37:BH:144:VAL:O | 37:BH:144:VAL:HG12 | 2.09 | 0.51 |
| 39:BN:13:TRP:O | 39:BN:135:PRO:HG2 | 2.09 | 0.51 |
| 39:BN:55:VAL:HG12 | 39:BN:126:PRO:HA | 1.92 | 0.51 |
| 42:BQ:134:ARG:O | 42:BQ:135:ASP:HB2 | 2.10 | 0.51 |
| 43:BR:10:LEU:HB3 | 43:BR:17:ARG:CZ | 2.39 | 0.51 |
| 44:BS:42:ASP:C | 44:BS:44:LYS:N | 2.69 | 0.51 |
| 45:BT:32:TYR:CB | 45:BT:81:PRO:HB2 | 2.40 | 0.51 |
| 45:BT:100:TYR:CD2 | 45:BT:103:ARG:NH2 | 2.71 | 0.51 |
| 45:BT:121:ILE:O | 45:BT:124:ASP:HB2 | 2.09 | 0.51 |
| 46:BU:75:ASN:HB2 | 46:BU:78:THR:HG1 | 1.70 | 0.51 |
| 48:BW:92:ARG:HG2 | 48:BW:92:ARG:NH1 | 2.25 | 0.51 |
| 51:BZ:161:VAL:HG12 | 51:BZ:161:VAL:O | 2.10 | 0.51 |
| 1:CA:1522:U:O2 | 1:CA:1523:G:C8 | 2.64 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 4:CD:79:PHE:CD1 | 4:CD:207:TYR:CD1 | 2.98 | 0.51 |
| 7:CG:92:SER:OG | 7:CG:93:PRO:HD2 | 2.09 | 0.51 |
| 8:CH:30:ARG:O | 8:CH:34:GLU:HG2 | 2.10 | 0.51 |
| 8:CH:86:ILE:HB | 8:CH:133:LEU:HD22 | 1.90 | 0.51 |
| 10:CJ:40:LEU:HD23 | 10:CJ:40:LEU:H | 1.75 | 0.51 |
| 23:D1:87:PRO:O | 23:D1:91:LYS:N | 2.31 | 0.51 |
| 24:D2:41:ILE:HG12 | 31:DA:94(A):G:N2 | 2.25 | 0.51 |
| 28:D6:42:TRP:CE3 | 28:D6:42:TRP:HA | 2.44 | 0.51 |
| 30:D8:39:LYS:O | 30:D8:39:LYS:HG2 | 2.09 | 0.51 |
| 31:DA:49:A:H4' | 31:DA:50:U:H5' | 1.91 | 0.51 |
| 31:DA:272(D):G:H1 | 31:DA:364:C:H42 | 1.58 | 0.51 |
| 31:DA:414:C:H2' | 31:DA:415:A:C8 | 2.46 | 0.51 |
| 31:DA:1494:A:H2' | 31:DA:1494:A:N3 | 2.24 | 0.51 |
| 31:DA:1568:G:H5' | 33:DD:60:ARG:HA | 1.92 | 0.51 |
| 31:DA:2000:G:OP2 | 43:DR:3:HIS:CE1 | 2.63 | 0.51 |
| 31:DA:2517:C:C6 | 31:DA:2542:A:C2 | 2.98 | 0.51 |
| 31:DA:2637:U:H2' | 31:DA:2638:G:C5' | 2.41 | 0.51 |
| 33:DD:85:ASP:HB2 | 33:DD:92:ILE:CG1 | 2.39 | 0.51 |
| 33:DD:142:VAL:HG21 | 33:DD:191:ALA:HB1 | 1.92 | 0.51 |
| 39:DN:82:LEU:HD12 | 39:DN:82:LEU:N | 2.24 | 0.51 |
| 40:DO:2:ILE:HG13 | 40:DO:8:LEU:HD11 | 1.93 | 0.51 |
| 41:DP:66:GLY:O | 41:DP:67:MET:C | 2.53 | 0.51 |
| 47:DV:25:LEU:CB | 47:DV:94:LEU:HD13 | 2.40 | 0.51 |
| 51:DZ:16:SER:HA | 51:DZ:19:ARG:HD2 | 1.91 | 0.51 |
| 1:AA:194:C:H2' | 1:AA:195:A:H5" | 1.91 | 0.51 |
| 1:AA:367:U:O2 | 1:AA:369:C:C6 | 2.62 | 0.51 |
| 1:AA:592:G:C2 | 1:AA:648:A:C2 | 2.99 | 0.51 |
| 1:AA:791:G:C6 | 1:AA:792:A:N7 | 2.78 | 0.51 |
| 1:AA:832:C:N4 | 1:AA:855:G:C6 | 2.79 | 0.51 |
| 1:AA:936:C:H2' | 1:AA:937:A:O4' | 2.10 | 0.51 |
| 8:AH:36:LEU:HD23 | 8:AH:39:LEU:HD23 | 1.92 | 0.51 |
| 31:BA:146:G:H8 | 31:BA:146:G:H5' | 1.74 | 0.51 |
| 31:BA:624:C:H2' | 31:BA:625:G:H5' | 1.91 | 0.51 |
| 31:BA:951:C:C2' | 31:BA:952:G:H5' | 2.40 | 0.51 |
| 31:BA:993:G:N3 | 47:BV:91:TYR:HE1 | 2.07 | 0.51 |
| 31:BA:1412:A:H2' | 31:BA:1413:G:C8 | 2.45 | 0.51 |
| 31:BA:1783:A:C2 | 31:BA:2587:A:C4 | 2.98 | 0.51 |
| 31:BA:1952:A:C5 | 40:BO:22:ILE:HD11 | 2.45 | 0.51 |
| 31:BA:2022:U:O2' | 31:BA:2617:C:H5' | 2.10 | 0.51 |
| 31:BA:2394:C:OP1 | 41:BP:63:PRO:CD | 2.40 | 0.51 |
| 33:BD:176:ARG:HG2 | 33:BD:176:ARG:NH1 | 2.24 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 34:BE:65:GLY:C | 34:BE:67:PHE:H | 2.18 | 0.51 |
| 34:BE:69:LYS:O | 34:BE:70:ALA:C | 2.52 | 0.51 |
| 39:BN:128:HIS:CD2 | 39:BN:128:HIS:O | 2.64 | 0.51 |
| 41:BP:16:ARG:O | 41:BP:18:ARG:N | 2.43 | 0.51 |
| 41:BP:148:LEU:N | 41:BP:148:LEU:HD13 | 2.24 | 0.51 |
| 41:BP:148:LEU:HD22 | 41:BP:148:LEU:O | 2.10 | 0.51 |
| 46:BU:10:ARG:HG2 | 46:BU:14:HIS:CE1 | 2.45 | 0.51 |
| 1:CA:723:U:H5" | 1:CA:724:G:OP2 | 2.10 | 0.51 |
| 1:CA:858:G:O6 | 1:CA:869:G:H3' | 2.10 | 0.51 |
| 1:CA:1037:C:H2' | 1:CA:1038:C:O4' | 2.10 | 0.51 |
| 1:CA:1290:G:N3 | 1:CA:1290:G:H2' | 2.25 | 0.51 |
| 7:CG:12:LEU:HD13 | 7:CG:24:THR:OG1 | 2.10 | 0.51 |
| 8:CH:26:VAL:HG22 | 8:CH:27:PRO:O | 2.10 | 0.51 |
| 18:CR:50:ILE:HD11 | 18:CR:70:ILE:CG2 | 2.36 | 0.51 |
| 20:CT:53:LEU:HD21 | 20:CT:92:LEU:HD11 | 1.92 | 0.51 |
| 27:D5:47:PRO:C | 27:D5:48:GLU:OE2 | 2.54 | 0.51 |
| 31:DA:92:A:O2' | 31:DA:93:G:H5' | 2.10 | 0.51 |
| 31:DA:271(Q):G:O2' | 31:DA:271(R):G:P | 2.69 | 0.51 |
| 31:DA:614(A):U:H4' | 31:DA:614(B):G:H5" | 1.91 | 0.51 |
| 31:DA:634:C:H2' | 31:DA:635:C:C6 | 2.45 | 0.51 |
| 31:DA:644:A:O2' | 31:DA:645:C:O2 | 2.27 | 0.51 |
| 31:DA:754:C:O4' | 31:DA:1618:A:H2 | 1.93 | 0.51 |
| 31:DA:1169:G:N2 | 31:DA:1181:C:C2 | 2.77 | 0.51 |
| 31:DA:1385:G:H4' | 31:DA:1386:C:OP1 | 2.09 | 0.51 |
| 31:DA:1489:U:O2' | 31:DA:1490:A:H5" | 2.10 | 0.51 |
| 31:DA:1685:C:C2' | 31:DA:1686:C:O5' | 2.59 | 0.51 |
| 31:DA:2473:U:N3 | 31:DA:2474:C:C6 | 2.78 | 0.51 |
| 32:DB:46:A:C5 | 32:DB:47:C:C4 | 2.98 | 0.51 |
| 33:DD:43:ARG:NH1 | 33:DD:44:ASN:ND2 | 2.59 | 0.51 |
| 33:DD:182:LEU:O | 33:DD:271:ILE:HD12 | 2.10 | 0.51 |
| 35:DF:63:LYS:HE2 | 35:DF:67:GLN:HB3 | 1.93 | 0.51 |
| 40:DO:107:ARG:NH1 | 40:DO:112:MET:HE1 | 2.25 | 0.51 |
| 41:DP:85:LEU:H | 41:DP:85:LEU:CD2 | 2.23 | 0.51 |
| 41:DP:89:ALA:O | 41:DP:90:ARG:C | 2.53 | 0.51 |
| 41:DP:148:LEU:HD22 | 41:DP:148:LEU:O | 2.09 | 0.51 |
| 44:DS:88:ASP:CG | 44:DS:89:ARG:N | 2.66 | 0.51 |
| 47:DV:54:GLY:C | 47:DV:56:SER:H | 2.16 | 0.51 |
| 51:DZ:67:LEU:H | 51:DZ:67:LEU:HD12 | 1.76 | 0.51 |
| 1:AA:180:U:H2' | 1:AA:181:G:H5' | 1.93 | 0.51 |
| 1:AA:250:A:C4' | 1:AA:251:G:O5' | 2.51 | 0.51 |
| 1:AA:719:C:C5 | 1:AA:720:C:C4 | 2.97 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:872:A:C2 | 1:AA:874:G:C6 | 2.98 | 0.51 |
| 1:AA:1128:C:H5' | 9:AI:16:ARG:CZ | 2.41 | 0.51 |
| 1:AA:1256:A:O3' | 1:AA:1257:U:H4' | 2.09 | 0.51 |
| 1:AA:1406:U:H2' | 1:AA:1407:C:O4' | 2.11 | 0.51 |
| 2:AB:97:TRP:CZ3 | 2:AB:173:ALA:HA | 2.45 | 0.51 |
| 7:AG:75:VAL:HG21 | 7:AG:144:MET:HB3 | 1.91 | 0.51 |
| 10:AJ:49:VAL:CG1 | 14:AN:41:ARG:HB2 | 2.39 | 0.51 |
| 13:AM:61:GLU:HA | 13:AM:66:LEU:HD11 | 1.93 | 0.51 |
| 16:AP:21:VAL:HG22 | 16:AP:34:GLU:O | 2.10 | 0.51 |
| 20:AT:24:LEU:HD13 | 20:AT:24:LEU:C | 2.34 | 0.51 |
| 24:B2:22:GLU:O | 24:B2:25:VAL:HG12 | 2.10 | 0.51 |
| 28:B6:11:LEU:O | 28:B6:23:THR:HA | 2.10 | 0.51 |
| 31:BA:143:G:H2' | 31:BA:143(A):C:C6 | 2.46 | 0.51 |
| 31:BA:185:U:H4' | 31:BA:218:A:H4' | 1.92 | 0.51 |
| 31:BA:271(Q):G:O2' | 31:BA:271(R):G:OP2 | 2.22 | 0.51 |
| 31:BA:992:C:H2' | 31:BA:993:G:O5' | 2.10 | 0.51 |
| 31:BA:1173:G:H5' | 31:BA:1174:A:OP2 | 2.10 | 0.51 |
| 31:BA:1509(A):A:H2' | 31:BA:1509(B):A:C8 | 2.45 | 0.51 |
| 31:BA:2206:G:H21 | 31:BA:2207:G:C4' | 2.23 | 0.51 |
| 31:BA:2473:U:C2 | 31:BA:2474:C:C6 | 2.99 | 0.51 |
| 31:BA:2753:A:O2' | 31:BA:2754:U:P | 2.69 | 0.51 |
| 32:BB:86:G:H1 | 32:BB:91:C:N4 | 2.08 | 0.51 |
| 33:BD:14:ARG:O | 33:BD:15:PHE:CD2 | 2.64 | 0.51 |
| 33:BD:233:HIS:CD2 | 33:BD:233:HIS:N | 2.76 | 0.51 |
| 34:BE:76:ARG:O | 34:BE:77:ILE:HG22 | 2.11 | 0.51 |
| 37:BH:89:ILE:O | 37:BH:90:LYS:CB | 2.57 | 0.51 |
| 38:BI:133:HIS:CB | 38:BI:134:PRO:CD | 2.76 | 0.51 |
| 39:BN:30:ILE:O | 39:BN:34:LEU:HD22 | 2.10 | 0.51 |
| 41:BP:70:GLN:HG3 | 41:BP:71:VAL:H | 1.76 | 0.51 |
| 43:BR:9:LYS:O | 43:BR:10:LEU:HD23 | 2.10 | 0.51 |
| 46:BU:64:ARG:NH2 | 46:BU:64:ARG:CA | 2.59 | 0.51 |
| 46:BU:83:LEU:CG | 46:BU:88:ILE:HG12 | 2.40 | 0.51 |
| 1:CA:949:A:C2 | 1:CA:1233:G:N3 | 2.79 | 0.51 |
| 1:CA:1052:U:H2' | 1:CA:1055:A:OP1 | 2.10 | 0.51 |
| 1:CA:1207:G:H2' | 1:CA:1208:C:C6 | 2.45 | 0.51 |
| 2:CB:97:TRP:HH2 | 2:CB:176:GLU:CG | 2.23 | 0.51 |
| 20:CT:78:ALA:HA | 20:CT:81:LYS:HD3 | 1.91 | 0.51 |
| 23:D1:15:ALA:HA | 23:D1:46:LEU:HD21 | 1.91 | 0.51 |
| 24:D2:12:GLU:CD | 24:D2:12:GLU:C | 2.78 | 0.51 |
| 24:D2:26:ARG:O | 24:D2:30:ARG:HD3 | 2.11 | 0.51 |
| 24:D2:50:ILE:O | 24:D2:51:ARG:HB3 | 2.11 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:573:G:C6 | 31:DA:2030:A:H3' | 2.46 | 0.51 |
| 31:DA:675:A:C6 | 31:DA:676:A:C6 | 2.98 | 0.51 |
| 31:DA:810:U:O2 | 41:DP:33:ARG:HD3 | 2.09 | 0.51 |
| 31:DA:947:G:N2 | 31:DA:971:C:C2 | 2.78 | 0.51 |
| 31:DA:953:A:C2 | 31:DA:954:G:C8 | 2.99 | 0.51 |
| 31:DA:993:G:N2 | 47:DV:91:TYR:OH | 2.43 | 0.51 |
| 31:DA:1658:C:H2' | 31:DA:1659:U:C6 | 2.45 | 0.51 |
| 31:DA:2308:G:C2 | 31:DA:2309:A:C6 | 2.98 | 0.51 |
| 31:DA:2808:U:O2' | 31:DA:2809:A:H5' | 2.11 | 0.51 |
| 33:DD:35:LYS:NZ | 33:DD:104:TYR:CD1 | 2.73 | 0.51 |
| 44:DS:12:PHE:CD1 | 44:DS:12:PHE:O | 2.64 | 0.51 |
| 47:DV:56:SER:O | 47:DV:57:VAL:HB | 2.10 | 0.51 |
| 1:AA:11:G:C5 | 1:AA:12:U:C5 | 2.98 | 0.51 |
| 1:AA:279:A:OP2 | 17:AQ:95:TYR:OH | 2.25 | 0.51 |
| 1:AA:617:G:N1 | 1:AA:618:C:C5 | 2.79 | 0.51 |
| 1:AA:1279:A:H2 | 10:AJ:43:ARG:NH1 | 2.08 | 0.51 |
| 1:AA:1288:A:H2 | 1:AA:1352:C:O2 | 1.94 | 0.51 |
| 2:AB:28:PHE:CD1 | 2:AB:190:THR:HA | 2.46 | 0.51 |
| 2:AB:90:MET:HE2 | 2:AB:90:MET:HA | 1.92 | 0.51 |
| 10:AJ:39:PRO:HB3 | 10:AJ:70:ARG:NH1 | 2.24 | 0.51 |
| 12:AL:86:ARG:HB2 | 12:AL:101:VAL:CG2 | 2.41 | 0.51 |
| 29:B7:43:THR:O | 29:B7:44:PRO:C | 2.52 | 0.51 |
| 31:BA:848:G:N9 | 31:BA:933:A:H8 | 2.07 | 0.51 |
| 31:BA:1598:C:H2' | 31:BA:1598:C:O2 | 2.10 | 0.51 |
| 31:BA:2801:A:H4' | 31:BA:2801(A):A:H5' | 1.92 | 0.51 |
| 31:BA:2859:G:C8 | 31:BA:2859:G:C3' | 2.93 | 0.51 |
| 32:BB:15:A:H2' | 32:BB:16:G:OP1 | 2.09 | 0.51 |
| 39:BN:78:TYR:HD1 | 39:BN:79:PRO:N | 2.09 | 0.51 |
| 44:BS:53:SER:O | 44:BS:56:LEU:N | 2.37 | 0.51 |
| 49:BX:52:VAL:HB | 49:BX:80:ILE:CG2 | 2.41 | 0.51 |
| 50:BY:28:LYS:O | 50:BY:29:GLU:C | 2.53 | 0.51 |
| 50:BY:52:SER:O | 50:BY:54:LYS:N | 2.43 | 0.51 |
| 50:BY:88:LYS:HZ3 | 50:BY:93:GLY:HA3 | 1.75 | 0.51 |
| 1:CA:20:U:H2' | 1:CA:21:G:O4' | 2.10 | 0.51 |
| 1:CA:401:C:OP1 | 4:CD:73:ARG:HD2 | 2.09 | 0.51 |
| 1:CA:577:G:C2 | 1:CA:578:C:C6 | 2.98 | 0.51 |
| 1:CA:611:A:N6 | 1:CA:629:G:H1 | 2.09 | 0.51 |
| 1:CA:938:A:N6 | 1:CA:939:G:C6 | 2.78 | 0.51 |
| 1:CA:1125:U:O4 | 10:CJ:5:ARG:HD2 | 2.10 | 0.51 |
| 1:CA:1321:C:C4 | 1:CA:1322:C:C4 | 2.99 | 0.51 |
| 1:CA:1350:A:OP1 | 9:CI:121:ARG:HG3 | 2.10 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:1490:C:O2' | 1:CA:1491:G:H5' | 2.11 | 0.51 |
| 2:CB:87:ARG:NE | 2:CB:233:SER:HB3 | 2.23 | 0.51 |
| 3:CC:7:PRO:O | 3:CC:11:ARG:HG2 | 2.11 | 0.51 |
| 10:CJ:48:THR:HA | 10:CJ:62:HIS:HB3 | 1.93 | 0.51 |
| 23:D1:64:ALA:O | 23:D1:65:SER:HB3 | 2.09 | 0.51 |
| 24:D2:44:LEU:HD12 | 24:D2:44:LEU:C | 2.34 | 0.51 |
| 31:DA:102:G:C8 | 31:DA:102:G:C5' | 2.82 | 0.51 |
| 31:DA:610:G:H2' | 31:DA:611:C:H6 | 1.74 | 0.51 |
| 31:DA:614:U:O2 | 31:DA:614:U:O4' | 2.26 | 0.51 |
| 31:DA:1001:A:H2' | 31:DA:1002:G:O4' | 2.10 | 0.51 |
| 31:DA:1497:U:C5' | 31:DA:1498:C:C5 | 2.90 | 0.51 |
| 31:DA:1790:C:C5' | 31:DA:1791:A:OP1 | 2.50 | 0.51 |
| 31:DA:2360:A:O2' | 31:DA:2361:A:C5' | 2.59 | 0.51 |
| 31:DA:2464:C:O2' | 31:DA:2465:C:OP2 | 2.28 | 0.51 |
| 32:DB:15:A:C2' | 32:DB:16:G:OP1 | 2.57 | 0.51 |
| 33:DD:16:MET:HE3 | 33:DD:206:LEU:O | 2.11 | 0.51 |
| 33:DD:53:PHE:CD1 | 33:DD:219:PRO:O | 2.64 | 0.51 |
| 33:DD:70:TRP:CZ3 | 33:DD:146:GLU:OE2 | 2.57 | 0.51 |
| 36:DG:115:ARG:NH1 | 36:DG:136:ARG:HG3 | 2.25 | 0.51 |
| 39:DN:99:LEU:O | 39:DN:103:VAL:HG23 | 2.11 | 0.51 |
| 43:DR:60:LEU:HD23 | 43:DR:61:HIS:N | 2.25 | 0.51 |
| 46:DU:10:ARG:O | 46:DU:11:ARG:C | 2.53 | 0.51 |
| 46:DU:33:ARG:O | 46:DU:36:ARG:N | 2.44 | 0.51 |
| 46:DU:92:ARG:HB3 | 46:DU:95:LEU:HD12 | 1.91 | 0.51 |
| 49:DX:21:PHE:N | 49:DX:21:PHE:CD1 | 2.79 | 0.51 |
| 49:DX:40:LYS:O | 49:DX:42:ALA:N | 2.35 | 0.51 |
| 50:DY:54:LYS:O | 50:DY:55:TYR:O | 2.28 | 0.51 |
| 50:DY:76:CYS:O | 50:DY:99:CYS:SG | 2.68 | 0.51 |
| 51:DZ:12:GLY:O | 51:DZ:13:GLU:HG3 | 2.10 | 0.51 |
| 1:AA:685:G:N2 | 1:AA:686:U:C4 | 2.78 | 0.51 |
| 1:AA:1206:G:O4' | 3:AC:194:GLY:HA2 | 2.10 | 0.51 |
| 1:AA:1277:C:O2' | 1:AA:1279:A:H1' | 2.10 | 0.51 |
| 1:AA:1290:G:H2' | 1:AA:1290:G:N3 | 2.25 | 0.51 |
| 2:AB:219:VAL:O | 2:AB:222:ILE:HB | 2.10 | 0.51 |
| 4:AD:203:VAL:HG12 | 4:AD:204:ILE:N | 2.24 | 0.51 |
| 13:AM:19:LEU:HB3 | 13:AM:25:ILE:HG21 | 1.93 | 0.51 |
| 15:AO:63:ARG:O | 15:AO:67:LEU:HB2 | 2.10 | 0.51 |
| 20:AT:67:ALA:HB2 | 20:AT:77:ALA:HB2 | 1.93 | 0.51 |
| 22:B0:19:LYS:HZ2 | 31:BA:2262:U:P | 2.33 | 0.51 |
| 27:B5:51:TYR:HD2 | 27:B5:52:TYR:CZ | 2.25 | 0.51 |
| 30:B8:4:MET:O | 30:B8:62:LEU:CD1 | 2.59 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:49:A:H3' | 31:BA:50:U:H5' | 1.93 | 0.51 |
| 31:BA:272(G):C:N4 | 31:BA:363(C):G:H1 | 2.07 | 0.51 |
| 31:BA:445:C:C2' | 31:BA:446:G:H5' | 2.40 | 0.51 |
| 31:BA:460:A:C2 | 31:BA:470:A:C5 | 2.98 | 0.51 |
| 31:BA:580:C:H2' | 31:BA:581:C:C6 | 2.45 | 0.51 |
| 31:BA:848:G:N3 | 31:BA:933:A:H1' | 2.25 | 0.51 |
| 31:BA:897:C:O2' | 31:BA:899:A:N7 | 2.39 | 0.51 |
| 31:BA:960:A:H61 | 42:BQ:82:ARG:HH21 | 1.58 | 0.51 |
| 31:BA:1204:A:C2 | 31:BA:1241:A:N1 | 2.79 | 0.51 |
| 31:BA:1349:A:N6 | 31:BA:1598:C:N4 | 2.58 | 0.51 |
| 31:BA:1434:A:H61 | 31:BA:1558:A:H62 | 1.58 | 0.51 |
| 31:BA:1719:G:O2' | 31:BA:1720:U:H5' | 2.11 | 0.51 |
| 31:BA:1856:G:H2' | 31:BA:1857:G:H5' | 1.93 | 0.51 |
| 31:BA:2094:G:H5' | 38:BI:25:TYR:CD2 | 2.46 | 0.51 |
| 32:BB:35:U:O2' | 32:BB:36:C:H5' | 2.10 | 0.51 |
| 33:BD:4:LYS:HZ1 | 33:BD:20:ASP:HA | 1.75 | 0.51 |
| 33:BD:232:PRO:C | 33:BD:233:HIS:O | 2.52 | 0.51 |
| 33:BD:266:SER:O | 33:BD:267:SER:HB2 | 2.10 | 0.51 |
| 37:BH:41:MET:CE | 37:BH:55:PRO:HD3 | 2.37 | 0.51 |
| 38:BI:15:VAL:C | 38:BI:17:GLN:H | 2.19 | 0.51 |
| 46:BU:31:SER:O | 46:BU:33:ARG:N | 2.43 | 0.51 |
| 47:BV:54:GLY:C | 47:BV:56:SER:N | 2.69 | 0.51 |
| 47:BV:83:ARG:NH1 | 47:BV:83:ARG:HG2 | 2.24 | 0.51 |
| 1:CA:438:G:H4' | 4:CD:123:HIS:ND1 | 2.26 | 0.51 |
| 1:CA:818:G:HO2' | 1:CA:820:U:H6 | 1.55 | 0.51 |
| 1:CA:827:U:N3 | 1:CA:870:U:C4 | 2.79 | 0.51 |
| 2:CB:23:ARG:O | 2:CB:23:ARG:HG2 | 2.11 | 0.51 |
| 6:CF:30:LEU:O | 6:CF:35:ALA:N | 2.42 | 0.51 |
| 7:CG:72:ARG:O | 7:CG:73:MET:HG3 | 2.09 | 0.51 |
| 28:D6:42:TRP:HA | 28:D6:42:TRP:HE3 | 1.74 | 0.51 |
| 30:D8:62:LEU:N | 30:D8:63:PRO:HD2 | 2.26 | 0.51 |
| 31:DA:142:A:H5" | 31:DA:142(A):C:C5 | 2.45 | 0.51 |
| 31:DA:271(K):U:H3' | 31:DA:271(L):U:H5' | 1.92 | 0.51 |
| 31:DA:594:U:H2' | 31:DA:595:C:C6 | 2.45 | 0.51 |
| 31:DA:667:U:H2' | 31:DA:668:G:H5' | 1.92 | 0.51 |
| 31:DA:719:C:O5' | 31:DA:719:C:H6 | 1.93 | 0.51 |
| 31:DA:870:A:C2 | 31:DA:908:C:C2 | 2.99 | 0.51 |
| 31:DA:993:G:N3 | 47:DV:91:TYR:CE1 | 2.78 | 0.51 |
| 31:DA:1171:G:N7 | 31:DA:1173:G:H1' | 2.26 | 0.51 |
| 31:DA:1288:U:C2 | 31:DA:1327:C:C2 | 2.99 | 0.51 |
| 31:DA:1484:G:H22 | 31:DA:1505:C:H5 | 1.57 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1569:A:O2' | 33:DD:38:LYS:HE2 | 2.11 | 0.51 |
| 31:DA:1797:C:O2' | 33:DD:259:THR:HB | 2.09 | 0.51 |
| 31:DA:2294:C:OP1 | 44:DS:92:TYR:HE1 | 1.92 | 0.51 |
| 31:DA:2830:G:N3 | 31:DA:2883:A:H2 | 2.09 | 0.51 |
| 31:DA:2870:C:H5" | 43:DR:65:LEU:HD21 | 1.91 | 0.51 |
| 34:DE:114:ALA:HB3 | 34:DE:160:TYR:HB3 | 1.93 | 0.51 |
| 37:DH:89:ILE:O | 37:DH:90:LYS:CB | 2.58 | 0.51 |
| 37:DH:105:LEU:H | 37:DH:105:LEU:HD13 | 1.75 | 0.51 |
| 39:DN:75:TYR:N | 39:DN:75:TYR:CD1 | 2.78 | 0.51 |
| 45:DT:65:LYS:HG3 | 45:DT:66:VAL:N | 2.25 | 0.51 |
| 47:DV:62:LEU:HB3 | 47:DV:98:GLU:HA | 1.92 | 0.51 |
| 51:DZ:51:ALA:O | 51:DZ:52:SER:HB3 | 2.10 | 0.51 |
| 1:AA:630:G:H2' | 1:AA:630:G:N3 | 2.25 | 0.51 |
| 1:AA:750:G:C2 | 1:AA:751:U:C6 | 2.99 | 0.51 |
| 1:AA:967:C:H5" | 1:AA:968:A:OP2 | 2.11 | 0.51 |
| 1:AA:1530:G:OP1 | 1:AA:1530:G:H4' | 2.10 | 0.51 |
| 2:AB:15:VAL:C | 2:AB:16:HIS:CG | 2.89 | 0.51 |
| 4:AD:75:PHE:CE2 | 4:AD:93:PHE:HZ | 2.29 | 0.51 |
| 8:AH:10:LEU:HD23 | 8:AH:10:LEU:H | 1.73 | 0.51 |
| 15:AO:25:THR:O | 15:AO:26:GLU:C | 2.54 | 0.51 |
| 19:AS:51:VAL:HG21 | 19:AS:71:LEU:HB3 | 1.93 | 0.51 |
| 20:AT:32:ALA:O | 20:AT:36:LEU:HD23 | 2.10 | 0.51 |
| 20:AT:74:LYS:C | 20:AT:76:ALA:H | 2.19 | 0.51 |
| 31:BA:139(A):G:H2' | 31:BA:140:G:O5' | 2.11 | 0.51 |
| 31:BA:613:G:N2 | 31:BA:614(C):A:O2' | 2.44 | 0.51 |
| 31:BA:1048:A:OP2 | 31:BA:1110:G:N1 | 2.44 | 0.51 |
| 31:BA:1276:A:C2 | 31:BA:1277:G:C8 | 2.98 | 0.51 |
| 31:BA:1857:G:C6 | 31:BA:1858:G:N1 | 2.78 | 0.51 |
| 31:BA:2400:G:C5 | 31:BA:2401:U:C5 | 2.98 | 0.51 |
| 31:BA:2803:C:H2' | 31:BA:2804:C:O4' | 2.11 | 0.51 |
| 31:BA:2808:U:C2' | 31:BA:2809:A:H5' | 2.40 | 0.51 |
| 33:BD:85:ASP:HB2 | 33:BD:92:ILE:CG1 | 2.41 | 0.51 |
| 35:BF:203:GLN:O | 35:BF:206:ILE:C | 2.54 | 0.51 |
| 37:BH:116:GLU:HG3 | 37:BH:117:PRO:HD2 | 1.92 | 0.51 |
| 37:BH:153:LYS:HG2 | 37:BH:154:PRO:N | 2.25 | 0.51 |
| 38:BI:38:LEU:C | 38:BI:40:THR:H | 2.17 | 0.51 |
| 39:BN:78:TYR:CD1 | 39:BN:79:PRO:CB | 2.94 | 0.51 |
| 39:BN:87:LEU:HD22 | 39:BN:87:LEU:O | 2.10 | 0.51 |
| 42:BQ:70:PRO:HA | 42:BQ:95:ALA:HB2 | 1.93 | 0.51 |
| 47:BV:18:LEU:HD22 | 47:BV:19:LYS:N | 2.25 | 0.51 |
| 1:CA:66:G:C2 | 1:CA:67:C:C6 | 2.98 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:630:G:N3 | 1:CA:630:G:H2' | 2.25 | 0.51 |
| 1:CA:1202:G:C2' | 1:CA:1203:C:H5' | 2.41 | 0.51 |
| 1:CA:1240:U:P | 7:CG:116:ALA:HB2 | 2.51 | 0.51 |
| 1:CA:1305:G:H5' | 21:CU:4:GLY:HA3 | 1.92 | 0.51 |
| 2:CB:97:TRP:CH2 | 2:CB:176:GLU:OE2 | 2.64 | 0.51 |
| 2:CB:130:ARG:HE | 2:CB:130:ARG:HA | 1.76 | 0.51 |
| 3:CC:206:GLU:HG2 | 3:CC:207:VAL:H | 1.76 | 0.51 |
| 7:CG:47:CYS:O | 7:CG:50:ILE:HB | 2.10 | 0.51 |
| 10:CJ:75:ILE:HG13 | 10:CJ:76:ASN:H | 1.75 | 0.51 |
| 12:CL:24:VAL:HG13 | 12:CL:98:TYR:HE2 | 1.75 | 0.51 |
| 27:D5:9:LYS:HA | 31:DA:2017:U:O2' | 2.11 | 0.51 |
| 31:DA:183:C:C2' | 31:DA:184:C:H5' | 2.41 | 0.51 |
| 31:DA:272(J):C:H42 | 31:DA:363(A):A:N6 | 2.08 | 0.51 |
| 31:DA:287:C:H42 | 31:DA:354:G:H1 | 1.58 | 0.51 |
| 31:DA:470:A:C2 | 31:DA:471:A:C4 | 2.98 | 0.51 |
| 31:DA:513:A:C2 | 31:DA:514:A:C5 | 2.99 | 0.51 |
| 31:DA:926:A:H5" | 31:DA:926:A:C8 | 2.45 | 0.51 |
| 31:DA:1366:A:H2' | 31:DA:1367:A:H5' | 1.92 | 0.51 |
| 31:DA:1441:G:H2' | 31:DA:1442:G:C8 | 2.46 | 0.51 |
| 31:DA:1822:G:C5' | 31:DA:1822:G:C8 | 2.94 | 0.51 |
| 31:DA:2187:G:N7 | 31:DA:2188:C:C4 | 2.79 | 0.51 |
| 31:DA:2632:A:O2' | 34:DE:61:ARG:NH2 | 2.44 | 0.51 |
| 37:DH:138:LYS:O | 37:DH:140:LYS:N | 2.44 | 0.51 |
| 38:DI:86:THR:HG23 | 38:DI:122:GLU:OE2 | 2.11 | 0.51 |
| 38:DI:120:ILE:HG22 | 38:DI:121:LYS:N | 2.26 | 0.51 |
| 41:DP:6:LEU:HD12 | 41:DP:8:PRO:HB2 | 1.92 | 0.51 |
| 42:DQ:69:PHE:CG | 42:DQ:70:PRO:HD2 | 2.46 | 0.51 |
| 43:DR:21:TYR:CZ | 43:DR:43:GLU:HG2 | 2.45 | 0.51 |
| 46:DU:91:ASP:O | 46:DU:95:LEU:HB2 | 2.11 | 0.51 |
| 47:DV:38:LEU:HG | 47:DV:39:LEU:H | 1.76 | 0.51 |
| 48:DW:92:ARG:O | 48:DW:93:ALA:HB3 | 2.09 | 0.51 |
| 49:DX:21:PHE:N | 49:DX:21:PHE:HD1 | 2.08 | 0.51 |
| 49:DX:83:VAL:O | 49:DX:84:ALA:CB | 2.58 | 0.51 |
| 1:AA:130:A:C8 | 17:AQ:63:ARG:HG3 | 2.46 | 0.51 |
| 1:AA:193:C:H2' | 1:AA:194:C:C6 | 2.43 | 0.51 |
| 1:AA:577:G:C2 | 1:AA:578:C:C6 | 2.99 | 0.51 |
| 1:AA:1233:G:H2' | 1:AA:1234:C:C6 | 2.46 | 0.51 |
| 1:AA:1271:G:H5' | 1:AA:1314:C:H5' | 1.92 | 0.51 |
| 1:AA:1418:A:H2 | 31:BA:1948:G:N3 | 2.08 | 0.51 |
| 3:AC:11:ARG:O | 3:AC:13:GLY:N | 2.43 | 0.51 |
| 5:AE:7:GLU:HG2 | 5:AE:112:LEU:HD22 | 1.92 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 5:AE:105:VAL:HG21 | 5:AE:128:PRO:HA | 1.92 | 0.51 |
| 6:AF:7:ASN:O | 6:AF:88:VAL:HA | 2.11 | 0.51 |
| 8:AH:5:PRO:O | 8:AH:8:ASP:HB3 | 2.11 | 0.51 |
| 10:AJ:26:ALA:HB1 | 10:AJ:29:ARG:NH2 | 2.24 | 0.51 |
| 10:AJ:44:VAL:HG22 | 10:AJ:66:ARG:HG2 | 1.92 | 0.51 |
| 11:AK:62:GLN:O | 11:AK:64:ALA:N | 2.43 | 0.51 |
| 11:AK:80:VAL:O | 11:AK:106:LYS:HB3 | 2.11 | 0.51 |
| 11:AK:126:ARG:O | 11:AK:127:LYS:C | 2.53 | 0.51 |
| 12:AL:5:PRO:HB2 | 12:AL:10:LEU:HD21 | 1.93 | 0.51 |
| 31:BA:1022:G:C5 | 31:BA:1140:C:N4 | 2.79 | 0.51 |
| 31:BA:1890:A:C2' | 31:BA:1891:G:H5' | 2.40 | 0.51 |
| 31:BA:2287:A:C5 | 31:BA:2289:G:C5 | 2.99 | 0.51 |
| 31:BA:2294:C:OP1 | 44:BS:92:TYR:HE1 | 1.93 | 0.51 |
| 31:BA:2536:G:C6 | 31:BA:2537:U:C4 | 2.98 | 0.51 |
| 32:BB:10:C:C2' | 32:BB:11:C:H5' | 2.41 | 0.51 |
| 32:BB:40:U:C2' | 32:BB:41:U:OP1 | 2.58 | 0.51 |
| 32:BB:41:U:C2' | 32:BB:42:C:OP1 | 2.59 | 0.51 |
| 35:BF:24:LEU:CB | 35:BF:25:PRO:HD2 | 2.35 | 0.51 |
| 38:BI:51:ILE:O | 38:BI:51:ILE:CG2 | 2.58 | 0.51 |
| 39:BN:30:ILE:HG22 | 39:BN:34:LEU:CD2 | 2.41 | 0.51 |
| 39:BN:62:VAL:CG2 | 39:BN:66:LYS:HG3 | 2.40 | 0.51 |
| 41:BP:149:GLU:O | 41:BP:149:GLU:HG3 | 2.10 | 0.51 |
| 45:BT:28:VAL:HG13 | 45:BT:46:GLU:HA | 1.92 | 0.51 |
| 47:BV:25:LEU:HD12 | 47:BV:94:LEU:HD22 | 1.93 | 0.51 |
| 47:BV:66:ARG:HD3 | 47:BV:67:GLY:N | 2.24 | 0.51 |
| 49:BX:73:ARG:O | 49:BX:74:PRO:O | 2.28 | 0.51 |
| 50:BY:46:LYS:O | 50:BY:47:LYS:NZ | 2.36 | 0.51 |
| 51:BZ:9:TYR:CE2 | 51:BZ:61:LEU:HD13 | 2.46 | 0.51 |
| 1:CA:63:C:O2' | 1:CA:380:G:H4' | 2.10 | 0.51 |
| 1:CA:355:C:C4 | 1:CA:356:A:N7 | 2.79 | 0.51 |
| 1:CA:1128:C:H5' | 9:CI:16:ARG:HH22 | 1.75 | 0.51 |
| 1:CA:1154:G:H2' | 1:CA:1155:G:H8 | 1.74 | 0.51 |
| 1:CA:1505:G:H4' | 1:CA:1506:U:C5' | 2.38 | 0.51 |
| 4:CD:31:CYS:SG | 4:CD:31:CYS:O | 2.69 | 0.51 |
| 11:CK:29:ILE:HB | 11:CK:44:SER:HB2 | 1.92 | 0.51 |
| 16:CP:16:HIS:O | 16:CP:17:TYR:O | 2.28 | 0.51 |
| 31:DA:386:G:H4' | 31:DA:387:U:OP2 | 2.11 | 0.51 |
| 31:DA:911:A:N9 | 42:DQ:9:TYR:OH | 2.43 | 0.51 |
| 31:DA:1173:G:H5' | 31:DA:1174:A:OP2 | 2.10 | 0.51 |
| 31:DA:1488:G:C2 | 31:DA:1489:U:O2 | 2.63 | 0.51 |
| 31:DA:1503:U:H6 | 31:DA:1503:U:H3' | 1.76 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1528:A:O2' | 31:DA:1528(A):A:C8 | 2.61 | 0.51 |
| 31:DA:1952:A:C6 | 31:DA:1953:A:N1 | 2.77 | 0.51 |
| 32:DB:6:C:C2 | 32:DB:116:G:N2 | 2.78 | 0.51 |
| 36:DG:43:LEU:HD23 | 36:DG:88:ILE:HG22 | 1.92 | 0.51 |
| 36:DG:71:THR:HB | 36:DG:89:GLY:HA3 | 1.92 | 0.51 |
| 39:DN:3:THR:O | 39:DN:4:TYR:CG | 2.64 | 0.51 |
| 45:DT:32:TYR:CD2 | 45:DT:81:PRO:O | 2.64 | 0.51 |
| 51:DZ:161:VAL:HG12 | 51:DZ:161:VAL:O | 2.09 | 0.51 |
| 1:AA:401:C:OP1 | 4:AD:73:ARG:HD2 | 2.11 | 0.51 |
| 1:AA:1287:A:C2 | 1:AA:1353:G:H1' | 2.45 | 0.51 |
| 1:AA:1388:C:H2' | 1:AA:1389:C:C6 | 2.45 | 0.51 |
| 8:AH:120:THR:H | 8:AH:123:GLU:HB2 | 1.75 | 0.51 |
| 14:AN:24:CYS:HB3 | 14:AN:27:CYS:O | 2.11 | 0.51 |
| 15:AO:74:ASP:OD2 | 15:AO:77:ARG:HG2 | 2.11 | 0.51 |
| 22:B0:48:GLY:HA3 | 22:B0:80:HIS:ND1 | 2.26 | 0.51 |
| 22:B0:55:ARG:HG3 | 31:BA:2365:G:OP1 | 2.11 | 0.51 |
| 24:B2:18:PRO:O | 24:B2:19:VAL:C | 2.53 | 0.51 |
| 31:BA:263:C:H2' | 31:BA:264:C:O4' | 2.11 | 0.51 |
| 31:BA:479:A:H4' | 31:BA:480:A:OP1 | 2.11 | 0.51 |
| 31:BA:1484:G:N2 | 31:BA:1505:C:C5 | 2.79 | 0.51 |
| 31:BA:2632:A:C2 | 34:BE:61:ARG:HD2 | 2.46 | 0.51 |
| 38:BI:82:ARG:O | 38:BI:89:TYR:HB2 | 2.10 | 0.51 |
| 38:BI:99:GLU:HG3 | 38:BI:103:ARG:CZ | 2.40 | 0.51 |
| 41:BP:88:LEU:O | 41:BP:89:ALA:C | 2.54 | 0.51 |
| 42:BQ:141:GLN:OXT | 51:BZ:98:MET:HB2 | 2.10 | 0.51 |
| 44:BS:34:HIS:CE1 | 44:BS:54:LEU:CB | 2.94 | 0.51 |
| 45:BT:128:GLU:O | 45:BT:130:ALA:N | 2.43 | 0.51 |
| 46:BU:31:SER:C | 46:BU:33:ARG:H | 2.19 | 0.51 |
| 46:BU:47:TYR:HA | 46:BU:50:ARG:NH2 | 2.26 | 0.51 |
| 50:BY:39:VAL:CG1 | 50:BY:40:GLU:H | 2.04 | 0.51 |
| 51:BZ:12:GLY:O | 51:BZ:13:GLU:HG3 | 2.11 | 0.51 |
| 51:BZ:39:VAL:HG23 | 51:BZ:40:ASP:N | 2.25 | 0.51 |
| 1:CA:504:C:H2' | 1:CA:504:C:O2 | 2.10 | 0.51 |
| 1:CA:658:G:C1' | 15:CO:22:THR:HB | 2.41 | 0.51 |
| 1:CA:706:A:H5" | 11:CK:22:HIS:CD2 | 2.46 | 0.51 |
| 8:CH:134:ILE:O | 8:CH:135:CYS:HB3 | 2.11 | 0.51 |
| 15:CO:66:LEU:O | 15:CO:69:TYR:HB3 | 2.11 | 0.51 |
| 17:CQ:65:ILE:HD12 | 17:CQ:65:ILE:N | 2.26 | 0.51 |
| 22:D0:50:ASN:O | 22:D0:62:LEU:HB2 | 2.11 | 0.51 |
| 24:D2:31:GLU:HG2 | 24:D2:37:PHE:CD1 | 2.41 | 0.51 |
| 24:D2:41:ILE:CG1 | 31:DA:94(A):G:N2 | 2.74 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 27:D5:36:CYS:SG | 27:D5:49:CYS:HB3 | 2.51 | 0.51 |
| 28:D6:24:GLU:HA | 28:D6:24:GLU:OE1 | 2.11 | 0.51 |
| 31:DA:271(Q):G:C2 | 31:DA:271(R):G:C5 | 2.99 | 0.51 |
| 31:DA:560:C:H5' | 46:DU:52:ARG:HH12 | 1.76 | 0.51 |
| 31:DA:1144:G:C6 | 31:DA:1145:C:N4 | 2.79 | 0.51 |
| 31:DA:1250:G:H4' | 31:DA:1251:C:OP2 | 2.10 | 0.51 |
| 31:DA:1788:C:H2' | 31:DA:1789:A:C8 | 2.46 | 0.51 |
| 31:DA:2543:G:H8 | 31:DA:2543:G:H5' | 1.76 | 0.51 |
| 31:DA:2740:A:C6 | 31:DA:2764:A:C8 | 2.99 | 0.51 |
| 31:DA:2808:U:H2' | 31:DA:2809:A:H5' | 1.91 | 0.51 |
| 31:DA:2808:U:N3 | 31:DA:2892:A:C6 | 2.78 | 0.51 |
| 33:DD:27:THR:O | 33:DD:28:GLU:HB2 | 2.10 | 0.51 |
| 34:DE:24:THR:HB | 34:DE:186:GLY:O | 2.11 | 0.51 |
| 35:DF:39:TRP:O | 35:DF:43:LYS:HG2 | 2.10 | 0.51 |
| 36:DG:82:LEU:C | 36:DG:83:ARG:HG3 | 2.35 | 0.51 |
| 37:DH:40:GLU:O | 37:DH:41:MET:CB | 2.57 | 0.51 |
| 37:DH:83:TYR:HA | 37:DH:135:GLY:O | 2.11 | 0.51 |
| 37:DH:144:VAL:HA | 37:DH:147:ASN:OD1 | 2.11 | 0.51 |
| 38:DI:93:THR:HG22 | 38:DI:119:PRO:HB3 | 1.91 | 0.51 |
| 41:DP:85:LEU:HB3 | 41:DP:114:ILE:HD12 | 1.92 | 0.51 |
| 42:DQ:63:LYS:HB2 | 42:DQ:63:LYS:HZ3 | 1.74 | 0.51 |
| 49:DX:92:LEU:O | 49:DX:93:GLU:HB3 | 2.11 | 0.51 |
| 1:AA:163:C:H2' | 1:AA:164:U:H6 | 1.71 | 0.51 |
| 1:AA:170:U:O2' | 1:AA:171:A:H5' | 2.11 | 0.51 |
| 1:AA:181:G:N2 | 1:AA:183:G:N2 | 2.59 | 0.51 |
| 1:AA:380:G:C2 | 1:AA:384:G:N1 | 2.79 | 0.51 |
| 1:AA:858:G:O6 | 1:AA:869:G:H3' | 2.10 | 0.51 |
| 1:AA:986:A:C6 | 1:AA:1220:G:C2 | 2.98 | 0.51 |
| 1:AA:1162:C:C2 | 1:AA:1175:G:N2 | 2.79 | 0.51 |
| 1:AA:1347:G:C8 | 9:AI:107:ARG:HB3 | 2.46 | 0.51 |
| 1:AA:1418:A:N3 | 31:BA:1959:G:H1' | 2.26 | 0.51 |
| 2:AB:97:TRP:HH2 | 2:AB:176:GLU:CG | 2.24 | 0.51 |
| 3:AC:6:HIS:NE2 | 3:AC:184:TYR:CE2 | 2.78 | 0.51 |
| 4:AD:2:GLY:O | 4:AD:3:ARG:C | 2.53 | 0.51 |
| 15:AO:67:LEU:CD2 | 15:AO:78:TYR:CE1 | 2.94 | 0.51 |
| 18:AR:50:ILE:HD11 | 18:AR:70:ILE:CG2 | 2.33 | 0.51 |
| 24:B2:32:LEU:O | 24:B2:33:MET:C | 2.54 | 0.51 |
| 28:B6:40:CYS:SG | 28:B6:45:LYS:CD | 2.99 | 0.51 |
| 30:B8:4:MET:HE1 | 31:BA:593:G:C1' | 2.41 | 0.51 |
| 31:BA:34:C:HO2' | 31:BA:35:G:P | 2.34 | 0.51 |
| 31:BA:52:A:C2' | 31:BA:53:A:H5' | 2.40 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:445:C:OP1 | 46:BU:2:PRO:HA | 2.11 | 0.51 |
| 31:BA:786:C:C2' | 31:BA:787:U:H5' | 2.40 | 0.51 |
| 31:BA:1374:G:C5 | 31:BA:1375:C:C4 | 2.99 | 0.51 |
| 31:BA:1385:G:H4' | 31:BA:1386:C:OP1 | 2.11 | 0.51 |
| 31:BA:2377:A:H4' | 44:BS:107:GLU:CG | 2.41 | 0.51 |
| 31:BA:2475:C:H5" | 31:BA:2475:C:C6 | 2.38 | 0.51 |
| 31:BA:2492:U:H2' | 31:BA:2493:U:H6 | 1.76 | 0.51 |
| 32:BB:2:C:H2' | 32:BB:3:C:C6 | 2.45 | 0.51 |
| 35:BF:119:ARG:HG2 | 35:BF:119:ARG:O | 2.10 | 0.51 |
| 36:BG:16:ARG:NE | 36:BG:31:VAL:HG11 | 2.26 | 0.51 |
| 37:BH:13:LYS:HE2 | 37:BH:13:LYS:HA | 1.93 | 0.51 |
| 37:BH:89:ILE:O | 37:BH:90:LYS:CG | 2.57 | 0.51 |
| 42:BQ:7:MET:O | 42:BQ:10:ARG:NE | 2.41 | 0.51 |
| 42:BQ:34:LEU:CD1 | 42:BQ:129:THR:HB | 2.41 | 0.51 |
| 48:BW:6:ILE:HA | 48:BW:103:ILE:O | 2.11 | 0.51 |
| 1:CA:17:U:C2 | 1:CA:18:C:C5 | 2.99 | 0.51 |
| 1:CA:343:U:H2' | 1:CA:346:G:O6 | 2.11 | 0.51 |
| 3:CC:199:LYS:HB3 | 3:CC:201:TYR:HE1 | 1.76 | 0.51 |
| 6:CF:18:GLN:O | 6:CF:21:LEU:HB2 | 2.11 | 0.51 |
| 6:CF:99:ALA:HB3 | 18:CR:29:PHE:CE2 | 2.46 | 0.51 |
| 8:CH:6:ILE:HD12 | 8:CH:6:ILE:N | 2.26 | 0.51 |
| 8:CH:13:ILE:O | 8:CH:17:THR:HG23 | 2.11 | 0.51 |
| 10:CJ:82:ILE:HG22 | 10:CJ:82:ILE:O | 2.09 | 0.51 |
| 12:CL:83:VAL:CG1 | 12:CL:100:ILE:HG23 | 2.41 | 0.51 |
| 12:CL:86:ARG:HB2 | 12:CL:101:VAL:HG23 | 1.93 | 0.51 |
| 23:D1:49:VAL:HB | 23:D1:64:ALA:HB2 | 1.92 | 0.51 |
| 30:D8:4:MET:O | 30:D8:62:LEU:HD12 | 2.10 | 0.51 |
| 30:D8:53:PRO:O | 30:D8:54:GLU:C | 2.54 | 0.51 |
| 31:DA:90:U:O4' | 31:DA:90:U:O2 | 2.27 | 0.51 |
| 31:DA:409:C:O2' | 31:DA:410:G:H5' | 2.11 | 0.51 |
| 31:DA:1494:A:N3 | 31:DA:1494:A:C2' | 2.74 | 0.51 |
| 31:DA:1670:C:O2 | 34:DE:129:HIS:CE1 | 2.57 | 0.51 |
| 31:DA:2315:G:H2' | 31:DA:2316:C:H6 | 1.73 | 0.51 |
| 31:DA:2335:A:C8 | 31:DA:2337:G:N7 | 2.79 | 0.51 |
| 31:DA:2377:A:H4' | 44:DS:107:GLU:CG | 2.41 | 0.51 |
| 31:DA:2830:G:C5' | 31:DA:2830:G:C8 | 2.93 | 0.51 |
| 32:DB:45:A:H2' | 32:DB:46:A:H5' | 1.93 | 0.51 |
| 33:DD:48:ARG:NH1 | 33:DD:48:ARG:HG3 | 2.26 | 0.51 |
| 33:DD:149:PRO:O | 33:DD:150:LYS:HB2 | 2.11 | 0.51 |
| 41:DP:45:LEU:HD22 | 41:DP:46:LYS:N | 2.25 | 0.51 |
| 42:DQ:75:THR:CA | 42:DQ:88:GLY:HA2 | 2.41 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 42:DQ:97:VAL:HG21 | 42:DQ:103:MET:CE | 2.41 | 0.51 |
| 46:DU:65:ILE:CG1 | 46:DU:96:ALA:HB3 | 2.41 | 0.51 |
| 51:DZ:14:LYS:HB2 | 51:DZ:17:ALA:HB3 | 1.93 | 0.51 |
| 1:AA:102:G:N3 | 1:AA:103:C:C6 | 2.78 | 0.51 |
| 1:AA:817:C:H4' | 1:AA:818:G:OP1 | 2.11 | 0.51 |
| 1:AA:963:G:H21 | 10:AJ:55:LYS:CD | 2.25 | 0.51 |
| 1:AA:1274:G:N2 | 1:AA:1275:A:N6 | 2.59 | 0.51 |
| 3:AC:61:ALA:O | 3:AC:62:ASP:HB2 | 2.09 | 0.51 |
| 3:AC:182:ILE:HG12 | 3:AC:203:PHE:HA | 1.92 | 0.51 |
| 4:AD:3:ARG:O | 4:AD:5:ILE:HG13 | 2.11 | 0.51 |
| 6:AF:53:ALA:O | 6:AF:54:LYS:HB2 | 2.10 | 0.51 |
| 13:AM:3:ARG:HH21 | 36:BG:146:TYR:HB2 | 1.76 | 0.51 |
| 16:AP:5:ARG:HB3 | 16:AP:67:THR:OG1 | 2.12 | 0.51 |
| 27:B5:25:LEU:HD11 | 48:BW:19:LEU:HB3 | 1.93 | 0.51 |
| 31:BA:528:A:C2 | 31:BA:2043:C:H5' | 2.46 | 0.51 |
| 31:BA:637:A:OP2 | 41:BP:115:LEU:HB2 | 2.11 | 0.51 |
| 31:BA:1721:G:C2 | 31:BA:1739:U:OP2 | 2.63 | 0.51 |
| 31:BA:2575:C:H2' | 31:BA:2578:G:O6 | 2.11 | 0.51 |
| 32:BB:66:A:C6 | 32:BB:109:C:C5 | 2.99 | 0.51 |
| 34:BE:60:ASN:OD1 | 34:BE:62:PRO:HD2 | 2.11 | 0.51 |
| 34:BE:70:ALA:O | 34:BE:72:VAL:C | 2.54 | 0.51 |
| 39:BN:78:TYR:HD1 | 39:BN:79:PRO:CG | 2.24 | 0.51 |
| 42:BQ:34:LEU:HD11 | 42:BQ:129:THR:CB | 2.41 | 0.51 |
| 42:BQ:37:LEU:HB2 | 42:BQ:128:LYS:O | 2.11 | 0.51 |
| 44:BS:14:VAL:HG13 | 44:BS:15:ARG:N | 2.23 | 0.51 |
| 44:BS:98:VAL:CG1 | 44:BS:100:ALA:H | 2.23 | 0.51 |
| 48:BW:62:HIS:O | 48:BW:63:ASP:C | 2.53 | 0.51 |
| 50:BY:27:VAL:HG12 | 50:BY:29:GLU:N | 2.19 | 0.51 |
| 50:BY:28:LYS:HG3 | 50:BY:37:VAL:HA | 1.92 | 0.51 |
| 50:BY:28:LYS:CB | 50:BY:37:VAL:HB | 2.32 | 0.51 |
| 51:BZ:70:LEU:HG | 51:BZ:91:LEU:HD11 | 1.93 | 0.51 |
| 51:BZ:175:VAL:HB | 51:BZ:176:PRO:CD | 2.40 | 0.51 |
| 1:CA:67:C:H2' | 1:CA:68:G:C8 | 2.45 | 0.51 |
| 1:CA:186:C:C2 | 1:CA:187:C:C5 | 2.98 | 0.51 |
| 1:CA:1105:A:H2' | 1:CA:1106:G:H8 | 1.76 | 0.51 |
| 1:CA:1468:A:H2' | 1:CA:1469:G:O4' | 2.10 | 0.51 |
| 2:CB:9:GLU:CD | 2:CB:9:GLU:H | 2.17 | 0.51 |
| 2:CB:12:GLU:HA | 2:CB:16:HIS:HB2 | 1.93 | 0.51 |
| 5:CE:7:GLU:HG2 | 5:CE:112:LEU:HD22 | 1.91 | 0.51 |
| 25:D3:1:MET:HB2 | 25:D3:38:GLU:HG3 | 1.92 | 0.51 |
| 25:D3:32:GLN:HB2 | 31:DA:1158:C:H4' | 1.93 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:82:G:O2' | 31:DA:83:G:H5' | 2.11 | 0.51 |
| 31:DA:139(A):G:C2' | 31:DA:140:G:O5' | 2.59 | 0.51 |
| 31:DA:574:C:N3 | 34:DE:145:LYS:HE2 | 2.25 | 0.51 |
| 31:DA:1159:U:O2' | 31:DA:1160:G:H5' | 2.11 | 0.51 |
| 31:DA:1679:U:C2' | 31:DA:1680:U:H5' | 2.40 | 0.51 |
| 31:DA:2287:A:H2 | 31:DA:2346:A:N1 | 2.09 | 0.51 |
| 31:DA:2309:A:C2 | 31:DA:2310:A:H2 | 2.29 | 0.51 |
| 31:DA:2408:U:H2' | 31:DA:2409:G:C8 | 2.45 | 0.51 |
| 31:DA:2492:U:H2' | 31:DA:2493:U:H6 | 1.76 | 0.51 |
| 33:DD:11:PRO:C | 33:DD:13:ARG:H | 2.19 | 0.51 |
| 34:DE:49:LEU:HD23 | 34:DE:81:ILE:HG12 | 1.92 | 0.51 |
| 38:DI:38:LEU:C | 38:DI:40:THR:H | 2.18 | 0.51 |
| 38:DI:108:THR:O | 38:DI:109:ILE:HG13 | 2.09 | 0.51 |
| 41:DP:148:LEU:HD13 | 41:DP:148:LEU:N | 2.26 | 0.51 |
| 46:DU:95:LEU:HD22 | 47:DV:4:ILE:HD13 | 1.92 | 0.51 |
| 1:AA:7:G:N2 | 5:AE:121:LYS:HG2 | 2.27 | 0.50 |
| 1:AA:321:A:H62 | 1:AA:328:C:H1' | 1.76 | 0.50 |
| 1:AA:397:A:H5" | 1:AA:397:A:N3 | 2.27 | 0.50 |
| 1:AA:832:C:H42 | 1:AA:854:G:H1 | 1.59 | 0.50 |
| 1:AA:1037:C:H2' | 1:AA:1038:C:O4' | 2.11 | 0.50 |
| 1:AA:1098:C:N3 | 1:AA:1099:G:C8 | 2.79 | 0.50 |
| 2:AB:116:GLU:HA | 2:AB:119:GLU:HB3 | 1.93 | 0.50 |
| 5:AE:40:ARG:NH2 | 5:AE:66:MET:HE3 | 2.26 | 0.50 |
| 10:AJ:33:GLN:O | 10:AJ:75:ILE:HG23 | 2.11 | 0.50 |
| 16:AP:34:GLU:OE2 | 16:AP:55:ARG:HD3 | 2.11 | 0.50 |
| 18:AR:59:SER:HB3 | 18:AR:62:GLU:CG | 2.38 | 0.50 |
| 22:B0:50:ASN:ND2 | 22:B0:83:PRO:HD3 | 2.25 | 0.50 |
| 30:B8:4:MET:HE3 | 30:B8:61:LEU:HD13 | 1.93 | 0.50 |
| 31:BA:259:G:H21 | 31:BA:621:A:H8 | 1.57 | 0.50 |
| 31:BA:329:G:H22 | 50:BY:19:LYS:HZ1 | 1.55 | 0.50 |
| 31:BA:588:U:H2' | 31:BA:589:C:H6 | 1.72 | 0.50 |
| 31:BA:1112:G:C1' | 31:BA:1113:U:OP1 | 2.59 | 0.50 |
| 31:BA:1336:A:H2' | 31:BA:1337:G:H8 | 1.74 | 0.50 |
| 31:BA:1381:G:C2' | 31:BA:1382:G:H5' | 2.42 | 0.50 |
| 31:BA:1484:G:H22 | 31:BA:1505:C:H5 | 1.57 | 0.50 |
| 31:BA:1786:A:H1' | 31:BA:1938:A:N6 | 2.26 | 0.50 |
| 31:BA:2012:G:C4' | 48:BW:96:ILE:HD11 | 2.27 | 0.50 |
| 31:BA:2050:C:H1' | 34:BE:156:MET:CE | 2.41 | 0.50 |
| 31:BA:2187:G:N7 | 31:BA:2188:C:C4 | 2.79 | 0.50 |
| 31:BA:2428:G:H5" | 31:BA:2429:G:O5' | 2.10 | 0.50 |
| 31:BA:2480:C:N4 | 31:BA:2481:G:C6 | 2.79 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 33:BD:142:VAL:HG21 | 33:BD:191:ALA:HB1 | 1.93 | 0.50 |
| 38:BI:68:LEU:HA | 38:BI:71:ILE:HG23 | 1.93 | 0.50 |
| 42:BQ:133:ARG:O | 42:BQ:134:ARG:HB2 | 2.10 | 0.50 |
| 43:BR:10:LEU:HB3 | 43:BR:17:ARG:CD | 2.41 | 0.50 |
| 43:BR:29:LEU:HB3 | 43:BR:75:LEU:HD11 | 1.93 | 0.50 |
| 47:BV:4:ILE:O | 47:BV:39:LEU:HB3 | 2.11 | 0.50 |
| 50:BY:7:VAL:HB | 50:BY:8:LYS:CD | 2.40 | 0.50 |
| 1:CA:143:A:N1 | 1:CA:220:G:O6 | 2.45 | 0.50 |
| 1:CA:181:G:N2 | 1:CA:183:G:N2 | 2.59 | 0.50 |
| 1:CA:193:C:H2' | 1:CA:194:C:C6 | 2.46 | 0.50 |
| 1:CA:570:G:H2' | 1:CA:571:U:C6 | 2.46 | 0.50 |
| 1:CA:655:A:C2 | 1:CA:754:C:C4 | 2.99 | 0.50 |
| 1:CA:668:G:O2' | 15:CO:46:HIS:HD2 | 1.94 | 0.50 |
| 1:CA:832:C:H42 | 1:CA:854:G:H1 | 1.59 | 0.50 |
| 1:CA:881:G:OP2 | 12:CL:12:ARG:NH2 | 2.44 | 0.50 |
| 1:CA:1252:A:H2' | 1:CA:1253:G:O4' | 2.11 | 0.50 |
| 1:CA:1321:C:H5' | 1:CA:1322:C:C5' | 2.40 | 0.50 |
| 6:CF:11:ASN:O | 6:CF:14:LEU:HG | 2.12 | 0.50 |
| 6:CF:76:ALA:O | 6:CF:80:ARG:HG3 | 2.09 | 0.50 |
| 12:CL:5:PRO:HB2 | 12:CL:10:LEU:HD21 | 1.91 | 0.50 |
| 13:CM:91:ARG:HB2 | 13:CM:98:VAL:CG2 | 2.38 | 0.50 |
| 16:CP:18:ARG:HD3 | 16:CP:35:LYS:HD2 | 1.92 | 0.50 |
| 23:D1:56:GLN:OE1 | 23:D1:56:GLN:HA | 2.05 | 0.50 |
| 23:D1:67:ILE:HD12 | 23:D1:67:ILE:H | 1.76 | 0.50 |
| 24:D2:12:GLU:C | 24:D2:14:ARG:N | 2.68 | 0.50 |
| 27:D5:40:LYS:HZ3 | 27:D5:46:CYS:CA | 2.24 | 0.50 |
| 30:D8:48:PHE:H | 30:D8:48:PHE:HD1 | 1.56 | 0.50 |
| 31:DA:415:A:H2' | 31:DA:416:C:C6 | 2.46 | 0.50 |
| 31:DA:661:C:O3' | 41:DP:18:ARG:HG2 | 2.11 | 0.50 |
| 31:DA:685:A:N7 | 31:DA:774:A:C5 | 2.79 | 0.50 |
| 31:DA:1834:U:O2 | 31:DA:1834:U:H2' | 2.11 | 0.50 |
| 31:DA:2228:G:C6 | 31:DA:2229:C:C4 | 3.00 | 0.50 |
| 31:DA:2712:U:O2' | 31:DA:2712(A):A:P | 2.69 | 0.50 |
| 32:DB:15:A:H2' | 32:DB:16:G:OP1 | 2.10 | 0.50 |
| 33:DD:138:VAL:HG21 | 33:DD:167:GLY:HA2 | 1.93 | 0.50 |
| 39:DN:2:LYS:HE3 | 47:DV:12:TYR:HA | 1.92 | 0.50 |
| 41:DP:21:ARG:O | 41:DP:21:ARG:CG | 2.57 | 0.50 |
| 41:DP:62:LEU:H | 41:DP:62:LEU:CD1 | 2.07 | 0.50 |
| 42:DQ:141:GLN:HG3 | 51:DZ:72:ARG:NH1 | 2.25 | 0.50 |
| 43:DR:67:LEU:CD1 | 43:DR:76:VAL:HG21 | 2.41 | 0.50 |
| 44:DS:34:HIS:CE1 | 44:DS:54:LEU:CB | 2.93 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:615:C:H2' | 1:AA:616:G:O4' | 2.11 | 0.50 |
| 1:AA:723:U:H5" | 1:AA:724:G:OP2 | 2.11 | 0.50 |
| 1:AA:814:A:N7 | 1:AA:816:A:C4 | 2.79 | 0.50 |
| 1:AA:1015:A:C6 | 1:AA:1016:A:C6 | 2.98 | 0.50 |
| 1:AA:1438:G:H2' | 1:AA:1439:C:H6 | 1.75 | 0.50 |
| 2:AB:111:ARG:HH11 | 2:AB:111:ARG:CG | 1.98 | 0.50 |
| 2:AB:130:ARG:HA | 2:AB:130:ARG:HE | 1.77 | 0.50 |
| 6:AF:55:ASP:HB2 | 6:AF:86:ARG:HH12 | 1.75 | 0.50 |
| 7:AG:15:ASP:O | 7:AG:19:GLY:HA2 | 2.11 | 0.50 |
| 12:AL:60:LEU:N | 12:AL:60:LEU:HD22 | 2.26 | 0.50 |
| 24:B2:33:MET:HG3 | 49:BX:10:ALA:HB1 | 1.93 | 0.50 |
| 31:BA:911:A:C4 | 42:BQ:9:TYR:OH | 2.54 | 0.50 |
| 31:BA:1158:C:O2' | 31:BA:1159:U:H5' | 2.11 | 0.50 |
| 31:BA:1169:G:C8 | 31:BA:1169:G:C3' | 2.95 | 0.50 |
| 31:BA:1171:G:N7 | 31:BA:1173:G:H1' | 2.26 | 0.50 |
| 31:BA:1262:A:C5 | 31:BA:1263:U:C5 | 2.99 | 0.50 |
| 31:BA:1493:C:O2 | 31:BA:1493:C:C2' | 2.56 | 0.50 |
| 31:BA:1678:G:N2 | 31:BA:1989:G:N2 | 2.50 | 0.50 |
| 31:BA:2506:U:H4' | 31:BA:2507:C:OP1 | 2.11 | 0.50 |
| 31:BA:2526:G:C5 | 31:BA:2527:C:C5 | 3.00 | 0.50 |
| 40:BO:61:VAL:O | 40:BO:61:VAL:HG13 | 2.11 | 0.50 |
| 42:BQ:85:LYS:O | 42:BQ:86:GLY:C | 2.53 | 0.50 |
| 42:BQ:141:GLN:HG3 | 51:BZ:72:ARG:NH1 | 2.26 | 0.50 |
| 1:CA:276:G:C2' | 1:CA:277:C:H5' | 2.41 | 0.50 |
| 1:CA:1240:U:OP2 | 7:CG:116:ALA:HB2 | 2.11 | 0.50 |
| 2:CB:219:VAL:O | 2:CB:222:ILE:HB | 2.10 | 0.50 |
| 3:CC:182:ILE:CG1 | 3:CC:203:PHE:HD1 | 2.21 | 0.50 |
| 4:CD:131:ARG:HD3 | 4:CD:131:ARG:H | 1.76 | 0.50 |
| 5:CE:40:ARG:NH2 | 5:CE:66:MET:HE3 | 2.26 | 0.50 |
| 31:DA:52:A:O2' | 31:DA:53:A:H5' | 2.10 | 0.50 |
| 31:DA:105:C:H2' | 31:DA:106:C:H6 | 1.76 | 0.50 |
| 31:DA:271(P):C:O2' | 31:DA:271(Q):G:H5' | 2.11 | 0.50 |
| 31:DA:414:C:O2' | 31:DA:415:A:H5' | 2.12 | 0.50 |
| 31:DA:633:A:H2' | 31:DA:634:C:H5' | 1.93 | 0.50 |
| 31:DA:975:C:H2' | 31:DA:975:C:O2 | 2.10 | 0.50 |
| 31:DA:1044:G:N2 | 31:DA:1112:G:O6 | 2.44 | 0.50 |
| 31:DA:2654:A:OP1 | 31:DA:2654:A:C8 | 2.57 | 0.50 |
| 31:DA:2801:A:O2' | 31:DA:2895:U:H4' | 2.12 | 0.50 |
| 31:DA:2846:G:H2' | 31:DA:2847:U:O4' | 2.11 | 0.50 |
| 31:DA:2880:C:O2' | 43:DR:90:ARG:HD3 | 2.10 | 0.50 |
| 34:DE:59:VAL:CG2 | 34:DE:63:LEU:HA | 2.36 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 35:DF:2:LYS:O | 35:DF:25:PRO:HG2 | 2.11 | 0.50 |
| 36:DG:129:GLY:C | 36:DG:130:ASN:CG | 2.79 | 0.50 |
| 37:DH:89:ILE:O | 37:DH:90:LYS:CG | 2.59 | 0.50 |
| 37:DH:149:ARG:HA | 37:DH:162:ILE:CG1 | 2.38 | 0.50 |
| 39:DN:27:ALA:CB | 39:DN:106:MET:CE | 2.88 | 0.50 |
| 41:DP:149:GLU:O | 41:DP:149:GLU:HG3 | 2.11 | 0.50 |
| 43:DR:81:ASP:O | 43:DR:85:PRO:HG2 | 2.11 | 0.50 |
| 44:DS:98:VAL:CG1 | 44:DS:100:ALA:H | 2.25 | 0.50 |
| 45:DT:128:GLU:O | 45:DT:130:ALA:N | 2.42 | 0.50 |
| 47:DV:19:LYS:HZ3 | 47:DV:20:LEU:CB | 2.18 | 0.50 |
| 51:DZ:29:TYR:HA | 51:DZ:33:LEU:O | 2.10 | 0.50 |
| 51:DZ:150:LEU:HD23 | 51:DZ:171:ILE:HD11 | 1.92 | 0.50 |
| 1:AA:66:G:C4’ | 1:AA:173:U:C5 | 2.94 | 0.50 |
| 1:AA:484:G:H4’ | 1:AA:485:G:O5’ | 2.11 | 0.50 |
| 3:AC:113:ALA:O | 3:AC:115:LEU:N | 2.44 | 0.50 |
| 6:AF:12:PRO:O | 6:AF:14:LEU:N | 2.44 | 0.50 |
| 12:AL:24:VAL:HG13 | 12:AL:98:TYR:HE2 | 1.76 | 0.50 |
| 12:AL:46:LYS:HB2 | 12:AL:92:ASP:O | 2.11 | 0.50 |
| 16:AP:39:TYR:C | 16:AP:39:TYR:CD1 | 2.87 | 0.50 |
| 30:B8:6:THR:CG2 | 31:BA:243:U:OP1 | 2.58 | 0.50 |
| 30:B8:46:ARG:NH2 | 41:BP:65:ARG:NH2 | 2.57 | 0.50 |
| 31:BA:197:A:N6 | 31:BA:2430:A:H2’ | 2.25 | 0.50 |
| 31:BA:285:C:N3 | 31:BA:286:C:C5 | 2.80 | 0.50 |
| 31:BA:309:G:N3 | 31:BA:329:G:O2’ | 2.43 | 0.50 |
| 31:BA:454:A:H4’ | 31:BA:455:C:OP2 | 2.12 | 0.50 |
| 31:BA:481:G:OP2 | 50:BY:47:LYS:CE | 2.60 | 0.50 |
| 31:BA:667:U:H2’ | 31:BA:668:G:C5’ | 2.41 | 0.50 |
| 31:BA:774:A:C2 | 31:BA:787:U:O2’ | 2.61 | 0.50 |
| 31:BA:774:A:H2 | 31:BA:787:U:O2’ | 1.93 | 0.50 |
| 31:BA:1649:G:C6 | 31:BA:2009:G:O6 | 2.65 | 0.50 |
| 35:BF:29:ASN:OD1 | 35:BF:29:ASN:C | 2.54 | 0.50 |
| 35:BF:144:LYS:C | 35:BF:146:ALA:H | 2.19 | 0.50 |
| 35:BF:164:ARG:HG2 | 35:BF:164:ARG:NH1 | 2.26 | 0.50 |
| 36:BG:76:SER:HB3 | 36:BG:84:LYS:H | 1.76 | 0.50 |
| 36:BG:106:LEU:HD12 | 36:BG:110:ALA:HB3 | 1.94 | 0.50 |
| 37:BH:43:VAL:HG23 | 37:BH:43:VAL:O | 2.11 | 0.50 |
| 37:BH:54:ARG:HD2 | 37:BH:56:SER:O | 2.11 | 0.50 |
| 37:BH:92:ILE:C | 37:BH:94:TYR:H | 2.19 | 0.50 |
| 38:BI:98:ALA:O | 38:BI:99:GLU:C | 2.53 | 0.50 |
| 39:BN:79:PRO:CD | 39:BN:80:GLY:H | 2.25 | 0.50 |
| 45:BT:28:VAL:HG22 | 45:BT:47:GLY:N | 2.27 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 45:BT:28:VAL:O | 45:BT:29:ARG:CD | 2.59 | 0.50 |
| 1:CA:409:G:C2' | 1:CA:410:G:H5' | 2.40 | 0.50 |
| 1:CA:710:G:H5" | 6:CF:54:LYS:HE3 | 1.94 | 0.50 |
| 1:CA:914:A:O2' | 1:CA:915:A:H5' | 2.10 | 0.50 |
| 1:CA:1084:G:C5 | 1:CA:1085:U:C4 | 2.99 | 0.50 |
| 4:CD:203:VAL:HG12 | 4:CD:204:ILE:N | 2.26 | 0.50 |
| 6:CF:60:PHE:C | 6:CF:61:LEU:HD12 | 2.36 | 0.50 |
| 11:CK:27:ASN:OD1 | 11:CK:28:THR:N | 2.44 | 0.50 |
| 16:CP:50:LYS:C | 16:CP:50:LYS:HD3 | 2.37 | 0.50 |
| 17:CQ:87:LYS:HE2 | 17:CQ:87:LYS:HA | 1.92 | 0.50 |
| 20:CT:26:ASN:O | 20:CT:30:LYS:HB2 | 2.11 | 0.50 |
| 24:D2:14:ARG:NH1 | 24:D2:57:ILE:HG22 | 2.26 | 0.50 |
| 24:D2:57:ILE:HG23 | 24:D2:57:ILE:O | 2.10 | 0.50 |
| 28:D6:39:TYR:HE1 | 31:DA:2347:C:HO2' | 1.59 | 0.50 |
| 29:D7:46:VAL:HB | 29:D7:48:LYS:NZ | 2.26 | 0.50 |
| 30:D8:61:LEU:HD22 | 31:DA:593:G:O3' | 2.11 | 0.50 |
| 31:DA:1225:G:P | 47:DV:88:ARG:HB3 | 2.51 | 0.50 |
| 31:DA:1266:G:O2' | 31:DA:2012:G:O6 | 2.28 | 0.50 |
| 31:DA:1337:G:H2' | 31:DA:1338:G:H8 | 1.75 | 0.50 |
| 31:DA:1469:A:H2' | 31:DA:1470:G:O5' | 2.10 | 0.50 |
| 31:DA:1512:U:O2' | 31:DA:1513:C:H5' | 2.12 | 0.50 |
| 31:DA:1884:A:O2' | 31:DA:1885:A:H5' | 2.11 | 0.50 |
| 31:DA:1987:G:C8 | 31:DA:1987:G:C5' | 2.94 | 0.50 |
| 31:DA:2308:G:H2' | 31:DA:2309:A:C8 | 2.47 | 0.50 |
| 31:DA:2661:G:H8 | 31:DA:2662:A:N3 | 2.10 | 0.50 |
| 33:DD:267:SER:O | 33:DD:268:ARG:CB | 2.57 | 0.50 |
| 35:DF:167:ALA:HB1 | 35:DF:173:VAL:HG11 | 1.92 | 0.50 |
| 37:DH:54:ARG:HD2 | 37:DH:56:SER:O | 2.11 | 0.50 |
| 39:DN:65:LYS:O | 39:DN:69:GLN:HG3 | 2.11 | 0.50 |
| 41:DP:35:HIS:O | 41:DP:36:LYS:CB | 2.59 | 0.50 |
| 45:DT:57:PHE:CG | 45:DT:58:ASN:N | 2.79 | 0.50 |
| 47:DV:50:PRO:C | 47:DV:51:VAL:HG23 | 2.36 | 0.50 |
| 50:DY:14:LEU:HD11 | 50:DY:22:GLY:HA2 | 1.93 | 0.50 |
| 1:AA:10:A:H2' | 1:AA:11:G:H8 | 1.75 | 0.50 |
| 1:AA:328:C:O2 | 1:AA:328:C:C2' | 2.58 | 0.50 |
| 1:AA:1216:G:H5" | 14:AN:5:ALA:HB2 | 1.92 | 0.50 |
| 1:AA:1371:G:C6 | 1:AA:1372:U:C4 | 3.00 | 0.50 |
| 4:AD:64:LEU:HD12 | 4:AD:64:LEU:O | 2.11 | 0.50 |
| 4:AD:119:GLN:O | 4:AD:123:HIS:HD2 | 1.94 | 0.50 |
| 6:AF:30:LEU:O | 6:AF:35:ALA:N | 2.42 | 0.50 |
| 8:AH:8:ASP:O | 8:AH:11:THR:N | 2.45 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 21:AU:22:ARG:N | 21:AU:23:PRO:HD3 | 2.26 | 0.50 |
| 23:B1:13:ILE:HG12 | 23:B1:14:VAL:CA | 2.41 | 0.50 |
| 31:BA:303:U:H2' | 31:BA:304:G:H8 | 1.76 | 0.50 |
| 31:BA:594:U:H2' | 31:BA:595:C:C6 | 2.46 | 0.50 |
| 31:BA:2684:U:O2' | 40:BO:68:GLU:HG3 | 2.11 | 0.50 |
| 33:BD:197:GLY:O | 33:BD:198:ASN:HB3 | 2.11 | 0.50 |
| 42:BQ:141:GLN:HE21 | 51:BZ:72:ARG:N | 2.09 | 0.50 |
| 43:BR:2:ARG:CD | 43:BR:2:ARG:N | 2.74 | 0.50 |
| 45:BT:31:SER:HA | 45:BT:32:TYR:CD2 | 2.47 | 0.50 |
| 45:BT:109:GLU:HB3 | 45:BT:113:LYS:CE | 2.41 | 0.50 |
| 50:BY:15:VAL:HG12 | 50:BY:16:ALA:H | 1.77 | 0.50 |
| 50:BY:37:VAL:O | 50:BY:38:ILE:HG12 | 2.11 | 0.50 |
| 1:CA:10:A:H2' | 1:CA:11:G:H8 | 1.76 | 0.50 |
| 1:CA:840:C:H4' | 1:CA:848:C:O2 | 2.12 | 0.50 |
| 1:CA:1095:U:H2' | 1:CA:1096:C:C6 | 2.45 | 0.50 |
| 2:CB:28:PHE:CD1 | 2:CB:190:THR:HA | 2.47 | 0.50 |
| 8:CH:14:ARG:O | 8:CH:18:ARG:HD3 | 2.11 | 0.50 |
| 10:CJ:8:LEU:HD22 | 10:CJ:20:ALA:HB2 | 1.94 | 0.50 |
| 12:CL:119:LYS:O | 12:CL:120:TYR:HB2 | 2.12 | 0.50 |
| 20:CT:26:ASN:CB | 20:CT:71:THR:HG23 | 2.41 | 0.50 |
| 23:D1:85:LEU:CA | 23:D1:87:PRO:HD3 | 2.42 | 0.50 |
| 30:D8:31:HIS:HB3 | 31:DA:2420:C:H41 | 1.76 | 0.50 |
| 30:D8:46:ARG:NH2 | 41:DP:65:ARG:NH2 | 2.59 | 0.50 |
| 31:DA:1364:G:H5" | 31:DA:1365:A:OP2 | 2.11 | 0.50 |
| 31:DA:1484:G:N2 | 31:DA:1505:C:C5 | 2.79 | 0.50 |
| 31:DA:1593:G:H2' | 31:DA:1594:G:H8 | 1.76 | 0.50 |
| 38:DI:53:ALA:HB2 | 38:DI:56:LYS:CG | 2.32 | 0.50 |
| 42:DQ:24:GLY:HA3 | 51:DZ:78:LYS:HD2 | 1.94 | 0.50 |
| 42:DQ:134:ARG:O | 42:DQ:135:ASP:HB2 | 2.11 | 0.50 |
| 43:DR:9:LYS:C | 43:DR:10:LEU:HG | 2.37 | 0.50 |
| 43:DR:13:HIS:HE1 | 43:DR:15:SER:OG | 1.94 | 0.50 |
| 49:DX:52:VAL:HB | 49:DX:80:ILE:CG2 | 2.42 | 0.50 |
| 49:DX:65:ARG:O | 49:DX:66:LEU:CB | 2.59 | 0.50 |
| 50:DY:2:ARG:C | 50:DY:4:LYS:N | 2.64 | 0.50 |
| 1:AA:594:G:H1 | 1:AA:645:C:H42 | 1.59 | 0.50 |
| 1:AA:1082:G:C6 | 1:AA:1083:U:N3 | 2.80 | 0.50 |
| 2:AB:61:LEU:HD21 | 2:AB:68:ILE:HD11 | 1.93 | 0.50 |
| 2:AB:163:PHE:HA | 2:AB:185:ILE:O | 2.10 | 0.50 |
| 7:AG:47:CYS:O | 7:AG:50:ILE:HB | 2.12 | 0.50 |
| 9:AI:48:GLU:N | 9:AI:49:PRO:CD | 2.74 | 0.50 |
| 13:AM:44:ARG:HB2 | 13:AM:47:ASP:OD1 | 2.11 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 20:AT:78:ALA:HA | 20:AT:81:LYS:HD3 | 1.94 | 0.50 |
| 23:B1:11:ARG:CB | 23:B1:12:PRO:CD | 2.89 | 0.50 |
| 24:B2:46:GLN:C | 24:B2:48:HIS:N | 2.67 | 0.50 |
| 29:B7:8:ASN:ND2 | 29:B7:11:LYS:HB3 | 2.26 | 0.50 |
| 29:B7:24:THR:OG1 | 29:B7:25:PRO:HD2 | 2.10 | 0.50 |
| 30:B8:26:LYS:HB2 | 30:B8:44:LYS:HG3 | 1.93 | 0.50 |
| 31:BA:139(A):G:C2' | 31:BA:140:G:O5' | 2.59 | 0.50 |
| 31:BA:142:A:H5" | 31:BA:142(A):C:C5 | 2.47 | 0.50 |
| 31:BA:228:A:H2' | 31:BA:230:U:C1' | 2.41 | 0.50 |
| 31:BA:309:G:O3' | 50:BY:18:GLY:HA3 | 2.10 | 0.50 |
| 31:BA:1109:C:C5 | 31:BA:1110:G:C5 | 2.95 | 0.50 |
| 31:BA:1188:U:O2' | 31:BA:1189:A:H5' | 2.11 | 0.50 |
| 31:BA:1495:A:H2' | 31:BA:1496:A:C2 | 2.46 | 0.50 |
| 31:BA:1973:G:H2' | 31:BA:1974:C:H6 | 1.75 | 0.50 |
| 31:BA:2504:U:O2 | 31:BA:2504:U:H2' | 2.12 | 0.50 |
| 34:BE:112:GLY:O | 34:BE:159:HIS:HA | 2.11 | 0.50 |
| 35:BF:84:VAL:C | 35:BF:86:GLY:N | 2.68 | 0.50 |
| 36:BG:29:TRP:C | 36:BG:31:VAL:N | 2.69 | 0.50 |
| 36:BG:82:LEU:C | 36:BG:83:ARG:HG3 | 2.36 | 0.50 |
| 39:BN:128:HIS:O | 39:BN:128:HIS:CG | 2.64 | 0.50 |
| 41:BP:62:LEU:H | 41:BP:62:LEU:CD1 | 2.09 | 0.50 |
| 44:BS:29:PHE:H | 44:BS:89:ARG:HG2 | 1.76 | 0.50 |
| 47:BV:50:PRO:C | 47:BV:51:VAL:HG23 | 2.36 | 0.50 |
| 48:BW:20:VAL:O | 48:BW:23:LEU:HB2 | 2.10 | 0.50 |
| 49:BX:18:TYR:O | 49:BX:20:GLY:N | 2.45 | 0.50 |
| 49:BX:37:THR:HG22 | 49:BX:37:THR:O | 2.12 | 0.50 |
| 51:BZ:27:VAL:HA | 51:BZ:37:VAL:HG23 | 1.94 | 0.50 |
| 1:CA:321:A:H62 | 1:CA:328:C:H1' | 1.77 | 0.50 |
| 1:CA:701:C:O2 | 1:CA:703:G:N1 | 2.45 | 0.50 |
| 1:CA:946:A:C2 | 1:CA:1236:A:C2 | 2.99 | 0.50 |
| 1:CA:1157:A:C4 | 1:CA:1181:G:N2 | 2.79 | 0.50 |
| 1:CA:1242:C:H5" | 21:CU:10:ARG:HH12 | 1.76 | 0.50 |
| 1:CA:1279:A:H2 | 10:CJ:43:ARG:NH1 | 2.09 | 0.50 |
| 2:CB:15:VAL:C | 2:CB:16:HIS:CG | 2.90 | 0.50 |
| 3:CC:61:ALA:O | 3:CC:62:ASP:HB2 | 2.10 | 0.50 |
| 5:CE:144:THR:O | 5:CE:148:VAL:HG23 | 2.11 | 0.50 |
| 6:CF:22:GLU:OE1 | 6:CF:84:ASN:HB2 | 2.12 | 0.50 |
| 10:CJ:24:VAL:O | 10:CJ:28:ARG:HG3 | 2.11 | 0.50 |
| 10:CJ:32:ALA:HB1 | 10:CJ:75:ILE:CG1 | 2.41 | 0.50 |
| 13:CM:19:LEU:HB3 | 13:CM:25:ILE:HG21 | 1.93 | 0.50 |
| 20:CT:81:LYS:C | 20:CT:83:ARG:N | 2.69 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 22:D0:50:ASN:C | 22:D0:62:LEU:HB2 | 2.36 | 0.50 |
| 22:D0:70:GLN:O | 22:D0:77:ARG:HA | 2.12 | 0.50 |
| 31:DA:11:G:C2' | 31:DA:12:U:H5' | 2.42 | 0.50 |
| 31:DA:329:G:C4' | 31:DA:330:A:OP2 | 2.58 | 0.50 |
| 31:DA:782:A:H5' | 31:DA:783:A:C2 | 2.46 | 0.50 |
| 31:DA:1212:G:C2 | 31:DA:1236:G:C4 | 3.00 | 0.50 |
| 31:DA:1247:A:OP1 | 35:DF:95:ARG:NH2 | 2.38 | 0.50 |
| 31:DA:1499:C:O2' | 31:DA:1500:G:H5' | 2.12 | 0.50 |
| 31:DA:1607:C:H4' | 31:DA:1608:A:O5' | 2.11 | 0.50 |
| 31:DA:1722:A:N6 | 31:DA:1741:A:C2 | 2.80 | 0.50 |
| 31:DA:2190:G:H2' | 31:DA:2191:G:H5' | 1.92 | 0.50 |
| 31:DA:2314:C:O2 | 31:DA:2314:C:H2' | 2.11 | 0.50 |
| 32:DB:75:G:H5' | 32:DB:75:G:C8 | 2.42 | 0.50 |
| 33:DD:106:ILE:HD13 | 33:DD:106:ILE:C | 2.36 | 0.50 |
| 33:DD:131:LEU:HD12 | 33:DD:131:LEU:N | 2.26 | 0.50 |
| 33:DD:181:GLU:O | 33:DD:182:LEU:HD23 | 2.11 | 0.50 |
| 40:DO:17:ARG:HG2 | 40:DO:47:ILE:HD13 | 1.93 | 0.50 |
| 41:DP:48:PRO:O | 41:DP:51:PHE:N | 2.44 | 0.50 |
| 42:DQ:141:GLN:HE21 | 51:DZ:72:ARG:N | 2.09 | 0.50 |
| 44:DS:95:HIS:ND1 | 44:DS:96:GLY:N | 2.59 | 0.50 |
| 47:DV:24:LYS:HB2 | 47:DV:92:THR:HG1 | 1.71 | 0.50 |
| 51:DZ:63:ASP:O | 51:DZ:65:GLN:N | 2.44 | 0.50 |
| 1:AA:184:G:N2 | 1:AA:194:C:C2 | 2.80 | 0.50 |
| 1:AA:258:G:N3 | 1:AA:259:G:C8 | 2.80 | 0.50 |
| 1:AA:607:A:H2' | 1:AA:608:A:O4' | 2.11 | 0.50 |
| 1:AA:620:C:H2' | 1:AA:621:A:O4' | 2.11 | 0.50 |
| 1:AA:734:G:C2 | 1:AA:735:C:C2 | 2.99 | 0.50 |
| 1:AA:1242:C:H5" | 21:AU:10:ARG:HH12 | 1.77 | 0.50 |
| 1:AA:1321:C:C4 | 1:AA:1322:C:C4 | 3.00 | 0.50 |
| 3:AC:206:GLU:HG2 | 3:AC:207:VAL:H | 1.76 | 0.50 |
| 5:AE:68:GLU:O | 5:AE:68:GLU:HG3 | 2.11 | 0.50 |
| 6:AF:94:GLN:NE2 | 18:AR:32:ARG:HD2 | 2.26 | 0.50 |
| 8:AH:29:SER:HB3 | 8:AH:32:LYS:HD2 | 1.93 | 0.50 |
| 11:AK:102:GLY:C | 11:AK:103:LEU:HD22 | 2.36 | 0.50 |
| 12:AL:32:PHE:HE1 | 12:AL:86:ARG:HG3 | 1.75 | 0.50 |
| 23:B1:79:GLY:O | 23:B1:80:LEU:HD23 | 2.12 | 0.50 |
| 23:B1:94:LEU:CD2 | 23:B1:95:LEU:N | 2.75 | 0.50 |
| 25:B3:52:HIS:ND1 | 25:B3:53:LEU:HG | 2.27 | 0.50 |
| 30:B8:4:MET:HE1 | 31:BA:593:G:H1' | 1.94 | 0.50 |
| 30:B8:46:ARG:HH22 | 41:BP:65:ARG:HH22 | 1.54 | 0.50 |
| 31:BA:271(P):C:C2' | 31:BA:271(Q):G:H5' | 2.42 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:384:U:H2' | 31:BA:385:C:C6 | 2.44 | 0.50 |
| 31:BA:953:A:H2' | 31:BA:954:G:H5' | 1.91 | 0.50 |
| 31:BA:996:A:C4' | 46:BU:92:ARG:HE | 2.22 | 0.50 |
| 31:BA:1722:A:N6 | 31:BA:1741:A:C2 | 2.79 | 0.50 |
| 31:BA:1902:C:H2' | 31:BA:1903:G:O5' | 2.12 | 0.50 |
| 31:BA:2203:U:C1' | 33:BD:151:LYS:HE2 | 2.40 | 0.50 |
| 34:BE:153:GLY:O | 34:BE:154:LYS:C | 2.54 | 0.50 |
| 39:BN:18:ALA:HB1 | 39:BN:21:LYS:HB2 | 1.94 | 0.50 |
| 39:BN:36:GLY:H | 39:BN:42:TRP:HZ3 | 1.58 | 0.50 |
| 43:BR:20:LEU:O | 43:BR:21:TYR:C | 2.55 | 0.50 |
| 45:BT:89:VAL:HG12 | 45:BT:91:ARG:HG2 | 1.94 | 0.50 |
| 47:BV:79:VAL:CG2 | 47:BV:82:ARG:HD2 | 2.41 | 0.50 |
| 51:BZ:125:LEU:HD23 | 51:BZ:126:VAL:N | 2.27 | 0.50 |
| 1:CA:152:A:N6 | 1:CA:170:U:C2 | 2.80 | 0.50 |
| 1:CA:559:A:C5' | 1:CA:560:U:H3' | 2.41 | 0.50 |
| 1:CA:685:G:O2' | 1:CA:686:U:C5' | 2.54 | 0.50 |
| 1:CA:936:C:H2' | 1:CA:937:A:O4' | 2.11 | 0.50 |
| 1:CA:1086:U:H2' | 1:CA:1087:G:H8 | 1.76 | 0.50 |
| 5:CE:45:PHE:CE2 | 5:CE:47:LYS:HD2 | 2.46 | 0.50 |
| 6:CF:96:PRO:HB3 | 18:CR:30:ASP:OD2 | 2.11 | 0.50 |
| 8:CH:38:ILE:HD11 | 8:CH:118:VAL:O | 2.12 | 0.50 |
| 12:CL:22:SER:O | 12:CL:24:VAL:N | 2.44 | 0.50 |
| 27:D5:55:ARG:O | 27:D5:56:LYS:HG3 | 2.11 | 0.50 |
| 31:DA:528:A:N1 | 31:DA:2043:C:O5' | 2.44 | 0.50 |
| 31:DA:686:G:N2 | 31:DA:788:A:H61 | 2.09 | 0.50 |
| 31:DA:1179:C:O2 | 31:DA:1179:C:C2' | 2.54 | 0.50 |
| 31:DA:1495:A:H2' | 31:DA:1496:A:C2 | 2.46 | 0.50 |
| 31:DA:1499:C:C2' | 31:DA:1500:G:H5' | 2.42 | 0.50 |
| 31:DA:1509(A):A:H2' | 31:DA:1509(B):A:C8 | 2.47 | 0.50 |
| 31:DA:1527:G:C5' | 31:DA:1528:A:OP1 | 2.57 | 0.50 |
| 31:DA:1592:C:C2' | 31:DA:1593:G:H5' | 2.41 | 0.50 |
| 31:DA:1720:U:H2' | 31:DA:1721:G:O4' | 2.11 | 0.50 |
| 31:DA:1833:U:C2 | 31:DA:1834:U:C6 | 3.00 | 0.50 |
| 31:DA:2552:U:C2 | 31:DA:2554:U:C5' | 2.94 | 0.50 |
| 32:DB:41:U:C2' | 32:DB:42:C:OP1 | 2.59 | 0.50 |
| 36:DG:123:ASN:O | 36:DG:126:ASP:HB2 | 2.10 | 0.50 |
| 38:DI:130:TYR:O | 38:DI:131:LYS:HG3 | 2.10 | 0.50 |
| 43:DR:13:HIS:O | 43:DR:14:SER:C | 2.54 | 0.50 |
| 46:DU:31:SER:C | 46:DU:33:ARG:H | 2.19 | 0.50 |
| 48:DW:86:LEU:HD12 | 48:DW:87:PRO:N | 2.26 | 0.50 |
| 50:DY:83:THR:CG2 | 50:DY:94:LYS:HB3 | 2.41 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 51:DZ:102:LEU:HG | 51:DZ:123:ASP:HA | 1.94 | 0.50 |
| 1:AA:159:G:H21 | 1:AA:161:A:H8 | 1.60 | 0.50 |
| 1:AA:831:U:O2' | 1:AA:832:C:H5' | 2.11 | 0.50 |
| 1:AA:865:A:C2 | 1:AA:918:A:H4' | 2.47 | 0.50 |
| 1:AA:1478:C:H2' | 1:AA:1479:C:C6 | 2.47 | 0.50 |
| 2:AB:141:GLU:O | 2:AB:145:LEU:HB2 | 2.12 | 0.50 |
| 3:AC:11:ARG:O | 3:AC:12:LEU:C | 2.53 | 0.50 |
| 5:AE:36:ASP:O | 5:AE:37:ARG:HB2 | 2.11 | 0.50 |
| 12:AL:105:TYR:C | 12:AL:107:ALA:H | 2.20 | 0.50 |
| 13:AM:78:ILE:HG22 | 13:AM:93:ARG:HH22 | 1.77 | 0.50 |
| 16:AP:19:ILE:HB | 16:AP:37:GLY:O | 2.11 | 0.50 |
| 17:AQ:45:HIS:O | 17:AQ:73:VAL:HG23 | 2.11 | 0.50 |
| 20:AT:81:LYS:C | 20:AT:83:ARG:N | 2.69 | 0.50 |
| 23:B1:67:ILE:N | 23:B1:68:PRO:CD | 2.74 | 0.50 |
| 29:B7:48:LYS:HD3 | 29:B7:48:LYS:N | 2.26 | 0.50 |
| 30:B8:62:LEU:N | 30:B8:63:PRO:HD2 | 2.27 | 0.50 |
| 31:BA:11:G:C2' | 31:BA:12:U:H5' | 2.42 | 0.50 |
| 31:BA:197:A:H2' | 31:BA:198:C:C5' | 2.42 | 0.50 |
| 31:BA:528:A:C2 | 31:BA:2043:C:C5' | 2.95 | 0.50 |
| 31:BA:746:A:N3 | 55:BA:3351:ZIT:H161 | 2.26 | 0.50 |
| 31:BA:995:C:N3 | 46:BU:57:PHE:CE2 | 2.80 | 0.50 |
| 31:BA:1027:A:C2 | 31:BA:2488:A:H5' | 2.47 | 0.50 |
| 31:BA:1044:G:C2 | 31:BA:1112:G:O6 | 2.65 | 0.50 |
| 31:BA:1433:U:O2' | 31:BA:1434:A:H5' | 2.12 | 0.50 |
| 31:BA:1495:A:C2 | 31:BA:1496:A:C2 | 2.98 | 0.50 |
| 31:BA:1563:G:C5 | 31:BA:1564:C:C5 | 3.00 | 0.50 |
| 31:BA:1578:U:O2 | 31:BA:1578:U:H2' | 2.11 | 0.50 |
| 31:BA:1861:G:C2 | 31:BA:1862:G:C8 | 3.00 | 0.50 |
| 31:BA:1952:A:C5 | 40:BO:22:ILE:CD1 | 2.95 | 0.50 |
| 31:BA:2795:G:N2 | 31:BA:2796:U:O2' | 2.45 | 0.50 |
| 36:BG:45:GLU:HB2 | 36:BG:47:LYS:HG3 | 1.94 | 0.50 |
| 36:BG:123:ASN:O | 36:BG:126:ASP:HB2 | 2.11 | 0.50 |
| 41:BP:82:GLY:HA2 | 41:BP:113:LYS:O | 2.12 | 0.50 |
| 46:BU:28:ARG:HG2 | 46:BU:38:THR:OG1 | 2.11 | 0.50 |
| 49:BX:18:TYR:O | 49:BX:19:ALA:C | 2.55 | 0.50 |
| 49:BX:82:GLN:NE2 | 49:BX:83:VAL:HG22 | 2.25 | 0.50 |
| 49:BX:93:GLU:O | 49:BX:93:GLU:HG3 | 2.12 | 0.50 |
| 50:BY:28:LYS:C | 50:BY:38:ILE:HB | 2.35 | 0.50 |
| 1:CA:102:G:N3 | 1:CA:103:C:C6 | 2.79 | 0.50 |
| 1:CA:354:G:C6 | 1:CA:355:C:N4 | 2.80 | 0.50 |
| 1:CA:923:A:O2' | 1:CA:1399:C:OP2 | 2.27 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:1081:G:N2 | 1:CA:1082:G:H1' | 2.26 | 0.50 |
| 1:CA:1128:C:H5' | 9:CI:16:ARG:CZ | 2.42 | 0.50 |
| 1:CA:1233:G:H2' | 1:CA:1234:C:C6 | 2.47 | 0.50 |
| 1:CA:1490:C:H2' | 1:CA:1491:G:O4' | 2.11 | 0.50 |
| 2:CB:21:ARG:O | 2:CB:21:ARG:HG3 | 2.09 | 0.50 |
| 2:CB:141:GLU:O | 2:CB:145:LEU:HB2 | 2.12 | 0.50 |
| 3:CC:11:ARG:O | 3:CC:14:ILE:O | 2.30 | 0.50 |
| 8:CH:5:PRO:O | 8:CH:8:ASP:HB3 | 2.12 | 0.50 |
| 10:CJ:33:GLN:O | 10:CJ:75:ILE:HG23 | 2.12 | 0.50 |
| 11:CK:125:PHE:N | 11:CK:125:PHE:HD1 | 2.09 | 0.50 |
| 22:D0:13:GLY:O | 22:D0:14:ARG:CB | 2.40 | 0.50 |
| 23:D1:33:LYS:HB3 | 31:DA:2395:C:O2' | 2.11 | 0.50 |
| 24:D2:22:GLU:O | 24:D2:23:LYS:C | 2.55 | 0.50 |
| 28:D6:40:CYS:SG | 28:D6:45:LYS:HD2 | 2.52 | 0.50 |
| 30:D8:2:PRO:N | 31:DA:591:C:O2 | 2.45 | 0.50 |
| 30:D8:39:LYS:HE3 | 30:D8:42:ARG:NH1 | 2.27 | 0.50 |
| 31:DA:139(A):G:H2' | 31:DA:140:G:O5' | 2.11 | 0.50 |
| 31:DA:197:A:H2' | 31:DA:198:C:O5' | 2.12 | 0.50 |
| 31:DA:926:A:H5" | 31:DA:926:A:H8 | 1.77 | 0.50 |
| 31:DA:1225:G:H8 | 31:DA:1225:G:O5' | 1.94 | 0.50 |
| 31:DA:1380:G:N2 | 31:DA:1570:A:C2 | 2.79 | 0.50 |
| 31:DA:1517:G:C2' | 31:DA:1518:U:H5' | 2.42 | 0.50 |
| 31:DA:1637:A:H4' | 31:DA:2711:A:O2' | 2.11 | 0.50 |
| 31:DA:1664:A:N6 | 31:DA:1665:A:N6 | 2.59 | 0.50 |
| 31:DA:1813:G:H2' | 31:DA:1814:G:H5' | 1.92 | 0.50 |
| 31:DA:2395:C:H5' | 31:DA:2395:C:C6 | 2.35 | 0.50 |
| 31:DA:2662:A:H4' | 31:DA:2663:G:C8 | 2.46 | 0.50 |
| 31:DA:2680:C:H5' | 34:DE:189:PRO:HA | 1.92 | 0.50 |
| 35:DF:65:TRP:CZ3 | 35:DF:73:ALA:O | 2.63 | 0.50 |
| 36:DG:52:ILE:HG22 | 36:DG:54:GLU:HG3 | 1.92 | 0.50 |
| 36:DG:57:ALA:O | 36:DG:60:LEU:HB3 | 2.12 | 0.50 |
| 44:DS:61:ASN:ND2 | 44:DS:64:GLU:OE2 | 2.44 | 0.50 |
| 1:AA:66:G:C6 | 1:AA:67:C:C5 | 3.00 | 0.50 |
| 1:AA:264:U:O2' | 17:AQ:64:PRO:HB2 | 2.12 | 0.50 |
| 1:AA:340:U:H2' | 1:AA:341:C:O4' | 2.11 | 0.50 |
| 1:AA:963:G:H21 | 10:AJ:55:LYS:HD3 | 1.75 | 0.50 |
| 3:AC:19:GLU:HG2 | 3:AC:19:GLU:O | 2.12 | 0.50 |
| 4:AD:150:GLU:HG2 | 4:AD:151:LYS:H | 1.77 | 0.50 |
| 8:AH:80:ILE:O | 8:AH:80:ILE:HG22 | 2.12 | 0.50 |
| 11:AK:69:ALA:O | 11:AK:73:MET:HG2 | 2.12 | 0.50 |
| 12:AL:32:PHE:CE1 | 12:AL:86:ARG:HG3 | 2.47 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 12:AL:60:LEU:HD22 | 12:AL:60:LEU:H | 1.77 | 0.50 |
| 16:AP:8:ARG:HG2 | 16:AP:9:PHE:H | 1.77 | 0.50 |
| 23:B1:44:PRO:HA | 31:BA:2231:C:OP1 | 2.12 | 0.50 |
| 31:BA:470:A:OP1 | 35:BF:59:TYR:HE2 | 1.95 | 0.50 |
| 31:BA:558:G:OP1 | 39:BN:111:PRO:HD2 | 2.12 | 0.50 |
| 31:BA:607:U:H3 | 31:BA:621:A:H2 | 1.50 | 0.50 |
| 31:BA:614:U:O2 | 31:BA:614:U:O4' | 2.29 | 0.50 |
| 31:BA:958:U:OP2 | 42:BQ:14:ARG:NH1 | 2.44 | 0.50 |
| 31:BA:1171:G:H3' | 31:BA:1173:G:O4' | 2.12 | 0.50 |
| 31:BA:1291:C:O2' | 31:BA:1292:U:H5' | 2.12 | 0.50 |
| 31:BA:1531:C:H3' | 31:BA:1532:C:C4' | 2.42 | 0.50 |
| 31:BA:1899:G:N2 | 31:BA:1902:C:H5 | 1.95 | 0.50 |
| 31:BA:1952:A:C6 | 40:BO:22:ILE:HD12 | 2.47 | 0.50 |
| 31:BA:2186:G:H3' | 31:BA:2187:G:H5" | 1.93 | 0.50 |
| 31:BA:2186:G:C2' | 31:BA:2187:G:H5" | 2.41 | 0.50 |
| 31:BA:2309:A:N3 | 31:BA:2310:A:H2 | 2.09 | 0.50 |
| 31:BA:2808:U:O2' | 31:BA:2809:A:H5' | 2.12 | 0.50 |
| 32:BB:13:A:N1 | 32:BB:69:G:O2' | 2.43 | 0.50 |
| 33:BD:85:ASP:OD2 | 33:BD:88:ARG:HD2 | 2.12 | 0.50 |
| 33:BD:248:SER:C | 33:BD:250:TRP:N | 2.69 | 0.50 |
| 34:BE:134:ILE:H | 34:BE:134:ILE:CD1 | 2.24 | 0.50 |
| 34:BE:143:ASN:HB2 | 34:BE:147:PRO:HD2 | 1.93 | 0.50 |
| 36:BG:57:ALA:O | 36:BG:60:LEU:HB3 | 2.11 | 0.50 |
| 38:BI:88:ILE:HG22 | 38:BI:89:TYR:N | 2.27 | 0.50 |
| 39:BN:58:ASP:C | 39:BN:60:ILE:H | 2.19 | 0.50 |
| 46:BU:33:ARG:O | 46:BU:36:ARG:N | 2.45 | 0.50 |
| 48:BW:17:VAL:HG22 | 48:BW:76:VAL:HG11 | 1.92 | 0.50 |
| 49:BX:33:LYS:CA | 49:BX:35:THR:HG22 | 2.42 | 0.50 |
| 50:BY:88:LYS:O | 50:BY:89:PHE:CB | 2.60 | 0.50 |
| 1:CA:367:U:O2 | 1:CA:369:C:C6 | 2.65 | 0.50 |
| 1:CA:373:A:C2 | 1:CA:482:A:N6 | 2.80 | 0.50 |
| 1:CA:1148:U:H2' | 1:CA:1149:C:O4' | 2.11 | 0.50 |
| 1:CA:1385:G:O2' | 1:CA:1386:G:H5' | 2.11 | 0.50 |
| 1:CA:1478:C:H2' | 1:CA:1479:C:C6 | 2.47 | 0.50 |
| 4:CD:52:SER:O | 4:CD:54:TYR:N | 2.45 | 0.50 |
| 4:CD:133:VAL:HG12 | 4:CD:135:LEU:H | 1.75 | 0.50 |
| 19:CS:79:THR:O | 19:CS:80:TYR:HB3 | 2.12 | 0.50 |
| 23:D1:94:LEU:CD2 | 23:D1:95:LEU:N | 2.75 | 0.50 |
| 27:D5:25:LEU:HD12 | 48:DW:19:LEU:O | 2.11 | 0.50 |
| 31:DA:624:C:C2' | 31:DA:625:G:H5' | 2.42 | 0.50 |
| 31:DA:743:G:H2' | 31:DA:744:G:H5' | 1.94 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:878:A:C6 | 31:DA:900:A:C8 | 3.00 | 0.50 |
| 31:DA:1648:C:C2' | 31:DA:1649:G:O5' | 2.59 | 0.50 |
| 31:DA:1719:G:H2' | 31:DA:1720:U:C5' | 2.40 | 0.50 |
| 31:DA:1788:C:H2' | 31:DA:1789:A:H8 | 1.77 | 0.50 |
| 31:DA:1797:C:H4' | 33:DD:257:LEU:O | 2.12 | 0.50 |
| 31:DA:1899:G:N2 | 31:DA:1902:C:C4 | 2.79 | 0.50 |
| 31:DA:2309:A:N3 | 31:DA:2310:A:H2 | 2.09 | 0.50 |
| 31:DA:2496:C:OP1 | 42:DQ:81:VAL:HG12 | 2.11 | 0.50 |
| 35:DF:34:TRP:CZ2 | 41:DP:12:ALA:HB2 | 2.47 | 0.50 |
| 35:DF:51:THR:OG1 | 35:DF:91:GLY:HA3 | 2.11 | 0.50 |
| 35:DF:53:THR:HG23 | 35:DF:55:GLY:N | 2.26 | 0.50 |
| 36:DG:16:ARG:NE | 36:DG:31:VAL:HG11 | 2.27 | 0.50 |
| 36:DG:51:ARG:HB3 | 36:DG:53:LEU:HD23 | 1.94 | 0.50 |
| 39:DN:9:VAL:CG1 | 39:DN:39:ARG:HH22 | 2.20 | 0.50 |
| 42:DQ:20:ALA:C | 42:DQ:22:LYS:N | 2.70 | 0.50 |
| 43:DR:96:ARG:HH21 | 43:DR:117:VAL:HG23 | 1.77 | 0.50 |
| 48:DW:24:ILE:HD12 | 48:DW:24:ILE:O | 2.12 | 0.50 |
| 48:DW:59:VAL:HG12 | 48:DW:60:ASN:H | 1.74 | 0.50 |
| 48:DW:64:MET:O | 48:DW:65:LEU:CB | 2.51 | 0.50 |
| 1:AA:445:G:C2 | 1:AA:446:G:C4 | 3.00 | 0.50 |
| 1:AA:473:G:H5' | 16:AP:81:ARG:HG3 | 1.93 | 0.50 |
| 1:AA:938:A:N6 | 1:AA:939:G:C6 | 2.80 | 0.50 |
| 1:AA:987:G:N2 | 1:AA:1219:U:C2 | 2.80 | 0.50 |
| 1:AA:1412:C:C2 | 1:AA:1489:G:N2 | 2.80 | 0.50 |
| 2:AB:19:HIS:CE1 | 2:AB:206:ASP:HB2 | 2.46 | 0.50 |
| 6:AF:18:GLN:HA | 6:AF:21:LEU:HD23 | 1.93 | 0.50 |
| 12:AL:119:LYS:O | 12:AL:120:TYR:HB2 | 2.10 | 0.50 |
| 13:AM:54:VAL:O | 13:AM:58:GLU:HG2 | 2.12 | 0.50 |
| 15:AO:54:ARG:O | 15:AO:57:LEU:HB2 | 2.11 | 0.50 |
| 19:AS:48:THR:HG22 | 19:AS:61:TYR:HA | 1.94 | 0.50 |
| 22:B0:43:THR:HG22 | 31:BA:2331:G:O2' | 2.10 | 0.50 |
| 23:B1:34:THR:HG21 | 31:BA:388:G:OP1 | 2.11 | 0.50 |
| 24:B2:25:VAL:C | 24:B2:27:GLU:H | 2.20 | 0.50 |
| 28:B6:39:TYR:HE1 | 31:BA:2347:C:HO2' | 1.58 | 0.50 |
| 31:BA:271(H):G:O6 | 31:BA:271(Q):G:C6 | 2.65 | 0.50 |
| 31:BA:848:G:H2' | 31:BA:849:A:H8 | 1.77 | 0.50 |
| 31:BA:855:G:C6 | 31:BA:856:C:N4 | 2.80 | 0.50 |
| 31:BA:1257:C:H4' | 35:BF:83:PHE:CE2 | 2.47 | 0.50 |
| 31:BA:1465:G:C4 | 31:BA:1466:G:C8 | 2.99 | 0.50 |
| 31:BA:1946:U:H2' | 31:BA:1947:C:C6 | 2.44 | 0.50 |
| 31:BA:2098:U:H2' | 31:BA:2099:U:H6 | 1.77 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2301:C:H2' | 31:BA:2302:G:O4' | 2.12 | 0.50 |
| 31:BA:2315:G:H2' | 31:BA:2316:C:H6 | 1.74 | 0.50 |
| 31:BA:2584:U:C2' | 31:BA:2585:U:H5' | 2.41 | 0.50 |
| 31:BA:2842:G:O2' | 31:BA:2843:G:H5' | 2.11 | 0.50 |
| 32:BB:71:C:O2 | 32:BB:71:C:C2' | 2.56 | 0.50 |
| 32:BB:110:G:N1 | 32:BB:111:G:C5 | 2.79 | 0.50 |
| 36:BG:120:LEU:HB2 | 36:BG:179:PRO:O | 2.11 | 0.50 |
| 39:BN:56:ASN:H | 39:BN:125:GLY:HA3 | 1.77 | 0.50 |
| 51:BZ:56:VAL:HG12 | 51:BZ:57:ILE:N | 2.26 | 0.50 |
| 1:CA:112:G:H2' | 1:CA:112:G:N3 | 2.27 | 0.50 |
| 1:CA:174:C:C5 | 1:CA:175:C:H5 | 2.29 | 0.50 |
| 1:CA:445:G:N3 | 1:CA:446:G:C8 | 2.80 | 0.50 |
| 1:CA:791:G:C5 | 1:CA:792:A:N7 | 2.79 | 0.50 |
| 1:CA:1288:A:H2 | 1:CA:1352:C:O2 | 1.94 | 0.50 |
| 4:CD:58:LEU:HD22 | 4:CD:62:GLN:HG2 | 1.93 | 0.50 |
| 4:CD:105:VAL:HG22 | 4:CD:146:ILE:HG21 | 1.93 | 0.50 |
| 5:CE:131:ILE:O | 5:CE:134:ALA:HB3 | 2.12 | 0.50 |
| 7:CG:25:ALA:O | 7:CG:29:LYS:HG2 | 2.12 | 0.50 |
| 8:CH:26:VAL:HG13 | 8:CH:59:LEU:HB2 | 1.93 | 0.50 |
| 9:CI:10:ARG:HG2 | 9:CI:104:ARG:O | 2.11 | 0.50 |
| 9:CI:48:GLU:N | 9:CI:49:PRO:CD | 2.75 | 0.50 |
| 13:CM:61:GLU:HA | 13:CM:66:LEU:HD11 | 1.93 | 0.50 |
| 24:D2:25:VAL:C | 24:D2:27:GLU:H | 2.20 | 0.50 |
| 31:DA:17:G:H2' | 31:DA:18:C:H6 | 1.75 | 0.50 |
| 31:DA:452:G:C4 | 31:DA:458:G:C6 | 3.00 | 0.50 |
| 31:DA:768:G:C6 | 31:DA:769:G:C5 | 3.00 | 0.50 |
| 31:DA:786:C:C2' | 31:DA:787:U:H5' | 2.41 | 0.50 |
| 31:DA:979:G:H3' | 31:DA:980:A:C5' | 2.42 | 0.50 |
| 31:DA:1525:G:H2' | 31:DA:1526:G:C8 | 2.47 | 0.50 |
| 31:DA:2808:U:H2' | 31:DA:2809:A:C5' | 2.41 | 0.50 |
| 33:DD:35:LYS:NZ | 33:DD:104:TYR:HD1 | 2.05 | 0.50 |
| 36:DG:6:ALA:HB3 | 36:DG:104:GLU:OE1 | 2.11 | 0.50 |
| 37:DH:70:THR:O | 37:DH:71:LEU:C | 2.55 | 0.50 |
| 38:DI:109:ILE:CG2 | 38:DI:130:TYR:OH | 2.60 | 0.50 |
| 39:DN:131:GLN:OE1 | 39:DN:134:ARG:HB3 | 2.11 | 0.50 |
| 46:DU:112:ARG:HH11 | 46:DU:112:ARG:HG3 | 1.76 | 0.50 |
| 49:DX:36:LYS:C | 49:DX:38:GLU:H | 2.20 | 0.50 |
| 51:DZ:54:HIS:HE1 | 51:DZ:123:ASP:OD2 | 1.95 | 0.50 |
| 1:AA:59:A:H3' | 1:AA:331:G:H22 | 1.77 | 0.49 |
| 1:AA:67:C:H2' | 1:AA:68:G:C8 | 2.47 | 0.49 |
| 1:AA:143:A:N1 | 1:AA:220:G:O6 | 2.44 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:414:A:C5 | 1:AA:431:A:C2 | 2.99 | 0.49 |
| 1:AA:685:G:C2 | 1:AA:686:U:C4 | 3.00 | 0.49 |
| 1:AA:754:C:H3' | 1:AA:754:C:O2 | 2.11 | 0.49 |
| 1:AA:939:G:H5" | 7:AG:102:ARG:NH1 | 2.26 | 0.49 |
| 1:AA:1252:A:H2' | 1:AA:1253:G:O4' | 2.12 | 0.49 |
| 2:AB:61:LEU:HD23 | 2:AB:68:ILE:HD11 | 1.94 | 0.49 |
| 3:AC:66:VAL:HG11 | 3:AC:91:LEU:HD13 | 1.95 | 0.49 |
| 4:AD:11:LEU:C | 4:AD:13:ARG:N | 2.64 | 0.49 |
| 8:AH:8:ASP:O | 8:AH:9:MET:C | 2.55 | 0.49 |
| 10:AJ:45:ARG:HB2 | 10:AJ:65:LEU:HB3 | 1.92 | 0.49 |
| 11:AK:62:GLN:O | 11:AK:63:LEU:C | 2.55 | 0.49 |
| 18:AR:36:ASN:HD22 | 18:AR:39:VAL:HG21 | 1.76 | 0.49 |
| 24:B2:39:ALA:O | 24:B2:40:SER:C | 2.54 | 0.49 |
| 24:B2:41:ILE:HG21 | 31:BA:95:G:H21 | 1.77 | 0.49 |
| 28:B6:13:CYS:HA | 28:B6:50:ARG:O | 2.11 | 0.49 |
| 31:BA:9:U:C5 | 31:BA:2629:A:N6 | 2.79 | 0.49 |
| 31:BA:494:G:C5' | 31:BA:494:G:C8 | 2.95 | 0.49 |
| 31:BA:662:G:OP1 | 41:BP:18:ARG:NH1 | 2.44 | 0.49 |
| 31:BA:794:G:H2' | 31:BA:795:C:C6 | 2.47 | 0.49 |
| 31:BA:971:C:H2' | 31:BA:972:G:H5' | 1.94 | 0.49 |
| 31:BA:1517:G:C2' | 31:BA:1518:U:H5' | 2.42 | 0.49 |
| 31:BA:1545:A:H2' | 31:BA:1546:C:C5' | 2.42 | 0.49 |
| 31:BA:1885:A:H2' | 31:BA:1886:C:O4' | 2.10 | 0.49 |
| 31:BA:2360:A:O2' | 31:BA:2361:A:O5' | 2.30 | 0.49 |
| 31:BA:2564:A:C2 | 31:BA:2647:U:H4' | 2.46 | 0.49 |
| 35:BF:34:TRP:CZ2 | 41:BP:12:ALA:HB2 | 2.46 | 0.49 |
| 47:BV:82:ARG:HG2 | 47:BV:82:ARG:NH1 | 2.16 | 0.49 |
| 49:BX:63:LYS:O | 49:BX:68:ARG:HA | 2.12 | 0.49 |
| 49:BX:92:LEU:O | 49:BX:93:GLU:HB3 | 2.12 | 0.49 |
| 50:BY:26:LYS:HG2 | 50:BY:27:VAL:H | 1.76 | 0.49 |
| 50:BY:27:VAL:HG12 | 50:BY:29:GLU:OE1 | 2.11 | 0.49 |
| 51:BZ:63:ASP:O | 51:BZ:65:GLN:N | 2.44 | 0.49 |
| 1:CA:607:A:H2' | 1:CA:608:A:O4' | 2.12 | 0.49 |
| 1:CA:865:A:C2 | 1:CA:918:A:H4' | 2.47 | 0.49 |
| 1:CA:1272:G:C6 | 1:CA:1273:G:C5 | 3.00 | 0.49 |
| 1:CA:1505:G:H5" | 1:CA:1506:U:OP1 | 2.12 | 0.49 |
| 4:CD:79:PHE:CD2 | 4:CD:79:PHE:C | 2.90 | 0.49 |
| 7:CG:146:GLU:HA | 7:CG:149:ARG:HB2 | 1.92 | 0.49 |
| 8:CH:21:LYS:O | 8:CH:22:GLU:C | 2.55 | 0.49 |
| 12:CL:83:VAL:HG13 | 12:CL:100:ILE:HG23 | 1.92 | 0.49 |
| 12:CL:93:LEU:O | 12:CL:94:PRO:C | 2.55 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 18:CR:52:PRO:O | 18:CR:56:THR:HG23 | 2.12 | 0.49 |
| 19:CS:48:THR:HG22 | 19:CS:61:TYR:HA | 1.94 | 0.49 |
| 20:CT:73:HIS:HB3 | 20:CT:74:LYS:HG2 | 1.94 | 0.49 |
| 23:D1:19:GLN:CD | 23:D1:44:PRO:CB | 2.85 | 0.49 |
| 23:D1:89:GLU:O | 23:D1:90:ILE:C | 2.54 | 0.49 |
| 24:D2:17:SER:O | 24:D2:20:GLU:HB2 | 2.12 | 0.49 |
| 31:DA:146:G:C5' | 31:DA:146:G:H8 | 2.25 | 0.49 |
| 31:DA:271(P):C:C2' | 31:DA:271(Q):G:H5' | 2.42 | 0.49 |
| 31:DA:481:G:OP2 | 50:DY:47:LYS:CE | 2.60 | 0.49 |
| 31:DA:864:G:C6 | 31:DA:865:C:N4 | 2.80 | 0.49 |
| 31:DA:884:C:O2' | 31:DA:892:G:C8 | 2.56 | 0.49 |
| 31:DA:1276:A:O2' | 43:DR:16:HIS:HE1 | 1.94 | 0.49 |
| 31:DA:1531:C:H3' | 31:DA:1532:C:C4' | 2.42 | 0.49 |
| 31:DA:2364:C:H2' | 31:DA:2365:G:O4' | 2.11 | 0.49 |
| 31:DA:2418:A:H2' | 31:DA:2419:U:H6 | 1.77 | 0.49 |
| 33:DD:35:LYS:HE2 | 33:DD:65:ILE:HG22 | 1.94 | 0.49 |
| 33:DD:89:SER:HB2 | 33:DD:159:ALA:HB2 | 1.94 | 0.49 |
| 34:DE:153:GLY:O | 34:DE:154:LYS:C | 2.55 | 0.49 |
| 34:DE:182:LEU:HD12 | 34:DE:182:LEU:C | 2.37 | 0.49 |
| 38:DI:94:ALA:CB | 38:DI:114:LEU:HD12 | 2.41 | 0.49 |
| 41:DP:144:GLU:N | 41:DP:145:PRO:CD | 2.75 | 0.49 |
| 42:DQ:20:ALA:CB | 42:DQ:99:PRO:HG2 | 2.42 | 0.49 |
| 43:DR:61:HIS:O | 43:DR:62:ALA:C | 2.55 | 0.49 |
| 45:DT:109:GLU:CA | 45:DT:112:ARG:HG3 | 2.40 | 0.49 |
| 48:DW:5:ALA:C | 48:DW:6:ILE:HG13 | 2.37 | 0.49 |
| 49:DX:36:LYS:C | 49:DX:38:GLU:N | 2.70 | 0.49 |
| 50:DY:27:VAL:HG12 | 50:DY:29:GLU:N | 2.18 | 0.49 |
| 50:DY:27:VAL:CB | 50:DY:29:GLU:OE1 | 2.59 | 0.49 |
| 1:AA:41:G:H2' | 1:AA:42:G:H8 | 1.77 | 0.49 |
| 1:AA:59:A:H1' | 1:AA:354:G:N2 | 2.28 | 0.49 |
| 1:AA:93:G:C6 | 1:AA:96:U:C4 | 3.00 | 0.49 |
| 1:AA:109:A:C6 | 1:AA:326:G:C6 | 3.00 | 0.49 |
| 1:AA:913:A:H4' | 1:AA:914:A:O5' | 2.12 | 0.49 |
| 1:AA:1086:U:H2' | 1:AA:1087:G:H8 | 1.77 | 0.49 |
| 1:AA:1148:U:H2' | 1:AA:1149:C:O4' | 2.11 | 0.49 |
| 1:AA:1160:G:N2 | 1:AA:1161:C:C6 | 2.80 | 0.49 |
| 1:AA:1452:C:H4' | 1:AA:1456:G:N3 | 2.27 | 0.49 |
| 5:AE:13:ILE:HA | 5:AE:29:GLY:O | 2.13 | 0.49 |
| 8:AH:69:ARG:HD3 | 8:AH:75:ARG:O | 2.12 | 0.49 |
| 9:AI:55:ALA:CB | 9:AI:58:ARG:HD2 | 2.41 | 0.49 |
| 19:AS:79:THR:O | 19:AS:80:TYR:HB3 | 2.12 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 20:AT:58:LYS:HG3 | 20:AT:62:LEU:HD12 | 1.94 | 0.49 |
| 23:B1:94:LEU:O | 23:B1:95:LEU:HG | 2.12 | 0.49 |
| 31:BA:82:G:O2' | 31:BA:83:G:H5' | 2.12 | 0.49 |
| 31:BA:272(D):G:H1 | 31:BA:364:C:H42 | 1.59 | 0.49 |
| 31:BA:814:C:N4 | 41:BP:27:HIS:CD2 | 2.77 | 0.49 |
| 31:BA:1568:G:H5' | 33:BD:60:ARG:HA | 1.94 | 0.49 |
| 31:BA:1747(A):G:C2' | 31:BA:1748:G:H5" | 2.42 | 0.49 |
| 31:BA:1836:C:H2' | 31:BA:1837:C:H6 | 1.77 | 0.49 |
| 31:BA:2205:C:O2 | 31:BA:2220:G:C2 | 2.65 | 0.49 |
| 31:BA:2286:A:O2' | 31:BA:2286:A:H8 | 1.95 | 0.49 |
| 31:BA:2309:A:C2 | 31:BA:2310:A:H2 | 2.29 | 0.49 |
| 31:BA:2836:U:C4 | 31:BA:2883:A:N6 | 2.80 | 0.49 |
| 32:BB:28:C:H2' | 32:BB:29:A:H8 | 1.77 | 0.49 |
| 33:BD:178:PRO:O | 33:BD:179:SER:C | 2.55 | 0.49 |
| 37:BH:19:VAL:HG21 | 37:BH:44:VAL:HA | 1.95 | 0.49 |
| 37:BH:43:VAL:CG1 | 37:BH:53:GLU:H | 2.25 | 0.49 |
| 46:BU:108:GLU:O | 46:BU:109:LEU:C | 2.53 | 0.49 |
| 47:BV:35:LEU:HB2 | 47:BV:59:ALA:HB1 | 1.93 | 0.49 |
| 50:BY:17:SER:HB2 | 50:BY:71:LYS:HE2 | 1.94 | 0.49 |
| 1:CA:328:C:O2 | 1:CA:328:C:C2' | 2.60 | 0.49 |
| 1:CA:473:G:H2' | 1:CA:474:G:H8 | 1.77 | 0.49 |
| 1:CA:833:U:H2' | 1:CA:834:C:C6 | 2.47 | 0.49 |
| 2:CB:97:TRP:CZ3 | 2:CB:173:ALA:HA | 2.46 | 0.49 |
| 3:CC:150:LYS:HE3 | 3:CC:167:TRP:HE1 | 1.77 | 0.49 |
| 4:CD:150:GLU:HG2 | 4:CD:151:LYS:H | 1.77 | 0.49 |
| 4:CD:202:LEU:N | 4:CD:202:LEU:HD23 | 2.27 | 0.49 |
| 16:CP:14:ASN:N | 16:CP:15:PRO:HD3 | 2.27 | 0.49 |
| 24:D2:27:GLU:O | 24:D2:30:ARG:HG2 | 2.11 | 0.49 |
| 27:D5:36:CYS:C | 27:D5:38:ALA:H | 2.20 | 0.49 |
| 30:D8:2:PRO:O | 30:D8:3:LYS:C | 2.55 | 0.49 |
| 30:D8:35:GLN:OE1 | 31:DA:2421:G:OP2 | 2.29 | 0.49 |
| 31:DA:309:G:O3' | 50:DY:18:GLY:CA | 2.60 | 0.49 |
| 31:DA:389:G:H22 | 41:DP:72:PRO:HD3 | 1.76 | 0.49 |
| 31:DA:460:A:C2 | 31:DA:470:A:C4 | 3.00 | 0.49 |
| 31:DA:613:G:N2 | 31:DA:614(C):A:O2' | 2.45 | 0.49 |
| 31:DA:767:U:O2' | 31:DA:768:G:H5' | 2.12 | 0.49 |
| 31:DA:848:G:H2' | 31:DA:849:A:H8 | 1.74 | 0.49 |
| 31:DA:1451:C:N3 | 31:DA:1459:G:O6 | 2.44 | 0.49 |
| 31:DA:2186:G:C2' | 31:DA:2187:G:H5" | 2.42 | 0.49 |
| 31:DA:2286:A:O2' | 31:DA:2286:A:H8 | 1.94 | 0.49 |
| 31:DA:2723:C:H5" | 43:DR:2:ARG:HD2 | 1.93 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2808:U:N3 | 31:DA:2892:A:C5 | 2.79 | 0.49 |
| 32:DB:7:G:H3' | 32:DB:8:U:C5' | 2.35 | 0.49 |
| 32:DB:31:C:O2 | 32:DB:52:A:H2 | 1.95 | 0.49 |
| 33:DD:233:HIS:N | 33:DD:233:HIS:CD2 | 2.81 | 0.49 |
| 33:DD:243:GLY:O | 33:DD:244:ARG:HB3 | 2.10 | 0.49 |
| 34:DE:60:ASN:N | 34:DE:60:ASN:HD22 | 2.08 | 0.49 |
| 34:DE:70:ALA:O | 34:DE:72:VAL:C | 2.55 | 0.49 |
| 35:DF:84:VAL:C | 35:DF:86:GLY:N | 2.69 | 0.49 |
| 36:DG:119:GLY:H | 36:DG:181:ARG:HH21 | 1.59 | 0.49 |
| 38:DI:79:ILE:CG2 | 38:DI:81:VAL:HG23 | 2.41 | 0.49 |
| 41:DP:92:GLU:HA | 41:DP:123:LEU:HD22 | 1.94 | 0.49 |
| 50:DY:8:LYS:HD2 | 50:DY:8:LYS:N | 2.26 | 0.49 |
| 1:AA:16:A:O2' | 1:AA:17:U:H5' | 2.12 | 0.49 |
| 1:AA:173:U:C2 | 1:AA:197:A:N1 | 2.80 | 0.49 |
| 1:AA:246:A:C2 | 1:AA:282:A:C5 | 3.00 | 0.49 |
| 1:AA:352:C:O2' | 1:AA:354:G:OP1 | 2.15 | 0.49 |
| 1:AA:883:C:C2' | 1:AA:884:U:H5' | 2.41 | 0.49 |
| 4:AD:119:GLN:CG | 4:AD:123:HIS:HD2 | 2.22 | 0.49 |
| 18:AR:53:ARG:C | 18:AR:55:ARG:N | 2.69 | 0.49 |
| 18:AR:72:ARG:O | 18:AR:76:LEU:HD23 | 2.11 | 0.49 |
| 31:BA:271(G):C:O2 | 31:BA:271(G):C:H2' | 2.12 | 0.49 |
| 31:BA:817:C:O2' | 31:BA:839:U:H5" | 2.13 | 0.49 |
| 31:BA:1131:G:OP2 | 31:BA:2515:C:H4' | 2.12 | 0.49 |
| 31:BA:1410:G:C5 | 31:BA:1411:C:C5 | 3.00 | 0.49 |
| 31:BA:1503:U:H3' | 31:BA:1503:U:H6 | 1.77 | 0.49 |
| 31:BA:1694:C:O2' | 31:BA:1695:G:C5 | 2.64 | 0.49 |
| 31:BA:2543:G:H8 | 31:BA:2543:G:H5' | 1.76 | 0.49 |
| 31:BA:2586:C:C5 | 31:BA:2608:G:N2 | 2.80 | 0.49 |
| 31:BA:2637:U:H2' | 31:BA:2638:G:C5' | 2.43 | 0.49 |
| 31:BA:2808:U:N3 | 31:BA:2892:A:C6 | 2.80 | 0.49 |
| 32:BB:7:G:H3' | 32:BB:8:U:C5' | 2.37 | 0.49 |
| 32:BB:31:C:H2' | 32:BB:32:C:H6 | 1.77 | 0.49 |
| 32:BB:58:A:H5' | 32:BB:59:A:OP2 | 2.12 | 0.49 |
| 33:BD:233:HIS:CD2 | 33:BD:233:HIS:H | 2.29 | 0.49 |
| 34:BE:201:THR:HG22 | 34:BE:203:LYS:N | 2.27 | 0.49 |
| 35:BF:199:TRP:CH2 | 35:BF:203:GLN:NE2 | 2.80 | 0.49 |
| 37:BH:158:HIS:HE2 | 37:BH:169:VAL:C | 2.21 | 0.49 |
| 37:BH:158:HIS:CD2 | 37:BH:170:ARG:HA | 2.47 | 0.49 |
| 41:BP:14:LYS:H | 41:BP:14:LYS:HD2 | 1.76 | 0.49 |
| 42:BQ:141:GLN:HG2 | 51:BZ:71:VAL:O | 2.12 | 0.49 |
| 48:BW:5:ALA:C | 48:BW:6:ILE:HG13 | 2.37 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 50:BY:45:VAL:HG13 | 50:BY:62:GLU:OE2 | 2.13 | 0.49 |
| 1:CA:445:G:C2 | 1:CA:446:G:C4 | 3.00 | 0.49 |
| 1:CA:585:G:N3 | 1:CA:879:C:H4' | 2.27 | 0.49 |
| 1:CA:674:G:O2' | 1:CA:675:A:H5' | 2.12 | 0.49 |
| 1:CA:692:U:O2' | 1:CA:694:A:N7 | 2.36 | 0.49 |
| 1:CA:797:C:O2' | 1:CA:798:G:H5' | 2.12 | 0.49 |
| 1:CA:1388:C:H2' | 1:CA:1389:C:C6 | 2.46 | 0.49 |
| 6:CF:10:LEU:N | 6:CF:10:LEU:CD1 | 2.75 | 0.49 |
| 13:CM:66:LEU:O | 13:CM:70:LEU:HB2 | 2.12 | 0.49 |
| 16:CP:81:ARG:C | 16:CP:82:GLN:HE21 | 2.20 | 0.49 |
| 27:D5:57:VAL:CG2 | 27:D5:58:LEU:H | 2.19 | 0.49 |
| 31:DA:150:C:H2' | 31:DA:151:C:C6 | 2.47 | 0.49 |
| 31:DA:662:G:OP1 | 41:DP:18:ARG:NH1 | 2.43 | 0.49 |
| 31:DA:993:G:C5' | 47:DV:75:PHE:CE2 | 2.90 | 0.49 |
| 31:DA:1171:G:H3' | 31:DA:1173:G:O4' | 2.12 | 0.49 |
| 31:DA:1377:G:H8 | 31:DA:1377:G:O5' | 1.95 | 0.49 |
| 31:DA:1628:G:H2' | 31:DA:1629:U:C6 | 2.47 | 0.49 |
| 31:DA:1635:G:H5' | 31:DA:1635:G:C8 | 2.47 | 0.49 |
| 31:DA:1771:C:H1' | 31:DA:1786:A:C8 | 2.48 | 0.49 |
| 31:DA:1858:G:H8 | 31:DA:1858:G:OP2 | 1.94 | 0.49 |
| 31:DA:1861:G:C2 | 31:DA:1862:G:C8 | 3.01 | 0.49 |
| 31:DA:2098:U:H2' | 31:DA:2099:U:H6 | 1.76 | 0.49 |
| 31:DA:2575:C:H2' | 31:DA:2578:G:O6 | 2.13 | 0.49 |
| 31:DA:2584:U:O2 | 31:DA:2584:U:O5' | 2.30 | 0.49 |
| 32:DB:28:C:N3 | 32:DB:29:A:N7 | 2.60 | 0.49 |
| 33:DD:30:GLU:HG3 | 33:DD:63:ARG:NE | 2.26 | 0.49 |
| 34:DE:119:ARG:HB3 | 34:DE:120:TRP:CD1 | 2.47 | 0.49 |
| 40:DO:61:VAL:O | 40:DO:63:VAL:HG12 | 2.12 | 0.49 |
| 41:DP:8:PRO:O | 41:DP:10:PRO:HD3 | 2.11 | 0.49 |
| 41:DP:66:GLY:O | 41:DP:67:MET:O | 2.29 | 0.49 |
| 42:DQ:75:THR:HA | 42:DQ:88:GLY:CA | 2.42 | 0.49 |
| 46:DU:8:VAL:HG11 | 46:DU:12:ARG:NH1 | 2.26 | 0.49 |
| 46:DU:92:ARG:HD2 | 47:DV:11:GLN:HG2 | 1.90 | 0.49 |
| 47:DV:91:TYR:CD2 | 47:DV:91:TYR:C | 2.90 | 0.49 |
| 50:DY:45:VAL:HG13 | 50:DY:62:GLU:OE2 | 2.12 | 0.49 |
| 51:DZ:44:PHE:CZ | 51:DZ:48:PHE:HD2 | 2.31 | 0.49 |
| 1:AA:146:G:O6 | 1:AA:176:C:N3 | 2.45 | 0.49 |
| 1:AA:186:C:C2 | 1:AA:187:C:C5 | 3.01 | 0.49 |
| 1:AA:373:A:C2 | 1:AA:482:A:N6 | 2.81 | 0.49 |
| 1:AA:441:A:H3' | 1:AA:442:C:C6 | 2.47 | 0.49 |
| 1:AA:515:G:N2 | 1:AA:537:G:C4 | 2.81 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:655:A:C2 | 1:AA:754:C:C4 | 3.00 | 0.49 |
| 1:AA:675:A:C6 | 1:AA:676:A:C5 | 3.01 | 0.49 |
| 1:AA:679:C:O2' | 1:AA:680:C:H5' | 2.13 | 0.49 |
| 1:AA:776:G:HO2' | 1:AA:777:A:P | 2.36 | 0.49 |
| 1:AA:986:A:N1 | 1:AA:1220:G:C2 | 2.80 | 0.49 |
| 1:AA:1030(D):A:H62 | 1:AA:1031:G:H21 | 1.60 | 0.49 |
| 1:AA:1464:G:OP2 | 45:BT:111:ARG:NH2 | 2.45 | 0.49 |
| 1:AA:1505:G:H5" | 1:AA:1506:U:OP1 | 2.13 | 0.49 |
| 6:AF:60:PHE:C | 6:AF:61:LEU:HD12 | 2.38 | 0.49 |
| 7:AG:146:GLU:HA | 7:AG:149:ARG:HB2 | 1.93 | 0.49 |
| 11:AK:29:ILE:HB | 11:AK:44:SER:HB2 | 1.94 | 0.49 |
| 11:AK:59:TYR:CE2 | 11:AK:63:LEU:HD12 | 2.48 | 0.49 |
| 12:AL:41:ARG:CG | 12:AL:42:THR:N | 2.74 | 0.49 |
| 22:B0:71:ASP:C | 22:B0:72:ARG:HG2 | 2.38 | 0.49 |
| 31:BA:38:A:H2' | 31:BA:39:C:C6 | 2.48 | 0.49 |
| 31:BA:107:C:H2' | 31:BA:108:U:C6 | 2.48 | 0.49 |
| 31:BA:386:G:H4' | 31:BA:387:U:OP2 | 2.11 | 0.49 |
| 31:BA:878:A:N6 | 31:BA:900:A:C8 | 2.80 | 0.49 |
| 31:BA:1489:U:O2' | 31:BA:1490:A:H5" | 2.12 | 0.49 |
| 31:BA:1497:U:C2' | 31:BA:1498:C:OP1 | 2.60 | 0.49 |
| 31:BA:1509(B):A:H2' | 31:BA:1510:G:C8 | 2.48 | 0.49 |
| 31:BA:2287:A:C4 | 31:BA:2289:G:N7 | 2.80 | 0.49 |
| 31:BA:2306:C:OP2 | 31:BA:2307:G:H8 | 1.95 | 0.49 |
| 31:BA:2312:U:C2' | 31:BA:2313:C:H5' | 2.43 | 0.49 |
| 34:BE:128:SER:O | 34:BE:130:GLY:N | 2.45 | 0.49 |
| 38:BI:8:PRO:O | 38:BI:9:LEU:HD23 | 2.12 | 0.49 |
| 38:BI:109:ILE:HG22 | 38:BI:130:TYR:HH | 1.75 | 0.49 |
| 39:BN:131:GLN:OE1 | 39:BN:134:ARG:HB3 | 2.10 | 0.49 |
| 40:BO:87:ILE:CG2 | 40:BO:88:ASN:N | 2.74 | 0.49 |
| 41:BP:8:PRO:O | 41:BP:10:PRO:HD3 | 2.12 | 0.49 |
| 41:BP:143:GLY:C | 41:BP:145:PRO:CD | 2.76 | 0.49 |
| 41:BP:144:GLU:N | 41:BP:145:PRO:CD | 2.75 | 0.49 |
| 43:BR:96:ARG:HH21 | 43:BR:117:VAL:HG23 | 1.78 | 0.49 |
| 47:BV:19:LYS:HE2 | 47:BV:20:LEU:HD12 | 1.93 | 0.49 |
| 48:BW:4:LYS:CB | 48:BW:106:ILE:HG22 | 2.40 | 0.49 |
| 1:CA:382:A:C2 | 1:CA:383:A:C4 | 3.00 | 0.49 |
| 1:CA:1099:G:H2' | 1:CA:1099:G:N3 | 2.27 | 0.49 |
| 1:CA:1271:G:H5' | 1:CA:1314:C:H5' | 1.94 | 0.49 |
| 2:CB:19:HIS:CE1 | 2:CB:206:ASP:HB2 | 2.46 | 0.49 |
| 2:CB:61:LEU:HD21 | 2:CB:68:ILE:HD11 | 1.94 | 0.49 |
| 7:CG:62:PHE:HA | 7:CG:124:LEU:HD22 | 1.93 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 8:CH:120:THR:H | 8:CH:123:GLU:HB2 | 1.77 | 0.49 |
| 10:CJ:44:VAL:HG22 | 10:CJ:66:ARG:HG2 | 1.94 | 0.49 |
| 23:D1:25:LYS:O | 23:D1:26:ARG:HB3 | 2.13 | 0.49 |
| 31:DA:745:G:H2' | 31:DA:746:A:H5' | 1.93 | 0.49 |
| 31:DA:1275:A:C8 | 43:DR:16:HIS:CD2 | 3.00 | 0.49 |
| 31:DA:1349:A:N6 | 31:DA:1598:C:N4 | 2.60 | 0.49 |
| 31:DA:1433:U:O2' | 31:DA:1434:A:H5' | 2.12 | 0.49 |
| 31:DA:2599:G:OP2 | 33:DD:236:GLY:N | 2.45 | 0.49 |
| 32:DB:30:C:H2' | 32:DB:31:C:H5' | 1.94 | 0.49 |
| 33:DD:155:LEU:HD23 | 33:DD:177:LEU:HD22 | 1.94 | 0.49 |
| 35:DF:53:THR:HG23 | 35:DF:56:GLU:H | 1.78 | 0.49 |
| 36:DG:25:TYR:CE2 | 36:DG:32:PRO:HD3 | 2.46 | 0.49 |
| 36:DG:174:GLU:HG3 | 36:DG:180:PHE:CD1 | 2.47 | 0.49 |
| 47:DV:4:ILE:O | 47:DV:39:LEU:HB3 | 2.11 | 0.49 |
| 51:DZ:175:VAL:HB | 51:DZ:176:PRO:CD | 2.43 | 0.49 |
| 1:AA:335:C:O2' | 1:AA:336:C:H5' | 2.13 | 0.49 |
| 1:AA:354:G:C4 | 1:AA:355:C:C5 | 3.00 | 0.49 |
| 1:AA:710:G:H5" | 6:AF:54:LYS:HE3 | 1.95 | 0.49 |
| 1:AA:1142:G:H3' | 1:AA:1143:G:H8 | 1.77 | 0.49 |
| 1:AA:1152:A:H5" | 10:AJ:13:HIS:HD2 | 1.74 | 0.49 |
| 2:AB:87:ARG:NE | 2:AB:233:SER:HB3 | 2.23 | 0.49 |
| 2:AB:185:ILE:HA | 2:AB:199:TYR:O | 2.12 | 0.49 |
| 8:AH:30:ARG:O | 8:AH:34:GLU:HG2 | 2.13 | 0.49 |
| 8:AH:134:ILE:O | 8:AH:135:CYS:HB3 | 2.13 | 0.49 |
| 9:AI:4:TYR:N | 9:AI:4:TYR:CD1 | 2.81 | 0.49 |
| 9:AI:113:LYS:H | 9:AI:119:ALA:HA | 1.77 | 0.49 |
| 10:AJ:8:LEU:HD22 | 10:AJ:20:ALA:HB2 | 1.94 | 0.49 |
| 11:AK:96:ARG:O | 11:AK:99:GLN:HG2 | 2.13 | 0.49 |
| 12:AL:22:SER:C | 12:AL:24:VAL:N | 2.71 | 0.49 |
| 20:AT:73:HIS:HB3 | 20:AT:74:LYS:HG2 | 1.93 | 0.49 |
| 31:BA:372:G:O2' | 31:BA:373:U:P | 2.71 | 0.49 |
| 31:BA:541:C:C4 | 31:BA:542:C:N4 | 2.81 | 0.49 |
| 31:BA:1557:C:H5" | 31:BA:1558:A:OP2 | 2.12 | 0.49 |
| 31:BA:1688:U:O2 | 31:BA:1700:A:H8 | 1.96 | 0.49 |
| 31:BA:2808:U:H2' | 31:BA:2809:A:C5' | 2.42 | 0.49 |
| 32:BB:94:C:H2' | 32:BB:95:C:H6 | 1.77 | 0.49 |
| 33:BD:24:ILE:CG2 | 33:BD:24:ILE:O | 2.59 | 0.49 |
| 33:BD:92:ILE:HD13 | 33:BD:104:TYR:CE2 | 2.48 | 0.49 |
| 33:BD:211:ARG:HA | 33:BD:214:TRP:CD2 | 2.47 | 0.49 |
| 38:BI:31:LEU:HD13 | 38:BI:31:LEU:N | 2.27 | 0.49 |
| 46:BU:55:ARG:HA | 46:BU:58:ARG:HD2 | 1.94 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 46:BU:57:PHE:O | 46:BU:58:ARG:C | 2.53 | 0.49 |
| 46:BU:66:ASN:ND2 | 46:BU:66:ASN:C | 2.70 | 0.49 |
| 49:BX:37:THR:HG23 | 49:BX:54:VAL:CB | 2.35 | 0.49 |
| 1:CA:658:G:C6 | 1:CA:749:C:N4 | 2.79 | 0.49 |
| 1:CA:960:U:H2' | 1:CA:960:U:O2 | 2.11 | 0.49 |
| 1:CA:1381:U:H2' | 1:CA:1382:C:H5' | 1.94 | 0.49 |
| 2:CB:90:MET:HE2 | 2:CB:90:MET:HA | 1.93 | 0.49 |
| 3:CC:19:GLU:HG2 | 3:CC:19:GLU:O | 2.13 | 0.49 |
| 5:CE:10:MET:HB2 | 5:CE:32:VAL:CG2 | 2.38 | 0.49 |
| 11:CK:108:ILE:HB | 18:CR:87:ARG:HA | 1.94 | 0.49 |
| 20:CT:37:SER:O | 20:CT:41:ILE:HG12 | 2.13 | 0.49 |
| 21:CU:22:ARG:N | 21:CU:23:PRO:HD3 | 2.27 | 0.49 |
| 22:D0:16:SER:HB2 | 31:DA:2262:U:OP2 | 2.11 | 0.49 |
| 22:D0:74:ARG:NH2 | 32:DB:13:A:C8 | 2.80 | 0.49 |
| 30:D8:35:GLN:CB | 30:D8:36:LYS:HZ3 | 2.25 | 0.49 |
| 31:DA:150:C:H2' | 31:DA:151:C:H6 | 1.78 | 0.49 |
| 31:DA:667:U:H2' | 31:DA:668:G:C5' | 2.43 | 0.49 |
| 31:DA:685:A:C8 | 31:DA:774:A:C6 | 3.00 | 0.49 |
| 31:DA:745:G:C2' | 31:DA:746:A:H5' | 2.42 | 0.49 |
| 31:DA:746:A:N3 | 55:DA:3311:ZIT:H161 | 2.25 | 0.49 |
| 31:DA:911:A:C4 | 42:DQ:9:TYR:OH | 2.58 | 0.49 |
| 31:DA:1204:A:H2 | 31:DA:1241:A:N1 | 2.09 | 0.49 |
| 31:DA:1688:U:O2 | 31:DA:1700:A:H8 | 1.95 | 0.49 |
| 31:DA:2074:U:H2' | 31:DA:2075:U:C6 | 2.47 | 0.49 |
| 34:DE:33:VAL:CG1 | 34:DE:90:THR:H | 2.24 | 0.49 |
| 36:DG:114:ILE:O | 36:DG:115:ARG:C | 2.56 | 0.49 |
| 38:DI:50:ARG:C | 38:DI:52:ARG:N | 2.70 | 0.49 |
| 46:DU:26:GLY:C | 46:DU:28:ARG:N | 2.70 | 0.49 |
| 46:DU:92:ARG:CB | 47:DV:11:GLN:NE2 | 2.70 | 0.49 |
| 47:DV:35:LEU:HB2 | 47:DV:59:ALA:HB1 | 1.93 | 0.49 |
| 49:DX:37:THR:C | 49:DX:38:GLU:OE1 | 2.55 | 0.49 |
| 1:AA:166:G:C4 | 1:AA:167:G:C8 | 3.00 | 0.49 |
| 1:AA:262:A:C6 | 1:AA:263:A:N6 | 2.80 | 0.49 |
| 1:AA:950:U:H2' | 1:AA:951:G:H8 | 1.78 | 0.49 |
| 1:AA:1128:C:H5' | 9:AI:16:ARG:HH22 | 1.76 | 0.49 |
| 4:AD:52:SER:O | 4:AD:53:ASP:C | 2.55 | 0.49 |
| 8:AH:38:ILE:HD11 | 8:AH:118:VAL:O | 2.13 | 0.49 |
| 23:B1:64:ALA:C | 23:B1:67:ILE:HD11 | 2.38 | 0.49 |
| 31:BA:856:C:C6 | 31:BA:856:C:H5" | 2.46 | 0.49 |
| 31:BA:1316:U:O2' | 31:BA:1317:A:H5' | 2.12 | 0.49 |
| 31:BA:1478:G:H2' | 31:BA:1479:G:H5' | 1.94 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1887:C:C3' | 31:BA:1888:G:H5' | 2.42 | 0.49 |
| 31:BA:2314:C:O2 | 31:BA:2314:C:H2' | 2.11 | 0.49 |
| 31:BA:2315:G:C6 | 31:BA:2316:C:N4 | 2.81 | 0.49 |
| 33:BD:43:ARG:NH1 | 33:BD:44:ASN:ND2 | 2.61 | 0.49 |
| 36:BG:56:ALA:HB2 | 36:BG:153:ARG:NH2 | 2.26 | 0.49 |
| 38:BI:79:ILE:CG2 | 38:BI:81:VAL:HG23 | 2.43 | 0.49 |
| 38:BI:86:THR:HG23 | 38:BI:122:GLU:OE2 | 2.12 | 0.49 |
| 41:BP:6:LEU:HD12 | 41:BP:8:PRO:HB2 | 1.94 | 0.49 |
| 41:BP:83:VAL:CG1 | 41:BP:112:LEU:HD21 | 2.42 | 0.49 |
| 42:BQ:63:LYS:HB2 | 42:BQ:63:LYS:HZ3 | 1.77 | 0.49 |
| 42:BQ:116:GLU:O | 42:BQ:117:ALA:C | 2.52 | 0.49 |
| 1:CA:279:A:OP2 | 17:CQ:95:TYR:OH | 2.26 | 0.49 |
| 1:CA:340:U:H2' | 1:CA:341:C:O4' | 2.12 | 0.49 |
| 1:CA:615:C:H2' | 1:CA:616:G:O4' | 2.11 | 0.49 |
| 1:CA:1498:U:H1' | 1:CA:1499:A:OP2 | 2.13 | 0.49 |
| 11:CK:111:ASP:HA | 18:CR:84:LYS:CE | 2.42 | 0.49 |
| 22:D0:55:ARG:HG3 | 31:DA:2365:G:OP1 | 2.11 | 0.49 |
| 24:D2:26:ARG:HG3 | 24:D2:29:LYS:NZ | 2.28 | 0.49 |
| 27:D5:47:PRO:O | 27:D5:48:GLU:HG3 | 2.12 | 0.49 |
| 31:DA:27:G:O2' | 31:DA:28:A:OP2 | 2.30 | 0.49 |
| 31:DA:372:G:O2' | 31:DA:373:U:P | 2.70 | 0.49 |
| 31:DA:558:G:P | 39:DN:111:PRO:HD2 | 2.52 | 0.49 |
| 31:DA:1568:G:H21 | 33:DD:58:HIS:HE1 | 1.61 | 0.49 |
| 31:DA:1693:U:H4' | 31:DA:1694:C:OP2 | 2.11 | 0.49 |
| 31:DA:2206:G:H21 | 31:DA:2207:G:C4' | 2.24 | 0.49 |
| 31:DA:2228:G:H2' | 31:DA:2229:C:C6 | 2.48 | 0.49 |
| 31:DA:2273:A:H2' | 31:DA:2274:A:C8 | 2.48 | 0.49 |
| 31:DA:2619:C:H2' | 31:DA:2620:C:H6 | 1.76 | 0.49 |
| 35:DF:21:ALA:C | 35:DF:23:ASP:H | 2.21 | 0.49 |
| 35:DF:24:LEU:CB | 35:DF:25:PRO:HD2 | 2.35 | 0.49 |
| 41:DP:45:LEU:CD2 | 41:DP:46:LYS:N | 2.76 | 0.49 |
| 47:DV:28:GLU:CG | 47:DV:29:PRO:HD3 | 2.41 | 0.49 |
| 48:DW:4:LYS:HE3 | 48:DW:6:ILE:HD11 | 1.95 | 0.49 |
| 49:DX:25:LYS:HG3 | 49:DX:26:TYR:CD1 | 2.46 | 0.49 |
| 50:DY:27:VAL:HG12 | 50:DY:29:GLU:OE1 | 2.12 | 0.49 |
| 50:DY:37:VAL:CG2 | 50:DY:67:LEU:HD23 | 2.42 | 0.49 |
| 1:AA:355:C:C4 | 1:AA:356:A:N7 | 2.80 | 0.49 |
| 1:AA:977:A:C2' | 1:AA:978:A:H5' | 2.43 | 0.49 |
| 1:AA:1321:C:H3' | 1:AA:1322:C:H5" | 1.94 | 0.49 |
| 1:AA:1530:G:OP1 | 1:AA:1530:G:C4' | 2.61 | 0.49 |
| 3:AC:27:LYS:HA | 3:AC:27:LYS:HZ3 | 1.78 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 5:AE:57:LYS:O | 5:AE:61:TYR:CD2 | 2.63 | 0.49 |
| 6:AF:8:ILE:HG22 | 6:AF:10:LEU:HD11 | 1.95 | 0.49 |
| 6:AF:18:GLN:O | 6:AF:21:LEU:HB2 | 2.12 | 0.49 |
| 17:AQ:59:ILE:HD13 | 17:AQ:73:VAL:HA | 1.93 | 0.49 |
| 27:B5:27:PRO:HB3 | 48:BW:23:LEU:CD1 | 2.42 | 0.49 |
| 31:BA:528:A:C2 | 31:BA:2043:C:H4' | 2.46 | 0.49 |
| 31:BA:860:U:C5 | 31:BA:2268:A:C8 | 3.01 | 0.49 |
| 31:BA:1498:C:N4 | 31:BA:1499:C:N4 | 2.60 | 0.49 |
| 31:BA:1673:U:C4 | 34:BE:129:HIS:HD2 | 2.30 | 0.49 |
| 31:BA:1679:U:H2' | 31:BA:1680:U:H5' | 1.95 | 0.49 |
| 31:BA:1884:A:O2' | 31:BA:1885:A:H5' | 2.12 | 0.49 |
| 31:BA:2740:A:C6 | 31:BA:2764:A:C8 | 3.01 | 0.49 |
| 32:BB:81:G:O6 | 32:BB:96:U:O2 | 2.30 | 0.49 |
| 33:BD:221:VAL:HG22 | 33:BD:226:MET:HE3 | 1.93 | 0.49 |
| 37:BH:145:ALA:HB1 | 37:BH:164:TYR:CE1 | 2.48 | 0.49 |
| 39:BN:2:LYS:HE3 | 47:BV:12:TYR:HA | 1.93 | 0.49 |
| 39:BN:18:ALA:O | 39:BN:21:LYS:N | 2.44 | 0.49 |
| 41:BP:125:VAL:O | 41:BP:145:PRO:HD2 | 2.13 | 0.49 |
| 45:BT:128:GLU:C | 45:BT:130:ALA:N | 2.67 | 0.49 |
| 46:BU:95:LEU:C | 46:BU:97:ASP:N | 2.71 | 0.49 |
| 1:CA:139:G:H2' | 1:CA:140:A:H8 | 1.77 | 0.49 |
| 1:CA:146:G:O6 | 1:CA:176:C:N3 | 2.45 | 0.49 |
| 1:CA:437:U:C5' | 4:CD:155:LEU:HD13 | 2.42 | 0.49 |
| 1:CA:592:G:C2 | 1:CA:648:A:C2 | 3.01 | 0.49 |
| 1:CA:617:G:N1 | 1:CA:618:C:C5 | 2.81 | 0.49 |
| 1:CA:706:A:C5 | 1:CA:707:C:C5 | 3.00 | 0.49 |
| 1:CA:914:A:C2' | 1:CA:915:A:H5' | 2.42 | 0.49 |
| 1:CA:1151:A:O2' | 1:CA:1152:A:H8 | 1.95 | 0.49 |
| 4:CD:119:GLN:O | 4:CD:123:HIS:HD2 | 1.96 | 0.49 |
| 5:CE:147:ASP:N | 5:CE:147:ASP:OD2 | 2.45 | 0.49 |
| 9:CI:77:ILE:O | 9:CI:81:ILE:HG12 | 2.12 | 0.49 |
| 11:CK:59:TYR:CE2 | 11:CK:63:LEU:HD12 | 2.48 | 0.49 |
| 19:CS:36:ARG:HD2 | 19:CS:52:TYR:O | 2.13 | 0.49 |
| 23:D1:11:ARG:HB3 | 23:D1:12:PRO:HD3 | 1.94 | 0.49 |
| 24:D2:14:ARG:NH2 | 24:D2:57:ILE:HG22 | 2.28 | 0.49 |
| 24:D2:45:SER:O | 24:D2:48:HIS:HB2 | 2.13 | 0.49 |
| 28:D6:9:LEU:HD13 | 28:D6:9:LEU:C | 2.37 | 0.49 |
| 29:D7:43:THR:O | 29:D7:44:PRO:C | 2.56 | 0.49 |
| 31:DA:301:G:H1 | 31:DA:316:C:H42 | 1.60 | 0.49 |
| 31:DA:543:C:N4 | 31:DA:551:G:H1 | 2.10 | 0.49 |
| 31:DA:1112:G:C1' | 31:DA:1113:U:OP1 | 2.61 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1509(B):A:H2' | 31:DA:1510:G:C8 | 2.47 | 0.49 |
| 31:DA:1654:A:C1' | 31:DA:2823:A:H5' | 2.43 | 0.49 |
| 31:DA:2186:G:H3' | 31:DA:2187:G:H5" | 1.94 | 0.49 |
| 31:DA:2416:C:OP1 | 41:DP:64:LYS:O | 2.31 | 0.49 |
| 31:DA:2759:G:O2' | 31:DA:2760:C:H5' | 2.13 | 0.49 |
| 32:DB:2:C:C2 | 32:DB:3:C:C5 | 3.00 | 0.49 |
| 35:DF:84:VAL:O | 35:DF:85:GLY:C | 2.56 | 0.49 |
| 36:DG:11:TYR:OH | 36:DG:33:ARG:HA | 2.13 | 0.49 |
| 36:DG:32:PRO:HB3 | 36:DG:163:ALA:HB2 | 1.95 | 0.49 |
| 36:DG:106:LEU:O | 36:DG:111:LEU:HG | 2.13 | 0.49 |
| 37:DH:54:ARG:HB3 | 37:DH:65:HIS:HB2 | 1.93 | 0.49 |
| 37:DH:66:GLY:HA2 | 37:DH:69:ARG:HB2 | 1.94 | 0.49 |
| 45:DT:33:LYS:HA | 45:DT:33:LYS:NZ | 2.27 | 0.49 |
| 49:DX:65:ARG:NE | 49:DX:65:ARG:CA | 2.75 | 0.49 |
| 50:DY:75:ILE:O | 50:DY:76:CYS:HB2 | 2.12 | 0.49 |
| 1:AA:17:U:C2 | 1:AA:18:C:C5 | 3.01 | 0.49 |
| 1:AA:66:G:O4' | 1:AA:173:U:C4 | 2.66 | 0.49 |
| 1:AA:363:A:C5 | 12:AL:31:PRO:HD2 | 2.48 | 0.49 |
| 1:AA:533:A:C4' | 1:AA:534:U:OP1 | 2.60 | 0.49 |
| 1:AA:611:A:N6 | 1:AA:629:G:H1 | 2.10 | 0.49 |
| 1:AA:701:C:O2 | 1:AA:703:G:N1 | 2.46 | 0.49 |
| 1:AA:1305:G:H5' | 21:AU:4:GLY:HA3 | 1.95 | 0.49 |
| 1:AA:1398:A:N1 | 5:AE:19:MET:HE3 | 2.28 | 0.49 |
| 3:AC:182:ILE:CG1 | 3:AC:203:PHE:HD1 | 2.24 | 0.49 |
| 6:AF:91:VAL:HG12 | 6:AF:92:LYS:O | 2.12 | 0.49 |
| 7:AG:25:ALA:O | 7:AG:29:LYS:HG2 | 2.12 | 0.49 |
| 8:AH:109:ILE:HG22 | 8:AH:137:VAL:HB | 1.95 | 0.49 |
| 10:AJ:48:THR:HA | 10:AJ:62:HIS:HB3 | 1.95 | 0.49 |
| 22:B0:34:GLY:O | 22:B0:35:ASN:C | 2.55 | 0.49 |
| 31:BA:298:G:H5" | 31:BA:299:A:OP1 | 2.13 | 0.49 |
| 31:BA:415:A:H2' | 31:BA:416:C:C6 | 2.48 | 0.49 |
| 31:BA:675:A:C6 | 31:BA:676:A:C6 | 3.01 | 0.49 |
| 31:BA:848:G:C4 | 31:BA:933:A:H8 | 2.30 | 0.49 |
| 31:BA:993:G:N3 | 47:BV:91:TYR:CE1 | 2.81 | 0.49 |
| 31:BA:1050:A:C2 | 31:BA:2751:G:C5 | 3.01 | 0.49 |
| 31:BA:1279:G:H5' | 43:BR:34:ILE:HD11 | 1.95 | 0.49 |
| 31:BA:2680:C:H5' | 34:BE:189:PRO:HA | 1.94 | 0.49 |
| 33:BD:43:ARG:HH11 | 33:BD:44:ASN:ND2 | 2.09 | 0.49 |
| 33:BD:44:ASN:OD1 | 33:BD:44:ASN:N | 2.45 | 0.49 |
| 33:BD:84:TYR:HE2 | 33:BD:86:PRO:HB3 | 1.77 | 0.49 |
| 36:BG:135:LEU:HD12 | 36:BG:135:LEU:N | 2.27 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 37:BH:40:GLU:O | 37:BH:41:MET:CB | 2.57 | 0.49 |
| 37:BH:89:ILE:C | 37:BH:89:ILE:HD12 | 2.38 | 0.49 |
| 41:BP:112:LEU:CD2 | 41:BP:113:LYS:N | 2.76 | 0.49 |
| 45:BT:32:TYR:CD2 | 45:BT:81:PRO:O | 2.65 | 0.49 |
| 45:BT:78:LEU:O | 45:BT:79:HIS:CG | 2.65 | 0.49 |
| 47:BV:40:LEU:O | 47:BV:41:GLY:C | 2.56 | 0.49 |
| 47:BV:46:VAL:O | 47:BV:47:VAL:HB | 2.12 | 0.49 |
| 1:CA:93:G:C6 | 1:CA:96:U:C4 | 3.00 | 0.49 |
| 1:CA:299:G:C5 | 1:CA:300:A:C6 | 3.01 | 0.49 |
| 1:CA:327:A:C3' | 1:CA:328:C:H5" | 2.39 | 0.49 |
| 1:CA:664:G:P | 18:CR:64:ARG:HH21 | 2.36 | 0.49 |
| 1:CA:834:C:H2' | 1:CA:835:U:C6 | 2.48 | 0.49 |
| 1:CA:1142:G:H3' | 1:CA:1143:G:H8 | 1.77 | 0.49 |
| 1:CA:1160:G:N2 | 1:CA:1161:C:C6 | 2.81 | 0.49 |
| 1:CA:1187:G:C6 | 1:CA:1188:A:C6 | 3.00 | 0.49 |
| 1:CA:1255:G:H2' | 1:CA:1255:G:N3 | 2.28 | 0.49 |
| 1:CA:1452:C:H4' | 1:CA:1456:G:N3 | 2.28 | 0.49 |
| 2:CB:83:MET:O | 2:CB:85:ALA:N | 2.46 | 0.49 |
| 3:CC:11:ARG:O | 3:CC:12:LEU:C | 2.55 | 0.49 |
| 7:CG:108:ALA:O | 7:CG:111:ARG:HB2 | 2.12 | 0.49 |
| 12:CL:60:LEU:N | 12:CL:60:LEU:HD22 | 2.28 | 0.49 |
| 13:CM:14:ARG:HA | 13:CM:43:THR:O | 2.13 | 0.49 |
| 20:CT:79:ARG:HA | 20:CT:82:SER:OG | 2.13 | 0.49 |
| 28:D6:32:ASN:OD1 | 28:D6:33:LYS:N | 2.45 | 0.49 |
| 30:D8:4:MET:HE1 | 31:DA:593:G:C1' | 2.42 | 0.49 |
| 31:DA:34:C:HO2' | 31:DA:35:G:P | 2.36 | 0.49 |
| 31:DA:271(J):C:C3' | 31:DA:271(K):U:H5" | 2.43 | 0.49 |
| 31:DA:484:C:H2' | 31:DA:485:C:H6 | 1.78 | 0.49 |
| 31:DA:708:C:H42 | 31:DA:723:G:H1 | 1.59 | 0.49 |
| 31:DA:776:G:C8 | 31:DA:793:A:C2 | 3.01 | 0.49 |
| 31:DA:1312:U:OP2 | 49:DX:62:LYS:HE3 | 2.12 | 0.49 |
| 31:DA:1658:C:H2' | 31:DA:1659:U:H6 | 1.77 | 0.49 |
| 31:DA:1859:A:C2 | 31:DA:1884:A:H1' | 2.47 | 0.49 |
| 31:DA:2056:G:N3 | 31:DA:2056:G:H2' | 2.27 | 0.49 |
| 31:DA:2409:G:H2' | 31:DA:2410:G:O4' | 2.13 | 0.49 |
| 31:DA:2803:C:H2' | 31:DA:2804:C:O4' | 2.12 | 0.49 |
| 37:DH:85:LYS:CE | 37:DH:145:ALA:N | 2.76 | 0.49 |
| 37:DH:145:ALA:HB1 | 37:DH:164:TYR:CE1 | 2.48 | 0.49 |
| 38:DI:44:LEU:HD23 | 38:DI:44:LEU:HA | 1.39 | 0.49 |
| 39:DN:1:MET:CB | 47:DV:20:LEU:HD22 | 2.25 | 0.49 |
| 45:DT:57:PHE:O | 45:DT:58:ASN:C | 2.55 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 45:DT:109:GLU:HB3 | 45:DT:113:LYS:CE | 2.40 | 0.49 |
| 46:DU:66:ASN:HD21 | 46:DU:70:ARG:HE | 1.60 | 0.49 |
| 47:DV:73:SER:O | 47:DV:74:LYS:HB2 | 2.12 | 0.49 |
| 49:DX:36:LYS:HD2 | 49:DX:36:LYS:C | 2.36 | 0.49 |
| 49:DX:65:ARG:C | 49:DX:65:ARG:HD3 | 2.38 | 0.49 |
| 50:DY:79:CYS:O | 50:DY:80:GLY:O | 2.31 | 0.49 |
| 50:DY:86:ARG:HG2 | 50:DY:87:LYS:H | 1.77 | 0.49 |
| 1:AA:44:G:N2 | 1:AA:399:G:C4 | 2.80 | 0.49 |
| 1:AA:90:U:O2' | 1:AA:91:C:C5 | 2.65 | 0.49 |
| 1:AA:327:A:C3' | 1:AA:328:C:H5" | 2.40 | 0.49 |
| 1:AA:382:A:H2' | 1:AA:383:A:C8 | 2.48 | 0.49 |
| 1:AA:448:A:OP2 | 1:AA:485:G:N2 | 2.36 | 0.49 |
| 1:AA:961:U:OP2 | 1:AA:1223:C:H4' | 2.13 | 0.49 |
| 1:AA:1478:C:O2' | 1:AA:1479:C:H5' | 2.13 | 0.49 |
| 2:AB:23:ARG:HG2 | 2:AB:23:ARG:O | 2.13 | 0.49 |
| 2:AB:239:VAL:HG12 | 2:AB:239:VAL:O | 2.13 | 0.49 |
| 3:AC:11:ARG:O | 3:AC:14:ILE:O | 2.30 | 0.49 |
| 5:AE:45:PHE:CE2 | 5:AE:47:LYS:HD2 | 2.48 | 0.49 |
| 12:AL:90:VAL:C | 12:AL:92:ASP:H | 2.19 | 0.49 |
| 20:AT:79:ARG:HA | 20:AT:82:SER:OG | 2.13 | 0.49 |
| 27:B5:57:VAL:CG2 | 27:B5:58:LEU:H | 2.18 | 0.49 |
| 28:B6:34:LEU:HD22 | 28:B6:50:ARG:NH1 | 2.27 | 0.49 |
| 30:B8:2:PRO:O | 30:B8:3:LYS:C | 2.55 | 0.49 |
| 30:B8:43:GLN:C | 30:B8:44:LYS:HD2 | 2.38 | 0.49 |
| 31:BA:601:C:O2 | 31:BA:605:C:H4' | 2.13 | 0.49 |
| 31:BA:719:C:H6 | 31:BA:719:C:O5' | 1.95 | 0.49 |
| 31:BA:923:C:H2' | 31:BA:924:C:H6 | 1.78 | 0.49 |
| 31:BA:1697:G:H8 | 31:BA:1697:G:C5' | 2.11 | 0.49 |
| 31:BA:1890:A:H2' | 31:BA:1891:G:H5' | 1.93 | 0.49 |
| 31:BA:1991:U:C2' | 31:BA:1992:G:H5" | 2.42 | 0.49 |
| 31:BA:2286:A:H8 | 31:BA:2286:A:HO2' | 1.55 | 0.49 |
| 31:BA:2887:U:O2' | 31:BA:2888:C:H5' | 2.13 | 0.49 |
| 33:BD:28:GLU:HB2 | 33:BD:29:PRO:CD | 2.42 | 0.49 |
| 36:BG:128:ARG:O | 36:BG:129:GLY:C | 2.55 | 0.49 |
| 39:BN:78:TYR:N | 39:BN:79:PRO:CD | 2.71 | 0.49 |
| 43:BR:72:ASP:HB3 | 43:BR:75:LEU:CB | 2.42 | 0.49 |
| 44:BS:19:LYS:O | 44:BS:19:LYS:HG2 | 2.11 | 0.49 |
| 45:BT:106:SER:O | 45:BT:107:ASP:CB | 2.60 | 0.49 |
| 46:BU:91:ASP:O | 46:BU:92:ARG:HB3 | 2.13 | 0.49 |
| 47:BV:38:LEU:HG | 47:BV:39:LEU:H | 1.78 | 0.49 |
| 48:BW:18:ARG:NH1 | 48:BW:18:ARG:CG | 2.75 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 51:BZ:141:VAL:O | 51:BZ:141:VAL:HG13 | 2.13 | 0.49 |
| 1:CA:682:G:H1 | 1:CA:708:C:H42 | 1.61 | 0.49 |
| 1:CA:693:G:H1' | 7:CG:82:GLY:HA3 | 1.94 | 0.49 |
| 1:CA:1082:G:C6 | 1:CA:1083:U:N3 | 2.81 | 0.49 |
| 1:CA:1206:G:O4' | 3:CC:194:GLY:HA2 | 2.13 | 0.49 |
| 8:CH:65:TYR:N | 8:CH:65:TYR:CD1 | 2.81 | 0.49 |
| 14:CN:44:LEU:C | 14:CN:44:LEU:HD12 | 2.37 | 0.49 |
| 16:CP:8:ARG:HG2 | 16:CP:9:PHE:H | 1.77 | 0.49 |
| 17:CQ:83:ASP:O | 17:CQ:87:LYS:HG2 | 2.13 | 0.49 |
| 24:D2:30:ARG:H | 24:D2:30:ARG:CD | 2.12 | 0.49 |
| 24:D2:47:ASN:HB2 | 24:D2:51:ARG:HD2 | 1.94 | 0.49 |
| 25:D3:46:ASN:O | 25:D3:50:VAL:HG22 | 2.12 | 0.49 |
| 25:D3:50:VAL:O | 25:D3:51:ALA:C | 2.55 | 0.49 |
| 25:D3:52:HIS:CD2 | 32:DB:83:G:C5' | 2.95 | 0.49 |
| 27:D5:40:LYS:HZ3 | 27:D5:46:CYS:CB | 2.25 | 0.49 |
| 30:D8:4:MET:HE3 | 30:D8:61:LEU:HD13 | 1.94 | 0.49 |
| 31:DA:675:A:N6 | 31:DA:676:A:N6 | 2.61 | 0.49 |
| 31:DA:1227:G:C2' | 31:DA:1228:G:H5' | 2.43 | 0.49 |
| 31:DA:1349:A:C2' | 31:DA:1350:C:OP1 | 2.61 | 0.49 |
| 31:DA:1836:C:H2' | 31:DA:1837:C:H6 | 1.78 | 0.49 |
| 31:DA:2410:G:C2 | 31:DA:2411:A:H1' | 2.48 | 0.49 |
| 31:DA:2747:G:C6 | 31:DA:2754:U:C5 | 3.00 | 0.49 |
| 31:DA:2881:C:H2' | 31:DA:2882:A:O4' | 2.13 | 0.49 |
| 33:DD:28:GLU:HB2 | 33:DD:29:PRO:HD3 | 1.95 | 0.49 |
| 37:DH:43:VAL:CG1 | 37:DH:53:GLU:H | 2.26 | 0.49 |
| 39:DN:65:LYS:HD3 | 39:DN:65:LYS:C | 2.36 | 0.49 |
| 41:DP:39:LYS:C | 41:DP:41:ARG:H | 2.20 | 0.49 |
| 42:DQ:20:ALA:HB2 | 42:DQ:99:PRO:HG2 | 1.93 | 0.49 |
| 43:DR:21:TYR:OH | 43:DR:43:GLU:HG2 | 2.11 | 0.49 |
| 44:DS:91:PRO:O | 44:DS:93:LYS:N | 2.46 | 0.49 |
| 46:DU:26:GLY:C | 46:DU:28:ARG:H | 2.20 | 0.49 |
| 48:DW:17:VAL:O | 48:DW:18:ARG:C | 2.55 | 0.49 |
| 49:DX:25:LYS:HE3 | 49:DX:25:LYS:HA | 1.94 | 0.49 |
| 49:DX:35:THR:CB | 49:DX:75:ASP:OD2 | 2.59 | 0.49 |
| 50:DY:39:VAL:O | 50:DY:40:GLU:CD | 2.56 | 0.49 |
| 1:AA:438:G:H4' | 4:AD:123:HIS:ND1 | 2.28 | 0.49 |
| 1:AA:523:A:H61 | 12:AL:53:ARG:NH1 | 2.11 | 0.49 |
| 1:AA:721:G:H4' | 1:AA:722:A:O4' | 2.12 | 0.49 |
| 1:AA:1272:G:C6 | 1:AA:1273:G:C5 | 3.01 | 0.49 |
| 1:AA:1327:C:H2' | 1:AA:1328:C:C6 | 2.48 | 0.49 |
| 1:AA:1385:G:O2' | 1:AA:1386:G:H5' | 2.13 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:1490:C:H2' | 1:AA:1491:G:O4' | 2.13 | 0.49 |
| 1:AA:1505:G:H4' | 1:AA:1506:U:C5' | 2.38 | 0.49 |
| 2:AB:135:GLN:O | 2:AB:139:LYS:HB2 | 2.12 | 0.49 |
| 8:AH:14:ARG:O | 8:AH:18:ARG:HD3 | 2.12 | 0.49 |
| 9:AI:28:VAL:HG13 | 9:AI:65:VAL:HG12 | 1.95 | 0.49 |
| 11:AK:105:VAL:O | 11:AK:105:VAL:HG23 | 2.13 | 0.49 |
| 17:AQ:83:ASP:O | 17:AQ:87:LYS:HG2 | 2.12 | 0.49 |
| 24:B2:23:LYS:CB | 49:BX:5:TYR:CE1 | 2.95 | 0.49 |
| 24:B2:34:GLU:OE2 | 24:B2:34:GLU:N | 2.37 | 0.49 |
| 25:B3:32:GLN:HB2 | 31:BA:1158:C:H4' | 1.94 | 0.49 |
| 28:B6:39:TYR:HB3 | 28:B6:49:HIS:ND1 | 2.27 | 0.49 |
| 30:B8:27:THR:HA | 41:BP:62:LEU:HD11 | 1.94 | 0.49 |
| 31:BA:214:G:H1' | 31:BA:216:A:O2' | 2.13 | 0.49 |
| 31:BA:271(S):G:C5 | 31:BA:271(T):C:C5 | 3.01 | 0.49 |
| 31:BA:848:G:C4 | 31:BA:933:A:C8 | 3.01 | 0.49 |
| 31:BA:1654:A:C1' | 31:BA:2823:A:H5' | 2.43 | 0.49 |
| 31:BA:2335:A:C8 | 31:BA:2337:G:C5 | 3.00 | 0.49 |
| 31:BA:2483:C:N3 | 42:BQ:124:LYS:NZ | 2.60 | 0.49 |
| 31:BA:2507:C:C2 | 31:BA:2508:G:C8 | 3.00 | 0.49 |
| 31:BA:2552:U:C2 | 31:BA:2554:U:C5' | 2.96 | 0.49 |
| 31:BA:2801(A):A:C3' | 31:BA:2802:G:H5' | 2.43 | 0.49 |
| 33:BD:224:ALA:O | 33:BD:225:ALA:HB2 | 2.13 | 0.49 |
| 34:BE:128:SER:OG | 34:BE:129:HIS:N | 2.41 | 0.49 |
| 35:BF:139:PHE:CD2 | 35:BF:139:PHE:C | 2.91 | 0.49 |
| 36:BG:60:LEU:O | 36:BG:63:ILE:HG13 | 2.13 | 0.49 |
| 41:BP:16:ARG:HG3 | 41:BP:17:LYS:H | 1.76 | 0.49 |
| 42:BQ:43:THR:HG1 | 42:BQ:46:GLN:HG3 | 1.77 | 0.49 |
| 45:BT:30:VAL:HG21 | 45:BT:84:GLN:H | 1.78 | 0.49 |
| 47:BV:53:GLU:O | 47:BV:53:GLU:HG3 | 2.13 | 0.49 |
| 50:BY:27:VAL:CB | 50:BY:29:GLU:OE1 | 2.60 | 0.49 |
| 1:CA:262:A:C6 | 1:CA:263:A:N6 | 2.81 | 0.49 |
| 1:CA:356:A:H1' | 1:CA:368:U:O2' | 2.12 | 0.49 |
| 1:CA:693:G:O2' | 1:CA:694:A:H5' | 2.13 | 0.49 |
| 1:CA:1076:C:N3 | 1:CA:1082:G:C2 | 2.81 | 0.49 |
| 2:CB:97:TRP:HH2 | 2:CB:176:GLU:CD | 2.21 | 0.49 |
| 3:CC:66:VAL:HG11 | 3:CC:91:LEU:HD13 | 1.95 | 0.49 |
| 12:CL:55:VAL:HG13 | 12:CL:68:ALA:O | 2.13 | 0.49 |
| 13:CM:44:ARG:HB2 | 13:CM:47:ASP:OD1 | 2.13 | 0.49 |
| 23:D1:89:GLU:O | 23:D1:93:GLU:N | 2.41 | 0.49 |
| 30:D8:35:GLN:HE21 | 30:D8:36:LYS:NZ | 2.11 | 0.49 |
| 31:DA:1043:C:OP2 | 31:DA:1043:C:C6 | 2.66 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1410:G:C5 | 31:DA:1411:C:C5 | 3.00 | 0.49 |
| 31:DA:1651:G:N2 | 31:DA:2007:C:C2 | 2.81 | 0.49 |
| 31:DA:1794:U:H2' | 31:DA:1795:C:H6 | 1.78 | 0.49 |
| 31:DA:1952:A:C5 | 40:DO:22:ILE:HD11 | 2.48 | 0.49 |
| 31:DA:2314:C:C2' | 31:DA:2315:G:H5' | 2.42 | 0.49 |
| 31:DA:2328:A:H2' | 31:DA:2329:G:O4' | 2.12 | 0.49 |
| 31:DA:2400:G:H2' | 31:DA:2400:G:N3 | 2.27 | 0.49 |
| 31:DA:2801(A):A:C3' | 31:DA:2802:G:H5' | 2.43 | 0.49 |
| 33:DD:28:GLU:HB2 | 33:DD:29:PRO:CD | 2.42 | 0.49 |
| 36:DG:131:TYR:HB3 | 36:DG:159:VAL:CG1 | 2.40 | 0.49 |
| 39:DN:79:PRO:CD | 39:DN:80:GLY:H | 2.26 | 0.49 |
| 39:DN:79:PRO:HD2 | 39:DN:81:GLY:H | 1.78 | 0.49 |
| 41:DP:88:LEU:O | 41:DP:89:ALA:C | 2.56 | 0.49 |
| 42:DQ:16:ARG:HB2 | 42:DQ:16:ARG:NH1 | 2.25 | 0.49 |
| 50:DY:88:LYS:O | 50:DY:89:PHE:CB | 2.61 | 0.49 |
| 51:DZ:56:VAL:HG12 | 51:DZ:57:ILE:N | 2.28 | 0.49 |
| 1:AA:70:G:H2' | 1:AA:71:C:C6 | 2.48 | 0.48 |
| 1:AA:189(F):U:C2 | 17:AQ:72:ARG:NH1 | 2.81 | 0.48 |
| 1:AA:366:C:O2' | 1:AA:367:U:H5" | 2.12 | 0.48 |
| 1:AA:437:U:C5' | 4:AD:155:LEU:HD13 | 2.43 | 0.48 |
| 1:AA:1501:C:H5" | 1:AA:1502:A:OP2 | 2.13 | 0.48 |
| 7:AG:49:ILE:HG22 | 7:AG:49:ILE:O | 2.13 | 0.48 |
| 7:AG:108:ALA:O | 7:AG:111:ARG:HB2 | 2.13 | 0.48 |
| 10:AJ:54:PHE:CZ | 10:AJ:55:LYS:NZ | 2.81 | 0.48 |
| 31:BA:38:A:C2 | 31:BA:442:G:C2 | 3.00 | 0.48 |
| 31:BA:330:A:H2 | 31:BA:1210:A:HO2' | 1.58 | 0.48 |
| 31:BA:580:C:H2' | 31:BA:581:C:H6 | 1.77 | 0.48 |
| 31:BA:620:G:N3 | 31:BA:620:G:H5' | 2.28 | 0.48 |
| 31:BA:1486:A:H2' | 31:BA:1487:G:H8 | 1.78 | 0.48 |
| 31:BA:1827:C:OP2 | 33:BD:222:ARG:HD2 | 2.13 | 0.48 |
| 31:BA:1886:C:O5' | 31:BA:1886:C:H6 | 1.96 | 0.48 |
| 31:BA:2243:U:O2 | 31:BA:2434:A:C2 | 2.66 | 0.48 |
| 31:BA:2245:U:H5' | 31:BA:2246:G:H5' | 1.94 | 0.48 |
| 31:BA:2801:A:O2' | 31:BA:2895:U:H4' | 2.13 | 0.48 |
| 33:BD:118:VAL:HG22 | 33:BD:119:ALA:H | 1.76 | 0.48 |
| 33:BD:183:ARG:HD2 | 33:BD:270:ILE:CG2 | 2.43 | 0.48 |
| 36:BG:89:GLY:O | 36:BG:90:LEU:C | 2.56 | 0.48 |
| 38:BI:88:ILE:HG12 | 38:BI:122:GLU:N | 2.28 | 0.48 |
| 40:BO:17:ARG:HG2 | 40:BO:47:ILE:HD13 | 1.95 | 0.48 |
| 42:BQ:133:ARG:O | 42:BQ:134:ARG:CB | 2.61 | 0.48 |
| 44:BS:93:LYS:O | 44:BS:93:LYS:HG3 | 2.12 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 46:BU:83:LEU:CB | 46:BU:88:ILE:HD11 | 2.35 | 0.48 |
| 47:BV:5:VAL:HB | 47:BV:60:GLU:OE1 | 2.13 | 0.48 |
| 47:BV:19:LYS:HG3 | 47:BV:20:LEU:O | 2.12 | 0.48 |
| 48:BW:45:TYR:CZ | 48:BW:49:LYS:HE3 | 2.48 | 0.48 |
| 48:BW:57:ASN:O | 48:BW:58:ALA:C | 2.56 | 0.48 |
| 50:BY:45:VAL:CG1 | 50:BY:60:PHE:O | 2.54 | 0.48 |
| 50:BY:83:THR:HG21 | 50:BY:94:LYS:HD2 | 1.95 | 0.48 |
| 1:CA:132:C:O2' | 1:CA:133:U:H5' | 2.13 | 0.48 |
| 1:CA:189(F):U:C2 | 17:CQ:72:ARG:NH1 | 2.81 | 0.48 |
| 1:CA:255:G:O6 | 1:CA:266:G:O6 | 2.31 | 0.48 |
| 1:CA:658:G:C5 | 1:CA:659:U:C5 | 3.01 | 0.48 |
| 1:CA:868:C:H2' | 1:CA:869:G:O4' | 2.13 | 0.48 |
| 1:CA:964:A:N3 | 1:CA:969:A:O2' | 2.37 | 0.48 |
| 1:CA:1067:A:H1' | 1:CA:1068:G:H8 | 1.75 | 0.48 |
| 1:CA:1248:A:C2' | 1:CA:1249:C:H5' | 2.43 | 0.48 |
| 1:CA:1366:C:OP1 | 9:CI:117:HIS:HE1 | 1.96 | 0.48 |
| 1:CA:1399:C:H4' | 1:CA:1400:C:C5' | 2.43 | 0.48 |
| 1:CA:1496:C:H2' | 1:CA:1497:G:O4' | 2.12 | 0.48 |
| 2:CB:167:PRO:HG3 | 2:CB:188:ALA:HB2 | 1.95 | 0.48 |
| 2:CB:185:ILE:HA | 2:CB:199:TYR:O | 2.13 | 0.48 |
| 8:CH:9:MET:O | 8:CH:12:ARG:HB2 | 2.13 | 0.48 |
| 10:CJ:54:PHE:HZ | 10:CJ:55:LYS:HZ1 | 1.59 | 0.48 |
| 15:CO:37:ASN:N | 15:CO:37:ASN:ND2 | 2.58 | 0.48 |
| 31:DA:272(B):G:O2' | 31:DA:272(C):G:C5' | 2.61 | 0.48 |
| 31:DA:924:C:H2' | 31:DA:925:C:H6 | 1.77 | 0.48 |
| 31:DA:941:A:C2 | 31:DA:942:G:C4 | 3.00 | 0.48 |
| 31:DA:1856:G:C2' | 31:DA:1857:G:H5' | 2.43 | 0.48 |
| 31:DA:1899:G:O2' | 31:DA:1900:A:H5" | 2.12 | 0.48 |
| 31:DA:2206:G:N2 | 31:DA:2207:G:C5' | 2.45 | 0.48 |
| 31:DA:2259:G:C2 | 31:DA:2282:G:N1 | 2.81 | 0.48 |
| 31:DA:2753:A:H1' | 31:DA:2754:U:H5' | 1.95 | 0.48 |
| 34:DE:34:VAL:HG22 | 34:DE:48:GLN:HE21 | 1.78 | 0.48 |
| 35:DF:203:GLN:O | 35:DF:206:ILE:C | 2.56 | 0.48 |
| 39:DN:27:ALA:CB | 39:DN:106:MET:HE2 | 2.43 | 0.48 |
| 40:DO:23:ARG:NH1 | 40:DO:23:ARG:CG | 2.71 | 0.48 |
| 41:DP:55:ARG:O | 41:DP:56:SER:OG | 2.31 | 0.48 |
| 45:DT:23:ARG:CB | 45:DT:24:PRO:CD | 2.89 | 0.48 |
| 51:DZ:143:GLY:O | 51:DZ:144:LEU:HD13 | 2.13 | 0.48 |
| 1:AA:355:C:O2' | 1:AA:356:A:H5' | 2.12 | 0.48 |
| 1:AA:414:A:C4 | 1:AA:415:A:C8 | 3.00 | 0.48 |
| 1:AA:663:A:O2' | 1:AA:664:G:H5' | 2.14 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:735:C:H1' | 18:AR:75:ILE:HD11 | 1.95 | 0.48 |
| 1:AA:865:A:H2 | 1:AA:918:A:H4' | 1.79 | 0.48 |
| 1:AA:1255:G:H2' | 1:AA:1255:G:N3 | 2.28 | 0.48 |
| 3:AC:102:ASN:O | 3:AC:103:VAL:HG23 | 2.13 | 0.48 |
| 4:AD:43:HIS:HA | 4:AD:46:LYS:HE3 | 1.94 | 0.48 |
| 4:AD:79:PHE:C | 4:AD:79:PHE:CD2 | 2.92 | 0.48 |
| 6:AF:76:ALA:O | 6:AF:80:ARG:HG3 | 2.13 | 0.48 |
| 9:AI:8:GLY:HA2 | 9:AI:79:LEU:HD12 | 1.95 | 0.48 |
| 9:AI:10:ARG:HG2 | 9:AI:104:ARG:O | 2.13 | 0.48 |
| 11:AK:108:ILE:HB | 18:AR:87:ARG:HA | 1.95 | 0.48 |
| 16:AP:20:VAL:HG22 | 16:AP:21:VAL:H | 1.78 | 0.48 |
| 27:B5:8:LYS:C | 27:B5:9:LYS:HG2 | 2.37 | 0.48 |
| 28:B6:12:GLU:CA | 28:B6:23:THR:HA | 2.35 | 0.48 |
| 31:BA:695:G:C2 | 31:BA:768:G:C5 | 3.02 | 0.48 |
| 31:BA:846:C:H4' | 31:BA:847:U:O5' | 2.13 | 0.48 |
| 31:BA:864:G:C6 | 31:BA:865:C:N4 | 2.81 | 0.48 |
| 31:BA:1043:C:OP2 | 31:BA:1043:C:C6 | 2.67 | 0.48 |
| 31:BA:1047:G:C2 | 31:BA:1111:A:N6 | 2.81 | 0.48 |
| 31:BA:1141:U:H4' | 31:BA:1142(A):A:O4' | 2.13 | 0.48 |
| 31:BA:1352:U:O2' | 31:BA:1353:A:H5' | 2.14 | 0.48 |
| 31:BA:1858:G:H8 | 31:BA:1858:G:OP2 | 1.95 | 0.48 |
| 31:BA:2317:C:O2' | 31:BA:2318:G:H5' | 2.12 | 0.48 |
| 32:BB:65:C:N4 | 32:BB:109:C:C2 | 2.81 | 0.48 |
| 33:BD:145:VAL:HG11 | 33:BD:175:LEU:HD11 | 1.95 | 0.48 |
| 35:BF:18:ARG:HG2 | 35:BF:19:GLU:N | 2.17 | 0.48 |
| 36:BG:114:ILE:CD1 | 36:BG:140:ILE:HD12 | 2.43 | 0.48 |
| 37:BH:40:GLU:O | 37:BH:41:MET:HE3 | 2.12 | 0.48 |
| 42:BQ:93:TYR:N | 42:BQ:93:TYR:CD1 | 2.81 | 0.48 |
| 47:BV:1:MET:HA | 47:BV:1:MET:CE | 2.42 | 0.48 |
| 47:BV:72:VAL:O | 47:BV:73:SER:OG | 2.26 | 0.48 |
| 49:BX:77:LYS:CD | 49:BX:78:LYS:HG3 | 2.42 | 0.48 |
| 1:CA:247:G:C6 | 1:CA:248:C:C5 | 3.01 | 0.48 |
| 1:CA:579:G:C6 | 1:CA:580:U:C4 | 3.01 | 0.48 |
| 1:CA:693:G:O2' | 7:CG:82:GLY:HA3 | 2.14 | 0.48 |
| 1:CA:890:G:O2' | 1:CA:906:G:O6 | 2.25 | 0.48 |
| 1:CA:1068:G:N3 | 1:CA:1191:A:C2 | 2.81 | 0.48 |
| 1:CA:1371:G:C6 | 1:CA:1372:U:C4 | 3.01 | 0.48 |
| 1:CA:1412:C:C2 | 1:CA:1489:G:N2 | 2.81 | 0.48 |
| 5:CE:78:HIS:CE1 | 5:CE:142:LEU:HA | 2.45 | 0.48 |
| 5:CE:105:VAL:HG21 | 5:CE:128:PRO:HA | 1.94 | 0.48 |
| 5:CE:137:GLU:O | 5:CE:141:GLN:HG3 | 2.13 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 15:CO:9:GLN:O | 15:CO:10:LYS:C | 2.55 | 0.48 |
| 16:CP:20:VAL:HG22 | 16:CP:21:VAL:H | 1.77 | 0.48 |
| 18:CR:53:ARG:C | 18:CR:55:ARG:H | 2.20 | 0.48 |
| 30:D8:35:GLN:NE2 | 30:D8:36:LYS:HZ2 | 2.10 | 0.48 |
| 31:DA:142:A:H5" | 31:DA:142(A):C:H5 | 1.78 | 0.48 |
| 31:DA:794:G:H2' | 31:DA:795:C:H6 | 1.77 | 0.48 |
| 31:DA:1163:G:O2' | 31:DA:1164:G:H5' | 2.13 | 0.48 |
| 31:DA:1859:A:C6 | 31:DA:1884:A:C8 | 3.01 | 0.48 |
| 31:DA:2428:G:H5" | 31:DA:2429:G:O5' | 2.13 | 0.48 |
| 31:DA:2500:U:H5" | 31:DA:2501:C:OP2 | 2.13 | 0.48 |
| 31:DA:2537:U:H2' | 31:DA:2538:C:C6 | 2.48 | 0.48 |
| 33:DD:266:SER:O | 33:DD:267:SER:HB2 | 2.12 | 0.48 |
| 36:DG:60:LEU:O | 36:DG:63:ILE:HG13 | 2.12 | 0.48 |
| 36:DG:141:PHE:CE2 | 36:DG:155:MET:HE1 | 2.48 | 0.48 |
| 38:DI:138:ILE:HD12 | 38:DI:138:ILE:N | 2.28 | 0.48 |
| 40:DO:10:VAL:HG13 | 40:DO:17:ARG:O | 2.13 | 0.48 |
| 40:DO:87:ILE:CG2 | 40:DO:88:ASN:N | 2.76 | 0.48 |
| 41:DP:91:PHE:CE2 | 41:DP:95:VAL:HG12 | 2.45 | 0.48 |
| 43:DR:17:ARG:O | 43:DR:20:LEU:HB3 | 2.13 | 0.48 |
| 44:DS:74:ALA:HB1 | 44:DS:103:GLU:HB3 | 1.94 | 0.48 |
| 45:DT:106:SER:O | 45:DT:107:ASP:CB | 2.61 | 0.48 |
| 47:DV:19:LYS:HG3 | 47:DV:20:LEU:O | 2.13 | 0.48 |
| 48:DW:9:TYR:N | 48:DW:102:HIS:CD2 | 2.74 | 0.48 |
| 50:DY:44:ILE:H | 50:DY:44:ILE:CD1 | 2.16 | 0.48 |
| 1:AA:579:G:C6 | 1:AA:580:U:C4 | 3.02 | 0.48 |
| 1:AA:658:G:C1' | 15:AO:22:THR:HB | 2.43 | 0.48 |
| 1:AA:1189:C:O2' | 3:AC:176:HIS:CD2 | 2.66 | 0.48 |
| 2:AB:42:ILE:HD11 | 2:AB:202:PRO:HB2 | 1.94 | 0.48 |
| 11:AK:125:PHE:HD1 | 11:AK:125:PHE:N | 2.10 | 0.48 |
| 12:AL:62:SER:O | 12:AL:64:TYR:N | 2.46 | 0.48 |
| 15:AO:37:ASN:N | 15:AO:37:ASN:HD22 | 2.09 | 0.48 |
| 16:AP:81:ARG:C | 16:AP:82:GLN:HE21 | 2.21 | 0.48 |
| 19:AS:10:PHE:HZ | 19:AS:70:LYS:CE | 2.26 | 0.48 |
| 22:B0:41:ARG:HB2 | 31:BA:2330:G:O2' | 2.13 | 0.48 |
| 28:B6:32:ASN:O | 28:B6:33:LYS:CB | 2.61 | 0.48 |
| 31:BA:307:G:H21 | 31:BA:330:A:H62 | 1.61 | 0.48 |
| 31:BA:481:G:C4 | 31:BA:507:A:C2 | 3.01 | 0.48 |
| 31:BA:661:C:H4' | 41:BP:16:ARG:HH12 | 1.76 | 0.48 |
| 31:BA:1047:G:H21 | 31:BA:1111:A:N6 | 2.10 | 0.48 |
| 31:BA:2068:U:N3 | 31:BA:2430:A:C2 | 2.57 | 0.48 |
| 31:BA:2496:C:P | 42:BQ:81:VAL:HG12 | 2.53 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2523:G:C2' | 31:BA:2524:G:H5" | 2.43 | 0.48 |
| 33:BD:83:GLU:HB2 | 33:BD:92:ILE:HD11 | 1.93 | 0.48 |
| 38:BI:90:GLY:O | 38:BI:91:SER:O | 2.32 | 0.48 |
| 39:BN:65:LYS:HD3 | 39:BN:65:LYS:C | 2.37 | 0.48 |
| 41:BP:40:SER:O | 41:BP:41:ARG:HD2 | 2.13 | 0.48 |
| 43:BR:9:LYS:C | 43:BR:10:LEU:HG | 2.38 | 0.48 |
| 43:BR:13:HIS:HE1 | 43:BR:15:SER:OG | 1.97 | 0.48 |
| 43:BR:18:LEU:C | 43:BR:18:LEU:HD22 | 2.38 | 0.48 |
| 45:BT:106:SER:C | 45:BT:107:ASP:CG | 2.81 | 0.48 |
| 1:CA:70:G:H2' | 1:CA:71:C:C6 | 2.48 | 0.48 |
| 1:CA:78:G:H22 | 1:CA:91:C:N4 | 2.10 | 0.48 |
| 1:CA:264:U:O2' | 17:CQ:64:PRO:HB2 | 2.12 | 0.48 |
| 1:CA:355:C:C2 | 1:CA:356:A:C8 | 3.00 | 0.48 |
| 1:CA:444:C:C2 | 1:CA:445:G:C8 | 3.01 | 0.48 |
| 1:CA:490:G:OP2 | 4:CD:132:ARG:NH2 | 2.46 | 0.48 |
| 1:CA:533:A:C4' | 1:CA:534:U:OP1 | 2.61 | 0.48 |
| 1:CA:620:C:H2' | 1:CA:621:A:O4' | 2.12 | 0.48 |
| 1:CA:828:A:H5" | 1:CA:859:A:C2 | 2.48 | 0.48 |
| 1:CA:986:A:C6 | 1:CA:1220:G:C2 | 3.02 | 0.48 |
| 1:CA:1169:A:C2' | 1:CA:1170:A:H8 | 2.20 | 0.48 |
| 1:CA:1327:C:H2' | 1:CA:1328:C:C6 | 2.48 | 0.48 |
| 2:CB:42:ILE:HD11 | 2:CB:202:PRO:HB2 | 1.95 | 0.48 |
| 6:CF:5:GLU:HG2 | 6:CF:62:TRP:HZ2 | 1.76 | 0.48 |
| 16:CP:19:ILE:HB | 16:CP:37:GLY:O | 2.12 | 0.48 |
| 17:CQ:59:ILE:HD13 | 17:CQ:73:VAL:HA | 1.95 | 0.48 |
| 24:D2:26:ARG:NE | 24:D2:29:LYS:HE2 | 2.27 | 0.48 |
| 28:D6:34:LEU:HD22 | 28:D6:50:ARG:NH1 | 2.28 | 0.48 |
| 31:DA:65:C:H2' | 31:DA:66:C:C6 | 2.48 | 0.48 |
| 31:DA:719:C:O2' | 31:DA:720:C:H5' | 2.13 | 0.48 |
| 31:DA:971:C:H2' | 31:DA:972:G:H5' | 1.93 | 0.48 |
| 31:DA:996:A:N6 | 31:DA:1160:G:C6 | 2.81 | 0.48 |
| 31:DA:1210:A:C4' | 31:DA:1211:U:OP2 | 2.60 | 0.48 |
| 31:DA:1339:G:H21 | 31:DA:1603:A:H1' | 1.78 | 0.48 |
| 31:DA:1508:A:O2' | 31:DA:1509:C:P | 2.71 | 0.48 |
| 31:DA:1682:G:H2' | 31:DA:1683:C:C6 | 2.48 | 0.48 |
| 31:DA:1902:C:C2' | 31:DA:1903:G:O5' | 2.61 | 0.48 |
| 33:DD:17:THR:OG1 | 33:DD:204:ILE:HA | 2.12 | 0.48 |
| 33:DD:35:LYS:HA | 33:DD:64:ILE:CG2 | 2.42 | 0.48 |
| 41:DP:71:VAL:CG1 | 41:DP:72:PRO:HD3 | 2.39 | 0.48 |
| 42:DQ:141:GLN:C | 51:DZ:70:LEU:HD13 | 2.39 | 0.48 |
| 43:DR:41:ALA:O | 43:DR:44:LEU:N | 2.45 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 45:DT:32:TYR:HD2 | 45:DT:81:PRO:O | 1.95 | 0.48 |
| 47:DV:15:GLU:CB | 47:DV:16:PRO:HD2 | 2.35 | 0.48 |
| 48:DW:29:LEU:O | 48:DW:33:ARG:HD2 | 2.12 | 0.48 |
| 1:AA:78:G:H22 | 1:AA:91:C:N4 | 2.11 | 0.48 |
| 1:AA:235:C:H5' | 17:AQ:70:ARG:HG2 | 1.95 | 0.48 |
| 1:AA:394:G:H2' | 1:AA:395:C:H6 | 1.78 | 0.48 |
| 1:AA:948:C:OP1 | 13:AM:107:ALA:HA | 2.13 | 0.48 |
| 1:AA:1157:A:C4 | 1:AA:1181:G:N2 | 2.81 | 0.48 |
| 1:AA:1381:U:H2' | 1:AA:1382:C:H5' | 1.95 | 0.48 |
| 2:AB:114:ARG:HA | 2:AB:117:GLU:HB2 | 1.95 | 0.48 |
| 3:AC:199:LYS:HB3 | 3:AC:201:TYR:HE1 | 1.78 | 0.48 |
| 4:AD:21:LEU:O | 4:AD:113:SER:HB2 | 2.12 | 0.48 |
| 9:AI:45:ALA:O | 9:AI:48:GLU:HB2 | 2.13 | 0.48 |
| 12:AL:18:VAL:O | 12:AL:19:ARG:HB3 | 2.13 | 0.48 |
| 12:AL:83:VAL:HG13 | 12:AL:100:ILE:HG23 | 1.94 | 0.48 |
| 13:AM:3:ARG:HA | 13:AM:9:ILE:HG13 | 1.93 | 0.48 |
| 17:AQ:40:LYS:HD2 | 17:AQ:42:TYR:CZ | 2.47 | 0.48 |
| 23:B1:37:ILE:O | 23:B1:37:ILE:HG23 | 2.14 | 0.48 |
| 23:B1:64:ALA:O | 23:B1:65:SER:HB3 | 2.13 | 0.48 |
| 31:BA:810:U:O2 | 41:BP:33:ARG:HD3 | 2.13 | 0.48 |
| 31:BA:1822:G:H5" | 31:BA:1822:G:H8 | 1.79 | 0.48 |
| 31:BA:2305:A:C5' | 36:BG:134:GLY:HA3 | 2.27 | 0.48 |
| 33:BD:75:ILE:CG2 | 33:BD:99:ASP:HB2 | 2.41 | 0.48 |
| 35:BF:29:ASN:O | 35:BF:30:PRO:C | 2.55 | 0.48 |
| 38:BI:144:VAL:HG12 | 38:BI:145:VAL:N | 2.24 | 0.48 |
| 39:BN:3:THR:O | 39:BN:4:TYR:CD2 | 2.66 | 0.48 |
| 39:BN:39:ARG:NE | 39:BN:41:ASP:HB2 | 2.28 | 0.48 |
| 42:BQ:114:ALA:O | 42:BQ:115:MET:C | 2.56 | 0.48 |
| 43:BR:60:LEU:HD23 | 43:BR:61:HIS:N | 2.28 | 0.48 |
| 47:BV:25:LEU:H | 47:BV:94:LEU:HD12 | 1.79 | 0.48 |
| 48:BW:45:TYR:O | 48:BW:46:PHE:C | 2.55 | 0.48 |
| 49:BX:36:LYS:C | 49:BX:38:GLU:H | 2.20 | 0.48 |
| 50:BY:26:LYS:O | 50:BY:27:VAL:C | 2.56 | 0.48 |
| 50:BY:100:ALA:O | 50:BY:101:LYS:CB | 2.61 | 0.48 |
| 1:CA:7:G:H21 | 5:CE:121:LYS:HG2 | 1.78 | 0.48 |
| 1:CA:438:G:O2' | 1:CA:493:G:C2 | 2.63 | 0.48 |
| 1:CA:668:G:O2' | 15:CO:46:HIS:CD2 | 2.66 | 0.48 |
| 1:CA:961:U:OP2 | 1:CA:1223:C:H4' | 2.14 | 0.48 |
| 1:CA:1075:C:H4' | 1:CA:1101:A:N6 | 2.27 | 0.48 |
| 1:CA:1310:G:N2 | 1:CA:1328:C:C2 | 2.81 | 0.48 |
| 3:CC:125:GLU:HA | 3:CC:191:THR:HG22 | 1.95 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 10:CJ:54:PHE:CZ | 10:CJ:55:LYS:NZ | 2.82 | 0.48 |
| 13:CM:54:VAL:O | 13:CM:58:GLU:HG2 | 2.13 | 0.48 |
| 14:CN:12:ARG:C | 14:CN:14:PRO:HD2 | 2.39 | 0.48 |
| 15:CO:39:LEU:HD22 | 15:CO:43:LEU:HG | 1.94 | 0.48 |
| 18:CR:25:THR:HG22 | 18:CR:42:ARG:HH11 | 1.78 | 0.48 |
| 20:CT:29:LYS:O | 20:CT:33:ILE:HG12 | 2.12 | 0.48 |
| 22:D0:53:MET:CE | 22:D0:57:PHE:CD1 | 2.96 | 0.48 |
| 24:D2:40:SER:O | 24:D2:44:LEU:HB3 | 2.14 | 0.48 |
| 29:D7:12:ARG:HG3 | 31:DA:686:G:O6 | 2.13 | 0.48 |
| 31:DA:197:A:H2' | 31:DA:198:C:C5' | 2.42 | 0.48 |
| 31:DA:828:U:H3' | 31:DA:828:U:O2 | 2.13 | 0.48 |
| 31:DA:1497:U:C2' | 31:DA:1498:C:OP1 | 2.61 | 0.48 |
| 31:DA:1803:A:H4' | 33:DD:259:THR:HG23 | 1.96 | 0.48 |
| 31:DA:1889:A:H2' | 31:DA:1890:A:C8 | 2.48 | 0.48 |
| 31:DA:2036:C:H5' | 31:DA:2036:C:C6 | 2.40 | 0.48 |
| 31:DA:2319:G:C2 | 31:DA:2320:A:N1 | 2.81 | 0.48 |
| 31:DA:2655:G:O2' | 31:DA:2656:U:H5 | 1.96 | 0.48 |
| 32:DB:31:C:H2' | 32:DB:32:C:C6 | 2.48 | 0.48 |
| 33:DD:133:LEU:HD23 | 33:DD:136:ILE:HD12 | 1.95 | 0.48 |
| 34:DE:108:SER:HB3 | 34:DE:165:VAL:HG21 | 1.94 | 0.48 |
| 37:DH:127:GLU:HA | 37:DH:127:GLU:OE1 | 2.13 | 0.48 |
| 38:DI:15:VAL:C | 38:DI:17:GLN:H | 2.19 | 0.48 |
| 41:DP:61:ARG:HD2 | 41:DP:61:ARG:H | 1.79 | 0.48 |
| 43:DR:84:ALA:HB3 | 43:DR:85:PRO:HD3 | 1.96 | 0.48 |
| 49:DX:9:LEU:HB2 | 49:DX:29:TRP:O | 2.12 | 0.48 |
| 50:DY:39:VAL:CG1 | 50:DY:40:GLU:H | 2.11 | 0.48 |
| 51:DZ:144:LEU:HD22 | 51:DZ:144:LEU:N | 2.29 | 0.48 |
| 1:AA:448:A:H2' | 1:AA:449:C:C6 | 2.48 | 0.48 |
| 1:AA:498:U:H2' | 1:AA:498:U:O2 | 2.12 | 0.48 |
| 1:AA:1064:G:H5' | 1:AA:1066:C:H1' | 1.95 | 0.48 |
| 5:AE:78:HIS:CE1 | 5:AE:142:LEU:HA | 2.44 | 0.48 |
| 12:AL:93:LEU:O | 12:AL:94:PRO:C | 2.56 | 0.48 |
| 13:AM:66:LEU:O | 13:AM:70:LEU:HB2 | 2.12 | 0.48 |
| 15:AO:37:ASN:N | 15:AO:37:ASN:ND2 | 2.61 | 0.48 |
| 19:AS:36:ARG:HD2 | 19:AS:52:TYR:O | 2.13 | 0.48 |
| 23:B1:87:PRO:HB2 | 23:B1:91:LYS:NZ | 2.29 | 0.48 |
| 31:BA:142:A:H8 | 31:BA:1595:G:N2 | 2.08 | 0.48 |
| 31:BA:183:C:C2' | 31:BA:184:C:H5' | 2.43 | 0.48 |
| 31:BA:303:U:H2' | 31:BA:304:G:C8 | 2.48 | 0.48 |
| 31:BA:466:A:C3' | 31:BA:467:G:H5' | 2.44 | 0.48 |
| 31:BA:727:A:H2 | 33:BD:9:TYR:CD2 | 2.30 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:754:C:H2' | 31:BA:755:C:H6 | 1.78 | 0.48 |
| 31:BA:822:U:O2' | 31:BA:823:G:H5' | 2.13 | 0.48 |
| 31:BA:1327:C:H2' | 31:BA:1328:G:O4' | 2.13 | 0.48 |
| 31:BA:1362:C:C2' | 31:BA:1363:C:H5' | 2.43 | 0.48 |
| 31:BA:1987:G:H2' | 31:BA:1988:C:H6 | 1.79 | 0.48 |
| 31:BA:2637:U:H2' | 31:BA:2638:G:O5' | 2.14 | 0.48 |
| 32:BB:31:C:H2' | 32:BB:32:C:C6 | 2.48 | 0.48 |
| 33:BD:25:THR:HB | 33:BD:82:ILE:H | 1.77 | 0.48 |
| 33:BD:28:GLU:HB2 | 33:BD:29:PRO:HD3 | 1.95 | 0.48 |
| 35:BF:65:TRP:O | 35:BF:66:PRO:C | 2.57 | 0.48 |
| 35:BF:178:PRO:C | 35:BF:180:GLY:H | 2.22 | 0.48 |
| 36:BG:129:GLY:C | 36:BG:130:ASN:CG | 2.81 | 0.48 |
| 38:BI:9:LEU:N | 38:BI:13:GLY:HA3 | 2.12 | 0.48 |
| 41:BP:13:ASN:H | 41:BP:13:ASN:ND2 | 2.10 | 0.48 |
| 42:BQ:20:ALA:HB2 | 42:BQ:99:PRO:HG2 | 1.94 | 0.48 |
| 42:BQ:23:GLY:O | 42:BQ:99:PRO:O | 2.31 | 0.48 |
| 46:BU:28:ARG:HD3 | 46:BU:38:THR:OG1 | 2.12 | 0.48 |
| 46:BU:88:ILE:H | 46:BU:88:ILE:HD12 | 1.78 | 0.48 |
| 47:BV:73:SER:O | 47:BV:74:LYS:HB2 | 2.13 | 0.48 |
| 50:BY:16:ALA:HA | 50:BY:21:LYS:CD | 2.43 | 0.48 |
| 1:CA:448:A:H2' | 1:CA:449:C:C6 | 2.48 | 0.48 |
| 1:CA:872:A:C4 | 1:CA:874:G:N7 | 2.81 | 0.48 |
| 1:CA:1321:C:H5" | 1:CA:1322:C:H5" | 1.93 | 0.48 |
| 2:CB:168:THR:CG2 | 2:CB:192:SER:HA | 2.43 | 0.48 |
| 12:CL:110:VAL:CG2 | 12:CL:120:TYR:HB3 | 2.43 | 0.48 |
| 23:D1:25:LYS:O | 23:D1:26:ARG:CB | 2.61 | 0.48 |
| 23:D1:37:ILE:HD12 | 31:DA:2079:U:HO2' | 1.77 | 0.48 |
| 23:D1:46:LEU:HA | 31:DA:396:G:O3' | 2.13 | 0.48 |
| 24:D2:52:ASP:H | 24:D2:55:ARG:HB2 | 1.78 | 0.48 |
| 27:D5:9:LYS:HE3 | 31:DA:2019:A:N7 | 2.28 | 0.48 |
| 31:DA:196:A:C4 | 31:DA:805:G:C6 | 3.02 | 0.48 |
| 31:DA:1208:C:C2' | 31:DA:1209:G:H5' | 2.43 | 0.48 |
| 31:DA:1213:A:H1' | 31:DA:1238:G:N3 | 2.28 | 0.48 |
| 31:DA:1327:C:H2' | 31:DA:1328:G:O4' | 2.13 | 0.48 |
| 31:DA:1387:C:C2 | 31:DA:1388:G:C8 | 3.01 | 0.48 |
| 31:DA:1497:U:O2 | 31:DA:1497:U:H2' | 2.13 | 0.48 |
| 31:DA:1819:A:H5" | 33:DD:158:ALA:CB | 2.44 | 0.48 |
| 31:DA:2275:C:H5' | 31:DA:2275:C:H6 | 1.78 | 0.48 |
| 31:DA:2301:C:H2' | 31:DA:2302:G:O4' | 2.13 | 0.48 |
| 31:DA:2312:U:C2' | 31:DA:2313:C:H5' | 2.44 | 0.48 |
| 31:DA:2641:G:H8 | 31:DA:2641:G:H5" | 1.79 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 33:DD:94:LEU:HB2 | 33:DD:104:TYR:CD2 | 2.48 | 0.48 |
| 34:DE:21:VAL:HG23 | 34:DE:21:VAL:O | 2.14 | 0.48 |
| 34:DE:116:VAL:HG11 | 34:DE:122:PHE:CD2 | 2.48 | 0.48 |
| 34:DE:134:ILE:H | 34:DE:134:ILE:HD13 | 1.78 | 0.48 |
| 36:DG:13:GLU:O | 36:DG:13:GLU:HG3 | 2.12 | 0.48 |
| 37:DH:44:VAL:O | 37:DH:46:GLU:OE2 | 2.32 | 0.48 |
| 41:DP:82:GLY:HA2 | 41:DP:113:LYS:O | 2.14 | 0.48 |
| 42:DQ:52:VAL:HA | 42:DQ:55:VAL:HG13 | 1.95 | 0.48 |
| 43:DR:41:ALA:O | 43:DR:42:LYS:C | 2.57 | 0.48 |
| 44:DS:29:PHE:HD2 | 44:DS:29:PHE:C | 2.18 | 0.48 |
| 45:DT:78:LEU:O | 45:DT:79:HIS:ND1 | 2.47 | 0.48 |
| 46:DU:12:ARG:HA | 46:DU:15:LYS:HG2 | 1.96 | 0.48 |
| 48:DW:9:TYR:N | 48:DW:102:HIS:HD2 | 1.94 | 0.48 |
| 48:DW:107:LEU:HD12 | 48:DW:107:LEU:HA | 1.67 | 0.48 |
| 49:DX:89:ILE:O | 49:DX:89:ILE:CG2 | 2.61 | 0.48 |
| 1:AA:130:A:H1' | 1:AA:263:A:O2' | 2.13 | 0.48 |
| 1:AA:134:A:H61 | 16:AP:25:ARG:NH1 | 2.11 | 0.48 |
| 1:AA:197:A:N6 | 1:AA:221:C:H5' | 2.28 | 0.48 |
| 1:AA:371:G:H5" | 1:AA:372:C:OP2 | 2.13 | 0.48 |
| 1:AA:402:G:C6 | 1:AA:403:C:C4 | 3.01 | 0.48 |
| 1:AA:473:G:H2' | 1:AA:474:G:H8 | 1.77 | 0.48 |
| 1:AA:677:U:H3 | 1:AA:714:G:H22 | 1.61 | 0.48 |
| 1:AA:682:G:H1 | 1:AA:708:C:H42 | 1.61 | 0.48 |
| 1:AA:1128:C:H5' | 9:AI:16:ARG:NH1 | 2.29 | 0.48 |
| 1:AA:1321:C:H5" | 1:AA:1322:C:H5" | 1.94 | 0.48 |
| 1:AA:1446:U:O2' | 1:AA:1447:A:H8 | 1.90 | 0.48 |
| 2:AB:168:THR:CG2 | 2:AB:192:SER:HA | 2.43 | 0.48 |
| 3:AC:7:PRO:O | 3:AC:11:ARG:HG2 | 2.13 | 0.48 |
| 31:BA:17:G:H4' | 46:BU:25:TRP:CZ2 | 2.49 | 0.48 |
| 31:BA:142:A:H5" | 31:BA:142(A):C:H5 | 1.78 | 0.48 |
| 31:BA:466:A:H2' | 31:BA:467:G:H5' | 1.96 | 0.48 |
| 31:BA:471:A:H8 | 31:BA:471:A:O5' | 1.95 | 0.48 |
| 31:BA:901:A:H2' | 31:BA:901:A:N3 | 2.27 | 0.48 |
| 31:BA:1499:C:C2' | 31:BA:1500:G:H5' | 2.44 | 0.48 |
| 31:BA:1658:C:H2' | 31:BA:1659:U:C6 | 2.49 | 0.48 |
| 31:BA:2662:A:H4' | 31:BA:2663:G:C8 | 2.48 | 0.48 |
| 31:BA:2875:C:C4' | 45:BT:5:ALA:HB2 | 2.41 | 0.48 |
| 32:BB:8:U:H5' | 32:BB:8:U:C6 | 2.37 | 0.48 |
| 33:BD:35:LYS:NZ | 33:BD:64:ILE:C | 2.71 | 0.48 |
| 36:BG:51:ARG:HB3 | 36:BG:53:LEU:HD23 | 1.95 | 0.48 |
| 37:BH:156:ALA:O | 37:BH:158:HIS:N | 2.46 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 41:BP:101:VAL:HB | 41:BP:107:LYS:H | 1.79 | 0.48 |
| 48:BW:17:VAL:HG21 | 48:BW:103:ILE:HD13 | 1.95 | 0.48 |
| 51:BZ:144:LEU:HD22 | 51:BZ:144:LEU:N | 2.28 | 0.48 |
| 1:CA:577:G:H1' | 1:CA:816:A:C4 | 2.48 | 0.48 |
| 1:CA:586:C:C2' | 1:CA:587:G:H5' | 2.44 | 0.48 |
| 1:CA:716:A:N3 | 11:CK:118:GLY:HA2 | 2.29 | 0.48 |
| 1:CA:825:G:N2 | 8:CH:11:THR:HG21 | 2.29 | 0.48 |
| 1:CA:832:C:O2' | 1:CA:833:U:P | 2.71 | 0.48 |
| 1:CA:862:C:O2' | 1:CA:863:U:H5' | 2.14 | 0.48 |
| 1:CA:967:C:H5" | 1:CA:968:A:OP2 | 2.13 | 0.48 |
| 1:CA:1015:A:C6 | 1:CA:1016:A:C6 | 3.01 | 0.48 |
| 1:CA:1256:A:H5' | 1:CA:1257:U:OP1 | 2.13 | 0.48 |
| 1:CA:1438:G:H2' | 1:CA:1439:C:H6 | 1.78 | 0.48 |
| 2:CB:135:GLN:O | 2:CB:139:LYS:HB2 | 2.13 | 0.48 |
| 3:CC:134:ILE:HG22 | 3:CC:168:ALA:HB3 | 1.96 | 0.48 |
| 4:CD:64:LEU:O | 4:CD:64:LEU:HD12 | 2.13 | 0.48 |
| 13:CM:31:LYS:HA | 13:CM:34:LEU:HD12 | 1.94 | 0.48 |
| 19:CS:42:PRO:O | 19:CS:43:GLU:CB | 2.55 | 0.48 |
| 23:D1:42:GLN:HG2 | 23:D1:43:TYR:N | 2.28 | 0.48 |
| 31:DA:38:A:C2 | 31:DA:442:G:C2 | 3.01 | 0.48 |
| 31:DA:228:A:H2' | 31:DA:230:U:C1' | 2.43 | 0.48 |
| 31:DA:661:C:H4' | 41:DP:16:ARG:HH12 | 1.79 | 0.48 |
| 31:DA:706:A:H2' | 31:DA:707:G:O4' | 2.13 | 0.48 |
| 31:DA:813:U:H2' | 31:DA:814:C:C6 | 2.48 | 0.48 |
| 31:DA:1234:U:O2' | 31:DA:1235:G:H5' | 2.14 | 0.48 |
| 31:DA:1366:A:C2' | 31:DA:1367:A:H5' | 2.43 | 0.48 |
| 31:DA:1639:U:H4' | 31:DA:2699:C:H4' | 1.96 | 0.48 |
| 31:DA:2056:G:C2 | 31:DA:2057:A:C8 | 3.01 | 0.48 |
| 31:DA:2684:U:O2' | 40:DO:68:GLU:HG3 | 2.14 | 0.48 |
| 32:DB:110:G:N1 | 32:DB:111:G:C5 | 2.82 | 0.48 |
| 33:DD:71:ASP:CB | 33:DD:103:ARG:HH22 | 2.26 | 0.48 |
| 33:DD:92:ILE:HD13 | 33:DD:104:TYR:CE2 | 2.48 | 0.48 |
| 34:DE:103:ASP:OD1 | 34:DE:201:THR:HA | 2.14 | 0.48 |
| 35:DF:178:PRO:HG2 | 35:DF:179:GLU:OE1 | 2.13 | 0.48 |
| 36:DG:45:GLU:HB2 | 36:DG:47:LYS:HG3 | 1.94 | 0.48 |
| 36:DG:114:ILE:CD1 | 36:DG:140:ILE:HD12 | 2.43 | 0.48 |
| 37:DH:19:VAL:HB | 37:DH:44:VAL:HG13 | 1.96 | 0.48 |
| 37:DH:116:GLU:HG3 | 37:DH:117:PRO:HD2 | 1.95 | 0.48 |
| 39:DN:67:LEU:O | 39:DN:69:GLN:N | 2.46 | 0.48 |
| 39:DN:82:LEU:H | 39:DN:82:LEU:CD1 | 2.25 | 0.48 |
| 41:DP:30:THR:CG2 | 41:DP:31:ALA:N | 2.75 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 45:DT:83:ILE:HG13 | 45:DT:84:GLN:N | 2.20 | 0.48 |
| 1:AA:376:G:P | 16:AP:67:THR:HG21 | 2.53 | 0.48 |
| 1:AA:542:G:H5' | 4:AD:41:GLY:HA3 | 1.96 | 0.48 |
| 1:AA:1095:U:H2' | 1:AA:1096:C:H6 | 1.78 | 0.48 |
| 4:AD:150:GLU:CD | 4:AD:150:GLU:H | 2.21 | 0.48 |
| 7:AG:100:ALA:O | 7:AG:104:LEU:HD23 | 2.13 | 0.48 |
| 17:AQ:65:ILE:HD12 | 17:AQ:65:ILE:H | 1.77 | 0.48 |
| 17:AQ:81:ARG:NH2 | 17:AQ:83:ASP:OD2 | 2.42 | 0.48 |
| 24:B2:12:GLU:C | 24:B2:14:ARG:N | 2.71 | 0.48 |
| 31:BA:236:C:H2' | 31:BA:237:C:H6 | 1.77 | 0.48 |
| 31:BA:941:A:C2 | 31:BA:942:G:C4 | 3.02 | 0.48 |
| 31:BA:1685:C:C2' | 31:BA:1686:C:O5' | 2.62 | 0.48 |
| 31:BA:1693:U:H4' | 31:BA:1694:C:OP2 | 2.13 | 0.48 |
| 32:BB:46:A:C5 | 32:BB:47:C:C4 | 3.02 | 0.48 |
| 33:BD:50:THR:O | 33:BD:51:VAL:HG23 | 2.14 | 0.48 |
| 33:BD:248:SER:C | 33:BD:250:TRP:H | 2.20 | 0.48 |
| 34:BE:111:ARG:CZ | 43:BR:2:ARG:NH2 | 2.77 | 0.48 |
| 35:BF:57:VAL:HG13 | 35:BF:58:ALA:N | 2.28 | 0.48 |
| 38:BI:52:ARG:HA | 38:BI:52:ARG:HE | 1.78 | 0.48 |
| 38:BI:90:GLY:O | 38:BI:91:SER:C | 2.56 | 0.48 |
| 38:BI:94:ALA:CB | 38:BI:114:LEU:HD12 | 2.42 | 0.48 |
| 43:BR:67:LEU:CD1 | 43:BR:76:VAL:HG21 | 2.43 | 0.48 |
| 44:BS:74:ALA:HB1 | 44:BS:103:GLU:HB3 | 1.94 | 0.48 |
| 45:BT:33:LYS:HA | 45:BT:33:LYS:CE | 2.43 | 0.48 |
| 1:CA:33:A:H2' | 1:CA:34:C:C6 | 2.49 | 0.48 |
| 1:CA:108:G:C6 | 20:CT:15:ARG:HD2 | 2.48 | 0.48 |
| 1:CA:414:A:C5 | 1:CA:431:A:C2 | 3.01 | 0.48 |
| 1:CA:659:U:O2' | 1:CA:660:G:H5' | 2.13 | 0.48 |
| 1:CA:857:C:H2' | 1:CA:858:G:O4' | 2.13 | 0.48 |
| 1:CA:872:A:C2 | 1:CA:874:G:C6 | 3.02 | 0.48 |
| 1:CA:1097:C:O2 | 1:CA:1169:A:H2 | 1.97 | 0.48 |
| 1:CA:1406:U:H2' | 1:CA:1407:C:O4' | 2.14 | 0.48 |
| 1:CA:1442(B):A:N1 | 45:DT:118:ARG:CZ | 2.76 | 0.48 |
| 5:CE:12:LEU:HD13 | 5:CE:31:LEU:HB2 | 1.96 | 0.48 |
| 6:CF:18:GLN:HA | 6:CF:21:LEU:HD23 | 1.94 | 0.48 |
| 9:CI:45:ALA:O | 9:CI:48:GLU:HB2 | 2.14 | 0.48 |
| 14:CN:12:ARG:C | 14:CN:14:PRO:CD | 2.87 | 0.48 |
| 23:D1:18:ILE:HD12 | 23:D1:18:ILE:N | 2.29 | 0.48 |
| 23:D1:47:GLN:CG | 31:DA:2230:G:H1' | 2.30 | 0.48 |
| 23:D1:73:LEU:HD23 | 23:D1:73:LEU:HA | 1.64 | 0.48 |
| 24:D2:29:LYS:O | 24:D2:32:LEU:N | 2.47 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 24:D2:37:PHE:HE2 | 24:D2:40:SER:CA | 2.26 | 0.48 |
| 26:D4:23:GLU:O | 26:D4:24:THR:CB | 2.61 | 0.48 |
| 29:D7:15:THR:HG22 | 29:D7:16:HIS:CG | 2.48 | 0.48 |
| 31:DA:60:G:C8 | 31:DA:63:U:C5 | 3.01 | 0.48 |
| 31:DA:586:A:N1 | 31:DA:809:G:O2' | 2.42 | 0.48 |
| 31:DA:777:A:C2 | 31:DA:778:G:C4 | 3.01 | 0.48 |
| 31:DA:836:G:N7 | 31:DA:837:C:N4 | 2.62 | 0.48 |
| 31:DA:1241:A:C2' | 31:DA:1242:A:O5' | 2.62 | 0.48 |
| 31:DA:1902:C:OP1 | 33:DD:242:ARG:HD2 | 2.13 | 0.48 |
| 31:DA:2391:G:O6 | 31:DA:2425:A:H8 | 1.97 | 0.48 |
| 31:DA:2480:C:N4 | 31:DA:2481:G:C6 | 2.82 | 0.48 |
| 31:DA:2596:U:C2' | 31:DA:2597:G:H5' | 2.44 | 0.48 |
| 31:DA:2748:A:C6 | 31:DA:2749:A:C5 | 3.02 | 0.48 |
| 32:DB:45:A:C2' | 32:DB:46:A:H5' | 2.44 | 0.48 |
| 32:DB:45:A:C4 | 32:DB:46:A:C8 | 3.02 | 0.48 |
| 34:DE:52:LEU:O | 34:DE:74:PRO:CA | 2.54 | 0.48 |
| 38:DI:99:GLU:O | 38:DI:103:ARG:HG3 | 2.14 | 0.48 |
| 39:DN:18:ALA:HB1 | 39:DN:21:LYS:HB2 | 1.95 | 0.48 |
| 39:DN:62:VAL:CG2 | 39:DN:66:LYS:HG3 | 2.42 | 0.48 |
| 41:DP:16:ARG:HD3 | 41:DP:16:ARG:C | 2.38 | 0.48 |
| 41:DP:64:LYS:O | 41:DP:65:ARG:C | 2.54 | 0.48 |
| 42:DQ:141:GLN:OXT | 51:DZ:98:MET:HB2 | 2.14 | 0.48 |
| 43:DR:103:ARG:HD2 | 43:DR:108:GLY:O | 2.13 | 0.48 |
| 44:DS:28:VAL:HG11 | 44:DS:97:ARG:CZ | 2.43 | 0.48 |
| 44:DS:29:PHE:H | 44:DS:89:ARG:CG | 2.26 | 0.48 |
| 48:DW:36:LEU:CD1 | 48:DW:48:ALA:HA | 2.44 | 0.48 |
| 1:AA:299:G:C5 | 1:AA:300:A:C6 | 3.01 | 0.48 |
| 1:AA:340:U:O2' | 1:AA:341:C:H5' | 2.14 | 0.48 |
| 1:AA:447:G:C6 | 1:AA:485:G:H1' | 2.49 | 0.48 |
| 1:AA:458:C:H2' | 1:AA:460:G:C8 | 2.49 | 0.48 |
| 1:AA:502:G:H4' | 1:AA:550:G:H4' | 1.95 | 0.48 |
| 1:AA:1096:C:H2' | 1:AA:1097:C:C6 | 2.46 | 0.48 |
| 1:AA:1151:A:O2' | 1:AA:1152:A:H8 | 1.97 | 0.48 |
| 1:AA:1478:C:H2' | 1:AA:1479:C:H6 | 1.79 | 0.48 |
| 4:AD:80:GLU:O | 4:AD:84:LYS:HG2 | 2.12 | 0.48 |
| 4:AD:133:VAL:HG12 | 4:AD:135:LEU:H | 1.79 | 0.48 |
| 8:AH:65:TYR:N | 8:AH:65:TYR:CD1 | 2.81 | 0.48 |
| 12:AL:40:VAL:HG12 | 12:AL:40:VAL:O | 2.13 | 0.48 |
| 14:AN:12:ARG:C | 14:AN:14:PRO:CD | 2.87 | 0.48 |
| 20:AT:81:LYS:C | 20:AT:83:ARG:H | 2.21 | 0.48 |
| 27:B5:40:LYS:HZ2 | 27:B5:46:CYS:N | 2.12 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 28:B6:18:ARG:HG3 | 28:B6:19:ARG:HG2 | 1.95 | 0.48 |
| 30:B8:48:PHE:CD1 | 30:B8:48:PHE:N | 2.80 | 0.48 |
| 31:BA:975:C:O2 | 31:BA:975:C:H2' | 2.14 | 0.48 |
| 31:BA:1528(A):A:N7 | 31:BA:1529:G:H8 | 2.05 | 0.48 |
| 31:BA:1607:C:H4' | 31:BA:1608:A:O5' | 2.13 | 0.48 |
| 31:BA:2364:C:H2' | 31:BA:2365:G:O4' | 2.13 | 0.48 |
| 31:BA:2544:G:H8 | 31:BA:2544:G:O5' | 1.96 | 0.48 |
| 31:BA:2753:A:O2' | 31:BA:2754:U:C5' | 2.61 | 0.48 |
| 32:BB:13:A:H2' | 32:BB:70:C:O2' | 2.13 | 0.48 |
| 32:BB:41:U:C4 | 36:BG:70:VAL:O | 2.67 | 0.48 |
| 37:BH:22:GLY:C | 37:BH:23:ARG:HG3 | 2.39 | 0.48 |
| 39:BN:13:TRP:CH2 | 39:BN:130:HIS:HE1 | 2.31 | 0.48 |
| 42:BQ:24:GLY:HA3 | 51:BZ:78:LYS:HD2 | 1.95 | 0.48 |
| 47:BV:82:ARG:HD3 | 47:BV:82:ARG:C | 2.38 | 0.48 |
| 49:BX:35:THR:CB | 49:BX:75:ASP:OD2 | 2.60 | 0.48 |
| 50:BY:86:ARG:HG2 | 50:BY:87:LYS:H | 1.79 | 0.48 |
| 1:CA:105:G:H2' | 1:CA:106:C:H6 | 1.77 | 0.48 |
| 1:CA:184:G:N2 | 1:CA:194:C:C2 | 2.82 | 0.48 |
| 1:CA:184:G:O2' | 1:CA:185:A:H5' | 2.13 | 0.48 |
| 1:CA:677:U:H3 | 1:CA:714:G:H22 | 1.61 | 0.48 |
| 1:CA:1530:G:OP1 | 1:CA:1530:G:C4' | 2.61 | 0.48 |
| 4:CD:17:VAL:HG11 | 4:CD:197:PRO:HB2 | 1.95 | 0.48 |
| 4:CD:100:ARG:NH1 | 4:CD:137:SER:HA | 2.29 | 0.48 |
| 7:CG:146:GLU:OE2 | 7:CG:149:ARG:HD2 | 2.14 | 0.48 |
| 12:CL:60:LEU:CD2 | 12:CL:66:VAL:HG22 | 2.41 | 0.48 |
| 12:CL:119:LYS:HB2 | 12:CL:120:TYR:CD1 | 2.49 | 0.48 |
| 23:D1:9:GLY:O | 23:D1:10:LYS:CB | 2.61 | 0.48 |
| 23:D1:34:THR:CG2 | 31:DA:387:U:O3' | 2.62 | 0.48 |
| 25:D3:52:HIS:ND1 | 25:D3:53:LEU:HG | 2.29 | 0.48 |
| 27:D5:55:ARG:CD | 27:D5:56:LYS:N | 2.75 | 0.48 |
| 31:DA:449:A:OP1 | 35:DF:84:VAL:O | 2.32 | 0.48 |
| 31:DA:669:G:H8 | 31:DA:669:G:O2' | 1.90 | 0.48 |
| 31:DA:774:A:C2 | 31:DA:787:U:O2' | 2.64 | 0.48 |
| 31:DA:817:C:O2' | 31:DA:839:U:H5" | 2.13 | 0.48 |
| 31:DA:846:C:H4' | 31:DA:847:U:O5' | 2.14 | 0.48 |
| 31:DA:996:A:C4' | 46:DU:92:ARG:HE | 2.21 | 0.48 |
| 31:DA:1037:G:H1 | 31:DA:1118:C:N4 | 2.12 | 0.48 |
| 31:DA:1862:G:O2' | 31:DA:1863:G:H5' | 2.13 | 0.48 |
| 32:DB:31:C:H2' | 32:DB:32:C:H6 | 1.77 | 0.48 |
| 33:DD:224:ALA:O | 33:DD:225:ALA:HB2 | 2.13 | 0.48 |
| 35:DF:119:ARG:HG2 | 35:DF:119:ARG:O | 2.13 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 38:DI:33:ARG:O | 38:DI:35:LEU:HD23 | 2.14 | 0.48 |
| 41:DP:112:LEU:CD2 | 41:DP:113:LYS:N | 2.77 | 0.48 |
| 47:DV:1:MET:HA | 47:DV:1:MET:CE | 2.41 | 0.48 |
| 47:DV:63:GLY:O | 47:DV:64:HIS:HB3 | 2.13 | 0.48 |
| 49:DX:59:VAL:O | 49:DX:60:ARG:O | 2.32 | 0.48 |
| 51:DZ:100:VAL:O | 51:DZ:123:ASP:HB2 | 2.14 | 0.48 |
| 1:AA:49:U:C2 | 1:AA:361:G:N2 | 2.81 | 0.48 |
| 1:AA:184:G:O2' | 1:AA:185:A:H5' | 2.13 | 0.48 |
| 1:AA:373:A:C2 | 1:AA:374:A:C8 | 3.01 | 0.48 |
| 1:AA:515:G:H2' | 1:AA:516:U:O4' | 2.13 | 0.48 |
| 1:AA:674:G:N2 | 11:AK:116:HIS:HB2 | 2.29 | 0.48 |
| 1:AA:827:U:H2' | 1:AA:859:A:H61 | 1.79 | 0.48 |
| 1:AA:1147:C:C5 | 1:AA:1148:U:C4 | 3.02 | 0.48 |
| 1:AA:1151:A:O2' | 1:AA:1152:A:C8 | 2.66 | 0.48 |
| 3:AC:58:GLU:H | 3:AC:65:ALA:CB | 2.26 | 0.48 |
| 8:AH:109:ILE:CG2 | 8:AH:137:VAL:HB | 2.44 | 0.48 |
| 12:AL:22:SER:C | 12:AL:24:VAL:H | 2.21 | 0.48 |
| 20:AT:94:ALA:O | 20:AT:95:ALA:HB3 | 2.13 | 0.48 |
| 23:B1:49:VAL:HB | 23:B1:64:ALA:HB2 | 1.95 | 0.48 |
| 30:B8:8:LYS:HE2 | 31:BA:243:U:OP2 | 2.14 | 0.48 |
| 31:BA:68:G:H2' | 31:BA:69:C:C6 | 2.49 | 0.48 |
| 31:BA:265:A:H1' | 31:BA:266:G:O4' | 2.13 | 0.48 |
| 31:BA:428:A:H8 | 31:BA:428:A:OP2 | 1.96 | 0.48 |
| 31:BA:1317:A:H2' | 31:BA:1318:C:C6 | 2.47 | 0.48 |
| 31:BA:1508:A:O2' | 31:BA:1509:C:P | 2.72 | 0.48 |
| 31:BA:1515:G:H2' | 31:BA:1516:C:H6 | 1.78 | 0.48 |
| 31:BA:1527:G:C5' | 31:BA:1528:A:OP1 | 2.58 | 0.48 |
| 31:BA:1696:G:C2' | 31:BA:1697:G:C5' | 2.81 | 0.48 |
| 31:BA:2751:G:H3' | 31:BA:2752:C:H6 | 1.78 | 0.48 |
| 32:BB:31:C:N4 | 44:BS:32:LEU:CD2 | 2.77 | 0.48 |
| 35:BF:129:PHE:CD2 | 35:BF:163:VAL:HG21 | 2.48 | 0.48 |
| 36:BG:71:THR:CG2 | 36:BG:72:ARG:N | 2.77 | 0.48 |
| 38:BI:75:LEU:HD11 | 38:BI:105:HIS:HE1 | 1.79 | 0.48 |
| 39:BN:125:GLY:HA2 | 39:BN:126:PRO:O | 2.13 | 0.48 |
| 41:BP:45:LEU:HD22 | 41:BP:46:LYS:N | 2.29 | 0.48 |
| 41:BP:48:PRO:HG2 | 41:BP:49:ARG:H | 1.79 | 0.48 |
| 46:BU:91:ASP:O | 46:BU:92:ARG:O | 2.32 | 0.48 |
| 46:BU:112:ARG:HH11 | 46:BU:112:ARG:HG3 | 1.74 | 0.48 |
| 1:CA:24:U:O2' | 1:CA:25:C:H5' | 2.13 | 0.48 |
| 1:CA:346:G:H5" | 45:DT:41:ARG:CZ | 2.44 | 0.48 |
| 1:CA:349:A:O2' | 1:CA:350:G:H5' | 2.13 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:394:G:H2' | 1:CA:395:C:H6 | 1.78 | 0.48 |
| 1:CA:451:A:C5 | 1:CA:481:G:C5 | 3.02 | 0.48 |
| 1:CA:920:U:H1' | 1:CA:1080:A:C2 | 2.49 | 0.48 |
| 4:CD:104:VAL:O | 4:CD:108:LEU:HD13 | 2.13 | 0.48 |
| 19:CS:51:VAL:HG21 | 19:CS:71:LEU:HB3 | 1.95 | 0.48 |
| 22:D0:84:LEU:HD12 | 22:D0:84:LEU:N | 2.29 | 0.48 |
| 24:D2:34:GLU:OE2 | 24:D2:34:GLU:N | 2.35 | 0.48 |
| 25:D3:28:LEU:HA | 25:D3:33:GLN:OE1 | 2.13 | 0.48 |
| 31:DA:271(P):C:H2' | 31:DA:271(Q):G:H5' | 1.95 | 0.48 |
| 31:DA:344:G:O2' | 31:DA:345:A:H5' | 2.14 | 0.48 |
| 31:DA:352:G:O2' | 31:DA:353:G:O5' | 2.23 | 0.48 |
| 31:DA:378:C:C2' | 31:DA:379:G:H5' | 2.44 | 0.48 |
| 31:DA:394:A:C6 | 31:DA:395:U:C4 | 3.02 | 0.48 |
| 31:DA:883:G:H1 | 31:DA:893:C:H41 | 1.62 | 0.48 |
| 31:DA:892:G:H3' | 31:DA:892:G:N3 | 2.28 | 0.48 |
| 31:DA:1740:G:H4' | 31:DA:1741:A:OP1 | 2.14 | 0.48 |
| 31:DA:1804:C:O5' | 31:DA:1804:C:H6 | 1.96 | 0.48 |
| 31:DA:2012:G:C4' | 48:DW:96:ILE:HD11 | 2.24 | 0.48 |
| 31:DA:2061:G:H5" | 31:DA:2503:A:C2 | 2.49 | 0.48 |
| 31:DA:2394:C:H2' | 31:DA:2395:C:C5' | 2.43 | 0.48 |
| 31:DA:2416:C:H6 | 31:DA:2416:C:O5' | 1.96 | 0.48 |
| 31:DA:2543:G:H2' | 31:DA:2544:G:C8 | 2.49 | 0.48 |
| 31:DA:2662:A:H4' | 31:DA:2663:G:O4' | 2.13 | 0.48 |
| 31:DA:2772:C:H2' | 31:DA:2773:C:H6 | 1.79 | 0.48 |
| 31:DA:2842:G:C2' | 31:DA:2843:G:H5' | 2.44 | 0.48 |
| 32:DB:40:U:C2' | 32:DB:41:U:OP1 | 2.62 | 0.48 |
| 32:DB:52:A:O2' | 32:DB:53:A:C8 | 2.63 | 0.48 |
| 32:DB:78:A:C2 | 32:DB:100:A:C4 | 3.02 | 0.48 |
| 35:DF:1:MET:O | 35:DF:2:LYS:O | 2.32 | 0.48 |
| 36:DG:56:ALA:HB2 | 36:DG:153:ARG:NH2 | 2.28 | 0.48 |
| 36:DG:82:LEU:O | 36:DG:83:ARG:HG3 | 2.13 | 0.48 |
| 42:DQ:38:GLU:HB2 | 42:DQ:127:ILE:CG2 | 2.44 | 0.48 |
| 42:DQ:70:PRO:HA | 42:DQ:95:ALA:HB2 | 1.96 | 0.48 |
| 44:DS:35:ILE:O | 44:DS:35:ILE:HG23 | 2.13 | 0.48 |
| 45:DT:45:PHE:HE2 | 45:DT:63:VAL:HG22 | 1.79 | 0.48 |
| 46:DU:95:LEU:HD11 | 47:DV:11:GLN:HG3 | 1.95 | 0.48 |
| 47:DV:46:VAL:O | 47:DV:47:VAL:HB | 2.13 | 0.48 |
| 48:DW:57:ASN:O | 48:DW:58:ALA:C | 2.56 | 0.48 |
| 1:AA:105:G:H2' | 1:AA:106:C:H6 | 1.75 | 0.48 |
| 1:AA:139:G:H2' | 1:AA:140:A:H8 | 1.79 | 0.48 |
| 1:AA:262:A:C6 | 1:AA:263:A:C6 | 3.02 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:344:A:H4' | 1:AA:345:C:OP2 | 2.13 | 0.48 |
| 1:AA:353:A:H8 | 1:AA:353:A:C5' | 2.21 | 0.48 |
| 1:AA:382:A:C2 | 1:AA:383:A:C4 | 3.02 | 0.48 |
| 1:AA:389:A:H2' | 1:AA:390:C:H5' | 1.95 | 0.48 |
| 1:AA:391:G:C6 | 1:AA:392:G:C5 | 3.02 | 0.48 |
| 1:AA:562:C:N4 | 1:AA:884:U:C6 | 2.82 | 0.48 |
| 1:AA:674:G:O2' | 1:AA:675:A:H5' | 2.14 | 0.48 |
| 1:AA:930:C:C4 | 1:AA:931:C:C5 | 3.02 | 0.48 |
| 1:AA:1187:G:C6 | 1:AA:1188:A:C6 | 3.02 | 0.48 |
| 1:AA:1256:A:H5' | 1:AA:1257:U:OP1 | 2.14 | 0.48 |
| 1:AA:1372:U:OP1 | 9:AI:71:SER:HB3 | 2.13 | 0.48 |
| 3:AC:83:ARG:O | 3:AC:86:VAL:HG22 | 2.14 | 0.48 |
| 3:AC:108:ASN:HB3 | 3:AC:111:LEU:HB2 | 1.96 | 0.48 |
| 4:AD:31:CYS:C | 4:AD:33:MET:N | 2.61 | 0.48 |
| 4:AD:170:VAL:HG22 | 4:AD:171:GLY:N | 2.29 | 0.48 |
| 5:AE:60:TYR:HE1 | 5:AE:64:ARG:HH21 | 1.58 | 0.48 |
| 20:AT:37:SER:O | 20:AT:41:ILE:HG12 | 2.14 | 0.48 |
| 22:B0:13:GLY:O | 22:B0:14:ARG:CB | 2.38 | 0.48 |
| 24:B2:50:ILE:O | 24:B2:51:ARG:HB3 | 2.13 | 0.48 |
| 30:B8:51:ALA:N | 30:B8:53:PRO:HD2 | 2.29 | 0.48 |
| 31:BA:384:U:O2' | 31:BA:385:C:H5' | 2.13 | 0.48 |
| 31:BA:414:C:O2' | 31:BA:415:A:H5' | 2.14 | 0.48 |
| 31:BA:953:A:C2 | 31:BA:954:G:C8 | 3.01 | 0.48 |
| 31:BA:1106:A:O2' | 31:BA:1107:G:O4' | 2.27 | 0.48 |
| 31:BA:1252:G:C2 | 31:BA:1253:A:C2 | 3.02 | 0.48 |
| 31:BA:2273:A:H2' | 31:BA:2274:A:C8 | 2.48 | 0.48 |
| 31:BA:2464:C:O2' | 31:BA:2465:C:O5' | 2.31 | 0.48 |
| 31:BA:2584:U:O2 | 31:BA:2584:U:O4' | 2.31 | 0.48 |
| 31:BA:2606:C:C2' | 31:BA:2607:G:H5' | 2.43 | 0.48 |
| 31:BA:2661:G:H8 | 31:BA:2662:A:N3 | 2.12 | 0.48 |
| 31:BA:2759:G:C2' | 31:BA:2760:C:H5' | 2.44 | 0.48 |
| 31:BA:2808:U:H2' | 31:BA:2809:A:H5' | 1.94 | 0.48 |
| 32:BB:7:G:H2' | 32:BB:8:U:H5" | 1.95 | 0.48 |
| 33:BD:228:PRO:HD3 | 33:BD:234:GLY:C | 2.39 | 0.48 |
| 34:BE:3:GLY:O | 34:BE:4:ILE:HB | 2.13 | 0.48 |
| 34:BE:49:LEU:HD23 | 34:BE:81:ILE:HG12 | 1.96 | 0.48 |
| 35:BF:21:ALA:C | 35:BF:23:ASP:H | 2.20 | 0.48 |
| 39:BN:66:LYS:HA | 39:BN:69:GLN:HB2 | 1.95 | 0.48 |
| 41:BP:100:LEU:HD12 | 41:BP:100:LEU:HA | 1.69 | 0.48 |
| 45:BT:32:TYR:HD2 | 45:BT:81:PRO:O | 1.97 | 0.48 |
| 45:BT:41:ARG:HH11 | 45:BT:43:GLN:HA | 1.78 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 51:BZ:14:LYS:HB2 | 51:BZ:17:ALA:HB3 | 1.95 | 0.48 |
| 1:CA:977:A:C2' | 1:CA:978:A:H5' | 2.43 | 0.48 |
| 1:CA:1097:C:O2' | 1:CA:1098:C:H5' | 2.14 | 0.48 |
| 1:CA:1142:G:H3' | 1:CA:1143:G:C8 | 2.48 | 0.48 |
| 1:CA:1443:G:N2 | 1:CA:1460:A:H1' | 2.28 | 0.48 |
| 1:CA:1507:A:C2 | 1:CA:1508:G:C4 | 3.02 | 0.48 |
| 4:CD:15:GLU:HG2 | 4:CD:63:LYS:HG3 | 1.96 | 0.48 |
| 5:CE:31:LEU:HD22 | 5:CE:43:LEU:HD11 | 1.96 | 0.48 |
| 6:CF:46:ARG:NH1 | 18:CR:37:VAL:HG21 | 2.24 | 0.48 |
| 6:CF:75:LEU:HD23 | 6:CF:79:LEU:HD21 | 1.96 | 0.48 |
| 9:CI:51:ARG:HE | 9:CI:56:LEU:CD1 | 2.26 | 0.48 |
| 12:CL:46:LYS:HB2 | 12:CL:92:ASP:O | 2.13 | 0.48 |
| 22:D0:50:ASN:ND2 | 22:D0:83:PRO:HD3 | 2.28 | 0.48 |
| 24:D2:23:LYS:O | 24:D2:24:LEU:C | 2.57 | 0.48 |
| 24:D2:33:MET:HG3 | 49:DX:10:ALA:HB1 | 1.96 | 0.48 |
| 27:D5:8:LYS:C | 27:D5:9:LYS:HG2 | 2.38 | 0.48 |
| 28:D6:12:GLU:CA | 28:D6:23:THR:HA | 2.32 | 0.48 |
| 28:D6:32:ASN:O | 28:D6:33:LYS:CB | 2.61 | 0.48 |
| 28:D6:39:TYR:HB3 | 28:D6:49:HIS:ND1 | 2.29 | 0.48 |
| 30:D8:48:PHE:CD1 | 30:D8:48:PHE:N | 2.80 | 0.48 |
| 31:DA:118:A:H3' | 31:DA:119:A:C5' | 2.44 | 0.48 |
| 31:DA:428:A:H8 | 31:DA:428:A:OP2 | 1.97 | 0.48 |
| 31:DA:559:G:H22 | 46:DU:49:HIS:CD2 | 2.32 | 0.48 |
| 31:DA:671:C:H2' | 31:DA:672:C:C6 | 2.48 | 0.48 |
| 31:DA:952:G:C6 | 31:DA:966:G:C6 | 3.02 | 0.48 |
| 31:DA:1299:G:H5" | 31:DA:1300:U:OP1 | 2.14 | 0.48 |
| 31:DA:1488:G:C5 | 31:DA:1489:U:N3 | 2.82 | 0.48 |
| 31:DA:1646:C:H5" | 31:DA:1647:G:H5" | 1.94 | 0.48 |
| 31:DA:1822:G:H5' | 31:DA:1822:G:C8 | 2.48 | 0.48 |
| 31:DA:2567:G:H2' | 31:DA:2568:C:H6 | 1.78 | 0.48 |
| 31:DA:2569:G:C2' | 31:DA:2570:G:H5' | 2.44 | 0.48 |
| 31:DA:2655:G:HO2' | 31:DA:2656:U:H5 | 1.60 | 0.48 |
| 33:DD:108:PRO:HD2 | 33:DD:111:LEU:HG | 1.96 | 0.48 |
| 33:DD:228:PRO:HD3 | 33:DD:234:GLY:C | 2.38 | 0.48 |
| 36:DG:45:GLU:HG2 | 36:DG:47:LYS:H | 1.79 | 0.48 |
| 36:DG:109:VAL:C | 36:DG:112:PRO:HD2 | 2.39 | 0.48 |
| 37:DH:19:VAL:HG21 | 37:DH:44:VAL:HA | 1.96 | 0.48 |
| 37:DH:153:LYS:N | 37:DH:153:LYS:CD | 2.75 | 0.48 |
| 42:DQ:38:GLU:HB2 | 42:DQ:127:ILE:HG22 | 1.95 | 0.48 |
| 1:AA:67:C:O2' | 1:AA:171:A:H1' | 2.14 | 0.47 |
| 1:AA:192:U:C1' | 20:AT:103:GLY:HA2 | 2.43 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:504:C:O2 | 1:AA:504:C:H2' | 2.13 | 0.47 |
| 1:AA:542:G:O2' | 1:AA:543:C:H5' | 2.14 | 0.47 |
| 1:AA:1159:U:C5 | 1:AA:1182:G:C4 | 3.02 | 0.47 |
| 1:AA:1310:G:N2 | 1:AA:1328:C:C2 | 2.83 | 0.47 |
| 1:AA:1366:C:OP1 | 9:AI:117:HIS:HE1 | 1.95 | 0.47 |
| 1:AA:1452:C:H4' | 1:AA:1456:G:O5' | 2.15 | 0.47 |
| 2:AB:69:LEU:HB3 | 2:AB:162:ILE:HG22 | 1.96 | 0.47 |
| 4:AD:68:TYR:N | 4:AD:68:TYR:CD1 | 2.81 | 0.47 |
| 4:AD:108:LEU:HD12 | 4:AD:174:LEU:HD13 | 1.95 | 0.47 |
| 11:AK:41:THR:HG22 | 11:AK:42:TRP:N | 2.28 | 0.47 |
| 16:AP:54:GLU:O | 16:AP:57:ARG:HB2 | 2.14 | 0.47 |
| 19:AS:16:LEU:O | 19:AS:20:LEU:HB2 | 2.14 | 0.47 |
| 20:AT:26:ASN:CB | 20:AT:71:THR:HG23 | 2.43 | 0.47 |
| 20:AT:26:ASN:O | 20:AT:30:LYS:HB2 | 2.13 | 0.47 |
| 22:B0:74:ARG:NH2 | 32:BB:13:A:C8 | 2.82 | 0.47 |
| 30:B8:38:GLY:C | 30:B8:40:GLU:N | 2.71 | 0.47 |
| 31:BA:356:G:C2 | 31:BA:357:A:C4 | 3.02 | 0.47 |
| 31:BA:363(A):A:N3 | 31:BA:363(A):A:H2' | 2.29 | 0.47 |
| 31:BA:543:C:N4 | 31:BA:551:G:H1 | 2.11 | 0.47 |
| 31:BA:829:A:N7 | 31:BA:2248:C:H5' | 2.29 | 0.47 |
| 31:BA:979:G:H3' | 31:BA:980:A:C5' | 2.43 | 0.47 |
| 31:BA:2321:G:H5" | 31:BA:2322:A:OP2 | 2.14 | 0.47 |
| 31:BA:2463:C:O2' | 31:BA:2464:C:H5' | 2.14 | 0.47 |
| 33:BD:35:LYS:HE2 | 33:BD:65:ILE:HG22 | 1.95 | 0.47 |
| 33:BD:173:VAL:HG23 | 33:BD:174:ILE:N | 2.29 | 0.47 |
| 33:BD:231:HIS:CG | 33:BD:232:PRO:HD2 | 2.49 | 0.47 |
| 35:BF:132:VAL:C | 35:BF:134:GLY:H | 2.21 | 0.47 |
| 35:BF:160:ASN:ND2 | 35:BF:162:LEU:H | 2.11 | 0.47 |
| 38:BI:64:GLU:N | 38:BI:64:GLU:CD | 2.72 | 0.47 |
| 39:BN:79:PRO:HD2 | 39:BN:81:GLY:H | 1.78 | 0.47 |
| 40:BO:66:LYS:H | 40:BO:82:ASN:HD21 | 1.58 | 0.47 |
| 42:BQ:20:ALA:C | 42:BQ:22:LYS:N | 2.71 | 0.47 |
| 44:BS:53:SER:OG | 44:BS:54:LEU:N | 2.47 | 0.47 |
| 45:BT:28:VAL:HG22 | 45:BT:46:GLU:HG3 | 1.95 | 0.47 |
| 47:BV:15:GLU:CB | 47:BV:16:PRO:HD2 | 2.33 | 0.47 |
| 50:BY:22:GLY:O | 50:BY:23:ARG:HG2 | 2.13 | 0.47 |
| 50:BY:40:GLU:HA | 50:BY:40:GLU:OE2 | 2.14 | 0.47 |
| 1:CA:1372:U:OP1 | 9:CI:71:SER:HB3 | 2.14 | 0.47 |
| 3:CC:83:ARG:O | 3:CC:86:VAL:HG22 | 2.14 | 0.47 |
| 9:CI:118:LYS:O | 9:CI:119:ALA:HB3 | 2.14 | 0.47 |
| 13:CM:3:ARG:HA | 13:CM:9:ILE:HG13 | 1.95 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 15:CO:67:LEU:CD2 | 15:CO:78:TYR:CE1 | 2.97 | 0.47 |
| 20:CT:96:GLY:O | 20:CT:97:ALA:HB3 | 2.13 | 0.47 |
| 20:CT:100:ILE:HD12 | 20:CT:100:ILE:N | 2.28 | 0.47 |
| 31:DA:30:G:H2' | 31:DA:31:C:C6 | 2.49 | 0.47 |
| 31:DA:197:A:N6 | 31:DA:2430:A:H2' | 2.29 | 0.47 |
| 31:DA:565:C:H5' | 31:DA:1253:A:N6 | 2.29 | 0.47 |
| 31:DA:1044:G:C2 | 31:DA:1112:G:O6 | 2.67 | 0.47 |
| 31:DA:1131:G:H21 | 39:DN:73:THR:CG2 | 2.27 | 0.47 |
| 31:DA:1489:U:C2' | 31:DA:1490:A:OP2 | 2.61 | 0.47 |
| 31:DA:2347:C:H2' | 31:DA:2348:U:C6 | 2.48 | 0.47 |
| 31:DA:2615:U:H2' | 31:DA:2616:C:C6 | 2.49 | 0.47 |
| 31:DA:2682:U:C5 | 34:DE:11:MET:HE2 | 2.49 | 0.47 |
| 35:DF:78:ILE:H | 35:DF:78:ILE:HD13 | 1.79 | 0.47 |
| 41:DP:62:LEU:N | 41:DP:62:LEU:CD2 | 2.68 | 0.47 |
| 42:DQ:17:LEU:HD21 | 42:DQ:41:TRP:HE1 | 1.79 | 0.47 |
| 45:DT:32:TYR:CB | 45:DT:81:PRO:HB2 | 2.43 | 0.47 |
| 46:DU:57:PHE:O | 46:DU:58:ARG:C | 2.57 | 0.47 |
| 47:DV:51:VAL:CG1 | 47:DV:52:VAL:N | 2.76 | 0.47 |
| 47:DV:64:HIS:O | 47:DV:64:HIS:ND1 | 2.47 | 0.47 |
| 48:DW:12:ILE:HG13 | 48:DW:42:ARG:NH1 | 2.29 | 0.47 |
| 49:DX:37:THR:HG23 | 49:DX:54:VAL:CB | 2.37 | 0.47 |
| 49:DX:63:LYS:O | 49:DX:68:ARG:HA | 2.13 | 0.47 |
| 50:DY:28:LYS:HG3 | 50:DY:37:VAL:HA | 1.95 | 0.47 |
| 1:AA:445:G:N3 | 1:AA:446:G:C8 | 2.81 | 0.47 |
| 1:AA:1068:G:N3 | 1:AA:1191:A:C2 | 2.82 | 0.47 |
| 1:AA:1076:C:N3 | 1:AA:1082:G:C2 | 2.83 | 0.47 |
| 1:AA:1496:C:H2' | 1:AA:1497:G:O4' | 2.14 | 0.47 |
| 2:AB:111:ARG:O | 2:AB:145:LEU:HD11 | 2.14 | 0.47 |
| 4:AD:68:TYR:H | 4:AD:68:TYR:HD1 | 1.61 | 0.47 |
| 5:AE:69:VAL:HG12 | 5:AE:71:LEU:CD2 | 2.44 | 0.47 |
| 5:AE:147:ASP:N | 5:AE:147:ASP:OD2 | 2.46 | 0.47 |
| 6:AF:72:VAL:CG2 | 6:AF:90:VAL:HG11 | 2.43 | 0.47 |
| 8:AH:21:LYS:O | 8:AH:22:GLU:C | 2.58 | 0.47 |
| 18:AR:25:THR:HG22 | 18:AR:42:ARG:HH11 | 1.80 | 0.47 |
| 18:AR:31:LEU:H | 18:AR:31:LEU:CD2 | 2.24 | 0.47 |
| 22:B0:27:GLU:HG3 | 22:B0:68:GLU:HA | 1.96 | 0.47 |
| 23:B1:33:LYS:HB3 | 31:BA:2395:C:O2' | 2.14 | 0.47 |
| 29:B7:43:THR:HG23 | 29:B7:44:PRO:CD | 2.44 | 0.47 |
| 30:B8:48:PHE:HD1 | 30:B8:48:PHE:H | 1.56 | 0.47 |
| 31:BA:606:U:H4' | 31:BA:658:C:H4' | 1.95 | 0.47 |
| 31:BA:1208:C:H2' | 31:BA:1209:G:H5' | 1.97 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1503:U:H2' | 31:BA:1504:C:C6 | 2.48 | 0.47 |
| 31:BA:1568:G:OP2 | 33:BD:63:ARG:NH2 | 2.46 | 0.47 |
| 31:BA:2305:A:O2' | 36:BG:136:ARG:HD3 | 2.14 | 0.47 |
| 32:BB:30:C:H2' | 32:BB:31:C:H5' | 1.95 | 0.47 |
| 33:BD:71:ASP:OD2 | 33:BD:103:ARG:NH2 | 2.47 | 0.47 |
| 33:BD:79:VAL:HG12 | 33:BD:79:VAL:O | 2.13 | 0.47 |
| 36:BG:37:VAL:CG2 | 36:BG:99:MET:HG3 | 2.44 | 0.47 |
| 37:BH:158:HIS:CE1 | 37:BH:170:ARG:HA | 2.48 | 0.47 |
| 38:BI:138:ILE:HD12 | 38:BI:138:ILE:N | 2.23 | 0.47 |
| 42:BQ:104:PHE:CE1 | 42:BQ:125:LEU:HD11 | 2.42 | 0.47 |
| 43:BR:37:THR:OG1 | 43:BR:40:LYS:HB2 | 2.14 | 0.47 |
| 44:BS:14:VAL:O | 44:BS:15:ARG:C | 2.56 | 0.47 |
| 45:BT:108:ARG:O | 45:BT:111:ARG:HB2 | 2.14 | 0.47 |
| 49:BX:59:VAL:HG23 | 49:BX:74:PRO:CD | 2.41 | 0.47 |
| 50:BY:8:LYS:HE3 | 50:BY:72:VAL:HG23 | 1.94 | 0.47 |
| 50:BY:45:VAL:CG2 | 50:BY:62:GLU:N | 2.78 | 0.47 |
| 51:BZ:143:GLY:O | 51:BZ:144:LEU:HD13 | 2.14 | 0.47 |
| 1:CA:66:G:C4' | 1:CA:173:U:C5 | 2.96 | 0.47 |
| 1:CA:109:A:C6 | 1:CA:326:G:C6 | 3.02 | 0.47 |
| 1:CA:235:C:H5' | 17:CQ:70:ARG:HG2 | 1.95 | 0.47 |
| 1:CA:671:G:C5 | 1:CA:672:U:C5 | 3.02 | 0.47 |
| 1:CA:920:U:H2' | 1:CA:921:U:C6 | 2.48 | 0.47 |
| 1:CA:954:G:C2 | 1:CA:955:U:C2 | 3.02 | 0.47 |
| 1:CA:1202:G:C6 | 14:CN:42:ILE:HG21 | 2.49 | 0.47 |
| 12:CL:105:TYR:C | 12:CL:107:ALA:H | 2.21 | 0.47 |
| 13:CM:14:ARG:CZ | 13:CM:42:ALA:HA | 2.43 | 0.47 |
| 16:CP:54:GLU:O | 16:CP:57:ARG:HB2 | 2.14 | 0.47 |
| 28:D6:11:LEU:O | 28:D6:23:THR:HA | 2.14 | 0.47 |
| 28:D6:27:LYS:HE3 | 31:DA:2285:C:H5 | 1.78 | 0.47 |
| 31:DA:343:C:H2' | 31:DA:343:C:O2 | 2.14 | 0.47 |
| 31:DA:528:A:C2 | 31:DA:2043:C:H4' | 2.49 | 0.47 |
| 31:DA:1272:A:H3' | 31:DA:1273:U:H5" | 1.96 | 0.47 |
| 31:DA:1353:A:H4' | 33:DD:38:LYS:NZ | 2.29 | 0.47 |
| 31:DA:1505:C:C2' | 31:DA:1506:C:O5' | 2.62 | 0.47 |
| 31:DA:1572:A:O5' | 31:DA:1572:A:H8 | 1.96 | 0.47 |
| 31:DA:1887:C:C3' | 31:DA:1888:G:H5' | 2.43 | 0.47 |
| 31:DA:2315:G:C6 | 31:DA:2316:C:N4 | 2.81 | 0.47 |
| 31:DA:2515:C:C2' | 31:DA:2516:G:H5' | 2.45 | 0.47 |
| 31:DA:2645:G:C3' | 31:DA:2646:C:H5' | 2.43 | 0.47 |
| 31:DA:2748:A:N6 | 31:DA:2749:A:C6 | 2.83 | 0.47 |
| 31:DA:2753:A:O2' | 31:DA:2754:U:C5' | 2.61 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 32:DB:88:C:C4 | 32:DB:89:G:C6 | 3.02 | 0.47 |
| 33:DD:14:ARG:O | 33:DD:15:PHE:CD2 | 2.67 | 0.47 |
| 33:DD:52:ARG:CZ | 33:DD:53:PHE:HE2 | 2.27 | 0.47 |
| 33:DD:109:ASP:HB2 | 33:DD:197:GLY:CA | 2.45 | 0.47 |
| 34:DE:144:ARG:HB3 | 34:DE:145:LYS:H | 1.47 | 0.47 |
| 35:DF:53:THR:C | 35:DF:55:GLY:N | 2.70 | 0.47 |
| 36:DG:71:THR:CG2 | 36:DG:72:ARG:N | 2.78 | 0.47 |
| 39:DN:66:LYS:HA | 39:DN:69:GLN:HB2 | 1.95 | 0.47 |
| 41:DP:90:ARG:HD2 | 41:DP:91:PHE:HD1 | 1.78 | 0.47 |
| 43:DR:38:VAL:HB | 43:DR:39:PRO:HD3 | 1.96 | 0.47 |
| 43:DR:116:LEU:HD23 | 43:DR:116:LEU:HA | 1.57 | 0.47 |
| 46:DU:91:ASP:O | 46:DU:92:ARG:HB3 | 2.13 | 0.47 |
| 47:DV:72:VAL:HG13 | 47:DV:88:ARG:NH2 | 2.29 | 0.47 |
| 49:DX:23:GLU:O | 49:DX:25:LYS:N | 2.47 | 0.47 |
| 49:DX:59:VAL:HG23 | 49:DX:74:PRO:CD | 2.41 | 0.47 |
| 50:DY:26:LYS:HG2 | 50:DY:27:VAL:H | 1.78 | 0.47 |
| 50:DY:46:LYS:O | 50:DY:47:LYS:CB | 2.62 | 0.47 |
| 51:DZ:74:VAL:HG22 | 51:DZ:86:VAL:CG1 | 2.44 | 0.47 |
| 1:AA:61:G:OP1 | 20:AT:10:LEU:HD11 | 2.13 | 0.47 |
| 1:AA:79:G:H4' | 1:AA:80:G:OP1 | 2.15 | 0.47 |
| 1:AA:276:G:C2' | 1:AA:277:C:H5' | 2.44 | 0.47 |
| 1:AA:343:U:H2' | 1:AA:346:G:O6 | 2.13 | 0.47 |
| 1:AA:673:G:O3' | 6:AF:87:ARG:NH2 | 2.47 | 0.47 |
| 1:AA:764:C:H2' | 1:AA:765:G:H8 | 1.78 | 0.47 |
| 1:AA:804:U:H5" | 1:AA:805:C:OP2 | 2.14 | 0.47 |
| 1:AA:825:G:N2 | 8:AH:11:THR:HG21 | 2.29 | 0.47 |
| 1:AA:1074:G:N3 | 1:AA:1102:A:C2 | 2.82 | 0.47 |
| 1:AA:1399:C:H4' | 1:AA:1400:C:C5' | 2.44 | 0.47 |
| 1:AA:1432:G:H8 | 1:AA:1432:G:O5' | 1.98 | 0.47 |
| 2:AB:75:LYS:C | 2:AB:77:ALA:H | 2.22 | 0.47 |
| 4:AD:108:LEU:O | 4:AD:110:PHE:N | 2.46 | 0.47 |
| 4:AD:159:ARG:HA | 4:AD:162:LEU:HB2 | 1.95 | 0.47 |
| 8:AH:26:VAL:HG13 | 8:AH:59:LEU:HB2 | 1.96 | 0.47 |
| 9:AI:118:LYS:O | 9:AI:119:ALA:HB3 | 2.14 | 0.47 |
| 20:AT:12:ALA:O | 20:AT:15:ARG:HB2 | 2.13 | 0.47 |
| 20:AT:57:ARG:HD3 | 20:AT:103:GLY:H | 1.79 | 0.47 |
| 22:B0:18:ALA:HB1 | 31:BA:2271:G:OP1 | 2.14 | 0.47 |
| 22:B0:36:ILE:HD13 | 22:B0:36:ILE:C | 2.39 | 0.47 |
| 22:B0:50:ASN:C | 22:B0:62:LEU:HB2 | 2.39 | 0.47 |
| 25:B3:8:LEU:HD13 | 25:B3:31:LEU:HA | 1.96 | 0.47 |
| 31:BA:530:G:O6 | 31:BA:2023:G:OP1 | 2.32 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:670:A:H5" | 31:BA:670:A:H8 | 1.79 | 0.47 |
| 31:BA:777:A:C2 | 31:BA:778:G:C8 | 3.03 | 0.47 |
| 31:BA:870:A:C2 | 31:BA:908:C:C2 | 3.02 | 0.47 |
| 31:BA:1268:A:H2' | 31:BA:1269:A:O4' | 2.14 | 0.47 |
| 31:BA:1525:G:H2' | 31:BA:1526:G:C8 | 2.50 | 0.47 |
| 31:BA:2056:G:C2 | 31:BA:2057:A:C8 | 3.02 | 0.47 |
| 31:BA:2199:A:C8 | 31:BA:2200:C:C5 | 3.02 | 0.47 |
| 31:BA:2752:C:H2' | 31:BA:2753:A:H5' | 1.95 | 0.47 |
| 33:BD:54:ARG:O | 33:BD:218:ARG:HD3 | 2.14 | 0.47 |
| 33:BD:155:LEU:HD23 | 33:BD:177:LEU:HD22 | 1.96 | 0.47 |
| 34:BE:37:ARG:NH1 | 34:BE:80:GLU:OE2 | 2.46 | 0.47 |
| 34:BE:161:GLY:O | 34:BE:162:ALA:HB3 | 2.14 | 0.47 |
| 38:BI:99:GLU:O | 38:BI:103:ARG:HG3 | 2.14 | 0.47 |
| 41:BP:31:ALA:C | 41:BP:33:ARG:H | 2.22 | 0.47 |
| 42:BQ:75:THR:HA | 42:BQ:88:GLY:CA | 2.41 | 0.47 |
| 45:BT:50:ILE:HA | 45:BT:99:LEU:HD11 | 1.95 | 0.47 |
| 50:BY:54:LYS:O | 50:BY:55:TYR:O | 2.31 | 0.47 |
| 1:CA:380:G:C2 | 1:CA:384:G:N1 | 2.82 | 0.47 |
| 1:CA:382:A:H2' | 1:CA:383:A:C8 | 2.50 | 0.47 |
| 1:CA:582:U:C2 | 1:CA:760:G:C6 | 3.02 | 0.47 |
| 1:CA:625:G:C4 | 1:CA:626:U:C5 | 3.02 | 0.47 |
| 1:CA:671:G:C4 | 1:CA:672:U:C6 | 3.02 | 0.47 |
| 1:CA:892:A:H2' | 1:CA:893:C:H6 | 1.78 | 0.47 |
| 1:CA:1385:G:C2' | 1:CA:1386:G:H5' | 2.44 | 0.47 |
| 2:CB:75:LYS:C | 2:CB:77:ALA:H | 2.22 | 0.47 |
| 2:CB:114:ARG:HA | 2:CB:117:GLU:HB2 | 1.96 | 0.47 |
| 4:CD:36:ARG:HG2 | 4:CD:38:TYR:CZ | 2.50 | 0.47 |
| 5:CE:36:ASP:O | 5:CE:37:ARG:HB2 | 2.14 | 0.47 |
| 5:CE:60:TYR:HE1 | 5:CE:64:ARG:HH21 | 1.56 | 0.47 |
| 9:CI:113:LYS:H | 9:CI:119:ALA:HA | 1.79 | 0.47 |
| 17:CQ:45:HIS:O | 17:CQ:73:VAL:HG23 | 2.14 | 0.47 |
| 23:D1:73:LEU:HD13 | 23:D1:90:ILE:CG2 | 2.43 | 0.47 |
| 31:DA:250:G:C6 | 31:DA:251:A:C6 | 3.03 | 0.47 |
| 31:DA:695:G:C2 | 31:DA:768:G:C5 | 3.02 | 0.47 |
| 31:DA:707:G:C6 | 31:DA:708:C:C4 | 3.03 | 0.47 |
| 31:DA:1374:G:H2' | 31:DA:1375:C:C6 | 2.49 | 0.47 |
| 31:DA:1591:G:C5' | 31:DA:1591:G:H8 | 2.26 | 0.47 |
| 31:DA:1892:C:H6 | 31:DA:1892:C:O5' | 1.97 | 0.47 |
| 31:DA:2026:C:H2' | 31:DA:2027:G:O5' | 2.13 | 0.47 |
| 31:DA:2202:C:O2 | 33:DD:151:LYS:NZ | 2.40 | 0.47 |
| 31:DA:2473:U:C4 | 31:DA:2474:C:C5 | 3.03 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 32:DB:10:C:C2' | 32:DB:11:C:H5' | 2.44 | 0.47 |
| 33:DD:109:ASP:HB2 | 33:DD:197:GLY:HA2 | 1.96 | 0.47 |
| 34:DE:10:GLY:HA2 | 34:DE:192:ASN:ND2 | 2.28 | 0.47 |
| 34:DE:201:THR:HG22 | 34:DE:203:LYS:N | 2.30 | 0.47 |
| 43:DR:10:LEU:HD22 | 43:DR:17:ARG:CD | 2.41 | 0.47 |
| 44:DS:56:LEU:O | 44:DS:57:LYS:CB | 2.62 | 0.47 |
| 47:DV:54:GLY:C | 47:DV:56:SER:N | 2.72 | 0.47 |
| 51:DZ:126:VAL:HG12 | 51:DZ:163:LEU:HA | 1.96 | 0.47 |
| 1:AA:20:U:H2' | 1:AA:21:G:O4' | 2.14 | 0.47 |
| 1:AA:352:C:H4' | 1:AA:354:G:OP1 | 2.13 | 0.47 |
| 1:AA:693:G:O2' | 1:AA:694:A:H5' | 2.15 | 0.47 |
| 1:AA:726:C:O2' | 1:AA:727:G:H5' | 2.14 | 0.47 |
| 1:AA:834:C:H2' | 1:AA:835:U:C6 | 2.49 | 0.47 |
| 1:AA:1034:G:N2 | 1:AA:1035:A:N6 | 2.63 | 0.47 |
| 1:AA:1070:U:C2 | 1:AA:1071:C:C5 | 3.02 | 0.47 |
| 1:AA:1099:G:N3 | 1:AA:1099:G:H2' | 2.30 | 0.47 |
| 5:AE:31:LEU:HD22 | 5:AE:43:LEU:HD11 | 1.96 | 0.47 |
| 19:AS:22:LEU:O | 19:AS:26:GLY:HA2 | 2.15 | 0.47 |
| 20:AT:82:SER:O | 20:AT:86:ARG:CB | 2.62 | 0.47 |
| 22:B0:53:MET:CE | 22:B0:57:PHE:CD1 | 2.94 | 0.47 |
| 23:B1:10:LYS:HB2 | 23:B1:14:VAL:CA | 2.44 | 0.47 |
| 31:BA:65:C:H2' | 31:BA:66:C:C6 | 2.49 | 0.47 |
| 31:BA:105:C:H2' | 31:BA:106:C:H6 | 1.79 | 0.47 |
| 31:BA:380:U:O2 | 31:BA:381:G:C8 | 2.68 | 0.47 |
| 31:BA:745:G:H2' | 31:BA:746:A:H5' | 1.97 | 0.47 |
| 31:BA:768:G:C6 | 31:BA:769:G:C5 | 3.03 | 0.47 |
| 31:BA:828:U:O2 | 31:BA:828:U:C2' | 2.61 | 0.47 |
| 31:BA:1366:A:C2' | 31:BA:1367:A:H5' | 2.44 | 0.47 |
| 31:BA:1572:A:H8 | 31:BA:1572:A:O5' | 1.97 | 0.47 |
| 31:BA:1684:C:C2' | 31:BA:1685:C:H5' | 2.44 | 0.47 |
| 31:BA:2228:G:H2' | 31:BA:2229:C:C6 | 2.50 | 0.47 |
| 31:BA:2243:U:O2' | 31:BA:2244:U:H5' | 2.14 | 0.47 |
| 31:BA:2360:A:O2' | 31:BA:2361:A:C5' | 2.63 | 0.47 |
| 31:BA:2395:C:H6 | 31:BA:2395:C:C5' | 2.21 | 0.47 |
| 31:BA:2688:U:O4' | 31:BA:2688:U:O2 | 2.29 | 0.47 |
| 31:BA:2859:G:O2' | 31:BA:2860:A:P | 2.72 | 0.47 |
| 33:BD:210:GLY:HA2 | 33:BD:213:ARG:HG2 | 1.97 | 0.47 |
| 34:BE:152:LYS:HE2 | 39:BN:78:TYR:HD2 | 1.79 | 0.47 |
| 38:BI:120:ILE:HG22 | 38:BI:121:LYS:N | 2.29 | 0.47 |
| 44:BS:28:VAL:CG1 | 44:BS:29:PHE:N | 2.77 | 0.47 |
| 45:BT:51:ARG:HG2 | 45:BT:52:ILE:N | 2.27 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:344:A:H4' | 1:CA:345:C:OP2 | 2.14 | 0.47 |
| 1:CA:363:A:C5 | 12:CL:31:PRO:HD2 | 2.48 | 0.47 |
| 1:CA:411:A:C8 | 1:CA:413:G:C8 | 3.02 | 0.47 |
| 1:CA:832:C:N4 | 1:CA:854:G:H1 | 2.12 | 0.47 |
| 1:CA:1098:C:N3 | 1:CA:1099:G:C8 | 2.82 | 0.47 |
| 1:CA:1147:C:C5 | 1:CA:1148:U:C4 | 3.03 | 0.47 |
| 9:CI:28:VAL:HG13 | 9:CI:65:VAL:HG12 | 1.96 | 0.47 |
| 11:CK:62:GLN:O | 11:CK:64:ALA:N | 2.47 | 0.47 |
| 12:CL:20:LYS:HD3 | 12:CL:20:LYS:H | 1.79 | 0.47 |
| 12:CL:62:SER:O | 12:CL:64:TYR:N | 2.47 | 0.47 |
| 18:CR:36:ASN:HD22 | 18:CR:39:VAL:HG21 | 1.80 | 0.47 |
| 20:CT:58:LYS:HG3 | 20:CT:62:LEU:HD12 | 1.96 | 0.47 |
| 23:D1:47:GLN:CD | 31:DA:2090:G:H21 | 2.23 | 0.47 |
| 30:D8:38:GLY:C | 30:D8:40:GLU:N | 2.71 | 0.47 |
| 31:DA:271(S):G:C5 | 31:DA:271(T):C:C5 | 3.02 | 0.47 |
| 31:DA:901:A:H2' | 31:DA:901:A:N3 | 2.30 | 0.47 |
| 31:DA:1291:C:H2' | 31:DA:1292:U:C6 | 2.49 | 0.47 |
| 31:DA:1503:U:C6 | 31:DA:1504:C:C5 | 3.02 | 0.47 |
| 31:DA:1517:G:O2' | 31:DA:1518:U:H5' | 2.14 | 0.47 |
| 31:DA:1557:C:H5" | 31:DA:1558:A:OP2 | 2.15 | 0.47 |
| 31:DA:1623:G:C2 | 31:DA:1624:G:C8 | 3.03 | 0.47 |
| 31:DA:1722:A:O2' | 31:DA:1739:U:C5' | 2.62 | 0.47 |
| 31:DA:2306:C:OP2 | 31:DA:2307:G:H8 | 1.97 | 0.47 |
| 31:DA:2358:G:C5 | 31:DA:2359:C:C5 | 3.03 | 0.47 |
| 31:DA:2516:G:C6 | 31:DA:2517:C:N4 | 2.82 | 0.47 |
| 32:DB:37:C:C5 | 32:DB:38:C:C4 | 3.02 | 0.47 |
| 32:DB:79:C:C2' | 32:DB:80:U:H5' | 2.44 | 0.47 |
| 33:DD:9:TYR:O | 33:DD:10:THR:HG22 | 2.13 | 0.47 |
| 33:DD:210:GLY:HA2 | 33:DD:213:ARG:HG2 | 1.96 | 0.47 |
| 34:DE:119:ARG:HD2 | 34:DE:120:TRP:CE2 | 2.49 | 0.47 |
| 35:DF:8:GLN:CB | 35:DF:126:VAL:HA | 2.43 | 0.47 |
| 35:DF:132:VAL:CG2 | 35:DF:133:ASN:N | 2.77 | 0.47 |
| 35:DF:132:VAL:O | 35:DF:134:GLY:N | 2.46 | 0.47 |
| 42:DQ:93:TYR:N | 42:DQ:93:TYR:CD1 | 2.82 | 0.47 |
| 44:DS:87:PHE:O | 44:DS:88:ASP:CB | 2.62 | 0.47 |
| 46:DU:92:ARG:NH2 | 47:DV:10:LYS:HB3 | 2.29 | 0.47 |
| 47:DV:66:ARG:HD2 | 47:DV:67:GLY:N | 2.29 | 0.47 |
| 47:DV:70:ILE:HA | 47:DV:90:PRO:HB2 | 1.96 | 0.47 |
| 50:DY:40:GLU:OE2 | 50:DY:40:GLU:HA | 2.14 | 0.47 |
| 1:AA:59:A:C5' | 1:AA:60:A:H5" | 2.14 | 0.47 |
| 1:AA:626:U:C2 | 1:AA:627:G:N7 | 2.82 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:738:C:H2' | 1:AA:739:C:C6 | 2.49 | 0.47 |
| 1:AA:827:U:C4 | 1:AA:870:U:N3 | 2.82 | 0.47 |
| 1:AA:833:U:H2' | 1:AA:834:C:C6 | 2.46 | 0.47 |
| 1:AA:1142:G:H3' | 1:AA:1143:G:C8 | 2.49 | 0.47 |
| 1:AA:1264:C:H2' | 1:AA:1265:G:H8 | 1.78 | 0.47 |
| 2:AB:12:GLU:HA | 2:AB:16:HIS:HB2 | 1.97 | 0.47 |
| 2:AB:21:ARG:HB2 | 2:AB:38:GLY:O | 2.14 | 0.47 |
| 2:AB:100:GLY:N | 2:AB:176:GLU:OE2 | 2.42 | 0.47 |
| 3:AC:125:GLU:HA | 3:AC:191:THR:HG22 | 1.95 | 0.47 |
| 5:AE:101:ILE:CD1 | 5:AE:119:LEU:HD23 | 2.44 | 0.47 |
| 6:AF:97:PHE:O | 18:AR:31:LEU:HD23 | 2.14 | 0.47 |
| 9:AI:28:VAL:HA | 9:AI:63:ILE:O | 2.15 | 0.47 |
| 16:AP:59:TRP:O | 16:AP:64:ALA:HB3 | 2.13 | 0.47 |
| 19:AS:42:PRO:O | 19:AS:43:GLU:CB | 2.55 | 0.47 |
| 20:AT:53:LEU:HD21 | 20:AT:92:LEU:HD11 | 1.96 | 0.47 |
| 20:AT:89:ARG:HH22 | 20:AT:104:LEU:HD21 | 1.77 | 0.47 |
| 23:B1:19:GLN:CD | 23:B1:44:PRO:CB | 2.88 | 0.47 |
| 24:B2:47:ASN:HB2 | 24:B2:51:ARG:HD2 | 1.97 | 0.47 |
| 29:B7:34:ARG:HB2 | 29:B7:42:LEU:HD22 | 1.96 | 0.47 |
| 31:BA:118:A:H3' | 31:BA:119:A:C5' | 2.45 | 0.47 |
| 31:BA:150:C:H2' | 31:BA:151:C:C6 | 2.50 | 0.47 |
| 31:BA:543:C:H5 | 31:BA:547:A:H62 | 1.62 | 0.47 |
| 31:BA:745:G:C2' | 31:BA:746:A:H5' | 2.45 | 0.47 |
| 31:BA:943:U:O2' | 31:BA:944:G:H5' | 2.15 | 0.47 |
| 31:BA:1353:A:H4' | 33:BD:38:LYS:NZ | 2.30 | 0.47 |
| 31:BA:1500:G:C5 | 31:BA:1501:C:C4 | 3.03 | 0.47 |
| 31:BA:2394:C:H2' | 31:BA:2395:C:C5' | 2.43 | 0.47 |
| 31:BA:2602:A:H4' | 31:BA:2603:G:O5' | 2.14 | 0.47 |
| 36:BG:103:LEU:HD23 | 36:BG:106:LEU:HD23 | 1.95 | 0.47 |
| 36:BG:146:TYR:O | 36:BG:149:VAL:HG22 | 2.15 | 0.47 |
| 38:BI:88:ILE:HD11 | 38:BI:122:GLU:C | 2.38 | 0.47 |
| 39:BN:3:THR:O | 39:BN:4:TYR:CG | 2.66 | 0.47 |
| 39:BN:67:LEU:O | 39:BN:69:GLN:N | 2.48 | 0.47 |
| 41:BP:8:PRO:C | 41:BP:10:PRO:HD3 | 2.39 | 0.47 |
| 41:BP:16:ARG:HD3 | 41:BP:16:ARG:C | 2.40 | 0.47 |
| 42:BQ:141:GLN:CG | 51:BZ:72:ARG:HA | 2.44 | 0.47 |
| 42:BQ:141:GLN:C | 51:BZ:70:LEU:HD13 | 2.39 | 0.47 |
| 44:BS:29:PHE:HD2 | 44:BS:29:PHE:C | 2.21 | 0.47 |
| 46:BU:112:ARG:NH1 | 46:BU:112:ARG:HG3 | 2.27 | 0.47 |
| 49:BX:21:PHE:N | 49:BX:21:PHE:CD1 | 2.83 | 0.47 |
| 50:BY:75:ILE:HD11 | 50:BY:80:GLY:N | 2.29 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:59:A:H3' | 1:CA:331:G:H22 | 1.80 | 0.47 |
| 1:CA:63:C:H42 | 1:CA:104:G:H1 | 1.62 | 0.47 |
| 1:CA:343:U:C2' | 1:CA:346:G:O6 | 2.63 | 0.47 |
| 1:CA:355:C:O2' | 1:CA:356:A:H5' | 2.14 | 0.47 |
| 1:CA:1160:G:C2 | 1:CA:1161:C:C6 | 3.03 | 0.47 |
| 2:CB:59:GLU:O | 2:CB:63:MET:HG2 | 2.14 | 0.47 |
| 2:CB:185:ILE:HG12 | 2:CB:185:ILE:O | 2.13 | 0.47 |
| 6:CF:9:VAL:HA | 6:CF:59:TYR:O | 2.14 | 0.47 |
| 7:CG:150:ALA:O | 11:CK:57:THR:HG21 | 2.14 | 0.47 |
| 8:CH:29:SER:HB3 | 8:CH:32:LYS:CD | 2.45 | 0.47 |
| 9:CI:79:LEU:HD13 | 9:CI:79:LEU:C | 2.39 | 0.47 |
| 19:CS:10:PHE:HZ | 19:CS:70:LYS:CE | 2.27 | 0.47 |
| 23:D1:28:GLY:C | 23:D1:30:VAL:H | 2.22 | 0.47 |
| 24:D2:18:PRO:O | 24:D2:22:GLU:HB2 | 2.15 | 0.47 |
| 27:D5:28:PRO:HD2 | 48:DW:35:ILE:HG23 | 1.96 | 0.47 |
| 31:DA:196:A:C4 | 31:DA:805:G:O6 | 2.68 | 0.47 |
| 31:DA:315:G:H2' | 31:DA:316:C:O4' | 2.14 | 0.47 |
| 31:DA:506:G:O3' | 31:DA:507:A:H8 | 1.97 | 0.47 |
| 31:DA:1003:G:N2 | 31:DA:1153:C:C2 | 2.83 | 0.47 |
| 31:DA:1721:G:C6 | 31:DA:1739:U:H5' | 2.49 | 0.47 |
| 31:DA:2335:A:N7 | 31:DA:2337:G:C5 | 2.83 | 0.47 |
| 31:DA:2478:A:C2' | 31:DA:2479:G:H5' | 2.44 | 0.47 |
| 33:DD:31:LYS:C | 33:DD:32:SER:O | 2.56 | 0.47 |
| 34:DE:65:GLY:C | 34:DE:67:PHE:H | 2.22 | 0.47 |
| 34:DE:78:LEU:N | 34:DE:78:LEU:HD23 | 2.30 | 0.47 |
| 35:DF:63:LYS:CE | 35:DF:67:GLN:CB | 2.88 | 0.47 |
| 41:DP:30:THR:CG2 | 41:DP:31:ALA:H | 2.28 | 0.47 |
| 45:DT:27:THR:O | 45:DT:28:VAL:CG2 | 2.62 | 0.47 |
| 45:DT:29:ARG:HE | 45:DT:84:GLN:CD | 2.22 | 0.47 |
| 47:DV:2:PHE:HE1 | 47:DV:13:ARG:NH2 | 2.11 | 0.47 |
| 47:DV:90:PRO:HG2 | 47:DV:91:TYR:N | 2.25 | 0.47 |
| 49:DX:93:GLU:O | 49:DX:93:GLU:HG3 | 2.15 | 0.47 |
| 50:DY:8:LYS:HE3 | 50:DY:72:VAL:HG23 | 1.92 | 0.47 |
| 50:DY:75:ILE:HD13 | 50:DY:75:ILE:HA | 1.57 | 0.47 |
| 1:AA:429:U:H4' | 1:AA:430:A:O5' | 2.12 | 0.47 |
| 1:AA:655:A:C2 | 1:AA:754:C:N4 | 2.83 | 0.47 |
| 1:AA:1085:U:C2 | 1:AA:1094:G:O6 | 2.68 | 0.47 |
| 1:AA:1443:G:N2 | 1:AA:1460:A:H1' | 2.30 | 0.47 |
| 1:AA:1510:U:H2' | 1:AA:1511:G:C8 | 2.50 | 0.47 |
| 8:AH:51:VAL:HB | 8:AH:52:ASP:H | 1.51 | 0.47 |
| 20:AT:96:GLY:O | 20:AT:97:ALA:HB3 | 2.15 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 25:B3:1:MET:HB2 | 25:B3:38:GLU:HG3 | 1.97 | 0.47 |
| 31:BA:28:A:C5 | 31:BA:29:U:C5 | 3.02 | 0.47 |
| 31:BA:393:C:C4 | 31:BA:394:A:N7 | 2.83 | 0.47 |
| 31:BA:869:G:C4 | 31:BA:870:A:C8 | 3.03 | 0.47 |
| 31:BA:1429:G:H2' | 31:BA:1430:C:H6 | 1.76 | 0.47 |
| 31:BA:1444:G:N2 | 31:BA:1548:C:C2 | 2.83 | 0.47 |
| 31:BA:1504:C:HO2' | 31:BA:1505:C:C5' | 2.26 | 0.47 |
| 31:BA:1911:U:C2 | 31:BA:1918:A:C2 | 3.03 | 0.47 |
| 31:BA:1948:G:C2' | 31:BA:1949:G:H5' | 2.44 | 0.47 |
| 31:BA:2291:U:C5' | 31:BA:2380:C:O2 | 2.62 | 0.47 |
| 31:BA:2309:A:C2 | 31:BA:2310:A:C2 | 3.02 | 0.47 |
| 33:BD:44:ASN:HB2 | 33:BD:48:ARG:O | 2.14 | 0.47 |
| 36:BG:11:TYR:OH | 36:BG:33:ARG:HA | 2.15 | 0.47 |
| 36:BG:106:LEU:O | 36:BG:111:LEU:HG | 2.14 | 0.47 |
| 37:BH:70:THR:O | 37:BH:73:ALA:N | 2.48 | 0.47 |
| 39:BN:41:ASP:O | 39:BN:42:TRP:C | 2.57 | 0.47 |
| 39:BN:45:ASN:HD22 | 39:BN:45:ASN:N | 1.90 | 0.47 |
| 43:BR:60:LEU:O | 43:BR:60:LEU:HG | 2.12 | 0.47 |
| 44:BS:85:VAL:HG23 | 44:BS:106:ARG:CB | 2.21 | 0.47 |
| 46:BU:39:LEU:HD23 | 46:BU:39:LEU:HA | 1.68 | 0.47 |
| 48:BW:86:LEU:C | 48:BW:86:LEU:CD1 | 2.87 | 0.47 |
| 49:BX:88:LYS:O | 49:BX:89:ILE:HB | 2.13 | 0.47 |
| 1:CA:60:A:C4' | 1:CA:61:G:O5' | 2.63 | 0.47 |
| 1:CA:200:G:H1 | 1:CA:217:C:N4 | 2.05 | 0.47 |
| 1:CA:276:G:H2' | 1:CA:277:C:H5' | 1.96 | 0.47 |
| 1:CA:391:G:C6 | 1:CA:392:G:C5 | 3.03 | 0.47 |
| 1:CA:473:G:C2 | 1:CA:474:G:C8 | 3.02 | 0.47 |
| 1:CA:538:G:OP2 | 12:CL:115:LYS:CG | 2.62 | 0.47 |
| 1:CA:564:C:H2' | 1:CA:565:U:H5' | 1.97 | 0.47 |
| 1:CA:626:U:H4' | 16:CP:38:TYR:CZ | 2.49 | 0.47 |
| 1:CA:781:A:C3' | 1:CA:782:A:H5' | 2.45 | 0.47 |
| 1:CA:946:A:O2' | 1:CA:1333:A:N3 | 2.42 | 0.47 |
| 1:CA:1001(A):G:H2' | 1:CA:1002:G:O4' | 2.15 | 0.47 |
| 5:CE:93:PRO:HA | 5:CE:118:ILE:HD12 | 1.97 | 0.47 |
| 6:CF:6:VAL:HG12 | 6:CF:7:ASN:N | 2.30 | 0.47 |
| 6:CF:72:VAL:CG2 | 6:CF:90:VAL:HG11 | 2.44 | 0.47 |
| 10:CJ:22:LYS:O | 10:CJ:22:LYS:HD2 | 2.15 | 0.47 |
| 11:CK:48:ILE:HG21 | 11:CK:63:LEU:HD13 | 1.96 | 0.47 |
| 11:CK:81:ASP:CG | 11:CK:106:LYS:HG2 | 2.39 | 0.47 |
| 13:CM:78:ILE:HG22 | 13:CM:93:ARG:HH22 | 1.78 | 0.47 |
| 30:D8:8:LYS:HE2 | 31:DA:243:U:OP2 | 2.15 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:778:G:C5 | 31:DA:779:U:C4 | 3.02 | 0.47 |
| 31:DA:1948:G:C2' | 31:DA:1949:G:H5' | 2.45 | 0.47 |
| 32:DB:43:C:H4' | 36:DG:66:GLN:HE22 | 1.78 | 0.47 |
| 32:DB:82:G:O2' | 32:DB:83:G:H5' | 2.14 | 0.47 |
| 33:DD:253:GLN:HB3 | 33:DD:255:LYS:NZ | 2.30 | 0.47 |
| 34:DE:92:THR:H | 34:DE:95:ILE:CD1 | 2.20 | 0.47 |
| 34:DE:111:ARG:CZ | 43:DR:2:ARG:NH2 | 2.77 | 0.47 |
| 35:DF:9:ILE:HG12 | 35:DF:14:PRO:HA | 1.97 | 0.47 |
| 37:DH:159:GLU:HA | 37:DH:159:GLU:OE1 | 2.14 | 0.47 |
| 41:DP:13:ASN:H | 41:DP:13:ASN:ND2 | 2.13 | 0.47 |
| 45:DT:120:ARG:HA | 45:DT:123:GLN:HG2 | 1.95 | 0.47 |
| 48:DW:8:ARG:HB3 | 48:DW:9:TYR:CD1 | 2.50 | 0.47 |
| 51:DZ:9:TYR:CE2 | 51:DZ:61:LEU:HD13 | 2.50 | 0.47 |
| 1:AA:112:G:N3 | 1:AA:112:G:H2' | 2.30 | 0.47 |
| 1:AA:115:G:C2 | 1:AA:289:G:N7 | 2.82 | 0.47 |
| 1:AA:343:U:C2' | 1:AA:346:G:O6 | 2.63 | 0.47 |
| 1:AA:355:C:N3 | 1:AA:356:A:N7 | 2.63 | 0.47 |
| 1:AA:691:G:H1' | 1:AA:696:A:N6 | 2.29 | 0.47 |
| 1:AA:946:A:N1 | 1:AA:1236:A:C2 | 2.83 | 0.47 |
| 1:AA:1128:C:O2 | 1:AA:1130:A:N7 | 2.48 | 0.47 |
| 1:AA:1203:C:H2' | 1:AA:1204:A:O4' | 2.14 | 0.47 |
| 1:AA:1320:C:H5' | 19:AS:70:LYS:HG3 | 1.96 | 0.47 |
| 2:AB:83:MET:O | 2:AB:85:ALA:N | 2.47 | 0.47 |
| 4:AD:126:ILE:HG22 | 4:AD:127:THR:N | 2.29 | 0.47 |
| 5:AE:122:GLU:OE1 | 5:AE:131:ILE:HG13 | 2.15 | 0.47 |
| 6:AF:30:LEU:H | 6:AF:30:LEU:HD23 | 1.79 | 0.47 |
| 7:AG:37:ASN:HD21 | 9:AI:40:LEU:CD2 | 2.28 | 0.47 |
| 7:AG:72:ARG:O | 7:AG:73:MET:HG3 | 2.15 | 0.47 |
| 8:AH:97:VAL:HG13 | 8:AH:98:LYS:HG3 | 1.97 | 0.47 |
| 11:AK:110:ASP:O | 18:AR:84:LYS:HD2 | 2.13 | 0.47 |
| 13:AM:91:ARG:HB2 | 13:AM:98:VAL:CG2 | 2.38 | 0.47 |
| 14:AN:44:LEU:C | 14:AN:44:LEU:HD12 | 2.39 | 0.47 |
| 17:AQ:59:ILE:HG22 | 17:AQ:71:PHE:CD1 | 2.46 | 0.47 |
| 18:AR:56:THR:CB | 18:AR:58:LEU:HD13 | 2.44 | 0.47 |
| 18:AR:74:ARG:HB3 | 18:AR:81:PHE:CE1 | 2.49 | 0.47 |
| 22:B0:14:ARG:HB2 | 22:B0:14:ARG:HE | 1.51 | 0.47 |
| 22:B0:73:GLY:O | 22:B0:75:LEU:N | 2.48 | 0.47 |
| 23:B1:89:GLU:O | 23:B1:90:ILE:C | 2.58 | 0.47 |
| 27:B5:47:PRO:O | 27:B5:48:GLU:HG3 | 2.15 | 0.47 |
| 31:BA:49:A:H4' | 31:BA:50:U:H5' | 1.95 | 0.47 |
| 31:BA:52:A:O2' | 31:BA:53:A:H5' | 2.15 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:142:A:H8 | 31:BA:1408:C:H1' | 1.73 | 0.47 |
| 31:BA:151:C:C2' | 31:BA:152:G:H5' | 2.45 | 0.47 |
| 31:BA:151:C:C2 | 31:BA:176:G:N2 | 2.82 | 0.47 |
| 31:BA:271(Q):G:C2 | 31:BA:271(R):G:C5 | 3.02 | 0.47 |
| 31:BA:301:G:H1 | 31:BA:316:C:H42 | 1.62 | 0.47 |
| 31:BA:322:A:OP2 | 35:BF:169:ASN:HB2 | 2.15 | 0.47 |
| 31:BA:344:G:O2' | 31:BA:345:A:H5' | 2.15 | 0.47 |
| 31:BA:483:A:C8 | 31:BA:484:C:C5 | 3.03 | 0.47 |
| 31:BA:570:G:H2' | 31:BA:2030:A:N6 | 2.29 | 0.47 |
| 31:BA:659:C:C6 | 31:BA:659:C:C4' | 2.97 | 0.47 |
| 31:BA:743:G:H2' | 31:BA:744:G:H5' | 1.96 | 0.47 |
| 31:BA:906:G:C2' | 31:BA:907:U:O5' | 2.63 | 0.47 |
| 31:BA:926:A:C8 | 31:BA:926:A:H5" | 2.49 | 0.47 |
| 31:BA:971:C:C2' | 31:BA:972:G:H5' | 2.45 | 0.47 |
| 31:BA:1163:G:O2' | 31:BA:1164:G:H5' | 2.15 | 0.47 |
| 31:BA:1272:A:H3' | 31:BA:1273:U:H5" | 1.96 | 0.47 |
| 31:BA:1489:U:C2' | 31:BA:1490:A:OP2 | 2.62 | 0.47 |
| 31:BA:1505:C:C2' | 31:BA:1506:C:O5' | 2.63 | 0.47 |
| 31:BA:1591:G:C5' | 31:BA:1591:G:H8 | 2.28 | 0.47 |
| 31:BA:1683:C:H2' | 31:BA:1684:C:C6 | 2.49 | 0.47 |
| 31:BA:1740:G:H4' | 31:BA:1741:A:OP1 | 2.13 | 0.47 |
| 31:BA:1767:C:C2' | 31:BA:1768:U:H5' | 2.44 | 0.47 |
| 31:BA:1857:G:C6 | 31:BA:1858:G:C2 | 3.03 | 0.47 |
| 31:BA:1859:A:C6 | 31:BA:1884:A:C8 | 3.03 | 0.47 |
| 31:BA:2537:U:H2' | 31:BA:2538:C:C6 | 2.50 | 0.47 |
| 31:BA:2557:G:H2' | 31:BA:2558:C:C6 | 2.49 | 0.47 |
| 31:BA:2619:C:H2' | 31:BA:2620:C:H6 | 1.80 | 0.47 |
| 31:BA:2631:G:N3 | 31:BA:2810:A:C2 | 2.82 | 0.47 |
| 31:BA:2646:C:H2' | 31:BA:2647:U:O4' | 2.14 | 0.47 |
| 31:BA:2762:G:C2' | 31:BA:2763:G:H5' | 2.44 | 0.47 |
| 31:BA:2870:C:H2' | 31:BA:2871:C:C5' | 2.45 | 0.47 |
| 32:BB:37:C:H2' | 32:BB:38:C:H5' | 1.96 | 0.47 |
| 33:BD:71:ASP:CB | 33:BD:103:ARG:HH22 | 2.27 | 0.47 |
| 33:BD:211:ARG:HA | 33:BD:214:TRP:CG | 2.50 | 0.47 |
| 33:BD:253:GLN:HB3 | 33:BD:255:LYS:CE | 2.45 | 0.47 |
| 34:BE:31:CYS:HA | 34:BE:32:PRO:HD3 | 1.66 | 0.47 |
| 34:BE:52:LEU:O | 34:BE:74:PRO:CA | 2.55 | 0.47 |
| 34:BE:75:VAL:C | 34:BE:77:ILE:H | 2.21 | 0.47 |
| 35:BF:28:ILE:HA | 35:BF:112:MET:HG2 | 1.96 | 0.47 |
| 36:BG:63:ILE:HD13 | 36:BG:141:PHE:CZ | 2.49 | 0.47 |
| 36:BG:109:VAL:C | 36:BG:112:PRO:HD2 | 2.40 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 36:BG:141:PHE:CE2 | 36:BG:155:MET:HE1 | 2.49 | 0.47 |
| 37:BH:54:ARG:HB3 | 37:BH:65:HIS:HB2 | 1.96 | 0.47 |
| 37:BH:154:PRO:O | 37:BH:156:ALA:N | 2.39 | 0.47 |
| 41:BP:21:ARG:O | 41:BP:21:ARG:CG | 2.63 | 0.47 |
| 42:BQ:23:GLY:HA3 | 42:BQ:99:PRO:O | 2.15 | 0.47 |
| 42:BQ:39:PRO:HA | 42:BQ:97:VAL:O | 2.15 | 0.47 |
| 42:BQ:79:LEU:C | 42:BQ:80:GLU:CD | 2.83 | 0.47 |
| 43:BR:41:ALA:O | 43:BR:43:GLU:N | 2.48 | 0.47 |
| 44:BS:38:GLN:HB3 | 44:BS:47:THR:CG2 | 2.45 | 0.47 |
| 45:BT:57:PHE:CG | 45:BT:58:ASN:N | 2.82 | 0.47 |
| 46:BU:8:VAL:HG11 | 46:BU:12:ARG:NH1 | 2.29 | 0.47 |
| 46:BU:25:TRP:CD1 | 46:BU:26:GLY:H | 2.30 | 0.47 |
| 47:BV:63:GLY:O | 47:BV:64:HIS:HB3 | 2.13 | 0.47 |
| 48:BW:24:ILE:HD12 | 48:BW:24:ILE:C | 2.39 | 0.47 |
| 49:BX:36:LYS:NZ | 49:BX:38:GLU:C | 2.71 | 0.47 |
| 49:BX:65:ARG:NH1 | 49:BX:66:LEU:N | 2.62 | 0.47 |
| 50:BY:39:VAL:O | 50:BY:40:GLU:CD | 2.57 | 0.47 |
| 50:BY:81:LYS:HD3 | 50:BY:97:ARG:O | 2.14 | 0.47 |
| 51:BZ:26:GLY:HA2 | 51:BZ:85:HIS:CD2 | 2.50 | 0.47 |
| 51:BZ:98:MET:HE2 | 51:BZ:100:VAL:HG23 | 1.96 | 0.47 |
| 51:BZ:111:VAL:CG1 | 51:BZ:112:ARG:N | 2.78 | 0.47 |
| 1:CA:35:G:H2' | 1:CA:36:C:C6 | 2.49 | 0.47 |
| 1:CA:67:C:O2' | 1:CA:171:A:H1' | 2.15 | 0.47 |
| 1:CA:114:U:O2' | 1:CA:115:G:H5' | 2.15 | 0.47 |
| 1:CA:130:A:C8 | 17:CQ:63:ARG:HG3 | 2.49 | 0.47 |
| 1:CA:247:G:C2 | 1:CA:248:C:C6 | 3.02 | 0.47 |
| 1:CA:444:C:H2' | 1:CA:445:G:C8 | 2.36 | 0.47 |
| 1:CA:518:C:H4' | 1:CA:519:C:H5" | 1.95 | 0.47 |
| 1:CA:577:G:H1' | 1:CA:816:A:N3 | 2.29 | 0.47 |
| 1:CA:629:G:C6 | 1:CA:630:G:N7 | 2.83 | 0.47 |
| 1:CA:853:G:O2' | 1:CA:854:G:H5' | 2.14 | 0.47 |
| 1:CA:865:A:H2 | 1:CA:918:A:H4' | 1.80 | 0.47 |
| 1:CA:948:C:OP1 | 13:CM:107:ALA:HA | 2.14 | 0.47 |
| 1:CA:983:A:H5' | 1:CA:984:C:OP2 | 2.15 | 0.47 |
| 1:CA:985:C:H2' | 1:CA:986:A:C8 | 2.50 | 0.47 |
| 1:CA:1071:C:H5" | 5:CE:49:PRO:HG3 | 1.97 | 0.47 |
| 1:CA:1074:G:N3 | 1:CA:1102:A:C2 | 2.83 | 0.47 |
| 1:CA:1371:G:O3' | 9:CI:69:GLY:HA3 | 2.15 | 0.47 |
| 1:CA:1398:A:N1 | 5:CE:19:MET:HE3 | 2.30 | 0.47 |
| 1:CA:1504:G:H4' | 1:CA:1505:G:O4' | 2.15 | 0.47 |
| 2:CB:153:ARG:O | 2:CB:154:LEU:C | 2.57 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 2:CB:239:VAL:O | 2:CB:239:VAL:HG12 | 2.15 | 0.47 |
| 3:CC:123:GLN:O | 3:CC:128:PHE:HB2 | 2.15 | 0.47 |
| 4:CD:159:ARG:HA | 4:CD:162:LEU:HB2 | 1.96 | 0.47 |
| 4:CD:159:ARG:O | 4:CD:163:GLU:N | 2.48 | 0.47 |
| 5:CE:57:LYS:O | 5:CE:61:TYR:CD2 | 2.62 | 0.47 |
| 8:CH:69:ARG:HD3 | 8:CH:75:ARG:O | 2.12 | 0.47 |
| 9:CI:4:TYR:N | 9:CI:4:TYR:CD1 | 2.81 | 0.47 |
| 9:CI:8:GLY:HA2 | 9:CI:79:LEU:HD12 | 1.96 | 0.47 |
| 9:CI:53:VAL:CG1 | 9:CI:95:LYS:HE3 | 2.45 | 0.47 |
| 12:CL:24:VAL:HG13 | 12:CL:98:TYR:CE2 | 2.50 | 0.47 |
| 14:CN:41:ARG:HG3 | 14:CN:42:ILE:N | 2.28 | 0.47 |
| 16:CP:20:VAL:HG21 | 16:CP:32:TYR:CB | 2.45 | 0.47 |
| 16:CP:20:VAL:HG21 | 16:CP:32:TYR:CD2 | 2.49 | 0.47 |
| 16:CP:50:LYS:HD3 | 16:CP:51:VAL:N | 2.30 | 0.47 |
| 18:CR:56:THR:CB | 18:CR:58:LEU:HD13 | 2.45 | 0.47 |
| 20:CT:81:LYS:C | 20:CT:83:ARG:H | 2.22 | 0.47 |
| 24:D2:39:ALA:O | 24:D2:40:SER:C | 2.57 | 0.47 |
| 24:D2:46:GLN:C | 24:D2:48:HIS:N | 2.72 | 0.47 |
| 30:D8:50:LEU:HD12 | 30:D8:51:ALA:H | 1.79 | 0.47 |
| 30:D8:51:ALA:N | 30:D8:53:PRO:HD2 | 2.30 | 0.47 |
| 31:DA:9:U:C4 | 31:DA:2629:A:C6 | 3.03 | 0.47 |
| 31:DA:151:C:C2’ | 31:DA:152:G:H5’ | 2.45 | 0.47 |
| 31:DA:287:C:C4 | 31:DA:288:C:C5 | 3.02 | 0.47 |
| 31:DA:792:G:H5” | 31:DA:793:A:H5’ | 1.96 | 0.47 |
| 31:DA:824:A:C2’ | 31:DA:825:C:H5’ | 2.45 | 0.47 |
| 31:DA:848:G:C2 | 31:DA:933:A:H1’ | 2.50 | 0.47 |
| 31:DA:971:C:C2’ | 31:DA:972:G:H5’ | 2.45 | 0.47 |
| 31:DA:1019:U:N3 | 31:DA:1142(A):A:N6 | 2.51 | 0.47 |
| 31:DA:1027:A:C2 | 31:DA:2488:A:H5’ | 2.50 | 0.47 |
| 31:DA:1040:C:HO2’ | 31:DA:1041:C:P | 2.34 | 0.47 |
| 31:DA:1131:G:H8 | 31:DA:2025:C:H4’ | 1.80 | 0.47 |
| 31:DA:1190:G:H5’ | 41:DP:35:HIS:HA | 1.96 | 0.47 |
| 31:DA:1322:A:C5 | 31:DA:1323:U:C5 | 3.02 | 0.47 |
| 31:DA:1403:C:H2’ | 31:DA:1404:C:O5’ | 2.15 | 0.47 |
| 31:DA:1464:C:C2’ | 31:DA:1528:A:H8 | 2.28 | 0.47 |
| 31:DA:1952:A:C6 | 40:DO:22:ILE:HD12 | 2.49 | 0.47 |
| 31:DA:1988:C:O2’ | 31:DA:1989:G:H5’ | 2.14 | 0.47 |
| 31:DA:2006:C:O2’ | 31:DA:2823:A:N3 | 2.47 | 0.47 |
| 31:DA:2053:G:N2 | 31:DA:2054:A:C4 | 2.82 | 0.47 |
| 31:DA:2337:G:C2 | 31:DA:2338:G:C8 | 3.03 | 0.47 |
| 31:DA:2346:A:H5” | 31:DA:2383:G:H1’ | 1.97 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2699:C:H2' | 31:DA:2700:C:O4' | 2.14 | 0.47 |
| 31:DA:2884:U:H2' | 31:DA:2885:C:H5' | 1.97 | 0.47 |
| 32:DB:71:C:O2 | 32:DB:71:C:C2' | 2.61 | 0.47 |
| 33:DD:44:ASN:OD1 | 33:DD:44:ASN:N | 2.46 | 0.47 |
| 37:DH:13:LYS:HE2 | 37:DH:13:LYS:HA | 1.97 | 0.47 |
| 37:DH:158:HIS:CE1 | 37:DH:170:ARG:HA | 2.48 | 0.47 |
| 41:DP:13:ASN:O | 41:DP:15:ARG:N | 2.46 | 0.47 |
| 42:DQ:141:GLN:CG | 51:DZ:72:ARG:HA | 2.44 | 0.47 |
| 44:DS:25:ARG:HH11 | 44:DS:25:ARG:CB | 2.26 | 0.47 |
| 44:DS:87:PHE:CG | 44:DS:88:ASP:N | 2.83 | 0.47 |
| 45:DT:28:VAL:HG22 | 45:DT:46:GLU:HG3 | 1.93 | 0.47 |
| 45:DT:31:SER:HA | 45:DT:32:TYR:CD2 | 2.49 | 0.47 |
| 45:DT:33:LYS:HA | 45:DT:33:LYS:CE | 2.44 | 0.47 |
| 45:DT:55:ASN:H | 45:DT:59:THR:HB | 1.79 | 0.47 |
| 45:DT:88:ILE:HG22 | 45:DT:89:VAL:N | 2.30 | 0.47 |
| 46:DU:112:ARG:NH1 | 46:DU:112:ARG:HG3 | 2.28 | 0.47 |
| 50:DY:17:SER:CA | 50:DY:71:LYS:HD2 | 2.44 | 0.47 |
| 50:DY:28:LYS:HE3 | 50:DY:37:VAL:HG12 | 1.95 | 0.47 |
| 50:DY:50:ARG:HB3 | 50:DY:51:VAL:H | 1.59 | 0.47 |
| 51:DZ:37:VAL:O | 51:DZ:38:TYR:HB3 | 2.14 | 0.47 |
| 1:AA:303:A:C4 | 1:AA:304:U:C6 | 3.03 | 0.47 |
| 1:AA:629:G:C6 | 1:AA:630:G:N7 | 2.83 | 0.47 |
| 1:AA:655:A:N1 | 1:AA:754:C:N4 | 2.62 | 0.47 |
| 1:AA:665:A:N6 | 1:AA:725:G:O6 | 2.47 | 0.47 |
| 1:AA:832:C:N4 | 1:AA:854:G:H1 | 2.13 | 0.47 |
| 1:AA:853:G:O2' | 1:AA:854:G:H5' | 2.14 | 0.47 |
| 1:AA:882:C:O2' | 1:AA:883:C:H5' | 2.15 | 0.47 |
| 1:AA:983:A:H3' | 1:AA:983:A:N3 | 2.29 | 0.47 |
| 1:AA:1031:G:H2' | 1:AA:1032:G:C8 | 2.50 | 0.47 |
| 9:AI:51:ARG:HE | 9:AI:56:LEU:CD1 | 2.27 | 0.47 |
| 10:AJ:51:ARG:HG3 | 10:AJ:61:GLU:N | 2.30 | 0.47 |
| 14:AN:41:ARG:HG3 | 14:AN:42:ILE:N | 2.29 | 0.47 |
| 16:AP:43:LYS:HG2 | 16:AP:48:TRP:CE3 | 2.50 | 0.47 |
| 23:B1:82:LEU:HG | 23:B1:83:GLU:N | 2.30 | 0.47 |
| 24:B2:20:GLU:O | 24:B2:23:LYS:N | 2.47 | 0.47 |
| 24:B2:23:LYS:CA | 49:BX:5:TYR:CD1 | 2.98 | 0.47 |
| 24:B2:25:VAL:HG22 | 24:B2:26:ARG:HH11 | 1.80 | 0.47 |
| 30:B8:51:ALA:HB1 | 30:B8:54:GLU:OE1 | 2.15 | 0.47 |
| 30:B8:62:LEU:C | 30:B8:64:TYR:H | 2.22 | 0.47 |
| 31:BA:607:U:C2 | 31:BA:621:A:N1 | 2.83 | 0.47 |
| 31:BA:705:A:C8 | 31:BA:727:A:C2 | 3.03 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:810:U:O2' | 41:BP:33:ARG:CZ | 2.63 | 0.47 |
| 31:BA:1225:G:P | 47:BV:88:ARG:HB3 | 2.54 | 0.47 |
| 31:BA:1623:G:C2 | 31:BA:1624:G:C8 | 3.03 | 0.47 |
| 31:BA:1689:A:H62 | 31:BA:1698:A:H2 | 1.62 | 0.47 |
| 31:BA:1862:G:O2' | 31:BA:1863:G:H5' | 2.15 | 0.47 |
| 31:BA:2187:G:C5 | 31:BA:2188:C:C2 | 3.03 | 0.47 |
| 31:BA:2328:A:H2' | 31:BA:2329:G:O4' | 2.12 | 0.47 |
| 31:BA:2536:G:C5 | 31:BA:2537:U:C5 | 3.02 | 0.47 |
| 32:BB:88:C:C4 | 32:BB:89:G:C6 | 3.03 | 0.47 |
| 33:BD:52:ARG:NH1 | 33:BD:53:PHE:HE2 | 2.12 | 0.47 |
| 34:BE:174:ASP:OD2 | 34:BE:175:VAL:N | 2.46 | 0.47 |
| 35:BF:1:MET:O | 35:BF:2:LYS:O | 2.32 | 0.47 |
| 35:BF:132:VAL:C | 35:BF:134:GLY:N | 2.73 | 0.47 |
| 35:BF:140:LEU:HD13 | 35:BF:140:LEU:HA | 1.75 | 0.47 |
| 40:BO:35:VAL:HG13 | 40:BO:65:THR:CG2 | 2.45 | 0.47 |
| 45:BT:29:ARG:HE | 45:BT:84:GLN:CD | 2.23 | 0.47 |
| 49:BX:36:LYS:C | 49:BX:38:GLU:N | 2.71 | 0.47 |
| 50:BY:18:GLY:O | 50:BY:19:LYS:C | 2.57 | 0.47 |
| 51:BZ:16:SER:HA | 51:BZ:19:ARG:HD2 | 1.96 | 0.47 |
| 1:CA:192:U:C1' | 20:CT:103:GLY:HA2 | 2.44 | 0.47 |
| 1:CA:199:G:O2' | 1:CA:200:G:H5' | 2.14 | 0.47 |
| 1:CA:429:U:H4' | 1:CA:430:A:O5' | 2.14 | 0.47 |
| 1:CA:655:A:C2 | 1:CA:754:C:N4 | 2.83 | 0.47 |
| 1:CA:657:G:C2 | 1:CA:750:G:C5 | 3.03 | 0.47 |
| 1:CA:721:G:H4' | 1:CA:722:A:O4' | 2.14 | 0.47 |
| 1:CA:1189:C:O2' | 3:CC:176:HIS:CD2 | 2.67 | 0.47 |
| 2:CB:21:ARG:HB2 | 2:CB:38:GLY:O | 2.15 | 0.47 |
| 4:CD:21:LEU:O | 4:CD:113:SER:HB2 | 2.14 | 0.47 |
| 5:CE:10:MET:HG3 | 5:CE:13:ILE:CD1 | 2.45 | 0.47 |
| 10:CJ:40:LEU:HB2 | 10:CJ:41:PRO:CD | 2.39 | 0.47 |
| 11:CK:62:GLN:O | 11:CK:63:LEU:C | 2.58 | 0.47 |
| 22:D0:25:ARG:HG3 | 22:D0:29:GLN:HE21 | 1.80 | 0.47 |
| 28:D6:25:LYS:O | 31:DA:2286:A:H2 | 1.98 | 0.47 |
| 31:DA:322:A:OP2 | 35:DF:169:ASN:HB2 | 2.15 | 0.47 |
| 31:DA:475:U:O2 | 31:DA:505:A:H2 | 1.97 | 0.47 |
| 31:DA:696:G:O2' | 31:DA:697:C:H5' | 2.15 | 0.47 |
| 31:DA:1178:C:C5 | 31:DA:1179:C:H5 | 2.33 | 0.47 |
| 31:DA:1227:G:O2' | 31:DA:1228:G:H5' | 2.15 | 0.47 |
| 31:DA:1341:U:N3 | 49:DX:77:LYS:HE2 | 2.30 | 0.47 |
| 31:DA:1983:C:O2' | 31:DA:1984:G:H5' | 2.14 | 0.47 |
| 31:DA:2063:C:C4 | 31:DA:2064:C:C4 | 3.03 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2199:A:C4 | 31:DA:2200:C:C6 | 3.03 | 0.47 |
| 31:DA:2317:C:O2' | 31:DA:2318:G:H5' | 2.15 | 0.47 |
| 31:DA:2515:C:O2 | 31:DA:2570:G:C2 | 2.68 | 0.47 |
| 31:DA:2701:C:C3' | 31:DA:2702:U:C5' | 2.68 | 0.47 |
| 33:DD:35:LYS:CD | 33:DD:63:ARG:C | 2.87 | 0.47 |
| 33:DD:253:GLN:HB3 | 33:DD:255:LYS:CE | 2.45 | 0.47 |
| 35:DF:132:VAL:C | 35:DF:134:GLY:H | 2.22 | 0.47 |
| 35:DF:199:TRP:CH2 | 35:DF:203:GLN:NE2 | 2.83 | 0.47 |
| 39:DN:39:ARG:NE | 39:DN:41:ASP:HB2 | 2.29 | 0.47 |
| 39:DN:79:PRO:HD2 | 39:DN:80:GLY:H | 1.79 | 0.47 |
| 41:DP:88:LEU:O | 41:DP:90:ARG:N | 2.48 | 0.47 |
| 43:DR:65:LEU:HA | 43:DR:65:LEU:HD12 | 1.58 | 0.47 |
| 43:DR:72:ASP:HB3 | 43:DR:75:LEU:CB | 2.44 | 0.47 |
| 49:DX:12:VAL:HG11 | 49:DX:27:THR:HG23 | 1.97 | 0.47 |
| 49:DX:33:LYS:CA | 49:DX:35:THR:HG22 | 2.44 | 0.47 |
| 50:DY:22:GLY:O | 50:DY:23:ARG:CG | 2.63 | 0.47 |
| 50:DY:47:LYS:HB3 | 50:DY:47:LYS:HZ2 | 1.76 | 0.47 |
| 51:DZ:141:VAL:O | 51:DZ:141:VAL:HG13 | 2.15 | 0.47 |
| 1:AA:303:A:C5 | 1:AA:304:U:C5 | 3.03 | 0.47 |
| 1:AA:414:A:H2' | 1:AA:415:A:O4' | 2.15 | 0.47 |
| 1:AA:664:G:P | 18:AR:64:ARG:HH21 | 2.38 | 0.47 |
| 1:AA:1203:C:H2' | 1:AA:1204:A:C8 | 2.50 | 0.47 |
| 2:AB:76:GLN:O | 2:AB:208:ILE:HG12 | 2.15 | 0.47 |
| 2:AB:167:PRO:HG3 | 2:AB:188:ALA:HB2 | 1.97 | 0.47 |
| 4:AD:93:PHE:O | 4:AD:97:LEU:HB2 | 2.15 | 0.47 |
| 5:AE:79:GLU:HB3 | 5:AE:92:LYS:HG2 | 1.97 | 0.47 |
| 11:AK:122:LYS:HB3 | 11:AK:122:LYS:HE2 | 1.66 | 0.47 |
| 12:AL:58:VAL:O | 12:AL:65:GLU:HA | 2.15 | 0.47 |
| 13:AM:3:ARG:HG2 | 13:AM:9:ILE:HD11 | 1.95 | 0.47 |
| 16:AP:50:LYS:HD3 | 16:AP:50:LYS:C | 2.40 | 0.47 |
| 17:AQ:22:LEU:HD11 | 17:AQ:39:SER:HB2 | 1.97 | 0.47 |
| 23:B1:15:ALA:HA | 23:B1:46:LEU:HD21 | 1.95 | 0.47 |
| 24:B2:26:ARG:CG | 49:BX:5:TYR:O | 2.63 | 0.47 |
| 25:B3:47:VAL:HG11 | 25:B3:56:VAL:HG21 | 1.96 | 0.47 |
| 25:B3:52:HIS:NE2 | 32:BB:83:G:H5" | 2.29 | 0.47 |
| 27:B5:55:ARG:CD | 27:B5:56:LYS:N | 2.75 | 0.47 |
| 30:B8:4:MET:SD | 30:B8:61:LEU:CD1 | 3.02 | 0.47 |
| 31:BA:64:A:O3' | 49:BX:68:ARG:O | 2.33 | 0.47 |
| 31:BA:69:C:O2 | 31:BA:69:C:C2' | 2.55 | 0.47 |
| 31:BA:271(J):C:C3' | 31:BA:271(K):U:H5" | 2.44 | 0.47 |
| 31:BA:271(S):G:C4 | 31:BA:271(T):C:C6 | 3.03 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:995:C:C2 | 39:BN:4:TYR:OH | 2.43 | 0.47 |
| 31:BA:1175:U:H4' | 31:BA:1176:G:H2' | 1.97 | 0.47 |
| 31:BA:1517:G:O2' | 31:BA:1518:U:H5' | 2.15 | 0.47 |
| 31:BA:2281:C:O2' | 31:BA:2282:G:H5' | 2.15 | 0.47 |
| 31:BA:2584:U:O2 | 31:BA:2584:U:O5' | 2.32 | 0.47 |
| 31:BA:2681:C:C5 | 31:BA:2725:A:N6 | 2.64 | 0.47 |
| 31:BA:2881:C:H2' | 31:BA:2882:A:O4' | 2.15 | 0.47 |
| 34:BE:103:ASP:OD1 | 34:BE:201:THR:HA | 2.15 | 0.47 |
| 35:BF:34:TRP:CD1 | 41:BP:11:GLY:HA2 | 2.50 | 0.47 |
| 36:BG:174:GLU:HG3 | 36:BG:180:PHE:CD1 | 2.48 | 0.47 |
| 37:BH:97:ARG:O | 37:BH:98:LEU:C | 2.58 | 0.47 |
| 37:BH:111:HIS:CG | 37:BH:112:PRO:HD2 | 2.50 | 0.47 |
| 38:BI:129:THR:HG22 | 38:BI:130:TYR:O | 2.14 | 0.47 |
| 40:BO:87:ILE:HG23 | 40:BO:88:ASN:N | 2.30 | 0.47 |
| 49:BX:25:LYS:CG | 49:BX:26:TYR:N | 2.61 | 0.47 |
| 49:BX:47:PHE:O | 49:BX:48:LYS:C | 2.58 | 0.47 |
| 50:BY:91:GLU:HB3 | 50:BY:92:ASN:H | 1.54 | 0.47 |
| 1:CA:149:A:O2' | 1:CA:150:C:P | 2.73 | 0.47 |
| 1:CA:738:C:H2' | 1:CA:739:C:C6 | 2.49 | 0.47 |
| 1:CA:983:A:H3' | 1:CA:983:A:N3 | 2.30 | 0.47 |
| 4:CD:75:PHE:CE2 | 4:CD:93:PHE:HZ | 2.33 | 0.47 |
| 5:CE:6:PHE:CD2 | 5:CE:36:ASP:HB3 | 2.50 | 0.47 |
| 10:CJ:38:ILE:O | 10:CJ:38:ILE:HG13 | 2.15 | 0.47 |
| 11:CK:21:ILE:HD12 | 11:CK:21:ILE:N | 2.30 | 0.47 |
| 22:D0:16:SER:CB | 31:DA:2262:U:OP2 | 2.63 | 0.47 |
| 23:D1:44:PRO:HA | 31:DA:2231:C:OP1 | 2.15 | 0.47 |
| 31:DA:68:G:H2' | 31:DA:69:C:H6 | 1.79 | 0.47 |
| 31:DA:733:G:H8 | 31:DA:733:G:O5' | 1.98 | 0.47 |
| 31:DA:847:U:OP2 | 31:DA:928:G:O6 | 2.33 | 0.47 |
| 31:DA:1503:U:C6 | 31:DA:1504:C:H5 | 2.33 | 0.47 |
| 31:DA:1600:C:O2' | 31:DA:1601:G:H5' | 2.14 | 0.47 |
| 31:DA:1719:G:O2' | 31:DA:1720:U:H5' | 2.14 | 0.47 |
| 31:DA:1890:A:C2' | 31:DA:1891:G:H5' | 2.45 | 0.47 |
| 31:DA:1946:U:C2 | 31:DA:1947:C:C5 | 3.02 | 0.47 |
| 31:DA:1997:G:O2' | 31:DA:1998:G:H5' | 2.14 | 0.47 |
| 31:DA:2309:A:C2 | 31:DA:2310:A:C2 | 3.02 | 0.47 |
| 31:DA:2504:U:O2 | 31:DA:2504:U:H2' | 2.14 | 0.47 |
| 31:DA:2681:C:C5 | 31:DA:2725:A:N6 | 2.62 | 0.47 |
| 31:DA:2726:U:O2' | 31:DA:2727:G:H8 | 1.97 | 0.47 |
| 34:DE:96:PHE:HA | 34:DE:100:GLU:OE1 | 2.14 | 0.47 |
| 34:DE:101:ARG:HB3 | 34:DE:201:THR:OG1 | 2.15 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 34:DE:188:VAL:HB | 34:DE:189:PRO:HD2 | 1.96 | 0.47 |
| 35:DF:160:ASN:ND2 | 35:DF:162:LEU:H | 2.13 | 0.47 |
| 37:DH:41:MET:HB3 | 37:DH:43:VAL:HG22 | 1.97 | 0.47 |
| 37:DH:46:GLU:HG3 | 37:DH:51:ARG:HB3 | 1.96 | 0.47 |
| 37:DH:83:TYR:HB3 | 37:DH:134:SER:HA | 1.97 | 0.47 |
| 38:DI:98:ALA:O | 38:DI:99:GLU:C | 2.58 | 0.47 |
| 39:DN:78:TYR:CD1 | 39:DN:79:PRO:HD3 | 2.49 | 0.47 |
| 41:DP:24:GLY:CA | 41:DP:33:ARG:HE | 2.28 | 0.47 |
| 43:DR:77:ARG:O | 43:DR:78:LYS:C | 2.57 | 0.47 |
| 44:DS:95:HIS:O | 44:DS:98:VAL:HG23 | 2.15 | 0.47 |
| 45:DT:50:ILE:HA | 45:DT:99:LEU:HD11 | 1.96 | 0.47 |
| 45:DT:89:VAL:HG12 | 45:DT:91:ARG:HG2 | 1.97 | 0.47 |
| 46:DU:92:ARG:NH2 | 47:DV:10:LYS:CB | 2.78 | 0.47 |
| 47:DV:90:PRO:CG | 47:DV:91:TYR:N | 2.74 | 0.47 |
| 50:DY:7:VAL:HB | 50:DY:8:LYS:CD | 2.42 | 0.47 |
| 1:AA:124:G:C5 | 1:AA:125:U:C4 | 3.02 | 0.47 |
| 1:AA:177:C:OP1 | 20:AT:65:LYS:HD3 | 2.14 | 0.47 |
| 1:AA:451:A:C5 | 1:AA:481:G:C5 | 3.03 | 0.47 |
| 1:AA:512:U:H2' | 1:AA:513:C:H6 | 1.80 | 0.47 |
| 1:AA:581:G:N2 | 1:AA:582:U:C4 | 2.83 | 0.47 |
| 1:AA:586:C:C2' | 1:AA:587:G:H5' | 2.45 | 0.47 |
| 1:AA:625:G:C4 | 1:AA:626:U:C5 | 3.02 | 0.47 |
| 1:AA:706:A:C5 | 1:AA:707:C:C5 | 3.03 | 0.47 |
| 1:AA:960:U:O2 | 1:AA:960:U:H2' | 2.13 | 0.47 |
| 2:AB:55:PHE:C | 2:AB:57:PHE:N | 2.72 | 0.47 |
| 6:AF:39:LYS:O | 6:AF:40:VAL:HB | 2.15 | 0.47 |
| 16:AP:16:HIS:O | 16:AP:17:TYR:O | 2.33 | 0.47 |
| 30:B8:35:GLN:HG2 | 31:BA:2420:C:OP1 | 2.13 | 0.47 |
| 30:B8:50:LEU:HD12 | 30:B8:51:ALA:H | 1.80 | 0.47 |
| 31:BA:221:A:N1 | 31:BA:265:A:O2' | 2.48 | 0.47 |
| 31:BA:1001:A:H2' | 31:BA:1002:G:O4' | 2.15 | 0.47 |
| 31:BA:1290:C:H2' | 31:BA:1291:C:H6 | 1.78 | 0.47 |
| 31:BA:1503:U:O2' | 31:BA:1504:C:H5' | 2.15 | 0.47 |
| 31:BA:2646:C:H6 | 31:BA:2646:C:O5' | 1.97 | 0.47 |
| 33:BD:246:PRO:C | 33:BD:254:THR:HG22 | 2.40 | 0.47 |
| 34:BE:48:GLN:NE2 | 34:BE:78:LEU:HD13 | 2.30 | 0.47 |
| 34:BE:96:PHE:HA | 34:BE:100:GLU:OE1 | 2.15 | 0.47 |
| 38:BI:93:THR:HG22 | 38:BI:119:PRO:HB3 | 1.95 | 0.47 |
| 40:BO:120:GLU:OE2 | 40:BO:122:LEU:HD21 | 2.14 | 0.47 |
| 41:BP:83:VAL:HG12 | 41:BP:112:LEU:HD21 | 1.95 | 0.47 |
| 41:BP:90:ARG:HD2 | 41:BP:91:PHE:HD1 | 1.79 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 43:BR:25:ALA:O | 43:BR:26:LYS:C | 2.58 | 0.47 |
| 44:BS:24:LEU:HD13 | 44:BS:24:LEU:HA | 1.76 | 0.47 |
| 45:BT:27:THR:OG1 | 45:BT:28:VAL:N | 2.47 | 0.47 |
| 46:BU:104:GLN:O | 46:BU:105:VAL:C | 2.57 | 0.47 |
| 48:BW:82:LEU:HG | 48:BW:84:ARG:NH2 | 2.30 | 0.47 |
| 1:CA:674:G:H2' | 1:CA:675:A:C8 | 2.48 | 0.47 |
| 1:CA:675:A:C6 | 1:CA:676:A:C5 | 3.03 | 0.47 |
| 2:CB:111:ARG:O | 2:CB:145:LEU:HD11 | 2.14 | 0.47 |
| 2:CB:144:ARG:HG3 | 2:CB:145:LEU:N | 2.30 | 0.47 |
| 4:CD:11:LEU:C | 4:CD:13:ARG:N | 2.62 | 0.47 |
| 4:CD:68:TYR:CD1 | 4:CD:68:TYR:N | 2.83 | 0.47 |
| 7:CG:155:ARG:O | 7:CG:156:TRP:HD1 | 1.98 | 0.47 |
| 12:CL:27:LEU:C | 12:CL:29:GLY:H | 2.22 | 0.47 |
| 12:CL:32:PHE:HE1 | 12:CL:86:ARG:HG3 | 1.80 | 0.47 |
| 12:CL:117:ARG:NH2 | 12:CL:124:LYS:HB2 | 2.30 | 0.47 |
| 25:D3:47:VAL:CG1 | 25:D3:56:VAL:HG21 | 2.44 | 0.47 |
| 30:D8:4:MET:HE1 | 31:DA:593:G:H1' | 1.96 | 0.47 |
| 30:D8:35:GLN:CA | 31:DA:2420:C:OP1 | 2.63 | 0.47 |
| 30:D8:37:SER:HB2 | 30:D8:38:GLY:H | 1.41 | 0.47 |
| 31:DA:259:G:H21 | 31:DA:621:A:H8 | 1.63 | 0.47 |
| 31:DA:267:C:H6 | 31:DA:267:C:H5" | 1.80 | 0.47 |
| 31:DA:477:A:H2' | 31:DA:478:A:C8 | 2.50 | 0.47 |
| 31:DA:629:G:H4' | 31:DA:650:C:O2 | 2.14 | 0.47 |
| 31:DA:644:A:O2' | 31:DA:645:C:H5" | 2.14 | 0.47 |
| 31:DA:1175:U:H4' | 31:DA:1176:G:H2' | 1.97 | 0.47 |
| 31:DA:1327:C:C2' | 31:DA:1328:G:H5' | 2.45 | 0.47 |
| 31:DA:1356:G:C6 | 31:DA:1357:U:C4 | 3.03 | 0.47 |
| 31:DA:1839:G:N3 | 31:DA:1839:G:H2' | 2.29 | 0.47 |
| 31:DA:2795:G:N2 | 31:DA:2796:U:O2' | 2.47 | 0.47 |
| 32:DB:87:G:H5" | 32:DB:88:C:OP2 | 2.15 | 0.47 |
| 32:DB:115:G:C3' | 32:DB:116:G:H5" | 2.45 | 0.47 |
| 33:DD:71:ASP:CB | 33:DD:103:ARG:NH2 | 2.77 | 0.47 |
| 33:DD:118:VAL:HG22 | 33:DD:119:ALA:N | 2.30 | 0.47 |
| 39:DN:3:THR:O | 39:DN:4:TYR:CD2 | 2.68 | 0.47 |
| 40:DO:17:ARG:HD3 | 40:DO:17:ARG:HA | 1.60 | 0.47 |
| 43:DR:20:LEU:O | 43:DR:21:TYR:C | 2.58 | 0.47 |
| 43:DR:51:LEU:CD1 | 43:DR:70:LEU:HD21 | 2.45 | 0.47 |
| 43:DR:74:LYS:O | 43:DR:75:LEU:C | 2.58 | 0.47 |
| 44:DS:53:SER:OG | 44:DS:54:LEU:N | 2.47 | 0.47 |
| 45:DT:26:ASP:HB3 | 45:DT:89:VAL:O | 2.15 | 0.47 |
| 46:DU:6:THR:HG21 | 46:DU:10:ARG:CZ | 2.44 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 49:DX:9:LEU:HD12 | 49:DX:30:VAL:C | 2.39 | 0.47 |
| 50:DY:28:LYS:O | 50:DY:29:GLU:C | 2.58 | 0.47 |
| 50:DY:41:GLY:O | 50:DY:43:ASN:OD1 | 2.32 | 0.47 |
| 50:DY:52:SER:C | 50:DY:54:LYS:N | 2.72 | 0.47 |
| 51:DZ:153:SER:O | 51:DZ:154:ASP:OD2 | 2.33 | 0.47 |
| 1:AA:191:G:N9 | 20:AT:105:SER:HB3 | 2.30 | 0.46 |
| 1:AA:303:A:H2' | 1:AA:304:U:O4' | 2.15 | 0.46 |
| 1:AA:677:U:H2' | 1:AA:678:U:H6 | 1.79 | 0.46 |
| 1:AA:708:C:O2' | 1:AA:709:G:H5' | 2.14 | 0.46 |
| 1:AA:709:G:H2' | 1:AA:710:G:C8 | 2.47 | 0.46 |
| 1:AA:1001(A):G:H2' | 1:AA:1002:G:O4' | 2.15 | 0.46 |
| 1:AA:1226:C:OP1 | 13:AM:91:ARG:NH1 | 2.47 | 0.46 |
| 1:AA:1248:A:C2' | 1:AA:1249:C:H5' | 2.44 | 0.46 |
| 1:AA:1368:G:OP2 | 9:AI:112:LYS:HD2 | 2.15 | 0.46 |
| 1:AA:1385:G:C2' | 1:AA:1386:G:H5' | 2.45 | 0.46 |
| 2:AB:185:ILE:HG12 | 2:AB:185:ILE:O | 2.16 | 0.46 |
| 5:AE:93:PRO:HA | 5:AE:118:ILE:HD12 | 1.98 | 0.46 |
| 8:AH:13:ILE:O | 8:AH:17:THR:HG23 | 2.15 | 0.46 |
| 13:AM:31:LYS:HA | 13:AM:34:LEU:HD12 | 1.97 | 0.46 |
| 20:AT:82:SER:O | 20:AT:86:ARG:HB2 | 2.14 | 0.46 |
| 24:B2:28:LYS:HE3 | 24:B2:28:LYS:HA | 1.97 | 0.46 |
| 27:B5:19:ARG:NH2 | 31:BA:1264:G:OP1 | 2.44 | 0.46 |
| 28:B6:16:CYS:O | 28:B6:17:LYS:HB2 | 2.14 | 0.46 |
| 30:B8:35:GLN:HB3 | 30:B8:36:LYS:H | 1.42 | 0.46 |
| 31:BA:315:G:H2' | 31:BA:316:C:O4' | 2.14 | 0.46 |
| 31:BA:528:A:N1 | 31:BA:2043:C:O5' | 2.48 | 0.46 |
| 31:BA:824:A:O2' | 31:BA:825:C:H5' | 2.15 | 0.46 |
| 31:BA:892:G:H3' | 31:BA:892:G:N3 | 2.30 | 0.46 |
| 31:BA:2011:U:H2' | 31:BA:2012:G:H5' | 1.97 | 0.46 |
| 31:BA:2199:A:H1' | 38:BI:28:ASN:HD22 | 1.80 | 0.46 |
| 31:BA:2516:G:C6 | 31:BA:2517:C:N4 | 2.83 | 0.46 |
| 31:BA:2701:C:C3' | 31:BA:2702:U:C5' | 2.69 | 0.46 |
| 31:BA:2808:U:N3 | 31:BA:2892:A:C5 | 2.82 | 0.46 |
| 34:BE:160:TYR:CD2 | 34:BE:161:GLY:N | 2.82 | 0.46 |
| 35:BF:183:VAL:O | 35:BF:184:TYR:C | 2.57 | 0.46 |
| 37:BH:54:ARG:N | 37:BH:65:HIS:HD2 | 2.13 | 0.46 |
| 38:BI:91:SER:HB3 | 38:BI:121:LYS:CB | 2.43 | 0.46 |
| 43:BR:18:LEU:HD13 | 43:BR:19:ALA:N | 2.30 | 0.46 |
| 44:BS:106:ARG:HE | 44:BS:106:ARG:HB3 | 1.39 | 0.46 |
| 45:BT:3:ARG:CB | 45:BT:6:LEU:HB3 | 2.37 | 0.46 |
| 49:BX:21:PHE:N | 49:BX:21:PHE:HD1 | 2.13 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 49:BX:83:VAL:O | 49:BX:83:VAL:HG23 | 2.15 | 0.46 |
| 1:CA:262:A:C6 | 1:CA:263:A:C6 | 3.03 | 0.46 |
| 1:CA:342:C:H2' | 1:CA:343:U:O4' | 2.14 | 0.46 |
| 1:CA:373:A:H2' | 1:CA:374:A:H8 | 1.80 | 0.46 |
| 1:CA:552:U:O2' | 12:CL:86:ARG:O | 2.33 | 0.46 |
| 4:CD:128:VAL:O | 4:CD:129:ASN:C | 2.58 | 0.46 |
| 4:CD:173:TRP:HA | 4:CD:187:ARG:NH1 | 2.30 | 0.46 |
| 9:CI:21:PRO:HA | 9:CI:58:ARG:O | 2.15 | 0.46 |
| 10:CJ:49:VAL:CG1 | 14:CN:41:ARG:HB2 | 2.44 | 0.46 |
| 15:CO:63:ARG:O | 15:CO:67:LEU:HB2 | 2.14 | 0.46 |
| 17:CQ:81:ARG:NH2 | 17:CQ:83:ASP:OD2 | 2.44 | 0.46 |
| 20:CT:84:LEU:C | 20:CT:84:LEU:HD13 | 2.40 | 0.46 |
| 30:D8:4:MET:CE | 31:DA:593:G:O4' | 2.64 | 0.46 |
| 31:DA:790:C:H6 | 31:DA:790:C:H2' | 1.58 | 0.46 |
| 31:DA:912:C:N3 | 31:DA:913:U:C5 | 2.84 | 0.46 |
| 31:DA:958:U:OP2 | 42:DQ:14:ARG:NH1 | 2.48 | 0.46 |
| 31:DA:1503:U:H2' | 31:DA:1504:C:C6 | 2.50 | 0.46 |
| 31:DA:1857:G:C6 | 31:DA:1858:G:C2 | 3.04 | 0.46 |
| 31:DA:1952:A:C5 | 40:DO:22:ILE:CD1 | 2.99 | 0.46 |
| 31:DA:2017:U:H5" | 31:DA:2018:G:P | 2.54 | 0.46 |
| 31:DA:2052:G:C4' | 34:DE:142:GLY:HA3 | 2.44 | 0.46 |
| 31:DA:2199:A:H3' | 31:DA:2200:C:H6 | 1.79 | 0.46 |
| 31:DA:2350:C:H2' | 31:DA:2351:G:O4' | 2.15 | 0.46 |
| 31:DA:2463:C:O2' | 31:DA:2464:C:H5' | 2.15 | 0.46 |
| 32:DB:41:U:C4 | 36:DG:70:VAL:O | 2.69 | 0.46 |
| 32:DB:46:A:H2' | 32:DB:47:C:H6 | 1.80 | 0.46 |
| 32:DB:50:G:P | 44:DS:63:THR:HG23 | 2.54 | 0.46 |
| 32:DB:86:G:H1 | 32:DB:91:C:N4 | 2.13 | 0.46 |
| 36:DG:128:ARG:O | 36:DG:129:GLY:C | 2.57 | 0.46 |
| 36:DG:135:LEU:HD12 | 36:DG:135:LEU:N | 2.30 | 0.46 |
| 38:DI:11:ASN:C | 38:DI:12:LEU:HD23 | 2.40 | 0.46 |
| 42:DQ:12:GLN:HG2 | 42:DQ:73:PRO:HD2 | 1.96 | 0.46 |
| 43:DR:97:VAL:HG22 | 43:DR:114:VAL:HG22 | 1.97 | 0.46 |
| 51:DZ:96:VAL:N | 51:DZ:128:VAL:O | 2.48 | 0.46 |
| 51:DZ:98:MET:HE2 | 51:DZ:100:VAL:HG23 | 1.96 | 0.46 |
| 51:DZ:152:ALA:HB2 | 51:DZ:168:GLU:CA | 2.44 | 0.46 |
| 1:AA:271:C:C2 | 1:AA:272:C:C5 | 3.03 | 0.46 |
| 1:AA:353:A:C2' | 1:AA:354:G:OP2 | 2.63 | 0.46 |
| 1:AA:561:U:HO2' | 1:AA:562:C:P | 2.38 | 0.46 |
| 1:AA:659:U:O2' | 1:AA:660:G:H5' | 2.15 | 0.46 |
| 1:AA:782:A:H4' | 1:AA:1514:C:O2' | 2.15 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:828:A:H5" | 1:AA:859:A:C2 | 2.50 | 0.46 |
| 4:AD:119:GLN:O | 4:AD:123:HIS:CD2 | 2.68 | 0.46 |
| 6:AF:82:ARG:HB2 | 6:AF:85:VAL:HG23 | 1.96 | 0.46 |
| 7:AG:152:ALA:O | 7:AG:155:ARG:HG3 | 2.14 | 0.46 |
| 10:AJ:75:ILE:HG13 | 10:AJ:76:ASN:H | 1.77 | 0.46 |
| 17:AQ:99:SER:O | 17:AQ:100:LYS:HD3 | 2.15 | 0.46 |
| 20:AT:76:ALA:O | 20:AT:77:ALA:C | 2.58 | 0.46 |
| 22:B0:25:ARG:HG3 | 22:B0:29:GLN:HE21 | 1.79 | 0.46 |
| 22:B0:72:ARG:O | 22:B0:75:LEU:HB2 | 2.14 | 0.46 |
| 27:B5:55:ARG:HD2 | 27:B5:56:LYS:N | 2.21 | 0.46 |
| 31:BA:271(P):C:H2' | 31:BA:271(Q):G:H5' | 1.96 | 0.46 |
| 31:BA:623:G:H2' | 31:BA:624:C:C6 | 2.51 | 0.46 |
| 31:BA:878:A:C6 | 31:BA:900:A:C8 | 3.03 | 0.46 |
| 31:BA:1362:C:O2' | 31:BA:1363:C:H5' | 2.15 | 0.46 |
| 31:BA:1607:C:N4 | 31:BA:1622:G:OP2 | 2.36 | 0.46 |
| 31:BA:1859:A:C2 | 31:BA:1884:A:H1' | 2.50 | 0.46 |
| 31:BA:1899:G:N2 | 31:BA:1902:C:C4 | 2.82 | 0.46 |
| 31:BA:2400:G:N3 | 31:BA:2400:G:H2' | 2.30 | 0.46 |
| 31:BA:2688:U:O2 | 31:BA:2688:U:O5' | 2.33 | 0.46 |
| 33:BD:25:THR:CG2 | 33:BD:82:ILE:H | 2.29 | 0.46 |
| 33:BD:202:LYS:HG2 | 33:BD:203:ASN:OD1 | 2.16 | 0.46 |
| 36:BG:91:ARG:O | 36:BG:91:ARG:HD2 | 2.15 | 0.46 |
| 37:BH:19:VAL:HB | 37:BH:44:VAL:HG13 | 1.98 | 0.46 |
| 39:BN:79:PRO:HD2 | 39:BN:80:GLY:H | 1.80 | 0.46 |
| 39:BN:125:GLY:CA | 39:BN:126:PRO:O | 2.63 | 0.46 |
| 40:BO:29:ASN:O | 40:BO:30:ALA:C | 2.58 | 0.46 |
| 40:BO:104:ARG:C | 40:BO:106:LEU:N | 2.73 | 0.46 |
| 44:BS:99:LYS:HD3 | 44:BS:99:LYS:N | 2.30 | 0.46 |
| 45:BT:124:ASP:C | 45:BT:126:ALA:N | 2.73 | 0.46 |
| 46:BU:76:TYR:CD2 | 46:BU:76:TYR:C | 2.94 | 0.46 |
| 47:BV:2:PHE:O | 47:BV:3:ALA:HB3 | 2.15 | 0.46 |
| 49:BX:25:LYS:HA | 49:BX:25:LYS:HE3 | 1.98 | 0.46 |
| 1:CA:16:A:C2 | 1:CA:17:U:C6 | 3.04 | 0.46 |
| 1:CA:109:A:H2' | 1:CA:326:G:H21 | 1.79 | 0.46 |
| 1:CA:373:A:C2 | 1:CA:374:A:C8 | 3.03 | 0.46 |
| 1:CA:414:A:C4 | 1:CA:415:A:C8 | 3.03 | 0.46 |
| 1:CA:458:C:H2' | 1:CA:460:G:C8 | 2.50 | 0.46 |
| 1:CA:678:U:H2' | 1:CA:679:C:C6 | 2.50 | 0.46 |
| 1:CA:719:C:H5 | 1:CA:720:C:N3 | 2.13 | 0.46 |
| 1:CA:735:C:H1' | 18:CR:75:ILE:HD11 | 1.96 | 0.46 |
| 1:CA:974:A:P | 14:CN:41:ARG:HH12 | 2.38 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:1157:A:H5' | 1:CA:1158:C:C6 | 2.50 | 0.46 |
| 1:CA:1203:C:H2' | 1:CA:1204:A:O4' | 2.16 | 0.46 |
| 5:CE:102:ALA:HB1 | 5:CE:106:PRO:HG2 | 1.96 | 0.46 |
| 6:CF:97:PHE:O | 18:CR:31:LEU:HD23 | 2.14 | 0.46 |
| 8:CH:64:LYS:CG | 8:CH:79:VAL:HG21 | 2.45 | 0.46 |
| 12:CL:10:LEU:HB3 | 17:CQ:32:TYR:CE1 | 2.50 | 0.46 |
| 12:CL:18:VAL:O | 12:CL:19:ARG:HB3 | 2.15 | 0.46 |
| 12:CL:25:PRO:C | 12:CL:27:LEU:N | 2.72 | 0.46 |
| 16:CP:5:ARG:HB3 | 16:CP:67:THR:OG1 | 2.16 | 0.46 |
| 24:D2:28:LYS:HA | 24:D2:28:LYS:HE3 | 1.98 | 0.46 |
| 27:D5:4:HIS:CB | 27:D5:5:PRO:CD | 2.91 | 0.46 |
| 27:D5:55:ARG:HD3 | 27:D5:55:ARG:HA | 1.41 | 0.46 |
| 31:DA:93:G:HO2' | 50:DY:51:VAL:HG21 | 1.79 | 0.46 |
| 31:DA:239:U:H2' | 31:DA:240:G:O4' | 2.16 | 0.46 |
| 31:DA:363(A):A:H2' | 31:DA:363(A):A:N3 | 2.30 | 0.46 |
| 31:DA:814:C:C5 | 41:DP:27:HIS:NE2 | 2.83 | 0.46 |
| 31:DA:954:G:C4 | 31:DA:955:C:C6 | 3.03 | 0.46 |
| 32:DB:79:C:O2' | 32:DB:80:U:H5' | 2.16 | 0.46 |
| 32:DB:119:G:H2' | 32:DB:119:G:N3 | 2.31 | 0.46 |
| 33:DD:183:ARG:HD2 | 33:DD:270:ILE:CG2 | 2.45 | 0.46 |
| 33:DD:209:ALA:C | 33:DD:210:GLY:O | 2.55 | 0.46 |
| 37:DH:141:VAL:HG12 | 37:DH:142:GLY:N | 2.30 | 0.46 |
| 38:DI:73:GLU:OE1 | 38:DI:137:PRO:HD2 | 2.15 | 0.46 |
| 42:DQ:34:LEU:CD1 | 42:DQ:129:THR:HB | 2.44 | 0.46 |
| 43:DR:2:ARG:HD2 | 43:DR:2:ARG:N | 2.30 | 0.46 |
| 43:DR:72:ASP:CG | 43:DR:75:LEU:H | 2.23 | 0.46 |
| 47:DV:52:VAL:O | 47:DV:53:GLU:CB | 2.62 | 0.46 |
| 47:DV:69:LYS:O | 47:DV:70:ILE:HB | 2.15 | 0.46 |
| 50:DY:100:ALA:O | 50:DY:101:LYS:CB | 2.63 | 0.46 |
| 51:DZ:127:LYS:HD3 | 51:DZ:162:GLU:OE1 | 2.15 | 0.46 |
| 1:AA:276:G:H2' | 1:AA:277:C:H5' | 1.96 | 0.46 |
| 1:AA:648:A:H2' | 1:AA:649:G:H8 | 1.81 | 0.46 |
| 1:AA:671:G:C4 | 1:AA:672:U:C6 | 3.04 | 0.46 |
| 1:AA:954:G:C2 | 1:AA:955:U:C2 | 3.02 | 0.46 |
| 1:AA:983:A:H5' | 1:AA:984:C:OP2 | 2.15 | 0.46 |
| 1:AA:1255:G:HO2' | 1:AA:1258:G:HO2' | 1.64 | 0.46 |
| 2:AB:97:TRP:CH2 | 2:AB:176:GLU:OE2 | 2.69 | 0.46 |
| 4:AD:148:VAL:CG1 | 4:AD:149:ALA:H | 2.24 | 0.46 |
| 5:AE:12:LEU:HD13 | 5:AE:31:LEU:HB2 | 1.96 | 0.46 |
| 24:B2:14:ARG:NH2 | 24:B2:57:ILE:HG22 | 2.29 | 0.46 |
| 27:B5:32:PRO:O | 27:B5:33:CYS:CB | 2.59 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:7:G:C2' | 31:BA:8:A:O4' | 2.63 | 0.46 |
| 31:BA:154:G:O6 | 31:BA:172:C:N4 | 2.48 | 0.46 |
| 31:BA:272(B):G:O2' | 31:BA:272(C):G:C5' | 2.63 | 0.46 |
| 31:BA:286:C:H2' | 31:BA:286:C:O2 | 2.16 | 0.46 |
| 31:BA:560:C:H5' | 46:BU:52:ARG:HH12 | 1.80 | 0.46 |
| 31:BA:814:C:N4 | 41:BP:27:HIS:NE2 | 2.63 | 0.46 |
| 31:BA:1331:A:H2' | 31:BA:1333:C:C5 | 2.51 | 0.46 |
| 31:BA:1434:A:C2 | 31:BA:1435:G:C4 | 3.04 | 0.46 |
| 31:BA:1952:A:C6 | 31:BA:1953:A:N1 | 2.83 | 0.46 |
| 31:BA:2274:A:C5 | 31:BA:2276:G:C8 | 3.03 | 0.46 |
| 31:BA:2394:C:C2' | 31:BA:2395:C:H5' | 2.45 | 0.46 |
| 31:BA:2895:U:C5 | 31:BA:2896:C:C5 | 3.04 | 0.46 |
| 33:BD:70:TRP:CZ2 | 33:BD:150:LYS:HD3 | 2.51 | 0.46 |
| 37:BH:153:LYS:N | 37:BH:153:LYS:CD | 2.76 | 0.46 |
| 38:BI:94:ALA:HB2 | 38:BI:116:LEU:HD23 | 1.97 | 0.46 |
| 39:BN:18:ALA:HB1 | 39:BN:21:LYS:CB | 2.44 | 0.46 |
| 39:BN:66:LYS:H | 39:BN:66:LYS:HG2 | 1.59 | 0.46 |
| 40:BO:10:VAL:HG13 | 40:BO:17:ARG:C | 2.40 | 0.46 |
| 41:BP:7:ARG:HD2 | 41:BP:7:ARG:HA | 1.66 | 0.46 |
| 41:BP:21:ARG:O | 41:BP:23:PRO:HD3 | 2.16 | 0.46 |
| 41:BP:45:LEU:HD22 | 41:BP:46:LYS:H | 1.79 | 0.46 |
| 44:BS:35:ILE:O | 44:BS:35:ILE:HG23 | 2.15 | 0.46 |
| 45:BT:109:GLU:CA | 45:BT:112:ARG:HG3 | 2.43 | 0.46 |
| 45:BT:115:ARG:HB3 | 45:BT:116:ALA:H | 1.55 | 0.46 |
| 47:BV:25:LEU:C | 47:BV:27:ALA:N | 2.68 | 0.46 |
| 48:BW:88:ARG:NH1 | 48:BW:94:ASP:OD1 | 2.42 | 0.46 |
| 49:BX:82:GLN:HB3 | 49:BX:85:PRO:CG | 2.41 | 0.46 |
| 1:CA:66:G:C6 | 1:CA:67:C:C5 | 3.03 | 0.46 |
| 1:CA:473:G:N3 | 1:CA:474:G:C8 | 2.83 | 0.46 |
| 1:CA:679:C:O2' | 1:CA:680:C:H5' | 2.15 | 0.46 |
| 1:CA:1277:C:H2' | 1:CA:1278:U:H5' | 1.97 | 0.46 |
| 2:CB:187:LEU:HA | 2:CB:201:ILE:HB | 1.98 | 0.46 |
| 4:CD:8:VAL:O | 4:CD:10:ARG:N | 2.48 | 0.46 |
| 4:CD:173:TRP:O | 4:CD:174:LEU:HD23 | 2.15 | 0.46 |
| 5:CE:48:ALA:HB1 | 5:CE:49:PRO:HD2 | 1.98 | 0.46 |
| 5:CE:69:VAL:HG12 | 5:CE:71:LEU:CD2 | 2.44 | 0.46 |
| 7:CG:106:GLN:O | 7:CG:110:GLN:HG3 | 2.15 | 0.46 |
| 8:CH:112:LEU:HD12 | 8:CH:112:LEU:O | 2.16 | 0.46 |
| 10:CJ:82:ILE:O | 10:CJ:86:MET:HB3 | 2.16 | 0.46 |
| 11:CK:69:ALA:HB1 | 11:CK:103:LEU:CD2 | 2.45 | 0.46 |
| 31:DA:271(N):U:C6 | 31:DA:271(N):U:OP1 | 2.69 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:272(J):C:O2' | 31:DA:274:G:OP1 | 2.29 | 0.46 |
| 31:DA:685:A:C2 | 31:DA:689:A:C6 | 3.04 | 0.46 |
| 31:DA:700:G:H2' | 31:DA:701:G:O4' | 2.15 | 0.46 |
| 31:DA:706:A:H8 | 31:DA:706:A:O5' | 1.99 | 0.46 |
| 31:DA:1169:G:C8 | 31:DA:1169:G:C3' | 2.98 | 0.46 |
| 31:DA:1274:A:N3 | 31:DA:1297:C:H1' | 2.30 | 0.46 |
| 31:DA:1671:U:O2' | 31:DA:1673:U:H5 | 1.98 | 0.46 |
| 31:DA:1783:A:N1 | 31:DA:2587:A:N3 | 2.63 | 0.46 |
| 31:DA:2584:U:O2 | 31:DA:2584:U:O4' | 2.33 | 0.46 |
| 31:DA:2810:A:H2' | 34:DE:61:ARG:CZ | 2.43 | 0.46 |
| 33:DD:30:GLU:CG | 33:DD:63:ARG:CZ | 2.93 | 0.46 |
| 33:DD:48:ARG:HG3 | 33:DD:48:ARG:HH11 | 1.81 | 0.46 |
| 35:DF:23:ASP:O | 35:DF:24:LEU:HD22 | 2.15 | 0.46 |
| 35:DF:157:VAL:HB | 35:DF:194:MET:HB3 | 1.98 | 0.46 |
| 36:DG:63:ILE:O | 36:DG:63:ILE:HD12 | 2.15 | 0.46 |
| 38:DI:14:ASP:O | 38:DI:15:VAL:C | 2.58 | 0.46 |
| 38:DI:90:GLY:O | 38:DI:91:SER:O | 2.33 | 0.46 |
| 39:DN:19:GLU:HG3 | 39:DN:20:GLY:N | 2.28 | 0.46 |
| 40:DO:66:LYS:H | 40:DO:82:ASN:HD21 | 1.55 | 0.46 |
| 41:DP:7:ARG:HD2 | 41:DP:7:ARG:HA | 1.65 | 0.46 |
| 41:DP:8:PRO:C | 41:DP:10:PRO:HD3 | 2.41 | 0.46 |
| 41:DP:125:VAL:O | 41:DP:125:VAL:HG22 | 2.15 | 0.46 |
| 42:DQ:7:MET:O | 42:DQ:10:ARG:NH2 | 2.48 | 0.46 |
| 42:DQ:87:LYS:HE3 | 42:DQ:87:LYS:HB2 | 1.77 | 0.46 |
| 45:DT:28:VAL:HG13 | 45:DT:46:GLU:HA | 1.96 | 0.46 |
| 45:DT:82:LEU:N | 45:DT:82:LEU:CD1 | 2.73 | 0.46 |
| 45:DT:129:ARG:HH11 | 45:DT:131:ALA:HB3 | 1.80 | 0.46 |
| 47:DV:62:LEU:HB3 | 47:DV:98:GLU:CA | 2.44 | 0.46 |
| 48:DW:62:HIS:O | 48:DW:63:ASP:C | 2.58 | 0.46 |
| 50:DY:17:SER:HB2 | 50:DY:71:LYS:HE2 | 1.96 | 0.46 |
| 51:DZ:69:THR:HG22 | 51:DZ:90:VAL:CA | 2.35 | 0.46 |
| 1:AA:24:U:O2' | 1:AA:25:C:H5' | 2.16 | 0.46 |
| 1:AA:149:A:O2' | 1:AA:150:C:P | 2.73 | 0.46 |
| 1:AA:544:G:H2' | 1:AA:545:C:H6 | 1.81 | 0.46 |
| 1:AA:724:G:N3 | 1:AA:725:G:C8 | 2.83 | 0.46 |
| 2:AB:9:GLU:HA | 2:AB:12:GLU:OE1 | 2.15 | 0.46 |
| 3:AC:152:ILE:HB | 3:AC:199:LYS:HB2 | 1.96 | 0.46 |
| 4:AD:173:TRP:HA | 4:AD:187:ARG:NH1 | 2.31 | 0.46 |
| 5:AE:6:PHE:CD2 | 5:AE:36:ASP:HB3 | 2.51 | 0.46 |
| 5:AE:6:PHE:CE1 | 5:AE:66:MET:HE1 | 2.51 | 0.46 |
| 9:AI:53:VAL:CG1 | 9:AI:95:LYS:HE3 | 2.45 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 12:AL:25:PRO:C | 12:AL:27:LEU:N | 2.74 | 0.46 |
| 18:AR:53:ARG:C | 18:AR:55:ARG:H | 2.24 | 0.46 |
| 23:B1:94:LEU:HD22 | 23:B1:95:LEU:N | 2.30 | 0.46 |
| 31:BA:107:C:C2 | 31:BA:108:U:C5 | 3.03 | 0.46 |
| 31:BA:481:G:OP2 | 50:BY:47:LYS:HE2 | 2.15 | 0.46 |
| 31:BA:775:G:C5 | 31:BA:794:G:C8 | 3.04 | 0.46 |
| 31:BA:1142(A):A:C4 | 31:BA:1144:G:C8 | 3.04 | 0.46 |
| 31:BA:1176:G:H1' | 31:BA:1177:A:OP1 | 2.15 | 0.46 |
| 31:BA:1721:G:N2 | 31:BA:1739:U:OP2 | 2.48 | 0.46 |
| 31:BA:1777:U:HO2' | 31:BA:1778:U:H5' | 1.77 | 0.46 |
| 31:BA:2102:U:C6 | 31:BA:2187:G:O6 | 2.68 | 0.46 |
| 31:BA:2190:G:C2' | 31:BA:2191:G:H5' | 2.45 | 0.46 |
| 31:BA:2324:C:H5" | 31:BA:2325:G:H5' | 1.98 | 0.46 |
| 31:BA:2346:A:H5" | 31:BA:2383:G:H1' | 1.97 | 0.46 |
| 31:BA:2394:C:P | 41:BP:63:PRO:HD2 | 2.51 | 0.46 |
| 31:BA:2494:G:H2' | 31:BA:2495:G:O5' | 2.15 | 0.46 |
| 31:BA:2699:C:H2' | 31:BA:2700:C:O4' | 2.16 | 0.46 |
| 33:BD:248:SER:HB2 | 33:BD:249:PRO:HD2 | 1.96 | 0.46 |
| 38:BI:33:ARG:O | 38:BI:35:LEU:HD23 | 2.15 | 0.46 |
| 39:BN:100:GLU:O | 39:BN:101:HIS:C | 2.57 | 0.46 |
| 40:BO:22:ILE:HD13 | 40:BO:22:ILE:HA | 1.63 | 0.46 |
| 45:BT:29:ARG:HD2 | 45:BT:29:ARG:HA | 1.46 | 0.46 |
| 47:BV:47:VAL:CG1 | 47:BV:48:GLY:H | 2.10 | 0.46 |
| 51:BZ:44:PHE:CZ | 51:BZ:48:PHE:HD2 | 2.34 | 0.46 |
| 51:BZ:112:ARG:C | 51:BZ:114:GLY:H | 2.22 | 0.46 |
| 1:CA:397:A:N7 | 1:CA:548:G:C8 | 2.84 | 0.46 |
| 1:CA:451:A:C5 | 1:CA:481:G:C6 | 3.03 | 0.46 |
| 1:CA:544:G:H2' | 1:CA:545:C:H6 | 1.80 | 0.46 |
| 1:CA:691:G:H1' | 1:CA:696:A:N6 | 2.30 | 0.46 |
| 1:CA:801:U:H2' | 1:CA:802:A:C8 | 2.48 | 0.46 |
| 1:CA:923:A:H61 | 1:CA:1393:U:H3 | 1.62 | 0.46 |
| 1:CA:950:U:H2' | 1:CA:951:G:H8 | 1.80 | 0.46 |
| 1:CA:1522:U:C2 | 1:CA:1523:G:C8 | 3.03 | 0.46 |
| 3:CC:124:ILE:HG13 | 3:CC:130:VAL:HG22 | 1.98 | 0.46 |
| 4:CD:68:TYR:HD1 | 4:CD:68:TYR:H | 1.63 | 0.46 |
| 5:CE:139:LEU:HD23 | 5:CE:142:LEU:HD11 | 1.98 | 0.46 |
| 13:CM:68:GLY:O | 13:CM:70:LEU:N | 2.49 | 0.46 |
| 15:CO:60:VAL:HG11 | 31:DA:715:G:O4' | 2.15 | 0.46 |
| 23:D1:10:LYS:O | 23:D1:13:ILE:CG2 | 2.63 | 0.46 |
| 23:D1:20:ARG:CD | 23:D1:41:ARG:HD3 | 2.45 | 0.46 |
| 31:DA:271(G):C:O2 | 31:DA:271(G):C:H2' | 2.15 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:513:A:C2 | 31:DA:514:A:C8 | 3.04 | 0.46 |
| 31:DA:922:U:H2' | 31:DA:923:C:C6 | 2.50 | 0.46 |
| 31:DA:1142(A):A:C4 | 31:DA:1144:G:C8 | 3.03 | 0.46 |
| 31:DA:1235:G:C6 | 31:DA:1236:G:N1 | 2.83 | 0.46 |
| 31:DA:1601:G:O2' | 31:DA:1602:U:H5' | 2.15 | 0.46 |
| 31:DA:2006:C:O5' | 31:DA:2006:C:H6 | 1.98 | 0.46 |
| 31:DA:2094:G:H5' | 38:DI:25:TYR:CD2 | 2.50 | 0.46 |
| 31:DA:2273:A:C2' | 31:DA:2274:A:H5' | 2.45 | 0.46 |
| 31:DA:2483:C:O2 | 31:DA:2483:C:C2' | 2.63 | 0.46 |
| 31:DA:2689:U:H5" | 31:DA:2690:C:H5' | 1.97 | 0.46 |
| 32:DB:31:C:N4 | 44:DS:32:LEU:CD2 | 2.78 | 0.46 |
| 32:DB:35:U:O2' | 32:DB:36:C:H5' | 2.14 | 0.46 |
| 33:DD:4:LYS:HZ1 | 33:DD:20:ASP:HA | 1.79 | 0.46 |
| 35:DF:32:LEU:CD1 | 35:DF:105:VAL:HG13 | 2.41 | 0.46 |
| 41:DP:23:PRO:O | 41:DP:33:ARG:NE | 2.48 | 0.46 |
| 46:DU:66:ASN:ND2 | 46:DU:66:ASN:C | 2.74 | 0.46 |
| 47:DV:66:ARG:HG2 | 47:DV:94:LEU:CG | 2.43 | 0.46 |
| 51:DZ:44:PHE:CZ | 51:DZ:48:PHE:CD2 | 3.03 | 0.46 |
| 51:DZ:44:PHE:CE1 | 51:DZ:48:PHE:HB2 | 2.51 | 0.46 |
| 51:DZ:67:LEU:HD12 | 51:DZ:67:LEU:N | 2.30 | 0.46 |
| 1:AA:255:G:O6 | 1:AA:266:G:O6 | 2.33 | 0.46 |
| 1:AA:390:C:O2' | 1:AA:391:G:H5' | 2.16 | 0.46 |
| 1:AA:503:C:C6 | 1:AA:504:C:H5 | 2.34 | 0.46 |
| 1:AA:577:G:H1' | 1:AA:816:A:N3 | 2.30 | 0.46 |
| 1:AA:945:G:H2' | 1:AA:945:G:N3 | 2.30 | 0.46 |
| 1:AA:1371:G:O3' | 9:AI:69:GLY:HA3 | 2.15 | 0.46 |
| 1:AA:1452:C:H5' | 1:AA:1456:G:C5 | 2.51 | 0.46 |
| 4:AD:36:ARG:HG2 | 4:AD:38:TYR:CZ | 2.51 | 0.46 |
| 4:AD:92:VAL:O | 4:AD:96:LEU:CD2 | 2.64 | 0.46 |
| 6:AF:5:GLU:HG2 | 6:AF:62:TRP:HZ2 | 1.80 | 0.46 |
| 6:AF:40:VAL:HA | 6:AF:62:TRP:O | 2.15 | 0.46 |
| 8:AH:29:SER:HB3 | 8:AH:32:LYS:CD | 2.46 | 0.46 |
| 12:AL:27:LEU:C | 12:AL:29:GLY:H | 2.23 | 0.46 |
| 13:AM:14:ARG:HA | 13:AM:43:THR:O | 2.15 | 0.46 |
| 22:B0:2:ALA:HA | 31:BA:2494:G:OP1 | 2.15 | 0.46 |
| 23:B1:67:ILE:O | 23:B1:70:VAL:HB | 2.15 | 0.46 |
| 26:B4:25:TYR:C | 26:B4:27:THR:N | 2.74 | 0.46 |
| 31:BA:83:G:N1 | 31:BA:102:G:H2' | 2.28 | 0.46 |
| 31:BA:108:U:C2 | 31:BA:109:G:C8 | 3.04 | 0.46 |
| 31:BA:466:A:C2' | 31:BA:467:G:H5' | 2.45 | 0.46 |
| 31:BA:547:A:H8 | 31:BA:549:G:O6 | 1.98 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:633:A:H2' | 31:BA:634:C:H5' | 1.98 | 0.46 |
| 31:BA:1022:G:N2 | 31:BA:1142(A):A:H2 | 2.12 | 0.46 |
| 31:BA:1106:A:C2' | 31:BA:1107:G:C8 | 2.98 | 0.46 |
| 31:BA:1403:C:H2' | 31:BA:1404:C:O5' | 2.16 | 0.46 |
| 31:BA:1628:G:H2' | 31:BA:1629:U:C6 | 2.50 | 0.46 |
| 31:BA:2093:G:O5' | 38:BI:24:GLY:HA3 | 2.16 | 0.46 |
| 31:BA:2636:U:O2' | 31:BA:2637:U:H5' | 2.15 | 0.46 |
| 31:BA:2637:U:C2' | 31:BA:2638:G:O5' | 2.64 | 0.46 |
| 31:BA:2734:A:H8 | 31:BA:2734:A:H5' | 1.80 | 0.46 |
| 32:BB:40:U:O2' | 32:BB:41:U:OP1 | 2.26 | 0.46 |
| 32:BB:87:G:H5" | 32:BB:88:C:OP2 | 2.15 | 0.46 |
| 33:BD:181:GLU:O | 33:BD:182:LEU:HD23 | 2.15 | 0.46 |
| 35:BF:9:ILE:HG12 | 35:BF:14:PRO:HA | 1.97 | 0.46 |
| 35:BF:132:VAL:O | 35:BF:134:GLY:N | 2.48 | 0.46 |
| 43:BR:34:ILE:HA | 43:BR:34:ILE:HD12 | 1.55 | 0.46 |
| 43:BR:77:ARG:O | 43:BR:78:LYS:C | 2.58 | 0.46 |
| 46:BU:50:ARG:NH2 | 47:BV:75:PHE:HD2 | 2.13 | 0.46 |
| 1:CA:432:A:N7 | 1:CA:433:C:C2 | 2.84 | 0.46 |
| 1:CA:501:C:H2' | 1:CA:502:G:H8 | 1.81 | 0.46 |
| 1:CA:559:A:H4' | 1:CA:560:U:O5' | 2.15 | 0.46 |
| 1:CA:731:G:OP1 | 1:CA:766:A:H1' | 2.15 | 0.46 |
| 1:CA:782:A:H4' | 1:CA:1514:C:O2' | 2.15 | 0.46 |
| 1:CA:976:G:P | 14:CN:32:SER:H | 2.37 | 0.46 |
| 1:CA:1312:G:N2 | 1:CA:1326:C:C2 | 2.84 | 0.46 |
| 1:CA:1446:U:O2 | 1:CA:1456:G:O6 | 2.33 | 0.46 |
| 2:CB:67:THR:HG21 | 2:CB:155:LEU:CD2 | 2.44 | 0.46 |
| 2:CB:69:LEU:HB3 | 2:CB:162:ILE:HG22 | 1.98 | 0.46 |
| 3:CC:186:PHE:CE2 | 3:CC:188:LEU:HD21 | 2.51 | 0.46 |
| 4:CD:92:VAL:O | 4:CD:96:LEU:CD2 | 2.63 | 0.46 |
| 6:CF:91:VAL:HG12 | 6:CF:92:LYS:O | 2.15 | 0.46 |
| 7:CG:100:ALA:O | 7:CG:104:LEU:HD23 | 2.14 | 0.46 |
| 16:CP:43:LYS:CG | 16:CP:48:TRP:CE3 | 2.98 | 0.46 |
| 18:CR:44:LEU:HD23 | 18:CR:48:GLY:O | 2.15 | 0.46 |
| 19:CS:22:LEU:O | 19:CS:26:GLY:HA2 | 2.16 | 0.46 |
| 20:CT:87:LYS:HE3 | 20:CT:91:LEU:HD11 | 1.98 | 0.46 |
| 23:D1:79:GLY:O | 23:D1:80:LEU:HD23 | 2.16 | 0.46 |
| 27:D5:40:LYS:NZ | 27:D5:49:CYS:SG | 2.67 | 0.46 |
| 31:DA:9:U:C5 | 31:DA:2629:A:N6 | 2.84 | 0.46 |
| 31:DA:272(G):C:N4 | 31:DA:363(C):G:H1 | 2.13 | 0.46 |
| 31:DA:892:G:H2' | 31:DA:893:C:O4' | 2.16 | 0.46 |
| 31:DA:930:U:O4' | 31:DA:930:U:O2 | 2.31 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1047:G:C2 | 31:DA:1111:A:N6 | 2.84 | 0.46 |
| 31:DA:1176:G:C1' | 31:DA:1177:A:OP1 | 2.63 | 0.46 |
| 31:DA:1190:G:H5' | 41:DP:35:HIS:HB3 | 1.97 | 0.46 |
| 31:DA:1378:A:O2' | 31:DA:1379:A:C5' | 2.54 | 0.46 |
| 31:DA:1407:C:H2' | 31:DA:1407:C:O2 | 2.15 | 0.46 |
| 31:DA:1486:A:H2' | 31:DA:1487:G:H8 | 1.80 | 0.46 |
| 31:DA:1531:C:H5" | 31:DA:1532:C:H6 | 1.80 | 0.46 |
| 31:DA:2467:C:H2' | 31:DA:2468:G:O4' | 2.16 | 0.46 |
| 31:DA:2562:U:C2' | 31:DA:2563:U:H5' | 2.45 | 0.46 |
| 31:DA:2734:A:C5' | 31:DA:2734:A:C8 | 2.99 | 0.46 |
| 32:DB:46:A:C4 | 32:DB:47:C:C5 | 3.04 | 0.46 |
| 33:DD:221:VAL:HG22 | 33:DD:226:MET:HE3 | 1.98 | 0.46 |
| 34:DE:36:ARG:NH1 | 34:DE:85:ASN:ND2 | 2.64 | 0.46 |
| 37:DH:56:SER:OG | 37:DH:61:HIS:ND1 | 2.49 | 0.46 |
| 38:DI:8:PRO:O | 38:DI:9:LEU:HD23 | 2.15 | 0.46 |
| 39:DN:30:ILE:HG23 | 39:DN:52:VAL:HG11 | 1.97 | 0.46 |
| 39:DN:78:TYR:N | 39:DN:79:PRO:CD | 2.70 | 0.46 |
| 44:DS:95:HIS:CG | 44:DS:96:GLY:N | 2.80 | 0.46 |
| 45:DT:28:VAL:HG22 | 45:DT:47:GLY:N | 2.31 | 0.46 |
| 47:DV:25:LEU:C | 47:DV:27:ALA:N | 2.70 | 0.46 |
| 49:DX:12:VAL:HG13 | 49:DX:12:VAL:O | 2.14 | 0.46 |
| 49:DX:75:ASP:O | 49:DX:76:ARG:HB3 | 2.15 | 0.46 |
| 49:DX:77:LYS:CD | 49:DX:78:LYS:HG3 | 2.45 | 0.46 |
| 51:DZ:48:PHE:C | 51:DZ:48:PHE:CD1 | 2.93 | 0.46 |
| 51:DZ:139:VAL:C | 51:DZ:141:VAL:H | 2.23 | 0.46 |
| 1:AA:35:G:O2' | 12:AL:121:GLY:HA2 | 2.15 | 0.46 |
| 1:AA:247:G:C6 | 1:AA:248:C:C5 | 3.04 | 0.46 |
| 1:AA:327:A:C2 | 1:AA:329:A:C4 | 3.03 | 0.46 |
| 1:AA:407:G:H4' | 4:AD:115:ARG:O | 2.15 | 0.46 |
| 1:AA:504:C:H1' | 1:AA:510:A:C4 | 2.51 | 0.46 |
| 1:AA:514:C:O2' | 1:AA:515:G:H5' | 2.16 | 0.46 |
| 1:AA:674:G:H2' | 1:AA:675:A:C8 | 2.48 | 0.46 |
| 1:AA:1097:C:O2 | 1:AA:1169:A:H2 | 1.98 | 0.46 |
| 2:AB:14:GLY:C | 2:AB:15:VAL:HG22 | 2.40 | 0.46 |
| 2:AB:51:LEU:HD21 | 2:AB:214:ILE:HD12 | 1.97 | 0.46 |
| 9:AI:77:ILE:O | 9:AI:81:ILE:HG12 | 2.15 | 0.46 |
| 11:AK:81:ASP:CG | 11:AK:106:LYS:HG2 | 2.40 | 0.46 |
| 16:AP:20:VAL:HG21 | 16:AP:32:TYR:CB | 2.46 | 0.46 |
| 20:AT:26:ASN:HA | 20:AT:29:LYS:HG2 | 1.98 | 0.46 |
| 22:B0:50:ASN:O | 22:B0:62:LEU:HB2 | 2.16 | 0.46 |
| 24:B2:27:GLU:O | 24:B2:30:ARG:HG2 | 2.15 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 27:B5:55:ARG:HD3 | 27:B5:55:ARG:HA | 1.43 | 0.46 |
| 28:B6:33:LYS:HD3 | 28:B6:33:LYS:HA | 1.54 | 0.46 |
| 29:B7:19:ARG:HG2 | 29:B7:19:ARG:NH1 | 2.31 | 0.46 |
| 30:B8:32:LEU:CA | 30:B8:34:TRP:H | 2.29 | 0.46 |
| 31:BA:125:G:H4' | 31:BA:126:A:OP2 | 2.16 | 0.46 |
| 31:BA:174:C:C3' | 31:BA:175:G:H5" | 2.45 | 0.46 |
| 31:BA:213:A:H2' | 31:BA:214:G:O4' | 2.15 | 0.46 |
| 31:BA:527:C:OP2 | 31:BA:2779:U:C5 | 2.57 | 0.46 |
| 31:BA:883:G:H1 | 31:BA:893:C:H41 | 1.63 | 0.46 |
| 31:BA:1439:A:C2 | 31:BA:1553:A:C5 | 3.03 | 0.46 |
| 31:BA:1608:A:HO2' | 31:BA:1610:A:P | 2.39 | 0.46 |
| 31:BA:1834:U:O2 | 31:BA:1834:U:H2' | 2.15 | 0.46 |
| 31:BA:1892:C:O5' | 31:BA:1892:C:H6 | 1.99 | 0.46 |
| 31:BA:2531:A:H2 | 31:BA:2658:C:O2 | 1.99 | 0.46 |
| 31:BA:2664:G:C2' | 31:BA:2665:A:O5' | 2.64 | 0.46 |
| 33:BD:25:THR:HG21 | 33:BD:81:ALA:CA | 2.46 | 0.46 |
| 33:BD:79:VAL:HG12 | 33:BD:113:VAL:HA | 1.98 | 0.46 |
| 33:BD:147:LEU:HD13 | 33:BD:155:LEU:HD11 | 1.96 | 0.46 |
| 35:BF:66:PRO:O | 35:BF:67:GLN:CB | 2.56 | 0.46 |
| 35:BF:161:GLU:O | 35:BF:165:ARG:HG2 | 2.16 | 0.46 |
| 36:BG:45:GLU:HB2 | 36:BG:47:LYS:CD | 2.46 | 0.46 |
| 37:BH:55:PRO:O | 37:BH:56:SER:O | 2.34 | 0.46 |
| 37:BH:153:LYS:CB | 37:BH:154:PRO:CD | 2.93 | 0.46 |
| 39:BN:78:TYR:CD1 | 39:BN:79:PRO:HD3 | 2.50 | 0.46 |
| 41:BP:16:ARG:NH1 | 41:BP:16:ARG:O | 2.41 | 0.46 |
| 44:BS:95:HIS:O | 44:BS:96:GLY:C | 2.58 | 0.46 |
| 48:BW:86:LEU:HD12 | 48:BW:87:PRO:N | 2.30 | 0.46 |
| 49:BX:9:LEU:HB2 | 49:BX:29:TRP:O | 2.15 | 0.46 |
| 49:BX:12:VAL:HG13 | 49:BX:17:ALA:HB2 | 1.97 | 0.46 |
| 50:BY:10:GLY:C | 50:BY:27:VAL:HG22 | 2.40 | 0.46 |
| 50:BY:14:LEU:HD12 | 50:BY:23:ARG:H | 1.81 | 0.46 |
| 51:BZ:8:TYR:O | 51:BZ:37:VAL:HG12 | 2.16 | 0.46 |
| 1:CA:586:C:H2' | 1:CA:587:G:H5' | 1.98 | 0.46 |
| 1:CA:685:G:N2 | 1:CA:686:U:N3 | 2.64 | 0.46 |
| 1:CA:827:U:C4 | 1:CA:870:U:N3 | 2.84 | 0.46 |
| 1:CA:863:U:H2' | 1:CA:865:A:OP2 | 2.16 | 0.46 |
| 1:CA:1064:G:OP2 | 1:CA:1386:G:H4' | 2.16 | 0.46 |
| 1:CA:1377:A:H2' | 7:CG:7:ALA:CB | 2.45 | 0.46 |
| 1:CA:1386:G:C2 | 1:CA:1387:G:C8 | 3.04 | 0.46 |
| 1:CA:1478:C:H2' | 1:CA:1479:C:H6 | 1.80 | 0.46 |
| 3:CC:108:ASN:HB3 | 3:CC:111:LEU:HB2 | 1.97 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 4:CD:2:GLY:O | 4:CD:4:TYR:N | 2.48 | 0.46 |
| 4:CD:126:ILE:HG22 | 4:CD:127:THR:N | 2.27 | 0.46 |
| 7:CG:49:ILE:HG22 | 7:CG:49:ILE:O | 2.15 | 0.46 |
| 8:CH:107:LEU:N | 8:CH:107:LEU:HD23 | 2.30 | 0.46 |
| 12:CL:34:ARG:HG2 | 12:CL:35:GLY:N | 2.30 | 0.46 |
| 16:CP:21:VAL:O | 16:CP:33:ILE:HB | 2.16 | 0.46 |
| 16:CP:43:LYS:HG2 | 16:CP:48:TRP:CE3 | 2.50 | 0.46 |
| 24:D2:18:PRO:O | 24:D2:19:VAL:C | 2.58 | 0.46 |
| 31:DA:218:A:H2 | 31:DA:235:U:H4' | 1.81 | 0.46 |
| 31:DA:272(B):G:O2' | 31:DA:272(C):G:O4' | 2.26 | 0.46 |
| 31:DA:535:C:H2' | 31:DA:536:A:H5' | 1.95 | 0.46 |
| 31:DA:670:A:H8 | 31:DA:670:A:H5" | 1.81 | 0.46 |
| 31:DA:854:G:C4 | 31:DA:855:G:C8 | 3.03 | 0.46 |
| 31:DA:1176:G:H1' | 31:DA:1177:A:OP1 | 2.15 | 0.46 |
| 31:DA:1767:C:C2' | 31:DA:1768:U:H5' | 2.46 | 0.46 |
| 31:DA:2243:U:O2 | 31:DA:2434:A:C2 | 2.69 | 0.46 |
| 31:DA:2544:G:H8 | 31:DA:2544:G:O5' | 1.98 | 0.46 |
| 31:DA:2753:A:O2' | 31:DA:2754:U:P | 2.74 | 0.46 |
| 31:DA:2842:G:O2' | 31:DA:2843:G:H5' | 2.15 | 0.46 |
| 31:DA:2843:G:C2' | 31:DA:2844:G:O5' | 2.64 | 0.46 |
| 32:DB:2:C:C2 | 32:DB:3:C:C6 | 3.03 | 0.46 |
| 32:DB:60:C:C2 | 32:DB:61:G:C8 | 3.04 | 0.46 |
| 32:DB:66:A:C6 | 32:DB:109:C:C6 | 3.03 | 0.46 |
| 33:DD:11:PRO:C | 33:DD:13:ARG:N | 2.72 | 0.46 |
| 33:DD:25:THR:O | 33:DD:27:THR:CB | 2.54 | 0.46 |
| 34:DE:11:MET:HB3 | 34:DE:24:THR:HA | 1.97 | 0.46 |
| 34:DE:174:ASP:OD2 | 34:DE:175:VAL:N | 2.48 | 0.46 |
| 35:DF:144:LYS:C | 35:DF:146:ALA:H | 2.23 | 0.46 |
| 35:DF:164:ARG:HG2 | 35:DF:164:ARG:NH1 | 2.27 | 0.46 |
| 35:DF:196:LEU:H | 35:DF:196:LEU:HG | 1.61 | 0.46 |
| 36:DG:37:VAL:CG2 | 36:DG:99:MET:HG3 | 2.45 | 0.46 |
| 41:DP:31:ALA:C | 41:DP:33:ARG:H | 2.22 | 0.46 |
| 44:DS:17:ARG:HA | 44:DS:20:ARG:HG2 | 1.98 | 0.46 |
| 45:DT:108:ARG:O | 45:DT:111:ARG:HB2 | 2.15 | 0.46 |
| 46:DU:104:GLN:O | 46:DU:105:VAL:C | 2.59 | 0.46 |
| 47:DV:40:LEU:O | 47:DV:41:GLY:C | 2.58 | 0.46 |
| 48:DW:37:ARG:HG2 | 48:DW:38:TYR:CE2 | 2.51 | 0.46 |
| 49:DX:23:GLU:CG | 49:DX:24:GLY:H | 2.28 | 0.46 |
| 50:DY:26:LYS:O | 50:DY:27:VAL:C | 2.58 | 0.46 |
| 50:DY:88:LYS:HZ3 | 50:DY:93:GLY:HA3 | 1.80 | 0.46 |
| 1:AA:427:U:OP1 | 4:AD:13:ARG:NH2 | 2.49 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:542:G:C4 | 1:AA:543:C:C5 | 3.04 | 0.46 |
| 1:AA:542:G:C2 | 1:AA:543:C:C5 | 3.03 | 0.46 |
| 1:AA:685:G:N2 | 1:AA:686:U:N3 | 2.64 | 0.46 |
| 1:AA:719:C:H5 | 1:AA:720:C:N3 | 2.13 | 0.46 |
| 1:AA:857:C:H2' | 1:AA:858:G:O4' | 2.15 | 0.46 |
| 1:AA:1270:C:H2' | 1:AA:1271:G:O4' | 2.16 | 0.46 |
| 2:AB:67:THR:HG21 | 2:AB:155:LEU:CD2 | 2.45 | 0.46 |
| 2:AB:219:VAL:HA | 2:AB:222:ILE:CD1 | 2.46 | 0.46 |
| 4:AD:43:HIS:ND1 | 4:AD:46:LYS:NZ | 2.55 | 0.46 |
| 4:AD:127:THR:HB | 4:AD:132:ARG:HA | 1.97 | 0.46 |
| 5:AE:131:ILE:O | 5:AE:134:ALA:HB3 | 2.16 | 0.46 |
| 6:AF:11:ASN:O | 6:AF:14:LEU:HG | 2.15 | 0.46 |
| 13:AM:74:VAL:O | 13:AM:78:ILE:HG13 | 2.16 | 0.46 |
| 17:AQ:63:ARG:HG2 | 17:AQ:64:PRO:HD2 | 1.98 | 0.46 |
| 23:B1:9:GLY:O | 23:B1:10:LYS:CB | 2.63 | 0.46 |
| 24:B2:26:ARG:NE | 24:B2:29:LYS:HE2 | 2.28 | 0.46 |
| 29:B7:21:ARG:O | 29:B7:27:GLY:HA3 | 2.15 | 0.46 |
| 30:B8:8:LYS:CE | 31:BA:243:U:OP2 | 2.64 | 0.46 |
| 30:B8:62:LEU:C | 30:B8:64:TYR:N | 2.73 | 0.46 |
| 31:BA:34:C:O2' | 31:BA:35:G:OP1 | 2.32 | 0.46 |
| 31:BA:430:G:H5" | 31:BA:431:U:OP2 | 2.16 | 0.46 |
| 31:BA:535:C:H2' | 31:BA:536:A:H5' | 1.97 | 0.46 |
| 31:BA:639:U:O2' | 31:BA:640:C:H5' | 2.16 | 0.46 |
| 31:BA:705:A:C2 | 31:BA:706:A:C4 | 3.04 | 0.46 |
| 31:BA:1322:A:C5 | 31:BA:1323:U:C5 | 3.04 | 0.46 |
| 31:BA:1387:C:C2 | 31:BA:1388:G:C8 | 3.04 | 0.46 |
| 31:BA:1529:G:H21 | 31:BA:1530:C:C5' | 2.10 | 0.46 |
| 31:BA:1664:A:N6 | 31:BA:1665:A:N6 | 2.64 | 0.46 |
| 31:BA:1667:G:H4' | 31:BA:1668:A:OP1 | 2.14 | 0.46 |
| 31:BA:1694:C:H2' | 31:BA:1694:C:O2 | 2.15 | 0.46 |
| 31:BA:1824:G:OP1 | 33:BD:52:ARG:NH1 | 2.49 | 0.46 |
| 32:BB:48:A:OP1 | 44:BS:93:LYS:HB3 | 2.16 | 0.46 |
| 33:BD:59:LYS:HG3 | 33:BD:60:ARG:N | 2.31 | 0.46 |
| 33:BD:94:LEU:HB2 | 33:BD:104:TYR:CD2 | 2.51 | 0.46 |
| 33:BD:106:ILE:HD11 | 33:BD:196:VAL:HG13 | 1.98 | 0.46 |
| 35:BF:18:ARG:CZ | 35:BF:199:TRP:HZ3 | 2.28 | 0.46 |
| 36:BG:114:ILE:O | 36:BG:115:ARG:C | 2.57 | 0.46 |
| 37:BH:159:GLU:HA | 37:BH:159:GLU:OE1 | 2.16 | 0.46 |
| 41:BP:146:VAL:HG22 | 41:BP:147:LEU:H | 1.78 | 0.46 |
| 42:BQ:75:THR:HG21 | 42:BQ:85:LYS:HE3 | 1.98 | 0.46 |
| 42:BQ:140:ALA:CB | 51:BZ:99:TYR:HB2 | 2.43 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 44:BS:101:LEU:O | 44:BS:102:ALA:O | 2.33 | 0.46 |
| 50:BY:22:GLY:O | 50:BY:23:ARG:CG | 2.63 | 0.46 |
| 51:BZ:52:SER:OG | 51:BZ:53:ILE:N | 2.42 | 0.46 |
| 1:CA:134:A:H61 | 16:CP:25:ARG:NH1 | 2.14 | 0.46 |
| 1:CA:159:G:H21 | 1:CA:161:A:H8 | 1.62 | 0.46 |
| 1:CA:182:U:OP2 | 1:CA:183:G:OP2 | 2.33 | 0.46 |
| 1:CA:258:G:N3 | 1:CA:259:G:C8 | 2.84 | 0.46 |
| 1:CA:602:A:C2 | 1:CA:603:U:C2 | 3.04 | 0.46 |
| 1:CA:708:C:O2' | 1:CA:709:G:H5' | 2.15 | 0.46 |
| 1:CA:716:A:C2' | 1:CA:717:C:O5' | 2.64 | 0.46 |
| 1:CA:724:G:N3 | 1:CA:725:G:C8 | 2.84 | 0.46 |
| 1:CA:817:C:H4' | 1:CA:818:G:OP1 | 2.16 | 0.46 |
| 1:CA:926:G:C6 | 1:CA:1505:G:C6 | 3.04 | 0.46 |
| 1:CA:1442(B):A:C2 | 45:DT:118:ARG:NH2 | 2.83 | 0.46 |
| 2:CB:9:GLU:HA | 2:CB:12:GLU:OE1 | 2.15 | 0.46 |
| 2:CB:14:GLY:C | 2:CB:15:VAL:HG22 | 2.40 | 0.46 |
| 2:CB:116:GLU:HA | 2:CB:119:GLU:HB3 | 1.97 | 0.46 |
| 4:CD:194:LEU:HD22 | 4:CD:194:LEU:N | 2.30 | 0.46 |
| 6:CF:40:VAL:HA | 6:CF:62:TRP:O | 2.16 | 0.46 |
| 8:CH:8:ASP:O | 8:CH:11:THR:N | 2.49 | 0.46 |
| 8:CH:112:LEU:HD12 | 8:CH:114:THR:HG22 | 1.97 | 0.46 |
| 10:CJ:86:MET:O | 10:CJ:86:MET:HG3 | 2.15 | 0.46 |
| 12:CL:92:ASP:C | 12:CL:93:LEU:HD23 | 2.41 | 0.46 |
| 15:CO:32:LEU:O | 15:CO:33:THR:C | 2.58 | 0.46 |
| 19:CS:37:ARG:H | 19:CS:37:ARG:HG3 | 1.61 | 0.46 |
| 30:D8:35:GLN:HB3 | 30:D8:36:LYS:HZ3 | 1.80 | 0.46 |
| 31:DA:125:G:H4' | 31:DA:126:A:OP2 | 2.15 | 0.46 |
| 31:DA:773:U:H2' | 31:DA:774:A:H5' | 1.98 | 0.46 |
| 31:DA:921:G:C6 | 31:DA:922:U:C4 | 3.03 | 0.46 |
| 31:DA:992:C:H2' | 31:DA:993:G:O5' | 2.16 | 0.46 |
| 31:DA:1503:U:O2' | 31:DA:1504:C:H5' | 2.16 | 0.46 |
| 31:DA:2102:U:C6 | 31:DA:2187:G:O6 | 2.68 | 0.46 |
| 31:DA:2642:G:C2 | 31:DA:2773:C:C2 | 3.04 | 0.46 |
| 32:DB:7:G:H2' | 32:DB:8:U:H5" | 1.97 | 0.46 |
| 32:DB:75:G:C5' | 32:DB:75:G:C8 | 2.96 | 0.46 |
| 33:DD:45:ASN:CG | 33:DD:46:GLN:N | 2.70 | 0.46 |
| 33:DD:172:TYR:HD1 | 33:DD:185:VAL:C | 2.23 | 0.46 |
| 35:DF:18:ARG:CZ | 35:DF:199:TRP:HZ3 | 2.28 | 0.46 |
| 35:DF:28:ILE:HA | 35:DF:112:MET:HG2 | 1.96 | 0.46 |
| 35:DF:129:PHE:HE1 | 35:DF:142:TRP:CH2 | 2.33 | 0.46 |
| 35:DF:161:GLU:O | 35:DF:165:ARG:HG2 | 2.15 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 35:DF:185:ASP:OD1 | 35:DF:188:ARG:NH1 | 2.43 | 0.46 |
| 36:DG:141:PHE:HE2 | 36:DG:155:MET:HE1 | 1.81 | 0.46 |
| 38:DI:10:GLU:O | 38:DI:12:LEU:N | 2.49 | 0.46 |
| 40:DO:3:GLN:CB | 40:DO:4:PRO:HD2 | 2.45 | 0.46 |
| 40:DO:87:ILE:HG23 | 40:DO:88:ASN:N | 2.31 | 0.46 |
| 41:DP:5:ASP:CG | 41:DP:6:LEU:HD23 | 2.41 | 0.46 |
| 43:DR:107:ASP:OD2 | 43:DR:107:ASP:C | 2.58 | 0.46 |
| 45:DT:99:LEU:CD1 | 45:DT:99:LEU:O | 2.64 | 0.46 |
| 50:DY:65:ALA:HA | 50:DY:66:PRO:HD2 | 1.72 | 0.46 |
| 50:DY:75:ILE:CD1 | 50:DY:79:CYS:C | 2.85 | 0.46 |
| 1:AA:109:A:H2' | 1:AA:326:G:H21 | 1.80 | 0.46 |
| 1:AA:182:U:OP2 | 1:AA:183:G:OP2 | 2.33 | 0.46 |
| 1:AA:342:C:H2' | 1:AA:343:U:O4' | 2.15 | 0.46 |
| 1:AA:985:C:H2' | 1:AA:986:A:C8 | 2.51 | 0.46 |
| 1:AA:1446:U:O2 | 1:AA:1456:G:O6 | 2.33 | 0.46 |
| 2:AB:142:LEU:HD23 | 2:AB:142:LEU:C | 2.41 | 0.46 |
| 2:AB:204:ASN:HD22 | 2:AB:206:ASP:H | 1.64 | 0.46 |
| 6:AF:75:LEU:HD23 | 6:AF:79:LEU:HD21 | 1.98 | 0.46 |
| 8:AH:31:PHE:O | 8:AH:34:GLU:HB2 | 2.16 | 0.46 |
| 13:AM:14:ARG:CZ | 13:AM:42:ALA:HA | 2.45 | 0.46 |
| 13:AM:81:LEU:HB3 | 13:AM:89:GLY:CA | 2.39 | 0.46 |
| 16:AP:9:PHE:CE2 | 16:AP:18:ARG:HB2 | 2.51 | 0.46 |
| 17:AQ:68:ARG:O | 17:AQ:68:ARG:HG3 | 2.15 | 0.46 |
| 19:AS:58:VAL:O | 19:AS:58:VAL:HG23 | 2.15 | 0.46 |
| 23:B1:56:GLN:OE1 | 23:B1:56:GLN:HA | 2.08 | 0.46 |
| 30:B8:32:LEU:H | 30:B8:32:LEU:CD1 | 2.29 | 0.46 |
| 31:BA:27:G:O2' | 31:BA:28:A:OP2 | 2.32 | 0.46 |
| 31:BA:216:A:C4 | 31:BA:432:A:C2 | 3.04 | 0.46 |
| 31:BA:774:A:O2' | 31:BA:775:G:H5" | 2.16 | 0.46 |
| 31:BA:1349:A:C2' | 31:BA:1350:C:OP1 | 2.64 | 0.46 |
| 31:BA:1722:A:O2' | 31:BA:1739:U:C5' | 2.64 | 0.46 |
| 31:BA:1833:U:O2' | 31:BA:1969:A:N1 | 2.45 | 0.46 |
| 31:BA:1862:G:H2' | 31:BA:1863:G:O5' | 2.16 | 0.46 |
| 31:BA:1970:A:H5" | 31:BA:1971:A:OP1 | 2.16 | 0.46 |
| 31:BA:2026:C:H2' | 31:BA:2027:G:O5' | 2.16 | 0.46 |
| 31:BA:2100:G:H1 | 31:BA:2189:U:H3 | 1.63 | 0.46 |
| 31:BA:2183:C:H2' | 31:BA:2184:G:C8 | 2.50 | 0.46 |
| 31:BA:2263:C:O2' | 31:BA:2264:C:H5' | 2.16 | 0.46 |
| 31:BA:2598:A:O5' | 33:BD:236:GLY:HA3 | 2.16 | 0.46 |
| 33:BD:138:VAL:HG21 | 33:BD:167:GLY:HA2 | 1.98 | 0.46 |
| 33:BD:141:VAL:O | 33:BD:141:VAL:HG23 | 2.14 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 33:BD:172:TYR:HD1 | 33:BD:185:VAL:C | 2.23 | 0.46 |
| 35:BF:126:VAL:HG11 | 35:BF:142:TRP:HH2 | 1.80 | 0.46 |
| 37:BH:45:VAL:O | 37:BH:45:VAL:HG12 | 2.16 | 0.46 |
| 38:BI:31:LEU:N | 38:BI:31:LEU:CD1 | 2.78 | 0.46 |
| 46:BU:66:ASN:HA | 46:BU:76:TYR:HB2 | 1.98 | 0.46 |
| 48:BW:23:LEU:HD13 | 48:BW:23:LEU:HA | 1.73 | 0.46 |
| 50:BY:7:VAL:CB | 50:BY:8:LYS:NZ | 2.74 | 0.46 |
| 51:BZ:61:LEU:HB3 | 51:BZ:62:PRO:HD2 | 1.97 | 0.46 |
| 1:CA:265:G:H4' | 17:CQ:66:SER:HA | 1.97 | 0.46 |
| 1:CA:734:G:C6 | 1:CA:735:C:C4 | 3.04 | 0.46 |
| 1:CA:827:U:H2' | 1:CA:859:A:H61 | 1.80 | 0.46 |
| 1:CA:854:G:H3' | 1:CA:871:U:O4 | 2.15 | 0.46 |
| 1:CA:1030(D):A:H62 | 1:CA:1031:G:H21 | 1.62 | 0.46 |
| 1:CA:1128:C:O2 | 1:CA:1130:A:N7 | 2.48 | 0.46 |
| 1:CA:1226:C:OP1 | 13:CM:91:ARG:NH1 | 2.49 | 0.46 |
| 1:CA:1242:C:H6 | 1:CA:1242:C:O5' | 1.99 | 0.46 |
| 1:CA:1368:G:OP2 | 9:CI:112:LYS:HD2 | 2.16 | 0.46 |
| 2:CB:61:LEU:HD23 | 2:CB:68:ILE:HD11 | 1.94 | 0.46 |
| 2:CB:76:GLN:O | 2:CB:208:ILE:HG12 | 2.15 | 0.46 |
| 2:CB:118:LEU:HD11 | 2:CB:141:GLU:HG2 | 1.97 | 0.46 |
| 2:CB:167:PRO:HG2 | 2:CB:192:SER:HB3 | 1.98 | 0.46 |
| 5:CE:101:ILE:CD1 | 5:CE:119:LEU:HD23 | 2.46 | 0.46 |
| 10:CJ:29:ARG:HH22 | 10:CJ:84:GLN:HG2 | 1.81 | 0.46 |
| 12:CL:56:ALA:O | 12:CL:68:ALA:N | 2.49 | 0.46 |
| 18:CR:22:VAL:HG12 | 18:CR:22:VAL:O | 2.16 | 0.46 |
| 20:CT:94:ALA:O | 20:CT:95:ALA:HB3 | 2.15 | 0.46 |
| 23:D1:10:LYS:HB2 | 23:D1:14:VAL:CA | 2.45 | 0.46 |
| 23:D1:82:LEU:HG | 23:D1:83:GLU:N | 2.31 | 0.46 |
| 30:D8:4:MET:HE2 | 30:D8:4:MET:HB2 | 1.50 | 0.46 |
| 31:DA:27:G:H22 | 31:DA:512:G:H1' | 1.79 | 0.46 |
| 31:DA:128:C:O2' | 31:DA:129:C:P | 2.74 | 0.46 |
| 31:DA:190:A:P | 31:DA:205:G:H22 | 2.39 | 0.46 |
| 31:DA:271(Q):G:O2' | 31:DA:271(R):G:OP2 | 2.27 | 0.46 |
| 31:DA:572:A:H5" | 31:DA:573:G:OP2 | 2.16 | 0.46 |
| 31:DA:681:G:H2' | 31:DA:682:G:O4' | 2.16 | 0.46 |
| 31:DA:770:G:N3 | 31:DA:1354:A:H2 | 2.13 | 0.46 |
| 31:DA:828:U:O2 | 31:DA:828:U:C2' | 2.63 | 0.46 |
| 31:DA:1012:U:O4 | 39:DN:25:ARG:HA | 2.16 | 0.46 |
| 31:DA:1028:A:N6 | 31:DA:1125:G:H2' | 2.31 | 0.46 |
| 31:DA:1359:A:N7 | 31:DA:1372:U:O4 | 2.49 | 0.46 |
| 34:DE:11:MET:HB2 | 34:DE:23:VAL:O | 2.15 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 37:DH:153:LYS:CB | 37:DH:154:PRO:CD | 2.94 | 0.46 |
| 42:DQ:21:THR:O | 51:DZ:78:LYS:HE2 | 2.16 | 0.46 |
| 47:DV:89:GLN:NE2 | 47:DV:91:TYR:HD1 | 2.13 | 0.46 |
| 49:DX:37:THR:HG22 | 49:DX:37:THR:O | 2.15 | 0.46 |
| 1:AA:406:G:H5' | 4:AD:5:ILE:HD13 | 1.97 | 0.46 |
| 1:AA:409:G:H5' | 4:AD:24:GLU:OE1 | 2.16 | 0.46 |
| 1:AA:868:C:H2' | 1:AA:869:G:O4' | 2.16 | 0.46 |
| 1:AA:1103:C:H5" | 2:AB:98:LEU:HD13 | 1.98 | 0.46 |
| 1:AA:1160:G:C2 | 1:AA:1161:C:C6 | 3.03 | 0.46 |
| 1:AA:1163:C:C2 | 1:AA:1174:G:N2 | 2.84 | 0.46 |
| 1:AA:1350:A:N6 | 1:AA:1373:G:N2 | 2.64 | 0.46 |
| 2:AB:79:ASP:C | 2:AB:81:VAL:H | 2.24 | 0.46 |
| 2:AB:97:TRP:HH2 | 2:AB:176:GLU:CD | 2.24 | 0.46 |
| 4:AD:14:ARG:HA | 4:AD:39:PRO:CB | 2.46 | 0.46 |
| 4:AD:141:ARG:HB3 | 4:AD:142:PRO:HD3 | 1.98 | 0.46 |
| 5:AE:10:MET:HG3 | 5:AE:13:ILE:CD1 | 2.46 | 0.46 |
| 5:AE:100:VAL:O | 5:AE:107:ARG:NH2 | 2.41 | 0.46 |
| 5:AE:139:LEU:C | 5:AE:141:GLN:H | 2.22 | 0.46 |
| 10:AJ:86:MET:O | 10:AJ:86:MET:HG3 | 2.16 | 0.46 |
| 13:AM:17:VAL:O | 13:AM:20:THR:HB | 2.16 | 0.46 |
| 18:AR:86:VAL:O | 18:AR:87:ARG:HB3 | 2.15 | 0.46 |
| 22:B0:43:THR:O | 22:B0:45:PHE:N | 2.48 | 0.46 |
| 24:B2:15:LYS:O | 24:B2:16:LEU:HB2 | 2.13 | 0.46 |
| 24:B2:52:ASP:H | 24:B2:55:ARG:HB2 | 1.81 | 0.46 |
| 25:B3:54:VAL:CG1 | 25:B3:55:ARG:H | 2.29 | 0.46 |
| 28:B6:27:LYS:HD2 | 31:BA:2285:C:C5 | 2.51 | 0.46 |
| 29:B7:5:TRP:CZ3 | 31:BA:464:U:H4' | 2.51 | 0.46 |
| 30:B8:58:ILE:HG22 | 41:BP:49:ARG:CD | 2.45 | 0.46 |
| 31:BA:272(J):C:O2' | 31:BA:274:G:OP1 | 2.28 | 0.46 |
| 31:BA:958:U:H5" | 42:BQ:14:ARG:HD3 | 1.98 | 0.46 |
| 31:BA:1176:G:C1' | 31:BA:1177:A:OP1 | 2.63 | 0.46 |
| 31:BA:1198:U:H2' | 31:BA:1199:U:C6 | 2.51 | 0.46 |
| 31:BA:1380:G:N2 | 31:BA:1570:A:C2 | 2.84 | 0.46 |
| 31:BA:1528:A:O2' | 31:BA:1528(A):A:P | 2.73 | 0.46 |
| 31:BA:1721:G:C6 | 31:BA:1739:U:H5' | 2.50 | 0.46 |
| 31:BA:1856:G:C2' | 31:BA:1857:G:H5' | 2.46 | 0.46 |
| 31:BA:2016:U:H2' | 31:BA:2017:U:H6 | 1.78 | 0.46 |
| 31:BA:2864:G:H2' | 31:BA:2865:U:O4' | 2.15 | 0.46 |
| 32:BB:7:G:O5' | 44:BS:29:PHE:HE1 | 1.99 | 0.46 |
| 33:BD:62:TYR:CE1 | 33:BD:64:ILE:HA | 2.51 | 0.46 |
| 34:BE:23:VAL:HA | 34:BE:186:GLY:N | 2.29 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 34:BE:24:THR:CG2 | 34:BE:184:VAL:HG23 | 2.43 | 0.46 |
| 41:BP:92:GLU:HA | 41:BP:123:LEU:HD22 | 1.98 | 0.46 |
| 44:BS:34:HIS:HD2 | 44:BS:34:HIS:N | 2.14 | 0.46 |
| 44:BS:56:LEU:O | 44:BS:57:LYS:CB | 2.64 | 0.46 |
| 46:BU:74:LEU:N | 46:BU:74:LEU:CD1 | 2.78 | 0.46 |
| 47:BV:4:ILE:O | 47:BV:39:LEU:CB | 2.64 | 0.46 |
| 47:BV:66:ARG:HD2 | 47:BV:67:GLY:CA | 2.45 | 0.46 |
| 47:BV:83:ARG:HH11 | 47:BV:83:ARG:HG2 | 1.74 | 0.46 |
| 49:BX:36:LYS:HD2 | 49:BX:36:LYS:C | 2.41 | 0.46 |
| 50:BY:81:LYS:NZ | 50:BY:97:ARG:HG3 | 2.30 | 0.46 |
| 50:BY:84:ARG:C | 50:BY:85:VAL:CG2 | 2.89 | 0.46 |
| 51:BZ:48:PHE:O | 51:BZ:49:ARG:C | 2.58 | 0.46 |
| 1:CA:61:G:OP1 | 20:CT:10:LEU:HD11 | 2.16 | 0.46 |
| 1:CA:177:C:OP1 | 20:CT:65:LYS:HD3 | 2.16 | 0.46 |
| 1:CA:427:U:OP1 | 4:CD:13:ARG:NH2 | 2.49 | 0.46 |
| 1:CA:1296:C:H5' | 1:CA:1297:C:OP2 | 2.16 | 0.46 |
| 1:CA:1381:U:C2' | 1:CA:1382:C:H5' | 2.46 | 0.46 |
| 2:CB:51:LEU:HD21 | 2:CB:214:ILE:HD12 | 1.96 | 0.46 |
| 3:CC:102:ASN:O | 3:CC:103:VAL:HG23 | 2.16 | 0.46 |
| 5:CE:78:HIS:HE1 | 5:CE:142:LEU:HD23 | 1.81 | 0.46 |
| 5:CE:78:HIS:CE1 | 5:CE:142:LEU:HD23 | 2.51 | 0.46 |
| 6:CF:81:ILE:O | 6:CF:82:ARG:C | 2.59 | 0.46 |
| 12:CL:41:ARG:HG2 | 12:CL:42:THR:N | 2.30 | 0.46 |
| 22:D0:46:LYS:HB3 | 22:D0:47:PRO:HD2 | 1.98 | 0.46 |
| 22:D0:72:ARG:O | 22:D0:73:GLY:C | 2.58 | 0.46 |
| 27:D5:20:ARG:HG2 | 27:D5:23:HIS:CD2 | 2.51 | 0.46 |
| 30:D8:32:LEU:CG | 30:D8:35:GLN:H | 2.29 | 0.46 |
| 31:DA:128:C:H4' | 31:DA:129:C:OP1 | 2.16 | 0.46 |
| 31:DA:301:G:C4 | 31:DA:302:C:C5 | 3.04 | 0.46 |
| 31:DA:543:C:H5 | 31:DA:547:A:H62 | 1.62 | 0.46 |
| 31:DA:769:G:O2' | 31:DA:770:G:H5' | 2.15 | 0.46 |
| 31:DA:869:G:C5 | 31:DA:870:A:N7 | 2.84 | 0.46 |
| 31:DA:924:C:C2 | 31:DA:925:C:C5 | 3.04 | 0.46 |
| 31:DA:995:C:O2 | 46:DU:57:PHE:CG | 2.69 | 0.46 |
| 31:DA:1149:G:H2' | 31:DA:1150:C:C6 | 2.51 | 0.46 |
| 31:DA:1497:U:C2' | 31:DA:1497:U:O2 | 2.64 | 0.46 |
| 31:DA:1701:A:C2' | 31:DA:1702:G:H5' | 2.46 | 0.46 |
| 31:DA:2552:U:C2 | 31:DA:2554:U:H5' | 2.51 | 0.46 |
| 32:DB:15:A:O2' | 32:DB:110:G:C8 | 2.55 | 0.46 |
| 33:DD:62:TYR:HE1 | 33:DD:64:ILE:HA | 1.80 | 0.46 |
| 37:DH:157:TYR:O | 37:DH:158:HIS:HB2 | 2.15 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 41:DP:146:VAL:HG22 | 41:DP:147:LEU:H | 1.81 | 0.46 |
| 42:DQ:14:ARG:HG2 | 42:DQ:41:TRP:HH2 | 1.80 | 0.46 |
| 44:DS:24:LEU:HD13 | 44:DS:24:LEU:HA | 1.74 | 0.46 |
| 51:DZ:112:ARG:C | 51:DZ:114:GLY:H | 2.23 | 0.46 |
| 1:AA:15:G:C4 | 1:AA:16:A:C8 | 3.04 | 0.46 |
| 1:AA:124:G:C6 | 1:AA:125:U:C4 | 3.03 | 0.46 |
| 1:AA:199:G:O2' | 1:AA:200:G:H5' | 2.15 | 0.46 |
| 1:AA:411:A:C8 | 1:AA:413:G:C8 | 3.03 | 0.46 |
| 1:AA:1081:G:N2 | 1:AA:1082:G:H1' | 2.31 | 0.46 |
| 1:AA:1202:G:C6 | 14:AN:42:ILE:HG21 | 2.50 | 0.46 |
| 1:AA:1322:C:OP1 | 1:AA:1322:C:H6 | 1.99 | 0.46 |
| 4:AD:74:GLN:O | 4:AD:78:LEU:HG | 2.15 | 0.46 |
| 5:AE:101:ILE:HA | 5:AE:107:ARG:NH2 | 2.31 | 0.46 |
| 8:AH:9:MET:O | 8:AH:12:ARG:HB2 | 2.16 | 0.46 |
| 10:AJ:82:ILE:O | 10:AJ:86:MET:HB3 | 2.16 | 0.46 |
| 12:AL:10:LEU:HB3 | 17:AQ:32:TYR:CE1 | 2.51 | 0.46 |
| 12:AL:117:ARG:NH2 | 12:AL:124:LYS:HB2 | 2.31 | 0.46 |
| 16:AP:9:PHE:CD2 | 16:AP:18:ARG:HB2 | 2.51 | 0.46 |
| 22:B0:42:GLY:HA2 | 22:B0:57:PHE:CD2 | 2.50 | 0.46 |
| 24:B2:55:ARG:HG2 | 24:B2:55:ARG:O | 2.16 | 0.46 |
| 31:BA:203:C:C3' | 31:BA:204:A:H5" | 2.46 | 0.46 |
| 31:BA:449:A:OP1 | 35:BF:84:VAL:O | 2.34 | 0.46 |
| 31:BA:511:U:O4 | 31:BA:512:G:N1 | 2.49 | 0.46 |
| 31:BA:719:C:H2' | 31:BA:720:C:C6 | 2.51 | 0.46 |
| 31:BA:847:U:OP2 | 31:BA:928:G:O6 | 2.34 | 0.46 |
| 31:BA:892:G:H2' | 31:BA:893:C:O4' | 2.15 | 0.46 |
| 31:BA:893:C:C2' | 31:BA:894:C:O5' | 2.64 | 0.46 |
| 31:BA:1497:U:C5' | 31:BA:1498:C:C5 | 2.94 | 0.46 |
| 31:BA:1783:A:N1 | 31:BA:2587:A:C4 | 2.84 | 0.46 |
| 31:BA:2615:U:H2' | 31:BA:2616:C:C6 | 2.51 | 0.46 |
| 31:BA:2788:C:O2' | 31:BA:2809:A:N3 | 2.41 | 0.46 |
| 32:BB:70:C:C4 | 32:BB:71:C:C5 | 3.04 | 0.46 |
| 34:BE:66:HIS:O | 34:BE:66:HIS:CD2 | 2.69 | 0.46 |
| 35:BF:39:TRP:O | 35:BF:43:LYS:HG2 | 2.15 | 0.46 |
| 38:BI:83:ALA:CB | 38:BI:88:ILE:HA | 2.46 | 0.46 |
| 38:BI:145:VAL:HG12 | 38:BI:146:ALA:N | 2.31 | 0.46 |
| 43:BR:14:SER:OG | 43:BR:15:SER:N | 2.49 | 0.46 |
| 44:BS:34:HIS:ND1 | 44:BS:54:LEU:HB2 | 2.31 | 0.46 |
| 45:BT:26:ASP:HB3 | 45:BT:89:VAL:O | 2.15 | 0.46 |
| 50:BY:96:ILE:CB | 50:BY:99:CYS:HB3 | 2.46 | 0.46 |
| 1:CA:20:U:O2' | 1:CA:21:G:H5' | 2.16 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:327:A:C6 | 1:CA:329:A:C5 | 3.04 | 0.46 |
| 1:CA:409:G:H5' | 4:CD:24:GLU:OE1 | 2.16 | 0.46 |
| 1:CA:502:G:C2 | 1:CA:503:C:C2 | 3.04 | 0.46 |
| 1:CA:533:A:OP1 | 1:CA:533:A:H3' | 2.15 | 0.46 |
| 1:CA:681:C:C2 | 1:CA:710:G:N2 | 2.84 | 0.46 |
| 1:CA:950:U:H3' | 13:CM:102:ARG:HH12 | 1.80 | 0.46 |
| 1:CA:1429:C:H2' | 1:CA:1430:C:C6 | 2.51 | 0.46 |
| 2:CB:30:ARG:HG3 | 2:CB:31:TYR:CE2 | 2.51 | 0.46 |
| 2:CB:204:ASN:ND2 | 2:CB:205:ASP:N | 2.64 | 0.46 |
| 4:CD:170:VAL:HG22 | 4:CD:171:GLY:N | 2.30 | 0.46 |
| 5:CE:122:GLU:OE1 | 5:CE:131:ILE:HG13 | 2.16 | 0.46 |
| 11:CK:69:ALA:O | 11:CK:73:MET:HG2 | 2.16 | 0.46 |
| 11:CK:111:ASP:CA | 18:CR:84:LYS:HE2 | 2.44 | 0.46 |
| 16:CP:20:VAL:CG2 | 16:CP:32:TYR:HB2 | 2.46 | 0.46 |
| 17:CQ:59:ILE:HG22 | 17:CQ:71:PHE:CD1 | 2.50 | 0.46 |
| 20:CT:89:ARG:HH22 | 20:CT:104:LEU:HD21 | 1.78 | 0.46 |
| 23:D1:67:ILE:O | 23:D1:70:VAL:HB | 2.15 | 0.46 |
| 30:D8:34:TRP:HZ3 | 30:D8:41:ILE:CG2 | 2.29 | 0.46 |
| 30:D8:61:LEU:HD23 | 30:D8:61:LEU:HA | 1.61 | 0.46 |
| 31:DA:58:G:H2' | 31:DA:59:U:H6 | 1.81 | 0.46 |
| 31:DA:478:A:C6 | 31:DA:480:A:C6 | 3.03 | 0.46 |
| 31:DA:483:A:C8 | 31:DA:484:C:C5 | 3.03 | 0.46 |
| 31:DA:796:C:C2 | 31:DA:797:C:C5 | 3.04 | 0.46 |
| 31:DA:1495:A:C2 | 31:DA:1496:A:C2 | 3.03 | 0.46 |
| 31:DA:1711:C:O2' | 31:DA:1712:C:H5' | 2.16 | 0.46 |
| 31:DA:1771:C:O2' | 31:DA:1786:A:H8 | 1.76 | 0.46 |
| 31:DA:2100:G:H1 | 31:DA:2189:U:H3 | 1.64 | 0.46 |
| 31:DA:2290:G:O2' | 31:DA:2381:C:H1' | 2.16 | 0.46 |
| 31:DA:2631:G:N3 | 31:DA:2810:A:C2 | 2.83 | 0.46 |
| 31:DA:2782:G:H8 | 31:DA:2782:G:O5' | 1.98 | 0.46 |
| 31:DA:2843:G:H2' | 31:DA:2844:G:O5' | 2.16 | 0.46 |
| 32:DB:40:U:H1' | 32:DB:45:A:N6 | 2.31 | 0.46 |
| 33:DD:142:VAL:HG21 | 33:DD:191:ALA:CB | 2.46 | 0.46 |
| 38:DI:5:LEU:HD11 | 38:DI:19:VAL:HG11 | 1.98 | 0.46 |
| 38:DI:92:VAL:O | 38:DI:92:VAL:HG13 | 2.16 | 0.46 |
| 39:DN:78:TYR:CD1 | 39:DN:79:PRO:CB | 2.99 | 0.46 |
| 41:DP:91:PHE:HE2 | 41:DP:95:VAL:CG1 | 2.25 | 0.46 |
| 41:DP:147:LEU:HB2 | 41:DP:148:LEU:H | 1.48 | 0.46 |
| 44:DS:14:VAL:HG13 | 44:DS:15:ARG:N | 2.31 | 0.46 |
| 1:AA:295:C:H2' | 1:AA:296:U:C6 | 2.51 | 0.45 |
| 1:AA:509:A:O2' | 1:AA:510:A:C5' | 2.64 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:1308:U:H5" | 13:AM:98:VAL:N | 2.31 | 0.45 |
| 1:AA:1312:G:N2 | 1:AA:1326:C:C2 | 2.84 | 0.45 |
| 2:AB:167:PRO:HG2 | 2:AB:192:SER:HB3 | 1.97 | 0.45 |
| 7:AG:106:GLN:O | 7:AG:110:GLN:HG3 | 2.16 | 0.45 |
| 15:AO:24:SER:O | 15:AO:28:GLN:HG3 | 2.15 | 0.45 |
| 20:AT:92:LEU:C | 20:AT:94:ALA:H | 2.24 | 0.45 |
| 25:B3:8:LEU:HD12 | 25:B3:31:LEU:HA | 1.98 | 0.45 |
| 25:B3:18:ASP:HB2 | 25:B3:49:LYS:CE | 2.46 | 0.45 |
| 31:BA:995:C:N3 | 46:BU:57:PHE:CZ | 2.85 | 0.45 |
| 31:BA:1049:C:O2 | 31:BA:1050:A:N7 | 2.49 | 0.45 |
| 31:BA:1158:C:C2' | 31:BA:1159:U:H5' | 2.46 | 0.45 |
| 31:BA:1599:C:OP1 | 49:BX:36:LYS:HA | 2.15 | 0.45 |
| 31:BA:2409:G:H2' | 31:BA:2410:G:O4' | 2.15 | 0.45 |
| 31:BA:2632:A:O2' | 34:BE:61:ARG:NH2 | 2.49 | 0.45 |
| 31:BA:2658:C:C5' | 31:BA:2659:G:OP2 | 2.64 | 0.45 |
| 31:BA:2862:G:C5 | 31:BA:2863:C:C5 | 3.04 | 0.45 |
| 34:BE:56:PRO:O | 34:BE:58:ARG:N | 2.49 | 0.45 |
| 38:BI:10:GLU:O | 38:BI:12:LEU:N | 2.49 | 0.45 |
| 43:BR:52:ILE:O | 43:BR:53:HIS:C | 2.57 | 0.45 |
| 45:BT:57:PHE:O | 45:BT:58:ASN:ND2 | 2.49 | 0.45 |
| 48:BW:37:ARG:HG2 | 48:BW:38:TYR:CE2 | 2.51 | 0.45 |
| 49:BX:60:ARG:HG2 | 49:BX:74:PRO:HD2 | 1.94 | 0.45 |
| 49:BX:70:LEU:O | 49:BX:71:GLY:C | 2.59 | 0.45 |
| 50:BY:27:VAL:O | 50:BY:29:GLU:OE1 | 2.34 | 0.45 |
| 51:BZ:19:ARG:HG2 | 51:BZ:19:ARG:NH1 | 2.31 | 0.45 |
| 1:CA:272:C:H2' | 1:CA:273:A:C8 | 2.51 | 0.45 |
| 1:CA:283:C:H2' | 1:CA:284:G:O4' | 2.16 | 0.45 |
| 1:CA:320:C:H2' | 1:CA:321:A:O4' | 2.16 | 0.45 |
| 1:CA:407:G:H4' | 4:CD:115:ARG:O | 2.16 | 0.45 |
| 1:CA:1079:G:O3' | 5:CE:14:ARG:NH2 | 2.49 | 0.45 |
| 2:CB:25:ASN:OD1 | 2:CB:25:ASN:C | 2.58 | 0.45 |
| 4:CD:108:LEU:HD12 | 4:CD:174:LEU:HD13 | 1.98 | 0.45 |
| 7:CG:152:ALA:O | 7:CG:155:ARG:HG3 | 2.15 | 0.45 |
| 8:CH:20:TYR:CD1 | 8:CH:65:TYR:CD2 | 2.89 | 0.45 |
| 13:CM:13:LYS:O | 13:CM:45:VAL:HG23 | 2.16 | 0.45 |
| 23:D1:94:LEU:HD22 | 23:D1:95:LEU:N | 2.31 | 0.45 |
| 28:D6:18:ARG:HG3 | 28:D6:19:ARG:HG2 | 1.98 | 0.45 |
| 30:D8:35:GLN:HG2 | 31:DA:2420:C:OP1 | 2.16 | 0.45 |
| 31:DA:271(K):U:H2' | 31:DA:271(M):G:N2 | 2.30 | 0.45 |
| 31:DA:869:G:C6 | 31:DA:870:A:N7 | 2.84 | 0.45 |
| 31:DA:874:G:N2 | 31:DA:904:C:C2 | 2.84 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1022:G:C5 | 31:DA:1140:C:C4 | 3.05 | 0.45 |
| 31:DA:1464:C:O2' | 31:DA:1528:A:H1' | 2.15 | 0.45 |
| 31:DA:1550:C:O2' | 31:DA:1551:C:H5' | 2.16 | 0.45 |
| 31:DA:1907:G:O2' | 31:DA:1908:C:H5' | 2.17 | 0.45 |
| 31:DA:2187:G:C5 | 31:DA:2188:C:C2 | 3.04 | 0.45 |
| 31:DA:2478:A:H2' | 31:DA:2479:G:H5' | 1.98 | 0.45 |
| 31:DA:2664:G:C2' | 31:DA:2665:A:O5' | 2.65 | 0.45 |
| 31:DA:2733:A:C3' | 31:DA:2734:A:H5" | 2.46 | 0.45 |
| 31:DA:2864:G:H2' | 31:DA:2865:U:O4' | 2.16 | 0.45 |
| 33:DD:33:LEU:HB2 | 33:DD:34:VAL:H | 1.67 | 0.45 |
| 33:DD:75:ILE:CG2 | 33:DD:99:ASP:HB2 | 2.43 | 0.45 |
| 33:DD:85:ASP:OD2 | 33:DD:88:ARG:HD2 | 2.16 | 0.45 |
| 33:DD:228:PRO:HD3 | 33:DD:235:GLY:CA | 2.42 | 0.45 |
| 34:DE:51:PHE:CD1 | 34:DE:52:LEU:N | 2.84 | 0.45 |
| 35:DF:160:ASN:HD21 | 35:DF:162:LEU:HB2 | 1.81 | 0.45 |
| 35:DF:178:PRO:C | 35:DF:180:GLY:H | 2.25 | 0.45 |
| 38:DI:88:ILE:HG12 | 38:DI:122:GLU:N | 2.30 | 0.45 |
| 40:DO:35:VAL:HA | 40:DO:62:VAL:CG1 | 2.42 | 0.45 |
| 41:DP:48:PRO:HG2 | 41:DP:49:ARG:H | 1.81 | 0.45 |
| 41:DP:100:LEU:HD12 | 41:DP:100:LEU:HA | 1.67 | 0.45 |
| 43:DR:113:LEU:HA | 43:DR:113:LEU:HD12 | 1.67 | 0.45 |
| 46:DU:28:ARG:HD3 | 46:DU:38:THR:OG1 | 2.16 | 0.45 |
| 47:DV:17:GLY:O | 47:DV:18:LEU:HB3 | 2.16 | 0.45 |
| 50:DY:10:GLY:C | 50:DY:27:VAL:HG22 | 2.40 | 0.45 |
| 1:AA:60:A:C4' | 1:AA:61:G:O5' | 2.64 | 0.45 |
| 1:AA:114:U:O2' | 1:AA:115:G:H5' | 2.16 | 0.45 |
| 1:AA:132:C:H2' | 1:AA:133:U:H6 | 1.81 | 0.45 |
| 1:AA:166:G:O2' | 1:AA:167:G:H5' | 2.16 | 0.45 |
| 1:AA:586:C:H2' | 1:AA:587:G:H5' | 1.98 | 0.45 |
| 1:AA:716:A:C2' | 1:AA:717:C:O5' | 2.64 | 0.45 |
| 1:AA:1350:A:H2' | 1:AA:1351:U:C6 | 2.52 | 0.45 |
| 1:AA:1442(A):G:C3' | 1:AA:1442(B):A:C5' | 2.85 | 0.45 |
| 2:AB:54:THR:HG21 | 2:AB:201:ILE:HD11 | 1.98 | 0.45 |
| 5:AE:20:GLN:O | 5:AE:21:ALA:C | 2.58 | 0.45 |
| 6:AF:46:ARG:NH1 | 18:AR:37:VAL:HG21 | 2.27 | 0.45 |
| 6:AF:81:ILE:O | 6:AF:82:ARG:C | 2.60 | 0.45 |
| 8:AH:51:VAL:O | 8:AH:52:ASP:HB2 | 2.16 | 0.45 |
| 11:AK:52:GLY:H | 11:AK:55:LYS:HG3 | 1.81 | 0.45 |
| 11:AK:111:ASP:HA | 18:AR:84:LYS:CE | 2.44 | 0.45 |
| 16:AP:21:VAL:O | 16:AP:33:ILE:HB | 2.16 | 0.45 |
| 16:AP:49:LEU:HG | 16:AP:50:LYS:N | 2.31 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 18:AR:74:ARG:HG3 | 18:AR:79:LEU:HB3 | 1.98 | 0.45 |
| 19:AS:10:PHE:HZ | 19:AS:70:LYS:HZ3 | 1.64 | 0.45 |
| 27:B5:28:PRO:HD2 | 48:BW:35:ILE:HG23 | 1.98 | 0.45 |
| 29:B7:39:ARG:HD3 | 31:BA:458:G:O2' | 2.17 | 0.45 |
| 29:B7:46:VAL:HB | 29:B7:48:LYS:NZ | 2.31 | 0.45 |
| 31:BA:128:C:H4' | 31:BA:129:C:OP1 | 2.17 | 0.45 |
| 31:BA:250:G:C6 | 31:BA:251:A:C6 | 3.05 | 0.45 |
| 31:BA:536:A:H2' | 31:BA:537:C:H6 | 1.78 | 0.45 |
| 31:BA:675:A:C8 | 31:BA:804:A:N1 | 2.84 | 0.45 |
| 31:BA:1178:C:C5 | 31:BA:1179:C:H5 | 2.34 | 0.45 |
| 31:BA:1213:A:O2' | 31:BA:1214:A:H5' | 2.16 | 0.45 |
| 31:BA:1475:G:H5' | 31:BA:1476:C:OP2 | 2.16 | 0.45 |
| 31:BA:1862:G:C2' | 31:BA:1863:G:O5' | 2.64 | 0.45 |
| 32:BB:31:C:O2 | 32:BB:52:A:H2 | 1.99 | 0.45 |
| 33:BD:11:PRO:C | 33:BD:13:ARG:N | 2.74 | 0.45 |
| 36:BG:45:GLU:HG2 | 36:BG:47:LYS:H | 1.80 | 0.45 |
| 37:BH:158:HIS:CE1 | 37:BH:169:VAL:O | 2.69 | 0.45 |
| 39:BN:78:TYR:CD1 | 39:BN:79:PRO:CG | 3.00 | 0.45 |
| 39:BN:108:PRO:O | 39:BN:113:GLY:HA3 | 2.17 | 0.45 |
| 41:BP:115:LEU:HA | 41:BP:134:ALA:CB | 2.46 | 0.45 |
| 42:BQ:141:GLN:O | 51:BZ:70:LEU:HD13 | 2.15 | 0.45 |
| 44:BS:29:PHE:H | 44:BS:89:ARG:CG | 2.29 | 0.45 |
| 44:BS:95:HIS:CG | 44:BS:96:GLY:N | 2.76 | 0.45 |
| 45:BT:80:SER:CB | 45:BT:81:PRO:HD3 | 2.47 | 0.45 |
| 45:BT:82:LEU:N | 45:BT:82:LEU:CD1 | 2.79 | 0.45 |
| 48:BW:31:GLU:O | 48:BW:35:ILE:HG13 | 2.17 | 0.45 |
| 49:BX:23:GLU:CG | 49:BX:24:GLY:H | 2.29 | 0.45 |
| 1:CA:90:U:H6 | 1:CA:90:U:H3' | 1.82 | 0.45 |
| 1:CA:297:G:H4' | 1:CA:557:G:H4' | 1.98 | 0.45 |
| 1:CA:375:U:C4 | 1:CA:376:G:N7 | 2.84 | 0.45 |
| 1:CA:505:G:C5 | 1:CA:535:A:C2 | 3.04 | 0.45 |
| 1:CA:594:G:O2' | 1:CA:595:G:H5' | 2.16 | 0.45 |
| 1:CA:883:C:C2' | 1:CA:884:U:H5' | 2.46 | 0.45 |
| 1:CA:951:G:OP2 | 13:CM:102:ARG:NH2 | 2.50 | 0.45 |
| 4:CD:102:ASP:HB3 | 4:CD:136:PRO:HA | 1.98 | 0.45 |
| 25:D3:54:VAL:CG1 | 25:D3:55:ARG:H | 2.27 | 0.45 |
| 27:D5:19:ARG:NH2 | 31:DA:1264:G:OP1 | 2.44 | 0.45 |
| 30:D8:4:MET:HE1 | 31:DA:593:G:O4' | 2.16 | 0.45 |
| 31:DA:2086:U:H2' | 31:DA:2087:G:C8 | 2.51 | 0.45 |
| 31:DA:2305:A:C5' | 36:DG:134:GLY:HA3 | 2.30 | 0.45 |
| 31:DA:2680:C:OP2 | 34:DE:111:ARG:NH2 | 2.49 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 33:DD:6:PHE:N | 33:DD:6:PHE:CD1 | 2.84 | 0.45 |
| 33:DD:53:PHE:HD1 | 33:DD:219:PRO:O | 1.98 | 0.45 |
| 33:DD:143:HIS:HD2 | 33:DD:144:ALA:CB | 2.28 | 0.45 |
| 34:DE:23:VAL:HA | 34:DE:184:VAL:O | 2.16 | 0.45 |
| 34:DE:51:PHE:H | 34:DE:74:PRO:CB | 2.29 | 0.45 |
| 35:DF:160:ASN:CG | 35:DF:163:VAL:HG23 | 2.41 | 0.45 |
| 36:DG:45:GLU:HB2 | 36:DG:47:LYS:CD | 2.47 | 0.45 |
| 37:DH:111:HIS:CG | 37:DH:112:PRO:HD2 | 2.51 | 0.45 |
| 37:DH:158:HIS:HE2 | 37:DH:169:VAL:C | 2.24 | 0.45 |
| 41:DP:115:LEU:HA | 41:DP:134:ALA:CB | 2.45 | 0.45 |
| 41:DP:146:VAL:HG13 | 41:DP:147:LEU:N | 2.29 | 0.45 |
| 42:DQ:26:TYR:CE1 | 42:DQ:28:ALA:HB2 | 2.51 | 0.45 |
| 42:DQ:75:THR:HG21 | 42:DQ:85:LYS:HE3 | 1.98 | 0.45 |
| 43:DR:71:GLN:NE2 | 43:DR:71:GLN:HA | 2.31 | 0.45 |
| 46:DU:90:VAL:HG12 | 46:DU:91:ASP:N | 2.25 | 0.45 |
| 46:DU:91:ASP:O | 46:DU:92:ARG:O | 2.34 | 0.45 |
| 47:DV:25:LEU:N | 47:DV:94:LEU:HD13 | 2.30 | 0.45 |
| 48:DW:45:TYR:CZ | 48:DW:49:LYS:HE3 | 2.51 | 0.45 |
| 49:DX:72:LYS:HB2 | 49:DX:74:PRO:HD3 | 1.96 | 0.45 |
| 1:AA:538:G:OP2 | 12:AL:115:LYS:CG | 2.63 | 0.45 |
| 1:AA:577:G:H1’ | 1:AA:816:A:C4 | 2.51 | 0.45 |
| 1:AA:602:A:C2 | 1:AA:603:U:C2 | 3.05 | 0.45 |
| 1:AA:617:G:C2 | 1:AA:618:C:C5 | 3.04 | 0.45 |
| 1:AA:659:U:C2 | 1:AA:660:G:C8 | 3.04 | 0.45 |
| 1:AA:976:G:P | 14:AN:32:SER:H | 2.38 | 0.45 |
| 1:AA:1199:U:H4’ | 10:AJ:54:PHE:CE2 | 2.50 | 0.45 |
| 1:AA:1494:G:N2 | 31:BA:1912:A:C2 | 2.84 | 0.45 |
| 2:AB:30:ARG:HG3 | 2:AB:31:TYR:CE2 | 2.52 | 0.45 |
| 5:AE:41:VAL:HG13 | 5:AE:113:ALA:HA | 1.99 | 0.45 |
| 6:AF:3:ARG:HD3 | 6:AF:64:GLN:OE1 | 2.15 | 0.45 |
| 18:AR:22:VAL:O | 18:AR:22:VAL:HG12 | 2.17 | 0.45 |
| 20:AT:30:LYS:HA | 20:AT:30:LYS:HD2 | 1.73 | 0.45 |
| 23:B1:18:ILE:HD12 | 23:B1:18:ILE:N | 2.31 | 0.45 |
| 24:B2:48:HIS:O | 24:B2:49:LYS:HD3 | 2.15 | 0.45 |
| 24:B2:57:ILE:O | 24:B2:57:ILE:HG23 | 2.16 | 0.45 |
| 28:B6:9:LEU:HD22 | 28:B6:9:LEU:C | 2.38 | 0.45 |
| 28:B6:26:ASN:ND2 | 28:B6:32:ASN:ND2 | 2.65 | 0.45 |
| 31:BA:271(N):U:C6 | 31:BA:271(N):U:OP1 | 2.69 | 0.45 |
| 31:BA:362:U:H6 | 31:BA:362:U:H3’ | 1.82 | 0.45 |
| 31:BA:675:A:C8 | 31:BA:804:A:C6 | 3.03 | 0.45 |
| 31:BA:869:G:C6 | 31:BA:870:A:N7 | 2.85 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:940:G:H2' | 31:BA:941:A:O4' | 2.17 | 0.45 |
| 31:BA:1241:A:C2' | 31:BA:1242:A:O5' | 2.64 | 0.45 |
| 31:BA:1299:G:H5" | 31:BA:1300:U:P | 2.57 | 0.45 |
| 31:BA:2406:U:O5' | 31:BA:2406:U:H2' | 2.16 | 0.45 |
| 31:BA:2492:U:H2' | 31:BA:2493:U:C6 | 2.52 | 0.45 |
| 31:BA:2663:G:C5 | 31:BA:2664:G:C5 | 3.05 | 0.45 |
| 31:BA:2830:G:N3 | 31:BA:2883:A:H2 | 2.15 | 0.45 |
| 33:BD:71:ASP:CB | 33:BD:103:ARG:NH2 | 2.79 | 0.45 |
| 33:BD:246:PRO:O | 33:BD:254:THR:HG22 | 2.16 | 0.45 |
| 36:BG:115:ARG:HB2 | 36:BG:116:ASP:H | 1.59 | 0.45 |
| 38:BI:73:GLU:OE1 | 38:BI:137:PRO:HD2 | 2.16 | 0.45 |
| 42:BQ:35:VAL:CG1 | 42:BQ:130:LYS:HB3 | 2.42 | 0.45 |
| 49:BX:40:LYS:C | 49:BX:42:ALA:H | 2.23 | 0.45 |
| 1:CA:159:G:N3 | 1:CA:161:A:OP2 | 2.50 | 0.45 |
| 1:CA:303:A:C5 | 1:CA:304:U:C5 | 3.05 | 0.45 |
| 1:CA:626:U:C2 | 1:CA:627:G:N7 | 2.85 | 0.45 |
| 1:CA:747:C:C5 | 1:CA:748:C:C4 | 3.04 | 0.45 |
| 1:CA:930:C:C2' | 1:CA:931:C:H5' | 2.46 | 0.45 |
| 1:CA:1034:G:N2 | 1:CA:1035:A:N6 | 2.64 | 0.45 |
| 1:CA:1118:C:H42 | 1:CA:1155:G:H1 | 1.64 | 0.45 |
| 1:CA:1413:A:C2 | 1:CA:1414:U:C2 | 3.05 | 0.45 |
| 2:CB:91:PRO:N | 2:CB:154:LEU:HD12 | 2.32 | 0.45 |
| 4:CD:109:GLY:O | 4:CD:110:PHE:C | 2.59 | 0.45 |
| 6:CF:75:LEU:CD2 | 6:CF:79:LEU:HD21 | 2.47 | 0.45 |
| 10:CJ:39:PRO:CB | 10:CJ:70:ARG:HH12 | 2.29 | 0.45 |
| 11:CK:41:THR:HG22 | 11:CK:42:TRP:N | 2.31 | 0.45 |
| 18:CR:31:LEU:H | 18:CR:31:LEU:CD2 | 2.24 | 0.45 |
| 18:CR:45:SER:HB3 | 18:CR:51:LEU:HD21 | 1.98 | 0.45 |
| 20:CT:82:SER:O | 20:CT:86:ARG:CB | 2.64 | 0.45 |
| 22:D0:27:GLU:HG3 | 22:D0:68:GLU:HA | 1.98 | 0.45 |
| 26:D4:19:GLY:C | 26:D4:21:VAL:N | 2.74 | 0.45 |
| 28:D6:46:HIS:ND1 | 31:DA:2371:G:O2' | 2.48 | 0.45 |
| 30:D8:62:LEU:C | 30:D8:64:TYR:H | 2.24 | 0.45 |
| 31:DA:154:G:O6 | 31:DA:172:C:N4 | 2.49 | 0.45 |
| 31:DA:253:C:C2' | 31:DA:254:G:H5' | 2.46 | 0.45 |
| 31:DA:280:C:C2' | 31:DA:281:G:O5' | 2.64 | 0.45 |
| 31:DA:494:G:C8 | 31:DA:494:G:C5' | 3.00 | 0.45 |
| 31:DA:701:G:N2 | 31:DA:732:C:C2 | 2.84 | 0.45 |
| 31:DA:705:A:C2 | 31:DA:706:A:C4 | 3.03 | 0.45 |
| 31:DA:775:G:C4 | 31:DA:794:G:C8 | 3.04 | 0.45 |
| 31:DA:879:G:O6 | 31:DA:899:A:C2 | 2.69 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1132:A:H1' | 39:DN:73:THR:HG21 | 1.99 | 0.45 |
| 31:DA:1276:A:C2 | 31:DA:1277:G:C8 | 3.04 | 0.45 |
| 31:DA:2025:C:H2' | 31:DA:2026:C:C6 | 2.51 | 0.45 |
| 31:DA:2438:U:O2' | 31:DA:2440:C:OP1 | 2.34 | 0.45 |
| 31:DA:2660:A:H5' | 31:DA:2661:G:N3 | 2.32 | 0.45 |
| 33:DD:70:TRP:CZ2 | 33:DD:150:LYS:HD3 | 2.52 | 0.45 |
| 35:DF:53:THR:CG2 | 35:DF:56:GLU:H | 2.28 | 0.45 |
| 35:DF:65:TRP:O | 35:DF:66:PRO:C | 2.59 | 0.45 |
| 36:DG:28:VAL:O | 36:DG:31:VAL:HB | 2.17 | 0.45 |
| 36:DG:37:VAL:HG23 | 36:DG:99:MET:HG3 | 1.98 | 0.45 |
| 37:DH:54:ARG:N | 37:DH:65:HIS:HD2 | 2.14 | 0.45 |
| 39:DN:72:TYR:HE1 | 39:DN:87:LEU:HD23 | 1.81 | 0.45 |
| 41:DP:17:LYS:O | 41:DP:19:VAL:HG23 | 2.16 | 0.45 |
| 41:DP:32:THR:HG21 | 41:DP:37:GLY:HA2 | 1.98 | 0.45 |
| 42:DQ:79:LEU:C | 42:DQ:80:GLU:CD | 2.84 | 0.45 |
| 42:DQ:85:LYS:O | 42:DQ:86:GLY:C | 2.58 | 0.45 |
| 42:DQ:104:PHE:CE1 | 42:DQ:125:LEU:HD11 | 2.43 | 0.45 |
| 44:DS:28:VAL:CG1 | 44:DS:29:PHE:N | 2.79 | 0.45 |
| 44:DS:34:HIS:ND1 | 44:DS:54:LEU:HB2 | 2.31 | 0.45 |
| 44:DS:41:ASP:HB3 | 44:DS:48:LEU:HD11 | 1.98 | 0.45 |
| 45:DT:27:THR:OG1 | 45:DT:28:VAL:N | 2.49 | 0.45 |
| 47:DV:5:VAL:HB | 47:DV:60:GLU:OE1 | 2.16 | 0.45 |
| 49:DX:18:TYR:C | 49:DX:20:GLY:N | 2.74 | 0.45 |
| 50:DY:99:CYS:SG | 50:DY:99:CYS:O | 2.74 | 0.45 |
| 1:AA:97:G:O2' | 1:AA:98:G:O5' | 2.35 | 0.45 |
| 1:AA:382:A:H2' | 1:AA:383:A:H8 | 1.80 | 0.45 |
| 1:AA:1025:U:O2' | 1:AA:1026:G:H8 | 1.99 | 0.45 |
| 1:AA:1277:C:H2' | 1:AA:1278:U:H5' | 1.98 | 0.45 |
| 1:AA:1381:U:C2' | 1:AA:1382:C:H5' | 2.47 | 0.45 |
| 2:AB:22:LYS:HZ3 | 2:AB:40:HIS:HE1 | 1.64 | 0.45 |
| 3:AC:66:VAL:HG11 | 3:AC:91:LEU:CD1 | 2.46 | 0.45 |
| 6:AF:6:VAL:HG12 | 6:AF:7:ASN:N | 2.31 | 0.45 |
| 10:AJ:22:LYS:O | 10:AJ:22:LYS:HD2 | 2.16 | 0.45 |
| 16:AP:43:LYS:CG | 16:AP:48:TRP:CE3 | 2.99 | 0.45 |
| 20:AT:56:MET:HE2 | 20:AT:56:MET:HB3 | 1.75 | 0.45 |
| 29:B7:35:ARG:HD3 | 31:BA:54:G:O2' | 2.16 | 0.45 |
| 31:BA:191:A:O2' | 31:BA:678:C:O2' | 2.32 | 0.45 |
| 31:BA:343:C:C2' | 31:BA:344:G:H5' | 2.46 | 0.45 |
| 31:BA:513:A:C2 | 31:BA:514:A:C4 | 3.04 | 0.45 |
| 31:BA:536:A:C2' | 31:BA:537:C:O5' | 2.65 | 0.45 |
| 31:BA:892:G:C8 | 31:BA:893:C:C4 | 3.04 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1117:G:H2' | 31:BA:1118:C:O4' | 2.16 | 0.45 |
| 31:BA:1130:U:O2 | 31:BA:2025:C:H5" | 2.17 | 0.45 |
| 31:BA:1187:G:H8 | 31:BA:1187:G:O5' | 1.98 | 0.45 |
| 31:BA:1449:A:O3' | 31:BA:1530:C:N4 | 2.50 | 0.45 |
| 31:BA:1497:U:O2 | 31:BA:1497:U:H2' | 2.15 | 0.45 |
| 31:BA:1503:U:C6 | 31:BA:1504:C:H5 | 2.35 | 0.45 |
| 31:BA:1512:U:O2' | 31:BA:1513:C:H5' | 2.17 | 0.45 |
| 31:BA:1549:C:H2' | 31:BA:1550:C:H6 | 1.82 | 0.45 |
| 31:BA:1820:U:C2 | 33:BD:202:LYS:HB3 | 2.52 | 0.45 |
| 31:BA:1996:C:H4' | 31:BA:1997:G:OP1 | 2.16 | 0.45 |
| 31:BA:2523:G:C2' | 31:BA:2524:G:H5' | 2.36 | 0.45 |
| 31:BA:2660:A:C5' | 31:BA:2661:G:N3 | 2.80 | 0.45 |
| 31:BA:2866:U:O4' | 31:BA:2866:U:O2 | 2.33 | 0.45 |
| 34:BE:50:GLY:HA3 | 34:BE:74:PRO:HG3 | 1.99 | 0.45 |
| 34:BE:89:ASP:O | 34:BE:90:THR:CB | 2.65 | 0.45 |
| 36:BG:63:ILE:HD12 | 36:BG:63:ILE:O | 2.16 | 0.45 |
| 36:BG:82:LEU:O | 36:BG:83:ARG:HG3 | 2.16 | 0.45 |
| 37:BH:89:ILE:C | 37:BH:89:ILE:CD1 | 2.89 | 0.45 |
| 41:BP:45:LEU:CD2 | 41:BP:46:LYS:N | 2.80 | 0.45 |
| 44:BS:17:ARG:C | 44:BS:19:LYS:N | 2.74 | 0.45 |
| 44:BS:83:LYS:CE | 44:BS:105:ALA:HB2 | 2.42 | 0.45 |
| 47:BV:18:LEU:O | 47:BV:19:LYS:CB | 2.65 | 0.45 |
| 1:CA:356:A:H2' | 1:CA:357:G:O5' | 2.16 | 0.45 |
| 1:CA:357:G:C2 | 1:CA:358:U:C5 | 3.05 | 0.45 |
| 1:CA:945:G:H2' | 1:CA:945:G:N3 | 2.31 | 0.45 |
| 1:CA:1070:U:C2 | 1:CA:1071:C:C5 | 3.04 | 0.45 |
| 1:CA:1163:C:C2 | 1:CA:1174:G:N2 | 2.84 | 0.45 |
| 1:CA:1320:C:H5' | 19:CS:70:LYS:HG3 | 1.97 | 0.45 |
| 4:CD:171:GLY:HA2 | 4:CD:172:PRO:HD3 | 1.86 | 0.45 |
| 6:CF:3:ARG:HD3 | 6:CF:64:GLN:OE1 | 2.16 | 0.45 |
| 6:CF:23:LYS:HB3 | 6:CF:23:LYS:HE2 | 1.76 | 0.45 |
| 6:CF:39:LYS:O | 6:CF:40:VAL:HB | 2.17 | 0.45 |
| 22:D0:48:GLY:O | 22:D0:49:LYS:C | 2.59 | 0.45 |
| 23:D1:9:GLY:O | 23:D1:10:LYS:HB3 | 2.16 | 0.45 |
| 24:D2:15:LYS:O | 24:D2:16:LEU:HB2 | 2.12 | 0.45 |
| 24:D2:32:LEU:O | 24:D2:32:LEU:HD12 | 2.16 | 0.45 |
| 28:D6:40:CYS:SG | 28:D6:45:LYS:NZ | 2.72 | 0.45 |
| 30:D8:32:LEU:H | 30:D8:32:LEU:CD1 | 2.30 | 0.45 |
| 31:DA:142:A:H8 | 31:DA:1595:G:N2 | 2.10 | 0.45 |
| 31:DA:299:A:H5" | 50:DY:84:ARG:HH21 | 1.81 | 0.45 |
| 31:DA:659:C:C6 | 31:DA:659:C:C4' | 3.00 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:893:C:C2' | 31:DA:894:C:O5' | 2.64 | 0.45 |
| 31:DA:1504:C:HO2' | 31:DA:1505:C:C5' | 2.26 | 0.45 |
| 31:DA:1794:U:H2' | 31:DA:1795:C:C6 | 2.51 | 0.45 |
| 31:DA:1853:A:C6 | 31:DA:1889:A:C5 | 3.04 | 0.45 |
| 31:DA:2291:U:C5' | 31:DA:2380:C:O2 | 2.65 | 0.45 |
| 31:DA:2473:U:O2 | 31:DA:2473:U:H2' | 2.17 | 0.45 |
| 31:DA:2702:U:OP1 | 31:DA:2702:U:O4' | 2.34 | 0.45 |
| 31:DA:2734:A:H5' | 31:DA:2734:A:C8 | 2.50 | 0.45 |
| 31:DA:2836:U:C4 | 31:DA:2883:A:N6 | 2.85 | 0.45 |
| 31:DA:2849:U:H4' | 31:DA:2868:A:C2 | 2.51 | 0.45 |
| 34:DE:36:ARG:NH1 | 34:DE:85:ASN:HD21 | 2.15 | 0.45 |
| 36:DG:63:ILE:HD13 | 36:DG:141:PHE:CZ | 2.50 | 0.45 |
| 40:DO:87:ILE:HD13 | 40:DO:87:ILE:HA | 1.51 | 0.45 |
| 43:DR:13:HIS:CE1 | 43:DR:15:SER:OG | 2.70 | 0.45 |
| 44:DS:13:ARG:HB2 | 44:DS:14:VAL:H | 1.55 | 0.45 |
| 45:DT:82:LEU:HD12 | 45:DT:82:LEU:H | 1.79 | 0.45 |
| 48:DW:13:SER:O | 48:DW:14:PRO:C | 2.59 | 0.45 |
| 49:DX:84:ALA:C | 49:DX:86:GLY:H | 2.25 | 0.45 |
| 51:DZ:40:ASP:OD1 | 51:DZ:42:VAL:HG12 | 2.17 | 0.45 |
| 1:AA:90:U:H6 | 1:AA:90:U:H3' | 1.82 | 0.45 |
| 1:AA:327:A:C6 | 1:AA:329:A:C5 | 3.04 | 0.45 |
| 1:AA:328:C:H4' | 1:AA:329:A:H5' | 1.98 | 0.45 |
| 1:AA:600:C:H2' | 1:AA:601:C:H6 | 1.77 | 0.45 |
| 1:AA:658:G:C5 | 1:AA:659:U:C5 | 3.05 | 0.45 |
| 1:AA:681:C:C2 | 1:AA:710:G:N2 | 2.84 | 0.45 |
| 1:AA:797:C:O2' | 1:AA:798:G:H5' | 2.16 | 0.45 |
| 1:AA:950:U:H2' | 1:AA:951:G:C8 | 2.51 | 0.45 |
| 1:AA:954:G:N2 | 1:AA:1227:A:H62 | 1.99 | 0.45 |
| 1:AA:1098:C:C4 | 1:AA:1099:G:C8 | 3.05 | 0.45 |
| 1:AA:1188:A:H2' | 1:AA:1189:C:H5' | 1.99 | 0.45 |
| 2:AB:44:LEU:O | 2:AB:47:THR:HB | 2.16 | 0.45 |
| 3:AC:134:ILE:HG22 | 3:AC:168:ALA:HB3 | 1.98 | 0.45 |
| 4:AD:8:VAL:O | 4:AD:10:ARG:N | 2.48 | 0.45 |
| 4:AD:127:THR:OG1 | 4:AD:128:VAL:N | 2.48 | 0.45 |
| 5:AE:137:GLU:O | 5:AE:141:GLN:HG3 | 2.14 | 0.45 |
| 10:AJ:29:ARG:HH22 | 10:AJ:84:GLN:HG2 | 1.81 | 0.45 |
| 14:AN:12:ARG:C | 14:AN:14:PRO:HD2 | 2.41 | 0.45 |
| 17:AQ:18:THR:CG2 | 17:AQ:69:LYS:HE3 | 2.44 | 0.45 |
| 19:AS:19:VAL:HG12 | 19:AS:19:VAL:O | 2.16 | 0.45 |
| 22:B0:19:LYS:NZ | 31:BA:2262:U:P | 2.90 | 0.45 |
| 23:B1:30:VAL:O | 23:B1:30:VAL:CG1 | 2.63 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 23:B1:87:PRO:CG | 23:B1:88:LYS:H | 2.29 | 0.45 |
| 27:B5:5:PRO:C | 27:B5:6:VAL:HG23 | 2.42 | 0.45 |
| 27:B5:46:CYS:SG | 27:B5:47:PRO:CG | 3.04 | 0.45 |
| 31:BA:221:A:H4' | 31:BA:222:A:O5' | 2.12 | 0.45 |
| 31:BA:287:C:H42 | 31:BA:354:G:H1 | 1.64 | 0.45 |
| 31:BA:389:G:H1 | 41:BP:71:VAL:H | 1.64 | 0.45 |
| 31:BA:414:C:H2' | 31:BA:415:A:C8 | 2.50 | 0.45 |
| 31:BA:576:U:H2' | 31:BA:577:G:C8 | 2.51 | 0.45 |
| 31:BA:684:G:N1 | 31:BA:774:A:C2 | 2.84 | 0.45 |
| 31:BA:995:C:N3 | 39:BN:4:TYR:HE1 | 2.15 | 0.45 |
| 31:BA:1147:C:C2' | 31:BA:1148:A:O5' | 2.65 | 0.45 |
| 31:BA:1319:G:C6 | 31:BA:1320:C:N4 | 2.84 | 0.45 |
| 31:BA:1503:U:C6 | 31:BA:1504:C:C5 | 3.03 | 0.45 |
| 31:BA:1563:G:H2' | 31:BA:1564:C:H6 | 1.80 | 0.45 |
| 31:BA:1670:C:O2 | 34:BE:129:HIS:CE1 | 2.64 | 0.45 |
| 31:BA:1851:U:C2' | 31:BA:1852:C:H5' | 2.47 | 0.45 |
| 31:BA:2010:G:H5" | 48:BW:42:ARG:HB2 | 1.99 | 0.45 |
| 31:BA:2731:G:C6 | 31:BA:2732:G:O6 | 2.70 | 0.45 |
| 31:BA:2843:G:C2' | 31:BA:2844:G:O5' | 2.65 | 0.45 |
| 32:BB:40:U:H1' | 32:BB:45:A:N6 | 2.32 | 0.45 |
| 32:BB:60:C:C2 | 32:BB:61:G:C8 | 3.05 | 0.45 |
| 33:BD:228:PRO:HD2 | 33:BD:235:GLY:HA3 | 1.98 | 0.45 |
| 33:BD:246:PRO:HB2 | 33:BD:255:LYS:HG3 | 1.97 | 0.45 |
| 34:BE:105:THR:HG22 | 34:BE:106:GLY:N | 2.32 | 0.45 |
| 36:BG:7:LEU:HD23 | 36:BG:176:LEU:HD22 | 1.98 | 0.45 |
| 38:BI:114:LEU:O | 38:BI:115:ALA:C | 2.57 | 0.45 |
| 39:BN:95:PRO:HD2 | 39:BN:96:GLU:OE2 | 2.17 | 0.45 |
| 40:BO:31:LYS:C | 40:BO:32:TYR:HD1 | 2.22 | 0.45 |
| 41:BP:5:ASP:CG | 41:BP:6:LEU:HD23 | 2.41 | 0.45 |
| 43:BR:56:LYS:HD2 | 43:BR:88:ARG:H | 1.80 | 0.45 |
| 44:BS:25:ARG:HH11 | 44:BS:25:ARG:CB | 2.29 | 0.45 |
| 49:BX:53:LYS:N | 49:BX:80:ILE:HG22 | 2.32 | 0.45 |
| 49:BX:90:GLU:C | 49:BX:92:LEU:H | 2.25 | 0.45 |
| 50:BY:37:VAL:HG21 | 50:BY:67:LEU:HD23 | 1.99 | 0.45 |
| 50:BY:66:PRO:O | 50:BY:67:LEU:HB3 | 2.17 | 0.45 |
| 51:BZ:166:SER:HB2 | 51:BZ:167:PRO:C | 2.40 | 0.45 |
| 1:CA:327:A:C2 | 1:CA:329:A:C4 | 3.04 | 0.45 |
| 1:CA:769:G:H4' | 1:CA:1513:A:H4' | 1.98 | 0.45 |
| 1:CA:831:U:O2' | 1:CA:832:C:H5' | 2.17 | 0.45 |
| 1:CA:1064:G:H5' | 1:CA:1066:C:H1' | 1.98 | 0.45 |
| 1:CA:1142:G:H2' | 1:CA:1143:G:O4' | 2.16 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:1199:U:H4' | 10:CJ:54:PHE:CE2 | 2.52 | 0.45 |
| 1:CA:1202:G:H2' | 1:CA:1203:C:O4' | 2.16 | 0.45 |
| 1:CA:1248:A:H2' | 1:CA:1249:C:H5' | 1.99 | 0.45 |
| 1:CA:1452:C:H4' | 1:CA:1456:G:O5' | 2.16 | 0.45 |
| 8:CH:51:VAL:O | 8:CH:52:ASP:HB2 | 2.16 | 0.45 |
| 9:CI:126:SER:C | 9:CI:128:ARG:H | 2.24 | 0.45 |
| 17:CQ:22:LEU:HD11 | 17:CQ:39:SER:HB2 | 1.97 | 0.45 |
| 17:CQ:68:ARG:HG3 | 17:CQ:68:ARG:O | 2.14 | 0.45 |
| 23:D1:34:THR:HG21 | 31:DA:387:U:O3' | 2.15 | 0.45 |
| 29:D7:43:THR:HG23 | 29:D7:44:PRO:CD | 2.46 | 0.45 |
| 30:D8:3:LYS:HE3 | 31:DA:242:G:O5' | 2.17 | 0.45 |
| 31:DA:55:G:O2' | 31:DA:127:A:N1 | 2.48 | 0.45 |
| 31:DA:69:C:O2 | 31:DA:73:A:O2' | 2.33 | 0.45 |
| 31:DA:83:G:N1 | 31:DA:102:G:H2' | 2.24 | 0.45 |
| 31:DA:855:G:C6 | 31:DA:856:C:N4 | 2.85 | 0.45 |
| 31:DA:972:G:C6 | 31:DA:973:A:C6 | 3.04 | 0.45 |
| 31:DA:1496:A:H8 | 31:DA:1577:C:O2' | 2.00 | 0.45 |
| 31:DA:1649:G:C6 | 31:DA:2009:G:C6 | 3.04 | 0.45 |
| 31:DA:1826:G:H2' | 31:DA:1827:C:H6 | 1.81 | 0.45 |
| 31:DA:1890:A:H2' | 31:DA:1891:G:H5' | 1.97 | 0.45 |
| 31:DA:1905:C:H2' | 31:DA:1905:C:O2 | 2.16 | 0.45 |
| 31:DA:2590:A:OP2 | 33:DD:238:GLY:O | 2.34 | 0.45 |
| 31:DA:2660:A:C2 | 31:DA:2661:G:H4' | 2.52 | 0.45 |
| 31:DA:2876:G:H4' | 45:DT:3:ARG:CD | 2.45 | 0.45 |
| 33:DD:222:ARG:O | 33:DD:225:ALA:HB3 | 2.16 | 0.45 |
| 34:DE:37:ARG:HD3 | 34:DE:44:TYR:CE2 | 2.52 | 0.45 |
| 34:DE:82:ARG:HA | 34:DE:82:ARG:HD3 | 1.77 | 0.45 |
| 35:DF:66:PRO:O | 35:DF:67:GLN:CB | 2.64 | 0.45 |
| 35:DF:132:VAL:C | 35:DF:134:GLY:N | 2.73 | 0.45 |
| 36:DG:19:LEU:HG | 36:DG:175:LEU:HD12 | 1.98 | 0.45 |
| 38:DI:25:TYR:CE1 | 38:DI:30:LEU:HD21 | 2.52 | 0.45 |
| 38:DI:133:HIS:ND1 | 38:DI:134:PRO:CD | 2.79 | 0.45 |
| 43:DR:25:ALA:O | 43:DR:26:LYS:C | 2.59 | 0.45 |
| 44:DS:95:HIS:O | 44:DS:96:GLY:C | 2.59 | 0.45 |
| 1:AA:101:A:C2' | 1:AA:102:G:H5' | 2.46 | 0.45 |
| 1:AA:265:G:H4' | 17:AQ:66:SER:HA | 1.98 | 0.45 |
| 1:AA:321:A:C2 | 1:AA:333:G:C2 | 3.04 | 0.45 |
| 1:AA:544:G:H2' | 1:AA:545:C:C6 | 2.52 | 0.45 |
| 1:AA:736:C:H2' | 1:AA:737:A:C8 | 2.51 | 0.45 |
| 1:AA:781:A:C3' | 1:AA:782:A:H5' | 2.47 | 0.45 |
| 4:AD:12:CYS:SG | 4:AD:19:LEU:O | 2.74 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 13:AM:32:GLU:OE2 | 13:AM:64:TRP:CH2 | 2.70 | 0.45 |
| 13:AM:68:GLY:O | 13:AM:70:LEU:N | 2.50 | 0.45 |
| 15:AO:43:LEU:CD1 | 15:AO:56:LEU:HD22 | 2.47 | 0.45 |
| 24:B2:17:SER:O | 24:B2:20:GLU:HB2 | 2.16 | 0.45 |
| 29:B7:15:THR:HG22 | 29:B7:16:HIS:CG | 2.51 | 0.45 |
| 31:BA:343:C:H2' | 31:BA:344:G:H5' | 1.97 | 0.45 |
| 31:BA:706:A:H2' | 31:BA:707:G:O4' | 2.17 | 0.45 |
| 31:BA:776:G:C8 | 31:BA:793:A:C2 | 3.04 | 0.45 |
| 31:BA:806:C:OP2 | 41:BP:39:LYS:CG | 2.64 | 0.45 |
| 31:BA:1225:G:H8 | 31:BA:1225:G:O5' | 2.00 | 0.45 |
| 31:BA:1531:C:H5" | 31:BA:1532:C:H6 | 1.82 | 0.45 |
| 31:BA:1649:G:C2' | 31:BA:1650:G:H5' | 2.47 | 0.45 |
| 31:BA:2756:U:C4' | 31:BA:2757:A:OP1 | 2.59 | 0.45 |
| 31:BA:2787:C:O2 | 34:BE:61:ARG:NH1 | 2.50 | 0.45 |
| 31:BA:2884:U:C5 | 31:BA:2885:C:C5 | 3.05 | 0.45 |
| 32:BB:52:A:O2' | 32:BB:53:A:C8 | 2.66 | 0.45 |
| 33:BD:35:LYS:HE3 | 33:BD:63:ARG:O | 2.16 | 0.45 |
| 33:BD:84:TYR:CD2 | 33:BD:84:TYR:C | 2.95 | 0.45 |
| 33:BD:148:GLU:HB2 | 33:BD:151:LYS:HD2 | 1.99 | 0.45 |
| 34:BE:82:ARG:HA | 34:BE:82:ARG:HD3 | 1.80 | 0.45 |
| 34:BE:120:TRP:O | 34:BE:121:ASN:HB2 | 2.15 | 0.45 |
| 35:BF:78:ILE:HD13 | 35:BF:78:ILE:H | 1.80 | 0.45 |
| 36:BG:15:VAL:HG13 | 36:BG:175:LEU:CD1 | 2.47 | 0.45 |
| 36:BG:39:ILE:HG23 | 36:BG:92:VAL:CG1 | 2.46 | 0.45 |
| 38:BI:11:ASN:C | 38:BI:12:LEU:HD23 | 2.42 | 0.45 |
| 40:BO:3:GLN:CB | 40:BO:4:PRO:HD2 | 2.47 | 0.45 |
| 40:BO:106:LEU:HA | 40:BO:106:LEU:HD23 | 1.47 | 0.45 |
| 41:BP:30:THR:O | 41:BP:31:ALA:C | 2.58 | 0.45 |
| 42:BQ:8:LYS:HG3 | 42:BQ:9:TYR:H | 1.80 | 0.45 |
| 44:BS:69:VAL:HG13 | 44:BS:70:GLY:H | 1.82 | 0.45 |
| 46:BU:52:ARG:HD3 | 46:BU:52:ARG:HA | 1.61 | 0.45 |
| 47:BV:2:PHE:HB3 | 47:BV:42:GLY:HA3 | 1.97 | 0.45 |
| 47:BV:25:LEU:O | 47:BV:27:ALA:N | 2.49 | 0.45 |
| 48:BW:18:ARG:O | 48:BW:19:LEU:C | 2.59 | 0.45 |
| 50:BY:6:HIS:N | 50:BY:6:HIS:ND1 | 2.64 | 0.45 |
| 50:BY:75:ILE:HA | 50:BY:75:ILE:HD13 | 1.52 | 0.45 |
| 51:BZ:54:HIS:HE1 | 51:BZ:123:ASP:OD2 | 1.99 | 0.45 |
| 1:CA:25:C:H5" | 1:CA:524:G:H1' | 1.99 | 0.45 |
| 1:CA:159:G:C4 | 1:CA:161:A:OP2 | 2.69 | 0.45 |
| 1:CA:414:A:H2' | 1:CA:415:A:C8 | 2.52 | 0.45 |
| 2:CB:142:LEU:HD23 | 2:CB:142:LEU:C | 2.42 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 4:CD:127:THR:HB | 4:CD:132:ARG:HA | 1.98 | 0.45 |
| 4:CD:189:PRO:HB2 | 4:CD:194:LEU:CD2 | 2.38 | 0.45 |
| 8:CH:17:THR:O | 8:CH:20:TYR:N | 2.48 | 0.45 |
| 8:CH:80:ILE:O | 8:CH:80:ILE:HG22 | 2.16 | 0.45 |
| 11:CK:102:GLY:O | 11:CK:103:LEU:HD13 | 2.17 | 0.45 |
| 12:CL:82:VAL:H | 12:CL:106:ASP:HB2 | 1.81 | 0.45 |
| 16:CP:9:PHE:CD2 | 16:CP:18:ARG:HB2 | 2.52 | 0.45 |
| 20:CT:26:ASN:HA | 20:CT:29:LYS:HG2 | 1.98 | 0.45 |
| 27:D5:16:ARG:HG2 | 27:D5:16:ARG:NH1 | 2.07 | 0.45 |
| 27:D5:25:LEU:HD11 | 48:DW:19:LEU:HB3 | 1.98 | 0.45 |
| 31:DA:190:A:H2 | 31:DA:679:C:O2 | 1.99 | 0.45 |
| 31:DA:236:C:H2' | 31:DA:237:C:H6 | 1.80 | 0.45 |
| 31:DA:271(S):G:C4 | 31:DA:271(T):C:C6 | 3.04 | 0.45 |
| 31:DA:307:G:H21 | 31:DA:330:A:H62 | 1.65 | 0.45 |
| 31:DA:547:A:H2' | 31:DA:547:A:N3 | 2.32 | 0.45 |
| 31:DA:637:A:OP2 | 41:DP:115:LEU:HB2 | 2.15 | 0.45 |
| 31:DA:682:G:N2 | 31:DA:796:C:C2 | 2.85 | 0.45 |
| 31:DA:945:A:C4 | 31:DA:2448:A:C2 | 3.05 | 0.45 |
| 31:DA:1268:A:C6 | 31:DA:2013:A:C8 | 3.04 | 0.45 |
| 31:DA:1321:A:H2' | 31:DA:1322:A:O4' | 2.17 | 0.45 |
| 31:DA:1862:G:H2' | 31:DA:1863:G:O5' | 2.17 | 0.45 |
| 31:DA:2190:G:C2' | 31:DA:2191:G:H5' | 2.47 | 0.45 |
| 31:DA:2584:U:C2' | 31:DA:2585:U:H5' | 2.47 | 0.45 |
| 31:DA:2658:C:C5' | 31:DA:2659:G:OP2 | 2.64 | 0.45 |
| 33:DD:25:THR:HG21 | 33:DD:81:ALA:CA | 2.47 | 0.45 |
| 33:DD:31:LYS:O | 33:DD:32:SER:O | 2.34 | 0.45 |
| 33:DD:248:SER:C | 33:DD:250:TRP:N | 2.72 | 0.45 |
| 36:DG:106:LEU:HA | 36:DG:110:ALA:HB3 | 1.99 | 0.45 |
| 38:DI:120:ILE:HG22 | 38:DI:122:GLU:H | 1.81 | 0.45 |
| 38:DI:129:THR:HG22 | 38:DI:130:TYR:N | 2.32 | 0.45 |
| 39:DN:58:ASP:C | 39:DN:60:ILE:N | 2.74 | 0.45 |
| 41:DP:125:VAL:O | 41:DP:145:PRO:HD2 | 2.16 | 0.45 |
| 43:DR:71:GLN:HE21 | 43:DR:71:GLN:HA | 1.80 | 0.45 |
| 44:DS:93:LYS:HG3 | 44:DS:93:LYS:O | 2.17 | 0.45 |
| 47:DV:2:PHE:HB3 | 47:DV:42:GLY:HA3 | 1.97 | 0.45 |
| 48:DW:23:LEU:O | 48:DW:27:LYS:HD2 | 2.16 | 0.45 |
| 48:DW:44:ALA:O | 48:DW:45:TYR:C | 2.59 | 0.45 |
| 49:DX:60:ARG:HG2 | 49:DX:74:PRO:HD2 | 1.93 | 0.45 |
| 51:DZ:100:VAL:C | 51:DZ:123:ASP:HB2 | 2.41 | 0.45 |
| 1:AA:219:C:C5 | 1:AA:220:G:C8 | 3.05 | 0.45 |
| 1:AA:297:G:H4' | 1:AA:557:G:H4' | 1.98 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:658:G:O2' | 1:AA:659:U:H5' | 2.17 | 0.45 |
| 1:AA:791:G:C5 | 1:AA:792:A:N7 | 2.84 | 0.45 |
| 1:AA:1125:U:C2' | 1:AA:1126:U:OP2 | 2.65 | 0.45 |
| 1:AA:1138:G:H3' | 1:AA:1138:G:N3 | 2.32 | 0.45 |
| 1:AA:1423:G:H2' | 1:AA:1424:C:O4' | 2.16 | 0.45 |
| 1:AA:1442(A):G:N7 | 45:BT:118:ARG:HD2 | 2.31 | 0.45 |
| 4:AD:65:ARG:HG3 | 4:AD:75:PHE:CD1 | 2.52 | 0.45 |
| 10:AJ:7:LYS:O | 10:AJ:96:ILE:HA | 2.17 | 0.45 |
| 20:AT:81:LYS:O | 20:AT:83:ARG:N | 2.50 | 0.45 |
| 23:B1:54:ALA:CB | 23:B1:56:GLN:O | 2.65 | 0.45 |
| 25:B3:46:ASN:O | 25:B3:50:VAL:HG22 | 2.16 | 0.45 |
| 31:BA:547:A:C8 | 31:BA:549:G:C6 | 3.03 | 0.45 |
| 31:BA:672:C:C2 | 31:BA:673:C:C5 | 3.05 | 0.45 |
| 31:BA:729:G:C5 | 33:BD:208:LYS:HB2 | 2.52 | 0.45 |
| 31:BA:1394:U:C4 | 31:BA:1395:A:C5 | 3.05 | 0.45 |
| 31:BA:1568:G:H21 | 33:BD:58:HIS:CE1 | 2.31 | 0.45 |
| 31:BA:1682:G:H2' | 31:BA:1683:C:C6 | 2.52 | 0.45 |
| 31:BA:1742:G:C8 | 31:BA:1742:G:C3' | 2.99 | 0.45 |
| 31:BA:1857:G:N2 | 31:BA:1886:C:N4 | 2.65 | 0.45 |
| 31:BA:1985:G:O2' | 31:BA:1986:A:H5' | 2.16 | 0.45 |
| 31:BA:1987:G:C8 | 31:BA:1987:G:C5' | 2.99 | 0.45 |
| 31:BA:2637:U:O2' | 31:BA:2638:G:H5' | 2.17 | 0.45 |
| 31:BA:2642:G:C2 | 31:BA:2773:C:C2 | 3.05 | 0.45 |
| 31:BA:2762:G:H2' | 31:BA:2763:G:H5' | 1.99 | 0.45 |
| 33:BD:209:ALA:C | 33:BD:210:GLY:O | 2.55 | 0.45 |
| 34:BE:51:PHE:H | 34:BE:74:PRO:CB | 2.30 | 0.45 |
| 35:BF:18:ARG:CG | 35:BF:19:GLU:N | 2.78 | 0.45 |
| 39:BN:1:MET:CB | 47:BV:20:LEU:HD22 | 2.28 | 0.45 |
| 46:BU:6:THR:HG21 | 46:BU:10:ARG:CZ | 2.46 | 0.45 |
| 49:BX:12:VAL:HG11 | 49:BX:27:THR:HG23 | 1.98 | 0.45 |
| 49:BX:29:TRP:CE3 | 49:BX:76:ARG:HB3 | 2.51 | 0.45 |
| 1:CA:59:A:H1' | 1:CA:354:G:N2 | 2.32 | 0.45 |
| 1:CA:293:G:C6 | 1:CA:294:U:C4 | 3.04 | 0.45 |
| 1:CA:406:G:H5' | 4:CD:5:ILE:HD13 | 1.97 | 0.45 |
| 1:CA:665:A:N6 | 1:CA:725:G:O6 | 2.49 | 0.45 |
| 1:CA:1159:U:C5 | 1:CA:1182:G:C4 | 3.05 | 0.45 |
| 1:CA:1250:A:H4' | 9:CI:68:GLY:H | 1.81 | 0.45 |
| 2:CB:238:LEU:HD23 | 2:CB:238:LEU:H | 1.82 | 0.45 |
| 3:CC:124:ILE:HD11 | 3:CC:130:VAL:HG13 | 1.97 | 0.45 |
| 4:CD:100:ARG:HH22 | 4:CD:118:ARG:HH22 | 1.64 | 0.45 |
| 4:CD:127:THR:OG1 | 4:CD:128:VAL:N | 2.50 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 4:CD:138:TYR:HD2 | 4:CD:139:ARG:N | 2.14 | 0.45 |
| 5:CE:6:PHE:HB2 | 5:CE:34:VAL:HG13 | 1.98 | 0.45 |
| 8:CH:33:GLU:O | 8:CH:34:GLU:C | 2.59 | 0.45 |
| 11:CK:52:GLY:H | 11:CK:55:LYS:HG3 | 1.81 | 0.45 |
| 11:CK:84:VAL:HG11 | 11:CK:95:ILE:HD12 | 1.98 | 0.45 |
| 12:CL:22:SER:C | 12:CL:24:VAL:N | 2.73 | 0.45 |
| 15:CO:43:LEU:CD1 | 15:CO:56:LEU:HD22 | 2.47 | 0.45 |
| 16:CP:75:ARG:C | 16:CP:77:ALA:H | 2.23 | 0.45 |
| 18:CR:44:LEU:O | 18:CR:45:SER:C | 2.60 | 0.45 |
| 18:CR:47:THR:HB | 18:CR:49:LYS:HG3 | 1.99 | 0.45 |
| 19:CS:16:LEU:O | 19:CS:20:LEU:HB2 | 2.16 | 0.45 |
| 23:D1:17:SER:C | 23:D1:18:ILE:HD12 | 2.42 | 0.45 |
| 24:D2:30:ARG:HH11 | 24:D2:30:ARG:CG | 2.30 | 0.45 |
| 26:D4:19:GLY:O | 26:D4:21:VAL:N | 2.48 | 0.45 |
| 31:DA:27:G:C2 | 31:DA:512:G:N3 | 2.85 | 0.45 |
| 31:DA:53:A:H61 | 31:DA:117:G:C2' | 2.29 | 0.45 |
| 31:DA:729:G:C5 | 33:DD:208:LYS:HB2 | 2.51 | 0.45 |
| 31:DA:780:G:C2 | 31:DA:782:A:C2 | 3.05 | 0.45 |
| 31:DA:900:A:H2' | 31:DA:900:A:N3 | 2.32 | 0.45 |
| 31:DA:933:A:H2' | 31:DA:934:G:C5' | 2.45 | 0.45 |
| 31:DA:995:C:N3 | 39:DN:4:TYR:HE1 | 2.15 | 0.45 |
| 31:DA:1525:G:H2' | 31:DA:1526:G:H8 | 1.80 | 0.45 |
| 31:DA:1564:C:O2' | 31:DA:1565:C:H5' | 2.17 | 0.45 |
| 31:DA:1649:G:C2' | 31:DA:1650:G:H5' | 2.47 | 0.45 |
| 31:DA:2094:G:P | 38:DI:22:LYS:HE2 | 2.57 | 0.45 |
| 31:DA:2183:C:H2' | 31:DA:2184:G:C8 | 2.52 | 0.45 |
| 31:DA:2473:U:C2 | 31:DA:2474:C:C6 | 3.04 | 0.45 |
| 31:DA:2591:C:H2' | 31:DA:2592:G:C8 | 2.52 | 0.45 |
| 33:DD:84:TYR:C | 33:DD:84:TYR:CD2 | 2.95 | 0.45 |
| 34:DE:8:LYS:NZ | 34:DE:188:VAL:O | 2.49 | 0.45 |
| 34:DE:111:ARG:CG | 43:DR:2:ARG:HG3 | 2.47 | 0.45 |
| 35:DF:115:ALA:O | 35:DF:116:ASP:C | 2.60 | 0.45 |
| 39:DN:13:TRP:CH2 | 39:DN:130:HIS:CE1 | 3.02 | 0.45 |
| 40:DO:3:GLN:HB2 | 40:DO:4:PRO:HD2 | 1.99 | 0.45 |
| 41:DP:14:LYS:HD2 | 41:DP:14:LYS:H | 1.80 | 0.45 |
| 41:DP:101:VAL:HB | 41:DP:107:LYS:H | 1.81 | 0.45 |
| 42:DQ:46:GLN:O | 42:DQ:47:ILE:C | 2.59 | 0.45 |
| 44:DS:17:ARG:O | 44:DS:18:ILE:CB | 2.62 | 0.45 |
| 44:DS:83:LYS:HE3 | 44:DS:105:ALA:CB | 2.45 | 0.45 |
| 45:DT:109:GLU:O | 45:DT:112:ARG:HG3 | 2.17 | 0.45 |
| 46:DU:83:LEU:CG | 46:DU:88:ILE:HG12 | 2.43 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 49:DX:83:VAL:O | 49:DX:83:VAL:HG23 | 2.16 | 0.45 |
| 1:AA:159:G:C4 | 1:AA:161:A:OP2 | 2.69 | 0.45 |
| 1:AA:258:G:H2' | 1:AA:259:G:H8 | 1.82 | 0.45 |
| 1:AA:356:A:H1' | 1:AA:368:U:O2' | 2.17 | 0.45 |
| 1:AA:461:A:C5 | 1:AA:471:G:C6 | 3.05 | 0.45 |
| 1:AA:533:A:OP1 | 1:AA:533:A:H3' | 2.16 | 0.45 |
| 1:AA:559:A:C5' | 1:AA:560:U:H3' | 2.46 | 0.45 |
| 1:AA:974:A:P | 14:AN:41:ARG:HH12 | 2.40 | 0.45 |
| 1:AA:1075:C:H4' | 1:AA:1101:A:N6 | 2.32 | 0.45 |
| 1:AA:1097:C:O2' | 1:AA:1098:C:H5' | 2.16 | 0.45 |
| 1:AA:1296:C:H5' | 1:AA:1297:C:OP2 | 2.16 | 0.45 |
| 2:AB:180:LEU:O | 2:AB:181:PHE:CB | 2.62 | 0.45 |
| 4:AD:58:LEU:HD22 | 4:AD:62:GLN:HG2 | 1.97 | 0.45 |
| 5:AE:78:HIS:HE1 | 5:AE:142:LEU:HD23 | 1.81 | 0.45 |
| 12:AL:64:TYR:O | 12:AL:65:GLU:HB2 | 2.16 | 0.45 |
| 16:AP:20:VAL:CG2 | 16:AP:32:TYR:HB2 | 2.47 | 0.45 |
| 18:AR:36:ASN:ND2 | 18:AR:39:VAL:HG21 | 2.32 | 0.45 |
| 24:B2:45:SER:O | 24:B2:48:HIS:HB2 | 2.17 | 0.45 |
| 27:B5:36:CYS:C | 27:B5:38:ALA:N | 2.74 | 0.45 |
| 30:B8:35:GLN:NE2 | 30:B8:36:LYS:HZ2 | 2.15 | 0.45 |
| 31:BA:287:C:H2' | 31:BA:288:C:O4' | 2.16 | 0.45 |
| 31:BA:511:U:O4 | 31:BA:512:G:C6 | 2.70 | 0.45 |
| 31:BA:671:C:H2' | 31:BA:672:C:C6 | 2.52 | 0.45 |
| 31:BA:762:U:H4' | 31:BA:763:G:C5' | 2.47 | 0.45 |
| 31:BA:1822:G:C8 | 31:BA:1822:G:H5' | 2.51 | 0.45 |
| 31:BA:1889:A:H2' | 31:BA:1890:A:C8 | 2.51 | 0.45 |
| 31:BA:2502:G:H5" | 31:BA:2503:A:H5" | 1.99 | 0.45 |
| 31:BA:2660:A:H5' | 31:BA:2661:G:N3 | 2.32 | 0.45 |
| 31:BA:2660:A:C2 | 31:BA:2661:G:H4' | 2.51 | 0.45 |
| 31:BA:2702:U:OP1 | 31:BA:2702:U:O4' | 2.34 | 0.45 |
| 31:BA:2802:G:OP2 | 31:BA:2803:C:OP2 | 2.35 | 0.45 |
| 33:BD:83:GLU:OE1 | 33:BD:104:TYR:OH | 2.27 | 0.45 |
| 34:BE:36:ARG:NH1 | 34:BE:85:ASN:ND2 | 2.65 | 0.45 |
| 34:BE:76:ARG:HG3 | 34:BE:195:LEU:CD1 | 2.42 | 0.45 |
| 34:BE:176:ILE:O | 34:BE:176:ILE:HG22 | 2.16 | 0.45 |
| 36:BG:13:GLU:O | 36:BG:13:GLU:HG3 | 2.16 | 0.45 |
| 36:BG:33:ARG:HB2 | 36:BG:162:THR:OG1 | 2.17 | 0.45 |
| 36:BG:63:ILE:HD13 | 36:BG:141:PHE:CE2 | 2.52 | 0.45 |
| 37:BH:89:ILE:HB | 37:BH:90:LYS:H | 1.54 | 0.45 |
| 38:BI:81:VAL:CG1 | 38:BI:88:ILE:HG23 | 2.42 | 0.45 |
| 40:BO:88:ASN:HB3 | 40:BO:92:GLU:O | 2.17 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 41:BP:141:ALA:HB3 | 25:D3:1:MET:HE1 | 1.97 | 0.45 |
| 43:BR:61:HIS:O | 43:BR:62:ALA:C | 2.60 | 0.45 |
| 44:BS:17:ARG:HA | 44:BS:20:ARG:HG2 | 1.99 | 0.45 |
| 45:BT:57:PHE:O | 45:BT:58:ASN:C | 2.58 | 0.45 |
| 46:BU:66:ASN:HD21 | 46:BU:70:ARG:HE | 1.65 | 0.45 |
| 49:BX:53:LYS:H | 49:BX:80:ILE:HG22 | 1.80 | 0.45 |
| 49:BX:80:ILE:O | 49:BX:81:VAL:HB | 2.17 | 0.45 |
| 1:CA:78:G:N2 | 1:CA:91:C:H42 | 2.15 | 0.45 |
| 1:CA:79:G:H4' | 1:CA:80:G:OP1 | 2.17 | 0.45 |
| 1:CA:254:G:O2' | 1:CA:255:G:H5' | 2.15 | 0.45 |
| 1:CA:335:C:O2' | 1:CA:336:C:H5' | 2.16 | 0.45 |
| 1:CA:514:C:O2' | 1:CA:515:G:H5' | 2.17 | 0.45 |
| 1:CA:515:G:H2' | 1:CA:516:U:O4' | 2.17 | 0.45 |
| 1:CA:600:C:H2' | 1:CA:601:C:H6 | 1.78 | 0.45 |
| 1:CA:632:A:H8 | 1:CA:633:G:C8 | 2.34 | 0.45 |
| 1:CA:764:C:C2 | 1:CA:765:G:C8 | 3.05 | 0.45 |
| 1:CA:938:A:C6 | 1:CA:939:G:C5 | 3.05 | 0.45 |
| 1:CA:1031:G:H2' | 1:CA:1032:G:C8 | 2.52 | 0.45 |
| 1:CA:1270:C:H2' | 1:CA:1271:G:O4' | 2.16 | 0.45 |
| 2:CB:44:LEU:O | 2:CB:47:THR:HB | 2.17 | 0.45 |
| 5:CE:79:GLU:HB3 | 5:CE:92:LYS:HG2 | 1.99 | 0.45 |
| 10:CJ:16:LEU:HD13 | 10:CJ:16:LEU:O | 2.17 | 0.45 |
| 12:CL:64:TYR:O | 12:CL:65:GLU:HB2 | 2.17 | 0.45 |
| 16:CP:9:PHE:CE2 | 16:CP:18:ARG:HB2 | 2.52 | 0.45 |
| 17:CQ:6:LEU:O | 17:CQ:58:GLU:HA | 2.16 | 0.45 |
| 23:D1:94:LEU:O | 23:D1:95:LEU:HG | 2.16 | 0.45 |
| 27:D5:41:PRO:O | 27:D5:44:THR:OG1 | 2.35 | 0.45 |
| 27:D5:46:CYS:O | 27:D5:48:GLU:N | 2.48 | 0.45 |
| 28:D6:45:LYS:HB3 | 31:DA:2371:G:H4' | 1.99 | 0.45 |
| 30:D8:62:LEU:C | 30:D8:64:TYR:N | 2.73 | 0.45 |
| 31:DA:707:G:C5 | 31:DA:708:C:C5 | 3.05 | 0.45 |
| 31:DA:774:A:O2' | 31:DA:775:G:H5" | 2.16 | 0.45 |
| 31:DA:829:A:N7 | 31:DA:2248:C:H5' | 2.31 | 0.45 |
| 31:DA:910:A:H2' | 31:DA:2264:C:O2' | 2.17 | 0.45 |
| 31:DA:1011:G:C2 | 31:DA:1151:G:C2 | 3.04 | 0.45 |
| 31:DA:1429:G:H2' | 31:DA:1430:C:H6 | 1.77 | 0.45 |
| 31:DA:1465:G:C4 | 31:DA:1466:G:C8 | 3.04 | 0.45 |
| 31:DA:2280:G:O2' | 31:DA:2281:C:H5' | 2.17 | 0.45 |
| 31:DA:2496:C:P | 42:DQ:81:VAL:HG12 | 2.57 | 0.45 |
| 32:DB:27:C:N4 | 32:DB:28:C:C4 | 2.84 | 0.45 |
| 33:DD:35:LYS:HG2 | 33:DD:64:ILE:HG23 | 1.99 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 34:DE:10:GLY:CA | 45:DT:8:LYS:HE3 | 2.46 | 0.45 |
| 34:DE:31:CYS:HA | 34:DE:32:PRO:HD3 | 1.70 | 0.45 |
| 34:DE:61:ARG:C | 34:DE:63:LEU:N | 2.72 | 0.45 |
| 35:DF:96:ASP:OD1 | 35:DF:96:ASP:C | 2.58 | 0.45 |
| 35:DF:117:ARG:HD3 | 35:DF:117:ARG:HA | 1.75 | 0.45 |
| 35:DF:139:PHE:CD2 | 35:DF:139:PHE:C | 2.95 | 0.45 |
| 36:DG:60:LEU:HD22 | 36:DG:63:ILE:HG13 | 1.99 | 0.45 |
| 37:DH:43:VAL:O | 37:DH:43:VAL:HG23 | 2.16 | 0.45 |
| 39:DN:18:ALA:HB3 | 39:DN:26:LEU:HD22 | 1.98 | 0.45 |
| 40:DO:44:LYS:HD3 | 40:DO:44:LYS:HA | 1.83 | 0.45 |
| 41:DP:79:ARG:NH2 | 41:DP:109:GLY:HA2 | 2.31 | 0.45 |
| 46:DU:93:LYS:H | 46:DU:93:LYS:CD | 2.25 | 0.45 |
| 46:DU:101:ARG:C | 46:DU:102:GLU:HG2 | 2.42 | 0.45 |
| 49:DX:47:PHE:O | 49:DX:48:LYS:C | 2.60 | 0.45 |
| 1:AA:93:G:H2' | 1:AA:96:U:H5' | 1.98 | 0.45 |
| 1:AA:175:C:N3 | 1:AA:176:C:C5 | 2.84 | 0.45 |
| 1:AA:702:A:H3' | 1:AA:703:G:C5' | 2.47 | 0.45 |
| 2:AB:144:ARG:HG3 | 2:AB:145:LEU:N | 2.32 | 0.45 |
| 2:AB:153:ARG:O | 2:AB:154:LEU:C | 2.58 | 0.45 |
| 4:AD:163:GLU:C | 4:AD:165:MET:N | 2.75 | 0.45 |
| 11:AK:48:ILE:HG21 | 11:AK:63:LEU:HD13 | 1.98 | 0.45 |
| 12:AL:18:VAL:HG23 | 12:AL:19:ARG:N | 2.32 | 0.45 |
| 16:AP:43:LYS:HG2 | 16:AP:48:TRP:CD2 | 2.52 | 0.45 |
| 18:AR:50:ILE:HD13 | 18:AR:50:ILE:HA | 1.83 | 0.45 |
| 23:B1:20:ARG:HD3 | 23:B1:41:ARG:HD3 | 1.99 | 0.45 |
| 23:B1:47:GLN:CD | 31:BA:2090:G:H21 | 2.25 | 0.45 |
| 27:B5:57:VAL:C | 27:B5:58:LEU:CG | 2.90 | 0.45 |
| 28:B6:27:LYS:HD2 | 31:BA:2285:C:C6 | 2.52 | 0.45 |
| 30:B8:35:GLN:HE21 | 30:B8:35:GLN:HB3 | 1.56 | 0.45 |
| 30:B8:36:LYS:HE2 | 30:B8:36:LYS:HB2 | 1.80 | 0.45 |
| 31:BA:157:U:H5' | 31:BA:171:G:N2 | 2.32 | 0.45 |
| 31:BA:157:U:OP2 | 31:BA:157:U:H6 | 2.00 | 0.45 |
| 31:BA:658:C:C3' | 31:BA:659:C:H5" | 2.46 | 0.45 |
| 31:BA:1242:A:N1 | 41:BP:8:PRO:HG3 | 2.32 | 0.45 |
| 31:BA:1983:C:C2' | 31:BA:1984:G:H5' | 2.47 | 0.45 |
| 31:BA:2517:C:C2 | 31:BA:2542:A:C6 | 3.05 | 0.45 |
| 34:BE:56:PRO:O | 34:BE:57:LYS:C | 2.60 | 0.45 |
| 38:BI:44:LEU:HD23 | 38:BI:44:LEU:HA | 1.39 | 0.45 |
| 38:BI:130:TYR:HB2 | 38:BI:138:ILE:HD11 | 1.99 | 0.45 |
| 45:BT:34:VAL:HG13 | 45:BT:39:ARG:HA | 1.99 | 0.45 |
| 45:BT:109:GLU:O | 45:BT:112:ARG:HG3 | 2.17 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 46:BU:24:TYR:CD1 | 46:BU:38:THR:HG21 | 2.52 | 0.45 |
| 46:BU:92:ARG:NH2 | 47:BV:10:LYS:HB3 | 2.32 | 0.45 |
| 46:BU:95:LEU:HD13 | 47:BV:11:GLN:HG3 | 1.96 | 0.45 |
| 47:BV:72:VAL:HG13 | 47:BV:88:ARG:NH2 | 2.32 | 0.45 |
| 49:BX:40:LYS:O | 49:BX:42:ALA:N | 2.36 | 0.45 |
| 51:BZ:96:VAL:N | 51:BZ:128:VAL:O | 2.50 | 0.45 |
| 1:CA:166:G:C5 | 1:CA:167:G:N7 | 2.85 | 0.45 |
| 1:CA:245:C:O2 | 1:CA:283:C:N3 | 2.49 | 0.45 |
| 1:CA:303:A:C4 | 1:CA:304:U:C6 | 3.05 | 0.45 |
| 1:CA:344:A:O2' | 1:CA:346:G:N7 | 2.24 | 0.45 |
| 1:CA:366:C:H1' | 1:CA:367:U:OP1 | 2.16 | 0.45 |
| 1:CA:407:G:O2' | 4:CD:116:GLN:HG3 | 2.17 | 0.45 |
| 1:CA:414:A:H2' | 1:CA:415:A:O4' | 2.17 | 0.45 |
| 1:CA:461:A:C5 | 1:CA:471:G:C6 | 3.05 | 0.45 |
| 1:CA:502:G:H4' | 1:CA:550:G:H4' | 1.99 | 0.45 |
| 1:CA:542:G:H5' | 4:CD:41:GLY:HA3 | 1.99 | 0.45 |
| 1:CA:849:C:H2' | 1:CA:850:U:O4' | 2.17 | 0.45 |
| 1:CA:1158:C:O2 | 1:CA:1158:C:C2' | 2.65 | 0.45 |
| 1:CA:1189:C:H5" | 3:CC:5:ILE:HG21 | 1.98 | 0.45 |
| 1:CA:1250:A:H2 | 1:CA:1370:G:H1' | 1.82 | 0.45 |
| 1:CA:1264:C:H2' | 1:CA:1265:G:H8 | 1.82 | 0.45 |
| 1:CA:1322:C:H6 | 1:CA:1322:C:OP1 | 2.00 | 0.45 |
| 2:CB:28:PHE:CD1 | 2:CB:28:PHE:O | 2.69 | 0.45 |
| 2:CB:171:ALA:HA | 2:CB:174:VAL:HG23 | 1.99 | 0.45 |
| 4:CD:101:LEU:C | 4:CD:103:ASN:N | 2.75 | 0.45 |
| 4:CD:150:GLU:CD | 4:CD:150:GLU:H | 2.25 | 0.45 |
| 4:CD:158:ILE:HG23 | 4:CD:162:LEU:CD1 | 2.47 | 0.45 |
| 6:CF:94:GLN:NE2 | 18:CR:32:ARG:HD2 | 2.31 | 0.45 |
| 6:CF:99:ALA:HB1 | 18:CR:23:LYS:NZ | 2.32 | 0.45 |
| 8:CH:8:ASP:O | 8:CH:9:MET:C | 2.59 | 0.45 |
| 10:CJ:30:SER:HB2 | 10:CJ:80:LYS:HG3 | 1.98 | 0.45 |
| 24:D2:20:GLU:O | 24:D2:23:LYS:N | 2.50 | 0.45 |
| 26:D4:28:LYS:CB | 36:DG:113:ARG:HH22 | 2.29 | 0.45 |
| 31:DA:285:C:N3 | 31:DA:286:C:C5 | 2.85 | 0.45 |
| 31:DA:329:G:H22 | 50:DY:19:LYS:HZ1 | 1.63 | 0.45 |
| 31:DA:628:G:C6 | 31:DA:636:G:C2 | 3.05 | 0.45 |
| 31:DA:777:A:C2 | 31:DA:778:G:C8 | 3.05 | 0.45 |
| 31:DA:1106:A:C2' | 31:DA:1107:G:C8 | 2.99 | 0.45 |
| 31:DA:1117:G:H2' | 31:DA:1118:C:O4' | 2.17 | 0.45 |
| 31:DA:1138:G:H1' | 39:DN:105:GLY:O | 2.17 | 0.45 |
| 31:DA:1244:G:C2' | 31:DA:1245:G:H5' | 2.47 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1291:C:O2' | 31:DA:1292:U:H5' | 2.16 | 0.45 |
| 31:DA:1468:C:C2 | 31:DA:1525:G:N2 | 2.84 | 0.45 |
| 31:DA:1470:G:N2 | 31:DA:1523:U:C4 | 2.85 | 0.45 |
| 31:DA:1668:A:H4' | 31:DA:1669:A:O5' | 2.16 | 0.45 |
| 31:DA:1813:G:C2' | 31:DA:1814:G:H5' | 2.47 | 0.45 |
| 31:DA:2031:A:OP1 | 31:DA:2031:A:H8 | 2.00 | 0.45 |
| 31:DA:2065:C:H2' | 31:DA:2066:C:C6 | 2.51 | 0.45 |
| 31:DA:2199:A:OP2 | 31:DA:2200:C:C5 | 2.70 | 0.45 |
| 31:DA:2252:G:H2' | 31:DA:2253:G:C8 | 2.52 | 0.45 |
| 31:DA:2291:U:H2' | 31:DA:2292:C:C6 | 2.52 | 0.45 |
| 31:DA:2483:C:N3 | 42:DQ:124:LYS:NZ | 2.65 | 0.45 |
| 31:DA:2685:G:O2' | 31:DA:2726:U:H5 | 2.00 | 0.45 |
| 32:DB:21:G:O2' | 32:DB:22:U:H6 | 1.98 | 0.45 |
| 33:DD:35:LYS:NZ | 33:DD:64:ILE:C | 2.75 | 0.45 |
| 33:DD:233:HIS:CD2 | 33:DD:233:HIS:H | 2.34 | 0.45 |
| 36:DG:139:LEU:C | 36:DG:141:PHE:H | 2.25 | 0.45 |
| 38:DI:91:SER:CB | 38:DI:121:LYS:HB2 | 2.44 | 0.45 |
| 39:DN:36:GLY:H | 39:DN:42:TRP:HZ3 | 1.63 | 0.45 |
| 40:DO:1:MET:HB2 | 40:DO:32:TYR:HD2 | 1.82 | 0.45 |
| 40:DO:68:GLU:HA | 40:DO:78:ARG:HB2 | 1.98 | 0.45 |
| 45:DT:47:GLY:HA3 | 45:DT:63:VAL:HG23 | 1.98 | 0.45 |
| 47:DV:85:LYS:C | 47:DV:87:HIS:N | 2.74 | 0.45 |
| 50:DY:6:HIS:ND1 | 50:DY:6:HIS:N | 2.64 | 0.45 |
| 1:AA:448:A:C2 | 1:AA:449:C:C4 | 3.05 | 0.45 |
| 1:AA:747:C:C5 | 1:AA:748:C:C4 | 3.05 | 0.45 |
| 1:AA:757:U:H2' | 1:AA:758:G:O4' | 2.17 | 0.45 |
| 1:AA:950:U:H3' | 13:AM:102:ARG:HH12 | 1.81 | 0.45 |
| 1:AA:1248:A:H2' | 1:AA:1249:C:H5' | 1.99 | 0.45 |
| 1:AA:1507:A:C2 | 1:AA:1508:G:C4 | 3.05 | 0.45 |
| 4:AD:121:VAL:O | 4:AD:134:ASP:HA | 2.17 | 0.45 |
| 4:AD:159:ARG:O | 4:AD:163:GLU:N | 2.50 | 0.45 |
| 5:AE:78:HIS:CE1 | 5:AE:142:LEU:HD23 | 2.52 | 0.45 |
| 8:AH:64:LYS:CG | 8:AH:79:VAL:HG21 | 2.47 | 0.45 |
| 10:AJ:38:ILE:O | 10:AJ:38:ILE:HG13 | 2.16 | 0.45 |
| 13:AM:105:THR:O | 13:AM:106:ASN:O | 2.34 | 0.45 |
| 18:AR:44:LEU:O | 18:AR:45:SER:C | 2.59 | 0.45 |
| 20:AT:63:ILE:O | 20:AT:66:ALA:HB3 | 2.17 | 0.45 |
| 25:B3:50:VAL:O | 25:B3:51:ALA:C | 2.59 | 0.45 |
| 28:B6:26:ASN:ND2 | 28:B6:32:ASN:HD21 | 2.15 | 0.45 |
| 31:BA:84:A:H5' | 50:BY:9:LYS:HB3 | 1.99 | 0.45 |
| 31:BA:128:C:O2' | 31:BA:129:C:P | 2.75 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:301:G:C4 | 31:BA:302:C:C5 | 3.05 | 0.45 |
| 31:BA:574:C:N3 | 34:BE:145:LYS:HE2 | 2.31 | 0.45 |
| 31:BA:778:G:C5 | 31:BA:779:U:C4 | 3.05 | 0.45 |
| 31:BA:919:G:H5' | 32:BB:81:G:H1' | 1.99 | 0.45 |
| 31:BA:1374:G:H2' | 31:BA:1375:C:C6 | 2.52 | 0.45 |
| 31:BA:1487:G:O2' | 31:BA:1488:G:H5' | 2.17 | 0.45 |
| 31:BA:1791:A:C5' | 31:BA:1792:G:OP2 | 2.65 | 0.45 |
| 31:BA:1926:U:O2 | 31:BA:1929:G:C2 | 2.70 | 0.45 |
| 31:BA:2228:G:C6 | 31:BA:2229:C:C4 | 3.05 | 0.45 |
| 31:BA:2251:G:C6 | 31:BA:2252:G:C5 | 3.05 | 0.45 |
| 31:BA:2590:A:OP2 | 33:BD:238:GLY:O | 2.35 | 0.45 |
| 31:BA:2843:G:H2' | 31:BA:2844:G:O5' | 2.16 | 0.45 |
| 33:BD:30:GLU:CD | 33:BD:63:ARG:HE | 2.25 | 0.45 |
| 33:BD:61:LEU:O | 33:BD:63:ARG:NH1 | 2.50 | 0.45 |
| 33:BD:186:HIS:HD2 | 33:BD:188:GLU:HB2 | 1.82 | 0.45 |
| 34:BE:8:LYS:NZ | 34:BE:188:VAL:O | 2.50 | 0.45 |
| 35:BF:13:SER:HA | 35:BF:14:PRO:HD3 | 1.84 | 0.45 |
| 36:BG:116:ASP:O | 36:BG:117:PHE:HB3 | 2.17 | 0.45 |
| 37:BH:43:VAL:HG11 | 37:BH:53:GLU:O | 2.16 | 0.45 |
| 39:BN:99:LEU:O | 39:BN:103:VAL:HG23 | 2.16 | 0.45 |
| 41:BP:141:ALA:CB | 25:D3:1:MET:SD | 2.96 | 0.45 |
| 44:BS:28:VAL:HG11 | 44:BS:97:ARG:CZ | 2.47 | 0.45 |
| 44:BS:89:ARG:O | 44:BS:90:GLY:C | 2.60 | 0.45 |
| 46:BU:65:ILE:CG1 | 46:BU:96:ALA:HB3 | 2.47 | 0.45 |
| 47:BV:5:VAL:HG22 | 47:BV:6:LYS:N | 2.31 | 0.45 |
| 48:BW:9:TYR:N | 48:BW:102:HIS:CD2 | 2.74 | 0.45 |
| 49:BX:82:GLN:CG | 49:BX:83:VAL:N | 2.79 | 0.45 |
| 51:BZ:127:LYS:HD3 | 51:BZ:162:GLU:OE1 | 2.16 | 0.45 |
| 1:CA:93:G:H2' | 1:CA:96:U:H5' | 1.98 | 0.45 |
| 1:CA:356:A:H2' | 1:CA:357:G:C8 | 2.52 | 0.45 |
| 1:CA:473:G:C2 | 1:CA:474:G:N7 | 2.85 | 0.45 |
| 1:CA:512:U:H2' | 1:CA:513:C:H6 | 1.80 | 0.45 |
| 1:CA:543:C:O2' | 1:CA:544:G:H5' | 2.16 | 0.45 |
| 1:CA:927:G:OP2 | 1:CA:1503:A:C5 | 2.70 | 0.45 |
| 1:CA:961:U:C4 | 1:CA:962:C:C4 | 3.05 | 0.45 |
| 1:CA:1128:C:H5' | 9:CI:16:ARG:NH1 | 2.32 | 0.45 |
| 1:CA:1168:A:N6 | 1:CA:1169:A:C6 | 2.85 | 0.45 |
| 1:CA:1285:A:C4' | 1:CA:1286:A:O5' | 2.65 | 0.45 |
| 2:CB:84:GLU:OE1 | 2:CB:219:VAL:HB | 2.16 | 0.45 |
| 3:CC:113:ALA:O | 3:CC:115:LEU:N | 2.49 | 0.45 |
| 7:CG:69:VAL:HG13 | 7:CG:134:ALA:O | 2.17 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 10:CJ:51:ARG:HG3 | 10:CJ:61:GLU:N | 2.31 | 0.45 |
| 12:CL:38:THR:HG21 | 12:CL:65:GLU:OE2 | 2.17 | 0.45 |
| 13:CM:83:ASP:CG | 13:CM:84:ILE:N | 2.75 | 0.45 |
| 18:CR:74:ARG:HG3 | 18:CR:79:LEU:HB3 | 2.00 | 0.45 |
| 19:CS:58:VAL:O | 19:CS:58:VAL:HG23 | 2.17 | 0.45 |
| 31:DA:146:G:H8 | 31:DA:146:G:H5' | 1.80 | 0.45 |
| 31:DA:300:A:P | 50:DY:84:ARG:HH22 | 2.39 | 0.45 |
| 31:DA:603:A:O2' | 31:DA:604:G:OP2 | 2.23 | 0.45 |
| 31:DA:993:G:H1' | 47:DV:91:TYR:HD1 | 1.80 | 0.45 |
| 31:DA:1348:G:C2' | 31:DA:1349:A:H5' | 2.46 | 0.45 |
| 31:DA:1374:G:C6 | 31:DA:1375:C:C4 | 3.05 | 0.45 |
| 31:DA:1487:G:O2' | 31:DA:1488:G:H5' | 2.17 | 0.45 |
| 31:DA:1500:G:C5 | 31:DA:1501:C:C4 | 3.04 | 0.45 |
| 31:DA:1862:G:C2' | 31:DA:1863:G:O5' | 2.65 | 0.45 |
| 31:DA:1911:U:C2 | 31:DA:1918:A:C2 | 3.05 | 0.45 |
| 31:DA:1994:C:O2' | 31:DA:1995:U:H5' | 2.17 | 0.45 |
| 31:DA:2887:U:O2' | 31:DA:2888:C:H5' | 2.17 | 0.45 |
| 33:DD:44:ASN:HB2 | 33:DD:48:ARG:O | 2.17 | 0.45 |
| 35:DF:70:THR:CG2 | 35:DF:72:ARG:HB2 | 2.47 | 0.45 |
| 39:DN:18:ALA:HB1 | 39:DN:21:LYS:CB | 2.46 | 0.45 |
| 39:DN:56:ASN:H | 39:DN:125:GLY:CA | 2.30 | 0.45 |
| 40:DO:104:ARG:C | 40:DO:106:LEU:N | 2.71 | 0.45 |
| 43:DR:56:LYS:HD2 | 43:DR:88:ARG:H | 1.82 | 0.45 |
| 44:DS:74:ALA:CB | 44:DS:103:GLU:HG3 | 2.47 | 0.45 |
| 46:DU:66:ASN:CB | 46:DU:76:TYR:HB2 | 2.47 | 0.45 |
| 47:DV:18:LEU:O | 47:DV:19:LYS:CB | 2.65 | 0.45 |
| 49:DX:29:TRP:CE3 | 49:DX:76:ARG:HB3 | 2.52 | 0.45 |
| 51:DZ:111:VAL:CG1 | 51:DZ:112:ARG:N | 2.80 | 0.45 |
| 1:AA:27:G:H2' | 1:AA:28:G:C8 | 2.53 | 0.44 |
| 1:AA:78:G:H1 | 1:AA:91:C:N4 | 2.14 | 0.44 |
| 1:AA:179:A:H2' | 1:AA:180:U:H6 | 1.82 | 0.44 |
| 1:AA:668:G:O2' | 15:AO:46:HIS:HD2 | 2.00 | 0.44 |
| 1:AA:872:A:C5 | 1:AA:874:G:C8 | 3.05 | 0.44 |
| 1:AA:1202:G:H2' | 1:AA:1203:C:O4' | 2.17 | 0.44 |
| 1:AA:1242:C:O5' | 1:AA:1242:C:H6 | 1.99 | 0.44 |
| 1:AA:1457:G:C2 | 1:AA:1458:G:C8 | 3.05 | 0.44 |
| 1:AA:1516:G:H2' | 1:AA:1518:A:OP2 | 2.17 | 0.44 |
| 3:AC:188:LEU:HB3 | 3:AC:189:ALA:H | 1.45 | 0.44 |
| 4:AD:138:TYR:HD2 | 4:AD:139:ARG:N | 2.15 | 0.44 |
| 7:AG:146:GLU:OE2 | 7:AG:149:ARG:HD2 | 2.17 | 0.44 |
| 8:AH:45:ILE:HG22 | 8:AH:63:LEU:HA | 1.99 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 9:AI:79:LEU:C | 9:AI:79:LEU:HD13 | 2.41 | 0.44 |
| 9:AI:122:ALA:HB1 | 9:AI:123:PRO:HD2 | 1.99 | 0.44 |
| 10:AJ:30:SER:HB2 | 10:AJ:80:LYS:HG3 | 1.99 | 0.44 |
| 12:AL:62:SER:O | 12:AL:64:TYR:HD1 | 2.00 | 0.44 |
| 15:AO:74:ASP:HA | 15:AO:75:PRO:HD2 | 1.83 | 0.44 |
| 16:AP:20:VAL:HG21 | 16:AP:32:TYR:CD2 | 2.50 | 0.44 |
| 18:AR:47:THR:HB | 18:AR:49:LYS:HG3 | 1.98 | 0.44 |
| 20:AT:100:ILE:HD12 | 20:AT:100:ILE:N | 2.33 | 0.44 |
| 28:B6:16:CYS:O | 28:B6:18:ARG:CZ | 2.65 | 0.44 |
| 30:B8:34:TRP:HD1 | 31:BA:2391:G:OP1 | 2.00 | 0.44 |
| 31:BA:58:G:H2' | 31:BA:59:U:H6 | 1.82 | 0.44 |
| 31:BA:183:C:H2' | 31:BA:184:C:H5' | 2.00 | 0.44 |
| 31:BA:287:C:C4 | 31:BA:288:C:C5 | 3.05 | 0.44 |
| 31:BA:452:G:C4 | 31:BA:458:G:C6 | 3.05 | 0.44 |
| 31:BA:824:A:C2' | 31:BA:825:C:H5' | 2.46 | 0.44 |
| 31:BA:933:A:H2' | 31:BA:934:G:C5' | 2.47 | 0.44 |
| 31:BA:1499:C:O2' | 31:BA:1500:G:H5' | 2.17 | 0.44 |
| 31:BA:2086:U:H2' | 31:BA:2087:G:C8 | 2.52 | 0.44 |
| 31:BA:2199:A:H3' | 31:BA:2200:C:H6 | 1.81 | 0.44 |
| 31:BA:2467:C:H2' | 31:BA:2468:G:O4' | 2.17 | 0.44 |
| 31:BA:2543:G:H2' | 31:BA:2544:G:C8 | 2.53 | 0.44 |
| 31:BA:2859:G:H3' | 31:BA:2859:G:H8 | 1.81 | 0.44 |
| 33:BD:35:LYS:HA | 33:BD:64:ILE:CG2 | 2.44 | 0.44 |
| 36:BG:33:ARG:N | 36:BG:162:THR:HB | 2.28 | 0.44 |
| 37:BH:55:PRO:O | 37:BH:56:SER:C | 2.59 | 0.44 |
| 44:BS:57:LYS:HG2 | 44:BS:58:LEU:H | 1.82 | 0.44 |
| 47:BV:79:VAL:HG23 | 47:BV:82:ARG:HD2 | 1.98 | 0.44 |
| 50:BY:83:THR:CG2 | 50:BY:94:LYS:HB3 | 2.46 | 0.44 |
| 51:BZ:27:VAL:HG22 | 51:BZ:35:ARG:O | 2.17 | 0.44 |
| 1:CA:295:C:H2' | 1:CA:296:U:C6 | 2.52 | 0.44 |
| 1:CA:559:A:H4' | 1:CA:560:U:H3' | 1.98 | 0.44 |
| 1:CA:617:G:C2 | 1:CA:618:C:C5 | 3.06 | 0.44 |
| 1:CA:709:G:H2' | 1:CA:710:G:C8 | 2.48 | 0.44 |
| 1:CA:1245:A:N1 | 1:CA:1293:G:C6 | 2.85 | 0.44 |
| 1:CA:1289:A:H2' | 1:CA:1290:G:O4' | 2.17 | 0.44 |
| 1:CA:1308:U:H5" | 13:CM:98:VAL:N | 2.31 | 0.44 |
| 4:CD:117:ALA:O | 4:CD:121:VAL:HG22 | 2.16 | 0.44 |
| 4:CD:128:VAL:HA | 4:CD:145:GLU:O | 2.16 | 0.44 |
| 12:CL:110:VAL:HG21 | 12:CL:120:TYR:HB3 | 1.99 | 0.44 |
| 16:CP:21:VAL:O | 16:CP:21:VAL:HG23 | 2.16 | 0.44 |
| 16:CP:43:LYS:HG2 | 16:CP:48:TRP:CD2 | 2.51 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 20:CT:92:LEU:C | 20:CT:94:ALA:H | 2.24 | 0.44 |
| 24:D2:49:LYS:NZ | 24:D2:53:LEU:HD12 | 2.32 | 0.44 |
| 24:D2:55:ARG:O | 24:D2:55:ARG:HG2 | 2.17 | 0.44 |
| 31:DA:18:C:H2' | 31:DA:19:C:C6 | 2.52 | 0.44 |
| 31:DA:49:A:H3' | 31:DA:50:U:C5' | 2.47 | 0.44 |
| 31:DA:108:U:C2 | 31:DA:109:G:C8 | 3.05 | 0.44 |
| 31:DA:196:A:C5 | 31:DA:805:G:C6 | 3.05 | 0.44 |
| 31:DA:327:G:H2' | 31:DA:328:U:C6 | 2.51 | 0.44 |
| 31:DA:391:G:C2 | 31:DA:411:G:C5 | 3.06 | 0.44 |
| 31:DA:513:A:C2 | 31:DA:514:A:C4 | 3.05 | 0.44 |
| 31:DA:565:C:H5' | 31:DA:1253:A:H61 | 1.82 | 0.44 |
| 31:DA:906:G:C2' | 31:DA:907:U:O5' | 2.65 | 0.44 |
| 31:DA:1005:C:C2 | 31:DA:1143:A:C5 | 3.04 | 0.44 |
| 31:DA:1052:C:H3' | 31:DA:1052:C:C6 | 2.51 | 0.44 |
| 31:DA:1241:A:H2' | 31:DA:1242:A:O5' | 2.17 | 0.44 |
| 31:DA:1410:G:C4 | 31:DA:1411:C:C5 | 3.04 | 0.44 |
| 31:DA:1742:G:C8 | 31:DA:1742:G:C3' | 3.00 | 0.44 |
| 31:DA:2311:A:O2' | 31:DA:2312:U:O4' | 2.34 | 0.44 |
| 31:DA:2394:C:C2' | 31:DA:2395:C:H5' | 2.46 | 0.44 |
| 31:DA:2660:A:C5' | 31:DA:2661:G:N3 | 2.80 | 0.44 |
| 31:DA:2862:G:C5 | 31:DA:2863:C:C5 | 3.05 | 0.44 |
| 33:DD:248:SER:HB2 | 33:DD:249:PRO:HD2 | 1.99 | 0.44 |
| 35:DF:102:PRO:HB2 | 35:DF:105:VAL:HG23 | 1.97 | 0.44 |
| 36:DG:146:TYR:O | 36:DG:149:VAL:HG22 | 2.17 | 0.44 |
| 39:DN:28:THR:HG23 | 39:DN:29:LYS:N | 2.33 | 0.44 |
| 39:DN:57:ALA:O | 39:DN:58:ASP:O | 2.34 | 0.44 |
| 41:DP:88:LEU:HD22 | 41:DP:114:ILE:HD12 | 1.99 | 0.44 |
| 42:DQ:63:LYS:HG2 | 42:DQ:65:PHE:CE2 | 2.52 | 0.44 |
| 42:DQ:140:ALA:CB | 51:DZ:99:TYR:HB2 | 2.43 | 0.44 |
| 42:DQ:141:GLN:HG2 | 51:DZ:71:VAL:O | 2.17 | 0.44 |
| 46:DU:80:ILE:HA | 46:DU:80:ILE:HD13 | 1.70 | 0.44 |
| 47:DV:1:MET:SD | 47:DV:46:VAL:HB | 2.57 | 0.44 |
| 47:DV:15:GLU:CB | 47:DV:16:PRO:CD | 2.88 | 0.44 |
| 47:DV:69:LYS:CG | 47:DV:70:ILE:N | 2.74 | 0.44 |
| 47:DV:79:VAL:CG2 | 47:DV:82:ARG:HD2 | 2.47 | 0.44 |
| 47:DV:82:ARG:NH1 | 47:DV:84:LYS:HD2 | 2.32 | 0.44 |
| 49:DX:65:ARG:NH1 | 49:DX:66:LEU:H | 2.15 | 0.44 |
| 50:DY:75:ILE:CD1 | 50:DY:79:CYS:HA | 2.41 | 0.44 |
| 51:DZ:14:LYS:HB2 | 51:DZ:17:ALA:CB | 2.46 | 0.44 |
| 1:AA:90:U:O3' | 1:AA:91:C:H6 | 2.01 | 0.44 |
| 1:AA:159:G:N3 | 1:AA:161:A:OP2 | 2.50 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:283:C:H2' | 1:AA:284:G:O4' | 2.17 | 0.44 |
| 1:AA:414:A:H2' | 1:AA:415:A:C8 | 2.49 | 0.44 |
| 1:AA:564:C:O2' | 1:AA:565:U:H5' | 2.17 | 0.44 |
| 1:AA:570:G:H2' | 1:AA:571:U:C6 | 2.52 | 0.44 |
| 1:AA:594:G:O2' | 1:AA:595:G:H5' | 2.17 | 0.44 |
| 1:AA:892:A:C6 | 1:AA:893:C:N4 | 2.85 | 0.44 |
| 1:AA:930:C:C2' | 1:AA:931:C:H5' | 2.47 | 0.44 |
| 1:AA:955:U:H1' | 1:AA:1227:A:H61 | 1.82 | 0.44 |
| 1:AA:1377:A:H2' | 7:AG:7:ALA:CB | 2.46 | 0.44 |
| 1:AA:1386:G:C2 | 1:AA:1387:G:C8 | 3.05 | 0.44 |
| 1:AA:1435:G:H2' | 1:AA:1436:U:H6 | 1.77 | 0.44 |
| 2:AB:12:GLU:H | 2:AB:12:GLU:HG3 | 1.48 | 0.44 |
| 4:AD:100:ARG:HH22 | 4:AD:118:ARG:HH22 | 1.64 | 0.44 |
| 5:AE:47:LYS:HD3 | 5:AE:47:LYS:N | 2.32 | 0.44 |
| 6:AF:16:GLN:CD | 6:AF:16:GLN:H | 2.25 | 0.44 |
| 12:AL:24:VAL:HG13 | 12:AL:98:TYR:CE2 | 2.51 | 0.44 |
| 16:AP:50:LYS:HD3 | 16:AP:51:VAL:N | 2.32 | 0.44 |
| 16:AP:75:ARG:C | 16:AP:77:ALA:H | 2.25 | 0.44 |
| 17:AQ:29:HIS:HA | 17:AQ:30:PRO:HD2 | 1.84 | 0.44 |
| 17:AQ:52:LYS:HE3 | 17:AQ:52:LYS:HB3 | 1.87 | 0.44 |
| 23:B1:11:ARG:HB3 | 23:B1:12:PRO:HD3 | 2.00 | 0.44 |
| 23:B1:47:GLN:CG | 31:BA:2230:G:H1' | 2.37 | 0.44 |
| 23:B1:78:LYS:NZ | 23:B1:93:GLU:HB2 | 2.33 | 0.44 |
| 27:B5:40:LYS:NZ | 27:B5:49:CYS:SG | 2.86 | 0.44 |
| 27:B5:55:ARG:HG2 | 43:BR:33:ARG:HH11 | 1.82 | 0.44 |
| 30:B8:4:MET:HE1 | 31:BA:593:G:O4' | 2.17 | 0.44 |
| 31:BA:154(A):C:O2 | 31:BA:154(A):C:O4' | 2.35 | 0.44 |
| 31:BA:553:G:C6 | 31:BA:554:U:C4 | 3.05 | 0.44 |
| 31:BA:649:G:C5 | 31:BA:650:C:C4 | 3.04 | 0.44 |
| 31:BA:1316:U:H2' | 31:BA:1317:A:H8 | 1.81 | 0.44 |
| 31:BA:1416:G:O2' | 31:BA:1417:C:P | 2.74 | 0.44 |
| 31:BA:1975:G:C2 | 31:BA:1976:U:O2 | 2.71 | 0.44 |
| 31:BA:2319:G:C2 | 31:BA:2320:A:N1 | 2.85 | 0.44 |
| 31:BA:2337:G:C2 | 31:BA:2338:G:C8 | 3.05 | 0.44 |
| 31:BA:2842:G:C2' | 31:BA:2843:G:H5' | 2.48 | 0.44 |
| 31:BA:2846:G:H2' | 31:BA:2847:U:O4' | 2.17 | 0.44 |
| 32:BB:43:C:H4' | 36:BG:66:GLN:HE22 | 1.82 | 0.44 |
| 33:BD:17:THR:OG1 | 33:BD:204:ILE:HA | 2.16 | 0.44 |
| 34:BE:24:THR:HB | 34:BE:186:GLY:O | 2.17 | 0.44 |
| 34:BE:37:ARG:HD3 | 34:BE:44:TYR:CZ | 2.52 | 0.44 |
| 35:BF:37:VAL:O | 35:BF:41:LEU:HG | 2.17 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 38:BI:31:LEU:HD13 | 38:BI:31:LEU:H | 1.82 | 0.44 |
| 39:BN:1:MET:HB3 | 47:BV:20:LEU:CD2 | 2.28 | 0.44 |
| 39:BN:5:VAL:HG13 | 39:BN:6:PRO:N | 2.32 | 0.44 |
| 39:BN:17:ASP:O | 39:BN:17:ASP:CG | 2.60 | 0.44 |
| 39:BN:97:ARG:O | 39:BN:100:GLU:N | 2.51 | 0.44 |
| 40:BO:87:ILE:HD13 | 40:BO:87:ILE:HA | 1.50 | 0.44 |
| 41:BP:85:LEU:HB3 | 41:BP:114:ILE:CD1 | 2.48 | 0.44 |
| 45:BT:78:LEU:C | 45:BT:79:HIS:HD1 | 2.25 | 0.44 |
| 46:BU:22:LYS:HD3 | 46:BU:22:LYS:HA | 1.71 | 0.44 |
| 47:BV:66:ARG:CD | 47:BV:66:ARG:C | 2.90 | 0.44 |
| 48:BW:40:ASN:O | 48:BW:41:LYS:HD3 | 2.18 | 0.44 |
| 1:CA:321:A:C2 | 1:CA:333:G:C2 | 3.05 | 0.44 |
| 1:CA:659:U:C2 | 1:CA:660:G:C8 | 3.04 | 0.44 |
| 1:CA:677:U:H3 | 1:CA:714:G:N2 | 2.16 | 0.44 |
| 1:CA:1326:C:H2' | 1:CA:1327:C:C6 | 2.52 | 0.44 |
| 1:CA:1432:G:OP1 | 45:DT:107:ASP:HB2 | 2.17 | 0.44 |
| 1:CA:1442:G:C5 | 1:CA:1442(B):A:C2 | 3.05 | 0.44 |
| 12:CL:32:PHE:CE1 | 12:CL:86:ARG:HG3 | 2.51 | 0.44 |
| 12:CL:60:LEU:HD22 | 12:CL:60:LEU:H | 1.82 | 0.44 |
| 16:CP:16:HIS:O | 16:CP:17:TYR:C | 2.61 | 0.44 |
| 19:CS:19:VAL:HG12 | 19:CS:19:VAL:O | 2.16 | 0.44 |
| 22:D0:72:ARG:HH21 | 22:D0:75:LEU:CD1 | 2.30 | 0.44 |
| 24:D2:23:LYS:CA | 49:DX:5:TYR:CD1 | 3.00 | 0.44 |
| 24:D2:28:LYS:HD3 | 24:D2:43:GLN:HB3 | 1.98 | 0.44 |
| 27:D5:42:PRO:CB | 27:D5:43:HIS:HD2 | 2.30 | 0.44 |
| 28:D6:45:LYS:HD3 | 28:D6:45:LYS:HA | 1.79 | 0.44 |
| 30:D8:30:ARG:O | 30:D8:31:HIS:O | 2.35 | 0.44 |
| 30:D8:34:TRP:HD1 | 31:DA:2391:G:OP1 | 2.01 | 0.44 |
| 31:DA:70:G:H21 | 31:DA:71:A:H62 | 1.66 | 0.44 |
| 31:DA:300:A:P | 50:DY:84:ARG:NH2 | 2.90 | 0.44 |
| 31:DA:479:A:H4' | 31:DA:480:A:OP1 | 2.16 | 0.44 |
| 31:DA:1187:G:H8 | 31:DA:1187:G:O5' | 2.00 | 0.44 |
| 31:DA:1316:U:H2' | 31:DA:1317:A:H8 | 1.82 | 0.44 |
| 31:DA:1375:C:H2' | 31:DA:1376:C:H6 | 1.81 | 0.44 |
| 31:DA:1475:G:C2 | 31:DA:1517:G:C2 | 3.05 | 0.44 |
| 31:DA:1853:A:H2' | 31:DA:1854:A:C8 | 2.53 | 0.44 |
| 31:DA:1858:G:H1' | 31:DA:1884:A:N6 | 2.32 | 0.44 |
| 31:DA:2287:A:C2 | 31:DA:2346:A:C2 | 3.05 | 0.44 |
| 31:DA:2592:G:H2' | 31:DA:2593:U:O4' | 2.17 | 0.44 |
| 31:DA:2593:U:C2 | 31:DA:2594:C:C5 | 3.05 | 0.44 |
| 31:DA:2637:U:O2' | 31:DA:2638:G:H5' | 2.16 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2644:G:O2' | 31:DA:2645:G:H5' | 2.18 | 0.44 |
| 31:DA:2877:G:O2' | 31:DA:2878:U:H5' | 2.17 | 0.44 |
| 32:DB:31:C:O2' | 32:DB:32:C:H5' | 2.16 | 0.44 |
| 32:DB:38:C:O4' | 44:DS:95:HIS:NE2 | 2.50 | 0.44 |
| 32:DB:55:U:O2 | 32:DB:56:G:C8 | 2.70 | 0.44 |
| 33:DD:232:PRO:C | 33:DD:233:HIS:O | 2.59 | 0.44 |
| 34:DE:128:SER:OG | 34:DE:129:HIS:N | 2.47 | 0.44 |
| 36:DG:15:VAL:HG13 | 36:DG:175:LEU:CD1 | 2.47 | 0.44 |
| 37:DH:22:GLY:C | 37:DH:23:ARG:HG3 | 2.42 | 0.44 |
| 38:DI:56:LYS:HZ2 | 38:DI:57:ARG:CA | 2.30 | 0.44 |
| 38:DI:91:SER:HB3 | 38:DI:121:LYS:CB | 2.41 | 0.44 |
| 39:DN:75:TYR:CD2 | 39:DN:83:LYS:NZ | 2.71 | 0.44 |
| 43:DR:34:ILE:HD12 | 43:DR:34:ILE:HA | 1.63 | 0.44 |
| 44:DS:14:VAL:O | 44:DS:15:ARG:C | 2.59 | 0.44 |
| 44:DS:66:ALA:HA | 44:DS:69:VAL:CG1 | 2.47 | 0.44 |
| 45:DT:117:ASP:O | 45:DT:118:ARG:C | 2.60 | 0.44 |
| 45:DT:128:GLU:OE1 | 45:DT:129:ARG:N | 2.50 | 0.44 |
| 47:DV:72:VAL:O | 47:DV:73:SER:OG | 2.26 | 0.44 |
| 49:DX:82:GLN:HB3 | 49:DX:85:PRO:CG | 2.42 | 0.44 |
| 50:DY:27:VAL:CG1 | 50:DY:29:GLU:OE1 | 2.65 | 0.44 |
| 1:AA:254:G:O2' | 1:AA:255:G:H5' | 2.18 | 0.44 |
| 1:AA:272:C:H2' | 1:AA:273:A:C8 | 2.51 | 0.44 |
| 1:AA:407:G:O2' | 4:AD:116:GLN:HG3 | 2.17 | 0.44 |
| 1:AA:444:C:C2 | 1:AA:445:G:C8 | 3.05 | 0.44 |
| 1:AA:513:C:H2' | 1:AA:513:C:O2 | 2.18 | 0.44 |
| 1:AA:682:G:C6 | 1:AA:683:G:N7 | 2.85 | 0.44 |
| 1:AA:715:A:O2' | 1:AA:716:A:H5' | 2.17 | 0.44 |
| 1:AA:758:G:H8 | 1:AA:758:G:O5' | 2.01 | 0.44 |
| 1:AA:1293:G:O2' | 1:AA:1294:G:H8 | 1.99 | 0.44 |
| 2:AB:28:PHE:CD1 | 2:AB:28:PHE:O | 2.70 | 0.44 |
| 3:AC:16:ARG:HB2 | 3:AC:16:ARG:NH1 | 2.33 | 0.44 |
| 7:AG:155:ARG:O | 7:AG:156:TRP:HD1 | 2.00 | 0.44 |
| 9:AI:21:PRO:HA | 9:AI:58:ARG:O | 2.17 | 0.44 |
| 9:AI:79:LEU:HD21 | 9:AI:83:ARG:HH21 | 1.82 | 0.44 |
| 9:AI:112:LYS:HA | 9:AI:119:ALA:HA | 1.99 | 0.44 |
| 16:AP:21:VAL:O | 16:AP:21:VAL:HG23 | 2.18 | 0.44 |
| 16:AP:43:LYS:O | 16:AP:44:THR:C | 2.60 | 0.44 |
| 23:B1:48:LYS:HD3 | 23:B1:48:LYS:HA | 1.49 | 0.44 |
| 24:B2:18:PRO:O | 24:B2:22:GLU:HB2 | 2.17 | 0.44 |
| 27:B5:41:PRO:O | 27:B5:42:PRO:C | 2.61 | 0.44 |
| 31:BA:58:G:H1 | 31:BA:69:C:H42 | 1.65 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:280:C:C2' | 31:BA:281:G:O5' | 2.64 | 0.44 |
| 31:BA:452:G:C2 | 31:BA:458:G:C5 | 3.06 | 0.44 |
| 31:BA:754:C:H2' | 31:BA:755:C:C6 | 2.52 | 0.44 |
| 31:BA:828:U:O2 | 31:BA:828:U:H3' | 2.17 | 0.44 |
| 31:BA:2295:C:O2 | 31:BA:2295:C:H2' | 2.16 | 0.44 |
| 31:BA:2500:U:H5" | 31:BA:2501:C:OP2 | 2.16 | 0.44 |
| 32:BB:45:A:C4 | 32:BB:46:A:C8 | 3.06 | 0.44 |
| 32:BB:79:C:C2' | 32:BB:80:U:H5' | 2.47 | 0.44 |
| 33:BD:9:TYR:C | 33:BD:10:THR:CG2 | 2.90 | 0.44 |
| 33:BD:35:LYS:HG2 | 33:BD:64:ILE:HG23 | 1.99 | 0.44 |
| 34:BE:11:MET:HB2 | 34:BE:23:VAL:O | 2.18 | 0.44 |
| 35:BF:104:LYS:O | 35:BF:108:LYS:HG2 | 2.17 | 0.44 |
| 39:BN:16:ILE:HD11 | 39:BN:26:LEU:HD11 | 1.97 | 0.44 |
| 40:BO:35:VAL:HA | 40:BO:62:VAL:CG1 | 2.42 | 0.44 |
| 40:BO:68:GLU:HB3 | 40:BO:78:ARG:NH1 | 2.33 | 0.44 |
| 41:BP:138:LEU:C | 41:BP:138:LEU:HD12 | 2.42 | 0.44 |
| 42:BQ:14:ARG:HG2 | 42:BQ:41:TRP:HH2 | 1.82 | 0.44 |
| 42:BQ:52:VAL:HA | 42:BQ:55:VAL:HG13 | 1.98 | 0.44 |
| 42:BQ:77:LYS:NZ | 42:BQ:84:GLY:O | 2.50 | 0.44 |
| 45:BT:29:ARG:CG | 45:BT:30:VAL:HG13 | 2.47 | 0.44 |
| 47:BV:62:LEU:HB3 | 47:BV:98:GLU:HA | 2.00 | 0.44 |
| 47:BV:64:HIS:O | 47:BV:64:HIS:ND1 | 2.50 | 0.44 |
| 47:BV:66:ARG:HD2 | 47:BV:66:ARG:C | 2.42 | 0.44 |
| 50:BY:17:SER:CA | 50:BY:71:LYS:HD2 | 2.42 | 0.44 |
| 1:CA:165:C:H2' | 1:CA:166:G:H8 | 1.82 | 0.44 |
| 1:CA:197:A:N6 | 1:CA:221:C:H5' | 2.33 | 0.44 |
| 1:CA:251:G:H4' | 1:CA:252:U:O5' | 2.16 | 0.44 |
| 1:CA:324:G:N2 | 1:CA:327:A:C8 | 2.86 | 0.44 |
| 1:CA:328:C:H4' | 1:CA:329:A:H5' | 1.99 | 0.44 |
| 1:CA:354:G:C4 | 1:CA:355:C:C5 | 3.06 | 0.44 |
| 1:CA:443:C:H2' | 1:CA:443:C:O2 | 2.17 | 0.44 |
| 1:CA:499:A:H4' | 1:CA:500:G:H5' | 2.00 | 0.44 |
| 1:CA:612:C:O2 | 1:CA:629:G:N2 | 2.50 | 0.44 |
| 1:CA:1265:G:C6 | 1:CA:1266:G:C6 | 3.06 | 0.44 |
| 1:CA:1334:G:OP2 | 1:CA:1334:G:C8 | 2.70 | 0.44 |
| 2:CB:97:TRP:CH2 | 2:CB:176:GLU:CD | 2.96 | 0.44 |
| 3:CC:206:GLU:O | 3:CC:207:VAL:C | 2.61 | 0.44 |
| 4:CD:14:ARG:C | 4:CD:16:GLY:H | 2.25 | 0.44 |
| 4:CD:119:GLN:O | 4:CD:123:HIS:CD2 | 2.70 | 0.44 |
| 5:CE:139:LEU:C | 5:CE:141:GLN:H | 2.24 | 0.44 |
| 7:CG:70:LYS:O | 7:CG:138:LYS:HE3 | 2.16 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 9:CI:99:LEU:O | 9:CI:100:GLY:C | 2.60 | 0.44 |
| 12:CL:22:SER:C | 12:CL:24:VAL:H | 2.26 | 0.44 |
| 22:D0:36:ILE:C | 22:D0:36:ILE:HD13 | 2.42 | 0.44 |
| 23:D1:64:ALA:C | 23:D1:67:ILE:HD11 | 2.42 | 0.44 |
| 24:D2:41:ILE:HG12 | 31:DA:94(A):G:H21 | 1.81 | 0.44 |
| 28:D6:11:LEU:CD1 | 28:D6:51:GLU:HB2 | 2.47 | 0.44 |
| 30:D8:8:LYS:CE | 31:DA:243:U:OP2 | 2.65 | 0.44 |
| 30:D8:32:LEU:CD2 | 30:D8:35:GLN:H | 2.28 | 0.44 |
| 31:DA:271(F):C:H2' | 31:DA:271(G):C:C6 | 2.36 | 0.44 |
| 31:DA:282:A:C4 | 31:DA:359:A:C2 | 3.05 | 0.44 |
| 31:DA:515:A:H1' | 31:DA:581:C:H1' | 1.99 | 0.44 |
| 31:DA:547:A:H8 | 31:DA:549:G:O6 | 1.99 | 0.44 |
| 31:DA:776:G:H4' | 31:DA:777:A:O5' | 2.17 | 0.44 |
| 31:DA:897:C:O2' | 31:DA:899:A:N7 | 2.39 | 0.44 |
| 31:DA:1412:A:H2' | 31:DA:1413:G:H8 | 1.82 | 0.44 |
| 31:DA:2334:G:H5' | 44:DS:13:ARG:HB3 | 2.00 | 0.44 |
| 31:DA:2759:G:C2' | 31:DA:2760:C:H5' | 2.47 | 0.44 |
| 31:DA:2895:U:C5 | 31:DA:2896:C:C5 | 3.06 | 0.44 |
| 32:DB:29:A:H5" | 44:DS:32:LEU:CD1 | 2.47 | 0.44 |
| 38:DI:29:TYR:HD2 | 38:DI:30:LEU:HD23 | 1.82 | 0.44 |
| 42:DQ:16:ARG:NH1 | 42:DQ:16:ARG:CB | 2.80 | 0.44 |
| 44:DS:58:LEU:HD21 | 44:DS:68:GLN:CB | 2.47 | 0.44 |
| 45:DT:29:ARG:HD2 | 45:DT:29:ARG:HA | 1.41 | 0.44 |
| 47:DV:53:GLU:O | 47:DV:54:GLY:C | 2.59 | 0.44 |
| 1:AA:189(F):U:C4 | 17:AQ:72:ARG:NH2 | 2.86 | 0.44 |
| 1:AA:320:C:H2' | 1:AA:321:A:O4' | 2.18 | 0.44 |
| 1:AA:537:G:OP1 | 12:AL:113:ARG:NH2 | 2.46 | 0.44 |
| 1:AA:685:G:O2' | 1:AA:686:U:C5' | 2.55 | 0.44 |
| 1:AA:783:C:C2' | 1:AA:784:C:H5' | 2.47 | 0.44 |
| 1:AA:1188:A:C2' | 1:AA:1189:C:H5' | 2.48 | 0.44 |
| 1:AA:1285:A:C4' | 1:AA:1286:A:O5' | 2.64 | 0.44 |
| 1:AA:1392:G:N2 | 1:AA:1502:A:C8 | 2.85 | 0.44 |
| 4:AD:75:PHE:CE2 | 4:AD:93:PHE:CZ | 3.06 | 0.44 |
| 5:AE:139:LEU:O | 5:AE:142:LEU:HG | 2.18 | 0.44 |
| 7:AG:75:VAL:O | 7:AG:75:VAL:HG23 | 2.18 | 0.44 |
| 12:AL:110:VAL:CG2 | 12:AL:120:TYR:HB3 | 2.47 | 0.44 |
| 16:AP:4:ILE:HD12 | 16:AP:4:ILE:N | 2.31 | 0.44 |
| 16:AP:16:HIS:O | 16:AP:17:TYR:C | 2.61 | 0.44 |
| 23:B1:20:ARG:CD | 23:B1:41:ARG:HD3 | 2.47 | 0.44 |
| 24:B2:33:MET:HG2 | 49:BX:11:PRO:HD2 | 1.99 | 0.44 |
| 28:B6:45:LYS:HB3 | 31:BA:2371:G:H4' | 1.99 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:27:G:C2 | 31:BA:512:G:N3 | 2.86 | 0.44 |
| 31:BA:197:A:H2' | 31:BA:198:C:H5' | 1.98 | 0.44 |
| 31:BA:685:A:C8 | 31:BA:774:A:C6 | 3.05 | 0.44 |
| 31:BA:813:U:H2' | 31:BA:814:C:C6 | 2.52 | 0.44 |
| 31:BA:818:G:H4' | 31:BA:838:C:O3' | 2.18 | 0.44 |
| 31:BA:926:A:H5" | 31:BA:926:A:H8 | 1.82 | 0.44 |
| 31:BA:933:A:H2' | 31:BA:934:G:H5' | 2.00 | 0.44 |
| 31:BA:1340:U:OP2 | 49:BX:76:ARG:NH2 | 2.50 | 0.44 |
| 31:BA:1491:G:O2' | 33:BD:101:GLU:HB2 | 2.18 | 0.44 |
| 31:BA:1528(A):A:H2' | 31:BA:1529:G:O4' | 2.17 | 0.44 |
| 31:BA:1657:C:H2' | 31:BA:1658:C:H6 | 1.82 | 0.44 |
| 31:BA:2593:U:C2 | 31:BA:2594:C:C5 | 3.05 | 0.44 |
| 31:BA:2654:A:OP1 | 31:BA:2654:A:C8 | 2.63 | 0.44 |
| 31:BA:2855:C:H2' | 31:BA:2856:C:H6 | 1.82 | 0.44 |
| 32:BB:13:A:O2' | 32:BB:15:A:O5' | 2.35 | 0.44 |
| 33:BD:11:PRO:C | 33:BD:13:ARG:H | 2.24 | 0.44 |
| 33:BD:231:HIS:CD2 | 33:BD:232:PRO:HD2 | 2.52 | 0.44 |
| 34:BE:70:ALA:O | 34:BE:72:VAL:O | 2.35 | 0.44 |
| 36:BG:91:ARG:HD2 | 36:BG:91:ARG:C | 2.43 | 0.44 |
| 37:BH:105:LEU:HD13 | 37:BH:105:LEU:H | 1.82 | 0.44 |
| 41:BP:23:PRO:O | 41:BP:33:ARG:NE | 2.50 | 0.44 |
| 44:BS:20:ARG:HD3 | 44:BS:20:ARG:HA | 1.53 | 0.44 |
| 44:BS:41:ASP:HB3 | 44:BS:48:LEU:HD11 | 1.98 | 0.44 |
| 45:BT:88:ILE:HG22 | 45:BT:89:VAL:N | 2.32 | 0.44 |
| 49:BX:23:GLU:O | 49:BX:25:LYS:N | 2.51 | 0.44 |
| 49:BX:59:VAL:O | 49:BX:60:ARG:O | 2.35 | 0.44 |
| 51:BZ:9:TYR:HE2 | 51:BZ:61:LEU:HD13 | 1.82 | 0.44 |
| 51:BZ:118:GLN:O | 51:BZ:120:ILE:N | 2.49 | 0.44 |
| 51:BZ:139:VAL:C | 51:BZ:141:VAL:H | 2.24 | 0.44 |
| 1:CA:41:G:H2' | 1:CA:42:G:H8 | 1.80 | 0.44 |
| 1:CA:255:G:H2' | 1:CA:256:U:C6 | 2.52 | 0.44 |
| 1:CA:303:A:H2' | 1:CA:304:U:O4' | 2.18 | 0.44 |
| 1:CA:355:C:N3 | 1:CA:356:A:N7 | 2.66 | 0.44 |
| 1:CA:389:A:H2' | 1:CA:390:C:H5' | 1.99 | 0.44 |
| 1:CA:577:G:C8 | 1:CA:816:A:N1 | 2.86 | 0.44 |
| 1:CA:726:C:O2' | 1:CA:727:G:H5' | 2.17 | 0.44 |
| 1:CA:736:C:H2' | 1:CA:737:A:C8 | 2.53 | 0.44 |
| 2:CB:67:THR:HG21 | 2:CB:155:LEU:CG | 2.46 | 0.44 |
| 2:CB:194:PRO:HB2 | 2:CB:195:ASP:H | 1.61 | 0.44 |
| 3:CC:66:VAL:HG11 | 3:CC:91:LEU:CD1 | 2.47 | 0.44 |
| 4:CD:18:LYS:HD2 | 4:CD:33:MET:HG2 | 2.00 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 5:CE:41:VAL:HG13 | 5:CE:113:ALA:HA | 1.99 | 0.44 |
| 12:CL:58:VAL:O | 12:CL:65:GLU:HA | 2.16 | 0.44 |
| 21:CU:12:LYS:HB3 | 21:CU:22:ARG:HD2 | 2.00 | 0.44 |
| 22:D0:27:GLU:HB2 | 22:D0:69:PHE:CD1 | 2.52 | 0.44 |
| 26:D4:11:PRO:C | 26:D4:13:ARG:N | 2.75 | 0.44 |
| 31:DA:21:A:H2' | 31:DA:22:C:C6 | 2.52 | 0.44 |
| 31:DA:41:C:H2' | 31:DA:42:G:O4' | 2.17 | 0.44 |
| 31:DA:384:U:O2' | 31:DA:385:C:H5' | 2.18 | 0.44 |
| 31:DA:518:G:H4' | 48:DW:18:ARG:HH12 | 1.81 | 0.44 |
| 31:DA:844:C:H2' | 31:DA:845:G:H5' | 1.99 | 0.44 |
| 31:DA:1011:G:C5 | 31:DA:1013:C:C5 | 3.06 | 0.44 |
| 31:DA:2399:G:C6 | 31:DA:2400:G:C5 | 3.05 | 0.44 |
| 31:DA:2468:G:C6 | 31:DA:2481:G:C4 | 3.05 | 0.44 |
| 31:DA:2648:C:H2' | 31:DA:2649:U:C6 | 2.52 | 0.44 |
| 32:DB:9:G:OP1 | 44:DS:17:ARG:HD2 | 2.18 | 0.44 |
| 32:DB:70:C:C4 | 32:DB:71:C:C5 | 3.05 | 0.44 |
| 34:DE:37:ARG:HD3 | 34:DE:44:TYR:CZ | 2.52 | 0.44 |
| 34:DE:105:THR:HG22 | 34:DE:106:GLY:N | 2.31 | 0.44 |
| 36:DG:107:LEU:HD23 | 36:DG:111:LEU:HD12 | 1.99 | 0.44 |
| 37:DH:87:LEU:HD23 | 37:DH:164:TYR:HD1 | 1.83 | 0.44 |
| 40:DO:10:VAL:HG22 | 40:DO:10:VAL:O | 2.17 | 0.44 |
| 41:DP:133:SER:O | 41:DP:137:LYS:HG2 | 2.18 | 0.44 |
| 42:DQ:23:GLY:O | 42:DQ:100:GLY:CA | 2.54 | 0.44 |
| 45:DT:124:ASP:C | 45:DT:126:ALA:N | 2.75 | 0.44 |
| 47:DV:78:LYS:HD3 | 47:DV:78:LYS:C | 2.42 | 0.44 |
| 1:AA:66:G:H4' | 1:AA:173:U:H5 | 1.78 | 0.44 |
| 1:AA:124:G:C6 | 1:AA:125:U:N3 | 2.86 | 0.44 |
| 1:AA:490:G:OP2 | 4:AD:132:ARG:NH2 | 2.49 | 0.44 |
| 1:AA:590:C:O2' | 1:AA:591:U:H5' | 2.17 | 0.44 |
| 1:AA:707:C:O2' | 1:AA:708:C:C5' | 2.60 | 0.44 |
| 1:AA:1157:A:N3 | 1:AA:1181:G:N3 | 2.65 | 0.44 |
| 1:AA:1168:A:N6 | 1:AA:1169:A:C6 | 2.84 | 0.44 |
| 1:AA:1184:G:OP1 | 1:AA:1184:G:H8 | 2.01 | 0.44 |
| 2:AB:59:GLU:O | 2:AB:63:MET:HG2 | 2.17 | 0.44 |
| 2:AB:67:THR:HG21 | 2:AB:155:LEU:CG | 2.45 | 0.44 |
| 2:AB:193:ASP:O | 2:AB:194:PRO:O | 2.35 | 0.44 |
| 4:AD:158:ILE:HG23 | 4:AD:162:LEU:CD1 | 2.47 | 0.44 |
| 4:AD:189:PRO:HB2 | 4:AD:194:LEU:CD2 | 2.36 | 0.44 |
| 5:AE:11:ILE:HB | 5:AE:31:LEU:HB3 | 2.00 | 0.44 |
| 8:AH:30:ARG:HB3 | 8:AH:30:ARG:NH1 | 2.33 | 0.44 |
| 9:AI:13:ALA:HA | 9:AI:67:GLY:O | 2.18 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 12:AL:62:SER:C | 12:AL:64:TYR:N | 2.75 | 0.44 |
| 30:B8:17:THR:OG1 | 30:B8:18:ALA:N | 2.45 | 0.44 |
| 31:BA:53:A:H61 | 31:BA:117:G:C2' | 2.30 | 0.44 |
| 31:BA:286:C:N3 | 31:BA:287:C:C5 | 2.85 | 0.44 |
| 31:BA:814:C:O2' | 31:BA:815:C:H5' | 2.18 | 0.44 |
| 31:BA:877:U:O2' | 31:BA:878:A:H5" | 2.18 | 0.44 |
| 31:BA:1321:A:H2' | 31:BA:1322:A:O4' | 2.17 | 0.44 |
| 31:BA:1363:C:C2' | 31:BA:1364:G:O5' | 2.66 | 0.44 |
| 31:BA:1505:C:H2' | 31:BA:1506:C:O4' | 2.18 | 0.44 |
| 31:BA:1777:U:C2' | 31:BA:1778:U:H5' | 2.46 | 0.44 |
| 31:BA:2206:G:N3 | 31:BA:2207:G:H5' | 2.33 | 0.44 |
| 31:BA:2652:C:H2' | 31:BA:2653:U:H5' | 1.98 | 0.44 |
| 33:BD:30:GLU:HG3 | 33:BD:63:ARG:NE | 2.32 | 0.44 |
| 34:BE:3:GLY:HA3 | 34:BE:81:ILE:HG21 | 2.00 | 0.44 |
| 34:BE:109:LYS:HD3 | 43:BR:2:ARG:HH12 | 1.82 | 0.44 |
| 36:BG:47:LYS:HG3 | 36:BG:82:LEU:CD1 | 2.47 | 0.44 |
| 36:BG:139:LEU:C | 36:BG:141:PHE:H | 2.26 | 0.44 |
| 38:BI:79:ILE:HA | 38:BI:80:PRO:HD3 | 1.78 | 0.44 |
| 43:BR:8:ARG:HE | 43:BR:8:ARG:CA | 2.15 | 0.44 |
| 43:BR:44:LEU:O | 43:BR:45:ARG:C | 2.59 | 0.44 |
| 43:BR:111:LEU:HD23 | 43:BR:111:LEU:HA | 1.74 | 0.44 |
| 46:BU:92:ARG:HB3 | 46:BU:95:LEU:HD12 | 1.98 | 0.44 |
| 48:BW:27:LYS:O | 48:BW:28:SER:C | 2.60 | 0.44 |
| 51:BZ:51:ALA:O | 51:BZ:52:SER:HB3 | 2.18 | 0.44 |
| 51:BZ:74:VAL:HG22 | 51:BZ:86:VAL:CG1 | 2.46 | 0.44 |
| 51:BZ:126:VAL:HG12 | 51:BZ:163:LEU:HA | 1.98 | 0.44 |
| 51:BZ:152:ALA:HB2 | 51:BZ:168:GLU:CA | 2.48 | 0.44 |
| 1:CA:102:G:C5 | 1:CA:103:C:C5 | 3.06 | 0.44 |
| 1:CA:137:C:N4 | 1:CA:226:G:H1 | 2.14 | 0.44 |
| 1:CA:189(F):U:C4 | 17:CQ:72:ARG:NH2 | 2.86 | 0.44 |
| 1:CA:196:A:N3 | 1:CA:222:U:H1' | 2.33 | 0.44 |
| 1:CA:353:A:C2' | 1:CA:354:G:OP2 | 2.65 | 0.44 |
| 1:CA:500:G:N2 | 1:CA:546:G:H1' | 2.32 | 0.44 |
| 1:CA:542:G:O2' | 1:CA:543:C:H5' | 2.17 | 0.44 |
| 1:CA:544:G:H2' | 1:CA:545:C:C6 | 2.53 | 0.44 |
| 1:CA:562:C:N4 | 1:CA:884:U:C6 | 2.85 | 0.44 |
| 1:CA:601:C:O2' | 1:CA:602:A:H5' | 2.17 | 0.44 |
| 1:CA:677:U:H2' | 1:CA:678:U:H6 | 1.81 | 0.44 |
| 1:CA:859:A:C8 | 1:CA:860:A:C8 | 3.05 | 0.44 |
| 1:CA:930:C:C4 | 1:CA:931:C:C5 | 3.06 | 0.44 |
| 1:CA:1107:C:C4 | 1:CA:1108:G:C8 | 3.04 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:1138:G:H3' | 1:CA:1138:G:N3 | 2.33 | 0.44 |
| 2:CB:219:VAL:HA | 2:CB:222:ILE:CD1 | 2.47 | 0.44 |
| 4:CD:92:VAL:O | 4:CD:96:LEU:HD22 | 2.17 | 0.44 |
| 5:CE:103:GLY:O | 5:CE:104:ALA:C | 2.61 | 0.44 |
| 9:CI:111:ARG:HG3 | 14:CN:61:TRP:HE1 | 1.83 | 0.44 |
| 10:CJ:7:LYS:O | 10:CJ:96:ILE:HA | 2.17 | 0.44 |
| 15:CO:32:LEU:O | 15:CO:36:ILE:HG13 | 2.17 | 0.44 |
| 17:CQ:5:VAL:O | 17:CQ:6:LEU:HD23 | 2.18 | 0.44 |
| 20:CT:63:ILE:O | 20:CT:66:ALA:HB3 | 2.17 | 0.44 |
| 20:CT:82:SER:O | 20:CT:86:ARG:HB2 | 2.17 | 0.44 |
| 24:D2:32:LEU:HA | 24:D2:37:PHE:HB3 | 2.00 | 0.44 |
| 30:D8:26:LYS:HD2 | 30:D8:47:LYS:HZ2 | 1.81 | 0.44 |
| 31:DA:34:C:O2' | 31:DA:35:G:OP1 | 2.33 | 0.44 |
| 31:DA:181:A:H2 | 31:DA:434:U:O4' | 2.01 | 0.44 |
| 31:DA:192:C:C2' | 31:DA:193:U:H5' | 2.48 | 0.44 |
| 31:DA:213:A:H2' | 31:DA:214:G:O4' | 2.18 | 0.44 |
| 31:DA:252:G:P | 41:DP:50:ARG:HH11 | 2.40 | 0.44 |
| 31:DA:649:G:C5 | 31:DA:650:C:C4 | 3.05 | 0.44 |
| 31:DA:675:A:C5 | 31:DA:804:A:C2 | 3.05 | 0.44 |
| 31:DA:684:G:N1 | 31:DA:774:A:C2 | 2.86 | 0.44 |
| 31:DA:705:A:C8 | 31:DA:727:A:C2 | 3.05 | 0.44 |
| 31:DA:828:U:O2 | 31:DA:828:U:C3' | 2.65 | 0.44 |
| 31:DA:943:U:O2' | 31:DA:944:G:H5' | 2.17 | 0.44 |
| 31:DA:1131:G:OP1 | 39:DN:80:GLY:HA2 | 2.16 | 0.44 |
| 31:DA:1323:U:OP1 | 48:DW:98:LYS:NZ | 2.51 | 0.44 |
| 31:DA:1444:G:N2 | 31:DA:1548:C:C2 | 2.86 | 0.44 |
| 31:DA:1529:G:H21 | 31:DA:1530:C:C5' | 2.11 | 0.44 |
| 31:DA:1886:C:H6 | 31:DA:1886:C:O5' | 1.99 | 0.44 |
| 31:DA:1904:G:H2' | 31:DA:1905:C:O5' | 2.18 | 0.44 |
| 31:DA:1985:G:O2' | 31:DA:1986:A:H5' | 2.18 | 0.44 |
| 31:DA:2494:G:H2' | 31:DA:2495:G:O5' | 2.15 | 0.44 |
| 31:DA:2870:C:H2' | 31:DA:2871:C:C5' | 2.46 | 0.44 |
| 32:DB:13:A:O2' | 32:DB:15:A:O5' | 2.34 | 0.44 |
| 33:DD:235:GLY:C | 33:DD:237:GLU:HG2 | 2.39 | 0.44 |
| 33:DD:248:SER:C | 33:DD:250:TRP:H | 2.25 | 0.44 |
| 34:DE:1:MET:HE2 | 34:DE:1:MET:N | 2.33 | 0.44 |
| 34:DE:33:VAL:O | 34:DE:33:VAL:HG13 | 2.17 | 0.44 |
| 34:DE:66:HIS:O | 34:DE:66:HIS:CD2 | 2.70 | 0.44 |
| 34:DE:89:ASP:O | 34:DE:90:THR:CB | 2.65 | 0.44 |
| 35:DF:118:ALA:C | 35:DF:120:GLU:H | 2.25 | 0.44 |
| 41:DP:101:VAL:C | 41:DP:103:ALA:H | 2.25 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 42:DQ:83:MET:O | 42:DQ:83:MET:CG | 2.62 | 0.44 |
| 45:DT:52:ILE:HG22 | 45:DT:53:ARG:N | 2.33 | 0.44 |
| 46:DU:8:VAL:O | 46:DU:9:VAL:C | 2.60 | 0.44 |
| 1:AA:432:A:N7 | 1:AA:433:C:C2 | 2.85 | 0.44 |
| 1:AA:438:G:O2' | 1:AA:493:G:C2 | 2.66 | 0.44 |
| 1:AA:451:A:C5 | 1:AA:481:G:C6 | 3.05 | 0.44 |
| 1:AA:501:C:H2' | 1:AA:502:G:H8 | 1.83 | 0.44 |
| 1:AA:518:C:H4' | 1:AA:519:C:H5" | 1.99 | 0.44 |
| 1:AA:1054:C:H3' | 1:AA:1054:C:O2 | 2.17 | 0.44 |
| 1:AA:1425:U:H2' | 1:AA:1426:C:H6 | 1.83 | 0.44 |
| 1:AA:1429:C:H2' | 1:AA:1430:C:C6 | 2.53 | 0.44 |
| 3:AC:186:PHE:CE2 | 3:AC:188:LEU:HD21 | 2.52 | 0.44 |
| 3:AC:206:GLU:O | 3:AC:207:VAL:C | 2.60 | 0.44 |
| 4:AD:17:VAL:HG12 | 4:AD:18:LYS:N | 2.32 | 0.44 |
| 4:AD:100:ARG:O | 4:AD:101:LEU:C | 2.60 | 0.44 |
| 4:AD:128:VAL:O | 4:AD:129:ASN:C | 2.60 | 0.44 |
| 6:AF:10:LEU:N | 6:AF:10:LEU:CD1 | 2.80 | 0.44 |
| 8:AH:112:LEU:HD12 | 8:AH:114:THR:HG22 | 2.00 | 0.44 |
| 11:AK:124:LYS:HB3 | 11:AK:125:PHE:HD1 | 1.82 | 0.44 |
| 15:AO:32:LEU:O | 15:AO:33:THR:C | 2.61 | 0.44 |
| 23:B1:79:GLY:O | 23:B1:80:LEU:HG | 2.18 | 0.44 |
| 23:B1:91:LYS:O | 23:B1:92:LYS:HD2 | 2.18 | 0.44 |
| 26:B4:11:PRO:C | 26:B4:13:ARG:N | 2.76 | 0.44 |
| 26:B4:19:GLY:O | 26:B4:21:VAL:N | 2.51 | 0.44 |
| 29:B7:34:ARG:NH1 | 31:BA:466:A:OP1 | 2.50 | 0.44 |
| 30:B8:4:MET:O | 30:B8:62:LEU:HD12 | 2.18 | 0.44 |
| 30:B8:25:MET:HE3 | 30:B8:25:MET:HB3 | 1.85 | 0.44 |
| 31:BA:465:G:C6 | 31:BA:466:A:N6 | 2.85 | 0.44 |
| 31:BA:565:C:H5' | 31:BA:1253:A:N6 | 2.33 | 0.44 |
| 31:BA:719:C:H2' | 31:BA:720:C:H6 | 1.82 | 0.44 |
| 31:BA:1037:G:H1 | 31:BA:1118:C:N4 | 2.14 | 0.44 |
| 31:BA:1353:A:H4' | 33:BD:38:LYS:HZ1 | 1.82 | 0.44 |
| 31:BA:1488:G:N2 | 31:BA:1502:C:C6 | 2.85 | 0.44 |
| 31:BA:1504:C:HO2' | 31:BA:1505:C:P | 2.40 | 0.44 |
| 31:BA:1668:A:H4' | 31:BA:1669:A:O5' | 2.17 | 0.44 |
| 31:BA:2779:U:O2 | 31:BA:2779:U:O4' | 2.35 | 0.44 |
| 33:BD:34:VAL:C | 33:BD:35:LYS:O | 2.59 | 0.44 |
| 33:BD:142:VAL:HG21 | 33:BD:191:ALA:CB | 2.47 | 0.44 |
| 35:BF:203:GLN:CA | 35:BF:206:ILE:O | 2.63 | 0.44 |
| 36:BG:96:ARG:O | 36:BG:99:MET:HB3 | 2.17 | 0.44 |
| 39:BN:18:ALA:O | 39:BN:19:GLU:C | 2.60 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 39:BN:82:LEU:H | 39:BN:82:LEU:CD1 | 2.24 | 0.44 |
| 42:BQ:8:LYS:CD | 42:BQ:9:TYR:N | 2.75 | 0.44 |
| 42:BQ:26:TYR:CE1 | 42:BQ:28:ALA:HB2 | 2.52 | 0.44 |
| 42:BQ:82:ARG:HG2 | 42:BQ:82:ARG:NH1 | 2.33 | 0.44 |
| 43:BR:18:LEU:O | 43:BR:19:ALA:C | 2.61 | 0.44 |
| 43:BR:104:ARG:HD2 | 43:BR:111:LEU:HD11 | 2.00 | 0.44 |
| 44:BS:58:LEU:HD21 | 44:BS:68:GLN:CB | 2.47 | 0.44 |
| 44:BS:83:LYS:HE3 | 44:BS:105:ALA:CB | 2.41 | 0.44 |
| 45:BT:16:ARG:HD3 | 45:BT:16:ARG:HA | 1.52 | 0.44 |
| 49:BX:27:THR:OG1 | 49:BX:77:LYS:HA | 2.16 | 0.44 |
| 51:BZ:40:ASP:OD1 | 51:BZ:42:VAL:HG12 | 2.18 | 0.44 |
| 1:CA:66:G:O4' | 1:CA:173:U:C4 | 2.70 | 0.44 |
| 1:CA:101:A:C2' | 1:CA:102:G:H5' | 2.47 | 0.44 |
| 1:CA:522:C:H5" | 12:CL:120:TYR:OH | 2.18 | 0.44 |
| 1:CA:581:G:N2 | 1:CA:582:U:C4 | 2.86 | 0.44 |
| 1:CA:748:C:C4' | 1:CA:749:C:O5' | 2.64 | 0.44 |
| 1:CA:790:A:N6 | 1:CA:791:G:C6 | 2.86 | 0.44 |
| 1:CA:1112:C:N3 | 3:CC:178:LEU:HD23 | 2.33 | 0.44 |
| 1:CA:1184:G:OP1 | 1:CA:1184:G:H8 | 2.00 | 0.44 |
| 1:CA:1206:G:C6 | 1:CA:1207:G:C6 | 3.06 | 0.44 |
| 1:CA:1260:C:H4' | 1:CA:1284:C:H5' | 2.00 | 0.44 |
| 1:CA:1293:G:O2' | 1:CA:1294:G:P | 2.76 | 0.44 |
| 2:CB:79:ASP:C | 2:CB:81:VAL:H | 2.26 | 0.44 |
| 5:CE:139:LEU:C | 5:CE:141:GLN:N | 2.75 | 0.44 |
| 8:CH:31:PHE:O | 8:CH:34:GLU:HB2 | 2.18 | 0.44 |
| 9:CI:13:ALA:HA | 9:CI:67:GLY:O | 2.18 | 0.44 |
| 11:CK:122:LYS:HE2 | 11:CK:122:LYS:HB3 | 1.66 | 0.44 |
| 13:CM:105:THR:O | 13:CM:106:ASN:O | 2.35 | 0.44 |
| 20:CT:81:LYS:O | 20:CT:83:ARG:N | 2.51 | 0.44 |
| 24:D2:52:ASP:O | 24:D2:55:ARG:N | 2.42 | 0.44 |
| 27:D5:4:HIS:O | 27:D5:5:PRO:C | 2.57 | 0.44 |
| 27:D5:51:TYR:HD2 | 27:D5:52:TYR:CZ | 2.33 | 0.44 |
| 31:DA:110:G:C2 | 31:DA:111:A:C8 | 3.05 | 0.44 |
| 31:DA:448:U:H1' | 35:DF:84:VAL:CG1 | 2.47 | 0.44 |
| 31:DA:721:C:O2 | 31:DA:721:C:C2' | 2.60 | 0.44 |
| 31:DA:1262:A:C5 | 31:DA:1263:U:C5 | 3.05 | 0.44 |
| 31:DA:1688:U:H5' | 31:DA:1689:A:OP1 | 2.18 | 0.44 |
| 31:DA:2243:U:H2' | 31:DA:2244:U:C6 | 2.52 | 0.44 |
| 31:DA:2280:G:O2' | 31:DA:2388:A:N1 | 2.39 | 0.44 |
| 31:DA:2295:C:H2' | 31:DA:2295:C:O2 | 2.16 | 0.44 |
| 31:DA:2305:A:O2' | 36:DG:136:ARG:HD3 | 2.17 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2319:G:OP2 | 31:DA:2319:G:H4' | 2.18 | 0.44 |
| 31:DA:2468:G:C5' | 42:DQ:120:ILE:HD12 | 2.48 | 0.44 |
| 31:DA:2598:A:H3' | 33:DD:236:GLY:HA2 | 2.00 | 0.44 |
| 33:DD:34:VAL:O | 33:DD:34:VAL:HG13 | 2.17 | 0.44 |
| 33:DD:62:TYR:CD1 | 33:DD:62:TYR:C | 2.96 | 0.44 |
| 33:DD:228:PRO:HD3 | 33:DD:234:GLY:O | 2.18 | 0.44 |
| 35:DF:53:THR:HG23 | 35:DF:55:GLY:CA | 2.47 | 0.44 |
| 36:DG:7:LEU:HD23 | 36:DG:176:LEU:HD22 | 1.97 | 0.44 |
| 36:DG:89:GLY:O | 36:DG:90:LEU:C | 2.60 | 0.44 |
| 37:DH:70:THR:O | 37:DH:73:ALA:N | 2.50 | 0.44 |
| 38:DI:85:GLU:O | 38:DI:123:LEU:HD12 | 2.16 | 0.44 |
| 39:DN:22:THR:HA | 39:DN:61:ARG:O | 2.17 | 0.44 |
| 43:DR:44:LEU:HD23 | 43:DR:44:LEU:HA | 1.84 | 0.44 |
| 43:DR:111:LEU:HD23 | 43:DR:111:LEU:HA | 1.71 | 0.44 |
| 44:DS:64:GLU:OE2 | 44:DS:64:GLU:N | 2.50 | 0.44 |
| 46:DU:20:LEU:HB3 | 46:DU:39:LEU:HD11 | 1.98 | 0.44 |
| 46:DU:66:ASN:HA | 46:DU:76:TYR:HB2 | 1.99 | 0.44 |
| 46:DU:83:LEU:O | 46:DU:88:ILE:HG13 | 2.17 | 0.44 |
| 50:DY:66:PRO:O | 50:DY:67:LEU:HB3 | 2.18 | 0.44 |
| 1:AA:20:U:O2' | 1:AA:21:G:H5' | 2.17 | 0.44 |
| 1:AA:600:C:O2' | 1:AA:601:C:H5' | 2.17 | 0.44 |
| 1:AA:638:G:O2' | 1:AA:639:G:H5' | 2.17 | 0.44 |
| 1:AA:1142:G:H2' | 1:AA:1143:G:O4' | 2.16 | 0.44 |
| 1:AA:1245:A:N1 | 1:AA:1293:G:C6 | 2.85 | 0.44 |
| 1:AA:1505:G:C4' | 1:AA:1506:U:H5" | 2.41 | 0.44 |
| 7:AG:54:THR:C | 7:AG:56:GLN:H | 2.24 | 0.44 |
| 11:AK:57:THR:HG23 | 11:AK:58:PRO:HD2 | 1.99 | 0.44 |
| 13:AM:20:THR:C | 13:AM:22:ILE:H | 2.26 | 0.44 |
| 16:AP:3:LYS:O | 16:AP:21:VAL:HA | 2.17 | 0.44 |
| 18:AR:44:LEU:HD23 | 18:AR:48:GLY:O | 2.17 | 0.44 |
| 23:B1:28:GLY:C | 23:B1:30:VAL:H | 2.25 | 0.44 |
| 24:B2:33:MET:CG | 49:BX:10:ALA:HB1 | 2.48 | 0.44 |
| 27:B5:41:PRO:HG2 | 27:B5:44:THR:OG1 | 2.17 | 0.44 |
| 31:BA:41:C:H2' | 31:BA:42:G:O4' | 2.18 | 0.44 |
| 31:BA:108:U:H2' | 31:BA:109:G:H8 | 1.83 | 0.44 |
| 31:BA:529:A:N6 | 31:BA:2041:U:C2 | 2.86 | 0.44 |
| 31:BA:1204:A:H2 | 31:BA:1241:A:N1 | 2.16 | 0.44 |
| 31:BA:1244:G:C2' | 31:BA:1245:G:H5' | 2.48 | 0.44 |
| 31:BA:1481:U:H5' | 31:BA:1482:G:P | 2.58 | 0.44 |
| 31:BA:1673:U:C4 | 34:BE:129:HIS:CD2 | 3.06 | 0.44 |
| 31:BA:2243:U:H2' | 31:BA:2244:U:C6 | 2.52 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2877:G:O2' | 31:BA:2878:U:H5' | 2.16 | 0.44 |
| 33:BD:78:LYS:HE3 | 33:BD:78:LYS:HB2 | 1.85 | 0.44 |
| 34:BE:111:ARG:CG | 43:BR:2:ARG:HG3 | 2.46 | 0.44 |
| 36:BG:141:PHE:HE2 | 36:BG:155:MET:HE1 | 1.83 | 0.44 |
| 37:BH:85:LYS:CE | 37:BH:145:ALA:CA | 2.96 | 0.44 |
| 38:BI:25:TYR:CE1 | 38:BI:30:LEU:HD21 | 2.53 | 0.44 |
| 41:BP:146:VAL:CG2 | 41:BP:147:LEU:N | 2.75 | 0.44 |
| 45:BT:11:GLU:CD | 45:BT:11:GLU:H | 2.26 | 0.44 |
| 45:BT:65:LYS:HG3 | 45:BT:66:VAL:H | 1.82 | 0.44 |
| 47:BV:67:GLY:O | 47:BV:69:LYS:N | 2.50 | 0.44 |
| 51:BZ:10:ARG:HH21 | 51:BZ:26:GLY:N | 2.12 | 0.44 |
| 51:BZ:48:PHE:C | 51:BZ:48:PHE:CD1 | 2.96 | 0.44 |
| 1:CA:90:U:H3' | 1:CA:90:U:C6 | 2.53 | 0.44 |
| 1:CA:97:G:O2' | 1:CA:98:G:O5' | 2.34 | 0.44 |
| 1:CA:173:U:O4' | 1:CA:197:A:C4 | 2.71 | 0.44 |
| 1:CA:339:C:OP2 | 40:DO:97:ARG:NH1 | 2.51 | 0.44 |
| 1:CA:366:C:C1' | 1:CA:367:U:OP1 | 2.66 | 0.44 |
| 1:CA:445:G:C6 | 1:CA:490:G:C6 | 3.06 | 0.44 |
| 1:CA:955:U:H1' | 1:CA:1227:A:H61 | 1.81 | 0.44 |
| 1:CA:1096:C:H2' | 1:CA:1097:C:C6 | 2.48 | 0.44 |
| 1:CA:1191:A:H5" | 3:CC:4:LYS:HZ2 | 1.83 | 0.44 |
| 4:CD:108:LEU:CD1 | 4:CD:174:LEU:HD13 | 2.48 | 0.44 |
| 5:CE:100:VAL:O | 5:CE:107:ARG:NH2 | 2.41 | 0.44 |
| 6:CF:82:ARG:HB2 | 6:CF:85:VAL:HG23 | 1.99 | 0.44 |
| 7:CG:145:ALA:O | 7:CG:146:GLU:HB2 | 2.17 | 0.44 |
| 10:CJ:22:LYS:C | 10:CJ:24:VAL:H | 2.26 | 0.44 |
| 15:CO:55:GLY:O | 15:CO:59:MET:HG3 | 2.17 | 0.44 |
| 20:CT:84:LEU:O | 20:CT:88:VAL:HG23 | 2.17 | 0.44 |
| 28:D6:15:GLU:OE1 | 28:D6:18:ARG:HD2 | 2.17 | 0.44 |
| 29:D7:32:LYS:HE3 | 31:DA:180:G:OP1 | 2.18 | 0.44 |
| 31:DA:389:G:H1 | 41:DP:71:VAL:H | 1.66 | 0.44 |
| 31:DA:639:U:O2' | 31:DA:640:C:H5' | 2.17 | 0.44 |
| 31:DA:1327:C:O2' | 31:DA:1328:G:H5' | 2.18 | 0.44 |
| 31:DA:1844:C:OP1 | 33:DD:257:LEU:HD23 | 2.18 | 0.44 |
| 31:DA:2203:U:O2' | 33:DD:151:LYS:HG2 | 2.17 | 0.44 |
| 31:DA:2287:A:H2 | 31:DA:2346:A:C2 | 2.35 | 0.44 |
| 31:DA:2464:C:O2' | 31:DA:2465:C:O5' | 2.36 | 0.44 |
| 31:DA:2649:U:C2 | 31:DA:2672:G:N2 | 2.85 | 0.44 |
| 32:DB:58:A:H5' | 32:DB:59:A:OP2 | 2.18 | 0.44 |
| 35:DF:25:PRO:HB3 | 35:DF:119:ARG:HG3 | 1.99 | 0.44 |
| 38:DI:88:ILE:HD11 | 38:DI:122:GLU:C | 2.42 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 39:DN:78:TYR:CD1 | 39:DN:78:TYR:C | 2.94 | 0.44 |
| 39:DN:91:LEU:O | 39:DN:95:PRO:HB3 | 2.18 | 0.44 |
| 41:DP:21:ARG:O | 41:DP:21:ARG:HG2 | 2.17 | 0.44 |
| 42:DQ:34:LEU:HD11 | 42:DQ:129:THR:CB | 2.46 | 0.44 |
| 43:DR:59:ASP:OD1 | 43:DR:61:HIS:HB3 | 2.17 | 0.44 |
| 45:DT:11:GLU:CD | 45:DT:11:GLU:H | 2.26 | 0.44 |
| 45:DT:61:PHE:CZ | 45:DT:76:PHE:HB2 | 2.53 | 0.44 |
| 46:DU:66:ASN:HB2 | 46:DU:76:TYR:HB2 | 1.99 | 0.44 |
| 46:DU:92:ARG:NH2 | 47:DV:10:LYS:HG2 | 2.33 | 0.44 |
| 46:DU:92:ARG:HH22 | 47:DV:10:LYS:HB3 | 1.83 | 0.44 |
| 47:DV:54:GLY:O | 47:DV:56:SER:N | 2.42 | 0.44 |
| 49:DX:7:VAL:HG12 | 49:DX:30:VAL:HG12 | 1.99 | 0.44 |
| 50:DY:83:THR:HG21 | 50:DY:94:LYS:HD2 | 1.98 | 0.44 |
| 1:AA:174:C:O5’ | 1:AA:174:C:H6 | 2.01 | 0.44 |
| 1:AA:346:G:H5” | 45:BT:41:ARG:HE | 1.83 | 0.44 |
| 1:AA:380:G:N1 | 1:AA:384:G:C6 | 2.86 | 0.44 |
| 1:AA:413:G:N2 | 1:AA:429:U:OP2 | 2.47 | 0.44 |
| 1:AA:525:C:H2’ | 1:AA:526:C:H6 | 1.79 | 0.44 |
| 1:AA:582:U:C2 | 1:AA:760:G:C6 | 3.06 | 0.44 |
| 1:AA:657:G:C2 | 1:AA:750:G:C5 | 3.05 | 0.44 |
| 1:AA:702:A:H3’ | 1:AA:703:G:H5’ | 1.99 | 0.44 |
| 1:AA:1100:C:OP2 | 2:AB:96:ARG:HG2 | 2.18 | 0.44 |
| 1:AA:1497:G:C2’ | 1:AA:1498:U:H5’ | 2.44 | 0.44 |
| 1:AA:1498:U:H1’ | 1:AA:1499:A:OP2 | 2.18 | 0.44 |
| 3:AC:124:ILE:HD11 | 3:AC:130:VAL:HG13 | 1.99 | 0.44 |
| 4:AD:108:LEU:CD1 | 4:AD:174:LEU:HD13 | 2.48 | 0.44 |
| 5:AE:81:GLU:HG2 | 5:AE:90:VAL:HG22 | 2.00 | 0.44 |
| 12:AL:91:LYS:O | 12:AL:91:LYS:CG | 2.66 | 0.44 |
| 13:AM:83:ASP:CG | 13:AM:84:ILE:N | 2.75 | 0.44 |
| 16:AP:81:ARG:HD3 | 16:AP:83:GLU:OE2 | 2.17 | 0.44 |
| 21:AU:21:TYR:O | 21:AU:22:ARG:HB2 | 2.18 | 0.44 |
| 22:B0:82:ARG:HA | 22:B0:83:PRO:HD2 | 1.86 | 0.44 |
| 27:B5:40:LYS:NZ | 27:B5:46:CYS:O | 2.49 | 0.44 |
| 28:B6:19:ARG:HD2 | 28:B6:43:CYS:SG | 2.57 | 0.44 |
| 30:B8:61:LEU:HD13 | 31:BA:593:G:H4’ | 2.00 | 0.44 |
| 31:BA:118:A:N3 | 31:BA:178:G:H1’ | 2.33 | 0.44 |
| 31:BA:142:A:C8 | 31:BA:1595:G:N2 | 2.66 | 0.44 |
| 31:BA:299:A:H5” | 50:BY:84:ARG:HH21 | 1.83 | 0.44 |
| 31:BA:385:C:C2’ | 31:BA:386:G:OP2 | 2.65 | 0.44 |
| 31:BA:540:C:H2’ | 31:BA:541:C:O5’ | 2.18 | 0.44 |
| 31:BA:838:C:H2’ | 31:BA:839:U:H6 | 1.83 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1410:G:C4 | 31:BA:1411:C:C5 | 3.06 | 0.44 |
| 31:BA:1439:A:C2 | 31:BA:1553:A:C4 | 3.06 | 0.44 |
| 31:BA:1461:G:C2' | 31:BA:1462:C:H5' | 2.48 | 0.44 |
| 31:BA:1496:A:H8 | 31:BA:1577:C:O2' | 1.99 | 0.44 |
| 31:BA:2050:C:H1' | 34:BE:156:MET:HE2 | 2.00 | 0.44 |
| 31:BA:2473:U:O2 | 31:BA:2473:U:H2' | 2.17 | 0.44 |
| 32:BB:55:U:O2 | 32:BB:56:G:C8 | 2.71 | 0.44 |
| 36:BG:51:ARG:HD3 | 36:BG:53:LEU:HD21 | 2.00 | 0.44 |
| 37:BH:83:TYR:HB3 | 37:BH:134:SER:HA | 1.98 | 0.44 |
| 42:BQ:20:ALA:HB2 | 42:BQ:99:PRO:CG | 2.48 | 0.44 |
| 43:BR:29:LEU:HD12 | 43:BR:29:LEU:HA | 1.86 | 0.44 |
| 44:BS:64:GLU:OE2 | 44:BS:64:GLU:N | 2.51 | 0.44 |
| 45:BT:78:LEU:HD22 | 45:BT:79:HIS:CE1 | 2.52 | 0.44 |
| 46:BU:50:ARG:NH2 | 47:BV:75:PHE:CD2 | 2.86 | 0.44 |
| 51:BZ:119:GLU:C | 51:BZ:121:HIS:N | 2.76 | 0.44 |
| 51:BZ:157:LEU:HA | 51:BZ:158:PRO:HD2 | 1.90 | 0.44 |
| 1:CA:67:C:H2' | 1:CA:68:G:H8 | 1.82 | 0.44 |
| 1:CA:976:G:OP1 | 14:CN:32:SER:N | 2.41 | 0.44 |
| 1:CA:1072:G:C6 | 1:CA:1073:U:O4 | 2.70 | 0.44 |
| 1:CA:1442:G:C8 | 1:CA:1442(B):A:H2 | 2.35 | 0.44 |
| 1:CA:1510:U:H2' | 1:CA:1511:G:C8 | 2.52 | 0.44 |
| 1:CA:1522:U:H2' | 1:CA:1523:G:H8 | 1.83 | 0.44 |
| 2:CB:185:ILE:CG2 | 2:CB:199:TYR:HB2 | 2.23 | 0.44 |
| 5:CE:120:THR:O | 5:CE:121:LYS:HB2 | 2.18 | 0.44 |
| 8:CH:30:ARG:HB3 | 8:CH:30:ARG:NH1 | 2.33 | 0.44 |
| 11:CK:96:ARG:O | 11:CK:99:GLN:HG2 | 2.17 | 0.44 |
| 23:D1:48:LYS:HD3 | 23:D1:48:LYS:HA | 1.60 | 0.44 |
| 23:D1:78:LYS:NZ | 23:D1:93:GLU:HB2 | 2.33 | 0.44 |
| 30:D8:35:GLN:HG2 | 31:DA:2420:C:P | 2.58 | 0.44 |
| 31:DA:17:G:H4' | 46:DU:25:TRP:CZ2 | 2.52 | 0.44 |
| 31:DA:73:A:H2' | 31:DA:74:A:OP2 | 2.18 | 0.44 |
| 31:DA:198:C:H5' | 31:DA:2244:U:OP1 | 2.17 | 0.44 |
| 31:DA:216:A:C4 | 31:DA:432:A:C2 | 3.05 | 0.44 |
| 31:DA:1147:C:C2' | 31:DA:1148:A:O5' | 2.66 | 0.44 |
| 31:DA:1176:G:C4' | 31:DA:1177:A:OP1 | 2.66 | 0.44 |
| 31:DA:1210:A:H5" | 31:DA:1211:U:C3' | 2.46 | 0.44 |
| 31:DA:1340:U:OP2 | 49:DX:76:ARG:NH2 | 2.51 | 0.44 |
| 31:DA:2082:A:H2' | 31:DA:2083:G:O4' | 2.17 | 0.44 |
| 31:DA:2646:C:O5' | 31:DA:2646:C:H6 | 2.01 | 0.44 |
| 32:DB:33:G:N2 | 32:DB:50:G:C4 | 2.86 | 0.44 |
| 32:DB:117:G:H4' | 44:DS:55:ALA:HB1 | 1.98 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 34:DE:75:VAL:C | 34:DE:77:ILE:H | 2.25 | 0.44 |
| 37:DH:97:ARG:O | 37:DH:98:LEU:C | 2.61 | 0.44 |
| 37:DH:154:PRO:O | 37:DH:156:ALA:N | 2.40 | 0.44 |
| 38:DI:72:LEU:HD12 | 38:DI:138:ILE:HG23 | 2.00 | 0.44 |
| 39:DN:72:TYR:CE1 | 39:DN:87:LEU:HD23 | 2.53 | 0.44 |
| 43:DR:12:ARG:HB3 | 43:DR:16:HIS:HB3 | 2.00 | 0.44 |
| 45:DT:34:VAL:HG13 | 45:DT:39:ARG:HA | 1.99 | 0.44 |
| 46:DU:108:GLU:O | 46:DU:109:LEU:C | 2.59 | 0.44 |
| 49:DX:35:THR:O | 49:DX:39:ILE:HG23 | 2.18 | 0.44 |
| 49:DX:82:GLN:CG | 49:DX:83:VAL:N | 2.80 | 0.44 |
| 50:DY:32:PRO:C | 50:DY:34:LYS:N | 2.75 | 0.44 |
| 1:AA:173:U:O4’ | 1:AA:197:A:C4 | 2.71 | 0.44 |
| 1:AA:373:A:C4 | 1:AA:482:A:N7 | 2.85 | 0.44 |
| 1:AA:543:C:N3 | 1:AA:544:G:N7 | 2.66 | 0.44 |
| 1:AA:626:U:H4’ | 16:AP:38:TYR:CZ | 2.52 | 0.44 |
| 1:AA:859:A:C8 | 1:AA:860:A:C8 | 3.05 | 0.44 |
| 1:AA:1071:C:H5” | 5:AE:49:PRO:HG3 | 1.99 | 0.44 |
| 2:AB:91:PRO:N | 2:AB:154:LEU:HD12 | 2.33 | 0.44 |
| 2:AB:118:LEU:HD11 | 2:AB:141:GLU:HG2 | 2.00 | 0.44 |
| 3:AC:138:VAL:CG2 | 3:AC:151:VAL:HG23 | 2.47 | 0.44 |
| 4:AD:2:GLY:O | 4:AD:4:TYR:N | 2.50 | 0.44 |
| 7:AG:145:ALA:O | 7:AG:146:GLU:HB2 | 2.18 | 0.44 |
| 8:AH:64:LYS:C | 8:AH:65:TYR:CD1 | 2.96 | 0.44 |
| 9:AI:126:SER:C | 9:AI:128:ARG:H | 2.25 | 0.44 |
| 11:AK:62:GLN:C | 11:AK:64:ALA:N | 2.75 | 0.44 |
| 12:AL:38:THR:HG21 | 12:AL:65:GLU:OE2 | 2.18 | 0.44 |
| 15:AO:4:THR:HG1 | 15:AO:7:GLU:HB2 | 1.81 | 0.44 |
| 16:AP:58:TYR:C | 16:AP:60:LEU:N | 2.75 | 0.44 |
| 19:AS:10:PHE:HZ | 19:AS:70:LYS:HE2 | 1.83 | 0.44 |
| 31:BA:1839:G:H2’ | 31:BA:1839:G:N3 | 2.31 | 0.44 |
| 31:BA:2080:G:N2 | 31:BA:2241:A:C4 | 2.86 | 0.44 |
| 31:BA:2248:C:O2 | 31:BA:2248:C:H2’ | 2.18 | 0.44 |
| 31:BA:2432:A:H2’ | 31:BA:2433:A:C8 | 2.53 | 0.44 |
| 31:BA:2680:C:OP2 | 34:BE:111:ARG:NH2 | 2.51 | 0.44 |
| 31:BA:2747:G:C6 | 31:BA:2754:U:C5 | 3.06 | 0.44 |
| 32:BB:2:C:C2 | 32:BB:3:C:C5 | 3.06 | 0.44 |
| 32:BB:115:G:C3’ | 32:BB:116:G:H5” | 2.48 | 0.44 |
| 33:BD:33:LEU:HB2 | 33:BD:34:VAL:H | 1.70 | 0.44 |
| 36:BG:153:ARG:CZ | 36:BG:153:ARG:HB3 | 2.48 | 0.44 |
| 40:BO:111:PHE:O | 40:BO:112:MET:C | 2.61 | 0.44 |
| 41:BP:103:ALA:O | 41:BP:104:GLY:C | 2.60 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 45:BT:92:GLY:O | 45:BT:94:ALA:N | 2.50 | 0.44 |
| 46:BU:66:ASN:C | 46:BU:66:ASN:HD22 | 2.25 | 0.44 |
| 46:BU:92:ARG:NH2 | 47:BV:10:LYS:CB | 2.80 | 0.44 |
| 47:BV:97:LYS:O | 47:BV:97:LYS:HG3 | 2.18 | 0.44 |
| 49:BX:57:LEU:HD11 | 49:BX:77:LYS:HD2 | 1.99 | 0.44 |
| 49:BX:90:GLU:C | 49:BX:92:LEU:N | 2.75 | 0.44 |
| 1:CA:6:G:H4' | 1:CA:298:A:H4' | 2.00 | 0.44 |
| 1:CA:112:G:C2 | 1:CA:113:G:C8 | 3.05 | 0.44 |
| 1:CA:397:A:H5" | 1:CA:397:A:N3 | 2.33 | 0.44 |
| 1:CA:503:C:C6 | 1:CA:504:C:H5 | 2.36 | 0.44 |
| 1:CA:520:A:H2 | 1:CA:536:C:O2 | 2.01 | 0.44 |
| 1:CA:590:C:O2' | 1:CA:591:U:H5' | 2.17 | 0.44 |
| 1:CA:603:U:O2' | 1:CA:604:G:H5' | 2.18 | 0.44 |
| 1:CA:663:A:O2' | 1:CA:664:G:H5' | 2.17 | 0.44 |
| 1:CA:1422:G:O2' | 1:CA:1423:G:H5' | 2.17 | 0.44 |
| 1:CA:1465:C:C4 | 1:CA:1466:C:C4 | 3.06 | 0.44 |
| 3:CC:152:ILE:HB | 3:CC:199:LYS:HB2 | 1.99 | 0.44 |
| 4:CD:43:HIS:HA | 4:CD:46:LYS:HE3 | 1.99 | 0.44 |
| 4:CD:188:LEU:HA | 4:CD:189:PRO:HD2 | 1.88 | 0.44 |
| 6:CF:91:VAL:HG11 | 18:CR:72:ARG:NH1 | 2.33 | 0.44 |
| 8:CH:10:LEU:HD23 | 8:CH:10:LEU:H | 1.78 | 0.44 |
| 8:CH:64:LYS:C | 8:CH:65:TYR:CD1 | 2.96 | 0.44 |
| 12:CL:69:TYR:HB3 | 12:CL:99:HIS:CD2 | 2.52 | 0.44 |
| 17:CQ:3:LYS:HB2 | 17:CQ:60:ILE:HD11 | 2.00 | 0.44 |
| 22:D0:77:ARG:NH2 | 31:DA:857:C:OP2 | 2.48 | 0.44 |
| 23:D1:77:ALA:C | 23:D1:79:GLY:H | 2.25 | 0.44 |
| 30:D8:58:ILE:HD13 | 30:D8:58:ILE:HA | 1.80 | 0.44 |
| 31:DA:90:U:C1' | 31:DA:92:A:H5" | 2.43 | 0.44 |
| 31:DA:183:C:H2' | 31:DA:184:C:H5' | 1.99 | 0.44 |
| 31:DA:272(B):G:C2' | 31:DA:272(C):G:O5' | 2.66 | 0.44 |
| 31:DA:352:G:HO2' | 31:DA:353:G:P | 2.40 | 0.44 |
| 31:DA:356:G:C2 | 31:DA:357:A:C4 | 3.06 | 0.44 |
| 31:DA:536:A:C2' | 31:DA:537:C:O5' | 2.66 | 0.44 |
| 31:DA:554:U:O2' | 31:DA:555:U:H5' | 2.17 | 0.44 |
| 31:DA:626:U:H5" | 31:DA:627:A:C5' | 2.48 | 0.44 |
| 31:DA:691:C:O2' | 31:DA:692:C:H5' | 2.17 | 0.44 |
| 31:DA:774:A:H2 | 31:DA:787:U:O2' | 1.98 | 0.44 |
| 31:DA:942:G:C6 | 31:DA:943:U:C4 | 3.06 | 0.44 |
| 31:DA:958:U:H5" | 42:DQ:14:ARG:HD3 | 1.99 | 0.44 |
| 31:DA:1050:A:C2 | 31:DA:2751:G:C5 | 3.06 | 0.44 |
| 31:DA:1408:C:C2 | 31:DA:1595:G:N2 | 2.86 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1779:U:C2 | 31:DA:1783:A:N7 | 2.86 | 0.44 |
| 31:DA:1839:G:C8 | 31:DA:1927:A:C1' | 3.01 | 0.44 |
| 31:DA:1999:C:OP1 | 31:DA:2723:C:O2' | 2.35 | 0.44 |
| 31:DA:2064:C:H2' | 31:DA:2065:C:C6 | 2.52 | 0.44 |
| 31:DA:2377:A:H2' | 31:DA:2378:A:C8 | 2.53 | 0.44 |
| 32:DB:14:U:O3' | 32:DB:108:U:O2' | 2.33 | 0.44 |
| 33:DD:25:THR:CG2 | 33:DD:82:ILE:H | 2.30 | 0.44 |
| 33:DD:28:GLU:CB | 33:DD:29:PRO:CD | 2.96 | 0.44 |
| 34:DE:1:MET:O | 34:DE:2:LYS:C | 2.60 | 0.44 |
| 34:DE:98:PRO:HD3 | 34:DE:175:VAL:CG1 | 2.48 | 0.44 |
| 35:DF:83:PHE:C | 35:DF:84:VAL:HG23 | 2.43 | 0.44 |
| 35:DF:170:LEU:HD22 | 35:DF:172:TRP:CZ2 | 2.53 | 0.44 |
| 36:DG:10:LYS:O | 36:DG:15:VAL:HG23 | 2.17 | 0.44 |
| 36:DG:29:TRP:C | 36:DG:31:VAL:H | 2.26 | 0.44 |
| 36:DG:101:ILE:HG23 | 36:DG:102:PHE:N | 2.32 | 0.44 |
| 36:DG:116:ASP:O | 36:DG:117:PHE:HB3 | 2.18 | 0.44 |
| 38:DI:90:GLY:O | 38:DI:91:SER:C | 2.59 | 0.44 |
| 42:DQ:20:ALA:HB2 | 42:DQ:99:PRO:CG | 2.47 | 0.44 |
| 44:DS:53:SER:O | 44:DS:56:LEU:N | 2.42 | 0.44 |
| 46:DU:12:ARG:O | 46:DU:13:LYS:C | 2.58 | 0.44 |
| 46:DU:92:ARG:O | 46:DU:94:ASN:N | 2.51 | 0.44 |
| 47:DV:18:LEU:O | 47:DV:97:LYS:HA | 2.17 | 0.44 |
| 48:DW:84:ARG:HG3 | 48:DW:98:LYS:HZ2 | 1.82 | 0.44 |
| 51:DZ:125:LEU:HD23 | 51:DZ:126:VAL:N | 2.33 | 0.44 |
| 1:AA:567:G:C2 | 1:AA:568:G:H1' | 2.53 | 0.43 |
| 1:AA:748:C:C4' | 1:AA:749:C:O5' | 2.65 | 0.43 |
| 1:AA:938:A:C6 | 1:AA:939:G:C5 | 3.06 | 0.43 |
| 1:AA:997:U:H2' | 1:AA:998:G:C8 | 2.53 | 0.43 |
| 1:AA:1504:G:H4' | 1:AA:1505:G:O4' | 2.18 | 0.43 |
| 7:AG:16:LEU:HD12 | 9:AI:41:VAL:O | 2.17 | 0.43 |
| 10:AJ:51:ARG:HG3 | 10:AJ:60:ARG:C | 2.43 | 0.43 |
| 10:AJ:54:PHE:CE1 | 10:AJ:55:LYS:HE3 | 2.53 | 0.43 |
| 12:AL:119:LYS:C | 12:AL:120:TYR:CD1 | 2.96 | 0.43 |
| 19:AS:6:LYS:H | 19:AS:6:LYS:HD2 | 1.83 | 0.43 |
| 20:AT:46:GLU:HG2 | 20:AT:48:LYS:HE2 | 2.00 | 0.43 |
| 20:AT:84:LEU:HD13 | 20:AT:84:LEU:C | 2.43 | 0.43 |
| 22:B0:53:MET:HA | 22:B0:58:THR:O | 2.18 | 0.43 |
| 27:B5:57:VAL:HB | 27:B5:58:LEU:CD1 | 2.46 | 0.43 |
| 28:B6:16:CYS:O | 28:B6:18:ARG:NH1 | 2.51 | 0.43 |
| 30:B8:38:GLY:O | 30:B8:39:LYS:HB3 | 2.18 | 0.43 |
| 31:BA:330:A:H2 | 31:BA:1210:A:C2' | 2.30 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:626:U:H5" | 31:BA:627:A:C5' | 2.48 | 0.43 |
| 31:BA:1377:G:H8 | 31:BA:1377:G:O5' | 2.00 | 0.43 |
| 31:BA:1940:U:C4 | 31:BA:1964:G:H4' | 2.53 | 0.43 |
| 31:BA:2558:C:C4 | 31:BA:2559:C:C5 | 3.06 | 0.43 |
| 31:BA:2774:C:H2' | 31:BA:2775:A:O4' | 2.18 | 0.43 |
| 33:BD:31:LYS:C | 33:BD:32:SER:O | 2.60 | 0.43 |
| 33:BD:155:LEU:O | 33:BD:156:ALA:C | 2.61 | 0.43 |
| 34:BE:10:GLY:HA2 | 34:BE:192:ASN:ND2 | 2.33 | 0.43 |
| 34:BE:92:THR:O | 34:BE:93:VAL:HG23 | 2.17 | 0.43 |
| 36:BG:11:TYR:HD2 | 36:BG:12:TYR:CE1 | 2.36 | 0.43 |
| 41:BP:122:PRO:HB3 | 25:D3:1:MET:CE | 2.48 | 0.43 |
| 42:BQ:38:GLU:HB2 | 42:BQ:127:ILE:CG2 | 2.48 | 0.43 |
| 44:BS:19:LYS:O | 44:BS:19:LYS:CG | 2.66 | 0.43 |
| 45:BT:117:ASP:O | 45:BT:118:ARG:C | 2.61 | 0.43 |
| 47:BV:18:LEU:O | 47:BV:97:LYS:HA | 2.18 | 0.43 |
| 47:BV:25:LEU:CB | 47:BV:94:LEU:HD13 | 2.48 | 0.43 |
| 47:BV:91:TYR:C | 47:BV:91:TYR:CD2 | 2.96 | 0.43 |
| 49:BX:80:ILE:HG23 | 49:BX:81:VAL:N | 2.31 | 0.43 |
| 51:BZ:45:ASP:O | 51:BZ:46:LYS:C | 2.59 | 0.43 |
| 1:CA:101:A:H2' | 1:CA:102:G:H5' | 2.00 | 0.43 |
| 1:CA:437:U:H2' | 1:CA:438:G:O4' | 2.18 | 0.43 |
| 1:CA:541:G:H2' | 1:CA:542:G:C8 | 2.52 | 0.43 |
| 1:CA:706:A:C5 | 1:CA:707:C:H5 | 2.36 | 0.43 |
| 1:CA:937:A:H1' | 1:CA:1379:G:N2 | 2.33 | 0.43 |
| 1:CA:997:U:H2' | 1:CA:998:G:C8 | 2.52 | 0.43 |
| 1:CA:1054:C:O2 | 1:CA:1054:C:H3' | 2.18 | 0.43 |
| 1:CA:1057:G:C5 | 1:CA:1204:A:C2 | 3.05 | 0.43 |
| 1:CA:1483:A:H8 | 1:CA:1483:A:O5' | 2.01 | 0.43 |
| 3:CC:58:GLU:H | 3:CC:65:ALA:CB | 2.26 | 0.43 |
| 3:CC:138:VAL:CG2 | 3:CC:151:VAL:HG23 | 2.48 | 0.43 |
| 4:CD:17:VAL:HG12 | 4:CD:18:LYS:N | 2.32 | 0.43 |
| 5:CE:47:LYS:HD3 | 5:CE:47:LYS:N | 2.33 | 0.43 |
| 13:CM:3:ARG:HG2 | 13:CM:9:ILE:HD11 | 1.99 | 0.43 |
| 15:CO:74:ASP:OD2 | 15:CO:77:ARG:HG2 | 2.17 | 0.43 |
| 17:CQ:99:SER:O | 17:CQ:100:LYS:HD3 | 2.18 | 0.43 |
| 22:D0:41:ARG:H | 22:D0:41:ARG:CD | 2.19 | 0.43 |
| 23:D1:48:LYS:O | 23:D1:49:VAL:HB | 2.18 | 0.43 |
| 30:D8:35:GLN:HE21 | 30:D8:35:GLN:HB3 | 1.53 | 0.43 |
| 31:DA:950:G:C6 | 31:DA:951:C:N3 | 2.86 | 0.43 |
| 31:DA:1207:C:H2' | 31:DA:1208:C:H6 | 1.82 | 0.43 |
| 31:DA:1651:G:OP1 | 43:DR:40:LYS:NZ | 2.50 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1683:C:H2' | 31:DA:1684:C:C6 | 2.52 | 0.43 |
| 31:DA:1803:A:H2 | 31:DA:1822:G:N3 | 2.15 | 0.43 |
| 31:DA:1895:C:C2 | 31:DA:1896:G:C8 | 3.06 | 0.43 |
| 31:DA:1970:A:H5" | 31:DA:1971:A:OP1 | 2.18 | 0.43 |
| 31:DA:2038:G:H2' | 31:DA:2039:C:O4' | 2.18 | 0.43 |
| 31:DA:2408:U:O2' | 31:DA:2409:G:H5' | 2.18 | 0.43 |
| 31:DA:2683:C:O2 | 40:DO:70:LYS:NZ | 2.49 | 0.43 |
| 31:DA:2687:U:C4 | 31:DA:2688:U:C5 | 3.06 | 0.43 |
| 33:DD:197:GLY:O | 33:DD:198:ASN:HB3 | 2.17 | 0.43 |
| 33:DD:246:PRO:HB2 | 33:DD:255:LYS:HG3 | 1.99 | 0.43 |
| 34:DE:12:THR:HG22 | 45:DT:58:ASN:OD1 | 2.18 | 0.43 |
| 34:DE:93:VAL:C | 34:DE:95:ILE:N | 2.75 | 0.43 |
| 34:DE:200:GLU:OE2 | 34:DE:200:GLU:N | 2.50 | 0.43 |
| 35:DF:33:LEU:O | 35:DF:37:VAL:HG23 | 2.17 | 0.43 |
| 36:DG:39:ILE:HG23 | 36:DG:92:VAL:CG1 | 2.48 | 0.43 |
| 36:DG:96:ARG:HB2 | 36:DG:97:ASP:H | 1.62 | 0.43 |
| 37:DH:152:ARG:HD2 | 37:DH:152:ARG:HA | 1.70 | 0.43 |
| 42:DQ:6:ARG:HD3 | 42:DQ:6:ARG:HA | 1.61 | 0.43 |
| 42:DQ:77:LYS:NZ | 42:DQ:84:GLY:O | 2.51 | 0.43 |
| 42:DQ:114:ALA:O | 42:DQ:115:MET:C | 2.61 | 0.43 |
| 46:DU:39:LEU:HD23 | 46:DU:39:LEU:HA | 1.72 | 0.43 |
| 46:DU:80:ILE:O | 46:DU:83:LEU:HB2 | 2.18 | 0.43 |
| 1:AA:585:G:N3 | 1:AA:879:C:H4' | 2.33 | 0.43 |
| 1:AA:1118:C:H42 | 1:AA:1155:G:H1 | 1.66 | 0.43 |
| 2:AB:74:LYS:NZ | 2:AB:76:GLN:HB2 | 2.33 | 0.43 |
| 3:AC:125:GLU:OE2 | 3:AC:189:ALA:HA | 2.18 | 0.43 |
| 4:AD:92:VAL:O | 4:AD:96:LEU:HD22 | 2.17 | 0.43 |
| 4:AD:190:ASP:O | 4:AD:191:ARG:C | 2.62 | 0.43 |
| 4:AD:192:GLU:CD | 4:AD:192:GLU:H | 2.26 | 0.43 |
| 6:AF:75:LEU:CD2 | 6:AF:79:LEU:HD21 | 2.48 | 0.43 |
| 11:AK:19:ALA:HA | 11:AK:32:ILE:HA | 2.00 | 0.43 |
| 12:AL:32:PHE:CB | 12:AL:84:LEU:HD21 | 2.48 | 0.43 |
| 12:AL:64:TYR:HB3 | 12:AL:65:GLU:H | 1.63 | 0.43 |
| 15:AO:9:GLN:O | 15:AO:10:LYS:C | 2.59 | 0.43 |
| 17:AQ:3:LYS:HB2 | 17:AQ:60:ILE:HD11 | 1.99 | 0.43 |
| 23:B1:77:ALA:C | 23:B1:79:GLY:H | 2.25 | 0.43 |
| 24:B2:23:LYS:O | 24:B2:27:GLU:HB2 | 2.18 | 0.43 |
| 27:B5:2:ALA:N | 31:BA:747:U:C2 | 2.86 | 0.43 |
| 28:B6:40:CYS:SG | 28:B6:45:LYS:HD2 | 2.58 | 0.43 |
| 28:B6:46:HIS:ND1 | 31:BA:2371:G:O2' | 2.51 | 0.43 |
| 30:B8:62:LEU:H | 30:B8:62:LEU:HG | 1.55 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:271(S):G:H2' | 31:BA:271(T):C:O4' | 2.18 | 0.43 |
| 31:BA:272(B):G:O2' | 31:BA:272(C):G:O4' | 2.28 | 0.43 |
| 31:BA:797:C:OP2 | 35:BF:62:ARG:HG3 | 2.17 | 0.43 |
| 31:BA:2416:C:OP1 | 41:BP:64:LYS:O | 2.36 | 0.43 |
| 31:BA:2562:U:C2' | 31:BA:2563:U:H5' | 2.48 | 0.43 |
| 33:BD:174:ILE:O | 33:BD:174:ILE:HG22 | 2.18 | 0.43 |
| 33:BD:206:LEU:HD23 | 33:BD:206:LEU:HA | 1.43 | 0.43 |
| 34:BE:11:MET:HB3 | 34:BE:24:THR:HA | 2.00 | 0.43 |
| 35:BF:8:GLN:CB | 35:BF:126:VAL:HA | 2.44 | 0.43 |
| 37:BH:149:ARG:O | 37:BH:152:ARG:O | 2.36 | 0.43 |
| 38:BI:5:LEU:HA | 38:BI:36:ALA:HB2 | 2.00 | 0.43 |
| 38:BI:73:GLU:O | 38:BI:73:GLU:HG3 | 2.18 | 0.43 |
| 42:BQ:16:ARG:NH1 | 42:BQ:16:ARG:CB | 2.81 | 0.43 |
| 44:BS:66:ALA:HA | 44:BS:69:VAL:CG1 | 2.48 | 0.43 |
| 48:BW:8:ARG:HB3 | 48:BW:9:TYR:CD1 | 2.53 | 0.43 |
| 49:BX:57:LEU:HD12 | 49:BX:76:ARG:HG3 | 2.00 | 0.43 |
| 50:BY:27:VAL:CG1 | 50:BY:29:GLU:OE1 | 2.66 | 0.43 |
| 50:BY:97:ARG:O | 50:BY:98:VAL:C | 2.59 | 0.43 |
| 51:BZ:15:PRO:O | 51:BZ:19:ARG:HD2 | 2.18 | 0.43 |
| 1:CA:92:C:H2' | 1:CA:93:G:C8 | 2.54 | 0.43 |
| 1:CA:124:G:C5 | 1:CA:125:U:C4 | 3.06 | 0.43 |
| 1:CA:509:A:O2' | 1:CA:510:A:C5' | 2.66 | 0.43 |
| 1:CA:750:G:C2 | 1:CA:751:U:C6 | 3.06 | 0.43 |
| 1:CA:986:A:N1 | 1:CA:1220:G:C2 | 2.86 | 0.43 |
| 1:CA:1125:U:C2' | 1:CA:1126:U:OP2 | 2.65 | 0.43 |
| 1:CA:1305:G:C5' | 21:CU:4:GLY:HA3 | 2.48 | 0.43 |
| 1:CA:1316:G:H1 | 19:CS:5:LEU:HD21 | 1.84 | 0.43 |
| 1:CA:1429:C:H2' | 1:CA:1430:C:H6 | 1.82 | 0.43 |
| 2:CB:21:ARG:HB2 | 2:CB:39:ILE:HA | 2.00 | 0.43 |
| 4:CD:101:LEU:C | 4:CD:103:ASN:H | 2.26 | 0.43 |
| 4:CD:148:VAL:CG1 | 4:CD:149:ALA:H | 2.22 | 0.43 |
| 10:CJ:80:LYS:HB2 | 10:CJ:80:LYS:NZ | 2.33 | 0.43 |
| 11:CK:106:LYS:O | 11:CK:107:SER:HB3 | 2.17 | 0.43 |
| 14:CN:13:THR:N | 14:CN:14:PRO:CD | 2.80 | 0.43 |
| 16:CP:43:LYS:O | 16:CP:44:THR:C | 2.60 | 0.43 |
| 22:D0:18:ALA:HB1 | 31:DA:2271:G:OP1 | 2.17 | 0.43 |
| 22:D0:36:ILE:HG12 | 22:D0:37:LEU:N | 2.34 | 0.43 |
| 27:D5:57:VAL:HB | 27:D5:58:LEU:CD1 | 2.47 | 0.43 |
| 30:D8:4:MET:SD | 30:D8:61:LEU:CD1 | 3.06 | 0.43 |
| 31:DA:7:G:C2' | 31:DA:8:A:O4' | 2.63 | 0.43 |
| 31:DA:141:A:H2' | 31:DA:1408:C:O2' | 2.18 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:174:C:C3' | 31:DA:175:G:H5" | 2.48 | 0.43 |
| 31:DA:271(A):A:N7 | 31:DA:271(W):G:N2 | 2.67 | 0.43 |
| 31:DA:481:G:OP2 | 50:DY:47:LYS:HE2 | 2.18 | 0.43 |
| 31:DA:518:G:C4 | 31:DA:519:U:C5 | 3.07 | 0.43 |
| 31:DA:1140:C:OP1 | 39:DN:23:LEU:O | 2.35 | 0.43 |
| 31:DA:1281:G:C2 | 31:DA:1290:C:C2 | 3.06 | 0.43 |
| 31:DA:1378:A:C4' | 31:DA:1379:A:OP1 | 2.56 | 0.43 |
| 31:DA:2681:C:H2' | 31:DA:2681:C:O2 | 2.16 | 0.43 |
| 33:DD:5:LYS:HD2 | 33:DD:5:LYS:H | 1.82 | 0.43 |
| 33:DD:54:ARG:O | 33:DD:218:ARG:HD3 | 2.17 | 0.43 |
| 33:DD:65:ILE:HD13 | 33:DD:65:ILE:C | 2.43 | 0.43 |
| 34:DE:24:THR:CG2 | 34:DE:184:VAL:HG23 | 2.43 | 0.43 |
| 37:DH:156:ALA:O | 37:DH:158:HIS:N | 2.51 | 0.43 |
| 38:DI:64:GLU:CD | 38:DI:64:GLU:N | 2.76 | 0.43 |
| 39:DN:41:ASP:O | 39:DN:42:TRP:C | 2.60 | 0.43 |
| 41:DP:92:GLU:HG3 | 41:DP:123:LEU:CD2 | 2.47 | 0.43 |
| 41:DP:113:LYS:HA | 41:DP:129:ALA:O | 2.17 | 0.43 |
| 43:DR:28:LEU:HD22 | 43:DR:28:LEU:O | 2.18 | 0.43 |
| 43:DR:118:GLU:HA | 43:DR:118:GLU:OE1 | 2.18 | 0.43 |
| 47:DV:75:PHE:HE1 | 47:DV:89:GLN:CB | 2.15 | 0.43 |
| 48:DW:5:ALA:CB | 48:DW:50:VAL:HG23 | 2.48 | 0.43 |
| 51:DZ:44:PHE:CE1 | 51:DZ:48:PHE:CD2 | 3.05 | 0.43 |
| 1:AA:329:A:C2 | 1:AA:332:G:C4 | 3.05 | 0.43 |
| 1:AA:434:U:C4 | 1:AA:435:C:N4 | 2.86 | 0.43 |
| 1:AA:601:C:O2' | 1:AA:602:A:H5' | 2.18 | 0.43 |
| 1:AA:951:G:OP2 | 13:AM:102:ARG:NH2 | 2.50 | 0.43 |
| 1:AA:960:U:C5 | 1:AA:1225:A:H1' | 2.54 | 0.43 |
| 1:AA:1418:A:C2 | 1:AA:1483:A:C2 | 3.06 | 0.43 |
| 1:AA:1499:A:H1' | 1:AA:1520:G:H5' | 2.00 | 0.43 |
| 2:AB:17:PHE:CD2 | 2:AB:17:PHE:C | 2.94 | 0.43 |
| 2:AB:187:LEU:HA | 2:AB:201:ILE:HB | 1.99 | 0.43 |
| 2:AB:204:ASN:ND2 | 2:AB:205:ASP:N | 2.66 | 0.43 |
| 4:AD:127:THR:HA | 4:AD:132:ARG:HA | 2.01 | 0.43 |
| 15:AO:2:PRO:HB2 | 15:AO:3:ILE:H | 1.61 | 0.43 |
| 24:B2:49:LYS:NZ | 24:B2:53:LEU:HD12 | 2.32 | 0.43 |
| 30:B8:60:LEU:HB3 | 30:B8:63:PRO:HG2 | 2.00 | 0.43 |
| 31:BA:70:G:H21 | 31:BA:71:A:H62 | 1.66 | 0.43 |
| 31:BA:146:G:H5' | 31:BA:146:G:C8 | 2.53 | 0.43 |
| 31:BA:547:A:H2' | 31:BA:547:A:N3 | 2.33 | 0.43 |
| 31:BA:996:A:OP2 | 46:BU:92:ARG:NH2 | 2.51 | 0.43 |
| 31:BA:1364:G:H5" | 31:BA:1365:A:OP2 | 2.18 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1562:A:C2 | 31:BA:1563:G:C4 | 3.07 | 0.43 |
| 31:BA:1899:G:C2' | 31:BA:1900:A:OP2 | 2.67 | 0.43 |
| 31:BA:2259:G:C2 | 31:BA:2282:G:N1 | 2.86 | 0.43 |
| 31:BA:2422:A:C4 | 31:BA:2424:C:C5 | 3.06 | 0.43 |
| 31:BA:2515:C:O2' | 31:BA:2516:G:H5' | 2.18 | 0.43 |
| 31:BA:2876:G:H4' | 45:BT:3:ARG:CD | 2.48 | 0.43 |
| 32:BB:27:C:C4 | 32:BB:28:C:C4 | 3.07 | 0.43 |
| 33:BD:31:LYS:O | 33:BD:32:SER:O | 2.37 | 0.43 |
| 33:BD:45:ASN:CG | 33:BD:46:GLN:H | 2.26 | 0.43 |
| 33:BD:127:VAL:HA | 33:BD:193:VAL:CG2 | 2.42 | 0.43 |
| 34:BE:12:THR:HG22 | 45:BT:58:ASN:OD1 | 2.17 | 0.43 |
| 35:BF:70:THR:CG2 | 35:BF:72:ARG:HB2 | 2.49 | 0.43 |
| 35:BF:108:LYS:HD3 | 35:BF:108:LYS:HA | 1.66 | 0.43 |
| 35:BF:136:THR:O | 35:BF:137:LYS:C | 2.61 | 0.43 |
| 35:BF:160:ASN:CG | 35:BF:163:VAL:HG23 | 2.43 | 0.43 |
| 36:BG:37:VAL:HA | 36:BG:158:ALA:O | 2.18 | 0.43 |
| 37:BH:87:LEU:HD13 | 37:BH:148:ILE:HG21 | 2.00 | 0.43 |
| 38:BI:133:HIS:O | 38:BI:135:GLU:HG3 | 2.18 | 0.43 |
| 42:BQ:22:LYS:NZ | 42:BQ:25:ASP:OD1 | 2.50 | 0.43 |
| 43:BR:42:LYS:O | 43:BR:45:ARG:HG3 | 2.18 | 0.43 |
| 45:BT:68:TYR:O | 45:BT:70:VAL:N | 2.51 | 0.43 |
| 46:BU:87:GLY:O | 47:BV:52:VAL:HG13 | 2.19 | 0.43 |
| 51:BZ:166:SER:HB2 | 51:BZ:167:PRO:CA | 2.49 | 0.43 |
| 1:CA:32:A:C2 | 1:CA:33:A:C4 | 3.07 | 0.43 |
| 1:CA:451:A:C6 | 1:CA:481:G:C5 | 3.06 | 0.43 |
| 1:CA:499:A:C4' | 1:CA:500:G:H5' | 2.49 | 0.43 |
| 1:CA:537:G:OP1 | 12:CL:113:ARG:NH2 | 2.47 | 0.43 |
| 1:CA:545:C:H5" | 4:CD:72:GLU:HG2 | 2.00 | 0.43 |
| 1:CA:567:G:C2 | 1:CA:568:G:H1' | 2.53 | 0.43 |
| 1:CA:1025:U:O2' | 1:CA:1026:G:H8 | 2.00 | 0.43 |
| 1:CA:1292:U:O2' | 1:CA:1293:G:H5' | 2.19 | 0.43 |
| 5:CE:20:GLN:HE21 | 5:CE:20:GLN:HB3 | 1.61 | 0.43 |
| 7:CG:54:THR:C | 7:CG:56:GLN:H | 2.26 | 0.43 |
| 7:CG:75:VAL:HG23 | 7:CG:75:VAL:O | 2.18 | 0.43 |
| 11:CK:124:LYS:HB3 | 11:CK:125:PHE:HD1 | 1.83 | 0.43 |
| 23:D1:20:ARG:HG3 | 31:DA:381:G:OP1 | 2.18 | 0.43 |
| 23:D1:34:THR:HG22 | 31:DA:388:G:OP1 | 2.18 | 0.43 |
| 28:D6:27:LYS:HD2 | 31:DA:2285:C:C6 | 2.52 | 0.43 |
| 29:D7:46:VAL:HB | 29:D7:48:LYS:HZ3 | 1.84 | 0.43 |
| 31:DA:14:A:C6 | 31:DA:526:A:C2 | 3.05 | 0.43 |
| 31:DA:128:C:H2' | 31:DA:129:C:H6 | 1.82 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:157:U:H5' | 31:DA:171:G:N2 | 2.32 | 0.43 |
| 31:DA:725:G:C6 | 31:DA:726:G:N1 | 2.86 | 0.43 |
| 31:DA:777:A:N3 | 31:DA:778:G:C8 | 2.86 | 0.43 |
| 31:DA:812:C:H5" | 31:DA:1250:G:O2' | 2.18 | 0.43 |
| 31:DA:1685:C:H2' | 31:DA:1686:C:O5' | 2.19 | 0.43 |
| 31:DA:2494:G:C4 | 31:DA:2495:G:C8 | 3.06 | 0.43 |
| 31:DA:2855:C:H2' | 31:DA:2856:C:H6 | 1.83 | 0.43 |
| 33:DD:3:VAL:O | 33:DD:3:VAL:HG12 | 2.18 | 0.43 |
| 33:DD:30:GLU:CG | 33:DD:63:ARG:NE | 2.81 | 0.43 |
| 33:DD:94:LEU:HD22 | 33:DD:94:LEU:C | 2.42 | 0.43 |
| 33:DD:228:PRO:HD2 | 33:DD:235:GLY:HA3 | 1.98 | 0.43 |
| 34:DE:56:PRO:O | 34:DE:57:LYS:C | 2.61 | 0.43 |
| 35:DF:29:ASN:O | 35:DF:30:PRO:C | 2.60 | 0.43 |
| 37:DH:145:ALA:O | 37:DH:146:ALA:C | 2.61 | 0.43 |
| 37:DH:149:ARG:O | 37:DH:152:ARG:O | 2.35 | 0.43 |
| 38:DI:83:ALA:CB | 38:DI:88:ILE:HA | 2.48 | 0.43 |
| 38:DI:130:TYR:HB2 | 38:DI:138:ILE:HD11 | 2.00 | 0.43 |
| 42:DQ:70:PRO:CA | 42:DQ:95:ALA:HB2 | 2.48 | 0.43 |
| 1:AA:96:U:HO2' | 1:AA:97:G:P | 2.39 | 0.43 |
| 1:AA:255:G:H2' | 1:AA:256:U:C6 | 2.54 | 0.43 |
| 1:AA:375:U:N3 | 1:AA:376:G:N7 | 2.67 | 0.43 |
| 1:AA:994:A:C2 | 14:AN:5:ALA:HA | 2.54 | 0.43 |
| 1:AA:1265:G:C6 | 1:AA:1266:G:C6 | 3.06 | 0.43 |
| 1:AA:1334:G:OP2 | 1:AA:1334:G:C8 | 2.71 | 0.43 |
| 2:AB:25:ASN:HA | 2:AB:26:PRO:HD2 | 1.75 | 0.43 |
| 4:AD:100:ARG:NH1 | 4:AD:137:SER:HA | 2.33 | 0.43 |
| 5:AE:103:GLY:O | 5:AE:104:ALA:C | 2.61 | 0.43 |
| 8:AH:20:TYR:HA | 8:AH:65:TYR:HE2 | 1.79 | 0.43 |
| 10:AJ:39:PRO:CB | 10:AJ:70:ARG:HH12 | 2.31 | 0.43 |
| 11:AK:126:ARG:O | 11:AK:126:ARG:HG2 | 2.18 | 0.43 |
| 17:AQ:6:LEU:O | 17:AQ:58:GLU:HA | 2.19 | 0.43 |
| 24:B2:47:ASN:C | 24:B2:49:LYS:H | 2.25 | 0.43 |
| 25:B3:6:VAL:HG22 | 25:B3:56:VAL:HG22 | 1.99 | 0.43 |
| 28:B6:34:LEU:HD22 | 28:B6:50:ARG:HH12 | 1.83 | 0.43 |
| 29:B7:39:ARG:NH2 | 31:BA:468:G:N7 | 2.60 | 0.43 |
| 30:B8:58:ILE:O | 30:B8:61:LEU:CG | 2.64 | 0.43 |
| 31:BA:192:C:H2' | 31:BA:193:U:H5' | 2.01 | 0.43 |
| 31:BA:194:G:H2' | 31:BA:195:A:O4' | 2.18 | 0.43 |
| 31:BA:478:A:N1 | 31:BA:500:G:H4' | 2.34 | 0.43 |
| 31:BA:547:A:HO2' | 31:BA:548:A:P | 2.42 | 0.43 |
| 31:BA:548:A:HO2' | 31:BA:549:G:P | 2.38 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:700:G:H2' | 31:BA:701:G:O4' | 2.18 | 0.43 |
| 31:BA:773:U:H2' | 31:BA:774:A:H5' | 2.00 | 0.43 |
| 31:BA:838:C:C4 | 31:BA:839:U:C5 | 3.06 | 0.43 |
| 31:BA:1004:C:C2' | 31:BA:1005:C:OP2 | 2.66 | 0.43 |
| 31:BA:1144:G:C6 | 31:BA:1145:C:N4 | 2.86 | 0.43 |
| 31:BA:1312:U:OP2 | 49:BX:62:LYS:HE3 | 2.18 | 0.43 |
| 31:BA:1408:C:C2 | 31:BA:1595:G:N2 | 2.86 | 0.43 |
| 31:BA:1658:C:OP1 | 34:BE:132:HIS:O | 2.36 | 0.43 |
| 31:BA:2015:A:H2' | 31:BA:2016:U:H5' | 2.01 | 0.43 |
| 33:BD:44:ASN:ND2 | 33:BD:47:GLY:O | 2.52 | 0.43 |
| 33:BD:94:LEU:HD22 | 33:BD:95:LEU:N | 2.34 | 0.43 |
| 33:BD:109:ASP:HB2 | 33:BD:197:GLY:CA | 2.49 | 0.43 |
| 33:BD:136:ILE:HA | 33:BD:137:PRO:HD3 | 1.81 | 0.43 |
| 36:BG:67:LYS:HA | 36:BG:68:PRO:HD2 | 1.90 | 0.43 |
| 37:BH:149:ARG:HA | 37:BH:162:ILE:CG1 | 2.40 | 0.43 |
| 39:BN:40:PRO:O | 46:BU:64:ARG:NH2 | 2.52 | 0.43 |
| 41:BP:23:PRO:HB2 | 41:BP:33:ARG:HG3 | 2.01 | 0.43 |
| 41:BP:73:GLY:O | 41:BP:74:GLU:O | 2.36 | 0.43 |
| 44:BS:95:HIS:O | 44:BS:98:VAL:HG23 | 2.17 | 0.43 |
| 45:BT:26:ASP:O | 45:BT:26:ASP:OD2 | 2.36 | 0.43 |
| 45:BT:41:ARG:NH1 | 45:BT:43:GLN:HB2 | 2.34 | 0.43 |
| 46:BU:114:LYS:H | 46:BU:114:LYS:HG2 | 1.62 | 0.43 |
| 47:BV:1:MET:SD | 47:BV:46:VAL:HB | 2.58 | 0.43 |
| 50:BY:2:ARG:O | 50:BY:4:LYS:N | 2.51 | 0.43 |
| 51:BZ:56:VAL:CG1 | 51:BZ:57:ILE:N | 2.81 | 0.43 |
| 1:CA:509:A:H3' | 1:CA:509:A:OP2 | 2.19 | 0.43 |
| 1:CA:525:C:H2' | 1:CA:526:C:H6 | 1.83 | 0.43 |
| 1:CA:629:G:C5 | 1:CA:630:G:N7 | 2.86 | 0.43 |
| 1:CA:658:G:O2' | 1:CA:659:U:H5' | 2.17 | 0.43 |
| 1:CA:758:G:O5' | 1:CA:758:G:H8 | 2.01 | 0.43 |
| 1:CA:950:U:H2' | 1:CA:951:G:C8 | 2.53 | 0.43 |
| 1:CA:994:A:C2 | 14:CN:5:ALA:HA | 2.54 | 0.43 |
| 1:CA:1141:C:H3' | 1:CA:1141:C:H6 | 1.84 | 0.43 |
| 1:CA:1221:G:H4' | 19:CS:53:ASN:O | 2.18 | 0.43 |
| 1:CA:1332:A:H8 | 1:CA:1332:A:O5' | 2.01 | 0.43 |
| 1:CA:1479:C:O2' | 1:CA:1480:G:H5' | 2.17 | 0.43 |
| 3:CC:29:TYR:O | 3:CC:29:TYR:CD2 | 2.72 | 0.43 |
| 3:CC:79:ARG:O | 3:CC:79:ARG:HG3 | 2.18 | 0.43 |
| 4:CD:88:VAL:O | 4:CD:92:VAL:HG23 | 2.19 | 0.43 |
| 4:CD:196:LEU:HB3 | 4:CD:197:PRO:HD2 | 2.01 | 0.43 |
| 7:CG:16:LEU:HD12 | 9:CI:41:VAL:O | 2.18 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 7:CG:50:ILE:O | 7:CG:54:THR:O | 2.37 | 0.43 |
| 9:CI:79:LEU:HD21 | 9:CI:83:ARG:HH21 | 1.83 | 0.43 |
| 10:CJ:54:PHE:CE1 | 10:CJ:55:LYS:HE3 | 2.53 | 0.43 |
| 19:CS:10:PHE:HZ | 19:CS:70:LYS:HE2 | 1.83 | 0.43 |
| 20:CT:14:LYS:HB2 | 20:CT:17:ARG:NH2 | 2.33 | 0.43 |
| 20:CT:57:ARG:HD3 | 20:CT:103:GLY:H | 1.83 | 0.43 |
| 21:CU:21:TYR:O | 21:CU:22:ARG:HB2 | 2.18 | 0.43 |
| 23:D1:75:GLU:O | 23:D1:76:ARG:HD3 | 2.17 | 0.43 |
| 28:D6:24:GLU:HB3 | 28:D6:25:LYS:H | 1.57 | 0.43 |
| 29:D7:5:TRP:CZ3 | 31:DA:464:U:C4’ | 3.02 | 0.43 |
| 29:D7:38:GLY:O | 29:D7:39:ARG:C | 2.61 | 0.43 |
| 31:DA:241:A:H5’ | 31:DA:243:U:O4’ | 2.18 | 0.43 |
| 31:DA:375:C:H2’ | 31:DA:376:C:C6 | 2.53 | 0.43 |
| 31:DA:445:C:C2’ | 31:DA:446:G:H5’ | 2.48 | 0.43 |
| 31:DA:814:C:N4 | 41:DP:27:HIS:CD2 | 2.82 | 0.43 |
| 31:DA:927:G:H2’ | 31:DA:928:G:O4’ | 2.18 | 0.43 |
| 31:DA:959:A:N1 | 31:DA:960:A:C2 | 2.86 | 0.43 |
| 31:DA:996:A:H4’ | 46:DU:92:ARG:CD | 2.48 | 0.43 |
| 31:DA:1109:C:C5 | 31:DA:1110:G:C5 | 2.97 | 0.43 |
| 31:DA:1189:A:N7 | 31:DA:1190:G:C8 | 2.87 | 0.43 |
| 31:DA:1359:A:N7 | 31:DA:1372:U:C4 | 2.85 | 0.43 |
| 31:DA:1528(A):A:H2’ | 31:DA:1529:G:O4’ | 2.17 | 0.43 |
| 31:DA:2471:C:O2 | 31:DA:2471:C:H2’ | 2.19 | 0.43 |
| 31:DA:2564:A:C6 | 31:DA:2565:A:C6 | 3.07 | 0.43 |
| 31:DA:2816:C:C2 | 31:DA:2831:G:C2 | 3.06 | 0.43 |
| 35:DF:8:GLN:HB2 | 35:DF:124:LEU:HD11 | 2.00 | 0.43 |
| 36:DG:63:ILE:HD13 | 36:DG:141:PHE:CE2 | 2.53 | 0.43 |
| 39:DN:78:TYR:CD1 | 39:DN:79:PRO:CD | 2.88 | 0.43 |
| 41:DP:83:VAL:HG12 | 41:DP:112:LEU:CD2 | 2.48 | 0.43 |
| 42:DQ:80:GLU:HA | 42:DQ:80:GLU:OE2 | 2.17 | 0.43 |
| 44:DS:19:LYS:HG2 | 44:DS:19:LYS:O | 2.17 | 0.43 |
| 44:DS:83:LYS:CE | 44:DS:105:ALA:HB2 | 2.44 | 0.43 |
| 44:DS:93:LYS:O | 44:DS:94:TYR:C | 2.61 | 0.43 |
| 44:DS:101:LEU:O | 44:DS:102:ALA:O | 2.37 | 0.43 |
| 45:DT:27:THR:O | 45:DT:47:GLY:O | 2.36 | 0.43 |
| 46:DU:51:LYS:O | 46:DU:53:ARG:N | 2.51 | 0.43 |
| 49:DX:12:VAL:O | 49:DX:12:VAL:CG1 | 2.66 | 0.43 |
| 50:DY:2:ARG:O | 50:DY:4:LYS:N | 2.52 | 0.43 |
| 51:DZ:166:SER:HB2 | 51:DZ:167:PRO:C | 2.44 | 0.43 |
| 1:AA:101:A:H2’ | 1:AA:102:G:H5’ | 2.00 | 0.43 |
| 1:AA:112:G:C2 | 1:AA:113:G:C8 | 3.07 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:251:G:N2 | 1:AA:253:U:C5 | 2.87 | 0.43 |
| 1:AA:258:G:C2 | 1:AA:259:G:C8 | 3.06 | 0.43 |
| 1:AA:437:U:H2' | 1:AA:438:G:O4' | 2.18 | 0.43 |
| 1:AA:499:A:H4' | 1:AA:500:G:H5' | 2.00 | 0.43 |
| 1:AA:629:G:C5 | 1:AA:630:G:N7 | 2.87 | 0.43 |
| 1:AA:1095:U:P | 1:AA:1108:G:H1 | 2.41 | 0.43 |
| 4:AD:17:VAL:HG11 | 4:AD:197:PRO:CB | 2.49 | 0.43 |
| 5:AE:146:ALA:O | 5:AE:148:VAL:N | 2.52 | 0.43 |
| 12:AL:34:ARG:HG2 | 12:AL:35:GLY:N | 2.33 | 0.43 |
| 13:AM:75:ALA:O | 13:AM:79:LYS:HG3 | 2.19 | 0.43 |
| 17:AQ:95:TYR:O | 17:AQ:97:SER:N | 2.51 | 0.43 |
| 24:B2:26:ARG:HG3 | 24:B2:29:LYS:NZ | 2.32 | 0.43 |
| 27:B5:4:HIS:CB | 27:B5:5:PRO:CD | 2.89 | 0.43 |
| 28:B6:14:THR:C | 28:B6:15:GLU:O | 2.60 | 0.43 |
| 31:BA:271(D):G:C5 | 31:BA:271(E):U:C5 | 3.07 | 0.43 |
| 31:BA:312:G:H5' | 31:BA:331:A:O2' | 2.19 | 0.43 |
| 31:BA:541:C:N3 | 31:BA:542:C:N4 | 2.67 | 0.43 |
| 31:BA:696:G:O2' | 31:BA:697:C:H5' | 2.18 | 0.43 |
| 31:BA:954:G:H2' | 31:BA:955:C:O5' | 2.19 | 0.43 |
| 31:BA:1486:A:H2' | 31:BA:1487:G:C8 | 2.53 | 0.43 |
| 31:BA:1613:G:C2 | 31:BA:1619:G:C5 | 3.06 | 0.43 |
| 31:BA:1711:C:O2' | 31:BA:1712:C:H5' | 2.18 | 0.43 |
| 31:BA:2056:G:H2' | 31:BA:2056:G:N3 | 2.33 | 0.43 |
| 31:BA:2103:C:H3' | 31:BA:2104:G:H5" | 1.99 | 0.43 |
| 31:BA:2837:G:C2' | 31:BA:2838:G:H5' | 2.48 | 0.43 |
| 32:BB:28:C:H2' | 32:BB:29:A:C8 | 2.53 | 0.43 |
| 32:BB:45:A:H2' | 32:BB:46:A:H5' | 2.00 | 0.43 |
| 34:BE:23:VAL:HA | 34:BE:184:VAL:O | 2.18 | 0.43 |
| 37:BH:138:LYS:C | 37:BH:140:LYS:H | 2.26 | 0.43 |
| 41:BP:135:LEU:HD22 | 41:BP:135:LEU:HA | 1.89 | 0.43 |
| 43:BR:72:ASP:CG | 43:BR:75:LEU:H | 2.27 | 0.43 |
| 44:BS:25:ARG:NH1 | 44:BS:42:ASP:OD1 | 2.52 | 0.43 |
| 44:BS:59:LYS:NZ | 44:BS:68:GLN:HE22 | 2.16 | 0.43 |
| 45:BT:23:ARG:CB | 45:BT:24:PRO:HD2 | 2.17 | 0.43 |
| 45:BT:58:ASN:C | 45:BT:58:ASN:ND2 | 2.73 | 0.43 |
| 45:BT:91:ARG:HB2 | 45:BT:92:GLY:H | 1.59 | 0.43 |
| 47:BV:24:LYS:HE3 | 47:BV:68:LYS:HE3 | 2.01 | 0.43 |
| 49:BX:7:VAL:HG12 | 49:BX:30:VAL:HG12 | 2.00 | 0.43 |
| 1:CA:175:C:N3 | 1:CA:176:C:C5 | 2.86 | 0.43 |
| 1:CA:219:C:C5 | 1:CA:220:G:C8 | 3.07 | 0.43 |
| 1:CA:448:A:C2 | 1:CA:449:C:C4 | 3.06 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:472:A:O2' | 16:CP:81:ARG:HA | 2.19 | 0.43 |
| 1:CA:604:G:N7 | 1:CA:605:U:C5 | 2.87 | 0.43 |
| 1:CA:648:A:H2' | 1:CA:649:G:H8 | 1.82 | 0.43 |
| 1:CA:776:G:HO2' | 1:CA:777:A:P | 2.42 | 0.43 |
| 1:CA:853:G:H2' | 1:CA:854:G:C8 | 2.52 | 0.43 |
| 1:CA:946:A:N1 | 1:CA:1236:A:C2 | 2.87 | 0.43 |
| 1:CA:1157:A:N3 | 1:CA:1181:G:N3 | 2.66 | 0.43 |
| 1:CA:1425:U:H2' | 1:CA:1426:C:C6 | 2.54 | 0.43 |
| 1:CA:1499:A:H1' | 1:CA:1520:G:H5' | 1.99 | 0.43 |
| 1:CA:1516:G:H2' | 1:CA:1518:A:OP2 | 2.18 | 0.43 |
| 6:CF:45:LEU:HG | 6:CF:46:ARG:H | 1.83 | 0.43 |
| 7:CG:37:ASN:HD21 | 9:CI:40:LEU:CD2 | 2.29 | 0.43 |
| 8:CH:58:TYR:C | 8:CH:59:LEU:HD23 | 2.44 | 0.43 |
| 16:CP:58:TYR:C | 16:CP:60:LEU:N | 2.76 | 0.43 |
| 24:D2:22:GLU:O | 24:D2:25:VAL:HG12 | 2.17 | 0.43 |
| 24:D2:29:LYS:O | 24:D2:33:MET:SD | 2.77 | 0.43 |
| 24:D2:33:MET:HG2 | 49:DX:11:PRO:HD2 | 2.00 | 0.43 |
| 29:D7:21:ARG:O | 29:D7:27:GLY:HA3 | 2.19 | 0.43 |
| 31:DA:171:G:H2' | 31:DA:172:C:C1' | 2.46 | 0.43 |
| 31:DA:372:G:HO2' | 31:DA:373:U:P | 2.41 | 0.43 |
| 31:DA:584:C:N4 | 31:DA:585:G:C6 | 2.86 | 0.43 |
| 31:DA:589:C:O2' | 31:DA:590:A:H5' | 2.18 | 0.43 |
| 31:DA:613:G:C2 | 31:DA:615:G:C6 | 3.07 | 0.43 |
| 31:DA:1131:G:C8 | 31:DA:2025:C:H4' | 2.54 | 0.43 |
| 31:DA:1336:A:O2' | 31:DA:1337:G:H5' | 2.19 | 0.43 |
| 31:DA:1528:A:O2' | 31:DA:1528(A):A:P | 2.76 | 0.43 |
| 31:DA:1858:G:OP2 | 31:DA:1858:G:C8 | 2.72 | 0.43 |
| 31:DA:2013:A:N3 | 31:DA:2013:A:H2' | 2.34 | 0.43 |
| 31:DA:2077:A:O2' | 31:DA:2434:A:O2' | 2.33 | 0.43 |
| 31:DA:2784:C:O2 | 34:DE:37:ARG:NH2 | 2.50 | 0.43 |
| 32:DB:51:G:H5' | 32:DB:52:A:OP2 | 2.19 | 0.43 |
| 33:DD:35:LYS:HZ2 | 33:DD:65:ILE:HA | 1.81 | 0.43 |
| 35:DF:13:SER:HA | 35:DF:14:PRO:HD3 | 1.84 | 0.43 |
| 36:DG:11:TYR:HD2 | 36:DG:12:TYR:CE1 | 2.37 | 0.43 |
| 37:DH:103:LEU:CD2 | 37:DH:115:VAL:HB | 2.49 | 0.43 |
| 38:DI:31:LEU:N | 38:DI:31:LEU:HD13 | 2.33 | 0.43 |
| 41:DP:106:LEU:HD12 | 41:DP:106:LEU:HA | 1.75 | 0.43 |
| 42:DQ:97:VAL:HG11 | 42:DQ:103:MET:HE3 | 2.01 | 0.43 |
| 43:DR:76:VAL:O | 43:DR:77:ARG:C | 2.61 | 0.43 |
| 47:DV:2:PHE:CB | 47:DV:42:GLY:HA3 | 2.48 | 0.43 |
| 50:DY:81:LYS:NZ | 50:DY:97:ARG:HG3 | 2.33 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 51:DZ:70:LEU:HG | 51:DZ:91:LEU:HD11 | 2.00 | 0.43 |
| 1:AA:564:C:H2' | 1:AA:565:U:H5' | 1.99 | 0.43 |
| 1:AA:671:G:C5 | 1:AA:672:U:C5 | 3.07 | 0.43 |
| 1:AA:687:A:C2 | 1:AA:704:A:C6 | 3.06 | 0.43 |
| 1:AA:769:G:H4' | 1:AA:1513:A:H4' | 2.00 | 0.43 |
| 1:AA:818:G:C2 | 1:AA:820:U:O2' | 2.70 | 0.43 |
| 1:AA:1086:U:H2' | 1:AA:1087:G:O4' | 2.18 | 0.43 |
| 1:AA:1157:A:H5' | 1:AA:1158:C:C6 | 2.53 | 0.43 |
| 1:AA:1250:A:H4' | 9:AI:68:GLY:H | 1.83 | 0.43 |
| 1:AA:1289:A:H2' | 1:AA:1290:G:O4' | 2.19 | 0.43 |
| 2:AB:129:GLU:HB3 | 2:AB:130:ARG:H | 1.61 | 0.43 |
| 3:AC:14:ILE:O | 3:AC:15:THR:CB | 2.66 | 0.43 |
| 6:AF:23:LYS:HE2 | 6:AF:23:LYS:HB3 | 1.75 | 0.43 |
| 12:AL:86:ARG:HB2 | 12:AL:101:VAL:HG23 | 2.00 | 0.43 |
| 22:B0:48:GLY:O | 22:B0:49:LYS:C | 2.62 | 0.43 |
| 22:B0:49:LYS:O | 22:B0:50:ASN:HB2 | 2.18 | 0.43 |
| 23:B1:17:SER:C | 23:B1:18:ILE:HD12 | 2.43 | 0.43 |
| 23:B1:42:GLN:HG2 | 23:B1:43:TYR:N | 2.34 | 0.43 |
| 24:B2:49:LYS:O | 24:B2:51:ARG:O | 2.37 | 0.43 |
| 27:B5:33:CYS:SG | 27:B5:49:CYS:HB2 | 2.58 | 0.43 |
| 27:B5:51:TYR:CD2 | 27:B5:52:TYR:OH | 2.71 | 0.43 |
| 30:B8:4:MET:CE | 31:BA:593:G:O4' | 2.66 | 0.43 |
| 30:B8:34:TRP:HZ3 | 30:B8:41:ILE:CG2 | 2.32 | 0.43 |
| 30:B8:35:GLN:HG2 | 31:BA:2420:C:P | 2.58 | 0.43 |
| 30:B8:58:ILE:HD13 | 30:B8:58:ILE:HA | 1.84 | 0.43 |
| 31:BA:26:G:C6 | 31:BA:27:G:N1 | 2.87 | 0.43 |
| 31:BA:190:A:H2 | 31:BA:679:C:O2 | 2.01 | 0.43 |
| 31:BA:231:C:O2' | 31:BA:232:G:H5' | 2.18 | 0.43 |
| 31:BA:513:A:C2 | 31:BA:514:A:C8 | 3.06 | 0.43 |
| 31:BA:562:U:C4 | 31:BA:2036:C:O4' | 2.72 | 0.43 |
| 31:BA:565:C:H2' | 31:BA:566:U:O4' | 2.18 | 0.43 |
| 31:BA:762:U:H4' | 31:BA:763:G:H5" | 2.00 | 0.43 |
| 31:BA:777:A:C2 | 31:BA:778:G:C4 | 3.06 | 0.43 |
| 31:BA:1173:G:H5' | 31:BA:1174:A:P | 2.59 | 0.43 |
| 31:BA:1342:A:O2' | 31:BA:1344:G:OP2 | 2.35 | 0.43 |
| 31:BA:1400:G:H2' | 31:BA:1401:G:C8 | 2.54 | 0.43 |
| 31:BA:1461:G:O2' | 31:BA:1462:C:H5' | 2.18 | 0.43 |
| 31:BA:1700:A:H2' | 31:BA:1701:A:H5' | 2.00 | 0.43 |
| 31:BA:1997:G:O2' | 31:BA:1998:G:H5' | 2.18 | 0.43 |
| 31:BA:2038:G:H2' | 31:BA:2039:C:O4' | 2.18 | 0.43 |
| 31:BA:2314:C:N3 | 31:BA:2315:G:N7 | 2.67 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 32:BB:10:C:O2' | 32:BB:11:C:H5' | 2.18 | 0.43 |
| 33:BD:243:GLY:O | 33:BD:244:ARG:HB3 | 2.17 | 0.43 |
| 34:BE:36:ARG:NH1 | 34:BE:85:ASN:HD21 | 2.16 | 0.43 |
| 35:BF:19:GLU:O | 35:BF:20:LEU:HB2 | 2.18 | 0.43 |
| 39:BN:82:LEU:O | 39:BN:83:LYS:C | 2.61 | 0.43 |
| 40:BO:55:GLY:O | 40:BO:56:ASP:C | 2.62 | 0.43 |
| 42:BQ:46:GLN:O | 42:BQ:47:ILE:C | 2.62 | 0.43 |
| 42:BQ:70:PRO:CA | 42:BQ:95:ALA:HB2 | 2.48 | 0.43 |
| 42:BQ:87:LYS:HB2 | 42:BQ:87:LYS:HE3 | 1.79 | 0.43 |
| 43:BR:118:GLU:HA | 43:BR:118:GLU:OE1 | 2.18 | 0.43 |
| 46:BU:40:PHE:HB3 | 47:BV:78:LYS:HD2 | 2.01 | 0.43 |
| 48:BW:21:VAL:HG21 | 48:BW:76:VAL:HG12 | 2.00 | 0.43 |
| 1:CA:78:G:H1 | 1:CA:91:C:N4 | 2.13 | 0.43 |
| 1:CA:413:G:N2 | 1:CA:429:U:OP2 | 2.47 | 0.43 |
| 1:CA:740:U:O2' | 1:CA:741:G:H5' | 2.18 | 0.43 |
| 1:CA:960:U:C5 | 1:CA:1225:A:H1' | 2.53 | 0.43 |
| 1:CA:1203:C:H2' | 1:CA:1204:A:C8 | 2.53 | 0.43 |
| 2:CB:17:PHE:CD2 | 2:CB:17:PHE:C | 2.94 | 0.43 |
| 2:CB:145:LEU:CD1 | 2:CB:149:LEU:HD12 | 2.49 | 0.43 |
| 3:CC:19:GLU:HA | 3:CC:54:ARG:NH2 | 2.33 | 0.43 |
| 8:CH:36:LEU:HD12 | 8:CH:59:LEU:HD12 | 2.00 | 0.43 |
| 10:CJ:40:LEU:CB | 10:CJ:41:PRO:HD2 | 2.40 | 0.43 |
| 11:CK:84:VAL:C | 11:CK:85:ARG:HG3 | 2.44 | 0.43 |
| 11:CK:92:GLU:O | 11:CK:95:ILE:HG12 | 2.18 | 0.43 |
| 12:CL:26:ALA:O | 12:CL:27:LEU:HB2 | 2.19 | 0.43 |
| 20:CT:56:MET:HE2 | 20:CT:56:MET:HB3 | 1.73 | 0.43 |
| 27:D5:20:ARG:HA | 27:D5:23:HIS:CD2 | 2.53 | 0.43 |
| 28:D6:19:ARG:CG | 28:D6:20:ASN:H | 2.18 | 0.43 |
| 31:DA:271(H):G:C6 | 31:DA:271(Q):G:N1 | 2.87 | 0.43 |
| 31:DA:822:U:O2' | 31:DA:823:G:H5' | 2.18 | 0.43 |
| 31:DA:1173:G:H5' | 31:DA:1174:A:P | 2.58 | 0.43 |
| 31:DA:1505:C:H2' | 31:DA:1506:C:O4' | 2.18 | 0.43 |
| 31:DA:1817:G:P | 33:DD:88:ARG:HH22 | 2.41 | 0.43 |
| 31:DA:1824:G:OP1 | 33:DD:52:ARG:NH1 | 2.50 | 0.43 |
| 31:DA:2321:G:H5" | 31:DA:2322:A:OP2 | 2.19 | 0.43 |
| 31:DA:2563:U:H4' | 40:DO:28:SER:HA | 2.01 | 0.43 |
| 31:DA:2652:C:H2' | 31:DA:2653:U:H5' | 2.01 | 0.43 |
| 31:DA:2689:U:C4' | 31:DA:2690:C:H5' | 2.49 | 0.43 |
| 31:DA:2717:G:H2' | 31:DA:2718:G:H5" | 2.01 | 0.43 |
| 31:DA:2859:G:H3' | 31:DA:2859:G:H8 | 1.81 | 0.43 |
| 33:DD:71:ASP:CG | 33:DD:103:ARG:HH22 | 2.27 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 33:DD:231:HIS:HB3 | 33:DD:233:HIS:O | 2.18 | 0.43 |
| 34:DE:120:TRP:CE3 | 34:DE:155:LYS:HD3 | 2.54 | 0.43 |
| 36:DG:15:VAL:HG22 | 36:DG:175:LEU:HB3 | 2.01 | 0.43 |
| 42:DQ:141:GLN:O | 51:DZ:70:LEU:HD13 | 2.17 | 0.43 |
| 46:DU:83:LEU:C | 46:DU:88:ILE:HD11 | 2.43 | 0.43 |
| 49:DX:73:ARG:O | 49:DX:74:PRO:O | 2.36 | 0.43 |
| 1:AA:391:G:O6 | 1:AA:392:G:C6 | 2.72 | 0.43 |
| 1:AA:484:G:C4' | 1:AA:485:G:O5' | 2.66 | 0.43 |
| 1:AA:520:A:H2 | 1:AA:536:C:O2 | 2.02 | 0.43 |
| 1:AA:1030(B):C:N4 | 1:AA:1030(C):G:C2 | 2.86 | 0.43 |
| 1:AA:1053:G:N7 | 1:AA:1200:C:H5' | 2.34 | 0.43 |
| 1:AA:1128:C:H1' | 1:AA:1146:A:H61 | 1.84 | 0.43 |
| 1:AA:1293:G:O2' | 1:AA:1294:G:P | 2.77 | 0.43 |
| 1:AA:1425:U:H2' | 1:AA:1426:C:C6 | 2.53 | 0.43 |
| 4:AD:15:GLU:HG2 | 4:AD:63:LYS:HG3 | 2.00 | 0.43 |
| 8:AH:20:TYR:CE1 | 8:AH:78:GLN:NE2 | 2.87 | 0.43 |
| 10:AJ:22:LYS:C | 10:AJ:24:VAL:H | 2.26 | 0.43 |
| 12:AL:83:VAL:HG12 | 12:AL:84:LEU:N | 2.32 | 0.43 |
| 21:AU:12:LYS:HB3 | 21:AU:22:ARG:HD2 | 2.01 | 0.43 |
| 22:B0:40:GLN:OE1 | 22:B0:44:ARG:HB2 | 2.18 | 0.43 |
| 24:B2:34:GLU:O | 24:B2:35:LEU:C | 2.61 | 0.43 |
| 30:B8:8:LYS:NZ | 31:BA:243:U:OP2 | 2.49 | 0.43 |
| 31:BA:266:G:H2' | 31:BA:267:C:H5" | 2.00 | 0.43 |
| 31:BA:269:U:O2 | 31:BA:269:U:C2' | 2.67 | 0.43 |
| 31:BA:271(K):U:H3' | 31:BA:271(L):U:C5' | 2.49 | 0.43 |
| 31:BA:271(K):U:H2' | 31:BA:271(M):G:N2 | 2.33 | 0.43 |
| 31:BA:610:G:H2' | 31:BA:611:C:H6 | 1.81 | 0.43 |
| 31:BA:806:C:P | 41:BP:39:LYS:HG3 | 2.58 | 0.43 |
| 31:BA:828:U:O2 | 31:BA:828:U:C3' | 2.67 | 0.43 |
| 31:BA:844:C:H2' | 31:BA:845:G:H5' | 2.00 | 0.43 |
| 31:BA:900:A:H5" | 31:BA:901:A:OP2 | 2.19 | 0.43 |
| 31:BA:1052:C:H3' | 31:BA:1052:C:C6 | 2.53 | 0.43 |
| 31:BA:1383:C:O5' | 31:BA:1383:C:H6 | 2.02 | 0.43 |
| 31:BA:1608:A:O2' | 31:BA:1610:A:OP2 | 2.36 | 0.43 |
| 31:BA:1804:C:H6 | 31:BA:1804:C:O5' | 2.02 | 0.43 |
| 31:BA:1833:U:C2 | 31:BA:1834:U:C6 | 3.07 | 0.43 |
| 31:BA:2250:G:C6 | 42:BQ:82:ARG:CD | 3.02 | 0.43 |
| 31:BA:2684:U:H1' | 40:BO:70:LYS:HD2 | 2.01 | 0.43 |
| 31:BA:2747:G:C2 | 31:BA:2756:U:C5 | 3.06 | 0.43 |
| 32:BB:21:G:O2' | 32:BB:22:U:H6 | 2.02 | 0.43 |
| 32:BB:46:A:H2' | 32:BB:47:C:H6 | 1.83 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 33:BD:248:SER:O | 33:BD:250:TRP:N | 2.52 | 0.43 |
| 33:BD:253:GLN:HB3 | 33:BD:255:LYS:NZ | 2.33 | 0.43 |
| 34:BE:33:VAL:O | 34:BE:33:VAL:HG22 | 2.18 | 0.43 |
| 36:BG:32:PRO:HB3 | 36:BG:163:ALA:HB2 | 1.99 | 0.43 |
| 36:BG:135:LEU:HD23 | 36:BG:140:ILE:HD11 | 2.01 | 0.43 |
| 37:BH:64:LEU:HD23 | 37:BH:64:LEU:HA | 1.85 | 0.43 |
| 38:BI:14:ASP:O | 38:BI:17:GLN:HB3 | 2.19 | 0.43 |
| 38:BI:53:ALA:HB1 | 38:BI:57:ARG:H | 1.84 | 0.43 |
| 41:BP:39:LYS:HZ2 | 41:BP:39:LYS:HG2 | 1.73 | 0.43 |
| 44:BS:12:PHE:O | 44:BS:12:PHE:HD1 | 2.00 | 0.43 |
| 46:BU:8:VAL:HG12 | 46:BU:9:VAL:N | 2.32 | 0.43 |
| 47:BV:69:LYS:O | 47:BV:70:ILE:HB | 2.19 | 0.43 |
| 50:BY:96:ILE:CG1 | 50:BY:99:CYS:HB3 | 2.48 | 0.43 |
| 51:BZ:100:VAL:O | 51:BZ:123:ASP:HB2 | 2.18 | 0.43 |
| 1:CA:9:G:OP1 | 5:CE:122:GLU:HB2 | 2.18 | 0.43 |
| 1:CA:44:G:N2 | 1:CA:399:G:C4 | 2.87 | 0.43 |
| 1:CA:397:A:H5’ | 1:CA:398:C:P | 2.58 | 0.43 |
| 1:CA:429:U:H1’ | 1:CA:430:A:H5” | 2.00 | 0.43 |
| 1:CA:434:U:C4 | 1:CA:435:C:N4 | 2.87 | 0.43 |
| 1:CA:438:G:OP1 | 4:CD:125:HIS:HE1 | 2.02 | 0.43 |
| 1:CA:738:C:OP1 | 6:CF:2:ARG:NH1 | 2.51 | 0.43 |
| 1:CA:747:C:H5 | 1:CA:748:C:N4 | 2.17 | 0.43 |
| 1:CA:1097:C:C2’ | 1:CA:1098:C:H5’ | 2.49 | 0.43 |
| 1:CA:1505:G:C4’ | 1:CA:1506:U:H5” | 2.41 | 0.43 |
| 2:CB:55:PHE:C | 2:CB:57:PHE:N | 2.71 | 0.43 |
| 3:CC:16:ARG:NH1 | 3:CC:16:ARG:HB2 | 2.34 | 0.43 |
| 3:CC:159:GLY:HA2 | 3:CC:193:TYR:CD1 | 2.54 | 0.43 |
| 5:CE:132:ALA:O | 5:CE:133:TYR:C | 2.61 | 0.43 |
| 11:CK:127:LYS:HA | 11:CK:127:LYS:CE | 2.49 | 0.43 |
| 20:CT:23:ARG:O | 20:CT:27:LYS:HB2 | 2.18 | 0.43 |
| 22:D0:77:ARG:HH22 | 31:DA:857:C:H5’ | 1.84 | 0.43 |
| 24:D2:45:SER:O | 24:D2:48:HIS:CB | 2.67 | 0.43 |
| 27:D5:55:ARG:HG2 | 43:DR:33:ARG:HH11 | 1.83 | 0.43 |
| 27:D5:57:VAL:C | 27:D5:58:LEU:CG | 2.91 | 0.43 |
| 31:DA:271(P):C:H2’ | 31:DA:271(Q):G:C5’ | 2.49 | 0.43 |
| 31:DA:271(S):G:H2’ | 31:DA:271(T):C:O4’ | 2.19 | 0.43 |
| 31:DA:494:G:H5” | 31:DA:494:G:C8 | 2.53 | 0.43 |
| 31:DA:541:C:H2’ | 31:DA:542:C:C5 | 2.54 | 0.43 |
| 31:DA:777:A:O2’ | 31:DA:778:G:H5’ | 2.19 | 0.43 |
| 31:DA:867:C:C6 | 31:DA:868:U:C5 | 3.07 | 0.43 |
| 31:DA:1400:G:H2’ | 31:DA:1401:G:C8 | 2.53 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1486:A:H2' | 31:DA:1487:G:C8 | 2.54 | 0.43 |
| 31:DA:1575:C:H2' | 31:DA:1576:U:C6 | 2.53 | 0.43 |
| 31:DA:1588:C:H2' | 31:DA:1588:C:O2 | 2.19 | 0.43 |
| 31:DA:1721:G:N2 | 31:DA:1739:U:OP2 | 2.51 | 0.43 |
| 31:DA:1907:G:C2' | 31:DA:1908:C:H5' | 2.49 | 0.43 |
| 31:DA:2056:G:OP2 | 31:DA:2057:A:OP2 | 2.36 | 0.43 |
| 31:DA:2632:A:N3 | 34:DE:61:ARG:HD2 | 2.34 | 0.43 |
| 31:DA:2730:C:H4' | 34:DE:168:MET:O | 2.19 | 0.43 |
| 31:DA:2847:U:OP1 | 45:DT:98:LYS:HD3 | 2.19 | 0.43 |
| 32:DB:18:G:H2' | 32:DB:19:G:O4' | 2.19 | 0.43 |
| 32:DB:21:G:H2' | 32:DB:21:G:N3 | 2.32 | 0.43 |
| 33:DD:206:LEU:HA | 33:DD:206:LEU:HD23 | 1.49 | 0.43 |
| 33:DD:270:ILE:HD12 | 33:DD:270:ILE:O | 2.17 | 0.43 |
| 34:DE:70:ALA:O | 34:DE:72:VAL:O | 2.36 | 0.43 |
| 35:DF:22:ALA:CA | 35:DF:26:ALA:CB | 2.96 | 0.43 |
| 35:DF:29:ASN:OD1 | 35:DF:29:ASN:C | 2.61 | 0.43 |
| 36:DG:51:ARG:HD3 | 36:DG:53:LEU:HD21 | 2.00 | 0.43 |
| 37:DH:98:LEU:HD12 | 37:DH:102:ALA:O | 2.18 | 0.43 |
| 38:DI:56:LYS:HA | 38:DI:59:ALA:HB3 | 2.00 | 0.43 |
| 39:DN:82:LEU:N | 39:DN:82:LEU:CD1 | 2.82 | 0.43 |
| 40:DO:88:ASN:HB3 | 40:DO:92:GLU:O | 2.19 | 0.43 |
| 41:DP:16:ARG:NH1 | 41:DP:16:ARG:HB2 | 2.33 | 0.43 |
| 41:DP:16:ARG:NH1 | 41:DP:16:ARG:O | 2.40 | 0.43 |
| 43:DR:53:HIS:CD2 | 43:DR:94:TYR:OH | 2.67 | 0.43 |
| 45:DT:3:ARG:CB | 45:DT:6:LEU:HB3 | 2.38 | 0.43 |
| 45:DT:41:ARG:HH11 | 45:DT:43:GLN:HA | 1.83 | 0.43 |
| 47:DV:82:ARG:O | 47:DV:82:ARG:HD3 | 2.19 | 0.43 |
| 49:DX:90:GLU:C | 49:DX:92:LEU:N | 2.76 | 0.43 |
| 51:DZ:145:GLU:CG | 51:DZ:146:ILE:H | 2.32 | 0.43 |
| 1:AA:375:U:C4 | 1:AA:376:G:N7 | 2.86 | 0.43 |
| 1:AA:612:C:O2 | 1:AA:629:G:N2 | 2.52 | 0.43 |
| 1:AA:682:G:C2 | 1:AA:709:G:C6 | 3.06 | 0.43 |
| 1:AA:849:C:H2' | 1:AA:850:U:O4' | 2.18 | 0.43 |
| 1:AA:1037:C:O5' | 1:AA:1037:C:H6 | 2.02 | 0.43 |
| 1:AA:1112:C:N3 | 3:AC:178:LEU:HD23 | 2.33 | 0.43 |
| 1:AA:1202:G:H2' | 1:AA:1203:C:H5' | 2.01 | 0.43 |
| 1:AA:1343:G:H2' | 1:AA:1344:C:C6 | 2.54 | 0.43 |
| 1:AA:1479:C:O2' | 1:AA:1480:G:H5' | 2.19 | 0.43 |
| 2:AB:75:LYS:HA | 2:AB:78:GLN:HG3 | 2.01 | 0.43 |
| 4:AD:14:ARG:C | 4:AD:16:GLY:H | 2.25 | 0.43 |
| 4:AD:58:LEU:HD23 | 4:AD:58:LEU:HA | 1.90 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 10:AJ:16:LEU:HD13 | 10:AJ:16:LEU:O | 2.18 | 0.43 |
| 11:AK:13:GLN:HB3 | 11:AK:75:TYR:O | 2.19 | 0.43 |
| 22:B0:72:ARG:HH21 | 22:B0:75:LEU:HD12 | 1.83 | 0.43 |
| 22:B0:73:GLY:C | 22:B0:75:LEU:N | 2.75 | 0.43 |
| 24:B2:26:ARG:O | 24:B2:30:ARG:HD3 | 2.18 | 0.43 |
| 24:B2:44:LEU:HD12 | 24:B2:44:LEU:C | 2.30 | 0.43 |
| 26:B4:28:LYS:CB | 36:BG:113:ARG:HH22 | 2.31 | 0.43 |
| 28:B6:44:ARG:O | 28:B6:45:LYS:CG | 2.66 | 0.43 |
| 31:BA:27:G:H22 | 31:BA:512:G:H1' | 1.80 | 0.43 |
| 31:BA:50:U:OP2 | 31:BA:50:U:H4' | 2.18 | 0.43 |
| 31:BA:212:G:HO2' | 31:BA:213:A:H5' | 1.83 | 0.43 |
| 31:BA:288:C:O2 | 31:BA:288:C:H2' | 2.19 | 0.43 |
| 31:BA:394:A:O2' | 31:BA:395:U:H5' | 2.19 | 0.43 |
| 31:BA:719:C:O2' | 31:BA:720:C:H5' | 2.19 | 0.43 |
| 31:BA:985:C:H2' | 31:BA:986:C:C6 | 2.54 | 0.43 |
| 31:BA:1276:A:O2' | 43:BR:16:HIS:CE1 | 2.72 | 0.43 |
| 31:BA:1694:C:O2 | 31:BA:1694:C:C2' | 2.67 | 0.43 |
| 31:BA:2772:C:H2' | 31:BA:2773:C:H6 | 1.84 | 0.43 |
| 31:BA:2848:G:H3' | 45:BT:95:ARG:O | 2.18 | 0.43 |
| 32:BB:18:G:H2' | 32:BB:19:G:O4' | 2.19 | 0.43 |
| 32:BB:33:G:C2 | 32:BB:50:G:C2 | 3.07 | 0.43 |
| 32:BB:45:A:C2' | 32:BB:46:A:H5' | 2.49 | 0.43 |
| 32:BB:51:G:H5' | 32:BB:52:A:OP2 | 2.19 | 0.43 |
| 32:BB:79:C:O2' | 32:BB:80:U:H5' | 2.19 | 0.43 |
| 34:BE:14:ILE:CG1 | 34:BE:21:VAL:CG2 | 2.93 | 0.43 |
| 35:BF:36:VAL:O | 35:BF:39:TRP:HB3 | 2.19 | 0.43 |
| 35:BF:115:ALA:O | 35:BF:116:ASP:C | 2.60 | 0.43 |
| 36:BG:15:VAL:HG22 | 36:BG:175:LEU:HB3 | 1.99 | 0.43 |
| 37:BH:43:VAL:CG2 | 37:BH:52:VAL:HG13 | 2.36 | 0.43 |
| 37:BH:94:TYR:N | 37:BH:94:TYR:CD1 | 2.87 | 0.43 |
| 37:BH:157:TYR:O | 37:BH:158:HIS:HB2 | 2.19 | 0.43 |
| 43:BR:21:TYR:CZ | 43:BR:43:GLU:HG2 | 2.54 | 0.43 |
| 43:BR:107:ASP:C | 43:BR:107:ASP:OD2 | 2.60 | 0.43 |
| 43:BR:113:LEU:HA | 43:BR:113:LEU:HD12 | 1.63 | 0.43 |
| 44:BS:30:ARG:HD2 | 44:BS:31:SER:O | 2.19 | 0.43 |
| 46:BU:80:ILE:O | 46:BU:83:LEU:HB2 | 2.19 | 0.43 |
| 48:BW:12:ILE:HB | 48:BW:42:ARG:HH12 | 1.83 | 0.43 |
| 49:BX:13:LEU:HD23 | 49:BX:13:LEU:N | 2.34 | 0.43 |
| 50:BY:98:VAL:O | 50:BY:99:CYS:HB2 | 2.18 | 0.43 |
| 51:BZ:44:PHE:CE1 | 51:BZ:48:PHE:HB2 | 2.54 | 0.43 |
| 51:BZ:100:VAL:C | 51:BZ:123:ASP:HB2 | 2.44 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:90:U:O2' | 1:CA:91:C:C5 | 2.66 | 0.43 |
| 1:CA:271:C:C2 | 1:CA:272:C:C5 | 3.07 | 0.43 |
| 1:CA:328:C:H4' | 1:CA:329:A:C5' | 2.49 | 0.43 |
| 1:CA:504:C:H1' | 1:CA:510:A:C4 | 2.54 | 0.43 |
| 1:CA:510:A:H5" | 1:CA:511:C:P | 2.59 | 0.43 |
| 1:CA:524:G:C6 | 1:CA:525:C:N4 | 2.87 | 0.43 |
| 1:CA:682:G:H1 | 1:CA:708:C:N4 | 2.17 | 0.43 |
| 1:CA:715:A:O2' | 1:CA:716:A:H5' | 2.18 | 0.43 |
| 1:CA:1085:U:C2 | 1:CA:1094:G:O6 | 2.71 | 0.43 |
| 1:CA:1086:U:H2' | 1:CA:1087:G:O4' | 2.19 | 0.43 |
| 1:CA:1293:G:O2' | 1:CA:1294:G:H8 | 2.00 | 0.43 |
| 2:CB:75:LYS:HA | 2:CB:78:GLN:HG3 | 2.00 | 0.43 |
| 2:CB:100:GLY:O | 2:CB:104:ASN:N | 2.50 | 0.43 |
| 2:CB:144:ARG:O | 2:CB:145:LEU:C | 2.61 | 0.43 |
| 4:CD:93:PHE:O | 4:CD:97:LEU:HB2 | 2.18 | 0.43 |
| 4:CD:100:ARG:O | 4:CD:101:LEU:C | 2.61 | 0.43 |
| 4:CD:132:ARG:H | 4:CD:132:ARG:HG3 | 1.51 | 0.43 |
| 5:CE:31:LEU:HD11 | 5:CE:129:ILE:HA | 2.00 | 0.43 |
| 23:D1:86:SER:C | 23:D1:89:GLU:HG2 | 2.44 | 0.43 |
| 25:D3:52:HIS:NE2 | 32:DB:83:G:H5" | 2.33 | 0.43 |
| 31:DA:288:C:N4 | 31:DA:353:G:H1 | 2.09 | 0.43 |
| 31:DA:934:G:H2' | 31:DA:935:C:C6 | 2.53 | 0.43 |
| 31:DA:1568:G:OP2 | 33:DD:63:ARG:NH2 | 2.49 | 0.43 |
| 31:DA:1783:A:C2 | 31:DA:2587:A:C6 | 3.07 | 0.43 |
| 31:DA:2564:A:OP1 | 31:DA:2648:C:H4' | 2.18 | 0.43 |
| 31:DA:2645:G:H3' | 31:DA:2646:C:H5' | 2.00 | 0.43 |
| 31:DA:2886:G:H2' | 31:DA:2887:U:H6 | 1.82 | 0.43 |
| 33:DD:34:VAL:C | 33:DD:35:LYS:O | 2.60 | 0.43 |
| 34:DE:27:LEU:HD13 | 45:DT:1:MET:HE2 | 2.01 | 0.43 |
| 35:DF:65:TRP:CZ3 | 35:DF:72:ARG:HB3 | 2.53 | 0.43 |
| 36:DG:19:LEU:HG | 36:DG:175:LEU:CD1 | 2.49 | 0.43 |
| 36:DG:98:ARG:O | 36:DG:101:ILE:HG22 | 2.19 | 0.43 |
| 36:DG:109:VAL:O | 36:DG:112:PRO:HD2 | 2.19 | 0.43 |
| 37:DH:30:LYS:HZ2 | 37:DH:81:GLU:H | 1.67 | 0.43 |
| 39:DN:123:TYR:CZ | 39:DN:130:HIS:HD2 | 2.36 | 0.43 |
| 44:DS:38:GLN:HB3 | 44:DS:47:THR:CG2 | 2.48 | 0.43 |
| 45:DT:8:LYS:O | 45:DT:9:LEU:C | 2.62 | 0.43 |
| 45:DT:16:ARG:HD3 | 45:DT:16:ARG:HA | 1.53 | 0.43 |
| 45:DT:114:LEU:O | 45:DT:115:ARG:O | 2.36 | 0.43 |
| 47:DV:82:ARG:C | 47:DV:83:ARG:HG3 | 2.41 | 0.43 |
| 49:DX:57:LEU:HD12 | 49:DX:76:ARG:HG3 | 2.00 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 51:DZ:129:SER:OG | 51:DZ:131:ARG:HG3 | 2.19 | 0.43 |
| 1:AA:81:U:H2' | 1:AA:82:U:H5 | 1.84 | 0.43 |
| 1:AA:175:C:C2 | 1:AA:176:C:C6 | 3.07 | 0.43 |
| 1:AA:192:U:C4' | 20:AT:103:GLY:HA2 | 2.49 | 0.43 |
| 1:AA:392:G:C4 | 1:AA:393:A:C8 | 3.07 | 0.43 |
| 1:AA:429:U:H1' | 1:AA:430:A:H5" | 2.01 | 0.43 |
| 1:AA:515:G:C2 | 1:AA:537:G:C2 | 3.07 | 0.43 |
| 1:AA:738:C:OP1 | 6:AF:2:ARG:NH1 | 2.50 | 0.43 |
| 1:AA:800:G:H8 | 1:AA:800:G:O5' | 2.02 | 0.43 |
| 1:AA:926:G:C6 | 1:AA:1505:G:C6 | 3.07 | 0.43 |
| 1:AA:928:G:C2 | 1:AA:1390:U:O2 | 2.72 | 0.43 |
| 1:AA:985:C:O5' | 1:AA:985:C:H6 | 2.01 | 0.43 |
| 1:AA:1006:C:H42 | 1:AA:1024:G:H21 | 1.66 | 0.43 |
| 1:AA:1030(D):A:N7 | 1:AA:1031:G:N3 | 2.66 | 0.43 |
| 1:AA:1189:C:H5" | 3:AC:5:ILE:HG21 | 1.99 | 0.43 |
| 1:AA:1250:A:H2 | 1:AA:1370:G:H1' | 1.83 | 0.43 |
| 2:AB:221:LEU:HD13 | 2:AB:221:LEU:O | 2.18 | 0.43 |
| 4:AD:109:GLY:O | 4:AD:110:PHE:C | 2.61 | 0.43 |
| 4:AD:122:ARG:HD3 | 4:AD:122:ARG:O | 2.19 | 0.43 |
| 4:AD:202:LEU:N | 4:AD:202:LEU:HD23 | 2.33 | 0.43 |
| 7:AG:50:ILE:O | 7:AG:54:THR:O | 2.37 | 0.43 |
| 8:AH:25:ASP:HA | 8:AH:59:LEU:O | 2.19 | 0.43 |
| 9:AI:49:PRO:O | 9:AI:53:VAL:HG13 | 2.19 | 0.43 |
| 15:AO:51:HIS:O | 15:AO:54:ARG:HB3 | 2.17 | 0.43 |
| 16:AP:68:ASP:C | 16:AP:70:ALA:H | 2.27 | 0.43 |
| 17:AQ:33:GLY:O | 17:AQ:34:LYS:O | 2.37 | 0.43 |
| 27:B5:57:VAL:CB | 27:B5:58:LEU:HD12 | 2.46 | 0.43 |
| 31:BA:458:G:O2' | 31:BA:469:G:O6 | 2.37 | 0.43 |
| 31:BA:477:A:H2' | 31:BA:478:A:C8 | 2.53 | 0.43 |
| 31:BA:869:G:C5 | 31:BA:870:A:N7 | 2.86 | 0.43 |
| 31:BA:1042:G:H3' | 31:BA:1043:C:O4' | 2.18 | 0.43 |
| 31:BA:1176:G:C4' | 31:BA:1177:A:OP1 | 2.67 | 0.43 |
| 31:BA:2517:C:C6 | 31:BA:2542:A:N1 | 2.87 | 0.43 |
| 31:BA:2584:U:H2' | 31:BA:2585:U:H5' | 2.00 | 0.43 |
| 31:BA:2592:G:H2' | 31:BA:2593:U:O4' | 2.19 | 0.43 |
| 32:BB:15:A:HO2' | 32:BB:110:G:H8 | 1.58 | 0.43 |
| 34:BE:201:THR:HG22 | 34:BE:203:LYS:H | 1.83 | 0.43 |
| 35:BF:53:THR:CG2 | 35:BF:56:GLU:H | 2.31 | 0.43 |
| 36:BG:37:VAL:HG23 | 36:BG:99:MET:HG3 | 1.99 | 0.43 |
| 36:BG:60:LEU:HD22 | 36:BG:63:ILE:CG1 | 2.49 | 0.43 |
| 38:BI:14:ASP:O | 38:BI:15:VAL:C | 2.61 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 38:BI:133:HIS:ND1 | 38:BI:134:PRO:CD | 2.78 | 0.43 |
| 39:BN:33:LEU:HD12 | 39:BN:33:LEU:HA | 1.85 | 0.43 |
| 42:BQ:83:MET:HB3 | 42:BQ:83:MET:HE2 | 1.80 | 0.43 |
| 44:BS:69:VAL:HG13 | 44:BS:70:GLY:N | 2.33 | 0.43 |
| 47:BV:90:PRO:CD | 47:BV:91:TYR:H | 2.32 | 0.43 |
| 48:BW:78:GLU:HG3 | 48:BW:79:GLY:N | 2.33 | 0.43 |
| 50:BY:14:LEU:CD1 | 50:BY:22:GLY:HA2 | 2.48 | 0.43 |
| 50:BY:32:PRO:C | 50:BY:34:LYS:N | 2.76 | 0.43 |
| 50:BY:88:LYS:HZ2 | 50:BY:93:GLY:HA3 | 1.82 | 0.43 |
| 51:BZ:14:LYS:HB2 | 51:BZ:17:ALA:CB | 2.48 | 0.43 |
| 51:BZ:37:VAL:O | 51:BZ:38:TYR:HB3 | 2.18 | 0.43 |
| 1:CA:193:C:O2' | 1:CA:194:C:H5' | 2.17 | 0.43 |
| 1:CA:410:G:OP2 | 4:CD:25:ARG:HG3 | 2.18 | 0.43 |
| 1:CA:760:G:C2' | 1:CA:761:G:H5' | 2.49 | 0.43 |
| 1:CA:882:C:O2' | 1:CA:883:C:H5' | 2.18 | 0.43 |
| 1:CA:1502:A:C2 | 1:CA:1505:G:N1 | 2.62 | 0.43 |
| 2:CB:22:LYS:HZ3 | 2:CB:40:HIS:HE1 | 1.63 | 0.43 |
| 3:CC:73:PRO:HA | 3:CC:76:VAL:CG1 | 2.49 | 0.43 |
| 3:CC:125:GLU:OE2 | 3:CC:189:ALA:HA | 2.19 | 0.43 |
| 5:CE:146:ALA:C | 5:CE:148:VAL:H | 2.27 | 0.43 |
| 9:CI:28:VAL:HA | 9:CI:63:ILE:O | 2.18 | 0.43 |
| 12:CL:13:LYS:N | 12:CL:13:LYS:HD2 | 2.34 | 0.43 |
| 13:CM:17:VAL:O | 13:CM:20:THR:HB | 2.18 | 0.43 |
| 13:CM:20:THR:C | 13:CM:22:ILE:H | 2.26 | 0.43 |
| 19:CS:19:VAL:CG1 | 19:CS:44:MET:HE2 | 2.49 | 0.43 |
| 20:CT:12:ALA:O | 20:CT:15:ARG:HB2 | 2.19 | 0.43 |
| 22:D0:34:GLY:O | 22:D0:35:ASN:C | 2.61 | 0.43 |
| 23:D1:13:ILE:CG1 | 23:D1:14:VAL:N | 2.58 | 0.43 |
| 23:D1:69:LYS:O | 23:D1:70:VAL:C | 2.60 | 0.43 |
| 30:D8:13:ARG:NH2 | 31:DA:250:G:OP2 | 2.52 | 0.43 |
| 31:DA:207:A:H2' | 31:DA:208:C:O4' | 2.19 | 0.43 |
| 31:DA:271(D):G:H1 | 31:DA:271(T):C:H42 | 1.67 | 0.43 |
| 31:DA:489:G:C5 | 31:DA:1284:A:C2 | 3.07 | 0.43 |
| 31:DA:543:C:N4 | 31:DA:551:G:N1 | 2.67 | 0.43 |
| 31:DA:621:A:H2' | 31:DA:622:G:C5' | 2.49 | 0.43 |
| 31:DA:685:A:N3 | 31:DA:689:A:C6 | 2.87 | 0.43 |
| 31:DA:1142(A):A:C5 | 31:DA:1144:G:C5 | 3.07 | 0.43 |
| 31:DA:1143:A:OP1 | 39:DN:25:ARG:NH2 | 2.52 | 0.43 |
| 31:DA:1394:U:C4 | 31:DA:1395:A:C5 | 3.07 | 0.43 |
| 31:DA:1449:A:O3' | 31:DA:1530:C:N4 | 2.52 | 0.43 |
| 31:DA:2224:G:OP1 | 33:DD:268:ARG:NH1 | 2.52 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2544:G:H1' | 31:DA:2646:C:H4' | 2.01 | 0.43 |
| 31:DA:2828:C:O2' | 31:DA:2829:C:H5' | 2.18 | 0.43 |
| 31:DA:2840:C:H2' | 31:DA:2841:C:H6 | 1.84 | 0.43 |
| 32:DB:46:A:H2' | 32:DB:47:C:C6 | 2.54 | 0.43 |
| 33:DD:10:THR:C | 33:DD:11:PRO:O | 2.61 | 0.43 |
| 33:DD:89:SER:HB2 | 33:DD:159:ALA:CB | 2.48 | 0.43 |
| 33:DD:142:VAL:CG2 | 33:DD:191:ALA:HB1 | 2.49 | 0.43 |
| 34:DE:13:ARG:HA | 34:DE:21:VAL:O | 2.19 | 0.43 |
| 34:DE:137:HIS:HB3 | 34:DE:138:PRO:HD2 | 2.00 | 0.43 |
| 34:DE:181:LEU:HB3 | 34:DE:182:LEU:H | 1.69 | 0.43 |
| 38:DI:67:ARG:O | 38:DI:68:LEU:CB | 2.65 | 0.43 |
| 39:DN:58:ASP:O | 39:DN:60:ILE:N | 2.50 | 0.43 |
| 40:DO:31:LYS:HB3 | 40:DO:32:TYR:HD1 | 1.77 | 0.43 |
| 49:DX:90:GLU:C | 49:DX:92:LEU:H | 2.26 | 0.43 |
| 50:DY:55:TYR:O | 50:DY:56:PRO:C | 2.62 | 0.43 |
| 1:AA:342:C:C2' | 1:AA:343:U:H5' | 2.49 | 0.43 |
| 1:AA:356:A:H2' | 1:AA:357:G:O5' | 2.18 | 0.43 |
| 1:AA:499:A:C4' | 1:AA:500:G:H5' | 2.49 | 0.43 |
| 1:AA:682:G:H1 | 1:AA:708:C:N4 | 2.17 | 0.43 |
| 1:AA:1054:C:O2' | 1:AA:1055:A:H5" | 2.19 | 0.43 |
| 1:AA:1173:G:H2' | 1:AA:1174:G:H8 | 1.83 | 0.43 |
| 1:AA:1242:C:H5" | 21:AU:10:ARG:NH1 | 2.34 | 0.43 |
| 1:AA:1326:C:H2' | 1:AA:1327:C:C6 | 2.53 | 0.43 |
| 1:AA:1332:A:H8 | 1:AA:1332:A:O5' | 2.01 | 0.43 |
| 4:AD:8:VAL:HB | 4:AD:21:LEU:CD1 | 2.47 | 0.43 |
| 4:AD:101:LEU:C | 4:AD:103:ASN:N | 2.76 | 0.43 |
| 5:AE:146:ALA:C | 5:AE:148:VAL:N | 2.76 | 0.43 |
| 10:AJ:32:ALA:HB1 | 10:AJ:75:ILE:CD1 | 2.48 | 0.43 |
| 12:AL:119:LYS:C | 12:AL:120:TYR:HD1 | 2.27 | 0.43 |
| 15:AO:55:GLY:O | 15:AO:56:LEU:C | 2.62 | 0.43 |
| 17:AQ:95:TYR:O | 17:AQ:96:GLU:C | 2.62 | 0.43 |
| 20:AT:46:GLU:CD | 20:AT:48:LYS:HE2 | 2.44 | 0.43 |
| 24:B2:23:LYS:HA | 49:BX:5:TYR:HD1 | 1.81 | 0.43 |
| 24:B2:51:ARG:O | 24:B2:52:ASP:HB2 | 2.17 | 0.43 |
| 26:B4:29:PRO:C | 26:B4:31:ILE:N | 2.77 | 0.43 |
| 30:B8:3:LYS:HE3 | 31:BA:242:G:O5' | 2.19 | 0.43 |
| 31:BA:84:A:N1 | 31:BA:98:G:O2' | 2.36 | 0.43 |
| 31:BA:239:U:H2' | 31:BA:240:G:O4' | 2.18 | 0.43 |
| 31:BA:675:A:N6 | 31:BA:676:A:N6 | 2.66 | 0.43 |
| 31:BA:912:C:N3 | 31:BA:913:U:C5 | 2.86 | 0.43 |
| 31:BA:942:G:C6 | 31:BA:943:U:C4 | 3.07 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1210:A:C5' | 31:BA:1212:G:H5' | 2.49 | 0.43 |
| 31:BA:1813:G:C2' | 31:BA:1814:G:H5' | 2.48 | 0.43 |
| 31:BA:2070:G:C2 | 31:BA:2442:C:C2 | 3.07 | 0.43 |
| 31:BA:2267:A:H5" | 31:BA:2268:A:H5' | 2.01 | 0.43 |
| 31:BA:2393:A:H2' | 31:BA:2394:C:O4' | 2.19 | 0.43 |
| 31:BA:2575:C:H6 | 31:BA:2575:C:O5' | 2.02 | 0.43 |
| 31:BA:2592:G:C5 | 31:BA:2593:U:C5 | 3.07 | 0.43 |
| 35:BF:28:ILE:HG12 | 35:BF:119:ARG:HH21 | 1.83 | 0.43 |
| 35:BF:53:THR:C | 35:BF:55:GLY:H | 2.27 | 0.43 |
| 36:BG:98:ARG:O | 36:BG:101:ILE:HG22 | 2.19 | 0.43 |
| 38:BI:29:TYR:HD2 | 38:BI:30:LEU:HD23 | 1.83 | 0.43 |
| 40:BO:17:ARG:HD3 | 40:BO:17:ARG:HA | 1.60 | 0.43 |
| 45:BT:54:ARG:HA | 45:BT:59:THR:CB | 2.43 | 0.43 |
| 45:BT:114:LEU:O | 45:BT:115:ARG:O | 2.37 | 0.43 |
| 45:BT:128:GLU:C | 45:BT:128:GLU:OE1 | 2.61 | 0.43 |
| 46:BU:80:ILE:HD13 | 46:BU:80:ILE:HA | 1.74 | 0.43 |
| 47:BV:19:LYS:HZ1 | 47:BV:20:LEU:CB | 2.23 | 0.43 |
| 47:BV:50:PRO:HB2 | 47:BV:51:VAL:H | 1.52 | 0.43 |
| 49:BX:8:ILE:N | 49:BX:8:ILE:HD12 | 2.34 | 0.43 |
| 49:BX:32:PRO:CG | 49:BX:72:LYS:HD3 | 2.49 | 0.43 |
| 49:BX:75:ASP:O | 49:BX:76:ARG:HB3 | 2.19 | 0.43 |
| 49:BX:84:ALA:C | 49:BX:86:GLY:H | 2.26 | 0.43 |
| 1:CA:25:C:C5' | 1:CA:524:G:H1' | 2.49 | 0.43 |
| 1:CA:39:G:C6 | 1:CA:40:C:C5 | 3.07 | 0.43 |
| 1:CA:90:U:O3' | 1:CA:91:C:H6 | 2.02 | 0.43 |
| 1:CA:448:A:OP2 | 1:CA:485:G:N2 | 2.38 | 0.43 |
| 1:CA:510:A:H5" | 1:CA:511:C:OP1 | 2.18 | 0.43 |
| 1:CA:734:G:C6 | 1:CA:735:C:N3 | 2.87 | 0.43 |
| 4:CD:43:HIS:CE1 | 4:CD:46:LYS:HZ2 | 2.34 | 0.43 |
| 4:CD:131:ARG:HD3 | 4:CD:131:ARG:N | 2.34 | 0.43 |
| 5:CE:13:ILE:HA | 5:CE:29:GLY:O | 2.18 | 0.43 |
| 5:CE:139:LEU:O | 5:CE:142:LEU:HG | 2.19 | 0.43 |
| 8:CH:9:MET:HE1 | 8:CH:32:LYS:HA | 2.01 | 0.43 |
| 10:CJ:78:ASN:O | 10:CJ:82:ILE:HG12 | 2.19 | 0.43 |
| 11:CK:59:TYR:O | 11:CK:62:GLN:HB3 | 2.19 | 0.43 |
| 13:CM:81:LEU:HD11 | 13:CM:88:ARG:HH12 | 1.84 | 0.43 |
| 15:CO:74:ASP:HA | 15:CO:75:PRO:HD2 | 1.82 | 0.43 |
| 16:CP:58:TYR:O | 16:CP:61:SER:N | 2.52 | 0.43 |
| 17:CQ:67:LYS:O | 17:CQ:69:LYS:N | 2.52 | 0.43 |
| 19:CS:4:SER:O | 19:CS:5:LEU:HB2 | 2.17 | 0.43 |
| 19:CS:6:LYS:HD2 | 19:CS:6:LYS:H | 1.83 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 20:CT:57:ARG:HB2 | 20:CT:57:ARG:NH1 | 2.34 | 0.43 |
| 22:D0:70:GLN:OE1 | 22:D0:72:ARG:HD3 | 2.18 | 0.43 |
| 23:D1:56:GLN:HB3 | 23:D1:57:GLU:H | 1.73 | 0.43 |
| 29:D7:40:TRP:CG | 31:DA:459:U:H5" | 2.53 | 0.43 |
| 31:DA:394:A:O2' | 31:DA:395:U:H5' | 2.18 | 0.43 |
| 31:DA:540:C:H2' | 31:DA:541:C:O5' | 2.19 | 0.43 |
| 31:DA:562:U:C4 | 31:DA:2036:C:O4' | 2.72 | 0.43 |
| 31:DA:607:U:C2 | 31:DA:621:A:N1 | 2.86 | 0.43 |
| 31:DA:1010:A:H5' | 46:DU:62:ILE:HG21 | 2.01 | 0.43 |
| 31:DA:1042:G:H3' | 31:DA:1043:C:O4' | 2.17 | 0.43 |
| 31:DA:1399:C:O2' | 31:DA:1400:G:H5' | 2.19 | 0.43 |
| 31:DA:1416:G:O2' | 31:DA:1417:C:P | 2.77 | 0.43 |
| 31:DA:1461:G:C2' | 31:DA:1462:C:H5' | 2.48 | 0.43 |
| 31:DA:1488:G:N2 | 31:DA:1502:C:C6 | 2.87 | 0.43 |
| 31:DA:1670:C:OP2 | 31:DA:2550:G:OP1 | 2.37 | 0.43 |
| 31:DA:1831:G:H2' | 31:DA:1832:C:C6 | 2.54 | 0.43 |
| 31:DA:2192:G:H2' | 31:DA:2193:G:H5' | 2.01 | 0.43 |
| 31:DA:2274:A:C6 | 31:DA:2276:G:C8 | 3.07 | 0.43 |
| 31:DA:2588:G:C2' | 31:DA:2589:A:H5' | 2.49 | 0.43 |
| 31:DA:2637:U:H2' | 31:DA:2638:G:O5' | 2.19 | 0.43 |
| 31:DA:2733:A:H2' | 31:DA:2734:A:C5' | 2.40 | 0.43 |
| 33:DD:59:LYS:HG3 | 33:DD:60:ARG:N | 2.34 | 0.43 |
| 33:DD:67:PHE:CE1 | 33:DD:157:ARG:CZ | 3.02 | 0.43 |
| 34:DE:92:THR:O | 34:DE:93:VAL:HG23 | 2.19 | 0.43 |
| 36:DG:60:LEU:HD22 | 36:DG:63:ILE:CG1 | 2.48 | 0.43 |
| 36:DG:91:ARG:HD2 | 36:DG:91:ARG:C | 2.44 | 0.43 |
| 36:DG:153:ARG:HB3 | 36:DG:153:ARG:CZ | 2.49 | 0.43 |
| 38:DI:73:GLU:O | 38:DI:73:GLU:HG3 | 2.18 | 0.43 |
| 38:DI:75:LEU:HD11 | 38:DI:105:HIS:HE1 | 1.84 | 0.43 |
| 39:DN:97:ARG:O | 39:DN:100:GLU:HB2 | 2.18 | 0.43 |
| 40:DO:43:VAL:HG12 | 40:DO:54:GLU:HA | 2.00 | 0.43 |
| 40:DO:61:VAL:O | 40:DO:61:VAL:HG13 | 2.18 | 0.43 |
| 41:DP:30:THR:O | 41:DP:31:ALA:C | 2.62 | 0.43 |
| 41:DP:88:LEU:HD12 | 41:DP:88:LEU:HA | 1.76 | 0.43 |
| 43:DR:18:LEU:O | 43:DR:19:ALA:C | 2.61 | 0.43 |
| 45:DT:80:SER:CB | 45:DT:81:PRO:HD3 | 2.46 | 0.43 |
| 45:DT:99:LEU:O | 45:DT:99:LEU:HD13 | 2.19 | 0.43 |
| 46:DU:95:LEU:C | 46:DU:97:ASP:N | 2.76 | 0.43 |
| 46:DU:114:LYS:H | 46:DU:114:LYS:HG2 | 1.58 | 0.43 |
| 47:DV:4:ILE:O | 47:DV:39:LEU:HB2 | 2.18 | 0.43 |
| 48:DW:28:SER:C | 48:DW:30:GLU:N | 2.74 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 49:DX:12:VAL:CG1 | 49:DX:27:THR:HG23 | 2.48 | 0.43 |
| 49:DX:65:ARG:NH1 | 49:DX:66:LEU:HA | 2.34 | 0.43 |
| 51:DZ:10:ARG:HH21 | 51:DZ:26:GLY:N | 2.17 | 0.43 |
| 51:DZ:42:VAL:HG13 | 51:DZ:43:GLU:H | 1.83 | 0.43 |
| 1:AA:39:G:C6 | 1:AA:40:C:C5 | 3.07 | 0.42 |
| 1:AA:116:A:OP2 | 1:AA:116:A:C8 | 2.72 | 0.42 |
| 1:AA:165:C:H2' | 1:AA:166:G:H8 | 1.80 | 0.42 |
| 1:AA:166:G:C5 | 1:AA:167:G:N7 | 2.87 | 0.42 |
| 1:AA:473:G:C2 | 1:AA:474:G:C8 | 3.06 | 0.42 |
| 1:AA:502:G:C2 | 1:AA:503:C:C2 | 3.06 | 0.42 |
| 1:AA:559:A:H4' | 1:AA:560:U:O5' | 2.19 | 0.42 |
| 1:AA:939:G:H5" | 7:AG:102:ARG:HH22 | 1.84 | 0.42 |
| 1:AA:981:U:C2 | 1:AA:982:U:C4 | 3.07 | 0.42 |
| 1:AA:1141:C:H6 | 1:AA:1141:C:H3' | 1.83 | 0.42 |
| 1:AA:1498:U:O2' | 1:AA:1499:A:OP2 | 2.31 | 0.42 |
| 2:AB:21:ARG:HB2 | 2:AB:39:ILE:HA | 1.98 | 0.42 |
| 2:AB:238:LEU:H | 2:AB:238:LEU:HD23 | 1.83 | 0.42 |
| 4:AD:101:LEU:HD13 | 4:AD:140:VAL:HG22 | 2.00 | 0.42 |
| 4:AD:170:VAL:HG22 | 4:AD:174:LEU:HD12 | 2.00 | 0.42 |
| 10:AJ:97:GLU:C | 10:AJ:98:ILE:HD12 | 2.45 | 0.42 |
| 11:AK:99:GLN:CD | 11:AK:105:VAL:HG11 | 2.43 | 0.42 |
| 14:AN:13:THR:N | 14:AN:14:PRO:CD | 2.82 | 0.42 |
| 19:AS:27:GLU:HB3 | 19:AS:28:LYS:H | 1.56 | 0.42 |
| 26:B4:6:HIS:HA | 36:BG:67:LYS:HG3 | 2.00 | 0.42 |
| 31:BA:219:G:O2' | 31:BA:220:G:H5' | 2.19 | 0.42 |
| 31:BA:280:C:H2' | 31:BA:281:G:C5' | 2.49 | 0.42 |
| 31:BA:375:C:H2' | 31:BA:376:C:C6 | 2.53 | 0.42 |
| 31:BA:464:U:C2' | 31:BA:465:G:H5' | 2.48 | 0.42 |
| 31:BA:644:A:O2' | 31:BA:645:C:H5" | 2.19 | 0.42 |
| 31:BA:667:U:C2' | 31:BA:668:G:C5' | 2.97 | 0.42 |
| 31:BA:900:A:N3 | 31:BA:900:A:H2' | 2.34 | 0.42 |
| 31:BA:1004:C:O5' | 31:BA:1004:C:H6 | 2.01 | 0.42 |
| 31:BA:1190:G:H5' | 41:BP:35:HIS:HB3 | 2.01 | 0.42 |
| 31:BA:1286:A:O2' | 31:BA:1288:U:P | 2.76 | 0.42 |
| 31:BA:1337:G:H2' | 31:BA:1338:G:H8 | 1.83 | 0.42 |
| 31:BA:1858:G:H1' | 31:BA:1884:A:N6 | 2.33 | 0.42 |
| 31:BA:2011:U:C2' | 31:BA:2012:G:H5' | 2.48 | 0.42 |
| 31:BA:2399:G:C6 | 31:BA:2400:G:C5 | 3.07 | 0.42 |
| 31:BA:2606:C:H2' | 31:BA:2607:G:H5' | 2.01 | 0.42 |
| 31:BA:2849:U:H4' | 31:BA:2868:A:C2 | 2.54 | 0.42 |
| 32:BB:17:C:H2' | 32:BB:18:G:H5' | 2.01 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 32:BB:46:A:C4 | 32:BB:47:C:C5 | 3.07 | 0.42 |
| 33:BD:16:MET:HA | 33:BD:205:VAL:O | 2.19 | 0.42 |
| 33:BD:28:GLU:CB | 33:BD:29:PRO:CD | 2.96 | 0.42 |
| 36:BG:35:GLU:O | 36:BG:35:GLU:HG2 | 2.19 | 0.42 |
| 36:BG:58:GLN:O | 36:BG:61:ALA:HB3 | 2.19 | 0.42 |
| 37:BH:158:HIS:HD2 | 37:BH:170:ARG:O | 1.98 | 0.42 |
| 42:BQ:51:ARG:O | 42:BQ:52:VAL:C | 2.62 | 0.42 |
| 43:BR:38:VAL:HB | 43:BR:39:PRO:HD3 | 2.01 | 0.42 |
| 46:BU:25:TRP:HD1 | 46:BU:26:GLY:H | 1.66 | 0.42 |
| 47:BV:38:LEU:HG | 47:BV:39:LEU:N | 2.35 | 0.42 |
| 50:BY:77:PRO:O | 50:BY:78:ALA:CB | 2.67 | 0.42 |
| 1:CA:15:G:C4 | 1:CA:16:A:C8 | 3.06 | 0.42 |
| 1:CA:72:C:H2' | 1:CA:73:G:C8 | 2.54 | 0.42 |
| 1:CA:191:G:N9 | 20:CT:105:SER:HB3 | 2.32 | 0.42 |
| 1:CA:542:G:C4 | 1:CA:543:C:C5 | 3.06 | 0.42 |
| 1:CA:663:A:O3' | 18:CR:64:ARG:NH2 | 2.48 | 0.42 |
| 1:CA:1053:G:C4 | 1:CA:1199:U:C5 | 3.07 | 0.42 |
| 1:CA:1164:G:N2 | 1:CA:1165:C:C2 | 2.87 | 0.42 |
| 1:CA:1188:A:H2' | 1:CA:1189:C:H5' | 2.00 | 0.42 |
| 1:CA:1272:G:C5 | 1:CA:1273:G:C8 | 3.07 | 0.42 |
| 1:CA:1350:A:H2' | 1:CA:1351:U:C6 | 2.54 | 0.42 |
| 2:CB:54:THR:HG21 | 2:CB:201:ILE:HD11 | 2.00 | 0.42 |
| 2:CB:165:VAL:HG23 | 2:CB:166:ASP:N | 2.32 | 0.42 |
| 4:CD:6:GLY:O | 4:CD:7:PRO:C | 2.61 | 0.42 |
| 12:CL:34:ARG:CG | 12:CL:35:GLY:N | 2.82 | 0.42 |
| 13:CM:40:ASN:HA | 13:CM:41:PRO:HD3 | 1.78 | 0.42 |
| 15:CO:2:PRO:HB2 | 15:CO:3:ILE:H | 1.58 | 0.42 |
| 16:CP:39:TYR:CE1 | 16:CP:41:PRO:HA | 2.54 | 0.42 |
| 17:CQ:31:LEU:HG | 17:CQ:32:TYR:CE2 | 2.54 | 0.42 |
| 20:CT:38:LYS:O | 20:CT:39:LYS:C | 2.62 | 0.42 |
| 20:CT:46:GLU:CD | 20:CT:48:LYS:HE2 | 2.44 | 0.42 |
| 28:D6:14:THR:C | 28:D6:15:GLU:O | 2.61 | 0.42 |
| 30:D8:5:LYS:HE2 | 31:DA:254:G:N7 | 2.33 | 0.42 |
| 31:DA:26:G:C6 | 31:DA:27:G:N1 | 2.86 | 0.42 |
| 31:DA:118:A:H3' | 31:DA:119:A:H5" | 1.99 | 0.42 |
| 31:DA:738:G:C6 | 31:DA:739:G:C2 | 3.07 | 0.42 |
| 31:DA:1140:C:O3' | 39:DN:25:ARG:NH1 | 2.51 | 0.42 |
| 31:DA:1190:G:H5' | 41:DP:35:HIS:CA | 2.49 | 0.42 |
| 31:DA:1439:A:C2 | 31:DA:1553:A:C5 | 3.07 | 0.42 |
| 31:DA:1478:G:H2' | 31:DA:1479:G:H5' | 1.99 | 0.42 |
| 31:DA:1833:U:C4 | 31:DA:1834:U:C5 | 3.07 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1851:U:C2' | 31:DA:1852:C:H5' | 2.49 | 0.42 |
| 31:DA:1940:U:C4 | 31:DA:1964:G:H4' | 2.54 | 0.42 |
| 32:DB:7:G:O5' | 44:DS:29:PHE:HE1 | 2.00 | 0.42 |
| 33:DD:24:ILE:O | 33:DD:24:ILE:HG23 | 2.18 | 0.42 |
| 33:DD:186:HIS:HD2 | 33:DD:188:GLU:HB2 | 1.81 | 0.42 |
| 34:DE:55:ASN:H | 34:DE:72:VAL:HG11 | 1.84 | 0.42 |
| 35:DF:126:VAL:HG11 | 35:DF:142:TRP:HH2 | 1.83 | 0.42 |
| 36:DG:35:GLU:O | 36:DG:35:GLU:HG2 | 2.19 | 0.42 |
| 36:DG:96:ARG:O | 36:DG:99:MET:HB3 | 2.18 | 0.42 |
| 37:DH:85:LYS:HE3 | 37:DH:141:VAL:O | 2.19 | 0.42 |
| 38:DI:10:GLU:C | 38:DI:12:LEU:N | 2.77 | 0.42 |
| 38:DI:114:LEU:O | 38:DI:115:ALA:C | 2.60 | 0.42 |
| 40:DO:111:PHE:O | 40:DO:112:MET:C | 2.62 | 0.42 |
| 42:DQ:83:MET:HB3 | 42:DQ:83:MET:HE2 | 1.84 | 0.42 |
| 43:DR:42:LYS:O | 43:DR:45:ARG:HG3 | 2.20 | 0.42 |
| 43:DR:48:VAL:O | 43:DR:49:ASP:C | 2.63 | 0.42 |
| 46:DU:69:CYS:HB3 | 46:DU:106:PHE:HZ | 1.83 | 0.42 |
| 46:DU:83:LEU:HD12 | 46:DU:83:LEU:HA | 1.76 | 0.42 |
| 47:DV:78:LYS:HG2 | 47:DV:79:VAL:N | 2.34 | 0.42 |
| 48:DW:35:ILE:O | 48:DW:36:LEU:C | 2.61 | 0.42 |
| 49:DX:13:LEU:N | 49:DX:13:LEU:HD23 | 2.34 | 0.42 |
| 49:DX:82:GLN:CD | 49:DX:83:VAL:N | 2.77 | 0.42 |
| 50:DY:7:VAL:CB | 50:DY:8:LYS:NZ | 2.79 | 0.42 |
| 50:DY:20:TYR:CD2 | 50:DY:41:GLY:CA | 3.00 | 0.42 |
| 51:DZ:15:PRO:O | 51:DZ:19:ARG:HD2 | 2.19 | 0.42 |
| 1:AA:44:G:H2' | 1:AA:45:U:O4' | 2.19 | 0.42 |
| 1:AA:90:U:H3' | 1:AA:90:U:C6 | 2.53 | 0.42 |
| 1:AA:196:A:N3 | 1:AA:222:U:H1' | 2.33 | 0.42 |
| 1:AA:488:C:O5' | 1:AA:488:C:H6 | 2.02 | 0.42 |
| 1:AA:491:G:C2 | 1:AA:492:G:C4 | 3.07 | 0.42 |
| 1:AA:515:G:N3 | 1:AA:537:G:C2 | 2.88 | 0.42 |
| 1:AA:668:G:O2' | 15:AO:46:HIS:CD2 | 2.73 | 0.42 |
| 1:AA:677:U:H3 | 1:AA:714:G:N2 | 2.17 | 0.42 |
| 1:AA:720:C:H5" | 1:AA:721:G:O5' | 2.19 | 0.42 |
| 1:AA:763:G:C5 | 1:AA:764:C:C5 | 3.07 | 0.42 |
| 1:AA:946:A:O2' | 1:AA:1333:A:N3 | 2.43 | 0.42 |
| 1:AA:1061:G:C4 | 1:AA:1197:G:N2 | 2.88 | 0.42 |
| 1:AA:1320:C:H5' | 19:AS:70:LYS:CG | 2.49 | 0.42 |
| 2:AB:97:TRP:HH2 | 2:AB:176:GLU:HG3 | 1.84 | 0.42 |
| 2:AB:144:ARG:O | 2:AB:145:LEU:C | 2.62 | 0.42 |
| 3:AC:159:GLY:HA2 | 3:AC:193:TYR:CD1 | 2.54 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 4:AD:132:ARG:H | 4:AD:132:ARG:HG3 | 1.50 | 0.42 |
| 5:AE:108:ALA:O | 5:AE:109:ILE:C | 2.62 | 0.42 |
| 8:AH:36:LEU:HA | 8:AH:39:LEU:HD23 | 2.01 | 0.42 |
| 12:AL:55:VAL:HG13 | 12:AL:68:ALA:O | 2.20 | 0.42 |
| 14:AN:29:ARG:HD3 | 14:AN:40:CYS:SG | 2.59 | 0.42 |
| 18:AR:66:LEU:HD11 | 18:AR:70:ILE:HD11 | 2.00 | 0.42 |
| 20:AT:23:ARG:O | 20:AT:27:LYS:HB2 | 2.19 | 0.42 |
| 24:B2:24:LEU:HA | 24:B2:24:LEU:HD12 | 1.78 | 0.42 |
| 28:B6:28:ARG:O | 28:B6:32:ASN:HB3 | 2.18 | 0.42 |
| 31:BA:92:A:C2' | 31:BA:93:G:O4' | 2.67 | 0.42 |
| 31:BA:150:C:H2' | 31:BA:151:C:H6 | 1.84 | 0.42 |
| 31:BA:207:A:H2' | 31:BA:208:C:O4' | 2.18 | 0.42 |
| 31:BA:343:C:H2' | 31:BA:343:C:O2 | 2.18 | 0.42 |
| 31:BA:460:A:C2 | 31:BA:470:A:C4 | 3.07 | 0.42 |
| 31:BA:705:A:O2' | 31:BA:706:A:H5' | 2.18 | 0.42 |
| 31:BA:959:A:N1 | 31:BA:960:A:C2 | 2.87 | 0.42 |
| 31:BA:1210:A:H5" | 31:BA:1211:U:C3' | 2.48 | 0.42 |
| 31:BA:1438:U:O2' | 31:BA:1439:A:H5' | 2.20 | 0.42 |
| 31:BA:1510:G:O2' | 31:BA:1511:C:H5' | 2.19 | 0.42 |
| 31:BA:1639:U:H4' | 31:BA:2699:C:H4' | 2.01 | 0.42 |
| 31:BA:2017:U:H5" | 31:BA:2018:G:P | 2.59 | 0.42 |
| 31:BA:2306:C:OP2 | 31:BA:2307:G:C8 | 2.72 | 0.42 |
| 31:BA:2321:G:N3 | 31:BA:2321:G:H2' | 2.34 | 0.42 |
| 31:BA:2410:G:C2 | 31:BA:2411:A:H1' | 2.55 | 0.42 |
| 31:BA:2655:G:O2' | 31:BA:2656:U:H5 | 2.01 | 0.42 |
| 33:BD:154:LYS:C | 33:BD:155:LEU:HD12 | 2.44 | 0.42 |
| 33:BD:228:PRO:HD3 | 33:BD:235:GLY:CA | 2.46 | 0.42 |
| 34:BE:36:ARG:NH2 | 34:BE:86:PRO:HD2 | 2.33 | 0.42 |
| 34:BE:61:ARG:N | 34:BE:62:PRO:HD2 | 2.14 | 0.42 |
| 34:BE:97:LYS:O | 34:BE:100:GLU:HG3 | 2.19 | 0.42 |
| 34:BE:132:HIS:CG | 34:BE:132:HIS:O | 2.72 | 0.42 |
| 37:BH:32:GLU:C | 37:BH:33:LEU:HD23 | 2.44 | 0.42 |
| 37:BH:89:ILE:CD1 | 37:BH:129:THR:HB | 2.41 | 0.42 |
| 37:BH:157:TYR:CD1 | 37:BH:170:ARG:O | 2.72 | 0.42 |
| 38:BI:67:ARG:O | 38:BI:68:LEU:CB | 2.67 | 0.42 |
| 38:BI:69:LYS:HG2 | 38:BI:69:LYS:O | 2.20 | 0.42 |
| 41:BP:5:ASP:HB3 | 41:BP:6:LEU:H | 1.67 | 0.42 |
| 42:BQ:55:VAL:HG22 | 42:BQ:56:ARG:N | 2.34 | 0.42 |
| 44:BS:28:VAL:O | 44:BS:29:PHE:CB | 2.67 | 0.42 |
| 45:BT:50:ILE:HD13 | 45:BT:64:ARG:HB3 | 2.01 | 0.42 |
| 46:BU:92:ARG:NH2 | 47:BV:10:LYS:HG2 | 2.34 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 47:BV:19:LYS:NZ | 47:BV:20:LEU:H | 2.16 | 0.42 |
| 50:BY:19:LYS:HB3 | 50:BY:20:TYR:HD1 | 1.82 | 0.42 |
| 51:BZ:5:LEU:HD21 | 51:BZ:43:GLU:HB3 | 2.00 | 0.42 |
| 1:CA:371:G:H5" | 1:CA:372:C:OP2 | 2.19 | 0.42 |
| 1:CA:390:C:H3' | 1:CA:390:C:H6 | 1.84 | 0.42 |
| 1:CA:488:C:H6 | 1:CA:488:C:O5' | 2.02 | 0.42 |
| 1:CA:552:U:H4' | 12:CL:86:ARG:HG2 | 2.01 | 0.42 |
| 1:CA:771:G:C6 | 1:CA:772:U:C4 | 3.08 | 0.42 |
| 1:CA:783:C:C2' | 1:CA:784:C:H5' | 2.48 | 0.42 |
| 1:CA:929:G:C6 | 1:CA:930:C:C4 | 3.06 | 0.42 |
| 1:CA:1030(A):G:O2' | 1:CA:1030(C):G:N7 | 2.46 | 0.42 |
| 1:CA:1346:A:C8 | 1:CA:1348:U:O2 | 2.72 | 0.42 |
| 1:CA:1452:C:H5' | 1:CA:1456:G:C5 | 2.54 | 0.42 |
| 1:CA:1457:G:C2 | 1:CA:1458:G:C8 | 3.07 | 0.42 |
| 5:CE:75:THR:OG1 | 5:CE:76:ILE:N | 2.51 | 0.42 |
| 9:CI:45:ALA:O | 9:CI:78:LYS:HE3 | 2.19 | 0.42 |
| 11:CK:62:GLN:C | 11:CK:64:ALA:N | 2.77 | 0.42 |
| 11:CK:99:GLN:CD | 11:CK:105:VAL:HG11 | 2.44 | 0.42 |
| 12:CL:92:ASP:O | 12:CL:93:LEU:HD23 | 2.19 | 0.42 |
| 13:CM:66:LEU:HB2 | 13:CM:67:GLU:H | 1.53 | 0.42 |
| 17:CQ:5:VAL:HG23 | 17:CQ:60:ILE:HG13 | 2.00 | 0.42 |
| 18:CR:74:ARG:HE | 18:CR:81:PHE:HA | 1.82 | 0.42 |
| 19:CS:12:ASP:OD1 | 19:CS:37:ARG:HD2 | 2.19 | 0.42 |
| 24:D2:23:LYS:N | 49:DX:5:TYR:HD1 | 2.17 | 0.42 |
| 25:D3:19:GLN:O | 25:D3:23:LEU:HD12 | 2.19 | 0.42 |
| 31:DA:271(T):C:C2 | 31:DA:271(U):G:C8 | 3.07 | 0.42 |
| 31:DA:518:G:H4' | 48:DW:18:ARG:CZ | 2.48 | 0.42 |
| 31:DA:768:G:C4 | 31:DA:769:G:C8 | 3.07 | 0.42 |
| 31:DA:877:U:O2' | 31:DA:878:A:H5" | 2.19 | 0.42 |
| 31:DA:1262:A:C4 | 31:DA:1263:U:C5 | 3.06 | 0.42 |
| 31:DA:1987:G:H2' | 31:DA:1988:C:H6 | 1.83 | 0.42 |
| 31:DA:2199:A:H1' | 38:DI:28:ASN:HD22 | 1.82 | 0.42 |
| 31:DA:2290:G:N1 | 31:DA:2343:C:O2 | 2.52 | 0.42 |
| 31:DA:2308:G:H21 | 36:DG:79:ASN:ND2 | 2.18 | 0.42 |
| 31:DA:2723:C:O5' | 31:DA:2723:C:H6 | 2.02 | 0.42 |
| 31:DA:2813:A:C6 | 31:DA:2814:C:C4 | 3.07 | 0.42 |
| 32:DB:21:G:C6 | 32:DB:63:G:C6 | 3.06 | 0.42 |
| 32:DB:48:A:OP1 | 44:DS:93:LYS:HB3 | 2.19 | 0.42 |
| 33:DD:25:THR:HB | 33:DD:82:ILE:H | 1.83 | 0.42 |
| 35:DF:53:THR:HG23 | 35:DF:55:GLY:H | 1.84 | 0.42 |
| 35:DF:64:ILE:HG21 | 35:DF:64:ILE:HD13 | 1.80 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 38:DI:145:VAL:HG12 | 38:DI:146:ALA:N | 2.34 | 0.42 |
| 39:DN:16:ILE:HD11 | 39:DN:26:LEU:HD11 | 2.01 | 0.42 |
| 39:DN:46:VAL:CG1 | 39:DN:48:MET:HG3 | 2.48 | 0.42 |
| 41:DP:13:ASN:N | 41:DP:13:ASN:ND2 | 2.66 | 0.42 |
| 42:DQ:62:GLY:O | 51:DZ:178:GLU:HG2 | 2.19 | 0.42 |
| 43:DR:55:ALA:HB2 | 43:DR:79:LEU:HD13 | 2.01 | 0.42 |
| 45:DT:91:ARG:HB2 | 45:DT:92:GLY:H | 1.61 | 0.42 |
| 46:DU:87:GLY:C | 47:DV:52:VAL:HG13 | 2.44 | 0.42 |
| 47:DV:53:GLU:O | 47:DV:53:GLU:HG3 | 2.19 | 0.42 |
| 49:DX:84:ALA:C | 49:DX:86:GLY:N | 2.76 | 0.42 |
| 50:DY:76:CYS:CB | 50:DY:77:PRO:HD2 | 2.49 | 0.42 |
| 51:DZ:76:LEU:HD23 | 51:DZ:76:LEU:HA | 1.82 | 0.42 |
| 1:AA:52:G:H2' | 1:AA:53:A:H5' | 1.98 | 0.42 |
| 1:AA:102:G:C5 | 1:AA:103:C:C5 | 3.07 | 0.42 |
| 1:AA:231:G:C2' | 1:AA:232:G:H5' | 2.50 | 0.42 |
| 1:AA:233:C:N3 | 1:AA:234:C:C5 | 2.86 | 0.42 |
| 1:AA:328:C:H4' | 1:AA:329:A:C5' | 2.48 | 0.42 |
| 1:AA:451:A:C6 | 1:AA:481:G:C5 | 3.07 | 0.42 |
| 1:AA:510:A:H5" | 1:AA:511:C:OP1 | 2.20 | 0.42 |
| 1:AA:760:G:C2' | 1:AA:761:G:H5' | 2.49 | 0.42 |
| 1:AA:1173:G:C5 | 1:AA:1174:G:N7 | 2.87 | 0.42 |
| 1:AA:1413:A:C2 | 1:AA:1414:U:C2 | 3.06 | 0.42 |
| 1:AA:1486:G:H2' | 1:AA:1487:G:O4' | 2.19 | 0.42 |
| 2:AB:11:LEU:HD11 | 2:AB:217:ARG:HH22 | 1.84 | 0.42 |
| 2:AB:102:LEU:N | 2:AB:102:LEU:CD1 | 2.81 | 0.42 |
| 2:AB:140:HIS:O | 2:AB:143:GLU:HB2 | 2.19 | 0.42 |
| 3:AC:79:ARG:O | 3:AC:79:ARG:HG3 | 2.20 | 0.42 |
| 3:AC:138:VAL:HG23 | 3:AC:151:VAL:HG23 | 2.02 | 0.42 |
| 4:AD:79:PHE:CZ | 4:AD:204:ILE:HD13 | 2.55 | 0.42 |
| 4:AD:101:LEU:C | 4:AD:103:ASN:H | 2.27 | 0.42 |
| 4:AD:188:LEU:HA | 4:AD:189:PRO:HD2 | 1.88 | 0.42 |
| 5:AE:6:PHE:HB2 | 5:AE:34:VAL:HG13 | 2.00 | 0.42 |
| 8:AH:107:LEU:HD23 | 8:AH:107:LEU:N | 2.34 | 0.42 |
| 15:AO:40:SER:O | 15:AO:44:LYS:HB2 | 2.19 | 0.42 |
| 15:AO:43:LEU:O | 15:AO:44:LYS:C | 2.62 | 0.42 |
| 18:AR:46:GLU:OE2 | 18:AR:46:GLU:HA | 2.19 | 0.42 |
| 22:B0:84:LEU:HD12 | 22:B0:84:LEU:N | 2.34 | 0.42 |
| 23:B1:16:ASN:CB | 23:B1:46:LEU:HG | 2.33 | 0.42 |
| 28:B6:37:ARG:O | 28:B6:48:VAL:O | 2.37 | 0.42 |
| 31:BA:355:G:C2 | 31:BA:356:G:C8 | 3.07 | 0.42 |
| 31:BA:494:G:H21 | 48:BW:57:ASN:HD21 | 1.67 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:770:G:N3 | 31:BA:1354:A:H2 | 2.17 | 0.42 |
| 31:BA:877:U:C2' | 31:BA:878:A:H5" | 2.49 | 0.42 |
| 31:BA:1275:A:C8 | 43:BR:16:HIS:CD2 | 3.07 | 0.42 |
| 31:BA:1646:C:H5" | 31:BA:1647:G:H5" | 2.01 | 0.42 |
| 31:BA:1851:U:H2' | 31:BA:1852:C:H5' | 2.01 | 0.42 |
| 31:BA:2074:U:H2' | 31:BA:2075:U:C6 | 2.54 | 0.42 |
| 31:BA:2287:A:H2 | 31:BA:2346:A:N1 | 2.16 | 0.42 |
| 31:BA:2395:C:H5' | 31:BA:2395:C:C6 | 2.41 | 0.42 |
| 31:BA:2632:A:N3 | 34:BE:61:ARG:HD2 | 2.35 | 0.42 |
| 32:BB:10:C:H2' | 32:BB:11:C:H5' | 2.02 | 0.42 |
| 32:BB:18:G:H1 | 32:BB:65:C:H42 | 1.66 | 0.42 |
| 33:BD:183:ARG:HD2 | 33:BD:270:ILE:HG22 | 2.01 | 0.42 |
| 35:BF:23:ASP:O | 35:BF:24:LEU:HD22 | 2.19 | 0.42 |
| 35:BF:96:ASP:OD1 | 35:BF:96:ASP:C | 2.62 | 0.42 |
| 36:BG:15:VAL:HG13 | 36:BG:175:LEU:HD13 | 2.00 | 0.42 |
| 36:BG:19:LEU:HD13 | 36:BG:32:PRO:HG2 | 2.02 | 0.42 |
| 37:BH:125:VAL:HG12 | 37:BH:127:GLU:O | 2.20 | 0.42 |
| 39:BN:89:LYS:O | 39:BN:93:THR:HG22 | 2.18 | 0.42 |
| 41:BP:30:THR:O | 41:BP:33:ARG:N | 2.49 | 0.42 |
| 41:BP:101:VAL:HB | 41:BP:107:LYS:N | 2.35 | 0.42 |
| 44:BS:67:ARG:N | 44:BS:69:VAL:HG12 | 2.31 | 0.42 |
| 46:BU:20:LEU:HB3 | 46:BU:39:LEU:HD11 | 2.01 | 0.42 |
| 49:BX:9:LEU:HD12 | 49:BX:30:VAL:C | 2.44 | 0.42 |
| 49:BX:65:ARG:C | 49:BX:65:ARG:CD | 2.91 | 0.42 |
| 50:BY:45:VAL:HG21 | 50:BY:62:GLU:N | 2.34 | 0.42 |
| 1:CA:189(K):U:O5' | 1:CA:189(K):U:H6 | 2.02 | 0.42 |
| 1:CA:229:U:C2' | 1:CA:230:G:H5' | 2.50 | 0.42 |
| 1:CA:376:G:C4' | 16:CP:5:ARG:HH11 | 2.18 | 0.42 |
| 1:CA:428:G:OP2 | 4:CD:7:PRO:HG2 | 2.19 | 0.42 |
| 1:CA:561:U:HO2' | 1:CA:562:C:P | 2.40 | 0.42 |
| 1:CA:604:G:C5 | 1:CA:605:U:C5 | 3.07 | 0.42 |
| 1:CA:720:C:O5' | 1:CA:720:C:H6 | 2.02 | 0.42 |
| 1:CA:764:C:H2' | 1:CA:765:G:C8 | 2.52 | 0.42 |
| 1:CA:928:G:C2 | 1:CA:1390:U:O2 | 2.72 | 0.42 |
| 1:CA:1030(D):A:N7 | 1:CA:1031:G:N3 | 2.66 | 0.42 |
| 1:CA:1086:U:C6 | 1:CA:1087:G:C8 | 3.07 | 0.42 |
| 1:CA:1095:U:H2' | 1:CA:1096:C:H6 | 1.84 | 0.42 |
| 2:CB:97:TRP:HH2 | 2:CB:176:GLU:HG3 | 1.84 | 0.42 |
| 2:CB:102:LEU:N | 2:CB:102:LEU:CD1 | 2.83 | 0.42 |
| 2:CB:139:LYS:O | 2:CB:143:GLU:HG2 | 2.19 | 0.42 |
| 2:CB:140:HIS:O | 2:CB:143:GLU:HB2 | 2.19 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 4:CD:118:ARG:C | 4:CD:120:LEU:N | 2.76 | 0.42 |
| 4:CD:192:GLU:CD | 4:CD:192:GLU:H | 2.28 | 0.42 |
| 5:CE:91:LEU:HD12 | 5:CE:120:THR:HG22 | 2.01 | 0.42 |
| 8:CH:97:VAL:HG13 | 8:CH:98:LYS:HG3 | 2.00 | 0.42 |
| 10:CJ:49:VAL:O | 10:CJ:60:ARG:HB2 | 2.20 | 0.42 |
| 10:CJ:97:GLU:C | 10:CJ:98:ILE:HD12 | 2.44 | 0.42 |
| 11:CK:23:ALA:HB1 | 11:CK:88:GLY:HA3 | 2.01 | 0.42 |
| 12:CL:18:VAL:HG23 | 12:CL:19:ARG:N | 2.33 | 0.42 |
| 12:CL:41:ARG:CG | 12:CL:42:THR:N | 2.77 | 0.42 |
| 14:CN:21:TYR:CD2 | 14:CN:22:THR:O | 2.72 | 0.42 |
| 16:CP:49:LEU:HG | 16:CP:50:LYS:N | 2.34 | 0.42 |
| 22:D0:2:ALA:HA | 31:DA:2494:G:OP1 | 2.19 | 0.42 |
| 24:D2:46:GLN:NE2 | 24:D2:47:ASN:CA | 2.82 | 0.42 |
| 25:D3:18:ASP:HB2 | 25:D3:49:LYS:CE | 2.48 | 0.42 |
| 28:D6:16:CYS:O | 28:D6:17:LYS:CB | 2.67 | 0.42 |
| 28:D6:36:LEU:O | 28:D6:37:ARG:HD2 | 2.18 | 0.42 |
| 31:DA:68:G:H2' | 31:DA:69:C:O4' | 2.19 | 0.42 |
| 31:DA:154:G:N1 | 31:DA:154(A):C:N4 | 2.67 | 0.42 |
| 31:DA:267:C:H6 | 31:DA:267:C:C5' | 2.33 | 0.42 |
| 31:DA:573:G:H1 | 31:DA:2030:A:H3' | 1.82 | 0.42 |
| 31:DA:1004:C:H6 | 31:DA:1004:C:O5' | 2.01 | 0.42 |
| 31:DA:1027:A:N7 | 31:DA:1126:A:C2 | 2.87 | 0.42 |
| 31:DA:1242:A:N1 | 41:DP:8:PRO:HG3 | 2.34 | 0.42 |
| 31:DA:1504:C:HO2' | 31:DA:1505:C:P | 2.42 | 0.42 |
| 31:DA:1570:A:H2' | 31:DA:1571:A:C8 | 2.54 | 0.42 |
| 31:DA:1637:A:C2 | 31:DA:1638:C:C2 | 3.08 | 0.42 |
| 31:DA:1638:C:H5" | 31:DA:2710:C:O2' | 2.19 | 0.42 |
| 31:DA:1667:G:H4' | 31:DA:1668:A:OP1 | 2.20 | 0.42 |
| 31:DA:2206:G:N3 | 31:DA:2207:G:H5' | 2.33 | 0.42 |
| 31:DA:2332:U:H5' | 31:DA:2336:A:N6 | 2.33 | 0.42 |
| 31:DA:2772:C:H2' | 31:DA:2773:C:C6 | 2.55 | 0.42 |
| 31:DA:2818:G:N2 | 31:DA:2819:G:C4 | 2.87 | 0.42 |
| 31:DA:2887:U:H2' | 31:DA:2888:C:C6 | 2.54 | 0.42 |
| 32:DB:15:A:H5' | 32:DB:16:G:N7 | 2.29 | 0.42 |
| 33:DD:143:HIS:CD2 | 33:DD:144:ALA:N | 2.87 | 0.42 |
| 36:DG:58:GLN:O | 36:DG:61:ALA:HB3 | 2.19 | 0.42 |
| 39:DN:28:THR:N | 39:DN:106:MET:HE3 | 2.34 | 0.42 |
| 40:DO:16:ALA:HA | 40:DO:46:ALA:CB | 2.49 | 0.42 |
| 41:DP:21:ARG:O | 41:DP:23:PRO:HD3 | 2.20 | 0.42 |
| 42:DQ:23:GLY:C | 42:DQ:100:GLY:HA3 | 2.41 | 0.42 |
| 43:DR:18:LEU:HD13 | 43:DR:19:ALA:N | 2.35 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 44:DS:77:ALA:O | 44:DS:78:LEU:C | 2.62 | 0.42 |
| 46:DU:3:ARG:NH2 | 46:DU:5:LYS:HG2 | 2.34 | 0.42 |
| 46:DU:52:ARG:HD3 | 46:DU:52:ARG:HA | 1.58 | 0.42 |
| 47:DV:11:GLN:C | 47:DV:12:TYR:CD2 | 2.97 | 0.42 |
| 47:DV:83:ARG:NH1 | 47:DV:83:ARG:HG2 | 2.34 | 0.42 |
| 48:DW:74:ALA:O | 48:DW:75:TYR:HB3 | 2.20 | 0.42 |
| 51:DZ:130:PRO:HA | 51:DZ:133:ILE:HD11 | 2.01 | 0.42 |
| 1:AA:131:C:H2' | 1:AA:132:C:H6 | 1.81 | 0.42 |
| 1:AA:233:C:C2 | 1:AA:234:C:C5 | 3.08 | 0.42 |
| 1:AA:274:A:H4' | 1:AA:275:G:OP1 | 2.18 | 0.42 |
| 1:AA:473:G:N3 | 1:AA:474:G:C8 | 2.87 | 0.42 |
| 1:AA:522:C:H5" | 12:AL:120:TYR:OH | 2.19 | 0.42 |
| 1:AA:640:A:N3 | 8:AH:115:SER:CB | 2.83 | 0.42 |
| 1:AA:937:A:H1' | 1:AA:1379:G:N2 | 2.34 | 0.42 |
| 1:AA:1054:C:OP1 | 1:AA:1197:G:OP2 | 2.37 | 0.42 |
| 1:AA:1352:C:OP1 | 21:AU:3:LYS:NZ | 2.47 | 0.42 |
| 2:AB:84:GLU:OE1 | 2:AB:219:VAL:HB | 2.18 | 0.42 |
| 5:AE:128:PRO:O | 5:AE:129:ILE:C | 2.62 | 0.42 |
| 7:AG:12:LEU:HD13 | 7:AG:24:THR:OG1 | 2.19 | 0.42 |
| 10:AJ:52:GLY:C | 10:AJ:54:PHE:H | 2.27 | 0.42 |
| 15:AO:39:LEU:O | 15:AO:40:SER:C | 2.63 | 0.42 |
| 15:AO:39:LEU:HD23 | 15:AO:39:LEU:HA | 1.74 | 0.42 |
| 16:AP:72:ARG:HH21 | 16:AP:73:LEU:CD2 | 2.26 | 0.42 |
| 19:AS:4:SER:O | 19:AS:5:LEU:HB2 | 2.18 | 0.42 |
| 23:B1:37:ILE:O | 23:B1:38:SER:HB2 | 2.20 | 0.42 |
| 24:B2:41:ILE:CG1 | 31:BA:94(A):G:N2 | 2.82 | 0.42 |
| 25:B3:13:ILE:HD12 | 31:BA:989:G:N7 | 2.34 | 0.42 |
| 27:B5:38:ALA:HB1 | 27:B5:39:MET:H | 1.59 | 0.42 |
| 30:B8:39:LYS:O | 30:B8:39:LYS:CG | 2.67 | 0.42 |
| 30:B8:61:LEU:HD23 | 30:B8:61:LEU:HA | 1.62 | 0.42 |
| 31:BA:378:C:C2' | 31:BA:379:G:H5' | 2.49 | 0.42 |
| 31:BA:475:U:O2 | 31:BA:505:A:H2 | 2.02 | 0.42 |
| 31:BA:478:A:C6 | 31:BA:480:A:C6 | 3.07 | 0.42 |
| 31:BA:513:A:N1 | 31:BA:514:A:C5 | 2.87 | 0.42 |
| 31:BA:776:G:H4' | 31:BA:777:A:O5' | 2.19 | 0.42 |
| 31:BA:790:C:O2' | 31:BA:791:C:C5' | 2.67 | 0.42 |
| 31:BA:933:A:C5 | 31:BA:934:G:C8 | 3.08 | 0.42 |
| 31:BA:1291:C:H2' | 31:BA:1292:U:C6 | 2.54 | 0.42 |
| 31:BA:1318:C:O2 | 31:BA:1318:C:H2' | 2.19 | 0.42 |
| 31:BA:1348:G:C2' | 31:BA:1349:A:H5' | 2.48 | 0.42 |
| 31:BA:1464:C:C2' | 31:BA:1528:A:H8 | 2.33 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:1637:A:H4' | 31:BA:2711:A:O2' | 2.18 | 0.42 |
| 31:BA:1665:A:C2' | 31:BA:1666:G:H5' | 2.50 | 0.42 |
| 31:BA:1988:C:O2' | 31:BA:1989:G:H5' | 2.20 | 0.42 |
| 31:BA:2504:U:O2 | 31:BA:2504:U:C2' | 2.65 | 0.42 |
| 31:BA:2552:U:C2 | 31:BA:2554:U:H5' | 2.55 | 0.42 |
| 31:BA:2552:U:H2' | 31:BA:2554:U:OP2 | 2.20 | 0.42 |
| 31:BA:2759:G:O2' | 31:BA:2760:C:H5' | 2.19 | 0.42 |
| 32:BB:79:C:H2' | 32:BB:80:U:O4' | 2.19 | 0.42 |
| 33:BD:25:THR:O | 33:BD:27:THR:CB | 2.52 | 0.42 |
| 33:BD:48:ARG:HH11 | 33:BD:48:ARG:CG | 2.30 | 0.42 |
| 33:BD:235:GLY:C | 33:BD:237:GLU:HG2 | 2.42 | 0.42 |
| 35:BF:53:THR:HG23 | 35:BF:55:GLY:CA | 2.50 | 0.42 |
| 35:BF:63:LYS:HA | 35:BF:76:GLY:O | 2.19 | 0.42 |
| 35:BF:88:VAL:HG11 | 35:BF:91:GLY:HA3 | 2.01 | 0.42 |
| 38:BI:15:VAL:CG2 | 38:BI:16:GLY:N | 2.82 | 0.42 |
| 40:BO:16:ALA:HA | 40:BO:46:ALA:CB | 2.50 | 0.42 |
| 43:BR:13:HIS:CE1 | 43:BR:15:SER:OG | 2.73 | 0.42 |
| 45:BT:35:LYS:C | 45:BT:37:GLY:N | 2.77 | 0.42 |
| 46:BU:8:VAL:HG11 | 46:BU:12:ARG:NE | 2.31 | 0.42 |
| 46:BU:92:ARG:O | 46:BU:95:LEU:N | 2.52 | 0.42 |
| 47:BV:75:PHE:CE1 | 47:BV:89:GLN:CB | 2.85 | 0.42 |
| 49:BX:33:LYS:C | 49:BX:35:THR:HG22 | 2.44 | 0.42 |
| 49:BX:82:GLN:CD | 49:BX:83:VAL:N | 2.77 | 0.42 |
| 49:BX:89:ILE:HD12 | 49:BX:92:LEU:HD12 | 2.01 | 0.42 |
| 50:BY:81:LYS:NZ | 50:BY:97:ARG:O | 2.42 | 0.42 |
| 1:CA:37:U:O2' | 1:CA:500:G:H4' | 2.20 | 0.42 |
| 1:CA:262:A:N6 | 1:CA:263:A:N6 | 2.67 | 0.42 |
| 1:CA:391:G:O6 | 1:CA:392:G:C6 | 2.72 | 0.42 |
| 1:CA:587:G:C2 | 1:CA:755:G:C5 | 3.07 | 0.42 |
| 1:CA:976:G:C5' | 1:CA:1358:U:O2' | 2.64 | 0.42 |
| 1:CA:1037:C:H6 | 1:CA:1037:C:O5' | 2.02 | 0.42 |
| 1:CA:1500:A:OP2 | 1:CA:1505:G:OP1 | 2.37 | 0.42 |
| 3:CC:138:VAL:HG23 | 3:CC:151:VAL:HG23 | 2.00 | 0.42 |
| 4:CD:146:ILE:HD12 | 4:CD:146:ILE:H | 1.84 | 0.42 |
| 5:CE:72:GLN:O | 5:CE:73:ASN:CB | 2.58 | 0.42 |
| 5:CE:108:ALA:O | 5:CE:109:ILE:C | 2.62 | 0.42 |
| 8:CH:100:ILE:HA | 8:CH:101:PRO:HD3 | 1.83 | 0.42 |
| 9:CI:56:LEU:O | 9:CI:56:LEU:HD23 | 2.20 | 0.42 |
| 11:CK:21:ILE:HA | 11:CK:30:VAL:HG12 | 2.00 | 0.42 |
| 12:CL:90:VAL:C | 12:CL:92:ASP:H | 2.22 | 0.42 |
| 13:CM:74:VAL:O | 13:CM:78:ILE:HG13 | 2.19 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 22:D0:20:ARG:HG3 | 31:DA:2356:C:H4' | 2.02 | 0.42 |
| 22:D0:51:VAL:C | 22:D0:62:LEU:HD12 | 2.44 | 0.42 |
| 24:D2:47:ASN:N | 24:D2:47:ASN:HD22 | 2.18 | 0.42 |
| 26:D4:1:MET:H2 | 36:DG:67:LYS:HZ1 | 1.68 | 0.42 |
| 28:D6:44:ARG:O | 28:D6:45:LYS:CG | 2.66 | 0.42 |
| 30:D8:31:HIS:CB | 31:DA:2420:C:H41 | 2.31 | 0.42 |
| 31:DA:197:A:C2' | 31:DA:198:C:O5' | 2.68 | 0.42 |
| 31:DA:271(K):U:H3' | 31:DA:271(L):U:C5' | 2.50 | 0.42 |
| 31:DA:638:G:C5 | 31:DA:639:U:C5 | 3.07 | 0.42 |
| 31:DA:762:U:H4' | 31:DA:763:G:H5" | 2.01 | 0.42 |
| 31:DA:762:U:H4' | 31:DA:763:G:C5' | 2.50 | 0.42 |
| 31:DA:1224:C:O3' | 47:DV:88:ARG:HB3 | 2.18 | 0.42 |
| 31:DA:1275:A:N1 | 31:DA:1295:C:O2' | 2.51 | 0.42 |
| 31:DA:1340:U:H4' | 31:DA:1341:U:OP2 | 2.18 | 0.42 |
| 31:DA:1419:A:C8 | 31:DA:1421:G:C6 | 3.08 | 0.42 |
| 31:DA:1472:A:O2' | 31:DA:1473:G:H5' | 2.19 | 0.42 |
| 31:DA:2392:A:N1 | 31:DA:2424:C:N3 | 2.67 | 0.42 |
| 31:DA:2598:A:O5' | 33:DD:236:GLY:HA3 | 2.19 | 0.42 |
| 31:DA:2655:G:C2 | 31:DA:2664:G:O6 | 2.72 | 0.42 |
| 31:DA:2747:G:C2 | 31:DA:2756:U:C5 | 3.07 | 0.42 |
| 31:DA:2753:A:O2' | 31:DA:2754:U:C6 | 2.71 | 0.42 |
| 31:DA:2774:C:H2' | 31:DA:2775:A:O4' | 2.19 | 0.42 |
| 32:DB:15:A:C5' | 32:DB:16:G:H8 | 2.19 | 0.42 |
| 32:DB:17:C:H2' | 32:DB:18:G:H5' | 2.02 | 0.42 |
| 32:DB:45:A:H1' | 36:DG:95:ARG:NH2 | 2.34 | 0.42 |
| 33:DD:11:PRO:O | 33:DD:13:ARG:N | 2.52 | 0.42 |
| 33:DD:35:LYS:N | 33:DD:64:ILE:HG23 | 2.34 | 0.42 |
| 34:DE:10:GLY:HA2 | 34:DE:192:ASN:HD21 | 1.83 | 0.42 |
| 35:DF:165:ARG:HA | 35:DF:168:ARG:HD3 | 2.00 | 0.42 |
| 35:DF:170:LEU:HD23 | 35:DF:172:TRP:HE1 | 1.81 | 0.42 |
| 36:DG:48:GLU:O | 36:DG:49:ASP:CB | 2.68 | 0.42 |
| 36:DG:106:LEU:O | 36:DG:110:ALA:HB3 | 2.20 | 0.42 |
| 37:DH:40:GLU:O | 37:DH:41:MET:HE3 | 2.20 | 0.42 |
| 38:DI:31:LEU:N | 38:DI:31:LEU:CD1 | 2.82 | 0.42 |
| 38:DI:53:ALA:HB1 | 38:DI:56:LYS:HZ2 | 1.84 | 0.42 |
| 39:DN:82:LEU:O | 39:DN:83:LYS:C | 2.62 | 0.42 |
| 40:DO:10:VAL:HG13 | 40:DO:17:ARG:C | 2.43 | 0.42 |
| 40:DO:106:LEU:HD23 | 40:DO:106:LEU:HA | 1.42 | 0.42 |
| 42:DQ:21:THR:O | 42:DQ:21:THR:HG22 | 2.18 | 0.42 |
| 43:DR:104:ARG:HD3 | 43:DR:109:ALA:HB3 | 2.02 | 0.42 |
| 44:DS:67:ARG:N | 44:DS:69:VAL:HG12 | 2.31 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 45:DT:30:VAL:O | 45:DT:30:VAL:CG2 | 2.62 | 0.42 |
| 46:DU:15:LYS:HG3 | 46:DU:16:LYS:N | 2.34 | 0.42 |
| 49:DX:80:ILE:HG23 | 49:DX:81:VAL:N | 2.34 | 0.42 |
| 50:DY:16:ALA:HA | 50:DY:21:LYS:CD | 2.50 | 0.42 |
| 1:AA:37:U:O2' | 1:AA:500:G:H4' | 2.19 | 0.42 |
| 1:AA:230:G:H2' | 1:AA:231:G:O5' | 2.20 | 0.42 |
| 1:AA:324:G:N2 | 1:AA:327:A:C8 | 2.87 | 0.42 |
| 1:AA:445:G:C4 | 1:AA:446:G:C8 | 3.07 | 0.42 |
| 1:AA:500:G:N2 | 1:AA:546:G:H1' | 2.33 | 0.42 |
| 1:AA:519:C:H2' | 1:AA:520:A:O5' | 2.20 | 0.42 |
| 1:AA:622:A:N7 | 1:AA:623:C:C5 | 2.87 | 0.42 |
| 1:AA:666:G:C2 | 1:AA:741:G:C4 | 3.08 | 0.42 |
| 1:AA:690:G:O2' | 1:AA:691:G:H5' | 2.19 | 0.42 |
| 1:AA:1134:G:N2 | 1:AA:1141:C:C2 | 2.87 | 0.42 |
| 1:AA:1157:A:C2 | 1:AA:1181:G:N3 | 2.87 | 0.42 |
| 1:AA:1158:C:O2 | 1:AA:1158:C:C2' | 2.67 | 0.42 |
| 4:AD:194:LEU:N | 4:AD:194:LEU:HD22 | 2.35 | 0.42 |
| 6:AF:15:ASP:C | 6:AF:17:SER:N | 2.77 | 0.42 |
| 7:AG:16:LEU:HD13 | 9:AI:45:ALA:HB2 | 2.01 | 0.42 |
| 9:AI:99:LEU:O | 9:AI:100:GLY:C | 2.62 | 0.42 |
| 11:AK:84:VAL:C | 11:AK:85:ARG:HG3 | 2.45 | 0.42 |
| 11:AK:127:LYS:HA | 11:AK:127:LYS:CE | 2.49 | 0.42 |
| 12:AL:117:ARG:O | 12:AL:118:SER:C | 2.63 | 0.42 |
| 18:AR:51:LEU:HD23 | 18:AR:51:LEU:HA | 1.89 | 0.42 |
| 23:B1:63:ALA:O | 23:B1:65:SER:HB3 | 2.19 | 0.42 |
| 23:B1:73:LEU:HD23 | 23:B1:73:LEU:HA | 1.64 | 0.42 |
| 28:B6:28:ARG:HA | 28:B6:32:ASN:ND2 | 2.15 | 0.42 |
| 29:B7:12:ARG:HG3 | 31:BA:686:G:O6 | 2.19 | 0.42 |
| 31:BA:271(P):C:H2' | 31:BA:271(Q):G:C5' | 2.49 | 0.42 |
| 31:BA:573:G:H1 | 31:BA:2030:A:H3' | 1.81 | 0.42 |
| 31:BA:879:G:O6 | 31:BA:899:A:C2 | 2.72 | 0.42 |
| 31:BA:1262:A:C4 | 31:BA:1263:U:C5 | 3.07 | 0.42 |
| 31:BA:1275:A:N1 | 31:BA:1295:C:O2' | 2.51 | 0.42 |
| 31:BA:1407:C:O2 | 31:BA:1407:C:H2' | 2.19 | 0.42 |
| 31:BA:1412:A:H2' | 31:BA:1413:G:H8 | 1.83 | 0.42 |
| 31:BA:1469:A:C2' | 31:BA:1470:G:O5' | 2.67 | 0.42 |
| 31:BA:1670:C:OP2 | 31:BA:2550:G:OP1 | 2.38 | 0.42 |
| 31:BA:1853:A:H2' | 31:BA:1854:A:C8 | 2.54 | 0.42 |
| 31:BA:1856:G:C2 | 31:BA:1887:C:N3 | 2.87 | 0.42 |
| 31:BA:1902:C:C2' | 31:BA:1903:G:O5' | 2.67 | 0.42 |
| 31:BA:1905:C:O2 | 31:BA:1905:C:H2' | 2.20 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2610:C:C4' | 31:BA:2611:U:OP2 | 2.65 | 0.42 |
| 31:BA:2655:G:N3 | 31:BA:2655:G:C2' | 2.80 | 0.42 |
| 33:BD:133:LEU:HA | 33:BD:136:ILE:HD12 | 2.01 | 0.42 |
| 35:BF:18:ARG:C | 35:BF:19:GLU:HG2 | 2.44 | 0.42 |
| 35:BF:117:ARG:NH2 | 41:BP:5:ASP:N | 2.67 | 0.42 |
| 35:BF:157:VAL:HB | 35:BF:194:MET:HB3 | 2.01 | 0.42 |
| 36:BG:48:GLU:O | 36:BG:49:ASP:CB | 2.67 | 0.42 |
| 38:BI:10:GLU:CD | 38:BI:11:ASN:N | 2.77 | 0.42 |
| 38:BI:79:ILE:HA | 38:BI:79:ILE:HD13 | 1.77 | 0.42 |
| 39:BN:19:GLU:CG | 39:BN:20:GLY:N | 2.82 | 0.42 |
| 41:BP:133:SER:O | 41:BP:137:LYS:HG2 | 2.19 | 0.42 |
| 42:BQ:63:LYS:HG2 | 42:BQ:65:PHE:CE2 | 2.54 | 0.42 |
| 43:BR:116:LEU:HD23 | 43:BR:116:LEU:HA | 1.56 | 0.42 |
| 45:BT:28:VAL:HG22 | 45:BT:46:GLU:C | 2.44 | 0.42 |
| 45:BT:65:LYS:CG | 45:BT:66:VAL:N | 2.82 | 0.42 |
| 45:BT:129:ARG:HH11 | 45:BT:131:ALA:HB3 | 1.82 | 0.42 |
| 46:BU:25:TRP:O | 46:BU:26:GLY:O | 2.38 | 0.42 |
| 47:BV:82:ARG:CG | 47:BV:82:ARG:NH1 | 2.42 | 0.42 |
| 49:BX:12:VAL:CG1 | 49:BX:27:THR:HG23 | 2.50 | 0.42 |
| 1:CA:69:G:H2' | 1:CA:70:G:H8 | 1.84 | 0.42 |
| 1:CA:438:G:H4' | 4:CD:123:HIS:CE1 | 2.54 | 0.42 |
| 1:CA:1006:C:H42 | 1:CA:1024:G:H21 | 1.66 | 0.42 |
| 1:CA:1030(B):C:N4 | 1:CA:1030(C):G:C2 | 2.88 | 0.42 |
| 1:CA:1158:C:O2 | 1:CA:1158:C:H2' | 2.18 | 0.42 |
| 1:CA:1188:A:C2' | 1:CA:1189:C:H5' | 2.50 | 0.42 |
| 1:CA:1498:U:C1' | 1:CA:1499:A:OP2 | 2.68 | 0.42 |
| 4:CD:14:ARG:HA | 4:CD:39:PRO:CB | 2.49 | 0.42 |
| 4:CD:52:SER:O | 4:CD:53:ASP:C | 2.62 | 0.42 |
| 4:CD:106:TYR:CE1 | 4:CD:112:VAL:O | 2.73 | 0.42 |
| 5:CE:48:ALA:HB1 | 5:CE:49:PRO:CD | 2.50 | 0.42 |
| 6:CF:16:GLN:CD | 6:CF:16:GLN:H | 2.28 | 0.42 |
| 7:CG:75:VAL:CG2 | 7:CG:144:MET:HB3 | 2.50 | 0.42 |
| 11:CK:38:ASN:O | 11:CK:39:PRO:C | 2.62 | 0.42 |
| 11:CK:57:THR:HG23 | 11:CK:58:PRO:HD2 | 2.01 | 0.42 |
| 11:CK:126:ARG:O | 11:CK:126:ARG:HG2 | 2.18 | 0.42 |
| 19:CS:9:VAL:O | 19:CS:10:PHE:CD1 | 2.72 | 0.42 |
| 22:D0:72:ARG:HH21 | 22:D0:75:LEU:HD12 | 1.83 | 0.42 |
| 24:D2:26:ARG:CG | 49:DX:5:TYR:O | 2.65 | 0.42 |
| 26:D4:29:PRO:C | 26:D4:31:ILE:N | 2.76 | 0.42 |
| 27:D5:7:PRO:HA | 31:DA:2615:U:N1 | 2.34 | 0.42 |
| 27:D5:33:CYS:HA | 27:D5:34:PRO:HD2 | 1.90 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 28:D6:26:ASN:ND2 | 28:D6:32:ASN:HD21 | 2.17 | 0.42 |
| 30:D8:62:LEU:N | 30:D8:63:PRO:CD | 2.81 | 0.42 |
| 31:DA:28:A:C4 | 31:DA:29:U:C6 | 3.07 | 0.42 |
| 31:DA:142:A:H3' | 31:DA:142(A):C:H5' | 2.01 | 0.42 |
| 31:DA:157:U:OP2 | 31:DA:157:U:H6 | 2.01 | 0.42 |
| 31:DA:184:C:C2 | 31:DA:185:U:C5 | 3.08 | 0.42 |
| 31:DA:639:U:C2 | 31:DA:640:C:C5 | 3.07 | 0.42 |
| 31:DA:815:C:C2 | 31:DA:1193:G:C2 | 3.07 | 0.42 |
| 31:DA:940:G:H2' | 31:DA:941:A:O4' | 2.20 | 0.42 |
| 31:DA:1011:G:C4 | 31:DA:1013:C:C6 | 3.07 | 0.42 |
| 31:DA:1049:C:O2 | 31:DA:1050:A:N7 | 2.52 | 0.42 |
| 31:DA:1052:C:C6 | 31:DA:1052:C:C3' | 3.02 | 0.42 |
| 31:DA:1155:A:OP2 | 46:DU:58:ARG:NH1 | 2.52 | 0.42 |
| 31:DA:1344:G:H4' | 31:DA:1384:A:C5 | 2.55 | 0.42 |
| 31:DA:1533:G:O2' | 31:DA:1543:C:OP2 | 2.33 | 0.42 |
| 31:DA:1607:C:N4 | 31:DA:1622:G:OP2 | 2.39 | 0.42 |
| 31:DA:2228:G:C5 | 31:DA:2229:C:C5 | 3.07 | 0.42 |
| 31:DA:2324:C:H5" | 31:DA:2325:G:H5' | 2.01 | 0.42 |
| 31:DA:2392:A:H1' | 41:DP:60:MET:HG2 | 2.01 | 0.42 |
| 31:DA:2516:G:C6 | 31:DA:2517:C:C4 | 3.07 | 0.42 |
| 31:DA:2636:U:H4' | 34:DE:80:GLU:CD | 2.44 | 0.42 |
| 31:DA:2688:U:O2 | 31:DA:2688:U:O5' | 2.37 | 0.42 |
| 31:DA:2796:U:O2 | 31:DA:2796:U:O4' | 2.38 | 0.42 |
| 33:DD:52:ARG:CZ | 33:DD:53:PHE:CE2 | 3.02 | 0.42 |
| 35:DF:19:GLU:O | 35:DF:20:LEU:CB | 2.67 | 0.42 |
| 35:DF:46:ARG:HG2 | 35:DF:46:ARG:NH1 | 2.15 | 0.42 |
| 37:DH:37:VAL:HG12 | 37:DH:38:SER:N | 2.34 | 0.42 |
| 43:DR:44:LEU:O | 43:DR:45:ARG:C | 2.63 | 0.42 |
| 44:DS:28:VAL:O | 44:DS:29:PHE:CB | 2.67 | 0.42 |
| 45:DT:31:SER:OG | 45:DT:32:TYR:N | 2.53 | 0.42 |
| 48:DW:43:GLY:O | 48:DW:44:ALA:C | 2.62 | 0.42 |
| 49:DX:53:LYS:N | 49:DX:80:ILE:HG22 | 2.34 | 0.42 |
| 51:DZ:56:VAL:C | 51:DZ:57:ILE:HG12 | 2.45 | 0.42 |
| 1:AA:233:C:C2 | 1:AA:234:C:C6 | 3.08 | 0.42 |
| 1:AA:251:G:H4' | 1:AA:252:U:O5' | 2.19 | 0.42 |
| 1:AA:343:U:O2 | 1:AA:347:G:C2 | 2.72 | 0.42 |
| 1:AA:408:A:N1 | 1:AA:409:G:C4 | 2.88 | 0.42 |
| 1:AA:678:U:H2' | 1:AA:679:C:C6 | 2.54 | 0.42 |
| 1:AA:832:C:HO2' | 1:AA:833:U:P | 2.42 | 0.42 |
| 1:AA:945:G:C6 | 1:AA:1337:G:C6 | 3.08 | 0.42 |
| 1:AA:1316:G:O3' | 14:AN:18:VAL:HG22 | 2.20 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 2:AB:25:ASN:OD1 | 2:AB:25:ASN:C | 2.62 | 0.42 |
| 2:AB:139:LYS:O | 2:AB:143:GLU:HG2 | 2.19 | 0.42 |
| 2:AB:171:ALA:HA | 2:AB:174:VAL:HG23 | 2.02 | 0.42 |
| 3:AC:87:LEU:O | 3:AC:91:LEU:HG | 2.19 | 0.42 |
| 4:AD:131:ARG:HD3 | 4:AD:131:ARG:N | 2.34 | 0.42 |
| 5:AE:48:ALA:HB1 | 5:AE:49:PRO:HD2 | 2.00 | 0.42 |
| 9:AI:63:ILE:HD12 | 9:AI:63:ILE:N | 2.35 | 0.42 |
| 10:AJ:78:ASN:HB3 | 10:AJ:80:LYS:H | 1.84 | 0.42 |
| 12:AL:34:ARG:CG | 12:AL:35:GLY:N | 2.83 | 0.42 |
| 12:AL:69:TYR:HB3 | 12:AL:99:HIS:CD2 | 2.55 | 0.42 |
| 13:AM:40:ASN:HA | 13:AM:41:PRO:HD3 | 1.80 | 0.42 |
| 17:AQ:78:GLU:O | 17:AQ:78:GLU:HG3 | 2.19 | 0.42 |
| 19:AS:63:THR:O | 19:AS:66:MET:HG2 | 2.20 | 0.42 |
| 22:B0:72:ARG:HH21 | 22:B0:75:LEU:CD1 | 2.33 | 0.42 |
| 22:B0:77:ARG:NH2 | 31:BA:857:C:OP2 | 2.51 | 0.42 |
| 24:B2:26:ARG:HG2 | 49:BX:5:TYR:CB | 2.48 | 0.42 |
| 24:B2:41:ILE:HG12 | 31:BA:94(A):G:N2 | 2.34 | 0.42 |
| 24:B2:46:GLN:NE2 | 24:B2:47:ASN:CA | 2.82 | 0.42 |
| 26:B4:19:GLY:C | 26:B4:21:VAL:N | 2.75 | 0.42 |
| 31:BA:49:A:H3' | 31:BA:50:U:C5' | 2.49 | 0.42 |
| 31:BA:271(H):G:C6 | 31:BA:271(Q):G:N1 | 2.87 | 0.42 |
| 31:BA:292:C:H42 | 31:BA:348:G:H1 | 1.67 | 0.42 |
| 31:BA:707:G:C5 | 31:BA:708:C:C5 | 3.07 | 0.42 |
| 31:BA:921:G:C6 | 31:BA:922:U:C4 | 3.08 | 0.42 |
| 31:BA:966:G:C5 | 31:BA:967:C:C5 | 3.08 | 0.42 |
| 31:BA:993:G:H1' | 47:BV:91:TYR:HD1 | 1.84 | 0.42 |
| 31:BA:997:G:O2' | 31:BA:998:C:H5' | 2.20 | 0.42 |
| 31:BA:1288:U:C2 | 31:BA:1327:C:C2 | 3.07 | 0.42 |
| 31:BA:1411:C:O2' | 31:BA:1412:A:H5' | 2.19 | 0.42 |
| 31:BA:1679:U:C2' | 31:BA:1680:U:H5' | 2.49 | 0.42 |
| 31:BA:2052:G:C2 | 31:BA:2053:G:C8 | 3.07 | 0.42 |
| 31:BA:2313:C:H2' | 31:BA:2314:C:H6 | 1.85 | 0.42 |
| 31:BA:2660:A:N3 | 31:BA:2660:A:C3' | 2.78 | 0.42 |
| 31:BA:2766:G:H2' | 31:BA:2766:G:N3 | 2.34 | 0.42 |
| 31:BA:2818:G:N2 | 31:BA:2819:G:C4 | 2.88 | 0.42 |
| 32:BB:119:G:N3 | 32:BB:119:G:H2' | 2.33 | 0.42 |
| 33:BD:30:GLU:CG | 33:BD:63:ARG:CZ | 2.97 | 0.42 |
| 33:BD:62:TYR:CD1 | 33:BD:62:TYR:C | 2.97 | 0.42 |
| 33:BD:270:ILE:O | 33:BD:270:ILE:HD12 | 2.19 | 0.42 |
| 35:BF:19:GLU:O | 35:BF:20:LEU:CB | 2.66 | 0.42 |
| 35:BF:32:LEU:CD1 | 35:BF:105:VAL:HG13 | 2.44 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 36:BG:19:LEU:HG | 36:BG:175:LEU:HD12 | 2.02 | 0.42 |
| 37:BH:87:LEU:HD23 | 37:BH:164:TYR:HD1 | 1.85 | 0.42 |
| 38:BI:92:VAL:O | 38:BI:92:VAL:HG13 | 2.20 | 0.42 |
| 41:BP:102:ARG:O | 41:BP:103:ALA:CB | 2.67 | 0.42 |
| 42:BQ:7:MET:O | 42:BQ:10:ARG:NH2 | 2.50 | 0.42 |
| 46:BU:92:ARG:HB2 | 47:BV:11:GLN:CD | 2.42 | 0.42 |
| 50:BY:9:LYS:HA | 50:BY:30:VAL:HG21 | 2.01 | 0.42 |
| 51:BZ:44:PHE:CZ | 51:BZ:48:PHE:CD2 | 3.07 | 0.42 |
| 51:BZ:117:LEU:HG | 51:BZ:118:GLN:N | 2.34 | 0.42 |
| 1:CA:35:G:O2' | 12:CL:121:GLY:HA2 | 2.20 | 0.42 |
| 1:CA:44:G:H2' | 1:CA:45:U:O4' | 2.19 | 0.42 |
| 1:CA:328:C:H4' | 1:CA:329:A:O5' | 2.19 | 0.42 |
| 1:CA:380:G:C2 | 1:CA:384:G:C6 | 3.08 | 0.42 |
| 1:CA:1003:G:C2' | 1:CA:1004:A:H4' | 2.42 | 0.42 |
| 1:CA:1128:C:H1' | 1:CA:1146:A:H61 | 1.85 | 0.42 |
| 1:CA:1160:G:H5' | 2:CB:132:LYS:HE3 | 2.01 | 0.42 |
| 2:CB:80:ILE:HD12 | 2:CB:211:ILE:HB | 2.00 | 0.42 |
| 2:CB:193:ASP:O | 2:CB:194:PRO:O | 2.37 | 0.42 |
| 2:CB:204:ASN:HD22 | 2:CB:206:ASP:H | 1.66 | 0.42 |
| 3:CC:134:ILE:HG13 | 3:CC:153:VAL:CG2 | 2.49 | 0.42 |
| 3:CC:178:LEU:C | 3:CC:180:ALA:H | 2.28 | 0.42 |
| 4:CD:65:ARG:HG3 | 4:CD:75:PHE:CD1 | 2.55 | 0.42 |
| 4:CD:74:GLN:O | 4:CD:78:LEU:HG | 2.20 | 0.42 |
| 4:CD:148:VAL:CG1 | 4:CD:149:ALA:N | 2.81 | 0.42 |
| 4:CD:190:ASP:HB3 | 4:CD:193:ASP:OD2 | 2.20 | 0.42 |
| 15:CO:18:PHE:O | 15:CO:19:PRO:C | 2.63 | 0.42 |
| 16:CP:59:TRP:O | 16:CP:64:ALA:CB | 2.67 | 0.42 |
| 20:CT:28:ALA:O | 20:CT:29:LYS:C | 2.61 | 0.42 |
| 22:D0:70:GLN:OE1 | 22:D0:72:ARG:CD | 2.67 | 0.42 |
| 27:D5:2:ALA:HB3 | 31:DA:747:U:C6 | 2.55 | 0.42 |
| 28:D6:25:LYS:HB2 | 28:D6:25:LYS:HE3 | 1.90 | 0.42 |
| 31:DA:17:G:C4 | 31:DA:18:C:C5 | 3.08 | 0.42 |
| 31:DA:142:A:O2' | 31:DA:1407:C:H2' | 2.20 | 0.42 |
| 31:DA:491:G:H2' | 31:DA:492:A:C8 | 2.55 | 0.42 |
| 31:DA:633:A:H2' | 31:DA:634:C:C5' | 2.49 | 0.42 |
| 31:DA:764:A:O4' | 33:DD:213:ARG:HG3 | 2.20 | 0.42 |
| 31:DA:825:C:H3' | 31:DA:825:C:C6 | 2.54 | 0.42 |
| 31:DA:851:U:C2 | 31:DA:927:G:C2 | 3.07 | 0.42 |
| 31:DA:1510:G:C2 | 31:DA:1511:C:C2 | 3.07 | 0.42 |
| 31:DA:2323:G:H2' | 31:DA:2324:C:O4' | 2.18 | 0.42 |
| 34:DE:14:ILE:HD11 | 34:DE:173:VAL:HG11 | 2.02 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 35:DF:93:LYS:HB3 | 35:DF:94:PRO:HD2 | 2.01 | 0.42 |
| 35:DF:117:ARG:NH2 | 41:DP:5:ASP:N | 2.67 | 0.42 |
| 36:DG:33:ARG:HB2 | 36:DG:162:THR:OG1 | 2.18 | 0.42 |
| 37:DH:89:ILE:C | 37:DH:89:ILE:HD12 | 2.44 | 0.42 |
| 38:DI:14:ASP:O | 38:DI:17:GLN:HB3 | 2.19 | 0.42 |
| 38:DI:88:ILE:CG2 | 38:DI:89:TYR:H | 2.21 | 0.42 |
| 40:DO:40:VAL:HG12 | 40:DO:41:ALA:N | 2.35 | 0.42 |
| 42:DQ:18:LYS:O | 42:DQ:19:GLY:O | 2.38 | 0.42 |
| 42:DQ:39:PRO:HA | 42:DQ:97:VAL:O | 2.19 | 0.42 |
| 44:DS:59:LYS:NZ | 44:DS:68:GLN:HE22 | 2.17 | 0.42 |
| 48:DW:12:ILE:HG13 | 48:DW:42:ARG:HH11 | 1.85 | 0.42 |
| 49:DX:57:LEU:HD11 | 49:DX:77:LYS:HD2 | 2.02 | 0.42 |
| 50:DY:96:ILE:HG13 | 50:DY:99:CYS:O | 2.19 | 0.42 |
| 51:DZ:5:LEU:HD21 | 51:DZ:43:GLU:HB3 | 2.01 | 0.42 |
| 1:AA:6:G:H4' | 1:AA:298:A:H4' | 2.00 | 0.42 |
| 1:AA:193:C:O2' | 1:AA:194:C:H5' | 2.20 | 0.42 |
| 1:AA:443:C:O2 | 1:AA:443:C:H2' | 2.19 | 0.42 |
| 1:AA:565:U:OP2 | 1:AA:566:G:O2' | 2.30 | 0.42 |
| 1:AA:812:C:OP1 | 1:AA:903:G:H1' | 2.20 | 0.42 |
| 1:AA:827:U:C2 | 1:AA:870:U:C4 | 3.07 | 0.42 |
| 1:AA:953:G:C6 | 1:AA:1229:A:C6 | 3.08 | 0.42 |
| 1:AA:1218:C:H2' | 1:AA:1219:U:C5 | 2.54 | 0.42 |
| 1:AA:1221:G:H4' | 19:AS:53:ASN:O | 2.18 | 0.42 |
| 1:AA:1423:G:H5" | 40:BO:49:ARG:HH21 | 1.83 | 0.42 |
| 2:AB:79:ASP:C | 2:AB:81:VAL:N | 2.78 | 0.42 |
| 2:AB:165:VAL:HG23 | 2:AB:166:ASP:N | 2.30 | 0.42 |
| 3:AC:29:TYR:O | 3:AC:29:TYR:CD2 | 2.72 | 0.42 |
| 3:AC:155:GLY:O | 3:AC:156:ARG:CB | 2.68 | 0.42 |
| 4:AD:106:TYR:CE1 | 4:AD:113:SER:C | 2.98 | 0.42 |
| 5:AE:91:LEU:HD12 | 5:AE:120:THR:HG22 | 2.02 | 0.42 |
| 5:AE:139:LEU:C | 5:AE:141:GLN:N | 2.74 | 0.42 |
| 6:AF:22:GLU:OE1 | 6:AF:84:ASN:HB2 | 2.19 | 0.42 |
| 7:AG:111:ARG:HB2 | 7:AG:119:ARG:HD2 | 2.00 | 0.42 |
| 11:AK:84:VAL:HG11 | 11:AK:95:ILE:HD12 | 2.01 | 0.42 |
| 11:AK:106:LYS:O | 11:AK:107:SER:HB3 | 2.20 | 0.42 |
| 12:AL:115:LYS:O | 12:AL:117:ARG:HG3 | 2.19 | 0.42 |
| 23:B1:9:GLY:O | 23:B1:10:LYS:HB3 | 2.20 | 0.42 |
| 23:B1:26:ARG:CB | 23:B1:34:THR:CA | 2.95 | 0.42 |
| 23:B1:46:LEU:HA | 31:BA:396:G:O3' | 2.19 | 0.42 |
| 23:B1:75:GLU:O | 23:B1:76:ARG:HD3 | 2.19 | 0.42 |
| 24:B2:32:LEU:HA | 24:B2:37:PHE:HB3 | 2.01 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 24:B2:57:ILE:HG13 | 24:B2:58:ALA:O | 2.19 | 0.42 |
| 30:B8:51:ALA:HA | 30:B8:54:GLU:HG2 | 2.02 | 0.42 |
| 31:BA:49:A:C3' | 31:BA:50:U:C5' | 2.97 | 0.42 |
| 31:BA:271(F):C:C4 | 31:BA:271(G):C:C5 | 3.08 | 0.42 |
| 31:BA:1004:C:C6 | 31:BA:1004:C:C3' | 3.03 | 0.42 |
| 31:BA:1208:C:C2' | 31:BA:1209:G:H5' | 2.49 | 0.42 |
| 31:BA:1368:G:C2 | 31:BA:1369:G:C8 | 3.08 | 0.42 |
| 31:BA:1600:C:O2' | 31:BA:1601:G:H5' | 2.19 | 0.42 |
| 31:BA:1797:C:H4' | 33:BD:257:LEU:O | 2.19 | 0.42 |
| 31:BA:1803:A:H4' | 33:BD:259:THR:HG23 | 2.00 | 0.42 |
| 31:BA:1891:G:C5 | 31:BA:1892:C:C4 | 3.08 | 0.42 |
| 31:BA:2418:A:H2' | 31:BA:2419:U:C6 | 2.54 | 0.42 |
| 31:BA:2782:G:O5' | 31:BA:2782:G:H8 | 2.03 | 0.42 |
| 32:BB:21:G:N3 | 32:BB:21:G:H2' | 2.34 | 0.42 |
| 32:BB:115:G:H2' | 32:BB:116:G:H5" | 2.01 | 0.42 |
| 33:BD:132:PRO:HG3 | 33:BD:190:TYR:CE1 | 2.55 | 0.42 |
| 34:BE:77:ILE:HG12 | 34:BE:78:LEU:H | 1.85 | 0.42 |
| 34:BE:93:VAL:C | 34:BE:95:ILE:N | 2.78 | 0.42 |
| 35:BF:8:GLN:HA | 35:BF:8:GLN:OE1 | 2.19 | 0.42 |
| 35:BF:84:VAL:C | 35:BF:86:GLY:H | 2.26 | 0.42 |
| 35:BF:122:LYS:HA | 35:BF:122:LYS:HD3 | 1.78 | 0.42 |
| 38:BI:129:THR:HG22 | 38:BI:130:TYR:N | 2.33 | 0.42 |
| 43:BR:55:ALA:HB2 | 43:BR:79:LEU:HD13 | 2.00 | 0.42 |
| 44:BS:87:PHE:CG | 44:BS:88:ASP:N | 2.86 | 0.42 |
| 45:BT:56:GLY:C | 45:BT:59:THR:HG22 | 2.45 | 0.42 |
| 45:BT:106:SER:O | 45:BT:107:ASP:OD1 | 2.37 | 0.42 |
| 46:BU:52:ARG:O | 46:BU:53:ARG:C | 2.62 | 0.42 |
| 47:BV:18:LEU:O | 47:BV:19:LYS:HB2 | 2.20 | 0.42 |
| 50:BY:52:SER:C | 50:BY:54:LYS:N | 2.74 | 0.42 |
| 51:BZ:44:PHE:CE1 | 51:BZ:48:PHE:CD2 | 3.08 | 0.42 |
| 1:CA:124:G:C6 | 1:CA:125:U:C4 | 3.08 | 0.42 |
| 1:CA:167:G:C2' | 1:CA:168:G:H5' | 2.50 | 0.42 |
| 1:CA:273:A:N6 | 1:CA:274:A:N6 | 2.67 | 0.42 |
| 1:CA:295:C:H2' | 1:CA:296:U:H6 | 1.85 | 0.42 |
| 1:CA:329:A:C2 | 1:CA:332:G:C4 | 3.08 | 0.42 |
| 1:CA:812:C:OP1 | 1:CA:903:G:H1' | 2.19 | 0.42 |
| 1:CA:1320:C:H5' | 19:CS:70:LYS:CG | 2.49 | 0.42 |
| 1:CA:1376:U:H2' | 1:CA:1377:A:C8 | 2.54 | 0.42 |
| 1:CA:1415:G:C6 | 1:CA:1486:G:C6 | 3.08 | 0.42 |
| 3:CC:182:ILE:HG23 | 3:CC:202:ILE:C | 2.45 | 0.42 |
| 4:CD:62:GLN:HE22 | 4:CD:65:ARG:HH21 | 1.68 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 5:CE:36:ASP:OD1 | 5:CE:37:ARG:N | 2.52 | 0.42 |
| 8:CH:36:LEU:HA | 8:CH:39:LEU:HD23 | 2.00 | 0.42 |
| 11:CK:15:ALA:HA | 11:CK:77:MET:HA | 2.01 | 0.42 |
| 12:CL:55:VAL:CG1 | 12:CL:56:ALA:N | 2.82 | 0.42 |
| 15:CO:24:SER:O | 15:CO:28:GLN:HG3 | 2.19 | 0.42 |
| 16:CP:66:PRO:O | 16:CP:67:THR:C | 2.61 | 0.42 |
| 17:CQ:31:LEU:HG | 17:CQ:32:TYR:CD2 | 2.55 | 0.42 |
| 24:D2:12:GLU:C | 24:D2:14:ARG:H | 2.27 | 0.42 |
| 24:D2:51:ARG:O | 24:D2:52:ASP:HB2 | 2.17 | 0.42 |
| 29:D7:5:TRP:CH2 | 31:DA:686:G:N7 | 2.88 | 0.42 |
| 31:DA:64:A:C2' | 31:DA:65:C:H5' | 2.49 | 0.42 |
| 31:DA:332:A:C2 | 31:DA:335:C:C5 | 3.07 | 0.42 |
| 31:DA:354:G:H8 | 31:DA:354:G:O5' | 2.02 | 0.42 |
| 31:DA:536:A:H2' | 31:DA:537:C:H6 | 1.85 | 0.42 |
| 31:DA:565:C:H2' | 31:DA:566:U:O4' | 2.19 | 0.42 |
| 31:DA:648:G:C2' | 31:DA:649:G:H5' | 2.49 | 0.42 |
| 31:DA:775:G:C5 | 31:DA:794:G:C8 | 3.07 | 0.42 |
| 31:DA:1251:C:O5' | 46:DU:10:ARG:HG3 | 2.20 | 0.42 |
| 31:DA:1363:C:C2' | 31:DA:1364:G:O5' | 2.68 | 0.42 |
| 31:DA:1598:C:O2 | 31:DA:1598:C:C2' | 2.67 | 0.42 |
| 31:DA:2037:G:C6 | 31:DA:2038:G:C6 | 3.08 | 0.42 |
| 31:DA:2251:G:C6 | 31:DA:2252:G:C5 | 3.08 | 0.42 |
| 31:DA:2314:C:N3 | 31:DA:2315:G:N7 | 2.67 | 0.42 |
| 33:DD:58:HIS:CD2 | 33:DD:59:LYS:H | 2.31 | 0.42 |
| 35:DF:104:LYS:O | 35:DF:108:LYS:HG2 | 2.19 | 0.42 |
| 36:DG:39:ILE:HD12 | 36:DG:40:ASN:N | 2.34 | 0.42 |
| 37:DH:46:GLU:O | 37:DH:47:GLU:CB | 2.68 | 0.42 |
| 38:DI:79:ILE:HD13 | 38:DI:79:ILE:HA | 1.79 | 0.42 |
| 39:DN:1:MET:HE2 | 47:DV:20:LEU:HD21 | 2.01 | 0.42 |
| 39:DN:17:ASP:CG | 39:DN:56:ASN:HB2 | 2.45 | 0.42 |
| 43:DR:56:LYS:HE3 | 43:DR:94:TYR:OH | 2.20 | 0.42 |
| 43:DR:92:GLY:HA2 | 43:DR:94:TYR:CZ | 2.55 | 0.42 |
| 44:DS:58:LEU:HD12 | 44:DS:58:LEU:HA | 1.86 | 0.42 |
| 47:DV:38:LEU:HG | 47:DV:39:LEU:N | 2.34 | 0.42 |
| 48:DW:71:VAL:HA | 48:DW:107:LEU:HD12 | 2.01 | 0.42 |
| 50:DY:98:VAL:O | 50:DY:99:CYS:HB2 | 2.20 | 0.42 |
| 51:DZ:27:VAL:HA | 51:DZ:37:VAL:HG23 | 2.00 | 0.42 |
| 1:AA:69:G:H2' | 1:AA:70:G:H8 | 1.84 | 0.42 |
| 1:AA:101:A:C6 | 1:AA:102:G:N7 | 2.87 | 0.42 |
| 1:AA:152:A:C8 | 1:AA:153:C:C5 | 3.08 | 0.42 |
| 1:AA:533:A:H4' | 1:AA:534:U:OP1 | 2.19 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:716:A:N3 | 11:AK:118:GLY:HA2 | 2.34 | 0.42 |
| 1:AA:1126:U:P | 1:AA:1126:U:H6 | 2.42 | 0.42 |
| 1:AA:1260:C:H4' | 1:AA:1284:C:H5' | 2.01 | 0.42 |
| 1:AA:1346:A:C8 | 1:AA:1348:U:O2 | 2.72 | 0.42 |
| 1:AA:1429:C:H2' | 1:AA:1430:C:H6 | 1.84 | 0.42 |
| 2:AB:97:TRP:CZ2 | 2:AB:102:LEU:HD13 | 2.51 | 0.42 |
| 4:AD:208:SER:O | 4:AD:209:ARG:C | 2.62 | 0.42 |
| 5:AE:71:LEU:O | 5:AE:72:GLN:CG | 2.66 | 0.42 |
| 7:AG:79:ARG:HA | 7:AG:84:ASN:ND2 | 2.35 | 0.42 |
| 8:AH:28:ALA:HB3 | 8:AH:57:PRO:HB2 | 2.02 | 0.42 |
| 8:AH:33:GLU:O | 8:AH:34:GLU:C | 2.63 | 0.42 |
| 8:AH:58:TYR:C | 8:AH:59:LEU:HD23 | 2.45 | 0.42 |
| 9:AI:106:ALA:O | 9:AI:108:VAL:N | 2.53 | 0.42 |
| 11:AK:23:ALA:HB1 | 11:AK:88:GLY:HA3 | 2.02 | 0.42 |
| 15:AO:67:LEU:HD22 | 15:AO:78:TYR:CE1 | 2.52 | 0.42 |
| 23:B1:64:ALA:O | 23:B1:65:SER:CB | 2.67 | 0.42 |
| 23:B1:87:PRO:CD | 23:B1:88:LYS:N | 2.51 | 0.42 |
| 24:B2:14:ARG:NH2 | 24:B2:57:ILE:CG2 | 2.83 | 0.42 |
| 30:B8:29:LYS:HB3 | 30:B8:29:LYS:HE2 | 1.83 | 0.42 |
| 30:B8:34:TRP:CH2 | 30:B8:37:SER:C | 2.98 | 0.42 |
| 31:BA:271(D):G:H1 | 31:BA:271(T):C:H42 | 1.67 | 0.42 |
| 31:BA:287:C:H2' | 31:BA:288:C:H6 | 1.85 | 0.42 |
| 31:BA:356:G:H2' | 31:BA:357:A:C8 | 2.55 | 0.42 |
| 31:BA:661:C:O2' | 41:BP:16:ARG:O | 2.33 | 0.42 |
| 31:BA:1022:G:C6 | 31:BA:1141:U:C5 | 3.08 | 0.42 |
| 31:BA:1174:A:OP1 | 31:BA:1175:U:H5" | 2.19 | 0.42 |
| 31:BA:2102:U:C4 | 31:BA:2103:C:N4 | 2.88 | 0.42 |
| 31:BA:2444:G:OP2 | 35:BF:68:LYS:HE2 | 2.20 | 0.42 |
| 31:BA:2483:C:O2 | 31:BA:2483:C:C2' | 2.67 | 0.42 |
| 31:BA:2648:C:H2' | 31:BA:2649:U:C6 | 2.55 | 0.42 |
| 31:BA:2689:U:C5' | 31:BA:2690:C:H5' | 2.49 | 0.42 |
| 31:BA:2822:G:OP2 | 34:BE:110:GLY:O | 2.38 | 0.42 |
| 32:BB:28:C:N3 | 32:BB:29:A:N7 | 2.67 | 0.42 |
| 32:BB:38:C:O4' | 44:BS:95:HIS:NE2 | 2.51 | 0.42 |
| 33:BD:257:LEU:HA | 33:BD:257:LEU:HD23 | 1.84 | 0.42 |
| 34:BE:61:ARG:C | 34:BE:63:LEU:N | 2.74 | 0.42 |
| 34:BE:136:ARG:CG | 34:BE:136:ARG:NH1 | 2.77 | 0.42 |
| 36:BG:29:TRP:C | 36:BG:31:VAL:H | 2.28 | 0.42 |
| 36:BG:99:MET:O | 36:BG:100:TRP:C | 2.62 | 0.42 |
| 36:BG:129:GLY:O | 36:BG:130:ASN:CB | 2.67 | 0.42 |
| 38:BI:56:LYS:HZ2 | 38:BI:57:ARG:CA | 2.32 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 38:BI:124:GLY:N | 38:BI:142:VAL:HG23 | 2.35 | 0.42 |
| 40:BO:4:PRO:O | 40:BO:5:GLN:CB | 2.65 | 0.42 |
| 42:BQ:23:GLY:C | 42:BQ:100:GLY:C | 2.88 | 0.42 |
| 42:BQ:80:GLU:OE2 | 42:BQ:80:GLU:HA | 2.19 | 0.42 |
| 44:BS:17:ARG:NH2 | 44:BS:89:ARG:NH2 | 2.68 | 0.42 |
| 44:BS:77:ALA:O | 44:BS:78:LEU:C | 2.62 | 0.42 |
| 45:BT:36:GLU:HB3 | 45:BT:38:ASN:OD1 | 2.19 | 0.42 |
| 47:BV:2:PHE:CE1 | 47:BV:13:ARG:NH1 | 2.88 | 0.42 |
| 47:BV:53:GLU:O | 47:BV:54:GLY:C | 2.62 | 0.42 |
| 1:CA:192:U:C4' | 20:CT:103:GLY:HA2 | 2.50 | 0.42 |
| 1:CA:258:G:H2' | 1:CA:259:G:H8 | 1.84 | 0.42 |
| 1:CA:491:G:C2 | 1:CA:492:G:C4 | 3.08 | 0.42 |
| 1:CA:600:C:O2' | 1:CA:601:C:H5' | 2.18 | 0.42 |
| 1:CA:785:G:C2' | 1:CA:786:G:H5' | 2.50 | 0.42 |
| 1:CA:878:G:C5' | 8:CH:89:PRO:HG2 | 2.45 | 0.42 |
| 1:CA:920:U:C1' | 1:CA:1080:A:C2 | 3.02 | 0.42 |
| 1:CA:1073:U:H2' | 1:CA:1074:G:C8 | 2.45 | 0.42 |
| 1:CA:1134:G:N2 | 1:CA:1141:C:C2 | 2.88 | 0.42 |
| 1:CA:1242:C:H5" | 21:CU:10:ARG:NH1 | 2.34 | 0.42 |
| 1:CA:1497:G:C2' | 1:CA:1498:U:H5' | 2.47 | 0.42 |
| 4:CD:78:LEU:O | 4:CD:79:PHE:C | 2.62 | 0.42 |
| 5:CE:146:ALA:C | 5:CE:148:VAL:N | 2.78 | 0.42 |
| 6:CF:15:ASP:C | 6:CF:17:SER:N | 2.78 | 0.42 |
| 7:CG:85:TYR:CD1 | 7:CG:154:TYR:HE1 | 2.38 | 0.42 |
| 7:CG:111:ARG:HB2 | 7:CG:119:ARG:HD2 | 2.01 | 0.42 |
| 8:CH:121:ASP:OD1 | 8:CH:121:ASP:N | 2.53 | 0.42 |
| 11:CK:120:ARG:HA | 11:CK:121:PRO:HD3 | 1.85 | 0.42 |
| 18:CR:44:LEU:O | 18:CR:45:SER:O | 2.38 | 0.42 |
| 18:CR:58:LEU:HB3 | 18:CR:62:GLU:CB | 2.43 | 0.42 |
| 28:D6:27:LYS:HD2 | 31:DA:2285:C:C5 | 2.55 | 0.42 |
| 29:D7:19:ARG:HH11 | 29:D7:19:ARG:HG2 | 1.85 | 0.42 |
| 31:DA:55:G:N3 | 31:DA:127:A:H2 | 2.16 | 0.42 |
| 31:DA:58:G:OP1 | 49:DX:72:LYS:HA | 2.19 | 0.42 |
| 31:DA:107:C:N3 | 31:DA:108:U:C5 | 2.88 | 0.42 |
| 31:DA:269:U:O2 | 31:DA:269:U:C2' | 2.68 | 0.42 |
| 31:DA:287:C:H2' | 31:DA:288:C:O4' | 2.18 | 0.42 |
| 31:DA:465:G:C6 | 31:DA:466:A:N6 | 2.88 | 0.42 |
| 31:DA:794:G:C6 | 31:DA:795:C:C4 | 3.08 | 0.42 |
| 31:DA:811:U:O2 | 31:DA:1250:G:C3' | 2.65 | 0.42 |
| 31:DA:818:G:H4' | 31:DA:838:C:O3' | 2.19 | 0.42 |
| 31:DA:1381:G:H2' | 31:DA:1382:G:C5' | 2.48 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1439:A:C2 | 31:DA:1553:A:C4 | 3.06 | 0.42 |
| 31:DA:1491:G:O2' | 33:DD:101:GLU:HB2 | 2.20 | 0.42 |
| 31:DA:1549:C:H2' | 31:DA:1550:C:H6 | 1.85 | 0.42 |
| 31:DA:2199:A:H3' | 31:DA:2200:C:C6 | 2.54 | 0.42 |
| 31:DA:2287:A:C2 | 31:DA:2346:A:N1 | 2.87 | 0.42 |
| 31:DA:2580:U:H4' | 34:DE:130:GLY:CA | 2.50 | 0.42 |
| 35:DF:140:LEU:HD13 | 35:DF:140:LEU:HA | 1.71 | 0.42 |
| 38:DI:10:GLU:CD | 38:DI:11:ASN:N | 2.78 | 0.42 |
| 39:DN:75:TYR:N | 39:DN:75:TYR:HD1 | 2.16 | 0.42 |
| 41:DP:23:PRO:HB2 | 41:DP:33:ARG:HG3 | 2.01 | 0.42 |
| 43:DR:2:ARG:CD | 43:DR:2:ARG:N | 2.82 | 0.42 |
| 43:DR:37:THR:OG1 | 43:DR:40:LYS:HB2 | 2.20 | 0.42 |
| 44:DS:42:ASP:C | 44:DS:44:LYS:N | 2.71 | 0.42 |
| 45:DT:56:GLY:C | 45:DT:59:THR:HG22 | 2.45 | 0.42 |
| 46:DU:17:ILE:HD13 | 46:DU:39:LEU:HD12 | 2.02 | 0.42 |
| 47:DV:1:MET:HE3 | 47:DV:44:LYS:CB | 2.28 | 0.42 |
| 47:DV:47:VAL:CG2 | 47:DV:49:THR:HB | 2.50 | 0.42 |
| 47:DV:94:LEU:HD23 | 47:DV:94:LEU:C | 2.44 | 0.42 |
| 48:DW:75:TYR:CZ | 48:DW:104:THR:HG21 | 2.55 | 0.42 |
| 49:DX:65:ARG:C | 49:DX:65:ARG:CD | 2.92 | 0.42 |
| 51:DZ:45:ASP:O | 51:DZ:46:LYS:C | 2.62 | 0.42 |
| 1:AA:374:A:C2 | 1:AA:375:U:C2 | 3.08 | 0.42 |
| 1:AA:505:G:C5 | 1:AA:535:A:C2 | 3.08 | 0.42 |
| 1:AA:716:A:H2' | 1:AA:717:C:O5' | 2.19 | 0.42 |
| 1:AA:1014:A:C2 | 19:AS:34:TRP:CE2 | 3.07 | 0.42 |
| 1:AA:1064:G:OP2 | 1:AA:1386:G:H4' | 2.20 | 0.42 |
| 2:AB:17:PHE:O | 2:AB:18:GLY:O | 2.38 | 0.42 |
| 2:AB:124:SER:OG | 2:AB:125:PRO:HD2 | 2.19 | 0.42 |
| 3:AC:19:GLU:HA | 3:AC:54:ARG:NH2 | 2.34 | 0.42 |
| 5:AE:132:ALA:O | 5:AE:133:TYR:C | 2.61 | 0.42 |
| 8:AH:56:LYS:HA | 8:AH:57:PRO:HD2 | 1.77 | 0.42 |
| 8:AH:121:ASP:N | 8:AH:121:ASP:OD1 | 2.53 | 0.42 |
| 9:AI:45:ALA:O | 9:AI:78:LYS:HE3 | 2.19 | 0.42 |
| 16:AP:4:ILE:CD1 | 16:AP:64:ALA:HB1 | 2.50 | 0.42 |
| 16:AP:59:TRP:O | 16:AP:64:ALA:CB | 2.68 | 0.42 |
| 19:AS:37:ARG:H | 19:AS:37:ARG:HG3 | 1.61 | 0.42 |
| 23:B1:87:PRO:O | 23:B1:88:LYS:C | 2.62 | 0.42 |
| 27:B5:40:LYS:HE2 | 27:B5:40:LYS:HB3 | 1.87 | 0.42 |
| 29:B7:22:MET:HE2 | 29:B7:31:LEU:CD1 | 2.49 | 0.42 |
| 30:B8:29:LYS:O | 30:B8:31:HIS:N | 2.53 | 0.42 |
| 30:B8:36:LYS:HZ3 | 30:B8:36:LYS:HG3 | 1.67 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:228:A:H5' | 31:BA:229:A:OP2 | 2.19 | 0.42 |
| 31:BA:389:G:H22 | 41:BP:72:PRO:HD3 | 1.84 | 0.42 |
| 31:BA:848:G:C2 | 31:BA:933:A:H1' | 2.55 | 0.42 |
| 31:BA:1019:U:N3 | 31:BA:1142(A):A:N6 | 2.56 | 0.42 |
| 31:BA:1131:G:H8 | 31:BA:2025:C:H4' | 1.85 | 0.42 |
| 31:BA:1141:U:O2' | 31:BA:1142:U:OP2 | 2.37 | 0.42 |
| 31:BA:1155:A:OP2 | 46:BU:58:ARG:NH1 | 2.53 | 0.42 |
| 31:BA:1212:G:C2 | 31:BA:1236:G:C4 | 3.08 | 0.42 |
| 31:BA:1441:G:H2' | 31:BA:1442:G:H8 | 1.84 | 0.42 |
| 31:BA:1464:C:O2' | 31:BA:1528:A:H1' | 2.19 | 0.42 |
| 31:BA:1601:G:O2' | 31:BA:1602:U:H5' | 2.20 | 0.42 |
| 31:BA:1973:G:C5 | 31:BA:1974:C:C4 | 3.08 | 0.42 |
| 31:BA:2031:A:OP1 | 31:BA:2031:A:H8 | 2.03 | 0.42 |
| 31:BA:2266:A:H4' | 31:BA:2267:A:C4 | 2.55 | 0.42 |
| 31:BA:2287:A:C2 | 31:BA:2346:A:C2 | 3.08 | 0.42 |
| 31:BA:2347:C:H2' | 31:BA:2348:U:C6 | 2.54 | 0.42 |
| 31:BA:2636:U:C2' | 31:BA:2637:U:H5' | 2.50 | 0.42 |
| 31:BA:2681:C:O2 | 31:BA:2681:C:C2' | 2.61 | 0.42 |
| 31:BA:2785:C:H2' | 31:BA:2786:U:O4' | 2.19 | 0.42 |
| 31:BA:2801(A):A:H4' | 31:BA:2802:G:H2' | 2.01 | 0.42 |
| 33:BD:58:HIS:CD2 | 33:BD:59:LYS:H | 2.27 | 0.42 |
| 33:BD:142:VAL:O | 33:BD:142:VAL:HG13 | 2.19 | 0.42 |
| 36:BG:56:ALA:HA | 36:BG:59:GLU:OE1 | 2.19 | 0.42 |
| 37:BH:46:GLU:O | 37:BH:47:GLU:CB | 2.65 | 0.42 |
| 39:BN:58:ASP:O | 39:BN:59:LYS:HB2 | 2.20 | 0.42 |
| 39:BN:99:LEU:HD23 | 39:BN:99:LEU:HA | 1.44 | 0.42 |
| 41:BP:143:GLY:CA | 41:BP:145:PRO:HD3 | 2.48 | 0.42 |
| 42:BQ:21:THR:O | 42:BQ:21:THR:HG22 | 2.20 | 0.42 |
| 42:BQ:23:GLY:C | 42:BQ:100:GLY:HA3 | 2.39 | 0.42 |
| 43:BR:10:LEU:HD22 | 43:BR:17:ARG:CD | 2.44 | 0.42 |
| 43:BR:59:ASP:OD1 | 43:BR:61:HIS:HB3 | 2.20 | 0.42 |
| 44:BS:54:LEU:HD21 | 44:BS:59:LYS:O | 2.20 | 0.42 |
| 44:BS:98:VAL:HG13 | 44:BS:100:ALA:N | 2.35 | 0.42 |
| 46:BU:27:LEU:C | 46:BU:29:SER:N | 2.77 | 0.42 |
| 46:BU:76:TYR:CZ | 46:BU:80:ILE:HG13 | 2.55 | 0.42 |
| 47:BV:2:PHE:CB | 47:BV:42:GLY:HA3 | 2.50 | 0.42 |
| 47:BV:19:LYS:C | 47:BV:20:LEU:HD12 | 2.45 | 0.42 |
| 47:BV:47:VAL:CG2 | 47:BV:49:THR:HB | 2.50 | 0.42 |
| 49:BX:18:TYR:C | 49:BX:20:GLY:N | 2.78 | 0.42 |
| 50:BY:39:VAL:CG1 | 50:BY:40:GLU:N | 2.62 | 0.42 |
| 50:BY:71:LYS:HZ2 | 50:BY:71:LYS:HG3 | 1.67 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:93:G:C2' | 1:CA:96:U:H5' | 2.50 | 0.42 |
| 1:CA:501:C:H2' | 1:CA:502:G:C8 | 2.55 | 0.42 |
| 1:CA:682:G:C2 | 1:CA:709:G:C6 | 3.08 | 0.42 |
| 1:CA:757:U:H2' | 1:CA:758:G:O4' | 2.20 | 0.42 |
| 1:CA:834:C:H2' | 1:CA:835:U:H6 | 1.84 | 0.42 |
| 1:CA:929:G:O6 | 1:CA:930:C:N4 | 2.53 | 0.42 |
| 1:CA:1151:A:O2' | 1:CA:1152:A:C8 | 2.65 | 0.42 |
| 1:CA:1157:A:C2 | 1:CA:1181:G:N3 | 2.88 | 0.42 |
| 1:CA:1241:G:H2' | 1:CA:1242:C:C6 | 2.55 | 0.42 |
| 2:CB:97:TRP:CZ2 | 2:CB:102:LEU:HD13 | 2.49 | 0.42 |
| 2:CB:158:LEU:HD12 | 2:CB:158:LEU:N | 2.15 | 0.42 |
| 5:CE:11:ILE:HB | 5:CE:31:LEU:HB3 | 2.02 | 0.42 |
| 7:CG:92:SER:CB | 7:CG:94:ARG:HH21 | 2.33 | 0.42 |
| 7:CG:101:LEU:O | 7:CG:105:VAL:HG23 | 2.20 | 0.42 |
| 9:CI:25:LYS:HG2 | 9:CI:60:ASP:OD1 | 2.20 | 0.42 |
| 9:CI:122:ALA:HB1 | 9:CI:123:PRO:HD2 | 2.01 | 0.42 |
| 15:CO:67:LEU:HD22 | 15:CO:78:TYR:CE1 | 2.54 | 0.42 |
| 15:CO:75:PRO:O | 15:CO:78:TYR:HB3 | 2.20 | 0.42 |
| 20:CT:22:ARG:O | 20:CT:23:ARG:C | 2.63 | 0.42 |
| 25:D3:46:ASN:ND2 | 31:DA:851:U:C4' | 2.83 | 0.42 |
| 26:D4:6:HIS:HA | 36:DG:67:LYS:HG3 | 2.02 | 0.42 |
| 28:D6:34:LEU:HD22 | 28:D6:50:ARG:HH12 | 1.85 | 0.42 |
| 31:DA:343:C:C2' | 31:DA:344:G:H5' | 2.50 | 0.42 |
| 31:DA:678:C:H2' | 31:DA:679:C:C6 | 2.55 | 0.42 |
| 31:DA:690:G:O2' | 31:DA:691:C:H5' | 2.19 | 0.42 |
| 31:DA:963:U:H1' | 31:DA:2250:G:O6 | 2.20 | 0.42 |
| 31:DA:966:G:C5 | 31:DA:967:C:C5 | 3.07 | 0.42 |
| 31:DA:997:G:C2' | 31:DA:998:C:H5' | 2.50 | 0.42 |
| 31:DA:1023:U:O4 | 31:DA:1142(A):A:N1 | 2.53 | 0.42 |
| 31:DA:1481:U:H5' | 31:DA:1482:G:P | 2.60 | 0.42 |
| 31:DA:1857:G:N2 | 31:DA:1886:C:N4 | 2.68 | 0.42 |
| 31:DA:2346:A:C2 | 31:DA:2383:G:C2 | 3.08 | 0.42 |
| 31:DA:2404:C:H2' | 31:DA:2405:G:H5' | 2.01 | 0.42 |
| 32:DB:10:C:O2' | 32:DB:11:C:H5' | 2.19 | 0.42 |
| 32:DB:27:C:N4 | 32:DB:28:C:N4 | 2.68 | 0.42 |
| 32:DB:33:G:C2 | 32:DB:50:G:C2 | 3.08 | 0.42 |
| 34:DE:52:LEU:HA | 34:DE:53:PRO:HD3 | 1.76 | 0.42 |
| 34:DE:132:HIS:CG | 34:DE:132:HIS:O | 2.73 | 0.42 |
| 34:DE:134:ILE:O | 34:DE:134:ILE:CG1 | 2.66 | 0.42 |
| 34:DE:176:ILE:O | 34:DE:176:ILE:HG22 | 2.20 | 0.42 |
| 36:DG:91:ARG:HD2 | 36:DG:91:ARG:O | 2.19 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 36:DG:129:GLY:O | 36:DG:130:ASN:CB | 2.67 | 0.42 |
| 37:DH:43:VAL:CG1 | 37:DH:53:GLU:O | 2.67 | 0.42 |
| 38:DI:22:LYS:O | 38:DI:23:PRO:C | 2.63 | 0.42 |
| 38:DI:133:HIS:O | 38:DI:135:GLU:HG3 | 2.20 | 0.42 |
| 39:DN:43:THR:HG22 | 39:DN:45:ASN:ND2 | 2.35 | 0.42 |
| 45:DT:16:ARG:H | 45:DT:79:HIS:CD2 | 2.37 | 0.42 |
| 45:DT:78:LEU:O | 45:DT:79:HIS:CG | 2.73 | 0.42 |
| 46:DU:50:ARG:NH2 | 47:DV:75:PHE:HD2 | 2.18 | 0.42 |
| 47:DV:97:LYS:O | 47:DV:97:LYS:HG3 | 2.20 | 0.42 |
| 48:DW:18:ARG:NH1 | 48:DW:18:ARG:CG | 2.74 | 0.42 |
| 49:DX:44:GLU:HB2 | 49:DX:49:VAL:O | 2.20 | 0.42 |
| 51:DZ:119:GLU:C | 51:DZ:121:HIS:N | 2.78 | 0.42 |
| 51:DZ:163:LEU:C | 51:DZ:165:VAL:H | 2.28 | 0.42 |
| 1:AA:22:G:H4' | 1:AA:885:G:C8 | 2.55 | 0.42 |
| 1:AA:273:A:N6 | 1:AA:274:A:N6 | 2.68 | 0.42 |
| 1:AA:340:U:C2 | 1:AA:350:G:N2 | 2.88 | 0.42 |
| 1:AA:373:A:N3 | 1:AA:374:A:C8 | 2.88 | 0.42 |
| 1:AA:380:G:C2 | 1:AA:384:G:C6 | 3.07 | 0.42 |
| 1:AA:472:A:O2' | 16:AP:81:ARG:HA | 2.19 | 0.42 |
| 1:AA:677:U:O2' | 1:AA:678:U:H5' | 2.20 | 0.42 |
| 1:AA:740:U:O2' | 1:AA:741:G:H5' | 2.20 | 0.42 |
| 1:AA:747:C:H5 | 1:AA:748:C:N4 | 2.18 | 0.42 |
| 1:AA:784:C:H4' | 31:BA:1837:C:OP1 | 2.19 | 0.42 |
| 1:AA:790:A:N6 | 1:AA:791:G:C6 | 2.88 | 0.42 |
| 1:AA:1316:G:H1 | 19:AS:5:LEU:HD21 | 1.85 | 0.42 |
| 3:AC:123:GLN:O | 3:AC:128:PHE:HB2 | 2.20 | 0.42 |
| 4:AD:105:VAL:HG22 | 4:AD:146:ILE:HG21 | 2.02 | 0.42 |
| 5:AE:139:LEU:HD23 | 5:AE:142:LEU:HD11 | 2.01 | 0.42 |
| 9:AI:36:TYR:CE2 | 9:AI:37:PHE:CE2 | 3.08 | 0.42 |
| 9:AI:79:LEU:HD11 | 9:AI:83:ARG:CZ | 2.50 | 0.42 |
| 9:AI:104:ARG:O | 9:AI:104:ARG:HG2 | 2.20 | 0.42 |
| 10:AJ:78:ASN:O | 10:AJ:82:ILE:HG12 | 2.19 | 0.42 |
| 12:AL:105:TYR:O | 12:AL:107:ALA:N | 2.53 | 0.42 |
| 12:AL:110:VAL:HG21 | 12:AL:120:TYR:HB3 | 2.02 | 0.42 |
| 16:AP:39:TYR:CE1 | 16:AP:41:PRO:HA | 2.54 | 0.42 |
| 23:B1:86:SER:C | 23:B1:89:GLU:HG2 | 2.44 | 0.42 |
| 30:B8:30:ARG:NH2 | 41:BP:62:LEU:HB2 | 2.17 | 0.42 |
| 30:B8:35:GLN:HB3 | 30:B8:36:LYS:HZ3 | 1.84 | 0.42 |
| 30:B8:62:LEU:N | 30:B8:63:PRO:CD | 2.82 | 0.42 |
| 31:BA:64:A:O2' | 31:BA:65:C:H5' | 2.19 | 0.42 |
| 31:BA:171:G:H2' | 31:BA:172:C:C1' | 2.44 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:708:C:H42 | 31:BA:723:G:H1 | 1.68 | 0.42 |
| 31:BA:825:C:C6 | 31:BA:825:C:H3' | 2.54 | 0.42 |
| 31:BA:922:U:H2' | 31:BA:923:C:C6 | 2.55 | 0.42 |
| 31:BA:1020:A:N1 | 31:BA:1141:U:O2' | 2.53 | 0.42 |
| 31:BA:1176:G:H1' | 31:BA:1177:A:P | 2.60 | 0.42 |
| 31:BA:1179:C:N3 | 31:BA:1180:C:C6 | 2.88 | 0.42 |
| 31:BA:1342:A:C2 | 31:BA:1345:C:C5 | 3.08 | 0.42 |
| 31:BA:1566:A:OP1 | 33:BD:211:ARG:NH1 | 2.53 | 0.42 |
| 31:BA:2591:C:H2' | 31:BA:2592:G:C8 | 2.55 | 0.42 |
| 33:BD:83:GLU:HB2 | 33:BD:92:ILE:HD12 | 1.99 | 0.42 |
| 34:BE:132:HIS:ND1 | 34:BE:132:HIS:O | 2.53 | 0.42 |
| 34:BE:160:TYR:CD2 | 34:BE:160:TYR:C | 2.98 | 0.42 |
| 35:BF:135:LYS:HB3 | 35:BF:138:GLU:HG3 | 2.02 | 0.42 |
| 35:BF:150:GLY:HA2 | 35:BF:172:TRP:CD2 | 2.54 | 0.42 |
| 36:BG:127:GLY:HA2 | 36:BG:166:ASP:CB | 2.38 | 0.42 |
| 45:BT:31:SER:CA | 45:BT:32:TYR:CD2 | 3.02 | 0.42 |
| 45:BT:98:LYS:HB3 | 45:BT:100:TYR:CE1 | 2.55 | 0.42 |
| 48:BW:36:LEU:CD1 | 48:BW:48:ALA:HA | 2.50 | 0.42 |
| 50:BY:75:ILE:CD1 | 50:BY:79:CYS:C | 2.92 | 0.42 |
| 51:BZ:28:MET:HG2 | 51:BZ:37:VAL:HG21 | 2.01 | 0.42 |
| 1:CA:432:A:C8 | 1:CA:433:C:C5 | 3.08 | 0.42 |
| 1:CA:484:G:C4' | 1:CA:485:G:O5' | 2.68 | 0.42 |
| 1:CA:781:A:H5' | 1:CA:782:A:OP2 | 2.20 | 0.42 |
| 1:CA:1053:G:N7 | 1:CA:1200:C:H5' | 2.35 | 0.42 |
| 2:CB:19:HIS:CD2 | 2:CB:20:GLU:HG2 | 2.55 | 0.42 |
| 5:CE:129:ILE:O | 5:CE:133:TYR:HD1 | 2.03 | 0.42 |
| 6:CF:15:ASP:OD1 | 6:CF:17:SER:HB2 | 2.19 | 0.42 |
| 9:CI:106:ALA:O | 9:CI:108:VAL:N | 2.53 | 0.42 |
| 11:CK:84:VAL:O | 11:CK:85:ARG:HG3 | 2.19 | 0.42 |
| 12:CL:32:PHE:CB | 12:CL:84:LEU:HD21 | 2.49 | 0.42 |
| 16:CP:9:PHE:HB3 | 16:CP:10:GLY:H | 1.68 | 0.42 |
| 17:CQ:31:LEU:HD12 | 17:CQ:31:LEU:O | 2.20 | 0.42 |
| 23:D1:20:ARG:HD3 | 23:D1:41:ARG:HD3 | 2.02 | 0.42 |
| 23:D1:37:ILE:CG2 | 31:DA:2080:G:O5' | 2.68 | 0.42 |
| 24:D2:23:LYS:CB | 49:DX:5:TYR:CE1 | 3.02 | 0.42 |
| 25:D3:22:ALA:O | 25:D3:26:LEU:HG | 2.19 | 0.42 |
| 28:D6:26:ASN:ND2 | 28:D6:32:ASN:ND2 | 2.68 | 0.42 |
| 29:D7:40:TRP:CZ3 | 31:DA:459:U:H4' | 2.55 | 0.42 |
| 30:D8:36:LYS:O | 30:D8:37:SER:O | 2.37 | 0.42 |
| 31:DA:28:A:C2 | 31:DA:513:A:C8 | 3.07 | 0.42 |
| 31:DA:329:G:H22 | 50:DY:19:LYS:HZ2 | 1.65 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:775:G:C2 | 31:DA:777:A:N6 | 2.88 | 0.42 |
| 31:DA:792:G:C4' | 31:DA:793:A:H5' | 2.50 | 0.42 |
| 31:DA:812:C:H2' | 31:DA:813:U:H6 | 1.84 | 0.42 |
| 31:DA:892:G:C8 | 31:DA:893:C:C4 | 3.07 | 0.42 |
| 31:DA:905:U:C2' | 31:DA:906:G:H5' | 2.49 | 0.42 |
| 31:DA:1176:G:H1' | 31:DA:1177:A:P | 2.60 | 0.42 |
| 31:DA:1815:A:OP1 | 31:DA:1815:A:H8 | 2.03 | 0.42 |
| 31:DA:1820:U:C2 | 33:DD:202:LYS:HB3 | 2.55 | 0.42 |
| 31:DA:1853:A:C5 | 31:DA:1889:A:C6 | 3.07 | 0.42 |
| 31:DA:2017:U:H5" | 31:DA:2018:G:OP2 | 2.20 | 0.42 |
| 31:DA:2087:G:H8 | 31:DA:2087:G:O5' | 2.03 | 0.42 |
| 31:DA:2103:C:H3' | 31:DA:2104:G:H5" | 2.01 | 0.42 |
| 31:DA:2406:U:H2' | 31:DA:2406:U:O5' | 2.19 | 0.42 |
| 31:DA:2489:G:C6 | 31:DA:2490:G:N1 | 2.88 | 0.42 |
| 31:DA:2580:U:H4' | 34:DE:130:GLY:HA2 | 2.02 | 0.42 |
| 31:DA:2606:C:C2' | 31:DA:2607:G:H5' | 2.50 | 0.42 |
| 34:DE:76:ARG:HG3 | 34:DE:195:LEU:CD1 | 2.42 | 0.42 |
| 34:DE:101:ARG:HD3 | 34:DE:101:ARG:HA | 1.82 | 0.42 |
| 34:DE:120:TRP:O | 34:DE:121:ASN:HB2 | 2.20 | 0.42 |
| 37:DH:44:VAL:CG1 | 37:DH:45:VAL:N | 2.80 | 0.42 |
| 38:DI:114:LEU:HA | 38:DI:114:LEU:HD23 | 1.77 | 0.42 |
| 38:DI:124:GLY:N | 38:DI:142:VAL:HG23 | 2.35 | 0.42 |
| 40:DO:4:PRO:O | 40:DO:5:GLN:CB | 2.66 | 0.42 |
| 40:DO:11:ALA:O | 40:DO:98:VAL:HG23 | 2.19 | 0.42 |
| 41:DP:59:LEU:HA | 41:DP:61:ARG:CZ | 2.39 | 0.42 |
| 45:DT:78:LEU:HD22 | 45:DT:79:HIS:CE1 | 2.55 | 0.42 |
| 45:DT:100:TYR:HD2 | 45:DT:103:ARG:NH2 | 2.12 | 0.42 |
| 49:DX:76:ARG:O | 49:DX:77:LYS:CB | 2.64 | 0.42 |
| 49:DX:81:VAL:O | 49:DX:81:VAL:HG12 | 2.20 | 0.42 |
| 50:DY:14:LEU:HD12 | 50:DY:23:ARG:H | 1.85 | 0.42 |
| 50:DY:45:VAL:CG2 | 50:DY:62:GLU:N | 2.83 | 0.42 |
| 51:DZ:76:LEU:HD23 | 51:DZ:83:PRO:HA | 2.01 | 0.42 |
| 1:AA:27:G:H2' | 1:AA:28:G:H8 | 1.85 | 0.41 |
| 1:AA:221:C:H2' | 1:AA:222:U:H6 | 1.85 | 0.41 |
| 1:AA:237:C:O2' | 1:AA:238:G:H5' | 2.20 | 0.41 |
| 1:AA:364:A:C2 | 1:AA:365:U:O4 | 2.73 | 0.41 |
| 1:AA:376:G:C6 | 1:AA:389:A:N6 | 2.88 | 0.41 |
| 1:AA:410:G:N2 | 1:AA:431:A:C8 | 2.88 | 0.41 |
| 1:AA:552:U:C2' | 1:AA:553:A:H5' | 2.50 | 0.41 |
| 1:AA:552:U:O2' | 12:AL:86:ARG:O | 2.37 | 0.41 |
| 1:AA:623:C:H2' | 1:AA:624:C:H5' | 2.02 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:720:C:H6 | 1:AA:720:C:O5' | 2.02 | 0.41 |
| 1:AA:785:G:C2' | 1:AA:786:G:H5' | 2.50 | 0.41 |
| 1:AA:920:U:H2' | 1:AA:921:U:C6 | 2.55 | 0.41 |
| 1:AA:965:A:C2 | 1:AA:969:A:N1 | 2.88 | 0.41 |
| 1:AA:1097:C:C2' | 1:AA:1098:C:H5' | 2.49 | 0.41 |
| 1:AA:1205:U:H4' | 3:AC:195:VAL:HG21 | 2.02 | 0.41 |
| 1:AA:1281:U:C5' | 1:AA:1282:C:H5 | 2.32 | 0.41 |
| 1:AA:1366:C:H2' | 1:AA:1367:C:H6 | 1.85 | 0.41 |
| 4:AD:8:VAL:HA | 4:AD:11:LEU:HD21 | 2.02 | 0.41 |
| 4:AD:79:PHE:CE1 | 4:AD:204:ILE:HD13 | 2.55 | 0.41 |
| 8:AH:112:LEU:CB | 8:AH:133:LEU:HA | 2.50 | 0.41 |
| 10:AJ:80:LYS:HB2 | 10:AJ:80:LYS:NZ | 2.35 | 0.41 |
| 12:AL:90:VAL:C | 12:AL:92:ASP:N | 2.76 | 0.41 |
| 13:AM:97:PRO:C | 13:AM:98:VAL:HG13 | 2.45 | 0.41 |
| 22:B0:27:GLU:HB2 | 22:B0:69:PHE:HD1 | 1.85 | 0.41 |
| 22:B0:51:VAL:C | 22:B0:62:LEU:HD12 | 2.45 | 0.41 |
| 28:B6:11:LEU:CD1 | 28:B6:51:GLU:HB2 | 2.50 | 0.41 |
| 28:B6:45:LYS:HD3 | 28:B6:45:LYS:HA | 1.85 | 0.41 |
| 31:BA:184:C:C2 | 31:BA:185:U:C5 | 3.08 | 0.41 |
| 31:BA:543:C:N4 | 31:BA:551:G:N1 | 2.68 | 0.41 |
| 31:BA:607:U:O2 | 31:BA:621:A:N1 | 2.53 | 0.41 |
| 31:BA:880:G:N2 | 31:BA:898:C:C4 | 2.88 | 0.41 |
| 31:BA:970:C:H2' | 31:BA:971:C:C6 | 2.55 | 0.41 |
| 31:BA:1107:G:C6 | 31:BA:1108:U:N3 | 2.88 | 0.41 |
| 31:BA:1131:G:H21 | 39:BN:73:THR:CG2 | 2.33 | 0.41 |
| 31:BA:1181:C:H2' | 31:BA:1182:A:H5' | 2.02 | 0.41 |
| 31:BA:1484:G:N2 | 31:BA:1505:C:H5 | 2.16 | 0.41 |
| 31:BA:1485:G:N3 | 31:BA:1505:C:N4 | 2.68 | 0.41 |
| 31:BA:1833:U:C4 | 31:BA:1834:U:C5 | 3.09 | 0.41 |
| 31:BA:1853:A:C6 | 31:BA:1889:A:C5 | 3.08 | 0.41 |
| 31:BA:2206:G:N2 | 31:BA:2207:G:C5' | 2.46 | 0.41 |
| 31:BA:2611:U:H5' | 31:BA:2611:U:H6 | 1.84 | 0.41 |
| 31:BA:2702:U:H6 | 31:BA:2702:U:H2' | 1.71 | 0.41 |
| 31:BA:2784:C:O2 | 34:BE:37:ARG:NH2 | 2.53 | 0.41 |
| 32:BB:45:A:H1' | 36:BG:95:ARG:NH2 | 2.34 | 0.41 |
| 32:BB:54:G:C2 | 32:BB:55:U:C6 | 3.08 | 0.41 |
| 33:BD:34:VAL:O | 33:BD:34:VAL:HG13 | 2.19 | 0.41 |
| 33:BD:53:PHE:CD1 | 33:BD:220:HIS:HA | 2.55 | 0.41 |
| 34:BE:197:ILE:HG21 | 34:BE:197:ILE:HD13 | 1.81 | 0.41 |
| 36:BG:120:LEU:HD12 | 36:BG:179:PRO:HG2 | 2.01 | 0.41 |
| 38:BI:38:LEU:HD12 | 38:BI:38:LEU:N | 2.30 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 38:BI:75:LEU:HD21 | 38:BI:105:HIS:CE1 | 2.55 | 0.41 |
| 38:BI:83:ALA:HA | 38:BI:89:TYR:CE1 | 2.53 | 0.41 |
| 38:BI:88:ILE:CG2 | 38:BI:89:TYR:H | 2.26 | 0.41 |
| 39:BN:96:GLU:O | 39:BN:100:GLU:HG3 | 2.20 | 0.41 |
| 45:BT:114:LEU:HD23 | 45:BT:114:LEU:HA | 1.89 | 0.41 |
| 47:BV:1:MET:HE3 | 47:BV:44:LYS:CB | 2.32 | 0.41 |
| 48:BW:73:ALA:HB3 | 48:BW:106:ILE:CD1 | 2.50 | 0.41 |
| 1:CA:115:G:C2 | 1:CA:289:G:C5 | 3.08 | 0.41 |
| 1:CA:230:G:H2' | 1:CA:231:G:O5' | 2.20 | 0.41 |
| 1:CA:266:G:C5' | 1:CA:268:C:H41 | 2.27 | 0.41 |
| 1:CA:390:C:O2' | 1:CA:391:G:H5' | 2.19 | 0.41 |
| 1:CA:814:A:N7 | 1:CA:816:A:C5 | 2.87 | 0.41 |
| 1:CA:1400:C:H4' | 1:CA:1401:G:OP2 | 2.19 | 0.41 |
| 1:CA:1458:G:H2' | 1:CA:1459:C:C6 | 2.55 | 0.41 |
| 3:CC:87:LEU:O | 3:CC:91:LEU:HG | 2.20 | 0.41 |
| 4:CD:172:PRO:C | 4:CD:187:ARG:HH12 | 2.28 | 0.41 |
| 5:CE:146:ALA:O | 5:CE:148:VAL:N | 2.53 | 0.41 |
| 8:CH:126:LYS:C | 8:CH:128:GLY:N | 2.78 | 0.41 |
| 10:CJ:50:ILE:HA | 10:CJ:60:ARG:HB2 | 2.02 | 0.41 |
| 10:CJ:70:ARG:HG3 | 10:CJ:70:ARG:NH1 | 2.35 | 0.41 |
| 13:CM:32:GLU:OE2 | 13:CM:64:TRP:CH2 | 2.73 | 0.41 |
| 17:CQ:46:ASP:C | 17:CQ:48:GLU:H | 2.27 | 0.41 |
| 17:CQ:63:ARG:HG2 | 17:CQ:64:PRO:HD2 | 2.02 | 0.41 |
| 18:CR:86:VAL:O | 18:CR:87:ARG:HB3 | 2.20 | 0.41 |
| 22:D0:27:GLU:HB2 | 22:D0:69:PHE:HD1 | 1.84 | 0.41 |
| 23:D1:11:ARG:HB2 | 23:D1:12:PRO:CD | 2.50 | 0.41 |
| 23:D1:94:LEU:HB3 | 23:D1:95:LEU:H | 1.55 | 0.41 |
| 24:D2:14:ARG:NH2 | 24:D2:57:ILE:CG2 | 2.83 | 0.41 |
| 30:D8:58:ILE:O | 30:D8:61:LEU:CG | 2.66 | 0.41 |
| 31:DA:271(F):C:O5' | 31:DA:271(F):C:H6 | 2.03 | 0.41 |
| 31:DA:271(R):G:O2' | 31:DA:271(S):G:H5' | 2.20 | 0.41 |
| 31:DA:389:G:H22 | 41:DP:71:VAL:HG12 | 1.85 | 0.41 |
| 31:DA:513:A:N1 | 31:DA:514:A:C5 | 2.87 | 0.41 |
| 31:DA:1301:A:C8 | 31:DA:1303:G:C8 | 3.08 | 0.41 |
| 31:DA:1330:C:O2' | 31:DA:1331:A:H5' | 2.20 | 0.41 |
| 31:DA:1416:G:O2' | 31:DA:1417:C:OP2 | 2.33 | 0.41 |
| 31:DA:1469:A:C2' | 31:DA:1470:G:O5' | 2.67 | 0.41 |
| 31:DA:1608:A:H1' | 31:DA:1610:A:OP2 | 2.20 | 0.41 |
| 31:DA:2061:G:N3 | 31:DA:2063:C:C5 | 2.88 | 0.41 |
| 31:DA:2219:G:H8 | 31:DA:2219:G:OP2 | 2.03 | 0.41 |
| 31:DA:2262:U:H4' | 31:DA:2328:A:C2 | 2.55 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2306:C:OP2 | 31:DA:2307:G:C8 | 2.73 | 0.41 |
| 31:DA:2465:C:C2' | 31:DA:2466:C:H5' | 2.50 | 0.41 |
| 32:DB:87:G:C5' | 32:DB:88:C:OP2 | 2.68 | 0.41 |
| 33:DD:70:TRP:O | 33:DD:71:ASP:C | 2.63 | 0.41 |
| 34:DE:132:HIS:ND1 | 34:DE:132:HIS:O | 2.53 | 0.41 |
| 35:DF:19:GLU:O | 35:DF:20:LEU:HB2 | 2.19 | 0.41 |
| 35:DF:20:LEU:O | 35:DF:23:ASP:HB2 | 2.20 | 0.41 |
| 35:DF:65:TRP:C | 35:DF:67:GLN:N | 2.78 | 0.41 |
| 35:DF:136:THR:O | 35:DF:137:LYS:C | 2.63 | 0.41 |
| 36:DG:19:LEU:HD13 | 36:DG:32:PRO:HG2 | 2.02 | 0.41 |
| 36:DG:56:ALA:HA | 36:DG:59:GLU:OE1 | 2.20 | 0.41 |
| 39:DN:78:TYR:HD1 | 39:DN:79:PRO:CG | 2.31 | 0.41 |
| 40:DO:118:ALA:HA | 40:DO:119:PRO:HD2 | 1.63 | 0.41 |
| 41:DP:35:HIS:O | 41:DP:36:LYS:HB2 | 2.19 | 0.41 |
| 42:DQ:103:MET:HB2 | 42:DQ:104:PHE:CD1 | 2.55 | 0.41 |
| 44:DS:98:VAL:HG13 | 44:DS:100:ALA:N | 2.35 | 0.41 |
| 49:DX:27:THR:OG1 | 49:DX:77:LYS:HA | 2.20 | 0.41 |
| 49:DX:53:LYS:H | 49:DX:80:ILE:HG22 | 1.85 | 0.41 |
| 50:DY:20:TYR:CD1 | 50:DY:20:TYR:N | 2.88 | 0.41 |
| 50:DY:76:CYS:O | 50:DY:77:PRO:C | 2.63 | 0.41 |
| 51:DZ:165:VAL:CG1 | 51:DZ:166:SER:N | 2.76 | 0.41 |
| 1:AA:53:A:C2 | 1:AA:359:U:O2 | 2.73 | 0.41 |
| 1:AA:108:G:C6 | 20:AT:15:ARG:HD2 | 2.55 | 0.41 |
| 1:AA:191:G:C6 | 1:AA:192:U:C4 | 3.08 | 0.41 |
| 1:AA:246:A:C5 | 1:AA:279:A:C6 | 3.08 | 0.41 |
| 1:AA:364:A:H2' | 1:AA:365:U:O2 | 2.19 | 0.41 |
| 1:AA:432:A:C8 | 1:AA:433:C:C5 | 3.08 | 0.41 |
| 1:AA:516:U:C4 | 1:AA:517:G:C6 | 3.08 | 0.41 |
| 1:AA:559:A:H4' | 1:AA:560:U:H3' | 2.01 | 0.41 |
| 1:AA:1072:G:C6 | 1:AA:1073:U:O4 | 2.73 | 0.41 |
| 1:AA:1271:G:H5' | 1:AA:1314:C:C5' | 2.51 | 0.41 |
| 1:AA:1422:G:O3' | 40:BO:49:ARG:NH2 | 2.52 | 0.41 |
| 2:AB:97:TRP:CH2 | 2:AB:176:GLU:CD | 2.98 | 0.41 |
| 2:AB:181:PHE:O | 2:AB:183:PRO:HD3 | 2.20 | 0.41 |
| 3:AC:134:ILE:CG2 | 3:AC:151:VAL:HB | 2.48 | 0.41 |
| 4:AD:14:ARG:HA | 4:AD:39:PRO:HG3 | 2.02 | 0.41 |
| 4:AD:26:CYS:HA | 4:AD:31:CYS:HB2 | 2.02 | 0.41 |
| 4:AD:31:CYS:O | 4:AD:33:MET:N | 2.51 | 0.41 |
| 4:AD:101:LEU:HB2 | 4:AD:138:TYR:O | 2.20 | 0.41 |
| 4:AD:126:ILE:C | 4:AD:127:THR:HG22 | 2.45 | 0.41 |
| 4:AD:128:VAL:HA | 4:AD:145:GLU:O | 2.20 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 5:AE:72:GLN:O | 5:AE:73:ASN:CB | 2.58 | 0.41 |
| 5:AE:146:ALA:C | 5:AE:148:VAL:H | 2.26 | 0.41 |
| 6:AF:9:VAL:HA | 6:AF:59:TYR:O | 2.20 | 0.41 |
| 8:AH:126:LYS:C | 8:AH:128:GLY:N | 2.78 | 0.41 |
| 11:AK:111:ASP:CA | 18:AR:84:LYS:HE2 | 2.46 | 0.41 |
| 15:AO:32:LEU:O | 15:AO:36:ILE:HG13 | 2.20 | 0.41 |
| 15:AO:39:LEU:O | 15:AO:42:HIS:N | 2.53 | 0.41 |
| 18:AR:31:LEU:HD12 | 18:AR:66:LEU:HB2 | 2.02 | 0.41 |
| 18:AR:45:SER:HB3 | 18:AR:51:LEU:HD21 | 2.02 | 0.41 |
| 19:AS:7:LYS:HA | 19:AS:7:LYS:HD3 | 1.86 | 0.41 |
| 20:AT:75:ASN:O | 20:AT:79:ARG:HB2 | 2.20 | 0.41 |
| 22:B0:56:ASP:O | 22:B0:57:PHE:HB2 | 2.20 | 0.41 |
| 22:B0:68:GLU:CG | 22:B0:80:HIS:HB2 | 2.50 | 0.41 |
| 24:B2:14:ARG:HH11 | 24:B2:57:ILE:N | 2.18 | 0.41 |
| 24:B2:29:LYS:O | 24:B2:30:ARG:C | 2.62 | 0.41 |
| 30:B8:34:TRP:CZ3 | 30:B8:41:ILE:HG22 | 2.55 | 0.41 |
| 31:BA:17:G:H2' | 31:BA:18:C:H6 | 1.85 | 0.41 |
| 31:BA:218:A:H2 | 31:BA:235:U:H4' | 1.84 | 0.41 |
| 31:BA:598:G:H5' | 41:BP:15:ARG:HB2 | 2.02 | 0.41 |
| 31:BA:708:C:O2 | 31:BA:708:C:C2' | 2.63 | 0.41 |
| 31:BA:910:A:H2' | 31:BA:2264:C:O2' | 2.19 | 0.41 |
| 31:BA:954:G:C4 | 31:BA:955:C:C6 | 3.08 | 0.41 |
| 31:BA:1702:G:H2' | 31:BA:1703:G:O5' | 2.20 | 0.41 |
| 31:BA:1964:G:N2 | 31:BA:1967:C:C2 | 2.88 | 0.41 |
| 31:BA:2061:G:N3 | 31:BA:2063:C:C5 | 2.89 | 0.41 |
| 31:BA:2590:A:H2' | 31:BA:2591:C:C6 | 2.55 | 0.41 |
| 32:BB:86:G:H1 | 32:BB:91:C:H42 | 1.68 | 0.41 |
| 32:BB:116:G:C8 | 32:BB:116:G:C4' | 3.03 | 0.41 |
| 33:BD:25:THR:CG2 | 33:BD:82:ILE:N | 2.83 | 0.41 |
| 34:BE:1:MET:N | 34:BE:1:MET:HE2 | 2.35 | 0.41 |
| 34:BE:10:GLY:CA | 45:BT:8:LYS:HE3 | 2.50 | 0.41 |
| 34:BE:144:ARG:HB3 | 34:BE:145:LYS:H | 1.48 | 0.41 |
| 36:BG:96:ARG:O | 36:BG:97:ASP:C | 2.63 | 0.41 |
| 37:BH:29:PRO:C | 37:BH:30:LYS:HG2 | 2.44 | 0.41 |
| 37:BH:41:MET:O | 37:BH:42:ARG:C | 2.63 | 0.41 |
| 38:BI:33:ARG:O | 38:BI:35:LEU:CD2 | 2.68 | 0.41 |
| 39:BN:82:LEU:N | 39:BN:82:LEU:CD1 | 2.82 | 0.41 |
| 41:BP:85:LEU:HD13 | 41:BP:114:ILE:HD11 | 2.01 | 0.41 |
| 41:BP:95:VAL:C | 41:BP:96:THR:CG2 | 2.93 | 0.41 |
| 41:BP:112:LEU:HD22 | 41:BP:112:LEU:C | 2.44 | 0.41 |
| 45:BT:27:THR:O | 45:BT:47:GLY:O | 2.38 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 46:BU:104:GLN:O | 46:BU:107:ALA:HB3 | 2.21 | 0.41 |
| 51:BZ:76:LEU:HD23 | 51:BZ:83:PRO:HA | 2.02 | 0.41 |
| 1:CA:342:C:C2' | 1:CA:343:U:H5' | 2.50 | 0.41 |
| 1:CA:382:A:H2' | 1:CA:383:A:H8 | 1.84 | 0.41 |
| 1:CA:533:A:H4' | 1:CA:534:U:OP1 | 2.20 | 0.41 |
| 1:CA:1254:C:H2' | 1:CA:1255:G:H8 | 1.84 | 0.41 |
| 1:CA:1352:C:OP1 | 21:CU:3:LYS:NZ | 2.48 | 0.41 |
| 1:CA:1442:G:C5 | 1:CA:1442(B):A:N1 | 2.88 | 0.41 |
| 2:CB:178:ARG:HH21 | 8:CH:74:PRO:HG3 | 1.82 | 0.41 |
| 4:CD:3:ARG:O | 4:CD:3:ARG:HD3 | 2.20 | 0.41 |
| 9:CI:104:ARG:O | 9:CI:104:ARG:HG2 | 2.20 | 0.41 |
| 10:CJ:78:ASN:HB3 | 10:CJ:80:LYS:H | 1.84 | 0.41 |
| 12:CL:83:VAL:HG12 | 12:CL:84:LEU:N | 2.35 | 0.41 |
| 14:CN:51:GLY:C | 14:CN:53:LEU:N | 2.78 | 0.41 |
| 16:CP:72:ARG:HH21 | 16:CP:73:LEU:CD2 | 2.23 | 0.41 |
| 16:CP:81:ARG:HD3 | 16:CP:83:GLU:OE2 | 2.20 | 0.41 |
| 27:D5:36:CYS:C | 27:D5:38:ALA:N | 2.77 | 0.41 |
| 31:DA:84:A:H5' | 50:DY:9:LYS:HB3 | 2.01 | 0.41 |
| 31:DA:105:C:H2' | 31:DA:106:C:C6 | 2.55 | 0.41 |
| 31:DA:196:A:H2' | 31:DA:196:A:N3 | 2.35 | 0.41 |
| 31:DA:362:U:H3' | 31:DA:362:U:H6 | 1.85 | 0.41 |
| 31:DA:511:U:O4 | 31:DA:512:G:C6 | 2.73 | 0.41 |
| 31:DA:549:G:H2' | 31:DA:551:G:H5" | 2.03 | 0.41 |
| 31:DA:877:U:C2' | 31:DA:878:A:H5" | 2.49 | 0.41 |
| 31:DA:996:A:OP2 | 46:DU:92:ARG:NH2 | 2.53 | 0.41 |
| 31:DA:1655:A:C8 | 31:DA:1656:C:C5 | 3.08 | 0.41 |
| 31:DA:1665:A:C2' | 31:DA:1666:G:H5' | 2.51 | 0.41 |
| 31:DA:1996:C:H4' | 31:DA:1997:G:OP1 | 2.20 | 0.41 |
| 31:DA:2026:C:C2' | 31:DA:2027:G:O5' | 2.68 | 0.41 |
| 31:DA:2092:U:H5 | 31:DA:2226:C:OP1 | 2.03 | 0.41 |
| 31:DA:2690:C:OP1 | 43:DR:17:ARG:NH1 | 2.53 | 0.41 |
| 31:DA:2885:C:N3 | 31:DA:2886:G:H1' | 2.35 | 0.41 |
| 32:DB:27:C:C4 | 32:DB:28:C:C5 | 3.08 | 0.41 |
| 32:DB:47:C:C2' | 32:DB:48:A:H5' | 2.51 | 0.41 |
| 32:DB:52:A:HO2' | 32:DB:53:A:H8 | 1.57 | 0.41 |
| 33:DD:143:HIS:CD2 | 33:DD:144:ALA:HB2 | 2.51 | 0.41 |
| 33:DD:147:LEU:HD11 | 33:DD:183:ARG:NH1 | 2.35 | 0.41 |
| 34:DE:116:VAL:HG23 | 34:DE:157:ALA:N | 2.35 | 0.41 |
| 37:DH:46:GLU:CG | 37:DH:51:ARG:HB3 | 2.50 | 0.41 |
| 39:DN:125:GLY:HA2 | 39:DN:126:PRO:O | 2.20 | 0.41 |
| 41:DP:101:VAL:HG12 | 41:DP:106:LEU:HD23 | 2.02 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 43:DR:52:ILE:O | 43:DR:55:ALA:N | 2.53 | 0.41 |
| 43:DR:54:LEU:HA | 43:DR:54:LEU:HD12 | 1.74 | 0.41 |
| 46:DU:81:HIS:O | 46:DU:84:LYS:HB3 | 2.20 | 0.41 |
| 47:DV:86:GLY:O | 47:DV:87:HIS:CG | 2.73 | 0.41 |
| 49:DX:23:GLU:HG3 | 49:DX:24:GLY:H | 1.85 | 0.41 |
| 50:DY:7:VAL:CB | 50:DY:8:LYS:HD2 | 2.45 | 0.41 |
| 50:DY:91:GLU:HB3 | 50:DY:92:ASN:H | 1.53 | 0.41 |
| 1:AA:499:A:C4' | 1:AA:500:G:OP1 | 2.61 | 0.41 |
| 1:AA:524:G:C6 | 1:AA:525:C:N4 | 2.88 | 0.41 |
| 1:AA:541:G:C4 | 1:AA:542:G:C8 | 3.09 | 0.41 |
| 1:AA:1073:U:H2' | 1:AA:1074:G:C8 | 2.46 | 0.41 |
| 1:AA:1272:G:C5 | 1:AA:1273:G:C8 | 3.08 | 0.41 |
| 2:AB:80:ILE:HD12 | 2:AB:211:ILE:HB | 2.02 | 0.41 |
| 2:AB:174:VAL:O | 2:AB:178:ARG:HB2 | 2.20 | 0.41 |
| 4:AD:23:GLY:O | 4:AD:27:TYR:CD1 | 2.73 | 0.41 |
| 4:AD:108:LEU:C | 4:AD:110:PHE:H | 2.28 | 0.41 |
| 7:AG:101:LEU:O | 7:AG:105:VAL:HG23 | 2.20 | 0.41 |
| 8:AH:6:ILE:O | 8:AH:7:ALA:C | 2.64 | 0.41 |
| 9:AI:56:LEU:O | 9:AI:56:LEU:HD23 | 2.20 | 0.41 |
| 10:AJ:50:ILE:HA | 10:AJ:60:ARG:HB2 | 2.01 | 0.41 |
| 13:AM:108:ARG:HA | 13:AM:108:ARG:HD2 | 1.90 | 0.41 |
| 16:AP:74:LEU:HD23 | 16:AP:74:LEU:HA | 1.82 | 0.41 |
| 18:AR:50:ILE:HG12 | 18:AR:74:ARG:HH12 | 1.86 | 0.41 |
| 19:AS:12:ASP:OD1 | 19:AS:37:ARG:HD2 | 2.20 | 0.41 |
| 20:AT:30:LYS:HE3 | 20:AT:34:LYS:HE3 | 2.01 | 0.41 |
| 22:B0:46:LYS:HB3 | 22:B0:47:PRO:HD2 | 2.01 | 0.41 |
| 24:B2:45:SER:O | 24:B2:47:ASN:N | 2.52 | 0.41 |
| 25:B3:1:MET:O | 25:B3:3:ARG:N | 2.52 | 0.41 |
| 30:B8:34:TRP:HZ3 | 30:B8:41:ILE:HG22 | 1.85 | 0.41 |
| 30:B8:35:GLN:O | 30:B8:37:SER:N | 2.52 | 0.41 |
| 31:BA:327:G:H2' | 31:BA:328:U:C6 | 2.54 | 0.41 |
| 31:BA:538:G:H2' | 31:BA:539:G:H8 | 1.85 | 0.41 |
| 31:BA:1142(A):A:C5 | 31:BA:1144:G:C5 | 3.08 | 0.41 |
| 31:BA:1174:A:OP1 | 31:BA:1175:U:OP1 | 2.38 | 0.41 |
| 31:BA:1301:A:C4 | 31:BA:1303:G:N7 | 2.88 | 0.41 |
| 31:BA:1359:A:N7 | 31:BA:1372:U:O4 | 2.53 | 0.41 |
| 31:BA:1771:C:C1' | 31:BA:1786:A:C8 | 3.03 | 0.41 |
| 31:BA:1815:A:H8 | 31:BA:1815:A:OP1 | 2.02 | 0.41 |
| 31:BA:1894:C:H2' | 31:BA:1895:C:H6 | 1.86 | 0.41 |
| 31:BA:2061:G:N3 | 31:BA:2063:C:C4 | 2.89 | 0.41 |
| 31:BA:2244:U:H2' | 31:BA:2245:U:O4' | 2.20 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2264:C:C2' | 31:BA:2265:U:H5' | 2.50 | 0.41 |
| 31:BA:2275:C:O2' | 42:BQ:83:MET:HA | 2.21 | 0.41 |
| 31:BA:2316:C:C5 | 31:BA:2317:C:H5 | 2.38 | 0.41 |
| 31:BA:2350:C:H2' | 31:BA:2351:G:O4' | 2.19 | 0.41 |
| 31:BA:2733:A:C3' | 31:BA:2734:A:H5" | 2.51 | 0.41 |
| 33:BD:270:ILE:O | 33:BD:271:ILE:HG13 | 2.20 | 0.41 |
| 34:BE:55:ASN:H | 34:BE:72:VAL:HG11 | 1.84 | 0.41 |
| 36:BG:105:LYS:HB2 | 36:BG:105:LYS:NZ | 2.36 | 0.41 |
| 36:BG:107:LEU:HD23 | 36:BG:111:LEU:HD12 | 2.01 | 0.41 |
| 37:BH:61:HIS:C | 37:BH:63:SER:N | 2.78 | 0.41 |
| 39:BN:33:LEU:HD12 | 39:BN:38:HIS:CE1 | 2.55 | 0.41 |
| 40:BO:3:GLN:HB2 | 40:BO:4:PRO:HD2 | 2.02 | 0.41 |
| 41:BP:121:LYS:HB3 | 41:BP:123:LEU:HD23 | 2.01 | 0.41 |
| 42:BQ:23:GLY:HA2 | 42:BQ:101:ARG:N | 2.35 | 0.41 |
| 43:BR:74:LYS:O | 43:BR:75:LEU:C | 2.64 | 0.41 |
| 48:BW:23:LEU:O | 48:BW:27:LYS:HD2 | 2.19 | 0.41 |
| 48:BW:34:ASN:HD22 | 48:BW:34:ASN:HA | 1.69 | 0.41 |
| 51:BZ:133:ILE:HA | 51:BZ:134:PRO:HD2 | 1.90 | 0.41 |
| 1:CA:274:A:H4' | 1:CA:275:G:OP1 | 2.19 | 0.41 |
| 1:CA:374:A:C2 | 1:CA:375:U:C2 | 3.08 | 0.41 |
| 1:CA:452:A:C2 | 1:CA:453:A:C4 | 3.09 | 0.41 |
| 1:CA:925:G:H1' | 1:CA:1502:A:C4 | 2.54 | 0.41 |
| 1:CA:955:U:H2' | 1:CA:956:U:C6 | 2.55 | 0.41 |
| 1:CA:1014:A:H2 | 1:CA:1219:U:O2 | 2.03 | 0.41 |
| 1:CA:1017:G:O5' | 1:CA:1017:G:H8 | 2.03 | 0.41 |
| 1:CA:1060:C:O2' | 1:CA:1061:G:H5' | 2.21 | 0.41 |
| 1:CA:1205:U:H4' | 3:CC:195:VAL:HG21 | 2.02 | 0.41 |
| 1:CA:1342:C:O2' | 1:CA:1343:G:H5' | 2.21 | 0.41 |
| 2:CB:121:LEU:O | 2:CB:127:ILE:HD11 | 2.20 | 0.41 |
| 4:CD:163:GLU:C | 4:CD:165:MET:N | 2.78 | 0.41 |
| 5:CE:129:ILE:O | 5:CE:132:ALA:HB3 | 2.20 | 0.41 |
| 6:CF:79:LEU:HD12 | 6:CF:88:VAL:HG11 | 2.03 | 0.41 |
| 8:CH:6:ILE:O | 8:CH:7:ALA:C | 2.63 | 0.41 |
| 16:CP:20:VAL:HG22 | 16:CP:21:VAL:N | 2.35 | 0.41 |
| 20:CT:74:LYS:C | 20:CT:76:ALA:N | 2.77 | 0.41 |
| 23:D1:87:PRO:CG | 23:D1:88:LYS:H | 2.33 | 0.41 |
| 31:DA:115:C:O2' | 31:DA:116:C:H5' | 2.20 | 0.41 |
| 31:DA:271(H):G:C4 | 31:DA:271(I):G:C8 | 3.09 | 0.41 |
| 31:DA:280:C:H2' | 31:DA:281:G:C5' | 2.50 | 0.41 |
| 31:DA:286:C:H2' | 31:DA:287:C:C5' | 2.41 | 0.41 |
| 31:DA:363(E):U:H2' | 31:DA:363(F):A:O4' | 2.20 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1155:A:O2' | 31:DA:1156:A:H2' | 2.20 | 0.41 |
| 31:DA:1319:G:C2 | 31:DA:1334:G:C5 | 3.08 | 0.41 |
| 31:DA:1486:A:C6 | 31:DA:1504:C:N4 | 2.76 | 0.41 |
| 31:DA:2500:U:O2 | 31:DA:2504:U:C5 | 2.74 | 0.41 |
| 31:DA:2625:G:H2' | 31:DA:2626:C:C6 | 2.55 | 0.41 |
| 31:DA:2663:G:C5 | 31:DA:2664:G:C5 | 3.08 | 0.41 |
| 31:DA:2762:G:H8 | 31:DA:2762:G:C5' | 2.33 | 0.41 |
| 31:DA:2802:G:OP2 | 31:DA:2803:C:OP2 | 2.38 | 0.41 |
| 32:DB:75:G:N1 | 32:DB:103:G:N2 | 2.68 | 0.41 |
| 32:DB:75:G:N2 | 51:DZ:87:ASP:OD2 | 2.52 | 0.41 |
| 34:DE:92:THR:O | 34:DE:93:VAL:CB | 2.68 | 0.41 |
| 34:DE:97:LYS:O | 34:DE:98:PRO:C | 2.63 | 0.41 |
| 39:DN:24:GLY:O | 39:DN:28:THR:HG22 | 2.21 | 0.41 |
| 40:DO:24:VAL:HG23 | 40:DO:33:ALA:HB2 | 2.02 | 0.41 |
| 41:DP:143:GLY:C | 41:DP:145:PRO:CD | 2.78 | 0.41 |
| 42:DQ:23:GLY:HA2 | 42:DQ:101:ARG:N | 2.35 | 0.41 |
| 42:DQ:41:TRP:HB3 | 42:DQ:94:VAL:HB | 2.01 | 0.41 |
| 44:DS:57:LYS:HG2 | 44:DS:58:LEU:H | 1.85 | 0.41 |
| 45:DT:28:VAL:O | 45:DT:28:VAL:HG12 | 2.19 | 0.41 |
| 45:DT:33:LYS:O | 45:DT:40:THR:O | 2.37 | 0.41 |
| 45:DT:106:SER:C | 45:DT:107:ASP:CG | 2.85 | 0.41 |
| 46:DU:8:VAL:HG12 | 46:DU:9:VAL:N | 2.31 | 0.41 |
| 46:DU:87:GLY:O | 47:DV:52:VAL:HG13 | 2.19 | 0.41 |
| 47:DV:24:LYS:HE3 | 47:DV:68:LYS:HE3 | 2.03 | 0.41 |
| 48:DW:12:ILE:HB | 48:DW:42:ARG:HH12 | 1.85 | 0.41 |
| 48:DW:86:LEU:C | 48:DW:86:LEU:CD1 | 2.89 | 0.41 |
| 50:DY:31:LEU:HD13 | 50:DY:31:LEU:HA | 1.76 | 0.41 |
| 50:DY:56:PRO:HB2 | 50:DY:57:GLN:H | 1.63 | 0.41 |
| 1:AA:72:C:H2' | 1:AA:73:G:C8 | 2.56 | 0.41 |
| 1:AA:167:G:C2' | 1:AA:168:G:H5' | 2.50 | 0.41 |
| 1:AA:421:U:C4 | 3:AC:127:ARG:NH1 | 2.88 | 0.41 |
| 1:AA:448:A:H2' | 1:AA:449:C:H6 | 1.84 | 0.41 |
| 1:AA:541:G:H2' | 1:AA:542:G:C8 | 2.55 | 0.41 |
| 1:AA:543:C:O2' | 1:AA:544:G:H5' | 2.20 | 0.41 |
| 1:AA:734:G:C6 | 1:AA:735:C:C4 | 3.09 | 0.41 |
| 1:AA:781:A:H5' | 1:AA:782:A:OP2 | 2.20 | 0.41 |
| 1:AA:961:U:C4 | 1:AA:962:C:C4 | 3.08 | 0.41 |
| 1:AA:991:U:O2' | 1:AA:992:U:P | 2.78 | 0.41 |
| 1:AA:1067:A:C4' | 1:AA:1068:G:O5' | 2.68 | 0.41 |
| 4:AD:78:LEU:O | 4:AD:79:PHE:C | 2.64 | 0.41 |
| 4:AD:190:ASP:HB3 | 4:AD:193:ASP:OD2 | 2.20 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 15:AO:75:PRO:O | 15:AO:76:GLU:C | 2.63 | 0.41 |
| 16:AP:66:PRO:O | 16:AP:67:THR:C | 2.64 | 0.41 |
| 19:AS:9:VAL:O | 19:AS:10:PHE:CD1 | 2.73 | 0.41 |
| 20:AT:89:ARG:HB2 | 20:AT:104:LEU:CD1 | 2.51 | 0.41 |
| 22:B0:27:GLU:HB2 | 22:B0:69:PHE:CD1 | 2.55 | 0.41 |
| 23:B1:85:LEU:HD22 | 23:B1:85:LEU:HA | 1.83 | 0.41 |
| 27:B5:26:THR:O | 27:B5:26:THR:HG23 | 2.20 | 0.41 |
| 31:BA:154:G:N1 | 31:BA:154(A):C:N4 | 2.67 | 0.41 |
| 31:BA:252:G:P | 41:BP:50:ARG:HH11 | 2.43 | 0.41 |
| 31:BA:620:G:H4' | 31:BA:621:A:H5' | 2.02 | 0.41 |
| 31:BA:685:A:C2 | 31:BA:689:A:C6 | 3.07 | 0.41 |
| 31:BA:1190:G:H5' | 41:BP:35:HIS:HA | 2.01 | 0.41 |
| 31:BA:1202:C:N4 | 31:BA:1203:G:C6 | 2.89 | 0.41 |
| 31:BA:1213:A:H1' | 31:BA:1238:G:N3 | 2.35 | 0.41 |
| 31:BA:1235:G:C6 | 31:BA:1236:G:N1 | 2.88 | 0.41 |
| 31:BA:1356:G:C6 | 31:BA:1357:U:C4 | 3.08 | 0.41 |
| 31:BA:1394:U:C4 | 31:BA:1395:A:C6 | 3.08 | 0.41 |
| 31:BA:1575:C:H2' | 31:BA:1576:U:C6 | 2.56 | 0.41 |
| 31:BA:1814:G:C6 | 31:BA:1815:A:C6 | 3.08 | 0.41 |
| 31:BA:1952:A:OP1 | 40:BO:44:LYS:HE2 | 2.21 | 0.41 |
| 31:BA:2319:G:OP2 | 31:BA:2319:G:H4' | 2.20 | 0.41 |
| 33:BD:246:PRO:HD2 | 33:BD:255:LYS:HG2 | 2.01 | 0.41 |
| 34:BE:37:ARG:HD3 | 34:BE:44:TYR:CE2 | 2.55 | 0.41 |
| 35:BF:3:GLU:CD | 35:BF:21:ALA:H | 2.28 | 0.41 |
| 35:BF:170:LEU:HD23 | 35:BF:172:TRP:HE1 | 1.84 | 0.41 |
| 36:BG:89:GLY:O | 36:BG:90:LEU:O | 2.38 | 0.41 |
| 39:BN:28:THR:HG23 | 39:BN:29:LYS:N | 2.33 | 0.41 |
| 39:BN:112:LEU:O | 39:BN:112:LEU:HD12 | 2.21 | 0.41 |
| 41:BP:61:ARG:HD2 | 41:BP:61:ARG:N | 2.33 | 0.41 |
| 43:BR:2:ARG:HB2 | 43:BR:5:LYS:NZ | 2.35 | 0.41 |
| 43:BR:51:LEU:CD1 | 43:BR:70:LEU:HD21 | 2.51 | 0.41 |
| 43:BR:104:ARG:HD3 | 43:BR:109:ALA:HB3 | 2.03 | 0.41 |
| 45:BT:23:ARG:CB | 45:BT:24:PRO:CD | 2.88 | 0.41 |
| 45:BT:29:ARG:HG2 | 45:BT:84:GLN:HG3 | 2.00 | 0.41 |
| 45:BT:31:SER:C | 45:BT:32:TYR:CG | 2.96 | 0.41 |
| 45:BT:82:LEU:HD12 | 45:BT:82:LEU:H | 1.82 | 0.41 |
| 47:BV:2:PHE:HB3 | 47:BV:3:ALA:H | 1.31 | 0.41 |
| 49:BX:33:LYS:O | 49:BX:34:ALA:C | 2.64 | 0.41 |
| 49:BX:77:LYS:HG2 | 49:BX:78:LYS:N | 2.36 | 0.41 |
| 49:BX:89:ILE:O | 49:BX:89:ILE:HG22 | 2.19 | 0.41 |
| 51:BZ:145:GLU:CG | 51:BZ:146:ILE:H | 2.33 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:376:G:C6 | 1:CA:389:A:N6 | 2.88 | 0.41 |
| 1:CA:445:G:C4 | 1:CA:446:G:C8 | 3.07 | 0.41 |
| 1:CA:543:C:N3 | 1:CA:544:G:N7 | 2.69 | 0.41 |
| 1:CA:568:G:O6 | 12:CL:5:PRO:HD3 | 2.19 | 0.41 |
| 1:CA:716:A:H2' | 1:CA:717:C:O5' | 2.20 | 0.41 |
| 1:CA:783:C:C2 | 1:CA:784:C:C5 | 3.07 | 0.41 |
| 1:CA:1030(C):G:H2' | 1:CA:1030(D):A:C8 | 2.55 | 0.41 |
| 1:CA:1316:G:O3' | 14:CN:18:VAL:HG22 | 2.20 | 0.41 |
| 2:CB:204:ASN:HB3 | 2:CB:210:SER:HB3 | 2.01 | 0.41 |
| 3:CC:3:ASN:O | 3:CC:4:LYS:O | 2.38 | 0.41 |
| 3:CC:14:ILE:O | 3:CC:15:THR:CB | 2.68 | 0.41 |
| 4:CD:190:ASP:O | 4:CD:191:ARG:C | 2.63 | 0.41 |
| 4:CD:194:LEU:HB3 | 4:CD:196:LEU:HD11 | 2.02 | 0.41 |
| 6:CF:72:VAL:CG1 | 6:CF:73:ASN:N | 2.82 | 0.41 |
| 10:CJ:51:ARG:HG3 | 10:CJ:60:ARG:C | 2.46 | 0.41 |
| 11:CK:73:MET:SD | 11:CK:103:LEU:HD21 | 2.61 | 0.41 |
| 16:CP:18:ARG:HD3 | 16:CP:35:LYS:CD | 2.51 | 0.41 |
| 17:CQ:77:VAL:O | 17:CQ:78:GLU:CB | 2.69 | 0.41 |
| 19:CS:75:ALA:HA | 19:CS:76:PRO:HD2 | 1.93 | 0.41 |
| 23:D1:64:ALA:O | 23:D1:65:SER:CB | 2.68 | 0.41 |
| 23:D1:85:LEU:HD22 | 23:D1:85:LEU:HA | 1.84 | 0.41 |
| 24:D2:25:VAL:HG22 | 24:D2:26:ARG:HH11 | 1.84 | 0.41 |
| 27:D5:7:PRO:HA | 31:DA:2615:U:C2 | 2.56 | 0.41 |
| 28:D6:28:ARG:O | 28:D6:32:ASN:HB3 | 2.20 | 0.41 |
| 28:D6:39:TYR:O | 28:D6:49:HIS:HE1 | 2.03 | 0.41 |
| 31:DA:271(R):G:H2' | 31:DA:271(S):G:H8 | 1.84 | 0.41 |
| 31:DA:292:C:H42 | 31:DA:348:G:H1 | 1.68 | 0.41 |
| 31:DA:763:G:C4 | 31:DA:765:G:C8 | 3.08 | 0.41 |
| 31:DA:806:C:OP1 | 31:DA:831:G:H5" | 2.19 | 0.41 |
| 31:DA:869:G:H2' | 31:DA:870:A:O4' | 2.21 | 0.41 |
| 31:DA:953:A:H2' | 31:DA:954:G:H5' | 2.00 | 0.41 |
| 31:DA:1303:G:H1' | 31:DA:1641:A:N1 | 2.35 | 0.41 |
| 31:DA:1531:C:H3' | 31:DA:1532:C:O4' | 2.21 | 0.41 |
| 31:DA:1668:A:N3 | 31:DA:1670:C:C4 | 2.88 | 0.41 |
| 31:DA:1773:A:H2' | 31:DA:1774:C:H5' | 2.03 | 0.41 |
| 31:DA:2394:C:C2' | 31:DA:2395:C:C5' | 2.98 | 0.41 |
| 31:DA:2528:U:H2' | 31:DA:2530:A:O5' | 2.21 | 0.41 |
| 31:DA:2540:C:H2' | 31:DA:2541:A:O4' | 2.21 | 0.41 |
| 31:DA:2801(A):A:H4' | 31:DA:2802:G:H2' | 2.02 | 0.41 |
| 31:DA:2884:U:C2' | 31:DA:2885:C:H5' | 2.51 | 0.41 |
| 32:DB:24:G:N7 | 32:DB:56:G:H2' | 2.35 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 32:DB:115:G:H2' | 32:DB:116:G:H5" | 2.02 | 0.41 |
| 33:DD:106:ILE:HD11 | 33:DD:196:VAL:HG13 | 2.03 | 0.41 |
| 33:DD:248:SER:O | 33:DD:250:TRP:N | 2.54 | 0.41 |
| 34:DE:177:PRO:O | 34:DE:178:GLU:C | 2.63 | 0.41 |
| 38:DI:83:ALA:HA | 38:DI:89:TYR:CE1 | 2.55 | 0.41 |
| 45:DT:27:THR:C | 45:DT:28:VAL:HG23 | 2.43 | 0.41 |
| 45:DT:41:ARG:NH1 | 45:DT:43:GLN:HB2 | 2.35 | 0.41 |
| 46:DU:66:ASN:OD1 | 46:DU:76:TYR:HB3 | 2.21 | 0.41 |
| 48:DW:13:SER:O | 48:DW:16:LYS:HB2 | 2.20 | 0.41 |
| 50:DY:18:GLY:O | 50:DY:19:LYS:C | 2.63 | 0.41 |
| 50:DY:28:LYS:HB2 | 50:DY:37:VAL:CB | 2.32 | 0.41 |
| 1:AA:32:A:C2 | 1:AA:33:A:C4 | 3.08 | 0.41 |
| 1:AA:93:G:C2' | 1:AA:96:U:H5' | 2.50 | 0.41 |
| 1:AA:172:A:N7 | 1:AA:174:C:N3 | 2.69 | 0.41 |
| 1:AA:366:C:H1' | 1:AA:367:U:OP1 | 2.20 | 0.41 |
| 1:AA:764:C:H2' | 1:AA:765:G:C8 | 2.55 | 0.41 |
| 1:AA:923:A:C2 | 1:AA:1395:C:N3 | 2.89 | 0.41 |
| 1:AA:955:U:H2' | 1:AA:956:U:C6 | 2.55 | 0.41 |
| 1:AA:1116:C:C4 | 1:AA:1117:G:C8 | 3.09 | 0.41 |
| 4:AD:3:ARG:HD3 | 4:AD:5:ILE:HG13 | 2.03 | 0.41 |
| 6:AF:8:ILE:HG22 | 6:AF:10:LEU:HD12 | 2.01 | 0.41 |
| 6:AF:8:ILE:HA | 6:AF:87:ARG:O | 2.21 | 0.41 |
| 8:AH:17:THR:O | 8:AH:20:TYR:N | 2.51 | 0.41 |
| 11:AK:112:THR:CG2 | 11:AK:113:PRO:HD2 | 2.50 | 0.41 |
| 15:AO:39:LEU:HD22 | 15:AO:43:LEU:HG | 2.02 | 0.41 |
| 20:AT:87:LYS:HE3 | 20:AT:91:LEU:HD11 | 2.02 | 0.41 |
| 22:B0:38:VAL:HG12 | 22:B0:40:GLN:HG2 | 2.03 | 0.41 |
| 25:B3:52:HIS:CD2 | 32:BB:83:G:C4' | 3.01 | 0.41 |
| 31:BA:146:G:C5' | 31:BA:146:G:C8 | 3.01 | 0.41 |
| 31:BA:322:A:C5 | 31:BA:340:A:C2 | 3.09 | 0.41 |
| 31:BA:372:G:HO2' | 31:BA:373:U:P | 2.44 | 0.41 |
| 31:BA:448:U:H3' | 31:BA:449:A:H5' | 2.01 | 0.41 |
| 31:BA:699:A:H2' | 31:BA:700:G:O4' | 2.20 | 0.41 |
| 31:BA:790:C:H6 | 31:BA:790:C:H2' | 1.58 | 0.41 |
| 31:BA:1047:G:N3 | 31:BA:1111:A:N6 | 2.69 | 0.41 |
| 31:BA:1531:C:C3' | 31:BA:1532:C:C5' | 2.98 | 0.41 |
| 31:BA:1649:G:N1 | 31:BA:2009:G:C6 | 2.88 | 0.41 |
| 31:BA:1722:A:N6 | 31:BA:1741:A:N1 | 2.67 | 0.41 |
| 31:BA:1762:A:O5' | 31:BA:1762:A:C8 | 2.56 | 0.41 |
| 31:BA:2203:U:O2' | 33:BD:151:LYS:HG2 | 2.21 | 0.41 |
| 31:BA:2469:A:O2' | 42:BQ:56:ARG:CG | 2.67 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2677:G:H2' | 31:BA:2678:C:C6 | 2.56 | 0.41 |
| 31:BA:2690:C:OP1 | 43:BR:17:ARG:NH1 | 2.52 | 0.41 |
| 31:BA:2758:A:C3' | 31:BA:2759:G:H5" | 2.51 | 0.41 |
| 32:BB:58:A:C5' | 32:BB:59:A:OP2 | 2.68 | 0.41 |
| 33:BD:205:VAL:O | 33:BD:206:LEU:C | 2.64 | 0.41 |
| 34:BE:119:ARG:HD2 | 34:BE:120:TRP:CE2 | 2.55 | 0.41 |
| 35:BF:83:PHE:O | 35:BF:84:VAL:HG23 | 2.21 | 0.41 |
| 36:BG:60:LEU:HD22 | 36:BG:63:ILE:HG13 | 2.02 | 0.41 |
| 37:BH:46:GLU:HG3 | 37:BH:51:ARG:HB3 | 2.02 | 0.41 |
| 40:BO:10:VAL:O | 40:BO:10:VAL:HG22 | 2.19 | 0.41 |
| 42:BQ:23:GLY:C | 42:BQ:100:GLY:CA | 2.94 | 0.41 |
| 42:BQ:88:GLY:O | 42:BQ:89:ASN:HB2 | 2.20 | 0.41 |
| 43:BR:9:LYS:O | 43:BR:10:LEU:HG | 2.21 | 0.41 |
| 44:BS:74:ALA:CB | 44:BS:103:GLU:HG3 | 2.50 | 0.41 |
| 45:BT:49:VAL:HA | 45:BT:63:VAL:HA | 2.02 | 0.41 |
| 47:BV:27:ALA:O | 47:BV:28:GLU:C | 2.63 | 0.41 |
| 49:BX:21:PHE:CE1 | 49:BX:26:TYR:CG | 3.08 | 0.41 |
| 50:BY:55:TYR:O | 50:BY:56:PRO:C | 2.63 | 0.41 |
| 51:BZ:76:LEU:HD23 | 51:BZ:76:LEU:HA | 1.82 | 0.41 |
| 1:CA:10:A:OP2 | 5:CE:126:ARG:HD3 | 2.21 | 0.41 |
| 1:CA:149:A:HO2' | 1:CA:150:C:H6 | 1.60 | 0.41 |
| 1:CA:246:A:C5 | 1:CA:279:A:C6 | 3.09 | 0.41 |
| 1:CA:985:C:O5' | 1:CA:985:C:H6 | 2.02 | 0.41 |
| 4:CD:49:ARG:HA | 4:CD:49:ARG:NE | 2.36 | 0.41 |
| 6:CF:9:VAL:C | 6:CF:10:LEU:HD12 | 2.46 | 0.41 |
| 9:CI:36:TYR:CE2 | 9:CI:37:PHE:CE2 | 3.08 | 0.41 |
| 9:CI:112:LYS:HA | 9:CI:119:ALA:HA | 2.01 | 0.41 |
| 23:D1:37:ILE:O | 23:D1:38:SER:HB2 | 2.21 | 0.41 |
| 31:DA:214:G:H21 | 31:DA:216:A:H1' | 1.86 | 0.41 |
| 31:DA:238:C:C4 | 31:DA:239:U:C5 | 3.09 | 0.41 |
| 31:DA:243:U:O2' | 31:DA:244:A:H5' | 2.20 | 0.41 |
| 31:DA:286:C:N3 | 31:DA:287:C:C5 | 2.89 | 0.41 |
| 31:DA:827:U:O2 | 31:DA:2246:G:H4' | 2.20 | 0.41 |
| 31:DA:917:A:C2 | 31:DA:918:A:C4 | 3.09 | 0.41 |
| 31:DA:986:C:O2' | 31:DA:987:G:H5' | 2.19 | 0.41 |
| 31:DA:1047:G:N3 | 31:DA:1111:A:N6 | 2.69 | 0.41 |
| 31:DA:1485:G:N3 | 31:DA:1505:C:N4 | 2.68 | 0.41 |
| 31:DA:1751:C:O2' | 31:DA:2861:G:O2' | 2.34 | 0.41 |
| 31:DA:1777:U:C2' | 31:DA:1778:U:H5' | 2.50 | 0.41 |
| 31:DA:2020:A:O2' | 31:DA:2021:C:H5' | 2.20 | 0.41 |
| 31:DA:2462:U:H2' | 31:DA:2463:C:O4' | 2.20 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2590:A:O2' | 31:DA:2591:C:H5' | 2.20 | 0.41 |
| 31:DA:2592:G:C5 | 31:DA:2593:U:C5 | 3.08 | 0.41 |
| 31:DA:2631:G:N2 | 34:DE:61:ARG:NH1 | 2.67 | 0.41 |
| 31:DA:2639:A:N7 | 31:DA:2775:A:H2 | 2.18 | 0.41 |
| 31:DA:2738:A:N1 | 31:DA:2739:U:C2 | 2.88 | 0.41 |
| 33:DD:78:LYS:HE3 | 33:DD:78:LYS:HB2 | 1.83 | 0.41 |
| 33:DD:133:LEU:HA | 33:DD:136:ILE:HD12 | 2.02 | 0.41 |
| 33:DD:231:HIS:CG | 33:DD:232:PRO:HD2 | 2.56 | 0.41 |
| 34:DE:136:ARG:NH1 | 34:DE:136:ARG:CG | 2.79 | 0.41 |
| 35:DF:36:VAL:O | 35:DF:39:TRP:HB3 | 2.19 | 0.41 |
| 35:DF:70:THR:HG22 | 35:DF:72:ARG:H | 1.86 | 0.41 |
| 35:DF:101:LEU:HD12 | 35:DF:101:LEU:HA | 1.77 | 0.41 |
| 36:DG:105:LYS:HB2 | 36:DG:105:LYS:NZ | 2.35 | 0.41 |
| 38:DI:14:ASP:O | 38:DI:17:GLN:CB | 2.68 | 0.41 |
| 39:DN:1:MET:HE2 | 47:DV:20:LEU:CD2 | 2.51 | 0.41 |
| 39:DN:112:LEU:HD12 | 39:DN:112:LEU:O | 2.20 | 0.41 |
| 44:DS:101:LEU:HD22 | 44:DS:102:ALA:N | 2.35 | 0.41 |
| 45:DT:128:GLU:C | 45:DT:130:ALA:N | 2.66 | 0.41 |
| 46:DU:66:ASN:C | 46:DU:66:ASN:HD22 | 2.28 | 0.41 |
| 47:DV:1:MET:N | 47:DV:44:LYS:HD2 | 2.35 | 0.41 |
| 49:DX:60:ARG:CB | 49:DX:72:LYS:O | 2.53 | 0.41 |
| 50:DY:41:GLY:O | 50:DY:42:VAL:C | 2.64 | 0.41 |
| 50:DY:62:GLU:O | 50:DY:63:LYS:HG3 | 2.20 | 0.41 |
| 50:DY:96:ILE:CB | 50:DY:99:CYS:HB3 | 2.48 | 0.41 |
| 1:AA:293:G:C6 | 1:AA:294:U:C4 | 3.08 | 0.41 |
| 1:AA:510:A:O2' | 1:AA:542:G:H1' | 2.21 | 0.41 |
| 1:AA:658:G:C2 | 1:AA:749:C:N3 | 2.89 | 0.41 |
| 1:AA:745:C:H1' | 1:AA:836:G:O2' | 2.21 | 0.41 |
| 1:AA:748:C:C1' | 1:AA:749:C:OP2 | 2.68 | 0.41 |
| 1:AA:749:C:O2' | 1:AA:750:G:H5' | 2.21 | 0.41 |
| 1:AA:1158:C:O2 | 1:AA:1158:C:H2' | 2.21 | 0.41 |
| 1:AA:1241:G:H2' | 1:AA:1242:C:C6 | 2.55 | 0.41 |
| 1:AA:1254:C:H2' | 1:AA:1255:G:H8 | 1.85 | 0.41 |
| 1:AA:1291:G:O3' | 9:AI:39:GLY:HA3 | 2.20 | 0.41 |
| 3:AC:188:LEU:O | 3:AC:189:ALA:CB | 2.67 | 0.41 |
| 5:AE:139:LEU:O | 5:AE:141:GLN:N | 2.54 | 0.41 |
| 6:AF:99:ALA:HB1 | 18:AR:23:LYS:NZ | 2.34 | 0.41 |
| 11:AK:15:ALA:HA | 11:AK:77:MET:HA | 2.03 | 0.41 |
| 14:AN:21:TYR:CD2 | 14:AN:22:THR:O | 2.74 | 0.41 |
| 15:AO:18:PHE:O | 15:AO:19:PRO:C | 2.64 | 0.41 |
| 18:AR:74:ARG:HE | 18:AR:81:PHE:HA | 1.85 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 19:AS:36:ARG:NH1 | 19:AS:75:ALA:HB3 | 2.26 | 0.41 |
| 19:AS:58:VAL:HA | 19:AS:59:PRO:HD2 | 1.97 | 0.41 |
| 22:B0:26:TYR:O | 22:B0:27:GLU:C | 2.64 | 0.41 |
| 23:B1:34:THR:HG22 | 31:BA:388:G:OP1 | 2.18 | 0.41 |
| 24:B2:18:PRO:C | 24:B2:20:GLU:N | 2.78 | 0.41 |
| 27:B5:4:HIS:O | 27:B5:5:PRO:C | 2.60 | 0.41 |
| 27:B5:9:LYS:O | 27:B5:10:LYS:C | 2.63 | 0.41 |
| 31:BA:30:G:H2' | 31:BA:31:C:C6 | 2.56 | 0.41 |
| 31:BA:64:A:C2' | 31:BA:65:C:H5' | 2.51 | 0.41 |
| 31:BA:271(R):G:H2' | 31:BA:271(S):G:H8 | 1.86 | 0.41 |
| 31:BA:358:U:C5 | 31:BA:359:A:N7 | 2.88 | 0.41 |
| 31:BA:667:U:C2' | 31:BA:668:G:H5" | 2.50 | 0.41 |
| 31:BA:792:G:H5" | 31:BA:793:A:H5' | 2.03 | 0.41 |
| 31:BA:814:C:C5 | 41:BP:27:HIS:CD2 | 3.07 | 0.41 |
| 31:BA:924:C:H2' | 31:BA:925:C:H6 | 1.84 | 0.41 |
| 31:BA:1241:A:H2' | 31:BA:1242:A:O5' | 2.21 | 0.41 |
| 31:BA:1497:U:C2' | 31:BA:1497:U:O2 | 2.68 | 0.41 |
| 31:BA:1858:G:OP2 | 31:BA:1858:G:C8 | 2.73 | 0.41 |
| 31:BA:1862:G:C2 | 31:BA:1863:G:C8 | 3.08 | 0.41 |
| 31:BA:1902:C:OP1 | 33:BD:242:ARG:HD2 | 2.20 | 0.41 |
| 31:BA:2252:G:H2' | 31:BA:2253:G:C8 | 2.56 | 0.41 |
| 31:BA:2287:A:H2 | 31:BA:2346:A:C2 | 2.39 | 0.41 |
| 31:BA:2308:G:H21 | 36:BG:79:ASN:ND2 | 2.19 | 0.41 |
| 31:BA:2394:C:C2' | 31:BA:2395:C:C5' | 2.99 | 0.41 |
| 31:BA:2644:G:C2' | 31:BA:2645:G:H5' | 2.50 | 0.41 |
| 31:BA:2753:A:H1' | 31:BA:2754:U:H5' | 2.02 | 0.41 |
| 32:BB:37:C:C5 | 32:BB:38:C:C4 | 3.09 | 0.41 |
| 33:BD:70:TRP:CD1 | 33:BD:71:ASP:N | 2.89 | 0.41 |
| 34:BE:1:MET:HB2 | 34:BE:83:ASP:O | 2.20 | 0.41 |
| 34:BE:92:THR:H | 34:BE:95:ILE:CD1 | 2.20 | 0.41 |
| 34:BE:177:PRO:O | 34:BE:178:GLU:C | 2.63 | 0.41 |
| 34:BE:200:GLU:O | 34:BE:201:THR:O | 2.39 | 0.41 |
| 35:BF:46:ARG:HG2 | 35:BF:46:ARG:NH1 | 2.13 | 0.41 |
| 36:BG:109:VAL:O | 36:BG:112:PRO:HD2 | 2.21 | 0.41 |
| 37:BH:148:ILE:O | 37:BH:149:ARG:C | 2.63 | 0.41 |
| 37:BH:152:ARG:HD2 | 37:BH:152:ARG:HA | 1.69 | 0.41 |
| 38:BI:126:TYR:O | 38:BI:140:LEU:N | 2.48 | 0.41 |
| 39:BN:75:TYR:HD1 | 39:BN:75:TYR:H | 1.69 | 0.41 |
| 42:BQ:12:GLN:HG2 | 42:BQ:73:PRO:HD2 | 2.00 | 0.41 |
| 42:BQ:47:ILE:HG12 | 42:BQ:68:ILE:CD1 | 2.50 | 0.41 |
| 45:BT:11:GLU:N | 45:BT:11:GLU:OE2 | 2.54 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 45:BT:26:ASP:OD2 | 45:BT:26:ASP:C | 2.64 | 0.41 |
| 45:BT:93:ARG:O | 45:BT:94:ALA:O | 2.38 | 0.41 |
| 46:BU:12:ARG:O | 46:BU:13:LYS:C | 2.61 | 0.41 |
| 46:BU:87:GLY:C | 47:BV:52:VAL:HG13 | 2.45 | 0.41 |
| 47:BV:66:ARG:HD3 | 47:BV:67:GLY:C | 2.46 | 0.41 |
| 48:BW:19:LEU:HA | 48:BW:19:LEU:HD12 | 1.62 | 0.41 |
| 1:CA:327:A:O2' | 1:CA:329:A:H8 | 2.03 | 0.41 |
| 1:CA:355:C:H2' | 1:CA:356:A:H5' | 2.01 | 0.41 |
| 1:CA:523:A:N1 | 12:CL:92:ASP:HB2 | 2.36 | 0.41 |
| 1:CA:1486:G:H2' | 1:CA:1487:G:O4' | 2.19 | 0.41 |
| 4:CD:33:MET:HE2 | 4:CD:37:PRO:HA | 2.02 | 0.41 |
| 5:CE:139:LEU:O | 5:CE:141:GLN:N | 2.53 | 0.41 |
| 6:CF:82:ARG:NH1 | 6:CF:82:ARG:HB3 | 2.35 | 0.41 |
| 8:CH:28:ALA:HB3 | 8:CH:57:PRO:HB2 | 2.02 | 0.41 |
| 10:CJ:52:GLY:C | 10:CJ:54:PHE:H | 2.28 | 0.41 |
| 12:CL:62:SER:C | 12:CL:64:TYR:N | 2.76 | 0.41 |
| 15:CO:17:ARG:HD3 | 15:CO:26:GLU:HG3 | 2.02 | 0.41 |
| 16:CP:14:ASN:OD1 | 16:CP:16:HIS:HE1 | 2.03 | 0.41 |
| 17:CQ:95:TYR:O | 17:CQ:97:SER:N | 2.54 | 0.41 |
| 23:D1:37:ILE:HA | 23:D1:37:ILE:HD13 | 1.77 | 0.41 |
| 29:D7:31:LEU:O | 29:D7:32:LYS:C | 2.63 | 0.41 |
| 29:D7:41:ARG:C | 29:D7:43:THR:N | 2.79 | 0.41 |
| 31:DA:286:C:H2' | 31:DA:286:C:O2 | 2.19 | 0.41 |
| 31:DA:476:G:N1 | 31:DA:479:A:OP2 | 2.54 | 0.41 |
| 31:DA:735:A:H3' | 31:DA:736:C:H6 | 1.85 | 0.41 |
| 31:DA:784:A:C5 | 33:DD:229:VAL:HG21 | 2.56 | 0.41 |
| 31:DA:795:C:C2 | 31:DA:796:C:C5 | 3.08 | 0.41 |
| 31:DA:900:A:H5" | 31:DA:901:A:OP2 | 2.20 | 0.41 |
| 31:DA:1315:C:N4 | 31:DA:1337:G:H1 | 2.18 | 0.41 |
| 31:DA:1448:G:O2' | 31:DA:1528(A):A:N1 | 2.38 | 0.41 |
| 31:DA:1497:U:H2' | 31:DA:1498:C:OP1 | 2.20 | 0.41 |
| 31:DA:1508:A:O2' | 31:DA:1509:C:OP1 | 2.38 | 0.41 |
| 31:DA:1694:C:O2 | 31:DA:1694:C:H2' | 2.21 | 0.41 |
| 31:DA:1722:A:N6 | 31:DA:1741:A:N1 | 2.69 | 0.41 |
| 31:DA:2762:G:C2' | 31:DA:2763:G:H5' | 2.51 | 0.41 |
| 33:DD:87:ASN:HD22 | 33:DD:87:ASN:H | 1.59 | 0.41 |
| 34:DE:93:VAL:C | 34:DE:95:ILE:H | 2.29 | 0.41 |
| 34:DE:179:GLU:O | 34:DE:180:ASN:HB2 | 2.20 | 0.41 |
| 35:DF:57:VAL:HG12 | 35:DF:59:TYR:H | 1.85 | 0.41 |
| 36:DG:135:LEU:HD23 | 36:DG:140:ILE:HD11 | 2.02 | 0.41 |
| 37:DH:55:PRO:O | 37:DH:56:SER:C | 2.64 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 38:DI:38:LEU:HD12 | 38:DI:38:LEU:N | 2.30 | 0.41 |
| 39:DN:13:TRP:HZ3 | 39:DN:130:HIS:CE1 | 2.35 | 0.41 |
| 40:DO:77:ILE:CD1 | 40:DO:122:LEU:HB3 | 2.51 | 0.41 |
| 42:DQ:58:PHE:HD1 | 42:DQ:58:PHE:O | 2.04 | 0.41 |
| 44:DS:34:HIS:HD2 | 44:DS:34:HIS:N | 2.17 | 0.41 |
| 44:DS:66:ALA:HA | 44:DS:69:VAL:HG12 | 2.03 | 0.41 |
| 47:DV:25:LEU:O | 47:DV:27:ALA:N | 2.54 | 0.41 |
| 49:DX:32:PRO:CG | 49:DX:72:LYS:HD3 | 2.51 | 0.41 |
| 51:DZ:100:VAL:HA | 51:DZ:101:PRO:HD3 | 1.96 | 0.41 |
| 51:DZ:108:PRO:C | 51:DZ:109:ALA:O | 2.63 | 0.41 |
| 1:AA:194:C:C2’ | 1:AA:195:A:H5” | 2.51 | 0.41 |
| 1:AA:253:U:H2’ | 1:AA:254:G:C8 | 2.52 | 0.41 |
| 1:AA:355:C:H2’ | 1:AA:356:A:H5’ | 2.02 | 0.41 |
| 1:AA:425:G:N2 | 1:AA:426:G:H1’ | 2.36 | 0.41 |
| 1:AA:627:G:N3 | 1:AA:628:G:C8 | 2.89 | 0.41 |
| 1:AA:662:G:H2’ | 1:AA:663:A:H8 | 1.85 | 0.41 |
| 1:AA:1047:G:H5” | 14:AN:4:LYS:HG2 | 2.02 | 0.41 |
| 1:AA:1238:A:N6 | 1:AA:1299:A:N6 | 2.58 | 0.41 |
| 1:AA:1376:U:H2’ | 1:AA:1377:A:C8 | 2.55 | 0.41 |
| 2:AB:111:ARG:NH1 | 2:AB:111:ARG:CG | 2.66 | 0.41 |
| 2:AB:238:LEU:O | 2:AB:240:GLN:N | 2.53 | 0.41 |
| 3:AC:136:GLN:HG2 | 3:AC:140:ARG:NH2 | 2.36 | 0.41 |
| 3:AC:182:ILE:HG23 | 3:AC:202:ILE:C | 2.46 | 0.41 |
| 7:AG:85:TYR:CD1 | 7:AG:154:TYR:HE1 | 2.38 | 0.41 |
| 7:AG:137:LYS:HB3 | 7:AG:137:LYS:HE2 | 1.75 | 0.41 |
| 10:AJ:61:GLU:OE2 | 14:AN:45:ARG:HD2 | 2.20 | 0.41 |
| 12:AL:82:VAL:H | 12:AL:106:ASP:HB2 | 1.86 | 0.41 |
| 16:AP:18:ARG:HD3 | 16:AP:35:LYS:CD | 2.49 | 0.41 |
| 18:AR:26:LEU:HD21 | 18:AR:42:ARG:HD2 | 2.02 | 0.41 |
| 28:B6:46:HIS:ND1 | 28:B6:46:HIS:O | 2.54 | 0.41 |
| 30:B8:4:MET:HB2 | 30:B8:4:MET:HE2 | 1.46 | 0.41 |
| 30:B8:39:LYS:HE3 | 30:B8:42:ARG:HH12 | 1.84 | 0.41 |
| 31:BA:363(D):G:H2’ | 31:BA:363(E):U:C6 | 2.55 | 0.41 |
| 31:BA:718:A:H2’ | 31:BA:719:C:H5’ | 2.03 | 0.41 |
| 31:BA:777:A:N3 | 31:BA:778:G:C8 | 2.89 | 0.41 |
| 31:BA:1274:A:N3 | 31:BA:1297:C:H1’ | 2.36 | 0.41 |
| 31:BA:1472:A:O2’ | 31:BA:1473:G:H5’ | 2.21 | 0.41 |
| 31:BA:1497:U:H2’ | 31:BA:1498:C:OP1 | 2.20 | 0.41 |
| 31:BA:1528:A:O2’ | 31:BA:1528(A):A:H8 | 2.04 | 0.41 |
| 31:BA:1590:U:C3’ | 31:BA:1591:G:H5” | 2.51 | 0.41 |
| 31:BA:1826:G:C6 | 31:BA:1827:C:C4 | 3.08 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2312:U:H2' | 31:BA:2313:C:H5' | 2.03 | 0.41 |
| 31:BA:2516:G:C6 | 31:BA:2517:C:C4 | 3.09 | 0.41 |
| 31:BA:2730:C:H4' | 34:BE:168:MET:O | 2.21 | 0.41 |
| 32:BB:14:U:O3' | 32:BB:108:U:O2' | 2.34 | 0.41 |
| 32:BB:21:G:C6 | 32:BB:63:G:C6 | 3.08 | 0.41 |
| 32:BB:24:G:N7 | 32:BB:56:G:H2' | 2.35 | 0.41 |
| 33:BD:35:LYS:HE3 | 33:BD:65:ILE:N | 2.35 | 0.41 |
| 35:BF:20:LEU:O | 35:BF:23:ASP:CB | 2.69 | 0.41 |
| 35:BF:20:LEU:O | 35:BF:23:ASP:HB2 | 2.21 | 0.41 |
| 35:BF:118:ALA:C | 35:BF:120:GLU:H | 2.28 | 0.41 |
| 36:BG:101:ILE:HG23 | 36:BG:102:PHE:N | 2.35 | 0.41 |
| 37:BH:103:LEU:HG | 37:BH:105:LEU:HD13 | 2.01 | 0.41 |
| 38:BI:125:GLU:OE1 | 38:BI:141:LYS:HG2 | 2.20 | 0.41 |
| 40:BO:68:GLU:HB3 | 40:BO:78:ARG:HH11 | 1.85 | 0.41 |
| 41:BP:106:LEU:HA | 41:BP:106:LEU:HD12 | 1.76 | 0.41 |
| 42:BQ:41:TRP:HB3 | 42:BQ:94:VAL:HB | 2.02 | 0.41 |
| 43:BR:75:LEU:C | 43:BR:75:LEU:CD1 | 2.94 | 0.41 |
| 44:BS:102:ALA:HB3 | 44:BS:103:GLU:HG2 | 2.01 | 0.41 |
| 50:BY:76:CYS:O | 50:BY:77:PRO:C | 2.63 | 0.41 |
| 51:BZ:53:ILE:H | 51:BZ:53:ILE:HG12 | 1.71 | 0.41 |
| 51:BZ:130:PRO:HA | 51:BZ:133:ILE:HD11 | 2.03 | 0.41 |
| 1:CA:342:C:O2' | 1:CA:343:U:H5' | 2.20 | 0.41 |
| 1:CA:380:G:N1 | 1:CA:384:G:C6 | 2.88 | 0.41 |
| 1:CA:542:G:C2 | 1:CA:543:C:C5 | 3.09 | 0.41 |
| 1:CA:944:G:H5" | 1:CA:945:G:OP2 | 2.21 | 0.41 |
| 1:CA:1047:G:H5" | 14:CN:4:LYS:HG2 | 2.02 | 0.41 |
| 1:CA:1350:A:N6 | 1:CA:1373:G:N2 | 2.69 | 0.41 |
| 1:CA:1423:G:H2' | 1:CA:1424:C:O4' | 2.20 | 0.41 |
| 2:CB:17:PHE:O | 2:CB:18:GLY:O | 2.38 | 0.41 |
| 3:CC:156:ARG:H | 3:CC:163:ALA:HA | 1.86 | 0.41 |
| 7:CG:135:VAL:O | 7:CG:139:GLU:HG3 | 2.20 | 0.41 |
| 10:CJ:22:LYS:HD2 | 10:CJ:22:LYS:C | 2.46 | 0.41 |
| 12:CL:64:TYR:HB3 | 12:CL:65:GLU:H | 1.68 | 0.41 |
| 17:CQ:65:ILE:HD12 | 17:CQ:65:ILE:H | 1.84 | 0.41 |
| 18:CR:76:LEU:CD2 | 18:CR:76:LEU:N | 2.83 | 0.41 |
| 18:CR:84:LYS:HA | 18:CR:84:LYS:HD3 | 1.77 | 0.41 |
| 23:D1:60:PHE:CZ | 23:D1:90:ILE:HG21 | 2.54 | 0.41 |
| 24:D2:52:ASP:OD1 | 31:DA:72:U:O2 | 2.39 | 0.41 |
| 24:D2:55:ARG:O | 24:D2:55:ARG:CG | 2.69 | 0.41 |
| 30:D8:35:GLN:HB3 | 30:D8:36:LYS:H | 1.35 | 0.41 |
| 31:DA:1161:C:O2' | 47:DV:8:GLY:HA2 | 2.21 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:1204:A:H61 | 31:DA:1240:U:H2' | 1.86 | 0.41 |
| 31:DA:1661:G:C6 | 31:DA:2000:G:C6 | 3.08 | 0.41 |
| 31:DA:1952:A:C6 | 31:DA:1953:A:C6 | 3.08 | 0.41 |
| 31:DA:2065:C:H2' | 31:DA:2066:C:H6 | 1.85 | 0.41 |
| 31:DA:2246:G:H1' | 31:DA:2426:A:C2 | 2.56 | 0.41 |
| 31:DA:2637:U:H2' | 31:DA:2638:G:H5' | 2.02 | 0.41 |
| 31:DA:2752:C:H2' | 31:DA:2753:A:H5' | 2.02 | 0.41 |
| 31:DA:2758:A:C3' | 31:DA:2759:G:H5" | 2.51 | 0.41 |
| 33:DD:70:TRP:CD1 | 33:DD:71:ASP:N | 2.89 | 0.41 |
| 33:DD:183:ARG:HD2 | 33:DD:270:ILE:HG22 | 2.03 | 0.41 |
| 36:DG:77:ILE:HD13 | 36:DG:77:ILE:HA | 1.88 | 0.41 |
| 38:DI:41:GLU:HA | 38:DI:44:LEU:HB2 | 2.03 | 0.41 |
| 38:DI:108:THR:C | 38:DI:109:ILE:CG1 | 2.93 | 0.41 |
| 39:DN:78:TYR:CD1 | 39:DN:79:PRO:CG | 3.04 | 0.41 |
| 42:DQ:86:GLY:C | 42:DQ:88:GLY:N | 2.76 | 0.41 |
| 43:DR:41:ALA:O | 43:DR:43:GLU:N | 2.53 | 0.41 |
| 43:DR:60:LEU:O | 43:DR:61:HIS:C | 2.62 | 0.41 |
| 44:DS:28:VAL:O | 44:DS:29:PHE:HB3 | 2.20 | 0.41 |
| 46:DU:8:VAL:HG11 | 46:DU:12:ARG:NE | 2.35 | 0.41 |
| 46:DU:25:TRP:O | 46:DU:26:GLY:O | 2.39 | 0.41 |
| 49:DX:70:LEU:O | 49:DX:71:GLY:C | 2.63 | 0.41 |
| 50:DY:13:VAL:HG12 | 50:DY:73:ARG:C | 2.45 | 0.41 |
| 50:DY:45:VAL:HG13 | 50:DY:62:GLU:CG | 2.50 | 0.41 |
| 50:DY:53:PRO:O | 50:DY:56:PRO:O | 2.38 | 0.41 |
| 51:DZ:56:VAL:CG1 | 51:DZ:57:ILE:N | 2.84 | 0.41 |
| 1:AA:78:G:N2 | 1:AA:91:C:H42 | 2.15 | 0.41 |
| 1:AA:328:C:H4' | 1:AA:329:A:O5' | 2.20 | 0.41 |
| 1:AA:545:C:H5" | 4:AD:72:GLU:HG2 | 2.02 | 0.41 |
| 1:AA:1030(C):G:H2' | 1:AA:1030(D):A:C8 | 2.55 | 0.41 |
| 1:AA:1107:C:C4 | 1:AA:1108:G:C8 | 3.09 | 0.41 |
| 1:AA:1191:A:H5" | 3:AC:4:LYS:NZ | 2.36 | 0.41 |
| 1:AA:1284:C:O5' | 1:AA:1284:C:H6 | 2.04 | 0.41 |
| 4:AD:146:ILE:HD12 | 4:AD:146:ILE:H | 1.86 | 0.41 |
| 7:AG:4:ARG:HB3 | 7:AG:5:ARG:HH11 | 1.86 | 0.41 |
| 7:AG:70:LYS:O | 7:AG:138:LYS:HE3 | 2.20 | 0.41 |
| 8:AH:87:SER:HA | 8:AH:93:VAL:CG2 | 2.51 | 0.41 |
| 12:AL:13:LYS:N | 12:AL:13:LYS:HD2 | 2.36 | 0.41 |
| 12:AL:26:ALA:O | 12:AL:27:LEU:HB2 | 2.21 | 0.41 |
| 12:AL:92:ASP:C | 12:AL:93:LEU:HD23 | 2.45 | 0.41 |
| 15:AO:28:GLN:O | 15:AO:29:VAL:C | 2.64 | 0.41 |
| 16:AP:20:VAL:HG22 | 16:AP:21:VAL:N | 2.36 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 20:AT:53:LEU:O | 20:AT:54:LYS:C | 2.64 | 0.41 |
| 28:B6:15:GLU:OE2 | 28:B6:18:ARG:HD2 | 2.20 | 0.41 |
| 31:BA:18:C:H2' | 31:BA:19:C:C6 | 2.55 | 0.41 |
| 31:BA:266:G:C6 | 31:BA:267:C:C4 | 3.08 | 0.41 |
| 31:BA:597:U:H2' | 31:BA:598:G:C8 | 2.56 | 0.41 |
| 31:BA:607:U:OP1 | 35:BF:102:PRO:HA | 2.21 | 0.41 |
| 31:BA:661:C:O3' | 41:BP:18:ARG:HG2 | 2.20 | 0.41 |
| 31:BA:1131:G:H21 | 39:BN:73:THR:HG22 | 1.86 | 0.41 |
| 31:BA:1159:U:O2' | 31:BA:1160:G:H5' | 2.21 | 0.41 |
| 31:BA:1470:G:C6 | 31:BA:1519:G:N7 | 2.89 | 0.41 |
| 31:BA:1494:A:O4' | 31:BA:1495:A:OP1 | 2.39 | 0.41 |
| 31:BA:1788:C:H2' | 31:BA:1789:A:C8 | 2.56 | 0.41 |
| 31:BA:1907:G:O2' | 31:BA:1908:C:H5' | 2.21 | 0.41 |
| 31:BA:2064:C:H2' | 31:BA:2065:C:C6 | 2.56 | 0.41 |
| 31:BA:2307:G:H3' | 31:BA:2307:G:N3 | 2.35 | 0.41 |
| 31:BA:2517:C:N3 | 31:BA:2542:A:N6 | 2.69 | 0.41 |
| 31:BA:2659:G:H5" | 31:BA:2659:G:H8 | 1.85 | 0.41 |
| 31:BA:2791:C:H4' | 31:BA:2792:G:O5' | 2.21 | 0.41 |
| 31:BA:2809:A:C2 | 31:BA:2892:A:N3 | 2.89 | 0.41 |
| 31:BA:2873:A:N3 | 43:BR:6:SER:HB2 | 2.35 | 0.41 |
| 32:BB:87:G:C5' | 32:BB:88:C:OP2 | 2.69 | 0.41 |
| 33:BD:35:LYS:CE | 33:BD:63:ARG:C | 2.94 | 0.41 |
| 35:BF:7:TYR:HB3 | 35:BF:16:GLY:C | 2.46 | 0.41 |
| 35:BF:9:ILE:HG12 | 35:BF:14:PRO:CA | 2.50 | 0.41 |
| 36:BG:19:LEU:HG | 36:BG:175:LEU:CD1 | 2.51 | 0.41 |
| 36:BG:71:THR:HB | 36:BG:89:GLY:CA | 2.50 | 0.41 |
| 36:BG:77:ILE:HD13 | 36:BG:77:ILE:HA | 1.85 | 0.41 |
| 39:BN:24:GLY:O | 39:BN:28:THR:HG22 | 2.21 | 0.41 |
| 47:BV:69:LYS:CG | 47:BV:70:ILE:H | 2.32 | 0.41 |
| 49:BX:84:ALA:C | 49:BX:86:GLY:N | 2.78 | 0.41 |
| 50:BY:7:VAL:HB | 50:BY:8:LYS:CE | 2.50 | 0.41 |
| 50:BY:75:ILE:CD1 | 50:BY:79:CYS:HA | 2.39 | 0.41 |
| 1:CA:152:A:C8 | 1:CA:153:C:C5 | 3.08 | 0.41 |
| 1:CA:189(F):U:N3 | 17:CQ:72:ARG:NH1 | 2.69 | 0.41 |
| 1:CA:484:G:H4' | 1:CA:485:G:O5' | 2.14 | 0.41 |
| 1:CA:638:G:O2' | 1:CA:639:G:H5' | 2.20 | 0.41 |
| 1:CA:690:G:O2' | 1:CA:691:G:H5' | 2.21 | 0.41 |
| 1:CA:814:A:C8 | 1:CA:816:A:C8 | 3.09 | 0.41 |
| 1:CA:1400:C:O5' | 1:CA:1400:C:H6 | 2.04 | 0.41 |
| 2:CB:84:GLU:OE1 | 2:CB:216:SER:HA | 2.20 | 0.41 |
| 6:CF:48:LEU:HD21 | 6:CF:60:PHE:CZ | 2.56 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 7:CG:69:VAL:HA | 7:CG:138:LYS:HD2 | 2.03 | 0.41 |
| 8:CH:21:LYS:O | 8:CH:63:LEU:HD23 | 2.21 | 0.41 |
| 11:CK:58:PRO:HA | 11:CK:90:GLY:HA2 | 2.03 | 0.41 |
| 13:CM:75:ALA:O | 13:CM:79:LYS:HG3 | 2.21 | 0.41 |
| 16:CP:3:LYS:C | 16:CP:4:ILE:HD12 | 2.46 | 0.41 |
| 18:CR:65:ILE:H | 18:CR:65:ILE:HG13 | 1.60 | 0.41 |
| 19:CS:12:ASP:HB2 | 19:CS:15:LEU:HD23 | 2.03 | 0.41 |
| 20:CT:30:LYS:HD2 | 20:CT:30:LYS:HA | 1.69 | 0.41 |
| 20:CT:89:ARG:HB2 | 20:CT:104:LEU:CD1 | 2.51 | 0.41 |
| 22:D0:43:THR:N | 31:DA:2331:G:H4' | 2.36 | 0.41 |
| 24:D2:26:ARG:HG2 | 49:DX:5:TYR:CB | 2.51 | 0.41 |
| 28:D6:19:ARG:HD2 | 28:D6:43:CYS:SG | 2.60 | 0.41 |
| 30:D8:25:MET:HE3 | 30:D8:25:MET:HB3 | 1.78 | 0.41 |
| 30:D8:32:LEU:CA | 30:D8:34:TRP:H | 2.31 | 0.41 |
| 31:DA:271(D):G:C5 | 31:DA:271(E):U:C5 | 3.09 | 0.41 |
| 31:DA:452:G:N3 | 31:DA:457:A:H2 | 2.19 | 0.41 |
| 31:DA:740:U:H2' | 31:DA:741:G:C8 | 2.56 | 0.41 |
| 31:DA:787:U:C5 | 31:DA:791:C:C4 | 3.08 | 0.41 |
| 31:DA:917:A:N1 | 31:DA:918:A:C4 | 2.89 | 0.41 |
| 31:DA:956:G:OP2 | 42:DQ:14:ARG:NH2 | 2.54 | 0.41 |
| 31:DA:1131:G:C2 | 31:DA:1132:A:C4 | 3.09 | 0.41 |
| 31:DA:1190:G:H4' | 41:DP:35:HIS:HB3 | 2.03 | 0.41 |
| 31:DA:1260:G:H2' | 31:DA:1261:C:C6 | 2.55 | 0.41 |
| 31:DA:1613:G:C2 | 31:DA:1619:G:C5 | 3.08 | 0.41 |
| 31:DA:1710:C:O2' | 31:DA:1711:C:H5' | 2.20 | 0.41 |
| 31:DA:1812:A:C2 | 31:DA:1813:G:C4 | 3.09 | 0.41 |
| 31:DA:2244:U:H2' | 31:DA:2245:U:O4' | 2.21 | 0.41 |
| 31:DA:2422:A:C4 | 31:DA:2424:C:C5 | 3.09 | 0.41 |
| 31:DA:2579:C:H4' | 34:DE:134:ILE:HD12 | 2.01 | 0.41 |
| 31:DA:2642:G:H4' | 39:DN:78:TYR:OH | 2.21 | 0.41 |
| 32:DB:92:C:H5" | 51:DZ:79:ARG:HH22 | 1.86 | 0.41 |
| 33:DD:143:HIS:CD2 | 33:DD:144:ALA:CB | 3.04 | 0.41 |
| 34:DE:52:LEU:O | 34:DE:75:VAL:N | 2.52 | 0.41 |
| 35:DF:70:THR:HG21 | 35:DF:72:ARG:HB2 | 2.03 | 0.41 |
| 36:DG:71:THR:HB | 36:DG:89:GLY:CA | 2.51 | 0.41 |
| 37:DH:41:MET:O | 37:DH:42:ARG:C | 2.63 | 0.41 |
| 38:DI:4:ILE:HD11 | 38:DI:44:LEU:CD2 | 2.50 | 0.41 |
| 38:DI:123:LEU:HA | 38:DI:142:VAL:HG21 | 2.02 | 0.41 |
| 39:DN:15:LEU:HD21 | 39:DN:55:VAL:CG2 | 2.50 | 0.41 |
| 41:DP:39:LYS:C | 41:DP:41:ARG:N | 2.78 | 0.41 |
| 41:DP:101:VAL:C | 41:DP:103:ALA:N | 2.79 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 41:DP:105:LEU:N | 41:DP:105:LEU:CD1 | 2.82 | 0.41 |
| 44:DS:49:VAL:CG2 | 44:DS:80:LEU:HD11 | 2.51 | 0.41 |
| 45:DT:36:GLU:HB3 | 45:DT:38:ASN:OD1 | 2.21 | 0.41 |
| 45:DT:129:ARG:HD2 | 45:DT:131:ALA:HB3 | 2.03 | 0.41 |
| 46:DU:88:ILE:H | 46:DU:88:ILE:HD12 | 1.85 | 0.41 |
| 49:DX:65:ARG:NH1 | 49:DX:66:LEU:CA | 2.84 | 0.41 |
| 1:AA:10:A:H2' | 1:AA:11:G:C8 | 2.54 | 0.41 |
| 1:AA:92:C:H2' | 1:AA:93:G:C8 | 2.56 | 0.41 |
| 1:AA:149:A:O2' | 1:AA:150:C:H6 | 2.02 | 0.41 |
| 1:AA:189:G:C6 | 1:AA:189(L):G:C6 | 3.09 | 0.41 |
| 1:AA:189(F):U:N3 | 17:AQ:72:ARG:NH1 | 2.69 | 0.41 |
| 1:AA:189(K):U:H6 | 1:AA:189(K):U:O5' | 2.04 | 0.41 |
| 1:AA:192:U:O2' | 1:AA:193:C:H5' | 2.21 | 0.41 |
| 1:AA:245:C:O2 | 1:AA:283:C:N3 | 2.54 | 0.41 |
| 1:AA:393:A:C2 | 1:AA:394:G:C8 | 3.08 | 0.41 |
| 1:AA:410:G:OP2 | 4:AD:25:ARG:HG3 | 2.20 | 0.41 |
| 1:AA:427:U:H3' | 1:AA:428:G:H2' | 2.03 | 0.41 |
| 1:AA:431:A:H2' | 1:AA:432:A:O4' | 2.20 | 0.41 |
| 1:AA:438:G:OP1 | 4:AD:125:HIS:HE1 | 2.04 | 0.41 |
| 1:AA:552:U:H4' | 12:AL:86:ARG:HG2 | 2.02 | 0.41 |
| 1:AA:862:C:C2' | 1:AA:863:U:H5' | 2.50 | 0.41 |
| 1:AA:872:A:C4 | 1:AA:874:G:C8 | 3.08 | 0.41 |
| 1:AA:1003:G:C2' | 1:AA:1004:A:H4' | 2.43 | 0.41 |
| 1:AA:1053:G:C4 | 1:AA:1199:U:C5 | 3.09 | 0.41 |
| 1:AA:1293:G:HO2' | 1:AA:1294:G:P | 2.44 | 0.41 |
| 1:AA:1318:A:N6 | 14:AN:18:VAL:HG11 | 2.36 | 0.41 |
| 1:AA:1392:G:H21 | 1:AA:1502:A:H8 | 1.67 | 0.41 |
| 2:AB:19:HIS:CD2 | 2:AB:20:GLU:HG2 | 2.56 | 0.41 |
| 2:AB:185:ILE:HG22 | 2:AB:199:TYR:CD1 | 2.56 | 0.41 |
| 3:AC:91:LEU:HB3 | 3:AC:99:VAL:HG21 | 2.03 | 0.41 |
| 4:AD:18:LYS:HD2 | 4:AD:33:MET:HG2 | 2.02 | 0.41 |
| 4:AD:102:ASP:HB3 | 4:AD:136:PRO:HA | 2.01 | 0.41 |
| 4:AD:173:TRP:O | 4:AD:174:LEU:HD23 | 2.20 | 0.41 |
| 5:AE:12:LEU:CD1 | 5:AE:31:LEU:HB2 | 2.51 | 0.41 |
| 7:AG:75:VAL:CG2 | 7:AG:144:MET:HB3 | 2.50 | 0.41 |
| 8:AH:73:ASP:OD2 | 8:AH:75:ARG:HG3 | 2.21 | 0.41 |
| 10:AJ:49:VAL:O | 10:AJ:60:ARG:HB2 | 2.20 | 0.41 |
| 11:AK:21:ILE:HD12 | 11:AK:21:ILE:N | 2.35 | 0.41 |
| 11:AK:21:ILE:HA | 11:AK:30:VAL:HG12 | 2.02 | 0.41 |
| 11:AK:102:GLY:O | 11:AK:103:LEU:HD13 | 2.21 | 0.41 |
| 17:AQ:31:LEU:HG | 17:AQ:32:TYR:CE2 | 2.56 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 17:AQ:68:ARG:N | 17:AQ:70:ARG:NH1 | 2.68 | 0.41 |
| 20:AT:57:ARG:NH1 | 20:AT:57:ARG:HB2 | 2.36 | 0.41 |
| 23:B1:54:ALA:HB1 | 23:B1:56:GLN:O | 2.21 | 0.41 |
| 24:B2:55:ARG:O | 24:B2:55:ARG:CG | 2.68 | 0.41 |
| 27:B5:6:VAL:HA | 27:B5:7:PRO:HD3 | 1.94 | 0.41 |
| 28:B6:39:TYR:O | 28:B6:49:HIS:HE1 | 2.04 | 0.41 |
| 29:B7:31:LEU:O | 29:B7:32:LYS:C | 2.64 | 0.41 |
| 31:BA:58:G:OP1 | 49:BX:72:LYS:HA | 2.20 | 0.41 |
| 31:BA:549:G:H2' | 31:BA:551:G:H5" | 2.03 | 0.41 |
| 31:BA:558:G:P | 39:BN:111:PRO:HD2 | 2.61 | 0.41 |
| 31:BA:645:C:O2 | 31:BA:645:C:C3' | 2.69 | 0.41 |
| 31:BA:681:G:H2' | 31:BA:682:G:O4' | 2.20 | 0.41 |
| 31:BA:685:A:N7 | 31:BA:774:A:C5 | 2.89 | 0.41 |
| 31:BA:755:C:H2' | 31:BA:756:C:H6 | 1.85 | 0.41 |
| 31:BA:842:G:N2 | 31:BA:937:U:C2 | 2.89 | 0.41 |
| 31:BA:868:U:C4 | 31:BA:869:G:N7 | 2.89 | 0.41 |
| 31:BA:972:G:C6 | 31:BA:973:A:C6 | 3.09 | 0.41 |
| 31:BA:1267:U:O2 | 31:BA:1267:U:H2' | 2.20 | 0.41 |
| 31:BA:1591:G:H2' | 31:BA:1592:C:O4' | 2.20 | 0.41 |
| 31:BA:1819:A:H5" | 33:BD:158:ALA:HB2 | 2.03 | 0.41 |
| 31:BA:1839:G:C8 | 31:BA:1927:A:C1' | 3.03 | 0.41 |
| 31:BA:2301:C:H6 | 31:BA:2301:C:H3' | 1.86 | 0.41 |
| 31:BA:2468:G:C6 | 31:BA:2481:G:C4 | 3.09 | 0.41 |
| 31:BA:2549:G:C2' | 31:BA:2550:G:H5' | 2.51 | 0.41 |
| 31:BA:2563:U:H4' | 40:BO:28:SER:HA | 2.03 | 0.41 |
| 31:BA:2655:G:C2 | 31:BA:2664:G:O6 | 2.74 | 0.41 |
| 31:BA:2753:A:O2' | 31:BA:2754:U:C6 | 2.72 | 0.41 |
| 31:BA:2863:C:O2' | 31:BA:2864:G:H5' | 2.21 | 0.41 |
| 32:BB:31:C:O2' | 32:BB:32:C:H5' | 2.21 | 0.41 |
| 32:BB:33:G:N2 | 32:BB:50:G:C4 | 2.88 | 0.41 |
| 33:BD:48:ARG:O | 33:BD:50:THR:HG23 | 2.20 | 0.41 |
| 33:BD:49:ILE:HG13 | 33:BD:50:THR:N | 2.36 | 0.41 |
| 33:BD:237:GLU:OE2 | 33:BD:239:ARG:HA | 2.21 | 0.41 |
| 34:BE:97:LYS:O | 34:BE:98:PRO:C | 2.62 | 0.41 |
| 35:BF:7:TYR:HB3 | 35:BF:16:GLY:N | 2.36 | 0.41 |
| 35:BF:68:LYS:O | 35:BF:68:LYS:CG | 2.69 | 0.41 |
| 35:BF:167:ALA:HB1 | 35:BF:173:VAL:HG11 | 2.03 | 0.41 |
| 36:BG:39:ILE:HD12 | 36:BG:40:ASN:N | 2.35 | 0.41 |
| 36:BG:40:ASN:HD22 | 36:BG:91:ARG:HB2 | 1.86 | 0.41 |
| 36:BG:111:LEU:CA | 36:BG:114:ILE:HG12 | 2.49 | 0.41 |
| 37:BH:164:TYR:O | 37:BH:165:ALA:HB2 | 2.20 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 38:BI:4:ILE:HD11 | 38:BI:44:LEU:CD2 | 2.49 | 0.41 |
| 38:BI:85:GLU:O | 38:BI:123:LEU:HD12 | 2.19 | 0.41 |
| 39:BN:1:MET:HE2 | 47:BV:20:LEU:CD2 | 2.51 | 0.41 |
| 42:BQ:54:MET:SD | 42:BQ:118:LEU:HD23 | 2.61 | 0.41 |
| 43:BR:74:LYS:HD2 | 43:BR:77:ARG:HH21 | 1.86 | 0.41 |
| 44:BS:16:ASN:ND2 | 44:BS:92:TYR:CE1 | 2.89 | 0.41 |
| 44:BS:89:ARG:O | 44:BS:90:GLY:O | 2.39 | 0.41 |
| 44:BS:93:LYS:O | 44:BS:94:TYR:C | 2.64 | 0.41 |
| 46:BU:92:ARG:HH22 | 47:BV:10:LYS:HB3 | 1.86 | 0.41 |
| 47:BV:1:MET:N | 47:BV:44:LYS:HD2 | 2.36 | 0.41 |
| 47:BV:2:PHE:HE1 | 47:BV:13:ARG:NH1 | 2.15 | 0.41 |
| 48:BW:14:PRO:O | 48:BW:15:ARG:C | 2.61 | 0.41 |
| 50:BY:79:CYS:O | 50:BY:80:GLY:O | 2.38 | 0.41 |
| 51:BZ:53:ILE:HG13 | 51:BZ:53:ILE:O | 2.21 | 0.41 |
| 1:CA:10:A:H2' | 1:CA:11:G:C8 | 2.55 | 0.41 |
| 1:CA:141:A:H4' | 1:CA:182:U:H1' | 2.02 | 0.41 |
| 1:CA:174:C:H6 | 1:CA:174:C:O5' | 2.03 | 0.41 |
| 1:CA:221:C:H2' | 1:CA:222:U:H6 | 1.86 | 0.41 |
| 1:CA:258:G:C2 | 1:CA:259:G:C8 | 3.09 | 0.41 |
| 1:CA:373:A:C4 | 1:CA:482:A:N7 | 2.89 | 0.41 |
| 1:CA:448:A:H2' | 1:CA:449:C:H6 | 1.84 | 0.41 |
| 1:CA:501:C:HO2' | 1:CA:502:G:H5' | 1.86 | 0.41 |
| 1:CA:503:C:C2 | 1:CA:504:C:C5 | 3.09 | 0.41 |
| 1:CA:790:A:C6 | 1:CA:791:G:C6 | 3.08 | 0.41 |
| 1:CA:827:U:C2 | 1:CA:870:U:C4 | 3.09 | 0.41 |
| 1:CA:939:G:H5" | 7:CG:102:ARG:HH22 | 1.86 | 0.41 |
| 1:CA:991:U:O2' | 1:CA:992:U:P | 2.78 | 0.41 |
| 1:CA:1067:A:C4' | 1:CA:1068:G:O5' | 2.69 | 0.41 |
| 1:CA:1288:A:O2' | 1:CA:1289:A:H5' | 2.21 | 0.41 |
| 1:CA:1333:A:H2' | 1:CA:1334:G:O4' | 2.21 | 0.41 |
| 1:CA:1495:U:H2' | 1:CA:1496:C:C6 | 2.56 | 0.41 |
| 2:CB:74:LYS:NZ | 2:CB:76:GLN:HB2 | 2.35 | 0.41 |
| 2:CB:101:MET:O | 2:CB:105:PHE:HA | 2.21 | 0.41 |
| 3:CC:136:GLN:HG2 | 3:CC:140:ARG:NH2 | 2.35 | 0.41 |
| 3:CC:155:GLY:O | 3:CC:156:ARG:CB | 2.68 | 0.41 |
| 4:CD:4:TYR:O | 4:CD:5:ILE:HB | 2.21 | 0.41 |
| 4:CD:17:VAL:HG11 | 4:CD:197:PRO:CB | 2.51 | 0.41 |
| 4:CD:43:HIS:O | 4:CD:45:GLN:N | 2.54 | 0.41 |
| 4:CD:79:PHE:CZ | 4:CD:204:ILE:HD13 | 2.56 | 0.41 |
| 4:CD:151:LYS:HG2 | 4:CD:151:LYS:O | 2.20 | 0.41 |
| 6:CF:8:ILE:HA | 6:CF:87:ARG:O | 2.20 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 7:CG:69:VAL:HG12 | 7:CG:100:ALA:HA | 2.03 | 0.41 |
| 7:CG:79:ARG:HA | 7:CG:84:ASN:ND2 | 2.36 | 0.41 |
| 8:CH:88:LYS:CB | 8:CH:89:PRO:HD2 | 2.50 | 0.41 |
| 9:CI:43:ALA:HA | 9:CI:74:ILE:HD13 | 2.02 | 0.41 |
| 9:CI:79:LEU:HD11 | 9:CI:83:ARG:CZ | 2.51 | 0.41 |
| 10:CJ:48:THR:HA | 10:CJ:62:HIS:CB | 2.50 | 0.41 |
| 11:CK:19:ALA:HA | 11:CK:32:ILE:HA | 2.03 | 0.41 |
| 13:CM:97:PRO:C | 13:CM:98:VAL:HG13 | 2.45 | 0.41 |
| 15:CO:82:ILE:HD13 | 15:CO:82:ILE:C | 2.46 | 0.41 |
| 18:CR:40:LEU:O | 18:CR:43:PHE:N | 2.53 | 0.41 |
| 19:CS:9:VAL:O | 19:CS:9:VAL:HG12 | 2.21 | 0.41 |
| 20:CT:18:GLN:O | 20:CT:19:SER:C | 2.64 | 0.41 |
| 20:CT:53:LEU:HD21 | 20:CT:92:LEU:CD1 | 2.51 | 0.41 |
| 22:D0:40:GLN:OE1 | 22:D0:44:ARG:HB2 | 2.20 | 0.41 |
| 22:D0:52:GLY:N | 22:D0:62:LEU:HD12 | 2.36 | 0.41 |
| 23:D1:87:PRO:HB2 | 23:D1:91:LYS:HD2 | 2.03 | 0.41 |
| 25:D3:30:ARG:O | 25:D3:33:GLN:HB3 | 2.21 | 0.41 |
| 30:D8:32:LEU:HB2 | 30:D8:35:GLN:H | 1.81 | 0.41 |
| 31:DA:58:G:H1 | 31:DA:69:C:H42 | 1.69 | 0.41 |
| 31:DA:103:A:C2' | 31:DA:104:U:H5' | 2.51 | 0.41 |
| 31:DA:105:C:H6 | 31:DA:105:C:O5' | 2.04 | 0.41 |
| 31:DA:213:A:O2' | 31:DA:214:G:H5' | 2.21 | 0.41 |
| 31:DA:228:A:H5' | 31:DA:229:A:OP2 | 2.21 | 0.41 |
| 31:DA:231:C:O2' | 31:DA:232:G:H5' | 2.21 | 0.41 |
| 31:DA:475:U:C5 | 31:DA:481:G:O6 | 2.74 | 0.41 |
| 31:DA:856:C:H4' | 31:DA:857:C:OP1 | 2.19 | 0.41 |
| 31:DA:856:C:O2' | 31:DA:857:C:P | 2.79 | 0.41 |
| 31:DA:1210:A:H5' | 31:DA:1212:G:O4' | 2.21 | 0.41 |
| 31:DA:1221(A):C:C2 | 31:DA:1229:G:C2 | 3.09 | 0.41 |
| 31:DA:1284:A:H2' | 31:DA:1285:G:O4' | 2.21 | 0.41 |
| 31:DA:1290:C:H2' | 31:DA:1291:C:C6 | 2.55 | 0.41 |
| 31:DA:1300:U:O2' | 31:DA:1626:G:N2 | 2.53 | 0.41 |
| 31:DA:1459:G:C8 | 31:DA:1461:G:H1' | 2.56 | 0.41 |
| 31:DA:1470:G:C6 | 31:DA:1519:G:N7 | 2.88 | 0.41 |
| 31:DA:1480:G:C6 | 31:DA:1481:U:N3 | 2.89 | 0.41 |
| 31:DA:1511:C:C2' | 31:DA:1512:U:O5' | 2.69 | 0.41 |
| 31:DA:1614:A:N1 | 48:DW:91:GLY:HA2 | 2.36 | 0.41 |
| 31:DA:1809:A:C6 | 31:DA:1810:A:C6 | 3.09 | 0.41 |
| 31:DA:1826:G:H2' | 31:DA:1827:C:C6 | 2.56 | 0.41 |
| 31:DA:1847:A:H4' | 31:DA:1848:A:OP2 | 2.21 | 0.41 |
| 31:DA:2321:G:N3 | 31:DA:2321:G:H2' | 2.35 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:2393:A:H2' | 31:DA:2394:C:O4' | 2.21 | 0.41 |
| 31:DA:2463:C:O2 | 31:DA:2488:A:C2 | 2.74 | 0.41 |
| 31:DA:2468:G:H5" | 42:DQ:120:ILE:HD12 | 2.03 | 0.41 |
| 31:DA:2523:G:H2' | 31:DA:2524:G:H5" | 1.93 | 0.41 |
| 31:DA:2768:C:N4 | 31:DA:2769:C:N4 | 2.69 | 0.41 |
| 31:DA:2776:A:C2 | 31:DA:2778:A:C4 | 3.08 | 0.41 |
| 31:DA:2787:C:O2 | 34:DE:61:ARG:NH1 | 2.54 | 0.41 |
| 31:DA:2848:G:H3' | 45:DT:95:ARG:O | 2.21 | 0.41 |
| 31:DA:2884:U:C5 | 31:DA:2885:C:C5 | 3.09 | 0.41 |
| 32:DB:29:A:H5" | 44:DS:32:LEU:HD11 | 2.03 | 0.41 |
| 32:DB:116:G:C8 | 32:DB:116:G:C4' | 3.04 | 0.41 |
| 33:DD:5:LYS:HD2 | 33:DD:5:LYS:N | 2.35 | 0.41 |
| 33:DD:48:ARG:HH11 | 33:DD:48:ARG:CG | 2.33 | 0.41 |
| 33:DD:61:LEU:O | 33:DD:63:ARG:NH1 | 2.53 | 0.41 |
| 33:DD:231:HIS:CD2 | 33:DD:232:PRO:HD2 | 2.56 | 0.41 |
| 34:DE:37:ARG:NH1 | 34:DE:80:GLU:OE2 | 2.52 | 0.41 |
| 35:DF:9:ILE:HG12 | 35:DF:14:PRO:CA | 2.50 | 0.41 |
| 35:DF:9:ILE:O | 35:DF:128:ALA:HB2 | 2.21 | 0.41 |
| 35:DF:20:LEU:O | 35:DF:23:ASP:CB | 2.68 | 0.41 |
| 35:DF:108:LYS:HA | 35:DF:108:LYS:HD3 | 1.67 | 0.41 |
| 36:DG:15:VAL:HG13 | 36:DG:175:LEU:HD13 | 2.02 | 0.41 |
| 36:DG:37:VAL:HA | 36:DG:158:ALA:O | 2.21 | 0.41 |
| 36:DG:40:ASN:HD22 | 36:DG:91:ARG:HB2 | 1.85 | 0.41 |
| 36:DG:42:GLY:O | 36:DG:44:GLY:N | 2.54 | 0.41 |
| 37:DH:89:ILE:HB | 37:DH:90:LYS:H | 1.54 | 0.41 |
| 39:DN:100:GLU:O | 39:DN:101:HIS:C | 2.61 | 0.41 |
| 40:DO:68:GLU:HB3 | 40:DO:78:ARG:NH1 | 2.36 | 0.41 |
| 41:DP:101:VAL:HB | 41:DP:107:LYS:N | 2.36 | 0.41 |
| 42:DQ:78:PRO:O | 42:DQ:79:LEU:CG | 2.69 | 0.41 |
| 42:DQ:133:ARG:O | 42:DQ:134:ARG:HB2 | 2.21 | 0.41 |
| 43:DR:52:ILE:O | 43:DR:53:HIS:C | 2.64 | 0.41 |
| 44:DS:54:LEU:HD21 | 44:DS:59:LYS:O | 2.20 | 0.41 |
| 44:DS:77:ALA:O | 44:DS:80:LEU:HD12 | 2.21 | 0.41 |
| 45:DT:7:ILE:HG22 | 45:DT:11:GLU:OE1 | 2.21 | 0.41 |
| 45:DT:31:SER:CA | 45:DT:32:TYR:CD2 | 3.04 | 0.41 |
| 45:DT:32:TYR:CD2 | 45:DT:81:PRO:HB2 | 2.55 | 0.41 |
| 46:DU:44:ASN:O | 46:DU:45:TYR:C | 2.63 | 0.41 |
| 46:DU:95:LEU:HD13 | 47:DV:11:GLN:HG3 | 2.01 | 0.41 |
| 47:DV:49:THR:HA | 47:DV:50:PRO:HD3 | 1.81 | 0.41 |
| 49:DX:80:ILE:O | 49:DX:81:VAL:HB | 2.19 | 0.41 |
| 50:DY:80:GLY:O | 50:DY:81:LYS:CB | 2.68 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 51:DZ:10:ARG:NH2 | 51:DZ:26:GLY:O | 2.54 | 0.41 |
| 51:DZ:157:LEU:HA | 51:DZ:158:PRO:HD2 | 1.89 | 0.41 |
| 1:AA:167:G:O2' | 1:AA:168:G:H5' | 2.21 | 0.41 |
| 1:AA:191:G:C6 | 1:AA:192:U:N3 | 2.89 | 0.41 |
| 1:AA:327:A:O2' | 1:AA:329:A:H8 | 2.04 | 0.41 |
| 1:AA:503:C:C2 | 1:AA:504:C:C5 | 3.09 | 0.41 |
| 1:AA:520:A:N1 | 1:AA:536:C:H1' | 2.36 | 0.41 |
| 3:AC:70:VAL:HG12 | 3:AC:71:ALA:N | 2.36 | 0.41 |
| 3:AC:124:ILE:HG13 | 3:AC:130:VAL:HG22 | 2.02 | 0.41 |
| 3:AC:178:LEU:C | 3:AC:180:ALA:H | 2.29 | 0.41 |
| 8:AH:88:LYS:CB | 8:AH:89:PRO:HD2 | 2.51 | 0.41 |
| 9:AI:111:ARG:HG3 | 14:AN:61:TRP:HE1 | 1.86 | 0.41 |
| 12:AL:28:LYS:CE | 12:AL:33:ARG:HH12 | 2.34 | 0.41 |
| 15:AO:39:LEU:CD1 | 15:AO:56:LEU:HB2 | 2.51 | 0.41 |
| 16:AP:9:PHE:HB3 | 16:AP:10:GLY:H | 1.69 | 0.41 |
| 18:AR:65:ILE:H | 18:AR:65:ILE:HG13 | 1.58 | 0.41 |
| 22:B0:43:THR:CG2 | 31:BA:2331:G:O3' | 2.68 | 0.41 |
| 23:B1:56:GLN:HB3 | 23:B1:57:GLU:H | 1.67 | 0.41 |
| 31:BA:764:A:C6 | 31:BA:781:A:C2 | 3.10 | 0.41 |
| 31:BA:794:G:H2' | 31:BA:795:C:H6 | 1.82 | 0.41 |
| 31:BA:1330:C:O2' | 31:BA:1331:A:H5' | 2.21 | 0.41 |
| 31:BA:1831:G:H2' | 31:BA:1832:C:C6 | 2.55 | 0.41 |
| 31:BA:1975:G:C2 | 31:BA:1976:U:C2 | 3.09 | 0.41 |
| 31:BA:2290:G:O2' | 31:BA:2381:C:H1' | 2.21 | 0.41 |
| 31:BA:2494:G:C2' | 31:BA:2495:G:O5' | 2.69 | 0.41 |
| 31:BA:2663:G:C6 | 31:BA:2664:G:C5 | 3.09 | 0.41 |
| 33:BD:70:TRP:CZ3 | 33:BD:150:LYS:HA | 2.56 | 0.41 |
| 33:BD:98:VAL:HG23 | 33:BD:98:VAL:H | 1.57 | 0.41 |
| 33:BD:133:LEU:O | 33:BD:134:ARG:C | 2.64 | 0.41 |
| 35:BF:144:LYS:C | 35:BF:146:ALA:N | 2.79 | 0.41 |
| 36:BG:107:LEU:HD11 | 36:BG:178:PHE:CD1 | 2.56 | 0.41 |
| 36:BG:118:ARG:HB2 | 36:BG:181:ARG:CZ | 2.51 | 0.41 |
| 37:BH:136:ILE:H | 37:BH:136:ILE:HG13 | 1.76 | 0.41 |
| 38:BI:25:TYR:CE1 | 38:BI:30:LEU:HD11 | 2.55 | 0.41 |
| 38:BI:123:LEU:HA | 38:BI:142:VAL:HG21 | 2.03 | 0.41 |
| 40:BO:14:THR:HG22 | 40:BO:16:ALA:H | 1.86 | 0.41 |
| 40:BO:40:VAL:HG12 | 40:BO:41:ALA:N | 2.36 | 0.41 |
| 41:BP:73:GLY:O | 41:BP:74:GLU:C | 2.64 | 0.41 |
| 41:BP:103:ALA:O | 41:BP:104:GLY:O | 2.39 | 0.41 |
| 42:BQ:97:VAL:HG11 | 42:BQ:103:MET:HE3 | 2.02 | 0.41 |
| 43:BR:97:VAL:HG22 | 43:BR:114:VAL:HG22 | 2.01 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 44:BS:105:ALA:C | 44:BS:107:GLU:N | 2.76 | 0.41 |
| 47:BV:89:GLN:NE2 | 47:BV:91:TYR:HD1 | 2.18 | 0.41 |
| 1:CA:175:C:C2 | 1:CA:176:C:C6 | 3.08 | 0.41 |
| 1:CA:451:A:N7 | 1:CA:481:G:C6 | 2.89 | 0.41 |
| 2:CB:35:GLU:HA | 2:CB:39:ILE:O | 2.21 | 0.41 |
| 4:CD:14:ARG:HA | 4:CD:39:PRO:HG3 | 2.03 | 0.41 |
| 4:CD:101:LEU:HB2 | 4:CD:138:TYR:O | 2.21 | 0.41 |
| 4:CD:106:TYR:HE1 | 4:CD:113:SER:CA | 2.34 | 0.41 |
| 5:CE:12:LEU:CD1 | 5:CE:31:LEU:HB2 | 2.51 | 0.41 |
| 5:CE:127:ASN:O | 5:CE:128:PRO:C | 2.64 | 0.41 |
| 6:CF:89:MET:HG2 | 6:CF:89:MET:O | 2.21 | 0.41 |
| 13:CM:90:LEU:C | 13:CM:92:HIS:H | 2.28 | 0.41 |
| 16:CP:74:LEU:HD23 | 16:CP:74:LEU:HA | 1.79 | 0.41 |
| 17:CQ:78:GLU:O | 17:CQ:78:GLU:HG3 | 2.20 | 0.41 |
| 20:CT:87:LYS:HE3 | 20:CT:91:LEU:HD21 | 2.03 | 0.41 |
| 22:D0:43:THR:O | 22:D0:44:ARG:C | 2.64 | 0.41 |
| 28:D6:9:LEU:HD22 | 28:D6:9:LEU:C | 2.43 | 0.41 |
| 31:DA:991:C:O2 | 31:DA:1164:G:C2 | 2.74 | 0.41 |
| 31:DA:1207:C:H6 | 31:DA:1207:C:O5' | 2.04 | 0.41 |
| 31:DA:1322:A:N6 | 31:DA:1331:A:H61 | 2.18 | 0.41 |
| 31:DA:1764:G:O2' | 31:DA:1765:C:H5' | 2.21 | 0.41 |
| 31:DA:2078:C:C4 | 31:DA:2079:U:C4 | 3.09 | 0.41 |
| 31:DA:2225:A:H1' | 31:DA:2226:C:OP2 | 2.20 | 0.41 |
| 31:DA:2552:U:H2' | 31:DA:2554:U:OP2 | 2.22 | 0.41 |
| 31:DA:2590:A:H2' | 31:DA:2591:C:C6 | 2.56 | 0.41 |
| 34:DE:33:VAL:O | 34:DE:33:VAL:HG22 | 2.20 | 0.41 |
| 34:DE:50:GLY:HA3 | 34:DE:74:PRO:HG3 | 2.02 | 0.41 |
| 35:DF:34:TRP:HB2 | 41:DP:10:PRO:O | 2.21 | 0.41 |
| 35:DF:57:VAL:HG11 | 35:DF:59:TYR:HD1 | 1.84 | 0.41 |
| 36:DG:127:GLY:HA2 | 36:DG:166:ASP:CB | 2.40 | 0.41 |
| 38:DI:45:LYS:O | 38:DI:48:GLU:N | 2.54 | 0.41 |
| 38:DI:126:TYR:O | 38:DI:140:LEU:N | 2.47 | 0.41 |
| 38:DI:128:LEU:HB3 | 38:DI:129:THR:H | 1.77 | 0.41 |
| 38:DI:130:TYR:CD2 | 38:DI:136:VAL:HG13 | 2.56 | 0.41 |
| 38:DI:130:TYR:HD2 | 38:DI:136:VAL:HG13 | 1.86 | 0.41 |
| 38:DI:130:TYR:HD2 | 38:DI:136:VAL:CG1 | 2.34 | 0.41 |
| 39:DN:18:ALA:O | 39:DN:21:LYS:N | 2.48 | 0.41 |
| 41:DP:102:ARG:O | 41:DP:103:ALA:CB | 2.68 | 0.41 |
| 41:DP:126:VAL:O | 41:DP:127:ALA:HB2 | 2.21 | 0.41 |
| 43:DR:67:LEU:HD13 | 43:DR:67:LEU:HA | 1.82 | 0.41 |
| 45:DT:50:ILE:HA | 45:DT:99:LEU:CD1 | 2.51 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 45:DT:106:SER:CB | 45:DT:110:ILE:HD11 | 2.51 | 0.41 |
| 46:DU:83:LEU:HD13 | 46:DU:113:ALA:HB2 | 2.02 | 0.41 |
| 50:DY:14:LEU:CD1 | 50:DY:22:GLY:HA2 | 2.50 | 0.41 |
| 1:AA:126:G:H4' | 1:AA:634:C:O2 | 2.21 | 0.40 |
| 1:AA:246:A:C2 | 1:AA:282:A:C6 | 3.09 | 0.40 |
| 1:AA:263:A:OP2 | 20:AT:79:ARG:NH1 | 2.54 | 0.40 |
| 1:AA:587:G:C2 | 1:AA:755:G:C5 | 3.09 | 0.40 |
| 1:AA:763:G:H2' | 1:AA:764:C:H6 | 1.86 | 0.40 |
| 1:AA:919:A:C2' | 1:AA:920:U:O5' | 2.70 | 0.40 |
| 1:AA:1117:G:O6 | 1:AA:1156:G:N2 | 2.51 | 0.40 |
| 1:AA:1248:A:O2' | 9:AI:36:TYR:HE1 | 2.03 | 0.40 |
| 4:AD:43:HIS:CE1 | 4:AD:46:LYS:HZ2 | 2.36 | 0.40 |
| 11:AK:21:ILE:HG13 | 11:AK:30:VAL:CG1 | 2.51 | 0.40 |
| 11:AK:61:ALA:CB | 11:AK:90:GLY:HA3 | 2.51 | 0.40 |
| 16:AP:82:GLN:N | 16:AP:82:GLN:NE2 | 2.63 | 0.40 |
| 22:B0:43:THR:H | 31:BA:2331:G:H4' | 1.85 | 0.40 |
| 28:B6:36:LEU:O | 28:B6:37:ARG:HD2 | 2.21 | 0.40 |
| 31:BA:108:U:H2' | 31:BA:109:G:C8 | 2.57 | 0.40 |
| 31:BA:209:C:O5' | 31:BA:209:C:H6 | 2.04 | 0.40 |
| 31:BA:253:C:C2' | 31:BA:254:G:H5' | 2.51 | 0.40 |
| 31:BA:282:A:C4 | 31:BA:359:A:C2 | 3.09 | 0.40 |
| 31:BA:484:C:O2' | 31:BA:485:C:H5' | 2.21 | 0.40 |
| 31:BA:542:C:N4 | 31:BA:543:C:H42 | 2.18 | 0.40 |
| 31:BA:586:A:C2 | 31:BA:1254:A:C2 | 3.10 | 0.40 |
| 31:BA:594:U:C2 | 31:BA:595:C:C5 | 3.08 | 0.40 |
| 31:BA:639:U:C2 | 31:BA:640:C:C5 | 3.09 | 0.40 |
| 31:BA:733:G:H8 | 31:BA:733:G:O5' | 2.04 | 0.40 |
| 31:BA:1048:A:O2' | 31:BA:1049:C:OP2 | 2.36 | 0.40 |
| 31:BA:1468:C:C2 | 31:BA:1525:G:N2 | 2.89 | 0.40 |
| 31:BA:1817:G:P | 33:BD:88:ARG:HH22 | 2.44 | 0.40 |
| 31:BA:1948:G:H2' | 31:BA:1949:G:H5' | 2.03 | 0.40 |
| 31:BA:2075:U:C4 | 31:BA:2238:G:C6 | 3.08 | 0.40 |
| 31:BA:2358:G:C5 | 31:BA:2359:C:C5 | 3.09 | 0.40 |
| 31:BA:2426:A:H3' | 31:BA:2427:C:H5' | 2.03 | 0.40 |
| 31:BA:2674:G:H2' | 31:BA:2675:A:O4' | 2.21 | 0.40 |
| 31:BA:2734:A:C5' | 31:BA:2734:A:C8 | 3.01 | 0.40 |
| 32:BB:59:A:H2' | 32:BB:60:C:H6 | 1.87 | 0.40 |
| 34:BE:37:ARG:O | 34:BE:45:THR:HA | 2.21 | 0.40 |
| 34:BE:137:HIS:CB | 34:BE:138:PRO:CD | 2.99 | 0.40 |
| 35:BF:20:LEU:HD22 | 35:BF:23:ASP:OD2 | 2.21 | 0.40 |
| 35:BF:53:THR:HG23 | 35:BF:56:GLU:H | 1.85 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 36:BG:175:LEU:HD23 | 36:BG:175:LEU:HA | 1.92 | 0.40 |
| 40:BO:24:VAL:CG2 | 40:BO:33:ALA:HB2 | 2.51 | 0.40 |
| 41:BP:121:LYS:HB3 | 41:BP:123:LEU:CD2 | 2.52 | 0.40 |
| 43:BR:8:ARG:NE | 43:BR:8:ARG:CA | 2.80 | 0.40 |
| 43:BR:48:VAL:O | 43:BR:49:ASP:C | 2.64 | 0.40 |
| 44:BS:15:ARG:C | 44:BS:17:ARG:O | 2.64 | 0.40 |
| 44:BS:67:ARG:C | 44:BS:69:VAL:N | 2.77 | 0.40 |
| 45:BT:47:GLY:HA3 | 45:BT:63:VAL:HG23 | 2.02 | 0.40 |
| 47:BV:75:PHE:CD1 | 47:BV:75:PHE:N | 2.89 | 0.40 |
| 48:BW:5:ALA:CB | 48:BW:50:VAL:HG23 | 2.51 | 0.40 |
| 49:BX:24:GLY:HA3 | 49:BX:80:ILE:HG12 | 2.00 | 0.40 |
| 50:BY:41:GLY:O | 50:BY:43:ASN:OD1 | 2.39 | 0.40 |
| 50:BY:62:GLU:O | 50:BY:63:LYS:HG3 | 2.21 | 0.40 |
| 1:CA:364:A:H2' | 1:CA:365:U:O2 | 2.20 | 0.40 |
| 1:CA:565:U:C6 | 1:CA:566:G:C8 | 3.09 | 0.40 |
| 1:CA:622:A:N7 | 1:CA:623:C:C5 | 2.89 | 0.40 |
| 1:CA:709:G:O2' | 1:CA:710:G:H5' | 2.19 | 0.40 |
| 1:CA:748:C:C1' | 1:CA:749:C:OP2 | 2.66 | 0.40 |
| 1:CA:763:G:C5 | 1:CA:764:C:C5 | 3.08 | 0.40 |
| 1:CA:1061:G:C4 | 1:CA:1197:G:N2 | 2.88 | 0.40 |
| 1:CA:1173:G:C5 | 1:CA:1174:G:N7 | 2.89 | 0.40 |
| 1:CA:1320:C:O2' | 19:CS:73:GLU:HG2 | 2.21 | 0.40 |
| 1:CA:1352:C:O2 | 1:CA:1371:G:C2 | 2.73 | 0.40 |
| 2:CB:104:ASN:O | 2:CB:108:ILE:HG12 | 2.22 | 0.40 |
| 2:CB:154:LEU:HD23 | 2:CB:154:LEU:N | 2.20 | 0.40 |
| 4:CD:106:TYR:CE1 | 4:CD:113:SER:C | 2.99 | 0.40 |
| 7:CG:129:GLU:OE1 | 7:CG:131:LYS:HE2 | 2.20 | 0.40 |
| 9:CI:53:VAL:HB | 9:CI:92:TYR:HE2 | 1.86 | 0.40 |
| 12:CL:62:SER:O | 12:CL:64:TYR:HD1 | 2.03 | 0.40 |
| 12:CL:119:LYS:C | 12:CL:120:TYR:CD1 | 2.99 | 0.40 |
| 15:CO:28:GLN:O | 15:CO:29:VAL:C | 2.64 | 0.40 |
| 17:CQ:59:ILE:HG22 | 17:CQ:60:ILE:N | 2.35 | 0.40 |
| 20:CT:20:LEU:O | 20:CT:21:LYS:C | 2.64 | 0.40 |
| 20:CT:76:ALA:O | 20:CT:77:ALA:C | 2.62 | 0.40 |
| 22:D0:53:MET:HA | 22:D0:58:THR:O | 2.21 | 0.40 |
| 23:D1:34:THR:HG21 | 31:DA:388:G:OP1 | 2.11 | 0.40 |
| 23:D1:61:ARG:HH22 | 31:DA:1364:G:P | 2.44 | 0.40 |
| 24:D2:38:GLN:HE21 | 24:D2:38:GLN:HB3 | 1.64 | 0.40 |
| 30:D8:34:TRP:CZ3 | 30:D8:41:ILE:HG22 | 2.55 | 0.40 |
| 31:DA:64:A:C2 | 31:DA:65:C:C2 | 3.09 | 0.40 |
| 31:DA:154(A):C:O2 | 31:DA:154(A):C:O4' | 2.38 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:311:A:C6 | 31:DA:328:U:C4 | 3.09 | 0.40 |
| 31:DA:736:C:O2' | 31:DA:737:C:H5' | 2.21 | 0.40 |
| 31:DA:790:C:O2' | 31:DA:791:C:C5' | 2.68 | 0.40 |
| 31:DA:814:C:H5" | 47:DV:86:GLY:CA | 2.45 | 0.40 |
| 31:DA:880:G:N2 | 31:DA:898:C:C4 | 2.89 | 0.40 |
| 31:DA:1034:G:H2' | 31:DA:1035:U:H6 | 1.86 | 0.40 |
| 31:DA:1461:G:O2' | 31:DA:1462:C:H5' | 2.21 | 0.40 |
| 31:DA:1578:U:C2' | 31:DA:1579:A:H5" | 2.51 | 0.40 |
| 31:DA:1657:C:H2' | 31:DA:1658:C:H6 | 1.86 | 0.40 |
| 31:DA:1697:G:C8 | 31:DA:1697:G:C5' | 2.92 | 0.40 |
| 31:DA:2282:G:H4' | 31:DA:2389:G:O2' | 2.20 | 0.40 |
| 31:DA:2552:U:O5' | 31:DA:2552:U:H6 | 2.04 | 0.40 |
| 33:DD:31:LYS:HG2 | 33:DD:34:VAL:HG13 | 2.02 | 0.40 |
| 33:DD:86:PRO:C | 33:DD:87:ASN:ND2 | 2.79 | 0.40 |
| 38:DI:15:VAL:CG2 | 38:DI:16:GLY:N | 2.84 | 0.40 |
| 38:DI:69:LYS:O | 38:DI:69:LYS:HG2 | 2.21 | 0.40 |
| 42:DQ:76:LYS:N | 42:DQ:88:GLY:HA2 | 2.36 | 0.40 |
| 43:DR:29:LEU:HD12 | 43:DR:29:LEU:HA | 1.84 | 0.40 |
| 43:DR:74:LYS:HD2 | 43:DR:77:ARG:HH21 | 1.86 | 0.40 |
| 44:DS:93:LYS:HD2 | 44:DS:93:LYS:HA | 1.72 | 0.40 |
| 45:DT:31:SER:C | 45:DT:32:TYR:CG | 2.99 | 0.40 |
| 47:DV:2:PHE:HB3 | 47:DV:3:ALA:H | 1.26 | 0.40 |
| 47:DV:13:ARG:CG | 47:DV:13:ARG:O | 2.69 | 0.40 |
| 47:DV:56:SER:C | 47:DV:58:VAL:H | 2.29 | 0.40 |
| 47:DV:75:PHE:CD1 | 47:DV:75:PHE:N | 2.88 | 0.40 |
| 51:DZ:102:LEU:CD2 | 51:DZ:137:ILE:HB | 2.51 | 0.40 |
| 1:AA:189:G:O6 | 1:AA:189(L):G:C6 | 2.74 | 0.40 |
| 1:AA:390:C:H3' | 1:AA:390:C:H6 | 1.85 | 0.40 |
| 1:AA:633:G:H5' | 1:AA:634:C:OP2 | 2.21 | 0.40 |
| 1:AA:635:G:O2' | 1:AA:636:U:H5' | 2.21 | 0.40 |
| 1:AA:974:A:OP1 | 14:AN:31:ARG:HD3 | 2.21 | 0.40 |
| 1:AA:986:A:C6 | 1:AA:1220:G:N1 | 2.90 | 0.40 |
| 1:AA:1159:U:C5 | 1:AA:1182:G:N3 | 2.89 | 0.40 |
| 1:AA:1305:G:C5' | 21:AU:4:GLY:HA3 | 2.51 | 0.40 |
| 1:AA:1368:G:OP2 | 9:AI:112:LYS:CD | 2.70 | 0.40 |
| 4:AD:150:GLU:HG2 | 4:AD:151:LYS:N | 2.37 | 0.40 |
| 5:AE:78:HIS:CE1 | 5:AE:143:ARG:H | 2.39 | 0.40 |
| 5:AE:129:ILE:O | 5:AE:132:ALA:HB3 | 2.22 | 0.40 |
| 6:AF:48:LEU:HD21 | 6:AF:60:PHE:CZ | 2.57 | 0.40 |
| 8:AH:63:LEU:HB2 | 8:AH:65:TYR:CE1 | 2.56 | 0.40 |
| 9:AI:113:LYS:HD3 | 9:AI:119:ALA:O | 2.21 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 16:AP:58:TYR:O | 16:AP:61:SER:N | 2.55 | 0.40 |
| 23:B1:37:ILE:HD13 | 23:B1:37:ILE:HA | 1.71 | 0.40 |
| 23:B1:87:PRO:CG | 23:B1:88:LYS:N | 2.84 | 0.40 |
| 28:B6:24:GLU:HB3 | 28:B6:25:LYS:H | 1.57 | 0.40 |
| 30:B8:37:SER:HB2 | 30:B8:38:GLY:H | 1.41 | 0.40 |
| 31:BA:540:C:C2' | 31:BA:541:C:O5' | 2.69 | 0.40 |
| 31:BA:541:C:H2' | 31:BA:542:C:C5 | 2.56 | 0.40 |
| 31:BA:783:A:O2' | 31:BA:785:G:OP1 | 2.29 | 0.40 |
| 31:BA:945:A:C4 | 31:BA:2448:A:C2 | 3.09 | 0.40 |
| 31:BA:1027:A:N6 | 31:BA:1126:A:C4 | 2.89 | 0.40 |
| 31:BA:1112:G:N2 | 31:BA:1113:U:O2 | 2.54 | 0.40 |
| 31:BA:1140:C:OP1 | 39:BN:23:LEU:O | 2.38 | 0.40 |
| 31:BA:1149:G:H2' | 31:BA:1150:C:C6 | 2.56 | 0.40 |
| 31:BA:1459:G:C8 | 31:BA:1461:G:H1' | 2.56 | 0.40 |
| 31:BA:1475:G:C2 | 31:BA:1517:G:C2 | 3.08 | 0.40 |
| 31:BA:2475:C:H6 | 31:BA:2475:C:C5' | 2.28 | 0.40 |
| 31:BA:2582:G:C2 | 31:BA:2583:G:C8 | 3.09 | 0.40 |
| 31:BA:2663:G:C5 | 31:BA:2664:G:N7 | 2.89 | 0.40 |
| 31:BA:2691:C:O3' | 31:BA:2871:C:H4' | 2.21 | 0.40 |
| 31:BA:2747:G:O6 | 31:BA:2755:C:H5" | 2.21 | 0.40 |
| 33:BD:147:LEU:HD11 | 33:BD:183:ARG:NH1 | 2.37 | 0.40 |
| 34:BE:59:VAL:HG21 | 34:BE:63:LEU:HD23 | 2.03 | 0.40 |
| 34:BE:101:ARG:HA | 34:BE:101:ARG:HD3 | 1.80 | 0.40 |
| 34:BE:119:ARG:HG2 | 34:BE:160:TYR:HB2 | 2.02 | 0.40 |
| 36:BG:10:LYS:O | 36:BG:15:VAL:HG23 | 2.21 | 0.40 |
| 37:BH:41:MET:HB3 | 37:BH:43:VAL:HG22 | 2.03 | 0.40 |
| 39:BN:79:PRO:CG | 39:BN:80:GLY:N | 2.84 | 0.40 |
| 39:BN:128:HIS:O | 39:BN:129:PRO:C | 2.61 | 0.40 |
| 41:BP:110:TYR:CZ | 41:BP:111:ARG:HD2 | 2.55 | 0.40 |
| 42:BQ:52:VAL:O | 42:BQ:53:ALA:C | 2.65 | 0.40 |
| 43:BR:65:LEU:HD12 | 43:BR:65:LEU:HA | 1.68 | 0.40 |
| 44:BS:77:ALA:O | 44:BS:80:LEU:HD12 | 2.21 | 0.40 |
| 47:BV:94:LEU:HD23 | 47:BV:94:LEU:C | 2.45 | 0.40 |
| 51:BZ:108:PRO:O | 51:BZ:109:ALA:C | 2.64 | 0.40 |
| 1:CA:47:C:H5" | 1:CA:365:U:C6 | 2.55 | 0.40 |
| 1:CA:66:G:C2 | 1:CA:67:C:C5 | 3.09 | 0.40 |
| 1:CA:186:C:O2' | 1:CA:187:C:H5' | 2.22 | 0.40 |
| 1:CA:310:G:C5' | 16:CP:31:LYS:HB2 | 2.50 | 0.40 |
| 1:CA:310:G:H5" | 16:CP:31:LYS:HB2 | 2.03 | 0.40 |
| 1:CA:373:A:N3 | 1:CA:374:A:C8 | 2.90 | 0.40 |
| 1:CA:579:G:C5 | 1:CA:580:U:C4 | 3.10 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:CA:719:C:C5 | 1:CA:720:C:N3 | 2.90 | 0.40 |
| 1:CA:784:C:C4 | 1:CA:799:G:N1 | 2.89 | 0.40 |
| 1:CA:980:C:H5' | 1:CA:981:U:C5 | 2.56 | 0.40 |
| 1:CA:1098:C:C4 | 1:CA:1099:G:C8 | 3.08 | 0.40 |
| 1:CA:1173:G:H2' | 1:CA:1174:G:H8 | 1.86 | 0.40 |
| 1:CA:1291:G:O3' | 9:CI:39:GLY:HA3 | 2.21 | 0.40 |
| 1:CA:1316:G:N2 | 1:CA:1319:A:OP2 | 2.54 | 0.40 |
| 1:CA:1425:U:H2' | 1:CA:1426:C:H6 | 1.86 | 0.40 |
| 3:CC:27:LYS:HA | 3:CC:27:LYS:HZ3 | 1.86 | 0.40 |
| 3:CC:29:TYR:OH | 14:CN:54:PRO:HD2 | 2.21 | 0.40 |
| 4:CD:31:CYS:C | 4:CD:33:MET:N | 2.61 | 0.40 |
| 4:CD:101:LEU:HD13 | 4:CD:140:VAL:HG22 | 2.03 | 0.40 |
| 9:CI:49:PRO:O | 9:CI:53:VAL:HG13 | 2.22 | 0.40 |
| 10:CJ:32:ALA:HB1 | 10:CJ:75:ILE:CD1 | 2.50 | 0.40 |
| 15:CO:3:ILE:O | 15:CO:4:THR:C | 2.65 | 0.40 |
| 15:CO:51:HIS:O | 15:CO:54:ARG:HB3 | 2.21 | 0.40 |
| 15:CO:55:GLY:O | 15:CO:56:LEU:C | 2.65 | 0.40 |
| 17:CQ:18:THR:CG2 | 17:CQ:69:LYS:HE3 | 2.42 | 0.40 |
| 24:D2:14:ARG:HE | 24:D2:14:ARG:HB3 | 1.67 | 0.40 |
| 24:D2:50:ILE:O | 24:D2:51:ARG:CB | 2.69 | 0.40 |
| 24:D2:57:ILE:HG13 | 24:D2:58:ALA:O | 2.20 | 0.40 |
| 25:D3:8:LEU:HD12 | 25:D3:31:LEU:HA | 2.00 | 0.40 |
| 27:D5:46:CYS:SG | 27:D5:47:PRO:CG | 3.09 | 0.40 |
| 28:D6:16:CYS:O | 28:D6:18:ARG:NH1 | 2.54 | 0.40 |
| 31:DA:45:C:H2' | 31:DA:47:C:C6 | 2.56 | 0.40 |
| 31:DA:118:A:N3 | 31:DA:178:G:H1' | 2.36 | 0.40 |
| 31:DA:485:C:H2' | 31:DA:486:C:C6 | 2.56 | 0.40 |
| 31:DA:874:G:O2' | 51:DZ:120:ILE:HD12 | 2.21 | 0.40 |
| 31:DA:1368:G:C2 | 31:DA:1369:G:C8 | 3.09 | 0.40 |
| 31:DA:1646:C:H5" | 31:DA:1647:G:C5' | 2.51 | 0.40 |
| 31:DA:1658:C:OP1 | 34:DE:132:HIS:O | 2.38 | 0.40 |
| 31:DA:1833:U:O2 | 31:DA:1969:A:H2 | 2.04 | 0.40 |
| 31:DA:2248:C:O2 | 31:DA:2248:C:H2' | 2.21 | 0.40 |
| 31:DA:2575:C:O2' | 31:DA:2578:G:N7 | 2.52 | 0.40 |
| 31:DA:2715:C:O2' | 31:DA:2716:U:H5' | 2.22 | 0.40 |
| 31:DA:2781:A:C5' | 31:DA:2782:G:H5' | 2.49 | 0.40 |
| 32:DB:42:C:O2 | 36:DG:92:VAL:HA | 2.21 | 0.40 |
| 34:DE:3:GLY:HA3 | 34:DE:81:ILE:HG21 | 2.02 | 0.40 |
| 34:DE:97:LYS:O | 34:DE:100:GLU:HG3 | 2.21 | 0.40 |
| 35:DF:3:GLU:CD | 35:DF:21:ALA:H | 2.28 | 0.40 |
| 35:DF:164:ARG:HH11 | 35:DF:164:ARG:CG | 2.28 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 35:DF:183:VAL:O | 35:DF:184:TYR:C | 2.64 | 0.40 |
| 35:DF:203:GLN:CA | 35:DF:206:ILE:O | 2.66 | 0.40 |
| 37:DH:43:VAL:CG2 | 37:DH:52:VAL:HG13 | 2.39 | 0.40 |
| 38:DI:49:ALA:HA | 38:DI:52:ARG:HB2 | 2.02 | 0.40 |
| 42:DQ:54:MET:SD | 42:DQ:118:LEU:HD23 | 2.62 | 0.40 |
| 44:DS:30:ARG:HD2 | 44:DS:31:SER:O | 2.20 | 0.40 |
| 45:DT:30:VAL:HG21 | 45:DT:84:GLN:H | 1.86 | 0.40 |
| 47:DV:13:ARG:HG3 | 47:DV:13:ARG:O | 2.20 | 0.40 |
| 47:DV:62:LEU:CB | 47:DV:98:GLU:HB2 | 2.50 | 0.40 |
| 48:DW:40:ASN:O | 48:DW:41:LYS:HD3 | 2.20 | 0.40 |
| 49:DX:8:ILE:HD12 | 49:DX:8:ILE:N | 2.35 | 0.40 |
| 50:DY:88:LYS:HZ2 | 50:DY:93:GLY:HA3 | 1.86 | 0.40 |
| 51:DZ:9:TYR:HE2 | 51:DZ:61:LEU:HD13 | 1.85 | 0.40 |
| 51:DZ:48:PHE:O | 51:DZ:49:ARG:C | 2.63 | 0.40 |
| 1:AA:103:C:O2 | 1:AA:172:A:H2 | 2.03 | 0.40 |
| 1:AA:137:C:N4 | 1:AA:226:G:H1 | 2.20 | 0.40 |
| 1:AA:148:G:H2' | 1:AA:149:A:H8 | 1.85 | 0.40 |
| 1:AA:473:G:C2 | 1:AA:474:G:N7 | 2.90 | 0.40 |
| 1:AA:509:A:H3' | 1:AA:509:A:OP2 | 2.21 | 0.40 |
| 1:AA:519:C:C2' | 1:AA:520:A:O5' | 2.69 | 0.40 |
| 1:AA:631:G:H5" | 1:AA:632:A:OP1 | 2.21 | 0.40 |
| 1:AA:929:G:C6 | 1:AA:930:C:C4 | 3.09 | 0.40 |
| 1:AA:1250:A:C6 | 1:AA:1251:A:C6 | 3.10 | 0.40 |
| 3:AC:11:ARG:HE | 3:AC:180:ALA:HB3 | 1.87 | 0.40 |
| 5:AE:129:ILE:O | 5:AE:133:TYR:HD1 | 2.04 | 0.40 |
| 7:AG:69:VAL:HG13 | 7:AG:134:ALA:O | 2.21 | 0.40 |
| 7:AG:92:SER:CB | 7:AG:94:ARG:HH21 | 2.34 | 0.40 |
| 12:AL:62:SER:C | 12:AL:64:TYR:H | 2.30 | 0.40 |
| 13:AM:90:LEU:O | 13:AM:91:ARG:HB2 | 2.21 | 0.40 |
| 16:AP:22:THR:HB | 16:AP:23:ASP:H | 1.80 | 0.40 |
| 17:AQ:46:ASP:C | 17:AQ:48:GLU:H | 2.29 | 0.40 |
| 19:AS:9:VAL:O | 19:AS:9:VAL:HG12 | 2.21 | 0.40 |
| 19:AS:62:ILE:O | 19:AS:62:ILE:HG23 | 2.21 | 0.40 |
| 22:B0:20:ARG:HG3 | 31:BA:2356:C:H4' | 2.03 | 0.40 |
| 23:B1:40:ARG:NH2 | 31:BA:2082:A:H5' | 2.37 | 0.40 |
| 23:B1:48:LYS:HB3 | 23:B1:49:VAL:H | 1.67 | 0.40 |
| 24:B2:12:GLU:CD | 24:B2:13:ALA:N | 2.80 | 0.40 |
| 24:B2:30:ARG:HH11 | 24:B2:30:ARG:CG | 2.33 | 0.40 |
| 27:B5:46:CYS:SG | 27:B5:47:PRO:HG2 | 2.60 | 0.40 |
| 30:B8:8:LYS:O | 30:B8:12:LYS:HG3 | 2.21 | 0.40 |
| 30:B8:39:LYS:O | 30:B8:39:LYS:HE3 | 2.22 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:203:C:H2' | 31:BA:204:A:OP1 | 2.21 | 0.40 |
| 31:BA:569:U:C4 | 31:BA:570:G:C6 | 3.09 | 0.40 |
| 31:BA:1301:A:C8 | 31:BA:1303:G:C8 | 3.09 | 0.40 |
| 31:BA:1321:A:N6 | 31:BA:1322:A:C6 | 2.90 | 0.40 |
| 31:BA:1357:U:H2' | 31:BA:1358:G:O4' | 2.21 | 0.40 |
| 31:BA:1359:A:N7 | 31:BA:1372:U:C4 | 2.88 | 0.40 |
| 31:BA:1550:C:O2' | 31:BA:1551:C:H5' | 2.21 | 0.40 |
| 31:BA:1710:C:O2' | 31:BA:1711:C:H5' | 2.21 | 0.40 |
| 31:BA:1826:G:H2' | 31:BA:1827:C:H6 | 1.87 | 0.40 |
| 31:BA:2063:C:C4 | 31:BA:2064:C:C4 | 3.09 | 0.40 |
| 31:BA:2475:C:H42 | 31:BA:2529:G:N2 | 2.07 | 0.40 |
| 31:BA:2526:G:C6 | 31:BA:2527:C:C4 | 3.09 | 0.40 |
| 31:BA:2645:G:C3' | 31:BA:2646:C:H5' | 2.51 | 0.40 |
| 31:BA:2653:U:C3' | 31:BA:2654:A:H5" | 2.42 | 0.40 |
| 31:BA:2726:U:O2' | 31:BA:2727:G:H8 | 2.05 | 0.40 |
| 32:BB:92:C:H5" | 51:BZ:79:ARG:HH22 | 1.86 | 0.40 |
| 36:BG:52:ILE:HG22 | 36:BG:54:GLU:CG | 2.51 | 0.40 |
| 37:BH:30:LYS:NZ | 37:BH:81:GLU:CA | 2.83 | 0.40 |
| 40:BO:23:ARG:NH1 | 40:BO:23:ARG:CG | 2.67 | 0.40 |
| 41:BP:71:VAL:CG1 | 41:BP:72:PRO:CD | 2.98 | 0.40 |
| 42:BQ:78:PRO:O | 42:BQ:79:LEU:CG | 2.70 | 0.40 |
| 42:BQ:88:GLY:O | 42:BQ:89:ASN:CB | 2.66 | 0.40 |
| 43:BR:18:LEU:C | 43:BR:18:LEU:HD13 | 2.46 | 0.40 |
| 47:BV:6:LYS:HA | 47:BV:11:GLN:HA | 2.04 | 0.40 |
| 48:BW:36:LEU:HD13 | 48:BW:48:ALA:CA | 2.51 | 0.40 |
| 49:BX:26:TYR:OH | 49:BX:89:ILE:CG2 | 2.69 | 0.40 |
| 51:BZ:42:VAL:HG13 | 51:BZ:43:GLU:H | 1.87 | 0.40 |
| 51:BZ:98:MET:HE2 | 51:BZ:100:VAL:CG2 | 2.51 | 0.40 |
| 1:CA:63:C:H5' | 1:CA:64:G:OP2 | 2.21 | 0.40 |
| 1:CA:101:A:C6 | 1:CA:102:G:N7 | 2.90 | 0.40 |
| 1:CA:149:A:O2' | 1:CA:150:C:C6 | 2.68 | 0.40 |
| 1:CA:428:G:H5" | 4:CD:7:PRO:HB3 | 2.03 | 0.40 |
| 1:CA:689:C:O2 | 1:CA:689:C:H2' | 2.21 | 0.40 |
| 1:CA:720:C:H5" | 1:CA:721:G:O5' | 2.21 | 0.40 |
| 1:CA:862:C:C2' | 1:CA:863:U:H5' | 2.51 | 0.40 |
| 1:CA:965:A:C2 | 1:CA:969:A:N1 | 2.90 | 0.40 |
| 1:CA:1095:U:P | 1:CA:1108:G:H1 | 2.43 | 0.40 |
| 1:CA:1126:U:P | 1:CA:1126:U:H6 | 2.45 | 0.40 |
| 4:CD:68:TYR:O | 4:CD:69:GLY:C | 2.64 | 0.40 |
| 4:CD:104:VAL:HG21 | 4:CD:185:PHE:CD1 | 2.56 | 0.40 |
| 5:CE:78:HIS:CE1 | 5:CE:143:ARG:H | 2.40 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 8:CH:51:VAL:HB | 8:CH:52:ASP:H | 1.52 | 0.40 |
| 8:CH:87:SER:HA | 8:CH:93:VAL:CG2 | 2.50 | 0.40 |
| 8:CH:88:LYS:HB3 | 8:CH:89:PRO:HD2 | 2.03 | 0.40 |
| 9:CI:46:ALA:HA | 9:CI:78:LYS:HE3 | 2.02 | 0.40 |
| 11:CK:61:ALA:CB | 11:CK:90:GLY:HA3 | 2.51 | 0.40 |
| 13:CM:44:ARG:HB3 | 13:CM:46:LYS:HG2 | 2.04 | 0.40 |
| 15:CO:39:LEU:O | 15:CO:40:SER:C | 2.65 | 0.40 |
| 18:CR:36:ASN:ND2 | 18:CR:39:VAL:HG21 | 2.35 | 0.40 |
| 18:CR:50:ILE:HG12 | 18:CR:74:ARG:HH12 | 1.86 | 0.40 |
| 23:D1:80:LEU:HD23 | 23:D1:80:LEU:HA | 1.91 | 0.40 |
| 24:D2:33:MET:CG | 49:DX:11:PRO:HD2 | 2.52 | 0.40 |
| 25:D3:1:MET:O | 25:D3:3:ARG:N | 2.55 | 0.40 |
| 28:D6:46:HIS:ND1 | 28:D6:46:HIS:O | 2.55 | 0.40 |
| 30:D8:39:LYS:HE3 | 30:D8:42:ARG:HH12 | 1.86 | 0.40 |
| 31:DA:118:A:O5' | 31:DA:119:A:H5" | 2.20 | 0.40 |
| 31:DA:322:A:C5 | 31:DA:340:A:C2 | 3.10 | 0.40 |
| 31:DA:343:C:H2' | 31:DA:344:G:H5' | 2.02 | 0.40 |
| 31:DA:728:G:H4' | 33:DD:13:ARG:HH11 | 1.86 | 0.40 |
| 31:DA:755:C:H2' | 31:DA:756:C:H6 | 1.86 | 0.40 |
| 31:DA:1147:C:H2' | 31:DA:1148:A:O5' | 2.21 | 0.40 |
| 31:DA:1207:C:H2' | 31:DA:1208:C:C6 | 2.55 | 0.40 |
| 31:DA:1271:G:O2' | 31:DA:1618:A:OP1 | 2.27 | 0.40 |
| 31:DA:1470:G:H5" | 31:DA:1471:A:OP1 | 2.21 | 0.40 |
| 31:DA:1473:G:H2' | 31:DA:1474:C:O4' | 2.21 | 0.40 |
| 31:DA:1909:C:O2 | 31:DA:1909:C:H2' | 2.21 | 0.40 |
| 31:DA:1914:C:H2' | 31:DA:1915:U:O4' | 2.21 | 0.40 |
| 31:DA:2442:C:H2' | 31:DA:2443:C:H6 | 1.86 | 0.40 |
| 31:DA:2694:G:C6 | 31:DA:2695:C:C4 | 3.09 | 0.40 |
| 34:DE:23:VAL:HA | 34:DE:186:GLY:N | 2.34 | 0.40 |
| 36:DG:161:THR:HG21 | 36:DG:163:ALA:HB3 | 2.03 | 0.40 |
| 36:DG:181:ARG:HE | 36:DG:181:ARG:HB2 | 1.66 | 0.40 |
| 37:DH:148:ILE:O | 37:DH:162:ILE:HD11 | 2.22 | 0.40 |
| 37:DH:157:TYR:CD1 | 37:DH:170:ARG:O | 2.74 | 0.40 |
| 39:DN:99:LEU:HA | 39:DN:99:LEU:HD23 | 1.56 | 0.40 |
| 42:DQ:31:ASP:O | 42:DQ:133:ARG:O | 2.39 | 0.40 |
| 42:DQ:50:ALA:O | 42:DQ:53:ALA:HB3 | 2.22 | 0.40 |
| 44:DS:20:ARG:HA | 44:DS:20:ARG:HD3 | 1.52 | 0.40 |
| 44:DS:24:LEU:HB3 | 44:DS:85:VAL:HG13 | 2.02 | 0.40 |
| 44:DS:99:LYS:N | 44:DS:99:LYS:HD3 | 2.36 | 0.40 |
| 45:DT:11:GLU:OE2 | 45:DT:11:GLU:N | 2.54 | 0.40 |
| 45:DT:106:SER:O | 45:DT:107:ASP:OD1 | 2.39 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 47:DV:73:SER:OG | 47:DV:75:PHE:CE1 | 2.75 | 0.40 |
| 48:DW:24:ILE:HD12 | 48:DW:24:ILE:C | 2.47 | 0.40 |
| 51:DZ:98:MET:HE2 | 51:DZ:100:VAL:CG2 | 2.52 | 0.40 |
| 51:DZ:166:SER:HB2 | 51:DZ:167:PRO:CA | 2.51 | 0.40 |
| 1:AA:224:C:C2 | 1:AA:225:C:C5 | 3.10 | 0.40 |
| 1:AA:428:G:OP2 | 4:AD:7:PRO:HG2 | 2.22 | 0.40 |
| 1:AA:575:G:C5 | 1:AA:881:G:C2 | 3.10 | 0.40 |
| 1:AA:667:G:H4' | 15:AO:51:HIS:ND1 | 2.36 | 0.40 |
| 1:AA:872:A:C2 | 1:AA:874:G:C5 | 3.09 | 0.40 |
| 1:AA:952:U:H2' | 1:AA:953:G:C8 | 2.51 | 0.40 |
| 1:AA:964:A:N3 | 1:AA:969:A:O2' | 2.41 | 0.40 |
| 1:AA:1244:C:C2 | 1:AA:1294:G:N2 | 2.90 | 0.40 |
| 3:AC:3:ASN:O | 3:AC:4:LYS:O | 2.39 | 0.40 |
| 4:AD:163:GLU:C | 4:AD:165:MET:H | 2.28 | 0.40 |
| 5:AE:48:ALA:HB1 | 5:AE:49:PRO:CD | 2.51 | 0.40 |
| 6:AF:24:GLU:HG3 | 6:AF:28:ARG:CZ | 2.50 | 0.40 |
| 7:AG:42:ILE:HD13 | 7:AG:116:ALA:CB | 2.52 | 0.40 |
| 9:AI:46:ALA:HA | 9:AI:78:LYS:HE3 | 2.03 | 0.40 |
| 10:AJ:22:LYS:HD2 | 10:AJ:22:LYS:C | 2.46 | 0.40 |
| 10:AJ:48:THR:HA | 10:AJ:62:HIS:CB | 2.51 | 0.40 |
| 19:AS:12:ASP:HB2 | 19:AS:15:LEU:HD23 | 2.02 | 0.40 |
| 23:B1:73:LEU:HD13 | 23:B1:90:ILE:CG2 | 2.50 | 0.40 |
| 24:B2:50:ILE:O | 24:B2:51:ARG:CB | 2.70 | 0.40 |
| 29:B7:18:PHE:CD2 | 29:B7:18:PHE:C | 2.99 | 0.40 |
| 31:BA:196:A:H2' | 31:BA:196:A:N3 | 2.37 | 0.40 |
| 31:BA:363(E):U:H2' | 31:BA:363(F):A:O4' | 2.22 | 0.40 |
| 31:BA:628:G:C6 | 31:BA:629:G:C6 | 3.09 | 0.40 |
| 31:BA:695:G:C2 | 31:BA:768:G:C4 | 3.10 | 0.40 |
| 31:BA:705:A:C2' | 31:BA:706:A:H5' | 2.51 | 0.40 |
| 31:BA:1106:A:H2' | 31:BA:1107:G:C8 | 2.57 | 0.40 |
| 31:BA:1132:A:H1' | 39:BN:73:THR:HG21 | 2.03 | 0.40 |
| 31:BA:1177:A:H3' | 31:BA:1177:A:OP2 | 2.21 | 0.40 |
| 31:BA:1207:C:H2' | 31:BA:1208:C:H6 | 1.86 | 0.40 |
| 31:BA:1354:A:H2' | 31:BA:1355:G:O4' | 2.22 | 0.40 |
| 31:BA:1531:C:H3' | 31:BA:1532:C:O4' | 2.21 | 0.40 |
| 31:BA:1564:C:O2' | 31:BA:1565:C:H5' | 2.20 | 0.40 |
| 31:BA:1891:G:C6 | 31:BA:1892:C:C4 | 3.09 | 0.40 |
| 31:BA:1909:C:H2' | 31:BA:1909:C:O2 | 2.20 | 0.40 |
| 31:BA:2091:U:OP2 | 31:BA:2092:U:O2' | 2.37 | 0.40 |
| 31:BA:2197:U:O2' | 31:BA:2198:A:H2' | 2.21 | 0.40 |
| 31:BA:2641:G:H8 | 31:BA:2641:G:H5" | 1.86 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:BA:2772:C:H2' | 31:BA:2773:C:C6 | 2.55 | 0.40 |
| 33:BD:45:ASN:OD1 | 33:BD:46:GLN:N | 2.54 | 0.40 |
| 34:BE:146:THR:HA | 34:BE:147:PRO:C | 2.46 | 0.40 |
| 35:BF:22:ALA:CB | 35:BF:26:ALA:HB1 | 2.48 | 0.40 |
| 35:BF:117:ARG:HD3 | 35:BF:117:ARG:HA | 1.74 | 0.40 |
| 37:BH:56:SER:OG | 37:BH:61:HIS:ND1 | 2.53 | 0.40 |
| 39:BN:21:LYS:O | 39:BN:61:ARG:N | 2.52 | 0.40 |
| 39:BN:78:TYR:CD1 | 39:BN:79:PRO:CD | 2.89 | 0.40 |
| 40:BO:118:ALA:HA | 40:BO:119:PRO:HD2 | 1.66 | 0.40 |
| 40:BO:121:VAL:O | 40:BO:121:VAL:HG12 | 2.20 | 0.40 |
| 41:BP:58:THR:O | 41:BP:58:THR:HG22 | 2.21 | 0.40 |
| 42:BQ:16:ARG:CG | 42:BQ:17:LEU:N | 2.84 | 0.40 |
| 42:BQ:103:MET:HB2 | 42:BQ:104:PHE:CD1 | 2.56 | 0.40 |
| 43:BR:12:ARG:NE | 43:BR:20:LEU:HD22 | 2.36 | 0.40 |
| 43:BR:56:LYS:HE3 | 43:BR:94:TYR:OH | 2.21 | 0.40 |
| 45:BT:89:VAL:CG1 | 45:BT:91:ARG:HG2 | 2.51 | 0.40 |
| 47:BV:17:GLY:O | 47:BV:18:LEU:HB3 | 2.22 | 0.40 |
| 47:BV:86:GLY:O | 47:BV:87:HIS:CG | 2.75 | 0.40 |
| 50:BY:50:ARG:HB3 | 50:BY:51:VAL:H | 1.58 | 0.40 |
| 50:BY:80:GLY:O | 50:BY:81:LYS:CB | 2.69 | 0.40 |
| 51:BZ:10:ARG:NH2 | 51:BZ:26:GLY:O | 2.54 | 0.40 |
| 1:CA:166:G:O2' | 1:CA:167:G:H5' | 2.21 | 0.40 |
| 1:CA:410:G:N2 | 1:CA:431:A:C8 | 2.90 | 0.40 |
| 1:CA:577:G:C1' | 1:CA:816:A:C4 | 3.04 | 0.40 |
| 1:CA:965:A:C2 | 1:CA:969:A:C2 | 3.10 | 0.40 |
| 1:CA:1250:A:C6 | 1:CA:1251:A:C6 | 3.10 | 0.40 |
| 1:CA:1255:G:O2' | 1:CA:1258:G:O2' | 2.33 | 0.40 |
| 1:CA:1281:U:C5' | 1:CA:1282:C:H5 | 2.34 | 0.40 |
| 2:CB:238:LEU:O | 2:CB:240:GLN:N | 2.54 | 0.40 |
| 3:CC:19:GLU:O | 3:CC:40:ARG:NH2 | 2.54 | 0.40 |
| 3:CC:134:ILE:CG2 | 3:CC:151:VAL:HB | 2.48 | 0.40 |
| 4:CD:3:ARG:HD3 | 4:CD:5:ILE:HG13 | 2.02 | 0.40 |
| 4:CD:68:TYR:CD2 | 4:CD:97:LEU:HD22 | 2.57 | 0.40 |
| 4:CD:79:PHE:CG | 4:CD:207:TYR:HD1 | 2.39 | 0.40 |
| 4:CD:127:THR:HA | 4:CD:132:ARG:HA | 2.03 | 0.40 |
| 5:CE:69:VAL:HG12 | 5:CE:71:LEU:HD21 | 2.03 | 0.40 |
| 9:CI:56:LEU:HD23 | 9:CI:56:LEU:C | 2.47 | 0.40 |
| 14:CN:51:GLY:C | 14:CN:53:LEU:H | 2.29 | 0.40 |
| 24:D2:24:LEU:HA | 24:D2:24:LEU:HD12 | 1.85 | 0.40 |
| 24:D2:29:LYS:O | 24:D2:30:ARG:C | 2.65 | 0.40 |
| 30:D8:58:ILE:HG22 | 41:DP:49:ARG:CD | 2.48 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:50:U:OP2 | 31:DA:50:U:H4' | 2.21 | 0.40 |
| 31:DA:532:A:N7 | 31:DA:2021:C:O2' | 2.54 | 0.40 |
| 31:DA:540:C:C2' | 31:DA:541:C:O5' | 2.69 | 0.40 |
| 31:DA:585:G:C5 | 31:DA:1251:C:C4 | 3.08 | 0.40 |
| 31:DA:598:G:H5' | 41:DP:15:ARG:HB2 | 2.04 | 0.40 |
| 31:DA:646:A:H5' | 31:DA:646:A:N3 | 2.37 | 0.40 |
| 31:DA:814:C:C5 | 41:DP:27:HIS:CD2 | 3.10 | 0.40 |
| 31:DA:945:A:C6 | 31:DA:2448:A:C4 | 3.10 | 0.40 |
| 31:DA:1301:A:C4 | 31:DA:1303:G:N7 | 2.89 | 0.40 |
| 31:DA:1531:C:C3' | 31:DA:1532:C:C5' | 2.99 | 0.40 |
| 31:DA:1632:A:N6 | 31:DA:1633:G:N1 | 2.69 | 0.40 |
| 31:DA:1932:A:H2' | 31:DA:1933:G:H5' | 2.04 | 0.40 |
| 31:DA:1992:G:O5' | 31:DA:1992:G:C8 | 2.74 | 0.40 |
| 31:DA:2494:G:O2' | 31:DA:2495:G:H5' | 2.22 | 0.40 |
| 31:DA:2586:C:C5 | 31:DA:2608:G:N2 | 2.89 | 0.40 |
| 31:DA:2637:U:C2' | 31:DA:2638:G:H5' | 2.51 | 0.40 |
| 31:DA:2751:G:H3' | 31:DA:2752:C:C6 | 2.55 | 0.40 |
| 32:DB:52:A:O2' | 32:DB:53:A:H8 | 2.02 | 0.40 |
| 32:DB:79:C:H2' | 32:DB:80:U:O4' | 2.22 | 0.40 |
| 33:DD:105:ILE:HD12 | 33:DD:106:ILE:N | 2.37 | 0.40 |
| 33:DD:244:ARG:HA | 33:DD:245:PRO:HA | 1.77 | 0.40 |
| 34:DE:92:THR:O | 34:DE:93:VAL:HB | 2.21 | 0.40 |
| 35:DF:101:LEU:O | 35:DF:106:ARG:HD3 | 2.22 | 0.40 |
| 36:DG:19:LEU:O | 36:DG:20:ILE:C | 2.65 | 0.40 |
| 36:DG:47:LYS:HG3 | 36:DG:82:LEU:CD1 | 2.47 | 0.40 |
| 36:DG:120:LEU:HD12 | 36:DG:179:PRO:HG2 | 2.02 | 0.40 |
| 36:DG:173:LEU:HD13 | 36:DG:178:PHE:CZ | 2.57 | 0.40 |
| 37:DH:71:LEU:HA | 37:DH:71:LEU:HD12 | 1.86 | 0.40 |
| 37:DH:103:LEU:HG | 37:DH:105:LEU:HD13 | 2.03 | 0.40 |
| 37:DH:131:VAL:HG12 | 37:DH:132:ARG:N | 2.37 | 0.40 |
| 38:DI:33:ARG:O | 38:DI:35:LEU:CD2 | 2.69 | 0.40 |
| 38:DI:69:LYS:HG3 | 38:DI:136:VAL:HB | 2.04 | 0.40 |
| 40:DO:2:ILE:N | 40:DO:2:ILE:HD13 | 2.36 | 0.40 |
| 40:DO:22:ILE:HD13 | 40:DO:22:ILE:HA | 1.60 | 0.40 |
| 44:DS:15:ARG:C | 44:DS:17:ARG:O | 2.65 | 0.40 |
| 47:DV:27:ALA:O | 47:DV:28:GLU:C | 2.64 | 0.40 |
| 50:DY:77:PRO:O | 50:DY:78:ALA:CB | 2.67 | 0.40 |
| 1:AA:66:G:C2 | 1:AA:67:C:C5 | 3.09 | 0.40 |
| 1:AA:66:G:C5 | 1:AA:67:C:C5 | 3.09 | 0.40 |
| 1:AA:109:A:C5 | 1:AA:326:G:C2 | 3.10 | 0.40 |
| 1:AA:174:C:C5 | 1:AA:175:C:C5 | 3.10 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 1:AA:292:G:N3 | 1:AA:309:G:C2 | 2.89 | 0.40 |
| 1:AA:329:A:C2 | 1:AA:332:G:C8 | 3.10 | 0.40 |
| 1:AA:408:A:C6 | 1:AA:409:G:C5 | 3.09 | 0.40 |
| 1:AA:516:U:C5 | 1:AA:517:G:C6 | 3.09 | 0.40 |
| 1:AA:725:G:H2' | 1:AA:726:C:H6 | 1.87 | 0.40 |
| 1:AA:737:A:C4 | 1:AA:738:C:C5 | 3.10 | 0.40 |
| 1:AA:801:U:H2' | 1:AA:802:A:C8 | 2.54 | 0.40 |
| 1:AA:1277:C:H3' | 1:AA:1277:C:H6 | 1.87 | 0.40 |
| 1:AA:1352:C:O2 | 1:AA:1371:G:C2 | 2.74 | 0.40 |
| 1:AA:1379:G:O2' | 1:AA:1380:U:H5' | 2.22 | 0.40 |
| 1:AA:1421:G:C4 | 1:AA:1480:G:N2 | 2.89 | 0.40 |
| 2:AB:58:ILE:HD13 | 2:AB:58:ILE:HA | 1.82 | 0.40 |
| 4:AD:54:TYR:O | 4:AD:55:ALA:C | 2.65 | 0.40 |
| 4:AD:170:VAL:HG22 | 4:AD:171:GLY:H | 1.86 | 0.40 |
| 4:AD:181:MET:HE2 | 4:AD:181:MET:HB3 | 1.97 | 0.40 |
| 5:AE:120:THR:O | 5:AE:121:LYS:HB2 | 2.20 | 0.40 |
| 7:AG:45:ASP:O | 7:AG:49:ILE:HG12 | 2.22 | 0.40 |
| 8:AH:100:ILE:HA | 8:AH:101:PRO:HD3 | 1.80 | 0.40 |
| 10:AJ:70:ARG:HG3 | 10:AJ:70:ARG:NH1 | 2.35 | 0.40 |
| 19:AS:56:GLN:HB2 | 19:AS:57:HIS:H | 1.71 | 0.40 |
| 20:AT:84:LEU:O | 20:AT:88:VAL:HG23 | 2.20 | 0.40 |
| 22:B0:16:SER:CB | 31:BA:2262:U:OP2 | 2.70 | 0.40 |
| 23:B1:47:GLN:OE1 | 23:B1:47:GLN:C | 2.64 | 0.40 |
| 27:B5:20:ARG:HA | 27:B5:23:HIS:CD2 | 2.57 | 0.40 |
| 30:B8:32:LEU:CB | 30:B8:34:TRP:HB3 | 2.49 | 0.40 |
| 31:BA:128:C:O2' | 31:BA:129:C:O5' | 2.37 | 0.40 |
| 31:BA:196:A:C4 | 31:BA:805:G:C6 | 3.09 | 0.40 |
| 31:BA:648:G:C2' | 31:BA:649:G:H5' | 2.52 | 0.40 |
| 31:BA:709:U:H3 | 31:BA:722:A:H61 | 1.70 | 0.40 |
| 31:BA:855:G:C4 | 31:BA:856:C:C5 | 3.10 | 0.40 |
| 31:BA:1052:C:C6 | 31:BA:1052:C:C3' | 3.04 | 0.40 |
| 31:BA:2598:A:H3' | 33:BD:236:GLY:HA2 | 2.03 | 0.40 |
| 31:BA:2689:U:H4' | 31:BA:2690:C:H5' | 2.02 | 0.40 |
| 31:BA:2723:C:OP2 | 34:BE:109:LYS:NZ | 2.55 | 0.40 |
| 33:BD:35:LYS:CG | 33:BD:64:ILE:H | 2.29 | 0.40 |
| 33:BD:35:LYS:HZ2 | 33:BD:65:ILE:HA | 1.78 | 0.40 |
| 34:BE:24:THR:HG23 | 34:BE:184:VAL:CG2 | 2.46 | 0.40 |
| 34:BE:37:ARG:HD3 | 34:BE:44:TYR:OH | 2.21 | 0.40 |
| 35:BF:34:TRP:HB2 | 41:BP:10:PRO:O | 2.22 | 0.40 |
| 35:BF:196:LEU:H | 35:BF:196:LEU:HG | 1.60 | 0.40 |
| 36:BG:45:GLU:HB2 | 36:BG:47:LYS:CG | 2.51 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 37:BH:90:LYS:HD3 | 37:BH:90:LYS:HA | 1.91 | 0.40 |
| 37:BH:157:TYR:HD1 | 37:BH:170:ARG:O | 2.05 | 0.40 |
| 39:BN:65:LYS:O | 39:BN:69:GLN:CG | 2.69 | 0.40 |
| 40:BO:1:MET:HB2 | 40:BO:32:TYR:HD2 | 1.87 | 0.40 |
| 40:BO:26:LYS:HB3 | 40:BO:27:GLY:H | 1.65 | 0.40 |
| 41:BP:79:ARG:NH2 | 41:BP:109:GLY:HA2 | 2.34 | 0.40 |
| 44:BS:28:VAL:O | 44:BS:29:PHE:HB3 | 2.21 | 0.40 |
| 46:BU:88:ILE:C | 46:BU:88:ILE:CD1 | 2.94 | 0.40 |
| 47:BV:62:LEU:HB3 | 47:BV:98:GLU:CA | 2.50 | 0.40 |
| 50:BY:6:HIS:HB3 | 50:BY:35:TYR:CE1 | 2.57 | 0.40 |
| 50:BY:7:VAL:CB | 50:BY:8:LYS:HD2 | 2.46 | 0.40 |
| 1:CA:246:A:C2 | 1:CA:282:A:C5 | 3.10 | 0.40 |
| 1:CA:293:G:C5 | 1:CA:294:U:C4 | 3.10 | 0.40 |
| 1:CA:308:C:H2' | 1:CA:309:G:C8 | 2.54 | 0.40 |
| 1:CA:623:C:C2' | 1:CA:624:C:H5' | 2.52 | 0.40 |
| 1:CA:776:G:O2' | 1:CA:777:A:P | 2.80 | 0.40 |
| 1:CA:953:G:C6 | 1:CA:1229:A:C6 | 3.09 | 0.40 |
| 1:CA:1054:C:OP1 | 1:CA:1197:G:OP2 | 2.39 | 0.40 |
| 1:CA:1117:G:O6 | 1:CA:1156:G:N2 | 2.52 | 0.40 |
| 1:CA:1233:G:P | 9:CI:124:GLN:HB2 | 2.62 | 0.40 |
| 1:CA:1366:C:H2' | 1:CA:1367:C:H6 | 1.87 | 0.40 |
| 1:CA:1468:A:H8 | 1:CA:1468:A:O5' | 2.05 | 0.40 |
| 2:CB:139:LYS:O | 2:CB:142:LEU:HB3 | 2.22 | 0.40 |
| 4:CD:101:LEU:O | 4:CD:103:ASN:N | 2.55 | 0.40 |
| 5:CE:75:THR:HG23 | 5:CE:76:ILE:N | 2.35 | 0.40 |
| 6:CF:82:ARG:HH11 | 6:CF:82:ARG:CA | 2.33 | 0.40 |
| 8:CH:20:TYR:HA | 8:CH:65:TYR:HE2 | 1.81 | 0.40 |
| 9:CI:85:LEU:HD12 | 9:CI:85:LEU:C | 2.47 | 0.40 |
| 10:CJ:8:LEU:O | 10:CJ:16:LEU:HD21 | 2.21 | 0.40 |
| 12:CL:52:LEU:HD23 | 12:CL:52:LEU:HA | 1.96 | 0.40 |
| 16:CP:20:VAL:HG22 | 16:CP:32:TYR:HB2 | 2.04 | 0.40 |
| 16:CP:43:LYS:HG3 | 16:CP:48:TRP:CE3 | 2.56 | 0.40 |
| 23:D1:37:ILE:HG13 | 31:DA:2079:U:O3' | 2.21 | 0.40 |
| 31:DA:524:U:H2' | 31:DA:525:U:C6 | 2.56 | 0.40 |
| 31:DA:530:G:O6 | 31:DA:2023:G:OP1 | 2.38 | 0.40 |
| 31:DA:547:A:C8 | 31:DA:549:G:C6 | 3.05 | 0.40 |
| 31:DA:635:C:O2' | 31:DA:639:U:OP1 | 2.38 | 0.40 |
| 31:DA:675:A:C8 | 31:DA:804:A:C6 | 3.10 | 0.40 |
| 31:DA:778:G:C6 | 31:DA:779:U:C4 | 3.09 | 0.40 |
| 31:DA:784:A:C8 | 31:DA:792:G:C5 | 3.10 | 0.40 |
| 31:DA:907:U:O2' | 42:DQ:101:ARG:NH2 | 2.51 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 31:DA:970:C:H2' | 31:DA:971:C:C6 | 2.56 | 0.40 |
| 31:DA:1324:G:H4' | 31:DA:1616:A:C2 | 2.57 | 0.40 |
| 31:DA:1429:G:C5 | 31:DA:1568:G:C6 | 3.09 | 0.40 |
| 31:DA:1450(A):C:N4 | 31:DA:1451:C:H41 | 2.19 | 0.40 |
| 31:DA:1505:C:H2' | 31:DA:1506:C:O5' | 2.21 | 0.40 |
| 31:DA:1508:A:H2' | 31:DA:1509:C:OP1 | 2.22 | 0.40 |
| 31:DA:1721:G:O6 | 31:DA:1739:U:H5' | 2.22 | 0.40 |
| 31:DA:2275:C:H5' | 31:DA:2275:C:C6 | 2.56 | 0.40 |
| 31:DA:2352:A:C4 | 31:DA:2366:A:C2 | 3.10 | 0.40 |
| 31:DA:2419:U:H2' | 31:DA:2420:C:C6 | 2.57 | 0.40 |
| 31:DA:2452:C:N4 | 31:DA:2453:A:N6 | 2.70 | 0.40 |
| 31:DA:2494:G:C2' | 31:DA:2495:G:O5' | 2.70 | 0.40 |
| 31:DA:2619:C:H2' | 31:DA:2620:C:C6 | 2.54 | 0.40 |
| 32:DB:41:U:H2' | 32:DB:42:C:OP1 | 2.22 | 0.40 |
| 32:DB:74:U:H3' | 32:DB:75:G:H5" | 2.04 | 0.40 |
| 33:DD:25:THR:CG2 | 33:DD:82:ILE:N | 2.84 | 0.40 |
| 33:DD:218:ARG:HD3 | 33:DD:218:ARG:HH11 | 1.74 | 0.40 |
| 35:DF:160:ASN:ND2 | 35:DF:163:VAL:H | 2.19 | 0.40 |
| 38:DI:125:GLU:OE1 | 38:DI:141:LYS:HG2 | 2.21 | 0.40 |
| 38:DI:129:THR:CG2 | 38:DI:130:TYR:N | 2.84 | 0.40 |
| 41:DP:95:VAL:C | 41:DP:96:THR:CG2 | 2.94 | 0.40 |
| 42:DQ:9:TYR:C | 42:DQ:10:ARG:HG3 | 2.46 | 0.40 |
| 42:DQ:77:LYS:HA | 42:DQ:78:PRO:HD3 | 1.93 | 0.40 |
| 47:DV:18:LEU:O | 47:DV:19:LYS:HB2 | 2.22 | 0.40 |
| 47:DV:66:ARG:HD2 | 47:DV:66:ARG:C | 2.47 | 0.40 |
| 49:DX:33:LYS:C | 49:DX:35:THR:HG22 | 2.47 | 0.40 |

All (1) symmetry-related close contacts are listed below. The label for Atom-2 includes the symmetry operator and encoded unit-cell translations to be applied.

| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|---|---|---|---|
| 24:B2:12:GLU:CB | 31:DA:306:U:OP1[1_455] | 2.15 | 0.05 |

## 5.3 Torsion angles (i)

### 5.3.1 Protein backbone (i)

In the following table, the Percentiles column shows the percent Ramachandran outliers of the chain as a percentile score with respect to all X-ray entries followed by that with respect to entries of similar resolution.

The Analysed column shows the number of residues for which the backbone conformation was analysed, and the total number of residues.

| Mol | Chain | Analysed | Favoured | Allowed | Outliers | Percentiles | |
|---|---|---|---|---|---|---|---|
| 2 | AB | 233/256 (91%) | 178 (76%) | 38 (16%) | 17 (7%) | 1 | 4 |
| 2 | CB | 233/256 (91%) | 177 (76%) | 39 (17%) | 17 (7%) | 1 | 4 |
| 3 | AC | 205/239 (86%) | 155 (76%) | 36 (18%) | 14 (7%) | 1 | 5 |
| 3 | CC | 205/239 (86%) | 155 (76%) | 37 (18%) | 13 (6%) | 1 | 6 |
| 4 | AD | 206/209 (99%) | 138 (67%) | 52 (25%) | 16 (8%) | 1 | 4 |
| 4 | CD | 206/209 (99%) | 137 (66%) | 55 (27%) | 14 (7%) | 1 | 5 |
| 5 | AE | 149/162 (92%) | 105 (70%) | 31 (21%) | 13 (9%) | 0 | 3 |
| 5 | CE | 149/162 (92%) | 103 (69%) | 33 (22%) | 13 (9%) | 0 | 3 |
| 6 | AF | 99/101 (98%) | 76 (77%) | 15 (15%) | 8 (8%) | 1 | 3 |
| 6 | CF | 99/101 (98%) | 76 (77%) | 14 (14%) | 9 (9%) | 0 | 2 |
| 7 | AG | 153/156 (98%) | 130 (85%) | 19 (12%) | 4 (3%) | 4 | 23 |
| 7 | CG | 153/156 (98%) | 131 (86%) | 18 (12%) | 4 (3%) | 4 | 23 |
| 8 | AH | 136/138 (99%) | 98 (72%) | 31 (23%) | 7 (5%) | 1 | 10 |
| 8 | CH | 136/138 (99%) | 98 (72%) | 31 (23%) | 7 (5%) | 1 | 10 |
| 9 | AI | 123/128 (96%) | 92 (75%) | 24 (20%) | 7 (6%) | 1 | 8 |
| 9 | CI | 123/128 (96%) | 94 (76%) | 22 (18%) | 7 (6%) | 1 | 8 |
| 10 | AJ | 97/105 (92%) | 81 (84%) | 11 (11%) | 5 (5%) | 1 | 9 |
| 10 | CJ | 97/105 (92%) | 81 (84%) | 11 (11%) | 5 (5%) | 1 | 9 |
| 11 | AK | 117/129 (91%) | 87 (74%) | 26 (22%) | 4 (3%) | 3 | 17 |
| 11 | CK | 117/129 (91%) | 86 (74%) | 27 (23%) | 4 (3%) | 3 | 17 |
| 12 | AL | 123/135 (91%) | 82 (67%) | 31 (25%) | 10 (8%) | 1 | 3 |
| 12 | CL | 123/135 (91%) | 83 (68%) | 29 (24%) | 11 (9%) | 0 | 3 |
| 13 | AM | 107/126 (85%) | 84 (78%) | 17 (16%) | 6 (6%) | 1 | 8 |
| 13 | CM | 107/126 (85%) | 84 (78%) | 17 (16%) | 6 (6%) | 1 | 8 |
| 14 | AN | 58/61 (95%) | 45 (78%) | 11 (19%) | 2 (3%) | 3 | 17 |
| 14 | CN | 58/61 (95%) | 44 (76%) | 12 (21%) | 2 (3%) | 3 | 17 |
| 15 | AO | 86/89 (97%) | 62 (72%) | 19 (22%) | 5 (6%) | 1 | 8 |
| 15 | CO | 86/89 (97%) | 61 (71%) | 21 (24%) | 4 (5%) | 2 | 11 |
| 16 | AP | 82/88 (93%) | 48 (58%) | 27 (33%) | 7 (8%) | 0 | 3 |
| 16 | CP | 82/88 (93%) | 47 (57%) | 29 (35%) | 6 (7%) | 1 | 4 |

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| Mol | Chain | Analysed | Favoured | Allowed | Outliers | Percentiles |
|---|---|---|---|---|---|---|
| 17 | AQ | 98/105 (93%) | 74 (76%) | 18 (18%) | 6 (6%) | 1 7 |
| 17 | CQ | 98/105 (93%) | 73 (74%) | 19 (19%) | 6 (6%) | 1 7 |
| 18 | AR | 68/88 (77%) | 52 (76%) | 11 (16%) | 5 (7%) | 1 4 |
| 18 | CR | 68/88 (77%) | 51 (75%) | 13 (19%) | 4 (6%) | 1 7 |
| 19 | AS | 77/93 (83%) | 58 (75%) | 13 (17%) | 6 (8%) | 1 4 |
| 19 | CS | 77/93 (83%) | 59 (77%) | 12 (16%) | 6 (8%) | 1 4 |
| 20 | AT | 97/106 (92%) | 69 (71%) | 19 (20%) | 9 (9%) | 0 2 |
| 20 | CT | 97/106 (92%) | 65 (67%) | 23 (24%) | 9 (9%) | 0 2 |
| 21 | AU | 23/27 (85%) | 18 (78%) | 4 (17%) | 1 (4%) | 2 12 |
| 21 | CU | 23/27 (85%) | 18 (78%) | 4 (17%) | 1 (4%) | 2 12 |
| 22 | B0 | 83/85 (98%) | 65 (78%) | 14 (17%) | 4 (5%) | 2 11 |
| 22 | D0 | 83/85 (98%) | 64 (77%) | 15 (18%) | 4 (5%) | 2 11 |
| 23 | B1 | 87/98 (89%) | 48 (55%) | 17 (20%) | 22 (25%) | 0 0 |
| 23 | D1 | 87/98 (89%) | 45 (52%) | 19 (22%) | 23 (26%) | 0 0 |
| 24 | B2 | 49/72 (68%) | 23 (47%) | 19 (39%) | 7 (14%) | 0 1 |
| 24 | D2 | 49/72 (68%) | 23 (47%) | 18 (37%) | 8 (16%) | 0 0 |
| 25 | B3 | 58/60 (97%) | 52 (90%) | 4 (7%) | 2 (3%) | 3 17 |
| 25 | D3 | 58/60 (97%) | 51 (88%) | 5 (9%) | 2 (3%) | 3 17 |
| 26 | B4 | 30/71 (42%) | 5 (17%) | 11 (37%) | 14 (47%) | 0 0 |
| 26 | D4 | 30/71 (42%) | 5 (17%) | 10 (33%) | 15 (50%) | 0 0 |
| 27 | B5 | 57/60 (95%) | 38 (67%) | 11 (19%) | 8 (14%) | 0 1 |
| 27 | D5 | 57/60 (95%) | 36 (63%) | 14 (25%) | 7 (12%) | 0 1 |
| 28 | B6 | 41/54 (76%) | 21 (51%) | 6 (15%) | 14 (34%) | 0 0 |
| 28 | D6 | 41/54 (76%) | 19 (46%) | 8 (20%) | 14 (34%) | 0 0 |
| 29 | B7 | 47/49 (96%) | 41 (87%) | 4 (8%) | 2 (4%) | 2 12 |
| 29 | D7 | 47/49 (96%) | 40 (85%) | 4 (8%) | 3 (6%) | 1 6 |
| 30 | B8 | 62/65 (95%) | 42 (68%) | 11 (18%) | 9 (14%) | 0 1 |
| 30 | D8 | 62/65 (95%) | 41 (66%) | 12 (19%) | 9 (14%) | 0 1 |
| 33 | BD | 270/276 (98%) | 208 (77%) | 45 (17%) | 17 (6%) | 1 6 |
| 33 | DD | 270/276 (98%) | 207 (77%) | 47 (17%) | 16 (6%) | 1 7 |
| 34 | BE | 203/206 (98%) | 138 (68%) | 37 (18%) | 28 (14%) | 0 1 |

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| Mol | Chain | Analysed | Favoured | Allowed | Outliers | Percentiles | |
|---|---|---|---|---|---|---|---|
| 34 | DE | 203/206 (98%) | 138 (68%) | 38 (19%) | 27 (13%) | 0 | 1 |
| 35 | BF | 206/210 (98%) | 160 (78%) | 30 (15%) | 16 (8%) | 1 | 4 |
| 35 | DF | 206/210 (98%) | 156 (76%) | 33 (16%) | 17 (8%) | 0 | 3 |
| 36 | BG | 177/182 (97%) | 128 (72%) | 35 (20%) | 14 (8%) | 1 | 3 |
| 36 | DG | 177/182 (97%) | 127 (72%) | 36 (20%) | 14 (8%) | 1 | 3 |
| 37 | BH | 158/180 (88%) | 92 (58%) | 41 (26%) | 25 (16%) | 0 | 0 |
| 37 | DH | 158/180 (88%) | 93 (59%) | 39 (25%) | 26 (16%) | 0 | 0 |
| 38 | BI | 144/148 (97%) | 88 (61%) | 32 (22%) | 24 (17%) | 0 | 0 |
| 38 | DI | 144/148 (97%) | 87 (60%) | 35 (24%) | 22 (15%) | 0 | 0 |
| 39 | BN | 137/140 (98%) | 87 (64%) | 32 (23%) | 18 (13%) | 0 | 1 |
| 39 | DN | 137/140 (98%) | 88 (64%) | 32 (23%) | 17 (12%) | 0 | 1 |
| 40 | BO | 120/122 (98%) | 101 (84%) | 16 (13%) | 3 (2%) | 4 | 23 |
| 40 | DO | 120/122 (98%) | 99 (82%) | 17 (14%) | 4 (3%) | 3 | 17 |
| 41 | BP | 144/150 (96%) | 77 (54%) | 17 (12%) | 50 (35%) | 0 | 0 |
| 41 | DP | 144/150 (96%) | 76 (53%) | 18 (12%) | 50 (35%) | 0 | 0 |
| 42 | BQ | 134/141 (95%) | 92 (69%) | 28 (21%) | 14 (10%) | 0 | 2 |
| 42 | DQ | 134/141 (95%) | 96 (72%) | 23 (17%) | 15 (11%) | 0 | 1 |
| 43 | BR | 115/118 (98%) | 78 (68%) | 29 (25%) | 8 (7%) | 1 | 4 |
| 43 | DR | 115/118 (98%) | 82 (71%) | 24 (21%) | 9 (8%) | 1 | 4 |
| 44 | BS | 97/112 (87%) | 49 (50%) | 24 (25%) | 24 (25%) | 0 | 0 |
| 44 | DS | 97/112 (87%) | 49 (50%) | 23 (24%) | 25 (26%) | 0 | 0 |
| 45 | BT | 130/146 (89%) | 89 (68%) | 21 (16%) | 20 (15%) | 0 | 0 |
| 45 | DT | 130/146 (89%) | 90 (69%) | 21 (16%) | 19 (15%) | 0 | 0 |
| 46 | BU | 115/118 (98%) | 77 (67%) | 27 (24%) | 11 (10%) | 0 | 2 |
| 46 | DU | 115/118 (98%) | 74 (64%) | 29 (25%) | 12 (10%) | 0 | 2 |
| 47 | BV | 97/101 (96%) | 54 (56%) | 15 (16%) | 28 (29%) | 0 | 0 |
| 47 | DV | 97/101 (96%) | 52 (54%) | 18 (19%) | 27 (28%) | 0 | 0 |
| 48 | BW | 111/113 (98%) | 88 (79%) | 15 (14%) | 8 (7%) | 1 | 4 |
| 48 | DW | 111/113 (98%) | 89 (80%) | 15 (14%) | 7 (6%) | 1 | 6 |
| 49 | BX | 91/96 (95%) | 47 (52%) | 22 (24%) | 22 (24%) | 0 | 0 |
| 49 | DX | 91/96 (95%) | 48 (53%) | 22 (24%) | 21 (23%) | 0 | 0 |

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| Mol | Chain | Analysed | Favoured | Allowed | Outliers | Percentiles | |
|---|---|---|---|---|---|---|---|
| 50 | BY | 99/110 (90%) | 45 (46%) | 22 (22%) | 32 (32%) | 0 | 0 |
| 50 | DY | 99/110 (90%) | 46 (46%) | 21 (21%) | 32 (32%) | 0 | 0 |
| 51 | BZ | 175/206 (85%) | 113 (65%) | 43 (25%) | 19 (11%) | 0 | 1 |
| 51 | DZ | 175/206 (85%) | 113 (65%) | 44 (25%) | 18 (10%) | 0 | 2 |
| All | All | 11148/12060 (92%) | 7735 (69%) | 2187 (20%) | 1226 (11%) | 0 | 1 |

All (1226) Ramachandran outliers are listed below:

| Mol | Chain | Res | Type |
|---|---|---|---|
| 2 | AB | 15 | VAL |
| 2 | AB | 24 | TRP |
| 2 | AB | 154 | LEU |
| 2 | AB | 165 | VAL |
| 2 | AB | 194 | PRO |
| 2 | AB | 195 | ASP |
| 2 | AB | 226 | ARG |
| 3 | AC | 4 | LYS |
| 3 | AC | 12 | LEU |
| 3 | AC | 101 | LEU |
| 3 | AC | 189 | ALA |
| 4 | AD | 3 | ARG |
| 4 | AD | 4 | TYR |
| 4 | AD | 14 | ARG |
| 4 | AD | 53 | ASP |
| 4 | AD | 129 | ASN |
| 5 | AE | 73 | ASN |
| 5 | AE | 121 | LYS |
| 6 | AF | 39 | LYS |
| 6 | AF | 40 | VAL |
| 6 | AF | 81 | ILE |
| 7 | AG | 33 | ASP |
| 9 | AI | 23 | ASN |
| 9 | AI | 107 | ARG |
| 10 | AJ | 59 | SER |
| 11 | AK | 100 | ALA |
| 12 | AL | 47 | LYS |
| 12 | AL | 115 | LYS |
| 13 | AM | 12 | ASN |
| 13 | AM | 83 | ASP |
| 13 | AM | 106 | ASN |
| 16 | AP | 17 | TYR |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 16 | AP | 19 | ILE |
| 17 | AQ | 34 | LYS |
| 20 | AT | 9 | ASN |
| 20 | AT | 11 | SER |
| 20 | AT | 95 | ALA |
| 22 | B0 | 14 | ARG |
| 22 | B0 | 44 | ARG |
| 23 | B1 | 11 | ARG |
| 23 | B1 | 14 | VAL |
| 23 | B1 | 48 | LYS |
| 23 | B1 | 64 | ALA |
| 23 | B1 | 65 | SER |
| 23 | B1 | 81 | LYS |
| 24 | B2 | 16 | LEU |
| 24 | B2 | 35 | LEU |
| 24 | B2 | 46 | GLN |
| 24 | B2 | 52 | ASP |
| 26 | B4 | 6 | HIS |
| 26 | B4 | 7 | PRO |
| 26 | B4 | 10 | VAL |
| 26 | B4 | 16 | CYS |
| 26 | B4 | 24 | THR |
| 26 | B4 | 25 | TYR |
| 26 | B4 | 27 | THR |
| 26 | B4 | 29 | PRO |
| 27 | B5 | 4 | HIS |
| 27 | B5 | 47 | PRO |
| 27 | B5 | 49 | CYS |
| 27 | B5 | 57 | VAL |
| 28 | B6 | 20 | ASN |
| 28 | B6 | 31 | PRO |
| 28 | B6 | 33 | LYS |
| 28 | B6 | 49 | HIS |
| 28 | B6 | 52 | VAL |
| 30 | B8 | 32 | LEU |
| 30 | B8 | 35 | GLN |
| 30 | B8 | 36 | LYS |
| 30 | B8 | 37 | SER |
| 30 | B8 | 52 | LYS |
| 33 | BD | 26 | LYS |
| 33 | BD | 28 | GLU |
| 33 | BD | 32 | SER |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 33 | BD | 159 | ALA |
| 33 | BD | 225 | ALA |
| 33 | BD | 241 | PRO |
| 33 | BD | 272 | ALA |
| 34 | BE | 54 | GLN |
| 34 | BE | 60 | ASN |
| 34 | BE | 61 | ARG |
| 34 | BE | 77 | ILE |
| 34 | BE | 82 | ARG |
| 34 | BE | 88 | GLY |
| 34 | BE | 93 | VAL |
| 34 | BE | 118 | LYS |
| 34 | BE | 131 | ALA |
| 34 | BE | 185 | LYS |
| 34 | BE | 200 | GLU |
| 35 | BF | 2 | LYS |
| 35 | BF | 14 | PRO |
| 35 | BF | 66 | PRO |
| 35 | BF | 89 | VAL |
| 36 | BG | 14 | GLU |
| 36 | BG | 82 | LEU |
| 36 | BG | 86 | MET |
| 36 | BG | 87 | PRO |
| 36 | BG | 90 | LEU |
| 37 | BH | 41 | MET |
| 37 | BH | 47 | GLU |
| 37 | BH | 55 | PRO |
| 37 | BH | 56 | SER |
| 37 | BH | 83 | TYR |
| 37 | BH | 89 | ILE |
| 37 | BH | 90 | LYS |
| 37 | BH | 138 | LYS |
| 37 | BH | 153 | LYS |
| 37 | BH | 154 | PRO |
| 37 | BH | 155 | SER |
| 37 | BH | 156 | ALA |
| 37 | BH | 157 | TYR |
| 37 | BH | 165 | ALA |
| 38 | BI | 11 | ASN |
| 38 | BI | 68 | LEU |
| 38 | BI | 86 | THR |
| 38 | BI | 91 | SER |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 38 | BI | 133 | HIS |
| 38 | BI | 145 | VAL |
| 39 | BN | 58 | ASP |
| 39 | BN | 63 | THR |
| 39 | BN | 64 | GLY |
| 39 | BN | 74 | ARG |
| 39 | BN | 78 | TYR |
| 39 | BN | 79 | PRO |
| 39 | BN | 92 | ALA |
| 39 | BN | 130 | HIS |
| 40 | BO | 48 | PRO |
| 41 | BP | 12 | ALA |
| 41 | BP | 14 | LYS |
| 41 | BP | 15 | ARG |
| 41 | BP | 17 | LYS |
| 41 | BP | 18 | ARG |
| 41 | BP | 35 | HIS |
| 41 | BP | 36 | LYS |
| 41 | BP | 40 | SER |
| 41 | BP | 42 | SER |
| 41 | BP | 47 | ASP |
| 41 | BP | 49 | ARG |
| 41 | BP | 52 | GLU |
| 41 | BP | 56 | SER |
| 41 | BP | 57 | THR |
| 41 | BP | 58 | THR |
| 41 | BP | 74 | GLU |
| 41 | BP | 103 | ALA |
| 41 | BP | 106 | LEU |
| 41 | BP | 141 | ALA |
| 41 | BP | 147 | LEU |
| 42 | BQ | 8 | LYS |
| 42 | BQ | 13 | GLN |
| 42 | BQ | 30 | GLY |
| 42 | BQ | 79 | LEU |
| 42 | BQ | 90 | VAL |
| 42 | BQ | 134 | ARG |
| 43 | BR | 4 | LEU |
| 43 | BR | 5 | LYS |
| 43 | BR | 107 | ASP |
| 43 | BR | 117 | VAL |
| 44 | BS | 57 | LYS |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 44 | BS | 59 | LYS |
| 44 | BS | 67 | ARG |
| 44 | BS | 88 | ASP |
| 44 | BS | 89 | ARG |
| 44 | BS | 92 | TYR |
| 44 | BS | 102 | ALA |
| 45 | BT | 18 | ASP |
| 45 | BT | 24 | PRO |
| 45 | BT | 26 | ASP |
| 45 | BT | 31 | SER |
| 45 | BT | 33 | LYS |
| 45 | BT | 36 | GLU |
| 45 | BT | 80 | SER |
| 45 | BT | 94 | ALA |
| 45 | BT | 107 | ASP |
| 45 | BT | 115 | ARG |
| 46 | BU | 9 | VAL |
| 46 | BU | 32 | PHE |
| 47 | BV | 19 | LYS |
| 47 | BV | 23 | GLU |
| 47 | BV | 44 | LYS |
| 47 | BV | 47 | VAL |
| 47 | BV | 51 | VAL |
| 47 | BV | 68 | LYS |
| 47 | BV | 86 | GLY |
| 47 | BV | 90 | PRO |
| 49 | BX | 25 | LYS |
| 49 | BX | 34 | ALA |
| 49 | BX | 36 | LYS |
| 49 | BX | 37 | THR |
| 49 | BX | 40 | LYS |
| 49 | BX | 60 | ARG |
| 49 | BX | 65 | ARG |
| 49 | BX | 71 | GLY |
| 49 | BX | 77 | LYS |
| 49 | BX | 84 | ALA |
| 49 | BX | 86 | GLY |
| 49 | BX | 89 | ILE |
| 50 | BY | 3 | VAL |
| 50 | BY | 16 | ALA |
| 50 | BY | 17 | SER |
| 50 | BY | 38 | ILE |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 50 | BY | 42 | VAL |
| 50 | BY | 47 | LYS |
| 50 | BY | 56 | PRO |
| 50 | BY | 62 | GLU |
| 50 | BY | 66 | PRO |
| 50 | BY | 77 | PRO |
| 50 | BY | 78 | ALA |
| 50 | BY | 90 | LEU |
| 50 | BY | 99 | CYS |
| 50 | BY | 101 | LYS |
| 51 | BZ | 65 | GLN |
| 51 | BZ | 101 | PRO |
| 51 | BZ | 112 | ARG |
| 2 | CB | 15 | VAL |
| 2 | CB | 24 | TRP |
| 2 | CB | 84 | GLU |
| 2 | CB | 154 | LEU |
| 2 | CB | 165 | VAL |
| 2 | CB | 194 | PRO |
| 2 | CB | 195 | ASP |
| 2 | CB | 226 | ARG |
| 3 | CC | 4 | LYS |
| 3 | CC | 12 | LEU |
| 3 | CC | 101 | LEU |
| 3 | CC | 189 | ALA |
| 4 | CD | 3 | ARG |
| 4 | CD | 14 | ARG |
| 4 | CD | 53 | ASP |
| 4 | CD | 129 | ASN |
| 5 | CE | 73 | ASN |
| 5 | CE | 121 | LYS |
| 6 | CF | 13 | ASN |
| 6 | CF | 40 | VAL |
| 6 | CF | 81 | ILE |
| 7 | CG | 33 | ASP |
| 9 | CI | 23 | ASN |
| 9 | CI | 107 | ARG |
| 10 | CJ | 59 | SER |
| 11 | CK | 100 | ALA |
| 12 | CL | 47 | LYS |
| 12 | CL | 115 | LYS |
| 13 | CM | 12 | ASN |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 13 | CM | 83 | ASP |
| 13 | CM | 106 | ASN |
| 16 | CP | 17 | TYR |
| 16 | CP | 19 | ILE |
| 20 | CT | 11 | SER |
| 20 | CT | 95 | ALA |
| 22 | D0 | 5 | LYS |
| 22 | D0 | 14 | ARG |
| 22 | D0 | 44 | ARG |
| 23 | D1 | 10 | LYS |
| 23 | D1 | 11 | ARG |
| 23 | D1 | 14 | VAL |
| 23 | D1 | 48 | LYS |
| 23 | D1 | 64 | ALA |
| 23 | D1 | 65 | SER |
| 23 | D1 | 81 | LYS |
| 24 | D2 | 16 | LEU |
| 24 | D2 | 35 | LEU |
| 24 | D2 | 46 | GLN |
| 24 | D2 | 52 | ASP |
| 26 | D4 | 6 | HIS |
| 26 | D4 | 7 | PRO |
| 26 | D4 | 10 | VAL |
| 26 | D4 | 16 | CYS |
| 26 | D4 | 24 | THR |
| 26 | D4 | 25 | TYR |
| 26 | D4 | 27 | THR |
| 26 | D4 | 29 | PRO |
| 27 | D5 | 4 | HIS |
| 27 | D5 | 47 | PRO |
| 27 | D5 | 49 | CYS |
| 27 | D5 | 57 | VAL |
| 28 | D6 | 15 | GLU |
| 28 | D6 | 20 | ASN |
| 28 | D6 | 31 | PRO |
| 28 | D6 | 33 | LYS |
| 28 | D6 | 34 | LEU |
| 28 | D6 | 49 | HIS |
| 28 | D6 | 52 | VAL |
| 30 | D8 | 32 | LEU |
| 30 | D8 | 35 | GLN |
| 30 | D8 | 37 | SER |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 30 | D8 | 52 | LYS |
| 33 | DD | 26 | LYS |
| 33 | DD | 28 | GLU |
| 33 | DD | 32 | SER |
| 33 | DD | 159 | ALA |
| 33 | DD | 225 | ALA |
| 33 | DD | 241 | PRO |
| 33 | DD | 272 | ALA |
| 34 | DE | 4 | ILE |
| 34 | DE | 54 | GLN |
| 34 | DE | 61 | ARG |
| 34 | DE | 77 | ILE |
| 34 | DE | 82 | ARG |
| 34 | DE | 88 | GLY |
| 34 | DE | 93 | VAL |
| 34 | DE | 118 | LYS |
| 34 | DE | 131 | ALA |
| 34 | DE | 200 | GLU |
| 35 | DF | 2 | LYS |
| 35 | DF | 14 | PRO |
| 35 | DF | 20 | LEU |
| 35 | DF | 66 | PRO |
| 36 | DG | 14 | GLU |
| 36 | DG | 82 | LEU |
| 36 | DG | 86 | MET |
| 36 | DG | 87 | PRO |
| 36 | DG | 90 | LEU |
| 37 | DH | 41 | MET |
| 37 | DH | 47 | GLU |
| 37 | DH | 55 | PRO |
| 37 | DH | 83 | TYR |
| 37 | DH | 89 | ILE |
| 37 | DH | 90 | LYS |
| 37 | DH | 138 | LYS |
| 37 | DH | 153 | LYS |
| 37 | DH | 154 | PRO |
| 37 | DH | 155 | SER |
| 37 | DH | 156 | ALA |
| 37 | DH | 157 | TYR |
| 37 | DH | 165 | ALA |
| 38 | DI | 11 | ASN |
| 38 | DI | 68 | LEU |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 38 | DI | 86 | THR |
| 38 | DI | 91 | SER |
| 38 | DI | 133 | HIS |
| 38 | DI | 145 | VAL |
| 39 | DN | 58 | ASP |
| 39 | DN | 63 | THR |
| 39 | DN | 64 | GLY |
| 39 | DN | 74 | ARG |
| 39 | DN | 78 | TYR |
| 39 | DN | 79 | PRO |
| 39 | DN | 92 | ALA |
| 39 | DN | 130 | HIS |
| 40 | DO | 48 | PRO |
| 41 | DP | 12 | ALA |
| 41 | DP | 14 | LYS |
| 41 | DP | 15 | ARG |
| 41 | DP | 17 | LYS |
| 41 | DP | 18 | ARG |
| 41 | DP | 35 | HIS |
| 41 | DP | 36 | LYS |
| 41 | DP | 40 | SER |
| 41 | DP | 42 | SER |
| 41 | DP | 47 | ASP |
| 41 | DP | 49 | ARG |
| 41 | DP | 52 | GLU |
| 41 | DP | 56 | SER |
| 41 | DP | 57 | THR |
| 41 | DP | 58 | THR |
| 41 | DP | 74 | GLU |
| 41 | DP | 103 | ALA |
| 41 | DP | 104 | GLY |
| 41 | DP | 106 | LEU |
| 41 | DP | 141 | ALA |
| 41 | DP | 147 | LEU |
| 42 | DQ | 8 | LYS |
| 42 | DQ | 13 | GLN |
| 42 | DQ | 30 | GLY |
| 42 | DQ | 79 | LEU |
| 42 | DQ | 90 | VAL |
| 42 | DQ | 134 | ARG |
| 43 | DR | 4 | LEU |
| 43 | DR | 5 | LYS |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 43 | DR | 107 | ASP |
| 43 | DR | 117 | VAL |
| 44 | DS | 57 | LYS |
| 44 | DS | 59 | LYS |
| 44 | DS | 67 | ARG |
| 44 | DS | 88 | ASP |
| 44 | DS | 89 | ARG |
| 44 | DS | 92 | TYR |
| 44 | DS | 102 | ALA |
| 45 | DT | 18 | ASP |
| 45 | DT | 24 | PRO |
| 45 | DT | 26 | ASP |
| 45 | DT | 31 | SER |
| 45 | DT | 33 | LYS |
| 45 | DT | 35 | LYS |
| 45 | DT | 36 | GLU |
| 45 | DT | 80 | SER |
| 45 | DT | 107 | ASP |
| 45 | DT | 115 | ARG |
| 46 | DU | 9 | VAL |
| 46 | DU | 32 | PHE |
| 46 | DU | 52 | ARG |
| 46 | DU | 91 | ASP |
| 47 | DV | 19 | LYS |
| 47 | DV | 23 | GLU |
| 47 | DV | 44 | LYS |
| 47 | DV | 47 | VAL |
| 47 | DV | 51 | VAL |
| 47 | DV | 86 | GLY |
| 47 | DV | 90 | PRO |
| 49 | DX | 25 | LYS |
| 49 | DX | 34 | ALA |
| 49 | DX | 36 | LYS |
| 49 | DX | 37 | THR |
| 49 | DX | 40 | LYS |
| 49 | DX | 60 | ARG |
| 49 | DX | 77 | LYS |
| 49 | DX | 84 | ALA |
| 49 | DX | 86 | GLY |
| 49 | DX | 89 | ILE |
| 50 | DY | 3 | VAL |
| 50 | DY | 16 | ALA |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 50 | DY | 17 | SER |
| 50 | DY | 38 | ILE |
| 50 | DY | 42 | VAL |
| 50 | DY | 47 | LYS |
| 50 | DY | 56 | PRO |
| 50 | DY | 62 | GLU |
| 50 | DY | 66 | PRO |
| 50 | DY | 77 | PRO |
| 50 | DY | 78 | ALA |
| 50 | DY | 90 | LEU |
| 50 | DY | 99 | CYS |
| 50 | DY | 101 | LYS |
| 51 | DZ | 65 | GLN |
| 51 | DZ | 112 | ARG |
| 2 | AB | 18 | GLY |
| 2 | AB | 84 | GLU |
| 2 | AB | 232 | PRO |
| 2 | AB | 239 | VAL |
| 3 | AC | 18 | TRP |
| 3 | AC | 47 | LEU |
| 3 | AC | 100 | ALA |
| 3 | AC | 145 | GLY |
| 4 | AD | 10 | ARG |
| 4 | AD | 13 | ARG |
| 4 | AD | 17 | VAL |
| 4 | AD | 44 | GLY |
| 4 | AD | 47 | ARG |
| 4 | AD | 110 | PHE |
| 5 | AE | 27 | ARG |
| 5 | AE | 108 | ALA |
| 5 | AE | 132 | ALA |
| 5 | AE | 146 | ALA |
| 5 | AE | 153 | LYS |
| 6 | AF | 13 | ASN |
| 7 | AG | 7 | ALA |
| 8 | AH | 2 | LEU |
| 8 | AH | 51 | VAL |
| 8 | AH | 54 | ASP |
| 9 | AI | 100 | GLY |
| 9 | AI | 124 | GLN |
| 10 | AJ | 23 | ILE |
| 10 | AJ | 36 | GLY |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 11 | AK | 63 | LEU |
| 12 | AL | 23 | LYS |
| 12 | AL | 28 | LYS |
| 12 | AL | 91 | LYS |
| 12 | AL | 106 | ASP |
| 13 | AM | 100 | GLY |
| 16 | AP | 28 | ARG |
| 17 | AQ | 78 | GLU |
| 18 | AR | 45 | SER |
| 18 | AR | 54 | ARG |
| 19 | AS | 27 | GLU |
| 19 | AS | 28 | LYS |
| 20 | AT | 74 | LYS |
| 22 | B0 | 5 | LYS |
| 23 | B1 | 10 | LYS |
| 23 | B1 | 26 | ARG |
| 23 | B1 | 33 | LYS |
| 23 | B1 | 49 | VAL |
| 23 | B1 | 79 | GLY |
| 23 | B1 | 83 | GLU |
| 23 | B1 | 86 | SER |
| 23 | B1 | 87 | PRO |
| 24 | B2 | 42 | GLY |
| 24 | B2 | 49 | LYS |
| 26 | B4 | 20 | ASN |
| 27 | B5 | 48 | GLU |
| 28 | B6 | 15 | GLU |
| 28 | B6 | 23 | THR |
| 28 | B6 | 29 | ASN |
| 28 | B6 | 34 | LEU |
| 28 | B6 | 44 | ARG |
| 30 | B8 | 31 | HIS |
| 33 | BD | 3 | VAL |
| 33 | BD | 156 | ALA |
| 34 | BE | 4 | ILE |
| 34 | BE | 33 | VAL |
| 34 | BE | 71 | GLY |
| 34 | BE | 72 | VAL |
| 34 | BE | 76 | ARG |
| 35 | BF | 20 | LEU |
| 35 | BF | 133 | ASN |
| 35 | BF | 206 | ILE |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 36 | BG | 30 | GLU |
| 36 | BG | 47 | LYS |
| 36 | BG | 49 | ASP |
| 36 | BG | 96 | ARG |
| 36 | BG | 153 | ARG |
| 37 | BH | 84 | SER |
| 37 | BH | 93 | GLY |
| 37 | BH | 139 | GLN |
| 37 | BH | 158 | HIS |
| 38 | BI | 15 | VAL |
| 38 | BI | 88 | ILE |
| 38 | BI | 120 | ILE |
| 38 | BI | 144 | VAL |
| 39 | BN | 57 | ALA |
| 39 | BN | 73 | THR |
| 39 | BN | 83 | LYS |
| 39 | BN | 98 | VAL |
| 39 | BN | 125 | GLY |
| 40 | BO | 5 | GLN |
| 41 | BP | 10 | PRO |
| 41 | BP | 25 | SER |
| 41 | BP | 31 | ALA |
| 41 | BP | 34 | GLY |
| 41 | BP | 37 | GLY |
| 41 | BP | 98 | GLU |
| 41 | BP | 104 | GLY |
| 41 | BP | 107 | LYS |
| 41 | BP | 108 | LYS |
| 41 | BP | 109 | GLY |
| 41 | BP | 132 | LYS |
| 42 | BQ | 11 | LYS |
| 42 | BQ | 19 | GLY |
| 42 | BQ | 21 | THR |
| 42 | BQ | 62 | GLY |
| 42 | BQ | 83 | MET |
| 44 | BS | 58 | LEU |
| 44 | BS | 77 | ALA |
| 44 | BS | 85 | VAL |
| 44 | BS | 90 | GLY |
| 44 | BS | 94 | TYR |
| 44 | BS | 96 | GLY |
| 44 | BS | 100 | ALA |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 44 | BS | 106 | ARG |
| 45 | BT | 32 | TYR |
| 45 | BT | 35 | LYS |
| 45 | BT | 58 | ASN |
| 46 | BU | 26 | GLY |
| 46 | BU | 52 | ARG |
| 46 | BU | 89 | GLU |
| 46 | BU | 91 | ASP |
| 46 | BU | 92 | ARG |
| 47 | BV | 41 | GLY |
| 47 | BV | 50 | PRO |
| 47 | BV | 54 | GLY |
| 47 | BV | 55 | ALA |
| 47 | BV | 69 | LYS |
| 47 | BV | 70 | ILE |
| 47 | BV | 72 | VAL |
| 47 | BV | 74 | LYS |
| 48 | BW | 45 | TYR |
| 48 | BW | 63 | ASP |
| 48 | BW | 65 | LEU |
| 49 | BX | 19 | ALA |
| 49 | BX | 24 | GLY |
| 49 | BX | 59 | VAL |
| 49 | BX | 69 | TYR |
| 49 | BX | 81 | VAL |
| 50 | BY | 18 | GLY |
| 50 | BY | 27 | VAL |
| 50 | BY | 55 | TYR |
| 50 | BY | 80 | GLY |
| 50 | BY | 98 | VAL |
| 51 | BZ | 64 | GLY |
| 51 | BZ | 119 | GLU |
| 51 | BZ | 168 | GLU |
| 2 | CB | 18 | GLY |
| 2 | CB | 56 | ARG |
| 2 | CB | 232 | PRO |
| 2 | CB | 239 | VAL |
| 3 | CC | 15 | THR |
| 3 | CC | 18 | TRP |
| 3 | CC | 47 | LEU |
| 3 | CC | 100 | ALA |
| 3 | CC | 145 | GLY |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 4 | CD | 4 | TYR |
| 4 | CD | 10 | ARG |
| 4 | CD | 13 | ARG |
| 4 | CD | 44 | GLY |
| 4 | CD | 47 | ARG |
| 4 | CD | 110 | PHE |
| 5 | CE | 27 | ARG |
| 5 | CE | 108 | ALA |
| 5 | CE | 132 | ALA |
| 5 | CE | 140 | ARG |
| 5 | CE | 146 | ALA |
| 5 | CE | 153 | LYS |
| 6 | CF | 39 | LYS |
| 7 | CG | 7 | ALA |
| 8 | CH | 51 | VAL |
| 8 | CH | 54 | ASP |
| 8 | CH | 133 | LEU |
| 9 | CI | 100 | GLY |
| 10 | CJ | 23 | ILE |
| 10 | CJ | 36 | GLY |
| 11 | CK | 63 | LEU |
| 12 | CL | 23 | LYS |
| 12 | CL | 28 | LYS |
| 12 | CL | 91 | LYS |
| 12 | CL | 106 | ASP |
| 13 | CM | 100 | GLY |
| 17 | CQ | 34 | LYS |
| 17 | CQ | 78 | GLU |
| 18 | CR | 45 | SER |
| 18 | CR | 54 | ARG |
| 19 | CS | 27 | GLU |
| 19 | CS | 28 | LYS |
| 20 | CT | 9 | ASN |
| 20 | CT | 74 | LYS |
| 23 | D1 | 26 | ARG |
| 23 | D1 | 33 | LYS |
| 23 | D1 | 49 | VAL |
| 23 | D1 | 79 | GLY |
| 23 | D1 | 83 | GLU |
| 23 | D1 | 87 | PRO |
| 24 | D2 | 42 | GLY |
| 24 | D2 | 49 | LYS |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 25 | D3 | 13 | ILE |
| 26 | D4 | 20 | ASN |
| 28 | D6 | 23 | THR |
| 28 | D6 | 29 | ASN |
| 28 | D6 | 44 | ARG |
| 30 | D8 | 31 | HIS |
| 30 | D8 | 36 | LYS |
| 33 | DD | 3 | VAL |
| 33 | DD | 156 | ALA |
| 34 | DE | 29 | GLY |
| 34 | DE | 33 | VAL |
| 34 | DE | 60 | ASN |
| 34 | DE | 71 | GLY |
| 34 | DE | 72 | VAL |
| 34 | DE | 185 | LYS |
| 35 | DF | 89 | VAL |
| 35 | DF | 133 | ASN |
| 35 | DF | 176 | LEU |
| 36 | DG | 30 | GLU |
| 36 | DG | 47 | LYS |
| 36 | DG | 49 | ASP |
| 36 | DG | 96 | ARG |
| 36 | DG | 153 | ARG |
| 37 | DH | 56 | SER |
| 37 | DH | 84 | SER |
| 37 | DH | 93 | GLY |
| 37 | DH | 139 | GLN |
| 37 | DH | 158 | HIS |
| 38 | DI | 15 | VAL |
| 38 | DI | 52 | ARG |
| 38 | DI | 88 | ILE |
| 38 | DI | 120 | ILE |
| 38 | DI | 144 | VAL |
| 39 | DN | 57 | ALA |
| 39 | DN | 73 | THR |
| 39 | DN | 83 | LYS |
| 39 | DN | 98 | VAL |
| 39 | DN | 125 | GLY |
| 40 | DO | 5 | GLN |
| 41 | DP | 10 | PRO |
| 41 | DP | 25 | SER |
| 41 | DP | 31 | ALA |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 41 | DP | 34 | GLY |
| 41 | DP | 90 | ARG |
| 41 | DP | 98 | GLU |
| 41 | DP | 107 | LYS |
| 41 | DP | 108 | LYS |
| 41 | DP | 109 | GLY |
| 41 | DP | 132 | LYS |
| 42 | DQ | 11 | LYS |
| 42 | DQ | 19 | GLY |
| 42 | DQ | 21 | THR |
| 42 | DQ | 62 | GLY |
| 42 | DQ | 83 | MET |
| 44 | DS | 77 | ALA |
| 44 | DS | 85 | VAL |
| 44 | DS | 90 | GLY |
| 44 | DS | 94 | TYR |
| 44 | DS | 96 | GLY |
| 44 | DS | 100 | ALA |
| 45 | DT | 28 | VAL |
| 45 | DT | 32 | TYR |
| 45 | DT | 58 | ASN |
| 45 | DT | 94 | ALA |
| 46 | DU | 26 | GLY |
| 46 | DU | 34 | LYS |
| 46 | DU | 89 | GLU |
| 47 | DV | 41 | GLY |
| 47 | DV | 50 | PRO |
| 47 | DV | 54 | GLY |
| 47 | DV | 68 | LYS |
| 47 | DV | 70 | ILE |
| 47 | DV | 72 | VAL |
| 48 | DW | 45 | TYR |
| 48 | DW | 56 | ALA |
| 48 | DW | 63 | ASP |
| 48 | DW | 65 | LEU |
| 49 | DX | 19 | ALA |
| 49 | DX | 24 | GLY |
| 49 | DX | 59 | VAL |
| 49 | DX | 65 | ARG |
| 49 | DX | 71 | GLY |
| 49 | DX | 81 | VAL |
| 50 | DY | 18 | GLY |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 50 | DY | 27 | VAL |
| 50 | DY | 37 | VAL |
| 50 | DY | 55 | TYR |
| 50 | DY | 80 | GLY |
| 50 | DY | 81 | LYS |
| 50 | DY | 98 | VAL |
| 50 | DY | 100 | ALA |
| 51 | DZ | 64 | GLY |
| 51 | DZ | 119 | GLU |
| 51 | DZ | 168 | GLU |
| 2 | AB | 56 | ARG |
| 2 | AB | 181 | PHE |
| 3 | AC | 15 | THR |
| 3 | AC | 154 | SER |
| 5 | AE | 11 | ILE |
| 5 | AE | 140 | ARG |
| 8 | AH | 133 | LEU |
| 10 | AJ | 54 | PHE |
| 12 | AL | 65 | GLU |
| 12 | AL | 79 | GLU |
| 13 | AM | 90 | LEU |
| 15 | AO | 4 | THR |
| 15 | AO | 29 | VAL |
| 17 | AQ | 96 | GLU |
| 18 | AR | 20 | ALA |
| 19 | AS | 25 | LYS |
| 23 | B1 | 44 | PRO |
| 23 | B1 | 94 | LEU |
| 26 | B4 | 11 | PRO |
| 26 | B4 | 12 | ALA |
| 26 | B4 | 23 | GLU |
| 27 | B5 | 32 | PRO |
| 29 | B7 | 42 | LEU |
| 30 | B8 | 30 | ARG |
| 33 | BD | 33 | LEU |
| 34 | BE | 17 | ASP |
| 34 | BE | 29 | GLY |
| 34 | BE | 53 | PRO |
| 34 | BE | 57 | LYS |
| 34 | BE | 90 | THR |
| 34 | BE | 117 | MET |
| 34 | BE | 129 | HIS |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 34 | BE | 182 | LEU |
| 35 | BF | 11 | VAL |
| 35 | BF | 127 | GLU |
| 35 | BF | 144 | LYS |
| 35 | BF | 176 | LEU |
| 36 | BG | 129 | GLY |
| 37 | BH | 21 | PRO |
| 37 | BH | 71 | LEU |
| 38 | BI | 5 | LEU |
| 38 | BI | 34 | GLY |
| 38 | BI | 39 | ALA |
| 38 | BI | 78 | THR |
| 38 | BI | 134 | PRO |
| 41 | BP | 67 | MET |
| 41 | BP | 90 | ARG |
| 41 | BP | 146 | VAL |
| 42 | BQ | 15 | GLY |
| 42 | BQ | 82 | ARG |
| 43 | BR | 106 | GLY |
| 44 | BS | 87 | PHE |
| 44 | BS | 103 | GLU |
| 45 | BT | 28 | VAL |
| 45 | BT | 88 | ILE |
| 45 | BT | 129 | ARG |
| 46 | BU | 25 | TRP |
| 47 | BV | 18 | LEU |
| 47 | BV | 27 | ALA |
| 47 | BV | 28 | GLU |
| 48 | BW | 6 | ILE |
| 49 | BX | 4 | ALA |
| 49 | BX | 72 | LYS |
| 49 | BX | 88 | LYS |
| 50 | BY | 10 | GLY |
| 50 | BY | 31 | LEU |
| 50 | BY | 37 | VAL |
| 50 | BY | 48 | ALA |
| 50 | BY | 50 | ARG |
| 50 | BY | 54 | LYS |
| 50 | BY | 67 | LEU |
| 50 | BY | 81 | LYS |
| 50 | BY | 100 | ALA |
| 51 | BZ | 80 | ARG |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 51 | BZ | 85 | HIS |
| 51 | BZ | 93 | ASP |
| 51 | BZ | 111 | VAL |
| 2 | CB | 181 | PHE |
| 3 | CC | 154 | SER |
| 5 | CE | 11 | ILE |
| 8 | CH | 2 | LEU |
| 9 | CI | 124 | GLN |
| 10 | CJ | 54 | PHE |
| 13 | CM | 90 | LEU |
| 15 | CO | 4 | THR |
| 15 | CO | 29 | VAL |
| 16 | CP | 28 | ARG |
| 18 | CR | 20 | ALA |
| 19 | CS | 25 | LYS |
| 20 | CT | 101 | GLY |
| 23 | D1 | 86 | SER |
| 23 | D1 | 94 | LEU |
| 24 | D2 | 59 | ARG |
| 26 | D4 | 11 | PRO |
| 26 | D4 | 12 | ALA |
| 26 | D4 | 23 | GLU |
| 27 | D5 | 32 | PRO |
| 27 | D5 | 48 | GLU |
| 28 | D6 | 51 | GLU |
| 29 | D7 | 42 | LEU |
| 30 | D8 | 30 | ARG |
| 33 | DD | 25 | THR |
| 33 | DD | 33 | LEU |
| 33 | DD | 45 | ASN |
| 34 | DE | 17 | ASP |
| 34 | DE | 53 | PRO |
| 34 | DE | 76 | ARG |
| 34 | DE | 90 | THR |
| 34 | DE | 129 | HIS |
| 35 | DF | 11 | VAL |
| 35 | DF | 127 | GLU |
| 35 | DF | 144 | LYS |
| 35 | DF | 206 | ILE |
| 36 | DG | 129 | GLY |
| 37 | DH | 21 | PRO |
| 37 | DH | 71 | LEU |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 38 | DI | 30 | LEU |
| 38 | DI | 34 | GLY |
| 38 | DI | 78 | THR |
| 38 | DI | 101 | LEU |
| 38 | DI | 134 | PRO |
| 39 | DN | 68 | GLU |
| 40 | DO | 91 | LEU |
| 41 | DP | 32 | THR |
| 41 | DP | 65 | ARG |
| 41 | DP | 146 | VAL |
| 42 | DQ | 15 | GLY |
| 42 | DQ | 51 | ARG |
| 42 | DQ | 82 | ARG |
| 43 | DR | 106 | GLY |
| 44 | DS | 58 | LEU |
| 44 | DS | 87 | PHE |
| 44 | DS | 103 | GLU |
| 44 | DS | 107 | GLU |
| 45 | DT | 83 | ILE |
| 45 | DT | 88 | ILE |
| 45 | DT | 129 | ARG |
| 46 | DU | 25 | TRP |
| 46 | DU | 68 | ALA |
| 46 | DU | 92 | ARG |
| 47 | DV | 18 | LEU |
| 47 | DV | 27 | ALA |
| 47 | DV | 28 | GLU |
| 47 | DV | 69 | LYS |
| 47 | DV | 74 | LYS |
| 47 | DV | 87 | HIS |
| 48 | DW | 6 | ILE |
| 49 | DX | 4 | ALA |
| 49 | DX | 69 | TYR |
| 49 | DX | 72 | LYS |
| 49 | DX | 88 | LYS |
| 50 | DY | 22 | GLY |
| 50 | DY | 48 | ALA |
| 50 | DY | 50 | ARG |
| 50 | DY | 54 | LYS |
| 50 | DY | 67 | LEU |
| 51 | DZ | 78 | LYS |
| 51 | DZ | 80 | ARG |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 51 | DZ | 85 | HIS |
| 51 | DZ | 93 | ASP |
| 51 | DZ | 101 | PRO |
| 51 | DZ | 111 | VAL |
| 2 | AB | 83 | MET |
| 2 | AB | 209 | ARG |
| 2 | AB | 240 | GLN |
| 3 | AC | 206 | GLU |
| 3 | AC | 207 | VAL |
| 5 | AE | 8 | GLU |
| 8 | AH | 20 | TYR |
| 9 | AI | 24 | GLY |
| 9 | AI | 97 | LYS |
| 9 | AI | 103 | THR |
| 11 | AK | 62 | GLN |
| 12 | AL | 19 | ARG |
| 16 | AP | 67 | THR |
| 17 | AQ | 3 | LYS |
| 19 | AS | 43 | GLU |
| 20 | AT | 97 | ALA |
| 20 | AT | 98 | PRO |
| 22 | B0 | 83 | PRO |
| 23 | B1 | 28 | GLY |
| 23 | B1 | 85 | LEU |
| 24 | B2 | 59 | ARG |
| 25 | B3 | 3 | ARG |
| 25 | B3 | 13 | ILE |
| 26 | B4 | 31 | ILE |
| 27 | B5 | 33 | CYS |
| 28 | B6 | 51 | GLU |
| 30 | B8 | 64 | TYR |
| 33 | BD | 25 | THR |
| 33 | BD | 45 | ASN |
| 33 | BD | 242 | ARG |
| 34 | BE | 2 | LYS |
| 34 | BE | 86 | PRO |
| 35 | BF | 10 | PRO |
| 36 | BG | 21 | ARG |
| 36 | BG | 43 | LEU |
| 36 | BG | 115 | ARG |
| 37 | BH | 146 | ALA |
| 38 | BI | 7 | GLU |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 38 | BI | 30 | LEU |
| 38 | BI | 101 | LEU |
| 39 | BN | 68 | GLU |
| 40 | BO | 91 | LEU |
| 41 | BP | 32 | THR |
| 41 | BP | 65 | ARG |
| 41 | BP | 89 | ALA |
| 42 | BQ | 51 | ARG |
| 43 | BR | 102 | GLU |
| 44 | BS | 35 | ILE |
| 44 | BS | 98 | VAL |
| 44 | BS | 107 | GLU |
| 46 | BU | 34 | LYS |
| 46 | BU | 68 | ALA |
| 47 | BV | 2 | PHE |
| 47 | BV | 73 | SER |
| 48 | BW | 11 | ARG |
| 50 | BY | 30 | VAL |
| 50 | BY | 39 | VAL |
| 50 | BY | 53 | PRO |
| 51 | BZ | 78 | LYS |
| 51 | BZ | 121 | HIS |
| 51 | BZ | 147 | GLY |
| 2 | CB | 83 | MET |
| 2 | CB | 209 | ARG |
| 2 | CB | 240 | GLN |
| 3 | CC | 206 | GLU |
| 3 | CC | 207 | VAL |
| 4 | CD | 171 | GLY |
| 5 | CE | 8 | GLU |
| 5 | CE | 37 | ARG |
| 8 | CH | 20 | TYR |
| 9 | CI | 24 | GLY |
| 9 | CI | 97 | LYS |
| 9 | CI | 103 | THR |
| 12 | CL | 19 | ARG |
| 12 | CL | 65 | GLU |
| 12 | CL | 79 | GLU |
| 14 | CN | 16 | PHE |
| 17 | CQ | 3 | LYS |
| 17 | CQ | 96 | GLU |
| 19 | CS | 43 | GLU |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 20 | CT | 97 | ALA |
| 20 | CT | 98 | PRO |
| 22 | D0 | 83 | PRO |
| 23 | D1 | 22 | GLY |
| 23 | D1 | 28 | GLY |
| 23 | D1 | 44 | PRO |
| 23 | D1 | 85 | LEU |
| 26 | D4 | 31 | ILE |
| 27 | D5 | 33 | CYS |
| 30 | D8 | 17 | THR |
| 30 | D8 | 38 | GLY |
| 33 | DD | 242 | ARG |
| 34 | DE | 2 | LYS |
| 34 | DE | 86 | PRO |
| 35 | DF | 5 | ALA |
| 35 | DF | 10 | PRO |
| 36 | DG | 43 | LEU |
| 36 | DG | 115 | ARG |
| 37 | DH | 146 | ALA |
| 39 | DN | 80 | GLY |
| 41 | DP | 76 | LYS |
| 42 | DQ | 86 | GLY |
| 43 | DR | 102 | GLU |
| 44 | DS | 35 | ILE |
| 44 | DS | 106 | ARG |
| 45 | DT | 93 | ARG |
| 47 | DV | 55 | ALA |
| 47 | DV | 73 | SER |
| 48 | DW | 11 | ARG |
| 50 | DY | 31 | LEU |
| 50 | DY | 39 | VAL |
| 50 | DY | 53 | PRO |
| 2 | AB | 130 | ARG |
| 3 | AC | 54 | ARG |
| 4 | AD | 5 | ILE |
| 5 | AE | 37 | ARG |
| 6 | AF | 16 | GLN |
| 6 | AF | 42 | GLU |
| 6 | AF | 96 | PRO |
| 7 | AG | 83 | ALA |
| 8 | AH | 8 | ASP |
| 10 | AJ | 91 | PRO |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 11 | AK | 49 | GLY |
| 12 | AL | 63 | GLY |
| 13 | AM | 6 | GLY |
| 14 | AN | 16 | PHE |
| 15 | AO | 24 | SER |
| 15 | AO | 76 | GLU |
| 17 | AQ | 61 | GLU |
| 18 | AR | 87 | ARG |
| 19 | AS | 5 | LEU |
| 19 | AS | 80 | TYR |
| 20 | AT | 73 | HIS |
| 20 | AT | 82 | SER |
| 23 | B1 | 22 | GLY |
| 23 | B1 | 38 | SER |
| 26 | B4 | 14 | ILE |
| 28 | B6 | 18 | ARG |
| 30 | B8 | 38 | GLY |
| 33 | BD | 134 | ARG |
| 35 | BF | 160 | ASN |
| 37 | BH | 92 | ILE |
| 37 | BH | 117 | PRO |
| 37 | BH | 145 | ALA |
| 38 | BI | 13 | GLY |
| 38 | BI | 51 | ILE |
| 38 | BI | 122 | GLU |
| 38 | BI | 128 | LEU |
| 39 | BN | 19 | GLU |
| 39 | BN | 80 | GLY |
| 41 | BP | 6 | LEU |
| 41 | BP | 9 | ASN |
| 41 | BP | 33 | ARG |
| 41 | BP | 38 | GLN |
| 41 | BP | 43 | GLY |
| 41 | BP | 63 | PRO |
| 41 | BP | 110 | TYR |
| 43 | BR | 42 | LYS |
| 44 | BS | 13 | ARG |
| 44 | BS | 14 | VAL |
| 44 | BS | 29 | PHE |
| 45 | BT | 55 | ASN |
| 45 | BT | 81 | PRO |
| 45 | BT | 83 | ILE |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 45 | BT | 93 | ARG |
| 46 | BU | 51 | LYS |
| 47 | BV | 3 | ALA |
| 47 | BV | 26 | ASP |
| 47 | BV | 52 | VAL |
| 47 | BV | 64 | HIS |
| 48 | BW | 56 | ALA |
| 49 | BX | 74 | PRO |
| 49 | BX | 90 | GLU |
| 51 | BZ | 142 | SER |
| 51 | BZ | 169 | GLU |
| 2 | CB | 130 | ARG |
| 3 | CC | 54 | ARG |
| 4 | CD | 5 | ILE |
| 6 | CF | 16 | GLN |
| 6 | CF | 29 | ALA |
| 6 | CF | 42 | GLU |
| 6 | CF | 96 | PRO |
| 10 | CJ | 91 | PRO |
| 11 | CK | 62 | GLN |
| 12 | CL | 89 | ARG |
| 13 | CM | 6 | GLY |
| 15 | CO | 24 | SER |
| 15 | CO | 86 | GLY |
| 16 | CP | 63 | GLY |
| 16 | CP | 67 | THR |
| 17 | CQ | 61 | GLU |
| 18 | CR | 87 | ARG |
| 19 | CS | 5 | LEU |
| 20 | CT | 82 | SER |
| 23 | D1 | 89 | GLU |
| 24 | D2 | 47 | ASN |
| 26 | D4 | 14 | ILE |
| 28 | D6 | 17 | LYS |
| 28 | D6 | 18 | ARG |
| 28 | D6 | 28 | ARG |
| 33 | DD | 238 | GLY |
| 34 | DE | 117 | MET |
| 35 | DF | 7 | TYR |
| 36 | DG | 21 | ARG |
| 37 | DH | 92 | ILE |
| 37 | DH | 145 | ALA |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 38 | DI | 5 | LEU |
| 38 | DI | 7 | GLU |
| 38 | DI | 13 | GLY |
| 39 | DN | 42 | TRP |
| 40 | DO | 68 | GLU |
| 41 | DP | 6 | LEU |
| 41 | DP | 9 | ASN |
| 41 | DP | 33 | ARG |
| 41 | DP | 39 | LYS |
| 41 | DP | 63 | PRO |
| 41 | DP | 67 | MET |
| 41 | DP | 89 | ALA |
| 41 | DP | 110 | TYR |
| 41 | DP | 127 | ALA |
| 43 | DR | 20 | LEU |
| 43 | DR | 63 | ARG |
| 44 | DS | 29 | PHE |
| 44 | DS | 53 | SER |
| 46 | DU | 51 | LYS |
| 46 | DU | 64 | ARG |
| 47 | DV | 2 | PHE |
| 47 | DV | 3 | ALA |
| 47 | DV | 64 | HIS |
| 47 | DV | 91 | TYR |
| 51 | DZ | 121 | HIS |
| 51 | DZ | 142 | SER |
| 4 | AD | 171 | GLY |
| 4 | AD | 191 | ARG |
| 5 | AE | 133 | TYR |
| 6 | AF | 12 | PRO |
| 8 | AH | 86 | ILE |
| 16 | AP | 63 | GLY |
| 18 | AR | 41 | LYS |
| 20 | AT | 101 | GLY |
| 21 | AU | 22 | ARG |
| 23 | B1 | 90 | ILE |
| 28 | B6 | 17 | LYS |
| 28 | B6 | 28 | ARG |
| 33 | BD | 244 | ARG |
| 34 | BE | 32 | PRO |
| 34 | BE | 70 | ALA |
| 38 | BI | 85 | GLU |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 39 | BN | 94 | HIS |
| 41 | BP | 76 | LYS |
| 43 | BR | 8 | ARG |
| 47 | BV | 91 | TYR |
| 48 | BW | 58 | ALA |
| 51 | BZ | 120 | ILE |
| 51 | BZ | 165 | VAL |
| 4 | CD | 17 | VAL |
| 5 | CE | 133 | TYR |
| 6 | CF | 12 | PRO |
| 7 | CG | 14 | PRO |
| 7 | CG | 83 | ALA |
| 8 | CH | 8 | ASP |
| 8 | CH | 86 | ILE |
| 11 | CK | 49 | GLY |
| 17 | CQ | 80 | GLY |
| 19 | CS | 80 | TYR |
| 20 | CT | 73 | HIS |
| 21 | CU | 22 | ARG |
| 23 | D1 | 90 | ILE |
| 23 | D1 | 93 | GLU |
| 25 | D3 | 3 | ARG |
| 26 | D4 | 9 | LEU |
| 33 | DD | 244 | ARG |
| 34 | DE | 182 | LEU |
| 37 | DH | 13 | LYS |
| 37 | DH | 117 | PRO |
| 38 | DI | 128 | LEU |
| 41 | DP | 26 | GLY |
| 41 | DP | 43 | GLY |
| 44 | DS | 13 | ARG |
| 44 | DS | 14 | VAL |
| 45 | DT | 81 | PRO |
| 47 | DV | 57 | VAL |
| 49 | DX | 90 | GLU |
| 51 | DZ | 38 | TYR |
| 51 | DZ | 147 | GLY |
| 51 | DZ | 165 | VAL |
| 51 | DZ | 177 | PRO |
| 7 | AG | 14 | PRO |
| 15 | AO | 86 | GLY |
| 33 | BD | 232 | PRO |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 35 | BF | 24 | LEU |
| 35 | BF | 82 | ILE |
| 41 | BP | 8 | PRO |
| 47 | BV | 57 | VAL |
| 50 | BY | 22 | GLY |
| 5 | CE | 128 | PRO |
| 23 | D1 | 13 | ILE |
| 29 | D7 | 17 | GLY |
| 38 | DI | 51 | ILE |
| 44 | DS | 18 | ILE |
| 44 | DS | 98 | VAL |
| 47 | DV | 52 | VAL |
| 50 | DY | 10 | GLY |
| 5 | AE | 128 | PRO |
| 16 | AP | 51 | VAL |
| 17 | AQ | 80 | GLY |
| 23 | B1 | 13 | ILE |
| 33 | BD | 238 | GLY |
| 37 | BH | 44 | VAL |
| 41 | BP | 26 | GLY |
| 41 | BP | 144 | GLU |
| 44 | BS | 18 | ILE |
| 51 | BZ | 177 | PRO |
| 4 | CD | 141 | ARG |
| 12 | CL | 63 | GLY |
| 16 | CP | 51 | VAL |
| 29 | D7 | 44 | PRO |
| 33 | DD | 232 | PRO |
| 35 | DF | 24 | LEU |
| 37 | DH | 44 | VAL |
| 39 | DN | 94 | HIS |
| 41 | DP | 144 | GLU |
| 48 | DW | 59 | VAL |
| 3 | AC | 114 | PRO |
| 51 | BZ | 141 | VAL |
| 41 | DP | 24 | GLY |
| 51 | DZ | 141 | VAL |
| 4 | AD | 141 | ARG |
| 14 | AN | 13 | THR |
| 27 | B5 | 50 | GLY |
| 35 | BF | 9 | ILE |
| 38 | BI | 106 | GLY |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 41 | BP | 24 | GLY |
| 47 | BV | 79 | VAL |
| 48 | BW | 59 | VAL |
| 34 | DE | 32 | PRO |
| 35 | DF | 9 | ILE |
| 35 | DF | 82 | ILE |
| 38 | DI | 106 | GLY |
| 41 | DP | 8 | PRO |
| 41 | DP | 23 | PRO |
| 43 | DR | 32 | GLY |
| 50 | DY | 30 | VAL |
| 4 | AD | 109 | GLY |
| 16 | AP | 65 | GLN |
| 29 | B7 | 44 | PRO |
| 39 | BN | 77 | GLY |
| 14 | CN | 13 | THR |
| 34 | DE | 56 | PRO |
| 41 | BP | 48 | PRO |

### 5.3.2 Protein sidechains (i)

In the following table, the Percentiles column shows the percent sidechain outliers of the chain as a percentile score with respect to all X-ray entries followed by that with respect to entries of similar resolution.

The Analysed column shows the number of residues for which the sidechain conformation was analysed, and the total number of residues.

| Mol | Chain | Analysed | Rotameric | Outliers | Percentiles |
|---|---|---|---|---|---|
| 2 | AB | 202/220 (92%) | 171 (85%) | 31 (15%) | 3 14 |
| 2 | CB | 202/220 (92%) | 171 (85%) | 31 (15%) | 3 14 |
| 3 | AC | 160/188 (85%) | 151 (94%) | 9 (6%) | 19 52 |
| 3 | CC | 160/188 (85%) | 151 (94%) | 9 (6%) | 19 52 |
| 4 | AD | 180/181 (99%) | 153 (85%) | 27 (15%) | 3 14 |
| 4 | CD | 180/181 (99%) | 153 (85%) | 27 (15%) | 3 14 |
| 5 | AE | 115/123 (94%) | 96 (84%) | 19 (16%) | 2 12 |
| 5 | CE | 115/123 (94%) | 97 (84%) | 18 (16%) | 2 13 |
| 6 | AF | 90/90 (100%) | 79 (88%) | 11 (12%) | 5 21 |
| 6 | CF | 90/90 (100%) | 79 (88%) | 11 (12%) | 5 21 |

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| Mol | Chain | Analysed | Rotameric | Outliers | Percentiles |
|---|---|---|---|---|---|
| 7 | AG | 126/127 (99%) | 121 (96%) | 5 (4%) | 28 62 |
| 7 | CG | 126/127 (99%) | 121 (96%) | 5 (4%) | 28 62 |
| 8 | AH | 119/119 (100%) | 104 (87%) | 15 (13%) | 4 20 |
| 8 | CH | 119/119 (100%) | 104 (87%) | 15 (13%) | 4 20 |
| 9 | AI | 98/99 (99%) | 89 (91%) | 9 (9%) | 8 33 |
| 9 | CI | 98/99 (99%) | 89 (91%) | 9 (9%) | 8 33 |
| 10 | AJ | 88/92 (96%) | 83 (94%) | 5 (6%) | 18 52 |
| 10 | CJ | 88/92 (96%) | 83 (94%) | 5 (6%) | 18 52 |
| 11 | AK | 90/99 (91%) | 79 (88%) | 11 (12%) | 5 21 |
| 11 | CK | 90/99 (91%) | 78 (87%) | 12 (13%) | 4 18 |
| 12 | AL | 104/111 (94%) | 94 (90%) | 10 (10%) | 8 31 |
| 12 | CL | 104/111 (94%) | 94 (90%) | 10 (10%) | 8 31 |
| 13 | AM | 93/101 (92%) | 88 (95%) | 5 (5%) | 20 53 |
| 13 | CM | 93/101 (92%) | 88 (95%) | 5 (5%) | 20 53 |
| 14 | AN | 49/50 (98%) | 46 (94%) | 3 (6%) | 17 49 |
| 14 | CN | 49/50 (98%) | 46 (94%) | 3 (6%) | 17 49 |
| 15 | AO | 79/80 (99%) | 71 (90%) | 8 (10%) | 7 29 |
| 15 | CO | 79/80 (99%) | 70 (89%) | 9 (11%) | 5 24 |
| 16 | AP | 72/74 (97%) | 61 (85%) | 11 (15%) | 3 14 |
| 16 | CP | 72/74 (97%) | 61 (85%) | 11 (15%) | 3 14 |
| 17 | AQ | 94/97 (97%) | 89 (95%) | 5 (5%) | 20 54 |
| 17 | CQ | 94/97 (97%) | 89 (95%) | 5 (5%) | 20 54 |
| 18 | AR | 61/77 (79%) | 51 (84%) | 10 (16%) | 2 12 |
| 18 | CR | 61/77 (79%) | 51 (84%) | 10 (16%) | 2 12 |
| 19 | AS | 69/80 (86%) | 63 (91%) | 6 (9%) | 9 35 |
| 19 | CS | 69/80 (86%) | 63 (91%) | 6 (9%) | 9 35 |
| 20 | AT | 76/82 (93%) | 65 (86%) | 11 (14%) | 3 15 |
| 20 | CT | 76/82 (93%) | 64 (84%) | 12 (16%) | 2 13 |
| 21 | AU | 19/22 (86%) | 19 (100%) | 0 | 100 100 |
| 21 | CU | 19/22 (86%) | 19 (100%) | 0 | 100 100 |
| 22 | B0 | 61/67 (91%) | 47 (77%) | 14 (23%) | 1 5 |

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| Mol | Chain | Analysed | Rotameric | Outliers | Percentiles |
|---|---|---|---|---|---|
| 22 | D0 | 61/67 (91%) | 47 (77%) | 14 (23%) | 1 5 |
| 23 | B1 | 73/83 (88%) | 54 (74%) | 19 (26%) | 0 3 |
| 23 | D1 | 73/83 (88%) | 56 (77%) | 17 (23%) | 1 4 |
| 24 | B2 | 46/67 (69%) | 34 (74%) | 12 (26%) | 0 3 |
| 24 | D2 | 46/67 (69%) | 34 (74%) | 12 (26%) | 0 3 |
| 25 | B3 | 51/52 (98%) | 42 (82%) | 9 (18%) | 2 10 |
| 25 | D3 | 51/52 (98%) | 43 (84%) | 8 (16%) | 2 13 |
| 27 | B5 | 51/52 (98%) | 37 (72%) | 14 (28%) | 0 2 |
| 27 | D5 | 51/52 (98%) | 38 (74%) | 13 (26%) | 0 3 |
| 28 | B6 | 43/52 (83%) | 29 (67%) | 14 (33%) | 0 1 |
| 28 | D6 | 43/52 (83%) | 29 (67%) | 14 (33%) | 0 1 |
| 29 | B7 | 41/42 (98%) | 33 (80%) | 8 (20%) | 1 8 |
| 29 | D7 | 41/42 (98%) | 33 (80%) | 8 (20%) | 1 8 |
| 30 | B8 | 53/55 (96%) | 37 (70%) | 16 (30%) | 0 1 |
| 30 | D8 | 53/55 (96%) | 38 (72%) | 15 (28%) | 0 2 |
| 33 | BD | 213/218 (98%) | 150 (70%) | 63 (30%) | 0 2 |
| 33 | DD | 213/218 (98%) | 151 (71%) | 62 (29%) | 0 2 |
| 34 | BE | 165/166 (99%) | 124 (75%) | 41 (25%) | 1 3 |
| 34 | DE | 165/166 (99%) | 127 (77%) | 38 (23%) | 1 5 |
| 35 | BF | 165/166 (99%) | 128 (78%) | 37 (22%) | 1 5 |
| 35 | DF | 165/166 (99%) | 126 (76%) | 39 (24%) | 1 4 |
| 36 | BG | 155/156 (99%) | 126 (81%) | 29 (19%) | 1 9 |
| 36 | DG | 155/156 (99%) | 126 (81%) | 29 (19%) | 1 9 |
| 37 | BH | 132/148 (89%) | 106 (80%) | 26 (20%) | 1 8 |
| 37 | DH | 132/148 (89%) | 107 (81%) | 25 (19%) | 1 9 |
| 38 | BI | 122/124 (98%) | 98 (80%) | 24 (20%) | 1 8 |
| 38 | DI | 122/124 (98%) | 98 (80%) | 24 (20%) | 1 8 |
| 39 | BN | 117/119 (98%) | 88 (75%) | 29 (25%) | 1 3 |
| 39 | DN | 117/119 (98%) | 86 (74%) | 31 (26%) | 0 3 |
| 40 | BO | 100/100 (100%) | 71 (71%) | 29 (29%) | 0 2 |
| 40 | DO | 100/100 (100%) | 71 (71%) | 29 (29%) | 0 2 |

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| Mol | Chain | Analysed | Rotameric | Outliers | Percentiles |
|---|---|---|---|---|---|
| 41 | BP | 112/116 (97%) | 65 (58%) | 47 (42%) | 0 0 |
| 41 | DP | 112/116 (97%) | 66 (59%) | 46 (41%) | 0 0 |
| 42 | BQ | 106/111 (96%) | 82 (77%) | 24 (23%) | 1 5 |
| 42 | DQ | 106/111 (96%) | 83 (78%) | 23 (22%) | 1 6 |
| 43 | BR | 100/101 (99%) | 75 (75%) | 25 (25%) | 0 3 |
| 43 | DR | 100/101 (99%) | 75 (75%) | 25 (25%) | 0 3 |
| 44 | BS | 77/88 (88%) | 53 (69%) | 24 (31%) | 0 1 |
| 44 | DS | 77/88 (88%) | 55 (71%) | 22 (29%) | 0 2 |
| 45 | BT | 116/127 (91%) | 83 (72%) | 33 (28%) | 0 2 |
| 45 | DT | 116/127 (91%) | 83 (72%) | 33 (28%) | 0 2 |
| 46 | BU | 92/94 (98%) | 72 (78%) | 20 (22%) | 1 6 |
| 46 | DU | 92/94 (98%) | 71 (77%) | 21 (23%) | 1 5 |
| 47 | BV | 82/82 (100%) | 51 (62%) | 31 (38%) | 0 1 |
| 47 | DV | 82/82 (100%) | 50 (61%) | 32 (39%) | 0 1 |
| 48 | BW | 91/92 (99%) | 67 (74%) | 24 (26%) | 0 3 |
| 48 | DW | 91/92 (99%) | 67 (74%) | 24 (26%) | 0 3 |
| 49 | BX | 74/78 (95%) | 53 (72%) | 21 (28%) | 0 2 |
| 49 | DX | 74/78 (95%) | 53 (72%) | 21 (28%) | 0 2 |
| 50 | BY | 84/91 (92%) | 62 (74%) | 22 (26%) | 0 3 |
| 50 | DY | 84/91 (92%) | 63 (75%) | 21 (25%) | 0 3 |
| 51 | BZ | 155/179 (87%) | 128 (83%) | 27 (17%) | 2 10 |
| 51 | DZ | 155/179 (87%) | 128 (83%) | 27 (17%) | 2 10 |
| All | All | 9322/9876 (94%) | 7543 (81%) | 1779 (19%) | 1 8 |

All (1779) residues with a non-rotameric sidechain are listed below:

| Mol | Chain | Res | Type |
|---|---|---|---|
| 2 | AB | 9 | GLU |
| 2 | AB | 12 | GLU |
| 2 | AB | 15 | VAL |
| 2 | AB | 17 | PHE |
| 2 | AB | 22 | LYS |
| 2 | AB | 36 | ARG |
| 2 | AB | 42 | ILE |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 2 | AB | 44 | LEU |
| 2 | AB | 51 | LEU |
| 2 | AB | 59 | GLU |
| 2 | AB | 69 | LEU |
| 2 | AB | 71 | VAL |
| 2 | AB | 75 | LYS |
| 2 | AB | 79 | ASP |
| 2 | AB | 90 | MET |
| 2 | AB | 111 | ARG |
| 2 | AB | 130 | ARG |
| 2 | AB | 137 | ARG |
| 2 | AB | 145 | LEU |
| 2 | AB | 146 | GLN |
| 2 | AB | 154 | LEU |
| 2 | AB | 158 | LEU |
| 2 | AB | 179 | LYS |
| 2 | AB | 185 | ILE |
| 2 | AB | 187 | LEU |
| 2 | AB | 189 | ASP |
| 2 | AB | 196 | LEU |
| 2 | AB | 204 | ASN |
| 2 | AB | 205 | ASP |
| 2 | AB | 215 | LEU |
| 2 | AB | 221 | LEU |
| 3 | AC | 5 | ILE |
| 3 | AC | 12 | LEU |
| 3 | AC | 27 | LYS |
| 3 | AC | 69 | HIS |
| 3 | AC | 76 | VAL |
| 3 | AC | 104 | GLN |
| 3 | AC | 131 | ARG |
| 3 | AC | 138 | VAL |
| 3 | AC | 198 | VAL |
| 4 | AD | 3 | ARG |
| 4 | AD | 8 | VAL |
| 4 | AD | 9 | CYS |
| 4 | AD | 11 | LEU |
| 4 | AD | 12 | CYS |
| 4 | AD | 15 | GLU |
| 4 | AD | 19 | LEU |
| 4 | AD | 25 | ARG |
| 4 | AD | 36 | ARG |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 4 | AD | 58 | LEU |
| 4 | AD | 62 | GLN |
| 4 | AD | 68 | TYR |
| 4 | AD | 70 | ILE |
| 4 | AD | 94 | LEU |
| 4 | AD | 97 | LEU |
| 4 | AD | 104 | VAL |
| 4 | AD | 121 | VAL |
| 4 | AD | 127 | THR |
| 4 | AD | 129 | ASN |
| 4 | AD | 132 | ARG |
| 4 | AD | 135 | LEU |
| 4 | AD | 138 | TYR |
| 4 | AD | 158 | ILE |
| 4 | AD | 194 | LEU |
| 4 | AD | 196 | LEU |
| 4 | AD | 202 | LEU |
| 4 | AD | 203 | VAL |
| 5 | AE | 10 | MET |
| 5 | AE | 12 | LEU |
| 5 | AE | 13 | ILE |
| 5 | AE | 25 | ARG |
| 5 | AE | 31 | LEU |
| 5 | AE | 41 | VAL |
| 5 | AE | 51 | VAL |
| 5 | AE | 55 | VAL |
| 5 | AE | 73 | ASN |
| 5 | AE | 76 | ILE |
| 5 | AE | 78 | HIS |
| 5 | AE | 79 | GLU |
| 5 | AE | 82 | VAL |
| 5 | AE | 91 | LEU |
| 5 | AE | 98 | THR |
| 5 | AE | 101 | ILE |
| 5 | AE | 112 | LEU |
| 5 | AE | 115 | VAL |
| 5 | AE | 147 | ASP |
| 6 | AF | 10 | LEU |
| 6 | AF | 21 | LEU |
| 6 | AF | 30 | LEU |
| 6 | AF | 45 | LEU |
| 6 | AF | 46 | ARG |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 6 | AF | 52 | ILE |
| 6 | AF | 55 | ASP |
| 6 | AF | 63 | TYR |
| 6 | AF | 70 | ASP |
| 6 | AF | 92 | LYS |
| 6 | AF | 98 | LEU |
| 7 | AG | 27 | ILE |
| 7 | AG | 36 | LYS |
| 7 | AG | 79 | ARG |
| 7 | AG | 136 | LYS |
| 7 | AG | 151 | TYR |
| 8 | AH | 1 | MET |
| 8 | AH | 3 | THR |
| 8 | AH | 10 | LEU |
| 8 | AH | 24 | THR |
| 8 | AH | 39 | LEU |
| 8 | AH | 51 | VAL |
| 8 | AH | 63 | LEU |
| 8 | AH | 65 | TYR |
| 8 | AH | 81 | HIS |
| 8 | AH | 83 | ILE |
| 8 | AH | 91 | ARG |
| 8 | AH | 95 | VAL |
| 8 | AH | 102 | ARG |
| 8 | AH | 133 | LEU |
| 8 | AH | 137 | VAL |
| 9 | AI | 3 | GLN |
| 9 | AI | 10 | ARG |
| 9 | AI | 78 | LYS |
| 9 | AI | 95 | LYS |
| 9 | AI | 99 | LEU |
| 9 | AI | 101 | PHE |
| 9 | AI | 102 | LEU |
| 9 | AI | 114 | TYR |
| 9 | AI | 128 | ARG |
| 10 | AJ | 5 | ARG |
| 10 | AJ | 74 | ILE |
| 10 | AJ | 80 | LYS |
| 10 | AJ | 96 | ILE |
| 10 | AJ | 100 | THR |
| 11 | AK | 18 | ARG |
| 11 | AK | 22 | HIS |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 11 | AK | 29 | ILE |
| 11 | AK | 53 | SER |
| 11 | AK | 63 | LEU |
| 11 | AK | 92 | GLU |
| 11 | AK | 95 | ILE |
| 11 | AK | 114 | VAL |
| 11 | AK | 117 | ASN |
| 11 | AK | 125 | PHE |
| 11 | AK | 127 | LYS |
| 12 | AL | 24 | VAL |
| 12 | AL | 38 | THR |
| 12 | AL | 41 | ARG |
| 12 | AL | 42 | THR |
| 12 | AL | 62 | SER |
| 12 | AL | 75 | HIS |
| 12 | AL | 85 | ILE |
| 12 | AL | 89 | ARG |
| 12 | AL | 96 | VAL |
| 12 | AL | 102 | ARG |
| 13 | AM | 47 | ASP |
| 13 | AM | 66 | LEU |
| 13 | AM | 70 | LEU |
| 13 | AM | 93 | ARG |
| 13 | AM | 108 | ARG |
| 14 | AN | 18 | VAL |
| 14 | AN | 42 | ILE |
| 14 | AN | 44 | LEU |
| 15 | AO | 3 | ILE |
| 15 | AO | 26 | GLU |
| 15 | AO | 39 | LEU |
| 15 | AO | 41 | GLU |
| 15 | AO | 57 | LEU |
| 15 | AO | 65 | ARG |
| 15 | AO | 67 | LEU |
| 15 | AO | 82 | ILE |
| 16 | AP | 2 | VAL |
| 16 | AP | 27 | LYS |
| 16 | AP | 28 | ARG |
| 16 | AP | 47 | ASP |
| 16 | AP | 53 | VAL |
| 16 | AP | 65 | GLN |
| 16 | AP | 67 | THR |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 16 | AP | 69 | THR |
| 16 | AP | 74 | LEU |
| 16 | AP | 79 | VAL |
| 16 | AP | 82 | GLN |
| 17 | AQ | 5 | VAL |
| 17 | AQ | 52 | LYS |
| 17 | AQ | 63 | ARG |
| 17 | AQ | 74 | LEU |
| 17 | AQ | 89 | LEU |
| 18 | AR | 31 | LEU |
| 18 | AR | 47 | THR |
| 18 | AR | 53 | ARG |
| 18 | AR | 65 | ILE |
| 18 | AR | 70 | ILE |
| 18 | AR | 74 | ARG |
| 18 | AR | 76 | LEU |
| 18 | AR | 79 | LEU |
| 18 | AR | 82 | THR |
| 18 | AR | 87 | ARG |
| 19 | AS | 6 | LYS |
| 19 | AS | 7 | LYS |
| 19 | AS | 22 | LEU |
| 19 | AS | 31 | ILE |
| 19 | AS | 37 | ARG |
| 19 | AS | 49 | ILE |
| 20 | AT | 26 | ASN |
| 20 | AT | 27 | LYS |
| 20 | AT | 30 | LYS |
| 20 | AT | 35 | THR |
| 20 | AT | 41 | ILE |
| 20 | AT | 51 | GLU |
| 20 | AT | 56 | MET |
| 20 | AT | 57 | ARG |
| 20 | AT | 72 | LEU |
| 20 | AT | 73 | HIS |
| 20 | AT | 93 | GLU |
| 22 | B0 | 11 | ARG |
| 22 | B0 | 14 | ARG |
| 22 | B0 | 30 | VAL |
| 22 | B0 | 31 | VAL |
| 22 | B0 | 36 | ILE |
| 22 | B0 | 40 | GLN |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 22 | B0 | 41 | ARG |
| 22 | B0 | 53 | MET |
| 22 | B0 | 55 | ARG |
| 22 | B0 | 62 | LEU |
| 22 | B0 | 70 | GLN |
| 22 | B0 | 72 | ARG |
| 22 | B0 | 79 | VAL |
| 22 | B0 | 84 | LEU |
| 23 | B1 | 8 | SER |
| 23 | B1 | 13 | ILE |
| 23 | B1 | 14 | VAL |
| 23 | B1 | 25 | LYS |
| 23 | B1 | 26 | ARG |
| 23 | B1 | 34 | THR |
| 23 | B1 | 37 | ILE |
| 23 | B1 | 41 | ARG |
| 23 | B1 | 46 | LEU |
| 23 | B1 | 47 | GLN |
| 23 | B1 | 48 | LYS |
| 23 | B1 | 51 | VAL |
| 23 | B1 | 56 | GLN |
| 23 | B1 | 58 | ILE |
| 23 | B1 | 65 | SER |
| 23 | B1 | 67 | ILE |
| 23 | B1 | 74 | VAL |
| 23 | B1 | 85 | LEU |
| 23 | B1 | 89 | GLU |
| 24 | B2 | 12 | GLU |
| 24 | B2 | 14 | ARG |
| 24 | B2 | 22 | GLU |
| 24 | B2 | 28 | LYS |
| 24 | B2 | 30 | ARG |
| 24 | B2 | 33 | MET |
| 24 | B2 | 44 | LEU |
| 24 | B2 | 45 | SER |
| 24 | B2 | 46 | GLN |
| 24 | B2 | 47 | ASN |
| 24 | B2 | 50 | ILE |
| 24 | B2 | 56 | GLN |
| 25 | B3 | 8 | LEU |
| 25 | B3 | 9 | VAL |
| 25 | B3 | 10 | LYS |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 25 | B3 | 18 | ASP |
| 25 | B3 | 24 | LYS |
| 25 | B3 | 38 | GLU |
| 25 | B3 | 40 | THR |
| 25 | B3 | 47 | VAL |
| 25 | B3 | 59 | VAL |
| 27 | B5 | 4 | HIS |
| 27 | B5 | 9 | LYS |
| 27 | B5 | 11 | THR |
| 27 | B5 | 16 | ARG |
| 27 | B5 | 29 | THR |
| 27 | B5 | 32 | PRO |
| 27 | B5 | 36 | CYS |
| 27 | B5 | 44 | THR |
| 27 | B5 | 45 | VAL |
| 27 | B5 | 49 | CYS |
| 27 | B5 | 51 | TYR |
| 27 | B5 | 55 | ARG |
| 27 | B5 | 56 | LYS |
| 27 | B5 | 58 | LEU |
| 28 | B6 | 9 | LEU |
| 28 | B6 | 10 | LEU |
| 28 | B6 | 11 | LEU |
| 28 | B6 | 12 | GLU |
| 28 | B6 | 18 | ARG |
| 28 | B6 | 25 | LYS |
| 28 | B6 | 27 | LYS |
| 28 | B6 | 30 | THR |
| 28 | B6 | 37 | ARG |
| 28 | B6 | 42 | TRP |
| 28 | B6 | 46 | HIS |
| 28 | B6 | 47 | THR |
| 28 | B6 | 48 | VAL |
| 28 | B6 | 51 | GLU |
| 29 | B7 | 1 | MET |
| 29 | B7 | 8 | ASN |
| 29 | B7 | 9 | ARG |
| 29 | B7 | 15 | THR |
| 29 | B7 | 24 | THR |
| 29 | B7 | 42 | LEU |
| 29 | B7 | 43 | THR |
| 29 | B7 | 48 | LYS |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 30 | B8 | 4 | MET |
| 30 | B8 | 6 | THR |
| 30 | B8 | 13 | ARG |
| 30 | B8 | 16 | ILE |
| 30 | B8 | 32 | LEU |
| 30 | B8 | 37 | SER |
| 30 | B8 | 39 | LYS |
| 30 | B8 | 41 | ILE |
| 30 | B8 | 44 | LYS |
| 30 | B8 | 47 | LYS |
| 30 | B8 | 48 | PHE |
| 30 | B8 | 49 | VAL |
| 30 | B8 | 54 | GLU |
| 30 | B8 | 56 | GLU |
| 30 | B8 | 58 | ILE |
| 30 | B8 | 59 | LYS |
| 33 | BD | 3 | VAL |
| 33 | BD | 5 | LYS |
| 33 | BD | 10 | THR |
| 33 | BD | 17 | THR |
| 33 | BD | 24 | ILE |
| 33 | BD | 26 | LYS |
| 33 | BD | 27 | THR |
| 33 | BD | 31 | LYS |
| 33 | BD | 33 | LEU |
| 33 | BD | 34 | VAL |
| 33 | BD | 44 | ASN |
| 33 | BD | 46 | GLN |
| 33 | BD | 49 | ILE |
| 33 | BD | 59 | LYS |
| 33 | BD | 61 | LEU |
| 33 | BD | 63 | ARG |
| 33 | BD | 64 | ILE |
| 33 | BD | 65 | ILE |
| 33 | BD | 71 | ASP |
| 33 | BD | 72 | LYS |
| 33 | BD | 75 | ILE |
| 33 | BD | 87 | ASN |
| 33 | BD | 88 | ARG |
| 33 | BD | 89 | SER |
| 33 | BD | 92 | ILE |
| 33 | BD | 94 | LEU |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 33 | BD | 95 | LEU |
| 33 | BD | 98 | VAL |
| 33 | BD | 103 | ARG |
| 33 | BD | 105 | ILE |
| 33 | BD | 106 | ILE |
| 33 | BD | 111 | LEU |
| 33 | BD | 113 | VAL |
| 33 | BD | 117 | VAL |
| 33 | BD | 127 | VAL |
| 33 | BD | 131 | LEU |
| 33 | BD | 138 | VAL |
| 33 | BD | 147 | LEU |
| 33 | BD | 150 | LYS |
| 33 | BD | 157 | ARG |
| 33 | BD | 161 | THR |
| 33 | BD | 162 | SER |
| 33 | BD | 165 | ILE |
| 33 | BD | 166 | GLN |
| 33 | BD | 173 | VAL |
| 33 | BD | 174 | ILE |
| 33 | BD | 176 | ARG |
| 33 | BD | 182 | LEU |
| 33 | BD | 183 | ARG |
| 33 | BD | 192 | THR |
| 33 | BD | 193 | VAL |
| 33 | BD | 202 | LYS |
| 33 | BD | 212 | SER |
| 33 | BD | 217 | ARG |
| 33 | BD | 221 | VAL |
| 33 | BD | 229 | VAL |
| 33 | BD | 242 | ARG |
| 33 | BD | 255 | LYS |
| 33 | BD | 257 | LEU |
| 33 | BD | 259 | THR |
| 33 | BD | 266 | SER |
| 33 | BD | 270 | ILE |
| 33 | BD | 271 | ILE |
| 34 | BE | 1 | MET |
| 34 | BE | 7 | VAL |
| 34 | BE | 9 | VAL |
| 34 | BE | 12 | THR |
| 34 | BE | 19 | ARG |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 34 | BE | 23 | VAL |
| 34 | BE | 24 | THR |
| 34 | BE | 26 | ILE |
| 34 | BE | 33 | VAL |
| 34 | BE | 34 | VAL |
| 34 | BE | 49 | LEU |
| 34 | BE | 60 | ASN |
| 34 | BE | 63 | LEU |
| 34 | BE | 64 | LYS |
| 34 | BE | 69 | LYS |
| 34 | BE | 75 | VAL |
| 34 | BE | 76 | ARG |
| 34 | BE | 77 | ILE |
| 34 | BE | 78 | LEU |
| 34 | BE | 79 | ARG |
| 34 | BE | 82 | ARG |
| 34 | BE | 111 | ARG |
| 34 | BE | 116 | VAL |
| 34 | BE | 118 | LYS |
| 34 | BE | 119 | ARG |
| 34 | BE | 134 | ILE |
| 34 | BE | 136 | ARG |
| 34 | BE | 144 | ARG |
| 34 | BE | 145 | LYS |
| 34 | BE | 150 | VAL |
| 34 | BE | 154 | LYS |
| 34 | BE | 159 | HIS |
| 34 | BE | 175 | VAL |
| 34 | BE | 181 | LEU |
| 34 | BE | 182 | LEU |
| 34 | BE | 184 | VAL |
| 34 | BE | 188 | VAL |
| 34 | BE | 196 | VAL |
| 34 | BE | 197 | ILE |
| 34 | BE | 202 | LYS |
| 34 | BE | 203 | LYS |
| 35 | BF | 13 | SER |
| 35 | BF | 20 | LEU |
| 35 | BF | 23 | ASP |
| 35 | BF | 28 | ILE |
| 35 | BF | 33 | LEU |
| 35 | BF | 38 | ARG |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 35 | BF | 46 | ARG |
| 35 | BF | 57 | VAL |
| 35 | BF | 60 | SER |
| 35 | BF | 66 | PRO |
| 35 | BF | 67 | GLN |
| 35 | BF | 72 | ARG |
| 35 | BF | 74 | ARG |
| 35 | BF | 78 | ILE |
| 35 | BF | 82 | ILE |
| 35 | BF | 83 | PHE |
| 35 | BF | 84 | VAL |
| 35 | BF | 88 | VAL |
| 35 | BF | 106 | ARG |
| 35 | BF | 107 | LYS |
| 35 | BF | 110 | LEU |
| 35 | BF | 112 | MET |
| 35 | BF | 124 | LEU |
| 35 | BF | 127 | GLU |
| 35 | BF | 140 | LEU |
| 35 | BF | 162 | LEU |
| 35 | BF | 164 | ARG |
| 35 | BF | 165 | ARG |
| 35 | BF | 168 | ARG |
| 35 | BF | 170 | LEU |
| 35 | BF | 175 | THR |
| 35 | BF | 183 | VAL |
| 35 | BF | 191 | ARG |
| 35 | BF | 192 | LEU |
| 35 | BF | 196 | LEU |
| 35 | BF | 197 | ASP |
| 35 | BF | 206 | ILE |
| 36 | BG | 5 | VAL |
| 36 | BG | 7 | LEU |
| 36 | BG | 16 | ARG |
| 36 | BG | 21 | ARG |
| 36 | BG | 22 | ARG |
| 36 | BG | 28 | VAL |
| 36 | BG | 31 | VAL |
| 36 | BG | 35 | GLU |
| 36 | BG | 39 | ILE |
| 36 | BG | 45 | GLU |
| 36 | BG | 49 | ASP |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 36 | BG | 60 | LEU |
| 36 | BG | 63 | ILE |
| 36 | BG | 67 | LYS |
| 36 | BG | 74 | LYS |
| 36 | BG | 83 | ARG |
| 36 | BG | 86 | MET |
| 36 | BG | 104 | GLU |
| 36 | BG | 115 | ARG |
| 36 | BG | 125 | PHE |
| 36 | BG | 135 | LEU |
| 36 | BG | 139 | LEU |
| 36 | BG | 143 | GLU |
| 36 | BG | 155 | MET |
| 36 | BG | 156 | ASP |
| 36 | BG | 159 | VAL |
| 36 | BG | 161 | THR |
| 36 | BG | 166 | ASP |
| 36 | BG | 173 | LEU |
| 37 | BH | 13 | LYS |
| 37 | BH | 15 | VAL |
| 37 | BH | 21 | PRO |
| 37 | BH | 27 | LYS |
| 37 | BH | 34 | GLU |
| 37 | BH | 41 | MET |
| 37 | BH | 46 | GLU |
| 37 | BH | 53 | GLU |
| 37 | BH | 70 | THR |
| 37 | BH | 71 | LEU |
| 37 | BH | 87 | LEU |
| 37 | BH | 89 | ILE |
| 37 | BH | 92 | ILE |
| 37 | BH | 98 | LEU |
| 37 | BH | 103 | LEU |
| 37 | BH | 105 | LEU |
| 37 | BH | 122 | THR |
| 37 | BH | 134 | SER |
| 37 | BH | 136 | ILE |
| 37 | BH | 143 | GLN |
| 37 | BH | 149 | ARG |
| 37 | BH | 152 | ARG |
| 37 | BH | 153 | LYS |
| 37 | BH | 157 | TYR |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 37 | BH | 159 | GLU |
| 37 | BH | 170 | ARG |
| 38 | BI | 1 | MET |
| 38 | BI | 7 | GLU |
| 38 | BI | 15 | VAL |
| 38 | BI | 18 | VAL |
| 38 | BI | 20 | ASP |
| 38 | BI | 22 | LYS |
| 38 | BI | 31 | LEU |
| 38 | BI | 37 | VAL |
| 38 | BI | 51 | ILE |
| 38 | BI | 52 | ARG |
| 38 | BI | 56 | LYS |
| 38 | BI | 58 | LEU |
| 38 | BI | 64 | GLU |
| 38 | BI | 71 | ILE |
| 38 | BI | 72 | LEU |
| 38 | BI | 79 | ILE |
| 38 | BI | 81 | VAL |
| 38 | BI | 101 | LEU |
| 38 | BI | 109 | ILE |
| 38 | BI | 122 | GLU |
| 38 | BI | 136 | VAL |
| 38 | BI | 138 | ILE |
| 38 | BI | 142 | VAL |
| 38 | BI | 144 | VAL |
| 39 | BN | 5 | VAL |
| 39 | BN | 7 | LYS |
| 39 | BN | 9 | VAL |
| 39 | BN | 14 | VAL |
| 39 | BN | 16 | ILE |
| 39 | BN | 19 | GLU |
| 39 | BN | 23 | LEU |
| 39 | BN | 25 | ARG |
| 39 | BN | 33 | LEU |
| 39 | BN | 34 | LEU |
| 39 | BN | 43 | THR |
| 39 | BN | 45 | ASN |
| 39 | BN | 46 | VAL |
| 39 | BN | 48 | MET |
| 39 | BN | 55 | VAL |
| 39 | BN | 60 | ILE |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 39 | BN | 63 | THR |
| 39 | BN | 65 | LYS |
| 39 | BN | 66 | LYS |
| 39 | BN | 75 | TYR |
| 39 | BN | 78 | TYR |
| 39 | BN | 82 | LEU |
| 39 | BN | 87 | LEU |
| 39 | BN | 94 | HIS |
| 39 | BN | 99 | LEU |
| 39 | BN | 104 | LYS |
| 39 | BN | 120 | LEU |
| 39 | BN | 121 | LYS |
| 39 | BN | 131 | GLN |
| 40 | BO | 2 | ILE |
| 40 | BO | 3 | GLN |
| 40 | BO | 10 | VAL |
| 40 | BO | 19 | ILE |
| 40 | BO | 22 | ILE |
| 40 | BO | 23 | ARG |
| 40 | BO | 24 | VAL |
| 40 | BO | 28 | SER |
| 40 | BO | 32 | TYR |
| 40 | BO | 35 | VAL |
| 40 | BO | 42 | SER |
| 40 | BO | 47 | ILE |
| 40 | BO | 58 | VAL |
| 40 | BO | 63 | VAL |
| 40 | BO | 65 | THR |
| 40 | BO | 75 | SER |
| 40 | BO | 78 | ARG |
| 40 | BO | 80 | ASP |
| 40 | BO | 85 | VAL |
| 40 | BO | 87 | ILE |
| 40 | BO | 88 | ASN |
| 40 | BO | 89 | ASN |
| 40 | BO | 91 | LEU |
| 40 | BO | 94 | ARG |
| 40 | BO | 102 | VAL |
| 40 | BO | 108 | GLU |
| 40 | BO | 112 | MET |
| 40 | BO | 114 | ILE |
| 40 | BO | 121 | VAL |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 41 | BP | 9 | ASN |
| 41 | BP | 13 | ASN |
| 41 | BP | 16 | ARG |
| 41 | BP | 17 | LYS |
| 41 | BP | 18 | ARG |
| 41 | BP | 19 | VAL |
| 41 | BP | 29 | LYS |
| 41 | BP | 32 | THR |
| 41 | BP | 36 | LYS |
| 41 | BP | 39 | LYS |
| 41 | BP | 40 | SER |
| 41 | BP | 45 | LEU |
| 41 | BP | 47 | ASP |
| 41 | BP | 51 | PHE |
| 41 | BP | 52 | GLU |
| 41 | BP | 57 | THR |
| 41 | BP | 59 | LEU |
| 41 | BP | 60 | MET |
| 41 | BP | 61 | ARG |
| 41 | BP | 62 | LEU |
| 41 | BP | 64 | LYS |
| 41 | BP | 67 | MET |
| 41 | BP | 71 | VAL |
| 41 | BP | 75 | ILE |
| 41 | BP | 79 | ARG |
| 41 | BP | 81 | GLN |
| 41 | BP | 83 | VAL |
| 41 | BP | 85 | LEU |
| 41 | BP | 88 | LEU |
| 41 | BP | 95 | VAL |
| 41 | BP | 96 | THR |
| 41 | BP | 98 | GLU |
| 41 | BP | 100 | LEU |
| 41 | BP | 101 | VAL |
| 41 | BP | 102 | ARG |
| 41 | BP | 111 | ARG |
| 41 | BP | 112 | LEU |
| 41 | BP | 114 | ILE |
| 41 | BP | 115 | LEU |
| 41 | BP | 123 | LEU |
| 41 | BP | 125 | VAL |
| 41 | BP | 126 | VAL |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 41 | BP | 135 | LEU |
| 41 | BP | 138 | LEU |
| 41 | BP | 144 | GLU |
| 41 | BP | 147 | LEU |
| 41 | BP | 148 | LEU |
| 42 | BQ | 6 | ARG |
| 42 | BQ | 7 | MET |
| 42 | BQ | 9 | TYR |
| 42 | BQ | 13 | GLN |
| 42 | BQ | 17 | LEU |
| 42 | BQ | 22 | LYS |
| 42 | BQ | 25 | ASP |
| 42 | BQ | 35 | VAL |
| 42 | BQ | 42 | ILE |
| 42 | BQ | 45 | GLN |
| 42 | BQ | 52 | VAL |
| 42 | BQ | 54 | MET |
| 42 | BQ | 55 | VAL |
| 42 | BQ | 63 | LYS |
| 42 | BQ | 68 | ILE |
| 42 | BQ | 81 | VAL |
| 42 | BQ | 82 | ARG |
| 42 | BQ | 91 | GLU |
| 42 | BQ | 102 | VAL |
| 42 | BQ | 109 | VAL |
| 42 | BQ | 110 | THR |
| 42 | BQ | 115 | MET |
| 42 | BQ | 127 | ILE |
| 42 | BQ | 141 | GLN |
| 43 | BR | 2 | ARG |
| 43 | BR | 4 | LEU |
| 43 | BR | 5 | LYS |
| 43 | BR | 18 | LEU |
| 43 | BR | 28 | LEU |
| 43 | BR | 29 | LEU |
| 43 | BR | 34 | ILE |
| 43 | BR | 44 | LEU |
| 43 | BR | 45 | ARG |
| 43 | BR | 54 | LEU |
| 43 | BR | 56 | LYS |
| 43 | BR | 60 | LEU |
| 43 | BR | 64 | ARG |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 43 | BR | 65 | LEU |
| 43 | BR | 67 | LEU |
| 43 | BR | 71 | GLN |
| 43 | BR | 75 | LEU |
| 43 | BR | 79 | LEU |
| 43 | BR | 99 | LYS |
| 43 | BR | 100 | LEU |
| 43 | BR | 104 | ARG |
| 43 | BR | 113 | LEU |
| 43 | BR | 115 | GLU |
| 43 | BR | 116 | LEU |
| 43 | BR | 117 | VAL |
| 44 | BS | 11 | LYS |
| 44 | BS | 12 | PHE |
| 44 | BS | 14 | VAL |
| 44 | BS | 17 | ARG |
| 44 | BS | 18 | ILE |
| 44 | BS | 20 | ARG |
| 44 | BS | 24 | LEU |
| 44 | BS | 32 | LEU |
| 44 | BS | 33 | LYS |
| 44 | BS | 34 | HIS |
| 44 | BS | 36 | TYR |
| 44 | BS | 46 | VAL |
| 44 | BS | 50 | SER |
| 44 | BS | 56 | LEU |
| 44 | BS | 61 | ASN |
| 44 | BS | 69 | VAL |
| 44 | BS | 73 | LEU |
| 44 | BS | 80 | LEU |
| 44 | BS | 83 | LYS |
| 44 | BS | 84 | GLN |
| 44 | BS | 85 | VAL |
| 44 | BS | 92 | TYR |
| 44 | BS | 98 | VAL |
| 44 | BS | 101 | LEU |
| 45 | BT | 3 | ARG |
| 45 | BT | 7 | ILE |
| 45 | BT | 11 | GLU |
| 45 | BT | 13 | ARG |
| 45 | BT | 15 | VAL |
| 45 | BT | 16 | ARG |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 45 | BT | 17 | THR |
| 45 | BT | 19 | LEU |
| 45 | BT | 24 | PRO |
| 45 | BT | 29 | ARG |
| 45 | BT | 33 | LYS |
| 45 | BT | 38 | ASN |
| 45 | BT | 41 | ARG |
| 45 | BT | 49 | VAL |
| 45 | BT | 50 | ILE |
| 45 | BT | 51 | ARG |
| 45 | BT | 58 | ASN |
| 45 | BT | 59 | THR |
| 45 | BT | 62 | THR |
| 45 | BT | 63 | VAL |
| 45 | BT | 65 | LYS |
| 45 | BT | 72 | VAL |
| 45 | BT | 74 | ARG |
| 45 | BT | 82 | LEU |
| 45 | BT | 96 | ARG |
| 45 | BT | 99 | LEU |
| 45 | BT | 108 | ARG |
| 45 | BT | 110 | ILE |
| 45 | BT | 111 | ARG |
| 45 | BT | 112 | ARG |
| 45 | BT | 115 | ARG |
| 45 | BT | 121 | ILE |
| 45 | BT | 128 | GLU |
| 46 | BU | 17 | ILE |
| 46 | BU | 20 | LEU |
| 46 | BU | 27 | LEU |
| 46 | BU | 31 | SER |
| 46 | BU | 55 | ARG |
| 46 | BU | 56 | ASP |
| 46 | BU | 62 | ILE |
| 46 | BU | 64 | ARG |
| 46 | BU | 66 | ASN |
| 46 | BU | 74 | LEU |
| 46 | BU | 76 | TYR |
| 46 | BU | 83 | LEU |
| 46 | BU | 88 | ILE |
| 46 | BU | 89 | GLU |
| 46 | BU | 93 | LYS |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 46 | BU | 97 | ASP |
| 46 | BU | 100 | VAL |
| 46 | BU | 102 | GLU |
| 46 | BU | 112 | ARG |
| 46 | BU | 114 | LYS |
| 47 | BV | 1 | MET |
| 47 | BV | 2 | PHE |
| 47 | BV | 4 | ILE |
| 47 | BV | 6 | LYS |
| 47 | BV | 7 | THR |
| 47 | BV | 13 | ARG |
| 47 | BV | 14 | VAL |
| 47 | BV | 18 | LEU |
| 47 | BV | 19 | LYS |
| 47 | BV | 23 | GLU |
| 47 | BV | 28 | GLU |
| 47 | BV | 32 | THR |
| 47 | BV | 35 | LEU |
| 47 | BV | 37 | VAL |
| 47 | BV | 39 | LEU |
| 47 | BV | 40 | LEU |
| 47 | BV | 56 | SER |
| 47 | BV | 60 | GLU |
| 47 | BV | 62 | LEU |
| 47 | BV | 66 | ARG |
| 47 | BV | 71 | LEU |
| 47 | BV | 76 | LYS |
| 47 | BV | 78 | LYS |
| 47 | BV | 80 | GLN |
| 47 | BV | 82 | ARG |
| 47 | BV | 83 | ARG |
| 47 | BV | 88 | ARG |
| 47 | BV | 89 | GLN |
| 47 | BV | 92 | THR |
| 47 | BV | 96 | ILE |
| 47 | BV | 98 | GLU |
| 48 | BW | 6 | ILE |
| 48 | BW | 11 | ARG |
| 48 | BW | 15 | ARG |
| 48 | BW | 16 | LYS |
| 48 | BW | 17 | VAL |
| 48 | BW | 18 | ARG |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 48 | BW | 19 | LEU |
| 48 | BW | 20 | VAL |
| 48 | BW | 23 | LEU |
| 48 | BW | 27 | LYS |
| 48 | BW | 49 | LYS |
| 48 | BW | 50 | VAL |
| 48 | BW | 51 | LEU |
| 48 | BW | 52 | GLU |
| 48 | BW | 59 | VAL |
| 48 | BW | 60 | ASN |
| 48 | BW | 72 | LYS |
| 48 | BW | 76 | VAL |
| 48 | BW | 78 | GLU |
| 48 | BW | 83 | LYS |
| 48 | BW | 86 | LEU |
| 48 | BW | 103 | ILE |
| 48 | BW | 106 | ILE |
| 48 | BW | 107 | LEU |
| 49 | BX | 13 | LEU |
| 49 | BX | 15 | GLU |
| 49 | BX | 23 | GLU |
| 49 | BX | 25 | LYS |
| 49 | BX | 27 | THR |
| 49 | BX | 30 | VAL |
| 49 | BX | 33 | LYS |
| 49 | BX | 35 | THR |
| 49 | BX | 36 | LYS |
| 49 | BX | 38 | GLU |
| 49 | BX | 39 | ILE |
| 49 | BX | 43 | VAL |
| 49 | BX | 49 | VAL |
| 49 | BX | 57 | LEU |
| 49 | BX | 59 | VAL |
| 49 | BX | 65 | ARG |
| 49 | BX | 66 | LEU |
| 49 | BX | 73 | ARG |
| 49 | BX | 76 | ARG |
| 49 | BX | 78 | LYS |
| 49 | BX | 81 | VAL |
| 50 | BY | 2 | ARG |
| 50 | BY | 7 | VAL |
| 50 | BY | 9 | LYS |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 50 | BY | 13 | VAL |
| 50 | BY | 15 | VAL |
| 50 | BY | 32 | PRO |
| 50 | BY | 38 | ILE |
| 50 | BY | 42 | VAL |
| 50 | BY | 44 | ILE |
| 50 | BY | 47 | LYS |
| 50 | BY | 49 | VAL |
| 50 | BY | 66 | PRO |
| 50 | BY | 70 | SER |
| 50 | BY | 71 | LYS |
| 50 | BY | 75 | ILE |
| 50 | BY | 76 | CYS |
| 50 | BY | 85 | VAL |
| 50 | BY | 86 | ARG |
| 50 | BY | 89 | PHE |
| 50 | BY | 90 | LEU |
| 50 | BY | 96 | ILE |
| 50 | BY | 97 | ARG |
| 51 | BZ | 5 | LEU |
| 51 | BZ | 6 | LYS |
| 51 | BZ | 19 | ARG |
| 51 | BZ | 27 | VAL |
| 51 | BZ | 31 | ARG |
| 51 | BZ | 37 | VAL |
| 51 | BZ | 39 | VAL |
| 51 | BZ | 41 | LEU |
| 51 | BZ | 42 | VAL |
| 51 | BZ | 53 | ILE |
| 51 | BZ | 57 | ILE |
| 51 | BZ | 72 | ARG |
| 51 | BZ | 73 | GLN |
| 51 | BZ | 79 | ARG |
| 51 | BZ | 85 | HIS |
| 51 | BZ | 86 | VAL |
| 51 | BZ | 87 | ASP |
| 51 | BZ | 93 | ASP |
| 51 | BZ | 96 | VAL |
| 51 | BZ | 97 | GLU |
| 51 | BZ | 117 | LEU |
| 51 | BZ | 120 | ILE |
| 51 | BZ | 121 | HIS |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 51 | BZ | 124 | ILE |
| 51 | BZ | 140 | ASP |
| 51 | BZ | 150 | LEU |
| 51 | BZ | 161 | VAL |
| 2 | CB | 9 | GLU |
| 2 | CB | 12 | GLU |
| 2 | CB | 15 | VAL |
| 2 | CB | 17 | PHE |
| 2 | CB | 22 | LYS |
| 2 | CB | 36 | ARG |
| 2 | CB | 42 | ILE |
| 2 | CB | 44 | LEU |
| 2 | CB | 59 | GLU |
| 2 | CB | 69 | LEU |
| 2 | CB | 71 | VAL |
| 2 | CB | 75 | LYS |
| 2 | CB | 79 | ASP |
| 2 | CB | 90 | MET |
| 2 | CB | 111 | ARG |
| 2 | CB | 130 | ARG |
| 2 | CB | 137 | ARG |
| 2 | CB | 145 | LEU |
| 2 | CB | 146 | GLN |
| 2 | CB | 154 | LEU |
| 2 | CB | 158 | LEU |
| 2 | CB | 164 | VAL |
| 2 | CB | 179 | LYS |
| 2 | CB | 185 | ILE |
| 2 | CB | 187 | LEU |
| 2 | CB | 189 | ASP |
| 2 | CB | 196 | LEU |
| 2 | CB | 204 | ASN |
| 2 | CB | 205 | ASP |
| 2 | CB | 215 | LEU |
| 2 | CB | 221 | LEU |
| 3 | CC | 5 | ILE |
| 3 | CC | 12 | LEU |
| 3 | CC | 27 | LYS |
| 3 | CC | 69 | HIS |
| 3 | CC | 76 | VAL |
| 3 | CC | 104 | GLN |
| 3 | CC | 131 | ARG |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 3 | CC | 138 | VAL |
| 3 | CC | 198 | VAL |
| 4 | CD | 3 | ARG |
| 4 | CD | 8 | VAL |
| 4 | CD | 9 | CYS |
| 4 | CD | 11 | LEU |
| 4 | CD | 15 | GLU |
| 4 | CD | 19 | LEU |
| 4 | CD | 25 | ARG |
| 4 | CD | 26 | CYS |
| 4 | CD | 36 | ARG |
| 4 | CD | 58 | LEU |
| 4 | CD | 62 | GLN |
| 4 | CD | 68 | TYR |
| 4 | CD | 70 | ILE |
| 4 | CD | 94 | LEU |
| 4 | CD | 97 | LEU |
| 4 | CD | 104 | VAL |
| 4 | CD | 121 | VAL |
| 4 | CD | 127 | THR |
| 4 | CD | 129 | ASN |
| 4 | CD | 132 | ARG |
| 4 | CD | 135 | LEU |
| 4 | CD | 138 | TYR |
| 4 | CD | 158 | ILE |
| 4 | CD | 194 | LEU |
| 4 | CD | 196 | LEU |
| 4 | CD | 202 | LEU |
| 4 | CD | 203 | VAL |
| 5 | CE | 10 | MET |
| 5 | CE | 12 | LEU |
| 5 | CE | 20 | GLN |
| 5 | CE | 25 | ARG |
| 5 | CE | 31 | LEU |
| 5 | CE | 41 | VAL |
| 5 | CE | 55 | VAL |
| 5 | CE | 73 | ASN |
| 5 | CE | 76 | ILE |
| 5 | CE | 78 | HIS |
| 5 | CE | 79 | GLU |
| 5 | CE | 82 | VAL |
| 5 | CE | 91 | LEU |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 5 | CE | 98 | THR |
| 5 | CE | 101 | ILE |
| 5 | CE | 112 | LEU |
| 5 | CE | 115 | VAL |
| 5 | CE | 147 | ASP |
| 6 | CF | 10 | LEU |
| 6 | CF | 21 | LEU |
| 6 | CF | 30 | LEU |
| 6 | CF | 45 | LEU |
| 6 | CF | 46 | ARG |
| 6 | CF | 52 | ILE |
| 6 | CF | 55 | ASP |
| 6 | CF | 63 | TYR |
| 6 | CF | 70 | ASP |
| 6 | CF | 92 | LYS |
| 6 | CF | 98 | LEU |
| 7 | CG | 27 | ILE |
| 7 | CG | 36 | LYS |
| 7 | CG | 79 | ARG |
| 7 | CG | 136 | LYS |
| 7 | CG | 151 | TYR |
| 8 | CH | 1 | MET |
| 8 | CH | 3 | THR |
| 8 | CH | 10 | LEU |
| 8 | CH | 24 | THR |
| 8 | CH | 39 | LEU |
| 8 | CH | 51 | VAL |
| 8 | CH | 65 | TYR |
| 8 | CH | 81 | HIS |
| 8 | CH | 83 | ILE |
| 8 | CH | 91 | ARG |
| 8 | CH | 95 | VAL |
| 8 | CH | 102 | ARG |
| 8 | CH | 103 | VAL |
| 8 | CH | 133 | LEU |
| 8 | CH | 137 | VAL |
| 9 | CI | 3 | GLN |
| 9 | CI | 10 | ARG |
| 9 | CI | 78 | LYS |
| 9 | CI | 95 | LYS |
| 9 | CI | 99 | LEU |
| 9 | CI | 101 | PHE |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 9 | CI | 102 | LEU |
| 9 | CI | 114 | TYR |
| 9 | CI | 128 | ARG |
| 10 | CJ | 5 | ARG |
| 10 | CJ | 74 | ILE |
| 10 | CJ | 80 | LYS |
| 10 | CJ | 96 | ILE |
| 10 | CJ | 100 | THR |
| 11 | CK | 18 | ARG |
| 11 | CK | 22 | HIS |
| 11 | CK | 29 | ILE |
| 11 | CK | 53 | SER |
| 11 | CK | 63 | LEU |
| 11 | CK | 92 | GLU |
| 11 | CK | 95 | ILE |
| 11 | CK | 103 | LEU |
| 11 | CK | 114 | VAL |
| 11 | CK | 117 | ASN |
| 11 | CK | 125 | PHE |
| 11 | CK | 127 | LYS |
| 12 | CL | 24 | VAL |
| 12 | CL | 38 | THR |
| 12 | CL | 41 | ARG |
| 12 | CL | 42 | THR |
| 12 | CL | 62 | SER |
| 12 | CL | 75 | HIS |
| 12 | CL | 85 | ILE |
| 12 | CL | 89 | ARG |
| 12 | CL | 96 | VAL |
| 12 | CL | 102 | ARG |
| 13 | CM | 47 | ASP |
| 13 | CM | 66 | LEU |
| 13 | CM | 70 | LEU |
| 13 | CM | 93 | ARG |
| 13 | CM | 108 | ARG |
| 14 | CN | 18 | VAL |
| 14 | CN | 42 | ILE |
| 14 | CN | 44 | LEU |
| 15 | CO | 3 | ILE |
| 15 | CO | 26 | GLU |
| 15 | CO | 37 | ASN |
| 15 | CO | 39 | LEU |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 15 | CO | 41 | GLU |
| 15 | CO | 57 | LEU |
| 15 | CO | 65 | ARG |
| 15 | CO | 67 | LEU |
| 15 | CO | 82 | ILE |
| 16 | CP | 2 | VAL |
| 16 | CP | 27 | LYS |
| 16 | CP | 28 | ARG |
| 16 | CP | 47 | ASP |
| 16 | CP | 53 | VAL |
| 16 | CP | 65 | GLN |
| 16 | CP | 67 | THR |
| 16 | CP | 69 | THR |
| 16 | CP | 74 | LEU |
| 16 | CP | 79 | VAL |
| 16 | CP | 82 | GLN |
| 17 | CQ | 5 | VAL |
| 17 | CQ | 52 | LYS |
| 17 | CQ | 63 | ARG |
| 17 | CQ | 74 | LEU |
| 17 | CQ | 89 | LEU |
| 18 | CR | 31 | LEU |
| 18 | CR | 47 | THR |
| 18 | CR | 53 | ARG |
| 18 | CR | 65 | ILE |
| 18 | CR | 70 | ILE |
| 18 | CR | 74 | ARG |
| 18 | CR | 76 | LEU |
| 18 | CR | 79 | LEU |
| 18 | CR | 82 | THR |
| 18 | CR | 87 | ARG |
| 19 | CS | 6 | LYS |
| 19 | CS | 7 | LYS |
| 19 | CS | 22 | LEU |
| 19 | CS | 31 | ILE |
| 19 | CS | 37 | ARG |
| 19 | CS | 49 | ILE |
| 20 | CT | 26 | ASN |
| 20 | CT | 27 | LYS |
| 20 | CT | 30 | LYS |
| 20 | CT | 35 | THR |
| 20 | CT | 41 | ILE |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 20 | CT | 51 | GLU |
| 20 | CT | 56 | MET |
| 20 | CT | 57 | ARG |
| 20 | CT | 64 | ASP |
| 20 | CT | 72 | LEU |
| 20 | CT | 73 | HIS |
| 20 | CT | 93 | GLU |
| 22 | D0 | 11 | ARG |
| 22 | D0 | 14 | ARG |
| 22 | D0 | 30 | VAL |
| 22 | D0 | 31 | VAL |
| 22 | D0 | 36 | ILE |
| 22 | D0 | 40 | GLN |
| 22 | D0 | 41 | ARG |
| 22 | D0 | 53 | MET |
| 22 | D0 | 55 | ARG |
| 22 | D0 | 62 | LEU |
| 22 | D0 | 70 | GLN |
| 22 | D0 | 72 | ARG |
| 22 | D0 | 79 | VAL |
| 22 | D0 | 84 | LEU |
| 23 | D1 | 8 | SER |
| 23 | D1 | 13 | ILE |
| 23 | D1 | 14 | VAL |
| 23 | D1 | 25 | LYS |
| 23 | D1 | 26 | ARG |
| 23 | D1 | 34 | THR |
| 23 | D1 | 37 | ILE |
| 23 | D1 | 41 | ARG |
| 23 | D1 | 46 | LEU |
| 23 | D1 | 47 | GLN |
| 23 | D1 | 51 | VAL |
| 23 | D1 | 56 | GLN |
| 23 | D1 | 65 | SER |
| 23 | D1 | 67 | ILE |
| 23 | D1 | 74 | VAL |
| 23 | D1 | 85 | LEU |
| 23 | D1 | 89 | GLU |
| 24 | D2 | 12 | GLU |
| 24 | D2 | 14 | ARG |
| 24 | D2 | 22 | GLU |
| 24 | D2 | 28 | LYS |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 24 | D2 | 30 | ARG |
| 24 | D2 | 33 | MET |
| 24 | D2 | 44 | LEU |
| 24 | D2 | 45 | SER |
| 24 | D2 | 46 | GLN |
| 24 | D2 | 47 | ASN |
| 24 | D2 | 50 | ILE |
| 24 | D2 | 56 | GLN |
| 25 | D3 | 8 | LEU |
| 25 | D3 | 9 | VAL |
| 25 | D3 | 10 | LYS |
| 25 | D3 | 18 | ASP |
| 25 | D3 | 24 | LYS |
| 25 | D3 | 38 | GLU |
| 25 | D3 | 40 | THR |
| 25 | D3 | 59 | VAL |
| 27 | D5 | 4 | HIS |
| 27 | D5 | 9 | LYS |
| 27 | D5 | 11 | THR |
| 27 | D5 | 16 | ARG |
| 27 | D5 | 29 | THR |
| 27 | D5 | 36 | CYS |
| 27 | D5 | 44 | THR |
| 27 | D5 | 45 | VAL |
| 27 | D5 | 49 | CYS |
| 27 | D5 | 51 | TYR |
| 27 | D5 | 55 | ARG |
| 27 | D5 | 56 | LYS |
| 27 | D5 | 58 | LEU |
| 28 | D6 | 9 | LEU |
| 28 | D6 | 10 | LEU |
| 28 | D6 | 11 | LEU |
| 28 | D6 | 12 | GLU |
| 28 | D6 | 18 | ARG |
| 28 | D6 | 25 | LYS |
| 28 | D6 | 27 | LYS |
| 28 | D6 | 30 | THR |
| 28 | D6 | 37 | ARG |
| 28 | D6 | 42 | TRP |
| 28 | D6 | 46 | HIS |
| 28 | D6 | 47 | THR |
| 28 | D6 | 48 | VAL |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 28 | D6 | 51 | GLU |
| 29 | D7 | 1 | MET |
| 29 | D7 | 8 | ASN |
| 29 | D7 | 9 | ARG |
| 29 | D7 | 15 | THR |
| 29 | D7 | 24 | THR |
| 29 | D7 | 42 | LEU |
| 29 | D7 | 43 | THR |
| 29 | D7 | 48 | LYS |
| 30 | D8 | 4 | MET |
| 30 | D8 | 6 | THR |
| 30 | D8 | 13 | ARG |
| 30 | D8 | 16 | ILE |
| 30 | D8 | 32 | LEU |
| 30 | D8 | 37 | SER |
| 30 | D8 | 39 | LYS |
| 30 | D8 | 41 | ILE |
| 30 | D8 | 44 | LYS |
| 30 | D8 | 47 | LYS |
| 30 | D8 | 49 | VAL |
| 30 | D8 | 54 | GLU |
| 30 | D8 | 56 | GLU |
| 30 | D8 | 58 | ILE |
| 30 | D8 | 59 | LYS |
| 33 | DD | 3 | VAL |
| 33 | DD | 5 | LYS |
| 33 | DD | 10 | THR |
| 33 | DD | 24 | ILE |
| 33 | DD | 26 | LYS |
| 33 | DD | 27 | THR |
| 33 | DD | 31 | LYS |
| 33 | DD | 33 | LEU |
| 33 | DD | 44 | ASN |
| 33 | DD | 46 | GLN |
| 33 | DD | 49 | ILE |
| 33 | DD | 59 | LYS |
| 33 | DD | 61 | LEU |
| 33 | DD | 63 | ARG |
| 33 | DD | 64 | ILE |
| 33 | DD | 65 | ILE |
| 33 | DD | 71 | ASP |
| 33 | DD | 72 | LYS |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 33 | DD | 75 | ILE |
| 33 | DD | 87 | ASN |
| 33 | DD | 88 | ARG |
| 33 | DD | 89 | SER |
| 33 | DD | 92 | ILE |
| 33 | DD | 94 | LEU |
| 33 | DD | 95 | LEU |
| 33 | DD | 98 | VAL |
| 33 | DD | 103 | ARG |
| 33 | DD | 105 | ILE |
| 33 | DD | 106 | ILE |
| 33 | DD | 111 | LEU |
| 33 | DD | 113 | VAL |
| 33 | DD | 117 | VAL |
| 33 | DD | 127 | VAL |
| 33 | DD | 131 | LEU |
| 33 | DD | 138 | VAL |
| 33 | DD | 147 | LEU |
| 33 | DD | 150 | LYS |
| 33 | DD | 157 | ARG |
| 33 | DD | 161 | THR |
| 33 | DD | 162 | SER |
| 33 | DD | 165 | ILE |
| 33 | DD | 166 | GLN |
| 33 | DD | 173 | VAL |
| 33 | DD | 174 | ILE |
| 33 | DD | 176 | ARG |
| 33 | DD | 182 | LEU |
| 33 | DD | 183 | ARG |
| 33 | DD | 192 | THR |
| 33 | DD | 193 | VAL |
| 33 | DD | 202 | LYS |
| 33 | DD | 211 | ARG |
| 33 | DD | 212 | SER |
| 33 | DD | 217 | ARG |
| 33 | DD | 221 | VAL |
| 33 | DD | 229 | VAL |
| 33 | DD | 242 | ARG |
| 33 | DD | 255 | LYS |
| 33 | DD | 257 | LEU |
| 33 | DD | 259 | THR |
| 33 | DD | 266 | SER |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 33 | DD | 270 | ILE |
| 33 | DD | 271 | ILE |
| 34 | DE | 1 | MET |
| 34 | DE | 7 | VAL |
| 34 | DE | 9 | VAL |
| 34 | DE | 12 | THR |
| 34 | DE | 19 | ARG |
| 34 | DE | 23 | VAL |
| 34 | DE | 26 | ILE |
| 34 | DE | 34 | VAL |
| 34 | DE | 60 | ASN |
| 34 | DE | 63 | LEU |
| 34 | DE | 64 | LYS |
| 34 | DE | 69 | LYS |
| 34 | DE | 75 | VAL |
| 34 | DE | 76 | ARG |
| 34 | DE | 77 | ILE |
| 34 | DE | 78 | LEU |
| 34 | DE | 79 | ARG |
| 34 | DE | 82 | ARG |
| 34 | DE | 111 | ARG |
| 34 | DE | 116 | VAL |
| 34 | DE | 118 | LYS |
| 34 | DE | 119 | ARG |
| 34 | DE | 134 | ILE |
| 34 | DE | 136 | ARG |
| 34 | DE | 144 | ARG |
| 34 | DE | 145 | LYS |
| 34 | DE | 154 | LYS |
| 34 | DE | 159 | HIS |
| 34 | DE | 167 | VAL |
| 34 | DE | 175 | VAL |
| 34 | DE | 181 | LEU |
| 34 | DE | 182 | LEU |
| 34 | DE | 184 | VAL |
| 34 | DE | 188 | VAL |
| 34 | DE | 196 | VAL |
| 34 | DE | 197 | ILE |
| 34 | DE | 202 | LYS |
| 34 | DE | 203 | LYS |
| 35 | DF | 13 | SER |
| 35 | DF | 20 | LEU |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 35 | DF | 23 | ASP |
| 35 | DF | 28 | ILE |
| 35 | DF | 33 | LEU |
| 35 | DF | 38 | ARG |
| 35 | DF | 46 | ARG |
| 35 | DF | 51 | THR |
| 35 | DF | 56 | GLU |
| 35 | DF | 57 | VAL |
| 35 | DF | 66 | PRO |
| 35 | DF | 67 | GLN |
| 35 | DF | 74 | ARG |
| 35 | DF | 78 | ILE |
| 35 | DF | 82 | ILE |
| 35 | DF | 83 | PHE |
| 35 | DF | 84 | VAL |
| 35 | DF | 88 | VAL |
| 35 | DF | 106 | ARG |
| 35 | DF | 107 | LYS |
| 35 | DF | 110 | LEU |
| 35 | DF | 112 | MET |
| 35 | DF | 124 | LEU |
| 35 | DF | 127 | GLU |
| 35 | DF | 140 | LEU |
| 35 | DF | 160 | ASN |
| 35 | DF | 162 | LEU |
| 35 | DF | 164 | ARG |
| 35 | DF | 165 | ARG |
| 35 | DF | 168 | ARG |
| 35 | DF | 175 | THR |
| 35 | DF | 179 | GLU |
| 35 | DF | 183 | VAL |
| 35 | DF | 186 | ILE |
| 35 | DF | 191 | ARG |
| 35 | DF | 192 | LEU |
| 35 | DF | 196 | LEU |
| 35 | DF | 197 | ASP |
| 35 | DF | 206 | ILE |
| 36 | DG | 5 | VAL |
| 36 | DG | 7 | LEU |
| 36 | DG | 16 | ARG |
| 36 | DG | 21 | ARG |
| 36 | DG | 22 | ARG |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 36 | DG | 28 | VAL |
| 36 | DG | 31 | VAL |
| 36 | DG | 35 | GLU |
| 36 | DG | 39 | ILE |
| 36 | DG | 45 | GLU |
| 36 | DG | 49 | ASP |
| 36 | DG | 60 | LEU |
| 36 | DG | 63 | ILE |
| 36 | DG | 67 | LYS |
| 36 | DG | 74 | LYS |
| 36 | DG | 83 | ARG |
| 36 | DG | 86 | MET |
| 36 | DG | 115 | ARG |
| 36 | DG | 125 | PHE |
| 36 | DG | 130 | ASN |
| 36 | DG | 135 | LEU |
| 36 | DG | 139 | LEU |
| 36 | DG | 143 | GLU |
| 36 | DG | 155 | MET |
| 36 | DG | 156 | ASP |
| 36 | DG | 159 | VAL |
| 36 | DG | 161 | THR |
| 36 | DG | 166 | ASP |
| 36 | DG | 173 | LEU |
| 37 | DH | 13 | LYS |
| 37 | DH | 15 | VAL |
| 37 | DH | 21 | PRO |
| 37 | DH | 27 | LYS |
| 37 | DH | 34 | GLU |
| 37 | DH | 41 | MET |
| 37 | DH | 46 | GLU |
| 37 | DH | 53 | GLU |
| 37 | DH | 70 | THR |
| 37 | DH | 71 | LEU |
| 37 | DH | 89 | ILE |
| 37 | DH | 92 | ILE |
| 37 | DH | 98 | LEU |
| 37 | DH | 103 | LEU |
| 37 | DH | 105 | LEU |
| 37 | DH | 122 | THR |
| 37 | DH | 134 | SER |
| 37 | DH | 136 | ILE |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 37 | DH | 141 | VAL |
| 37 | DH | 143 | GLN |
| 37 | DH | 152 | ARG |
| 37 | DH | 153 | LYS |
| 37 | DH | 157 | TYR |
| 37 | DH | 159 | GLU |
| 37 | DH | 170 | ARG |
| 38 | DI | 1 | MET |
| 38 | DI | 7 | GLU |
| 38 | DI | 15 | VAL |
| 38 | DI | 18 | VAL |
| 38 | DI | 20 | ASP |
| 38 | DI | 22 | LYS |
| 38 | DI | 31 | LEU |
| 38 | DI | 37 | VAL |
| 38 | DI | 51 | ILE |
| 38 | DI | 56 | LYS |
| 38 | DI | 58 | LEU |
| 38 | DI | 64 | GLU |
| 38 | DI | 71 | ILE |
| 38 | DI | 72 | LEU |
| 38 | DI | 79 | ILE |
| 38 | DI | 81 | VAL |
| 38 | DI | 101 | LEU |
| 38 | DI | 109 | ILE |
| 38 | DI | 122 | GLU |
| 38 | DI | 134 | PRO |
| 38 | DI | 136 | VAL |
| 38 | DI | 138 | ILE |
| 38 | DI | 142 | VAL |
| 38 | DI | 144 | VAL |
| 39 | DN | 4 | TYR |
| 39 | DN | 5 | VAL |
| 39 | DN | 7 | LYS |
| 39 | DN | 9 | VAL |
| 39 | DN | 14 | VAL |
| 39 | DN | 16 | ILE |
| 39 | DN | 19 | GLU |
| 39 | DN | 23 | LEU |
| 39 | DN | 25 | ARG |
| 39 | DN | 33 | LEU |
| 39 | DN | 34 | LEU |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 39 | DN | 43 | THR |
| 39 | DN | 45 | ASN |
| 39 | DN | 46 | VAL |
| 39 | DN | 48 | MET |
| 39 | DN | 55 | VAL |
| 39 | DN | 60 | ILE |
| 39 | DN | 63 | THR |
| 39 | DN | 65 | LYS |
| 39 | DN | 66 | LYS |
| 39 | DN | 75 | TYR |
| 39 | DN | 78 | TYR |
| 39 | DN | 82 | LEU |
| 39 | DN | 85 | ILE |
| 39 | DN | 87 | LEU |
| 39 | DN | 94 | HIS |
| 39 | DN | 99 | LEU |
| 39 | DN | 104 | LYS |
| 39 | DN | 120 | LEU |
| 39 | DN | 121 | LYS |
| 39 | DN | 131 | GLN |
| 40 | DO | 2 | ILE |
| 40 | DO | 3 | GLN |
| 40 | DO | 10 | VAL |
| 40 | DO | 19 | ILE |
| 40 | DO | 22 | ILE |
| 40 | DO | 23 | ARG |
| 40 | DO | 24 | VAL |
| 40 | DO | 28 | SER |
| 40 | DO | 32 | TYR |
| 40 | DO | 35 | VAL |
| 40 | DO | 42 | SER |
| 40 | DO | 47 | ILE |
| 40 | DO | 58 | VAL |
| 40 | DO | 63 | VAL |
| 40 | DO | 65 | THR |
| 40 | DO | 75 | SER |
| 40 | DO | 78 | ARG |
| 40 | DO | 80 | ASP |
| 40 | DO | 85 | VAL |
| 40 | DO | 87 | ILE |
| 40 | DO | 88 | ASN |
| 40 | DO | 89 | ASN |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 40 | DO | 91 | LEU |
| 40 | DO | 94 | ARG |
| 40 | DO | 102 | VAL |
| 40 | DO | 108 | GLU |
| 40 | DO | 112 | MET |
| 40 | DO | 114 | ILE |
| 40 | DO | 121 | VAL |
| 41 | DP | 9 | ASN |
| 41 | DP | 10 | PRO |
| 41 | DP | 13 | ASN |
| 41 | DP | 16 | ARG |
| 41 | DP | 17 | LYS |
| 41 | DP | 18 | ARG |
| 41 | DP | 19 | VAL |
| 41 | DP | 32 | THR |
| 41 | DP | 36 | LYS |
| 41 | DP | 39 | LYS |
| 41 | DP | 40 | SER |
| 41 | DP | 45 | LEU |
| 41 | DP | 47 | ASP |
| 41 | DP | 51 | PHE |
| 41 | DP | 52 | GLU |
| 41 | DP | 57 | THR |
| 41 | DP | 59 | LEU |
| 41 | DP | 60 | MET |
| 41 | DP | 61 | ARG |
| 41 | DP | 62 | LEU |
| 41 | DP | 64 | LYS |
| 41 | DP | 67 | MET |
| 41 | DP | 71 | VAL |
| 41 | DP | 75 | ILE |
| 41 | DP | 79 | ARG |
| 41 | DP | 81 | GLN |
| 41 | DP | 83 | VAL |
| 41 | DP | 88 | LEU |
| 41 | DP | 95 | VAL |
| 41 | DP | 96 | THR |
| 41 | DP | 98 | GLU |
| 41 | DP | 100 | LEU |
| 41 | DP | 101 | VAL |
| 41 | DP | 102 | ARG |
| 41 | DP | 111 | ARG |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 41 | DP | 112 | LEU |
| 41 | DP | 114 | ILE |
| 41 | DP | 115 | LEU |
| 41 | DP | 123 | LEU |
| 41 | DP | 125 | VAL |
| 41 | DP | 126 | VAL |
| 41 | DP | 135 | LEU |
| 41 | DP | 138 | LEU |
| 41 | DP | 144 | GLU |
| 41 | DP | 147 | LEU |
| 41 | DP | 148 | LEU |
| 42 | DQ | 6 | ARG |
| 42 | DQ | 7 | MET |
| 42 | DQ | 9 | TYR |
| 42 | DQ | 13 | GLN |
| 42 | DQ | 17 | LEU |
| 42 | DQ | 22 | LYS |
| 42 | DQ | 25 | ASP |
| 42 | DQ | 35 | VAL |
| 42 | DQ | 45 | GLN |
| 42 | DQ | 52 | VAL |
| 42 | DQ | 54 | MET |
| 42 | DQ | 55 | VAL |
| 42 | DQ | 63 | LYS |
| 42 | DQ | 68 | ILE |
| 42 | DQ | 81 | VAL |
| 42 | DQ | 82 | ARG |
| 42 | DQ | 91 | GLU |
| 42 | DQ | 102 | VAL |
| 42 | DQ | 109 | VAL |
| 42 | DQ | 110 | THR |
| 42 | DQ | 115 | MET |
| 42 | DQ | 127 | ILE |
| 42 | DQ | 141 | GLN |
| 43 | DR | 2 | ARG |
| 43 | DR | 4 | LEU |
| 43 | DR | 5 | LYS |
| 43 | DR | 18 | LEU |
| 43 | DR | 28 | LEU |
| 43 | DR | 29 | LEU |
| 43 | DR | 34 | ILE |
| 43 | DR | 44 | LEU |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 43 | DR | 45 | ARG |
| 43 | DR | 54 | LEU |
| 43 | DR | 56 | LYS |
| 43 | DR | 60 | LEU |
| 43 | DR | 64 | ARG |
| 43 | DR | 65 | LEU |
| 43 | DR | 67 | LEU |
| 43 | DR | 71 | GLN |
| 43 | DR | 75 | LEU |
| 43 | DR | 79 | LEU |
| 43 | DR | 99 | LYS |
| 43 | DR | 100 | LEU |
| 43 | DR | 104 | ARG |
| 43 | DR | 113 | LEU |
| 43 | DR | 115 | GLU |
| 43 | DR | 116 | LEU |
| 43 | DR | 117 | VAL |
| 44 | DS | 11 | LYS |
| 44 | DS | 12 | PHE |
| 44 | DS | 14 | VAL |
| 44 | DS | 17 | ARG |
| 44 | DS | 18 | ILE |
| 44 | DS | 20 | ARG |
| 44 | DS | 32 | LEU |
| 44 | DS | 33 | LYS |
| 44 | DS | 34 | HIS |
| 44 | DS | 36 | TYR |
| 44 | DS | 46 | VAL |
| 44 | DS | 50 | SER |
| 44 | DS | 56 | LEU |
| 44 | DS | 61 | ASN |
| 44 | DS | 73 | LEU |
| 44 | DS | 80 | LEU |
| 44 | DS | 83 | LYS |
| 44 | DS | 84 | GLN |
| 44 | DS | 85 | VAL |
| 44 | DS | 92 | TYR |
| 44 | DS | 98 | VAL |
| 44 | DS | 101 | LEU |
| 45 | DT | 3 | ARG |
| 45 | DT | 7 | ILE |
| 45 | DT | 11 | GLU |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 45 | DT | 13 | ARG |
| 45 | DT | 15 | VAL |
| 45 | DT | 16 | ARG |
| 45 | DT | 17 | THR |
| 45 | DT | 19 | LEU |
| 45 | DT | 24 | PRO |
| 45 | DT | 29 | ARG |
| 45 | DT | 33 | LYS |
| 45 | DT | 38 | ASN |
| 45 | DT | 41 | ARG |
| 45 | DT | 49 | VAL |
| 45 | DT | 50 | ILE |
| 45 | DT | 51 | ARG |
| 45 | DT | 58 | ASN |
| 45 | DT | 59 | THR |
| 45 | DT | 62 | THR |
| 45 | DT | 63 | VAL |
| 45 | DT | 65 | LYS |
| 45 | DT | 72 | VAL |
| 45 | DT | 74 | ARG |
| 45 | DT | 82 | LEU |
| 45 | DT | 96 | ARG |
| 45 | DT | 99 | LEU |
| 45 | DT | 108 | ARG |
| 45 | DT | 110 | ILE |
| 45 | DT | 111 | ARG |
| 45 | DT | 112 | ARG |
| 45 | DT | 115 | ARG |
| 45 | DT | 121 | ILE |
| 45 | DT | 128 | GLU |
| 46 | DU | 17 | ILE |
| 46 | DU | 20 | LEU |
| 46 | DU | 27 | LEU |
| 46 | DU | 31 | SER |
| 46 | DU | 55 | ARG |
| 46 | DU | 56 | ASP |
| 46 | DU | 62 | ILE |
| 46 | DU | 64 | ARG |
| 46 | DU | 66 | ASN |
| 46 | DU | 74 | LEU |
| 46 | DU | 76 | TYR |
| 46 | DU | 80 | ILE |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 46 | DU | 83 | LEU |
| 46 | DU | 88 | ILE |
| 46 | DU | 89 | GLU |
| 46 | DU | 93 | LYS |
| 46 | DU | 97 | ASP |
| 46 | DU | 100 | VAL |
| 46 | DU | 102 | GLU |
| 46 | DU | 112 | ARG |
| 46 | DU | 114 | LYS |
| 47 | DV | 1 | MET |
| 47 | DV | 2 | PHE |
| 47 | DV | 4 | ILE |
| 47 | DV | 6 | LYS |
| 47 | DV | 7 | THR |
| 47 | DV | 11 | GLN |
| 47 | DV | 13 | ARG |
| 47 | DV | 14 | VAL |
| 47 | DV | 18 | LEU |
| 47 | DV | 19 | LYS |
| 47 | DV | 23 | GLU |
| 47 | DV | 28 | GLU |
| 47 | DV | 32 | THR |
| 47 | DV | 35 | LEU |
| 47 | DV | 37 | VAL |
| 47 | DV | 39 | LEU |
| 47 | DV | 40 | LEU |
| 47 | DV | 56 | SER |
| 47 | DV | 60 | GLU |
| 47 | DV | 62 | LEU |
| 47 | DV | 66 | ARG |
| 47 | DV | 71 | LEU |
| 47 | DV | 76 | LYS |
| 47 | DV | 78 | LYS |
| 47 | DV | 80 | GLN |
| 47 | DV | 82 | ARG |
| 47 | DV | 83 | ARG |
| 47 | DV | 88 | ARG |
| 47 | DV | 89 | GLN |
| 47 | DV | 92 | THR |
| 47 | DV | 96 | ILE |
| 47 | DV | 98 | GLU |
| 48 | DW | 6 | ILE |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 48 | DW | 11 | ARG |
| 48 | DW | 16 | LYS |
| 48 | DW | 17 | VAL |
| 48 | DW | 18 | ARG |
| 48 | DW | 19 | LEU |
| 48 | DW | 20 | VAL |
| 48 | DW | 23 | LEU |
| 48 | DW | 27 | LYS |
| 48 | DW | 49 | LYS |
| 48 | DW | 50 | VAL |
| 48 | DW | 51 | LEU |
| 48 | DW | 52 | GLU |
| 48 | DW | 59 | VAL |
| 48 | DW | 60 | ASN |
| 48 | DW | 76 | VAL |
| 48 | DW | 78 | GLU |
| 48 | DW | 82 | LEU |
| 48 | DW | 83 | LYS |
| 48 | DW | 86 | LEU |
| 48 | DW | 88 | ARG |
| 48 | DW | 103 | ILE |
| 48 | DW | 106 | ILE |
| 48 | DW | 107 | LEU |
| 49 | DX | 13 | LEU |
| 49 | DX | 15 | GLU |
| 49 | DX | 23 | GLU |
| 49 | DX | 25 | LYS |
| 49 | DX | 27 | THR |
| 49 | DX | 30 | VAL |
| 49 | DX | 33 | LYS |
| 49 | DX | 35 | THR |
| 49 | DX | 36 | LYS |
| 49 | DX | 38 | GLU |
| 49 | DX | 39 | ILE |
| 49 | DX | 43 | VAL |
| 49 | DX | 49 | VAL |
| 49 | DX | 57 | LEU |
| 49 | DX | 59 | VAL |
| 49 | DX | 65 | ARG |
| 49 | DX | 66 | LEU |
| 49 | DX | 73 | ARG |
| 49 | DX | 76 | ARG |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 49 | DX | 78 | LYS |
| 49 | DX | 81 | VAL |
| 50 | DY | 2 | ARG |
| 50 | DY | 7 | VAL |
| 50 | DY | 9 | LYS |
| 50 | DY | 13 | VAL |
| 50 | DY | 15 | VAL |
| 50 | DY | 38 | ILE |
| 50 | DY | 42 | VAL |
| 50 | DY | 44 | ILE |
| 50 | DY | 47 | LYS |
| 50 | DY | 49 | VAL |
| 50 | DY | 66 | PRO |
| 50 | DY | 70 | SER |
| 50 | DY | 71 | LYS |
| 50 | DY | 75 | ILE |
| 50 | DY | 76 | CYS |
| 50 | DY | 85 | VAL |
| 50 | DY | 86 | ARG |
| 50 | DY | 89 | PHE |
| 50 | DY | 90 | LEU |
| 50 | DY | 96 | ILE |
| 50 | DY | 97 | ARG |
| 51 | DZ | 5 | LEU |
| 51 | DZ | 6 | LYS |
| 51 | DZ | 19 | ARG |
| 51 | DZ | 27 | VAL |
| 51 | DZ | 31 | ARG |
| 51 | DZ | 37 | VAL |
| 51 | DZ | 39 | VAL |
| 51 | DZ | 41 | LEU |
| 51 | DZ | 42 | VAL |
| 51 | DZ | 53 | ILE |
| 51 | DZ | 57 | ILE |
| 51 | DZ | 72 | ARG |
| 51 | DZ | 73 | GLN |
| 51 | DZ | 79 | ARG |
| 51 | DZ | 85 | HIS |
| 51 | DZ | 86 | VAL |
| 51 | DZ | 87 | ASP |
| 51 | DZ | 93 | ASP |
| 51 | DZ | 96 | VAL |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 51 | DZ | 97 | GLU |
| 51 | DZ | 117 | LEU |
| 51 | DZ | 120 | ILE |
| 51 | DZ | 121 | HIS |
| 51 | DZ | 124 | ILE |
| 51 | DZ | 140 | ASP |
| 51 | DZ | 150 | LEU |
| 51 | DZ | 161 | VAL |

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (290) such sidechains are listed below:

| Mol | Chain | Res | Type |
|---|---|---|---|
| 2 | AB | 19 | HIS |
| 2 | AB | 40 | HIS |
| 2 | AB | 113 | HIS |
| 2 | AB | 146 | GLN |
| 2 | AB | 204 | ASN |
| 3 | AC | 28 | GLN |
| 3 | AC | 123 | GLN |
| 3 | AC | 170 | GLN |
| 4 | AD | 62 | GLN |
| 4 | AD | 123 | HIS |
| 4 | AD | 129 | ASN |
| 5 | AE | 20 | GLN |
| 5 | AE | 73 | ASN |
| 5 | AE | 78 | HIS |
| 6 | AF | 18 | GLN |
| 6 | AF | 27 | GLN |
| 6 | AF | 32 | ASN |
| 6 | AF | 84 | ASN |
| 6 | AF | 94 | GLN |
| 6 | AF | 100 | ASN |
| 7 | AG | 13 | GLN |
| 7 | AG | 37 | ASN |
| 7 | AG | 84 | ASN |
| 7 | AG | 106 | GLN |
| 8 | AH | 82 | HIS |
| 9 | AI | 34 | ASN |
| 9 | AI | 73 | GLN |
| 9 | AI | 87 | GLN |
| 9 | AI | 117 | HIS |
| 9 | AI | 124 | GLN |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 10 | AJ | 13 | HIS |
| 10 | AJ | 78 | ASN |
| 11 | AK | 26 | ASN |
| 11 | AK | 38 | ASN |
| 12 | AL | 8 | ASN |
| 12 | AL | 9 | GLN |
| 12 | AL | 49 | ASN |
| 13 | AM | 77 | ASN |
| 13 | AM | 92 | HIS |
| 15 | AO | 46 | HIS |
| 16 | AP | 76 | GLN |
| 16 | AP | 82 | GLN |
| 17 | AQ | 16 | GLN |
| 19 | AS | 53 | ASN |
| 20 | AT | 16 | HIS |
| 20 | AT | 18 | GLN |
| 22 | B0 | 29 | GLN |
| 22 | B0 | 35 | ASN |
| 22 | B0 | 70 | GLN |
| 23 | B1 | 19 | GLN |
| 23 | B1 | 45 | ASN |
| 23 | B1 | 66 | HIS |
| 24 | B2 | 38 | GLN |
| 24 | B2 | 46 | GLN |
| 24 | B2 | 47 | ASN |
| 24 | B2 | 56 | GLN |
| 25 | B3 | 19 | GLN |
| 25 | B3 | 32 | GLN |
| 25 | B3 | 46 | ASN |
| 25 | B3 | 52 | HIS |
| 27 | B5 | 4 | HIS |
| 27 | B5 | 23 | HIS |
| 27 | B5 | 43 | HIS |
| 28 | B6 | 20 | ASN |
| 28 | B6 | 26 | ASN |
| 28 | B6 | 32 | ASN |
| 28 | B6 | 49 | HIS |
| 29 | B7 | 8 | ASN |
| 29 | B7 | 36 | GLN |
| 30 | B8 | 33 | ASN |
| 30 | B8 | 35 | GLN |
| 33 | BD | 58 | HIS |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 33 | BD | 87 | ASN |
| 33 | BD | 96 | HIS |
| 33 | BD | 126 | GLN |
| 33 | BD | 143 | HIS |
| 33 | BD | 164 | GLN |
| 33 | BD | 166 | GLN |
| 33 | BD | 186 | HIS |
| 33 | BD | 198 | ASN |
| 34 | BE | 48 | GLN |
| 34 | BE | 54 | GLN |
| 34 | BE | 60 | ASN |
| 34 | BE | 66 | HIS |
| 34 | BE | 129 | HIS |
| 34 | BE | 132 | HIS |
| 34 | BE | 192 | ASN |
| 35 | BF | 69 | HIS |
| 35 | BF | 75 | HIS |
| 35 | BF | 160 | ASN |
| 35 | BF | 169 | ASN |
| 36 | BG | 66 | GLN |
| 37 | BH | 65 | HIS |
| 38 | BI | 28 | ASN |
| 38 | BI | 74 | ASN |
| 38 | BI | 105 | HIS |
| 38 | BI | 133 | HIS |
| 39 | BN | 45 | ASN |
| 39 | BN | 56 | ASN |
| 39 | BN | 101 | HIS |
| 39 | BN | 128 | HIS |
| 39 | BN | 130 | HIS |
| 40 | BO | 3 | GLN |
| 40 | BO | 82 | ASN |
| 41 | BP | 13 | ASN |
| 41 | BP | 81 | GLN |
| 41 | BP | 128 | HIS |
| 42 | BQ | 141 | GLN |
| 43 | BR | 13 | HIS |
| 43 | BR | 16 | HIS |
| 43 | BR | 23 | ASN |
| 43 | BR | 24 | GLN |
| 43 | BR | 53 | HIS |
| 43 | BR | 71 | GLN |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 43 | BR | 91 | GLN |
| 44 | BS | 34 | HIS |
| 44 | BS | 61 | ASN |
| 44 | BS | 68 | GLN |
| 45 | BT | 38 | ASN |
| 45 | BT | 43 | GLN |
| 45 | BT | 58 | ASN |
| 45 | BT | 90 | GLN |
| 45 | BT | 123 | GLN |
| 46 | BU | 14 | HIS |
| 46 | BU | 49 | HIS |
| 46 | BU | 71 | GLN |
| 46 | BU | 72 | HIS |
| 46 | BU | 94 | ASN |
| 47 | BV | 64 | HIS |
| 47 | BV | 89 | GLN |
| 48 | BW | 34 | ASN |
| 48 | BW | 57 | ASN |
| 48 | BW | 60 | ASN |
| 48 | BW | 61 | ASN |
| 48 | BW | 62 | HIS |
| 48 | BW | 102 | HIS |
| 48 | BW | 111 | HIS |
| 49 | BX | 31 | HIS |
| 49 | BX | 55 | ASN |
| 51 | BZ | 54 | HIS |
| 51 | BZ | 55 | HIS |
| 51 | BZ | 85 | HIS |
| 51 | BZ | 151 | HIS |
| 2 | CB | 19 | HIS |
| 2 | CB | 37 | ASN |
| 2 | CB | 40 | HIS |
| 2 | CB | 113 | HIS |
| 2 | CB | 146 | GLN |
| 2 | CB | 204 | ASN |
| 3 | CC | 28 | GLN |
| 3 | CC | 69 | HIS |
| 3 | CC | 123 | GLN |
| 3 | CC | 170 | GLN |
| 4 | CD | 62 | GLN |
| 4 | CD | 123 | HIS |
| 4 | CD | 129 | ASN |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 5 | CE | 20 | GLN |
| 5 | CE | 73 | ASN |
| 5 | CE | 78 | HIS |
| 6 | CF | 18 | GLN |
| 6 | CF | 27 | GLN |
| 6 | CF | 32 | ASN |
| 6 | CF | 94 | GLN |
| 6 | CF | 100 | ASN |
| 7 | CG | 13 | GLN |
| 7 | CG | 37 | ASN |
| 7 | CG | 84 | ASN |
| 7 | CG | 106 | GLN |
| 8 | CH | 82 | HIS |
| 9 | CI | 73 | GLN |
| 9 | CI | 87 | GLN |
| 9 | CI | 117 | HIS |
| 9 | CI | 124 | GLN |
| 10 | CJ | 13 | HIS |
| 10 | CJ | 78 | ASN |
| 11 | CK | 26 | ASN |
| 12 | CL | 8 | ASN |
| 12 | CL | 9 | GLN |
| 12 | CL | 49 | ASN |
| 13 | CM | 92 | HIS |
| 15 | CO | 37 | ASN |
| 15 | CO | 46 | HIS |
| 16 | CP | 16 | HIS |
| 16 | CP | 76 | GLN |
| 16 | CP | 82 | GLN |
| 17 | CQ | 16 | GLN |
| 17 | CQ | 94 | ASN |
| 19 | CS | 53 | ASN |
| 20 | CT | 16 | HIS |
| 20 | CT | 18 | GLN |
| 22 | D0 | 29 | GLN |
| 22 | D0 | 50 | ASN |
| 23 | D1 | 19 | GLN |
| 23 | D1 | 45 | ASN |
| 23 | D1 | 66 | HIS |
| 24 | D2 | 38 | GLN |
| 24 | D2 | 46 | GLN |
| 24 | D2 | 47 | ASN |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 24 | D2 | 56 | GLN |
| 25 | D3 | 19 | GLN |
| 25 | D3 | 32 | GLN |
| 25 | D3 | 46 | ASN |
| 25 | D3 | 52 | HIS |
| 27 | D5 | 4 | HIS |
| 27 | D5 | 23 | HIS |
| 27 | D5 | 43 | HIS |
| 28 | D6 | 20 | ASN |
| 28 | D6 | 26 | ASN |
| 28 | D6 | 32 | ASN |
| 28 | D6 | 49 | HIS |
| 29 | D7 | 8 | ASN |
| 29 | D7 | 36 | GLN |
| 30 | D8 | 33 | ASN |
| 30 | D8 | 35 | GLN |
| 33 | DD | 58 | HIS |
| 33 | DD | 87 | ASN |
| 33 | DD | 96 | HIS |
| 33 | DD | 126 | GLN |
| 33 | DD | 143 | HIS |
| 33 | DD | 164 | GLN |
| 33 | DD | 166 | GLN |
| 33 | DD | 186 | HIS |
| 33 | DD | 198 | ASN |
| 33 | DD | 220 | HIS |
| 34 | DE | 48 | GLN |
| 34 | DE | 54 | GLN |
| 34 | DE | 60 | ASN |
| 34 | DE | 66 | HIS |
| 34 | DE | 85 | ASN |
| 34 | DE | 129 | HIS |
| 34 | DE | 132 | HIS |
| 34 | DE | 192 | ASN |
| 35 | DF | 69 | HIS |
| 35 | DF | 75 | HIS |
| 35 | DF | 160 | ASN |
| 35 | DF | 169 | ASN |
| 36 | DG | 26 | GLN |
| 36 | DG | 66 | GLN |
| 37 | DH | 65 | HIS |
| 38 | DI | 28 | ASN |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 38 | DI | 74 | ASN |
| 38 | DI | 105 | HIS |
| 38 | DI | 133 | HIS |
| 39 | DN | 45 | ASN |
| 39 | DN | 56 | ASN |
| 39 | DN | 101 | HIS |
| 39 | DN | 128 | HIS |
| 39 | DN | 130 | HIS |
| 40 | DO | 3 | GLN |
| 40 | DO | 82 | ASN |
| 41 | DP | 13 | ASN |
| 41 | DP | 81 | GLN |
| 42 | DQ | 13 | GLN |
| 42 | DQ | 141 | GLN |
| 43 | DR | 13 | HIS |
| 43 | DR | 16 | HIS |
| 43 | DR | 23 | ASN |
| 43 | DR | 24 | GLN |
| 43 | DR | 50 | HIS |
| 43 | DR | 53 | HIS |
| 43 | DR | 71 | GLN |
| 43 | DR | 91 | GLN |
| 44 | DS | 34 | HIS |
| 44 | DS | 61 | ASN |
| 45 | DT | 38 | ASN |
| 45 | DT | 43 | GLN |
| 45 | DT | 58 | ASN |
| 45 | DT | 90 | GLN |
| 45 | DT | 123 | GLN |
| 46 | DU | 14 | HIS |
| 46 | DU | 49 | HIS |
| 46 | DU | 71 | GLN |
| 46 | DU | 94 | ASN |
| 47 | DV | 64 | HIS |
| 47 | DV | 89 | GLN |
| 48 | DW | 34 | ASN |
| 48 | DW | 57 | ASN |
| 48 | DW | 60 | ASN |
| 48 | DW | 61 | ASN |
| 48 | DW | 62 | HIS |
| 48 | DW | 102 | HIS |
| 48 | DW | 111 | HIS |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 49 | DX | 31 | HIS |
| 49 | DX | 41 | ASN |
| 49 | DX | 55 | ASN |
| 51 | DZ | 54 | HIS |
| 51 | DZ | 55 | HIS |
| 51 | DZ | 85 | HIS |
| 51 | DZ | 118 | GLN |
| 51 | DZ | 151 | HIS |

### 5.3.3 RNA

| Mol | Chain | Analysed | Backbone Outliers | Pucker Outliers |
|---|---|---|---|---|
| 1 | AA | 1503/1522 (98%) | 287 (19%) | 31 (2%) |
| 1 | CA | 1503/1522 (98%) | 288 (19%) | 31 (2%) |
| 31 | BA | 2723/2787 (97%) | 741 (27%) | 71 (2%) |
| 31 | DA | 2723/2787 (97%) | 735 (26%) | 70 (2%) |
| 32 | BB | 118/122 (96%) | 35 (29%) | 1 (0%) |
| 32 | DB | 118/122 (96%) | 35 (29%) | 0 |
| All | All | 8688/8862 (98%) | 2121 (24%) | 204 (2%) |

All (2121) RNA backbone outliers are listed below:

| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | AA | 9 | G |
| 1 | AA | 31 | G |
| 1 | AA | 32 | A |
| 1 | AA | 39 | G |
| 1 | AA | 47 | C |
| 1 | AA | 48 | C |
| 1 | AA | 50 | A |
| 1 | AA | 51 | A |
| 1 | AA | 53 | A |
| 1 | AA | 59 | A |
| 1 | AA | 61 | G |
| 1 | AA | 67 | C |
| 1 | AA | 70 | G |
| 1 | AA | 80 | G |
| 1 | AA | 81 | U |
| 1 | AA | 90 | U |
| 1 | AA | 91 | C |
| 1 | AA | 97 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | AA | 98 | G |
| 1 | AA | 101 | A |
| 1 | AA | 116 | A |
| 1 | AA | 120 | A |
| 1 | AA | 121 | C |
| 1 | AA | 122 | G |
| 1 | AA | 131 | C |
| 1 | AA | 147 | G |
| 1 | AA | 150 | C |
| 1 | AA | 157 | G |
| 1 | AA | 158 | G |
| 1 | AA | 163 | C |
| 1 | AA | 172 | A |
| 1 | AA | 173 | U |
| 1 | AA | 189(G) | G |
| 1 | AA | 189(H) | G |
| 1 | AA | 195 | A |
| 1 | AA | 197 | A |
| 1 | AA | 199 | G |
| 1 | AA | 216 | G |
| 1 | AA | 220 | G |
| 1 | AA | 231 | G |
| 1 | AA | 232 | G |
| 1 | AA | 243 | A |
| 1 | AA | 244 | U |
| 1 | AA | 247 | G |
| 1 | AA | 251 | G |
| 1 | AA | 266 | G |
| 1 | AA | 267 | C |
| 1 | AA | 270 | A |
| 1 | AA | 281 | G |
| 1 | AA | 289 | G |
| 1 | AA | 301 | G |
| 1 | AA | 321 | A |
| 1 | AA | 327 | A |
| 1 | AA | 328 | C |
| 1 | AA | 329 | A |
| 1 | AA | 332 | G |
| 1 | AA | 342 | C |
| 1 | AA | 343 | U |
| 1 | AA | 345 | C |
| 1 | AA | 352 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | AA | 353 | A |
| 1 | AA | 354 | G |
| 1 | AA | 357 | G |
| 1 | AA | 366 | C |
| 1 | AA | 367 | U |
| 1 | AA | 371 | G |
| 1 | AA | 372 | C |
| 1 | AA | 373 | A |
| 1 | AA | 384 | G |
| 1 | AA | 388 | G |
| 1 | AA | 390 | C |
| 1 | AA | 397 | A |
| 1 | AA | 398 | C |
| 1 | AA | 406 | G |
| 1 | AA | 412 | A |
| 1 | AA | 413 | G |
| 1 | AA | 414 | A |
| 1 | AA | 415 | A |
| 1 | AA | 420 | U |
| 1 | AA | 421 | U |
| 1 | AA | 422 | C |
| 1 | AA | 424 | G |
| 1 | AA | 428 | G |
| 1 | AA | 429 | U |
| 1 | AA | 430 | A |
| 1 | AA | 437 | U |
| 1 | AA | 439 | A |
| 1 | AA | 442 | C |
| 1 | AA | 448 | A |
| 1 | AA | 452 | A |
| 1 | AA | 461 | A |
| 1 | AA | 470 | C |
| 1 | AA | 472 | A |
| 1 | AA | 482 | A |
| 1 | AA | 484 | G |
| 1 | AA | 485 | G |
| 1 | AA | 487 | A |
| 1 | AA | 495 | A |
| 1 | AA | 496 | A |
| 1 | AA | 498 | U |
| 1 | AA | 500 | G |
| 1 | AA | 501 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | AA | 505 | G |
| 1 | AA | 509 | A |
| 1 | AA | 510 | A |
| 1 | AA | 511 | C |
| 1 | AA | 513 | C |
| 1 | AA | 518 | C |
| 1 | AA | 520 | A |
| 1 | AA | 527 | G |
| 1 | AA | 531 | U |
| 1 | AA | 532 | A |
| 1 | AA | 533 | A |
| 1 | AA | 534 | U |
| 1 | AA | 545 | C |
| 1 | AA | 547 | A |
| 1 | AA | 559 | A |
| 1 | AA | 560 | U |
| 1 | AA | 561 | U |
| 1 | AA | 562 | C |
| 1 | AA | 572 | A |
| 1 | AA | 573 | A |
| 1 | AA | 576 | G |
| 1 | AA | 577 | G |
| 1 | AA | 588 | G |
| 1 | AA | 596 | C |
| 1 | AA | 616 | G |
| 1 | AA | 618 | C |
| 1 | AA | 623 | C |
| 1 | AA | 630 | G |
| 1 | AA | 632 | A |
| 1 | AA | 653 | A |
| 1 | AA | 665 | A |
| 1 | AA | 687 | A |
| 1 | AA | 688 | G |
| 1 | AA | 693 | G |
| 1 | AA | 702 | A |
| 1 | AA | 703 | G |
| 1 | AA | 720 | C |
| 1 | AA | 721 | G |
| 1 | AA | 731 | G |
| 1 | AA | 733 | A |
| 1 | AA | 749 | C |
| 1 | AA | 753 | A |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | AA | 755 | G |
| 1 | AA | 766 | A |
| 1 | AA | 776 | G |
| 1 | AA | 777 | A |
| 1 | AA | 786 | G |
| 1 | AA | 792 | A |
| 1 | AA | 793 | U |
| 1 | AA | 794 | A |
| 1 | AA | 802 | A |
| 1 | AA | 816 | A |
| 1 | AA | 817 | C |
| 1 | AA | 819 | A |
| 1 | AA | 821 | G |
| 1 | AA | 827 | U |
| 1 | AA | 828 | A |
| 1 | AA | 833 | U |
| 1 | AA | 839 | U |
| 1 | AA | 840 | C |
| 1 | AA | 841 | U |
| 1 | AA | 848 | C |
| 1 | AA | 859 | A |
| 1 | AA | 864 | A |
| 1 | AA | 870 | U |
| 1 | AA | 872 | A |
| 1 | AA | 884 | U |
| 1 | AA | 885 | G |
| 1 | AA | 902 | G |
| 1 | AA | 914 | A |
| 1 | AA | 920 | U |
| 1 | AA | 926 | G |
| 1 | AA | 927 | G |
| 1 | AA | 932 | C |
| 1 | AA | 934 | C |
| 1 | AA | 935 | A |
| 1 | AA | 940 | C |
| 1 | AA | 960 | U |
| 1 | AA | 961 | U |
| 1 | AA | 968 | A |
| 1 | AA | 969 | A |
| 1 | AA | 971 | G |
| 1 | AA | 972 | C |
| 1 | AA | 974 | A |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | AA | 975 | A |
| 1 | AA | 976 | G |
| 1 | AA | 977 | A |
| 1 | AA | 978 | A |
| 1 | AA | 980 | C |
| 1 | AA | 991 | U |
| 1 | AA | 992 | U |
| 1 | AA | 993 | G |
| 1 | AA | 1005 | A |
| 1 | AA | 1026 | G |
| 1 | AA | 1050 | G |
| 1 | AA | 1054 | C |
| 1 | AA | 1064 | G |
| 1 | AA | 1065 | U |
| 1 | AA | 1066 | C |
| 1 | AA | 1068 | G |
| 1 | AA | 1081 | G |
| 1 | AA | 1094 | G |
| 1 | AA | 1095 | U |
| 1 | AA | 1101 | A |
| 1 | AA | 1117 | G |
| 1 | AA | 1118 | C |
| 1 | AA | 1124 | G |
| 1 | AA | 1125 | U |
| 1 | AA | 1126 | U |
| 1 | AA | 1129 | C |
| 1 | AA | 1136 | U |
| 1 | AA | 1137 | C |
| 1 | AA | 1138 | G |
| 1 | AA | 1139 | G |
| 1 | AA | 1146 | A |
| 1 | AA | 1152 | A |
| 1 | AA | 1159 | U |
| 1 | AA | 1160 | G |
| 1 | AA | 1190 | G |
| 1 | AA | 1196 | U |
| 1 | AA | 1197 | G |
| 1 | AA | 1202 | G |
| 1 | AA | 1212 | U |
| 1 | AA | 1213 | A |
| 1 | AA | 1225 | A |
| 1 | AA | 1227 | A |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | AA | 1238 | A |
| 1 | AA | 1239 | A |
| 1 | AA | 1249 | C |
| 1 | AA | 1255 | G |
| 1 | AA | 1256 | A |
| 1 | AA | 1257 | U |
| 1 | AA | 1258 | G |
| 1 | AA | 1270 | C |
| 1 | AA | 1280 | A |
| 1 | AA | 1281 | U |
| 1 | AA | 1286 | A |
| 1 | AA | 1287 | A |
| 1 | AA | 1290 | G |
| 1 | AA | 1294 | G |
| 1 | AA | 1300 | G |
| 1 | AA | 1301 | U |
| 1 | AA | 1302 | U |
| 1 | AA | 1305 | G |
| 1 | AA | 1312 | G |
| 1 | AA | 1317 | C |
| 1 | AA | 1320 | C |
| 1 | AA | 1322 | C |
| 1 | AA | 1323 | G |
| 1 | AA | 1331 | G |
| 1 | AA | 1335 | C |
| 1 | AA | 1338 | G |
| 1 | AA | 1347 | G |
| 1 | AA | 1363 | C |
| 1 | AA | 1363(A) | A |
| 1 | AA | 1364 | U |
| 1 | AA | 1370 | G |
| 1 | AA | 1388 | C |
| 1 | AA | 1397 | C |
| 1 | AA | 1402 | C |
| 1 | AA | 1419 | G |
| 1 | AA | 1442 | G |
| 1 | AA | 1442(A) | G |
| 1 | AA | 1442(B) | A |
| 1 | AA | 1443 | G |
| 1 | AA | 1447 | A |
| 1 | AA | 1452 | C |
| 1 | AA | 1456 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | AA | 1487 | G |
| 1 | AA | 1492 | A |
| 1 | AA | 1493 | A |
| 1 | AA | 1494 | G |
| 1 | AA | 1495 | U |
| 1 | AA | 1497 | G |
| 1 | AA | 1499 | A |
| 1 | AA | 1502 | A |
| 1 | AA | 1504 | G |
| 1 | AA | 1505 | G |
| 1 | AA | 1506 | U |
| 1 | AA | 1507 | A |
| 1 | AA | 1517 | G |
| 1 | AA | 1519 | A |
| 1 | AA | 1520 | G |
| 1 | AA | 1529 | G |
| 1 | AA | 1530 | G |
| 31 | BA | 9 | U |
| 31 | BA | 10 | G |
| 31 | BA | 11 | G |
| 31 | BA | 15 | G |
| 31 | BA | 23 | G |
| 31 | BA | 34 | C |
| 31 | BA | 35 | G |
| 31 | BA | 45 | C |
| 31 | BA | 50 | U |
| 31 | BA | 51 | G |
| 31 | BA | 55 | G |
| 31 | BA | 71 | A |
| 31 | BA | 72 | U |
| 31 | BA | 74 | A |
| 31 | BA | 75 | G |
| 31 | BA | 78 | A |
| 31 | BA | 83 | G |
| 31 | BA | 84 | A |
| 31 | BA | 90 | U |
| 31 | BA | 92 | A |
| 31 | BA | 94 | C |
| 31 | BA | 95 | G |
| 31 | BA | 99 | U |
| 31 | BA | 100 | G |
| 31 | BA | 102 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 103 | A |
| 31 | BA | 105 | C |
| 31 | BA | 117 | G |
| 31 | BA | 118 | A |
| 31 | BA | 119 | A |
| 31 | BA | 120 | U |
| 31 | BA | 129 | C |
| 31 | BA | 131 | G |
| 31 | BA | 137 | C |
| 31 | BA | 139(A) | G |
| 31 | BA | 140 | G |
| 31 | BA | 142 | A |
| 31 | BA | 142(A) | C |
| 31 | BA | 146 | G |
| 31 | BA | 149 | A |
| 31 | BA | 154(A) | C |
| 31 | BA | 157 | U |
| 31 | BA | 158 | U |
| 31 | BA | 171 | G |
| 31 | BA | 173 | G |
| 31 | BA | 175 | G |
| 31 | BA | 196 | A |
| 31 | BA | 197 | A |
| 31 | BA | 198 | C |
| 31 | BA | 199 | A |
| 31 | BA | 204 | A |
| 31 | BA | 205 | G |
| 31 | BA | 215 | G |
| 31 | BA | 216 | A |
| 31 | BA | 221 | A |
| 31 | BA | 222 | A |
| 31 | BA | 225 | A |
| 31 | BA | 228 | A |
| 31 | BA | 229 | A |
| 31 | BA | 233 | A |
| 31 | BA | 248 | G |
| 31 | BA | 250 | G |
| 31 | BA | 252 | G |
| 31 | BA | 267 | C |
| 31 | BA | 268 | C |
| 31 | BA | 269 | U |
| 31 | BA | 271(I) | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 271(J) | C |
| 31 | BA | 271(K) | U |
| 31 | BA | 271(L) | U |
| 31 | BA | 271(M) | G |
| 31 | BA | 271(N) | U |
| 31 | BA | 271(O) | C |
| 31 | BA | 271(P) | C |
| 31 | BA | 271(R) | G |
| 31 | BA | 271(T) | C |
| 31 | BA | 271(Y) | U |
| 31 | BA | 272 | G |
| 31 | BA | 272(B) | G |
| 31 | BA | 272(H) | C |
| 31 | BA | 272(I) | U |
| 31 | BA | 272(J) | C |
| 31 | BA | 274 | G |
| 31 | BA | 275 | G |
| 31 | BA | 279 | C |
| 31 | BA | 281 | G |
| 31 | BA | 283 | A |
| 31 | BA | 286 | C |
| 31 | BA | 287 | C |
| 31 | BA | 298 | G |
| 31 | BA | 311 | A |
| 31 | BA | 317 | G |
| 31 | BA | 327 | G |
| 31 | BA | 329 | G |
| 31 | BA | 330 | A |
| 31 | BA | 332 | A |
| 31 | BA | 343 | C |
| 31 | BA | 347 | A |
| 31 | BA | 352 | G |
| 31 | BA | 353 | G |
| 31 | BA | 362 | U |
| 31 | BA | 363 | G |
| 31 | BA | 363(B) | G |
| 31 | BA | 363(C) | G |
| 31 | BA | 363(E) | U |
| 31 | BA | 363(F) | A |
| 31 | BA | 366 | C |
| 31 | BA | 372 | G |
| 31 | BA | 386 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 388 | G |
| 31 | BA | 396 | G |
| 31 | BA | 405 | U |
| 31 | BA | 407 | G |
| 31 | BA | 411 | G |
| 31 | BA | 412 | A |
| 31 | BA | 415 | A |
| 31 | BA | 418 | G |
| 31 | BA | 428 | A |
| 31 | BA | 444 | C |
| 31 | BA | 448 | U |
| 31 | BA | 455 | C |
| 31 | BA | 457 | A |
| 31 | BA | 468 | G |
| 31 | BA | 470 | A |
| 31 | BA | 471 | A |
| 31 | BA | 472 | A |
| 31 | BA | 473 | G |
| 31 | BA | 474 | G |
| 31 | BA | 475 | U |
| 31 | BA | 481 | G |
| 31 | BA | 482 | A |
| 31 | BA | 494 | G |
| 31 | BA | 495 | G |
| 31 | BA | 504 | U |
| 31 | BA | 505 | A |
| 31 | BA | 508 | G |
| 31 | BA | 509 | C |
| 31 | BA | 511 | U |
| 31 | BA | 512 | G |
| 31 | BA | 518 | G |
| 31 | BA | 529 | A |
| 31 | BA | 530 | G |
| 31 | BA | 531 | C |
| 31 | BA | 532 | A |
| 31 | BA | 533 | G |
| 31 | BA | 536 | A |
| 31 | BA | 537 | C |
| 31 | BA | 541 | C |
| 31 | BA | 542 | C |
| 31 | BA | 543 | C |
| 31 | BA | 547 | A |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 548 | A |
| 31 | BA | 549 | G |
| 31 | BA | 551 | G |
| 31 | BA | 563 | G |
| 31 | BA | 571 | A |
| 31 | BA | 573 | G |
| 31 | BA | 575 | A |
| 31 | BA | 586 | A |
| 31 | BA | 588 | U |
| 31 | BA | 603 | A |
| 31 | BA | 604 | G |
| 31 | BA | 607 | U |
| 31 | BA | 614(B) | G |
| 31 | BA | 615 | G |
| 31 | BA | 619 | G |
| 31 | BA | 622 | G |
| 31 | BA | 626 | U |
| 31 | BA | 627 | A |
| 31 | BA | 634 | C |
| 31 | BA | 637 | A |
| 31 | BA | 645 | C |
| 31 | BA | 646 | A |
| 31 | BA | 647 | G |
| 31 | BA | 649 | G |
| 31 | BA | 650 | C |
| 31 | BA | 651 | G |
| 31 | BA | 652 | C |
| 31 | BA | 656 | G |
| 31 | BA | 657 | U |
| 31 | BA | 659 | C |
| 31 | BA | 668 | G |
| 31 | BA | 669 | G |
| 31 | BA | 670 | A |
| 31 | BA | 671 | C |
| 31 | BA | 686 | G |
| 31 | BA | 701 | G |
| 31 | BA | 708 | C |
| 31 | BA | 717 | G |
| 31 | BA | 721 | C |
| 31 | BA | 722 | A |
| 31 | BA | 726 | G |
| 31 | BA | 730 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 745 | G |
| 31 | BA | 752 | A |
| 31 | BA | 753 | C |
| 31 | BA | 774 | A |
| 31 | BA | 775 | G |
| 31 | BA | 776 | G |
| 31 | BA | 779 | U |
| 31 | BA | 782 | A |
| 31 | BA | 784 | A |
| 31 | BA | 785 | G |
| 31 | BA | 790 | C |
| 31 | BA | 791 | C |
| 31 | BA | 792 | G |
| 31 | BA | 805 | G |
| 31 | BA | 808 | G |
| 31 | BA | 812 | C |
| 31 | BA | 819 | A |
| 31 | BA | 820 | A |
| 31 | BA | 826 | U |
| 31 | BA | 827 | U |
| 31 | BA | 828 | U |
| 31 | BA | 829 | A |
| 31 | BA | 830 | G |
| 31 | BA | 831 | G |
| 31 | BA | 832 | G |
| 31 | BA | 834 | C |
| 31 | BA | 847 | U |
| 31 | BA | 848 | G |
| 31 | BA | 856 | C |
| 31 | BA | 857 | C |
| 31 | BA | 859 | G |
| 31 | BA | 867 | C |
| 31 | BA | 870 | A |
| 31 | BA | 872 | A |
| 31 | BA | 878 | A |
| 31 | BA | 879 | G |
| 31 | BA | 883 | G |
| 31 | BA | 884 | C |
| 31 | BA | 892 | G |
| 31 | BA | 894 | C |
| 31 | BA | 896 | A |
| 31 | BA | 897 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 898 | C |
| 31 | BA | 899 | A |
| 31 | BA | 900 | A |
| 31 | BA | 901 | A |
| 31 | BA | 902 | C |
| 31 | BA | 910 | A |
| 31 | BA | 911 | A |
| 31 | BA | 914 | C |
| 31 | BA | 917 | A |
| 31 | BA | 919 | G |
| 31 | BA | 923 | C |
| 31 | BA | 926 | A |
| 31 | BA | 932 | G |
| 31 | BA | 938 | G |
| 31 | BA | 941 | A |
| 31 | BA | 945 | A |
| 31 | BA | 946 | G |
| 31 | BA | 955 | C |
| 31 | BA | 958 | U |
| 31 | BA | 959 | A |
| 31 | BA | 961 | C |
| 31 | BA | 971 | C |
| 31 | BA | 974 | G |
| 31 | BA | 975 | C |
| 31 | BA | 975(A) | G |
| 31 | BA | 983 | A |
| 31 | BA | 990 | A |
| 31 | BA | 991 | C |
| 31 | BA | 993 | G |
| 31 | BA | 996 | A |
| 31 | BA | 1005 | C |
| 31 | BA | 1009 | A |
| 31 | BA | 1011 | G |
| 31 | BA | 1012 | U |
| 31 | BA | 1013 | C |
| 31 | BA | 1015 | G |
| 31 | BA | 1016 | G |
| 31 | BA | 1020 | A |
| 31 | BA | 1021 | A |
| 31 | BA | 1022 | G |
| 31 | BA | 1023 | U |
| 31 | BA | 1025 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 1026 | U |
| 31 | BA | 1031 | G |
| 31 | BA | 1032 | A |
| 31 | BA | 1033 | U |
| 31 | BA | 1038 | C |
| 31 | BA | 1039 | G |
| 31 | BA | 1041 | C |
| 31 | BA | 1042 | G |
| 31 | BA | 1043 | C |
| 31 | BA | 1044 | G |
| 31 | BA | 1045 | A |
| 31 | BA | 1047 | G |
| 31 | BA | 1048 | A |
| 31 | BA | 1049 | C |
| 31 | BA | 1051 | G |
| 31 | BA | 1052 | C |
| 31 | BA | 1053 | C |
| 31 | BA | 1106 | A |
| 31 | BA | 1107 | G |
| 31 | BA | 1110 | G |
| 31 | BA | 1111 | A |
| 31 | BA | 1112 | G |
| 31 | BA | 1113 | U |
| 31 | BA | 1116 | C |
| 31 | BA | 1118 | C |
| 31 | BA | 1130 | U |
| 31 | BA | 1135 | C |
| 31 | BA | 1136 | G |
| 31 | BA | 1139 | G |
| 31 | BA | 1142 | U |
| 31 | BA | 1148 | A |
| 31 | BA | 1155 | A |
| 31 | BA | 1156 | A |
| 31 | BA | 1159 | U |
| 31 | BA | 1169 | G |
| 31 | BA | 1171 | G |
| 31 | BA | 1173 | G |
| 31 | BA | 1174 | A |
| 31 | BA | 1175 | U |
| 31 | BA | 1176 | G |
| 31 | BA | 1177 | A |
| 31 | BA | 1178 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 1179 | C |
| 31 | BA | 1180 | C |
| 31 | BA | 1195 | G |
| 31 | BA | 1210 | A |
| 31 | BA | 1211 | U |
| 31 | BA | 1220 | A |
| 31 | BA | 1242 | A |
| 31 | BA | 1251 | C |
| 31 | BA | 1252 | G |
| 31 | BA | 1253 | A |
| 31 | BA | 1255 | U |
| 31 | BA | 1256 | G |
| 31 | BA | 1265 | A |
| 31 | BA | 1271 | G |
| 31 | BA | 1272 | A |
| 31 | BA | 1273 | U |
| 31 | BA | 1287 | A |
| 31 | BA | 1288 | U |
| 31 | BA | 1300 | U |
| 31 | BA | 1301 | A |
| 31 | BA | 1302 | A |
| 31 | BA | 1305 | C |
| 31 | BA | 1307 | A |
| 31 | BA | 1308 | A |
| 31 | BA | 1310 | G |
| 31 | BA | 1313 | U |
| 31 | BA | 1314 | C |
| 31 | BA | 1317 | A |
| 31 | BA | 1318 | C |
| 31 | BA | 1319 | G |
| 31 | BA | 1345 | C |
| 31 | BA | 1349 | A |
| 31 | BA | 1359 | A |
| 31 | BA | 1360 | A |
| 31 | BA | 1364 | G |
| 31 | BA | 1365 | A |
| 31 | BA | 1368 | G |
| 31 | BA | 1370 | C |
| 31 | BA | 1379 | A |
| 31 | BA | 1380 | G |
| 31 | BA | 1384 | A |
| 31 | BA | 1385 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 1386 | C |
| 31 | BA | 1389 | G |
| 31 | BA | 1404 | C |
| 31 | BA | 1416 | G |
| 31 | BA | 1417 | C |
| 31 | BA | 1420 | U |
| 31 | BA | 1421 | G |
| 31 | BA | 1427 | A |
| 31 | BA | 1428 | C |
| 31 | BA | 1437 | C |
| 31 | BA | 1445 | A |
| 31 | BA | 1449 | A |
| 31 | BA | 1450 | G |
| 31 | BA | 1450(A) | C |
| 31 | BA | 1455 | G |
| 31 | BA | 1458 | C |
| 31 | BA | 1461 | G |
| 31 | BA | 1467 | C |
| 31 | BA | 1469 | A |
| 31 | BA | 1470 | G |
| 31 | BA | 1471 | A |
| 31 | BA | 1475 | G |
| 31 | BA | 1478 | G |
| 31 | BA | 1479 | G |
| 31 | BA | 1480 | G |
| 31 | BA | 1481 | U |
| 31 | BA | 1482 | G |
| 31 | BA | 1484 | G |
| 31 | BA | 1485 | G |
| 31 | BA | 1490 | A |
| 31 | BA | 1491 | G |
| 31 | BA | 1493 | C |
| 31 | BA | 1494 | A |
| 31 | BA | 1495 | A |
| 31 | BA | 1497 | U |
| 31 | BA | 1498 | C |
| 31 | BA | 1502 | C |
| 31 | BA | 1503 | U |
| 31 | BA | 1505 | C |
| 31 | BA | 1508 | A |
| 31 | BA | 1509 | C |
| 31 | BA | 1509(A) | A |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 1511 | C |
| 31 | BA | 1512 | U |
| 31 | BA | 1520 | G |
| 31 | BA | 1526 | G |
| 31 | BA | 1528 | A |
| 31 | BA | 1528(A) | A |
| 31 | BA | 1529 | G |
| 31 | BA | 1530 | C |
| 31 | BA | 1531 | C |
| 31 | BA | 1532 | C |
| 31 | BA | 1533 | G |
| 31 | BA | 1543 | C |
| 31 | BA | 1545 | A |
| 31 | BA | 1546 | C |
| 31 | BA | 1554 | A |
| 31 | BA | 1558 | A |
| 31 | BA | 1559 | G |
| 31 | BA | 1569 | A |
| 31 | BA | 1578 | U |
| 31 | BA | 1579 | A |
| 31 | BA | 1580 | A |
| 31 | BA | 1581 | G |
| 31 | BA | 1584 | C |
| 31 | BA | 1586 | A |
| 31 | BA | 1588 | C |
| 31 | BA | 1591 | G |
| 31 | BA | 1598 | C |
| 31 | BA | 1603 | A |
| 31 | BA | 1608 | A |
| 31 | BA | 1609 | A |
| 31 | BA | 1610 | A |
| 31 | BA | 1617 | C |
| 31 | BA | 1618 | A |
| 31 | BA | 1632 | A |
| 31 | BA | 1635 | G |
| 31 | BA | 1636 | C |
| 31 | BA | 1640 | C |
| 31 | BA | 1646 | C |
| 31 | BA | 1648 | C |
| 31 | BA | 1652 | A |
| 31 | BA | 1653 | G |
| 31 | BA | 1654 | A |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 1674 | G |
| 31 | BA | 1675 | C |
| 31 | BA | 1686 | C |
| 31 | BA | 1687 | G |
| 31 | BA | 1694 | C |
| 31 | BA | 1695 | G |
| 31 | BA | 1696 | G |
| 31 | BA | 1697 | G |
| 31 | BA | 1700 | A |
| 31 | BA | 1703 | G |
| 31 | BA | 1722 | A |
| 31 | BA | 1739 | U |
| 31 | BA | 1741 | A |
| 31 | BA | 1742 | G |
| 31 | BA | 1744 | C |
| 31 | BA | 1745 | C |
| 31 | BA | 1746 | G |
| 31 | BA | 1748 | G |
| 31 | BA | 1750 | G |
| 31 | BA | 1756 | G |
| 31 | BA | 1758 | G |
| 31 | BA | 1763 | G |
| 31 | BA | 1764 | G |
| 31 | BA | 1773 | A |
| 31 | BA | 1780 | A |
| 31 | BA | 1782 | C |
| 31 | BA | 1791 | A |
| 31 | BA | 1799 | G |
| 31 | BA | 1800 | C |
| 31 | BA | 1801 | G |
| 31 | BA | 1816 | G |
| 31 | BA | 1819 | A |
| 31 | BA | 1820 | U |
| 31 | BA | 1822 | G |
| 31 | BA | 1829 | A |
| 31 | BA | 1835 | G |
| 31 | BA | 1838 | C |
| 31 | BA | 1839 | G |
| 31 | BA | 1847 | A |
| 31 | BA | 1858 | G |
| 31 | BA | 1865 | G |
| 31 | BA | 1866 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 1877 | A |
| 31 | BA | 1878 | G |
| 31 | BA | 1882 | C |
| 31 | BA | 1883 | G |
| 31 | BA | 1885 | A |
| 31 | BA | 1888 | G |
| 31 | BA | 1889 | A |
| 31 | BA | 1900 | A |
| 31 | BA | 1902 | C |
| 31 | BA | 1903 | G |
| 31 | BA | 1904 | G |
| 31 | BA | 1905 | C |
| 31 | BA | 1906 | G |
| 31 | BA | 1913 | A |
| 31 | BA | 1914 | C |
| 31 | BA | 1916 | A |
| 31 | BA | 1929 | G |
| 31 | BA | 1930 | G |
| 31 | BA | 1931 | U |
| 31 | BA | 1934 | C |
| 31 | BA | 1935 | G |
| 31 | BA | 1936 | A |
| 31 | BA | 1938 | A |
| 31 | BA | 1946 | U |
| 31 | BA | 1955 | U |
| 31 | BA | 1962 | C |
| 31 | BA | 1963 | U |
| 31 | BA | 1964 | G |
| 31 | BA | 1967 | C |
| 31 | BA | 1969 | A |
| 31 | BA | 1970 | A |
| 31 | BA | 1971 | A |
| 31 | BA | 1972 | A |
| 31 | BA | 1982 | C |
| 31 | BA | 1983 | C |
| 31 | BA | 1987 | G |
| 31 | BA | 1988 | C |
| 31 | BA | 1990 | C |
| 31 | BA | 1991 | U |
| 31 | BA | 1993 | U |
| 31 | BA | 1997 | G |
| 31 | BA | 2018 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 2020 | A |
| 31 | BA | 2023 | G |
| 31 | BA | 2030 | A |
| 31 | BA | 2031 | A |
| 31 | BA | 2033 | A |
| 31 | BA | 2036 | C |
| 31 | BA | 2043 | C |
| 31 | BA | 2055 | C |
| 31 | BA | 2056 | G |
| 31 | BA | 2060 | A |
| 31 | BA | 2061 | G |
| 31 | BA | 2062 | A |
| 31 | BA | 2069 | G |
| 31 | BA | 2093 | G |
| 31 | BA | 2096 | U |
| 31 | BA | 2100 | G |
| 31 | BA | 2103 | C |
| 31 | BA | 2104 | G |
| 31 | BA | 2187 | G |
| 31 | BA | 2190 | G |
| 31 | BA | 2191 | G |
| 31 | BA | 2192 | G |
| 31 | BA | 2197 | U |
| 31 | BA | 2198 | A |
| 31 | BA | 2199 | A |
| 31 | BA | 2200 | C |
| 31 | BA | 2201 | C |
| 31 | BA | 2203 | U |
| 31 | BA | 2206 | G |
| 31 | BA | 2207 | G |
| 31 | BA | 2208 | A |
| 31 | BA | 2218 | U |
| 31 | BA | 2219 | G |
| 31 | BA | 2225 | A |
| 31 | BA | 2226 | C |
| 31 | BA | 2227 | A |
| 31 | BA | 2234 | G |
| 31 | BA | 2238 | G |
| 31 | BA | 2239 | G |
| 31 | BA | 2240 | C |
| 31 | BA | 2246 | G |
| 31 | BA | 2263 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 2268 | A |
| 31 | BA | 2273 | A |
| 31 | BA | 2274 | A |
| 31 | BA | 2275 | C |
| 31 | BA | 2283 | C |
| 31 | BA | 2287 | A |
| 31 | BA | 2288 | A |
| 31 | BA | 2289 | G |
| 31 | BA | 2291 | U |
| 31 | BA | 2302 | G |
| 31 | BA | 2303 | G |
| 31 | BA | 2305 | A |
| 31 | BA | 2307 | G |
| 31 | BA | 2308 | G |
| 31 | BA | 2309 | A |
| 31 | BA | 2311 | A |
| 31 | BA | 2319 | G |
| 31 | BA | 2320 | A |
| 31 | BA | 2325 | G |
| 31 | BA | 2334 | G |
| 31 | BA | 2335 | A |
| 31 | BA | 2336 | A |
| 31 | BA | 2340 | G |
| 31 | BA | 2345 | G |
| 31 | BA | 2346 | A |
| 31 | BA | 2347 | C |
| 31 | BA | 2350 | C |
| 31 | BA | 2354 | G |
| 31 | BA | 2359 | C |
| 31 | BA | 2360 | A |
| 31 | BA | 2361 | A |
| 31 | BA | 2377 | A |
| 31 | BA | 2383 | G |
| 31 | BA | 2385 | C |
| 31 | BA | 2388 | A |
| 31 | BA | 2394 | C |
| 31 | BA | 2395 | C |
| 31 | BA | 2396 | G |
| 31 | BA | 2399 | G |
| 31 | BA | 2400 | G |
| 31 | BA | 2402 | C |
| 31 | BA | 2403 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 2405 | G |
| 31 | BA | 2406 | U |
| 31 | BA | 2410 | G |
| 31 | BA | 2411 | A |
| 31 | BA | 2422 | A |
| 31 | BA | 2423 | U |
| 31 | BA | 2425 | A |
| 31 | BA | 2427 | C |
| 31 | BA | 2429 | G |
| 31 | BA | 2430 | A |
| 31 | BA | 2435 | A |
| 31 | BA | 2439 | A |
| 31 | BA | 2440 | C |
| 31 | BA | 2441 | C |
| 31 | BA | 2448 | A |
| 31 | BA | 2464 | C |
| 31 | BA | 2465 | C |
| 31 | BA | 2468 | G |
| 31 | BA | 2469 | A |
| 31 | BA | 2470 | G |
| 31 | BA | 2471 | C |
| 31 | BA | 2474 | C |
| 31 | BA | 2475 | C |
| 31 | BA | 2476 | A |
| 31 | BA | 2477 | C |
| 31 | BA | 2478 | A |
| 31 | BA | 2482 | G |
| 31 | BA | 2484 | G |
| 31 | BA | 2489 | G |
| 31 | BA | 2494 | G |
| 31 | BA | 2502 | G |
| 31 | BA | 2504 | U |
| 31 | BA | 2505 | G |
| 31 | BA | 2518 | A |
| 31 | BA | 2520 | C |
| 31 | BA | 2524 | G |
| 31 | BA | 2529 | G |
| 31 | BA | 2535 | G |
| 31 | BA | 2543 | G |
| 31 | BA | 2544 | G |
| 31 | BA | 2550 | G |
| 31 | BA | 2554 | U |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 2556 | C |
| 31 | BA | 2566 | A |
| 31 | BA | 2567 | G |
| 31 | BA | 2569 | G |
| 31 | BA | 2572 | A |
| 31 | BA | 2573 | C |
| 31 | BA | 2577 | A |
| 31 | BA | 2586 | C |
| 31 | BA | 2602 | A |
| 31 | BA | 2603 | G |
| 31 | BA | 2608 | G |
| 31 | BA | 2609 | U |
| 31 | BA | 2610 | C |
| 31 | BA | 2611 | U |
| 31 | BA | 2612 | C |
| 31 | BA | 2615 | U |
| 31 | BA | 2630 | G |
| 31 | BA | 2636 | U |
| 31 | BA | 2637 | U |
| 31 | BA | 2638 | G |
| 31 | BA | 2654 | A |
| 31 | BA | 2655 | G |
| 31 | BA | 2658 | C |
| 31 | BA | 2659 | G |
| 31 | BA | 2660 | A |
| 31 | BA | 2661 | G |
| 31 | BA | 2662 | A |
| 31 | BA | 2663 | G |
| 31 | BA | 2673 | G |
| 31 | BA | 2680 | C |
| 31 | BA | 2686 | G |
| 31 | BA | 2690 | C |
| 31 | BA | 2691 | C |
| 31 | BA | 2702 | U |
| 31 | BA | 2703 | C |
| 31 | BA | 2707 | G |
| 31 | BA | 2712 | U |
| 31 | BA | 2712(A) | A |
| 31 | BA | 2713 | A |
| 31 | BA | 2714 | G |
| 31 | BA | 2718 | G |
| 31 | BA | 2726 | U |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 2733 | A |
| 31 | BA | 2734 | A |
| 31 | BA | 2752 | C |
| 31 | BA | 2754 | U |
| 31 | BA | 2756 | U |
| 31 | BA | 2757 | A |
| 31 | BA | 2758 | A |
| 31 | BA | 2759 | G |
| 31 | BA | 2762 | G |
| 31 | BA | 2764 | A |
| 31 | BA | 2765 | A |
| 31 | BA | 2766 | G |
| 31 | BA | 2778 | A |
| 31 | BA | 2779 | U |
| 31 | BA | 2780 | G |
| 31 | BA | 2781 | A |
| 31 | BA | 2782 | G |
| 31 | BA | 2789 | C |
| 31 | BA | 2790 | A |
| 31 | BA | 2791 | C |
| 31 | BA | 2794 | C |
| 31 | BA | 2795 | G |
| 31 | BA | 2801 | A |
| 31 | BA | 2801(A) | A |
| 31 | BA | 2802 | G |
| 31 | BA | 2803 | C |
| 31 | BA | 2808 | U |
| 31 | BA | 2813 | A |
| 31 | BA | 2818 | G |
| 31 | BA | 2820 | A |
| 31 | BA | 2821 | A |
| 31 | BA | 2830 | G |
| 31 | BA | 2833 | G |
| 31 | BA | 2835 | A |
| 31 | BA | 2838 | G |
| 31 | BA | 2844 | G |
| 31 | BA | 2845 | G |
| 31 | BA | 2860 | A |
| 31 | BA | 2872 | G |
| 31 | BA | 2875 | C |
| 31 | BA | 2880 | C |
| 31 | BA | 2892 | A |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 2894 | G |
| 31 | BA | 2895 | U |
| 32 | BB | 3 | C |
| 32 | BB | 7 | G |
| 32 | BB | 8 | U |
| 32 | BB | 13 | A |
| 32 | BB | 15 | A |
| 32 | BB | 16 | G |
| 32 | BB | 20 | C |
| 32 | BB | 22 | U |
| 32 | BB | 26 | A |
| 32 | BB | 27 | C |
| 32 | BB | 29 | A |
| 32 | BB | 33 | G |
| 32 | BB | 40 | U |
| 32 | BB | 42 | C |
| 32 | BB | 45 | A |
| 32 | BB | 46 | A |
| 32 | BB | 47 | C |
| 32 | BB | 52 | A |
| 32 | BB | 53 | A |
| 32 | BB | 56 | G |
| 32 | BB | 65 | C |
| 32 | BB | 67 | G |
| 32 | BB | 73 | A |
| 32 | BB | 75 | G |
| 32 | BB | 81 | G |
| 32 | BB | 85 | G |
| 32 | BB | 88 | C |
| 32 | BB | 90 | A |
| 32 | BB | 91 | C |
| 32 | BB | 102 | A |
| 32 | BB | 106 | G |
| 32 | BB | 110 | G |
| 32 | BB | 116 | G |
| 32 | BB | 118 | G |
| 32 | BB | 119 | G |
| 1 | CA | 9 | G |
| 1 | CA | 31 | G |
| 1 | CA | 32 | A |
| 1 | CA | 39 | G |
| 1 | CA | 47 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | CA | 48 | C |
| 1 | CA | 50 | A |
| 1 | CA | 51 | A |
| 1 | CA | 59 | A |
| 1 | CA | 61 | G |
| 1 | CA | 70 | G |
| 1 | CA | 80 | G |
| 1 | CA | 81 | U |
| 1 | CA | 90 | U |
| 1 | CA | 91 | C |
| 1 | CA | 97 | G |
| 1 | CA | 98 | G |
| 1 | CA | 101 | A |
| 1 | CA | 116 | A |
| 1 | CA | 120 | A |
| 1 | CA | 121 | C |
| 1 | CA | 122 | G |
| 1 | CA | 131 | C |
| 1 | CA | 147 | G |
| 1 | CA | 150 | C |
| 1 | CA | 157 | G |
| 1 | CA | 158 | G |
| 1 | CA | 163 | C |
| 1 | CA | 172 | A |
| 1 | CA | 173 | U |
| 1 | CA | 189(G) | G |
| 1 | CA | 189(H) | G |
| 1 | CA | 195 | A |
| 1 | CA | 197 | A |
| 1 | CA | 199 | G |
| 1 | CA | 216 | G |
| 1 | CA | 220 | G |
| 1 | CA | 231 | G |
| 1 | CA | 243 | A |
| 1 | CA | 244 | U |
| 1 | CA | 247 | G |
| 1 | CA | 251 | G |
| 1 | CA | 266 | G |
| 1 | CA | 267 | C |
| 1 | CA | 270 | A |
| 1 | CA | 281 | G |
| 1 | CA | 289 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | CA | 301 | G |
| 1 | CA | 321 | A |
| 1 | CA | 327 | A |
| 1 | CA | 328 | C |
| 1 | CA | 329 | A |
| 1 | CA | 332 | G |
| 1 | CA | 342 | C |
| 1 | CA | 343 | U |
| 1 | CA | 345 | C |
| 1 | CA | 352 | C |
| 1 | CA | 353 | A |
| 1 | CA | 354 | G |
| 1 | CA | 357 | G |
| 1 | CA | 366 | C |
| 1 | CA | 367 | U |
| 1 | CA | 371 | G |
| 1 | CA | 372 | C |
| 1 | CA | 373 | A |
| 1 | CA | 384 | G |
| 1 | CA | 388 | G |
| 1 | CA | 390 | C |
| 1 | CA | 397 | A |
| 1 | CA | 398 | C |
| 1 | CA | 406 | G |
| 1 | CA | 412 | A |
| 1 | CA | 413 | G |
| 1 | CA | 414 | A |
| 1 | CA | 415 | A |
| 1 | CA | 420 | U |
| 1 | CA | 421 | U |
| 1 | CA | 422 | C |
| 1 | CA | 424 | G |
| 1 | CA | 428 | G |
| 1 | CA | 429 | U |
| 1 | CA | 430 | A |
| 1 | CA | 435 | C |
| 1 | CA | 437 | U |
| 1 | CA | 439 | A |
| 1 | CA | 442 | C |
| 1 | CA | 448 | A |
| 1 | CA | 452 | A |
| 1 | CA | 461 | A |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | CA | 470 | C |
| 1 | CA | 472 | A |
| 1 | CA | 473 | G |
| 1 | CA | 482 | A |
| 1 | CA | 484 | G |
| 1 | CA | 485 | G |
| 1 | CA | 487 | A |
| 1 | CA | 495 | A |
| 1 | CA | 496 | A |
| 1 | CA | 498 | U |
| 1 | CA | 500 | G |
| 1 | CA | 501 | C |
| 1 | CA | 505 | G |
| 1 | CA | 509 | A |
| 1 | CA | 510 | A |
| 1 | CA | 511 | C |
| 1 | CA | 513 | C |
| 1 | CA | 518 | C |
| 1 | CA | 520 | A |
| 1 | CA | 527 | G |
| 1 | CA | 531 | U |
| 1 | CA | 532 | A |
| 1 | CA | 533 | A |
| 1 | CA | 534 | U |
| 1 | CA | 545 | C |
| 1 | CA | 547 | A |
| 1 | CA | 559 | A |
| 1 | CA | 560 | U |
| 1 | CA | 561 | U |
| 1 | CA | 562 | C |
| 1 | CA | 572 | A |
| 1 | CA | 573 | A |
| 1 | CA | 576 | G |
| 1 | CA | 577 | G |
| 1 | CA | 588 | G |
| 1 | CA | 596 | C |
| 1 | CA | 607 | A |
| 1 | CA | 616 | G |
| 1 | CA | 618 | C |
| 1 | CA | 623 | C |
| 1 | CA | 630 | G |
| 1 | CA | 632 | A |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | CA | 653 | A |
| 1 | CA | 665 | A |
| 1 | CA | 687 | A |
| 1 | CA | 688 | G |
| 1 | CA | 693 | G |
| 1 | CA | 702 | A |
| 1 | CA | 703 | G |
| 1 | CA | 720 | C |
| 1 | CA | 721 | G |
| 1 | CA | 731 | G |
| 1 | CA | 733 | A |
| 1 | CA | 749 | C |
| 1 | CA | 753 | A |
| 1 | CA | 755 | G |
| 1 | CA | 766 | A |
| 1 | CA | 775 | G |
| 1 | CA | 776 | G |
| 1 | CA | 777 | A |
| 1 | CA | 786 | G |
| 1 | CA | 792 | A |
| 1 | CA | 793 | U |
| 1 | CA | 794 | A |
| 1 | CA | 802 | A |
| 1 | CA | 816 | A |
| 1 | CA | 817 | C |
| 1 | CA | 819 | A |
| 1 | CA | 821 | G |
| 1 | CA | 827 | U |
| 1 | CA | 828 | A |
| 1 | CA | 833 | U |
| 1 | CA | 839 | U |
| 1 | CA | 840 | C |
| 1 | CA | 841 | U |
| 1 | CA | 848 | C |
| 1 | CA | 859 | A |
| 1 | CA | 864 | A |
| 1 | CA | 870 | U |
| 1 | CA | 872 | A |
| 1 | CA | 884 | U |
| 1 | CA | 885 | G |
| 1 | CA | 902 | G |
| 1 | CA | 914 | A |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | CA | 920 | U |
| 1 | CA | 922 | G |
| 1 | CA | 926 | G |
| 1 | CA | 927 | G |
| 1 | CA | 932 | C |
| 1 | CA | 934 | C |
| 1 | CA | 935 | A |
| 1 | CA | 940 | C |
| 1 | CA | 960 | U |
| 1 | CA | 961 | U |
| 1 | CA | 968 | A |
| 1 | CA | 969 | A |
| 1 | CA | 971 | G |
| 1 | CA | 972 | C |
| 1 | CA | 974 | A |
| 1 | CA | 975 | A |
| 1 | CA | 976 | G |
| 1 | CA | 977 | A |
| 1 | CA | 978 | A |
| 1 | CA | 980 | C |
| 1 | CA | 991 | U |
| 1 | CA | 992 | U |
| 1 | CA | 993 | G |
| 1 | CA | 1005 | A |
| 1 | CA | 1026 | G |
| 1 | CA | 1050 | G |
| 1 | CA | 1054 | C |
| 1 | CA | 1065 | U |
| 1 | CA | 1066 | C |
| 1 | CA | 1068 | G |
| 1 | CA | 1081 | G |
| 1 | CA | 1094 | G |
| 1 | CA | 1095 | U |
| 1 | CA | 1101 | A |
| 1 | CA | 1117 | G |
| 1 | CA | 1118 | C |
| 1 | CA | 1124 | G |
| 1 | CA | 1125 | U |
| 1 | CA | 1126 | U |
| 1 | CA | 1129 | C |
| 1 | CA | 1136 | U |
| 1 | CA | 1137 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | CA | 1138 | G |
| 1 | CA | 1139 | G |
| 1 | CA | 1146 | A |
| 1 | CA | 1152 | A |
| 1 | CA | 1159 | U |
| 1 | CA | 1160 | G |
| 1 | CA | 1190 | G |
| 1 | CA | 1196 | U |
| 1 | CA | 1197 | G |
| 1 | CA | 1202 | G |
| 1 | CA | 1212 | U |
| 1 | CA | 1213 | A |
| 1 | CA | 1225 | A |
| 1 | CA | 1227 | A |
| 1 | CA | 1238 | A |
| 1 | CA | 1239 | A |
| 1 | CA | 1249 | C |
| 1 | CA | 1255 | G |
| 1 | CA | 1256 | A |
| 1 | CA | 1257 | U |
| 1 | CA | 1258 | G |
| 1 | CA | 1270 | C |
| 1 | CA | 1280 | A |
| 1 | CA | 1281 | U |
| 1 | CA | 1286 | A |
| 1 | CA | 1287 | A |
| 1 | CA | 1290 | G |
| 1 | CA | 1294 | G |
| 1 | CA | 1300 | G |
| 1 | CA | 1301 | U |
| 1 | CA | 1302 | U |
| 1 | CA | 1305 | G |
| 1 | CA | 1312 | G |
| 1 | CA | 1317 | C |
| 1 | CA | 1320 | C |
| 1 | CA | 1322 | C |
| 1 | CA | 1323 | G |
| 1 | CA | 1331 | G |
| 1 | CA | 1335 | C |
| 1 | CA | 1338 | G |
| 1 | CA | 1347 | G |
| 1 | CA | 1363 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | CA | 1363(A) | A |
| 1 | CA | 1364 | U |
| 1 | CA | 1370 | G |
| 1 | CA | 1388 | C |
| 1 | CA | 1397 | C |
| 1 | CA | 1402 | C |
| 1 | CA | 1419 | G |
| 1 | CA | 1442 | G |
| 1 | CA | 1442(A) | G |
| 1 | CA | 1442(B) | A |
| 1 | CA | 1443 | G |
| 1 | CA | 1447 | A |
| 1 | CA | 1452 | C |
| 1 | CA | 1456 | G |
| 1 | CA | 1487 | G |
| 1 | CA | 1492 | A |
| 1 | CA | 1493 | A |
| 1 | CA | 1494 | G |
| 1 | CA | 1495 | U |
| 1 | CA | 1497 | G |
| 1 | CA | 1499 | A |
| 1 | CA | 1502 | A |
| 1 | CA | 1504 | G |
| 1 | CA | 1505 | G |
| 1 | CA | 1506 | U |
| 1 | CA | 1507 | A |
| 1 | CA | 1517 | G |
| 1 | CA | 1519 | A |
| 1 | CA | 1520 | G |
| 1 | CA | 1529 | G |
| 1 | CA | 1530 | G |
| 31 | DA | 9 | U |
| 31 | DA | 10 | G |
| 31 | DA | 11 | G |
| 31 | DA | 15 | G |
| 31 | DA | 23 | G |
| 31 | DA | 34 | C |
| 31 | DA | 35 | G |
| 31 | DA | 45 | C |
| 31 | DA | 50 | U |
| 31 | DA | 51 | G |
| 31 | DA | 55 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 71 | A |
| 31 | DA | 72 | U |
| 31 | DA | 74 | A |
| 31 | DA | 75 | G |
| 31 | DA | 78 | A |
| 31 | DA | 83 | G |
| 31 | DA | 84 | A |
| 31 | DA | 90 | U |
| 31 | DA | 92 | A |
| 31 | DA | 94 | C |
| 31 | DA | 95 | G |
| 31 | DA | 99 | U |
| 31 | DA | 100 | G |
| 31 | DA | 102 | G |
| 31 | DA | 103 | A |
| 31 | DA | 105 | C |
| 31 | DA | 117 | G |
| 31 | DA | 118 | A |
| 31 | DA | 119 | A |
| 31 | DA | 120 | U |
| 31 | DA | 129 | C |
| 31 | DA | 131 | G |
| 31 | DA | 137 | C |
| 31 | DA | 139(A) | G |
| 31 | DA | 140 | G |
| 31 | DA | 142 | A |
| 31 | DA | 142(A) | C |
| 31 | DA | 146 | G |
| 31 | DA | 149 | A |
| 31 | DA | 154(A) | C |
| 31 | DA | 157 | U |
| 31 | DA | 158 | U |
| 31 | DA | 171 | G |
| 31 | DA | 173 | G |
| 31 | DA | 175 | G |
| 31 | DA | 196 | A |
| 31 | DA | 197 | A |
| 31 | DA | 198 | C |
| 31 | DA | 199 | A |
| 31 | DA | 204 | A |
| 31 | DA | 205 | G |
| 31 | DA | 215 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 216 | A |
| 31 | DA | 221 | A |
| 31 | DA | 222 | A |
| 31 | DA | 225 | A |
| 31 | DA | 228 | A |
| 31 | DA | 229 | A |
| 31 | DA | 233 | A |
| 31 | DA | 248 | G |
| 31 | DA | 250 | G |
| 31 | DA | 252 | G |
| 31 | DA | 267 | C |
| 31 | DA | 268 | C |
| 31 | DA | 269 | U |
| 31 | DA | 271(I) | G |
| 31 | DA | 271(J) | C |
| 31 | DA | 271(K) | U |
| 31 | DA | 271(L) | U |
| 31 | DA | 271(M) | G |
| 31 | DA | 271(N) | U |
| 31 | DA | 271(O) | C |
| 31 | DA | 271(P) | C |
| 31 | DA | 271(R) | G |
| 31 | DA | 271(T) | C |
| 31 | DA | 271(Y) | U |
| 31 | DA | 272 | G |
| 31 | DA | 272(B) | G |
| 31 | DA | 272(C) | G |
| 31 | DA | 272(H) | C |
| 31 | DA | 272(I) | U |
| 31 | DA | 272(J) | C |
| 31 | DA | 274 | G |
| 31 | DA | 275 | G |
| 31 | DA | 279 | C |
| 31 | DA | 281 | G |
| 31 | DA | 283 | A |
| 31 | DA | 286 | C |
| 31 | DA | 287 | C |
| 31 | DA | 298 | G |
| 31 | DA | 310 | A |
| 31 | DA | 311 | A |
| 31 | DA | 317 | G |
| 31 | DA | 327 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 329 | G |
| 31 | DA | 330 | A |
| 31 | DA | 331 | A |
| 31 | DA | 332 | A |
| 31 | DA | 343 | C |
| 31 | DA | 347 | A |
| 31 | DA | 352 | G |
| 31 | DA | 353 | G |
| 31 | DA | 362 | U |
| 31 | DA | 363 | G |
| 31 | DA | 363(B) | G |
| 31 | DA | 363(C) | G |
| 31 | DA | 363(E) | U |
| 31 | DA | 363(F) | A |
| 31 | DA | 366 | C |
| 31 | DA | 372 | G |
| 31 | DA | 386 | G |
| 31 | DA | 388 | G |
| 31 | DA | 396 | G |
| 31 | DA | 405 | U |
| 31 | DA | 407 | G |
| 31 | DA | 411 | G |
| 31 | DA | 412 | A |
| 31 | DA | 415 | A |
| 31 | DA | 418 | G |
| 31 | DA | 428 | A |
| 31 | DA | 444 | C |
| 31 | DA | 448 | U |
| 31 | DA | 455 | C |
| 31 | DA | 457 | A |
| 31 | DA | 470 | A |
| 31 | DA | 472 | A |
| 31 | DA | 473 | G |
| 31 | DA | 474 | G |
| 31 | DA | 475 | U |
| 31 | DA | 481 | G |
| 31 | DA | 482 | A |
| 31 | DA | 494 | G |
| 31 | DA | 495 | G |
| 31 | DA | 504 | U |
| 31 | DA | 505 | A |
| 31 | DA | 508 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 509 | C |
| 31 | DA | 511 | U |
| 31 | DA | 512 | G |
| 31 | DA | 518 | G |
| 31 | DA | 527 | C |
| 31 | DA | 529 | A |
| 31 | DA | 530 | G |
| 31 | DA | 531 | C |
| 31 | DA | 532 | A |
| 31 | DA | 533 | G |
| 31 | DA | 536 | A |
| 31 | DA | 537 | C |
| 31 | DA | 541 | C |
| 31 | DA | 542 | C |
| 31 | DA | 543 | C |
| 31 | DA | 547 | A |
| 31 | DA | 548 | A |
| 31 | DA | 549 | G |
| 31 | DA | 551 | G |
| 31 | DA | 563 | G |
| 31 | DA | 571 | A |
| 31 | DA | 573 | G |
| 31 | DA | 575 | A |
| 31 | DA | 586 | A |
| 31 | DA | 588 | U |
| 31 | DA | 596 | G |
| 31 | DA | 603 | A |
| 31 | DA | 604 | G |
| 31 | DA | 607 | U |
| 31 | DA | 614(B) | G |
| 31 | DA | 615 | G |
| 31 | DA | 618 | C |
| 31 | DA | 622 | G |
| 31 | DA | 626 | U |
| 31 | DA | 627 | A |
| 31 | DA | 634 | C |
| 31 | DA | 637 | A |
| 31 | DA | 645 | C |
| 31 | DA | 646 | A |
| 31 | DA | 649 | G |
| 31 | DA | 650 | C |
| 31 | DA | 651 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 652 | C |
| 31 | DA | 656 | G |
| 31 | DA | 657 | U |
| 31 | DA | 659 | C |
| 31 | DA | 668 | G |
| 31 | DA | 669 | G |
| 31 | DA | 670 | A |
| 31 | DA | 671 | C |
| 31 | DA | 686 | G |
| 31 | DA | 708 | C |
| 31 | DA | 717 | G |
| 31 | DA | 722 | A |
| 31 | DA | 726 | G |
| 31 | DA | 730 | C |
| 31 | DA | 745 | G |
| 31 | DA | 752 | A |
| 31 | DA | 753 | C |
| 31 | DA | 774 | A |
| 31 | DA | 775 | G |
| 31 | DA | 776 | G |
| 31 | DA | 779 | U |
| 31 | DA | 782 | A |
| 31 | DA | 784 | A |
| 31 | DA | 785 | G |
| 31 | DA | 790 | C |
| 31 | DA | 791 | C |
| 31 | DA | 792 | G |
| 31 | DA | 805 | G |
| 31 | DA | 808 | G |
| 31 | DA | 812 | C |
| 31 | DA | 819 | A |
| 31 | DA | 820 | A |
| 31 | DA | 826 | U |
| 31 | DA | 827 | U |
| 31 | DA | 828 | U |
| 31 | DA | 829 | A |
| 31 | DA | 830 | G |
| 31 | DA | 831 | G |
| 31 | DA | 847 | U |
| 31 | DA | 848 | G |
| 31 | DA | 856 | C |
| 31 | DA | 857 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 859 | G |
| 31 | DA | 867 | C |
| 31 | DA | 870 | A |
| 31 | DA | 872 | A |
| 31 | DA | 878 | A |
| 31 | DA | 883 | G |
| 31 | DA | 884 | C |
| 31 | DA | 892 | G |
| 31 | DA | 894 | C |
| 31 | DA | 896 | A |
| 31 | DA | 897 | C |
| 31 | DA | 898 | C |
| 31 | DA | 899 | A |
| 31 | DA | 900 | A |
| 31 | DA | 901 | A |
| 31 | DA | 902 | C |
| 31 | DA | 910 | A |
| 31 | DA | 911 | A |
| 31 | DA | 914 | C |
| 31 | DA | 917 | A |
| 31 | DA | 919 | G |
| 31 | DA | 923 | C |
| 31 | DA | 926 | A |
| 31 | DA | 932 | G |
| 31 | DA | 938 | G |
| 31 | DA | 941 | A |
| 31 | DA | 945 | A |
| 31 | DA | 946 | G |
| 31 | DA | 958 | U |
| 31 | DA | 959 | A |
| 31 | DA | 961 | C |
| 31 | DA | 974 | G |
| 31 | DA | 975 | C |
| 31 | DA | 975(A) | G |
| 31 | DA | 983 | A |
| 31 | DA | 990 | A |
| 31 | DA | 991 | C |
| 31 | DA | 993 | G |
| 31 | DA | 996 | A |
| 31 | DA | 1005 | C |
| 31 | DA | 1011 | G |
| 31 | DA | 1012 | U |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 1013 | C |
| 31 | DA | 1015 | G |
| 31 | DA | 1016 | G |
| 31 | DA | 1020 | A |
| 31 | DA | 1021 | A |
| 31 | DA | 1022 | G |
| 31 | DA | 1023 | U |
| 31 | DA | 1025 | G |
| 31 | DA | 1026 | U |
| 31 | DA | 1031 | G |
| 31 | DA | 1032 | A |
| 31 | DA | 1033 | U |
| 31 | DA | 1038 | C |
| 31 | DA | 1039 | G |
| 31 | DA | 1041 | C |
| 31 | DA | 1042 | G |
| 31 | DA | 1043 | C |
| 31 | DA | 1044 | G |
| 31 | DA | 1045 | A |
| 31 | DA | 1047 | G |
| 31 | DA | 1048 | A |
| 31 | DA | 1049 | C |
| 31 | DA | 1051 | G |
| 31 | DA | 1052 | C |
| 31 | DA | 1053 | C |
| 31 | DA | 1106 | A |
| 31 | DA | 1107 | G |
| 31 | DA | 1110 | G |
| 31 | DA | 1112 | G |
| 31 | DA | 1113 | U |
| 31 | DA | 1116 | C |
| 31 | DA | 1118 | C |
| 31 | DA | 1130 | U |
| 31 | DA | 1135 | C |
| 31 | DA | 1136 | G |
| 31 | DA | 1139 | G |
| 31 | DA | 1142 | U |
| 31 | DA | 1148 | A |
| 31 | DA | 1155 | A |
| 31 | DA | 1156 | A |
| 31 | DA | 1159 | U |
| 31 | DA | 1169 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 1171 | G |
| 31 | DA | 1173 | G |
| 31 | DA | 1174 | A |
| 31 | DA | 1175 | U |
| 31 | DA | 1176 | G |
| 31 | DA | 1177 | A |
| 31 | DA | 1178 | C |
| 31 | DA | 1179 | C |
| 31 | DA | 1180 | C |
| 31 | DA | 1195 | G |
| 31 | DA | 1210 | A |
| 31 | DA | 1211 | U |
| 31 | DA | 1213 | A |
| 31 | DA | 1220 | A |
| 31 | DA | 1242 | A |
| 31 | DA | 1251 | C |
| 31 | DA | 1252 | G |
| 31 | DA | 1253 | A |
| 31 | DA | 1255 | U |
| 31 | DA | 1256 | G |
| 31 | DA | 1265 | A |
| 31 | DA | 1271 | G |
| 31 | DA | 1272 | A |
| 31 | DA | 1273 | U |
| 31 | DA | 1287 | A |
| 31 | DA | 1288 | U |
| 31 | DA | 1300 | U |
| 31 | DA | 1301 | A |
| 31 | DA | 1305 | C |
| 31 | DA | 1307 | A |
| 31 | DA | 1308 | A |
| 31 | DA | 1310 | G |
| 31 | DA | 1314 | C |
| 31 | DA | 1317 | A |
| 31 | DA | 1318 | C |
| 31 | DA | 1319 | G |
| 31 | DA | 1345 | C |
| 31 | DA | 1349 | A |
| 31 | DA | 1359 | A |
| 31 | DA | 1360 | A |
| 31 | DA | 1364 | G |
| 31 | DA | 1365 | A |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 1368 | G |
| 31 | DA | 1370 | C |
| 31 | DA | 1379 | A |
| 31 | DA | 1380 | G |
| 31 | DA | 1384 | A |
| 31 | DA | 1385 | G |
| 31 | DA | 1386 | C |
| 31 | DA | 1404 | C |
| 31 | DA | 1407 | C |
| 31 | DA | 1416 | G |
| 31 | DA | 1417 | C |
| 31 | DA | 1420 | U |
| 31 | DA | 1421 | G |
| 31 | DA | 1427 | A |
| 31 | DA | 1428 | C |
| 31 | DA | 1437 | C |
| 31 | DA | 1445 | A |
| 31 | DA | 1449 | A |
| 31 | DA | 1450 | G |
| 31 | DA | 1450(A) | C |
| 31 | DA | 1455 | G |
| 31 | DA | 1458 | C |
| 31 | DA | 1461 | G |
| 31 | DA | 1467 | C |
| 31 | DA | 1470 | G |
| 31 | DA | 1471 | A |
| 31 | DA | 1475 | G |
| 31 | DA | 1478 | G |
| 31 | DA | 1479 | G |
| 31 | DA | 1480 | G |
| 31 | DA | 1481 | U |
| 31 | DA | 1482 | G |
| 31 | DA | 1484 | G |
| 31 | DA | 1485 | G |
| 31 | DA | 1490 | A |
| 31 | DA | 1491 | G |
| 31 | DA | 1493 | C |
| 31 | DA | 1494 | A |
| 31 | DA | 1495 | A |
| 31 | DA | 1497 | U |
| 31 | DA | 1498 | C |
| 31 | DA | 1502 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 1503 | U |
| 31 | DA | 1505 | C |
| 31 | DA | 1508 | A |
| 31 | DA | 1509 | C |
| 31 | DA | 1509(A) | A |
| 31 | DA | 1511 | C |
| 31 | DA | 1512 | U |
| 31 | DA | 1520 | G |
| 31 | DA | 1526 | G |
| 31 | DA | 1528 | A |
| 31 | DA | 1528(A) | A |
| 31 | DA | 1529 | G |
| 31 | DA | 1530 | C |
| 31 | DA | 1531 | C |
| 31 | DA | 1532 | C |
| 31 | DA | 1533 | G |
| 31 | DA | 1543 | C |
| 31 | DA | 1545 | A |
| 31 | DA | 1546 | C |
| 31 | DA | 1554 | A |
| 31 | DA | 1558 | A |
| 31 | DA | 1559 | G |
| 31 | DA | 1569 | A |
| 31 | DA | 1578 | U |
| 31 | DA | 1579 | A |
| 31 | DA | 1580 | A |
| 31 | DA | 1581 | G |
| 31 | DA | 1584 | C |
| 31 | DA | 1586 | A |
| 31 | DA | 1588 | C |
| 31 | DA | 1591 | G |
| 31 | DA | 1593 | G |
| 31 | DA | 1598 | C |
| 31 | DA | 1603 | A |
| 31 | DA | 1608 | A |
| 31 | DA | 1609 | A |
| 31 | DA | 1610 | A |
| 31 | DA | 1617 | C |
| 31 | DA | 1618 | A |
| 31 | DA | 1632 | A |
| 31 | DA | 1635 | G |
| 31 | DA | 1636 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 1640 | C |
| 31 | DA | 1646 | C |
| 31 | DA | 1648 | C |
| 31 | DA | 1652 | A |
| 31 | DA | 1653 | G |
| 31 | DA | 1654 | A |
| 31 | DA | 1674 | G |
| 31 | DA | 1675 | C |
| 31 | DA | 1686 | C |
| 31 | DA | 1687 | G |
| 31 | DA | 1694 | C |
| 31 | DA | 1695 | G |
| 31 | DA | 1696 | G |
| 31 | DA | 1697 | G |
| 31 | DA | 1700 | A |
| 31 | DA | 1703 | G |
| 31 | DA | 1722 | A |
| 31 | DA | 1739 | U |
| 31 | DA | 1741 | A |
| 31 | DA | 1742 | G |
| 31 | DA | 1744 | C |
| 31 | DA | 1745 | C |
| 31 | DA | 1746 | G |
| 31 | DA | 1748 | G |
| 31 | DA | 1756 | G |
| 31 | DA | 1758 | G |
| 31 | DA | 1763 | G |
| 31 | DA | 1764 | G |
| 31 | DA | 1773 | A |
| 31 | DA | 1780 | A |
| 31 | DA | 1782 | C |
| 31 | DA | 1791 | A |
| 31 | DA | 1799 | G |
| 31 | DA | 1800 | C |
| 31 | DA | 1801 | G |
| 31 | DA | 1816 | G |
| 31 | DA | 1820 | U |
| 31 | DA | 1822 | G |
| 31 | DA | 1829 | A |
| 31 | DA | 1835 | G |
| 31 | DA | 1838 | C |
| 31 | DA | 1839 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 1847 | A |
| 31 | DA | 1858 | G |
| 31 | DA | 1865 | G |
| 31 | DA | 1866 | C |
| 31 | DA | 1877 | A |
| 31 | DA | 1878 | G |
| 31 | DA | 1882 | C |
| 31 | DA | 1883 | G |
| 31 | DA | 1885 | A |
| 31 | DA | 1888 | G |
| 31 | DA | 1889 | A |
| 31 | DA | 1900 | A |
| 31 | DA | 1902 | C |
| 31 | DA | 1903 | G |
| 31 | DA | 1904 | G |
| 31 | DA | 1905 | C |
| 31 | DA | 1906 | G |
| 31 | DA | 1913 | A |
| 31 | DA | 1914 | C |
| 31 | DA | 1916 | A |
| 31 | DA | 1929 | G |
| 31 | DA | 1930 | G |
| 31 | DA | 1931 | U |
| 31 | DA | 1934 | C |
| 31 | DA | 1935 | G |
| 31 | DA | 1936 | A |
| 31 | DA | 1938 | A |
| 31 | DA | 1946 | U |
| 31 | DA | 1955 | U |
| 31 | DA | 1962 | C |
| 31 | DA | 1963 | U |
| 31 | DA | 1964 | G |
| 31 | DA | 1967 | C |
| 31 | DA | 1969 | A |
| 31 | DA | 1970 | A |
| 31 | DA | 1971 | A |
| 31 | DA | 1972 | A |
| 31 | DA | 1982 | C |
| 31 | DA | 1983 | C |
| 31 | DA | 1987 | G |
| 31 | DA | 1988 | C |
| 31 | DA | 1990 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 1991 | U |
| 31 | DA | 1993 | U |
| 31 | DA | 1997 | G |
| 31 | DA | 2018 | G |
| 31 | DA | 2020 | A |
| 31 | DA | 2023 | G |
| 31 | DA | 2030 | A |
| 31 | DA | 2031 | A |
| 31 | DA | 2033 | A |
| 31 | DA | 2036 | C |
| 31 | DA | 2043 | C |
| 31 | DA | 2055 | C |
| 31 | DA | 2056 | G |
| 31 | DA | 2060 | A |
| 31 | DA | 2061 | G |
| 31 | DA | 2062 | A |
| 31 | DA | 2069 | G |
| 31 | DA | 2093 | G |
| 31 | DA | 2096 | U |
| 31 | DA | 2100 | G |
| 31 | DA | 2103 | C |
| 31 | DA | 2104 | G |
| 31 | DA | 2187 | G |
| 31 | DA | 2190 | G |
| 31 | DA | 2191 | G |
| 31 | DA | 2192 | G |
| 31 | DA | 2198 | A |
| 31 | DA | 2199 | A |
| 31 | DA | 2200 | C |
| 31 | DA | 2201 | C |
| 31 | DA | 2203 | U |
| 31 | DA | 2206 | G |
| 31 | DA | 2207 | G |
| 31 | DA | 2208 | A |
| 31 | DA | 2218 | U |
| 31 | DA | 2219 | G |
| 31 | DA | 2225 | A |
| 31 | DA | 2226 | C |
| 31 | DA | 2227 | A |
| 31 | DA | 2234 | G |
| 31 | DA | 2238 | G |
| 31 | DA | 2239 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 2240 | C |
| 31 | DA | 2246 | G |
| 31 | DA | 2252 | G |
| 31 | DA | 2263 | C |
| 31 | DA | 2268 | A |
| 31 | DA | 2273 | A |
| 31 | DA | 2274 | A |
| 31 | DA | 2275 | C |
| 31 | DA | 2283 | C |
| 31 | DA | 2287 | A |
| 31 | DA | 2288 | A |
| 31 | DA | 2289 | G |
| 31 | DA | 2291 | U |
| 31 | DA | 2302 | G |
| 31 | DA | 2303 | G |
| 31 | DA | 2305 | A |
| 31 | DA | 2307 | G |
| 31 | DA | 2308 | G |
| 31 | DA | 2309 | A |
| 31 | DA | 2311 | A |
| 31 | DA | 2319 | G |
| 31 | DA | 2320 | A |
| 31 | DA | 2325 | G |
| 31 | DA | 2334 | G |
| 31 | DA | 2336 | A |
| 31 | DA | 2340 | G |
| 31 | DA | 2345 | G |
| 31 | DA | 2346 | A |
| 31 | DA | 2347 | C |
| 31 | DA | 2350 | C |
| 31 | DA | 2354 | G |
| 31 | DA | 2359 | C |
| 31 | DA | 2360 | A |
| 31 | DA | 2361 | A |
| 31 | DA | 2377 | A |
| 31 | DA | 2383 | G |
| 31 | DA | 2385 | C |
| 31 | DA | 2388 | A |
| 31 | DA | 2394 | C |
| 31 | DA | 2395 | C |
| 31 | DA | 2396 | G |
| 31 | DA | 2399 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 2400 | G |
| 31 | DA | 2402 | C |
| 31 | DA | 2403 | C |
| 31 | DA | 2405 | G |
| 31 | DA | 2406 | U |
| 31 | DA | 2410 | G |
| 31 | DA | 2411 | A |
| 31 | DA | 2422 | A |
| 31 | DA | 2423 | U |
| 31 | DA | 2425 | A |
| 31 | DA | 2427 | C |
| 31 | DA | 2429 | G |
| 31 | DA | 2430 | A |
| 31 | DA | 2435 | A |
| 31 | DA | 2439 | A |
| 31 | DA | 2440 | C |
| 31 | DA | 2441 | C |
| 31 | DA | 2448 | A |
| 31 | DA | 2464 | C |
| 31 | DA | 2465 | C |
| 31 | DA | 2468 | G |
| 31 | DA | 2469 | A |
| 31 | DA | 2470 | G |
| 31 | DA | 2471 | C |
| 31 | DA | 2474 | C |
| 31 | DA | 2475 | C |
| 31 | DA | 2476 | A |
| 31 | DA | 2477 | C |
| 31 | DA | 2478 | A |
| 31 | DA | 2482 | G |
| 31 | DA | 2484 | G |
| 31 | DA | 2489 | G |
| 31 | DA | 2497 | A |
| 31 | DA | 2500 | U |
| 31 | DA | 2502 | G |
| 31 | DA | 2504 | U |
| 31 | DA | 2505 | G |
| 31 | DA | 2518 | A |
| 31 | DA | 2520 | C |
| 31 | DA | 2524 | G |
| 31 | DA | 2529 | G |
| 31 | DA | 2535 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 2543 | G |
| 31 | DA | 2544 | G |
| 31 | DA | 2550 | G |
| 31 | DA | 2554 | U |
| 31 | DA | 2556 | C |
| 31 | DA | 2566 | A |
| 31 | DA | 2567 | G |
| 31 | DA | 2569 | G |
| 31 | DA | 2570 | G |
| 31 | DA | 2572 | A |
| 31 | DA | 2573 | C |
| 31 | DA | 2577 | A |
| 31 | DA | 2586 | C |
| 31 | DA | 2602 | A |
| 31 | DA | 2603 | G |
| 31 | DA | 2608 | G |
| 31 | DA | 2609 | U |
| 31 | DA | 2610 | C |
| 31 | DA | 2611 | U |
| 31 | DA | 2612 | C |
| 31 | DA | 2615 | U |
| 31 | DA | 2630 | G |
| 31 | DA | 2636 | U |
| 31 | DA | 2637 | U |
| 31 | DA | 2638 | G |
| 31 | DA | 2646 | C |
| 31 | DA | 2654 | A |
| 31 | DA | 2655 | G |
| 31 | DA | 2658 | C |
| 31 | DA | 2659 | G |
| 31 | DA | 2660 | A |
| 31 | DA | 2661 | G |
| 31 | DA | 2662 | A |
| 31 | DA | 2663 | G |
| 31 | DA | 2673 | G |
| 31 | DA | 2680 | C |
| 31 | DA | 2686 | G |
| 31 | DA | 2689 | U |
| 31 | DA | 2690 | C |
| 31 | DA | 2691 | C |
| 31 | DA | 2702 | U |
| 31 | DA | 2703 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 2707 | G |
| 31 | DA | 2712 | U |
| 31 | DA | 2712(A) | A |
| 31 | DA | 2713 | A |
| 31 | DA | 2714 | G |
| 31 | DA | 2718 | G |
| 31 | DA | 2726 | U |
| 31 | DA | 2733 | A |
| 31 | DA | 2734 | A |
| 31 | DA | 2752 | C |
| 31 | DA | 2753 | A |
| 31 | DA | 2754 | U |
| 31 | DA | 2756 | U |
| 31 | DA | 2757 | A |
| 31 | DA | 2758 | A |
| 31 | DA | 2759 | G |
| 31 | DA | 2762 | G |
| 31 | DA | 2764 | A |
| 31 | DA | 2765 | A |
| 31 | DA | 2766 | G |
| 31 | DA | 2778 | A |
| 31 | DA | 2779 | U |
| 31 | DA | 2780 | G |
| 31 | DA | 2781 | A |
| 31 | DA | 2782 | G |
| 31 | DA | 2789 | C |
| 31 | DA | 2790 | A |
| 31 | DA | 2791 | C |
| 31 | DA | 2794 | C |
| 31 | DA | 2795 | G |
| 31 | DA | 2801 | A |
| 31 | DA | 2801(A) | A |
| 31 | DA | 2802 | G |
| 31 | DA | 2803 | C |
| 31 | DA | 2808 | U |
| 31 | DA | 2812 | G |
| 31 | DA | 2813 | A |
| 31 | DA | 2818 | G |
| 31 | DA | 2820 | A |
| 31 | DA | 2821 | A |
| 31 | DA | 2830 | G |
| 31 | DA | 2833 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 2835 | A |
| 31 | DA | 2844 | G |
| 31 | DA | 2845 | G |
| 31 | DA | 2860 | A |
| 31 | DA | 2872 | G |
| 31 | DA | 2875 | C |
| 31 | DA | 2880 | C |
| 31 | DA | 2892 | A |
| 31 | DA | 2894 | G |
| 31 | DA | 2895 | U |
| 32 | DB | 3 | C |
| 32 | DB | 7 | G |
| 32 | DB | 8 | U |
| 32 | DB | 13 | A |
| 32 | DB | 15 | A |
| 32 | DB | 16 | G |
| 32 | DB | 20 | C |
| 32 | DB | 22 | U |
| 32 | DB | 26 | A |
| 32 | DB | 27 | C |
| 32 | DB | 29 | A |
| 32 | DB | 33 | G |
| 32 | DB | 40 | U |
| 32 | DB | 42 | C |
| 32 | DB | 45 | A |
| 32 | DB | 46 | A |
| 32 | DB | 47 | C |
| 32 | DB | 52 | A |
| 32 | DB | 53 | A |
| 32 | DB | 56 | G |
| 32 | DB | 65 | C |
| 32 | DB | 67 | G |
| 32 | DB | 73 | A |
| 32 | DB | 75 | G |
| 32 | DB | 81 | G |
| 32 | DB | 85 | G |
| 32 | DB | 88 | C |
| 32 | DB | 90 | A |
| 32 | DB | 91 | C |
| 32 | DB | 102 | A |
| 32 | DB | 106 | G |
| 32 | DB | 110 | G |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 32 | DB | 116 | G |
| 32 | DB | 118 | G |
| 32 | DB | 119 | G |

All (204) RNA pucker outliers are listed below:

| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | AA | 60 | A |
| 1 | AA | 79 | G |
| 1 | AA | 115 | G |
| 1 | AA | 243 | A |
| 1 | AA | 250 | A |
| 1 | AA | 266 | G |
| 1 | AA | 328 | C |
| 1 | AA | 353 | A |
| 1 | AA | 366 | C |
| 1 | AA | 428 | G |
| 1 | AA | 429 | U |
| 1 | AA | 484 | G |
| 1 | AA | 499 | A |
| 1 | AA | 509 | A |
| 1 | AA | 533 | A |
| 1 | AA | 560 | U |
| 1 | AA | 687 | A |
| 1 | AA | 748 | C |
| 1 | AA | 776 | G |
| 1 | AA | 913 | A |
| 1 | AA | 991 | U |
| 1 | AA | 992 | U |
| 1 | AA | 1049 | U |
| 1 | AA | 1064 | G |
| 1 | AA | 1065 | U |
| 1 | AA | 1067 | A |
| 1 | AA | 1201 | A |
| 1 | AA | 1285 | A |
| 1 | AA | 1493 | A |
| 1 | AA | 1498 | U |
| 1 | AA | 1504 | G |
| 31 | BA | 34 | C |
| 31 | BA | 50 | U |
| 31 | BA | 71 | A |
| 31 | BA | 128 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 221 | A |
| 31 | BA | 272 | G |
| 31 | BA | 310 | A |
| 31 | BA | 331 | A |
| 31 | BA | 387 | U |
| 31 | BA | 472 | A |
| 31 | BA | 474 | G |
| 31 | BA | 481 | G |
| 31 | BA | 494 | G |
| 31 | BA | 512 | G |
| 31 | BA | 529 | A |
| 31 | BA | 542 | C |
| 31 | BA | 587 | C |
| 31 | BA | 603 | A |
| 31 | BA | 668 | G |
| 31 | BA | 669 | G |
| 31 | BA | 670 | A |
| 31 | BA | 746 | A |
| 31 | BA | 752 | A |
| 31 | BA | 774 | A |
| 31 | BA | 790 | C |
| 31 | BA | 827 | U |
| 31 | BA | 856 | C |
| 31 | BA | 974 | G |
| 31 | BA | 1022 | G |
| 31 | BA | 1112 | G |
| 31 | BA | 1176 | G |
| 31 | BA | 1210 | A |
| 31 | BA | 1250 | G |
| 31 | BA | 1251 | C |
| 31 | BA | 1299 | G |
| 31 | BA | 1300 | U |
| 31 | BA | 1301 | A |
| 31 | BA | 1378 | A |
| 31 | BA | 1379 | A |
| 31 | BA | 1427 | A |
| 31 | BA | 1494 | A |
| 31 | BA | 1533 | G |
| 31 | BA | 1544 | A |
| 31 | BA | 1558 | A |
| 31 | BA | 1608 | A |
| 31 | BA | 1617 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | BA | 1652 | A |
| 31 | BA | 1694 | C |
| 31 | BA | 1740 | G |
| 31 | BA | 1819 | A |
| 31 | BA | 1934 | C |
| 31 | BA | 1962 | C |
| 31 | BA | 1970 | A |
| 31 | BA | 1987 | G |
| 31 | BA | 1992 | G |
| 31 | BA | 2030 | A |
| 31 | BA | 2036 | C |
| 31 | BA | 2225 | A |
| 31 | BA | 2405 | G |
| 31 | BA | 2406 | U |
| 31 | BA | 2439 | A |
| 31 | BA | 2506 | U |
| 31 | BA | 2610 | C |
| 31 | BA | 2611 | U |
| 31 | BA | 2637 | U |
| 31 | BA | 2662 | A |
| 31 | BA | 2689 | U |
| 31 | BA | 2712 | U |
| 31 | BA | 2756 | U |
| 31 | BA | 2796 | U |
| 31 | BA | 2859 | G |
| 32 | BB | 81 | G |
| 1 | CA | 60 | A |
| 1 | CA | 79 | G |
| 1 | CA | 115 | G |
| 1 | CA | 243 | A |
| 1 | CA | 250 | A |
| 1 | CA | 266 | G |
| 1 | CA | 328 | C |
| 1 | CA | 353 | A |
| 1 | CA | 366 | C |
| 1 | CA | 428 | G |
| 1 | CA | 429 | U |
| 1 | CA | 484 | G |
| 1 | CA | 499 | A |
| 1 | CA | 509 | A |
| 1 | CA | 533 | A |
| 1 | CA | 560 | U |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 1 | CA | 687 | A |
| 1 | CA | 748 | C |
| 1 | CA | 776 | G |
| 1 | CA | 913 | A |
| 1 | CA | 991 | U |
| 1 | CA | 992 | U |
| 1 | CA | 1049 | U |
| 1 | CA | 1064 | G |
| 1 | CA | 1065 | U |
| 1 | CA | 1067 | A |
| 1 | CA | 1201 | A |
| 1 | CA | 1285 | A |
| 1 | CA | 1493 | A |
| 1 | CA | 1498 | U |
| 1 | CA | 1504 | G |
| 31 | DA | 34 | C |
| 31 | DA | 50 | U |
| 31 | DA | 71 | A |
| 31 | DA | 102 | G |
| 31 | DA | 128 | C |
| 31 | DA | 221 | A |
| 31 | DA | 272 | G |
| 31 | DA | 310 | A |
| 31 | DA | 331 | A |
| 31 | DA | 387 | U |
| 31 | DA | 472 | A |
| 31 | DA | 474 | G |
| 31 | DA | 481 | G |
| 31 | DA | 494 | G |
| 31 | DA | 512 | G |
| 31 | DA | 529 | A |
| 31 | DA | 542 | C |
| 31 | DA | 587 | C |
| 31 | DA | 603 | A |
| 31 | DA | 668 | G |
| 31 | DA | 669 | G |
| 31 | DA | 670 | A |
| 31 | DA | 752 | A |
| 31 | DA | 774 | A |
| 31 | DA | 790 | C |
| 31 | DA | 827 | U |
| 31 | DA | 856 | C |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 974 | G |
| 31 | DA | 1022 | G |
| 31 | DA | 1112 | G |
| 31 | DA | 1176 | G |
| 31 | DA | 1210 | A |
| 31 | DA | 1250 | G |
| 31 | DA | 1251 | C |
| 31 | DA | 1299 | G |
| 31 | DA | 1300 | U |
| 31 | DA | 1301 | A |
| 31 | DA | 1378 | A |
| 31 | DA | 1379 | A |
| 31 | DA | 1427 | A |
| 31 | DA | 1494 | A |
| 31 | DA | 1533 | G |
| 31 | DA | 1544 | A |
| 31 | DA | 1558 | A |
| 31 | DA | 1608 | A |
| 31 | DA | 1617 | C |
| 31 | DA | 1652 | A |
| 31 | DA | 1694 | C |
| 31 | DA | 1740 | G |
| 31 | DA | 1819 | A |
| 31 | DA | 1934 | C |
| 31 | DA | 1962 | C |
| 31 | DA | 1970 | A |
| 31 | DA | 1987 | G |
| 31 | DA | 1992 | G |
| 31 | DA | 2017 | U |
| 31 | DA | 2030 | A |
| 31 | DA | 2225 | A |
| 31 | DA | 2405 | G |
| 31 | DA | 2406 | U |
| 31 | DA | 2439 | A |
| 31 | DA | 2506 | U |
| 31 | DA | 2610 | C |
| 31 | DA | 2611 | U |
| 31 | DA | 2637 | U |
| 31 | DA | 2662 | A |
| 31 | DA | 2689 | U |
| 31 | DA | 2756 | U |
| 31 | DA | 2796 | U |

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| Mol | Chain | Res | Type |
|---|---|---|---|
| 31 | DA | 2859 | G |

## 5.4 Non-standard residues in protein, DNA, RNA chains (i)

There are no non-standard protein/DNA/RNA residues in this entry.

## 5.5 Carbohydrates (i)

There are no oligosaccharides in this entry.

## 5.6 Ligand geometry (i)

Of 802 ligands modelled in this entry, 800 are monoatomic - leaving 2 for Mogul analysis.

In the following table, the Counts columns list the number of bonds (or angles) for which Mogul statistics could be retrieved, the number of bonds (or angles) that are observed in the model and the number of bonds (or angles) that are defined in the Chemical Component Dictionary. The Link column lists molecule types, if any, to which the group is linked. The Z score for a bond length (or angle) is the number of standard deviations the observed value is removed from the expected value. A bond length (or angle) with $|Z| > 2$ is considered an outlier worth inspection. RMSZ is the root-mean-square of all Z scores of the bond lengths (or angles).

| Mol | Type | Chain | Res | Link | Bond lengths | | | Bond angles | | |
|---|---|---|---|---|---|---|---|---|---|---|
| | | | | | Counts | RMSZ | $\#\|Z\| > 2$ | Counts | RMSZ | $\#\|Z\| > 2$ |
| 55 | ZIT | BA | 3351 | - | 54,54,54 | 1.42 | 7 (12%) | 82,83,83 | 1.10 | 5 (6%) |
| 55 | ZIT | DA | 3311 | - | 54,54,54 | 1.42 | 6 (11%) | 82,83,83 | 1.10 | 5 (6%) |

In the following table, the Chirals column lists the number of chiral outliers, the number of chiral centers analysed, the number of these observed in the model and the number defined in the Chemical Component Dictionary. Similar counts are reported in the Torsion and Rings columns. '-' means no outliers of that kind were identified.

| Mol | Type | Chain | Res | Link | Chirals | Torsions | Rings |
|---|---|---|---|---|---|---|---|
| 55 | ZIT | BA | 3351 | - | - | 3/72/107/107 | 0/3/3/3 |
| 55 | ZIT | DA | 3311 | - | - | 3/72/107/107 | 0/3/3/3 |

All (13) bond length outliers are listed below:

| Mol | Chain | Res | Type | Atoms | Z | Observed(Å) | Ideal(Å) |
|---|---|---|---|---|---|---|---|
| 55 | DA | 3311 | ZIT | C13-C14 | 3.34 | 1.60 | 1.55 |

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| Mol | Chain | Res | Type | Atoms | Z | Observed(Å) | Ideal(Å) |
|---|---|---|---|---|---|---|---|
| 55 | BA | 3351 | ZIT | C22-C11 | 3.34 | 1.58 | 1.52 |
| 55 | DA | 3311 | ZIT | C22-C11 | 3.32 | 1.58 | 1.52 |
| 55 | BA | 3351 | ZIT | C13-C14 | 3.29 | 1.60 | 1.55 |
| 55 | DA | 3311 | ZIT | O13-C13 | 2.73 | 1.48 | 1.44 |
| 55 | BA | 3351 | ZIT | O13-C13 | 2.70 | 1.48 | 1.44 |
| 55 | BA | 3351 | ZIT | C6-C5 | 2.61 | 1.60 | 1.55 |
| 55 | DA | 3311 | ZIT | C6-C5 | 2.59 | 1.60 | 1.55 |
| 55 | BA | 3351 | ZIT | C13-C12 | 2.43 | 1.61 | 1.55 |
| 55 | DA | 3311 | ZIT | C13-C12 | 2.41 | 1.61 | 1.55 |
| 55 | BA | 3351 | ZIT | O6-C6 | 2.22 | 1.48 | 1.44 |
| 55 | DA | 3311 | ZIT | O6-C6 | 2.20 | 1.48 | 1.44 |
| 55 | BA | 3351 | ZIT | C19-C6 | 2.01 | 1.56 | 1.52 |

All (10) bond angle outliers are listed below:

| Mol | Chain | Res | Type | Atoms | Z | Observed($^o$) | Ideal($^o$) |
|---|---|---|---|---|---|---|---|
| 55 | BA | 3351 | ZIT | C9-N10-C11 | -3.08 | 106.95 | 112.06 |
| 55 | DA | 3311 | ZIT | C9-N10-C11 | -3.05 | 106.99 | 112.06 |
| 55 | DA | 3311 | ZIT | C7-C8-C9 | 2.83 | 116.09 | 112.10 |
| 55 | BA | 3351 | ZIT | C7-C8-C9 | 2.82 | 116.09 | 112.10 |
| 55 | DA | 3311 | ZIT | O6-C6-C7 | 2.20 | 113.84 | 108.34 |
| 55 | BA | 3351 | ZIT | O6-C6-C7 | 2.20 | 113.83 | 108.34 |
| 55 | BA | 3351 | ZIT | C4A-C3A-C2A | -2.16 | 106.91 | 110.02 |
| 55 | DA | 3311 | ZIT | C4A-C3A-C2A | -2.15 | 106.92 | 110.02 |
| 55 | BA | 3351 | ZIT | C23-C13-C14 | 2.09 | 114.09 | 111.26 |
| 55 | DA | 3311 | ZIT | C23-C13-C14 | 2.07 | 114.06 | 111.26 |

There are no chirality outliers.

All (6) torsion outliers are listed below:

| Mol | Chain | Res | Type | Atoms |
|---|---|---|---|---|
| 55 | BA | 3351 | ZIT | C12-C11-N10-C21 |
| 55 | BA | 3351 | ZIT | C22-C11-N10-C21 |
| 55 | DA | 3311 | ZIT | C12-C11-N10-C21 |
| 55 | DA | 3311 | ZIT | C22-C11-N10-C21 |
| 55 | BA | 3351 | ZIT | C12-C11-N10-C9 |
| 55 | DA | 3311 | ZIT | C12-C11-N10-C9 |

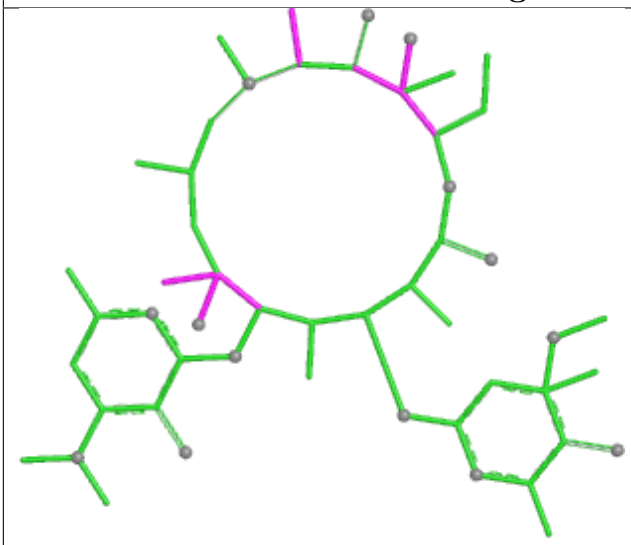
There are no ring outliers.

2 monomers are involved in 6 short contacts:

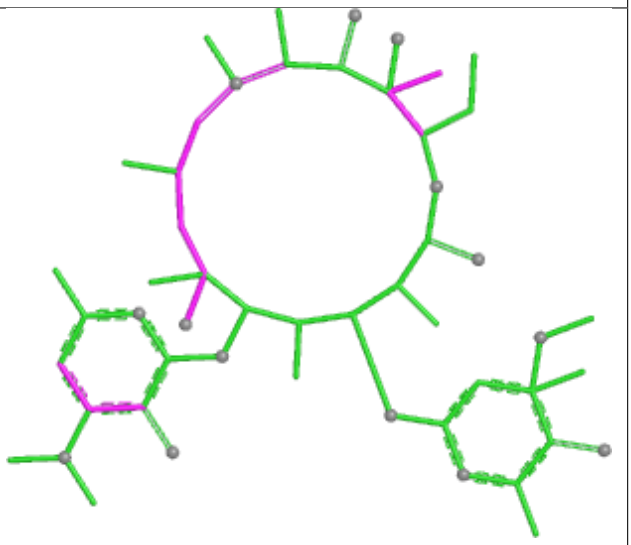
| Mol | Chain | Res | Type | Clashes | Symm-Clashes |
|---|---|---|---|---|---|
| 55 | BA | 3351 | ZIT | 3 | 0 |
| 55 | DA | 3311 | ZIT | 3 | 0 |

The following is a two-dimensional graphical depiction of Mogul quality analysis of bond lengths, bond angles, torsion angles, and ring geometry for all instances of the Ligand of Interest. In addition, ligands with molecular weight > 250 and outliers as shown on the validation Tables will also be included. For torsion angles, if less then 5% of the Mogul distribution of torsion angles is within 10 degrees of the torsion angle in question, then that torsion angle is considered an outlier. Any bond that is central to one or more torsion angles identified as an outlier by Mogul will be highlighted in the graph. For rings, the root-mean-square deviation (RMSD) between the ring in question and similar rings identified by Mogul is calculated over all ring torsion angles. If the average RMSD is greater than 60 degrees and the minimal RMSD between the ring in question and any Mogul-identified rings is also greater than 60 degrees, then that ring is considered an outlier. The outliers are highlighted in purple. The color gray indicates Mogul did not find sufficient equivalents in the CSD to analyse the geometry.

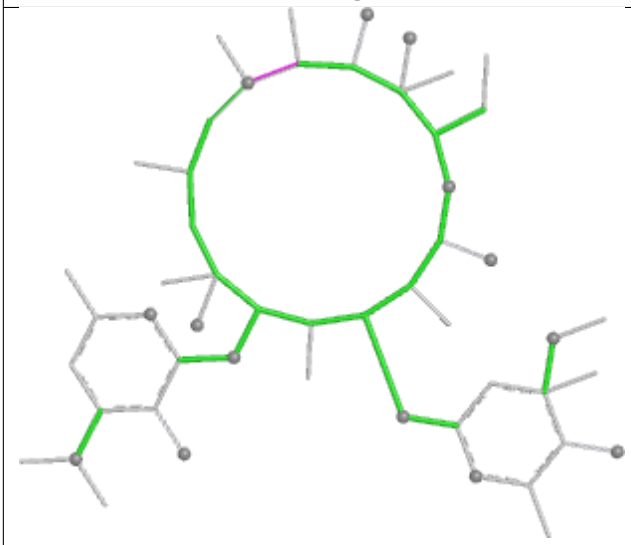
**Ligand ZIT BA 3351**



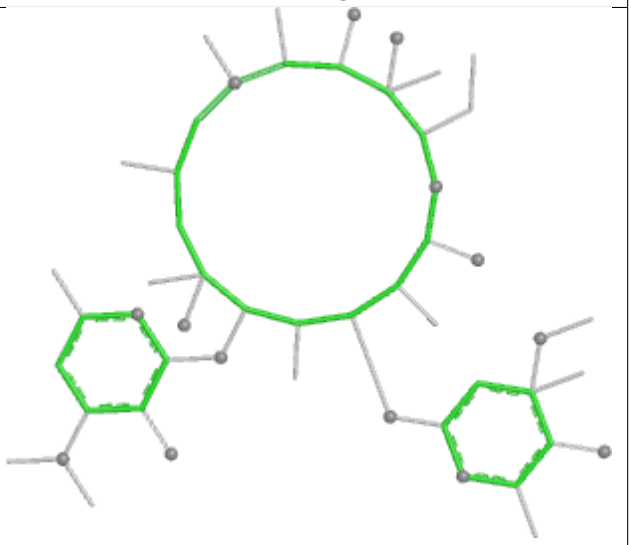
**Bond lengths**



**Bond angles**

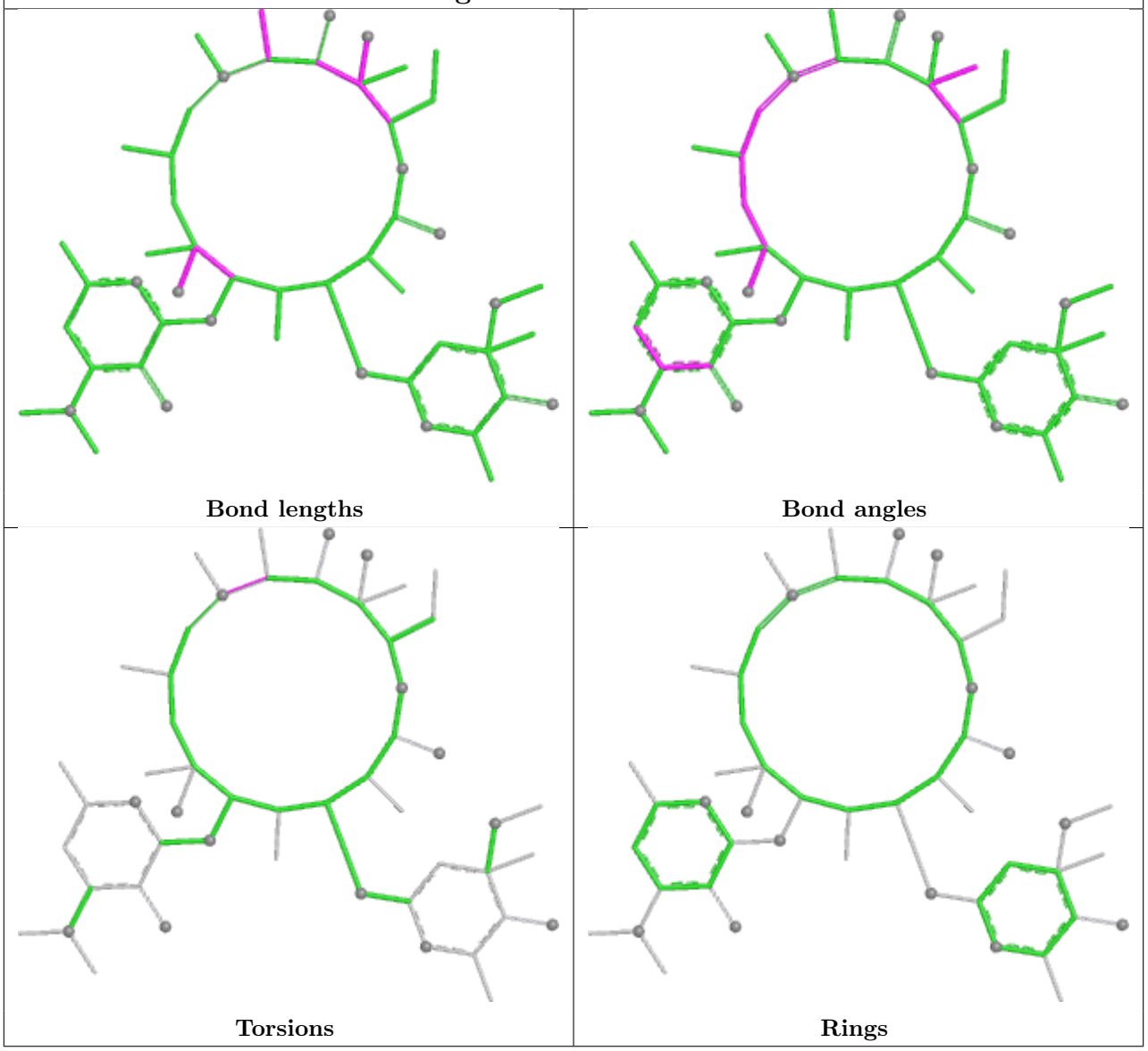


**Torsions**



**Rings**

**Ligand ZIT DA 3311**




## 5.7 Other polymers (i)

There are no such residues in this entry.

## 5.8 Polymer linkage issues (i)

The following chains have linkage breaks:

| Mol | Chain | Number of breaks |
|---|---|---|
| 13 | CM | 3 |
| 13 | AM | 3 |
| 36 | DG | 1 |

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| Mol | Chain | Number of breaks |
|---|---|---|
| 36 | BG | 1 |
| 28 | B6 | 1 |
| 28 | D6 | 1 |
| 9 | AI | 1 |
| 9 | CI | 1 |
| 47 | BV | 1 |
| 47 | DV | 1 |

All chain breaks are listed below:

| Model | Chain | Residue-1 | Atom-1 | Residue-2 | Atom-2 | Distance (Å) |
|---|---|---|---|---|---|---|
| 1 | CM | 69:GLU | C | 70:LEU | N | 5.29 |
| 1 | AM | 69:GLU | C | 70:LEU | N | 5.28 |
| 1 | DG | 112:PRO | C | 113:ARG | N | 4.77 |
| 1 | BG | 112:PRO | C | 113:ARG | N | 4.76 |
| 1 | AM | 112:GLY | C | 113:PRO | N | 4.20 |
| 1 | CM | 112:GLY | C | 113:PRO | N | 4.18 |
| 1 | AM | 97:PRO | C | 98:VAL | N | 4.12 |
| 1 | CM | 97:PRO | C | 98:VAL | N | 4.12 |
| 1 | B6 | 46:HIS | C | 47:THR | N | 4.06 |
| 1 | D6 | 46:HIS | C | 47:THR | N | 4.03 |
| 1 | AI | 53:VAL | C | 54:ASP | N | 3.98 |
| 1 | CI | 53:VAL | C | 54:ASP | N | 3.97 |
| 1 | BV | 80:GLN | C | 81:TYR | N | 3.43 |
| 1 | DV | 80:GLN | C | 81:TYR | N | 3.26 |

# 6 Fit of model and data

## 6.1 Protein, DNA and RNA chains

In the following table, the column labelled '#RSRZ> 2' contains the number (and percentage) of RSRZ outliers, followed by percent RSRZ outliers for the chain as percentile scores relative to all X-ray entries and entries of similar resolution. The OWAB column contains the minimum, median, $95^{th}$ percentile and maximum values of the occupancy-weighted average B-factor per residue. The column labelled 'Q< 0.9' lists the number of (and percentage) of residues with an average occupancy less than 0.9.

| Mol | Chain | Analysed | <RSRZ> | #RSRZ>2 | | | OWAB(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 1 | AA | 1504/1522 (98%) | 0.82 | 169 (11%) | 10 | 6 | 60, 125, 191, 194 | 0 |
| 1 | CA | 1504/1522 (98%) | 0.74 | 134 (8%) | 15 | 8 | 61, 125, 191, 194 | 0 |
| 2 | AB | 235/256 (91%) | 1.19 | 48 (20%) | 3 | 2 | 107, 156, 184, 191 | 0 |
| 2 | CB | 235/256 (91%) | 1.26 | 49 (20%) | 2 | 2 | 107, 158, 185, 191 | 0 |
| 3 | AC | 207/239 (86%) | 1.04 | 32 (15%) | 5 | 3 | 115, 163, 184, 189 | 0 |
| 3 | CC | 207/239 (86%) | 1.32 | 47 (22%) | 2 | 1 | 119, 166, 184, 191 | 0 |
| 4 | AD | 208/209 (99%) | 1.36 | 46 (22%) | 2 | 1 | 83, 131, 170, 181 | 0 |
| 4 | CD | 208/209 (99%) | 1.42 | 47 (22%) | 2 | 1 | 82, 131, 168, 182 | 0 |
| 5 | AE | 151/162 (93%) | 1.02 | 27 (17%) | 3 | 2 | 83, 116, 160, 188 | 0 |
| 5 | CE | 151/162 (93%) | 1.02 | 22 (14%) | 6 | 3 | 84, 117, 162, 189 | 0 |
| 6 | AF | 101/101 (100%) | 1.03 | 14 (13%) | 6 | 4 | 85, 132, 164, 180 | 0 |
| 6 | CF | 101/101 (100%) | 1.19 | 17 (16%) | 4 | 3 | 86, 132, 165, 182 | 0 |
| 7 | AG | 155/156 (99%) | 1.49 | 46 (29%) | 1 | 1 | 140, 171, 188, 191 | 0 |
| 7 | CG | 155/156 (99%) | 1.39 | 39 (25%) | 1 | 1 | 140, 171, 188, 190 | 0 |
| 8 | AH | 138/138 (100%) | 1.01 | 19 (13%) | 6 | 4 | 85, 121, 155, 164 | 0 |
| 8 | CH | 138/138 (100%) | 1.09 | 23 (16%) | 4 | 3 | 85, 123, 156, 162 | 0 |
| 9 | AI | 127/128 (99%) | 1.79 | 45 (35%) | 1 | 1 | 142, 182, 190, 192 | 0 |
| 9 | CI | 127/128 (99%) | 1.90 | 56 (44%) | 0 | 1 | 143, 183, 190, 191 | 0 |
| 10 | AJ | 99/105 (94%) | 1.90 | 37 (37%) | 1 | 1 | 130, 176, 189, 191 | 0 |
| 10 | CJ | 99/105 (94%) | 1.72 | 37 (37%) | 1 | 1 | 130, 177, 190, 193 | 0 |
| 11 | AK | 119/129 (92%) | 1.31 | 30 (25%) | 1 | 1 | 82, 123, 164, 187 | 0 |
| 11 | CK | 119/129 (92%) | 1.27 | 25 (21%) | 2 | 1 | 84, 123, 165, 186 | 0 |
| 12 | AL | 125/135 (92%) | 1.41 | 33 (26%) | 1 | 1 | 80, 108, 163, 189 | 0 |
| 12 | CL | 125/135 (92%) | 1.49 | 35 (28%) | 1 | 1 | 82, 109, 164, 188 | 0 |

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| Mol | Chain | Analysed | <RSRZ> | #RSRZ>2 | | | OWAB(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 13 | AM | 115/126 (91%) | 1.93 | 40 (34%) | 1 | 1 | 150, 185, 190, 193 | 0 |
| 13 | CM | 115/126 (91%) | 1.92 | 49 (42%) | 0 | 1 | 149, 185, 190, 192 | 0 |
| 14 | AN | 60/61 (98%) | 1.83 | 22 (36%) | 1 | 1 | 131, 168, 185, 189 | 0 |
| 14 | CN | 60/61 (98%) | 1.82 | 21 (35%) | 1 | 1 | 132, 170, 186, 189 | 0 |
| 15 | AO | 88/89 (98%) | 1.22 | 19 (21%) | 2 | 2 | 74, 111, 157, 162 | 0 |
| 15 | CO | 88/89 (98%) | 0.82 | 12 (13%) | 7 | 4 | 74, 112, 159, 165 | 0 |
| 16 | AP | 84/88 (95%) | 1.85 | 30 (35%) | 1 | 1 | 91, 118, 161, 179 | 0 |
| 16 | CP | 84/88 (95%) | 1.63 | 27 (32%) | 1 | 1 | 89, 116, 160, 180 | 0 |
| 17 | AQ | 100/105 (95%) | 1.14 | 19 (19%) | 3 | 2 | 80, 109, 153, 163 | 0 |
| 17 | CQ | 100/105 (95%) | 1.28 | 16 (16%) | 5 | 3 | 85, 110, 153, 159 | 0 |
| 18 | AR | 70/88 (79%) | 1.12 | 9 (12%) | 7 | 5 | 93, 121, 170, 183 | 0 |
| 18 | CR | 70/88 (79%) | 0.78 | 5 (7%) | 22 | 11 | 93, 122, 171, 183 | 0 |
| 19 | AS | 79/93 (84%) | 1.73 | 30 (37%) | 1 | 1 | 142, 186, 190, 191 | 0 |
| 19 | CS | 79/93 (84%) | 1.63 | 25 (31%) | 1 | 1 | 142, 186, 191, 192 | 0 |
| 20 | AT | 99/106 (93%) | 1.68 | 33 (33%) | 1 | 1 | 84, 119, 157, 177 | 0 |
| 20 | CT | 99/106 (93%) | 1.40 | 27 (27%) | 1 | 1 | 84, 119, 157, 179 | 0 |
| 21 | AU | 25/27 (92%) | 3.73 | 22 (88%) | 0 | 0 | 143, 174, 188, 190 | 0 |
| 21 | CU | 25/27 (92%) | 3.81 | 20 (80%) | 0 | 0 | 141, 172, 188, 189 | 0 |
| 22 | B0 | 85/85 (100%) | 1.14 | 15 (17%) | 4 | 3 | 49, 70, 175, 187 | 0 |
| 22 | D0 | 85/85 (100%) | 0.83 | 14 (16%) | 4 | 3 | 54, 74, 173, 188 | 0 |
| 23 | B1 | 89/98 (90%) | 1.92 | 31 (34%) | 1 | 1 | 50, 79, 150, 187 | 0 |
| 23 | D1 | 89/98 (90%) | 1.55 | 23 (25%) | 1 | 1 | 51, 81, 151, 190 | 0 |
| 24 | B2 | 51/72 (70%) | 2.81 | 31 (60%) | 0 | 0 | 59, 99, 175, 186 | 0 |
| 24 | D2 | 51/72 (70%) | 2.32 | 24 (47%) | 0 | 0 | 62, 100, 175, 188 | 0 |
| 25 | B3 | 60/60 (100%) | 0.92 | 10 (16%) | 4 | 3 | 46, 69, 132, 168 | 0 |
| 25 | D3 | 60/60 (100%) | 0.43 | 2 (3%) | 49 | 28 | 51, 72, 136, 161 | 0 |
| 26 | B4 | 32/71 (45%) | 1.96 | 13 (40%) | 0 | 1 | 133, 161, 182, 184 | 0 |
| 26 | D4 | 32/71 (45%) | 1.70 | 10 (31%) | 1 | 1 | 133, 164, 182, 186 | 0 |
| 27 | B5 | 58/60 (96%) | 1.00 | 11 (18%) | 3 | 2 | 34, 61, 165, 188 | 0 |
| 27 | D5 | 58/60 (96%) | 0.80 | 7 (12%) | 8 | 5 | 39, 63, 163, 190 | 0 |
| 28 | B6 | 45/54 (83%) | 2.30 | 17 (37%) | 1 | 1 | 49, 85, 141, 173 | 0 |

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| Mol | Chain | Analysed | <RSRZ> | #RSRZ>2 | OWAB(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|
| 28 | D6 | 45/54 (83%) | 1.79 | 17 (37%) 1 1 | 52, 87, 142, 172 | 0 |
| 29 | B7 | 49/49 (100%) | 0.48 | 6 (12%) 8 5 | 36, 45, 119, 172 | 0 |
| 29 | D7 | 49/49 (100%) | 0.50 | 6 (12%) 8 5 | 38, 49, 120, 173 | 0 |
| 30 | B8 | 64/65 (98%) | 1.79 | 21 (32%) 1 1 | 46, 68, 140, 165 | 0 |
| 30 | D8 | 64/65 (98%) | 1.48 | 20 (31%) 1 1 | 49, 73, 141, 169 | 0 |
| 31 | BA | 2725/2787 (97%) | 0.09 | 164 (6%) 27 14 | 33, 59, 153, 194 | 0 |
| 31 | DA | 2725/2787 (97%) | -0.01 | 105 (3%) 43 24 | 38, 64, 157, 194 | 0 |
| 32 | BB | 119/122 (97%) | 1.43 | 34 (28%) 1 1 | 50, 101, 149, 184 | 0 |
| 32 | DB | 119/122 (97%) | 1.12 | 24 (20%) 3 2 | 59, 105, 157, 184 | 0 |
| 33 | BD | 272/276 (98%) | 0.46 | 26 (9%) 13 7 | 37, 62, 120, 168 | 0 |
| 33 | DD | 272/276 (98%) | 0.45 | 20 (7%) 20 10 | 40, 65, 122, 165 | 0 |
| 34 | BE | 205/206 (99%) | 0.83 | 25 (12%) 8 5 | 36, 65, 153, 181 | 0 |
| 34 | DE | 205/206 (99%) | 0.71 | 26 (12%) 8 5 | 40, 69, 154, 182 | 0 |
| 35 | BF | 208/210 (99%) | 0.79 | 23 (11%) 10 6 | 35, 77, 175, 189 | 0 |
| 35 | DF | 208/210 (99%) | 0.62 | 20 (9%) 13 7 | 39, 79, 176, 188 | 0 |
| 36 | BG | 181/182 (99%) | 1.96 | 73 (40%) 0 1 | 100, 152, 186, 192 | 0 |
| 36 | DG | 181/182 (99%) | 1.79 | 65 (35%) 1 1 | 106, 159, 189, 191 | 0 |
| 37 | BH | 160/180 (88%) | 2.15 | 70 (43%) 0 1 | 69, 111, 151, 182 | 0 |
| 37 | DH | 160/180 (88%) | 1.15 | 22 (13%) 6 4 | 74, 114, 157, 185 | 0 |
| 38 | BI | 146/148 (98%) | 1.73 | 49 (33%) 1 1 | 67, 152, 187, 190 | 0 |
| 38 | DI | 146/148 (98%) | 1.63 | 41 (28%) 1 1 | 69, 156, 189, 191 | 0 |
| 39 | BN | 139/140 (99%) | 1.34 | 34 (24%) 2 1 | 45, 75, 143, 182 | 0 |
| 39 | DN | 139/140 (99%) | 0.75 | 17 (12%) 8 5 | 49, 78, 143, 183 | 0 |
| 40 | BO | 122/122 (100%) | 0.38 | 3 (2%) 58 35 | 45, 67, 123, 147 | 0 |
| 40 | DO | 122/122 (100%) | 0.35 | 8 (6%) 24 12 | 48, 69, 125, 149 | 0 |
| 41 | BP | 146/150 (97%) | 1.72 | 47 (32%) 1 1 | 29, 93, 149, 190 | 0 |
| 41 | DP | 146/150 (97%) | 1.24 | 38 (26%) 1 1 | 38, 95, 152, 188 | 0 |
| 42 | BQ | 136/141 (96%) | 1.66 | 30 (22%) 2 1 | 50, 77, 147, 183 | 0 |
| 42 | DQ | 136/141 (96%) | 1.19 | 17 (12%) 8 5 | 52, 79, 147, 183 | 0 |
| 43 | BR | 117/118 (99%) | 0.69 | 13 (11%) 10 6 | 40, 60, 130, 139 | 0 |
| 43 | DR | 117/118 (99%) | 0.55 | 17 (14%) 6 3 | 42, 62, 131, 140 | 0 |

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| Mol | Chain | Analysed | <RSRZ> | #RSRZ>2 | | | OWAB(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 44 | BS | 99/112 (88%) | 2.13 | 48 (48%) | 0 | 0 | 54, 111, 148, 165 | 0 |
| 44 | DS | 99/112 (88%) | 1.69 | 31 (31%) | 1 | 1 | 62, 113, 154, 170 | 0 |
| 45 | BT | 132/146 (90%) | 1.29 | 31 (23%) | 2 | 1 | 55, 87, 154, 181 | 0 |
| 45 | DT | 132/146 (90%) | 1.04 | 29 (21%) | 2 | 1 | 58, 90, 156, 179 | 0 |
| 46 | BU | 117/118 (99%) | 1.02 | 23 (19%) | 3 | 2 | 40, 62, 124, 176 | 0 |
| 46 | DU | 117/118 (99%) | 0.92 | 15 (12%) | 7 | 5 | 44, 67, 130, 175 | 0 |
| 47 | BV | 101/101 (100%) | 2.21 | 50 (49%) | 0 | 0 | 38, 103, 176, 188 | 0 |
| 47 | DV | 101/101 (100%) | 1.22 | 21 (20%) | 2 | 2 | 44, 109, 177, 188 | 0 |
| 48 | BW | 113/113 (100%) | 0.30 | 7 (6%) | 26 | 14 | 38, 51, 112, 179 | 0 |
| 48 | DW | 113/113 (100%) | 0.16 | 4 (3%) | 47 | 27 | 41, 54, 119, 181 | 0 |
| 49 | BX | 93/96 (96%) | 1.60 | 29 (31%) | 1 | 1 | 47, 74, 145, 179 | 0 |
| 49 | DX | 93/96 (96%) | 1.38 | 19 (20%) | 3 | 2 | 52, 76, 146, 179 | 0 |
| 50 | BY | 101/110 (91%) | 2.54 | 59 (58%) | 0 | 0 | 57, 107, 184, 192 | 0 |
| 50 | DY | 101/110 (91%) | 2.02 | 42 (41%) | 0 | 1 | 60, 108, 183, 193 | 0 |
| 51 | BZ | 177/206 (85%) | 1.61 | 53 (29%) | 1 | 1 | 68, 113, 158, 169 | 0 |
| 51 | DZ | 177/206 (85%) | 1.22 | 40 (22%) | 2 | 1 | 74, 117, 161, 168 | 0 |
| All | All | 20062/20922 (95%) | 0.87 | 3293 (16%) | 4 | 3 | 29, 99, 187, 194 | 0 |

All (3293) RSRZ outliers are listed below:

| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 42 | BQ | 140 | ALA | 31.8 |
| 42 | BQ | 141 | GLN | 22.8 |
| 39 | BN | 68 | GLU | 15.8 |
| 42 | DQ | 24 | GLY | 15.8 |
| 42 | BQ | 24 | GLY | 15.2 |
| 27 | B5 | 2 | ALA | 14.1 |
| 42 | DQ | 140 | ALA | 13.1 |
| 7 | AG | 5 | ARG | 12.5 |
| 17 | CQ | 101 | ARG | 12.4 |
| 42 | BQ | 139 | GLU | 11.9 |
| 31 | BA | 2101 | G | 11.8 |
| 42 | DQ | 141 | GLN | 11.8 |
| 24 | D2 | 42 | GLY | 11.6 |
| 37 | BH | 171 | LEU | 11.4 |
| 28 | B6 | 42 | TRP | 11.4 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 17 | CQ | 24 | GLU | 10.9 |
| 27 | D5 | 2 | ALA | 10.9 |
| 31 | BA | 652 | C | 10.9 |
| 31 | BA | 2189 | U | 10.7 |
| 49 | BX | 73 | ARG | 10.2 |
| 31 | BA | 897 | C | 10.1 |
| 1 | AA | 89 | C | 9.8 |
| 31 | DA | 2802 | G | 9.8 |
| 41 | BP | 107 | LYS | 9.8 |
| 39 | BN | 1 | MET | 9.7 |
| 21 | CU | 6 | ARG | 9.5 |
| 24 | B2 | 42 | GLY | 9.3 |
| 35 | BF | 24 | LEU | 9.0 |
| 1 | AA | 88 | A | 9.0 |
| 24 | B2 | 48 | HIS | 9.0 |
| 13 | AM | 101 | GLN | 9.0 |
| 50 | BY | 28 | LYS | 9.0 |
| 35 | BF | 56 | GLU | 8.9 |
| 9 | CI | 115 | GLY | 8.7 |
| 36 | BG | 88 | ILE | 8.7 |
| 3 | AC | 193 | TYR | 8.7 |
| 31 | BA | 2802 | G | 8.6 |
| 42 | BQ | 91 | GLU | 8.6 |
| 49 | DX | 73 | ARG | 8.6 |
| 50 | BY | 3 | VAL | 8.6 |
| 36 | BG | 35 | GLU | 8.5 |
| 24 | D2 | 43 | GLN | 8.5 |
| 42 | DQ | 139 | GLU | 8.3 |
| 50 | BY | 49 | VAL | 8.1 |
| 4 | CD | 7 | PRO | 8.1 |
| 4 | CD | 115 | ARG | 8.1 |
| 30 | D8 | 65 | GLU | 8.0 |
| 13 | AM | 102 | ARG | 8.0 |
| 21 | AU | 6 | ARG | 7.9 |
| 24 | D2 | 35 | LEU | 7.9 |
| 1 | CA | 83 | U | 7.8 |
| 50 | DY | 49 | VAL | 7.8 |
| 50 | DY | 28 | LYS | 7.8 |
| 23 | B1 | 93 | GLU | 7.8 |
| 30 | B8 | 65 | GLU | 7.7 |
| 37 | BH | 58 | GLU | 7.7 |
| 23 | B1 | 95 | LEU | 7.6 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 21 | CU | 10 | ARG | 7.6 |
| 49 | BX | 26 | TYR | 7.5 |
| 30 | B8 | 34 | TRP | 7.5 |
| 51 | BZ | 97 | GLU | 7.5 |
| 50 | DY | 61 | ILE | 7.5 |
| 50 | DY | 3 | VAL | 7.5 |
| 13 | CM | 102 | ARG | 7.5 |
| 24 | B2 | 54 | LYS | 7.5 |
| 41 | BP | 90 | ARG | 7.5 |
| 26 | B4 | 3 | GLU | 7.4 |
| 36 | BG | 86 | MET | 7.4 |
| 37 | BH | 147 | ASN | 7.4 |
| 50 | BY | 88 | LYS | 7.3 |
| 31 | DA | 2104 | G | 7.3 |
| 4 | AD | 4 | TYR | 7.3 |
| 7 | CG | 32 | ARG | 7.3 |
| 16 | AP | 29 | ASP | 7.3 |
| 35 | BF | 207 | GLY | 7.3 |
| 6 | CF | 90 | VAL | 7.3 |
| 14 | CN | 29 | ARG | 7.3 |
| 3 | CC | 158 | GLY | 7.2 |
| 49 | DX | 34 | ALA | 7.2 |
| 47 | BV | 53 | GLU | 7.2 |
| 33 | DD | 236 | GLY | 7.2 |
| 39 | DN | 109 | LYS | 7.2 |
| 33 | DD | 262 | ARG | 7.2 |
| 34 | BE | 76 | ARG | 7.1 |
| 9 | CI | 109 | VAL | 7.1 |
| 36 | DG | 28 | VAL | 7.1 |
| 30 | B8 | 37 | SER | 7.1 |
| 47 | BV | 47 | VAL | 7.1 |
| 1 | AA | 1286 | A | 7.1 |
| 25 | B3 | 30 | ARG | 7.1 |
| 44 | BS | 97 | ARG | 7.0 |
| 47 | DV | 76 | LYS | 7.0 |
| 34 | BE | 186 | GLY | 7.0 |
| 12 | CL | 16 | GLU | 7.0 |
| 13 | CM | 101 | GLN | 7.0 |
| 20 | AT | 12 | ALA | 6.9 |
| 28 | D6 | 42 | TRP | 6.9 |
| 46 | BU | 117 | GLN | 6.9 |
| 44 | BS | 54 | LEU | 6.9 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 38 | BI | 70 | GLU | 6.9 |
| 1 | AA | 1128 | C | 6.8 |
| 29 | D7 | 47 | ARG | 6.8 |
| 38 | DI | 65 | ALA | 6.7 |
| 3 | CC | 192 | THR | 6.7 |
| 4 | CD | 4 | TYR | 6.7 |
| 24 | B2 | 61 | LEU | 6.6 |
| 1 | CA | 1033 | G | 6.6 |
| 36 | BG | 87 | PRO | 6.6 |
| 41 | BP | 15 | ARG | 6.6 |
| 28 | B6 | 20 | ASN | 6.6 |
| 36 | DG | 35 | GLU | 6.6 |
| 49 | BX | 69 | TYR | 6.6 |
| 47 | BV | 68 | LYS | 6.5 |
| 37 | BH | 60 | ARG | 6.5 |
| 39 | BN | 131 | GLN | 6.5 |
| 31 | BA | 2477 | C | 6.5 |
| 9 | CI | 108 | VAL | 6.5 |
| 49 | BX | 34 | ALA | 6.5 |
| 29 | B7 | 47 | ARG | 6.5 |
| 45 | BT | 112 | ARG | 6.5 |
| 3 | AC | 192 | THR | 6.5 |
| 31 | BA | 2188 | C | 6.5 |
| 13 | AM | 27 | LYS | 6.5 |
| 23 | B1 | 78 | LYS | 6.4 |
| 23 | D1 | 50 | ARG | 6.4 |
| 31 | BA | 2104 | G | 6.4 |
| 20 | AT | 9 | ASN | 6.4 |
| 46 | DU | 97 | ASP | 6.4 |
| 23 | B1 | 19 | GLN | 6.4 |
| 47 | BV | 46 | VAL | 6.4 |
| 31 | BA | 1533 | G | 6.3 |
| 38 | BI | 121 | LYS | 6.3 |
| 24 | D2 | 37 | PHE | 6.3 |
| 32 | BB | 5 | C | 6.3 |
| 11 | AK | 90 | GLY | 6.3 |
| 36 | DG | 133 | LEU | 6.3 |
| 47 | BV | 60 | GLU | 6.3 |
| 51 | BZ | 171 | ILE | 6.2 |
| 6 | AF | 94 | GLN | 6.2 |
| 12 | AL | 91 | LYS | 6.2 |
| 11 | CK | 89 | ALA | 6.2 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 36 | BG | 142 | PRO | 6.2 |
| 23 | B1 | 50 | ARG | 6.2 |
| 31 | BA | 11 | G | 6.2 |
| 20 | AT | 10 | LEU | 6.2 |
| 37 | BH | 158 | HIS | 6.1 |
| 11 | CK | 128 | ALA | 6.1 |
| 10 | AJ | 58 | ASP | 6.1 |
| 1 | AA | 90 | U | 6.1 |
| 46 | BU | 97 | ASP | 6.1 |
| 2 | CB | 99 | GLY | 6.1 |
| 23 | D1 | 14 | VAL | 6.0 |
| 36 | DG | 134 | GLY | 6.0 |
| 12 | CL | 19 | ARG | 6.0 |
| 31 | DA | 2796 | U | 6.0 |
| 31 | BA | 508 | G | 6.0 |
| 36 | DG | 142 | PRO | 6.0 |
| 7 | CG | 2 | ALA | 6.0 |
| 3 | CC | 150 | LYS | 5.9 |
| 19 | CS | 27 | GLU | 5.9 |
| 39 | DN | 68 | GLU | 5.9 |
| 22 | B0 | 41 | ARG | 5.9 |
| 31 | BA | 1531 | C | 5.9 |
| 31 | DA | 652 | C | 5.9 |
| 32 | BB | 90 | A | 5.9 |
| 46 | BU | 118 | GLY | 5.9 |
| 46 | DU | 22 | LYS | 5.9 |
| 7 | AG | 32 | ARG | 5.9 |
| 47 | BV | 96 | ILE | 5.9 |
| 3 | AC | 195 | VAL | 5.9 |
| 33 | BD | 236 | GLY | 5.9 |
| 42 | DQ | 100 | GLY | 5.9 |
| 31 | BA | 1174 | A | 5.8 |
| 36 | BG | 63 | ILE | 5.8 |
| 9 | CI | 64 | THR | 5.8 |
| 31 | BA | 2402 | C | 5.8 |
| 45 | BT | 11 | GLU | 5.8 |
| 31 | BA | 1026 | U | 5.7 |
| 20 | CT | 9 | ASN | 5.7 |
| 12 | AL | 89 | ARG | 5.7 |
| 35 | DF | 207 | GLY | 5.7 |
| 37 | BH | 149 | ARG | 5.7 |
| 22 | B0 | 7 | LEU | 5.7 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 45 | BT | 109 | GLU | 5.7 |
| 13 | CM | 27 | LYS | 5.7 |
| 38 | DI | 100 | ALA | 5.7 |
| 37 | BH | 124 | GLU | 5.7 |
| 38 | BI | 73 | GLU | 5.7 |
| 31 | BA | 1544 | A | 5.7 |
| 42 | DQ | 21 | THR | 5.7 |
| 49 | DX | 33 | LYS | 5.7 |
| 22 | B0 | 1 | MET | 5.7 |
| 4 | AD | 3 | ARG | 5.7 |
| 20 | CT | 64 | ASP | 5.7 |
| 19 | AS | 49 | ILE | 5.7 |
| 31 | DA | 1533 | G | 5.6 |
| 21 | AU | 13 | ILE | 5.6 |
| 47 | BV | 28 | GLU | 5.6 |
| 51 | BZ | 123 | ASP | 5.6 |
| 30 | B8 | 63 | PRO | 5.6 |
| 30 | B8 | 64 | TYR | 5.6 |
| 21 | AU | 17 | THR | 5.6 |
| 21 | CU | 17 | THR | 5.6 |
| 21 | AU | 10 | ARG | 5.6 |
| 43 | BR | 2 | ARG | 5.6 |
| 5 | CE | 53 | LEU | 5.6 |
| 51 | BZ | 144 | LEU | 5.6 |
| 5 | AE | 88 | LYS | 5.6 |
| 23 | D1 | 15 | ALA | 5.6 |
| 36 | BG | 94 | LEU | 5.6 |
| 34 | BE | 61 | ARG | 5.6 |
| 51 | BZ | 80 | ARG | 5.6 |
| 31 | BA | 2105 | C | 5.6 |
| 51 | BZ | 169 | GLU | 5.5 |
| 7 | CG | 5 | ARG | 5.5 |
| 28 | D6 | 39 | TYR | 5.5 |
| 9 | AI | 81 | ILE | 5.5 |
| 50 | DY | 52 | SER | 5.5 |
| 23 | D1 | 95 | LEU | 5.5 |
| 46 | DU | 59 | ARG | 5.5 |
| 44 | BS | 89 | ARG | 5.5 |
| 45 | BT | 1 | MET | 5.5 |
| 8 | AH | 30 | ARG | 5.5 |
| 10 | AJ | 59 | SER | 5.5 |
| 50 | DY | 59 | GLY | 5.5 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 51 | BZ | 146 | ILE | 5.5 |
| 43 | DR | 2 | ARG | 5.5 |
| 50 | DY | 48 | ALA | 5.5 |
| 4 | CD | 6 | GLY | 5.5 |
| 2 | CB | 232 | PRO | 5.4 |
| 3 | CC | 159 | GLY | 5.4 |
| 9 | AI | 66 | ARG | 5.4 |
| 22 | B0 | 55 | ARG | 5.4 |
| 31 | BA | 2100 | G | 5.4 |
| 9 | AI | 106 | ALA | 5.4 |
| 32 | BB | 6 | C | 5.4 |
| 14 | CN | 31 | ARG | 5.4 |
| 45 | BT | 93 | ARG | 5.4 |
| 37 | BH | 59 | ARG | 5.4 |
| 46 | BU | 58 | ARG | 5.4 |
| 50 | DY | 60 | PHE | 5.4 |
| 24 | B2 | 49 | LYS | 5.4 |
| 24 | D2 | 54 | LYS | 5.4 |
| 49 | DX | 69 | TYR | 5.4 |
| 13 | AM | 63 | THR | 5.4 |
| 13 | CM | 107 | ALA | 5.4 |
| 3 | CC | 2 | GLY | 5.4 |
| 45 | BT | 113 | LYS | 5.4 |
| 2 | CB | 231 | GLU | 5.3 |
| 9 | CI | 62 | TYR | 5.3 |
| 15 | AO | 65 | ARG | 5.3 |
| 41 | DP | 65 | ARG | 5.3 |
| 37 | DH | 108 | GLY | 5.3 |
| 51 | DZ | 146 | ILE | 5.3 |
| 8 | CH | 99 | GLU | 5.3 |
| 1 | AA | 1280 | A | 5.3 |
| 1 | AA | 1287 | A | 5.3 |
| 47 | BV | 75 | PHE | 5.3 |
| 7 | CG | 80 | VAL | 5.3 |
| 32 | BB | 87 | G | 5.3 |
| 1 | CA | 344 | A | 5.3 |
| 21 | AU | 22 | ARG | 5.3 |
| 4 | CD | 134 | ASP | 5.3 |
| 47 | BV | 55 | ALA | 5.3 |
| 28 | B6 | 43 | CYS | 5.3 |
| 38 | BI | 91 | SER | 5.3 |
| 31 | BA | 271(K) | U | 5.3 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 16 | CP | 29 | ASP | 5.3 |
| 47 | DV | 81 | TYR | 5.3 |
| 10 | AJ | 10 | GLY | 5.2 |
| 16 | AP | 10 | GLY | 5.2 |
| 49 | BX | 35 | THR | 5.2 |
| 24 | B2 | 55 | ARG | 5.2 |
| 45 | BT | 39 | ARG | 5.2 |
| 50 | BY | 91 | GLU | 5.2 |
| 51 | BZ | 81 | ARG | 5.2 |
| 37 | BH | 85 | LYS | 5.2 |
| 1 | CA | 1149 | C | 5.2 |
| 44 | DS | 97 | ARG | 5.2 |
| 1 | CA | 84 | U | 5.2 |
| 24 | D2 | 60 | LEU | 5.2 |
| 42 | DQ | 23 | GLY | 5.2 |
| 45 | DT | 92 | GLY | 5.2 |
| 50 | BY | 43 | ASN | 5.2 |
| 47 | BV | 81 | TYR | 5.2 |
| 23 | D1 | 40 | ARG | 5.2 |
| 45 | BT | 3 | ARG | 5.2 |
| 8 | CH | 1 | MET | 5.2 |
| 32 | BB | 88 | C | 5.2 |
| 31 | DA | 2101 | G | 5.2 |
| 15 | AO | 3 | ILE | 5.1 |
| 21 | CU | 14 | TRP | 5.1 |
| 12 | AL | 16 | GLU | 5.1 |
| 39 | BN | 8 | GLN | 5.1 |
| 7 | CG | 27 | ILE | 5.1 |
| 36 | BG | 152 | LEU | 5.1 |
| 2 | AB | 70 | PHE | 5.1 |
| 31 | DA | 2189 | U | 5.1 |
| 4 | AD | 10 | ARG | 5.1 |
| 7 | CG | 39 | ALA | 5.1 |
| 19 | CS | 71 | LEU | 5.1 |
| 21 | CU | 13 | ILE | 5.1 |
| 30 | D8 | 56 | GLU | 5.1 |
| 32 | DB | 90 | A | 5.1 |
| 50 | BY | 33 | LYS | 5.1 |
| 13 | CM | 100 | GLY | 5.1 |
| 42 | BQ | 83 | MET | 5.1 |
| 44 | BS | 36 | TYR | 5.1 |
| 38 | DI | 68 | LEU | 5.1 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 37 | BH | 156 | ALA | 5.1 |
| 7 | CG | 33 | ASP | 5.1 |
| 50 | DY | 79 | CYS | 5.1 |
| 16 | AP | 28 | ARG | 5.1 |
| 24 | B2 | 59 | ARG | 5.1 |
| 9 | CI | 18 | PHE | 5.0 |
| 50 | BY | 57 | GLN | 5.0 |
| 37 | BH | 169 | VAL | 5.0 |
| 13 | AM | 100 | GLY | 5.0 |
| 31 | DA | 90 | U | 5.0 |
| 4 | CD | 73 | ARG | 5.0 |
| 12 | AL | 28 | LYS | 5.0 |
| 49 | DX | 77 | LYS | 5.0 |
| 50 | DY | 92 | ASN | 5.0 |
| 1 | CA | 1129 | C | 5.0 |
| 31 | BA | 1530 | C | 5.0 |
| 44 | BS | 107 | GLU | 5.0 |
| 26 | B4 | 4 | GLY | 5.0 |
| 51 | BZ | 72 | ARG | 5.0 |
| 14 | AN | 28 | GLY | 5.0 |
| 13 | CM | 97 | PRO | 5.0 |
| 51 | BZ | 53 | ILE | 5.0 |
| 4 | AD | 132 | ARG | 5.0 |
| 44 | BS | 11 | LYS | 5.0 |
| 9 | CI | 14 | VAL | 5.0 |
| 12 | CL | 18 | VAL | 5.0 |
| 13 | CM | 96 | LEU | 5.0 |
| 43 | DR | 10 | LEU | 5.0 |
| 14 | AN | 14 | PRO | 4.9 |
| 26 | D4 | 7 | PRO | 4.9 |
| 31 | BA | 603 | A | 4.9 |
| 30 | D8 | 40 | GLU | 4.9 |
| 45 | DT | 11 | GLU | 4.9 |
| 24 | B2 | 60 | LEU | 4.9 |
| 10 | CJ | 59 | SER | 4.9 |
| 13 | AM | 64 | TRP | 4.9 |
| 42 | BQ | 23 | GLY | 4.9 |
| 39 | BN | 129 | PRO | 4.9 |
| 9 | AI | 121 | ARG | 4.9 |
| 36 | BG | 73 | ALA | 4.9 |
| 43 | DR | 9 | LYS | 4.9 |
| 21 | AU | 21 | TYR | 4.9 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 8 | CH | 2 | LEU | 4.9 |
| 24 | B2 | 16 | LEU | 4.9 |
| 51 | DZ | 70 | LEU | 4.9 |
| 22 | B0 | 6 | GLY | 4.9 |
| 45 | BT | 2 | ASN | 4.9 |
| 1 | CA | 1286 | A | 4.9 |
| 31 | BA | 1049 | C | 4.9 |
| 50 | BY | 48 | ALA | 4.9 |
| 20 | CT | 73 | HIS | 4.9 |
| 51 | BZ | 179 | ASP | 4.9 |
| 4 | CD | 42 | GLN | 4.9 |
| 28 | B6 | 35 | GLU | 4.9 |
| 16 | CP | 13 | HIS | 4.9 |
| 31 | BA | 1532 | C | 4.9 |
| 31 | BA | 2803 | C | 4.9 |
| 31 | DA | 2402 | C | 4.9 |
| 34 | DE | 63 | LEU | 4.9 |
| 1 | AA | 1002 | G | 4.9 |
| 42 | BQ | 21 | THR | 4.8 |
| 16 | AP | 7 | ALA | 4.8 |
| 37 | BH | 12 | PRO | 4.8 |
| 13 | CM | 21 | TYR | 4.8 |
| 41 | BP | 110 | TYR | 4.8 |
| 21 | CU | 5 | ASP | 4.8 |
| 3 | CC | 160 | ALA | 4.8 |
| 16 | AP | 9 | PHE | 4.8 |
| 33 | BD | 33 | LEU | 4.8 |
| 44 | DS | 98 | VAL | 4.8 |
| 50 | DY | 6 | HIS | 4.8 |
| 3 | AC | 156 | ARG | 4.8 |
| 38 | BI | 7 | GLU | 4.8 |
| 12 | AL | 71 | PRO | 4.8 |
| 31 | BA | 2102 | U | 4.8 |
| 38 | DI | 121 | LYS | 4.8 |
| 1 | CA | 1224 | G | 4.8 |
| 11 | CK | 90 | GLY | 4.8 |
| 22 | B0 | 17 | GLN | 4.8 |
| 34 | BE | 17 | ASP | 4.8 |
| 49 | BX | 75 | ASP | 4.8 |
| 31 | BA | 12 | U | 4.8 |
| 2 | CB | 14 | GLY | 4.8 |
| 2 | AB | 71 | VAL | 4.8 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 9 | AI | 109 | VAL | 4.8 |
| 50 | BY | 17 | SER | 4.8 |
| 17 | CQ | 25 | ARG | 4.8 |
| 10 | CJ | 69 | ASN | 4.7 |
| 38 | DI | 71 | ILE | 4.7 |
| 14 | CN | 28 | GLY | 4.7 |
| 30 | D8 | 37 | SER | 4.7 |
| 32 | BB | 61 | G | 4.7 |
| 20 | CT | 65 | LYS | 4.7 |
| 30 | B8 | 40 | GLU | 4.7 |
| 31 | BA | 363(F) | A | 4.7 |
| 24 | B2 | 51 | ARG | 4.7 |
| 33 | BD | 262 | ARG | 4.7 |
| 41 | BP | 79 | ARG | 4.7 |
| 13 | AM | 20 | THR | 4.7 |
| 17 | CQ | 100 | LYS | 4.7 |
| 43 | BR | 105 | ARG | 4.7 |
| 37 | BH | 46 | GLU | 4.7 |
| 20 | AT | 64 | ASP | 4.7 |
| 1 | AA | 1224 | G | 4.7 |
| 12 | CL | 47 | LYS | 4.7 |
| 1 | CA | 82 | U | 4.7 |
| 9 | AI | 72 | GLY | 4.7 |
| 31 | BA | 656 | G | 4.7 |
| 37 | BH | 48 | GLY | 4.7 |
| 47 | BV | 86 | GLY | 4.7 |
| 16 | CP | 71 | ARG | 4.7 |
| 33 | BD | 244 | ARG | 4.7 |
| 37 | BH | 123 | PHE | 4.7 |
| 13 | CM | 98 | VAL | 4.7 |
| 12 | AL | 73 | GLU | 4.7 |
| 50 | DY | 44 | ILE | 4.6 |
| 49 | BX | 77 | LYS | 4.6 |
| 3 | AC | 2 | GLY | 4.6 |
| 20 | CT | 86 | ARG | 4.6 |
| 21 | CU | 15 | ARG | 4.6 |
| 34 | DE | 76 | ARG | 4.6 |
| 51 | BZ | 121 | HIS | 4.6 |
| 31 | DA | 274 | G | 4.6 |
| 51 | DZ | 97 | GLU | 4.6 |
| 31 | BA | 2892 | A | 4.6 |
| 1 | AA | 1149 | C | 4.6 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 32 | BB | 60 | C | 4.6 |
| 36 | BG | 3 | LEU | 4.6 |
| 22 | B0 | 2 | ALA | 4.6 |
| 39 | BN | 18 | ALA | 4.6 |
| 13 | AM | 32 | GLU | 4.6 |
| 38 | BI | 87 | LYS | 4.6 |
| 39 | DN | 74 | ARG | 4.6 |
| 45 | DT | 115 | ARG | 4.6 |
| 51 | DZ | 81 | ARG | 4.6 |
| 1 | AA | 1127 | G | 4.6 |
| 1 | CA | 1223 | C | 4.6 |
| 12 | AL | 101 | VAL | 4.6 |
| 16 | AP | 14 | ASN | 4.6 |
| 23 | D1 | 10 | LYS | 4.6 |
| 27 | B5 | 59 | GLU | 4.6 |
| 43 | BR | 11 | ASN | 4.6 |
| 3 | CC | 155 | GLY | 4.6 |
| 9 | CI | 17 | VAL | 4.6 |
| 13 | AM | 107 | ALA | 4.6 |
| 47 | BV | 76 | LYS | 4.6 |
| 42 | BQ | 16 | ARG | 4.6 |
| 30 | B8 | 35 | GLN | 4.6 |
| 50 | DY | 51 | VAL | 4.6 |
| 1 | CA | 1150 | U | 4.5 |
| 20 | AT | 60 | GLU | 4.5 |
| 28 | D6 | 24 | GLU | 4.5 |
| 44 | DS | 52 | SER | 4.5 |
| 31 | DA | 1544 | A | 4.5 |
| 30 | D8 | 48 | PHE | 4.5 |
| 43 | DR | 3 | HIS | 4.5 |
| 10 | AJ | 90 | LEU | 4.5 |
| 17 | AQ | 24 | GLU | 4.5 |
| 46 | BU | 89 | GLU | 4.5 |
| 51 | DZ | 179 | ASP | 4.5 |
| 44 | BS | 12 | PHE | 4.5 |
| 44 | BS | 14 | VAL | 4.5 |
| 7 | CG | 38 | LEU | 4.5 |
| 28 | D6 | 40 | CYS | 4.5 |
| 1 | AA | 84 | U | 4.5 |
| 21 | AU | 3 | LYS | 4.5 |
| 21 | CU | 9 | ARG | 4.5 |
| 36 | BG | 37 | VAL | 4.5 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 37 | BH | 137 | ASP | 4.5 |
| 50 | BY | 42 | VAL | 4.5 |
| 50 | BY | 89 | PHE | 4.5 |
| 1 | AA | 158 | G | 4.5 |
| 31 | BA | 2106 | G | 4.5 |
| 44 | BS | 16 | ASN | 4.5 |
| 47 | BV | 62 | LEU | 4.5 |
| 23 | D1 | 19 | GLN | 4.5 |
| 41 | DP | 50 | ARG | 4.5 |
| 31 | BA | 1109 | C | 4.4 |
| 35 | DF | 208 | GLY | 4.4 |
| 4 | CD | 5 | ILE | 4.4 |
| 1 | CA | 1034 | G | 4.4 |
| 31 | BA | 272(B) | G | 4.4 |
| 31 | BA | 2894 | G | 4.4 |
| 42 | DQ | 16 | ARG | 4.4 |
| 43 | BR | 71 | GLN | 4.4 |
| 24 | D2 | 61 | LEU | 4.4 |
| 11 | AK | 87 | THR | 4.4 |
| 1 | AA | 984 | C | 4.4 |
| 1 | CA | 1030(B) | C | 4.4 |
| 23 | D1 | 26 | ARG | 4.4 |
| 50 | BY | 2 | ARG | 4.4 |
| 9 | AI | 12 | GLU | 4.4 |
| 30 | B8 | 48 | PHE | 4.4 |
| 42 | BQ | 136 | ALA | 4.4 |
| 32 | BB | 89 | G | 4.4 |
| 50 | BY | 102 | CYS | 4.4 |
| 41 | BP | 27 | HIS | 4.4 |
| 51 | DZ | 88 | PHE | 4.4 |
| 30 | B8 | 32 | LEU | 4.4 |
| 21 | AU | 14 | TRP | 4.4 |
| 20 | AT | 58 | LYS | 4.4 |
| 21 | CU | 21 | TYR | 4.4 |
| 21 | CU | 11 | GLY | 4.4 |
| 16 | AP | 19 | ILE | 4.4 |
| 47 | BV | 38 | LEU | 4.4 |
| 1 | AA | 369 | C | 4.4 |
| 17 | AQ | 97 | SER | 4.4 |
| 4 | AD | 8 | VAL | 4.4 |
| 2 | AB | 163 | PHE | 4.3 |
| 3 | AC | 158 | GLY | 4.3 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 4 | CD | 156 | GLU | 4.3 |
| 37 | BH | 81 | GLU | 4.3 |
| 1 | AA | 1001(A) | G | 4.3 |
| 23 | B1 | 10 | LYS | 4.3 |
| 31 | BA | 651 | G | 4.3 |
| 31 | DA | 275 | G | 4.3 |
| 37 | BH | 138 | LYS | 4.3 |
| 22 | B0 | 56 | ASP | 4.3 |
| 28 | B6 | 23 | THR | 4.3 |
| 10 | AJ | 35 | SER | 4.3 |
| 28 | B6 | 26 | ASN | 4.3 |
| 39 | DN | 1 | MET | 4.3 |
| 42 | BQ | 137 | TYR | 4.3 |
| 17 | AQ | 43 | LEU | 4.3 |
| 36 | BG | 129 | GLY | 4.3 |
| 14 | AN | 29 | ARG | 4.3 |
| 22 | B0 | 3 | HIS | 4.3 |
| 31 | DA | 272(B) | G | 4.3 |
| 32 | DB | 89 | G | 4.3 |
| 50 | BY | 27 | VAL | 4.3 |
| 1 | AA | 1129 | C | 4.3 |
| 10 | AJ | 52 | GLY | 4.3 |
| 51 | DZ | 169 | GLU | 4.3 |
| 42 | BQ | 90 | VAL | 4.3 |
| 44 | BS | 98 | VAL | 4.3 |
| 8 | AH | 119 | LEU | 4.3 |
| 14 | AN | 41 | ARG | 4.3 |
| 31 | BA | 919 | G | 4.3 |
| 32 | DB | 91 | C | 4.3 |
| 37 | BH | 139 | GLN | 4.3 |
| 39 | BN | 73 | THR | 4.3 |
| 47 | DV | 45 | THR | 4.3 |
| 10 | CJ | 58 | ASP | 4.3 |
| 41 | BP | 74 | GLU | 4.3 |
| 16 | AP | 1 | MET | 4.2 |
| 32 | BB | 54 | G | 4.2 |
| 19 | AS | 71 | LEU | 4.2 |
| 28 | B6 | 39 | TYR | 4.2 |
| 46 | DU | 56 | ASP | 4.2 |
| 8 | AH | 127 | LEU | 4.2 |
| 21 | AU | 15 | ARG | 4.2 |
| 30 | D8 | 50 | LEU | 4.2 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 48 | DW | 92 | ARG | 4.2 |
| 8 | AH | 55 | GLY | 4.2 |
| 36 | BG | 134 | GLY | 4.2 |
| 1 | AA | 1139 | G | 4.2 |
| 1 | AA | 1190 | G | 4.2 |
| 31 | DA | 1049 | C | 4.2 |
| 18 | AR | 40 | LEU | 4.2 |
| 22 | D0 | 1 | MET | 4.2 |
| 11 | AK | 60 | ALA | 4.2 |
| 36 | DG | 157 | ILE | 4.2 |
| 35 | DF | 43 | LYS | 4.2 |
| 41 | BP | 144 | GLU | 4.2 |
| 13 | CM | 7 | VAL | 4.2 |
| 29 | D7 | 23 | ARG | 4.2 |
| 45 | DT | 1 | MET | 4.2 |
| 1 | CA | 1202 | G | 4.2 |
| 31 | BA | 1112 | G | 4.2 |
| 15 | AO | 87 | ILE | 4.2 |
| 12 | AL | 72 | GLY | 4.2 |
| 31 | DA | 1026 | U | 4.2 |
| 44 | BS | 27 | SER | 4.2 |
| 44 | DS | 12 | PHE | 4.2 |
| 1 | CA | 345 | C | 4.2 |
| 31 | BA | 2183 | C | 4.2 |
| 38 | DI | 118 | LYS | 4.2 |
| 31 | BA | 10 | G | 4.2 |
| 31 | DA | 11 | G | 4.2 |
| 44 | DS | 14 | VAL | 4.1 |
| 10 | CJ | 40 | LEU | 4.1 |
| 30 | B8 | 50 | LEU | 4.1 |
| 2 | CB | 163 | PHE | 4.1 |
| 7 | AG | 110 | GLN | 4.1 |
| 4 | AD | 12 | CYS | 4.1 |
| 29 | B7 | 48 | LYS | 4.1 |
| 23 | B1 | 20 | ARG | 4.1 |
| 23 | B1 | 40 | ARG | 4.1 |
| 21 | AU | 18 | TYR | 4.1 |
| 31 | BA | 1048 | A | 4.1 |
| 34 | BE | 59 | VAL | 4.1 |
| 37 | BH | 44 | VAL | 4.1 |
| 31 | BA | 1053 | C | 4.1 |
| 32 | BB | 62 | C | 4.1 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 42 | DQ | 80 | GLU | 4.1 |
| 24 | B2 | 41 | ILE | 4.1 |
| 38 | BI | 55 | ALA | 4.1 |
| 46 | BU | 91 | ASP | 4.1 |
| 13 | AM | 99 | ARG | 4.1 |
| 36 | DG | 72 | ARG | 4.1 |
| 33 | BD | 34 | VAL | 4.1 |
| 8 | CH | 36 | LEU | 4.1 |
| 1 | AA | 1503 | A | 4.1 |
| 1 | CA | 1285 | A | 4.1 |
| 2 | AB | 214 | ILE | 4.1 |
| 13 | AM | 62 | ASN | 4.1 |
| 23 | B1 | 26 | ARG | 4.1 |
| 28 | B6 | 37 | ARG | 4.1 |
| 34 | BE | 66 | HIS | 4.1 |
| 41 | DP | 27 | HIS | 4.1 |
| 2 | CB | 155 | LEU | 4.1 |
| 4 | AD | 31 | CYS | 4.1 |
| 50 | BY | 51 | VAL | 4.1 |
| 50 | BY | 79 | CYS | 4.1 |
| 25 | B3 | 1 | MET | 4.1 |
| 2 | CB | 133 | LYS | 4.1 |
| 21 | CU | 3 | LYS | 4.1 |
| 23 | B1 | 32 | LYS | 4.1 |
| 39 | BN | 115 | ARG | 4.1 |
| 50 | DY | 2 | ARG | 4.1 |
| 7 | AG | 20 | ASP | 4.1 |
| 13 | AM | 24 | GLY | 4.1 |
| 13 | CM | 62 | ASN | 4.1 |
| 4 | AD | 21 | LEU | 4.1 |
| 36 | DG | 155 | MET | 4.1 |
| 5 | AE | 122 | GLU | 4.1 |
| 14 | AN | 7 | ILE | 4.1 |
| 38 | BI | 97 | ILE | 4.1 |
| 20 | CT | 12 | ALA | 4.1 |
| 2 | AB | 137 | ARG | 4.1 |
| 17 | AQ | 68 | ARG | 4.1 |
| 32 | BB | 119 | G | 4.1 |
| 41 | DP | 7 | ARG | 4.1 |
| 41 | DP | 77 | ARG | 4.1 |
| 46 | BU | 59 | ARG | 4.1 |
| 31 | BA | 2796 | U | 4.1 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 9 | CI | 123 | PRO | 4.1 |
| 45 | BT | 25 | GLY | 4.1 |
| 45 | BT | 92 | GLY | 4.1 |
| 10 | AJ | 85 | LEU | 4.0 |
| 12 | CL | 21 | LYS | 4.0 |
| 19 | CS | 32 | LYS | 4.0 |
| 31 | DA | 897 | C | 4.0 |
| 36 | DG | 172 | LEU | 4.0 |
| 31 | BA | 1108 | U | 4.0 |
| 31 | BA | 1740 | G | 4.0 |
| 42 | BQ | 135 | ASP | 4.0 |
| 21 | CU | 8 | THR | 4.0 |
| 49 | DX | 26 | TYR | 4.0 |
| 1 | AA | 1201 | A | 4.0 |
| 31 | BA | 878 | A | 4.0 |
| 31 | DA | 1531 | C | 4.0 |
| 10 | CJ | 53 | PRO | 4.0 |
| 30 | D8 | 34 | TRP | 4.0 |
| 49 | BX | 33 | LYS | 4.0 |
| 50 | BY | 4 | LYS | 4.0 |
| 24 | B2 | 50 | ILE | 4.0 |
| 41 | BP | 33 | ARG | 4.0 |
| 51 | DZ | 99 | TYR | 4.0 |
| 24 | D2 | 12 | GLU | 4.0 |
| 1 | AA | 1026 | G | 4.0 |
| 2 | CB | 7 | VAL | 4.0 |
| 36 | BG | 70 | VAL | 4.0 |
| 23 | B1 | 15 | ALA | 4.0 |
| 26 | D4 | 32 | TYR | 4.0 |
| 30 | D8 | 64 | TYR | 4.0 |
| 3 | CC | 196 | LEU | 4.0 |
| 39 | BN | 118 | LYS | 4.0 |
| 44 | DS | 54 | LEU | 4.0 |
| 6 | AF | 90 | VAL | 4.0 |
| 46 | BU | 64 | ARG | 4.0 |
| 5 | AE | 89 | ILE | 4.0 |
| 24 | B2 | 57 | ILE | 4.0 |
| 38 | DI | 97 | ILE | 4.0 |
| 31 | BA | 2662 | A | 4.0 |
| 31 | BA | 2103 | C | 4.0 |
| 31 | DA | 2188 | C | 4.0 |
| 36 | BG | 76 | SER | 4.0 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 51 | BZ | 149 | SER | 4.0 |
| 7 | CG | 35 | LYS | 4.0 |
| 24 | D2 | 32 | LEU | 4.0 |
| 8 | AH | 93 | VAL | 3.9 |
| 21 | CU | 2 | GLY | 3.9 |
| 7 | AG | 31 | MET | 3.9 |
| 24 | B2 | 33 | MET | 3.9 |
| 46 | DU | 58 | ARG | 3.9 |
| 24 | D2 | 48 | HIS | 3.9 |
| 36 | BG | 39 | ILE | 3.9 |
| 50 | BY | 60 | PHE | 3.9 |
| 3 | AC | 3 | ASN | 3.9 |
| 42 | BQ | 130 | LYS | 3.9 |
| 46 | DU | 54 | LYS | 3.9 |
| 10 | AJ | 88 | LEU | 3.9 |
| 31 | BA | 2190 | G | 3.9 |
| 23 | B1 | 14 | VAL | 3.9 |
| 32 | BB | 25 | A | 3.9 |
| 4 | AD | 7 | PRO | 3.9 |
| 1 | AA | 218 | C | 3.9 |
| 1 | AA | 345 | C | 3.9 |
| 43 | BR | 91 | GLN | 3.9 |
| 10 | CJ | 20 | ALA | 3.9 |
| 14 | CN | 2 | ALA | 3.9 |
| 45 | BT | 85 | LYS | 3.9 |
| 26 | B4 | 20 | ASN | 3.9 |
| 49 | DX | 35 | THR | 3.9 |
| 45 | DT | 93 | ARG | 3.9 |
| 51 | DZ | 87 | ASP | 3.9 |
| 1 | CA | 88 | A | 3.9 |
| 1 | CA | 1032 | G | 3.9 |
| 31 | BA | 2602 | A | 3.9 |
| 32 | BB | 23 | G | 3.9 |
| 50 | BY | 44 | ILE | 3.9 |
| 19 | AS | 69 | HIS | 3.9 |
| 11 | CK | 127 | LYS | 3.9 |
| 14 | AN | 30 | ALA | 3.9 |
| 1 | AA | 1148 | U | 3.9 |
| 21 | CU | 18 | TYR | 3.9 |
| 28 | B6 | 21 | TYR | 3.9 |
| 45 | BT | 32 | TYR | 3.9 |
| 38 | BI | 135 | GLU | 3.9 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 47 | BV | 37 | VAL | 3.9 |
| 20 | CT | 75 | ASN | 3.9 |
| 26 | D4 | 20 | ASN | 3.9 |
| 35 | BF | 23 | ASP | 3.9 |
| 8 | AH | 116 | LYS | 3.9 |
| 18 | CR | 68 | LYS | 3.9 |
| 36 | BG | 75 | LYS | 3.9 |
| 37 | BH | 62 | LYS | 3.9 |
| 9 | CI | 106 | ALA | 3.9 |
| 22 | B0 | 74 | ARG | 3.9 |
| 24 | B2 | 30 | ARG | 3.9 |
| 1 | CA | 947 | G | 3.9 |
| 9 | AI | 108 | VAL | 3.9 |
| 13 | AM | 87 | TYR | 3.9 |
| 9 | CI | 89 | ASN | 3.9 |
| 31 | BA | 1175 | U | 3.9 |
| 31 | DA | 2102 | U | 3.9 |
| 12 | AL | 47 | LYS | 3.9 |
| 30 | B8 | 31 | HIS | 3.9 |
| 4 | CD | 3 | ARG | 3.8 |
| 13 | AM | 29 | ARG | 3.8 |
| 16 | AP | 42 | ARG | 3.8 |
| 41 | BP | 65 | ARG | 3.8 |
| 9 | CI | 114 | TYR | 3.8 |
| 11 | CK | 31 | THR | 3.8 |
| 31 | BA | 918 | A | 3.8 |
| 31 | DA | 2801(A) | A | 3.8 |
| 41 | BP | 104 | GLY | 3.8 |
| 10 | AJ | 47 | PHE | 3.8 |
| 4 | CD | 21 | LEU | 3.8 |
| 2 | CB | 136 | VAL | 3.8 |
| 8 | AH | 1 | MET | 3.8 |
| 9 | AI | 127 | LYS | 3.8 |
| 20 | CT | 74 | LYS | 3.8 |
| 37 | BH | 32 | GLU | 3.8 |
| 11 | AK | 49 | GLY | 3.8 |
| 12 | AL | 29 | GLY | 3.8 |
| 15 | AO | 2 | PRO | 3.8 |
| 39 | DN | 73 | THR | 3.8 |
| 45 | DT | 27 | THR | 3.8 |
| 9 | AI | 29 | ASN | 3.8 |
| 5 | AE | 53 | LEU | 3.8 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 11 | CK | 129 | SER | 3.8 |
| 15 | AO | 88 | ARG | 3.8 |
| 21 | AU | 7 | ARG | 3.8 |
| 44 | DS | 15 | ARG | 3.8 |
| 47 | BV | 51 | VAL | 3.8 |
| 23 | B1 | 39 | LYS | 3.8 |
| 24 | D2 | 49 | LYS | 3.8 |
| 29 | B7 | 2 | LYS | 3.8 |
| 44 | DS | 33 | LYS | 3.8 |
| 39 | DN | 75 | TYR | 3.8 |
| 4 | AD | 5 | ILE | 3.8 |
| 38 | DI | 138 | ILE | 3.8 |
| 41 | BP | 34 | GLY | 3.8 |
| 9 | CI | 66 | ARG | 3.8 |
| 23 | B1 | 82 | LEU | 3.8 |
| 43 | BR | 10 | LEU | 3.8 |
| 45 | BT | 6 | LEU | 3.8 |
| 37 | DH | 147 | ASN | 3.8 |
| 36 | DG | 41 | GLN | 3.8 |
| 12 | CL | 51 | ALA | 3.8 |
| 41 | BP | 87 | ASP | 3.8 |
| 51 | BZ | 54 | HIS | 3.8 |
| 30 | D8 | 39 | LYS | 3.8 |
| 44 | DS | 11 | LYS | 3.8 |
| 1 | AA | 1349 | A | 3.8 |
| 31 | BA | 860 | U | 3.8 |
| 1 | AA | 985 | C | 3.8 |
| 31 | DA | 2803 | C | 3.8 |
| 13 | AM | 97 | PRO | 3.8 |
| 41 | BP | 109 | GLY | 3.8 |
| 48 | BW | 45 | TYR | 3.8 |
| 2 | AB | 144 | ARG | 3.8 |
| 2 | CB | 67 | THR | 3.8 |
| 7 | CG | 12 | LEU | 3.8 |
| 12 | AL | 19 | ARG | 3.8 |
| 20 | AT | 80 | ARG | 3.8 |
| 35 | DF | 12 | LEU | 3.8 |
| 39 | DN | 67 | LEU | 3.8 |
| 33 | BD | 44 | ASN | 3.8 |
| 16 | AP | 12 | LYS | 3.8 |
| 19 | AS | 4 | SER | 3.8 |
| 46 | BU | 19 | LYS | 3.8 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 49 | BX | 72 | LYS | 3.8 |
| 49 | DX | 50 | LYS | 3.8 |
| 24 | B2 | 52 | ASP | 3.8 |
| 7 | AG | 49 | ILE | 3.8 |
| 20 | AT | 103 | GLY | 3.8 |
| 31 | DA | 1052 | C | 3.7 |
| 31 | DA | 1053 | C | 3.7 |
| 32 | BB | 49 | C | 3.7 |
| 28 | D6 | 20 | ASN | 3.7 |
| 1 | AA | 610 | G | 3.7 |
| 16 | CP | 8 | ARG | 3.7 |
| 38 | DI | 27 | ARG | 3.7 |
| 39 | BN | 74 | ARG | 3.7 |
| 10 | AJ | 41 | PRO | 3.7 |
| 36 | BG | 90 | LEU | 3.7 |
| 38 | DI | 77 | LEU | 3.7 |
| 41 | BP | 23 | PRO | 3.7 |
| 50 | BY | 63 | LYS | 3.7 |
| 9 | AI | 41 | VAL | 3.7 |
| 38 | BI | 107 | VAL | 3.7 |
| 41 | BP | 70 | GLN | 3.7 |
| 43 | DR | 11 | ASN | 3.7 |
| 37 | BH | 152 | ARG | 3.7 |
| 44 | BS | 15 | ARG | 3.7 |
| 8 | CH | 73 | ASP | 3.7 |
| 4 | AD | 6 | GLY | 3.7 |
| 38 | DI | 72 | LEU | 3.7 |
| 47 | BV | 41 | GLY | 3.7 |
| 10 | AJ | 37 | PRO | 3.7 |
| 12 | CL | 17 | LYS | 3.7 |
| 33 | BD | 4 | LYS | 3.7 |
| 1 | AA | 951 | G | 3.7 |
| 47 | BV | 45 | THR | 3.7 |
| 47 | BV | 58 | VAL | 3.7 |
| 28 | D6 | 13 | CYS | 3.7 |
| 34 | DE | 54 | GLN | 3.7 |
| 4 | CD | 66 | ARG | 3.7 |
| 22 | B0 | 12 | ASN | 3.7 |
| 43 | DR | 8 | ARG | 3.7 |
| 1 | AA | 1025 | U | 3.7 |
| 11 | CK | 42 | TRP | 3.7 |
| 50 | DY | 53 | PRO | 3.7 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 16 | CP | 84 | ALA | 3.7 |
| 10 | CJ | 24 | VAL | 3.7 |
| 20 | AT | 56 | MET | 3.7 |
| 47 | BV | 87 | HIS | 3.7 |
| 1 | CA | 158 | G | 3.7 |
| 7 | AG | 103 | TRP | 3.7 |
| 36 | BG | 133 | LEU | 3.7 |
| 36 | DG | 82 | LEU | 3.7 |
| 36 | DG | 108 | ASN | 3.7 |
| 2 | AB | 133 | LYS | 3.7 |
| 21 | AU | 2 | GLY | 3.7 |
| 26 | D4 | 8 | LYS | 3.7 |
| 45 | BT | 8 | LYS | 3.7 |
| 45 | DT | 69 | GLY | 3.7 |
| 45 | DT | 113 | LYS | 3.7 |
| 9 | AI | 126 | SER | 3.7 |
| 14 | AN | 32 | SER | 3.7 |
| 16 | CP | 11 | SER | 3.7 |
| 21 | AU | 23 | PRO | 3.7 |
| 10 | CJ | 17 | ASP | 3.7 |
| 44 | DS | 36 | TYR | 3.7 |
| 1 | CA | 1447 | A | 3.7 |
| 9 | CI | 41 | VAL | 3.7 |
| 44 | BS | 21 | THR | 3.7 |
| 1 | CA | 1141 | C | 3.7 |
| 33 | BD | 268 | ARG | 3.7 |
| 5 | CE | 20 | GLN | 3.7 |
| 51 | DZ | 76 | LEU | 3.7 |
| 2 | CB | 97 | TRP | 3.7 |
| 36 | BG | 108 | ASN | 3.7 |
| 46 | DU | 19 | LYS | 3.7 |
| 19 | CS | 82 | GLY | 3.7 |
| 8 | AH | 123 | GLU | 3.6 |
| 7 | AG | 9 | VAL | 3.6 |
| 38 | DI | 81 | VAL | 3.6 |
| 36 | DG | 98 | ARG | 3.6 |
| 41 | BP | 7 | ARG | 3.6 |
| 26 | B4 | 1 | MET | 3.6 |
| 2 | CB | 214 | ILE | 3.6 |
| 31 | DA | 2801 | A | 3.6 |
| 47 | DV | 70 | ILE | 3.6 |
| 51 | DZ | 163 | LEU | 3.6 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 41 | BP | 108 | LYS | 3.6 |
| 50 | BY | 8 | LYS | 3.6 |
| 50 | DY | 88 | LYS | 3.6 |
| 31 | DA | 2477 | C | 3.6 |
| 7 | CG | 26 | PHE | 3.6 |
| 35 | BF | 25 | PRO | 3.6 |
| 38 | BI | 132 | PRO | 3.6 |
| 4 | AD | 163 | GLU | 3.6 |
| 23 | B1 | 64 | ALA | 3.6 |
| 28 | B6 | 22 | ALA | 3.6 |
| 36 | BG | 92 | VAL | 3.6 |
| 46 | DU | 91 | ASP | 3.6 |
| 13 | AM | 4 | ILE | 3.6 |
| 15 | AO | 81 | LEU | 3.6 |
| 41 | DP | 105 | LEU | 3.6 |
| 45 | DT | 85 | LYS | 3.6 |
| 39 | BN | 130 | HIS | 3.6 |
| 44 | DS | 34 | HIS | 3.6 |
| 47 | DV | 87 | HIS | 3.6 |
| 13 | AM | 113 | PRO | 3.6 |
| 9 | CI | 16 | ARG | 3.6 |
| 14 | AN | 31 | ARG | 3.6 |
| 17 | CQ | 38 | ARG | 3.6 |
| 18 | AR | 32 | ARG | 3.6 |
| 31 | DA | 2799 | C | 3.6 |
| 32 | BB | 91 | C | 3.6 |
| 36 | DG | 160 | VAL | 3.6 |
| 49 | DX | 4 | ALA | 3.6 |
| 50 | BY | 65 | ALA | 3.6 |
| 9 | AI | 110 | GLU | 3.6 |
| 12 | CL | 127 | GLU | 3.6 |
| 16 | CP | 12 | LYS | 3.6 |
| 30 | B8 | 39 | LYS | 3.6 |
| 38 | BI | 72 | LEU | 3.6 |
| 50 | BY | 47 | LYS | 3.6 |
| 13 | CM | 112 | GLY | 3.6 |
| 5 | CE | 126 | ARG | 3.6 |
| 16 | AP | 71 | ARG | 3.6 |
| 36 | BG | 2 | PRO | 3.6 |
| 9 | AI | 122 | ALA | 3.6 |
| 14 | AN | 5 | ALA | 3.6 |
| 15 | CO | 11 | VAL | 3.6 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 1 | AA | 1202 | G | 3.6 |
| 4 | AD | 156 | GLU | 3.6 |
| 49 | BX | 93 | GLU | 3.6 |
| 1 | CA | 1503 | A | 3.6 |
| 4 | AD | 151 | LYS | 3.6 |
| 16 | AP | 17 | TYR | 3.6 |
| 36 | DG | 131 | TYR | 3.6 |
| 19 | AS | 15 | LEU | 3.6 |
| 41 | BP | 17 | LYS | 3.6 |
| 46 | DU | 16 | LYS | 3.6 |
| 10 | AJ | 19 | SER | 3.6 |
| 20 | AT | 55 | ILE | 3.6 |
| 32 | DB | 60 | C | 3.6 |
| 4 | CD | 122 | ARG | 3.6 |
| 9 | AI | 115 | GLY | 3.6 |
| 1 | CA | 1148 | U | 3.6 |
| 3 | CC | 3 | ASN | 3.6 |
| 6 | CF | 7 | ASN | 3.6 |
| 7 | AG | 29 | LYS | 3.6 |
| 12 | AL | 54 | LYS | 3.6 |
| 36 | DG | 75 | LYS | 3.6 |
| 19 | AS | 31 | ILE | 3.6 |
| 10 | CJ | 19 | SER | 3.6 |
| 11 | AK | 24 | SER | 3.6 |
| 20 | CT | 61 | SER | 3.6 |
| 24 | D2 | 62 | THR | 3.6 |
| 34 | BE | 174 | ASP | 3.6 |
| 51 | BZ | 148 | ASP | 3.6 |
| 1 | AA | 428 | G | 3.5 |
| 1 | CA | 1139 | G | 3.5 |
| 11 | CK | 91 | ARG | 3.5 |
| 30 | D8 | 46 | ARG | 3.5 |
| 31 | DA | 2655 | G | 3.5 |
| 34 | DE | 3 | GLY | 3.5 |
| 36 | DG | 100 | TRP | 3.5 |
| 50 | DY | 78 | ALA | 3.5 |
| 2 | AB | 187 | LEU | 3.5 |
| 28 | D6 | 34 | LEU | 3.5 |
| 7 | AG | 30 | ILE | 3.5 |
| 45 | BT | 104 | ASN | 3.5 |
| 47 | BV | 70 | ILE | 3.5 |
| 5 | AE | 25 | ARG | 3.5 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 16 | AP | 18 | ARG | 3.5 |
| 28 | D6 | 23 | THR | 3.5 |
| 33 | DD | 244 | ARG | 3.5 |
| 36 | DG | 33 | ARG | 3.5 |
| 38 | DI | 61 | ARG | 3.5 |
| 41 | DP | 70 | GLN | 3.5 |
| 23 | B1 | 33 | LYS | 3.5 |
| 29 | D7 | 48 | LYS | 3.5 |
| 43 | BR | 9 | LYS | 3.5 |
| 1 | AA | 1204 | A | 3.5 |
| 31 | DA | 829 | A | 3.5 |
| 5 | CE | 31 | LEU | 3.5 |
| 1 | AA | 105 | G | 3.5 |
| 31 | BA | 1107 | G | 3.5 |
| 32 | BB | 24 | G | 3.5 |
| 17 | CQ | 96 | GLU | 3.5 |
| 31 | DA | 1909 | C | 3.5 |
| 6 | CF | 4 | TYR | 3.5 |
| 13 | AM | 21 | TYR | 3.5 |
| 38 | BI | 74 | ASN | 3.5 |
| 21 | CU | 22 | ARG | 3.5 |
| 1 | AA | 961 | U | 3.5 |
| 22 | D0 | 7 | LEU | 3.5 |
| 23 | B1 | 80 | LEU | 3.5 |
| 24 | D2 | 50 | ILE | 3.5 |
| 35 | BF | 206 | ILE | 3.5 |
| 44 | BS | 18 | ILE | 3.5 |
| 7 | CG | 79 | ARG | 3.5 |
| 12 | CL | 73 | GLU | 3.5 |
| 28 | B6 | 12 | GLU | 3.5 |
| 1 | CA | 1130 | A | 3.5 |
| 31 | BA | 2801 | A | 3.5 |
| 37 | BH | 157 | TYR | 3.5 |
| 1 | CA | 381 | C | 3.5 |
| 31 | DA | 2103 | C | 3.5 |
| 47 | DV | 75 | PHE | 3.5 |
| 4 | CD | 31 | CYS | 3.5 |
| 13 | CM | 43 | THR | 3.5 |
| 21 | AU | 12 | LYS | 3.5 |
| 31 | BA | 2795 | G | 3.5 |
| 31 | DA | 2795 | G | 3.5 |
| 4 | CD | 152 | SER | 3.5 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 13 | AM | 112 | GLY | 3.5 |
| 14 | AN | 2 | ALA | 3.5 |
| 27 | D5 | 53 | ALA | 3.5 |
| 36 | BG | 163 | ALA | 3.5 |
| 2 | AB | 232 | PRO | 3.5 |
| 8 | AH | 2 | LEU | 3.5 |
| 34 | BE | 62 | PRO | 3.5 |
| 38 | DI | 58 | LEU | 3.5 |
| 19 | AS | 40 | ILE | 3.5 |
| 8 | CH | 22 | GLU | 3.5 |
| 38 | BI | 130 | TYR | 3.5 |
| 9 | AI | 118 | LYS | 3.5 |
| 13 | CM | 31 | LYS | 3.5 |
| 28 | D6 | 38 | LYS | 3.5 |
| 33 | DD | 255 | LYS | 3.5 |
| 8 | AH | 3 | THR | 3.5 |
| 24 | B2 | 62 | THR | 3.5 |
| 1 | CA | 1183 | A | 3.5 |
| 4 | CD | 23 | GLY | 3.5 |
| 36 | BG | 28 | VAL | 3.5 |
| 44 | BS | 49 | VAL | 3.5 |
| 28 | D6 | 43 | CYS | 3.5 |
| 36 | DG | 26 | GLN | 3.5 |
| 17 | AQ | 66 | SER | 3.5 |
| 37 | BH | 71 | LEU | 3.5 |
| 51 | DZ | 113 | ALA | 3.5 |
| 13 | CM | 113 | PRO | 3.5 |
| 31 | BA | 34 | C | 3.5 |
| 31 | BA | 1543 | C | 3.5 |
| 15 | AO | 64 | ARG | 3.5 |
| 36 | BG | 33 | ARG | 3.5 |
| 40 | BO | 97 | ARG | 3.5 |
| 50 | DY | 73 | ARG | 3.5 |
| 15 | AO | 57 | LEU | 3.4 |
| 19 | AS | 56 | GLN | 3.4 |
| 34 | BE | 63 | LEU | 3.4 |
| 3 | CC | 152 | ILE | 3.4 |
| 3 | CC | 154 | SER | 3.4 |
| 24 | D2 | 59 | ARG | 3.4 |
| 45 | BT | 118 | ARG | 3.4 |
| 2 | AB | 97 | TRP | 3.4 |
| 5 | CE | 47 | LYS | 3.4 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 47 | DV | 60 | GLU | 3.4 |
| 31 | BA | 2655 | G | 3.4 |
| 31 | BA | 2661 | G | 3.4 |
| 31 | DA | 2106 | G | 3.4 |
| 37 | BH | 45 | VAL | 3.4 |
| 7 | AG | 99 | LEU | 3.4 |
| 12 | AL | 68 | ALA | 3.4 |
| 2 | CB | 137 | ARG | 3.4 |
| 21 | CU | 24 | ARG | 3.4 |
| 34 | DE | 66 | HIS | 3.4 |
| 36 | DG | 88 | ILE | 3.4 |
| 13 | AM | 31 | LYS | 3.4 |
| 15 | CO | 48 | LYS | 3.4 |
| 50 | DY | 47 | LYS | 3.4 |
| 47 | DV | 53 | GLU | 3.4 |
| 1 | AA | 1183 | A | 3.4 |
| 1 | CA | 974 | A | 3.4 |
| 3 | CC | 201 | TYR | 3.4 |
| 35 | BF | 7 | TYR | 3.4 |
| 10 | AJ | 16 | LEU | 3.4 |
| 20 | CT | 10 | LEU | 3.4 |
| 3 | AC | 159 | GLY | 3.4 |
| 9 | CI | 107 | ARG | 3.4 |
| 13 | CM | 5 | ALA | 3.4 |
| 20 | AT | 25 | ARG | 3.4 |
| 47 | BV | 21 | ARG | 3.4 |
| 1 | CA | 1205 | U | 3.4 |
| 47 | BV | 92 | THR | 3.4 |
| 2 | AB | 222 | ILE | 3.4 |
| 38 | DI | 119 | PRO | 3.4 |
| 38 | DI | 134 | PRO | 3.4 |
| 22 | B0 | 4 | LYS | 3.4 |
| 46 | DU | 93 | LYS | 3.4 |
| 1 | AA | 1094 | G | 3.4 |
| 31 | DA | 1559 | G | 3.4 |
| 7 | CG | 77 | SER | 3.4 |
| 13 | AM | 16 | ASP | 3.4 |
| 51 | BZ | 168 | GLU | 3.4 |
| 13 | AM | 15 | VAL | 3.4 |
| 36 | BG | 82 | LEU | 3.4 |
| 4 | AD | 66 | ARG | 3.4 |
| 24 | D2 | 51 | ARG | 3.4 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 2 | CB | 13 | ALA | 3.4 |
| 9 | CI | 43 | ALA | 3.4 |
| 16 | CP | 10 | GLY | 3.4 |
| 51 | DZ | 164 | ALA | 3.4 |
| 3 | CC | 202 | ILE | 3.4 |
| 31 | BA | 917 | A | 3.4 |
| 31 | BA | 1177 | A | 3.4 |
| 38 | BI | 120 | ILE | 3.4 |
| 43 | BR | 3 | HIS | 3.4 |
| 50 | BY | 83 | THR | 3.4 |
| 36 | BG | 36 | LYS | 3.4 |
| 10 | CJ | 46 | ARG | 3.4 |
| 11 | AK | 111 | ASP | 3.4 |
| 38 | DI | 107 | VAL | 3.4 |
| 42 | BQ | 10 | ARG | 3.4 |
| 45 | DT | 39 | ARG | 3.4 |
| 1 | CA | 346 | G | 3.4 |
| 1 | CA | 380 | G | 3.4 |
| 4 | CD | 69 | GLY | 3.4 |
| 1 | CA | 1492 | A | 3.3 |
| 2 | CB | 221 | LEU | 3.3 |
| 11 | AK | 14 | VAL | 3.3 |
| 12 | CL | 43 | VAL | 3.3 |
| 20 | AT | 79 | ARG | 3.3 |
| 29 | B7 | 23 | ARG | 3.3 |
| 31 | BA | 2794 | C | 3.3 |
| 31 | DA | 279 | C | 3.3 |
| 34 | DE | 58 | ARG | 3.3 |
| 36 | BG | 91 | ARG | 3.3 |
| 31 | DA | 2348 | U | 3.3 |
| 51 | BZ | 145 | GLU | 3.3 |
| 38 | BI | 65 | ALA | 3.3 |
| 39 | BN | 78 | TYR | 3.3 |
| 13 | CM | 111 | LYS | 3.3 |
| 16 | AP | 13 | HIS | 3.3 |
| 39 | DN | 131 | GLN | 3.3 |
| 1 | AA | 1206 | G | 3.3 |
| 12 | CL | 89 | ARG | 3.3 |
| 13 | CM | 99 | ARG | 3.3 |
| 14 | CN | 35 | ARG | 3.3 |
| 33 | BD | 269 | PHE | 3.3 |
| 41 | BP | 41 | ARG | 3.3 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 4 | CD | 157 | LEU | 3.3 |
| 13 | CM | 60 | VAL | 3.3 |
| 37 | BH | 49 | VAL | 3.3 |
| 39 | BN | 82 | LEU | 3.3 |
| 9 | CI | 12 | GLU | 3.3 |
| 51 | BZ | 84 | GLU | 3.3 |
| 1 | CA | 1531 | A | 3.3 |
| 36 | BG | 105 | LYS | 3.3 |
| 9 | CI | 125 | TYR | 3.3 |
| 14 | CN | 7 | ILE | 3.3 |
| 47 | DV | 96 | ILE | 3.3 |
| 31 | BA | 1052 | C | 3.3 |
| 31 | BA | 1914 | C | 3.3 |
| 4 | CD | 29 | PRO | 3.3 |
| 39 | BN | 3 | THR | 3.3 |
| 39 | DN | 133 | GLN | 3.3 |
| 5 | AE | 18 | ARG | 3.3 |
| 9 | CI | 111 | ARG | 3.3 |
| 10 | AJ | 70 | ARG | 3.3 |
| 29 | B7 | 49 | ARG | 3.3 |
| 50 | DY | 50 | ARG | 3.3 |
| 34 | DE | 67 | PHE | 3.3 |
| 36 | BG | 130 | ASN | 3.3 |
| 23 | D1 | 33 | LYS | 3.3 |
| 41 | DP | 107 | LYS | 3.3 |
| 44 | BS | 99 | LYS | 3.3 |
| 1 | AA | 108 | G | 3.3 |
| 31 | DA | 651 | G | 3.3 |
| 31 | DA | 1107 | G | 3.3 |
| 32 | DB | 54 | G | 3.3 |
| 20 | AT | 70 | SER | 3.3 |
| 41 | DP | 116 | GLY | 3.3 |
| 44 | DS | 18 | ILE | 3.3 |
| 2 | AB | 148 | TYR | 3.3 |
| 9 | AI | 123 | PRO | 3.3 |
| 12 | CL | 44 | THR | 3.3 |
| 36 | DG | 96 | ARG | 3.3 |
| 42 | BQ | 133 | ARG | 3.3 |
| 43 | DR | 71 | GLN | 3.3 |
| 32 | BB | 4 | C | 3.3 |
| 32 | BB | 27 | C | 3.3 |
| 11 | AK | 127 | LYS | 3.3 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 14 | AN | 15 | LYS | 3.3 |
| 19 | CS | 60 | VAL | 3.3 |
| 29 | D7 | 2 | LYS | 3.3 |
| 39 | DN | 84 | LYS | 3.3 |
| 50 | BY | 87 | LYS | 3.3 |
| 3 | CC | 124 | ILE | 3.3 |
| 4 | AD | 179 | GLU | 3.3 |
| 38 | BI | 71 | ILE | 3.3 |
| 41 | DP | 54 | GLY | 3.3 |
| 51 | BZ | 147 | GLY | 3.3 |
| 10 | AJ | 5 | ARG | 3.3 |
| 1 | AA | 380 | G | 3.3 |
| 32 | BB | 116 | G | 3.3 |
| 44 | DS | 56 | LEU | 3.3 |
| 45 | DT | 40 | THR | 3.3 |
| 19 | CS | 9 | VAL | 3.3 |
| 23 | B1 | 96 | LYS | 3.3 |
| 35 | BF | 11 | VAL | 3.3 |
| 47 | DV | 51 | VAL | 3.3 |
| 1 | CA | 1325 | C | 3.3 |
| 3 | CC | 169 | ALA | 3.3 |
| 12 | CL | 72 | GLY | 3.3 |
| 38 | BI | 52 | ARG | 3.2 |
| 41 | DP | 41 | ARG | 3.2 |
| 34 | DE | 62 | PRO | 3.2 |
| 44 | BS | 58 | LEU | 3.2 |
| 7 | AG | 26 | PHE | 3.2 |
| 9 | AI | 3 | GLN | 3.2 |
| 15 | AO | 62 | GLN | 3.2 |
| 34 | BE | 159 | HIS | 3.2 |
| 43 | DR | 42 | LYS | 3.2 |
| 16 | CP | 1 | MET | 3.2 |
| 7 | CG | 83 | ALA | 3.2 |
| 44 | DS | 55 | ALA | 3.2 |
| 1 | CA | 1442(A) | G | 3.2 |
| 31 | BA | 90 | U | 3.2 |
| 31 | BA | 1110 | G | 3.2 |
| 1 | CA | 1001 | A | 3.2 |
| 3 | AC | 190 | ARG | 3.2 |
| 22 | D0 | 74 | ARG | 3.2 |
| 25 | B3 | 29 | ARG | 3.2 |
| 51 | DZ | 72 | ARG | 3.2 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 51 | DZ | 80 | ARG | 3.2 |
| 1 | CA | 1128 | C | 3.2 |
| 9 | AI | 113 | LYS | 3.2 |
| 9 | CI | 126 | SER | 3.2 |
| 12 | CL | 27 | LEU | 3.2 |
| 37 | BH | 64 | LEU | 3.2 |
| 38 | BI | 8 | PRO | 3.2 |
| 38 | BI | 69 | LYS | 3.2 |
| 38 | BI | 95 | LYS | 3.2 |
| 41 | DP | 64 | LYS | 3.2 |
| 51 | BZ | 125 | LEU | 3.2 |
| 4 | AD | 9 | CYS | 3.2 |
| 13 | CM | 16 | ASP | 3.2 |
| 46 | BU | 88 | ILE | 3.2 |
| 13 | CM | 76 | ALA | 3.2 |
| 37 | BH | 165 | ALA | 3.2 |
| 4 | CD | 118 | ARG | 3.2 |
| 19 | AS | 82 | GLY | 3.2 |
| 20 | CT | 101 | GLY | 3.2 |
| 30 | B8 | 46 | ARG | 3.2 |
| 34 | BE | 58 | ARG | 3.2 |
| 36 | DG | 118 | ARG | 3.2 |
| 41 | BP | 143 | GLY | 3.2 |
| 44 | BS | 17 | ARG | 3.2 |
| 3 | AC | 201 | TYR | 3.2 |
| 10 | CJ | 65 | LEU | 3.2 |
| 12 | CL | 91 | LYS | 3.2 |
| 14 | CN | 34 | TYR | 3.2 |
| 17 | AQ | 100 | LYS | 3.2 |
| 36 | BG | 135 | LEU | 3.2 |
| 36 | DG | 8 | LYS | 3.2 |
| 47 | BV | 19 | LYS | 3.2 |
| 51 | BZ | 150 | LEU | 3.2 |
| 1 | AA | 1257 | U | 3.2 |
| 14 | CN | 14 | PRO | 3.2 |
| 1 | AA | 1138 | G | 3.2 |
| 3 | CC | 153 | VAL | 3.2 |
| 31 | DA | 1114 | G | 3.2 |
| 31 | DA | 1921 | G | 3.2 |
| 31 | DA | 2661 | G | 3.2 |
| 24 | B2 | 43 | GLN | 3.2 |
| 1 | AA | 1145 | C | 3.2 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 11 | AK | 110 | ASP | 3.2 |
| 49 | DX | 27 | THR | 3.2 |
| 36 | BG | 96 | ARG | 3.2 |
| 36 | DG | 46 | ALA | 3.2 |
| 44 | BS | 20 | ARG | 3.2 |
| 46 | DU | 86 | ALA | 3.2 |
| 3 | AC | 167 | TRP | 3.2 |
| 23 | B1 | 79 | GLY | 3.2 |
| 36 | DG | 29 | TRP | 3.2 |
| 42 | BQ | 48 | GLU | 3.2 |
| 50 | BY | 92 | ASN | 3.2 |
| 19 | CS | 80 | TYR | 3.2 |
| 34 | BE | 56 | PRO | 3.2 |
| 47 | BV | 64 | HIS | 3.2 |
| 22 | D0 | 17 | GLN | 3.2 |
| 2 | AB | 101 | MET | 3.2 |
| 5 | CE | 11 | ILE | 3.2 |
| 24 | D2 | 55 | ARG | 3.2 |
| 36 | BG | 71 | THR | 3.2 |
| 46 | BU | 96 | ALA | 3.2 |
| 1 | AA | 1331 | G | 3.2 |
| 10 | CJ | 7 | LYS | 3.2 |
| 31 | BA | 325 | G | 3.2 |
| 31 | BA | 1959 | G | 3.2 |
| 46 | BU | 93 | LYS | 3.2 |
| 50 | BY | 59 | GLY | 3.2 |
| 50 | DY | 33 | LYS | 3.2 |
| 7 | CG | 22 | LEU | 3.2 |
| 9 | AI | 40 | LEU | 3.2 |
| 23 | D1 | 27 | GLU | 3.2 |
| 23 | D1 | 82 | LEU | 3.2 |
| 42 | BQ | 38 | GLU | 3.2 |
| 45 | DT | 36 | GLU | 3.2 |
| 45 | DT | 2 | ASN | 3.2 |
| 7 | AG | 118 | VAL | 3.2 |
| 39 | BN | 79 | PRO | 3.2 |
| 48 | BW | 46 | PHE | 3.2 |
| 14 | CN | 12 | ARG | 3.2 |
| 20 | AT | 18 | GLN | 3.2 |
| 40 | DO | 5 | GLN | 3.2 |
| 42 | BQ | 6 | ARG | 3.2 |
| 42 | DQ | 136 | ALA | 3.2 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 31 | DA | 362 | U | 3.1 |
| 44 | BS | 41 | ASP | 3.1 |
| 46 | BU | 54 | LYS | 3.1 |
| 47 | DV | 68 | LYS | 3.1 |
| 22 | B0 | 13 | GLY | 3.1 |
| 3 | AC | 196 | LEU | 3.1 |
| 41 | BP | 105 | LEU | 3.1 |
| 13 | CM | 69 | GLU | 3.1 |
| 25 | B3 | 48 | GLU | 3.1 |
| 37 | BH | 34 | GLU | 3.1 |
| 42 | DQ | 48 | GLU | 3.1 |
| 1 | AA | 1001 | A | 3.1 |
| 8 | CH | 118 | VAL | 3.1 |
| 20 | AT | 75 | ASN | 3.1 |
| 2 | CB | 91 | PRO | 3.1 |
| 10 | AJ | 53 | PRO | 3.1 |
| 1 | AA | 1223 | C | 3.1 |
| 9 | AI | 107 | ARG | 3.1 |
| 19 | CS | 81 | ARG | 3.1 |
| 43 | BR | 8 | ARG | 3.1 |
| 51 | DZ | 98 | MET | 3.1 |
| 33 | BD | 38 | LYS | 3.1 |
| 36 | BG | 93 | THR | 3.1 |
| 50 | BY | 52 | SER | 3.1 |
| 2 | AB | 155 | LEU | 3.1 |
| 12 | AL | 52 | LEU | 3.1 |
| 28 | B6 | 11 | LEU | 3.1 |
| 36 | DG | 135 | LEU | 3.1 |
| 37 | BH | 67 | LEU | 3.1 |
| 38 | BI | 77 | LEU | 3.1 |
| 39 | DN | 82 | LEU | 3.1 |
| 45 | DT | 105 | LEU | 3.1 |
| 49 | BX | 6 | ASP | 3.1 |
| 37 | BH | 47 | GLU | 3.1 |
| 37 | BH | 53 | GLU | 3.1 |
| 12 | CL | 32 | PHE | 3.1 |
| 27 | D5 | 32 | PRO | 3.1 |
| 41 | DP | 18 | ARG | 3.1 |
| 45 | DT | 32 | TYR | 3.1 |
| 11 | CK | 22 | HIS | 3.1 |
| 21 | AU | 20 | LYS | 3.1 |
| 28 | B6 | 53 | LYS | 3.1 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 34 | BE | 14 | ILE | 3.1 |
| 41 | BP | 86 | LYS | 3.1 |
| 7 | AG | 116 | ALA | 3.1 |
| 30 | B8 | 61 | LEU | 3.1 |
| 49 | BX | 27 | THR | 3.1 |
| 20 | AT | 61 | SER | 3.1 |
| 1 | CA | 973 | G | 3.1 |
| 1 | CA | 1001(A) | G | 3.1 |
| 1 | CA | 1036 | G | 3.1 |
| 3 | CC | 205 | GLY | 3.1 |
| 5 | AE | 22 | GLY | 3.1 |
| 31 | DA | 1910 | G | 3.1 |
| 14 | AN | 8 | GLU | 3.1 |
| 2 | CB | 81 | VAL | 3.1 |
| 2 | CB | 230 | VAL | 3.1 |
| 3 | AC | 207 | VAL | 3.1 |
| 4 | AD | 26 | CYS | 3.1 |
| 10 | CJ | 54 | PHE | 3.1 |
| 13 | CM | 17 | VAL | 3.1 |
| 8 | CH | 27 | PRO | 3.1 |
| 5 | CE | 133 | TYR | 3.1 |
| 51 | DZ | 29 | TYR | 3.1 |
| 1 | AA | 1240 | U | 3.1 |
| 1 | AA | 1364 | U | 3.1 |
| 9 | CI | 74 | ILE | 3.1 |
| 12 | CL | 20 | LYS | 3.1 |
| 33 | BD | 5 | LYS | 3.1 |
| 4 | AD | 42 | GLN | 3.1 |
| 24 | B2 | 13 | ALA | 3.1 |
| 34 | BE | 205 | ALA | 3.1 |
| 34 | DE | 204 | ALA | 3.1 |
| 49 | BX | 82 | GLN | 3.1 |
| 36 | DG | 152 | LEU | 3.1 |
| 22 | D0 | 13 | GLY | 3.1 |
| 31 | DA | 1106 | A | 3.1 |
| 31 | DA | 2660 | A | 3.1 |
| 32 | BB | 48 | A | 3.1 |
| 41 | DP | 104 | GLY | 3.1 |
| 16 | AP | 11 | SER | 3.1 |
| 1 | CA | 1027 | C | 3.1 |
| 4 | CD | 139 | ARG | 3.1 |
| 10 | AJ | 46 | ARG | 3.1 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 31 | DA | 1543 | C | 3.1 |
| 45 | BT | 16 | ARG | 3.1 |
| 50 | DY | 42 | VAL | 3.1 |
| 16 | AP | 15 | PRO | 3.1 |
| 50 | DY | 77 | PRO | 3.1 |
| 1 | CA | 347 | G | 3.1 |
| 31 | DA | 2833 | G | 3.1 |
| 39 | BN | 75 | TYR | 3.1 |
| 10 | CJ | 68 | HIS | 3.1 |
| 13 | AM | 2 | ALA | 3.1 |
| 13 | AM | 96 | LEU | 3.1 |
| 28 | D6 | 46 | HIS | 3.1 |
| 17 | AQ | 7 | THR | 3.1 |
| 41 | BP | 82 | GLY | 3.1 |
| 45 | DT | 25 | GLY | 3.1 |
| 41 | DP | 33 | ARG | 3.1 |
| 3 | CC | 195 | VAL | 3.1 |
| 7 | CG | 21 | VAL | 3.1 |
| 23 | D1 | 38 | SER | 3.1 |
| 16 | CP | 48 | TRP | 3.1 |
| 28 | D6 | 51 | GLU | 3.1 |
| 1 | CA | 946 | A | 3.1 |
| 41 | BP | 14 | LYS | 3.1 |
| 39 | BN | 41 | ASP | 3.1 |
| 2 | AB | 58 | ILE | 3.1 |
| 21 | CU | 23 | PRO | 3.1 |
| 38 | BI | 138 | ILE | 3.1 |
| 1 | AA | 1027 | C | 3.0 |
| 31 | BA | 1881 | C | 3.0 |
| 32 | DB | 62 | C | 3.0 |
| 14 | AN | 6 | LEU | 3.0 |
| 39 | DN | 132 | ALA | 3.0 |
| 50 | BY | 31 | LEU | 3.0 |
| 20 | AT | 73 | HIS | 3.0 |
| 3 | AC | 155 | GLY | 3.0 |
| 21 | AU | 4 | GLY | 3.0 |
| 31 | BA | 271(I) | G | 3.0 |
| 31 | BA | 2186 | G | 3.0 |
| 32 | DB | 23 | G | 3.0 |
| 10 | CJ | 48 | THR | 3.0 |
| 12 | CL | 122 | THR | 3.0 |
| 1 | CA | 950 | U | 3.0 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 1 | CA | 1126 | U | 3.0 |
| 7 | CG | 9 | VAL | 3.0 |
| 4 | CD | 151 | LYS | 3.0 |
| 37 | BH | 153 | LYS | 3.0 |
| 19 | AS | 76 | PRO | 3.0 |
| 37 | BH | 154 | PRO | 3.0 |
| 51 | BZ | 87 | ASP | 3.0 |
| 11 | CK | 25 | TYR | 3.0 |
| 3 | CC | 149 | ALA | 3.0 |
| 17 | CQ | 98 | LEU | 3.0 |
| 28 | B6 | 13 | CYS | 3.0 |
| 31 | BA | 92 | A | 3.0 |
| 31 | DA | 1919 | A | 3.0 |
| 31 | DA | 2602 | A | 3.0 |
| 32 | BB | 26 | A | 3.0 |
| 37 | DH | 156 | ALA | 3.0 |
| 46 | BU | 60 | LEU | 3.0 |
| 45 | DT | 118 | ARG | 3.0 |
| 9 | AI | 30 | GLY | 3.0 |
| 31 | BA | 271(O) | C | 3.0 |
| 44 | BS | 33 | LYS | 3.0 |
| 49 | BX | 62 | LYS | 3.0 |
| 44 | DS | 87 | PHE | 3.0 |
| 4 | CD | 81 | GLU | 3.0 |
| 5 | AE | 81 | GLU | 3.0 |
| 1 | AA | 1281 | U | 3.0 |
| 31 | BA | 1917 | U | 3.0 |
| 36 | DG | 68 | PRO | 3.0 |
| 1 | CA | 447 | G | 3.0 |
| 4 | AD | 157 | LEU | 3.0 |
| 35 | DF | 24 | LEU | 3.0 |
| 38 | BI | 35 | LEU | 3.0 |
| 4 | AD | 195 | ALA | 3.0 |
| 46 | BU | 56 | ASP | 3.0 |
| 16 | CP | 42 | ARG | 3.0 |
| 45 | DT | 3 | ARG | 3.0 |
| 46 | BU | 52 | ARG | 3.0 |
| 10 | AJ | 56 | HIS | 3.0 |
| 20 | AT | 16 | HIS | 3.0 |
| 39 | BN | 109 | LYS | 3.0 |
| 50 | DY | 57 | GLN | 3.0 |
| 1 | AA | 532 | A | 3.0 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 36 | DG | 64 | THR | 3.0 |
| 41 | DP | 74 | GLU | 3.0 |
| 47 | BV | 50 | PRO | 3.0 |
| 36 | DG | 107 | LEU | 3.0 |
| 43 | DR | 6 | SER | 3.0 |
| 45 | BT | 105 | LEU | 3.0 |
| 20 | CT | 25 | ARG | 3.0 |
| 36 | BG | 72 | ARG | 3.0 |
| 37 | DH | 94 | TYR | 3.0 |
| 39 | BN | 134 | ARG | 3.0 |
| 41 | BP | 16 | ARG | 3.0 |
| 44 | DS | 92 | TYR | 3.0 |
| 16 | CP | 68 | ASP | 3.0 |
| 51 | DZ | 55 | HIS | 3.0 |
| 50 | BY | 76 | CYS | 3.0 |
| 50 | DY | 76 | CYS | 3.0 |
| 15 | CO | 15 | PHE | 3.0 |
| 11 | CK | 87 | THR | 3.0 |
| 49 | BX | 3 | THR | 3.0 |
| 2 | AB | 201 | ILE | 3.0 |
| 15 | AO | 58 | MET | 3.0 |
| 24 | D2 | 41 | ILE | 3.0 |
| 4 | AD | 19 | LEU | 3.0 |
| 5 | CE | 83 | GLU | 3.0 |
| 19 | AS | 5 | LEU | 3.0 |
| 19 | AS | 16 | LEU | 3.0 |
| 21 | AU | 24 | ARG | 3.0 |
| 33 | DD | 268 | ARG | 3.0 |
| 38 | BI | 50 | ARG | 3.0 |
| 22 | D0 | 9 | SER | 3.0 |
| 30 | D8 | 51 | ALA | 3.0 |
| 51 | BZ | 166 | SER | 3.0 |
| 4 | CD | 22 | LYS | 3.0 |
| 11 | CK | 50 | TYR | 3.0 |
| 34 | BE | 151 | TYR | 3.0 |
| 35 | BF | 108 | LYS | 3.0 |
| 41 | DP | 110 | TYR | 3.0 |
| 2 | AB | 165 | VAL | 3.0 |
| 4 | AD | 43 | HIS | 3.0 |
| 6 | CF | 91 | VAL | 3.0 |
| 9 | CI | 44 | VAL | 3.0 |
| 36 | BG | 160 | VAL | 3.0 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 37 | BH | 133 | VAL | 3.0 |
| 1 | CA | 427 | U | 3.0 |
| 17 | CQ | 27 | PHE | 3.0 |
| 23 | B1 | 18 | ILE | 3.0 |
| 38 | DI | 76 | THR | 3.0 |
| 50 | BY | 61 | ILE | 3.0 |
| 2 | AB | 11 | LEU | 2.9 |
| 38 | DI | 75 | LEU | 2.9 |
| 2 | AB | 12 | GLU | 2.9 |
| 5 | AE | 126 | ARG | 2.9 |
| 24 | B2 | 12 | GLU | 2.9 |
| 24 | B2 | 34 | GLU | 2.9 |
| 27 | B5 | 47 | PRO | 2.9 |
| 30 | D8 | 54 | GLU | 2.9 |
| 41 | BP | 77 | ARG | 2.9 |
| 47 | DV | 21 | ARG | 2.9 |
| 1 | AA | 524 | G | 2.9 |
| 1 | CA | 1026 | G | 2.9 |
| 31 | BA | 1816 | G | 2.9 |
| 7 | AG | 2 | ALA | 2.9 |
| 10 | AJ | 7 | LYS | 2.9 |
| 20 | CT | 39 | LYS | 2.9 |
| 20 | CT | 66 | ALA | 2.9 |
| 9 | CI | 92 | TYR | 2.9 |
| 42 | DQ | 32 | TYR | 2.9 |
| 31 | BA | 548 | A | 2.9 |
| 31 | BA | 1111 | A | 2.9 |
| 42 | DQ | 25 | ASP | 2.9 |
| 48 | DW | 111 | HIS | 2.9 |
| 37 | BH | 143 | GLN | 2.9 |
| 41 | DP | 51 | PHE | 2.9 |
| 1 | AA | 589 | C | 2.9 |
| 1 | AA | 932 | C | 2.9 |
| 37 | DH | 136 | ILE | 2.9 |
| 38 | DI | 120 | ILE | 2.9 |
| 2 | AB | 98 | LEU | 2.9 |
| 9 | AI | 120 | ARG | 2.9 |
| 31 | BA | 2449 | U | 2.9 |
| 31 | BA | 2506 | U | 2.9 |
| 31 | DA | 2506 | U | 2.9 |
| 43 | DR | 105 | ARG | 2.9 |
| 45 | BT | 115 | ARG | 2.9 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 47 | BV | 40 | LEU | 2.9 |
| 51 | BZ | 82 | ARG | 2.9 |
| 6 | AF | 5 | GLU | 2.9 |
| 43 | DR | 99 | LYS | 2.9 |
| 44 | DS | 99 | LYS | 2.9 |
| 49 | DX | 72 | LYS | 2.9 |
| 37 | BH | 43 | VAL | 2.9 |
| 44 | BS | 28 | VAL | 2.9 |
| 50 | BY | 22 | GLY | 2.9 |
| 1 | AA | 1117 | G | 2.9 |
| 1 | AA | 1343 | G | 2.9 |
| 1 | CA | 631 | G | 2.9 |
| 32 | DB | 87 | G | 2.9 |
| 34 | BE | 67 | PHE | 2.9 |
| 41 | BP | 68 | GLN | 2.9 |
| 51 | BZ | 120 | ILE | 2.9 |
| 1 | AA | 611 | A | 2.9 |
| 9 | AI | 47 | LEU | 2.9 |
| 10 | CJ | 70 | ARG | 2.9 |
| 51 | DZ | 157 | LEU | 2.9 |
| 7 | AG | 37 | ASN | 2.9 |
| 9 | AI | 11 | LYS | 2.9 |
| 33 | BD | 26 | LYS | 2.9 |
| 1 | CA | 1140 | C | 2.9 |
| 1 | CA | 1260 | C | 2.9 |
| 36 | DG | 87 | PRO | 2.9 |
| 38 | BI | 111 | PRO | 2.9 |
| 38 | DI | 122 | GLU | 2.9 |
| 43 | DR | 102 | GLU | 2.9 |
| 12 | AL | 43 | VAL | 2.9 |
| 16 | AP | 38 | TYR | 2.9 |
| 37 | DH | 123 | PHE | 2.9 |
| 41 | BP | 51 | PHE | 2.9 |
| 51 | BZ | 52 | SER | 2.9 |
| 12 | CL | 99 | HIS | 2.9 |
| 6 | CF | 89 | MET | 2.9 |
| 22 | D0 | 55 | ARG | 2.9 |
| 36 | DG | 16 | ARG | 2.9 |
| 36 | DG | 77 | ILE | 2.9 |
| 37 | BH | 69 | ARG | 2.9 |
| 9 | AI | 116 | LYS | 2.9 |
| 9 | CI | 11 | LYS | 2.9 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 25 | D3 | 17 | LYS | 2.9 |
| 1 | CA | 1030(C) | G | 2.9 |
| 7 | CG | 84 | ASN | 2.9 |
| 31 | BA | 2833 | G | 2.9 |
| 35 | BF | 128 | ALA | 2.9 |
| 47 | BV | 34 | GLU | 2.9 |
| 1 | AA | 1447 | A | 2.9 |
| 32 | BB | 52 | A | 2.9 |
| 32 | DB | 25 | A | 2.9 |
| 1 | AA | 488 | C | 2.9 |
| 31 | BA | 272(J) | C | 2.9 |
| 1 | CA | 1040 | U | 2.9 |
| 7 | AG | 3 | ARG | 2.9 |
| 9 | CI | 37 | PHE | 2.9 |
| 37 | BH | 166 | GLY | 2.9 |
| 41 | BP | 26 | GLY | 2.9 |
| 21 | CU | 7 | ARG | 2.9 |
| 44 | DS | 17 | ARG | 2.9 |
| 4 | AD | 33 | MET | 2.9 |
| 38 | BI | 79 | ILE | 2.9 |
| 7 | CG | 36 | LYS | 2.9 |
| 25 | B3 | 24 | LYS | 2.9 |
| 50 | DY | 21 | LYS | 2.9 |
| 51 | DZ | 78 | LYS | 2.9 |
| 49 | DX | 82 | GLN | 2.9 |
| 13 | AM | 43 | THR | 2.9 |
| 36 | BG | 64 | THR | 2.9 |
| 37 | DH | 12 | PRO | 2.9 |
| 50 | DY | 56 | PRO | 2.9 |
| 6 | AF | 7 | ASN | 2.9 |
| 24 | B2 | 31 | GLU | 2.9 |
| 47 | DV | 5 | VAL | 2.9 |
| 50 | BY | 24 | VAL | 2.9 |
| 33 | BD | 273 | ARG | 2.9 |
| 34 | DE | 65 | GLY | 2.9 |
| 36 | BG | 89 | GLY | 2.9 |
| 37 | DH | 60 | ARG | 2.9 |
| 1 | AA | 1531 | A | 2.9 |
| 38 | BI | 29 | TYR | 2.9 |
| 44 | BS | 92 | TYR | 2.9 |
| 1 | CA | 698 | G | 2.9 |
| 19 | AS | 32 | LYS | 2.9 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 20 | AT | 104 | LEU | 2.9 |
| 31 | DA | 919 | G | 2.9 |
| 35 | BF | 20 | LEU | 2.9 |
| 38 | BI | 68 | LEU | 2.9 |
| 38 | DI | 79 | ILE | 2.9 |
| 19 | AS | 14 | HIS | 2.8 |
| 1 | AA | 219 | C | 2.8 |
| 31 | BA | 271(Z) | C | 2.8 |
| 31 | BA | 645 | C | 2.8 |
| 11 | AK | 81 | ASP | 2.8 |
| 13 | CM | 72 | ALA | 2.8 |
| 34 | BE | 18 | ASP | 2.8 |
| 2 | AB | 67 | THR | 2.8 |
| 23 | B1 | 27 | GLU | 2.8 |
| 28 | B6 | 24 | GLU | 2.8 |
| 50 | BY | 37 | VAL | 2.8 |
| 15 | AO | 68 | ARG | 2.8 |
| 41 | BP | 18 | ARG | 2.8 |
| 44 | DS | 30 | ARG | 2.8 |
| 46 | DU | 52 | ARG | 2.8 |
| 12 | AL | 115 | LYS | 2.8 |
| 20 | AT | 62 | LEU | 2.8 |
| 21 | AU | 16 | GLY | 2.8 |
| 34 | BE | 57 | LYS | 2.8 |
| 23 | B1 | 13 | ILE | 2.8 |
| 23 | B1 | 43 | TYR | 2.8 |
| 39 | BN | 128 | HIS | 2.8 |
| 31 | BA | 6 | A | 2.8 |
| 31 | BA | 1545 | A | 2.8 |
| 9 | AI | 43 | ALA | 2.8 |
| 49 | BX | 4 | ALA | 2.8 |
| 1 | AA | 64 | G | 2.8 |
| 1 | CA | 1124 | G | 2.8 |
| 1 | CA | 1160 | G | 2.8 |
| 31 | DA | 2190 | G | 2.8 |
| 32 | BB | 7 | G | 2.8 |
| 2 | AB | 107 | THR | 2.8 |
| 4 | CD | 132 | ARG | 2.8 |
| 28 | B6 | 52 | VAL | 2.8 |
| 32 | BB | 28 | C | 2.8 |
| 36 | BG | 128 | ARG | 2.8 |
| 49 | BX | 23 | GLU | 2.8 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 50 | DY | 86 | ARG | 2.8 |
| 51 | BZ | 20 | ARG | 2.8 |
| 11 | AK | 55 | LYS | 2.8 |
| 33 | DD | 4 | LYS | 2.8 |
| 34 | DE | 57 | LYS | 2.8 |
| 38 | BI | 112 | LYS | 2.8 |
| 43 | BR | 99 | LYS | 2.8 |
| 10 | AJ | 71 | LEU | 2.8 |
| 34 | DE | 88 | GLY | 2.8 |
| 37 | DH | 171 | LEU | 2.8 |
| 50 | DY | 55 | TYR | 2.8 |
| 38 | BI | 139 | GLN | 2.8 |
| 41 | DP | 25 | SER | 2.8 |
| 44 | BS | 50 | SER | 2.8 |
| 8 | CH | 84 | ARG | 2.8 |
| 14 | AN | 3 | ARG | 2.8 |
| 25 | B3 | 44 | ARG | 2.8 |
| 41 | DP | 15 | ARG | 2.8 |
| 46 | DU | 101 | ARG | 2.8 |
| 2 | AB | 136 | VAL | 2.8 |
| 25 | B3 | 36 | VAL | 2.8 |
| 31 | BA | 547 | A | 2.8 |
| 31 | BA | 896 | A | 2.8 |
| 31 | BA | 2629 | A | 2.8 |
| 31 | BA | 2657 | A | 2.8 |
| 8 | CH | 119 | LEU | 2.8 |
| 14 | CN | 61 | TRP | 2.8 |
| 20 | CT | 72 | LEU | 2.8 |
| 31 | BA | 504 | U | 2.8 |
| 44 | BS | 48 | LEU | 2.8 |
| 11 | AK | 26 | ASN | 2.8 |
| 27 | D5 | 54 | GLY | 2.8 |
| 50 | BY | 58 | GLY | 2.8 |
| 1 | AA | 63 | C | 2.8 |
| 1 | AA | 1029 | C | 2.8 |
| 2 | AB | 33 | TYR | 2.8 |
| 31 | BA | 271(J) | C | 2.8 |
| 31 | DA | 1185 | C | 2.8 |
| 31 | DA | 1546 | C | 2.8 |
| 2 | AB | 218 | ALA | 2.8 |
| 7 | CG | 25 | ALA | 2.8 |
| 14 | CN | 30 | ALA | 2.8 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 16 | CP | 28 | ARG | 2.8 |
| 12 | AL | 18 | VAL | 2.8 |
| 33 | DD | 59 | LYS | 2.8 |
| 34 | DE | 56 | PRO | 2.8 |
| 39 | BN | 55 | VAL | 2.8 |
| 41 | BP | 132 | LYS | 2.8 |
| 46 | BU | 22 | LYS | 2.8 |
| 9 | CI | 103 | THR | 2.8 |
| 10 | AJ | 40 | LEU | 2.8 |
| 13 | CM | 19 | LEU | 2.8 |
| 20 | AT | 84 | LEU | 2.8 |
| 38 | BI | 128 | LEU | 2.8 |
| 42 | BQ | 80 | GLU | 2.8 |
| 3 | CC | 186 | PHE | 2.8 |
| 11 | CK | 110 | ASP | 2.8 |
| 34 | DE | 83 | ASP | 2.8 |
| 7 | CG | 109 | ASN | 2.8 |
| 1 | CA | 1086 | U | 2.8 |
| 1 | CA | 1236 | A | 2.8 |
| 3 | AC | 184 | TYR | 2.8 |
| 7 | CG | 85 | TYR | 2.8 |
| 9 | CI | 5 | TYR | 2.8 |
| 31 | BA | 2801(A) | A | 2.8 |
| 36 | DG | 25 | TYR | 2.8 |
| 37 | BH | 164 | TYR | 2.8 |
| 37 | BH | 61 | HIS | 2.8 |
| 50 | BY | 99 | CYS | 2.8 |
| 24 | D2 | 26 | ARG | 2.8 |
| 1 | AA | 1244 | C | 2.8 |
| 1 | CA | 1097 | C | 2.8 |
| 50 | BY | 78 | ALA | 2.8 |
| 30 | B8 | 3 | LYS | 2.8 |
| 35 | BF | 107 | LYS | 2.8 |
| 1 | CA | 1068 | G | 2.8 |
| 11 | CK | 84 | VAL | 2.8 |
| 12 | CL | 94 | PRO | 2.8 |
| 31 | BA | 1436 | G | 2.8 |
| 31 | BA | 1526 | G | 2.8 |
| 31 | DA | 508 | G | 2.8 |
| 31 | DA | 1922 | G | 2.8 |
| 32 | BB | 56 | G | 2.8 |
| 32 | BB | 86 | G | 2.8 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 38 | DI | 21 | VAL | 2.8 |
| 51 | BZ | 139 | VAL | 2.8 |
| 13 | CM | 105 | THR | 2.8 |
| 34 | BE | 73 | GLU | 2.8 |
| 37 | BH | 89 | ILE | 2.8 |
| 38 | BI | 34 | GLY | 2.7 |
| 41 | DP | 26 | GLY | 2.7 |
| 49 | BX | 67 | GLY | 2.7 |
| 11 | CK | 26 | ASN | 2.7 |
| 44 | BS | 61 | ASN | 2.7 |
| 36 | BG | 25 | TYR | 2.7 |
| 4 | CD | 10 | ARG | 2.7 |
| 4 | CD | 35 | ARG | 2.7 |
| 9 | AI | 13 | ALA | 2.7 |
| 9 | CI | 127 | LYS | 2.7 |
| 13 | CM | 42 | ALA | 2.7 |
| 14 | CN | 50 | LYS | 2.7 |
| 23 | D1 | 64 | ALA | 2.7 |
| 41 | DP | 31 | ALA | 2.7 |
| 47 | BV | 97 | LYS | 2.7 |
| 1 | AA | 609 | A | 2.7 |
| 2 | CB | 15 | VAL | 2.7 |
| 3 | CC | 173 | VAL | 2.7 |
| 7 | AG | 105 | VAL | 2.7 |
| 14 | AN | 18 | VAL | 2.7 |
| 16 | CP | 51 | VAL | 2.7 |
| 31 | DA | 363(F) | A | 2.7 |
| 38 | BI | 81 | VAL | 2.7 |
| 6 | AF | 48 | LEU | 2.7 |
| 9 | AI | 99 | LEU | 2.7 |
| 36 | DG | 34 | LEU | 2.7 |
| 10 | AJ | 50 | ILE | 2.7 |
| 31 | BA | 1464 | C | 2.7 |
| 49 | DX | 23 | GLU | 2.7 |
| 35 | BF | 134 | GLY | 2.7 |
| 36 | BG | 29 | TRP | 2.7 |
| 44 | BS | 22 | GLY | 2.7 |
| 44 | BS | 90 | GLY | 2.7 |
| 1 | AA | 79 | G | 2.7 |
| 31 | BA | 2659 | G | 2.7 |
| 23 | B1 | 52 | ARG | 2.7 |
| 2 | CB | 132 | LYS | 2.7 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 6 | AF | 50 | TYR | 2.7 |
| 17 | CQ | 37 | LYS | 2.7 |
| 46 | BU | 15 | LYS | 2.7 |
| 50 | BY | 95 | LYS | 2.7 |
| 50 | BY | 101 | LYS | 2.7 |
| 24 | B2 | 56 | GLN | 2.7 |
| 38 | DI | 37 | VAL | 2.7 |
| 42 | BQ | 45 | GLN | 2.7 |
| 47 | BV | 14 | VAL | 2.7 |
| 49 | BX | 59 | VAL | 2.7 |
| 3 | AC | 87 | LEU | 2.7 |
| 8 | CH | 57 | PRO | 2.7 |
| 10 | CJ | 39 | PRO | 2.7 |
| 1 | AA | 957 | U | 2.7 |
| 1 | CA | 1065 | U | 2.7 |
| 1 | AA | 393 | A | 2.7 |
| 1 | AA | 1130 | A | 2.7 |
| 3 | AC | 15 | THR | 2.7 |
| 10 | AJ | 36 | GLY | 2.7 |
| 22 | D0 | 41 | ARG | 2.7 |
| 30 | B8 | 57 | ARG | 2.7 |
| 51 | BZ | 79 | ARG | 2.7 |
| 15 | AO | 47 | LYS | 2.7 |
| 36 | DG | 36 | LYS | 2.7 |
| 1 | AA | 163 | C | 2.7 |
| 1 | AA | 1066 | C | 2.7 |
| 18 | AR | 30 | ASP | 2.7 |
| 31 | DA | 2347 | C | 2.7 |
| 11 | AK | 25 | TYR | 2.7 |
| 27 | B5 | 57 | VAL | 2.7 |
| 7 | AG | 59 | LEU | 2.7 |
| 18 | AR | 79 | LEU | 2.7 |
| 23 | B1 | 85 | LEU | 2.7 |
| 33 | BD | 133 | LEU | 2.7 |
| 35 | BF | 12 | LEU | 2.7 |
| 36 | BG | 43 | LEU | 2.7 |
| 38 | BI | 6 | LEU | 2.7 |
| 1 | AA | 97 | G | 2.7 |
| 1 | AA | 933 | G | 2.7 |
| 1 | CA | 798 | G | 2.7 |
| 1 | CA | 1186 | G | 2.7 |
| 19 | CS | 76 | PRO | 2.7 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 36 | BG | 26 | GLN | 2.7 |
| 10 | AJ | 38 | ILE | 2.7 |
| 34 | DE | 77 | ILE | 2.7 |
| 36 | DG | 39 | ILE | 2.7 |
| 26 | B4 | 16 | CYS | 2.7 |
| 26 | B4 | 30 | GLU | 2.7 |
| 51 | DZ | 119 | GLU | 2.7 |
| 2 | AB | 175 | ARG | 2.7 |
| 7 | AG | 4 | ARG | 2.7 |
| 7 | AG | 115 | ARG | 2.7 |
| 11 | CK | 126 | ARG | 2.7 |
| 20 | CT | 8 | ARG | 2.7 |
| 3 | CC | 199 | LYS | 2.7 |
| 9 | CI | 116 | LYS | 2.7 |
| 10 | AJ | 57 | LYS | 2.7 |
| 12 | CL | 115 | LYS | 2.7 |
| 14 | AN | 9 | LYS | 2.7 |
| 19 | CS | 4 | SER | 2.7 |
| 21 | CU | 25 | LYS | 2.7 |
| 1 | CA | 161 | A | 2.7 |
| 31 | BA | 2268 | A | 2.7 |
| 37 | DH | 157 | TYR | 2.7 |
| 49 | BX | 91 | ALA | 2.7 |
| 51 | DZ | 152 | ALA | 2.7 |
| 13 | AM | 7 | VAL | 2.7 |
| 13 | CM | 15 | VAL | 2.7 |
| 19 | CS | 15 | LEU | 2.7 |
| 38 | DI | 92 | VAL | 2.7 |
| 36 | DG | 27 | ASN | 2.7 |
| 1 | AA | 91 | C | 2.7 |
| 1 | AA | 1018 | C | 2.7 |
| 1 | AA | 1030(B) | C | 2.7 |
| 1 | AA | 1267 | C | 2.7 |
| 9 | CI | 124 | GLN | 2.7 |
| 32 | DB | 88 | C | 2.7 |
| 26 | D4 | 5 | ILE | 2.7 |
| 9 | AI | 70 | LYS | 2.7 |
| 9 | AI | 100 | GLY | 2.7 |
| 11 | AK | 124 | LYS | 2.7 |
| 26 | B4 | 18 | CYS | 2.7 |
| 35 | BF | 208 | GLY | 2.7 |
| 41 | DP | 43 | GLY | 2.7 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 35 | BF | 175 | THR | 2.7 |
| 47 | BV | 32 | THR | 2.7 |
| 1 | AA | 391 | G | 2.7 |
| 1 | CA | 1131 | G | 2.7 |
| 31 | BA | 921 | G | 2.7 |
| 19 | AS | 35 | SER | 2.7 |
| 5 | AE | 82 | VAL | 2.7 |
| 10 | CJ | 88 | LEU | 2.7 |
| 12 | CL | 52 | LEU | 2.7 |
| 36 | BG | 34 | LEU | 2.7 |
| 47 | BV | 71 | LEU | 2.7 |
| 50 | DY | 90 | LEU | 2.7 |
| 51 | BZ | 99 | TYR | 2.7 |
| 51 | BZ | 163 | LEU | 2.7 |
| 36 | BG | 156 | ASP | 2.7 |
| 41 | DP | 87 | ASP | 2.7 |
| 45 | DT | 58 | ASN | 2.7 |
| 7 | AG | 93 | PRO | 2.6 |
| 41 | BP | 10 | PRO | 2.6 |
| 47 | DV | 90 | PRO | 2.6 |
| 37 | BH | 41 | MET | 2.6 |
| 50 | BY | 5 | MET | 2.6 |
| 41 | DP | 79 | ARG | 2.6 |
| 48 | BW | 92 | ARG | 2.6 |
| 3 | AC | 4 | LYS | 2.6 |
| 6 | AF | 54 | LYS | 2.6 |
| 7 | AG | 35 | LYS | 2.6 |
| 9 | CI | 113 | LYS | 2.6 |
| 7 | AG | 34 | GLY | 2.6 |
| 1 | AA | 330 | C | 2.6 |
| 31 | DA | 34 | C | 2.6 |
| 37 | BH | 159 | GLU | 2.6 |
| 27 | B5 | 36 | CYS | 2.6 |
| 49 | BX | 37 | THR | 2.6 |
| 36 | BG | 46 | ALA | 2.6 |
| 41 | DP | 88 | LEU | 2.6 |
| 9 | AI | 114 | TYR | 2.6 |
| 38 | BI | 133 | HIS | 2.6 |
| 1 | AA | 70 | G | 2.6 |
| 1 | AA | 159 | G | 2.6 |
| 1 | AA | 1290 | G | 2.6 |
| 1 | CA | 1185 | G | 2.6 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 31 | DA | 2659 | G | 2.6 |
| 37 | BH | 57 | ASP | 2.6 |
| 45 | DT | 86 | ILE | 2.6 |
| 10 | AJ | 39 | PRO | 2.6 |
| 1 | AA | 1205 | U | 2.6 |
| 7 | AG | 63 | LYS | 2.6 |
| 11 | AK | 27 | ASN | 2.6 |
| 11 | CK | 13 | GLN | 2.6 |
| 12 | CL | 28 | LYS | 2.6 |
| 17 | CQ | 93 | GLN | 2.6 |
| 20 | AT | 29 | LYS | 2.6 |
| 24 | D2 | 38 | GLN | 2.6 |
| 31 | BA | 158 | U | 2.6 |
| 31 | DA | 1033 | U | 2.6 |
| 41 | DP | 17 | LYS | 2.6 |
| 44 | BS | 29 | PHE | 2.6 |
| 44 | BS | 76 | LYS | 2.6 |
| 45 | DT | 104 | ASN | 2.6 |
| 14 | CN | 8 | GLU | 2.6 |
| 31 | BA | 2311 | A | 2.6 |
| 10 | CJ | 90 | LEU | 2.6 |
| 16 | CP | 6 | LEU | 2.6 |
| 17 | AQ | 98 | LEU | 2.6 |
| 19 | AS | 51 | VAL | 2.6 |
| 22 | D0 | 2 | ALA | 2.6 |
| 39 | BN | 67 | LEU | 2.6 |
| 48 | BW | 44 | ALA | 2.6 |
| 1 | AA | 165 | C | 2.6 |
| 1 | AA | 435 | C | 2.6 |
| 1 | AA | 1137 | C | 2.6 |
| 1 | AA | 1203 | C | 2.6 |
| 1 | CA | 1284 | C | 2.6 |
| 31 | DA | 1166 | C | 2.6 |
| 41 | BP | 25 | SER | 2.6 |
| 4 | CD | 20 | TYR | 2.6 |
| 3 | CC | 14 | ILE | 2.6 |
| 9 | AI | 104 | ARG | 2.6 |
| 12 | AL | 99 | HIS | 2.6 |
| 12 | CL | 23 | LYS | 2.6 |
| 17 | AQ | 69 | LYS | 2.6 |
| 17 | CQ | 91 | ARG | 2.6 |
| 20 | CT | 68 | LYS | 2.6 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 23 | D1 | 78 | LYS | 2.6 |
| 33 | DD | 217 | ARG | 2.6 |
| 41 | BP | 139 | LYS | 2.6 |
| 44 | BS | 30 | ARG | 2.6 |
| 47 | DV | 74 | LYS | 2.6 |
| 7 | AG | 33 | ASP | 2.6 |
| 7 | CG | 13 | GLN | 2.6 |
| 35 | DF | 133 | ASN | 2.6 |
| 37 | DH | 139 | GLN | 2.6 |
| 4 | CD | 44 | GLY | 2.6 |
| 25 | B3 | 38 | GLU | 2.6 |
| 48 | BW | 78 | GLU | 2.6 |
| 1 | AA | 306 | G | 2.6 |
| 1 | AA | 346 | G | 2.6 |
| 31 | DA | 1112 | G | 2.6 |
| 3 | AC | 188 | LEU | 2.6 |
| 3 | CC | 43 | LEU | 2.6 |
| 12 | AL | 27 | LEU | 2.6 |
| 24 | D2 | 16 | LEU | 2.6 |
| 3 | CC | 151 | VAL | 2.6 |
| 5 | CE | 21 | ALA | 2.6 |
| 6 | AF | 88 | VAL | 2.6 |
| 7 | CG | 40 | ALA | 2.6 |
| 26 | D4 | 12 | ALA | 2.6 |
| 36 | DG | 69 | ALA | 2.6 |
| 44 | BS | 51 | ALA | 2.6 |
| 31 | BA | 1490 | A | 2.6 |
| 31 | BA | 2665 | A | 2.6 |
| 32 | DB | 48 | A | 2.6 |
| 36 | BG | 98 | ARG | 2.6 |
| 41 | DP | 14 | LYS | 2.6 |
| 17 | CQ | 99 | SER | 2.6 |
| 19 | CS | 10 | PHE | 2.6 |
| 37 | DH | 109 | PHE | 2.6 |
| 1 | CA | 89 | C | 2.6 |
| 31 | DA | 2105 | C | 2.6 |
| 32 | DB | 5 | C | 2.6 |
| 16 | AP | 68 | ASP | 2.6 |
| 21 | AU | 5 | ASP | 2.6 |
| 4 | AD | 124 | GLY | 2.6 |
| 7 | CG | 82 | GLY | 2.6 |
| 18 | AR | 48 | GLY | 2.6 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 19 | AS | 68 | GLY | 2.6 |
| 47 | BV | 65 | GLY | 2.6 |
| 10 | CJ | 71 | LEU | 2.6 |
| 14 | CN | 6 | LEU | 2.6 |
| 15 | AO | 31 | LEU | 2.6 |
| 45 | BT | 36 | GLU | 2.6 |
| 11 | AK | 30 | VAL | 2.6 |
| 34 | DE | 68 | ALA | 2.6 |
| 39 | BN | 132 | ALA | 2.6 |
| 45 | DT | 34 | VAL | 2.6 |
| 50 | BY | 7 | VAL | 2.6 |
| 51 | BZ | 71 | VAL | 2.6 |
| 9 | AI | 9 | ARG | 2.6 |
| 15 | AO | 63 | ARG | 2.6 |
| 17 | AQ | 91 | ARG | 2.6 |
| 23 | D1 | 20 | ARG | 2.6 |
| 34 | DE | 69 | LYS | 2.6 |
| 5 | AE | 118 | ILE | 2.6 |
| 10 | AJ | 23 | ILE | 2.6 |
| 19 | CS | 49 | ILE | 2.6 |
| 27 | B5 | 49 | CYS | 2.6 |
| 44 | BS | 35 | ILE | 2.6 |
| 1 | AA | 973 | G | 2.6 |
| 1 | CA | 1003 | G | 2.6 |
| 2 | CB | 148 | TYR | 2.6 |
| 13 | AM | 59 | TYR | 2.6 |
| 5 | CE | 49 | PRO | 2.6 |
| 11 | CK | 24 | SER | 2.6 |
| 31 | BA | 1423 | G | 2.6 |
| 31 | BA | 1525 | G | 2.6 |
| 32 | DB | 119 | G | 2.6 |
| 50 | BY | 66 | PRO | 2.6 |
| 4 | AD | 201 | GLN | 2.6 |
| 6 | CF | 73 | ASN | 2.6 |
| 9 | CI | 102 | LEU | 2.6 |
| 15 | AO | 66 | LEU | 2.6 |
| 17 | AQ | 6 | LEU | 2.6 |
| 1 | AA | 92 | C | 2.6 |
| 1 | CA | 1145 | C | 2.6 |
| 5 | AE | 83 | GLU | 2.6 |
| 8 | AH | 99 | GLU | 2.6 |
| 31 | BA | 1546 | C | 2.6 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 31 | DA | 1532 | C | 2.6 |
| 34 | BE | 200 | GLU | 2.6 |
| 5 | AE | 134 | ALA | 2.6 |
| 7 | AG | 134 | ALA | 2.6 |
| 9 | AI | 111 | ARG | 2.6 |
| 11 | AK | 84 | VAL | 2.6 |
| 14 | CN | 4 | LYS | 2.6 |
| 16 | CP | 50 | LYS | 2.6 |
| 24 | B2 | 58 | ALA | 2.6 |
| 36 | BG | 38 | VAL | 2.6 |
| 47 | BV | 13 | ARG | 2.6 |
| 50 | BY | 72 | VAL | 2.6 |
| 26 | B4 | 2 | LYS | 2.6 |
| 33 | BD | 35 | LYS | 2.6 |
| 47 | BV | 74 | LYS | 2.6 |
| 36 | DG | 20 | ILE | 2.5 |
| 51 | BZ | 28 | MET | 2.5 |
| 2 | AB | 28 | PHE | 2.5 |
| 5 | CE | 45 | PHE | 2.5 |
| 4 | CD | 12 | CYS | 2.5 |
| 27 | D5 | 36 | CYS | 2.5 |
| 38 | DI | 130 | TYR | 2.5 |
| 47 | BV | 91 | TYR | 2.5 |
| 7 | CG | 34 | GLY | 2.5 |
| 15 | CO | 62 | GLN | 2.5 |
| 30 | D8 | 61 | LEU | 2.5 |
| 1 | AA | 1174 | G | 2.5 |
| 1 | CA | 485 | G | 2.5 |
| 1 | CA | 1127 | G | 2.5 |
| 2 | CB | 36 | ARG | 2.5 |
| 4 | CD | 166 | LYS | 2.5 |
| 9 | CI | 42 | ARG | 2.5 |
| 12 | CL | 119 | LYS | 2.5 |
| 13 | AM | 88 | ARG | 2.5 |
| 31 | BA | 1051 | G | 2.5 |
| 31 | DA | 2893 | G | 2.5 |
| 34 | DE | 75 | VAL | 2.5 |
| 50 | BY | 85 | VAL | 2.5 |
| 1 | AA | 344 | A | 2.5 |
| 19 | CS | 24 | ALA | 2.5 |
| 31 | DA | 2062 | A | 2.5 |
| 32 | BB | 59 | A | 2.5 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 12 | CL | 100 | ILE | 2.5 |
| 37 | BH | 151 | ILE | 2.5 |
| 41 | DP | 67 | MET | 2.5 |
| 1 | CA | 1098 | C | 2.5 |
| 31 | DA | 2896 | C | 2.5 |
| 27 | D5 | 4 | HIS | 2.5 |
| 1 | CA | 90 | U | 2.5 |
| 3 | AC | 42 | LEU | 2.5 |
| 4 | AD | 135 | LEU | 2.5 |
| 7 | AG | 77 | SER | 2.5 |
| 5 | AE | 20 | GLN | 2.5 |
| 13 | CM | 85 | GLY | 2.5 |
| 19 | CS | 5 | LEU | 2.5 |
| 27 | B5 | 54 | GLY | 2.5 |
| 31 | BA | 877 | U | 2.5 |
| 36 | BG | 53 | LEU | 2.5 |
| 47 | BV | 73 | SER | 2.5 |
| 45 | BT | 84 | GLN | 2.5 |
| 7 | CG | 78 | ARG | 2.5 |
| 13 | AM | 104 | ARG | 2.5 |
| 23 | D1 | 39 | LYS | 2.5 |
| 42 | BQ | 18 | LYS | 2.5 |
| 43 | DR | 68 | ARG | 2.5 |
| 6 | AF | 91 | VAL | 2.5 |
| 36 | DG | 38 | VAL | 2.5 |
| 47 | BV | 57 | VAL | 2.5 |
| 51 | DZ | 165 | VAL | 2.5 |
| 3 | AC | 160 | ALA | 2.5 |
| 6 | CF | 83 | ASP | 2.5 |
| 26 | B4 | 12 | ALA | 2.5 |
| 34 | BE | 68 | ALA | 2.5 |
| 36 | DG | 137 | GLU | 2.5 |
| 37 | DH | 81 | GLU | 2.5 |
| 23 | D1 | 13 | ILE | 2.5 |
| 37 | BH | 121 | ILE | 2.5 |
| 10 | AJ | 42 | THR | 2.5 |
| 19 | CS | 33 | THR | 2.5 |
| 19 | CS | 79 | THR | 2.5 |
| 10 | CJ | 47 | PHE | 2.5 |
| 24 | B2 | 37 | PHE | 2.5 |
| 1 | AA | 983 | A | 2.5 |
| 31 | DA | 2629 | A | 2.5 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 1 | AA | 104 | G | 2.5 |
| 31 | BA | 2382 | G | 2.5 |
| 31 | BA | 2630 | G | 2.5 |
| 5 | CE | 60 | TYR | 2.5 |
| 16 | CP | 38 | TYR | 2.5 |
| 45 | BT | 24 | PRO | 2.5 |
| 13 | CM | 66 | LEU | 2.5 |
| 1 | AA | 578 | C | 2.5 |
| 6 | CF | 92 | LYS | 2.5 |
| 7 | AG | 102 | ARG | 2.5 |
| 9 | AI | 10 | ARG | 2.5 |
| 16 | CP | 75 | ARG | 2.5 |
| 17 | AQ | 37 | LYS | 2.5 |
| 19 | AS | 30 | LEU | 2.5 |
| 31 | BA | 2313 | C | 2.5 |
| 31 | DA | 1547 | C | 2.5 |
| 32 | BB | 30 | C | 2.5 |
| 37 | DH | 59 | ARG | 2.5 |
| 42 | DQ | 133 | ARG | 2.5 |
| 49 | DX | 64 | LYS | 2.5 |
| 2 | AB | 150 | SER | 2.5 |
| 2 | AB | 29 | ALA | 2.5 |
| 23 | D1 | 77 | ALA | 2.5 |
| 34 | BE | 131 | ALA | 2.5 |
| 1 | AA | 65 | U | 2.5 |
| 1 | CA | 961 | U | 2.5 |
| 31 | DA | 1175 | U | 2.5 |
| 7 | CG | 15 | ASP | 2.5 |
| 8 | CH | 35 | ILE | 2.5 |
| 9 | CI | 7 | THR | 2.5 |
| 9 | CI | 33 | PHE | 2.5 |
| 13 | AM | 105 | THR | 2.5 |
| 41 | DP | 30 | THR | 2.5 |
| 45 | BT | 45 | PHE | 2.5 |
| 3 | CC | 29 | TYR | 2.5 |
| 3 | CC | 33 | LEU | 2.5 |
| 5 | AE | 92 | LYS | 2.5 |
| 6 | AF | 43 | LEU | 2.5 |
| 7 | CG | 16 | LEU | 2.5 |
| 13 | CM | 87 | TYR | 2.5 |
| 15 | CO | 2 | PRO | 2.5 |
| 38 | DI | 111 | PRO | 2.5 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 10 | CJ | 16 | LEU | 2.5 |
| 23 | D1 | 32 | LYS | 2.5 |
| 25 | B3 | 7 | LYS | 2.5 |
| 36 | DG | 94 | LEU | 2.5 |
| 44 | BS | 13 | ARG | 2.5 |
| 45 | DT | 112 | ARG | 2.5 |
| 50 | BY | 21 | LYS | 2.5 |
| 1 | AA | 397 | A | 2.5 |
| 1 | CA | 949 | A | 2.5 |
| 2 | AB | 72 | GLY | 2.5 |
| 31 | BA | 2753 | A | 2.5 |
| 32 | DB | 59 | A | 2.5 |
| 35 | DF | 131 | GLY | 2.5 |
| 15 | AO | 27 | VAL | 2.5 |
| 26 | B4 | 10 | VAL | 2.5 |
| 1 | AA | 1133 | G | 2.5 |
| 1 | CA | 963 | G | 2.5 |
| 2 | CB | 233 | SER | 2.5 |
| 3 | CC | 168 | ALA | 2.5 |
| 11 | AK | 61 | ALA | 2.5 |
| 31 | DA | 2206 | G | 2.5 |
| 10 | CJ | 23 | ILE | 2.5 |
| 13 | CM | 84 | ILE | 2.5 |
| 37 | BH | 40 | GLU | 2.5 |
| 37 | BH | 92 | ILE | 2.5 |
| 1 | AA | 1069 | C | 2.5 |
| 32 | DB | 27 | C | 2.5 |
| 16 | CP | 9 | PHE | 2.5 |
| 37 | DH | 57 | ASP | 2.5 |
| 51 | BZ | 88 | PHE | 2.5 |
| 1 | AA | 1065 | U | 2.5 |
| 1 | AA | 1090 | U | 2.5 |
| 1 | CA | 1393 | U | 2.5 |
| 11 | AK | 42 | TRP | 2.5 |
| 14 | AN | 22 | THR | 2.5 |
| 2 | AB | 213 | LEU | 2.5 |
| 9 | CI | 10 | ARG | 2.5 |
| 30 | D8 | 29 | LYS | 2.5 |
| 36 | DG | 67 | LYS | 2.5 |
| 37 | BH | 42 | ARG | 2.5 |
| 42 | BQ | 59 | ARG | 2.5 |
| 48 | BW | 49 | LYS | 2.5 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 30 | D8 | 63 | PRO | 2.5 |
| 51 | DZ | 62 | PRO | 2.5 |
| 4 | CD | 41 | GLY | 2.5 |
| 22 | D0 | 6 | GLY | 2.5 |
| 42 | BQ | 19 | GLY | 2.5 |
| 7 | CG | 105 | VAL | 2.5 |
| 23 | B1 | 30 | VAL | 2.5 |
| 47 | DV | 72 | VAL | 2.5 |
| 11 | AK | 97 | ALA | 2.4 |
| 35 | DF | 5 | ALA | 2.4 |
| 46 | BU | 4 | ALA | 2.4 |
| 40 | DO | 87 | ILE | 2.4 |
| 1 | AA | 958 | A | 2.4 |
| 24 | B2 | 22 | GLU | 2.4 |
| 31 | DA | 1237 | A | 2.4 |
| 31 | DA | 1545 | A | 2.4 |
| 47 | BV | 43 | GLU | 2.4 |
| 2 | AB | 195 | ASP | 2.4 |
| 2 | CB | 220 | ASP | 2.4 |
| 4 | CD | 84 | LYS | 2.4 |
| 8 | CH | 116 | LYS | 2.4 |
| 9 | CI | 105 | ASP | 2.4 |
| 13 | CM | 79 | LYS | 2.4 |
| 20 | CT | 80 | ARG | 2.4 |
| 36 | DG | 74 | LYS | 2.4 |
| 39 | BN | 83 | LYS | 2.4 |
| 40 | DO | 113 | LYS | 2.4 |
| 50 | DY | 8 | LYS | 2.4 |
| 1 | AA | 194 | C | 2.4 |
| 1 | CA | 1363 | C | 2.4 |
| 31 | BA | 364 | C | 2.4 |
| 31 | BA | 366 | C | 2.4 |
| 31 | BA | 407 | G | 2.4 |
| 32 | DB | 3 | C | 2.4 |
| 32 | DB | 24 | G | 2.4 |
| 1 | AA | 1086 | U | 2.4 |
| 31 | DA | 1113 | U | 2.4 |
| 37 | DH | 158 | HIS | 2.4 |
| 9 | AI | 44 | VAL | 2.4 |
| 33 | DD | 235 | GLY | 2.4 |
| 36 | DG | 159 | VAL | 2.4 |
| 37 | BH | 141 | VAL | 2.4 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 41 | BP | 142 | GLY | 2.4 |
| 41 | DP | 71 | VAL | 2.4 |
| 49 | DX | 83 | VAL | 2.4 |
| 44 | BS | 55 | ALA | 2.4 |
| 2 | CB | 80 | ILE | 2.4 |
| 36 | BG | 168 | GLU | 2.4 |
| 38 | DI | 60 | GLU | 2.4 |
| 2 | AB | 217 | ARG | 2.4 |
| 4 | AD | 122 | ARG | 2.4 |
| 5 | CE | 25 | ARG | 2.4 |
| 6 | CF | 54 | LYS | 2.4 |
| 18 | AR | 68 | LYS | 2.4 |
| 21 | CU | 12 | LYS | 2.4 |
| 41 | BP | 76 | LYS | 2.4 |
| 45 | BT | 29 | ARG | 2.4 |
| 50 | BY | 73 | ARG | 2.4 |
| 50 | DY | 4 | LYS | 2.4 |
| 13 | AM | 116 | THR | 2.4 |
| 13 | CM | 63 | THR | 2.4 |
| 26 | D4 | 27 | THR | 2.4 |
| 51 | BZ | 75 | ASN | 2.4 |
| 47 | BV | 26 | ASP | 2.4 |
| 2 | CB | 26 | PRO | 2.4 |
| 12 | CL | 31 | PRO | 2.4 |
| 7 | AG | 66 | VAL | 2.4 |
| 9 | AI | 14 | VAL | 2.4 |
| 15 | CO | 50 | HIS | 2.4 |
| 33 | BD | 235 | GLY | 2.4 |
| 44 | BS | 34 | HIS | 2.4 |
| 51 | DZ | 56 | VAL | 2.4 |
| 3 | CC | 189 | ALA | 2.4 |
| 7 | AG | 100 | ALA | 2.4 |
| 1 | AA | 947 | G | 2.4 |
| 1 | CA | 1025 | U | 2.4 |
| 1 | CA | 1030(A) | G | 2.4 |
| 1 | CA | 1361 | G | 2.4 |
| 1 | CA | 1456 | G | 2.4 |
| 11 | CK | 32 | ILE | 2.4 |
| 31 | BA | 1880 | C | 2.4 |
| 32 | BB | 41 | U | 2.4 |
| 36 | BG | 155 | MET | 2.4 |
| 4 | AD | 139 | ARG | 2.4 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 16 | CP | 31 | LYS | 2.4 |
| 30 | B8 | 42 | ARG | 2.4 |
| 41 | BP | 149 | GLU | 2.4 |
| 49 | BX | 68 | ARG | 2.4 |
| 51 | DZ | 79 | ARG | 2.4 |
| 12 | CL | 116 | SER | 2.4 |
| 23 | B1 | 38 | SER | 2.4 |
| 3 | CC | 175 | LEU | 2.4 |
| 7 | AG | 109 | ASN | 2.4 |
| 7 | CG | 37 | ASN | 2.4 |
| 37 | BH | 126 | PRO | 2.4 |
| 38 | DI | 74 | ASN | 2.4 |
| 50 | BY | 53 | PRO | 2.4 |
| 34 | DE | 17 | ASP | 2.4 |
| 42 | BQ | 25 | ASP | 2.4 |
| 51 | BZ | 74 | VAL | 2.4 |
| 8 | AH | 128 | GLY | 2.4 |
| 9 | CI | 72 | GLY | 2.4 |
| 9 | CI | 117 | HIS | 2.4 |
| 16 | AP | 30 | GLY | 2.4 |
| 26 | B4 | 6 | HIS | 2.4 |
| 1 | AA | 974 | A | 2.4 |
| 2 | AB | 42 | ILE | 2.4 |
| 4 | AD | 204 | ILE | 2.4 |
| 29 | D7 | 45 | ALA | 2.4 |
| 31 | BA | 1494 | A | 2.4 |
| 31 | DA | 911 | A | 2.4 |
| 36 | BG | 77 | ILE | 2.4 |
| 2 | CB | 83 | MET | 2.4 |
| 3 | CC | 156 | ARG | 2.4 |
| 8 | AH | 18 | ARG | 2.4 |
| 13 | AM | 65 | LYS | 2.4 |
| 27 | B5 | 37 | LYS | 2.4 |
| 38 | DI | 112 | LYS | 2.4 |
| 17 | AQ | 71 | PHE | 2.4 |
| 51 | DZ | 89 | PHE | 2.4 |
| 1 | AA | 71 | C | 2.4 |
| 1 | AA | 381 | C | 2.4 |
| 1 | AA | 972 | C | 2.4 |
| 1 | AA | 982 | U | 2.4 |
| 1 | AA | 1030 | C | 2.4 |
| 4 | AD | 24 | GLU | 2.4 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 16 | AP | 6 | LEU | 2.4 |
| 24 | B2 | 24 | LEU | 2.4 |
| 31 | BA | 543 | C | 2.4 |
| 31 | BA | 1468 | C | 2.4 |
| 31 | BA | 1962 | C | 2.4 |
| 31 | BA | 2799 | C | 2.4 |
| 32 | BB | 1 | U | 2.4 |
| 37 | BH | 119 | GLU | 2.4 |
| 38 | BI | 58 | LEU | 2.4 |
| 40 | DO | 25 | LEU | 2.4 |
| 44 | DS | 58 | LEU | 2.4 |
| 33 | DD | 12 | SER | 2.4 |
| 31 | BA | 274 | G | 2.4 |
| 31 | BA | 1529 | G | 2.4 |
| 11 | AK | 114 | VAL | 2.4 |
| 47 | DV | 14 | VAL | 2.4 |
| 5 | CE | 85 | GLY | 2.4 |
| 7 | AG | 15 | ASP | 2.4 |
| 49 | DX | 58 | HIS | 2.4 |
| 51 | DZ | 121 | HIS | 2.4 |
| 8 | AH | 109 | ILE | 2.4 |
| 10 | AJ | 75 | ILE | 2.4 |
| 12 | AL | 70 | ILE | 2.4 |
| 19 | CS | 31 | ILE | 2.4 |
| 30 | B8 | 51 | ALA | 2.4 |
| 44 | DS | 102 | ALA | 2.4 |
| 47 | DV | 99 | ILE | 2.4 |
| 51 | DZ | 53 | ILE | 2.4 |
| 6 | AF | 92 | LYS | 2.4 |
| 12 | CL | 114 | LYS | 2.4 |
| 17 | AQ | 87 | LYS | 2.4 |
| 21 | AU | 25 | LYS | 2.4 |
| 36 | DG | 84 | LYS | 2.4 |
| 39 | BN | 137 | LYS | 2.4 |
| 9 | CI | 128 | ARG | 2.4 |
| 20 | AT | 85 | MET | 2.4 |
| 36 | BG | 141 | PHE | 2.4 |
| 19 | AS | 20 | LEU | 2.4 |
| 20 | CT | 99 | LEU | 2.4 |
| 35 | DF | 33 | LEU | 2.4 |
| 38 | DI | 140 | LEU | 2.4 |
| 47 | BV | 95 | LEU | 2.4 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 50 | DY | 31 | LEU | 2.4 |
| 1 | AA | 1092 | A | 2.4 |
| 1 | CA | 1280 | A | 2.4 |
| 1 | CA | 1493 | A | 2.4 |
| 2 | CB | 170 | GLU | 2.4 |
| 5 | CE | 50 | GLU | 2.4 |
| 31 | DA | 2268 | A | 2.4 |
| 32 | DB | 26 | A | 2.4 |
| 50 | BY | 62 | GLU | 2.4 |
| 50 | DY | 91 | GLU | 2.4 |
| 19 | CS | 35 | SER | 2.4 |
| 20 | AT | 11 | SER | 2.4 |
| 44 | DS | 31 | SER | 2.4 |
| 1 | CA | 831 | U | 2.4 |
| 6 | CF | 88 | VAL | 2.4 |
| 8 | CH | 26 | VAL | 2.4 |
| 31 | BA | 1497 | U | 2.4 |
| 1 | AA | 386 | C | 2.3 |
| 1 | AA | 624 | C | 2.3 |
| 1 | AA | 1060 | C | 2.3 |
| 1 | AA | 1140 | C | 2.3 |
| 1 | CA | 924 | C | 2.3 |
| 8 | CH | 3 | THR | 2.3 |
| 9 | AI | 64 | THR | 2.3 |
| 13 | CM | 20 | THR | 2.3 |
| 16 | AP | 69 | THR | 2.3 |
| 31 | BA | 2474 | C | 2.3 |
| 33 | BD | 40 | THR | 2.3 |
| 41 | BP | 118 | GLY | 2.3 |
| 45 | BT | 27 | THR | 2.3 |
| 4 | CD | 70 | ILE | 2.3 |
| 10 | CJ | 26 | ALA | 2.3 |
| 11 | AK | 94 | ALA | 2.3 |
| 11 | CK | 11 | LYS | 2.3 |
| 15 | AO | 48 | LYS | 2.3 |
| 24 | B2 | 15 | LYS | 2.3 |
| 37 | BH | 111 | HIS | 2.3 |
| 21 | AU | 9 | ARG | 2.3 |
| 23 | D1 | 52 | ARG | 2.3 |
| 38 | DI | 109 | ILE | 2.3 |
| 51 | BZ | 133 | ILE | 2.3 |
| 36 | DG | 150 | ASP | 2.3 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 2 | CB | 48 | MET | 2.3 |
| 38 | DI | 1 | MET | 2.3 |
| 1 | CA | 1031 | G | 2.3 |
| 31 | BA | 281 | G | 2.3 |
| 31 | BA | 1910 | G | 2.3 |
| 31 | DA | 656 | G | 2.3 |
| 32 | DB | 61 | G | 2.3 |
| 2 | CB | 70 | PHE | 2.3 |
| 2 | CB | 138 | LEU | 2.3 |
| 4 | AD | 120 | LEU | 2.3 |
| 6 | CF | 60 | PHE | 2.3 |
| 9 | CI | 3 | GLN | 2.3 |
| 33 | DD | 33 | LEU | 2.3 |
| 36 | DG | 173 | LEU | 2.3 |
| 44 | BS | 38 | GLN | 2.3 |
| 20 | CT | 60 | GLU | 2.3 |
| 36 | BG | 54 | GLU | 2.3 |
| 38 | DI | 73 | GLU | 2.3 |
| 36 | DG | 32 | PRO | 2.3 |
| 1 | AA | 1288 | A | 2.3 |
| 30 | D8 | 52 | LYS | 2.3 |
| 31 | DA | 603 | A | 2.3 |
| 32 | DB | 53 | A | 2.3 |
| 44 | BS | 53 | SER | 2.3 |
| 7 | AG | 60 | LYS | 2.3 |
| 11 | AK | 122 | LYS | 2.3 |
| 20 | AT | 39 | LYS | 2.3 |
| 2 | AB | 80 | ILE | 2.3 |
| 5 | CE | 131 | ILE | 2.3 |
| 10 | AJ | 66 | ARG | 2.3 |
| 11 | AK | 126 | ARG | 2.3 |
| 14 | AN | 12 | ARG | 2.3 |
| 44 | BS | 63 | THR | 2.3 |
| 50 | BY | 86 | ARG | 2.3 |
| 50 | DY | 58 | GLY | 2.3 |
| 34 | BE | 204 | ALA | 2.3 |
| 34 | DE | 159 | HIS | 2.3 |
| 1 | AA | 164 | U | 2.3 |
| 2 | CB | 101 | MET | 2.3 |
| 37 | BH | 74 | ASN | 2.3 |
| 46 | DU | 76 | TYR | 2.3 |
| 1 | AA | 1243 | C | 2.3 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 31 | BA | 1957 | C | 2.3 |
| 31 | DA | 280 | C | 2.3 |
| 31 | DA | 1958 | C | 2.3 |
| 33 | DD | 66 | ASP | 2.3 |
| 2 | AB | 138 | LEU | 2.3 |
| 18 | AR | 31 | LEU | 2.3 |
| 34 | DE | 51 | PHE | 2.3 |
| 37 | BH | 103 | LEU | 2.3 |
| 38 | BI | 140 | LEU | 2.3 |
| 36 | DG | 66 | GLN | 2.3 |
| 35 | DF | 56 | GLU | 2.3 |
| 41 | DP | 144 | GLU | 2.3 |
| 1 | AA | 963 | G | 2.3 |
| 1 | AA | 1036 | G | 2.3 |
| 1 | AA | 1131 | G | 2.3 |
| 1 | CA | 306 | G | 2.3 |
| 1 | CA | 570 | G | 2.3 |
| 1 | CA | 855 | G | 2.3 |
| 4 | CD | 105 | VAL | 2.3 |
| 9 | AI | 28 | VAL | 2.3 |
| 10 | CJ | 44 | VAL | 2.3 |
| 17 | CQ | 5 | VAL | 2.3 |
| 31 | BA | 920 | G | 2.3 |
| 31 | DA | 1051 | G | 2.3 |
| 45 | BT | 34 | VAL | 2.3 |
| 51 | BZ | 175 | VAL | 2.3 |
| 41 | BP | 64 | LYS | 2.3 |
| 44 | BS | 19 | LYS | 2.3 |
| 51 | BZ | 14 | LYS | 2.3 |
| 4 | AD | 39 | PRO | 2.3 |
| 7 | AG | 112 | PRO | 2.3 |
| 8 | CH | 30 | ARG | 2.3 |
| 12 | AL | 113 | ARG | 2.3 |
| 27 | D5 | 19 | ARG | 2.3 |
| 4 | AD | 152 | SER | 2.3 |
| 11 | CK | 52 | GLY | 2.3 |
| 20 | AT | 101 | GLY | 2.3 |
| 5 | CE | 120 | THR | 2.3 |
| 10 | CJ | 27 | ALA | 2.3 |
| 15 | CO | 4 | THR | 2.3 |
| 51 | BZ | 113 | ALA | 2.3 |
| 51 | BZ | 153 | SER | 2.3 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 4 | CD | 181 | MET | 2.3 |
| 50 | DY | 5 | MET | 2.3 |
| 16 | AP | 39 | TYR | 2.3 |
| 19 | AS | 80 | TYR | 2.3 |
| 26 | D4 | 18 | CYS | 2.3 |
| 37 | BH | 83 | TYR | 2.3 |
| 3 | CC | 94 | LEU | 2.3 |
| 13 | AM | 34 | LEU | 2.3 |
| 38 | BI | 9 | LEU | 2.3 |
| 1 | AA | 1285 | A | 2.3 |
| 2 | CB | 193 | ASP | 2.3 |
| 48 | DW | 94 | ASP | 2.3 |
| 51 | BZ | 63 | ASP | 2.3 |
| 31 | DA | 12 | U | 2.3 |
| 36 | BG | 137 | GLU | 2.3 |
| 50 | BY | 64 | GLU | 2.3 |
| 1 | AA | 103 | C | 2.3 |
| 1 | CA | 194 | C | 2.3 |
| 1 | CA | 805 | C | 2.3 |
| 3 | CC | 45 | LYS | 2.3 |
| 5 | AE | 69 | VAL | 2.3 |
| 8 | AH | 129 | VAL | 2.3 |
| 28 | D6 | 52 | VAL | 2.3 |
| 30 | D8 | 59 | LYS | 2.3 |
| 31 | DA | 1957 | C | 2.3 |
| 44 | BS | 65 | VAL | 2.3 |
| 44 | DS | 69 | VAL | 2.3 |
| 47 | DV | 52 | VAL | 2.3 |
| 4 | AD | 36 | ARG | 2.3 |
| 7 | CG | 4 | ARG | 2.3 |
| 17 | CQ | 92 | ARG | 2.3 |
| 33 | BD | 176 | ARG | 2.3 |
| 41 | DP | 16 | ARG | 2.3 |
| 2 | AB | 202 | PRO | 2.3 |
| 2 | CB | 234 | PRO | 2.3 |
| 11 | CK | 48 | ILE | 2.3 |
| 13 | CM | 4 | ILE | 2.3 |
| 15 | CO | 3 | ILE | 2.3 |
| 16 | CP | 15 | PRO | 2.3 |
| 41 | DP | 23 | PRO | 2.3 |
| 22 | B0 | 54 | GLY | 2.3 |
| 50 | DY | 96 | ILE | 2.3 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 51 | DZ | 171 | ILE | 2.3 |
| 4 | AD | 117 | ALA | 2.3 |
| 33 | DD | 240 | ALA | 2.3 |
| 36 | BG | 162 | THR | 2.3 |
| 22 | D0 | 3 | HIS | 2.3 |
| 37 | DH | 111 | HIS | 2.3 |
| 44 | DS | 101 | LEU | 2.3 |
| 1 | AA | 1233 | G | 2.3 |
| 1 | CA | 1347 | G | 2.3 |
| 16 | CP | 17 | TYR | 2.3 |
| 31 | BA | 352 | G | 2.3 |
| 31 | DA | 281 | G | 2.3 |
| 31 | DA | 2793 | G | 2.3 |
| 36 | DG | 146 | TYR | 2.3 |
| 8 | CH | 31 | PHE | 2.3 |
| 49 | BX | 41 | ASN | 2.3 |
| 16 | CP | 65 | GLN | 2.3 |
| 17 | AQ | 93 | GLN | 2.3 |
| 33 | BD | 171 | ASP | 2.3 |
| 7 | AG | 70 | LYS | 2.3 |
| 15 | AO | 8 | LYS | 2.3 |
| 22 | D0 | 4 | LYS | 2.3 |
| 41 | BP | 137 | LYS | 2.3 |
| 1 | AA | 946 | A | 2.3 |
| 1 | AA | 1350 | A | 2.3 |
| 2 | AB | 96 | ARG | 2.3 |
| 3 | CC | 44 | GLU | 2.3 |
| 7 | CG | 17 | VAL | 2.3 |
| 9 | AI | 51 | ARG | 2.3 |
| 10 | AJ | 61 | GLU | 2.3 |
| 11 | AK | 91 | ARG | 2.3 |
| 17 | AQ | 70 | ARG | 2.3 |
| 19 | AS | 19 | VAL | 2.3 |
| 31 | DA | 547 | A | 2.3 |
| 47 | BV | 5 | VAL | 2.3 |
| 47 | BV | 72 | VAL | 2.3 |
| 1 | AA | 5 | U | 2.3 |
| 33 | DD | 105 | ILE | 2.3 |
| 45 | DT | 42 | ILE | 2.3 |
| 47 | BV | 90 | PRO | 2.3 |
| 3 | AC | 168 | ALA | 2.3 |
| 5 | CE | 86 | ALA | 2.3 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 10 | AJ | 20 | ALA | 2.3 |
| 16 | AP | 70 | ALA | 2.3 |
| 19 | CS | 26 | GLY | 2.3 |
| 23 | B1 | 77 | ALA | 2.3 |
| 41 | DP | 44 | GLY | 2.3 |
| 31 | BA | 1577 | C | 2.3 |
| 31 | DA | 2666 | C | 2.3 |
| 32 | DB | 28 | C | 2.3 |
| 40 | DO | 1 | MET | 2.3 |
| 2 | AB | 238 | LEU | 2.3 |
| 2 | CB | 61 | LEU | 2.3 |
| 13 | CM | 90 | LEU | 2.3 |
| 26 | D4 | 6 | HIS | 2.3 |
| 36 | BG | 165 | THR | 2.3 |
| 38 | DI | 93 | THR | 2.3 |
| 44 | DS | 48 | LEU | 2.3 |
| 47 | BV | 49 | THR | 2.3 |
| 49 | DX | 31 | HIS | 2.3 |
| 5 | AE | 45 | PHE | 2.3 |
| 5 | AE | 84 | PHE | 2.3 |
| 4 | AD | 22 | LYS | 2.3 |
| 12 | AL | 23 | LYS | 2.3 |
| 20 | AT | 74 | LYS | 2.3 |
| 41 | DP | 76 | LYS | 2.3 |
| 50 | BY | 19 | LYS | 2.3 |
| 2 | CB | 175 | ARG | 2.3 |
| 6 | CF | 86 | ARG | 2.3 |
| 29 | B7 | 41 | ARG | 2.3 |
| 35 | BF | 18 | ARG | 2.3 |
| 5 | AE | 90 | VAL | 2.3 |
| 8 | AH | 118 | VAL | 2.3 |
| 8 | CH | 121 | ASP | 2.3 |
| 8 | CH | 129 | VAL | 2.3 |
| 24 | D2 | 52 | ASP | 2.3 |
| 51 | DZ | 63 | ASP | 2.3 |
| 7 | AG | 74 | GLU | 2.2 |
| 23 | D1 | 75 | GLU | 2.2 |
| 34 | DE | 163 | GLU | 2.2 |
| 36 | DG | 18 | GLU | 2.2 |
| 1 | AA | 144 | G | 2.2 |
| 1 | AA | 347 | G | 2.2 |
| 1 | AA | 384 | G | 2.2 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 31 | BA | 892 | G | 2.2 |
| 3 | CC | 171 | GLY | 2.2 |
| 19 | CS | 59 | PRO | 2.2 |
| 6 | AF | 67 | MET | 2.2 |
| 33 | BD | 240 | ALA | 2.2 |
| 35 | BF | 130 | ALA | 2.2 |
| 40 | BO | 11 | ALA | 2.2 |
| 41 | BP | 43 | GLY | 2.2 |
| 49 | DX | 67 | GLY | 2.2 |
| 1 | AA | 1000 | U | 2.2 |
| 1 | AA | 1248 | A | 2.2 |
| 2 | AB | 215 | LEU | 2.2 |
| 4 | CD | 96 | LEU | 2.2 |
| 18 | AR | 44 | LEU | 2.2 |
| 18 | AR | 78 | LEU | 2.2 |
| 20 | AT | 99 | LEU | 2.2 |
| 31 | BA | 1578 | U | 2.2 |
| 31 | DA | 2476 | A | 2.2 |
| 51 | BZ | 33 | LEU | 2.2 |
| 3 | CC | 167 | TRP | 2.2 |
| 2 | CB | 140 | HIS | 2.2 |
| 5 | CE | 75 | THR | 2.2 |
| 19 | AS | 33 | THR | 2.2 |
| 28 | D6 | 49 | HIS | 2.2 |
| 4 | AD | 86 | LYS | 2.2 |
| 17 | AQ | 40 | LYS | 2.2 |
| 18 | CR | 34 | TYR | 2.2 |
| 39 | DN | 2 | LYS | 2.2 |
| 1 | AA | 187 | C | 2.2 |
| 1 | CA | 1028 | C | 2.2 |
| 12 | CL | 53 | ARG | 2.2 |
| 14 | AN | 19 | ARG | 2.2 |
| 31 | DA | 1530 | C | 2.2 |
| 31 | DA | 2791 | C | 2.2 |
| 38 | BI | 61 | ARG | 2.2 |
| 42 | BQ | 56 | ARG | 2.2 |
| 12 | CL | 83 | VAL | 2.2 |
| 24 | D2 | 47 | ASN | 2.2 |
| 39 | DN | 8 | GLN | 2.2 |
| 50 | DY | 99 | CYS | 2.2 |
| 2 | AB | 134 | GLU | 2.2 |
| 4 | AD | 150 | GLU | 2.2 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 16 | AP | 40 | ASP | 2.2 |
| 49 | DX | 93 | GLU | 2.2 |
| 13 | AM | 39 | ILE | 2.2 |
| 20 | CT | 100 | ILE | 2.2 |
| 3 | CC | 34 | LEU | 2.2 |
| 7 | AG | 16 | LEU | 2.2 |
| 7 | CG | 99 | LEU | 2.2 |
| 13 | CM | 2 | ALA | 2.2 |
| 19 | CS | 68 | GLY | 2.2 |
| 21 | AU | 11 | GLY | 2.2 |
| 36 | BG | 68 | PRO | 2.2 |
| 38 | DI | 36 | ALA | 2.2 |
| 50 | BY | 56 | PRO | 2.2 |
| 24 | B2 | 35 | LEU | 2.2 |
| 27 | B5 | 58 | LEU | 2.2 |
| 44 | BS | 73 | LEU | 2.2 |
| 50 | BY | 100 | ALA | 2.2 |
| 10 | AJ | 54 | PHE | 2.2 |
| 1 | CA | 541 | G | 2.2 |
| 1 | CA | 945 | G | 2.2 |
| 1 | CA | 951 | G | 2.2 |
| 1 | CA | 1182 | G | 2.2 |
| 1 | CA | 1355 | G | 2.2 |
| 3 | CC | 4 | LYS | 2.2 |
| 4 | CD | 85 | LYS | 2.2 |
| 12 | AL | 122 | THR | 2.2 |
| 16 | AP | 35 | LYS | 2.2 |
| 34 | DE | 132 | HIS | 2.2 |
| 36 | BG | 161 | THR | 2.2 |
| 37 | DH | 85 | LYS | 2.2 |
| 51 | DZ | 170 | THR | 2.2 |
| 1 | CA | 697 | U | 2.2 |
| 9 | CI | 83 | ARG | 2.2 |
| 9 | CI | 121 | ARG | 2.2 |
| 10 | CJ | 66 | ARG | 2.2 |
| 12 | AL | 98 | TYR | 2.2 |
| 28 | D6 | 21 | TYR | 2.2 |
| 31 | DA | 158 | U | 2.2 |
| 1 | AA | 959 | A | 2.2 |
| 1 | AA | 1236 | A | 2.2 |
| 31 | BA | 2062 | A | 2.2 |
| 31 | BA | 2660 | A | 2.2 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 29 | D7 | 46 | VAL | 2.2 |
| 37 | BH | 113 | VAL | 2.2 |
| 47 | BV | 52 | VAL | 2.2 |
| 2 | AB | 212 | GLN | 2.2 |
| 6 | CF | 94 | GLN | 2.2 |
| 36 | BG | 66 | GLN | 2.2 |
| 11 | AK | 48 | ILE | 2.2 |
| 28 | B6 | 51 | GLU | 2.2 |
| 36 | BG | 164 | GLU | 2.2 |
| 39 | BN | 88 | GLU | 2.2 |
| 40 | DO | 92 | GLU | 2.2 |
| 49 | BX | 89 | ILE | 2.2 |
| 8 | AH | 117 | GLY | 2.2 |
| 10 | CJ | 41 | PRO | 2.2 |
| 14 | AN | 39 | LEU | 2.2 |
| 36 | DG | 89 | GLY | 2.2 |
| 36 | DG | 175 | LEU | 2.2 |
| 48 | BW | 63 | ASP | 2.2 |
| 41 | BP | 54 | GLY | 2.2 |
| 51 | BZ | 12 | GLY | 2.2 |
| 11 | CK | 71 | LYS | 2.2 |
| 14 | AN | 58 | LYS | 2.2 |
| 39 | BN | 37 | LYS | 2.2 |
| 44 | BS | 87 | PHE | 2.2 |
| 49 | BX | 28 | PHE | 2.2 |
| 13 | CM | 108 | ARG | 2.2 |
| 19 | AS | 81 | ARG | 2.2 |
| 37 | BH | 51 | ARG | 2.2 |
| 11 | AK | 57 | THR | 2.2 |
| 38 | BI | 108 | THR | 2.2 |
| 19 | AS | 58 | VAL | 2.2 |
| 33 | DD | 34 | VAL | 2.2 |
| 43 | BR | 6 | SER | 2.2 |
| 31 | DA | 271(L) | U | 2.2 |
| 31 | DA | 860 | U | 2.2 |
| 10 | CJ | 4 | ILE | 2.2 |
| 1 | AA | 93 | G | 2.2 |
| 1 | AA | 941 | G | 2.2 |
| 1 | AA | 1003 | G | 2.2 |
| 1 | AA | 1030(A) | G | 2.2 |
| 1 | CA | 428 | G | 2.2 |
| 1 | CA | 509 | A | 2.2 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 1 | CA | 535 | A | 2.2 |
| 1 | CA | 1206 | G | 2.2 |
| 31 | BA | 2893 | G | 2.2 |
| 31 | DA | 1238 | G | 2.2 |
| 32 | BB | 99 | G | 2.2 |
| 26 | D4 | 9 | LEU | 2.2 |
| 30 | B8 | 62 | LEU | 2.2 |
| 31 | BA | 196 | A | 2.2 |
| 31 | BA | 271(M) | G | 2.2 |
| 31 | BA | 338 | G | 2.2 |
| 31 | BA | 646 | A | 2.2 |
| 31 | BA | 862 | G | 2.2 |
| 40 | DO | 88 | ASN | 2.2 |
| 31 | BA | 1508 | A | 2.2 |
| 47 | DV | 23 | GLU | 2.2 |
| 9 | CI | 13 | ALA | 2.2 |
| 16 | CP | 7 | ALA | 2.2 |
| 20 | CT | 69 | GLY | 2.2 |
| 34 | DE | 106 | GLY | 2.2 |
| 35 | DF | 167 | ALA | 2.2 |
| 37 | BH | 108 | GLY | 2.2 |
| 37 | DH | 120 | GLY | 2.2 |
| 38 | BI | 134 | PRO | 2.2 |
| 44 | BS | 105 | ALA | 2.2 |
| 44 | DS | 22 | GLY | 2.2 |
| 2 | CB | 179 | LYS | 2.2 |
| 7 | CG | 137 | LYS | 2.2 |
| 16 | AP | 47 | ASP | 2.2 |
| 33 | BD | 20 | ASP | 2.2 |
| 33 | BD | 200 | ASP | 2.2 |
| 6 | AF | 97 | PHE | 2.2 |
| 11 | CK | 12 | ARG | 2.2 |
| 42 | BQ | 82 | ARG | 2.2 |
| 42 | DQ | 56 | ARG | 2.2 |
| 43 | BR | 17 | ARG | 2.2 |
| 1 | CA | 536 | C | 2.2 |
| 31 | BA | 1467 | C | 2.2 |
| 31 | BA | 1547 | C | 2.2 |
| 31 | BA | 2306 | C | 2.2 |
| 3 | CC | 165 | THR | 2.2 |
| 10 | CJ | 42 | THR | 2.2 |
| 22 | B0 | 10 | THR | 2.2 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 3 | AC | 151 | VAL | 2.2 |
| 5 | AE | 133 | TYR | 2.2 |
| 8 | CH | 53 | VAL | 2.2 |
| 13 | AM | 60 | VAL | 2.2 |
| 9 | CI | 63 | ILE | 2.2 |
| 23 | D1 | 42 | GLN | 2.2 |
| 30 | B8 | 41 | ILE | 2.2 |
| 36 | BG | 78 | SER | 2.2 |
| 36 | BG | 157 | ILE | 2.2 |
| 36 | DG | 52 | ILE | 2.2 |
| 37 | BH | 155 | SER | 2.2 |
| 3 | CC | 42 | LEU | 2.2 |
| 11 | AK | 98 | LEU | 2.2 |
| 36 | DG | 3 | LEU | 2.2 |
| 36 | DG | 176 | LEU | 2.2 |
| 2 | AB | 170 | GLU | 2.2 |
| 2 | CB | 134 | GLU | 2.2 |
| 3 | AC | 169 | ALA | 2.2 |
| 7 | CG | 28 | ASN | 2.2 |
| 9 | CI | 84 | ALA | 2.2 |
| 9 | CI | 110 | GLU | 2.2 |
| 1 | CA | 1364 | U | 2.2 |
| 9 | CI | 21 | PRO | 2.2 |
| 12 | AL | 48 | PRO | 2.2 |
| 12 | AL | 94 | PRO | 2.2 |
| 12 | AL | 114 | LYS | 2.2 |
| 13 | AM | 41 | PRO | 2.2 |
| 13 | CM | 6 | GLY | 2.2 |
| 13 | CM | 13 | LYS | 2.2 |
| 17 | AQ | 67 | LYS | 2.2 |
| 20 | CT | 106 | ALA | 2.2 |
| 23 | D1 | 57 | GLU | 2.2 |
| 35 | BF | 133 | ASN | 2.2 |
| 35 | DF | 166 | ALA | 2.2 |
| 36 | BG | 69 | ALA | 2.2 |
| 37 | DH | 127 | GLU | 2.2 |
| 20 | CT | 103 | GLY | 2.2 |
| 22 | D0 | 5 | LYS | 2.2 |
| 24 | B2 | 28 | LYS | 2.2 |
| 43 | DR | 5 | LYS | 2.2 |
| 31 | BA | 272(I) | U | 2.2 |
| 47 | BV | 101 | GLY | 2.2 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 43 | DR | 57 | ARG | 2.2 |
| 3 | AC | 10 | PHE | 2.2 |
| 51 | DZ | 77 | ASP | 2.2 |
| 31 | BA | 900 | A | 2.2 |
| 31 | DA | 896 | A | 2.2 |
| 1 | AA | 44 | G | 2.2 |
| 1 | AA | 530 | G | 2.2 |
| 1 | AA | 711 | G | 2.2 |
| 1 | AA | 727 | G | 2.2 |
| 31 | BA | 1047 | G | 2.2 |
| 31 | DA | 1816 | G | 2.2 |
| 31 | DA | 2100 | G | 2.2 |
| 32 | DB | 118 | G | 2.2 |
| 39 | BN | 42 | TRP | 2.2 |
| 10 | CJ | 56 | HIS | 2.2 |
| 16 | AP | 16 | HIS | 2.2 |
| 6 | AF | 6 | VAL | 2.1 |
| 13 | AM | 109 | THR | 2.1 |
| 13 | CM | 109 | THR | 2.1 |
| 35 | DF | 175 | THR | 2.1 |
| 9 | AI | 125 | TYR | 2.1 |
| 12 | CL | 64 | TYR | 2.1 |
| 51 | DZ | 9 | TYR | 2.1 |
| 1 | AA | 1141 | C | 2.1 |
| 24 | D2 | 57 | ILE | 2.1 |
| 31 | DA | 271(O) | C | 2.1 |
| 36 | DG | 63 | ILE | 2.1 |
| 2 | CB | 44 | LEU | 2.1 |
| 44 | BS | 84 | GLN | 2.1 |
| 5 | AE | 136 | MET | 2.1 |
| 5 | CE | 19 | MET | 2.1 |
| 20 | CT | 29 | LYS | 2.1 |
| 26 | B4 | 28 | LYS | 2.1 |
| 30 | B8 | 11 | LYS | 2.1 |
| 36 | DG | 158 | ALA | 2.1 |
| 37 | BH | 84 | SER | 2.1 |
| 48 | DW | 110 | LYS | 2.1 |
| 5 | AE | 21 | ALA | 2.1 |
| 38 | BI | 36 | ALA | 2.1 |
| 47 | DV | 3 | ALA | 2.1 |
| 50 | BY | 70 | SER | 2.1 |
| 10 | CJ | 97 | GLU | 2.1 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 4 | AD | 23 | GLY | 2.1 |
| 8 | CH | 130 | GLY | 2.1 |
| 9 | CI | 9 | ARG | 2.1 |
| 20 | CT | 23 | ARG | 2.1 |
| 25 | D3 | 29 | ARG | 2.1 |
| 11 | CK | 117 | ASN | 2.1 |
| 12 | AL | 25 | PRO | 2.1 |
| 44 | BS | 104 | GLY | 2.1 |
| 1 | CA | 1159 | U | 2.1 |
| 51 | BZ | 77 | ASP | 2.1 |
| 3 | CC | 55 | VAL | 2.1 |
| 7 | AG | 17 | VAL | 2.1 |
| 16 | AP | 59 | TRP | 2.1 |
| 30 | D8 | 31 | HIS | 2.1 |
| 35 | BF | 126 | VAL | 2.1 |
| 35 | DF | 11 | VAL | 2.1 |
| 37 | BH | 115 | VAL | 2.1 |
| 1 | CA | 533 | A | 2.1 |
| 2 | AB | 200 | ILE | 2.1 |
| 3 | CC | 23 | TYR | 2.1 |
| 10 | CJ | 38 | ILE | 2.1 |
| 13 | AM | 78 | ILE | 2.1 |
| 31 | BA | 1960 | A | 2.1 |
| 36 | BG | 20 | ILE | 2.1 |
| 2 | CB | 115 | LEU | 2.1 |
| 3 | CC | 147 | LYS | 2.1 |
| 18 | CR | 76 | LEU | 2.1 |
| 18 | CR | 88 | LYS | 2.1 |
| 20 | AT | 24 | LEU | 2.1 |
| 23 | B1 | 91 | LYS | 2.1 |
| 25 | B3 | 20 | LYS | 2.1 |
| 35 | DF | 137 | LYS | 2.1 |
| 39 | BN | 65 | LYS | 2.1 |
| 49 | BX | 78 | LYS | 2.1 |
| 50 | DY | 63 | LYS | 2.1 |
| 1 | AA | 392 | G | 2.1 |
| 31 | BA | 859 | G | 2.1 |
| 31 | BA | 2206 | G | 2.1 |
| 31 | DA | 614(B) | G | 2.1 |
| 32 | BB | 117 | G | 2.1 |
| 32 | DB | 63 | G | 2.1 |
| 35 | DF | 194 | MET | 2.1 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 7 | AG | 117 | ALA | 2.1 |
| 13 | CM | 88 | ARG | 2.1 |
| 37 | BH | 101 | ARG | 2.1 |
| 41 | BP | 81 | GLN | 2.1 |
| 42 | DQ | 113 | GLN | 2.1 |
| 45 | DT | 90 | GLN | 2.1 |
| 3 | AC | 194 | GLY | 2.1 |
| 6 | CF | 17 | SER | 2.1 |
| 1 | AA | 48 | C | 2.1 |
| 1 | CA | 48 | C | 2.1 |
| 1 | CA | 1103 | C | 2.1 |
| 12 | CL | 48 | PRO | 2.1 |
| 17 | CQ | 97 | SER | 2.1 |
| 18 | CR | 48 | GLY | 2.1 |
| 20 | AT | 98 | PRO | 2.1 |
| 28 | D6 | 41 | PRO | 2.1 |
| 36 | DG | 168 | GLU | 2.1 |
| 31 | DA | 543 | C | 2.1 |
| 37 | BH | 118 | PRO | 2.1 |
| 45 | DT | 12 | SER | 2.1 |
| 49 | BX | 32 | PRO | 2.1 |
| 50 | DY | 64 | GLU | 2.1 |
| 51 | BZ | 68 | PRO | 2.1 |
| 51 | BZ | 83 | PRO | 2.1 |
| 4 | AD | 161 | ASN | 2.1 |
| 14 | AN | 61 | TRP | 2.1 |
| 35 | BF | 172 | TRP | 2.1 |
| 42 | BQ | 102 | VAL | 2.1 |
| 45 | BT | 107 | ASP | 2.1 |
| 49 | BX | 30 | VAL | 2.1 |
| 19 | AS | 57 | HIS | 2.1 |
| 1 | AA | 223 | U | 2.1 |
| 1 | AA | 367 | U | 2.1 |
| 3 | AC | 134 | ILE | 2.1 |
| 4 | AD | 85 | LYS | 2.1 |
| 7 | AG | 120 | ILE | 2.1 |
| 8 | CH | 111 | ILE | 2.1 |
| 12 | AL | 21 | LYS | 2.1 |
| 14 | CN | 58 | LYS | 2.1 |
| 16 | CP | 22 | THR | 2.1 |
| 22 | D0 | 10 | THR | 2.1 |
| 31 | BA | 2473 | U | 2.1 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 35 | DF | 140 | LEU | 2.1 |
| 39 | BN | 4 | TYR | 2.1 |
| 39 | BN | 84 | LYS | 2.1 |
| 39 | DN | 118 | LYS | 2.1 |
| 41 | BP | 124 | LYS | 2.1 |
| 46 | DU | 80 | ILE | 2.1 |
| 50 | DY | 95 | LYS | 2.1 |
| 51 | DZ | 33 | LEU | 2.1 |
| 14 | CN | 3 | ARG | 2.1 |
| 20 | AT | 22 | ARG | 2.1 |
| 34 | DE | 61 | ARG | 2.1 |
| 38 | BI | 67 | ARG | 2.1 |
| 7 | CG | 106 | GLN | 2.1 |
| 47 | BV | 27 | ALA | 2.1 |
| 7 | AG | 146 | GLU | 2.1 |
| 11 | AK | 118 | GLY | 2.1 |
| 13 | AM | 69 | GLU | 2.1 |
| 31 | BA | 357 | A | 2.1 |
| 31 | DA | 548 | A | 2.1 |
| 33 | DD | 188 | GLU | 2.1 |
| 45 | DT | 109 | GLU | 2.1 |
| 51 | BZ | 13 | GLU | 2.1 |
| 3 | AC | 154 | SER | 2.1 |
| 3 | CC | 49 | SER | 2.1 |
| 20 | CT | 11 | SER | 2.1 |
| 4 | CD | 161 | ASN | 2.1 |
| 46 | BU | 94 | ASN | 2.1 |
| 1 | AA | 1353 | G | 2.1 |
| 1 | AA | 1419 | G | 2.1 |
| 1 | CA | 348 | G | 2.1 |
| 1 | CA | 1017 | G | 2.1 |
| 1 | CA | 1024 | G | 2.1 |
| 1 | CA | 1293 | G | 2.1 |
| 2 | AB | 7 | VAL | 2.1 |
| 2 | CB | 93 | VAL | 2.1 |
| 4 | AD | 198 | VAL | 2.1 |
| 19 | AS | 9 | VAL | 2.1 |
| 31 | DA | 1525 | G | 2.1 |
| 31 | DA | 1740 | G | 2.1 |
| 47 | BV | 61 | VAL | 2.1 |
| 1 | AA | 186 | C | 2.1 |
| 1 | AA | 355 | C | 2.1 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 1 | AA | 1109 | C | 2.1 |
| 1 | CA | 824 | C | 2.1 |
| 31 | BA | 2695 | C | 2.1 |
| 2 | CB | 22 | LYS | 2.1 |
| 20 | AT | 21 | LYS | 2.1 |
| 3 | CC | 182 | ILE | 2.1 |
| 4 | AD | 102 | ASP | 2.1 |
| 33 | DD | 5 | LYS | 2.1 |
| 35 | DF | 172 | TRP | 2.1 |
| 36 | BG | 84 | LYS | 2.1 |
| 38 | BI | 45 | LYS | 2.1 |
| 38 | DI | 101 | LEU | 2.1 |
| 38 | DI | 133 | HIS | 2.1 |
| 51 | DZ | 36 | LYS | 2.1 |
| 4 | CD | 162 | LEU | 2.1 |
| 14 | CN | 42 | ILE | 2.1 |
| 16 | CP | 74 | LEU | 2.1 |
| 39 | BN | 71 | ILE | 2.1 |
| 40 | BO | 122 | LEU | 2.1 |
| 45 | DT | 114 | LEU | 2.1 |
| 51 | DZ | 133 | ILE | 2.1 |
| 7 | AG | 76 | ARG | 2.1 |
| 9 | CI | 104 | ARG | 2.1 |
| 24 | B2 | 26 | ARG | 2.1 |
| 4 | CD | 165 | MET | 2.1 |
| 9 | CI | 88 | TYR | 2.1 |
| 10 | CJ | 67 | THR | 2.1 |
| 12 | AL | 120 | TYR | 2.1 |
| 12 | CL | 42 | THR | 2.1 |
| 35 | BF | 72 | ARG | 2.1 |
| 44 | DS | 13 | ARG | 2.1 |
| 17 | CQ | 32 | TYR | 2.1 |
| 2 | CB | 225 | ALA | 2.1 |
| 13 | CM | 18 | ALA | 2.1 |
| 24 | D2 | 13 | ALA | 2.1 |
| 42 | DQ | 20 | ALA | 2.1 |
| 44 | BS | 102 | ALA | 2.1 |
| 44 | DS | 37 | ALA | 2.1 |
| 1 | CA | 1235 | U | 2.1 |
| 31 | BA | 1113 | U | 2.1 |
| 41 | DP | 68 | GLN | 2.1 |
| 43 | DR | 91 | GLN | 2.1 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 45 | BT | 90 | GLN | 2.1 |
| 9 | AI | 69 | GLY | 2.1 |
| 33 | DD | 216 | GLY | 2.1 |
| 36 | BG | 44 | GLY | 2.1 |
| 9 | AI | 2 | GLU | 2.1 |
| 23 | B1 | 89 | GLU | 2.1 |
| 33 | BD | 30 | GLU | 2.1 |
| 37 | DH | 36 | PRO | 2.1 |
| 2 | AB | 210 | SER | 2.1 |
| 19 | CS | 51 | VAL | 2.1 |
| 38 | DI | 145 | VAL | 2.1 |
| 43 | DR | 73 | VAL | 2.1 |
| 1 | CA | 532 | A | 2.1 |
| 3 | AC | 150 | LYS | 2.1 |
| 4 | AD | 166 | LYS | 2.1 |
| 8 | CH | 15 | ASN | 2.1 |
| 9 | CI | 118 | LYS | 2.1 |
| 10 | AJ | 55 | LYS | 2.1 |
| 10 | AJ | 69 | ASN | 2.1 |
| 19 | AS | 53 | ASN | 2.1 |
| 31 | BA | 1912 | A | 2.1 |
| 32 | BB | 78 | A | 2.1 |
| 32 | DB | 57 | A | 2.1 |
| 34 | DE | 145 | LYS | 2.1 |
| 45 | BT | 65 | LYS | 2.1 |
| 5 | AE | 43 | LEU | 2.1 |
| 5 | AE | 91 | LEU | 2.1 |
| 11 | AK | 21 | ILE | 2.1 |
| 16 | CP | 25 | ARG | 2.1 |
| 36 | DG | 101 | ILE | 2.1 |
| 38 | BI | 109 | ILE | 2.1 |
| 39 | DN | 71 | ILE | 2.1 |
| 44 | BS | 26 | LEU | 2.1 |
| 44 | DS | 35 | ILE | 2.1 |
| 47 | BV | 94 | LEU | 2.1 |
| 49 | BX | 60 | ARG | 2.1 |
| 50 | BY | 96 | ILE | 2.1 |
| 50 | BY | 6 | HIS | 2.1 |
| 12 | CL | 112 | ASP | 2.1 |
| 39 | BN | 17 | ASP | 2.1 |
| 7 | AG | 85 | TYR | 2.1 |
| 38 | BI | 93 | THR | 2.1 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 1 | AA | 1344 | C | 2.1 |
| 1 | AA | 1034 | G | 2.1 |
| 1 | AA | 1222 | G | 2.1 |
| 1 | CA | 1117 | G | 2.1 |
| 9 | CI | 61 | ALA | 2.1 |
| 13 | AM | 30 | ALA | 2.1 |
| 31 | BA | 1920 | C | 2.1 |
| 31 | BA | 1958 | C | 2.1 |
| 31 | BA | 2664 | G | 2.1 |
| 32 | BB | 98 | G | 2.1 |
| 36 | BG | 171 | ALA | 2.1 |
| 2 | CB | 224 | GLN | 2.1 |
| 4 | CD | 45 | GLN | 2.1 |
| 15 | CO | 86 | GLY | 2.1 |
| 41 | DP | 109 | GLY | 2.1 |
| 50 | BY | 41 | GLY | 2.1 |
| 10 | CJ | 37 | PRO | 2.1 |
| 19 | AS | 59 | PRO | 2.1 |
| 28 | D6 | 35 | GLU | 2.1 |
| 33 | BD | 237 | GLU | 2.1 |
| 37 | BH | 167 | GLU | 2.1 |
| 38 | BI | 60 | GLU | 2.1 |
| 41 | BP | 63 | PRO | 2.1 |
| 1 | AA | 82 | U | 2.0 |
| 1 | AA | 950 | U | 2.0 |
| 8 | AH | 95 | VAL | 2.0 |
| 9 | CI | 53 | VAL | 2.0 |
| 27 | B5 | 60 | VAL | 2.0 |
| 31 | BA | 271(L) | U | 2.0 |
| 31 | BA | 1033 | U | 2.0 |
| 46 | BU | 90 | VAL | 2.0 |
| 50 | DY | 85 | VAL | 2.0 |
| 51 | BZ | 86 | VAL | 2.0 |
| 51 | DZ | 161 | VAL | 2.0 |
| 5 | AE | 87 | SER | 2.0 |
| 14 | CN | 32 | SER | 2.0 |
| 14 | CN | 60 | SER | 2.0 |
| 2 | CB | 58 | ILE | 2.0 |
| 2 | CB | 108 | ILE | 2.0 |
| 3 | AC | 43 | LEU | 2.0 |
| 5 | AE | 64 | ARG | 2.0 |
| 5 | AE | 129 | ILE | 2.0 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 6 | CF | 81 | ILE | 2.0 |
| 7 | CG | 76 | ARG | 2.0 |
| 17 | AQ | 92 | ARG | 2.0 |
| 30 | D8 | 32 | LEU | 2.0 |
| 33 | DD | 49 | ILE | 2.0 |
| 35 | DF | 206 | ILE | 2.0 |
| 36 | BG | 60 | LEU | 2.0 |
| 37 | DH | 149 | ARG | 2.0 |
| 38 | BI | 33 | ARG | 2.0 |
| 38 | DI | 128 | LEU | 2.0 |
| 51 | DZ | 75 | ASN | 2.0 |
| 19 | AS | 34 | TRP | 2.0 |
| 10 | AJ | 86 | MET | 2.0 |
| 15 | CO | 58 | MET | 2.0 |
| 19 | CS | 44 | MET | 2.0 |
| 1 | CA | 1035 | A | 2.0 |
| 6 | CF | 63 | TYR | 2.0 |
| 26 | B4 | 24 | THR | 2.0 |
| 31 | BA | 1913 | A | 2.0 |
| 34 | BE | 44 | TYR | 2.0 |
| 5 | CE | 56 | GLN | 2.0 |
| 10 | CJ | 10 | GLY | 2.0 |
| 12 | AL | 9 | GLN | 2.0 |
| 4 | CD | 37 | PRO | 2.0 |
| 7 | CG | 93 | PRO | 2.0 |
| 51 | BZ | 176 | PRO | 2.0 |
| 1 | AA | 990 | C | 2.0 |
| 1 | AA | 1028 | C | 2.0 |
| 1 | AA | 1116 | C | 2.0 |
| 1 | AA | 1260 | C | 2.0 |
| 1 | CA | 1132 | C | 2.0 |
| 1 | CA | 1234 | C | 2.0 |
| 1 | CA | 1389 | C | 2.0 |
| 20 | AT | 14 | LYS | 2.0 |
| 33 | BD | 255 | LYS | 2.0 |
| 46 | BU | 84 | LYS | 2.0 |
| 51 | BZ | 11 | GLU | 2.0 |
| 51 | DZ | 168 | GLU | 2.0 |
| 37 | BH | 26 | VAL | 2.0 |
| 51 | BZ | 90 | VAL | 2.0 |
| 1 | AA | 939 | G | 2.0 |
| 1 | AA | 1047 | G | 2.0 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 1 | CA | 79 | G | 2.0 |
| 1 | CA | 159 | G | 2.0 |
| 1 | CA | 324 | G | 2.0 |
| 1 | CA | 1184 | G | 2.0 |
| 1 | CA | 1233 | G | 2.0 |
| 1 | CA | 1370 | G | 2.0 |
| 2 | AB | 153 | ARG | 2.0 |
| 2 | CB | 157 | ARG | 2.0 |
| 3 | AC | 175 | LEU | 2.0 |
| 4 | CD | 114 | ARG | 2.0 |
| 5 | CE | 12 | LEU | 2.0 |
| 6 | CF | 21 | LEU | 2.0 |
| 9 | CI | 47 | LEU | 2.0 |
| 15 | CO | 72 | ARG | 2.0 |
| 16 | AP | 8 | ARG | 2.0 |
| 19 | CS | 29 | ARG | 2.0 |
| 23 | B1 | 94 | LEU | 2.0 |
| 31 | BA | 275 | G | 2.0 |
| 31 | BA | 326 | G | 2.0 |
| 31 | DA | 879 | G | 2.0 |
| 35 | DF | 72 | ARG | 2.0 |
| 36 | BG | 7 | LEU | 2.0 |
| 44 | DS | 32 | LEU | 2.0 |
| 45 | DT | 6 | LEU | 2.0 |
| 3 | CC | 5 | ILE | 2.0 |
| 9 | AI | 74 | ILE | 2.0 |
| 12 | AL | 116 | SER | 2.0 |
| 33 | BD | 12 | SER | 2.0 |
| 13 | CM | 106 | ASN | 2.0 |
| 31 | BA | 1420 | U | 2.0 |
| 34 | BE | 55 | ASN | 2.0 |
| 46 | BU | 75 | ASN | 2.0 |
| 13 | CM | 92 | HIS | 2.0 |
| 2 | AB | 188 | ALA | 2.0 |
| 10 | CJ | 18 | ALA | 2.0 |
| 7 | AG | 154 | TYR | 2.0 |
| 15 | AO | 4 | THR | 2.0 |
| 41 | BP | 58 | THR | 2.0 |
| 51 | BZ | 172 | ALA | 2.0 |
| 42 | BQ | 9 | TYR | 2.0 |
| 36 | BG | 49 | ASP | 2.0 |
| 36 | DG | 116 | ASP | 2.0 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 4 | AD | 185 | PHE | 2.0 |
| 19 | AS | 26 | GLY | 2.0 |
| 3 | AC | 199 | LYS | 2.0 |
| 4 | CD | 26 | CYS | 2.0 |
| 36 | DG | 129 | GLY | 2.0 |
| 38 | DI | 90 | GLY | 2.0 |
| 44 | DS | 96 | GLY | 2.0 |
| 4 | CD | 46 | LYS | 2.0 |
| 14 | CN | 15 | LYS | 2.0 |
| 27 | B5 | 32 | PRO | 2.0 |
| 30 | D8 | 35 | GLN | 2.0 |
| 35 | BF | 66 | PRO | 2.0 |
| 36 | BG | 8 | LYS | 2.0 |
| 36 | BG | 41 | GLN | 2.0 |
| 39 | BN | 2 | LYS | 2.0 |
| 40 | DO | 53 | LYS | 2.0 |
| 1 | AA | 1332 | A | 2.0 |
| 1 | CA | 356 | A | 2.0 |
| 1 | CA | 983 | A | 2.0 |
| 3 | CC | 103 | VAL | 2.0 |
| 10 | AJ | 64 | GLU | 2.0 |
| 4 | CD | 107 | ARG | 2.0 |
| 8 | AH | 53 | VAL | 2.0 |
| 12 | AL | 41 | ARG | 2.0 |
| 13 | CM | 104 | ARG | 2.0 |
| 13 | CM | 110 | ARG | 2.0 |
| 15 | CO | 88 | ARG | 2.0 |
| 16 | AP | 2 | VAL | 2.0 |
| 42 | BQ | 112 | GLU | 2.0 |
| 43 | BR | 115 | GLU | 2.0 |
| 33 | DD | 48 | ARG | 2.0 |
| 2 | CB | 11 | LEU | 2.0 |
| 7 | AG | 101 | LEU | 2.0 |
| 13 | CM | 70 | LEU | 2.0 |
| 36 | DG | 90 | LEU | 2.0 |
| 39 | DN | 116 | LEU | 2.0 |
| 45 | BT | 114 | LEU | 2.0 |
| 37 | BH | 72 | ILE | 2.0 |
| 37 | BH | 148 | ILE | 2.0 |
| 50 | BY | 75 | ILE | 2.0 |
| 1 | AA | 1006 | C | 2.0 |
| 1 | CA | 110 | C | 2.0 |

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| Mol | Chain | Res | Type | RSRZ |
|---|---|---|---|---|
| 1 | CA | 962 | C | 2.0 |
| 31 | BA | 2804 | C | 2.0 |

## 6.2 Non-standard residues in protein, DNA, RNA chains (i)

There are no non-standard protein/DNA/RNA residues in this entry.

## 6.3 Carbohydrates (i)

There are no oligosaccharides in this entry.

## 6.4 Ligands (i)

In the following table, the Atoms column lists the number of modelled atoms in the group and the number defined in the chemical component dictionary. The B-factors column lists the minimum, median, $95^{th}$ percentile and maximum values of B factors of atoms in the group. The column labelled 'Q< 0.9' lists the number of atoms with occupancy less than 0.9.

| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | DA | 3291 | 1/1 | 0.53 | 0.32 | 86,86,86,86 | 0 |
| 52 | MG | BA | 3246 | 1/1 | 0.57 | 0.34 | 75,75,75,75 | 0 |
| 52 | MG | DA | 3126 | 1/1 | 0.64 | 0.35 | 73,73,73,73 | 0 |
| 52 | MG | DA | 3306 | 1/1 | 0.65 | 0.39 | 87,87,87,87 | 0 |
| 52 | MG | AA | 1604 | 1/1 | 0.66 | 0.33 | 95,95,95,95 | 0 |
| 52 | MG | CA | 1623 | 1/1 | 0.67 | 0.24 | 67,67,67,67 | 0 |
| 52 | MG | AA | 1646 | 1/1 | 0.68 | 0.50 | 82,82,82,82 | 0 |
| 52 | MG | DA | 3245 | 1/1 | 0.70 | 0.25 | 78,78,78,78 | 0 |
| 52 | MG | BA | 3312 | 1/1 | 0.70 | 0.32 | 87,87,87,87 | 0 |
| 52 | MG | DA | 3191 | 1/1 | 0.70 | 0.32 | 91,91,91,91 | 0 |
| 52 | MG | AA | 1640 | 1/1 | 0.71 | 0.34 | 83,83,83,83 | 0 |
| 52 | MG | DA | 3302 | 1/1 | 0.71 | 0.50 | 86,86,86,86 | 0 |
| 52 | MG | DA | 3255 | 1/1 | 0.71 | 0.22 | 75,75,75,75 | 0 |
| 52 | MG | DA | 3298 | 1/1 | 0.72 | 0.29 | 71,71,71,71 | 0 |
| 52 | MG | DA | 3261 | 1/1 | 0.72 | 0.31 | 95,95,95,95 | 0 |
| 52 | MG | DA | 3219 | 1/1 | 0.72 | 0.29 | 75,75,75,75 | 0 |
| 52 | MG | BA | 3303 | 1/1 | 0.73 | 0.27 | 68,68,68,68 | 0 |
| 52 | MG | DA | 3264 | 1/1 | 0.73 | 0.39 | 80,80,80,80 | 0 |
| 52 | MG | DA | 3223 | 1/1 | 0.73 | 0.22 | 65,65,65,65 | 0 |
| 52 | MG | AA | 1641 | 1/1 | 0.74 | 0.16 | 69,69,69,69 | 0 |
| 52 | MG | DA | 3216 | 1/1 | 0.74 | 0.52 | 85,85,85,85 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 54 | K | DA | 3310 | 1/1 | 0.74 | 0.42 | 106,106,106,106 | 0 |
| 52 | MG | CA | 1647 | 1/1 | 0.75 | 0.22 | 84,84,84,84 | 0 |
| 52 | MG | CA | 1611 | 1/1 | 0.75 | 0.40 | 81,81,81,81 | 0 |
| 52 | MG | DA | 3179 | 1/1 | 0.75 | 0.47 | 77,77,77,77 | 0 |
| 52 | MG | AA | 1622 | 1/1 | 0.75 | 0.44 | 75,75,75,75 | 0 |
| 52 | MG | DA | 3102 | 1/1 | 0.76 | 0.27 | 80,80,80,80 | 0 |
| 52 | MG | AA | 1645 | 1/1 | 0.77 | 0.28 | 81,81,81,81 | 0 |
| 52 | MG | CA | 1641 | 1/1 | 0.78 | 0.59 | 87,87,87,87 | 0 |
| 52 | MG | AA | 1627 | 1/1 | 0.78 | 0.25 | 66,66,66,66 | 0 |
| 52 | MG | CA | 1633 | 1/1 | 0.78 | 0.56 | 87,87,87,87 | 0 |
| 52 | MG | DA | 3105 | 1/1 | 0.78 | 0.35 | 47,47,47,47 | 0 |
| 52 | MG | DF | 301 | 1/1 | 0.78 | 0.27 | 92,92,92,92 | 0 |
| 52 | MG | DA | 3125 | 1/1 | 0.78 | 0.41 | 58,58,58,58 | 0 |
| 52 | MG | DA | 3020 | 1/1 | 0.79 | 0.29 | 64,64,64,64 | 0 |
| 52 | MG | BA | 3097 | 1/1 | 0.79 | 0.15 | 70,70,70,70 | 0 |
| 52 | MG | BA | 3190 | 1/1 | 0.79 | 0.20 | 53,53,53,53 | 0 |
| 52 | MG | DA | 3280 | 1/1 | 0.79 | 0.43 | 77,77,77,77 | 0 |
| 52 | MG | DA | 3289 | 1/1 | 0.79 | 0.41 | 92,92,92,92 | 0 |
| 52 | MG | CA | 1609 | 1/1 | 0.79 | 0.14 | 94,94,94,94 | 0 |
| 52 | MG | AA | 1633 | 1/1 | 0.79 | 0.16 | 81,81,81,81 | 0 |
| 52 | MG | DA | 3240 | 1/1 | 0.79 | 0.26 | 89,89,89,89 | 0 |
| 52 | MG | DA | 3146 | 1/1 | 0.79 | 0.24 | 69,69,69,69 | 0 |
| 52 | MG | DA | 3247 | 1/1 | 0.79 | 0.40 | 87,87,87,87 | 0 |
| 52 | MG | DA | 3250 | 1/1 | 0.79 | 0.17 | 75,75,75,75 | 0 |
| 52 | MG | CA | 1646 | 1/1 | 0.80 | 0.22 | 80,80,80,80 | 0 |
| 52 | MG | DA | 3294 | 1/1 | 0.80 | 0.15 | 67,67,67,67 | 0 |
| 52 | MG | BA | 3031 | 1/1 | 0.80 | 0.30 | 77,77,77,77 | 0 |
| 52 | MG | CA | 1632 | 1/1 | 0.80 | 0.34 | 79,79,79,79 | 0 |
| 52 | MG | BA | 3302 | 1/1 | 0.80 | 0.22 | 72,72,72,72 | 0 |
| 52 | MG | DA | 3190 | 1/1 | 0.80 | 0.30 | 63,63,63,63 | 0 |
| 52 | MG | CA | 1620 | 1/1 | 0.80 | 0.28 | 66,66,66,66 | 0 |
| 52 | MG | DA | 3111 | 1/1 | 0.81 | 0.37 | 71,71,71,71 | 0 |
| 52 | MG | BA | 3307 | 1/1 | 0.81 | 0.39 | 76,76,76,76 | 0 |
| 52 | MG | DA | 3308 | 1/1 | 0.81 | 0.31 | 81,81,81,81 | 0 |
| 52 | MG | DA | 3266 | 1/1 | 0.81 | 0.28 | 75,75,75,75 | 0 |
| 52 | MG | BA | 3195 | 1/1 | 0.81 | 0.46 | 58,58,58,58 | 0 |
| 52 | MG | BA | 3087 | 1/1 | 0.82 | 0.22 | 58,58,58,58 | 0 |
| 52 | MG | CA | 1625 | 1/1 | 0.82 | 0.40 | 74,74,74,74 | 0 |
| 52 | MG | BA | 3092 | 1/1 | 0.82 | 0.49 | 52,52,52,52 | 0 |
| 52 | MG | DA | 3265 | 1/1 | 0.82 | 0.34 | 79,79,79,79 | 0 |
| 52 | MG | DA | 3175 | 1/1 | 0.82 | 0.43 | 67,67,67,67 | 0 |
| 52 | MG | AA | 1651 | 1/1 | 0.82 | 0.25 | 75,75,75,75 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | CA | 1637 | 1/1 | 0.82 | 0.35 | 80,80,80,80 | 0 |
| 52 | MG | CA | 1638 | 1/1 | 0.82 | 0.27 | 71,71,71,71 | 0 |
| 52 | MG | BA | 3148 | 1/1 | 0.82 | 0.21 | 58,58,58,58 | 0 |
| 52 | MG | BA | 3341 | 1/1 | 0.82 | 0.22 | 63,63,63,63 | 0 |
| 52 | MG | CA | 1605 | 1/1 | 0.82 | 0.36 | 61,61,61,61 | 0 |
| 52 | MG | CA | 1607 | 1/1 | 0.82 | 0.35 | 82,82,82,82 | 0 |
| 52 | MG | AA | 1619 | 1/1 | 0.82 | 0.31 | 56,56,56,56 | 0 |
| 52 | MG | BA | 3039 | 1/1 | 0.82 | 0.36 | 60,60,60,60 | 0 |
| 54 | K | BA | 3350 | 1/1 | 0.82 | 0.34 | 95,95,95,95 | 0 |
| 52 | MG | BA | 3076 | 1/1 | 0.82 | 0.26 | 43,43,43,43 | 0 |
| 52 | MG | DA | 3208 | 1/1 | 0.83 | 0.38 | 62,62,62,62 | 0 |
| 52 | MG | DA | 3209 | 1/1 | 0.83 | 0.22 | 59,59,59,59 | 0 |
| 52 | MG | DA | 3013 | 1/1 | 0.83 | 0.44 | 77,77,77,77 | 0 |
| 52 | MG | AA | 1643 | 1/1 | 0.83 | 0.13 | 78,78,78,78 | 0 |
| 52 | MG | DA | 3100 | 1/1 | 0.83 | 0.22 | 50,50,50,50 | 0 |
| 52 | MG | DA | 3235 | 1/1 | 0.83 | 0.31 | 79,79,79,79 | 0 |
| 52 | MG | DA | 3166 | 1/1 | 0.83 | 0.28 | 46,46,46,46 | 0 |
| 52 | MG | BA | 3332 | 1/1 | 0.83 | 0.25 | 61,61,61,61 | 0 |
| 52 | MG | BA | 3336 | 1/1 | 0.83 | 0.31 | 65,65,65,65 | 0 |
| 52 | MG | DA | 3001 | 1/1 | 0.83 | 0.31 | 76,76,76,76 | 0 |
| 52 | MG | DA | 3124 | 1/1 | 0.83 | 0.40 | 83,83,83,83 | 0 |
| 52 | MG | DA | 3198 | 1/1 | 0.83 | 0.49 | 70,70,70,70 | 0 |
| 52 | MG | DA | 3207 | 1/1 | 0.83 | 0.38 | 78,78,78,78 | 0 |
| 52 | MG | DA | 3189 | 1/1 | 0.84 | 0.33 | 63,63,63,63 | 0 |
| 52 | MG | AA | 1644 | 1/1 | 0.84 | 0.31 | 99,99,99,99 | 0 |
| 52 | MG | DA | 3062 | 1/1 | 0.84 | 0.33 | 65,65,65,65 | 0 |
| 52 | MG | BA | 3321 | 1/1 | 0.84 | 0.35 | 75,75,75,75 | 0 |
| 52 | MG | CA | 1630 | 1/1 | 0.84 | 0.28 | 77,77,77,77 | 0 |
| 52 | MG | BA | 3330 | 1/1 | 0.84 | 0.41 | 71,71,71,71 | 0 |
| 52 | MG | DA | 3284 | 1/1 | 0.84 | 0.30 | 65,65,65,65 | 0 |
| 52 | MG | AA | 1636 | 1/1 | 0.84 | 0.24 | 63,63,63,63 | 0 |
| 52 | MG | BA | 3235 | 1/1 | 0.84 | 0.33 | 72,72,72,72 | 0 |
| 52 | MG | AA | 1603 | 1/1 | 0.84 | 0.32 | 62,62,62,62 | 0 |
| 52 | MG | BA | 3249 | 1/1 | 0.84 | 0.14 | 40,40,40,40 | 0 |
| 52 | MG | DA | 3233 | 1/1 | 0.84 | 0.31 | 68,68,68,68 | 0 |
| 52 | MG | BA | 3259 | 1/1 | 0.84 | 0.30 | 46,46,46,46 | 0 |
| 52 | MG | AA | 1650 | 1/1 | 0.84 | 0.20 | 68,68,68,68 | 0 |
| 52 | MG | BA | 3157 | 1/1 | 0.84 | 0.40 | 74,74,74,74 | 0 |
| 52 | MG | DA | 3178 | 1/1 | 0.84 | 0.22 | 65,65,65,65 | 0 |
| 52 | MG | BA | 3180 | 1/1 | 0.84 | 0.30 | 64,64,64,64 | 0 |
| 52 | MG | DA | 3067 | 1/1 | 0.85 | 0.28 | 81,81,81,81 | 0 |
| 52 | MG | DA | 3081 | 1/1 | 0.85 | 0.26 | 43,43,43,43 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | DA | 3094 | 1/1 | 0.85 | 0.31 | 56,56,56,56 | 0 |
| 52 | MG | BA | 3280 | 1/1 | 0.85 | 0.35 | 75,75,75,75 | 0 |
| 52 | MG | BA | 3289 | 1/1 | 0.85 | 0.22 | 55,55,55,55 | 0 |
| 52 | MG | CA | 1644 | 1/1 | 0.85 | 0.34 | 74,74,74,74 | 0 |
| 52 | MG | DA | 3279 | 1/1 | 0.85 | 0.21 | 64,64,64,64 | 0 |
| 52 | MG | BA | 3121 | 1/1 | 0.85 | 0.31 | 57,57,57,57 | 0 |
| 52 | MG | DA | 3115 | 1/1 | 0.85 | 0.42 | 72,72,72,72 | 0 |
| 52 | MG | DA | 3288 | 1/1 | 0.85 | 0.26 | 72,72,72,72 | 0 |
| 52 | MG | BA | 3240 | 1/1 | 0.85 | 0.34 | 60,60,60,60 | 0 |
| 52 | MG | DA | 3211 | 1/1 | 0.85 | 0.28 | 79,79,79,79 | 0 |
| 52 | MG | BA | 3082 | 1/1 | 0.85 | 0.33 | 49,49,49,49 | 0 |
| 52 | MG | CA | 1606 | 1/1 | 0.85 | 0.39 | 72,72,72,72 | 0 |
| 52 | MG | BA | 3192 | 1/1 | 0.85 | 0.32 | 58,58,58,58 | 0 |
| 52 | MG | DA | 3149 | 1/1 | 0.85 | 0.19 | 55,55,55,55 | 0 |
| 52 | MG | DA | 3153 | 1/1 | 0.85 | 0.29 | 59,59,59,59 | 0 |
| 52 | MG | DA | 3037 | 1/1 | 0.85 | 0.24 | 74,74,74,74 | 0 |
| 52 | MG | DA | 3167 | 1/1 | 0.85 | 0.25 | 48,48,48,48 | 0 |
| 52 | MG | BA | 3110 | 1/1 | 0.85 | 0.42 | 45,45,45,45 | 0 |
| 52 | MG | DA | 3246 | 1/1 | 0.86 | 0.29 | 87,87,87,87 | 0 |
| 52 | MG | DA | 3070 | 1/1 | 0.86 | 0.46 | 74,74,74,74 | 0 |
| 52 | MG | DA | 3248 | 1/1 | 0.86 | 0.39 | 74,74,74,74 | 0 |
| 52 | MG | DA | 3249 | 1/1 | 0.86 | 0.40 | 79,79,79,79 | 0 |
| 52 | MG | BA | 3263 | 1/1 | 0.86 | 0.40 | 62,62,62,62 | 0 |
| 52 | MG | BA | 3178 | 1/1 | 0.86 | 0.28 | 78,78,78,78 | 0 |
| 52 | MG | DA | 3182 | 1/1 | 0.86 | 0.13 | 55,55,55,55 | 0 |
| 52 | MG | CA | 1645 | 1/1 | 0.86 | 0.19 | 97,97,97,97 | 0 |
| 52 | MG | BA | 3074 | 1/1 | 0.86 | 0.31 | 36,36,36,36 | 0 |
| 52 | MG | AA | 1637 | 1/1 | 0.86 | 0.32 | 69,69,69,69 | 0 |
| 52 | MG | D5 | 102 | 1/1 | 0.86 | 0.27 | 79,79,79,79 | 0 |
| 52 | MG | BA | 3238 | 1/1 | 0.86 | 0.19 | 49,49,49,49 | 0 |
| 52 | MG | DA | 3120 | 1/1 | 0.86 | 0.30 | 71,71,71,71 | 0 |
| 52 | MG | BA | 3306 | 1/1 | 0.86 | 0.11 | 56,56,56,56 | 0 |
| 52 | MG | CA | 1634 | 1/1 | 0.86 | 0.51 | 87,87,87,87 | 0 |
| 52 | MG | DA | 3027 | 1/1 | 0.86 | 0.31 | 61,61,61,61 | 0 |
| 52 | MG | DA | 3145 | 1/1 | 0.86 | 0.40 | 88,88,88,88 | 0 |
| 52 | MG | CA | 1636 | 1/1 | 0.86 | 0.38 | 79,79,79,79 | 0 |
| 52 | MG | DA | 3300 | 1/1 | 0.86 | 0.13 | 75,75,75,75 | 0 |
| 52 | MG | DA | 3040 | 1/1 | 0.86 | 0.18 | 43,43,43,43 | 0 |
| 52 | MG | DA | 3303 | 1/1 | 0.86 | 0.33 | 49,49,49,49 | 0 |
| 52 | MG | DA | 3305 | 1/1 | 0.86 | 0.35 | 76,76,76,76 | 0 |
| 52 | MG | DA | 3043 | 1/1 | 0.86 | 0.26 | 35,35,35,35 | 0 |
| 52 | MG | DA | 3236 | 1/1 | 0.86 | 0.31 | 67,67,67,67 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | DA | 3309 | 1/1 | 0.86 | 0.29 | 84,84,84,84 | 0 |
| 52 | MG | CA | 1616 | 1/1 | 0.86 | 0.30 | 73,73,73,73 | 0 |
| 52 | MG | DA | 3243 | 1/1 | 0.86 | 0.30 | 70,70,70,70 | 0 |
| 52 | MG | CA | 1618 | 1/1 | 0.86 | 0.27 | 62,62,62,62 | 0 |
| 52 | MG | AA | 1613 | 1/1 | 0.87 | 0.19 | 70,70,70,70 | 0 |
| 52 | MG | BA | 3274 | 1/1 | 0.87 | 0.25 | 81,81,81,81 | 0 |
| 52 | MG | DA | 3152 | 1/1 | 0.87 | 0.26 | 43,43,43,43 | 0 |
| 52 | MG | BA | 3320 | 1/1 | 0.87 | 0.14 | 54,54,54,54 | 0 |
| 52 | MG | CA | 1640 | 1/1 | 0.87 | 0.23 | 68,68,68,68 | 0 |
| 52 | MG | BA | 3095 | 1/1 | 0.87 | 0.32 | 38,38,38,38 | 0 |
| 52 | MG | DA | 3169 | 1/1 | 0.87 | 0.36 | 51,51,51,51 | 0 |
| 52 | MG | DA | 3079 | 1/1 | 0.87 | 0.23 | 59,59,59,59 | 0 |
| 52 | MG | BA | 3066 | 1/1 | 0.87 | 0.25 | 43,43,43,43 | 0 |
| 52 | MG | DA | 3087 | 1/1 | 0.87 | 0.27 | 55,55,55,55 | 0 |
| 52 | MG | DA | 3089 | 1/1 | 0.87 | 0.37 | 47,47,47,47 | 0 |
| 52 | MG | DA | 3184 | 1/1 | 0.87 | 0.32 | 63,63,63,63 | 0 |
| 52 | MG | DA | 3090 | 1/1 | 0.87 | 0.32 | 76,76,76,76 | 0 |
| 52 | MG | DA | 3282 | 1/1 | 0.87 | 0.27 | 62,62,62,62 | 0 |
| 52 | MG | CA | 1619 | 1/1 | 0.87 | 0.34 | 75,75,75,75 | 0 |
| 52 | MG | BA | 3298 | 1/1 | 0.87 | 0.26 | 61,61,61,61 | 0 |
| 52 | MG | BA | 3334 | 1/1 | 0.87 | 0.27 | 53,53,53,53 | 0 |
| 52 | MG | BA | 3300 | 1/1 | 0.87 | 0.24 | 59,59,59,59 | 0 |
| 52 | MG | DA | 3110 | 1/1 | 0.87 | 0.33 | 73,73,73,73 | 0 |
| 52 | MG | CA | 1628 | 1/1 | 0.87 | 0.33 | 75,75,75,75 | 0 |
| 52 | MG | DA | 3299 | 1/1 | 0.87 | 0.21 | 66,66,66,66 | 0 |
| 52 | MG | DA | 3113 | 1/1 | 0.87 | 0.30 | 62,62,62,62 | 0 |
| 52 | MG | DA | 3114 | 1/1 | 0.87 | 0.31 | 65,65,65,65 | 0 |
| 52 | MG | AA | 1632 | 1/1 | 0.87 | 0.45 | 72,72,72,72 | 0 |
| 52 | MG | DA | 3222 | 1/1 | 0.87 | 0.29 | 67,67,67,67 | 0 |
| 52 | MG | BQ | 202 | 1/1 | 0.87 | 0.28 | 59,59,59,59 | 0 |
| 52 | MG | DA | 3025 | 1/1 | 0.87 | 0.40 | 50,50,50,50 | 0 |
| 52 | MG | BA | 3214 | 1/1 | 0.87 | 0.42 | 66,66,66,66 | 0 |
| 52 | MG | DA | 3029 | 1/1 | 0.87 | 0.23 | 87,87,87,87 | 0 |
| 52 | MG | DA | 3135 | 1/1 | 0.87 | 0.24 | 71,71,71,71 | 0 |
| 52 | MG | BA | 3229 | 1/1 | 0.87 | 0.19 | 50,50,50,50 | 0 |
| 52 | MG | BA | 3292 | 1/1 | 0.88 | 0.29 | 60,60,60,60 | 0 |
| 52 | MG | BA | 3080 | 1/1 | 0.88 | 0.31 | 34,34,34,34 | 0 |
| 52 | MG | DA | 3154 | 1/1 | 0.88 | 0.36 | 62,62,62,62 | 0 |
| 52 | MG | DA | 3160 | 1/1 | 0.88 | 0.14 | 59,59,59,59 | 0 |
| 52 | MG | DA | 3164 | 1/1 | 0.88 | 0.19 | 71,71,71,71 | 0 |
| 52 | MG | DA | 3083 | 1/1 | 0.88 | 0.22 | 47,47,47,47 | 0 |
| 52 | MG | BA | 3151 | 1/1 | 0.88 | 0.26 | 74,74,74,74 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | BA | 3232 | 1/1 | 0.88 | 0.31 | 70,70,70,70 | 0 |
| 52 | MG | DA | 3172 | 1/1 | 0.88 | 0.39 | 64,64,64,64 | 0 |
| 52 | MG | CA | 1615 | 1/1 | 0.88 | 0.38 | 67,67,67,67 | 0 |
| 52 | MG | DA | 3176 | 1/1 | 0.88 | 0.19 | 78,78,78,78 | 0 |
| 52 | MG | BA | 3027 | 1/1 | 0.88 | 0.34 | 42,42,42,42 | 0 |
| 52 | MG | DA | 3273 | 1/1 | 0.88 | 0.33 | 75,75,75,75 | 0 |
| 52 | MG | BA | 3237 | 1/1 | 0.88 | 0.18 | 61,61,61,61 | 0 |
| 52 | MG | BA | 3162 | 1/1 | 0.88 | 0.14 | 47,47,47,47 | 0 |
| 52 | MG | DA | 3281 | 1/1 | 0.88 | 0.11 | 30,30,30,30 | 0 |
| 52 | MG | BA | 3170 | 1/1 | 0.88 | 0.17 | 69,69,69,69 | 0 |
| 52 | MG | BA | 3171 | 1/1 | 0.88 | 0.32 | 62,62,62,62 | 0 |
| 52 | MG | BA | 3176 | 1/1 | 0.88 | 0.27 | 48,48,48,48 | 0 |
| 52 | MG | BA | 3104 | 1/1 | 0.88 | 0.27 | 37,37,37,37 | 0 |
| 52 | MG | AA | 1638 | 1/1 | 0.88 | 0.16 | 82,82,82,82 | 0 |
| 52 | MG | BA | 3270 | 1/1 | 0.88 | 0.12 | 50,50,50,50 | 0 |
| 52 | MG | DA | 3039 | 1/1 | 0.88 | 0.23 | 43,43,43,43 | 0 |
| 52 | MG | BA | 3272 | 1/1 | 0.88 | 0.23 | 51,51,51,51 | 0 |
| 52 | MG | BA | 3120 | 1/1 | 0.88 | 0.30 | 52,52,52,52 | 0 |
| 52 | MG | AA | 1639 | 1/1 | 0.88 | 0.26 | 95,95,95,95 | 0 |
| 52 | MG | DA | 3127 | 1/1 | 0.88 | 0.28 | 35,35,35,35 | 0 |
| 52 | MG | DA | 3129 | 1/1 | 0.88 | 0.18 | 87,87,87,87 | 0 |
| 52 | MG | CA | 1601 | 1/1 | 0.88 | 0.28 | 83,83,83,83 | 0 |
| 52 | MG | DA | 3226 | 1/1 | 0.88 | 0.28 | 64,64,64,64 | 0 |
| 52 | MG | BA | 3138 | 1/1 | 0.88 | 0.18 | 74,74,74,74 | 0 |
| 52 | MG | DA | 3075 | 1/1 | 0.88 | 0.32 | 55,55,55,55 | 0 |
| 52 | MG | DA | 3078 | 1/1 | 0.88 | 0.28 | 46,46,46,46 | 0 |
| 52 | MG | DA | 3150 | 1/1 | 0.88 | 0.51 | 77,77,77,77 | 0 |
| 52 | MG | DA | 3197 | 1/1 | 0.89 | 0.23 | 75,75,75,75 | 0 |
| 52 | MG | BA | 3183 | 1/1 | 0.89 | 0.23 | 72,72,72,72 | 0 |
| 52 | MG | DA | 3203 | 1/1 | 0.89 | 0.37 | 67,67,67,67 | 0 |
| 52 | MG | DA | 3267 | 1/1 | 0.89 | 0.42 | 72,72,72,72 | 0 |
| 52 | MG | BA | 3166 | 1/1 | 0.89 | 0.27 | 39,39,39,39 | 0 |
| 52 | MG | DA | 3278 | 1/1 | 0.89 | 0.31 | 68,68,68,68 | 0 |
| 52 | MG | CA | 1602 | 1/1 | 0.89 | 0.36 | 70,70,70,70 | 0 |
| 52 | MG | BA | 3310 | 1/1 | 0.89 | 0.14 | 47,47,47,47 | 0 |
| 52 | MG | DA | 3031 | 1/1 | 0.89 | 0.22 | 47,47,47,47 | 0 |
| 52 | MG | BA | 3060 | 1/1 | 0.89 | 0.24 | 40,40,40,40 | 0 |
| 52 | MG | DA | 3155 | 1/1 | 0.89 | 0.25 | 62,62,62,62 | 0 |
| 52 | MG | DA | 3159 | 1/1 | 0.89 | 0.24 | 61,61,61,61 | 0 |
| 52 | MG | BA | 3315 | 1/1 | 0.89 | 0.31 | 69,69,69,69 | 0 |
| 52 | MG | BA | 3288 | 1/1 | 0.89 | 0.24 | 72,72,72,72 | 0 |
| 52 | MG | BA | 3009 | 1/1 | 0.89 | 0.33 | 38,38,38,38 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | DA | 3297 | 1/1 | 0.89 | 0.32 | 74,74,74,74 | 0 |
| 52 | MG | DA | 3057 | 1/1 | 0.89 | 0.29 | 40,40,40,40 | 0 |
| 52 | MG | BA | 3071 | 1/1 | 0.89 | 0.30 | 47,47,47,47 | 0 |
| 52 | MG | DA | 3066 | 1/1 | 0.89 | 0.24 | 60,60,60,60 | 0 |
| 52 | MG | BA | 3252 | 1/1 | 0.89 | 0.22 | 50,50,50,50 | 0 |
| 52 | MG | BA | 3034 | 1/1 | 0.89 | 0.29 | 62,62,62,62 | 0 |
| 52 | MG | AA | 1649 | 1/1 | 0.89 | 0.20 | 86,86,86,86 | 0 |
| 52 | MG | BA | 3338 | 1/1 | 0.89 | 0.21 | 73,73,73,73 | 0 |
| 52 | MG | BA | 3181 | 1/1 | 0.89 | 0.20 | 51,51,51,51 | 0 |
| 52 | MG | BA | 3346 | 1/1 | 0.89 | 0.34 | 80,80,80,80 | 0 |
| 52 | MG | BA | 3349 | 1/1 | 0.89 | 0.21 | 61,61,61,61 | 0 |
| 52 | MG | DA | 3143 | 1/1 | 0.89 | 0.27 | 57,57,57,57 | 0 |
| 52 | MG | DA | 3084 | 1/1 | 0.89 | 0.17 | 54,54,54,54 | 0 |
| 52 | MG | BA | 3186 | 1/1 | 0.90 | 0.16 | 66,66,66,66 | 0 |
| 52 | MG | BA | 3279 | 1/1 | 0.90 | 0.18 | 50,50,50,50 | 0 |
| 52 | MG | DA | 3244 | 1/1 | 0.90 | 0.22 | 86,86,86,86 | 0 |
| 52 | MG | BA | 3116 | 1/1 | 0.90 | 0.23 | 51,51,51,51 | 0 |
| 52 | MG | BA | 3179 | 1/1 | 0.90 | 0.24 | 59,59,59,59 | 0 |
| 52 | MG | BA | 3090 | 1/1 | 0.90 | 0.18 | 38,38,38,38 | 0 |
| 52 | MG | BA | 3245 | 1/1 | 0.90 | 0.29 | 45,45,45,45 | 0 |
| 52 | MG | BA | 3200 | 1/1 | 0.90 | 0.42 | 59,59,59,59 | 0 |
| 52 | MG | DA | 3091 | 1/1 | 0.90 | 0.31 | 41,41,41,41 | 0 |
| 52 | MG | DA | 3093 | 1/1 | 0.90 | 0.29 | 64,64,64,64 | 0 |
| 52 | MG | DA | 3257 | 1/1 | 0.90 | 0.33 | 63,63,63,63 | 0 |
| 52 | MG | DA | 3260 | 1/1 | 0.90 | 0.42 | 73,73,73,73 | 0 |
| 52 | MG | DA | 3024 | 1/1 | 0.90 | 0.27 | 61,61,61,61 | 0 |
| 52 | MG | DA | 3262 | 1/1 | 0.90 | 0.25 | 77,77,77,77 | 0 |
| 52 | MG | DA | 3095 | 1/1 | 0.90 | 0.34 | 53,53,53,53 | 0 |
| 52 | MG | DA | 3173 | 1/1 | 0.90 | 0.24 | 65,65,65,65 | 0 |
| 52 | MG | DA | 3098 | 1/1 | 0.90 | 0.14 | 49,49,49,49 | 0 |
| 52 | MG | BA | 3073 | 1/1 | 0.90 | 0.27 | 53,53,53,53 | 0 |
| 52 | MG | DA | 3270 | 1/1 | 0.90 | 0.42 | 65,65,65,65 | 0 |
| 52 | MG | BA | 3343 | 1/1 | 0.90 | 0.24 | 58,58,58,58 | 0 |
| 52 | MG | DA | 3275 | 1/1 | 0.90 | 0.26 | 62,62,62,62 | 0 |
| 52 | MG | CA | 1626 | 1/1 | 0.90 | 0.29 | 78,78,78,78 | 0 |
| 52 | MG | DA | 3109 | 1/1 | 0.90 | 0.43 | 71,71,71,71 | 0 |
| 52 | MG | BA | 3344 | 1/1 | 0.90 | 0.32 | 48,48,48,48 | 0 |
| 52 | MG | DA | 3036 | 1/1 | 0.90 | 0.31 | 39,39,39,39 | 0 |
| 52 | MG | CA | 1629 | 1/1 | 0.90 | 0.15 | 82,82,82,82 | 0 |
| 52 | MG | BA | 3345 | 1/1 | 0.90 | 0.28 | 60,60,60,60 | 0 |
| 52 | MG | BA | 3219 | 1/1 | 0.90 | 0.05 | 38,38,38,38 | 0 |
| 52 | MG | DA | 3041 | 1/1 | 0.90 | 0.14 | 37,37,37,37 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | DA | 3290 | 1/1 | 0.90 | 0.27 | 52,52,52,52 | 0 |
| 52 | MG | BA | 3253 | 1/1 | 0.90 | 0.18 | 51,51,51,51 | 0 |
| 52 | MG | DA | 3205 | 1/1 | 0.90 | 0.38 | 54,54,54,54 | 0 |
| 52 | MG | DA | 3295 | 1/1 | 0.90 | 0.19 | 88,88,88,88 | 0 |
| 52 | MG | BA | 3254 | 1/1 | 0.90 | 0.18 | 53,53,53,53 | 0 |
| 52 | MG | BA | 3223 | 1/1 | 0.90 | 0.28 | 36,36,36,36 | 0 |
| 52 | MG | BA | 3182 | 1/1 | 0.90 | 0.22 | 68,68,68,68 | 0 |
| 52 | MG | CA | 1603 | 1/1 | 0.90 | 0.35 | 63,63,63,63 | 0 |
| 52 | MG | DA | 3132 | 1/1 | 0.90 | 0.29 | 43,43,43,43 | 0 |
| 52 | MG | DA | 3133 | 1/1 | 0.90 | 0.36 | 53,53,53,53 | 0 |
| 52 | MG | BA | 3230 | 1/1 | 0.90 | 0.30 | 38,38,38,38 | 0 |
| 52 | MG | DA | 3072 | 1/1 | 0.90 | 0.24 | 46,46,46,46 | 0 |
| 52 | MG | DA | 3074 | 1/1 | 0.90 | 0.24 | 53,53,53,53 | 0 |
| 52 | MG | DA | 3230 | 1/1 | 0.90 | 0.17 | 56,56,56,56 | 0 |
| 52 | MG | BA | 3081 | 1/1 | 0.90 | 0.16 | 37,37,37,37 | 0 |
| 52 | MG | DA | 3076 | 1/1 | 0.90 | 0.29 | 55,55,55,55 | 0 |
| 52 | MG | BA | 3319 | 1/1 | 0.90 | 0.17 | 40,40,40,40 | 0 |
| 52 | MG | BA | 3149 | 1/1 | 0.91 | 0.17 | 51,51,51,51 | 0 |
| 52 | MG | BA | 3293 | 1/1 | 0.91 | 0.25 | 55,55,55,55 | 0 |
| 52 | MG | BA | 3295 | 1/1 | 0.91 | 0.22 | 70,70,70,70 | 0 |
| 52 | MG | CA | 1604 | 1/1 | 0.91 | 0.21 | 86,86,86,86 | 0 |
| 52 | MG | DA | 3085 | 1/1 | 0.91 | 0.25 | 54,54,54,54 | 0 |
| 52 | MG | DA | 3158 | 1/1 | 0.91 | 0.20 | 61,61,61,61 | 0 |
| 52 | MG | D5 | 101 | 1/1 | 0.91 | 0.31 | 47,47,47,47 | 0 |
| 52 | MG | BA | 3085 | 1/1 | 0.91 | 0.21 | 40,40,40,40 | 0 |
| 52 | MG | DA | 3161 | 1/1 | 0.91 | 0.33 | 72,72,72,72 | 0 |
| 52 | MG | BA | 3049 | 1/1 | 0.91 | 0.37 | 41,41,41,41 | 0 |
| 52 | MG | DA | 3005 | 1/1 | 0.91 | 0.20 | 73,73,73,73 | 0 |
| 52 | MG | BA | 3158 | 1/1 | 0.91 | 0.20 | 49,49,49,49 | 0 |
| 52 | MG | DA | 3168 | 1/1 | 0.91 | 0.08 | 53,53,53,53 | 0 |
| 52 | MG | BA | 3058 | 1/1 | 0.91 | 0.29 | 39,39,39,39 | 0 |
| 52 | MG | AA | 1625 | 1/1 | 0.91 | 0.17 | 73,73,73,73 | 0 |
| 52 | MG | CA | 1614 | 1/1 | 0.91 | 0.33 | 76,76,76,76 | 0 |
| 52 | MG | BA | 3093 | 1/1 | 0.91 | 0.36 | 43,43,43,43 | 0 |
| 52 | MG | BA | 3008 | 1/1 | 0.91 | 0.29 | 34,34,34,34 | 0 |
| 52 | MG | BA | 3067 | 1/1 | 0.91 | 0.29 | 37,37,37,37 | 0 |
| 52 | MG | DA | 3032 | 1/1 | 0.91 | 0.35 | 69,69,69,69 | 0 |
| 52 | MG | BA | 3101 | 1/1 | 0.91 | 0.23 | 39,39,39,39 | 0 |
| 52 | MG | DA | 3274 | 1/1 | 0.91 | 0.17 | 63,63,63,63 | 0 |
| 52 | MG | BA | 3103 | 1/1 | 0.91 | 0.15 | 42,42,42,42 | 0 |
| 52 | MG | DA | 3188 | 1/1 | 0.91 | 0.17 | 48,48,48,48 | 0 |
| 52 | MG | AA | 1618 | 1/1 | 0.91 | 0.42 | 72,72,72,72 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | BA | 3010 | 1/1 | 0.91 | 0.26 | 38,38,38,38 | 0 |
| 52 | MG | BA | 3112 | 1/1 | 0.91 | 0.16 | 43,43,43,43 | 0 |
| 52 | MG | DA | 3195 | 1/1 | 0.91 | 0.28 | 56,56,56,56 | 0 |
| 52 | MG | DA | 3283 | 1/1 | 0.91 | 0.48 | 72,72,72,72 | 0 |
| 52 | MG | DA | 3117 | 1/1 | 0.91 | 0.07 | 59,59,59,59 | 0 |
| 52 | MG | AA | 1629 | 1/1 | 0.91 | 0.29 | 67,67,67,67 | 0 |
| 52 | MG | DA | 3047 | 1/1 | 0.91 | 0.29 | 45,45,45,45 | 0 |
| 52 | MG | BA | 3184 | 1/1 | 0.91 | 0.24 | 50,50,50,50 | 0 |
| 52 | MG | DA | 3059 | 1/1 | 0.91 | 0.23 | 53,53,53,53 | 0 |
| 52 | MG | DA | 3292 | 1/1 | 0.91 | 0.36 | 75,75,75,75 | 0 |
| 52 | MG | BA | 3119 | 1/1 | 0.91 | 0.23 | 52,52,52,52 | 0 |
| 52 | MG | BA | 3189 | 1/1 | 0.91 | 0.26 | 45,45,45,45 | 0 |
| 52 | MG | DA | 3296 | 1/1 | 0.91 | 0.13 | 60,60,60,60 | 0 |
| 52 | MG | AA | 1611 | 1/1 | 0.91 | 0.21 | 75,75,75,75 | 0 |
| 52 | MG | AA | 1614 | 1/1 | 0.91 | 0.12 | 77,77,77,77 | 0 |
| 52 | MG | DA | 3134 | 1/1 | 0.91 | 0.25 | 47,47,47,47 | 0 |
| 52 | MG | DA | 3221 | 1/1 | 0.91 | 0.23 | 53,53,53,53 | 0 |
| 52 | MG | AA | 1634 | 1/1 | 0.91 | 0.29 | 58,58,58,58 | 0 |
| 52 | MG | DA | 3141 | 1/1 | 0.91 | 0.35 | 61,61,61,61 | 0 |
| 52 | MG | BA | 3196 | 1/1 | 0.91 | 0.25 | 65,65,65,65 | 0 |
| 52 | MG | DA | 3228 | 1/1 | 0.91 | 0.10 | 60,60,60,60 | 0 |
| 52 | MG | DA | 3307 | 1/1 | 0.91 | 0.32 | 80,80,80,80 | 0 |
| 52 | MG | DA | 3229 | 1/1 | 0.91 | 0.07 | 45,45,45,45 | 0 |
| 52 | MG | BA | 3197 | 1/1 | 0.91 | 0.24 | 52,52,52,52 | 0 |
| 52 | MG | DB | 202 | 1/1 | 0.91 | 0.28 | 63,63,63,63 | 0 |
| 52 | MG | DB | 203 | 1/1 | 0.91 | 0.31 | 56,56,56,56 | 0 |
| 52 | MG | DA | 3231 | 1/1 | 0.91 | 0.24 | 79,79,79,79 | 0 |
| 52 | MG | DR | 201 | 1/1 | 0.91 | 0.09 | 43,43,43,43 | 0 |
| 52 | MG | BA | 3198 | 1/1 | 0.91 | 0.24 | 44,44,44,44 | 0 |
| 52 | MG | BA | 3047 | 1/1 | 0.91 | 0.21 | 21,21,21,21 | 0 |
| 52 | MG | BA | 3133 | 1/1 | 0.92 | 0.27 | 35,35,35,35 | 0 |
| 52 | MG | DA | 3162 | 1/1 | 0.92 | 0.25 | 69,69,69,69 | 0 |
| 52 | MG | BA | 3134 | 1/1 | 0.92 | 0.23 | 46,46,46,46 | 0 |
| 52 | MG | DA | 3096 | 1/1 | 0.92 | 0.17 | 45,45,45,45 | 0 |
| 52 | MG | BA | 3137 | 1/1 | 0.92 | 0.26 | 61,61,61,61 | 0 |
| 52 | MG | BA | 3007 | 1/1 | 0.92 | 0.26 | 48,48,48,48 | 0 |
| 52 | MG | BA | 3301 | 1/1 | 0.92 | 0.23 | 57,57,57,57 | 0 |
| 52 | MG | DA | 3104 | 1/1 | 0.92 | 0.44 | 48,48,48,48 | 0 |
| 52 | MG | DA | 3259 | 1/1 | 0.92 | 0.38 | 74,74,74,74 | 0 |
| 52 | MG | BA | 3140 | 1/1 | 0.92 | 0.22 | 40,40,40,40 | 0 |
| 52 | MG | DA | 3108 | 1/1 | 0.92 | 0.18 | 48,48,48,48 | 0 |
| 52 | MG | DA | 3042 | 1/1 | 0.92 | 0.23 | 47,47,47,47 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | BA | 3204 | 1/1 | 0.92 | 0.23 | 46,46,46,46 | 0 |
| 52 | MG | DA | 3045 | 1/1 | 0.92 | 0.30 | 39,39,39,39 | 0 |
| 52 | MG | DA | 3112 | 1/1 | 0.92 | 0.27 | 68,68,68,68 | 0 |
| 52 | MG | BA | 3305 | 1/1 | 0.92 | 0.25 | 54,54,54,54 | 0 |
| 52 | MG | DA | 3268 | 1/1 | 0.92 | 0.52 | 81,81,81,81 | 0 |
| 52 | MG | DA | 3185 | 1/1 | 0.92 | 0.26 | 61,61,61,61 | 0 |
| 52 | MG | DA | 3272 | 1/1 | 0.92 | 0.20 | 74,74,74,74 | 0 |
| 52 | MG | DA | 3056 | 1/1 | 0.92 | 0.27 | 60,60,60,60 | 0 |
| 52 | MG | BA | 3209 | 1/1 | 0.92 | 0.31 | 56,56,56,56 | 0 |
| 52 | MG | DA | 3116 | 1/1 | 0.92 | 0.24 | 41,41,41,41 | 0 |
| 52 | MG | DA | 3277 | 1/1 | 0.92 | 0.26 | 68,68,68,68 | 0 |
| 52 | MG | DA | 3058 | 1/1 | 0.92 | 0.18 | 58,58,58,58 | 0 |
| 52 | MG | BA | 3142 | 1/1 | 0.92 | 0.36 | 39,39,39,39 | 0 |
| 52 | MG | DA | 3196 | 1/1 | 0.92 | 0.19 | 51,51,51,51 | 0 |
| 52 | MG | BA | 3013 | 1/1 | 0.92 | 0.19 | 21,21,21,21 | 0 |
| 52 | MG | BA | 3041 | 1/1 | 0.92 | 0.23 | 29,29,29,29 | 0 |
| 52 | MG | DA | 3201 | 1/1 | 0.92 | 0.21 | 59,59,59,59 | 0 |
| 52 | MG | BA | 3114 | 1/1 | 0.92 | 0.25 | 56,56,56,56 | 0 |
| 52 | MG | DA | 3285 | 1/1 | 0.92 | 0.34 | 66,66,66,66 | 0 |
| 52 | MG | DA | 3287 | 1/1 | 0.92 | 0.08 | 51,51,51,51 | 0 |
| 52 | MG | DA | 3204 | 1/1 | 0.92 | 0.29 | 45,45,45,45 | 0 |
| 52 | MG | DA | 3069 | 1/1 | 0.92 | 0.36 | 51,51,51,51 | 0 |
| 52 | MG | BA | 3316 | 1/1 | 0.92 | 0.15 | 56,56,56,56 | 0 |
| 52 | MG | BA | 3152 | 1/1 | 0.92 | 0.32 | 61,61,61,61 | 0 |
| 52 | MG | CA | 1613 | 1/1 | 0.92 | 0.26 | 80,80,80,80 | 0 |
| 52 | MG | BA | 3275 | 1/1 | 0.92 | 0.27 | 47,47,47,47 | 0 |
| 52 | MG | DA | 3212 | 1/1 | 0.92 | 0.22 | 68,68,68,68 | 0 |
| 52 | MG | BA | 3277 | 1/1 | 0.92 | 0.34 | 62,62,62,62 | 0 |
| 52 | MG | DA | 3077 | 1/1 | 0.92 | 0.39 | 49,49,49,49 | 0 |
| 52 | MG | DA | 3142 | 1/1 | 0.92 | 0.34 | 41,41,41,41 | 0 |
| 52 | MG | AA | 1608 | 1/1 | 0.92 | 0.25 | 54,54,54,54 | 0 |
| 52 | MG | DA | 3144 | 1/1 | 0.92 | 0.18 | 47,47,47,47 | 0 |
| 52 | MG | BA | 3233 | 1/1 | 0.92 | 0.12 | 63,63,63,63 | 0 |
| 52 | MG | DA | 3227 | 1/1 | 0.92 | 0.24 | 74,74,74,74 | 0 |
| 52 | MG | DA | 3009 | 1/1 | 0.92 | 0.25 | 54,54,54,54 | 0 |
| 52 | MG | BA | 3284 | 1/1 | 0.92 | 0.25 | 71,71,71,71 | 0 |
| 52 | MG | DA | 3017 | 1/1 | 0.92 | 0.15 | 47,47,47,47 | 0 |
| 52 | MG | BA | 3068 | 1/1 | 0.92 | 0.25 | 54,54,54,54 | 0 |
| 52 | MG | CA | 1621 | 1/1 | 0.92 | 0.20 | 68,68,68,68 | 0 |
| 52 | MG | DA | 3088 | 1/1 | 0.92 | 0.29 | 34,34,34,34 | 0 |
| 52 | MG | CA | 1622 | 1/1 | 0.92 | 0.15 | 78,78,78,78 | 0 |
| 52 | MG | AA | 1626 | 1/1 | 0.92 | 0.33 | 76,76,76,76 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | DA | 3242 | 1/1 | 0.92 | 0.14 | 69,69,69,69 | 0 |
| 52 | MG | DX | 101 | 1/1 | 0.92 | 0.12 | 77,77,77,77 | 0 |
| 52 | MG | CA | 1624 | 1/1 | 0.92 | 0.30 | 65,65,65,65 | 0 |
| 52 | MG | BA | 3057 | 1/1 | 0.92 | 0.22 | 44,44,44,44 | 0 |
| 52 | MG | DA | 3048 | 1/1 | 0.93 | 0.17 | 40,40,40,40 | 0 |
| 52 | MG | DA | 3052 | 1/1 | 0.93 | 0.26 | 63,63,63,63 | 0 |
| 52 | MG | DA | 3054 | 1/1 | 0.93 | 0.18 | 36,36,36,36 | 0 |
| 52 | MG | DA | 3055 | 1/1 | 0.93 | 0.23 | 42,42,42,42 | 0 |
| 52 | MG | BA | 3187 | 1/1 | 0.93 | 0.28 | 42,42,42,42 | 0 |
| 52 | MG | DA | 3234 | 1/1 | 0.93 | 0.22 | 60,60,60,60 | 0 |
| 52 | MG | DA | 3140 | 1/1 | 0.93 | 0.19 | 49,49,49,49 | 0 |
| 52 | MG | BA | 3313 | 1/1 | 0.93 | 0.30 | 54,54,54,54 | 0 |
| 52 | MG | BA | 3005 | 1/1 | 0.93 | 0.22 | 47,47,47,47 | 0 |
| 52 | MG | BA | 3150 | 1/1 | 0.93 | 0.36 | 50,50,50,50 | 0 |
| 52 | MG | DA | 3061 | 1/1 | 0.93 | 0.28 | 57,57,57,57 | 0 |
| 52 | MG | AA | 1606 | 1/1 | 0.93 | 0.35 | 73,73,73,73 | 0 |
| 52 | MG | DA | 3064 | 1/1 | 0.93 | 0.23 | 68,68,68,68 | 0 |
| 52 | MG | DA | 3147 | 1/1 | 0.93 | 0.11 | 63,63,63,63 | 0 |
| 52 | MG | BA | 3258 | 1/1 | 0.93 | 0.22 | 61,61,61,61 | 0 |
| 52 | MG | BA | 3194 | 1/1 | 0.93 | 0.32 | 44,44,44,44 | 0 |
| 52 | MG | DA | 3151 | 1/1 | 0.93 | 0.31 | 72,72,72,72 | 0 |
| 52 | MG | BA | 3326 | 1/1 | 0.93 | 0.09 | 54,54,54,54 | 0 |
| 52 | MG | DA | 3251 | 1/1 | 0.93 | 0.36 | 63,63,63,63 | 0 |
| 52 | MG | DA | 3254 | 1/1 | 0.93 | 0.11 | 65,65,65,65 | 0 |
| 52 | MG | BA | 3328 | 1/1 | 0.93 | 0.20 | 65,65,65,65 | 0 |
| 52 | MG | BA | 3329 | 1/1 | 0.93 | 0.25 | 68,68,68,68 | 0 |
| 52 | MG | BA | 3262 | 1/1 | 0.93 | 0.26 | 75,75,75,75 | 0 |
| 52 | MG | AA | 1631 | 1/1 | 0.93 | 0.11 | 60,60,60,60 | 0 |
| 52 | MG | BA | 3155 | 1/1 | 0.93 | 0.19 | 41,41,41,41 | 0 |
| 52 | MG | BA | 3271 | 1/1 | 0.93 | 0.25 | 46,46,46,46 | 0 |
| 52 | MG | BA | 3156 | 1/1 | 0.93 | 0.28 | 53,53,53,53 | 0 |
| 52 | MG | BA | 3054 | 1/1 | 0.93 | 0.16 | 68,68,68,68 | 0 |
| 52 | MG | DA | 3163 | 1/1 | 0.93 | 0.27 | 68,68,68,68 | 0 |
| 52 | MG | CA | 1642 | 1/1 | 0.93 | 0.17 | 62,62,62,62 | 0 |
| 52 | MG | AA | 1607 | 1/1 | 0.93 | 0.28 | 74,74,74,74 | 0 |
| 52 | MG | BA | 3159 | 1/1 | 0.93 | 0.24 | 45,45,45,45 | 0 |
| 52 | MG | BA | 3160 | 1/1 | 0.93 | 0.26 | 39,39,39,39 | 0 |
| 52 | MG | DA | 3086 | 1/1 | 0.93 | 0.23 | 38,38,38,38 | 0 |
| 52 | MG | DA | 3171 | 1/1 | 0.93 | 0.26 | 43,43,43,43 | 0 |
| 52 | MG | BA | 3211 | 1/1 | 0.93 | 0.13 | 39,39,39,39 | 0 |
| 52 | MG | BA | 3282 | 1/1 | 0.93 | 0.18 | 67,67,67,67 | 0 |
| 52 | MG | BP | 201 | 1/1 | 0.93 | 0.14 | 13,13,13,13 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | D7 | 101 | 1/1 | 0.93 | 0.18 | 62,62,62,62 | 0 |
| 52 | MG | DA | 3177 | 1/1 | 0.93 | 0.35 | 61,61,61,61 | 0 |
| 52 | MG | AA | 1647 | 1/1 | 0.93 | 0.19 | 66,66,66,66 | 0 |
| 52 | MG | DA | 3004 | 1/1 | 0.93 | 0.27 | 49,49,49,49 | 0 |
| 52 | MG | BA | 3285 | 1/1 | 0.93 | 0.31 | 57,57,57,57 | 0 |
| 52 | MG | BA | 3088 | 1/1 | 0.93 | 0.10 | 10,10,10,10 | 0 |
| 52 | MG | BA | 3089 | 1/1 | 0.93 | 0.20 | 26,26,26,26 | 0 |
| 52 | MG | DA | 3014 | 1/1 | 0.93 | 0.25 | 68,68,68,68 | 0 |
| 52 | MG | DA | 3099 | 1/1 | 0.93 | 0.34 | 46,46,46,46 | 0 |
| 52 | MG | BA | 3122 | 1/1 | 0.93 | 0.29 | 40,40,40,40 | 0 |
| 52 | MG | DA | 3019 | 1/1 | 0.93 | 0.34 | 42,42,42,42 | 0 |
| 52 | MG | BA | 3126 | 1/1 | 0.93 | 0.30 | 50,50,50,50 | 0 |
| 52 | MG | DA | 3023 | 1/1 | 0.93 | 0.13 | 47,47,47,47 | 0 |
| 52 | MG | DA | 3106 | 1/1 | 0.93 | 0.30 | 76,76,76,76 | 0 |
| 52 | MG | BA | 3129 | 1/1 | 0.93 | 0.07 | 18,18,18,18 | 0 |
| 52 | MG | BA | 3011 | 1/1 | 0.93 | 0.13 | 7,7,7,7 | 0 |
| 52 | MG | DA | 3202 | 1/1 | 0.93 | 0.17 | 40,40,40,40 | 0 |
| 52 | MG | BA | 3234 | 1/1 | 0.93 | 0.25 | 45,45,45,45 | 0 |
| 52 | MG | CA | 1610 | 1/1 | 0.93 | 0.12 | 61,61,61,61 | 0 |
| 52 | MG | AA | 1648 | 1/1 | 0.93 | 0.38 | 62,62,62,62 | 0 |
| 52 | MG | DA | 3301 | 1/1 | 0.93 | 0.21 | 57,57,57,57 | 0 |
| 52 | MG | CA | 1612 | 1/1 | 0.93 | 0.14 | 77,77,77,77 | 0 |
| 52 | MG | BA | 3236 | 1/1 | 0.93 | 0.46 | 70,70,70,70 | 0 |
| 52 | MG | BA | 3023 | 1/1 | 0.93 | 0.19 | 34,34,34,34 | 0 |
| 52 | MG | BA | 3304 | 1/1 | 0.93 | 0.47 | 86,86,86,86 | 0 |
| 52 | MG | AA | 1605 | 1/1 | 0.93 | 0.23 | 75,75,75,75 | 0 |
| 52 | MG | DA | 3118 | 1/1 | 0.93 | 0.15 | 66,66,66,66 | 0 |
| 52 | MG | DA | 3218 | 1/1 | 0.93 | 0.35 | 78,78,78,78 | 0 |
| 52 | MG | CA | 1617 | 1/1 | 0.93 | 0.26 | 61,61,61,61 | 0 |
| 52 | MG | DA | 3121 | 1/1 | 0.93 | 0.11 | 37,37,37,37 | 0 |
| 52 | MG | DD | 301 | 1/1 | 0.93 | 0.06 | 35,35,35,35 | 0 |
| 52 | MG | AA | 1616 | 1/1 | 0.93 | 0.19 | 77,77,77,77 | 0 |
| 52 | MG | AA | 1609 | 1/1 | 0.93 | 0.27 | 51,51,51,51 | 0 |
| 52 | MG | DA | 3225 | 1/1 | 0.93 | 0.12 | 54,54,54,54 | 0 |
| 52 | MG | BA | 3309 | 1/1 | 0.93 | 0.28 | 61,61,61,61 | 0 |
| 52 | MG | B5 | 102 | 1/1 | 0.93 | 0.20 | 56,56,56,56 | 0 |
| 55 | ZIT | DA | 3311 | 52/52 | 0.93 | 0.20 | 100,100,100,100 | 0 |
| 52 | MG | DA | 3238 | 1/1 | 0.94 | 0.34 | 73,73,73,73 | 0 |
| 52 | MG | BB | 202 | 1/1 | 0.94 | 0.16 | 30,30,30,30 | 0 |
| 52 | MG | DA | 3241 | 1/1 | 0.94 | 0.18 | 60,60,60,60 | 0 |
| 52 | MG | BB | 203 | 1/1 | 0.94 | 0.20 | 41,41,41,41 | 0 |
| 52 | MG | BB | 205 | 1/1 | 0.94 | 0.18 | 78,78,78,78 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | BF | 301 | 1/1 | 0.94 | 0.07 | 62,62,62,62 | 0 |
| 52 | MG | DA | 3082 | 1/1 | 0.94 | 0.11 | 17,17,17,17 | 0 |
| 52 | MG | AA | 1635 | 1/1 | 0.94 | 0.41 | 63,63,63,63 | 0 |
| 52 | MG | AA | 1623 | 1/1 | 0.94 | 0.27 | 54,54,54,54 | 0 |
| 52 | MG | BA | 3296 | 1/1 | 0.94 | 0.40 | 67,67,67,67 | 0 |
| 52 | MG | BA | 3118 | 1/1 | 0.94 | 0.29 | 59,59,59,59 | 0 |
| 52 | MG | DA | 3012 | 1/1 | 0.94 | 0.12 | 23,23,23,23 | 0 |
| 52 | MG | AA | 1617 | 1/1 | 0.94 | 0.36 | 64,64,64,64 | 0 |
| 52 | MG | BA | 3188 | 1/1 | 0.94 | 0.29 | 62,62,62,62 | 0 |
| 52 | MG | AA | 1621 | 1/1 | 0.94 | 0.27 | 46,46,46,46 | 0 |
| 52 | MG | BA | 3241 | 1/1 | 0.94 | 0.24 | 79,79,79,79 | 0 |
| 52 | MG | DA | 3092 | 1/1 | 0.94 | 0.15 | 61,61,61,61 | 0 |
| 52 | MG | AA | 1615 | 1/1 | 0.94 | 0.34 | 76,76,76,76 | 0 |
| 52 | MG | BA | 3191 | 1/1 | 0.94 | 0.31 | 64,64,64,64 | 0 |
| 52 | MG | B7 | 101 | 1/1 | 0.94 | 0.07 | 37,37,37,37 | 0 |
| 52 | MG | BA | 3250 | 1/1 | 0.94 | 0.26 | 54,54,54,54 | 0 |
| 52 | MG | DA | 3097 | 1/1 | 0.94 | 0.18 | 44,44,44,44 | 0 |
| 52 | MG | DA | 3026 | 1/1 | 0.94 | 0.41 | 43,43,43,43 | 0 |
| 52 | MG | BA | 3096 | 1/1 | 0.94 | 0.21 | 55,55,55,55 | 0 |
| 52 | MG | BA | 3128 | 1/1 | 0.94 | 0.28 | 54,54,54,54 | 0 |
| 52 | MG | DA | 3269 | 1/1 | 0.94 | 0.09 | 61,61,61,61 | 0 |
| 52 | MG | BA | 3161 | 1/1 | 0.94 | 0.20 | 42,42,42,42 | 0 |
| 52 | MG | DA | 3181 | 1/1 | 0.94 | 0.20 | 50,50,50,50 | 0 |
| 52 | MG | BA | 3257 | 1/1 | 0.94 | 0.19 | 48,48,48,48 | 0 |
| 52 | MG | DA | 3033 | 1/1 | 0.94 | 0.21 | 44,44,44,44 | 0 |
| 52 | MG | DA | 3034 | 1/1 | 0.94 | 0.29 | 39,39,39,39 | 0 |
| 52 | MG | DA | 3276 | 1/1 | 0.94 | 0.16 | 70,70,70,70 | 0 |
| 52 | MG | DA | 3035 | 1/1 | 0.94 | 0.36 | 54,54,54,54 | 0 |
| 52 | MG | BA | 3014 | 1/1 | 0.94 | 0.27 | 32,32,32,32 | 0 |
| 52 | MG | BA | 3165 | 1/1 | 0.94 | 0.28 | 50,50,50,50 | 0 |
| 52 | MG | DA | 3038 | 1/1 | 0.94 | 0.24 | 48,48,48,48 | 0 |
| 52 | MG | BA | 3098 | 1/1 | 0.94 | 0.23 | 46,46,46,46 | 0 |
| 52 | MG | BA | 3167 | 1/1 | 0.94 | 0.39 | 52,52,52,52 | 0 |
| 52 | MG | BA | 3208 | 1/1 | 0.94 | 0.05 | 23,23,23,23 | 0 |
| 52 | MG | BA | 3324 | 1/1 | 0.94 | 0.24 | 59,59,59,59 | 0 |
| 52 | MG | BA | 3001 | 1/1 | 0.94 | 0.26 | 49,49,49,49 | 0 |
| 52 | MG | DA | 3286 | 1/1 | 0.94 | 0.23 | 58,58,58,58 | 0 |
| 52 | MG | BA | 3327 | 1/1 | 0.94 | 0.20 | 47,47,47,47 | 0 |
| 52 | MG | BA | 3210 | 1/1 | 0.94 | 0.20 | 37,37,37,37 | 0 |
| 52 | MG | DA | 3119 | 1/1 | 0.94 | 0.10 | 64,64,64,64 | 0 |
| 52 | MG | BA | 3003 | 1/1 | 0.94 | 0.11 | 43,43,43,43 | 0 |
| 52 | MG | DA | 3206 | 1/1 | 0.94 | 0.08 | 48,48,48,48 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | BA | 3174 | 1/1 | 0.94 | 0.40 | 57,57,57,57 | 0 |
| 52 | MG | CA | 1627 | 1/1 | 0.94 | 0.08 | 66,66,66,66 | 0 |
| 52 | MG | BA | 3331 | 1/1 | 0.94 | 0.21 | 46,46,46,46 | 0 |
| 52 | MG | BA | 3084 | 1/1 | 0.94 | 0.03 | 14,14,14,14 | 0 |
| 52 | MG | BA | 3107 | 1/1 | 0.94 | 0.11 | 34,34,34,34 | 0 |
| 52 | MG | DA | 3214 | 1/1 | 0.94 | 0.12 | 65,65,65,65 | 0 |
| 52 | MG | DA | 3215 | 1/1 | 0.94 | 0.27 | 59,59,59,59 | 0 |
| 52 | MG | BA | 3224 | 1/1 | 0.94 | 0.07 | 40,40,40,40 | 0 |
| 52 | MG | BA | 3337 | 1/1 | 0.94 | 0.34 | 58,58,58,58 | 0 |
| 52 | MG | BA | 3227 | 1/1 | 0.94 | 0.07 | 39,39,39,39 | 0 |
| 52 | MG | CA | 1635 | 1/1 | 0.94 | 0.07 | 86,86,86,86 | 0 |
| 52 | MG | BA | 3283 | 1/1 | 0.94 | 0.13 | 53,53,53,53 | 0 |
| 52 | MG | DA | 3137 | 1/1 | 0.94 | 0.26 | 38,38,38,38 | 0 |
| 52 | MG | DA | 3224 | 1/1 | 0.94 | 0.17 | 68,68,68,68 | 0 |
| 52 | MG | DA | 3138 | 1/1 | 0.94 | 0.21 | 50,50,50,50 | 0 |
| 52 | MG | DA | 3065 | 1/1 | 0.94 | 0.10 | 49,49,49,49 | 0 |
| 52 | MG | BA | 3342 | 1/1 | 0.94 | 0.19 | 49,49,49,49 | 0 |
| 52 | MG | BA | 3228 | 1/1 | 0.94 | 0.33 | 69,69,69,69 | 0 |
| 52 | MG | DA | 3068 | 1/1 | 0.94 | 0.30 | 49,49,49,49 | 0 |
| 52 | MG | BA | 3141 | 1/1 | 0.94 | 0.23 | 27,27,27,27 | 0 |
| 52 | MG | BA | 3063 | 1/1 | 0.94 | 0.23 | 48,48,48,48 | 0 |
| 52 | MG | BA | 3111 | 1/1 | 0.94 | 0.08 | 19,19,19,19 | 0 |
| 52 | MG | BA | 3347 | 1/1 | 0.94 | 0.20 | 66,66,66,66 | 0 |
| 52 | MG | BA | 3348 | 1/1 | 0.94 | 0.19 | 61,61,61,61 | 0 |
| 55 | ZIT | BA | 3351 | 52/52 | 0.94 | 0.20 | 100,100,100,100 | 0 |
| 52 | MG | AA | 1628 | 1/1 | 0.94 | 0.31 | 66,66,66,66 | 0 |
| 52 | MG | BA | 3339 | 1/1 | 0.95 | 0.20 | 41,41,41,41 | 0 |
| 52 | MG | CA | 1639 | 1/1 | 0.95 | 0.24 | 64,64,64,64 | 0 |
| 52 | MG | BA | 3105 | 1/1 | 0.95 | 0.32 | 46,46,46,46 | 0 |
| 52 | MG | BA | 3278 | 1/1 | 0.95 | 0.13 | 41,41,41,41 | 0 |
| 52 | MG | DA | 3237 | 1/1 | 0.95 | 0.12 | 53,53,53,53 | 0 |
| 52 | MG | BA | 3135 | 1/1 | 0.95 | 0.15 | 8,8,8,8 | 0 |
| 52 | MG | DA | 3239 | 1/1 | 0.95 | 0.17 | 59,59,59,59 | 0 |
| 52 | MG | CA | 1643 | 1/1 | 0.95 | 0.32 | 62,62,62,62 | 0 |
| 52 | MG | DA | 3071 | 1/1 | 0.95 | 0.21 | 40,40,40,40 | 0 |
| 52 | MG | BA | 3019 | 1/1 | 0.95 | 0.24 | 24,24,24,24 | 0 |
| 52 | MG | DA | 3073 | 1/1 | 0.95 | 0.21 | 42,42,42,42 | 0 |
| 52 | MG | BA | 3217 | 1/1 | 0.95 | 0.24 | 50,50,50,50 | 0 |
| 52 | MG | BA | 3109 | 1/1 | 0.95 | 0.30 | 34,34,34,34 | 0 |
| 52 | MG | BA | 3139 | 1/1 | 0.95 | 0.22 | 24,24,24,24 | 0 |
| 52 | MG | CA | 1648 | 1/1 | 0.95 | 0.21 | 79,79,79,79 | 0 |
| 52 | MG | DA | 3157 | 1/1 | 0.95 | 0.40 | 65,65,65,65 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | D0 | 101 | 1/1 | 0.95 | 0.18 | 62,62,62,62 | 0 |
| 52 | MG | D1 | 101 | 1/1 | 0.95 | 0.18 | 50,50,50,50 | 0 |
| 52 | MG | DA | 3080 | 1/1 | 0.95 | 0.30 | 40,40,40,40 | 0 |
| 52 | MG | DA | 3252 | 1/1 | 0.95 | 0.38 | 66,66,66,66 | 0 |
| 52 | MG | BA | 3175 | 1/1 | 0.95 | 0.18 | 49,49,49,49 | 0 |
| 52 | MG | BA | 3287 | 1/1 | 0.95 | 0.33 | 58,58,58,58 | 0 |
| 52 | MG | BA | 3226 | 1/1 | 0.95 | 0.14 | 32,32,32,32 | 0 |
| 52 | MG | DA | 3258 | 1/1 | 0.95 | 0.08 | 61,61,61,61 | 0 |
| 52 | MG | BA | 3043 | 1/1 | 0.95 | 0.10 | 36,36,36,36 | 0 |
| 52 | MG | BA | 3177 | 1/1 | 0.95 | 0.22 | 52,52,52,52 | 0 |
| 52 | MG | BA | 3045 | 1/1 | 0.95 | 0.20 | 26,26,26,26 | 0 |
| 52 | MG | BA | 3006 | 1/1 | 0.95 | 0.23 | 29,29,29,29 | 0 |
| 52 | MG | DA | 3263 | 1/1 | 0.95 | 0.19 | 67,67,67,67 | 0 |
| 52 | MG | DA | 3011 | 1/1 | 0.95 | 0.26 | 50,50,50,50 | 0 |
| 52 | MG | BA | 3144 | 1/1 | 0.95 | 0.19 | 53,53,53,53 | 0 |
| 52 | MG | BX | 101 | 1/1 | 0.95 | 0.11 | 61,61,61,61 | 0 |
| 52 | MG | BA | 3297 | 1/1 | 0.95 | 0.31 | 61,61,61,61 | 0 |
| 52 | MG | DA | 3174 | 1/1 | 0.95 | 0.39 | 63,63,63,63 | 0 |
| 52 | MG | BA | 3145 | 1/1 | 0.95 | 0.23 | 40,40,40,40 | 0 |
| 52 | MG | DA | 3018 | 1/1 | 0.95 | 0.30 | 32,32,32,32 | 0 |
| 52 | MG | BA | 3146 | 1/1 | 0.95 | 0.23 | 42,42,42,42 | 0 |
| 52 | MG | BA | 3113 | 1/1 | 0.95 | 0.25 | 26,26,26,26 | 0 |
| 52 | MG | DA | 3022 | 1/1 | 0.95 | 0.33 | 47,47,47,47 | 0 |
| 52 | MG | BA | 3025 | 1/1 | 0.95 | 0.20 | 54,54,54,54 | 0 |
| 52 | MG | BA | 3185 | 1/1 | 0.95 | 0.18 | 14,14,14,14 | 0 |
| 52 | MG | BA | 3050 | 1/1 | 0.95 | 0.23 | 38,38,38,38 | 0 |
| 52 | MG | BA | 3094 | 1/1 | 0.95 | 0.28 | 52,52,52,52 | 0 |
| 52 | MG | DA | 3186 | 1/1 | 0.95 | 0.19 | 63,63,63,63 | 0 |
| 52 | MG | BA | 3051 | 1/1 | 0.95 | 0.18 | 14,14,14,14 | 0 |
| 52 | MG | BA | 3242 | 1/1 | 0.95 | 0.16 | 48,48,48,48 | 0 |
| 52 | MG | BA | 3244 | 1/1 | 0.95 | 0.20 | 40,40,40,40 | 0 |
| 52 | MG | BA | 3153 | 1/1 | 0.95 | 0.26 | 25,25,25,25 | 0 |
| 52 | MG | DA | 3194 | 1/1 | 0.95 | 0.28 | 60,60,60,60 | 0 |
| 52 | MG | BA | 3311 | 1/1 | 0.95 | 0.15 | 46,46,46,46 | 0 |
| 52 | MG | BA | 3154 | 1/1 | 0.95 | 0.13 | 32,32,32,32 | 0 |
| 52 | MG | BA | 3077 | 1/1 | 0.95 | 0.26 | 28,28,28,28 | 0 |
| 52 | MG | AA | 1610 | 1/1 | 0.95 | 0.31 | 65,65,65,65 | 0 |
| 52 | MG | DA | 3199 | 1/1 | 0.95 | 0.13 | 47,47,47,47 | 0 |
| 52 | MG | AA | 1630 | 1/1 | 0.95 | 0.32 | 59,59,59,59 | 0 |
| 52 | MG | BA | 3317 | 1/1 | 0.95 | 0.10 | 41,41,41,41 | 0 |
| 52 | MG | BA | 3125 | 1/1 | 0.95 | 0.17 | 46,46,46,46 | 0 |
| 52 | MG | AA | 1612 | 1/1 | 0.95 | 0.24 | 66,66,66,66 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | BA | 3255 | 1/1 | 0.95 | 0.16 | 45,45,45,45 | 0 |
| 52 | MG | BA | 3323 | 1/1 | 0.95 | 0.11 | 64,64,64,64 | 0 |
| 52 | MG | BA | 3256 | 1/1 | 0.95 | 0.32 | 63,63,63,63 | 0 |
| 52 | MG | DA | 3044 | 1/1 | 0.95 | 0.24 | 46,46,46,46 | 0 |
| 52 | MG | BA | 3102 | 1/1 | 0.95 | 0.16 | 24,24,24,24 | 0 |
| 52 | MG | BA | 3038 | 1/1 | 0.95 | 0.31 | 25,25,25,25 | 0 |
| 52 | MG | DA | 3122 | 1/1 | 0.95 | 0.20 | 61,61,61,61 | 0 |
| 52 | MG | BA | 3132 | 1/1 | 0.95 | 0.26 | 55,55,55,55 | 0 |
| 52 | MG | DA | 3049 | 1/1 | 0.95 | 0.22 | 35,35,35,35 | 0 |
| 52 | MG | DA | 3304 | 1/1 | 0.95 | 0.22 | 52,52,52,52 | 0 |
| 52 | MG | DA | 3050 | 1/1 | 0.95 | 0.20 | 42,42,42,42 | 0 |
| 52 | MG | BA | 3201 | 1/1 | 0.95 | 0.20 | 29,29,29,29 | 0 |
| 52 | MG | DA | 3128 | 1/1 | 0.95 | 0.41 | 51,51,51,51 | 0 |
| 52 | MG | BA | 3202 | 1/1 | 0.95 | 0.18 | 41,41,41,41 | 0 |
| 52 | MG | BA | 3269 | 1/1 | 0.95 | 0.11 | 55,55,55,55 | 0 |
| 52 | MG | DB | 201 | 1/1 | 0.95 | 0.26 | 52,52,52,52 | 0 |
| 52 | MG | CA | 1631 | 1/1 | 0.95 | 0.43 | 70,70,70,70 | 0 |
| 52 | MG | BA | 3203 | 1/1 | 0.95 | 0.25 | 35,35,35,35 | 0 |
| 52 | MG | BA | 3333 | 1/1 | 0.95 | 0.27 | 80,80,80,80 | 0 |
| 52 | MG | DA | 3136 | 1/1 | 0.95 | 0.29 | 59,59,59,59 | 0 |
| 52 | MG | DP | 201 | 1/1 | 0.95 | 0.08 | 50,50,50,50 | 0 |
| 52 | MG | DQ | 201 | 1/1 | 0.95 | 0.27 | 78,78,78,78 | 0 |
| 52 | MG | BA | 3163 | 1/1 | 0.95 | 0.29 | 47,47,47,47 | 0 |
| 52 | MG | BA | 3205 | 1/1 | 0.95 | 0.21 | 55,55,55,55 | 0 |
| 52 | MG | DA | 3139 | 1/1 | 0.95 | 0.21 | 68,68,68,68 | 0 |
| 52 | MG | BA | 3164 | 1/1 | 0.95 | 0.20 | 47,47,47,47 | 0 |
| 52 | MG | BA | 3015 | 1/1 | 0.95 | 0.20 | 48,48,48,48 | 0 |
| 52 | MG | DA | 3232 | 1/1 | 0.95 | 0.16 | 72,72,72,72 | 0 |
| 52 | MG | BA | 3221 | 1/1 | 0.96 | 0.33 | 40,40,40,40 | 0 |
| 52 | MG | BA | 3059 | 1/1 | 0.96 | 0.23 | 39,39,39,39 | 0 |
| 52 | MG | AA | 1624 | 1/1 | 0.96 | 0.35 | 56,56,56,56 | 0 |
| 52 | MG | BA | 3026 | 1/1 | 0.96 | 0.11 | 49,49,49,49 | 0 |
| 52 | MG | BA | 3136 | 1/1 | 0.96 | 0.28 | 32,32,32,32 | 0 |
| 52 | MG | BA | 3308 | 1/1 | 0.96 | 0.34 | 64,64,64,64 | 0 |
| 52 | MG | BD | 301 | 1/1 | 0.96 | 0.12 | 43,43,43,43 | 0 |
| 52 | MG | BA | 3267 | 1/1 | 0.96 | 0.22 | 38,38,38,38 | 0 |
| 52 | MG | BA | 3268 | 1/1 | 0.96 | 0.09 | 40,40,40,40 | 0 |
| 52 | MG | DA | 3210 | 1/1 | 0.96 | 0.16 | 63,63,63,63 | 0 |
| 52 | MG | DA | 3148 | 1/1 | 0.96 | 0.21 | 48,48,48,48 | 0 |
| 52 | MG | BA | 3064 | 1/1 | 0.96 | 0.15 | 28,28,28,28 | 0 |
| 52 | MG | DA | 3213 | 1/1 | 0.96 | 0.24 | 36,36,36,36 | 0 |
| 52 | MG | BU | 201 | 1/1 | 0.96 | 0.17 | 26,26,26,26 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | BA | 3044 | 1/1 | 0.96 | 0.20 | 26,26,26,26 | 0 |
| 52 | MG | AA | 1601 | 1/1 | 0.96 | 0.21 | 58,58,58,58 | 0 |
| 52 | MG | DA | 3217 | 1/1 | 0.96 | 0.33 | 62,62,62,62 | 0 |
| 52 | MG | DA | 3046 | 1/1 | 0.96 | 0.20 | 28,28,28,28 | 0 |
| 52 | MG | BA | 3314 | 1/1 | 0.96 | 0.10 | 56,56,56,56 | 0 |
| 52 | MG | BA | 3231 | 1/1 | 0.96 | 0.33 | 52,52,52,52 | 0 |
| 52 | MG | DA | 3101 | 1/1 | 0.96 | 0.24 | 43,43,43,43 | 0 |
| 52 | MG | BA | 3273 | 1/1 | 0.96 | 0.25 | 58,58,58,58 | 0 |
| 52 | MG | BA | 3028 | 1/1 | 0.96 | 0.36 | 28,28,28,28 | 0 |
| 52 | MG | BA | 3069 | 1/1 | 0.96 | 0.21 | 18,18,18,18 | 0 |
| 52 | MG | BA | 3276 | 1/1 | 0.96 | 0.33 | 55,55,55,55 | 0 |
| 52 | MG | DA | 3107 | 1/1 | 0.96 | 0.27 | 38,38,38,38 | 0 |
| 52 | MG | CA | 1608 | 1/1 | 0.96 | 0.27 | 51,51,51,51 | 0 |
| 52 | MG | BA | 3115 | 1/1 | 0.96 | 0.28 | 49,49,49,49 | 0 |
| 52 | MG | BA | 3168 | 1/1 | 0.96 | 0.11 | 43,43,43,43 | 0 |
| 52 | MG | BA | 3143 | 1/1 | 0.96 | 0.18 | 30,30,30,30 | 0 |
| 52 | MG | DA | 3293 | 1/1 | 0.96 | 0.22 | 53,53,53,53 | 0 |
| 52 | MG | BA | 3070 | 1/1 | 0.96 | 0.22 | 35,35,35,35 | 0 |
| 52 | MG | AA | 1642 | 1/1 | 0.96 | 0.23 | 51,51,51,51 | 0 |
| 52 | MG | DA | 3006 | 1/1 | 0.96 | 0.28 | 39,39,39,39 | 0 |
| 52 | MG | DA | 3007 | 1/1 | 0.96 | 0.16 | 39,39,39,39 | 0 |
| 52 | MG | BA | 3239 | 1/1 | 0.96 | 0.23 | 48,48,48,48 | 0 |
| 52 | MG | DA | 3010 | 1/1 | 0.96 | 0.21 | 31,31,31,31 | 0 |
| 52 | MG | BA | 3072 | 1/1 | 0.96 | 0.16 | 24,24,24,24 | 0 |
| 52 | MG | BA | 3020 | 1/1 | 0.96 | 0.26 | 38,38,38,38 | 0 |
| 52 | MG | BA | 3036 | 1/1 | 0.96 | 0.05 | 0,0,0,0 | 0 |
| 52 | MG | BA | 3052 | 1/1 | 0.96 | 0.16 | 15,15,15,15 | 0 |
| 52 | MG | BA | 3100 | 1/1 | 0.96 | 0.15 | 21,21,21,21 | 0 |
| 52 | MG | DA | 3123 | 1/1 | 0.96 | 0.12 | 61,61,61,61 | 0 |
| 52 | MG | BA | 3022 | 1/1 | 0.96 | 0.23 | 49,49,49,49 | 0 |
| 52 | MG | BA | 3335 | 1/1 | 0.96 | 0.27 | 52,52,52,52 | 0 |
| 52 | MG | BA | 3127 | 1/1 | 0.96 | 0.11 | 50,50,50,50 | 0 |
| 52 | MG | BA | 3294 | 1/1 | 0.96 | 0.30 | 65,65,65,65 | 0 |
| 52 | MG | DA | 3187 | 1/1 | 0.96 | 0.11 | 52,52,52,52 | 0 |
| 52 | MG | BA | 3078 | 1/1 | 0.96 | 0.19 | 34,34,34,34 | 0 |
| 52 | MG | BA | 3251 | 1/1 | 0.96 | 0.07 | 35,35,35,35 | 0 |
| 52 | MG | DA | 3130 | 1/1 | 0.96 | 0.24 | 43,43,43,43 | 0 |
| 52 | MG | DE | 301 | 1/1 | 0.96 | 0.16 | 40,40,40,40 | 0 |
| 52 | MG | BA | 3213 | 1/1 | 0.96 | 0.28 | 32,32,32,32 | 0 |
| 52 | MG | B1 | 101 | 1/1 | 0.96 | 0.13 | 39,39,39,39 | 0 |
| 52 | MG | BA | 3299 | 1/1 | 0.96 | 0.34 | 58,58,58,58 | 0 |
| 52 | MG | DA | 3256 | 1/1 | 0.96 | 0.12 | 57,57,57,57 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | DU | 201 | 1/1 | 0.96 | 0.18 | 60,60,60,60 | 0 |
| 52 | MG | BA | 3216 | 1/1 | 0.96 | 0.32 | 43,43,43,43 | 0 |
| 53 | ZN | AD | 301 | 1/1 | 0.96 | 0.22 | 108,108,108,108 | 0 |
| 52 | MG | DA | 3030 | 1/1 | 0.96 | 0.28 | 42,42,42,42 | 0 |
| 52 | MG | BA | 3130 | 1/1 | 0.96 | 0.15 | 26,26,26,26 | 0 |
| 52 | MG | BA | 3040 | 1/1 | 0.96 | 0.38 | 51,51,51,51 | 0 |
| 52 | MG | DA | 3200 | 1/1 | 0.96 | 0.25 | 50,50,50,50 | 0 |
| 52 | MG | BP | 202 | 1/1 | 0.97 | 0.24 | 58,58,58,58 | 0 |
| 52 | MG | DA | 3165 | 1/1 | 0.97 | 0.14 | 52,52,52,52 | 0 |
| 52 | MG | BQ | 201 | 1/1 | 0.97 | 0.04 | 32,32,32,32 | 0 |
| 52 | MG | DA | 3220 | 1/1 | 0.97 | 0.17 | 62,62,62,62 | 0 |
| 52 | MG | DA | 3028 | 1/1 | 0.97 | 0.18 | 39,39,39,39 | 0 |
| 52 | MG | BA | 3322 | 1/1 | 0.97 | 0.12 | 20,20,20,20 | 0 |
| 52 | MG | BR | 201 | 1/1 | 0.97 | 0.10 | 20,20,20,20 | 0 |
| 52 | MG | DA | 3170 | 1/1 | 0.97 | 0.22 | 53,53,53,53 | 0 |
| 52 | MG | BA | 3086 | 1/1 | 0.97 | 0.12 | 18,18,18,18 | 0 |
| 52 | MG | BA | 3291 | 1/1 | 0.97 | 0.37 | 54,54,54,54 | 0 |
| 52 | MG | BA | 3325 | 1/1 | 0.97 | 0.26 | 43,43,43,43 | 0 |
| 52 | MG | BA | 3225 | 1/1 | 0.97 | 0.19 | 33,33,33,33 | 0 |
| 52 | MG | BA | 3018 | 1/1 | 0.97 | 0.10 | 26,26,26,26 | 0 |
| 52 | MG | AA | 1602 | 1/1 | 0.97 | 0.26 | 37,37,37,37 | 0 |
| 52 | MG | BA | 3199 | 1/1 | 0.97 | 0.31 | 49,49,49,49 | 0 |
| 52 | MG | BA | 3037 | 1/1 | 0.97 | 0.15 | 14,14,14,14 | 0 |
| 52 | MG | BA | 3123 | 1/1 | 0.97 | 0.20 | 22,22,22,22 | 0 |
| 52 | MG | DA | 3131 | 1/1 | 0.97 | 0.26 | 47,47,47,47 | 0 |
| 52 | MG | BA | 3124 | 1/1 | 0.97 | 0.06 | 42,42,42,42 | 0 |
| 52 | MG | DA | 3183 | 1/1 | 0.97 | 0.22 | 44,44,44,44 | 0 |
| 52 | MG | BA | 3264 | 1/1 | 0.97 | 0.15 | 35,35,35,35 | 0 |
| 52 | MG | BA | 3265 | 1/1 | 0.97 | 0.29 | 63,63,63,63 | 0 |
| 52 | MG | BA | 3046 | 1/1 | 0.97 | 0.19 | 37,37,37,37 | 0 |
| 52 | MG | BA | 3004 | 1/1 | 0.97 | 0.17 | 23,23,23,23 | 0 |
| 52 | MG | BA | 3062 | 1/1 | 0.97 | 0.30 | 44,44,44,44 | 0 |
| 52 | MG | BA | 3206 | 1/1 | 0.97 | 0.20 | 29,29,29,29 | 0 |
| 52 | MG | BA | 3207 | 1/1 | 0.97 | 0.10 | 23,23,23,23 | 0 |
| 52 | MG | DA | 3002 | 1/1 | 0.97 | 0.32 | 38,38,38,38 | 0 |
| 52 | MG | DA | 3192 | 1/1 | 0.97 | 0.26 | 55,55,55,55 | 0 |
| 52 | MG | DA | 3193 | 1/1 | 0.97 | 0.30 | 50,50,50,50 | 0 |
| 52 | MG | DA | 3003 | 1/1 | 0.97 | 0.28 | 56,56,56,56 | 0 |
| 52 | MG | BA | 3340 | 1/1 | 0.97 | 0.28 | 62,62,62,62 | 0 |
| 52 | MG | BA | 3048 | 1/1 | 0.97 | 0.25 | 30,30,30,30 | 0 |
| 52 | MG | DA | 3053 | 1/1 | 0.97 | 0.43 | 51,51,51,51 | 0 |
| 52 | MG | AA | 1620 | 1/1 | 0.97 | 0.15 | 73,73,73,73 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | BA | 3065 | 1/1 | 0.97 | 0.20 | 32,32,32,32 | 0 |
| 52 | MG | BA | 3029 | 1/1 | 0.97 | 0.14 | 26,26,26,26 | 0 |
| 52 | MG | BA | 3212 | 1/1 | 0.97 | 0.24 | 30,30,30,30 | 0 |
| 52 | MG | DA | 3103 | 1/1 | 0.97 | 0.48 | 70,70,70,70 | 0 |
| 52 | MG | B5 | 101 | 1/1 | 0.97 | 0.09 | 27,27,27,27 | 0 |
| 52 | MG | BA | 3243 | 1/1 | 0.97 | 0.10 | 58,58,58,58 | 0 |
| 52 | MG | BA | 3169 | 1/1 | 0.97 | 0.23 | 46,46,46,46 | 0 |
| 52 | MG | BA | 3042 | 1/1 | 0.97 | 0.17 | 15,15,15,15 | 0 |
| 52 | MG | DA | 3063 | 1/1 | 0.97 | 0.19 | 47,47,47,47 | 0 |
| 52 | MG | BB | 201 | 1/1 | 0.97 | 0.27 | 42,42,42,42 | 0 |
| 52 | MG | DA | 3156 | 1/1 | 0.97 | 0.33 | 44,44,44,44 | 0 |
| 52 | MG | BA | 3117 | 1/1 | 0.97 | 0.29 | 39,39,39,39 | 0 |
| 52 | MG | BA | 3218 | 1/1 | 0.97 | 0.25 | 33,33,33,33 | 0 |
| 52 | MG | BA | 3172 | 1/1 | 0.97 | 0.25 | 64,64,64,64 | 0 |
| 52 | MG | BA | 3220 | 1/1 | 0.97 | 0.23 | 27,27,27,27 | 0 |
| 53 | ZN | AN | 101 | 1/1 | 0.97 | 0.05 | 159,159,159,159 | 0 |
| 53 | ZN | CD | 301 | 1/1 | 0.97 | 0.17 | 93,93,93,93 | 0 |
| 53 | ZN | CN | 101 | 1/1 | 0.97 | 0.06 | 157,157,157,157 | 0 |
| 52 | MG | BE | 301 | 1/1 | 0.97 | 0.22 | 29,29,29,29 | 0 |
| 52 | MG | BA | 3173 | 1/1 | 0.97 | 0.21 | 24,24,24,24 | 0 |
| 52 | MG | BA | 3053 | 1/1 | 0.97 | 0.19 | 15,15,15,15 | 0 |
| 52 | MG | DA | 3271 | 1/1 | 0.97 | 0.08 | 46,46,46,46 | 0 |
| 52 | MG | BA | 3281 | 1/1 | 0.98 | 0.31 | 46,46,46,46 | 0 |
| 52 | MG | BA | 3091 | 1/1 | 0.98 | 0.15 | 9,9,9,9 | 0 |
| 52 | MG | DA | 3015 | 1/1 | 0.98 | 0.11 | 23,23,23,23 | 0 |
| 52 | MG | DA | 3016 | 1/1 | 0.98 | 0.34 | 56,56,56,56 | 0 |
| 52 | MG | BA | 3106 | 1/1 | 0.98 | 0.11 | 12,12,12,12 | 0 |
| 52 | MG | BA | 3261 | 1/1 | 0.98 | 0.13 | 38,38,38,38 | 0 |
| 52 | MG | BA | 3222 | 1/1 | 0.98 | 0.15 | 23,23,23,23 | 0 |
| 52 | MG | BA | 3030 | 1/1 | 0.98 | 0.11 | 17,17,17,17 | 0 |
| 52 | MG | DA | 3021 | 1/1 | 0.98 | 0.23 | 47,47,47,47 | 0 |
| 52 | MG | DA | 3051 | 1/1 | 0.98 | 0.19 | 26,26,26,26 | 0 |
| 52 | MG | BA | 3079 | 1/1 | 0.98 | 0.12 | 0,0,0,0 | 0 |
| 52 | MG | B0 | 101 | 1/1 | 0.98 | 0.16 | 34,34,34,34 | 0 |
| 52 | MG | BA | 3290 | 1/1 | 0.98 | 0.24 | 47,47,47,47 | 0 |
| 52 | MG | BA | 3032 | 1/1 | 0.98 | 0.20 | 15,15,15,15 | 0 |
| 52 | MG | DA | 3180 | 1/1 | 0.98 | 0.24 | 52,52,52,52 | 0 |
| 52 | MG | BA | 3012 | 1/1 | 0.98 | 0.13 | 22,22,22,22 | 0 |
| 52 | MG | BA | 3061 | 1/1 | 0.98 | 0.06 | 23,23,23,23 | 0 |
| 52 | MG | BA | 3247 | 1/1 | 0.98 | 0.03 | 40,40,40,40 | 0 |
| 52 | MG | BA | 3318 | 1/1 | 0.98 | 0.28 | 60,60,60,60 | 0 |
| 52 | MG | DA | 3060 | 1/1 | 0.98 | 0.04 | 34,34,34,34 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å$^2$) | Q<0.9 |
|---|---|---|---|---|---|---|---|---|
| 52 | MG | BA | 3248 | 1/1 | 0.98 | 0.05 | 47,47,47,47 | 0 |
| 52 | MG | BA | 3193 | 1/1 | 0.98 | 0.19 | 28,28,28,28 | 0 |
| 52 | MG | BA | 3035 | 1/1 | 0.98 | 0.11 | 21,21,21,21 | 0 |
| 52 | MG | DA | 3253 | 1/1 | 0.98 | 0.25 | 50,50,50,50 | 0 |
| 52 | MG | BA | 3099 | 1/1 | 0.98 | 0.15 | 26,26,26,26 | 0 |
| 52 | MG | BA | 3147 | 1/1 | 0.98 | 0.04 | 12,12,12,12 | 0 |
| 52 | MG | BA | 3021 | 1/1 | 0.98 | 0.14 | 16,16,16,16 | 0 |
| 52 | MG | BA | 3016 | 1/1 | 0.98 | 0.12 | 21,21,21,21 | 0 |
| 52 | MG | DA | 3008 | 1/1 | 0.98 | 0.41 | 52,52,52,52 | 0 |
| 52 | MG | BA | 3075 | 1/1 | 0.98 | 0.17 | 26,26,26,26 | 0 |
| 52 | MG | BA | 3002 | 1/1 | 0.98 | 0.20 | 20,20,20,20 | 0 |
| 52 | MG | BA | 3056 | 1/1 | 0.98 | 0.07 | 20,20,20,20 | 0 |
| 52 | MG | BB | 204 | 1/1 | 0.98 | 0.38 | 56,56,56,56 | 0 |
| 52 | MG | BA | 3033 | 1/1 | 0.99 | 0.06 | 20,20,20,20 | 0 |
| 52 | MG | BA | 3017 | 1/1 | 0.99 | 0.26 | 32,32,32,32 | 0 |
| 52 | MG | BA | 3215 | 1/1 | 0.99 | 0.18 | 36,36,36,36 | 0 |
| 52 | MG | BA | 3266 | 1/1 | 0.99 | 0.21 | 35,35,35,35 | 0 |
| 52 | MG | BA | 3131 | 1/1 | 0.99 | 0.11 | 45,45,45,45 | 0 |
| 52 | MG | BA | 3286 | 1/1 | 0.99 | 0.03 | 45,45,45,45 | 0 |
| 52 | MG | BA | 3055 | 1/1 | 0.99 | 0.19 | 19,19,19,19 | 0 |
| 52 | MG | BA | 3260 | 1/1 | 0.99 | 0.17 | 42,42,42,42 | 0 |
| 52 | MG | BA | 3083 | 1/1 | 0.99 | 0.06 | 5,5,5,5 | 0 |
| 52 | MG | BP | 203 | 1/1 | 0.99 | 0.09 | 0,0,0,0 | 0 |
| 52 | MG | BA | 3108 | 1/1 | 0.99 | 0.27 | 56,56,56,56 | 0 |
| 52 | MG | BA | 3024 | 1/1 | 1.00 | 0.03 | 2,2,2,2 | 0 |

## 6.5 Other polymers (i)

There are no such residues in this entry.