



Full wwPDB EM Validation Report ⓘ

Mar 8, 2026 – 03:47 PM UTC

PDB ID : 8UXF / pdb_00008uxf
EMDB ID : EMD-42762
Title : Structure of PKA phosphorylated human RyR2-R420W in the primed state
Authors : Miotto, M.C.; Marks, A.R.
Deposited on : 2023-11-09
Resolution : 3.13 Å(reported)
Based on initial model : 7UA5

This is a Full wwPDB EM 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/EMValidationReportHelp>
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>.

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

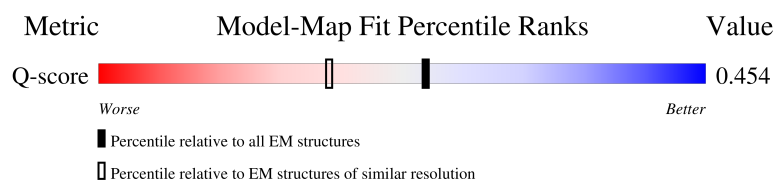
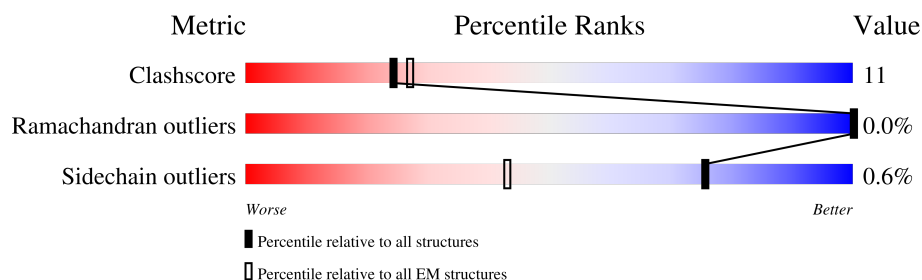
EMDB validation analysis : 0.0.1.dev132
Mogul : 2022.3.0, CSD as543be (2022)
MolProbity : 4-5-2 with Phenix2.0
Buster-report : wwPDB partial adaption of 1.1.7 (2018)
Percentile statistics : 20250101.v01 (using entries in the PDB archive January 1st 2025)
EM percentile statistics : 202505.v01 (Using data in the EMD archive up until May 2025)
MapQ : 1.9.13
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

The following experimental techniques were used to determine the structure:
ELECTRON MICROSCOPY

The reported resolution of this entry is 3.13 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)	EM structures (#Entries)	Similar EM resolution (#Entries, resolution range(Å))
Clashscore	229148	23984	-
Ramachandran outliers	224038	23583	-
Sidechain outliers	223484	23102	-
Q-score	-	25397	14478 (2.63 - 3.63)

The table below summarises the geometric issues observed across the polymeric chains and their fit to the map. The red, orange, yellow and green segments of the 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 $\leq 5\%$. The upper red bar (where present) indicates the fraction of residues that have poor fit to the EM map (all-atom inclusion $< 40\%$). The numeric value is given above the bar.

Mol	Chain	Length	Quality of chain
1	A	4967	 5% 65% 20% 15%
1	B	4967	 5% 65% 20% 15%
1	C	4967	 5% 65% 20% 15%
1	D	4967	 5% 65% 20% 15%

Continued on next page...

Continued from previous page...

Mol	Chain	Length	Quality of chain
2	E	108	 77%21%..
2	F	108	 73%24%..
2	G	108	 75%22%..
2	H	108	 74%24%..

2 Entry composition [i](#)

There are 4 unique types of molecules in this entry. The entry contains 138620 atoms, of which 0 are hydrogens and 0 are deuteriums.

In the tables below, 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 protein called Ryanodine receptor 2.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	A	4224	Total	C	N	O	S	2	0
			33774	21521	5743	6280	230		
1	B	4224	Total	C	N	O	S	2	0
			33774	21521	5743	6280	230		
1	C	4224	Total	C	N	O	S	2	0
			33774	21521	5743	6280	230		
1	D	4224	Total	C	N	O	S	2	0
			33774	21521	5743	6280	230		

There are 4 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
A	420	TRP	ARG	variant	UNP Q92736
B	420	TRP	ARG	variant	UNP Q92736
C	420	TRP	ARG	variant	UNP Q92736
D	420	TRP	ARG	variant	UNP Q92736

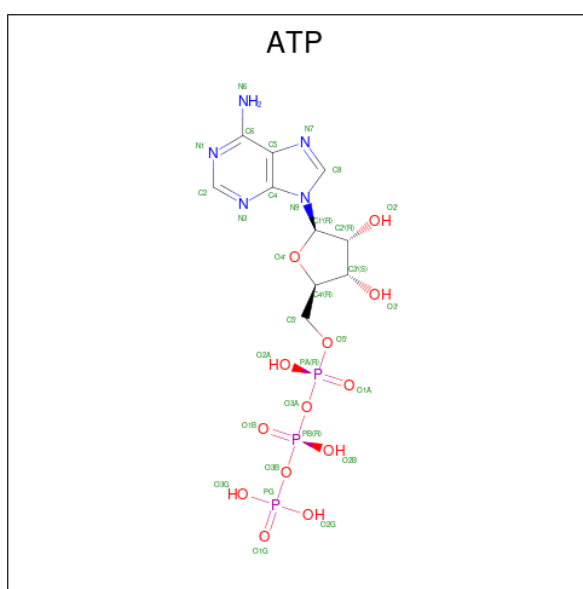
- Molecule 2 is a protein called Peptidyl-prolyl cis-trans isomerase FKBP1B.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	E	107	Total	C	N	O	S	0	0
			818	516	144	154	4		
2	F	107	Total	C	N	O	S	0	0
			818	516	144	154	4		
2	G	107	Total	C	N	O	S	0	0
			818	516	144	154	4		
2	H	107	Total	C	N	O	S	0	0
			818	516	144	154	4		

- Molecule 3 is ZINC ION (CCD ID: ZN) (formula: Zn) (labeled as "Ligand of Interest" by depositor).

Mol	Chain	Residues	Atoms		AltConf
3	A	1	Total	Zn	0
			1	1	
3	B	1	Total	Zn	0
			1	1	
3	C	1	Total	Zn	0
			1	1	
3	D	1	Total	Zn	0
			1	1	

- Molecule 4 is ADENOSINE-5'-TRIPHOSPHATE (CCD ID: ATP) (formula: $C_{10}H_{16}N_5O_{13}P_3$) (labeled as "Ligand of Interest" by depositor).



Mol	Chain	Residues	Atoms					AltConf
4	A	1	Total	C	N	O	P	0
			31	10	5	13	3	
4	A	1	Total	C	N	O	P	0
			31	10	5	13	3	
4	B	1	Total	C	N	O	P	0
			31	10	5	13	3	
4	B	1	Total	C	N	O	P	0
			31	10	5	13	3	
4	C	1	Total	C	N	O	P	0
			31	10	5	13	3	
4	C	1	Total	C	N	O	P	0
			31	10	5	13	3	
4	D	1	Total	C	N	O	P	0
			31	10	5	13	3	

Continued on next page...

Continued from previous page...

Mol	Chain	Residues	Atoms					AltConf
			Total	C	N	O	P	
4	D	1	31	10	5	13	3	0

3 Residue-property plots

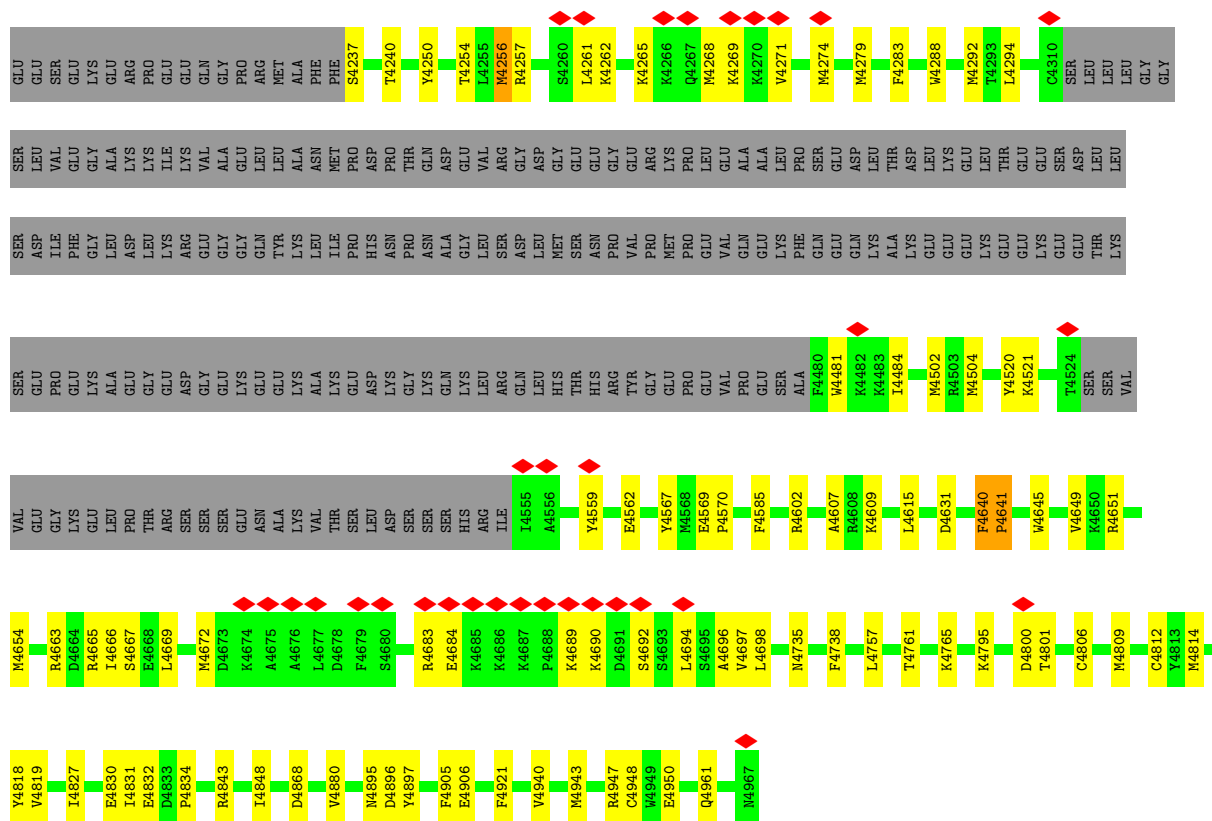
These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and atom inclusion in map 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 diamond above a residue indicates a poor fit to the EM map for this residue (all-atom inclusion < 40%). 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: Ryanodine receptor 2

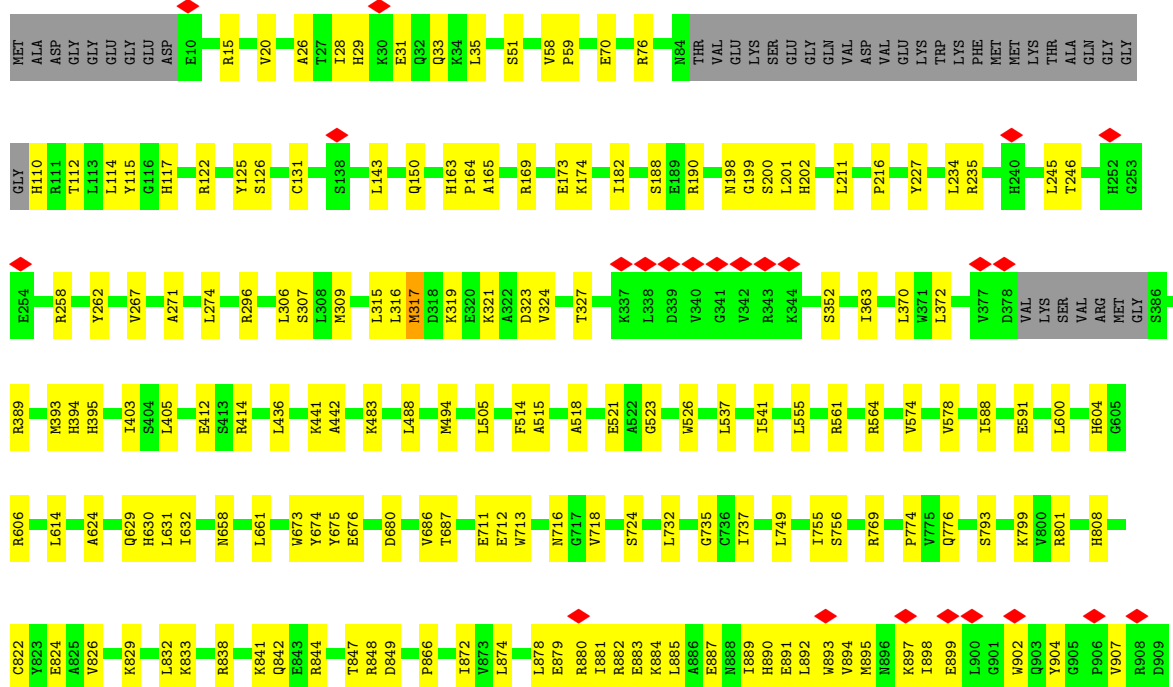


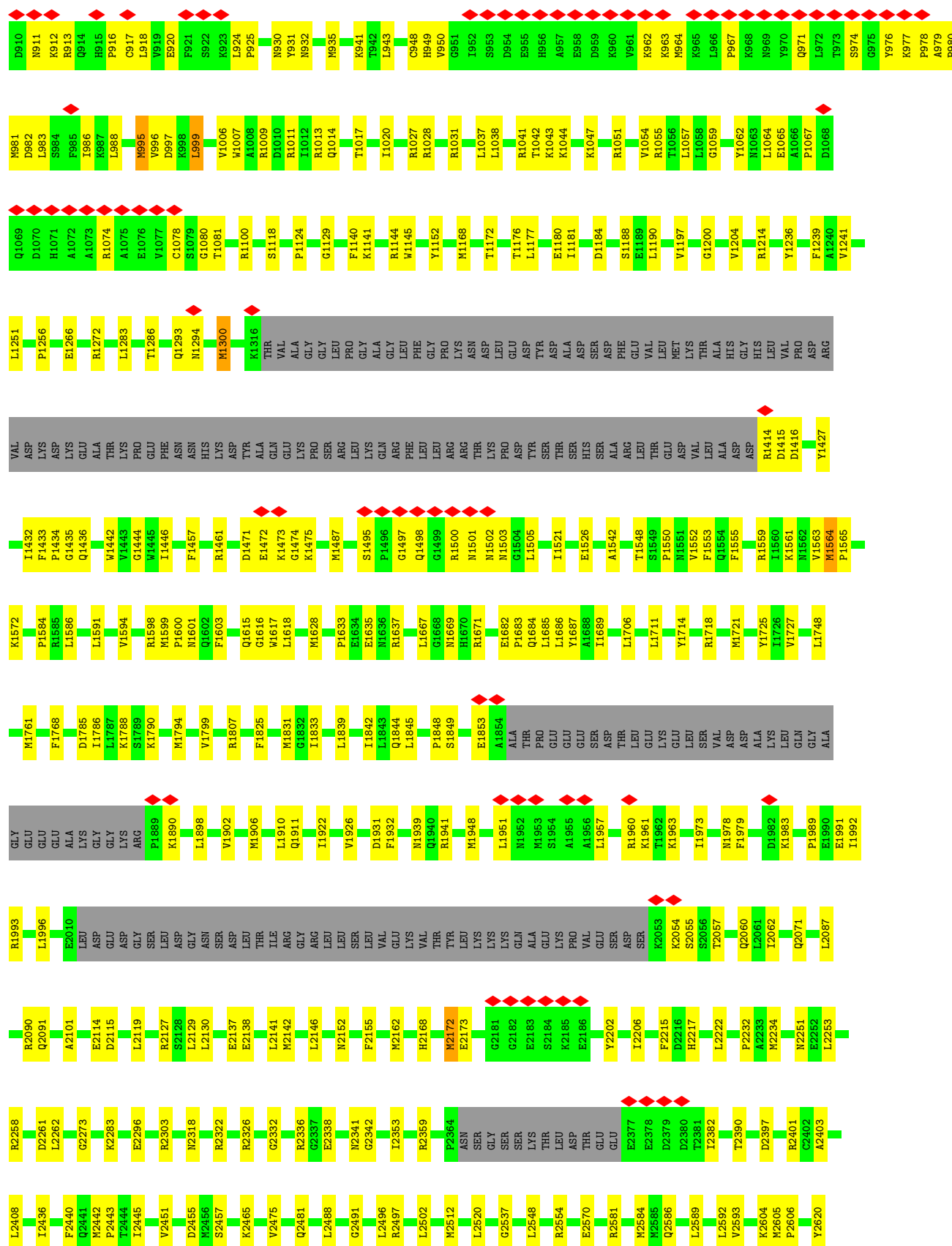




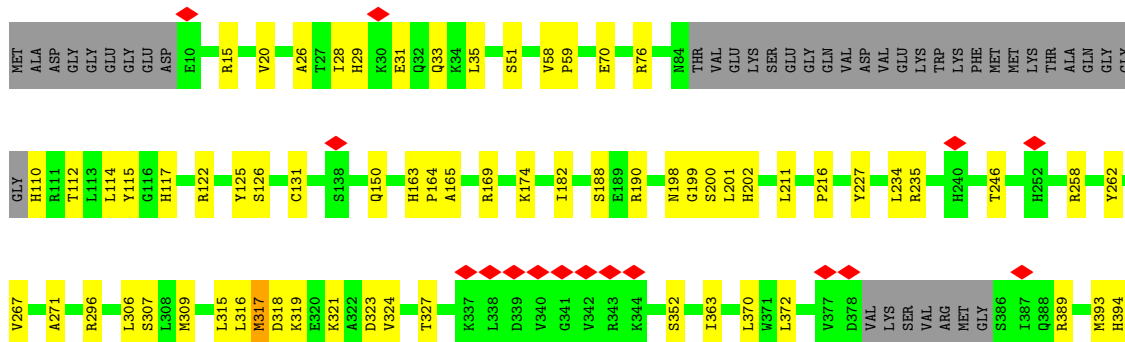


• Molecule 1: Ryanodine receptor 2











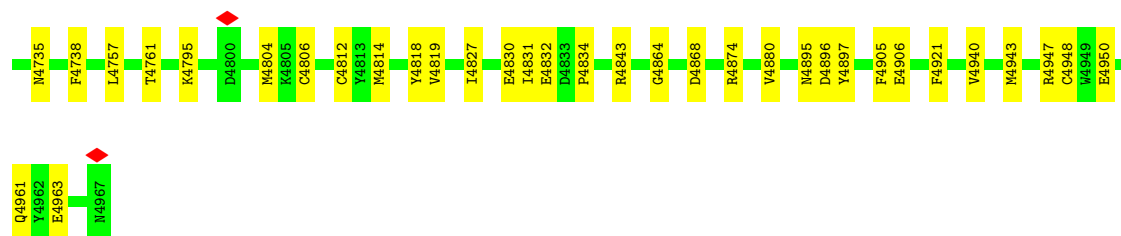






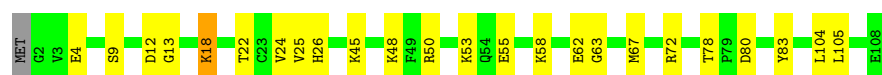






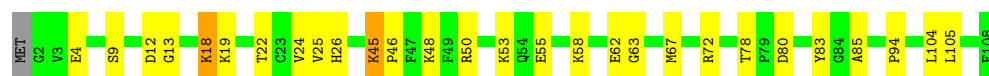
- Molecule 2: Peptidyl-prolyl cis-trans isomerase FKBP1B

Chain E: 77% 21% ..



- Molecule 2: Peptidyl-prolyl cis-trans isomerase FKBP1B

Chain F: 73% 24% ..



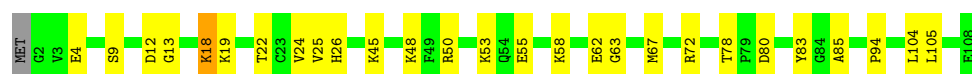
- Molecule 2: Peptidyl-prolyl cis-trans isomerase FKBP1B

Chain G: 75% 22% ..



- Molecule 2: Peptidyl-prolyl cis-trans isomerase FKBP1B

Chain H: 74% 24% ..



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C4	Depositor
Number of particles used	102478	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	58	Depositor
Minimum defocus (nm)	500	Depositor
Maximum defocus (nm)	1200	Depositor
Magnification	Not provided	
Image detector	GATAN K3 BIOQUANTUM (6k x 4k)	Depositor
Maximum map value	0.646	Depositor
Minimum map value	-0.013	Depositor
Average map value	0.011	Depositor
Map value standard deviation	0.031	Depositor
Recommended contour level	0.12	Depositor
Map size (Å)	430.848, 430.848, 430.848	wwPDB
Map dimensions	512, 512, 512	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	0.8415, 0.8415, 0.8415	Depositor

5 Model quality

5.1 Standard geometry

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

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	A	0.19	0/34516	0.41	4/46623 (0.0%)
1	B	0.19	0/34516	0.41	4/46623 (0.0%)
1	C	0.19	0/34516	0.41	4/46623 (0.0%)
1	D	0.19	0/34516	0.41	4/46623 (0.0%)
2	E	0.20	0/834	0.41	0/1123
2	F	0.20	0/834	0.41	0/1123
2	G	0.20	0/834	0.41	0/1123
2	H	0.20	0/834	0.41	0/1123
All	All	0.19	0/141400	0.41	16/190984 (0.0%)

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
1	A	0	2
1	B	0	2
1	C	0	2
1	D	0	2
All	All	0	8

There are no bond length outliers.

All (16) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	3215	MET	CB-CG-SD	5.48	129.15	112.70
1	C	3215	MET	CB-CG-SD	5.48	129.14	112.70
1	B	3215	MET	CB-CG-SD	5.47	129.12	112.70
1	D	3215	MET	CB-CG-SD	5.47	129.12	112.70

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	3310	VAL	CA-C-N	5.12	128.46	120.68
1	A	3310	VAL	C-N-CA	5.12	128.46	120.68
1	B	3310	VAL	CA-C-N	5.09	128.43	120.68
1	B	3310	VAL	C-N-CA	5.09	128.43	120.68
1	C	2884	LYS	CA-CB-CG	5.08	124.26	114.10
1	A	2884	LYS	CA-CB-CG	5.08	124.26	114.10
1	D	3310	VAL	CA-C-N	5.08	128.40	120.68
1	D	3310	VAL	C-N-CA	5.08	128.40	120.68
1	C	3310	VAL	CA-C-N	5.07	128.38	120.68
1	C	3310	VAL	C-N-CA	5.07	128.38	120.68
1	D	2884	LYS	CA-CB-CG	5.06	124.22	114.10
1	B	2884	LYS	CA-CB-CG	5.05	124.21	114.10

There are no chirality outliers.

All (8) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	A	3926	GLY	Peptide
1	A	4640	PHE	Peptide
1	B	3926	GLY	Peptide
1	B	4640	PHE	Peptide
1	C	3926	GLY	Peptide
1	C	4640	PHE	Peptide
1	D	3926	GLY	Peptide
1	D	4640	PHE	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	A	33774	0	33452	740	0
1	B	33774	0	33452	737	0
1	C	33774	0	33452	724	0
1	D	33774	0	33452	740	0
2	E	818	0	821	15	0
2	F	818	0	821	18	0
2	G	818	0	821	17	0

Continued on next page...

Continued from previous page...

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
2	H	818	0	821	17	0
3	A	1	0	0	0	0
3	B	1	0	0	0	0
3	C	1	0	0	0	0
3	D	1	0	0	0	0
4	A	62	0	24	0	0
4	B	62	0	24	0	0
4	C	62	0	24	0	0
4	D	62	0	24	0	0
All	All	138620	0	137188	2938	0

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 11.

All (2938) 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)
1:B:3235:MET:HA	1:B:3239:LEU:HD13	1.41	1.03
1:D:3235:MET:HA	1:D:3239:LEU:HD13	1.41	1.02
1:C:3235:MET:HA	1:C:3239:LEU:HD13	1.41	1.02
1:A:3235:MET:HA	1:A:3239:LEU:HD13	1.41	1.02
1:B:1564:MET:HE3	1:B:1565:PRO:HD2	1.51	0.93
1:A:1564:MET:HE3	1:A:1565:PRO:HD2	1.51	0.91
1:C:1564:MET:HE3	1:C:1565:PRO:HD2	1.51	0.91
1:D:1564:MET:HE3	1:D:1565:PRO:HD2	1.51	0.90
1:B:3152:ARG:HH21	1:B:3236:GLU:HB3	1.39	0.88
1:C:3152:ARG:HH21	1:C:3236:GLU:HB3	1.39	0.87
1:A:3270:SER:HA	1:A:3273:MET:HE2	1.57	0.87
2:H:50:ARG:HE	2:H:53:LYS:HZ3	1.23	0.86
1:A:3152:ARG:HH21	1:A:3236:GLU:HB3	1.39	0.86
1:A:4237:SER:N	1:A:4240:THR:HG1	1.74	0.85
1:C:3270:SER:HA	1:C:3273:MET:HE2	1.57	0.85
1:D:3270:SER:HA	1:D:3273:MET:HE2	1.57	0.85
2:G:50:ARG:HE	2:G:53:LYS:HZ3	1.24	0.85
1:C:4237:SER:N	1:C:4240:THR:HG1	1.74	0.85
1:B:3270:SER:HA	1:B:3273:MET:HE2	1.57	0.84
1:D:3152:ARG:HH21	1:D:3236:GLU:HB3	1.39	0.84
1:A:2798:MET:HE1	1:B:1497:GLY:HA3	1.59	0.84
2:F:50:ARG:HE	2:F:53:LYS:HZ3	1.24	0.83
1:B:4237:SER:N	1:B:4240:THR:HG1	1.77	0.82
2:E:50:ARG:HE	2:E:53:LYS:HZ3	1.24	0.81

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:4237:SER:N	1:D:4240:THR:HG1	1.77	0.81
1:D:894:VAL:HG12	1:D:897:LYS:HZ3	1.46	0.80
1:A:3197:LEU:HD23	1:A:3199:THR:H	1.47	0.80
1:D:3197:LEU:HD23	1:D:3199:THR:H	1.47	0.80
1:B:3197:LEU:HD23	1:B:3199:THR:H	1.47	0.79
1:B:2251:ASN:HD22	1:B:3817:LEU:HD11	1.48	0.78
1:C:3197:LEU:HD23	1:C:3199:THR:H	1.47	0.78
1:D:2251:ASN:HD22	1:D:3817:LEU:HD11	1.48	0.78
1:A:2251:ASN:HD22	1:A:3817:LEU:HD11	1.48	0.78
1:C:894:VAL:HG12	1:C:897:LYS:HZ3	1.48	0.78
1:A:894:VAL:HG12	1:A:897:LYS:HZ3	1.49	0.76
1:B:4665:ARG:HH21	1:B:4666:ILE:HD13	1.50	0.76
1:C:4665:ARG:HH21	1:C:4666:ILE:HD13	1.50	0.76
1:C:924:LEU:HD12	1:C:925:PRO:HD2	1.68	0.75
1:C:2251:ASN:HD22	1:C:3817:LEU:HD11	1.48	0.75
1:D:4665:ARG:HH21	1:D:4666:ILE:HD13	1.50	0.74
1:A:924:LEU:HD12	1:A:925:PRO:HD2	1.68	0.74
1:D:924:LEU:HD12	1:D:925:PRO:HD2	1.68	0.74
1:A:4665:ARG:HH21	1:A:4666:ILE:HD13	1.50	0.74
1:D:2693:SER:OG	1:D:2704:GLN:NE2	2.21	0.73
1:B:2693:SER:OG	1:B:2704:GLN:NE2	2.21	0.73
1:A:2693:SER:OG	1:A:2704:GLN:NE2	2.21	0.73
1:C:2693:SER:OG	1:C:2704:GLN:NE2	2.21	0.73
1:B:924:LEU:HD12	1:B:925:PRO:HD2	1.68	0.73
1:B:3013:VAL:O	1:B:3018:ARG:NH2	2.22	0.73
1:D:3013:VAL:O	1:D:3018:ARG:NH2	2.22	0.73
1:A:988:LEU:HD12	1:A:1055:ARG:HD3	1.71	0.72
1:C:988:LEU:HD12	1:C:1055:ARG:HD3	1.71	0.72
1:C:3013:VAL:O	1:C:3018:ARG:NH2	2.22	0.72
1:A:3013:VAL:O	1:A:3018:ARG:NH2	2.22	0.72
1:D:3269:ASN:HB3	1:D:3272:HIS:ND1	2.05	0.72
1:A:3269:ASN:HB3	1:A:3272:HIS:ND1	2.05	0.72
1:A:309:MET:HE3	1:A:315:LEU:HB3	1.72	0.72
1:D:309:MET:HE3	1:D:315:LEU:HB3	1.71	0.72
1:B:988:LEU:HD12	1:B:1055:ARG:HD3	1.71	0.71
1:D:1941:ARG:HH21	1:D:3609:TYR:HB2	1.55	0.71
1:A:1941:ARG:HH21	1:A:3609:TYR:HB2	1.55	0.71
1:A:2488:LEU:HD21	1:A:2548:LEU:HD22	1.72	0.71
1:C:1038:LEU:O	1:C:1043:LYS:NZ	2.24	0.71
1:C:309:MET:HE3	1:C:315:LEU:HB3	1.71	0.71
1:D:988:LEU:HD12	1:D:1055:ARG:HD3	1.71	0.71

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:2488:LEU:HD21	1:D:2548:LEU:HD22	1.72	0.71
1:B:3269:ASN:HB3	1:B:3272:HIS:ND1	2.05	0.71
1:B:309:MET:HE3	1:B:315:LEU:HB3	1.72	0.71
1:A:1038:LEU:O	1:A:1043:LYS:NZ	2.24	0.71
1:C:3269:ASN:HB3	1:C:3272:HIS:ND1	2.05	0.71
1:B:894:VAL:HG12	1:B:897:LYS:HZ3	1.56	0.70
1:B:1941:ARG:HH21	1:B:3609:TYR:HB2	1.55	0.70
1:C:1941:ARG:HH21	1:C:3609:TYR:HB2	1.55	0.70
1:C:3042:ALA:HB1	1:C:3121:LEU:HD13	1.73	0.70
1:B:3042:ALA:HB1	1:B:3121:LEU:HD13	1.73	0.70
1:C:2791:ARG:HH12	1:C:2795:GLY:HA3	1.57	0.70
1:D:2791:ARG:HH12	1:D:2795:GLY:HA3	1.57	0.70
1:D:894:VAL:HA	1:D:897:LYS:HG2	1.73	0.70
1:D:2723:LYS:HG2	1:D:2895:PHE:HZ	1.56	0.70
1:B:2791:ARG:HH12	1:B:2795:GLY:HA3	1.57	0.70
1:D:769:ARG:HG2	1:D:774:PRO:HA	1.74	0.70
1:A:363:ILE:HD11	1:A:403:ILE:HD13	1.74	0.70
1:A:894:VAL:HA	1:A:897:LYS:HG2	1.73	0.70
1:A:2791:ARG:HH12	1:A:2795:GLY:HA3	1.57	0.70
1:B:1038:LEU:O	1:B:1043:LYS:NZ	2.24	0.70
1:D:1038:LEU:O	1:D:1043:LYS:NZ	2.24	0.70
1:A:4481:TRP:HA	1:A:4484:ILE:HG22	1.74	0.70
1:B:1129:GLY:HA3	1:B:1145:TRP:HB3	1.74	0.70
1:B:2979:ARG:HH22	1:B:2983:LEU:HD22	1.57	0.69
1:B:2723:LYS:HG2	1:B:2895:PHE:HZ	1.56	0.69
1:A:1129:GLY:HA3	1:A:1145:TRP:HB3	1.74	0.69
1:A:3042:ALA:HB1	1:A:3121:LEU:HD13	1.73	0.69
1:B:2488:LEU:HD21	1:B:2548:LEU:HD22	1.72	0.69
1:A:2979:ARG:HH22	1:A:2983:LEU:HD22	1.57	0.69
1:C:769:ARG:HG2	1:C:774:PRO:HA	1.74	0.69
1:C:2488:LEU:HD21	1:C:2548:LEU:HD22	1.72	0.69
1:D:3042:ALA:HB1	1:D:3121:LEU:HD13	1.73	0.69
2:G:22:THR:HG22	2:G:50:ARG:HG2	1.74	0.69
1:B:769:ARG:HG2	1:B:774:PRO:HA	1.74	0.69
1:A:2723:LYS:HG2	1:A:2895:PHE:HZ	1.56	0.69
1:B:363:ILE:HD11	1:B:403:ILE:HD13	1.74	0.69
1:C:2723:LYS:HG2	1:C:2895:PHE:HZ	1.56	0.69
1:C:4481:TRP:HA	1:C:4484:ILE:HG22	1.74	0.69
1:C:894:VAL:HA	1:C:897:LYS:HG2	1.73	0.69
1:C:2604:LYS:NZ	1:C:2660:GLU:OE1	2.26	0.69
2:H:22:THR:HG22	2:H:50:ARG:HG2	1.74	0.69

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:4481:TRP:HA	1:D:4484:ILE:HG22	1.74	0.68
1:B:894:VAL:HA	1:B:897:LYS:HG2	1.73	0.68
1:B:2604:LYS:NZ	1:B:2660:GLU:OE1	2.26	0.68
1:B:3231:MET:HE3	1:B:3233:HIS:HB2	1.74	0.68
1:C:2979:ARG:HH22	1:C:2983:LEU:HD22	1.57	0.68
1:D:363:ILE:HD11	1:D:403:ILE:HD13	1.74	0.68
2:E:22:THR:HG22	2:E:50:ARG:HG2	1.74	0.68
1:B:3650:GLU:OE1	1:B:3660:ARG:NH1	2.27	0.68
1:B:4481:TRP:HA	1:B:4484:ILE:HG22	1.74	0.68
1:C:1129:GLY:HA3	1:C:1145:TRP:HB3	1.74	0.68
1:C:3231:MET:HE3	1:C:3233:HIS:HB2	1.74	0.68
1:A:3231:MET:HE3	1:A:3233:HIS:HB2	1.74	0.68
1:B:2627:TRP:HB2	1:B:2630:PHE:HB2	1.76	0.68
1:C:363:ILE:HD11	1:C:403:ILE:HD13	1.74	0.68
1:D:2604:LYS:NZ	1:D:2660:GLU:OE1	2.26	0.68
1:D:2979:ARG:HH22	1:D:2983:LEU:HD22	1.57	0.68
1:D:3650:GLU:OE1	1:D:3660:ARG:NH1	2.27	0.68
1:A:3650:GLU:OE1	1:A:3660:ARG:NH1	2.27	0.68
1:B:363:ILE:HD13	1:B:372:LEU:HD23	1.75	0.68
1:C:3650:GLU:OE1	1:C:3660:ARG:NH1	2.27	0.68
1:D:1129:GLY:HA3	1:D:1145:TRP:HB3	1.74	0.68
1:D:3778:LEU:HD13	1:D:3854:PHE:HD1	1.59	0.68
2:F:22:THR:HG22	2:F:50:ARG:HG2	1.74	0.68
1:A:769:ARG:HG2	1:A:774:PRO:HA	1.74	0.68
1:D:1635:GLU:OE1	1:D:1637:ARG:NH1	2.28	0.67
1:C:2627:TRP:HB2	1:C:2630:PHE:HB2	1.76	0.67
1:D:2882:LYS:HD2	1:D:2886:ARG:HH21	1.60	0.67
1:A:2627:TRP:HB2	1:A:2630:PHE:HB2	1.76	0.67
1:B:4654:MET:HE2	1:B:4663:ARG:HH11	1.60	0.67
1:C:3778:LEU:HD13	1:C:3854:PHE:HD1	1.59	0.67
1:A:4694:LEU:HD11	1:D:4268:MET:HG3	1.76	0.67
1:C:363:ILE:HD13	1:C:372:LEU:HD23	1.75	0.67
1:C:1009:ARG:HB2	1:C:1013:ARG:HH12	1.60	0.67
1:D:3231:MET:HE3	1:D:3233:HIS:HB2	1.74	0.67
1:A:1635:GLU:OE1	1:A:1637:ARG:NH1	2.28	0.67
1:A:2604:LYS:NZ	1:A:2660:GLU:OE1	2.26	0.67
1:D:1009:ARG:HB2	1:D:1013:ARG:HH12	1.60	0.67
1:B:1009:ARG:HB2	1:B:1013:ARG:HH12	1.60	0.67
1:B:1433:PHE:O	1:B:1500:ARG:NH2	2.28	0.67
1:B:3778:LEU:HD13	1:B:3854:PHE:HD1	1.59	0.67
1:D:363:ILE:HD13	1:D:372:LEU:HD23	1.75	0.67

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:4654:MET:HE2	1:D:4663:ARG:HH11	1.59	0.67
1:A:363:ILE:HD13	1:A:372:LEU:HD23	1.75	0.67
1:B:1635:GLU:OE1	1:B:1637:ARG:NH1	2.28	0.67
1:A:2882:LYS:HD2	1:A:2886:ARG:HH21	1.60	0.66
1:C:1635:GLU:OE1	1:C:1637:ARG:NH1	2.28	0.66
1:C:2882:LYS:HD2	1:C:2886:ARG:HH21	1.60	0.66
1:D:1433:PHE:O	1:D:1500:ARG:NH2	2.28	0.66
1:A:1009:ARG:HB2	1:A:1013:ARG:HH12	1.60	0.66
1:A:3778:LEU:HD13	1:A:3854:PHE:HD1	1.59	0.66
1:C:1433:PHE:O	1:C:1500:ARG:NH2	2.28	0.66
1:B:2882:LYS:HD2	1:B:2886:ARG:HH21	1.60	0.66
1:C:904:TYR:HB2	1:C:918:LEU:HD22	1.77	0.66
1:C:4072:THR:HG22	1:C:4078:LEU:HB3	1.78	0.66
1:D:4072:THR:HG22	1:D:4078:LEU:HB3	1.78	0.66
1:A:4484:ILE:HG13	1:D:4279:MET:HE2	1.78	0.66
1:B:4268:MET:HA	1:B:4271:VAL:HG22	1.77	0.66
1:A:2830:ASN:HB2	1:B:1434:PRO:O	1.96	0.66
1:A:1433:PHE:O	1:A:1500:ARG:NH2	2.28	0.66
1:A:4654:MET:HE2	1:A:4663:ARG:HH11	1.60	0.66
1:A:125:TYR:O	1:A:414:ARG:NH1	2.29	0.66
1:B:963:LYS:HZ1	1:B:977:LYS:HD3	1.61	0.66
1:C:4654:MET:HE2	1:C:4663:ARG:HH11	1.59	0.66
1:D:904:TYR:HB2	1:D:918:LEU:HD22	1.77	0.66
1:D:2627:TRP:HB2	1:D:2630:PHE:HB2	1.76	0.66
1:A:1768:PHE:O	2:E:83:TYR:OH	2.13	0.65
1:B:2657:TYR:OH	1:B:2663:LYS:NZ	2.30	0.65
1:B:4072:THR:HG22	1:B:4078:LEU:HB3	1.78	0.65
1:C:2657:TYR:OH	1:C:2663:LYS:NZ	2.30	0.65
1:C:4268:MET:HA	1:C:4271:VAL:HG22	1.77	0.65
1:A:904:TYR:HB2	1:A:918:LEU:HD22	1.77	0.65
1:A:4072:THR:HG22	1:A:4078:LEU:HB3	1.78	0.65
1:D:2657:TYR:OH	1:D:2663:LYS:NZ	2.30	0.65
1:A:1503:ASN:HB2	1:D:2824:ARG:HH22	1.60	0.65
1:A:2657:TYR:OH	1:A:2663:LYS:NZ	2.30	0.65
1:B:904:TYR:HB2	1:B:918:LEU:HD22	1.77	0.65
1:B:2860:LEU:HD11	1:B:2867:ASN:HA	1.79	0.65
1:C:2860:LEU:HD11	1:C:2867:ASN:HA	1.79	0.65
1:C:963:LYS:HZ1	1:C:977:LYS:HD3	1.61	0.65
1:D:125:TYR:O	1:D:414:ARG:NH1	2.29	0.65
1:A:3025:ASP:O	1:A:3028:SER:OG	2.14	0.65
1:D:881:ILE:HD12	1:D:884:LYS:HZ3	1.62	0.65

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3184:TYR:O	1:C:3192:ARG:NH2	2.30	0.65
1:D:963:LYS:HZ1	1:D:977:LYS:HD3	1.62	0.65
1:D:3025:ASP:O	1:D:3028:SER:OG	2.14	0.65
1:A:4268:MET:HA	1:A:4271:VAL:HG22	1.77	0.65
1:D:1432:ILE:HG22	1:D:1500:ARG:HH21	1.62	0.65
1:B:1432:ILE:HG22	1:B:1500:ARG:HH21	1.62	0.65
2:H:4:GLU:OE1	2:H:4:GLU:N	2.30	0.65
1:B:3179:ASN:O	1:B:3185:ASN:ND2	2.30	0.65
1:C:125:TYR:O	1:C:414:ARG:NH1	2.29	0.65
1:D:3184:TYR:O	1:D:3192:ARG:NH2	2.30	0.65
1:A:2232:PRO:HG3	1:A:2382:ILE:HD11	1.79	0.64
1:A:3184:TYR:O	1:A:3192:ARG:NH2	2.30	0.64
1:C:1911:GLN:OE1	1:C:2090:ARG:NH1	2.31	0.64
1:D:2860:LEU:HD11	1:D:2867:ASN:HA	1.78	0.64
1:B:3166:PHE:HE2	1:B:3168:VAL:HB	1.62	0.64
1:C:3166:PHE:HE2	1:C:3168:VAL:HB	1.62	0.64
1:C:4690:LYS:HG3	1:C:4692:SER:H	1.63	0.64
1:D:1911:GLN:OE1	1:D:2090:ARG:NH1	2.31	0.64
1:A:3166:PHE:HE2	1:A:3168:VAL:HB	1.62	0.64
1:B:2773:TRP:HB3	1:B:2774:PRO:HD3	1.79	0.64
1:C:2773:TRP:HB3	1:C:2774:PRO:HD3	1.79	0.64
1:C:3025:ASP:O	1:C:3028:SER:OG	2.14	0.64
1:C:3179:ASN:O	1:C:3185:ASN:ND2	2.30	0.64
1:A:1432:ILE:HG22	1:A:1500:ARG:HH21	1.62	0.64
1:A:1911:GLN:OE1	1:A:2090:ARG:NH1	2.31	0.64
1:A:2860:LEU:HD11	1:A:2867:ASN:HA	1.79	0.64
1:A:963:LYS:HZ1	1:A:977:LYS:HD3	1.62	0.64
1:B:125:TYR:O	1:B:414:ARG:NH1	2.29	0.64
1:D:2884:LYS:HD2	1:D:2884:LYS:C	2.23	0.64
1:D:3227:ARG:HD3	1:D:3229:THR:H	1.63	0.64
1:D:4268:MET:HA	1:D:4271:VAL:HG22	1.77	0.64
1:A:3179:ASN:O	1:A:3185:ASN:ND2	2.30	0.64
1:B:1911:GLN:OE1	1:B:2090:ARG:NH1	2.31	0.64
1:B:2884:LYS:HD2	1:B:2884:LYS:C	2.23	0.64
1:C:1432:ILE:HG22	1:C:1500:ARG:HH21	1.62	0.64
2:G:4:GLU:OE1	2:G:4:GLU:N	2.30	0.64
1:B:4690:LYS:HG3	1:B:4692:SER:H	1.63	0.64
1:C:2232:PRO:HG3	1:C:2382:ILE:HD11	1.79	0.64
1:A:3227:ARG:HD3	1:A:3229:THR:H	1.63	0.63
1:A:4262:LYS:HG3	1:B:4698:LEU:HD13	1.78	0.63
1:D:2773:TRP:HB3	1:D:2774:PRO:HD3	1.79	0.63

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:3901:GLN:O	1:B:3905:ASN:ND2	2.28	0.63
1:C:4268:MET:HG3	1:D:4694:LEU:HD11	1.81	0.63
1:D:3179:ASN:O	1:D:3185:ASN:ND2	2.30	0.63
1:D:3166:PHE:HE2	1:D:3168:VAL:HB	1.62	0.63
2:F:4:GLU:OE1	2:F:4:GLU:N	2.29	0.63
1:C:3227:ARG:HD3	1:C:3229:THR:H	1.63	0.63
1:C:3127:GLN:O	1:C:3131:TYR:HD2	1.82	0.63
1:D:2232:PRO:HG3	1:D:2382:ILE:HD11	1.79	0.63
1:A:3127:GLN:O	1:A:3131:TYR:HD2	1.82	0.63
1:B:3127:GLN:O	1:B:3131:TYR:HD2	1.82	0.63
1:D:4690:LYS:HG3	1:D:4692:SER:H	1.63	0.63
2:G:26:HIS:CD2	2:G:105:LEU:HD11	2.34	0.63
1:A:370:LEU:HB2	1:A:393:MET:HE3	1.81	0.63
1:A:2773:TRP:HB3	1:A:2774:PRO:HD3	1.79	0.63
1:A:2884:LYS:HD2	1:A:2884:LYS:C	2.23	0.63
1:A:4690:LYS:HG3	1:A:4692:SER:H	1.63	0.63
2:E:26:HIS:CD2	2:E:105:LEU:HD11	2.34	0.63
1:A:3797:MET:HE1	1:A:3870:ILE:HG23	1.81	0.63
1:B:2232:PRO:HG3	1:B:2382:ILE:HD11	1.79	0.63
1:D:3311:LYS:HB2	1:D:3314:LEU:HD13	1.80	0.63
1:A:2593:VAL:HG12	1:A:2644:LEU:HB2	1.81	0.62
1:B:3184:TYR:O	1:B:3192:ARG:NH2	2.30	0.62
1:C:3797:MET:HE1	1:C:3870:ILE:HG23	1.81	0.62
1:D:3127:GLN:O	1:D:3131:TYR:HD2	1.82	0.62
1:B:3797:MET:HE1	1:B:3870:ILE:HG23	1.81	0.62
1:C:2884:LYS:HD2	1:C:2884:LYS:C	2.23	0.62
1:D:2273:GLY:O	1:D:2336:ARG:NH2	2.32	0.62
1:D:3797:MET:HE1	1:D:3870:ILE:HG23	1.81	0.62
2:F:26:HIS:CD2	2:F:105:LEU:HD11	2.34	0.62
1:A:2273:GLY:O	1:A:2336:ARG:NH2	2.32	0.62
1:A:4268:MET:HG3	1:B:4694:LEU:HD11	1.81	0.62
2:E:4:GLU:N	2:E:4:GLU:OE1	2.29	0.62
1:B:3134:LEU:HB3	1:B:3162:PHE:CE2	2.35	0.62
1:C:872:ILE:O	1:C:941:LYS:NZ	2.32	0.62
1:A:872:ILE:O	1:A:941:LYS:NZ	2.32	0.62
1:A:2586:GLN:NE2	1:A:2636:GLU:OE1	2.33	0.62
1:D:2593:VAL:HG12	1:D:2644:LEU:HB2	1.81	0.62
1:D:3325:LYS:HE3	1:D:3328:LYS:HD2	1.81	0.62
1:A:3134:LEU:HB3	1:A:3162:PHE:CE2	2.35	0.62
1:A:3325:LYS:HE3	1:A:3328:LYS:HD2	1.81	0.62
1:B:3311:LYS:HB2	1:B:3314:LEU:HD13	1.80	0.62

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2824:ARG:HH22	1:C:1503:ASN:HB2	1.65	0.62
1:B:3325:LYS:HE3	1:B:3328:LYS:HD2	1.81	0.62
1:C:3311:LYS:HB2	1:C:3314:LEU:HD13	1.80	0.62
1:A:3311:LYS:HB2	1:A:3314:LEU:HD13	1.80	0.62
1:B:3025:ASP:O	1:B:3028:SER:OG	2.14	0.62
2:H:26:HIS:CD2	2:H:105:LEU:HD11	2.34	0.62
1:C:309:MET:HE1	1:C:317:MET:SD	2.40	0.62
1:C:370:LEU:HB2	1:C:393:MET:HE3	1.81	0.62
1:C:3134:LEU:HB3	1:C:3162:PHE:CE2	2.35	0.62
1:D:370:LEU:HB2	1:D:393:MET:HE3	1.81	0.62
1:D:872:ILE:O	1:D:941:LYS:NZ	2.32	0.62
1:A:309:MET:HE1	1:A:317:MET:SD	2.40	0.62
1:C:2593:VAL:HG12	1:C:2644:LEU:HB2	1.81	0.62
1:D:309:MET:HE1	1:D:317:MET:SD	2.40	0.62
1:D:3043:ARG:NE	1:D:3120:ASP:OD2	2.27	0.62
1:D:3134:LEU:HB3	1:D:3162:PHE:CE2	2.35	0.62
1:B:309:MET:HE1	1:B:317:MET:SD	2.40	0.61
1:B:872:ILE:O	1:B:941:LYS:NZ	2.32	0.61
1:B:2581:ARG:HD3	1:B:2584:MET:HE3	1.82	0.61
1:B:3227:ARG:HD3	1:B:3229:THR:H	1.63	0.61
1:C:3187:LYS:HZ3	1:C:3191:GLU:HB3	1.64	0.61
1:C:3325:LYS:HE3	1:C:3328:LYS:HD2	1.81	0.61
1:B:370:LEU:HB2	1:B:393:MET:HE3	1.81	0.61
1:B:1168:MET:HE2	1:B:1197:VAL:HG13	1.82	0.61
1:C:2273:GLY:O	1:C:2336:ARG:NH2	2.32	0.61
1:A:2581:ARG:HD3	1:A:2584:MET:HE3	1.82	0.61
1:B:3016:ARG:HG2	1:B:3017:HIS:CD2	2.36	0.61
1:C:1168:MET:HE2	1:C:1197:VAL:HG13	1.82	0.61
1:D:3016:ARG:HG2	1:D:3017:HIS:CD2	2.36	0.61
1:D:3901:GLN:O	1:D:3905:ASN:ND2	2.28	0.61
1:B:895:MET:SD	1:B:971:GLN:NE2	2.74	0.61
1:B:2273:GLY:O	1:B:2336:ARG:NH2	2.32	0.61
1:C:3901:GLN:O	1:C:3905:ASN:ND2	2.28	0.61
1:A:895:MET:SD	1:A:971:GLN:NE2	2.74	0.61
1:A:4665:ARG:HH22	1:A:4669:LEU:HD22	1.66	0.61
1:D:4187:GLU:OE2	1:D:4947:ARG:NH2	2.33	0.61
1:A:3016:ARG:HG2	1:A:3017:HIS:CD2	2.36	0.61
1:B:4665:ARG:HH22	1:B:4669:LEU:HD22	1.66	0.61
1:A:2824:ARG:HH22	1:B:1503:ASN:HB2	1.65	0.61
1:B:4609:LYS:HD2	1:B:4615:LEU:HD22	1.83	0.61
1:C:895:MET:SD	1:C:971:GLN:NE2	2.74	0.61

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2586:GLN:NE2	1:C:2636:GLU:OE1	2.33	0.61
1:B:4187:GLU:OE2	1:B:4947:ARG:NH2	2.33	0.61
1:D:4609:LYS:HD2	1:D:4615:LEU:HD22	1.83	0.61
1:A:4609:LYS:HD2	1:A:4615:LEU:HD22	1.83	0.61
1:C:3016:ARG:HG2	1:C:3017:HIS:CD2	2.36	0.61
1:D:2581:ARG:HD3	1:D:2584:MET:HE3	1.82	0.61
1:B:126:SER:HA	1:B:414:ARG:HH12	1.67	0.60
1:B:4010:VAL:HG11	1:B:4118:PHE:HZ	1.66	0.60
1:B:4874:ARG:NH1	1:C:4868:ASP:OD1	2.34	0.60
1:D:2831:VAL:HG12	1:D:2894:LYS:HD2	1.83	0.60
1:B:1910:LEU:HD13	1:B:2062:ILE:HG12	1.83	0.60
1:B:2593:VAL:HG12	1:B:2644:LEU:HB2	1.81	0.60
1:C:4010:VAL:HG11	1:C:4118:PHE:HZ	1.66	0.60
1:D:3313:GLN:HA	1:D:3316:LYS:NZ	2.17	0.60
1:A:3313:GLN:HA	1:A:3316:LYS:NZ	2.17	0.60
1:B:2831:VAL:HG12	1:B:2894:LYS:HD2	1.84	0.60
1:D:895:MET:SD	1:D:971:GLN:NE2	2.74	0.60
1:D:1168:MET:HE2	1:D:1197:VAL:HG13	1.82	0.60
1:C:4187:GLU:OE2	1:C:4947:ARG:NH2	2.33	0.60
1:C:1239:PHE:O	1:C:1807:ARG:NH2	2.35	0.60
1:C:4665:ARG:HH22	1:C:4669:LEU:HD22	1.66	0.60
1:D:2586:GLN:NE2	1:D:2636:GLU:OE1	2.33	0.60
1:D:3697:LYS:HA	1:D:3700:HIS:CD2	2.37	0.60
1:A:3059:ALA:O	1:A:3063:ASN:ND2	2.35	0.60
1:A:4010:VAL:HG11	1:A:4118:PHE:HZ	1.66	0.60
1:C:3137:LEU:HB2	1:C:3159:LEU:HD13	1.84	0.60
1:C:3313:GLN:HA	1:C:3316:LYS:NZ	2.17	0.60
1:D:1239:PHE:O	1:D:1807:ARG:NH2	2.35	0.60
1:A:126:SER:HA	1:A:414:ARG:HH12	1.66	0.60
1:A:1168:MET:HE2	1:A:1197:VAL:HG13	1.82	0.60
1:C:2831:VAL:HG12	1:C:2894:LYS:HD2	1.84	0.60
1:B:4279:MET:HE2	1:C:4484:ILE:HG13	1.82	0.60
1:C:2202:TYR:O	1:C:2206:ILE:HG12	2.02	0.60
1:C:2581:ARG:HD3	1:C:2584:MET:HE3	1.82	0.60
1:C:3313:GLN:HA	1:C:3316:LYS:HZ1	1.67	0.60
1:C:3697:LYS:HA	1:C:3700:HIS:CD2	2.36	0.60
1:D:2296:GLU:HG3	1:D:2390:THR:HG22	1.83	0.60
1:A:2831:VAL:HG12	1:A:2894:LYS:HD2	1.84	0.60
1:A:3697:LYS:HA	1:A:3700:HIS:CD2	2.37	0.60
1:B:2691:LYS:O	1:B:2694:SER:OG	2.17	0.60
1:C:2890:GLN:O	1:C:2894:LYS:HG2	2.02	0.60

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4609:LYS:HD2	1:C:4615:LEU:HD22	1.83	0.60
1:D:1251:LEU:HD23	1:D:1599:MET:HE2	1.84	0.60
1:A:1251:LEU:HD23	1:A:1599:MET:HE2	1.84	0.59
1:A:2202:TYR:O	1:A:2206:ILE:HG12	2.02	0.59
1:A:4187:GLU:OE2	1:A:4947:ARG:NH2	2.33	0.59
1:B:2586:GLN:NE2	1:B:2636:GLU:OE1	2.33	0.59
1:B:3313:GLN:HA	1:B:3316:LYS:NZ	2.17	0.59
1:C:126:SER:HA	1:C:414:ARG:HH12	1.67	0.59
1:A:1054:VAL:HA	1:A:1057:LEU:HD12	1.84	0.59
1:A:3137:LEU:HB2	1:A:3159:LEU:HD13	1.84	0.59
1:B:2202:TYR:O	1:B:2206:ILE:HG12	2.02	0.59
1:C:1910:LEU:HD13	1:C:2062:ILE:HG12	1.83	0.59
1:D:2202:TYR:O	1:D:2206:ILE:HG12	2.02	0.59
1:A:1239:PHE:O	1:A:1807:ARG:NH2	2.35	0.59
1:B:2890:GLN:O	1:B:2894:LYS:HG2	2.02	0.59
1:B:3697:LYS:HA	1:B:3700:HIS:CD2	2.36	0.59
1:B:4268:MET:HG3	1:C:4694:LEU:HD11	1.83	0.59
1:D:4665:ARG:HH22	1:D:4669:LEU:HD22	1.66	0.59
1:D:4689:LYS:HZ3	1:D:4696:ALA:HB2	1.66	0.59
1:A:2296:GLU:HG3	1:A:2390:THR:HG22	1.83	0.59
1:C:892:LEU:HD21	1:C:980:PRO:HD3	1.85	0.59
1:D:561:ARG:HB3	1:D:564:ARG:HD2	1.84	0.59
1:D:2436:ILE:HA	1:D:2465:LYS:HD3	1.85	0.59
1:B:3152:ARG:NH2	1:B:3236:GLU:HB3	2.16	0.59
1:C:949:HIS:ND1	1:C:1065:GLU:OE2	2.34	0.59
1:C:3288:LEU:HD11	1:C:3328:LYS:HD3	1.85	0.59
1:C:4010:VAL:HG11	1:C:4118:PHE:CZ	2.38	0.59
1:A:2436:ILE:HA	1:A:2465:LYS:HD3	1.85	0.59
1:A:2937:HIS:HB2	1:A:3014:LEU:HD21	1.85	0.59
1:B:1239:PHE:O	1:B:1807:ARG:NH2	2.35	0.59
1:C:3280:ILE:O	1:C:3284:ILE:HG12	2.03	0.59
1:A:188:SER:HB2	1:A:190:ARG:HH11	1.67	0.59
1:B:188:SER:HB2	1:B:190:ARG:HH11	1.67	0.59
1:B:561:ARG:HB3	1:B:564:ARG:HD2	1.84	0.59
1:B:3043:ARG:NE	1:B:3120:ASP:OD2	2.27	0.59
1:B:3175:LEU:HD12	1:B:3178:HIS:HD2	1.68	0.59
1:D:892:LEU:HD21	1:D:980:PRO:HD3	1.85	0.59
1:D:4010:VAL:HG11	1:D:4118:PHE:HZ	1.66	0.59
1:A:1910:LEU:HD13	1:A:2062:ILE:HG12	1.83	0.59
1:B:2436:ILE:HA	1:B:2465:LYS:HD3	1.85	0.59
1:B:3137:LEU:HB2	1:B:3159:LEU:HD13	1.84	0.59

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:3280:ILE:O	1:B:3284:ILE:HG12	2.03	0.59
1:C:2296:GLU:HG3	1:C:2390:THR:HG22	1.83	0.59
1:D:591:GLU:HG3	1:D:631:LEU:HD22	1.85	0.59
1:D:1910:LEU:HD13	1:D:2062:ILE:HG12	1.83	0.59
1:A:591:GLU:HG3	1:A:631:LEU:HD22	1.85	0.59
1:A:4237:SER:N	1:A:4240:THR:OG1	2.36	0.59
1:B:892:LEU:HD21	1:B:980:PRO:HD3	1.85	0.59
1:B:4010:VAL:HG11	1:B:4118:PHE:CZ	2.38	0.59
1:D:188:SER:HB2	1:D:190:ARG:HH11	1.67	0.59
1:A:561:ARG:HB3	1:A:564:ARG:HD2	1.84	0.59
1:A:881:ILE:HD12	1:A:884:LYS:HZ3	1.66	0.59
1:A:892:LEU:HD21	1:A:980:PRO:HD3	1.85	0.59
1:A:1825:PHE:CE1	1:A:1842:ILE:HG12	2.38	0.59
1:A:2890:GLN:O	1:A:2894:LYS:HG2	2.02	0.59
1:B:1009:ARG:HB2	1:B:1013:ARG:NH1	2.18	0.59
1:B:1100:ARG:HB3	1:B:1236:TYR:HA	1.85	0.59
1:B:2296:GLU:HG3	1:B:2390:THR:HG22	1.83	0.59
1:D:1009:ARG:HB2	1:D:1013:ARG:NH1	2.18	0.59
1:D:4271:VAL:HB	1:D:4274:MET:SD	2.43	0.59
1:A:15:ARG:NH2	1:A:110:HIS:O	2.36	0.58
1:B:4196:THR:HA	1:B:4199:GLU:HG2	1.85	0.58
1:C:3175:LEU:HD12	1:C:3178:HIS:CD2	2.38	0.58
1:D:2890:GLN:O	1:D:2894:LYS:HG2	2.02	0.58
1:D:3137:LEU:HB2	1:D:3159:LEU:HD13	1.84	0.58
1:D:4010:VAL:HG11	1:D:4118:PHE:CZ	2.38	0.58
1:B:1825:PHE:CE1	1:B:1842:ILE:HG12	2.38	0.58
1:C:1100:ARG:HB3	1:C:1236:TYR:HA	1.85	0.58
1:C:1251:LEU:HD23	1:C:1599:MET:HE2	1.84	0.58
1:D:983:LEU:O	1:D:1055:ARG:NH2	2.37	0.58
1:D:1825:PHE:CE1	1:D:1842:ILE:HG12	2.38	0.58
1:D:3280:ILE:O	1:D:3284:ILE:HG12	2.03	0.58
1:B:591:GLU:HG3	1:B:631:LEU:HD22	1.85	0.58
1:C:561:ARG:HB3	1:C:564:ARG:HD2	1.84	0.58
1:C:591:GLU:HG3	1:C:631:LEU:HD22	1.85	0.58
1:C:983:LEU:O	1:C:1055:ARG:NH2	2.37	0.58
1:C:1825:PHE:CE1	1:C:1842:ILE:HG12	2.38	0.58
1:C:2436:ILE:HA	1:C:2465:LYS:HD3	1.85	0.58
1:C:4196:THR:HA	1:C:4199:GLU:HG2	1.85	0.58
1:C:4271:VAL:HB	1:C:4274:MET:SD	2.43	0.58
1:D:3174:HIS:ND1	1:D:3175:LEU:HD23	2.18	0.58
1:B:1251:LEU:HD23	1:B:1599:MET:HE2	1.84	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:114:LEU:HB2	1:D:117:HIS:CE1	2.39	0.58
1:D:1100:ARG:HB3	1:D:1236:TYR:HA	1.85	0.58
1:D:3059:ALA:O	1:D:3063:ASN:ND2	2.35	0.58
1:A:3152:ARG:NH2	1:A:3236:GLU:HB3	2.16	0.58
1:A:3175:LEU:HD12	1:A:3178:HIS:CD2	2.38	0.58
1:A:4689:LYS:HZ3	1:A:4696:ALA:HB2	1.67	0.58
1:B:15:ARG:NH2	1:B:110:HIS:O	2.36	0.58
1:B:1686:LEU:HD22	1:B:1790:LYS:HZ1	1.67	0.58
1:B:2798:MET:HE1	1:C:1497:GLY:HA3	1.85	0.58
1:B:3174:HIS:ND1	1:B:3175:LEU:HD23	2.19	0.58
1:B:3288:LEU:HD11	1:B:3328:LYS:HD3	1.85	0.58
1:D:2791:ARG:HH12	1:D:2795:GLY:CA	2.17	0.58
1:B:114:LEU:HB2	1:B:117:HIS:CE1	2.39	0.58
1:C:114:LEU:HB2	1:C:117:HIS:CE1	2.39	0.58
1:C:3174:HIS:ND1	1:C:3175:LEU:HD23	2.19	0.58
1:D:126:SER:HA	1:D:414:ARG:HH12	1.66	0.58
1:D:889:ILE:O	1:D:893:TRP:HB2	2.04	0.58
1:D:3288:LEU:HD11	1:D:3328:LYS:HD3	1.85	0.58
1:A:114:LEU:HB2	1:A:117:HIS:CE1	2.38	0.58
1:A:3280:ILE:O	1:A:3284:ILE:HG12	2.03	0.58
1:B:3175:LEU:HD12	1:B:3178:HIS:CD2	2.38	0.58
1:C:2824:ARG:HH22	1:D:1503:ASN:HB2	1.66	0.58
1:D:15:ARG:NH2	1:D:110:HIS:O	2.36	0.58
1:D:1054:VAL:HA	1:D:1057:LEU:HD12	1.84	0.58
1:D:3246:MET:HE1	1:D:3273:MET:HA	1.86	0.58
1:C:188:SER:HB2	1:C:190:ARG:HH11	1.68	0.58
1:C:1009:ARG:HB2	1:C:1013:ARG:NH1	2.18	0.58
1:C:1054:VAL:HA	1:C:1057:LEU:HD12	1.84	0.58
1:C:2937:HIS:HB2	1:C:3014:LEU:HD21	1.85	0.58
1:D:3175:LEU:HD12	1:D:3178:HIS:HD2	1.68	0.58
1:A:3174:HIS:ND1	1:A:3175:LEU:HD23	2.19	0.58
1:A:3246:MET:HE1	1:A:3273:MET:HA	1.86	0.58
1:A:3901:GLN:O	1:A:3905:ASN:ND2	2.28	0.58
1:B:983:LEU:O	1:B:1055:ARG:NH2	2.37	0.58
1:B:2937:HIS:HB2	1:B:3014:LEU:HD21	1.85	0.58
1:B:4106:SER:HB2	1:B:4119:LEU:HD11	1.86	0.58
1:B:4237:SER:N	1:B:4240:THR:OG1	2.36	0.58
1:B:4271:VAL:HB	1:B:4274:MET:SD	2.43	0.58
1:C:15:ARG:NH2	1:C:110:HIS:O	2.36	0.58
1:C:3901:GLN:OE1	1:C:3904:ARG:NH1	2.37	0.58
1:D:2937:HIS:HB2	1:D:3014:LEU:HD21	1.85	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:4196:THR:HA	1:D:4199:GLU:HG2	1.85	0.58
1:A:2930:ILE:HG23	1:A:3010:LYS:HZ3	1.69	0.58
1:A:3175:LEU:HD12	1:A:3178:HIS:HD2	1.68	0.58
1:A:4106:SER:HB2	1:A:4119:LEU:HD11	1.86	0.58
1:B:2788:ARG:NH1	1:B:2905:ARG:O	2.37	0.58
1:B:3059:ALA:O	1:B:3063:ASN:ND2	2.35	0.58
1:B:3246:MET:HE1	1:B:3273:MET:HA	1.86	0.58
1:D:26:ALA:HB3	1:D:33:GLN:HB3	1.86	0.58
1:D:2999:LYS:O	1:D:3002:GLU:HG3	2.04	0.58
1:D:3175:LEU:HD12	1:D:3178:HIS:CD2	2.38	0.58
1:A:1100:ARG:HB3	1:A:1236:TYR:HA	1.85	0.57
1:C:515:ALA:HB2	1:C:523:GLY:HA3	1.86	0.57
1:C:3059:ALA:O	1:C:3063:ASN:ND2	2.35	0.57
1:C:4279:MET:HE2	1:D:4484:ILE:HG13	1.86	0.57
1:D:2455:ASP:OD2	1:D:2457:SER:OG	2.19	0.57
1:D:2788:ARG:NH1	1:D:2905:ARG:O	2.37	0.57
1:D:3045:VAL:O	1:D:3054:LYS:NZ	2.36	0.57
1:D:3160:ALA:N	1:D:3241:MET:HE1	2.19	0.57
1:A:2791:ARG:HH12	1:A:2795:GLY:CA	2.17	0.57
1:A:2999:LYS:O	1:A:3002:GLU:HG3	2.04	0.57
1:A:3288:LEU:HD11	1:A:3328:LYS:HD3	1.85	0.57
1:C:2788:ARG:NH1	1:C:2905:ARG:O	2.37	0.57
1:D:515:ALA:HB2	1:D:523:GLY:HA3	1.86	0.57
1:D:3152:ARG:NH2	1:D:3236:GLU:HB3	2.16	0.57
1:A:2788:ARG:NH1	1:A:2905:ARG:O	2.37	0.57
1:A:3160:ALA:N	1:A:3241:MET:HE1	2.19	0.57
1:C:889:ILE:O	1:C:893:TRP:HB2	2.04	0.57
1:C:2791:ARG:HH12	1:C:2795:GLY:CA	2.17	0.57
1:C:3175:LEU:HD12	1:C:3178:HIS:HD2	1.68	0.57
1:D:3187:LYS:NZ	1:D:3191:GLU:HB3	2.19	0.57
1:A:983:LEU:O	1:A:1055:ARG:NH2	2.37	0.57
1:A:4010:VAL:HG11	1:A:4118:PHE:CZ	2.38	0.57
1:B:2658:GLU:HB3	1:B:2661:LEU:HB3	1.87	0.57
1:B:2999:LYS:O	1:B:3002:GLU:HG3	2.04	0.57
1:C:26:ALA:HB3	1:C:33:GLN:HB3	1.86	0.57
1:C:3246:MET:HE1	1:C:3273:MET:HA	1.86	0.57
1:D:1521:ILE:HG22	1:D:1526:GLU:HG3	1.86	0.57
1:A:3044:THR:HA	1:A:3047:LYS:HZ3	1.70	0.57
1:A:4196:THR:HA	1:A:4199:GLU:HG2	1.85	0.57
1:A:4271:VAL:HB	1:A:4274:MET:SD	2.43	0.57
1:D:2658:GLU:HB3	1:D:2661:LEU:HB3	1.87	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:2975:PHE:HB3	1:D:3039:THR:HG21	1.86	0.57
1:D:3901:GLN:OE1	1:D:3904:ARG:NH1	2.37	0.57
1:A:515:ALA:HB2	1:A:523:GLY:HA3	1.86	0.57
1:A:1521:ILE:HG22	1:A:1526:GLU:HG3	1.87	0.57
1:B:3901:GLN:OE1	1:B:3904:ARG:NH1	2.37	0.57
1:B:4689:LYS:HZ3	1:B:4696:ALA:HB2	1.68	0.57
1:D:3817:LEU:HD22	1:D:3819:MET:HG3	1.87	0.57
1:A:1009:ARG:HB2	1:A:1013:ARG:NH1	2.18	0.57
1:A:3200:ASN:HB2	1:A:3203:ASP:HB2	1.87	0.57
1:B:1054:VAL:HA	1:B:1057:LEU:HD12	1.84	0.57
1:B:1521:ILE:HG22	1:B:1526:GLU:HG3	1.87	0.57
1:C:2999:LYS:O	1:C:3002:GLU:HG3	2.04	0.57
1:C:3200:ASN:HB2	1:C:3203:ASP:HB2	1.87	0.57
1:D:1283:LEU:HB2	1:D:1555:PHE:HB2	1.87	0.57
1:D:2442:MET:HE2	1:D:2502:LEU:HG	1.86	0.57
1:A:3901:GLN:OE1	1:A:3904:ARG:NH1	2.37	0.57
1:A:4049:HIS:CD2	1:A:4067:LEU:HD11	2.40	0.57
1:B:555:LEU:HD21	1:B:578:VAL:HG11	1.86	0.57
1:B:889:ILE:O	1:B:893:TRP:HB2	2.04	0.57
1:B:3152:ARG:NH2	1:B:3233:HIS:O	2.37	0.57
1:B:3200:ASN:HB2	1:B:3203:ASP:HB2	1.87	0.57
1:C:2658:GLU:HB3	1:C:2661:LEU:HB3	1.87	0.57
1:C:3152:ARG:NH2	1:C:3236:GLU:HB3	2.16	0.57
1:C:3172:GLU:OE1	1:C:3266:THR:OG1	2.22	0.57
1:C:3817:LEU:HD22	1:C:3819:MET:HG3	1.87	0.57
1:C:4049:HIS:CD2	1:C:4067:LEU:HD11	2.40	0.57
1:D:555:LEU:HD21	1:D:578:VAL:HG11	1.86	0.57
1:A:4279:MET:HE2	1:B:4484:ILE:HA	1.86	0.57
1:B:515:ALA:HB2	1:B:523:GLY:HA3	1.86	0.57
1:C:4093:ASP:OD1	1:C:4094:ILE:N	2.38	0.57
1:A:3045:VAL:O	1:A:3054:LYS:NZ	2.36	0.57
1:B:4049:HIS:CD2	1:B:4067:LEU:HD11	2.40	0.57
1:C:1521:ILE:HG22	1:C:1526:GLU:HG3	1.87	0.57
1:C:3187:LYS:NZ	1:C:3191:GLU:HB3	2.19	0.57
1:C:4106:SER:HB2	1:C:4119:LEU:HD11	1.86	0.57
1:D:2496:LEU:HD23	1:D:2520:LEU:HD13	1.87	0.57
1:D:3200:ASN:HB2	1:D:3203:ASP:HB2	1.87	0.57
1:D:4237:SER:N	1:D:4240:THR:OG1	2.36	0.57
1:A:889:ILE:O	1:A:893:TRP:HB2	2.04	0.56
1:B:1283:LEU:HB2	1:B:1555:PHE:HB2	1.87	0.56
1:A:555:LEU:HD21	1:A:578:VAL:HG11	1.86	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:3890:TRP:HB3	1:B:76:ARG:HG2	1.86	0.56
1:B:1844:GLN:NE2	1:B:1853:GLU:OE1	2.37	0.56
1:B:2791:ARG:HH12	1:B:2795:GLY:CA	2.17	0.56
1:C:555:LEU:HD21	1:C:578:VAL:HG11	1.86	0.56
1:C:3035:ILE:O	1:C:3039:THR:HG23	2.05	0.56
1:C:3043:ARG:NE	1:C:3120:ASP:OD2	2.27	0.56
1:C:4237:SER:N	1:C:4240:THR:OG1	2.36	0.56
1:D:555:LEU:HD12	1:D:588:ILE:HD11	1.87	0.56
1:D:4084:VAL:O	1:D:4088:HIS:CB	2.54	0.56
1:A:1686:LEU:HD22	1:A:1790:LYS:HZ1	1.70	0.56
1:A:2442:MET:HE2	1:A:2502:LEU:HG	1.86	0.56
1:A:3152:ARG:NH2	1:A:3233:HIS:O	2.37	0.56
1:A:3270:SER:O	1:A:3274:ASN:ND2	2.39	0.56
1:A:4084:VAL:O	1:A:4088:HIS:CB	2.54	0.56
1:A:4093:ASP:OD1	1:A:4094:ILE:N	2.38	0.56
1:B:1849:SER:O	1:B:2054:LYS:NZ	2.39	0.56
1:B:2057:THR:HB	1:B:2060:GLN:HG3	1.86	0.56
1:B:2975:PHE:HB3	1:B:3039:THR:HG21	1.86	0.56
1:B:3172:GLU:OE1	1:B:3266:THR:OG1	2.22	0.56
1:B:3817:LEU:HD22	1:B:3819:MET:HG3	1.87	0.56
1:C:3160:ALA:N	1:C:3241:MET:HE1	2.20	0.56
1:C:3227:ARG:NH1	1:C:3228:TYR:HB3	2.20	0.56
1:C:4177:VAL:HG11	1:C:4880:VAL:HA	1.87	0.56
1:D:1014:GLN:O	1:D:1027:ARG:NH2	2.39	0.56
1:D:1686:LEU:HD22	1:D:1790:LYS:HZ1	1.70	0.56
1:D:3227:ARG:NH1	1:D:3228:TYR:HB3	2.20	0.56
1:D:4106:SER:HB2	1:D:4119:LEU:HD11	1.86	0.56
1:A:2455:ASP:OD2	1:A:2457:SER:OG	2.20	0.56
1:B:3122:ILE:HG13	1:B:3126:VAL:HG23	1.88	0.56
1:B:3160:ALA:N	1:B:3241:MET:HE1	2.19	0.56
1:B:4177:VAL:HG11	1:B:4880:VAL:HA	1.87	0.56
1:C:1283:LEU:HB2	1:C:1555:PHE:HB2	1.87	0.56
1:C:2496:LEU:HD23	1:C:2520:LEU:HD13	1.87	0.56
1:C:2975:PHE:HB3	1:C:3039:THR:HG21	1.86	0.56
1:D:3035:ILE:O	1:D:3039:THR:HG23	2.06	0.56
1:A:4045:LYS:HD3	1:A:4072:THR:HG21	1.88	0.56
1:B:26:ALA:HB3	1:B:33:GLN:HB3	1.86	0.56
1:B:3270:SER:O	1:B:3274:ASN:ND2	2.39	0.56
1:B:4093:ASP:OD1	1:B:4094:ILE:N	2.38	0.56
1:C:2442:MET:HE2	1:C:2502:LEU:HG	1.86	0.56
1:D:891:GLU:HG3	1:D:976:TYR:CD2	2.41	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1286:THR:OG1	1:A:1550:PRO:O	2.19	0.56
1:A:1788:LYS:HD2	1:A:1833:ILE:HG22	1.87	0.56
1:A:2658:GLU:HB3	1:A:2661:LEU:HB3	1.87	0.56
1:B:1014:GLN:O	1:B:1027:ARG:NH2	2.39	0.56
1:B:4045:LYS:HD3	1:B:4072:THR:HG21	1.88	0.56
1:B:4084:VAL:O	1:B:4088:HIS:CB	2.54	0.56
1:D:2999:LYS:O	1:D:3003:MET:HG2	2.06	0.56
1:D:3323:MET:HB3	1:D:3327:LYS:NZ	2.21	0.56
1:D:4045:LYS:HD3	1:D:4072:THR:HG21	1.88	0.56
1:A:26:ALA:HB3	1:A:33:GLN:HB3	1.86	0.56
1:A:3817:LEU:HD22	1:A:3819:MET:HG3	1.87	0.56
1:B:894:VAL:HG12	1:B:897:LYS:NZ	2.21	0.56
1:C:1559:ARG:HD2	1:C:1565:PRO:HD3	1.88	0.56
1:C:2999:LYS:O	1:C:3003:MET:HG2	2.06	0.56
1:C:3270:SER:O	1:C:3274:ASN:ND2	2.39	0.56
1:D:2137:GLU:OE1	1:D:2137:GLU:N	2.32	0.56
1:D:3270:SER:O	1:D:3274:ASN:ND2	2.39	0.56
1:A:1014:GLN:O	1:A:1027:ARG:NH2	2.39	0.56
1:B:555:LEU:HD12	1:B:588:ILE:HD11	1.87	0.56
1:B:881:ILE:HD12	1:B:884:LYS:HZ3	1.69	0.56
1:B:1006:VAL:O	1:B:1009:ARG:HG2	2.06	0.56
1:B:2442:MET:HE2	1:B:2502:LEU:HG	1.86	0.56
1:B:3323:MET:HB3	1:B:3327:LYS:NZ	2.21	0.56
1:C:891:GLU:HG3	1:C:976:TYR:CD2	2.41	0.56
1:C:2057:THR:HB	1:C:2060:GLN:HG3	1.86	0.56
1:C:3045:VAL:O	1:C:3054:LYS:NZ	2.36	0.56
1:D:1788:LYS:HD2	1:D:1833:ILE:HG22	1.87	0.56
1:D:4093:ASP:OD1	1:D:4094:ILE:N	2.38	0.56
1:A:4640:PHE:CD2	1:A:4641:PRO:HD3	2.41	0.56
1:B:1559:ARG:HD2	1:B:1565:PRO:HD3	1.88	0.56
1:B:3187:LYS:NZ	1:B:3191:GLU:HB3	2.19	0.56
1:B:3238:ILE:HA	1:B:3241:MET:HB2	1.88	0.56
1:C:555:LEU:HD12	1:C:588:ILE:HD11	1.87	0.56
1:C:3293:GLY:HA3	1:C:3295:TRP:CZ3	2.41	0.56
1:A:4177:VAL:HG11	1:A:4880:VAL:HA	1.87	0.56
1:D:3152:ARG:NH2	1:D:3233:HIS:O	2.37	0.56
1:D:4177:VAL:HG11	1:D:4880:VAL:HA	1.87	0.56
1:A:3187:LYS:NZ	1:A:3191:GLU:HB3	2.20	0.55
1:A:4144:ARG:HB3	1:A:4961:GLN:HE22	1.72	0.55
1:B:2496:LEU:HD23	1:B:2520:LEU:HD13	1.87	0.55
1:B:4640:PHE:CD2	1:B:4641:PRO:HD3	2.41	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1849:SER:O	1:C:2054:LYS:NZ	2.39	0.55
1:C:4045:LYS:HD3	1:C:4072:THR:HG21	1.88	0.55
1:D:4049:HIS:CD2	1:D:4067:LEU:HD11	2.40	0.55
1:A:2057:THR:HB	1:A:2060:GLN:HG3	1.86	0.55
1:A:3227:ARG:NH1	1:A:3228:TYR:HB3	2.20	0.55
1:A:3323:MET:HB3	1:A:3327:LYS:NZ	2.21	0.55
1:C:3238:ILE:HA	1:C:3241:MET:HB2	1.88	0.55
1:C:3323:MET:HB3	1:C:3327:LYS:NZ	2.21	0.55
1:C:4640:PHE:CD2	1:C:4641:PRO:HD3	2.41	0.55
1:D:2057:THR:HB	1:D:2060:GLN:HG3	1.86	0.55
1:A:1006:VAL:O	1:A:1009:ARG:HG2	2.06	0.55
1:C:1788:LYS:HD2	1:C:1833:ILE:HG22	1.87	0.55
1:C:3152:ARG:NH2	1:C:3233:HIS:O	2.37	0.55
1:C:4084:VAL:O	1:C:4088:HIS:CB	2.54	0.55
1:D:3293:GLY:HA3	1:D:3295:TRP:CZ3	2.41	0.55
2:H:24:VAL:HG12	2:H:105:LEU:HD12	1.89	0.55
1:A:891:GLU:HG3	1:A:976:TYR:CD2	2.41	0.55
1:A:949:HIS:ND1	1:A:1065:GLU:OE2	2.34	0.55
1:B:878:LEU:HA	1:B:881:ILE:HG22	1.88	0.55
1:B:1788:LYS:HD2	1:B:1833:ILE:HG22	1.87	0.55
1:B:3035:ILE:O	1:B:3039:THR:HG23	2.05	0.55
1:B:3293:GLY:HA3	1:B:3295:TRP:CZ3	2.41	0.55
1:B:4144:ARG:HB3	1:B:4961:GLN:HE22	1.72	0.55
1:C:4689:LYS:HZ3	1:C:4696:ALA:HB2	1.71	0.55
1:D:1266:GLU:OE1	1:D:1266:GLU:N	2.30	0.55
1:D:1849:SER:O	1:D:2054:LYS:NZ	2.39	0.55
1:A:1266:GLU:OE1	1:A:1266:GLU:N	2.30	0.55
1:A:2496:LEU:HD23	1:A:2520:LEU:HD13	1.87	0.55
1:A:2975:PHE:HB3	1:A:3039:THR:HG21	1.86	0.55
1:A:3293:GLY:HA3	1:A:3295:TRP:CZ3	2.42	0.55
1:B:949:HIS:ND1	1:B:1065:GLU:OE2	2.34	0.55
1:C:878:LEU:HA	1:C:881:ILE:HG22	1.88	0.55
1:D:3238:ILE:HA	1:D:3241:MET:HB2	1.88	0.55
1:D:4179:GLU:N	1:D:4179:GLU:OE2	2.40	0.55
1:D:4832:GLU:O	1:D:4843:ARG:NH1	2.38	0.55
1:B:3045:VAL:O	1:B:3054:LYS:NZ	2.36	0.55
1:C:2703:PRO:HB2	1:C:2854:LYS:HG3	1.89	0.55
1:C:3122:ILE:HG13	1:C:3126:VAL:HG23	1.88	0.55
1:D:1006:VAL:O	1:D:1009:ARG:HG2	2.06	0.55
1:D:4144:ARG:HB3	1:D:4961:GLN:HE22	1.72	0.55
2:G:24:VAL:HG12	2:G:105:LEU:HD12	1.89	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:555:LEU:HD12	1:A:588:ILE:HD11	1.87	0.55
1:A:3238:ILE:HA	1:A:3241:MET:HB2	1.88	0.55
1:B:2832:THR:HG21	1:C:1290:PHE:HE2	1.71	0.55
1:B:3227:ARG:NH1	1:B:3228:TYR:HB3	2.20	0.55
1:C:1014:GLN:O	1:C:1027:ARG:NH2	2.39	0.55
1:C:4179:GLU:N	1:C:4179:GLU:OE2	2.40	0.55
1:A:3043:ARG:NE	1:A:3120:ASP:OD2	2.27	0.55
1:A:4179:GLU:N	1:A:4179:GLU:OE2	2.40	0.55
1:B:4179:GLU:N	1:B:4179:GLU:OE2	2.40	0.55
1:A:2930:ILE:HG23	1:A:3010:LYS:NZ	2.22	0.55
1:A:3035:ILE:O	1:A:3039:THR:HG23	2.05	0.55
1:B:891:GLU:HG3	1:B:976:TYR:CD2	2.41	0.55
1:B:1427:TYR:HB2	1:B:1563:VAL:HG11	1.89	0.55
1:C:370:LEU:HD22	1:C:395:HIS:HA	1.89	0.55
1:D:4640:PHE:CD2	1:D:4641:PRO:HD3	2.41	0.55
1:A:1283:LEU:HB2	1:A:1555:PHE:HB2	1.87	0.55
1:A:1559:ARG:HD2	1:A:1565:PRO:HD3	1.88	0.55
1:A:3016:ARG:O	1:A:3018:ARG:NE	2.38	0.55
1:C:1844:GLN:NE2	1:C:1853:GLU:OE1	2.37	0.55
1:D:3166:PHE:CE2	1:D:3168:VAL:HB	2.42	0.55
1:A:629:GLN:OE1	1:A:1669:ASN:ND2	2.40	0.54
1:A:1290:PHE:HE2	1:D:2832:THR:HG21	1.72	0.54
1:A:2703:PRO:HB2	1:A:2854:LYS:HG3	1.89	0.54
1:A:3699:CYS:SG	1:A:3731:LEU:HD12	2.48	0.54
1:B:2930:ILE:HG23	1:B:3010:LYS:NZ	2.22	0.54
1:B:2999:LYS:O	1:B:3003:MET:HG2	2.06	0.54
1:D:370:LEU:HD22	1:D:395:HIS:HA	1.89	0.54
1:D:2930:ILE:HG23	1:D:3010:LYS:HZ3	1.73	0.54
1:D:3122:ILE:HG13	1:D:3126:VAL:HG23	1.88	0.54
1:A:2772:ARG:HA	1:A:2775:ILE:HD12	1.90	0.54
1:B:2772:ARG:HA	1:B:2775:ILE:HD12	1.89	0.54
1:C:1006:VAL:O	1:C:1009:ARG:HG2	2.06	0.54
1:C:2832:THR:HG21	1:D:1290:PHE:HE2	1.72	0.54
1:C:3166:PHE:CE2	1:C:3168:VAL:HB	2.42	0.54
1:C:4832:GLU:O	1:C:4843:ARG:NH1	2.38	0.54
1:D:2703:PRO:HB2	1:D:2854:LYS:HG3	1.89	0.54
1:D:2772:ARG:HA	1:D:2775:ILE:HD12	1.90	0.54
1:C:1768:PHE:O	2:G:83:TYR:OH	2.20	0.54
1:C:2129:LEU:HD11	1:C:2141:LEU:HB2	1.89	0.54
1:D:3276:LEU:O	1:D:3280:ILE:HD12	2.08	0.54
1:A:2592:LEU:HD22	1:A:2606:PRO:HB3	1.90	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:3321:PRO:O	1:A:3325:LYS:HG2	2.08	0.54
1:B:2129:LEU:HD11	1:B:2141:LEU:HB2	1.89	0.54
1:B:3313:GLN:HA	1:B:3316:LYS:HZ1	1.72	0.54
1:B:3699:CYS:SG	1:B:3731:LEU:HD12	2.48	0.54
1:C:3276:LEU:O	1:C:3280:ILE:HD12	2.08	0.54
1:C:3699:CYS:SG	1:C:3731:LEU:HD12	2.48	0.54
1:D:2129:LEU:HD11	1:D:2141:LEU:HB2	1.89	0.54
2:F:24:VAL:HG12	2:F:105:LEU:HD12	1.89	0.54
1:A:1844:GLN:NE2	1:A:1853:GLU:OE1	2.37	0.54
1:A:4834:PRO:HB3	1:A:4843:ARG:HG2	1.89	0.54
2:E:24:VAL:HG12	2:E:105:LEU:HD12	1.89	0.54
1:B:902:TRP:CD1	1:B:913:ARG:HH21	2.26	0.54
1:B:2703:PRO:HB2	1:B:2854:LYS:HG3	1.89	0.54
1:B:3016:ARG:O	1:B:3018:ARG:NE	2.38	0.54
1:C:962:LYS:HZ1	1:C:982:ASP:H	1.55	0.54
1:C:2772:ARG:HA	1:C:2775:ILE:HD12	1.89	0.54
1:C:3214:LEU:O	1:C:3218:ILE:HG12	2.08	0.54
1:D:878:LEU:HA	1:D:881:ILE:HG22	1.88	0.54
1:A:2129:LEU:HD11	1:A:2141:LEU:HB2	1.89	0.54
1:A:3276:LEU:O	1:A:3280:ILE:HD12	2.08	0.54
1:B:2704:GLN:O	1:B:2704:GLN:HG2	2.08	0.54
1:B:3297:LYS:HE3	1:B:3334:VAL:HG13	1.90	0.54
1:C:3016:ARG:O	1:C:3018:ARG:NE	2.38	0.54
1:C:3823:GLU:HG3	1:C:3827:GLU:H	1.73	0.54
1:C:4144:ARG:HB3	1:C:4961:GLN:HE22	1.72	0.54
1:D:1559:ARG:HD2	1:D:1565:PRO:HD3	1.88	0.54
1:D:2592:LEU:HD22	1:D:2606:PRO:HB3	1.90	0.54
1:D:2704:GLN:O	1:D:2704:GLN:HG2	2.08	0.54
1:D:3699:CYS:SG	1:D:3731:LEU:HD12	2.48	0.54
2:G:63:GLY:O	2:G:67:MET:HG3	2.07	0.54
1:A:878:LEU:HA	1:A:881:ILE:HG22	1.88	0.54
1:A:1427:TYR:HB2	1:A:1563:VAL:HG11	1.89	0.54
1:C:1427:TYR:HB2	1:C:1563:VAL:HG11	1.89	0.54
1:C:3297:LYS:HE3	1:C:3334:VAL:HG13	1.90	0.54
1:D:2930:ILE:HG23	1:D:3010:LYS:NZ	2.22	0.54
2:F:63:GLY:O	2:F:67:MET:HG3	2.07	0.54
1:B:629:GLN:OE1	1:B:1669:ASN:ND2	2.40	0.54
1:B:3276:LEU:O	1:B:3280:ILE:HD12	2.08	0.54
1:C:629:GLN:OE1	1:C:1669:ASN:ND2	2.40	0.54
1:C:3018:ARG:HG3	1:C:3021:LEU:HD22	1.90	0.54
1:C:3321:PRO:O	1:C:3325:LYS:HG2	2.08	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4279:MET:HE2	1:D:4484:ILE:HA	1.89	0.54
1:C:4834:PRO:HB3	1:C:4843:ARG:HG2	1.89	0.54
1:D:629:GLN:OE1	1:D:1669:ASN:ND2	2.40	0.54
1:D:1272:ARG:HH12	1:D:1584:PRO:HA	1.72	0.54
1:D:1844:GLN:NE2	1:D:1853:GLU:OE1	2.37	0.54
1:A:1849:SER:O	1:A:2054:LYS:NZ	2.39	0.54
1:A:2704:GLN:HG2	1:A:2704:GLN:O	2.08	0.54
1:A:4257:ARG:O	1:A:4261:LEU:HG	2.08	0.54
1:B:3321:PRO:O	1:B:3325:LYS:HG2	2.08	0.54
1:B:4832:GLU:O	1:B:4843:ARG:NH1	2.38	0.54
1:C:2592:LEU:HD22	1:C:2606:PRO:HB3	1.90	0.54
1:C:2895:PHE:HA	1:C:2898:ILE:HG22	1.90	0.54
1:C:4906:GLU:OE1	1:D:4183:LYS:HE3	2.08	0.54
1:D:3321:PRO:O	1:D:3325:LYS:HG2	2.08	0.54
1:A:112:THR:HG21	1:A:174:LYS:HD3	1.91	0.53
1:A:2119:LEU:HB2	1:A:2152:ASN:HD22	1.73	0.53
1:A:2895:PHE:HA	1:A:2898:ILE:HG22	1.89	0.53
1:A:2999:LYS:O	1:A:3003:MET:HG2	2.06	0.53
1:B:2736:LYS:HD2	1:B:2757:MET:HE3	1.90	0.53
1:B:3823:GLU:HG3	1:B:3827:GLU:H	1.73	0.53
1:C:964:MET:HE2	1:C:979:ALA:HB1	1.91	0.53
1:C:1686:LEU:HD22	1:C:1790:LYS:NZ	2.24	0.53
1:C:2455:ASP:OD2	1:C:2457:SER:OG	2.20	0.53
1:C:2930:ILE:HG23	1:C:3010:LYS:NZ	2.22	0.53
1:D:2895:PHE:HA	1:D:2898:ILE:HG22	1.90	0.53
1:D:4834:PRO:HB3	1:D:4843:ARG:HG2	1.89	0.53
1:A:3122:ILE:HG13	1:A:3126:VAL:HG23	1.88	0.53
1:B:964:MET:HE2	1:B:979:ALA:HB1	1.91	0.53
1:B:2895:PHE:HA	1:B:2898:ILE:HG22	1.90	0.53
1:B:3214:LEU:O	1:B:3218:ILE:HG12	2.08	0.53
1:B:4257:ARG:O	1:B:4261:LEU:HG	2.08	0.53
1:C:902:TRP:CD1	1:C:913:ARG:HH21	2.26	0.53
1:D:112:THR:HG21	1:D:174:LYS:HD3	1.91	0.53
1:D:1685:LEU:O	1:D:1689:ILE:HG12	2.08	0.53
1:D:3198:PRO:HG2	1:D:3204:VAL:HA	1.91	0.53
2:H:63:GLY:O	2:H:67:MET:HG3	2.07	0.53
1:A:541:ILE:HD11	1:A:574:VAL:HG13	1.91	0.53
1:A:902:TRP:CD1	1:A:913:ARG:HH21	2.25	0.53
1:A:1685:LEU:O	1:A:1689:ILE:HG12	2.09	0.53
1:A:3172:GLU:OE1	1:A:3266:THR:OG1	2.22	0.53
2:E:63:GLY:O	2:E:67:MET:HG3	2.07	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2930:ILE:HG23	1:B:3010:LYS:HZ3	1.72	0.53
1:B:3198:PRO:HG2	1:B:3204:VAL:HA	1.91	0.53
1:D:949:HIS:ND1	1:D:1065:GLU:OE2	2.34	0.53
1:A:3166:PHE:CE2	1:A:3168:VAL:HB	2.42	0.53
1:A:3297:LYS:HE3	1:A:3334:VAL:HG13	1.90	0.53
1:A:4520:TYR:OH	1:A:4559:TYR:HA	2.09	0.53
1:B:4520:TYR:OH	1:B:4559:TYR:HA	2.09	0.53
1:C:3134:LEU:HB3	1:C:3162:PHE:HE2	1.74	0.53
1:C:4520:TYR:OH	1:C:4559:TYR:HA	2.09	0.53
1:D:1427:TYR:HB2	1:D:1563:VAL:HG11	1.89	0.53
1:D:3778:LEU:HD13	1:D:3854:PHE:CD1	2.41	0.53
1:D:4257:ARG:O	1:D:4261:LEU:HG	2.08	0.53
1:A:2734:MET:HE1	1:A:2823:PRO:HB2	1.91	0.53
1:A:3214:LEU:O	1:A:3218:ILE:HG12	2.08	0.53
1:B:541:ILE:HD11	1:B:574:VAL:HG13	1.91	0.53
1:B:2592:LEU:HD22	1:B:2606:PRO:HB3	1.89	0.53
1:B:3134:LEU:HB3	1:B:3162:PHE:HE2	1.74	0.53
1:B:4834:PRO:HB3	1:B:4843:ARG:HG2	1.89	0.53
1:C:2734:MET:HE1	1:C:2823:PRO:HB2	1.91	0.53
1:C:3068:LEU:O	1:C:3071:THR:OG1	2.25	0.53
1:D:3018:ARG:HG3	1:D:3021:LEU:HD22	1.90	0.53
1:D:3172:GLU:OE1	1:D:3266:THR:OG1	2.22	0.53
1:A:370:LEU:HD22	1:A:395:HIS:HA	1.89	0.53
1:A:1272:ARG:HH12	1:A:1584:PRO:HA	1.73	0.53
1:A:1957:LEU:HA	1:A:1960:ARG:NH1	2.24	0.53
1:C:1685:LEU:O	1:C:1689:ILE:HG12	2.08	0.53
1:C:2704:GLN:HG2	1:C:2704:GLN:O	2.08	0.53
1:C:3044:THR:HA	1:C:3047:LYS:HZ3	1.74	0.53
1:D:541:ILE:HD11	1:D:574:VAL:HG13	1.90	0.53
1:D:2440:PHE:CZ	1:D:2465:LYS:HE3	2.44	0.53
1:D:3214:LEU:O	1:D:3218:ILE:HG12	2.08	0.53
1:D:3297:LYS:HE3	1:D:3334:VAL:HG13	1.90	0.53
1:D:3882:GLN:HB2	1:D:3947:PHE:CE2	2.44	0.53
1:A:3198:PRO:HG2	1:A:3204:VAL:HA	1.91	0.53
1:A:3313:GLN:HA	1:A:3316:LYS:HZ1	1.74	0.53
1:A:4906:GLU:OE1	1:B:4183:LYS:HE3	2.09	0.53
1:B:3778:LEU:HD13	1:B:3854:PHE:CD1	2.41	0.53
1:C:2736:LYS:HD2	1:C:2757:MET:HE3	1.90	0.53
1:C:3882:GLN:HB2	1:C:3947:PHE:CE2	2.44	0.53
1:D:1957:LEU:HA	1:D:1960:ARG:NH1	2.24	0.53
1:D:2119:LEU:HB2	1:D:2152:ASN:HD22	1.73	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1686:LEU:HD22	1:A:1790:LYS:NZ	2.24	0.53
1:A:2055:SER:HB2	1:A:2060:GLN:HB2	1.91	0.53
1:A:3068:LEU:O	1:A:3071:THR:OG1	2.25	0.53
1:A:3071:THR:HA	1:A:3074:ASN:HD21	1.74	0.53
1:A:4832:GLU:O	1:A:4843:ARG:NH1	2.38	0.53
1:B:112:THR:HG21	1:B:174:LYS:HD3	1.91	0.53
1:C:829:LYS:NZ	1:C:1037:LEU:HB3	2.24	0.53
1:D:902:TRP:CD1	1:D:913:ARG:HH21	2.26	0.53
1:A:2440:PHE:CZ	1:A:2465:LYS:HE3	2.44	0.53
1:A:2570:GLU:HG2	1:A:2605:MET:HG3	1.91	0.53
1:B:370:LEU:HD22	1:B:395:HIS:HA	1.89	0.53
1:B:1685:LEU:O	1:B:1689:ILE:HG12	2.09	0.53
1:B:2734:MET:HE1	1:B:2823:PRO:HB2	1.91	0.53
1:B:3166:PHE:CE2	1:B:3168:VAL:HB	2.42	0.53
1:B:4651:ARG:HH11	1:B:4651:ARG:HG2	1.74	0.53
1:C:3198:PRO:HG2	1:C:3204:VAL:HA	1.91	0.53
1:D:829:LYS:NZ	1:D:1037:LEU:HB3	2.24	0.53
1:D:1686:LEU:HD22	1:D:1790:LYS:NZ	2.24	0.53
1:D:2570:GLU:HG2	1:D:2605:MET:HG3	1.91	0.53
1:D:3134:LEU:HB3	1:D:3162:PHE:HE2	1.74	0.53
1:D:3228:TYR:HA	1:D:3235:MET:CE	2.39	0.53
1:D:3697:LYS:HA	1:D:3700:HIS:NE2	2.24	0.53
1:D:3823:GLU:HG3	1:D:3827:GLU:H	1.73	0.53
1:A:514:PHE:HD2	1:A:526:TRP:HB2	1.74	0.53
1:A:3188:SER:OG	1:A:3191:GLU:OE1	2.28	0.53
1:B:3018:ARG:HG3	1:B:3021:LEU:HD22	1.90	0.53
1:B:3882:GLN:HB2	1:B:3947:PHE:CE2	2.44	0.53
1:C:112:THR:HG21	1:C:174:LYS:HD3	1.91	0.53
1:C:889:ILE:HA	1:C:892:LEU:HD12	1.91	0.53
1:C:1957:LEU:HA	1:C:1960:ARG:NH1	2.24	0.53
1:C:2119:LEU:HB2	1:C:2152:ASN:HD22	1.74	0.53
1:C:2440:PHE:CZ	1:C:2465:LYS:HE3	2.44	0.53
1:C:3228:TYR:HA	1:C:3235:MET:CE	2.39	0.53
1:D:2734:MET:HE1	1:D:2823:PRO:HB2	1.91	0.53
1:D:2736:LYS:HD2	1:D:2757:MET:HE3	1.90	0.53
1:A:2137:GLU:OE1	1:A:2137:GLU:N	2.32	0.52
1:A:3778:LEU:HD13	1:A:3854:PHE:CD1	2.41	0.52
1:B:1957:LEU:HA	1:B:1960:ARG:NH1	2.24	0.52
1:B:2055:SER:HB2	1:B:2060:GLN:HB2	1.91	0.52
1:C:3183:ILE:HG23	1:C:3184:TYR:HD1	1.74	0.52
1:C:4257:ARG:O	1:C:4261:LEU:HG	2.08	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:2691:LYS:O	1:D:2694:SER:OG	2.17	0.52
1:A:2736:LYS:HD2	1:A:2757:MET:HE3	1.90	0.52
1:A:4183:LYS:HE3	1:D:4906:GLU:OE1	2.09	0.52
1:B:35:LEU:HD11	1:B:201:LEU:HD13	1.92	0.52
1:B:2119:LEU:HB2	1:B:2152:ASN:HD22	1.73	0.52
1:B:2570:GLU:HG2	1:B:2605:MET:HG3	1.91	0.52
1:B:3044:THR:HA	1:B:3047:LYS:HZ3	1.74	0.52
1:B:3228:TYR:HA	1:B:3235:MET:CE	2.39	0.52
1:B:4903:HIS:CG	1:C:4183:LYS:HZ1	2.27	0.52
1:C:541:ILE:HD11	1:C:574:VAL:HG13	1.91	0.52
1:C:4654:MET:HE1	1:C:4663:ARG:HA	1.92	0.52
1:D:964:MET:HE2	1:D:979:ALA:HB1	1.91	0.52
1:D:4520:TYR:OH	1:D:4559:TYR:HA	2.09	0.52
1:A:3228:TYR:HA	1:A:3235:MET:CE	2.39	0.52
1:B:1272:ARG:HH12	1:B:1584:PRO:HA	1.72	0.52
1:B:2440:PHE:CZ	1:B:2465:LYS:HE3	2.44	0.52
1:B:3188:SER:OG	1:B:3191:GLU:OE1	2.27	0.52
1:C:2570:GLU:HG2	1:C:2605:MET:HG3	1.91	0.52
1:C:3697:LYS:HA	1:C:3700:HIS:NE2	2.24	0.52
1:D:3188:SER:OG	1:D:3191:GLU:OE1	2.27	0.52
1:A:165:ALA:HB1	1:A:211:LEU:HD22	1.92	0.52
1:A:894:VAL:HG12	1:A:897:LYS:NZ	2.21	0.52
1:A:964:MET:HE2	1:A:979:ALA:HB1	1.91	0.52
1:B:3071:THR:HA	1:B:3074:ASN:HD21	1.74	0.52
1:B:3129:SER:O	1:B:3133:ILE:HG13	2.10	0.52
1:C:1118:SER:HB3	1:C:1204:VAL:HG11	1.92	0.52
1:C:1286:THR:OG1	1:C:1550:PRO:O	2.19	0.52
1:C:2691:LYS:O	1:C:2694:SER:OG	2.17	0.52
1:C:3314:LEU:O	1:C:3318:HIS:ND1	2.42	0.52
1:A:3018:ARG:HG3	1:A:3021:LEU:HD22	1.90	0.52
1:A:3697:LYS:HA	1:A:3700:HIS:NE2	2.24	0.52
1:B:514:PHE:HD2	1:B:526:TRP:HB2	1.74	0.52
1:B:2926:LEU:HB3	1:B:3003:MET:HE2	1.92	0.52
1:C:4262:LYS:HG3	1:D:4698:LEU:HD13	1.91	0.52
1:D:35:LEU:HD11	1:D:201:LEU:HD13	1.92	0.52
1:D:889:ILE:HA	1:D:892:LEU:HD12	1.91	0.52
1:D:1118:SER:HB3	1:D:1204:VAL:HG11	1.92	0.52
1:D:4654:MET:HE1	1:D:4663:ARG:HA	1.92	0.52
1:A:35:LEU:HD11	1:A:201:LEU:HD13	1.91	0.52
1:B:3314:LEU:O	1:B:3318:HIS:ND1	2.42	0.52
1:C:35:LEU:HD11	1:C:201:LEU:HD13	1.92	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2926:LEU:HB3	1:C:3003:MET:HE2	1.92	0.52
1:C:4651:ARG:HG2	1:C:4651:ARG:HH11	1.74	0.52
1:A:829:LYS:NZ	1:A:1037:LEU:HB3	2.24	0.52
1:A:2926:LEU:HB3	1:A:3003:MET:HE2	1.92	0.52
1:A:3314:LEU:O	1:A:3318:HIS:ND1	2.42	0.52
1:C:514:PHE:HD2	1:C:526:TRP:HB2	1.74	0.52
1:C:2055:SER:HB2	1:C:2060:GLN:HB2	1.91	0.52
1:C:2137:GLU:OE1	1:C:2137:GLU:N	2.32	0.52
1:D:3016:ARG:O	1:D:3018:ARG:NE	2.38	0.52
1:A:2168:HIS:O	1:A:2172:MET:HB2	2.10	0.52
1:A:3072:MET:SD	1:A:3136:SER:HA	2.50	0.52
1:A:3183:ILE:HG23	1:A:3184:TYR:HD1	1.74	0.52
1:B:889:ILE:HA	1:B:892:LEU:HD12	1.92	0.52
1:D:165:ALA:HB1	1:D:211:LEU:HD22	1.92	0.52
1:D:2759:PRO:HD2	1:D:2762:LEU:HD22	1.92	0.52
1:D:3183:ILE:HG23	1:D:3184:TYR:HD1	1.74	0.52
1:B:829:LYS:NZ	1:B:1037:LEU:HB3	2.24	0.52
1:B:1118:SER:HB3	1:B:1204:VAL:HG11	1.92	0.52
1:B:3183:ILE:HG23	1:B:3184:TYR:HD1	1.74	0.52
1:B:4518:LEU:HD12	1:C:4809:MET:HG3	1.91	0.52
1:C:1272:ARG:HH12	1:C:1584:PRO:HA	1.73	0.52
1:C:2759:PRO:HD2	1:C:2762:LEU:HD22	1.92	0.52
1:C:3071:THR:HA	1:C:3074:ASN:HD21	1.74	0.52
1:D:1041:ARG:O	1:D:1044:LYS:HG3	2.10	0.52
1:D:2055:SER:HB2	1:D:2060:GLN:HB2	1.91	0.52
1:D:4651:ARG:HG2	1:D:4651:ARG:HH11	1.75	0.52
1:A:624:ALA:HB2	1:A:1667:LEU:HD12	1.92	0.52
1:A:948:CYS:HA	1:A:1067:PRO:HD3	1.92	0.52
1:A:1502:ASN:OD1	1:A:1503:ASN:N	2.43	0.52
1:B:1501:ASN:OD1	1:B:1502:ASN:N	2.44	0.52
1:B:2701:PHE:CE2	1:B:2703:PRO:HG3	2.45	0.52
1:B:3697:LYS:HA	1:B:3700:HIS:NE2	2.24	0.52
1:C:1041:ARG:O	1:C:1044:LYS:HG3	2.10	0.52
1:C:1501:ASN:OD1	1:C:1502:ASN:N	2.44	0.52
1:C:3188:SER:OG	1:C:3191:GLU:OE1	2.27	0.52
1:C:3778:LEU:HD13	1:C:3854:PHE:CD1	2.41	0.52
1:D:514:PHE:HD2	1:D:526:TRP:HB2	1.74	0.52
1:A:1041:ARG:O	1:A:1044:LYS:HG3	2.10	0.51
1:A:3070:LYS:O	1:A:3074:ASN:ND2	2.43	0.51
1:A:3882:GLN:HB2	1:A:3947:PHE:CE2	2.44	0.51
2:E:58:LYS:O	2:E:62:GLU:HG3	2.10	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1502:ASN:OD1	1:B:1503:ASN:N	2.43	0.51
1:B:3070:LYS:O	1:B:3074:ASN:ND2	2.43	0.51
1:C:624:ALA:HB2	1:C:1667:LEU:HD12	1.92	0.51
1:C:3070:LYS:O	1:C:3074:ASN:ND2	2.43	0.51
1:D:624:ALA:HB2	1:D:1667:LEU:HD12	1.92	0.51
1:D:2168:HIS:O	1:D:2172:MET:HB2	2.10	0.51
1:D:2926:LEU:HB3	1:D:3003:MET:HE2	1.92	0.51
1:D:3071:THR:HA	1:D:3074:ASN:HD21	1.74	0.51
1:A:3823:GLU:HG3	1:A:3827:GLU:H	1.73	0.51
1:A:4651:ARG:HG2	1:A:4651:ARG:HH11	1.74	0.51
1:A:4654:MET:HE1	1:A:4663:ARG:HA	1.92	0.51
1:B:2119:LEU:HB2	1:B:2152:ASN:ND2	2.25	0.51
1:B:3072:MET:SD	1:B:3136:SER:HA	2.50	0.51
1:D:1176:THR:HG22	1:D:1181:ILE:HA	1.93	0.51
1:D:1501:ASN:OD1	1:D:1502:ASN:N	2.44	0.51
1:D:3314:LEU:O	1:D:3318:HIS:ND1	2.42	0.51
1:A:849:ASP:OD1	1:A:1214:ARG:NE	2.44	0.51
1:A:889:ILE:HA	1:A:892:LEU:HD12	1.92	0.51
1:A:1118:SER:HB3	1:A:1204:VAL:HG11	1.92	0.51
1:A:4569:GLU:HB3	1:A:4570:PRO:HD3	1.92	0.51
1:B:892:LEU:HD21	1:B:980:PRO:HG3	1.93	0.51
1:B:1041:ARG:O	1:B:1044:LYS:HG3	2.10	0.51
1:B:1686:LEU:HD22	1:B:1790:LYS:NZ	2.24	0.51
1:B:4569:GLU:HB3	1:B:4570:PRO:HD3	1.92	0.51
1:B:4654:MET:HE1	1:B:4663:ARG:HA	1.92	0.51
1:C:881:ILE:HG12	1:C:1062:TYR:CE1	2.45	0.51
1:C:881:ILE:HD12	1:C:884:LYS:HZ3	1.75	0.51
1:D:881:ILE:HG12	1:D:1062:TYR:CE1	2.46	0.51
1:D:3882:GLN:HE21	1:D:3946:GLY:HA3	1.75	0.51
1:A:307:SER:OG	1:A:315:LEU:O	2.28	0.51
1:A:1184:ASP:OD1	1:A:1188:SER:OG	2.29	0.51
1:A:1414:ARG:O	1:A:1561:LYS:NZ	2.44	0.51
1:A:3129:SER:O	1:A:3133:ILE:HG13	2.10	0.51
1:A:3250:TRP:NE1	1:A:3256:ASN:HD21	2.09	0.51
1:A:3882:GLN:HE21	1:A:3946:GLY:HA3	1.75	0.51
1:B:881:ILE:HG13	1:B:885:LEU:HD23	1.93	0.51
1:C:3129:SER:O	1:C:3133:ILE:HG13	2.10	0.51
1:D:3068:LEU:O	1:D:3071:THR:OG1	2.25	0.51
2:F:58:LYS:O	2:F:62:GLU:HG3	2.10	0.51
1:A:1176:THR:HG22	1:A:1181:ILE:HA	1.93	0.51
1:A:3270:SER:HA	1:A:3273:MET:CE	2.37	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:881:ILE:HG12	1:B:1062:TYR:CE1	2.46	0.51
1:B:3209:PRO:HB3	1:B:3213:LYS:HE2	1.93	0.51
1:B:3250:TRP:NE1	1:B:3256:ASN:HD21	2.09	0.51
1:C:1436:GLN:O	1:C:1500:ARG:NH2	2.43	0.51
1:C:3250:TRP:NE1	1:C:3256:ASN:HD21	2.09	0.51
1:C:4569:GLU:HB3	1:C:4570:PRO:HD3	1.93	0.51
1:D:163:HIS:HB2	1:D:182:ILE:HG23	1.93	0.51
1:D:849:ASP:OD1	1:D:1214:ARG:NE	2.44	0.51
1:D:1502:ASN:OD1	1:D:1503:ASN:N	2.43	0.51
1:A:1436:GLN:O	1:A:1500:ARG:NH2	2.43	0.51
1:A:3228:TYR:CE2	1:A:3232:PRO:HB3	2.46	0.51
1:B:948:CYS:HA	1:B:1067:PRO:HD3	1.92	0.51
1:B:2168:HIS:O	1:B:2172:MET:HB2	2.10	0.51
1:C:1176:THR:HG22	1:C:1181:ILE:HA	1.93	0.51
1:C:2701:PHE:CE2	1:C:2703:PRO:HG3	2.45	0.51
1:C:2824:ARG:NH2	1:D:1502:ASN:O	2.44	0.51
1:D:894:VAL:HG12	1:D:897:LYS:NZ	2.21	0.51
1:D:2858:MET:N	1:D:2858:MET:HE2	2.26	0.51
1:A:2701:PHE:CE2	1:A:2703:PRO:HG3	2.45	0.51
1:A:3209:PRO:HB3	1:A:3213:LYS:HE2	1.93	0.51
1:B:235:ARG:NH2	1:B:412:GLU:OE2	2.27	0.51
1:B:624:ALA:HB2	1:B:1667:LEU:HD12	1.92	0.51
1:B:995:MET:O	1:B:999:LEU:HD23	2.11	0.51
1:B:1414:ARG:O	1:B:1561:LYS:NZ	2.44	0.51
1:B:2793:ARG:HH21	1:B:2796:ASP:HB3	1.76	0.51
1:C:307:SER:HB3	1:C:327:THR:HG22	1.93	0.51
1:C:892:LEU:HD21	1:C:980:PRO:HG3	1.93	0.51
1:D:1436:GLN:O	1:D:1500:ARG:NH2	2.43	0.51
1:D:2119:LEU:HB2	1:D:2152:ASN:ND2	2.25	0.51
1:D:2723:LYS:HG2	1:D:2895:PHE:CZ	2.43	0.51
1:A:884:LYS:HA	1:A:887:GLU:OE1	2.11	0.51
1:A:995:MET:O	1:A:999:LEU:HD23	2.11	0.51
1:B:165:ALA:HB1	1:B:211:LEU:HD22	1.92	0.51
1:B:307:SER:HB3	1:B:327:THR:HG22	1.93	0.51
1:B:884:LYS:HA	1:B:887:GLU:OE1	2.11	0.51
1:B:1184:ASP:OD1	1:B:1188:SER:OG	2.29	0.51
1:B:2858:MET:HE2	1:B:2858:MET:N	2.26	0.51
1:B:3228:TYR:CE2	1:B:3232:PRO:HB3	2.46	0.51
1:B:3882:GLN:HE21	1:B:3946:GLY:HA3	1.75	0.51
2:H:58:LYS:O	2:H:62:GLU:HG3	2.10	0.51
1:A:2119:LEU:HB2	1:A:2152:ASN:ND2	2.25	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:3145:SER:O	1:A:3149:GLU:HG2	2.11	0.51
1:B:1176:THR:HG22	1:B:1181:ILE:HA	1.93	0.51
1:B:1266:GLU:OE1	1:B:1266:GLU:N	2.30	0.51
1:B:1598:ARG:NH2	1:B:1601:ASN:OD1	2.44	0.51
1:C:881:ILE:HG13	1:C:885:LEU:HD23	1.93	0.51
1:C:1502:ASN:OD1	1:C:1503:ASN:N	2.43	0.51
1:C:2930:ILE:HG23	1:C:3010:LYS:HZ3	1.75	0.51
1:C:2930:ILE:HD12	1:C:3003:MET:HE3	1.92	0.51
1:C:3145:SER:O	1:C:3149:GLU:HG2	2.11	0.51
1:C:3882:GLN:HE21	1:C:3946:GLY:HA3	1.74	0.51
2:G:58:LYS:O	2:G:62:GLU:HG3	2.10	0.51
1:A:1501:ASN:OD1	1:A:1502:ASN:N	2.44	0.51
1:A:1598:ARG:NH2	1:A:1601:ASN:OD1	2.44	0.51
1:C:2338:GLU:N	1:C:2338:GLU:OE1	2.44	0.51
1:D:3070:LYS:O	1:D:3074:ASN:ND2	2.43	0.51
1:D:3072:MET:SD	1:D:3136:SER:HA	2.50	0.51
1:A:881:ILE:HG13	1:A:885:LEU:HD23	1.93	0.50
1:A:920:GLU:OE1	1:A:974:SER:OG	2.25	0.50
1:A:2759:PRO:HD2	1:A:2762:LEU:HD22	1.92	0.50
1:A:2793:ARG:HH21	1:A:2796:ASP:HB3	1.76	0.50
1:A:2930:ILE:HD12	1:A:3003:MET:HE3	1.92	0.50
1:B:1124:PRO:HD2	1:B:1594:VAL:HG23	1.93	0.50
1:B:1436:GLN:O	1:B:1500:ARG:NH2	2.43	0.50
1:B:2759:PRO:HD2	1:B:2762:LEU:HD22	1.92	0.50
1:B:4250:TYR:O	1:B:4254:THR:HG23	2.11	0.50
1:C:884:LYS:HA	1:C:887:GLU:OE1	2.11	0.50
1:C:995:MET:O	1:C:999:LEU:HD23	2.11	0.50
1:C:3072:MET:SD	1:C:3136:SER:HA	2.50	0.50
1:C:3209:PRO:HB3	1:C:3213:LYS:HE2	1.93	0.50
1:A:997:ASP:HA	1:A:1047:LYS:HZ2	1.74	0.50
1:A:2884:LYS:O	1:A:2887:GLU:HG3	2.11	0.50
1:A:3134:LEU:HB3	1:A:3162:PHE:HE2	1.74	0.50
1:B:3145:SER:O	1:B:3149:GLU:HG2	2.11	0.50
1:C:163:HIS:HB2	1:C:182:ILE:HG23	1.93	0.50
1:C:948:CYS:HA	1:C:1067:PRO:HD3	1.92	0.50
1:C:1414:ARG:O	1:C:1561:LYS:NZ	2.44	0.50
1:C:1960:ARG:HA	1:C:1963:LYS:HG2	1.94	0.50
1:C:3228:TYR:CE2	1:C:3232:PRO:HB3	2.46	0.50
1:A:307:SER:HB3	1:A:327:THR:HG22	1.93	0.50
1:A:881:ILE:HG12	1:A:1062:TYR:CE1	2.45	0.50
1:A:882:ARG:NH1	1:A:883:GLU:HB2	2.27	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:4943:MET:HE1	1:A:4950:GLU:HB2	1.94	0.50
1:B:849:ASP:OD1	1:B:1214:ARG:NE	2.44	0.50
1:B:1006:VAL:HA	1:B:1009:ARG:NE	2.27	0.50
1:B:4502:MET:SD	1:B:4585:PHE:HB3	2.52	0.50
1:C:894:VAL:HG12	1:C:897:LYS:NZ	2.21	0.50
1:C:2119:LEU:HB2	1:C:2152:ASN:ND2	2.25	0.50
1:C:4502:MET:SD	1:C:4585:PHE:HB3	2.52	0.50
1:D:1414:ARG:O	1:D:1561:LYS:NZ	2.43	0.50
1:D:4502:MET:SD	1:D:4585:PHE:HB3	2.52	0.50
1:A:163:HIS:HB2	1:A:182:ILE:HG23	1.93	0.50
1:B:776:GLN:NE2	1:B:1472:GLU:OE2	2.45	0.50
1:B:997:ASP:HA	1:B:1047:LYS:HZ2	1.76	0.50
1:C:1598:ARG:NH2	1:C:1601:ASN:OD1	2.44	0.50
1:C:2168:HIS:O	1:C:2172:MET:HB2	2.10	0.50
1:C:4806:CYS:HA	1:C:4812:CYS:HB2	1.93	0.50
1:D:3250:TRP:NE1	1:D:3256:ASN:HD21	2.09	0.50
1:A:2087:LEU:O	1:A:2091:GLN:HG2	2.12	0.50
1:A:4896:ASP:OD1	1:A:4897:TYR:N	2.45	0.50
1:C:165:ALA:HB1	1:C:211:LEU:HD22	1.92	0.50
1:C:323:ASP:OD1	1:C:324:VAL:N	2.45	0.50
1:C:776:GLN:NE2	1:C:1472:GLU:OE2	2.45	0.50
1:C:849:ASP:OD1	1:C:1214:ARG:NE	2.44	0.50
1:C:895:MET:O	1:C:899:GLU:HG2	2.12	0.50
1:C:4943:MET:HE1	1:C:4950:GLU:HB2	1.94	0.50
1:D:882:ARG:NH1	1:D:883:GLU:HB2	2.27	0.50
1:D:3209:PRO:HB3	1:D:3213:LYS:HE2	1.93	0.50
1:A:776:GLN:NE2	1:A:1472:GLU:OE2	2.45	0.50
1:A:892:LEU:HD21	1:A:980:PRO:HG3	1.92	0.50
1:A:2858:MET:N	1:A:2858:MET:HE2	2.26	0.50
1:B:658:ASN:ND2	1:B:833:LYS:HG2	2.27	0.50
1:B:1960:ARG:HA	1:B:1963:LYS:HG2	1.94	0.50
1:B:3234:VAL:HA	1:B:3238:ILE:HD12	1.94	0.50
1:C:1184:ASP:OD1	1:C:1188:SER:OG	2.29	0.50
1:C:2723:LYS:HG2	1:C:2895:PHE:CZ	2.43	0.50
1:C:2884:LYS:O	1:C:2887:GLU:HG3	2.11	0.50
1:D:884:LYS:HA	1:D:887:GLU:OE1	2.11	0.50
1:D:892:LEU:HD21	1:D:980:PRO:HG3	1.92	0.50
1:D:895:MET:O	1:D:899:GLU:HG2	2.12	0.50
1:D:1184:ASP:OD1	1:D:1188:SER:OG	2.29	0.50
1:D:1960:ARG:HA	1:D:1963:LYS:HG2	1.94	0.50
1:D:2701:PHE:CE2	1:D:2703:PRO:HG3	2.45	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:2884:LYS:O	1:D:2887:GLU:HG3	2.11	0.50
1:D:3129:SER:O	1:D:3133:ILE:HG13	2.10	0.50
1:D:4943:MET:HE1	1:D:4950:GLU:HB2	1.94	0.50
1:A:658:ASN:ND2	1:A:833:LYS:HG2	2.26	0.50
1:C:1006:VAL:HA	1:C:1009:ARG:NE	2.27	0.50
1:C:2858:MET:HE2	1:C:2858:MET:N	2.26	0.50
1:C:3650:GLU:HB2	1:C:3651:PRO:HD3	1.94	0.50
1:D:2087:LEU:O	1:D:2091:GLN:HG2	2.12	0.50
1:D:2338:GLU:OE1	1:D:2338:GLU:N	2.44	0.50
1:D:3986:LEU:HG	1:D:3999:MET:HE1	1.94	0.50
1:D:4569:GLU:HB3	1:D:4570:PRO:HD3	1.93	0.50
1:A:3650:GLU:HB2	1:A:3651:PRO:HD3	1.94	0.50
1:A:4250:TYR:O	1:A:4254:THR:HG23	2.11	0.50
1:A:4502:MET:SD	1:A:4585:PHE:HB3	2.52	0.50
1:B:28:ILE:HG22	1:B:29:HIS:HD2	1.77	0.50
1:C:28:ILE:HG22	1:C:29:HIS:HD2	1.77	0.50
1:C:1124:PRO:HD2	1:C:1594:VAL:HG23	1.93	0.50
1:C:4116:GLN:O	1:C:4120:GLU:HG2	2.12	0.50
1:D:3145:SER:O	1:D:3149:GLU:HG2	2.11	0.50
1:D:3228:TYR:CE2	1:D:3232:PRO:HB3	2.46	0.50
1:D:3650:GLU:HB2	1:D:3651:PRO:HD3	1.94	0.50
1:A:76:ARG:HG2	1:D:3890:TRP:HB3	1.93	0.50
1:A:235:ARG:NH2	1:A:412:GLU:OE2	2.27	0.50
1:B:2087:LEU:O	1:B:2091:GLN:HG2	2.12	0.50
1:B:2884:LYS:O	1:B:2887:GLU:HG3	2.11	0.50
1:B:3153:SER:O	1:B:3156:GLY:N	2.45	0.50
1:B:3650:GLU:HB2	1:B:3651:PRO:HD3	1.94	0.50
1:C:1686:LEU:HD22	1:C:1790:LYS:HZ1	1.76	0.50
1:C:2087:LEU:O	1:C:2091:GLN:HG2	2.12	0.50
1:C:2629:ASN:OD1	1:C:2630:PHE:N	2.45	0.50
1:C:4896:ASP:OD1	1:C:4897:TYR:N	2.45	0.50
1:D:881:ILE:HG13	1:D:885:LEU:HD23	1.93	0.50
1:D:1006:VAL:HA	1:D:1009:ARG:NE	2.27	0.50
1:B:323:ASP:OD1	1:B:324:VAL:N	2.45	0.49
1:B:2137:GLU:OE1	1:B:2137:GLU:N	2.32	0.49
1:B:3986:LEU:HG	1:B:3999:MET:HE1	1.93	0.49
1:B:4943:MET:HE1	1:B:4950:GLU:HB2	1.94	0.49
1:C:15:ARG:HH21	1:C:110:HIS:HB3	1.77	0.49
1:C:997:ASP:HA	1:C:1047:LYS:HZ2	1.76	0.49
1:C:2716:LYS:HD3	1:C:2791:ARG:HG3	1.94	0.49
1:C:4250:TYR:O	1:C:4254:THR:HG23	2.11	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:307:SER:HB3	1:D:327:THR:HG22	1.93	0.49
1:D:2930:ILE:HD12	1:D:3003:MET:HE3	1.92	0.49
1:D:4250:TYR:O	1:D:4254:THR:HG23	2.11	0.49
1:A:962:LYS:HZ1	1:A:982:ASP:H	1.59	0.49
1:A:1006:VAL:HA	1:A:1009:ARG:NE	2.27	0.49
1:A:1124:PRO:HD2	1:A:1594:VAL:HG23	1.93	0.49
1:A:1960:ARG:HA	1:A:1963:LYS:HG2	1.94	0.49
1:A:3234:VAL:HA	1:A:3238:ILE:HD12	1.94	0.49
1:A:3986:LEU:HG	1:A:3999:MET:HE1	1.94	0.49
2:E:24:VAL:HG22	2:E:48:LYS:HG2	1.94	0.49
1:B:882:ARG:NH1	1:B:883:GLU:HB2	2.27	0.49
1:B:2338:GLU:OE1	1:B:2338:GLU:N	2.44	0.49
1:B:2788:ARG:NH2	1:B:2906:GLY:O	2.45	0.49
1:C:1266:GLU:OE1	1:C:1266:GLU:N	2.30	0.49
1:D:323:ASP:OD1	1:D:324:VAL:N	2.45	0.49
1:D:948:CYS:HA	1:D:1067:PRO:HD3	1.92	0.49
1:D:4116:GLN:O	1:D:4120:GLU:HG2	2.12	0.49
1:D:4735:ASN:HB3	1:D:4738:PHE:CD2	2.47	0.49
1:A:1978:ASN:HB3	1:A:1983:LYS:NZ	2.27	0.49
1:A:2338:GLU:OE1	1:A:2338:GLU:N	2.44	0.49
1:B:163:HIS:HB2	1:B:182:ILE:HG23	1.93	0.49
1:B:4116:GLN:O	1:B:4120:GLU:HG2	2.12	0.49
1:B:4118:PHE:HA	1:B:4121:LEU:HD12	1.94	0.49
1:B:4896:ASP:OD1	1:B:4897:TYR:N	2.45	0.49
1:C:3234:VAL:HA	1:C:3238:ILE:HD12	1.94	0.49
1:C:3986:LEU:HG	1:C:3999:MET:HE1	1.94	0.49
1:D:1598:ARG:NH2	1:D:1601:ASN:OD1	2.44	0.49
1:D:2788:ARG:NH2	1:D:2906:GLY:O	2.45	0.49
1:D:3284:ILE:HG21	1:D:3336:GLU:HG2	1.94	0.49
1:A:3284:ILE:HG21	1:A:3336:GLU:HG2	1.94	0.49
1:B:891:GLU:O	1:B:895:MET:HG2	2.13	0.49
1:B:1436:GLN:H	1:B:1500:ARG:HH22	1.60	0.49
1:B:2930:ILE:HD12	1:B:3003:MET:HE3	1.92	0.49
1:B:4806:CYS:HA	1:B:4812:CYS:HB2	1.93	0.49
1:C:882:ARG:NH1	1:C:883:GLU:HB2	2.27	0.49
1:C:891:GLU:O	1:C:895:MET:HG2	2.13	0.49
1:D:658:ASN:ND2	1:D:833:LYS:HG2	2.27	0.49
1:D:995:MET:O	1:D:999:LEU:HD23	2.11	0.49
1:D:4038:ASP:OD2	1:D:4039:GLY:N	2.46	0.49
1:D:4806:CYS:HA	1:D:4812:CYS:HB2	1.93	0.49
2:H:78:THR:OG1	2:H:80:ASP:OD1	2.26	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:981:MET:N	1:A:981:MET:SD	2.86	0.49
1:B:2629:ASN:OD1	1:B:2630:PHE:N	2.45	0.49
1:C:307:SER:OG	1:C:315:LEU:O	2.28	0.49
1:C:981:MET:N	1:C:981:MET:SD	2.86	0.49
1:C:1727:VAL:HG11	1:C:1926:VAL:HG21	1.95	0.49
1:C:2793:ARG:HH21	1:C:2796:ASP:HB3	1.76	0.49
1:D:28:ILE:HG22	1:D:29:HIS:HD2	1.77	0.49
1:D:131:CYS:SG	1:D:150:GLN:HB2	2.53	0.49
1:A:1922:ILE:O	1:A:1926:VAL:HG23	2.13	0.49
1:A:2788:ARG:NH2	1:A:2906:GLY:O	2.45	0.49
1:A:4038:ASP:OD2	1:A:4039:GLY:N	2.46	0.49
1:A:4116:GLN:O	1:A:4120:GLU:HG2	2.12	0.49
1:B:2455:ASP:OD2	1:B:2457:SER:OG	2.20	0.49
1:B:4625:ASP:OD1	1:B:4625:ASP:N	2.44	0.49
1:B:4735:ASN:HB3	1:B:4738:PHE:CD2	2.47	0.49
1:C:131:CYS:SG	1:C:150:GLN:HB2	2.53	0.49
1:C:658:ASN:ND2	1:C:833:LYS:HG2	2.26	0.49
1:C:1436:GLN:H	1:C:1500:ARG:HH22	1.60	0.49
1:C:2788:ARG:NH2	1:C:2906:GLY:O	2.45	0.49
1:D:1768:PHE:O	2:H:83:TYR:OH	2.25	0.49
1:D:2261:ASP:OD1	1:D:2262:LEU:N	2.45	0.49
1:D:3234:VAL:HA	1:D:3238:ILE:HD12	1.94	0.49
2:G:24:VAL:HG22	2:G:48:LYS:HG2	1.94	0.49
1:A:164:PRO:HB3	1:A:169:ARG:HB2	1.95	0.49
1:A:1436:GLN:H	1:A:1500:ARG:HH22	1.60	0.49
1:A:1457:PHE:HA	1:A:1461:ARG:HH12	1.78	0.49
1:A:2629:ASN:OD1	1:A:2630:PHE:N	2.45	0.49
1:A:4806:CYS:HA	1:A:4812:CYS:HB2	1.93	0.49
1:B:307:SER:OG	1:B:315:LEU:O	2.28	0.49
1:B:4084:VAL:O	1:B:4088:HIS:HB3	2.13	0.49
1:B:4265:LYS:O	1:B:4269:LYS:HG2	2.13	0.49
1:C:1978:ASN:HB3	1:C:1983:LYS:NZ	2.27	0.49
1:C:3157:GLU:HG3	1:C:3302:PHE:HZ	1.77	0.49
1:D:1124:PRO:HD2	1:D:1594:VAL:HG23	1.93	0.49
1:D:1922:ILE:O	1:D:1926:VAL:HG23	2.13	0.49
1:D:3317:THR:H	1:D:3320:LEU:HD12	1.78	0.49
1:D:4943:MET:HG2	1:D:4948:CYS:HB3	1.95	0.49
1:A:323:ASP:OD1	1:A:324:VAL:N	2.45	0.49
1:A:675:TYR:HB3	1:A:822:CYS:SG	2.53	0.49
1:A:895:MET:O	1:A:899:GLU:HG2	2.12	0.49
1:A:2691:LYS:O	1:A:2694:SER:OG	2.17	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:3187:LYS:O	1:A:3188:SER:OG	2.28	0.49
1:B:894:VAL:O	1:B:898:ILE:HG12	2.13	0.49
1:B:1017:THR:O	1:B:1028:ARG:HA	2.13	0.49
1:B:1922:ILE:O	1:B:1926:VAL:HG23	2.13	0.49
1:B:4640:PHE:CG	1:B:4641:PRO:HD3	2.48	0.49
1:C:675:TYR:HB3	1:C:822:CYS:SG	2.53	0.49
1:C:4943:MET:HG2	1:C:4948:CYS:HB3	1.95	0.49
1:D:891:GLU:O	1:D:895:MET:HG2	2.13	0.49
1:D:920:GLU:N	1:D:974:SER:OG	2.46	0.49
1:D:1051:ARG:O	1:D:1055:ARG:HG2	2.13	0.49
1:D:1978:ASN:HB3	1:D:1983:LYS:NZ	2.27	0.49
1:D:2629:ASN:OD1	1:D:2630:PHE:N	2.45	0.49
1:D:3157:GLU:HG3	1:D:3302:PHE:HZ	1.77	0.49
1:D:4896:ASP:OD1	1:D:4897:TYR:N	2.45	0.49
1:A:131:CYS:SG	1:A:150:GLN:HB2	2.53	0.49
1:A:891:GLU:O	1:A:895:MET:HG2	2.13	0.49
1:A:920:GLU:N	1:A:974:SER:OG	2.46	0.49
1:A:3070:LYS:O	1:A:3073:GLU:HG3	2.13	0.49
1:A:4809:MET:HG3	1:D:4518:LEU:HD12	1.95	0.49
1:A:4943:MET:HG2	1:A:4948:CYS:HB3	1.95	0.49
1:B:131:CYS:SG	1:B:150:GLN:HB2	2.53	0.49
1:B:981:MET:SD	1:B:981:MET:N	2.86	0.49
1:B:1727:VAL:HG11	1:B:1926:VAL:HG21	1.95	0.49
1:B:3157:GLU:HG3	1:B:3302:PHE:HZ	1.77	0.49
1:B:4047:ASP:HA	1:B:4050:LYS:HG2	1.95	0.49
1:C:1457:PHE:HA	1:C:1461:ARG:HH12	1.78	0.49
1:C:1685:LEU:HD22	1:C:1706:LEU:HB2	1.95	0.49
1:C:2261:ASP:OD1	1:C:2262:LEU:N	2.46	0.49
1:C:3153:SER:O	1:C:3156:GLY:N	2.45	0.49
1:C:3317:THR:H	1:C:3320:LEU:HD12	1.78	0.49
1:C:4640:PHE:CG	1:C:4641:PRO:HD3	2.48	0.49
1:C:4735:ASN:HB3	1:C:4738:PHE:CD2	2.47	0.49
1:D:15:ARG:HH21	1:D:110:HIS:HB3	1.77	0.49
1:D:675:TYR:HB3	1:D:822:CYS:SG	2.53	0.49
1:D:776:GLN:NE2	1:D:1472:GLU:OE2	2.45	0.49
1:D:2716:LYS:HD3	1:D:2791:ARG:HG3	1.94	0.49
1:D:2918:GLU:HA	1:D:2923:TYR:CD1	2.48	0.49
1:D:3187:LYS:HZ2	1:D:3191:GLU:HB3	1.78	0.49
1:D:4640:PHE:CG	1:D:4641:PRO:HD3	2.48	0.49
1:A:1549:SER:HB2	1:D:2830:ASN:HB3	1.94	0.49
1:B:164:PRO:HB3	1:B:169:ARG:HB2	1.95	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:962:LYS:HZ1	1:B:982:ASP:H	1.61	0.49
1:B:1790:LYS:O	1:B:1794:MET:HG3	2.13	0.49
1:B:4038:ASP:OD2	1:B:4039:GLY:N	2.46	0.49
1:B:4113:THR:O	1:B:4117:THR:HG23	2.13	0.49
1:C:1051:ARG:O	1:C:1055:ARG:HG2	2.13	0.49
1:C:4038:ASP:OD2	1:C:4039:GLY:N	2.46	0.49
1:C:4113:THR:O	1:C:4117:THR:HG23	2.13	0.49
1:C:4118:PHE:HA	1:C:4121:LEU:HD12	1.94	0.49
1:D:1685:LEU:HD22	1:D:1706:LEU:HB2	1.95	0.49
1:D:3811:GLN:NE2	1:D:3828:LYS:HB3	2.28	0.49
1:D:4118:PHE:HA	1:D:4121:LEU:HD12	1.94	0.49
1:A:28:ILE:HG22	1:A:29:HIS:HD2	1.77	0.48
1:A:3811:GLN:NE2	1:A:3828:LYS:HB3	2.28	0.48
1:A:4113:THR:O	1:A:4117:THR:HG23	2.13	0.48
1:A:4118:PHE:HA	1:A:4121:LEU:HD12	1.94	0.48
1:B:920:GLU:OE1	1:B:974:SER:OG	2.25	0.48
1:B:1168:MET:HE1	1:B:1200:GLY:HA2	1.95	0.48
1:B:1978:ASN:HB3	1:B:1983:LYS:NZ	2.27	0.48
1:B:2770:ILE:HG13	1:B:2771:TYR:CD1	2.48	0.48
1:B:3317:THR:H	1:B:3320:LEU:HD12	1.78	0.48
1:C:894:VAL:O	1:C:898:ILE:HG12	2.13	0.48
1:C:920:GLU:N	1:C:974:SER:OG	2.45	0.48
1:C:1168:MET:HE1	1:C:1200:GLY:HA2	1.95	0.48
1:C:3284:ILE:HG21	1:C:3336:GLU:HG2	1.94	0.48
1:C:4654:MET:HE2	1:C:4663:ARG:HD3	1.95	0.48
1:D:307:SER:OG	1:D:315:LEU:O	2.28	0.48
1:D:2341:ASN:OD1	1:D:2342:GLY:N	2.46	0.48
1:D:2770:ILE:HG13	1:D:2771:TYR:CD1	2.48	0.48
1:A:1727:VAL:HG11	1:A:1926:VAL:HG21	1.95	0.48
1:A:4047:ASP:HA	1:A:4050:LYS:HG2	1.95	0.48
1:B:675:TYR:HB3	1:B:822:CYS:SG	2.53	0.48
1:B:2318:ASN:HB3	1:B:2322:ARG:HH12	1.78	0.48
1:B:2581:ARG:HB2	1:B:2584:MET:HG3	1.96	0.48
1:C:2918:GLU:HA	1:C:2923:TYR:CD1	2.48	0.48
1:D:1017:THR:O	1:D:1028:ARG:HA	2.13	0.48
1:D:2162:MET:HE2	1:D:2162:MET:N	2.28	0.48
1:D:2793:ARG:HH21	1:D:2796:ASP:HB3	1.76	0.48
1:D:4265:LYS:O	1:D:4269:LYS:HG2	2.13	0.48
2:F:24:VAL:HG22	2:F:48:LYS:HG2	1.94	0.48
2:H:24:VAL:HG22	2:H:48:LYS:HG2	1.94	0.48
1:A:881:ILE:HG13	1:A:885:LEU:CD2	2.44	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2581:ARG:HB2	1:A:2584:MET:HG3	1.96	0.48
1:A:2918:GLU:HA	1:A:2923:TYR:CD1	2.48	0.48
1:A:4279:MET:HE2	1:B:4484:ILE:HG13	1.94	0.48
1:A:4735:ASN:HB3	1:A:4738:PHE:CD2	2.47	0.48
1:B:1768:PHE:O	2:F:83:TYR:OH	2.21	0.48
1:B:2162:MET:HE2	1:B:2162:MET:N	2.28	0.48
1:B:2261:ASP:OD1	1:B:2262:LEU:N	2.46	0.48
1:B:2830:ASN:HB3	1:C:1549:SER:HB2	1.94	0.48
1:B:3070:LYS:O	1:B:3073:GLU:HG3	2.13	0.48
1:B:3284:ILE:HG21	1:B:3336:GLU:HG2	1.94	0.48
1:B:3692:ALA:HA	1:B:3695:MET:HE3	1.95	0.48
1:C:1017:THR:O	1:C:1028:ARG:HA	2.13	0.48
1:C:1922:ILE:O	1:C:1926:VAL:HG23	2.13	0.48
1:C:2341:ASN:OD1	1:C:2342:GLY:N	2.47	0.48
1:C:3070:LYS:O	1:C:3073:GLU:HG3	2.13	0.48
1:C:3811:GLN:NE2	1:C:3828:LYS:HB3	2.28	0.48
1:D:3070:LYS:O	1:D:3073:GLU:HG3	2.13	0.48
1:D:3187:LYS:O	1:D:3188:SER:OG	2.28	0.48
1:D:3692:ALA:HA	1:D:3695:MET:HE3	1.95	0.48
1:D:4047:ASP:HA	1:D:4050:LYS:HG2	1.94	0.48
1:A:258:ARG:NH1	1:A:317:MET:HA	2.28	0.48
1:A:894:VAL:O	1:A:898:ILE:HG12	2.13	0.48
1:A:2261:ASP:OD1	1:A:2262:LEU:N	2.46	0.48
1:A:2318:ASN:HB3	1:A:2322:ARG:HH12	1.78	0.48
1:A:2918:GLU:HG3	1:A:2923:TYR:CE1	2.49	0.48
1:A:3230:GLN:N	1:A:3230:GLN:OE1	2.46	0.48
1:A:3317:THR:H	1:A:3320:LEU:HD12	1.78	0.48
1:B:1051:ARG:O	1:B:1055:ARG:HG2	2.13	0.48
1:C:1446:ILE:HG12	1:C:1542:ALA:HB2	1.96	0.48
1:C:2162:MET:N	1:C:2162:MET:HE2	2.28	0.48
1:C:2318:ASN:HB3	1:C:2322:ARG:HH12	1.78	0.48
1:C:3260:ARG:HD2	1:C:3260:ARG:O	2.14	0.48
1:D:164:PRO:HB3	1:D:169:ARG:HB2	1.95	0.48
1:D:866:PRO:HA	1:D:1009:ARG:NH2	2.28	0.48
1:D:894:VAL:O	1:D:898:ILE:HG12	2.13	0.48
1:D:1286:THR:OG1	1:D:1550:PRO:O	2.19	0.48
1:D:2581:ARG:HB2	1:D:2584:MET:HG3	1.96	0.48
1:D:3260:ARG:HD2	1:D:3260:ARG:O	2.14	0.48
1:D:4654:MET:HE2	1:D:4663:ARG:HD3	1.95	0.48
1:A:866:PRO:HA	1:A:1009:ARG:NH2	2.28	0.48
1:A:2723:LYS:HG2	1:A:2895:PHE:CZ	2.43	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2768:LYS:O	1:A:2772:ARG:HG3	2.14	0.48
1:B:1446:ILE:HG12	1:B:1542:ALA:HB2	1.96	0.48
1:B:3811:GLN:NE2	1:B:3828:LYS:HB3	2.28	0.48
1:C:866:PRO:HA	1:C:1009:ARG:NH2	2.28	0.48
1:C:4047:ASP:HA	1:C:4050:LYS:HG2	1.95	0.48
1:D:1436:GLN:H	1:D:1500:ARG:HH22	1.60	0.48
1:D:1790:LYS:O	1:D:1794:MET:HG3	2.13	0.48
1:D:3153:SER:O	1:D:3156:GLY:N	2.45	0.48
1:D:3285:TYR:HA	1:D:3288:LEU:HD23	1.96	0.48
1:D:4084:VAL:O	1:D:4088:HIS:HB3	2.13	0.48
1:A:15:ARG:HH21	1:A:110:HIS:HB3	1.77	0.48
1:A:1017:THR:O	1:A:1028:ARG:HA	2.13	0.48
1:A:1168:MET:HE1	1:A:1200:GLY:HA2	1.95	0.48
1:A:2770:ILE:HG13	1:A:2771:TYR:CD1	2.48	0.48
1:A:3153:SER:O	1:A:3156:GLY:N	2.45	0.48
1:A:3273:MET:HE3	1:A:3273:MET:HB2	1.73	0.48
1:A:3316:LYS:O	1:A:3317:THR:OG1	2.31	0.48
1:B:15:ARG:HH21	1:B:110:HIS:HB3	1.77	0.48
1:B:895:MET:O	1:B:899:GLU:HG2	2.12	0.48
1:B:1457:PHE:HA	1:B:1461:ARG:HH12	1.78	0.48
1:B:3152:ARG:CZ	1:B:3233:HIS:HA	2.44	0.48
1:B:4654:MET:HE2	1:B:4663:ARG:HD3	1.95	0.48
1:C:4265:LYS:O	1:C:4269:LYS:HG2	2.13	0.48
1:D:881:ILE:HG13	1:D:885:LEU:CD2	2.44	0.48
1:D:943:LEU:HD23	1:D:999:LEU:HD21	1.96	0.48
1:A:1790:LYS:O	1:A:1794:MET:HG3	2.13	0.48
1:A:2716:LYS:HD3	1:A:2791:ARG:HG3	1.94	0.48
1:A:2791:ARG:HH12	1:A:2795:GLY:C	2.22	0.48
1:A:4265:LYS:O	1:A:4269:LYS:HG2	2.13	0.48
1:A:4521:LYS:HE3	1:A:4562:GLU:HG3	1.96	0.48
1:B:4943:MET:HG2	1:B:4948:CYS:HB3	1.95	0.48
1:C:2581:ARG:HB2	1:C:2584:MET:HG3	1.96	0.48
1:C:2791:ARG:HH12	1:C:2795:GLY:C	2.22	0.48
1:C:2830:ASN:HB3	1:D:1549:SER:HB2	1.95	0.48
1:C:3285:TYR:HA	1:C:3288:LEU:HD23	1.96	0.48
1:D:2944:ASP:O	1:D:2948:ARG:NH1	2.47	0.48
1:A:2162:MET:HE2	1:A:2162:MET:N	2.28	0.48
1:A:3152:ARG:CZ	1:A:3233:HIS:HA	2.44	0.48
1:A:3729:ALA:HA	1:A:3732:HIS:CD2	2.49	0.48
1:B:2918:GLU:HA	1:B:2923:TYR:CD1	2.48	0.48
1:B:3068:LEU:O	1:B:3071:THR:OG1	2.25	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:3729:ALA:HA	1:B:3732:HIS:CD2	2.49	0.48
1:C:881:ILE:HG13	1:C:885:LEU:CD2	2.44	0.48
1:C:2770:ILE:HG13	1:C:2771:TYR:CD1	2.48	0.48
1:C:3692:ALA:HA	1:C:3695:MET:HE3	1.95	0.48
1:D:885:LEU:O	1:D:889:ILE:HG12	2.14	0.48
1:D:1168:MET:HE1	1:D:1200:GLY:HA2	1.95	0.48
1:D:2497:ARG:HH22	1:D:2878:THR:HB	1.79	0.48
1:D:3114:GLN:HE22	1:D:3115:HIS:CE1	2.32	0.48
1:D:3230:GLN:N	1:D:3230:GLN:OE1	2.46	0.48
1:D:3298:ARG:HB3	1:D:3302:PHE:HE1	1.79	0.48
1:D:3729:ALA:HA	1:D:3732:HIS:CD2	2.49	0.48
1:D:4113:THR:O	1:D:4117:THR:HG23	2.13	0.48
1:A:1502:ASN:O	1:D:2824:ARG:NH2	2.47	0.48
1:A:2341:ASN:OD1	1:A:2342:GLY:N	2.46	0.48
1:A:2497:ARG:HH22	1:A:2878:THR:HB	1.79	0.48
1:A:2798:MET:SD	1:B:1498:GLN:HG3	2.54	0.48
1:A:3260:ARG:HD2	1:A:3260:ARG:O	2.14	0.48
1:A:3304:GLN:HA	1:A:3307:ILE:HG12	1.95	0.48
1:A:4084:VAL:O	1:A:4088:HIS:HB3	2.13	0.48
1:A:4848:ILE:HD11	1:D:4818:TYR:HD1	1.79	0.48
1:B:1685:LEU:HD22	1:B:1706:LEU:HB2	1.95	0.48
1:B:2341:ASN:OD1	1:B:2342:GLY:N	2.46	0.48
1:B:2791:ARG:HH12	1:B:2795:GLY:C	2.22	0.48
1:B:3270:SER:HA	1:B:3273:MET:CE	2.37	0.48
1:B:3273:MET:HE3	1:B:3273:MET:HB2	1.73	0.48
1:C:842:GLN:HB2	1:C:1603:PHE:HB2	1.96	0.48
1:C:1957:LEU:O	1:C:1961:LYS:HG2	2.14	0.48
1:C:4026:LEU:HD12	1:C:4055:HIS:ND1	2.29	0.48
1:A:842:GLN:HB2	1:A:1603:PHE:HB2	1.96	0.48
1:A:943:LEU:HD23	1:A:999:LEU:HD21	1.96	0.48
1:A:2927:GLN:HG2	1:A:3003:MET:HE1	1.96	0.48
1:B:3230:GLN:OE1	1:B:3230:GLN:N	2.46	0.48
1:B:3316:LYS:HE2	1:B:3316:LYS:HB2	1.54	0.48
1:C:2918:GLU:HG3	1:C:2923:TYR:CE1	2.49	0.48
1:C:2927:GLN:HG2	1:C:3003:MET:HE1	1.96	0.48
1:C:3152:ARG:CZ	1:C:3233:HIS:HA	2.44	0.48
1:D:981:MET:N	1:D:981:MET:SD	2.86	0.48
1:D:1727:VAL:HG11	1:D:1926:VAL:HG21	1.95	0.48
1:D:2791:ARG:HH12	1:D:2795:GLY:C	2.22	0.48
1:D:2918:GLU:HG3	1:D:2923:TYR:CE1	2.49	0.48
1:D:3325:LYS:O	1:D:3328:LYS:HG2	2.14	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:986:ILE:HG22	1:A:1055:ARG:NH1	2.29	0.47
1:B:2142:MET:HE3	1:B:2146:LEU:HG	1.96	0.47
1:B:2497:ARG:HH22	1:B:2878:THR:HB	1.79	0.47
1:B:2768:LYS:O	1:B:2772:ARG:HG3	2.14	0.47
1:B:3260:ARG:O	1:B:3260:ARG:HD2	2.14	0.47
1:B:4521:LYS:HE3	1:B:4562:GLU:HG3	1.96	0.47
1:C:943:LEU:HD23	1:C:999:LEU:HD21	1.96	0.47
1:C:3298:ARG:HB3	1:C:3302:PHE:HE1	1.79	0.47
1:C:3304:GLN:HA	1:C:3307:ILE:HG12	1.95	0.47
1:C:3325:LYS:O	1:C:3328:LYS:HG2	2.14	0.47
1:C:3729:ALA:HA	1:C:3732:HIS:CD2	2.49	0.47
1:C:4521:LYS:HE3	1:C:4562:GLU:HG3	1.96	0.47
1:C:4819:VAL:HG12	1:C:4830:GLU:HG3	1.96	0.47
1:D:258:ARG:NH1	1:D:317:MET:HA	2.28	0.47
1:D:986:ILE:HG22	1:D:1055:ARG:NH1	2.29	0.47
1:D:1957:LEU:O	1:D:1961:LYS:HG2	2.14	0.47
1:D:2318:ASN:HB3	1:D:2322:ARG:HH12	1.78	0.47
1:D:2768:LYS:O	1:D:2772:ARG:HG3	2.14	0.47
1:D:2844:MET:HE3	1:D:2844:MET:HB2	1.71	0.47
1:D:3238:ILE:O	1:D:3241:MET:HB2	2.14	0.47
1:A:317:MET:HG2	1:A:321:LYS:HE2	1.96	0.47
1:A:1051:ARG:O	1:A:1055:ARG:HG2	2.13	0.47
1:A:1931:ASP:OD1	1:A:1932:PHE:N	2.47	0.47
1:A:3114:GLN:HE22	1:A:3115:HIS:CE1	2.32	0.47
1:A:3157:GLU:HG3	1:A:3302:PHE:HZ	1.77	0.47
1:A:3692:ALA:HA	1:A:3695:MET:HE3	1.95	0.47
1:A:4026:LEU:HD12	1:A:4055:HIS:ND1	2.29	0.47
1:B:2716:LYS:HD3	1:B:2791:ARG:HG3	1.94	0.47
1:B:2918:GLU:HG3	1:B:2923:TYR:CE1	2.49	0.47
1:B:4026:LEU:HD12	1:B:4055:HIS:ND1	2.29	0.47
1:C:164:PRO:HB3	1:C:169:ARG:HB2	1.95	0.47
1:C:258:ARG:NH1	1:C:317:MET:HA	2.28	0.47
1:C:1790:LYS:O	1:C:1794:MET:HG3	2.13	0.47
1:C:2768:LYS:O	1:C:2772:ARG:HG3	2.14	0.47
1:C:3230:GLN:OE1	1:C:3230:GLN:N	2.46	0.47
1:D:3304:GLN:HA	1:D:3307:ILE:HG12	1.95	0.47
1:D:3313:GLN:HA	1:D:3316:LYS:HZ1	1.79	0.47
1:A:1497:GLY:HA3	1:D:2798:MET:HE1	1.96	0.47
1:A:4654:MET:HE2	1:A:4663:ARG:HD3	1.95	0.47
1:A:4818:TYR:HD1	1:B:4848:ILE:HD11	1.79	0.47
1:B:258:ARG:NH1	1:B:317:MET:HA	2.28	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:317:MET:HG2	1:B:321:LYS:HE2	1.96	0.47
1:B:881:ILE:HG13	1:B:885:LEU:CD2	2.44	0.47
1:B:920:GLU:N	1:B:974:SER:OG	2.46	0.47
1:B:1300:MET:HE2	1:B:1300:MET:HB3	1.71	0.47
1:B:1931:ASP:OD1	1:B:1932:PHE:N	2.47	0.47
1:B:3250:TRP:CE3	1:B:3268:LEU:HB3	2.50	0.47
1:B:3285:TYR:HA	1:B:3288:LEU:HD23	1.96	0.47
1:B:4819:VAL:HG12	1:B:4830:GLU:HG3	1.96	0.47
1:C:986:ILE:HG22	1:C:1055:ARG:NH1	2.29	0.47
1:C:3187:LYS:O	1:C:3188:SER:OG	2.28	0.47
1:C:3250:TRP:CE3	1:C:3268:LEU:HB3	2.50	0.47
1:D:2927:GLN:HG2	1:D:3003:MET:HE1	1.96	0.47
1:D:3323:MET:HB3	1:D:3327:LYS:HZ3	1.77	0.47
1:A:614:LEU:HD22	1:A:632:ILE:HG12	1.97	0.47
1:A:1080:GLY:O	1:A:1081:THR:OG1	2.32	0.47
1:A:2705:PRO:HD2	1:A:2854:LYS:HD2	1.97	0.47
1:A:2720:PHE:CE1	1:A:2896:LEU:HA	2.49	0.47
1:A:4640:PHE:CG	1:A:4641:PRO:HD3	2.48	0.47
1:B:514:PHE:CD2	1:B:526:TRP:HB2	2.50	0.47
1:B:885:LEU:O	1:B:889:ILE:HG12	2.14	0.47
1:B:2705:PRO:HD2	1:B:2854:LYS:HD2	1.97	0.47
1:C:3238:ILE:O	1:C:3241:MET:HB2	2.14	0.47
1:D:317:MET:HB2	1:D:317:MET:HE2	1.39	0.47
1:D:614:LEU:HD22	1:D:632:ILE:HG12	1.97	0.47
1:D:620:CYS:HG	1:D:621:HIS:HD1	1.62	0.47
1:D:842:GLN:HB2	1:D:1603:PHE:HB2	1.96	0.47
1:D:1941:ARG:NH2	1:D:3609:TYR:HB2	2.27	0.47
1:A:394:HIS:CD2	1:A:395:HIS:H	2.33	0.47
1:A:891:GLU:HB3	1:A:978:PRO:HB3	1.97	0.47
1:A:3325:LYS:O	1:A:3328:LYS:HG2	2.14	0.47
1:B:866:PRO:HA	1:B:1009:ARG:NH2	2.28	0.47
1:B:1957:LEU:O	1:B:1961:LYS:HG2	2.14	0.47
1:B:2723:LYS:HG2	1:B:2895:PHE:CZ	2.43	0.47
1:B:2732:TRP:HA	1:B:2735:ASP:OD1	2.15	0.47
1:C:317:MET:HG2	1:C:321:LYS:HE2	1.96	0.47
1:C:3114:GLN:HE22	1:C:3115:HIS:CE1	2.32	0.47
1:C:4084:VAL:O	1:C:4088:HIS:HB3	2.13	0.47
1:D:630:HIS:CE1	1:D:1671:ARG:HE	2.33	0.47
1:D:1457:PHE:HA	1:D:1461:ARG:HH12	1.78	0.47
1:D:2686:VAL:O	1:D:2688:MET:HE2	2.15	0.47
1:A:514:PHE:CD2	1:A:526:TRP:HB2	2.50	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:885:LEU:O	1:A:889:ILE:HG12	2.14	0.47
1:B:3042:ALA:O	1:B:3046:MET:HG2	2.15	0.47
1:C:1500:ARG:HG3	1:C:1505:LEU:HB2	1.97	0.47
1:C:1714:TYR:CZ	1:C:1761:MET:HB2	2.50	0.47
1:C:1931:ASP:OD1	1:C:1932:PHE:N	2.47	0.47
1:C:2142:MET:HE3	1:C:2146:LEU:HG	1.96	0.47
1:D:2720:PHE:CE1	1:D:2896:LEU:HA	2.49	0.47
1:A:1446:ILE:HG12	1:A:1542:ALA:HB2	1.96	0.47
1:A:1685:LEU:HD22	1:A:1706:LEU:HB2	1.95	0.47
1:A:1957:LEU:O	1:A:1961:LYS:HG2	2.14	0.47
1:A:2142:MET:HE3	1:A:2146:LEU:HG	1.96	0.47
1:A:3298:ARG:HB3	1:A:3302:PHE:HE1	1.79	0.47
1:A:4765:LYS:HD2	1:A:4765:LYS:HA	1.73	0.47
1:B:394:HIS:CD2	1:B:395:HIS:H	2.33	0.47
1:B:943:LEU:HD23	1:B:999:LEU:HD21	1.96	0.47
1:B:1725:TYR:HE1	1:B:2101:ALA:HB1	1.80	0.47
1:B:2720:PHE:CE1	1:B:2896:LEU:HA	2.49	0.47
1:B:2846:GLU:HB3	1:B:2874:TYR:CD2	2.50	0.47
1:B:2944:ASP:O	1:B:2948:ARG:NH1	2.47	0.47
1:B:3304:GLN:HA	1:B:3307:ILE:HG12	1.96	0.47
1:B:3325:LYS:O	1:B:3328:LYS:HG2	2.14	0.47
1:C:614:LEU:HD22	1:C:632:ILE:HG12	1.97	0.47
1:C:620:CYS:HG	1:C:621:HIS:HD1	1.62	0.47
1:C:1725:TYR:HE1	1:C:2101:ALA:HB1	1.80	0.47
1:C:3042:ALA:O	1:C:3046:MET:HG2	2.15	0.47
1:D:317:MET:HG2	1:D:321:LYS:HE2	1.96	0.47
1:D:394:HIS:CD2	1:D:395:HIS:H	2.33	0.47
1:D:891:GLU:HB3	1:D:978:PRO:HB3	1.97	0.47
1:D:897:LYS:HD3	1:D:902:TRP:HE3	1.80	0.47
1:D:1446:ILE:HG12	1:D:1542:ALA:HB2	1.96	0.47
1:D:2142:MET:HE3	1:D:2146:LEU:HG	1.96	0.47
1:A:2944:ASP:O	1:A:2948:ARG:NH1	2.47	0.47
1:B:614:LEU:HD22	1:B:632:ILE:HG12	1.97	0.47
1:B:891:GLU:HB3	1:B:978:PRO:HB3	1.97	0.47
1:B:1286:THR:OG1	1:B:1550:PRO:O	2.19	0.47
1:B:1500:ARG:HG3	1:B:1505:LEU:HB2	1.97	0.47
1:B:3114:GLN:HE22	1:B:3115:HIS:CE1	2.32	0.47
1:B:4683:ARG:NH1	1:B:4684:GLU:O	2.48	0.47
1:C:885:LEU:O	1:C:889:ILE:HG12	2.14	0.47
1:C:2705:PRO:HD2	1:C:2854:LYS:HD2	1.97	0.47
1:C:2732:TRP:HA	1:C:2735:ASP:OD1	2.15	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3323:MET:HB3	1:C:3327:LYS:HZ3	1.78	0.47
1:D:920:GLU:OE1	1:D:974:SER:OG	2.25	0.47
1:D:3152:ARG:CZ	1:D:3233:HIS:HA	2.44	0.47
1:D:3250:TRP:CE3	1:D:3268:LEU:HB3	2.50	0.47
1:D:3861:GLN:HB3	1:D:3864:ASN:HD22	1.80	0.47
1:A:2732:TRP:HA	1:A:2735:ASP:OD1	2.15	0.47
1:A:3250:TRP:CE3	1:A:3268:LEU:HB3	2.50	0.47
1:B:986:ILE:HG22	1:B:1055:ARG:NH1	2.29	0.47
1:B:1714:TYR:CZ	1:B:1761:MET:HB2	2.50	0.47
1:B:2927:GLN:HG2	1:B:3003:MET:HE1	1.96	0.47
1:B:3238:ILE:O	1:B:3241:MET:HB2	2.14	0.47
1:C:514:PHE:CD2	1:C:526:TRP:HB2	2.50	0.47
1:C:920:GLU:OE1	1:C:974:SER:OG	2.25	0.47
1:C:2497:ARG:HH22	1:C:2878:THR:HB	1.79	0.47
1:C:2944:ASP:O	1:C:2948:ARG:NH1	2.47	0.47
1:D:514:PHE:CD2	1:D:526:TRP:HB2	2.50	0.47
1:D:4521:LYS:HE3	1:D:4562:GLU:HG3	1.96	0.47
1:A:1500:ARG:HG3	1:A:1505:LEU:HB2	1.97	0.47
1:A:1725:TYR:HE1	1:A:2101:ALA:HB1	1.80	0.47
1:A:2801:TYR:OH	1:B:1495:SER:O	2.24	0.47
1:A:3238:ILE:O	1:A:3241:MET:HB2	2.14	0.47
1:A:3250:TRP:O	1:A:3256:ASN:ND2	2.49	0.47
1:A:4484:ILE:HA	1:D:4279:MET:HE2	1.96	0.47
1:B:1979:PHE:CZ	1:B:1996:LEU:HD23	2.50	0.47
1:C:317:MET:HB2	1:C:317:MET:HE2	1.40	0.47
1:C:891:GLU:HB3	1:C:978:PRO:HB3	1.97	0.47
1:C:2868:HIS:CD2	1:C:2870:LEU:HB2	2.50	0.47
1:C:4683:ARG:NH1	1:C:4684:GLU:O	2.48	0.47
1:D:1931:ASP:OD1	1:D:1932:PHE:N	2.47	0.47
1:D:4026:LEU:HD12	1:D:4055:HIS:ND1	2.29	0.47
1:A:59:PRO:HG3	1:A:296:ARG:CZ	2.45	0.46
1:A:630:HIS:CE1	1:A:1671:ARG:HE	2.33	0.46
1:A:1979:PHE:CZ	1:A:1996:LEU:HD23	2.50	0.46
1:A:3285:TYR:HA	1:A:3288:LEU:HD23	1.96	0.46
1:B:3298:ARG:HB3	1:B:3302:PHE:HE1	1.79	0.46
1:C:59:PRO:HG3	1:C:296:ARG:CZ	2.46	0.46
1:C:2686:VAL:O	1:C:2688:MET:HE2	2.15	0.46
1:C:3090:VAL:HA	1:C:3093:ILE:HG12	1.98	0.46
1:C:3861:GLN:HB3	1:C:3864:ASN:HD22	1.80	0.46
1:C:4055:HIS:CD2	1:C:4057:HIS:HD1	2.34	0.46
1:C:4567:TYR:O	1:C:4570:PRO:HD2	2.15	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3316:LYS:O	1:D:3317:THR:OG1	2.31	0.46
1:D:4567:TYR:O	1:D:4570:PRO:HD2	2.15	0.46
1:D:4683:ARG:NH1	1:D:4684:GLU:O	2.48	0.46
1:D:4819:VAL:HG12	1:D:4830:GLU:HG3	1.96	0.46
1:A:2716:LYS:NZ	1:A:2791:ARG:HB2	2.30	0.46
1:B:59:PRO:HG3	1:B:296:ARG:CZ	2.45	0.46
1:B:842:GLN:HB2	1:B:1603:PHE:HB2	1.96	0.46
1:C:1979:PHE:CZ	1:C:1996:LEU:HD23	2.50	0.46
1:C:2716:LYS:NZ	1:C:2791:ARG:HB2	2.31	0.46
1:C:2720:PHE:CE1	1:C:2896:LEU:HA	2.49	0.46
1:C:4625:ASP:OD1	1:C:4625:ASP:N	2.44	0.46
1:D:258:ARG:HH12	1:D:317:MET:HA	1.80	0.46
1:D:2705:PRO:HD2	1:D:2854:LYS:HD2	1.97	0.46
1:D:3250:TRP:O	1:D:3256:ASN:ND2	2.48	0.46
1:A:521:GLU:H	1:A:521:GLU:CD	2.24	0.46
1:A:3042:ALA:O	1:A:3046:MET:HG2	2.15	0.46
1:A:4055:HIS:CD2	1:A:4057:HIS:HD1	2.34	0.46
1:A:4819:VAL:HG12	1:A:4830:GLU:HG3	1.96	0.46
1:B:630:HIS:CE1	1:B:1671:ARG:HE	2.33	0.46
1:B:3316:LYS:O	1:B:3317:THR:OG1	2.31	0.46
1:C:3297:LYS:HZ2	1:C:3335:SER:H	1.62	0.46
1:C:3890:TRP:HB3	1:D:76:ARG:HG2	1.97	0.46
1:D:59:PRO:HG3	1:D:296:ARG:CZ	2.45	0.46
1:D:1500:ARG:HG3	1:D:1505:LEU:HB2	1.97	0.46
1:D:2732:TRP:HA	1:D:2735:ASP:OD1	2.15	0.46
1:D:2846:GLU:HB3	1:D:2874:TYR:CD2	2.50	0.46
1:D:3042:ALA:O	1:D:3046:MET:HG2	2.15	0.46
2:G:9:SER:HB3	2:G:72:ARG:HB3	1.97	0.46
1:A:897:LYS:HD3	1:A:902:TRP:HE3	1.80	0.46
1:A:1172:THR:HB	1:A:1190:LEU:HD22	1.98	0.46
1:A:2857:LYS:HG2	1:A:2861:GLU:OE2	2.16	0.46
1:A:2925:PHE:HZ	1:A:2970:LEU:HD13	1.80	0.46
1:A:3861:GLN:HB3	1:A:3864:ASN:HD22	1.80	0.46
1:B:1172:THR:HB	1:B:1190:LEU:HD22	1.98	0.46
1:B:2844:MET:HE3	1:B:2844:MET:HB2	1.71	0.46
1:B:4567:TYR:O	1:B:4570:PRO:HD2	2.15	0.46
1:B:4863:GLN:HG2	1:C:4860:ALA:HB2	1.98	0.46
1:C:897:LYS:HD3	1:C:902:TRP:HE3	1.80	0.46
1:C:1906:MET:HE3	1:C:1906:MET:HB3	1.83	0.46
1:D:1714:TYR:CZ	1:D:1761:MET:HB2	2.50	0.46
1:D:4055:HIS:CD2	1:D:4057:HIS:HD1	2.34	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:H:9:SER:HB3	2:H:72:ARG:HB3	1.97	0.46
1:A:2926:LEU:HD21	1:A:2975:PHE:HZ	1.81	0.46
1:B:2857:LYS:HG2	1:B:2861:GLU:OE2	2.16	0.46
1:C:28:ILE:O	1:C:31:GLU:HG3	2.15	0.46
1:C:1172:THR:HB	1:C:1190:LEU:HD22	1.98	0.46
1:C:2846:GLU:HB3	1:C:2874:TYR:CD2	2.50	0.46
1:D:776:GLN:HG2	1:D:1472:GLU:HA	1.98	0.46
1:D:1725:TYR:HE1	1:D:2101:ALA:HB1	1.80	0.46
1:D:2232:PRO:C	1:D:2234:MET:H	2.24	0.46
1:D:2857:LYS:HG2	1:D:2861:GLU:OE2	2.16	0.46
1:D:3270:SER:HA	1:D:3273:MET:CE	2.37	0.46
2:F:50:ARG:N	2:F:55:GLU:OE1	2.42	0.46
1:A:1682:GLU:HB3	1:A:1683:PRO:HD3	1.97	0.46
1:B:28:ILE:O	1:B:31:GLU:HG3	2.15	0.46
1:B:1682:GLU:HB3	1:B:1683:PRO:HD3	1.97	0.46
1:B:2071:GLN:CD	1:B:3648:GLY:HA3	2.41	0.46
1:C:394:HIS:CD2	1:C:395:HIS:H	2.33	0.46
1:C:2772:ARG:O	1:C:2776:LYS:HG3	2.16	0.46
1:C:2925:PHE:HZ	1:C:2970:LEU:HD13	1.80	0.46
1:D:521:GLU:H	1:D:521:GLU:CD	2.24	0.46
1:D:1172:THR:HB	1:D:1190:LEU:HD22	1.98	0.46
1:D:2716:LYS:NZ	1:D:2791:ARG:HB2	2.31	0.46
1:A:317:MET:HB2	1:A:317:MET:HE2	1.39	0.46
1:A:2846:GLU:HB3	1:A:2874:TYR:CD2	2.50	0.46
1:B:1080:GLY:O	1:B:1081:THR:OG1	2.32	0.46
1:B:2686:VAL:O	1:B:2688:MET:HE2	2.15	0.46
1:B:2776:LYS:HB3	1:B:2780:LYS:NZ	2.31	0.46
1:B:3250:TRP:O	1:B:3256:ASN:ND2	2.48	0.46
1:B:3861:GLN:HB3	1:B:3864:ASN:HD22	1.80	0.46
1:C:521:GLU:CD	1:C:521:GLU:H	2.24	0.46
1:C:801:ARG:NH1	1:C:1615:GLN:O	2.40	0.46
1:C:3250:TRP:O	1:C:3256:ASN:ND2	2.49	0.46
1:D:441:LYS:HD2	1:D:442:ALA:H	1.81	0.46
1:D:1241:VAL:HG12	1:D:1807:ARG:HH12	1.81	0.46
1:D:2620:TYR:HB2	1:D:2627:TRP:HD1	1.81	0.46
1:D:2868:HIS:CD2	1:D:2870:LEU:HB2	2.50	0.46
1:D:2925:PHE:HZ	1:D:2970:LEU:HD13	1.80	0.46
2:F:9:SER:HB3	2:F:72:ARG:HB3	1.97	0.46
1:A:28:ILE:HG22	1:A:29:HIS:CD2	2.51	0.46
1:A:776:GLN:HG2	1:A:1472:GLU:HA	1.98	0.46
1:A:1714:TYR:CE2	1:A:1718:ARG:HD2	2.51	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2686:VAL:O	1:A:2688:MET:HE2	2.15	0.46
2:E:78:THR:OG1	2:E:80:ASP:OD1	2.26	0.46
1:B:28:ILE:HG22	1:B:29:HIS:CD2	2.51	0.46
1:B:4055:HIS:CD2	1:B:4057:HIS:HD1	2.34	0.46
1:B:4689:LYS:NZ	1:B:4696:ALA:HB2	2.31	0.46
1:C:630:HIS:CE1	1:C:1671:ARG:HE	2.33	0.46
1:D:3090:VAL:HA	1:D:3093:ILE:HG12	1.98	0.46
2:H:50:ARG:N	2:H:55:GLU:OE1	2.42	0.46
1:A:1714:TYR:CZ	1:A:1761:MET:HB2	2.50	0.46
1:A:4683:ARG:NH1	1:A:4684:GLU:O	2.48	0.46
1:A:4831:ILE:HG13	1:A:4843:ARG:NH2	2.31	0.46
1:B:2716:LYS:NZ	1:B:2791:ARG:HB2	2.31	0.46
1:B:2975:PHE:O	1:B:2979:ARG:HB3	2.16	0.46
1:B:3090:VAL:HA	1:B:3093:ILE:HG12	1.98	0.46
1:C:258:ARG:HH12	1:C:317:MET:HA	1.80	0.46
1:C:2232:PRO:C	1:C:2234:MET:H	2.24	0.46
1:C:2620:TYR:HB2	1:C:2627:TRP:HD1	1.81	0.46
1:C:4827:ILE:O	1:C:4831:ILE:HG12	2.16	0.46
1:D:1682:GLU:HB3	1:D:1683:PRO:HD3	1.97	0.46
1:A:2481:GLN:NE2	1:A:2537:GLY:O	2.42	0.46
1:A:4256:MET:HE2	1:A:4256:MET:HB3	1.72	0.46
1:B:441:LYS:HD2	1:B:442:ALA:H	1.81	0.46
1:B:2772:ARG:O	1:B:2776:LYS:HG3	2.16	0.46
1:B:4827:ILE:O	1:B:4831:ILE:HG12	2.16	0.46
1:C:441:LYS:HD2	1:C:442:ALA:H	1.81	0.46
1:C:1241:VAL:HG12	1:C:1807:ARG:HH12	1.81	0.46
1:C:1682:GLU:HB3	1:C:1683:PRO:HD3	1.97	0.46
1:C:2857:LYS:HG2	1:C:2861:GLU:OE2	2.16	0.46
1:D:28:ILE:O	1:D:31:GLU:HG3	2.15	0.46
1:D:235:ARG:NH2	1:D:412:GLU:OE2	2.27	0.46
1:D:808:HIS:O	1:D:1616:GLY:HA2	2.16	0.46
1:A:28:ILE:O	1:A:31:GLU:HG3	2.15	0.45
1:A:258:ARG:HH12	1:A:317:MET:HA	1.80	0.45
1:A:2798:MET:HE1	1:B:1497:GLY:CA	2.39	0.45
1:A:4567:TYR:O	1:A:4570:PRO:HD2	2.15	0.45
1:B:1714:TYR:CE2	1:B:1718:ARG:HD2	2.51	0.45
1:B:1906:MET:HE3	1:B:1906:MET:HB3	1.83	0.45
1:B:2926:LEU:HD21	1:B:2975:PHE:HZ	1.81	0.45
1:C:1714:TYR:CE2	1:C:1718:ARG:HD2	2.51	0.45
1:C:2445:ILE:HA	1:C:2451:VAL:HA	1.98	0.45
1:C:4689:LYS:NZ	1:C:4696:ALA:HB2	2.31	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:749:LEU:HD22	1:D:755:ILE:HD11	1.98	0.45
1:D:1444:GLY:HA3	1:D:1487:MET:HA	1.98	0.45
1:A:713:TRP:CE2	1:A:1600:PRO:HD3	2.52	0.45
1:A:808:HIS:O	1:A:1616:GLY:HA2	2.16	0.45
1:A:1241:VAL:HG12	1:A:1807:ARG:HH12	1.81	0.45
1:A:2957:GLU:HA	1:A:2960:ILE:HG22	1.98	0.45
1:A:2966:VAL:HG12	1:A:2970:LEU:HD23	1.99	0.45
1:A:4827:ILE:O	1:A:4831:ILE:HG12	2.16	0.45
1:B:713:TRP:CE2	1:B:1600:PRO:HD3	2.52	0.45
1:B:1628:MET:HE2	1:B:1684:GLN:HB3	1.99	0.45
1:B:2137:GLU:H	1:B:2137:GLU:CD	2.22	0.45
1:B:3076:LYS:NZ	1:B:3140:LEU:HD23	2.31	0.45
1:B:3297:LYS:HZ2	1:B:3335:SER:H	1.65	0.45
1:C:1444:GLY:HA3	1:C:1487:MET:HA	1.99	0.45
1:C:3122:ILE:HG12	1:C:3127:GLN:HG2	1.99	0.45
1:D:981:MET:HE2	1:D:1059:GLY:C	2.41	0.45
1:D:4831:ILE:HG13	1:D:4843:ARG:NH2	2.31	0.45
1:A:2071:GLN:CD	1:A:3648:GLY:HA3	2.41	0.45
1:A:2670:SER:HB2	1:A:2973:GLN:HG2	1.98	0.45
1:A:2975:PHE:O	1:A:2979:ARG:HB3	2.16	0.45
1:B:1241:VAL:HG12	1:B:1807:ARG:HH12	1.81	0.45
1:C:606:ARG:HH21	1:C:1633:PRO:HD2	1.82	0.45
1:C:808:HIS:O	1:C:1616:GLY:HA2	2.16	0.45
1:C:2353:ILE:HG12	1:C:2359:ARG:HE	1.82	0.45
1:C:2975:PHE:O	1:C:2979:ARG:HB3	2.16	0.45
1:C:3076:LYS:NZ	1:C:3140:LEU:HD23	2.31	0.45
1:D:1979:PHE:CZ	1:D:1996:LEU:HD23	2.51	0.45
1:D:3076:LYS:NZ	1:D:3140:LEU:HD23	2.31	0.45
1:A:198:ASN:OD1	1:A:199:GLY:N	2.50	0.45
1:A:981:MET:HE2	1:A:1059:GLY:C	2.41	0.45
1:A:2137:GLU:H	1:A:2137:GLU:CD	2.22	0.45
1:A:2776:LYS:HB3	1:A:2780:LYS:NZ	2.31	0.45
1:A:3939:ARG:NH1	1:B:173:GLU:HG2	2.32	0.45
1:B:494:MET:HE3	1:B:494:MET:HB3	1.76	0.45
1:B:897:LYS:HD3	1:B:902:TRP:HE3	1.80	0.45
1:B:1043:LYS:O	1:B:1047:LYS:HB2	2.17	0.45
1:B:1444:GLY:HA3	1:B:1487:MET:HA	1.99	0.45
1:B:2353:ILE:HG12	1:B:2359:ARG:HE	1.82	0.45
1:C:713:TRP:CE2	1:C:1600:PRO:HD3	2.52	0.45
1:C:2071:GLN:CD	1:C:3648:GLY:HA3	2.41	0.45
1:C:2926:LEU:HD21	1:C:2975:PHE:HZ	1.80	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4874:ARG:NH1	1:D:4868:ASP:OD1	2.50	0.45
1:D:29:HIS:CE1	1:D:31:GLU:HG2	2.52	0.45
1:D:1031:ARG:HH11	1:D:1042:THR:HG21	1.81	0.45
1:D:1080:GLY:O	1:D:1081:THR:OG1	2.31	0.45
1:D:1957:LEU:HA	1:D:1960:ARG:CZ	2.47	0.45
1:D:2772:ARG:O	1:D:2776:LYS:HG3	2.16	0.45
1:D:2926:LEU:HD21	1:D:2975:PHE:HZ	1.81	0.45
1:A:1957:LEU:HA	1:A:1960:ARG:CZ	2.47	0.45
1:A:3025:ASP:O	1:A:3029:ILE:HD12	2.17	0.45
1:A:3076:LYS:NZ	1:A:3140:LEU:HD23	2.31	0.45
1:B:258:ARG:HH12	1:B:317:MET:HA	1.80	0.45
1:B:907:VAL:HG11	1:B:912:LYS:HD3	1.98	0.45
1:B:2674:GLY:HA2	1:B:2978:HIS:CE1	2.52	0.45
1:B:4906:GLU:OE1	1:C:4183:LYS:HE3	2.17	0.45
1:C:2326:ARG:HD3	1:C:2326:ARG:HA	1.76	0.45
1:C:3025:ASP:O	1:C:3029:ILE:HD12	2.17	0.45
1:C:3213:LYS:HA	1:C:3216:GLU:CD	2.42	0.45
1:D:661:LEU:HD13	1:D:673:TRP:CD1	2.52	0.45
1:D:713:TRP:CE2	1:D:1600:PRO:HD3	2.52	0.45
1:D:2445:ILE:HA	1:D:2451:VAL:HA	1.98	0.45
1:D:2776:LYS:HB3	1:D:2780:LYS:NZ	2.31	0.45
1:D:2975:PHE:O	1:D:2979:ARG:HB3	2.16	0.45
1:D:3025:ASP:O	1:D:3029:ILE:HD12	2.17	0.45
1:D:3316:LYS:HE2	1:D:3316:LYS:HB2	1.53	0.45
1:A:1628:MET:HE2	1:A:1684:GLN:HB3	1.99	0.45
1:A:2232:PRO:C	1:A:2234:MET:H	2.24	0.45
1:A:2620:TYR:HB2	1:A:2627:TRP:HD1	1.81	0.45
1:A:2772:ARG:O	1:A:2776:LYS:HG3	2.16	0.45
1:B:749:LEU:HD22	1:B:755:ILE:HD11	1.98	0.45
1:B:897:LYS:HE2	1:B:917:CYS:HB3	1.99	0.45
1:B:981:MET:HE2	1:B:1059:GLY:C	2.41	0.45
1:B:2620:TYR:HB2	1:B:2627:TRP:HD1	1.81	0.45
1:B:2957:GLU:HA	1:B:2960:ILE:HG22	1.98	0.45
1:C:2776:LYS:HB3	1:C:2780:LYS:NZ	2.31	0.45
1:D:1628:MET:HE3	1:D:1687:TYR:HD2	1.82	0.45
1:D:3124:GLU:C	1:D:3126:VAL:H	2.25	0.45
1:D:4667:SER:OG	1:D:4672:MET:O	2.34	0.45
1:A:712:GLU:OE1	1:A:1086:ARG:NH2	2.39	0.45
1:A:1628:MET:HE3	1:A:1687:TYR:HD2	1.82	0.45
1:A:3122:ILE:HG12	1:A:3127:GLN:HG2	1.99	0.45
1:A:4262:LYS:HE3	1:B:4698:LEU:HD22	1.99	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:776:GLN:HG2	1:B:1472:GLU:HA	1.98	0.45
1:B:2670:SER:HB2	1:B:2973:GLN:HG2	1.99	0.45
1:B:4831:ILE:HG13	1:B:4843:ARG:NH2	2.31	0.45
1:D:2670:SER:HB2	1:D:2973:GLN:HG2	1.98	0.45
1:D:3796:LEU:HD22	1:D:3835:PHE:HZ	1.82	0.45
1:A:271:ALA:HB2	1:A:488:LEU:HD22	1.99	0.45
1:A:674:TYR:CE1	1:A:756:SER:HB2	2.52	0.45
1:A:1444:GLY:HA3	1:A:1487:MET:HA	1.99	0.45
1:A:1500:ARG:HG3	1:A:1505:LEU:H	1.82	0.45
1:A:2868:HIS:CD2	1:A:2870:LEU:HB2	2.50	0.45
1:A:3042:ALA:HB3	1:A:3117:PHE:HD2	1.82	0.45
1:A:4795:LYS:HD2	1:A:4795:LYS:HA	1.81	0.45
1:B:521:GLU:H	1:B:521:GLU:CD	2.24	0.45
1:B:674:TYR:CE1	1:B:756:SER:HB2	2.52	0.45
1:B:801:ARG:NH1	1:B:1615:GLN:O	2.40	0.45
1:B:1500:ARG:HG3	1:B:1505:LEU:H	1.82	0.45
1:B:2724:TYR:CD1	1:B:2895:PHE:HD1	2.35	0.45
1:C:756:SER:OG	1:C:769:ARG:HB2	2.17	0.45
1:C:776:GLN:HG2	1:C:1472:GLU:HA	1.98	0.45
1:C:981:MET:HE2	1:C:1059:GLY:C	2.41	0.45
1:C:2957:GLU:HA	1:C:2960:ILE:HG22	1.98	0.45
1:C:3273:MET:HE3	1:C:3273:MET:HB2	1.73	0.45
1:D:907:VAL:HG11	1:D:912:LYS:HD3	1.98	0.45
1:D:1694:MET:HE2	1:D:1694:MET:HB2	1.87	0.45
1:D:2071:GLN:CD	1:D:3648:GLY:HA3	2.41	0.45
1:D:2353:ILE:HG12	1:D:2359:ARG:HE	1.82	0.45
1:D:2782:MET:HG2	1:D:2787:TRP:HE3	1.82	0.45
2:G:18:LYS:NZ	2:G:18:LYS:HB3	2.32	0.45
1:A:3090:VAL:HA	1:A:3093:ILE:HG12	1.98	0.45
1:A:3213:LYS:HA	1:A:3216:GLU:CD	2.42	0.45
1:A:3741:LEU:HD11	1:A:3777:MET:HG2	1.99	0.45
1:A:3796:LEU:HD22	1:A:3835:PHE:HZ	1.82	0.45
2:E:18:LYS:NZ	2:E:18:LYS:HB3	2.32	0.45
1:B:2445:ILE:HA	1:B:2451:VAL:HA	1.98	0.45
1:B:2868:HIS:CD2	1:B:2870:LEU:HB2	2.50	0.45
1:B:3042:ALA:HB3	1:B:3117:PHE:HD2	1.82	0.45
1:B:4055:HIS:CD2	1:B:4057:HIS:ND1	2.85	0.45
1:B:4667:SER:OG	1:B:4672:MET:O	2.34	0.45
1:C:70:GLU:OE2	1:C:122:ARG:HD3	2.17	0.45
1:C:1144:ARG:HB3	1:C:1152:TYR:HB3	1.99	0.45
1:C:1628:MET:HE3	1:C:1687:TYR:HD2	1.82	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2674:GLY:HA2	1:C:2978:HIS:CE1	2.52	0.45
1:C:3124:GLU:C	1:C:3126:VAL:H	2.25	0.45
1:C:3316:LYS:O	1:C:3317:THR:OG1	2.31	0.45
1:D:1714:TYR:CE2	1:D:1718:ARG:HD2	2.51	0.45
1:D:4897:TYR:OH	1:D:4963:GLU:OE2	2.32	0.45
1:A:2758:LYS:HE2	1:A:2763:LEU:HA	1.99	0.45
1:A:4054:SER:O	1:A:4056:LYS:HG3	2.17	0.45
1:B:606:ARG:HH21	1:B:1633:PRO:HD2	1.82	0.45
1:B:808:HIS:O	1:B:1616:GLY:HA2	2.16	0.45
1:B:950:VAL:HA	1:B:1064:LEU:HA	1.99	0.45
1:B:1031:ARG:HH11	1:B:1042:THR:HG21	1.81	0.45
1:B:2481:GLN:NE2	1:B:2537:GLY:O	2.42	0.45
1:B:2925:PHE:HZ	1:B:2970:LEU:HD13	1.80	0.45
1:B:3796:LEU:HD22	1:B:3835:PHE:HZ	1.82	0.45
1:C:29:HIS:CE1	1:C:31:GLU:HG2	2.52	0.45
1:C:1031:ARG:HH11	1:C:1042:THR:HG21	1.81	0.45
1:C:2724:TYR:CD1	1:C:2895:PHE:HD1	2.35	0.45
1:D:801:ARG:NH1	1:D:1615:GLN:O	2.40	0.45
1:D:826:VAL:HG21	1:D:832:LEU:HB2	1.99	0.45
1:D:1144:ARG:HB3	1:D:1152:TYR:HB3	1.99	0.45
1:D:1785:ASP:OD1	1:D:1786:ILE:N	2.48	0.45
1:D:3122:ILE:HG12	1:D:3127:GLN:HG2	1.99	0.45
1:D:4827:ILE:O	1:D:4831:ILE:HG12	2.16	0.45
2:G:50:ARG:N	2:G:55:GLU:OE1	2.42	0.45
1:A:29:HIS:CE1	1:A:31:GLU:HG2	2.52	0.44
1:A:441:LYS:HD2	1:A:442:ALA:H	1.81	0.44
1:A:2353:ILE:HG12	1:A:2359:ARG:HE	1.82	0.44
1:A:2674:GLY:HA2	1:A:2978:HIS:CE1	2.52	0.44
1:A:4607:ALA:HB1	1:A:4649:VAL:HG21	1.99	0.44
1:B:661:LEU:HD13	1:B:673:TRP:CD1	2.52	0.44
1:B:2966:VAL:HG12	1:B:2970:LEU:HD23	1.99	0.44
1:C:28:ILE:HG22	1:C:29:HIS:CD2	2.51	0.44
1:C:749:LEU:HD22	1:C:755:ILE:HD11	1.98	0.44
1:C:826:VAL:HG21	1:C:832:LEU:HB2	1.99	0.44
1:C:1007:TRP:HE1	1:C:1011:ARG:NH1	2.16	0.44
1:C:1628:MET:HE2	1:C:1684:GLN:HB3	1.99	0.44
1:C:3062:ASP:O	1:C:3066:GLU:OE1	2.35	0.44
1:C:3830:LEU:HB3	1:C:3833:ASP:OD2	2.18	0.44
1:C:4055:HIS:CD2	1:C:4057:HIS:ND1	2.85	0.44
1:C:4667:SER:OG	1:C:4672:MET:O	2.34	0.44
2:G:45:LYS:HA	2:G:46:PRO:HD3	1.82	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:661:LEU:HD13	1:A:673:TRP:CD1	2.52	0.44
1:A:2445:ILE:HA	1:A:2451:VAL:HA	1.98	0.44
1:A:4689:LYS:NZ	1:A:4696:ALA:HB2	2.31	0.44
1:B:2965:LYS:HA	1:B:2965:LYS:HD3	1.80	0.44
1:B:3062:ASP:O	1:B:3066:GLU:OE1	2.35	0.44
1:C:674:TYR:CE1	1:C:756:SER:HB2	2.52	0.44
1:C:895:MET:HE3	1:C:978:PRO:HG2	2.00	0.44
1:C:2758:LYS:HE2	1:C:2763:LEU:HA	1.99	0.44
1:C:3796:LEU:HD22	1:C:3835:PHE:HZ	1.82	0.44
1:D:70:GLU:OE2	1:D:122:ARG:HD3	2.17	0.44
1:D:3044:THR:HA	1:D:3047:LYS:HZ3	1.82	0.44
1:D:3213:LYS:HA	1:D:3216:GLU:CD	2.42	0.44
1:D:3830:LEU:HB3	1:D:3833:ASP:OD2	2.18	0.44
2:F:18:LYS:NZ	2:F:18:LYS:HB3	2.32	0.44
2:H:18:LYS:NZ	2:H:18:LYS:HB3	2.32	0.44
1:A:537:LEU:O	1:A:541:ILE:HG12	2.18	0.44
1:A:756:SER:OG	1:A:769:ARG:HB2	2.17	0.44
1:A:895:MET:HE3	1:A:978:PRO:HG2	2.00	0.44
1:A:1043:LYS:O	1:A:1047:LYS:HB2	2.17	0.44
1:A:2724:TYR:CD1	1:A:2895:PHE:HD1	2.35	0.44
2:E:9:SER:HB3	2:E:72:ARG:HB3	1.97	0.44
1:B:1785:ASP:OD1	1:B:1786:ILE:N	2.48	0.44
1:B:3044:THR:HA	1:B:3047:LYS:NZ	2.32	0.44
1:B:3830:LEU:HB3	1:B:3833:ASP:OD2	2.18	0.44
1:C:1043:LYS:O	1:C:1047:LYS:HB2	2.17	0.44
1:D:2884:LYS:HD2	1:D:2885:ASP:N	2.33	0.44
2:F:12:ASP:OD1	2:F:13:GLY:N	2.51	0.44
1:A:370:LEU:HB2	1:A:393:MET:CE	2.48	0.44
1:A:749:LEU:HD22	1:A:755:ILE:HD11	1.98	0.44
1:A:907:VAL:HG11	1:A:912:LYS:HD3	1.98	0.44
1:A:1697:LEU:HD23	1:A:1697:LEU:HA	1.87	0.44
1:A:2844:MET:HE3	1:A:2844:MET:HB2	1.71	0.44
1:A:2926:LEU:CB	1:A:3003:MET:HE2	2.47	0.44
1:A:3044:THR:HA	1:A:3047:LYS:NZ	2.32	0.44
1:A:3882:GLN:NE2	1:A:3946:GLY:HA3	2.32	0.44
1:A:4056:LYS:HE3	1:B:4660:PHE:O	2.17	0.44
1:B:70:GLU:OE2	1:B:122:ARG:HD3	2.17	0.44
1:B:756:SER:OG	1:B:769:ARG:HB2	2.17	0.44
1:B:3076:LYS:HZ3	1:B:3140:LEU:HD23	1.82	0.44
1:B:4054:SER:O	1:B:4056:LYS:HG3	2.17	0.44
1:C:235:ARG:NH2	1:C:412:GLU:OE2	2.27	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:537:LEU:O	1:C:541:ILE:HG12	2.18	0.44
1:C:897:LYS:HE2	1:C:917:CYS:HB3	1.99	0.44
1:C:963:LYS:NZ	1:C:979:ALA:HB3	2.33	0.44
1:D:28:ILE:HG22	1:D:29:HIS:CD2	2.51	0.44
1:D:370:LEU:HB2	1:D:393:MET:CE	2.47	0.44
1:D:537:LEU:O	1:D:541:ILE:HG12	2.18	0.44
1:D:606:ARG:HH21	1:D:1633:PRO:HD2	1.82	0.44
1:D:1991:GLU:HG2	1:D:1992:ILE:N	2.32	0.44
1:D:3044:THR:HA	1:D:3047:LYS:NZ	2.32	0.44
1:A:962:LYS:NZ	1:A:982:ASP:H	2.16	0.44
1:A:1031:ARG:HH11	1:A:1042:THR:HG21	1.81	0.44
1:A:1991:GLU:HG2	1:A:1992:ILE:N	2.32	0.44
1:A:2826:ILE:HG23	1:B:1501:ASN:O	2.17	0.44
1:A:3184:TYR:CE1	1:A:3197:LEU:HD21	2.53	0.44
1:B:962:LYS:NZ	1:B:982:ASP:H	2.16	0.44
1:B:1628:MET:HE3	1:B:1687:TYR:HD2	1.82	0.44
1:B:1957:LEU:HA	1:B:1960:ARG:CZ	2.47	0.44
1:B:2884:LYS:HD2	1:B:2885:ASP:N	2.33	0.44
1:B:3124:GLU:C	1:B:3126:VAL:H	2.25	0.44
1:C:200:SER:O	1:C:202:HIS:HD2	2.01	0.44
1:C:661:LEU:HD13	1:C:673:TRP:CD1	2.52	0.44
1:C:1848:PRO:HG3	1:C:1890:LYS:HD2	2.00	0.44
1:C:3178:HIS:CE1	1:C:3263:MET:HA	2.53	0.44
1:C:4194:GLU:HG2	1:C:4645:TRP:HZ3	1.82	0.44
1:D:897:LYS:HE2	1:D:917:CYS:HB3	1.99	0.44
1:D:2397:ASP:O	1:D:2401:ARG:HG3	2.18	0.44
1:D:2926:LEU:CB	1:D:3003:MET:HE2	2.48	0.44
1:D:2957:GLU:HA	1:D:2960:ILE:HG22	1.98	0.44
1:D:3184:TYR:CE1	1:D:3197:LEU:HD21	2.53	0.44
1:D:3321:PRO:HA	1:D:3324:GLU:HG2	2.00	0.44
1:D:4054:SER:O	1:D:4056:LYS:HG3	2.17	0.44
1:D:4055:HIS:CD2	1:D:4057:HIS:ND1	2.85	0.44
1:D:4795:LYS:HD2	1:D:4795:LYS:HA	1.81	0.44
1:A:70:GLU:OE2	1:A:122:ARG:HD3	2.17	0.44
1:A:826:VAL:HG21	1:A:832:LEU:HB2	1.99	0.44
1:A:1572:LYS:HD2	1:A:1572:LYS:N	2.33	0.44
1:A:2884:LYS:HD2	1:A:2885:ASP:N	2.33	0.44
1:A:4294:LEU:HD22	1:B:4719:PHE:CE2	2.52	0.44
2:E:50:ARG:N	2:E:55:GLU:OE1	2.42	0.44
1:B:370:LEU:HB2	1:B:393:MET:CE	2.48	0.44
1:B:1007:TRP:HE1	1:B:1011:ARG:NH1	2.16	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1991:GLU:HG2	1:B:1992:ILE:N	2.32	0.44
1:B:2758:LYS:HE2	1:B:2763:LEU:HA	2.00	0.44
1:B:2782:MET:HG2	1:B:2787:TRP:HE3	1.82	0.44
1:B:3025:ASP:O	1:B:3029:ILE:HD12	2.17	0.44
1:B:3102:LEU:HB2	1:B:3103:PRO:HD3	2.00	0.44
1:B:3187:LYS:HZ3	1:B:3191:GLU:HB3	1.83	0.44
1:B:4607:ALA:HB1	1:B:4649:VAL:HG21	1.99	0.44
1:C:494:MET:HE3	1:C:494:MET:HB3	1.76	0.44
1:C:890:HIS:NE2	1:C:924:LEU:HD22	2.33	0.44
1:C:907:VAL:HG11	1:C:912:LYS:HD3	1.98	0.44
1:C:950:VAL:HA	1:C:1064:LEU:HA	1.99	0.44
1:C:1941:ARG:NH2	1:C:3609:TYR:HB2	2.27	0.44
1:C:2798:MET:HE1	1:D:1497:GLY:HA3	1.98	0.44
1:C:3215:MET:CE	1:C:3215:MET:HA	2.48	0.44
1:D:115:TYR:CG	1:D:164:PRO:HG3	2.53	0.44
1:D:271:ALA:HB2	1:D:488:LEU:HD22	1.99	0.44
1:D:1293:GLN:O	1:D:1294:ASN:HB3	2.18	0.44
1:D:2674:GLY:HA2	1:D:2978:HIS:CE1	2.52	0.44
1:D:2758:LYS:HE2	1:D:2763:LEU:HA	2.00	0.44
2:G:12:ASP:OD1	2:G:13:GLY:N	2.51	0.44
2:G:25:VAL:HG12	2:G:104:LEU:HA	2.00	0.44
2:H:12:ASP:OD1	2:H:13:GLY:N	2.51	0.44
2:H:19:LYS:HB2	2:H:19:LYS:HE3	1.80	0.44
1:A:963:LYS:NZ	1:A:979:ALA:HB3	2.33	0.44
1:A:3062:ASP:O	1:A:3066:GLU:OE1	2.35	0.44
1:A:3323:MET:HB3	1:A:3327:LYS:HZ3	1.82	0.44
1:A:4055:HIS:CD2	1:A:4057:HIS:ND1	2.85	0.44
1:A:4667:SER:OG	1:A:4672:MET:O	2.34	0.44
2:E:12:ASP:OD1	2:E:13:GLY:N	2.51	0.44
1:B:537:LEU:O	1:B:541:ILE:HG12	2.18	0.44
1:B:2397:ASP:O	1:B:2401:ARG:HG3	2.18	0.44
1:B:3741:LEU:HD11	1:B:3777:MET:HG2	1.99	0.44
1:C:718:VAL:HG13	1:C:724:SER:OG	2.18	0.44
1:C:2397:ASP:O	1:C:2401:ARG:HG3	2.18	0.44
1:C:2670:SER:HB2	1:C:2973:GLN:HG2	1.98	0.44
1:C:3042:ALA:HB3	1:C:3117:PHE:HD2	1.82	0.44
1:D:756:SER:OG	1:D:769:ARG:HB2	2.17	0.44
1:D:890:HIS:NE2	1:D:924:LEU:HD22	2.33	0.44
1:D:895:MET:HE3	1:D:978:PRO:HG2	2.00	0.44
1:D:963:LYS:NZ	1:D:979:ALA:HB3	2.33	0.44
1:D:1415:ASP:OD2	1:D:1559:ARG:NH2	2.51	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1628:MET:HE2	1:D:1684:GLN:HB3	1.99	0.44
1:D:2318:ASN:HB3	1:D:2322:ARG:NH1	2.33	0.44
1:D:3042:ALA:HB3	1:D:3117:PHE:HD2	1.82	0.44
1:D:3062:ASP:O	1:D:3066:GLU:OE1	2.35	0.44
1:D:3741:LEU:HD11	1:D:3777:MET:HG2	1.99	0.44
1:D:4194:GLU:HG2	1:D:4645:TRP:HZ3	1.82	0.44
1:D:4804:MET:HE3	1:D:4804:MET:HB3	1.82	0.44
1:A:1074:ARG:HD3	1:A:1078:CYS:HB2	2.00	0.44
1:A:1293:GLN:O	1:A:1294:ASN:HB3	2.18	0.44
1:A:1694:MET:HE2	1:A:1694:MET:HB2	1.87	0.44
1:A:4288:TRP:CZ2	1:A:4292:MET:HE3	2.53	0.44
1:B:115:TYR:CG	1:B:164:PRO:HG3	2.52	0.44
1:B:1572:LYS:N	1:B:1572:LYS:HD2	2.33	0.44
1:B:3213:LYS:HA	1:B:3216:GLU:CD	2.42	0.44
1:B:3297:LYS:HE2	1:B:3297:LYS:HA	2.00	0.44
1:B:4084:VAL:O	1:B:4088:HIS:HB2	2.18	0.44
1:C:1074:ARG:HD3	1:C:1078:CYS:HB2	2.00	0.44
1:C:3044:THR:HA	1:C:3047:LYS:NZ	2.32	0.44
1:C:3882:GLN:NE2	1:C:3946:GLY:HA3	2.32	0.44
1:C:4795:LYS:HA	1:C:4795:LYS:HD2	1.81	0.44
1:D:306:LEU:HA	1:D:316:LEU:HD23	2.00	0.44
1:D:1043:LYS:O	1:D:1047:LYS:HB2	2.17	0.44
1:D:1572:LYS:HD2	1:D:1572:LYS:N	2.33	0.44
1:D:2966:VAL:HG12	1:D:2970:LEU:HD23	1.99	0.44
1:D:3304:GLN:OE1	1:D:3308:ASN:ND2	2.47	0.44
1:D:4023:LEU:HG	1:D:4084:VAL:HG12	1.99	0.44
1:A:3026:ALA:O	1:A:3030:VAL:HG23	2.18	0.44
1:A:3118:GLY:O	1:A:3122:ILE:HG22	2.18	0.44
1:A:3304:GLN:OE1	1:A:3308:ASN:ND2	2.48	0.44
1:A:3830:LEU:HB3	1:A:3833:ASP:OD2	2.18	0.44
1:B:1711:LEU:HB3	1:B:1831:MET:SD	2.58	0.44
1:B:1718:ARG:HD3	1:B:1831:MET:HA	2.00	0.44
1:B:3122:ILE:HG12	1:B:3127:GLN:HG2	1.99	0.44
1:B:3178:HIS:CE1	1:B:3263:MET:HA	2.53	0.44
1:C:712:GLU:OE1	1:C:1086:ARG:NH2	2.39	0.44
1:C:801:ARG:HG2	1:C:1618:LEU:HA	2.00	0.44
1:C:2954:PHE:CE2	1:C:2956:TYR:HB2	2.53	0.44
1:C:2966:VAL:HG12	1:C:2970:LEU:HD23	1.99	0.44
1:C:3017:HIS:O	1:C:3018:ARG:HD3	2.18	0.44
1:C:3297:LYS:HA	1:C:3297:LYS:HE2	2.00	0.44
1:C:4831:ILE:HG13	1:C:4843:ARG:NH2	2.31	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4867:ILE:HG12	1:D:4864:GLY:HA2	2.00	0.44
1:D:1007:TRP:HE1	1:D:1011:ARG:NH1	2.16	0.44
1:D:1500:ARG:HG3	1:D:1505:LEU:H	1.82	0.44
1:D:3215:MET:HA	1:D:3215:MET:CE	2.48	0.44
2:F:45:LYS:HA	2:F:46:PRO:HD3	1.82	0.44
2:F:78:THR:OG1	2:F:80:ASP:OD1	2.26	0.44
2:H:25:VAL:HG12	2:H:104:LEU:HA	2.00	0.44
1:A:950:VAL:HA	1:A:1064:LEU:HA	1.99	0.43
1:A:1304:LEU:HD23	1:A:1304:LEU:HA	1.84	0.43
1:A:2332:GLY:O	1:A:2336:ARG:HG3	2.18	0.43
1:A:4194:GLU:HG2	1:A:4645:TRP:HZ3	1.82	0.43
1:B:306:LEU:HA	1:B:316:LEU:HD23	2.00	0.43
1:B:718:VAL:HG13	1:B:724:SER:OG	2.18	0.43
1:B:2332:GLY:O	1:B:2336:ARG:HG3	2.18	0.43
1:B:3215:MET:HA	1:B:3215:MET:CE	2.48	0.43
1:C:198:ASN:OD1	1:C:199:GLY:N	2.50	0.43
1:C:370:LEU:HB2	1:C:393:MET:CE	2.47	0.43
1:C:505:LEU:HD23	1:C:505:LEU:HA	1.85	0.43
1:C:1711:LEU:HB3	1:C:1831:MET:SD	2.58	0.43
1:C:1957:LEU:HA	1:C:1960:ARG:CZ	2.47	0.43
1:C:1991:GLU:HG2	1:C:1992:ILE:N	2.32	0.43
1:C:2884:LYS:HD2	1:C:2885:ASP:N	2.33	0.43
1:D:674:TYR:CE1	1:D:756:SER:HB2	2.52	0.43
1:D:718:VAL:HG13	1:D:724:SER:OG	2.18	0.43
1:D:1074:ARG:HH11	1:D:1078:CYS:H	1.66	0.43
1:D:1848:PRO:HG3	1:D:1890:LYS:HD2	2.00	0.43
1:D:4036:ASP:OD1	1:D:4040:LYS:NZ	2.52	0.43
1:D:4288:TRP:CZ2	1:D:4292:MET:HE3	2.53	0.43
1:D:4689:LYS:NZ	1:D:4696:ALA:HB2	2.31	0.43
2:F:25:VAL:HG12	2:F:104:LEU:HA	2.00	0.43
1:A:718:VAL:HG13	1:A:724:SER:OG	2.18	0.43
1:A:1144:ARG:HB3	1:A:1152:TYR:HB3	1.99	0.43
1:A:1435:GLY:H	1:A:1500:ARG:HH12	1.66	0.43
1:A:1711:LEU:HB3	1:A:1831:MET:SD	2.58	0.43
1:A:1898:LEU:HD23	1:A:1902:VAL:HG12	2.00	0.43
1:A:2728:SER:HA	1:A:2731:LYS:NZ	2.33	0.43
1:A:2832:THR:OG1	1:B:1548:THR:O	2.33	0.43
1:A:4283:PHE:HE2	1:B:4495:PHE:HE2	1.66	0.43
1:B:29:HIS:CE1	1:B:31:GLU:HG2	2.52	0.43
1:B:826:VAL:HG21	1:B:832:LEU:HB2	1.99	0.43
1:B:895:MET:HE3	1:B:978:PRO:HG2	2.00	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2232:PRO:C	1:B:2234:MET:H	2.24	0.43
1:B:2926:LEU:CB	1:B:3003:MET:HE2	2.48	0.43
1:B:3017:HIS:O	1:B:3018:ARG:HD3	2.18	0.43
1:B:3184:TYR:CE1	1:B:3197:LEU:HD21	2.53	0.43
1:B:4602:ARG:NH1	1:B:4631:ASP:OD1	2.51	0.43
1:C:115:TYR:CG	1:C:164:PRO:HG3	2.53	0.43
1:C:271:ALA:HB2	1:C:488:LEU:HD22	1.99	0.43
1:C:306:LEU:HA	1:C:316:LEU:HD23	2.00	0.43
1:C:1300:MET:HE2	1:C:1300:MET:HB3	1.71	0.43
1:C:2137:GLU:H	1:C:2137:GLU:CD	2.22	0.43
1:C:3026:ALA:O	1:C:3030:VAL:HG23	2.18	0.43
1:C:3102:LEU:HB2	1:C:3103:PRO:HD3	2.00	0.43
1:C:3321:PRO:HA	1:C:3324:GLU:HG2	1.99	0.43
1:D:3118:GLY:O	1:D:3122:ILE:HG22	2.18	0.43
1:D:3178:HIS:CE1	1:D:3263:MET:HA	2.53	0.43
1:D:3273:MET:HE3	1:D:3273:MET:HB2	1.73	0.43
1:D:4602:ARG:NH1	1:D:4631:ASP:OD1	2.51	0.43
1:D:4607:ALA:HB1	1:D:4649:VAL:HG21	1.99	0.43
1:A:234:LEU:HD13	1:A:405:LEU:HD22	2.00	0.43
1:A:306:LEU:HA	1:A:316:LEU:HD23	2.00	0.43
1:A:801:ARG:HG2	1:A:1618:LEU:HA	2.00	0.43
1:A:1718:ARG:HD3	1:A:1831:MET:HA	2.00	0.43
1:A:2318:ASN:HB3	1:A:2322:ARG:NH1	2.33	0.43
1:A:2782:MET:HG2	1:A:2787:TRP:HE3	1.82	0.43
1:A:2954:PHE:CE2	1:A:2956:TYR:HB2	2.53	0.43
1:A:3017:HIS:O	1:A:3018:ARG:HD3	2.18	0.43
1:B:234:LEU:HD13	1:B:405:LEU:HD22	2.00	0.43
1:B:891:GLU:HG3	1:B:976:TYR:HD2	1.82	0.43
1:B:1074:ARG:HH11	1:B:1078:CYS:H	1.66	0.43
1:B:1415:ASP:OD2	1:B:1559:ARG:NH2	2.51	0.43
1:B:1898:LEU:HD23	1:B:1902:VAL:HG12	2.00	0.43
1:B:2954:PHE:CE2	1:B:2956:TYR:HB2	2.53	0.43
1:B:4036:ASP:OD1	1:B:4040:LYS:NZ	2.52	0.43
1:B:4085:LYS:HE3	1:B:4085:LYS:HB3	1.83	0.43
1:C:1442:TRP:HB3	1:C:1487:MET:HE2	2.01	0.43
1:C:1500:ARG:HG3	1:C:1505:LEU:H	1.82	0.43
1:C:1979:PHE:HZ	1:C:1996:LEU:HD23	1.84	0.43
1:C:4054:SER:O	1:C:4056:LYS:HG3	2.17	0.43
1:D:1435:GLY:H	1:D:1500:ARG:HH12	1.66	0.43
1:D:1898:LEU:HD23	1:D:1902:VAL:HG12	2.00	0.43
1:A:1434:PRO:O	1:D:2830:ASN:HB2	2.18	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1442:TRP:HB3	1:A:1487:MET:HE2	2.01	0.43
1:A:1848:PRO:HG3	1:A:1890:LYS:HD2	2.00	0.43
1:A:2831:VAL:HG22	1:B:1435:GLY:HA2	2.00	0.43
1:A:3297:LYS:HE2	1:A:3297:LYS:HA	2.00	0.43
1:A:4023:LEU:HG	1:A:4084:VAL:HG12	1.99	0.43
1:B:271:ALA:HB2	1:B:488:LEU:HD22	1.99	0.43
1:B:829:LYS:HZ2	1:B:1037:LEU:HB3	1.83	0.43
1:B:1038:LEU:HD23	1:B:1043:LYS:HG2	2.01	0.43
1:B:1848:PRO:HG3	1:B:1890:LYS:HD2	2.00	0.43
1:B:1979:PHE:HZ	1:B:1996:LEU:HD23	1.83	0.43
1:B:2130:LEU:HD21	1:B:2173:GLU:HG3	2.00	0.43
1:B:4023:LEU:HG	1:B:4084:VAL:HG12	1.99	0.43
1:C:891:GLU:HG3	1:C:976:TYR:HD2	1.82	0.43
1:C:1785:ASP:OD1	1:C:1786:ILE:N	2.48	0.43
1:C:2728:SER:HA	1:C:2731:LYS:NZ	2.33	0.43
1:D:234:LEU:HD13	1:D:405:LEU:HD22	2.00	0.43
1:D:950:VAL:HA	1:D:1064:LEU:HA	1.99	0.43
1:D:962:LYS:NZ	1:D:982:ASP:H	2.16	0.43
1:D:1074:ARG:HD3	1:D:1078:CYS:HB2	2.00	0.43
1:D:2696:ASP:O	1:D:2700:ASN:HA	2.19	0.43
1:D:3017:HIS:O	1:D:3018:ARG:HD3	2.18	0.43
1:A:274:LEU:HD23	1:A:274:LEU:HA	1.90	0.43
1:A:1007:TRP:HE1	1:A:1011:ARG:NH1	2.16	0.43
1:A:2397:ASP:O	1:A:2401:ARG:HG3	2.18	0.43
1:A:2759:PRO:HG2	1:A:2762:LEU:HD13	2.01	0.43
1:A:2782:MET:HG2	1:A:2787:TRP:CE3	2.54	0.43
1:A:2909:ASP:OD2	1:A:2912:LEU:HG	2.19	0.43
1:A:3124:GLU:C	1:A:3126:VAL:H	2.25	0.43
1:A:4036:ASP:OD1	1:A:4040:LYS:NZ	2.52	0.43
1:A:4084:VAL:O	1:A:4088:HIS:HB2	2.18	0.43
1:B:801:ARG:HG2	1:B:1618:LEU:HA	2.00	0.43
1:B:2782:MET:HG2	1:B:2787:TRP:CE3	2.54	0.43
1:B:3323:MET:HB3	1:B:3327:LYS:HZ1	1.83	0.43
1:B:3695:MET:HB3	1:B:3731:LEU:HD11	2.00	0.43
1:B:3882:GLN:NE2	1:B:3946:GLY:HA3	2.33	0.43
1:C:686:VAL:HG13	1:C:687:THR:HG23	2.00	0.43
1:C:1293:GLN:O	1:C:1294:ASN:HB3	2.18	0.43
1:C:2481:GLN:NE2	1:C:2537:GLY:O	2.42	0.43
1:C:2926:LEU:CB	1:C:3003:MET:HE2	2.47	0.43
1:C:3741:LEU:HD11	1:C:3777:MET:HG2	1.99	0.43
1:D:58:VAL:HG13	1:D:319:LYS:HG2	2.00	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:874:LEU:HD23	1:D:879:GLU:OE1	2.19	0.43
1:D:2137:GLU:H	1:D:2137:GLU:CD	2.22	0.43
1:D:2234:MET:HE2	1:D:2234:MET:HB3	1.90	0.43
1:D:2899:ASN:O	1:D:2899:ASN:ND2	2.52	0.43
1:D:2954:PHE:CE2	1:D:2956:TYR:HB2	2.53	0.43
1:D:3882:GLN:NE2	1:D:3946:GLY:HA3	2.32	0.43
1:A:891:GLU:HG3	1:A:976:TYR:HD2	1.82	0.43
1:A:897:LYS:HE2	1:A:917:CYS:HB3	1.99	0.43
1:A:1415:ASP:OD2	1:A:1559:ARG:NH2	2.51	0.43
1:B:1948:MET:HA	1:B:1951:LEU:HD23	2.00	0.43
1:C:880:ARG:HH12	1:C:881:ILE:HD13	1.84	0.43
1:C:1038:LEU:HD23	1:C:1043:LYS:HG2	2.01	0.43
1:C:1415:ASP:OD2	1:C:1559:ARG:NH2	2.51	0.43
1:C:2681:MET:HE2	1:C:2914:THR:HA	2.01	0.43
1:C:2782:MET:HG2	1:C:2787:TRP:HE3	1.82	0.43
1:C:3184:TYR:CE1	1:C:3197:LEU:HD21	2.53	0.43
1:C:3986:LEU:O	1:C:3989:ASN:HB2	2.19	0.43
1:C:4288:TRP:CZ2	1:C:4292:MET:HE3	2.53	0.43
1:C:4602:ARG:NH1	1:C:4631:ASP:OD1	2.51	0.43
1:D:200:SER:O	1:D:202:HIS:HD2	2.01	0.43
1:D:686:VAL:HG13	1:D:687:THR:HG23	2.00	0.43
1:D:801:ARG:HG2	1:D:1618:LEU:HA	2.00	0.43
1:D:2909:ASP:OD2	1:D:2912:LEU:HG	2.19	0.43
1:A:115:TYR:CG	1:A:164:PRO:HG3	2.53	0.43
1:A:718:VAL:HG23	1:A:793:SER:HB3	2.01	0.43
1:A:2888:LYS:HA	1:A:2891:ASP:OD2	2.19	0.43
1:B:930:ASN:OD1	1:B:931:TYR:N	2.52	0.43
1:B:963:LYS:NZ	1:B:979:ALA:HB3	2.33	0.43
1:B:1074:ARG:HD3	1:B:1078:CYS:HB2	2.00	0.43
1:B:1442:TRP:HB3	1:B:1487:MET:HE2	2.01	0.43
1:B:1939:ASN:ND2	1:B:1989:PRO:HD3	2.34	0.43
1:B:3026:ALA:O	1:B:3030:VAL:HG23	2.18	0.43
1:C:2332:GLY:O	1:C:2336:ARG:HG3	2.18	0.43
1:C:3695:MET:HB3	1:C:3731:LEU:HD11	2.00	0.43
1:C:3728:GLN:OE1	1:C:3770:ASN:ND2	2.52	0.43
1:C:4023:LEU:HG	1:C:4084:VAL:HG12	1.99	0.43
1:C:4036:ASP:OD1	1:C:4040:LYS:NZ	2.52	0.43
1:D:1718:ARG:HD3	1:D:1831:MET:HA	2.00	0.43
1:D:2215:PHE:CG	1:D:2253:LEU:HD22	2.54	0.43
1:D:2724:TYR:CD1	1:D:2895:PHE:HD1	2.35	0.43
1:D:3237:VAL:C	1:D:3240:PRO:HD2	2.44	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3297:LYS:HE2	1:D:3297:LYS:HA	2.00	0.43
1:A:200:SER:O	1:A:202:HIS:HD2	2.01	0.43
1:A:606:ARG:HH21	1:A:1633:PRO:HD2	1.82	0.43
1:A:930:ASN:OD1	1:A:931:TYR:N	2.52	0.43
1:A:1074:ARG:HH11	1:A:1078:CYS:H	1.67	0.43
1:B:200:SER:O	1:B:202:HIS:HD2	2.01	0.43
1:B:1144:ARG:HB3	1:B:1152:TYR:HB3	1.99	0.43
1:C:911:ASN:OD1	1:C:912:LYS:N	2.52	0.43
1:C:1074:ARG:HH11	1:C:1078:CYS:H	1.67	0.43
1:C:1140:PHE:CD2	1:C:1141:LYS:HE3	2.54	0.43
1:C:1718:ARG:HD3	1:C:1831:MET:HA	2.00	0.43
1:C:1939:ASN:ND2	1:C:1989:PRO:HD3	2.34	0.43
1:C:2234:MET:HE2	1:C:2234:MET:HB3	1.90	0.43
1:C:2741:TRP:CD1	1:C:2752:LYS:HZ3	2.36	0.43
1:C:2782:MET:HG2	1:C:2787:TRP:CE3	2.54	0.43
1:C:2909:ASP:OD2	1:C:2912:LEU:HG	2.19	0.43
1:C:3118:GLY:O	1:C:3122:ILE:HG22	2.18	0.43
1:C:3157:GLU:HG3	1:C:3302:PHE:CZ	2.54	0.43
1:C:3237:VAL:C	1:C:3240:PRO:HD2	2.44	0.43
1:D:494:MET:HE3	1:D:494:MET:HB3	1.75	0.43
1:D:997:ASP:HA	1:D:1047:LYS:HZ2	1.83	0.43
1:D:1140:PHE:CD2	1:D:1141:LYS:HE3	2.54	0.43
1:D:1442:TRP:HB3	1:D:1487:MET:HE2	2.01	0.43
1:D:1471:ASP:OD2	1:D:1474:GLY:N	2.51	0.43
1:D:2130:LEU:HD21	1:D:2173:GLU:HG3	2.00	0.43
1:D:2681:MET:HE2	1:D:2914:THR:HA	2.01	0.43
1:D:2692:GLN:H	1:D:2692:GLN:CD	2.27	0.43
1:D:2868:HIS:NE2	1:D:2870:LEU:HB2	2.34	0.43
1:A:505:LEU:HD23	1:A:505:LEU:HA	1.85	0.43
1:A:874:LEU:HD23	1:A:879:GLU:OE1	2.18	0.43
1:A:1948:MET:HA	1:A:1951:LEU:HD23	2.00	0.43
1:A:2130:LEU:HD21	1:A:2173:GLU:HG3	2.00	0.43
1:A:2222:LEU:HD23	1:A:2222:LEU:HA	1.88	0.43
1:B:246:THR:HG21	1:B:267:VAL:HG21	2.01	0.43
1:B:890:HIS:NE2	1:B:924:LEU:HD22	2.33	0.43
1:B:1293:GLN:O	1:B:1294:ASN:HB3	2.18	0.43
1:B:3118:GLY:O	1:B:3122:ILE:HG22	2.18	0.43
1:B:3611:LEU:HD23	1:B:3611:LEU:HA	1.89	0.43
1:B:3664:LEU:O	1:B:3668:ILE:HG13	2.19	0.43
1:B:3728:GLN:OE1	1:B:3770:ASN:ND2	2.52	0.43
1:B:3890:TRP:HB3	1:C:76:ARG:HG2	2.01	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:4194:GLU:HG2	1:B:4645:TRP:HZ3	1.82	0.43
1:B:4288:TRP:CZ2	1:B:4292:MET:HE3	2.53	0.43
1:B:4921:PHE:HE2	1:B:4940:VAL:HG11	1.84	0.43
1:C:711:GLU:OE1	1:C:716:ASN:ND2	2.52	0.43
1:C:874:LEU:HD23	1:C:879:GLU:OE1	2.18	0.43
1:C:4185:LYS:HG2	1:C:4186:MET:HE2	2.01	0.43
1:C:4607:ALA:HB1	1:C:4649:VAL:HG21	2.00	0.43
1:D:1475:LYS:HE3	1:D:1475:LYS:HB3	1.89	0.43
1:D:2589:LEU:O	1:D:2593:VAL:HG13	2.19	0.43
1:D:2782:MET:HG2	1:D:2787:TRP:CE3	2.54	0.43
1:A:58:VAL:HG13	1:A:319:LYS:HG2	2.00	0.43
1:A:880:ARG:HH12	1:A:881:ILE:HD13	1.84	0.43
1:A:890:HIS:NE2	1:A:924:LEU:HD22	2.33	0.43
1:A:1941:ARG:NH2	1:A:3609:TYR:HB2	2.27	0.43
1:A:3215:MET:HA	1:A:3215:MET:CE	2.48	0.43
1:A:3664:LEU:O	1:A:3668:ILE:HG13	2.19	0.43
1:B:686:VAL:HG13	1:B:687:THR:HG23	2.00	0.43
1:B:711:GLU:OE1	1:B:716:ASN:ND2	2.52	0.43
1:B:874:LEU:HD23	1:B:879:GLU:OE1	2.18	0.43
1:B:2929:LEU:HD21	1:B:2970:LEU:HD11	2.01	0.43
1:B:3321:PRO:HA	1:B:3324:GLU:HG2	2.00	0.43
1:C:246:THR:HG21	1:C:267:VAL:HG21	2.01	0.43
1:C:2130:LEU:HD21	1:C:2173:GLU:HG3	2.00	0.43
1:C:3312:PRO:HD2	1:C:3313:GLN:OE1	2.19	0.43
1:D:880:ARG:HH12	1:D:881:ILE:HD13	1.84	0.43
1:D:1979:PHE:HZ	1:D:1996:LEU:HD23	1.84	0.43
1:D:2332:GLY:O	1:D:2336:ARG:HG3	2.18	0.43
1:A:711:GLU:OE1	1:A:716:ASN:ND2	2.52	0.42
1:A:1038:LEU:HD23	1:A:1043:LYS:HG2	2.01	0.42
1:A:1979:PHE:HZ	1:A:1996:LEU:HD23	1.84	0.42
1:A:2696:ASP:O	1:A:2700:ASN:HA	2.19	0.42
1:A:3321:PRO:HA	1:A:3324:GLU:HG2	2.00	0.42
1:A:3649:ALA:C	1:A:3652:PRO:HD2	2.44	0.42
1:A:4085:LYS:HB3	1:A:4085:LYS:HE3	1.83	0.42
1:B:600:LEU:HD12	1:B:604:HIS:CD2	2.54	0.42
1:B:1031:ARG:HD2	1:B:1042:THR:HG21	2.01	0.42
1:B:2589:LEU:O	1:B:2593:VAL:HG13	2.19	0.42
1:B:2868:HIS:NE2	1:B:2870:LEU:HB2	2.34	0.42
1:B:3122:ILE:HG12	1:B:3127:GLN:CG	2.49	0.42
1:B:3986:LEU:O	1:B:3989:ASN:HB2	2.19	0.42
1:C:234:LEU:HD13	1:C:405:LEU:HD22	2.00	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:718:VAL:HG23	1:C:793:SER:HB3	2.01	0.42
1:C:930:ASN:OD1	1:C:931:TYR:N	2.52	0.42
1:C:2318:ASN:HB3	1:C:2322:ARG:NH1	2.33	0.42
1:C:2696:ASP:O	1:C:2700:ASN:HA	2.19	0.42
1:C:3316:LYS:HB2	1:C:3316:LYS:HE2	1.54	0.42
1:C:3326:LEU:HD21	1:C:3336:GLU:HB2	2.01	0.42
1:D:712:GLU:OE1	1:D:1086:ARG:NH2	2.39	0.42
1:D:930:ASN:OD1	1:D:931:TYR:N	2.52	0.42
1:D:967:PRO:O	1:D:971:GLN:HG2	2.19	0.42
1:D:2759:PRO:HG2	1:D:2762:LEU:HD13	2.01	0.42
1:D:3026:ALA:O	1:D:3030:VAL:HG23	2.18	0.42
1:A:971:GLN:HE21	1:A:978:PRO:HD2	1.84	0.42
1:A:4602:ARG:NH1	1:A:4631:ASP:OD1	2.51	0.42
2:E:25:VAL:HG12	2:E:104:LEU:HA	2.00	0.42
1:B:274:LEU:HD23	1:B:274:LEU:HA	1.89	0.42
1:B:505:LEU:HD23	1:B:505:LEU:HA	1.85	0.42
1:B:880:ARG:HH12	1:B:881:ILE:HD13	1.84	0.42
1:B:983:LEU:HD11	1:B:1055:ARG:HG3	2.00	0.42
1:B:2830:ASN:HB2	1:C:1434:PRO:O	2.19	0.42
1:B:3215:MET:HA	1:B:3215:MET:HE2	2.01	0.42
1:C:962:LYS:NZ	1:C:982:ASP:H	2.16	0.42
1:C:3304:GLN:OE1	1:C:3308:ASN:ND2	2.48	0.42
1:D:971:GLN:HE21	1:D:978:PRO:HD2	1.84	0.42
1:D:1038:LEU:HD23	1:D:1043:LYS:HG2	2.01	0.42
1:D:2436:ILE:HG22	1:D:2491:GLY:HA3	2.02	0.42
1:D:3195:LEU:HD22	1:D:3197:LEU:HB2	2.01	0.42
1:D:3201:VAL:O	1:D:3204:VAL:HG12	2.19	0.42
1:D:3213:LYS:HA	1:D:3216:GLU:OE2	2.19	0.42
1:D:3215:MET:HA	1:D:3215:MET:HE2	2.01	0.42
1:A:600:LEU:HD12	1:A:604:HIS:CD2	2.54	0.42
1:A:967:PRO:O	1:A:971:GLN:HG2	2.19	0.42
1:A:2681:MET:HE2	1:A:2914:THR:HA	2.01	0.42
1:A:3072:MET:HE2	1:A:3072:MET:HB2	1.79	0.42
1:A:3312:PRO:HD2	1:A:3313:GLN:OE1	2.19	0.42
1:A:3695:MET:HB3	1:A:3731:LEU:HD11	2.00	0.42
1:B:1140:PHE:CD2	1:B:1141:LYS:HE3	2.54	0.42
1:B:1572:LYS:HD2	1:B:1572:LYS:H	1.84	0.42
1:B:2554:ARG:HH11	1:B:2554:ARG:HG3	1.85	0.42
1:B:3213:LYS:HA	1:B:3216:GLU:OE2	2.19	0.42
1:C:58:VAL:HG13	1:C:319:LYS:HG2	2.00	0.42
1:C:600:LEU:HD12	1:C:604:HIS:CD2	2.54	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1471:ASP:OD2	1:C:1474:GLY:N	2.51	0.42
1:C:1948:MET:HA	1:C:1951:LEU:HD23	2.00	0.42
1:C:2868:HIS:NE2	1:C:2870:LEU:HB2	2.34	0.42
1:C:2874:TYR:CZ	1:C:2882:LYS:HD3	2.54	0.42
1:C:3649:ALA:C	1:C:3652:PRO:HD2	2.44	0.42
1:C:4084:VAL:O	1:C:4088:HIS:HB2	2.18	0.42
1:C:4814:MET:HA	1:C:4814:MET:HE3	2.02	0.42
1:D:1906:MET:HE3	1:D:1906:MET:HB3	1.83	0.42
1:D:2888:LYS:HA	1:D:2891:ASP:OD2	2.19	0.42
1:D:3102:LEU:HB2	1:D:3103:PRO:HD3	2.00	0.42
1:A:246:THR:HG21	1:A:267:VAL:HG21	2.01	0.42
1:A:1140:PHE:CD2	1:A:1141:LYS:HE3	2.54	0.42
1:A:2155:PHE:HD1	1:A:2162:MET:HG3	1.85	0.42
1:A:2692:GLN:H	1:A:2692:GLN:CD	2.27	0.42
1:A:2771:TYR:O	1:A:2774:PRO:HD2	2.20	0.42
1:A:3178:HIS:CE1	1:A:3263:MET:HA	2.53	0.42
1:A:3187:LYS:HZ3	1:A:3191:GLU:HB3	1.84	0.42
1:A:3201:VAL:O	1:A:3204:VAL:HG12	2.20	0.42
1:A:3326:LEU:HD21	1:A:3336:GLU:HB2	2.02	0.42
1:A:3712:LYS:O	1:A:3717:LYS:NZ	2.53	0.42
1:A:4814:MET:HA	1:A:4814:MET:HE3	2.02	0.42
1:B:911:ASN:OD1	1:B:912:LYS:N	2.52	0.42
1:B:2138:GLU:HA	1:B:2141:LEU:HD12	2.02	0.42
1:B:2318:ASN:HB3	1:B:2322:ARG:NH1	2.33	0.42
1:B:2696:ASP:O	1:B:2700:ASN:HA	2.19	0.42
1:B:2728:SER:HA	1:B:2731:LYS:NZ	2.33	0.42
1:B:4185:LYS:HG2	1:B:4186:MET:HE2	2.01	0.42
1:C:932:ASN:HA	1:C:935:MET:HG3	2.02	0.42
1:C:2403:ALA:HB2	1:C:2475:VAL:HG22	2.01	0.42
1:C:3195:LEU:HD22	1:C:3197:LEU:HB2	2.01	0.42
1:D:198:ASN:OD1	1:D:199:GLY:N	2.50	0.42
1:D:711:GLU:OE1	1:D:716:ASN:ND2	2.52	0.42
1:D:891:GLU:HG3	1:D:976:TYR:HD2	1.82	0.42
1:D:962:LYS:HZ1	1:D:982:ASP:H	1.66	0.42
1:D:983:LEU:HD11	1:D:1055:ARG:HG3	2.00	0.42
1:D:1711:LEU:HB3	1:D:1831:MET:SD	2.58	0.42
1:D:3712:LYS:O	1:D:3717:LYS:NZ	2.53	0.42
1:D:3986:LEU:O	1:D:3989:ASN:HB2	2.19	0.42
1:A:996:VAL:HG21	1:A:1051:ARG:HA	2.02	0.42
1:A:2666:LEU:HB3	1:A:2667:PRO:HD3	2.02	0.42
1:A:3215:MET:HA	1:A:3215:MET:HE2	2.01	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:58:VAL:HG13	1:B:319:LYS:HG2	2.00	0.42
1:B:227:TYR:CG	1:B:352:SER:HB3	2.55	0.42
1:B:718:VAL:HG23	1:B:793:SER:HB3	2.01	0.42
1:B:844:ARG:NH2	1:B:847:THR:HG21	2.35	0.42
1:B:1435:GLY:H	1:B:1500:ARG:HH12	1.66	0.42
1:B:2899:ASN:O	1:B:2899:ASN:ND2	2.52	0.42
1:B:3325:LYS:HE3	1:B:3325:LYS:HA	2.01	0.42
1:B:3712:LYS:O	1:B:3717:LYS:NZ	2.53	0.42
1:B:4814:MET:HA	1:B:4814:MET:HE3	2.02	0.42
1:C:2436:ILE:HG22	1:C:2491:GLY:HA3	2.02	0.42
1:C:2888:LYS:HA	1:C:2891:ASP:OD2	2.19	0.42
1:D:718:VAL:HG23	1:D:793:SER:HB3	2.01	0.42
1:D:2155:PHE:HD1	1:D:2162:MET:HG3	1.85	0.42
1:D:2554:ARG:HH11	1:D:2554:ARG:HG3	1.85	0.42
1:D:4084:VAL:O	1:D:4088:HIS:HB2	2.18	0.42
1:D:4085:LYS:HE3	1:D:4085:LYS:HB3	1.83	0.42
1:A:2436:ILE:HG22	1:A:2491:GLY:HA3	2.02	0.42
1:A:2589:LEU:O	1:A:2593:VAL:HG13	2.19	0.42
1:A:3237:VAL:C	1:A:3240:PRO:HD2	2.44	0.42
1:A:3325:LYS:HE3	1:A:3325:LYS:HA	2.01	0.42
1:B:198:ASN:OD1	1:B:199:GLY:N	2.50	0.42
1:B:1979:PHE:CG	1:B:1993:ARG:HG2	2.55	0.42
1:B:2888:LYS:HA	1:B:2891:ASP:OD2	2.19	0.42
1:C:1435:GLY:H	1:C:1500:ARG:HH12	1.66	0.42
1:C:2589:LEU:O	1:C:2593:VAL:HG13	2.19	0.42
1:C:2711:ILE:HD11	1:C:2783:LEU:HG	2.02	0.42
1:C:2929:LEU:HD21	1:C:2970:LEU:HD11	2.01	0.42
1:C:3201:VAL:O	1:C:3204:VAL:HG12	2.20	0.42
1:C:3215:MET:HA	1:C:3215:MET:HE2	2.01	0.42
1:C:3270:SER:HA	1:C:3273:MET:CE	2.37	0.42
1:C:3325:LYS:HE3	1:C:3325:LYS:HA	2.01	0.42
1:D:2666:LEU:HB3	1:D:2667:PRO:HD3	2.02	0.42
1:D:3325:LYS:HE3	1:D:3325:LYS:HA	2.01	0.42
1:A:686:VAL:HG13	1:A:687:THR:HG23	2.00	0.42
1:A:911:ASN:OD1	1:A:912:LYS:N	2.52	0.42
1:A:2146:LEU:HD23	1:A:2146:LEU:HA	1.87	0.42
1:A:2929:LEU:HD21	1:A:2970:LEU:HD11	2.01	0.42
1:A:3102:LEU:HB2	1:A:3103:PRO:HD3	2.00	0.42
1:A:3127:GLN:OE1	1:A:3183:ILE:HB	2.20	0.42
1:A:3213:LYS:HA	1:A:3216:GLU:OE2	2.19	0.42
1:B:881:ILE:O	1:B:885:LEU:HD23	2.20	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:932:ASN:HA	1:B:935:MET:HG3	2.02	0.42
1:B:996:VAL:HG21	1:B:1051:ARG:HA	2.02	0.42
1:B:2155:PHE:HD1	1:B:2162:MET:HG3	1.85	0.42
1:B:2408:LEU:HD23	1:B:2408:LEU:HA	1.90	0.42
1:B:2874:TYR:CZ	1:B:2882:LYS:HD3	2.54	0.42
1:B:2959:GLU:OE1	1:B:2959:GLU:N	2.53	0.42
1:B:3326:LEU:HD21	1:B:3336:GLU:HB2	2.02	0.42
1:B:4155:SER:O	1:B:4159:GLN:HG2	2.20	0.42
1:B:4262:LYS:HG3	1:C:4698:LEU:HD13	2.00	0.42
1:C:983:LEU:HD11	1:C:1055:ARG:HG3	2.01	0.42
1:C:2155:PHE:HD1	1:C:2162:MET:HG3	1.85	0.42
1:C:2554:ARG:HH11	1:C:2554:ARG:HG3	1.85	0.42
1:C:2708:THR:HB	1:C:2780:LYS:HB3	2.02	0.42
1:C:2759:PRO:HG2	1:C:2762:LEU:HD13	2.01	0.42
1:C:4155:SER:O	1:C:4159:GLN:HG2	2.20	0.42
1:D:1572:LYS:HD2	1:D:1572:LYS:H	1.85	0.42
1:D:1948:MET:HA	1:D:1951:LEU:HD23	2.00	0.42
1:D:2708:THR:HB	1:D:2780:LYS:HB3	2.01	0.42
1:D:2728:SER:HA	1:D:2731:LYS:NZ	2.33	0.42
1:A:844:ARG:NH2	1:A:847:THR:HG21	2.35	0.42
1:A:1572:LYS:HD2	1:A:1572:LYS:H	1.84	0.42
1:A:1939:ASN:ND2	1:A:1989:PRO:HD3	2.34	0.42
1:A:2215:PHE:CG	1:A:2253:LEU:HD22	2.54	0.42
1:A:2403:ALA:HB2	1:A:2475:VAL:HG22	2.01	0.42
1:A:2874:TYR:CZ	1:A:2882:LYS:HD3	2.54	0.42
1:A:3036:LEU:O	1:A:3040:LEU:HG	2.20	0.42
1:A:3756:ALA:O	1:A:3760:LYS:HG3	2.20	0.42
1:B:1064:LEU:HD23	1:B:1064:LEU:H	1.85	0.42
1:B:2403:ALA:HB2	1:B:2475:VAL:HG22	2.01	0.42
1:B:2909:ASP:OD2	1:B:2912:LEU:HG	2.19	0.42
1:B:3201:VAL:O	1:B:3204:VAL:HG12	2.20	0.42
1:C:844:ARG:NH2	1:C:847:THR:HG21	2.35	0.42
1:C:1064:LEU:H	1:C:1064:LEU:HD23	1.85	0.42
1:C:1572:LYS:H	1:C:1572:LYS:HD2	1.84	0.42
1:C:1572:LYS:HD2	1:C:1572:LYS:N	2.33	0.42
1:C:1979:PHE:CG	1:C:1993:ARG:HG2	2.55	0.42
1:C:2215:PHE:CG	1:C:2253:LEU:HD22	2.54	0.42
1:C:2692:GLN:CD	1:C:2692:GLN:H	2.27	0.42
1:C:2965:LYS:HA	1:C:2965:LYS:HD3	1.80	0.42
1:C:3213:LYS:HA	1:C:3216:GLU:OE2	2.19	0.42
1:C:3290:ILE:HG22	1:C:3291:ASP:OD2	2.20	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:600:LEU:HD12	1:D:604:HIS:CD2	2.54	0.42
1:D:2138:GLU:HA	1:D:2141:LEU:HD12	2.02	0.42
1:D:3312:PRO:HD2	1:D:3313:GLN:OE1	2.19	0.42
1:D:3649:ALA:C	1:D:3652:PRO:HD2	2.44	0.42
1:D:3695:MET:HB3	1:D:3731:LEU:HD11	2.00	0.42
1:D:3756:ALA:O	1:D:3760:LYS:HG3	2.20	0.42
2:F:19:LYS:HB2	2:F:19:LYS:HE3	1.80	0.42
1:A:2937:HIS:HA	1:A:3014:LEU:HD11	2.02	0.42
1:A:3157:GLU:HG3	1:A:3302:PHE:CZ	2.54	0.42
1:A:4921:PHE:HE2	1:A:4940:VAL:HG11	1.84	0.42
1:B:967:PRO:O	1:B:971:GLN:HG2	2.19	0.42
1:B:2681:MET:HE2	1:B:2914:THR:HA	2.01	0.42
1:B:2692:GLN:CD	1:B:2692:GLN:H	2.27	0.42
1:B:3237:VAL:C	1:B:3240:PRO:HD2	2.44	0.42
1:C:1080:GLY:O	1:C:1081:THR:OG1	2.32	0.42
1:C:1898:LEU:HD23	1:C:1902:VAL:HG12	2.00	0.42
1:C:2959:GLU:OE1	1:C:2959:GLU:N	2.53	0.42
1:C:3664:LEU:O	1:C:3668:ILE:HG13	2.19	0.42
1:D:227:TYR:CG	1:D:352:SER:HB3	2.55	0.42
1:D:844:ARG:NH2	1:D:847:THR:HG21	2.35	0.42
1:D:911:ASN:OD1	1:D:912:LYS:N	2.52	0.42
1:D:2711:ILE:HD11	1:D:2783:LEU:HG	2.02	0.42
1:D:3036:LEU:O	1:D:3040:LEU:HG	2.20	0.42
1:D:3290:ILE:HG22	1:D:3291:ASP:OD2	2.20	0.42
1:D:3326:LEU:HD21	1:D:3336:GLU:HB2	2.01	0.42
1:D:3728:GLN:OE1	1:D:3770:ASN:ND2	2.52	0.42
1:D:3981:MET:HE3	1:D:3981:MET:HB3	1.93	0.42
1:D:4921:PHE:HE2	1:D:4940:VAL:HG11	1.84	0.42
1:A:2138:GLU:HA	1:A:2141:LEU:HD12	2.02	0.42
1:A:4185:LYS:HG2	1:A:4186:MET:HE2	2.01	0.42
1:B:735:GLY:O	1:B:737:ILE:HD12	2.20	0.42
1:B:824:GLU:HA	1:B:1020:ILE:HD12	2.02	0.42
1:B:841:LYS:O	1:B:848:ARG:NH2	2.53	0.42
1:B:1591:LEU:HD12	1:B:1591:LEU:HA	1.93	0.42
1:B:1748:LEU:HD11	1:B:1853:GLU:HG3	2.02	0.42
1:B:2215:PHE:CG	1:B:2253:LEU:HD22	2.54	0.42
1:B:2919:LYS:HB2	1:B:2919:LYS:HE3	1.87	0.42
1:B:3312:PRO:HD2	1:B:3313:GLN:OE1	2.19	0.42
1:B:3649:ALA:C	1:B:3652:PRO:HD2	2.44	0.42
1:C:1031:ARG:HD2	1:C:1042:THR:HG21	2.01	0.42
1:C:2138:GLU:HA	1:C:2141:LEU:HD12	2.02	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4921:PHE:HE2	1:C:4940:VAL:HG11	1.84	0.42
1:D:932:ASN:HA	1:D:935:MET:HG3	2.02	0.42
1:D:1283:LEU:O	1:D:1555:PHE:N	2.49	0.42
1:D:2146:LEU:HD23	1:D:2146:LEU:HA	1.87	0.42
1:D:2326:ARG:HD3	1:D:2326:ARG:HA	1.76	0.42
1:D:2874:TYR:CZ	1:D:2882:LYS:HD3	2.54	0.42
1:D:3127:GLN:OE1	1:D:3183:ILE:HB	2.20	0.42
1:D:3664:LEU:O	1:D:3668:ILE:HG13	2.19	0.42
1:D:4061:SER:O	1:D:4064:GLU:HG3	2.20	0.42
1:D:4731:LEU:HD23	1:D:4731:LEU:HA	1.92	0.42
1:A:983:LEU:HD11	1:A:1055:ARG:HG3	2.00	0.41
1:A:2258:ARG:NE	1:A:2258:ARG:HA	2.35	0.41
1:A:2868:HIS:NE2	1:A:2870:LEU:HB2	2.34	0.41
1:A:3290:ILE:HG22	1:A:3291:ASP:OD2	2.20	0.41
1:A:4061:SER:O	1:A:4064:GLU:HG3	2.20	0.41
1:A:4155:SER:O	1:A:4159:GLN:HG2	2.20	0.41
1:B:674:TYR:OH	1:B:676:GLU:OE2	2.37	0.41
1:C:971:GLN:HE21	1:C:978:PRO:HD2	1.84	0.41
1:C:1721:MET:SD	1:C:2127:ARG:NH1	2.93	0.41
1:C:2114:GLU:HG2	1:C:2115:ASP:N	2.35	0.41
1:C:3036:LEU:O	1:C:3040:LEU:HG	2.20	0.41
1:C:3122:ILE:HG12	1:C:3127:GLN:CG	2.49	0.41
1:C:3717:LYS:HB2	1:C:3717:LYS:HE2	1.89	0.41
1:C:3981:MET:HE3	1:C:3981:MET:HB3	1.93	0.41
1:D:246:THR:HG21	1:D:267:VAL:HG21	2.01	0.41
1:D:881:ILE:O	1:D:885:LEU:HD23	2.20	0.41
1:D:1721:MET:SD	1:D:2127:ARG:NH1	2.93	0.41
1:D:1939:ASN:ND2	1:D:1989:PRO:HD3	2.34	0.41
1:D:2937:HIS:HA	1:D:3014:LEU:HD11	2.02	0.41
1:A:227:TYR:CG	1:A:352:SER:HB3	2.55	0.41
1:A:735:GLY:O	1:A:737:ILE:HD12	2.20	0.41
1:A:824:GLU:HA	1:A:1020:ILE:HD12	2.02	0.41
1:A:841:LYS:O	1:A:848:ARG:NH2	2.53	0.41
1:A:1748:LEU:HD11	1:A:1853:GLU:HG3	2.02	0.41
1:A:2554:ARG:HH11	1:A:2554:ARG:HG3	1.85	0.41
1:A:3728:GLN:OE1	1:A:3770:ASN:ND2	2.52	0.41
1:A:3986:LEU:O	1:A:3989:ASN:HB2	2.19	0.41
1:B:712:GLU:O	1:B:838:ARG:HG3	2.21	0.41
1:B:971:GLN:HE21	1:B:978:PRO:HD2	1.84	0.41
1:B:1721:MET:SD	1:B:2127:ARG:NH1	2.93	0.41
1:B:2711:ILE:HD11	1:B:2783:LEU:HG	2.02	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2759:PRO:HG2	1:B:2762:LEU:HD13	2.01	0.41
1:B:2967:VAL:O	1:B:2970:LEU:HG	2.20	0.41
1:B:3127:GLN:OE1	1:B:3183:ILE:HB	2.20	0.41
1:B:3213:LYS:O	1:B:3216:GLU:HG2	2.20	0.41
1:C:4061:SER:O	1:C:4064:GLU:HG3	2.20	0.41
1:C:4694:LEU:HA	1:C:4697:VAL:HG12	2.02	0.41
1:D:735:GLY:O	1:D:737:ILE:HD12	2.20	0.41
1:D:2172:MET:HG2	1:D:2217:HIS:HD2	1.85	0.41
1:D:2771:TYR:O	1:D:2774:PRO:HD2	2.20	0.41
1:D:4082:GLU:HA	1:D:4085:LYS:HE3	2.03	0.41
1:A:801:ARG:NH1	1:A:1615:GLN:O	2.40	0.41
1:A:4895:ASN:HA	1:A:4905:PHE:CD1	2.56	0.41
1:B:2436:ILE:HG22	1:B:2491:GLY:HA3	2.02	0.41
1:B:2728:SER:HA	1:B:2731:LYS:HZ2	1.86	0.41
1:B:2771:TYR:O	1:B:2774:PRO:HD2	2.20	0.41
1:B:3157:GLU:HG3	1:B:3302:PHE:CZ	2.54	0.41
1:B:4048:PHE:CD2	1:B:4052:MET:HE1	2.56	0.41
1:C:824:GLU:HA	1:C:1020:ILE:HD12	2.02	0.41
1:C:2258:ARG:NE	1:C:2258:ARG:HA	2.35	0.41
1:C:3213:LYS:O	1:C:3216:GLU:HG2	2.20	0.41
1:C:4651:ARG:HG2	1:C:4651:ARG:NH1	2.34	0.41
1:D:712:GLU:O	1:D:838:ARG:HG3	2.21	0.41
1:D:1255:LEU:HD12	1:D:1255:LEU:HA	1.93	0.41
1:D:2403:ALA:HB2	1:D:2475:VAL:HG22	2.01	0.41
1:D:2605:MET:HB3	1:D:2606:PRO:HD3	2.02	0.41
1:D:4735:ASN:HB3	1:D:4738:PHE:HD2	1.86	0.41
1:D:4814:MET:HA	1:D:4814:MET:HE3	2.02	0.41
1:A:114:LEU:HB2	1:A:117:HIS:ND1	2.35	0.41
1:A:829:LYS:HZ2	1:A:1037:LEU:HB3	1.85	0.41
1:A:1064:LEU:HD23	1:A:1064:LEU:H	1.85	0.41
1:A:2303:ARG:HE	1:A:2401:ARG:NE	2.19	0.41
1:A:2954:PHE:CD1	1:A:2955:PRO:HD2	2.56	0.41
1:A:3316:LYS:HE2	1:A:3316:LYS:HB2	1.54	0.41
1:A:4059:THR:OG1	1:A:4062:GLU:HG3	2.21	0.41
1:B:2114:GLU:HG2	1:B:2115:ASP:N	2.35	0.41
1:B:3290:ILE:HG22	1:B:3291:ASP:OD2	2.20	0.41
1:B:4059:THR:OG1	1:B:4062:GLU:HG3	2.21	0.41
1:B:4061:SER:O	1:B:4064:GLU:HG3	2.20	0.41
1:B:4651:ARG:HG2	1:B:4651:ARG:NH1	2.34	0.41
1:B:4694:LEU:HA	1:B:4697:VAL:HG12	2.02	0.41
1:C:2967:VAL:O	1:C:2970:LEU:HG	2.20	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4056:LYS:HE3	1:D:4660:PHE:O	2.20	0.41
1:C:4082:GLU:HA	1:C:4085:LYS:HE3	2.03	0.41
1:D:192:LEU:HD12	1:D:192:LEU:HA	1.91	0.41
1:D:996:VAL:HG21	1:D:1051:ARG:HA	2.02	0.41
1:D:4155:SER:O	1:D:4159:GLN:HG2	2.20	0.41
1:D:4185:LYS:HG2	1:D:4186:MET:HE2	2.01	0.41
1:A:483:LYS:HE2	1:A:483:LYS:HB2	1.80	0.41
1:A:1906:MET:HE3	1:A:1906:MET:HB3	1.83	0.41
1:A:2708:THR:HB	1:A:2780:LYS:HB3	2.02	0.41
1:A:4046:ARG:HG2	1:A:4076:GLU:OE2	2.21	0.41
1:B:114:LEU:HB2	1:B:117:HIS:ND1	2.35	0.41
1:B:2172:MET:HG2	1:B:2217:HIS:HD2	1.86	0.41
1:B:2258:ARG:HA	1:B:2258:ARG:NE	2.35	0.41
1:B:2303:ARG:HE	1:B:2401:ARG:NE	2.19	0.41
1:B:2326:ARG:HA	1:B:2326:ARG:HD3	1.76	0.41
1:B:3195:LEU:HD22	1:B:3197:LEU:HB2	2.01	0.41
1:B:4055:HIS:O	1:B:4057:HIS:ND1	2.54	0.41
1:C:735:GLY:O	1:C:737:ILE:HD12	2.20	0.41
1:C:2666:LEU:HB3	1:C:2667:PRO:HD3	2.02	0.41
1:C:4735:ASN:HB3	1:C:4738:PHE:HD2	1.85	0.41
1:D:1031:ARG:HD2	1:D:1042:THR:HG21	2.01	0.41
1:D:1064:LEU:HD23	1:D:1064:LEU:H	1.85	0.41
1:D:1748:LEU:HD11	1:D:1853:GLU:HG3	2.02	0.41
1:D:1979:PHE:CG	1:D:1993:ARG:HG2	2.55	0.41
1:D:2878:THR:HG23	1:D:2881:GLU:H	1.86	0.41
1:D:3157:GLU:HG3	1:D:3302:PHE:CZ	2.54	0.41
1:A:2172:MET:HG2	1:A:2217:HIS:HD2	1.86	0.41
1:A:2326:ARG:HA	1:A:2326:ARG:HD3	1.76	0.41
1:A:3173:THR:HA	1:A:3176:ASP:OD2	2.21	0.41
1:A:3195:LEU:HD22	1:A:3197:LEU:HB2	2.01	0.41
1:A:3674:THR:O	1:A:3679:LYS:NZ	2.44	0.41
1:A:4055:HIS:O	1:A:4057:HIS:ND1	2.54	0.41
1:A:4694:LEU:HA	1:A:4697:VAL:HG12	2.02	0.41
1:B:801:ARG:HA	1:B:1617:TRP:O	2.20	0.41
1:B:1177:LEU:O	1:B:1180:GLU:HG3	2.21	0.41
1:B:1941:ARG:NH2	1:B:3609:TYR:HB2	2.27	0.41
1:B:2878:THR:HG23	1:B:2881:GLU:H	1.86	0.41
1:B:2937:HIS:HA	1:B:3014:LEU:HD11	2.02	0.41
1:B:3173:THR:HA	1:B:3176:ASP:OD2	2.21	0.41
1:B:3756:ALA:O	1:B:3760:LYS:HG3	2.20	0.41
1:B:4735:ASN:HB3	1:B:4738:PHE:HD2	1.85	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2259:GLU:OE2	1:C:3806:ASN:ND2	2.54	0.41
1:C:3061:LEU:HD23	1:C:3061:LEU:HA	1.88	0.41
1:C:3127:GLN:OE1	1:C:3183:ILE:HB	2.20	0.41
1:C:3712:LYS:O	1:C:3717:LYS:NZ	2.53	0.41
1:C:4070:ALA:HB1	1:C:4078:LEU:HD13	2.03	0.41
1:D:274:LEU:HD23	1:D:274:LEU:HA	1.89	0.41
1:D:308:LEU:HD13	1:D:393:MET:HG3	2.03	0.41
1:D:932:ASN:O	1:D:935:MET:HG3	2.21	0.41
1:D:977:LYS:HA	1:D:978:PRO:HD3	1.89	0.41
1:D:2481:GLN:NE2	1:D:2537:GLY:O	2.42	0.41
1:D:2929:LEU:HD21	1:D:2970:LEU:HD11	2.01	0.41
1:D:3033:LEU:HD23	1:D:3033:LEU:HA	1.86	0.41
1:D:3061:LEU:HD23	1:D:3061:LEU:HA	1.88	0.41
1:D:3611:LEU:HD23	1:D:3611:LEU:HA	1.89	0.41
1:D:4048:PHE:CD2	1:D:4052:MET:HE1	2.56	0.41
1:A:712:GLU:O	1:A:838:ARG:HG3	2.21	0.41
1:A:801:ARG:HA	1:A:1617:TRP:O	2.20	0.41
1:A:881:ILE:O	1:A:885:LEU:HD23	2.20	0.41
1:A:2899:ASN:O	1:A:2899:ASN:ND2	2.52	0.41
1:A:3250:TRP:CD1	1:A:3256:ASN:HD21	2.39	0.41
1:A:4082:GLU:HA	1:A:4085:LYS:HE3	2.03	0.41
1:B:2708:THR:HB	1:B:2780:LYS:HB3	2.02	0.41
1:B:2954:PHE:CD1	1:B:2955:PRO:HD2	2.56	0.41
1:B:3036:LEU:O	1:B:3040:LEU:HG	2.20	0.41
1:B:4070:ALA:HB1	1:B:4078:LEU:HD13	2.03	0.41
1:C:841:LYS:O	1:C:848:ARG:NH2	2.53	0.41
1:C:967:PRO:O	1:C:971:GLN:HG2	2.20	0.41
1:C:996:VAL:HG21	1:C:1051:ARG:HA	2.02	0.41
1:C:3019:ILE:HG13	1:C:3020:SER:N	2.36	0.41
1:C:4048:PHE:CD2	1:C:4052:MET:HE1	2.55	0.41
1:D:314:LEU:HD23	1:D:314:LEU:HA	1.93	0.41
1:D:879:GLU:HA	1:D:882:ARG:HD2	2.03	0.41
1:D:1988:CYS:HA	1:D:1989:PRO:HD3	1.95	0.41
1:D:4757:LEU:O	1:D:4761:THR:HG23	2.21	0.41
1:A:844:ARG:HH11	1:A:844:ARG:HG3	1.86	0.41
1:A:879:GLU:HA	1:A:882:ARG:HD2	2.03	0.41
1:A:1586:LEU:HD12	1:A:1586:LEU:HA	1.94	0.41
1:A:1785:ASP:OD1	1:A:1786:ILE:N	2.48	0.41
1:A:1979:PHE:CG	1:A:1993:ARG:HG2	2.55	0.41
1:A:2222:LEU:HD12	1:A:2261:ASP:HB2	2.03	0.41
1:A:2259:GLU:OE2	1:A:3806:ASN:ND2	2.54	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2605:MET:HB3	1:A:2606:PRO:HD3	2.02	0.41
1:A:3122:ILE:HG12	1:A:3127:GLN:CG	2.50	0.41
1:A:4868:ASP:OD1	1:D:4874:ARG:NH1	2.53	0.41
1:B:20:VAL:HG12	1:B:216:PRO:HA	2.02	0.41
1:B:879:GLU:HA	1:B:882:ARG:HD2	2.03	0.41
1:B:1471:ASP:OD2	1:B:1474:GLY:N	2.51	0.41
1:B:2222:LEU:HD12	1:B:2261:ASP:HB2	2.03	0.41
1:B:3002:GLU:O	1:B:3005:THR:OG1	2.33	0.41
1:B:3929:THR:O	1:B:3933:GLN:HG2	2.21	0.41
1:C:227:TYR:CG	1:C:352:SER:HB3	2.55	0.41
1:C:436:LEU:HD13	1:C:518:ALA:HB2	2.03	0.41
1:C:881:ILE:O	1:C:885:LEU:HD23	2.20	0.41
1:C:1748:LEU:HD11	1:C:1853:GLU:HG3	2.02	0.41
1:C:2605:MET:HB3	1:C:2606:PRO:HD3	2.02	0.41
1:C:3072:MET:HE2	1:C:3072:MET:HB2	1.79	0.41
1:C:3279:ASN:O	1:C:3283:ILE:HG12	2.21	0.41
1:C:3756:ALA:O	1:C:3760:LYS:HG3	2.20	0.41
1:D:456:LEU:HD23	1:D:456:LEU:HA	1.96	0.41
1:D:2258:ARG:NE	1:D:2258:ARG:HA	2.35	0.41
1:D:2259:GLU:OE2	1:D:3806:ASN:ND2	2.54	0.41
1:D:3122:ILE:HG12	1:D:3127:GLN:CG	2.50	0.41
1:D:3965:ILE:HG22	1:D:3969:LYS:HE2	2.03	0.41
2:G:85:ALA:O	2:G:94:PRO:HB3	2.21	0.41
1:A:20:VAL:HG12	1:A:216:PRO:HA	2.02	0.41
1:A:459:LEU:HD23	1:A:459:LEU:HA	1.93	0.41
1:A:658:ASN:HB2	1:A:832:LEU:HD12	2.03	0.41
1:A:891:GLU:HG3	1:A:976:TYR:CE2	2.56	0.41
1:A:932:ASN:HA	1:A:935:MET:HG3	2.02	0.41
1:A:1177:LEU:O	1:A:1180:GLU:HG3	2.21	0.41
1:A:1721:MET:SD	1:A:2127:ARG:NH1	2.93	0.41
1:A:2167:MET:HE3	1:A:2199:PHE:HZ	1.86	0.41
1:A:2711:ILE:HD11	1:A:2783:LEU:HG	2.02	0.41
1:A:2736:LYS:HE2	1:A:2741:TRP:HB3	2.03	0.41
1:A:3187:LYS:HZ2	1:A:3191:GLU:HB3	1.84	0.41
1:A:4048:PHE:CD2	1:A:4052:MET:HE1	2.56	0.41
1:A:4070:ALA:HB1	1:A:4078:LEU:HD13	2.03	0.41
1:B:35:LEU:HD23	1:B:51:SER:HA	2.03	0.41
1:B:245:LEU:HD12	1:B:245:LEU:HA	1.96	0.41
1:B:1799:VAL:HG21	1:B:1845:LEU:HD22	2.03	0.41
1:B:1973:ILE:HB	1:B:3608:LEU:HD21	2.03	0.41
1:B:2222:LEU:HD23	1:B:2222:LEU:HA	1.88	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2283:LYS:HB3	1:B:2283:LYS:HE2	1.87	0.41
1:B:2694:SER:HA	1:B:2702:ASN:O	2.21	0.41
1:B:3250:TRP:CD1	1:B:3256:ASN:HD21	2.39	0.41
1:B:3279:ASN:O	1:B:3283:ILE:HG12	2.21	0.41
1:B:3981:MET:HE3	1:B:3981:MET:HB3	1.93	0.41
1:B:4093:ASP:OD1	1:B:4094:ILE:HG13	2.21	0.41
1:B:4279:MET:HE2	1:C:4484:ILE:HA	2.02	0.41
1:B:4895:ASN:HA	1:B:4905:PHE:CD1	2.56	0.41
1:C:317:MET:HB3	1:C:318:ASP:H	1.75	0.41
1:C:658:ASN:HB2	1:C:832:LEU:HD12	2.03	0.41
1:C:879:GLU:HA	1:C:882:ARG:HD2	2.03	0.41
1:C:1177:LEU:O	1:C:1180:GLU:HG3	2.21	0.41
1:C:1256:PRO:HB2	1:C:1591:LEU:HG	2.03	0.41
1:C:2694:SER:HA	1:C:2702:ASN:O	2.21	0.41
1:C:2736:LYS:HE2	1:C:2741:TRP:HB3	2.03	0.41
1:C:3929:THR:O	1:C:3933:GLN:HG2	2.21	0.41
1:C:3965:ILE:HG22	1:C:3969:LYS:HE2	2.03	0.41
1:C:4059:THR:OG1	1:C:4062:GLU:HG3	2.21	0.41
1:C:4256:MET:HE2	1:C:4256:MET:HB3	1.72	0.41
1:C:4804:MET:HE3	1:C:4804:MET:HB3	1.82	0.41
1:C:4895:ASN:HA	1:C:4905:PHE:CD1	2.56	0.41
1:D:317:MET:HB3	1:D:318:ASP:H	1.75	0.41
1:D:801:ARG:HA	1:D:1617:TRP:O	2.20	0.41
1:D:844:ARG:HG3	1:D:844:ARG:HH11	1.86	0.41
1:D:1304:LEU:HD23	1:D:1304:LEU:HA	1.84	0.41
1:D:1705:LEU:HD12	1:D:1705:LEU:HA	1.90	0.41
1:D:2303:ARG:HE	1:D:2401:ARG:NE	2.19	0.41
1:D:2728:SER:HA	1:D:2731:LYS:HZ2	1.86	0.41
1:D:3215:MET:O	1:D:3219:VAL:HG13	2.21	0.41
1:D:3279:ASN:O	1:D:3283:ILE:HG12	2.21	0.41
1:D:4070:ALA:HB1	1:D:4078:LEU:HD13	2.03	0.41
1:D:4093:ASP:OD1	1:D:4094:ILE:HG13	2.21	0.41
1:D:4651:ARG:HG2	1:D:4651:ARG:NH1	2.34	0.41
1:D:4895:ASN:HA	1:D:4905:PHE:CD1	2.56	0.41
2:F:85:ALA:O	2:F:94:PRO:HB3	2.21	0.41
1:A:713:TRP:HH2	1:A:1251:LEU:HD21	1.86	0.41
1:A:1471:ASP:OD2	1:A:1474:GLY:N	2.51	0.41
1:A:1552:VAL:HG12	1:A:1553:PHE:HD1	1.87	0.41
1:A:2426:LEU:HD23	1:B:143:LEU:HD22	2.02	0.41
1:A:3279:ASN:O	1:A:3283:ILE:HG12	2.21	0.41
1:B:907:VAL:O	1:B:916:PRO:HD3	2.21	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:963:LYS:HD2	1:B:979:ALA:O	2.21	0.41
1:B:1256:PRO:HB2	1:B:1591:LEU:HG	2.03	0.41
1:B:1839:LEU:HD12	1:B:1839:LEU:HA	1.93	0.41
1:B:2666:LEU:HB3	1:B:2667:PRO:HD3	2.02	0.41
1:B:2718:GLU:OE2	1:B:2722:ASN:ND2	2.54	0.41
1:B:4082:GLU:HA	1:B:4085:LYS:HE3	2.03	0.41
1:C:712:GLU:O	1:C:838:ARG:HG3	2.21	0.41
1:C:2878:THR:HG23	1:C:2881:GLU:H	1.86	0.41
1:C:2937:HIS:HA	1:C:3014:LEU:HD11	2.02	0.41
1:D:336:GLU:H	1:D:336:GLU:HG2	1.74	0.41
1:D:658:ASN:HB2	1:D:832:LEU:HD12	2.03	0.41
1:D:824:GLU:HA	1:D:1020:ILE:HD12	2.02	0.41
1:D:1415:ASP:OD1	1:D:1416:ASP:N	2.54	0.41
1:D:1973:ILE:HB	1:D:3608:LEU:HD21	2.03	0.41
1:D:2114:GLU:HG2	1:D:2115:ASP:N	2.35	0.41
1:D:2718:GLU:OE2	1:D:2722:ASN:ND2	2.54	0.41
1:D:2967:VAL:O	1:D:2970:LEU:HG	2.20	0.41
1:D:4694:LEU:HA	1:D:4697:VAL:HG12	2.02	0.41
1:A:35:LEU:HD23	1:A:51:SER:HA	2.03	0.40
1:A:308:LEU:HD13	1:A:393:MET:HG3	2.03	0.40
1:A:884:LYS:HA	1:A:887:GLU:CD	2.47	0.40
1:A:2443:PRO:HD3	1:A:2512:MET:HG2	2.03	0.40
1:A:2878:THR:HG23	1:A:2881:GLU:H	1.86	0.40
1:A:2967:VAL:O	1:A:2970:LEU:HG	2.20	0.40
1:A:3215:MET:O	1:A:3219:VAL:HG13	2.21	0.40
1:A:4757:LEU:O	1:A:4761:THR:HG23	2.21	0.40
1:B:483:LYS:HE2	1:B:483:LYS:HB2	1.80	0.40
1:B:844:ARG:HH11	1:B:844:ARG:HG3	1.86	0.40
1:B:1552:VAL:HG12	1:B:1553:PHE:HD1	1.87	0.40
1:B:2605:MET:HB3	1:B:2606:PRO:HD3	2.02	0.40
1:B:2764:SER:HB3	1:B:2767:GLU:HG3	2.03	0.40
1:B:3187:LYS:HZ2	1:B:3191:GLU:HB3	1.85	0.40
1:B:4757:LEU:O	1:B:4761:THR:HG23	2.21	0.40
1:B:4795:LYS:HA	1:B:4795:LYS:HD2	1.81	0.40
1:C:35:LEU:HD23	1:C:51:SER:HA	2.03	0.40
1:C:1799:VAL:HG21	1:C:1845:LEU:HD22	2.03	0.40
1:C:2443:PRO:HD3	1:C:2512:MET:HG2	2.03	0.40
1:C:3274:ASN:HA	1:C:3277:LEU:HG	2.03	0.40
1:C:4575:LEU:HD23	1:C:4575:LEU:HA	1.91	0.40
1:D:114:LEU:HB2	1:D:117:HIS:ND1	2.35	0.40
1:D:674:TYR:OH	1:D:676:GLU:OE2	2.37	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:891:GLU:HG3	1:D:976:TYR:CE2	2.56	0.40
1:D:1829:LEU:HD12	1:D:1829:LEU:HA	1.97	0.40
1:D:2443:PRO:HD3	1:D:2512:MET:HG2	2.03	0.40
2:G:19:LYS:HB2	2:G:19:LYS:HE3	1.80	0.40
1:A:233:VAL:HG12	1:A:274:LEU:HD22	2.04	0.40
1:A:244:CYS:SG	1:A:267:VAL:HG13	2.62	0.40
1:A:659:ILE:HD13	1:A:822:CYS:HB3	2.03	0.40
1:A:1031:ARG:HD2	1:A:1042:THR:HG21	2.01	0.40
1:A:1988:CYS:HA	1:A:1989:PRO:HD3	1.95	0.40
1:A:2114:GLU:HG2	1:A:2115:ASP:N	2.35	0.40
1:A:2764:SER:HB3	1:A:2767:GLU:HG3	2.03	0.40
1:A:3282:LYS:HA	1:A:3285:TYR:CD2	2.57	0.40
1:A:4800:ASP:OD1	1:A:4801:THR:N	2.55	0.40
1:B:658:ASN:HB2	1:B:832:LEU:HD12	2.03	0.40
1:B:732:LEU:HD23	1:B:732:LEU:HA	1.89	0.40
1:B:1586:LEU:HD12	1:B:1586:LEU:HA	1.94	0.40
1:B:3304:GLN:OE1	1:B:3308:ASN:ND2	2.48	0.40
1:B:3323:MET:HB3	1:B:3327:LYS:HZ3	1.86	0.40
1:B:3975:GLN:O	1:B:3979:VAL:HG23	2.21	0.40
1:C:114:LEU:HB2	1:C:117:HIS:ND1	2.35	0.40
1:C:419:ILE:HD13	1:C:492:GLU:HG3	2.03	0.40
1:C:891:GLU:HG3	1:C:976:TYR:CE2	2.56	0.40
1:C:963:LYS:HD2	1:C:979:ALA:O	2.21	0.40
1:C:2652:LEU:HD12	1:C:2652:LEU:HA	1.91	0.40
1:C:3074:ASN:OD1	1:C:3075:LEU:N	2.55	0.40
1:C:3213:LYS:O	1:C:3217:GLU:OE1	2.39	0.40
1:C:3319:PHE:O	1:C:3323:MET:HG2	2.21	0.40
1:C:4024:LYS:HB2	1:C:4088:HIS:CE1	2.57	0.40
1:C:4055:HIS:O	1:C:4057:HIS:ND1	2.54	0.40
1:D:436:LEU:HD13	1:D:518:ALA:HB2	2.03	0.40
1:D:1177:LEU:O	1:D:1180:GLU:HG3	2.21	0.40
1:D:1473:LYS:NZ	1:D:1475:LYS:HB2	2.36	0.40
1:D:1505:LEU:HD23	1:D:1506:GLU:N	2.37	0.40
1:D:2742:ILE:HG23	1:D:2753:VAL:HG23	2.04	0.40
1:D:2954:PHE:CD1	1:D:2955:PRO:HD2	2.56	0.40
1:D:3130:CYS:HB3	1:D:3162:PHE:CZ	2.56	0.40
1:D:3282:LYS:HA	1:D:3285:TYR:CD2	2.57	0.40
1:D:3929:THR:O	1:D:3933:GLN:HG2	2.21	0.40
1:A:892:LEU:HD21	1:A:980:PRO:CD	2.51	0.40
1:A:1272:ARG:HH22	1:A:1584:PRO:HA	1.87	0.40
1:A:1415:ASP:OD1	1:A:1416:ASP:N	2.54	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1973:ILE:HB	1:A:3608:LEU:HD21	2.03	0.40
1:A:2234:MET:HE2	1:A:2234:MET:HB3	1.90	0.40
1:A:2742:ILE:HG23	1:A:2753:VAL:HG23	2.03	0.40
1:A:3033:LEU:HD23	1:A:3033:LEU:HA	1.86	0.40
1:A:3213:LYS:O	1:A:3216:GLU:HG2	2.20	0.40
1:A:3981:MET:HE3	1:A:3981:MET:HB3	1.93	0.40
1:B:2443:PRO:HD3	1:B:2512:MET:HG2	2.03	0.40
1:B:3212:GLU:O	1:B:3216:GLU:OE1	2.40	0.40
1:C:801:ARG:HA	1:C:1617:TRP:O	2.20	0.40
1:C:907:VAL:O	1:C:916:PRO:HD3	2.21	0.40
1:C:1505:LEU:HD23	1:C:1506:GLU:N	2.37	0.40
1:C:1973:ILE:HB	1:C:3608:LEU:HD21	2.03	0.40
1:C:2303:ARG:HE	1:C:2401:ARG:NE	2.19	0.40
1:C:2573:LEU:HD12	1:C:2573:LEU:HA	1.88	0.40
1:C:2651:ALA:O	1:C:2655:LYS:HB2	2.21	0.40
1:C:3130:CYS:HB3	1:C:3162:PHE:CZ	2.56	0.40
1:D:244:CYS:SG	1:D:267:VAL:HG13	2.62	0.40
1:D:419:ILE:HD13	1:D:492:GLU:HG3	2.03	0.40
1:D:1256:PRO:HB2	1:D:1591:LEU:HG	2.03	0.40
1:D:2736:LYS:HE2	1:D:2741:TRP:HB3	2.03	0.40
1:D:2764:SER:HB3	1:D:2767:GLU:HG3	2.03	0.40
1:D:2965:LYS:HA	1:D:2965:LYS:HD3	1.80	0.40
1:D:3319:PHE:O	1:D:3323:MET:HG2	2.22	0.40
1:D:4575:LEU:HD23	1:D:4575:LEU:HA	1.90	0.40
1:A:192:LEU:HD12	1:A:192:LEU:HA	1.91	0.40
1:A:963:LYS:HD2	1:A:979:ALA:O	2.21	0.40
1:A:2790:GLU:O	1:A:2902:ALA:N	2.42	0.40
1:A:3130:CYS:HB3	1:A:3162:PHE:CZ	2.56	0.40
1:A:4038:ASP:HB3	1:A:4040:LYS:NZ	2.37	0.40
1:A:4196:THR:HG22	1:A:4200:MET:SD	2.62	0.40
1:A:4567:TYR:O	1:A:4567:TYR:CD2	2.75	0.40
1:A:4698:LEU:HD13	1:D:4262:LYS:HG3	2.03	0.40
1:B:680:ASP:N	1:B:799:LYS:O	2.54	0.40
1:B:891:GLU:HG3	1:B:976:TYR:CE2	2.56	0.40
1:B:1415:ASP:OD1	1:B:1416:ASP:N	2.54	0.40
1:B:2693:SER:O	1:B:2702:ASN:HB3	2.22	0.40
1:B:3215:MET:O	1:B:3219:VAL:HG13	2.21	0.40
1:B:3322:LEU:O	1:B:3326:LEU:HG	2.22	0.40
1:C:20:VAL:HG12	1:C:216:PRO:HA	2.02	0.40
1:C:262:TYR:HB2	1:C:389:ARG:HG3	2.04	0.40
1:C:932:ASN:O	1:C:935:MET:HG3	2.21	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2090:ARG:HE	1:C:2090:ARG:HB3	1.51	0.40
1:C:2167:MET:HE3	1:C:2199:PHE:HZ	1.86	0.40
1:C:2771:TYR:O	1:C:2774:PRO:HD2	2.20	0.40
1:C:2954:PHE:CD1	1:C:2955:PRO:HD2	2.56	0.40
1:C:4196:THR:HG22	1:C:4200:MET:SD	2.62	0.40
1:D:233:VAL:HG12	1:D:274:LEU:HD22	2.04	0.40
1:D:262:TYR:HB2	1:D:389:ARG:HG3	2.04	0.40
1:D:963:LYS:HD2	1:D:979:ALA:O	2.21	0.40
1:D:1799:VAL:HG21	1:D:1845:LEU:HD22	2.03	0.40
1:D:1948:MET:HA	1:D:1948:MET:HE2	2.04	0.40
1:D:2573:LEU:HD12	1:D:2573:LEU:HA	1.88	0.40
1:D:3213:LYS:O	1:D:3217:GLU:OE1	2.39	0.40
1:D:3274:ASN:HA	1:D:3277:LEU:HG	2.03	0.40
1:D:3322:LEU:O	1:D:3326:LEU:HG	2.22	0.40
1:D:4046:ARG:HG2	1:D:4076:GLU:OE2	2.21	0.40
1:A:1256:PRO:HB2	1:A:1591:LEU:HG	2.03	0.40
1:A:1300:MET:HE2	1:A:1300:MET:HB3	1.71	0.40
1:A:1473:LYS:NZ	1:A:1475:LYS:HB2	2.36	0.40
1:A:2487:LEU:HD12	1:A:2487:LEU:HA	1.94	0.40
1:A:2959:GLU:OE1	1:A:2959:GLU:N	2.53	0.40
1:A:3222:ALA:HA	1:A:3283:ILE:HG22	2.04	0.40
1:A:3965:ILE:HG22	1:A:3969:LYS:HE2	2.03	0.40
1:B:262:TYR:HB2	1:B:389:ARG:HG3	2.04	0.40
1:B:436:LEU:HD13	1:B:518:ALA:HB2	2.03	0.40
1:B:884:LYS:HA	1:B:887:GLU:CD	2.46	0.40
1:B:1272:ARG:HH22	1:B:1584:PRO:HA	1.87	0.40
1:B:1473:LYS:NZ	1:B:1475:LYS:HB2	2.36	0.40
1:B:2736:LYS:HE2	1:B:2741:TRP:HB3	2.03	0.40
1:B:4046:ARG:HG2	1:B:4076:GLU:OE2	2.21	0.40
1:C:844:ARG:HG3	1:C:844:ARG:HH11	1.86	0.40
1:C:1109:THR:O	1:C:1113:MET:HE1	2.22	0.40
1:C:1552:VAL:HG12	1:C:1553:PHE:HD1	1.87	0.40
1:C:2252:GLU:HB2	1:C:3819:MET:SD	2.62	0.40
1:C:3215:MET:O	1:C:3219:VAL:HG13	2.21	0.40
1:C:3975:GLN:O	1:C:3979:VAL:HG23	2.21	0.40
1:C:4800:ASP:OD1	1:C:4801:THR:N	2.55	0.40
1:D:2322:ARG:HA	1:D:2325:ILE:HG12	2.04	0.40
1:D:2651:ALA:O	1:D:2655:LYS:HB2	2.21	0.40
1:D:3173:THR:HA	1:D:3176:ASP:OD2	2.21	0.40
1:D:3213:LYS:O	1:D:3216:GLU:HG2	2.20	0.40
1:D:3295:TRP:O	1:D:3299:LEU:HG	2.22	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3994:THR:O	1:D:3998:GLN:HG3	2.22	0.40
1:D:4059:THR:OG1	1:D:4062:GLU:HG3	2.21	0.40
2:H:85:ALA:O	2:H:94:PRO:HB3	2.21	0.40

There are no symmetry-related clashes.

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 PDB entries followed by that with respect to all EM entries.

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	
1	A	4198/4967 (84%)	4097 (98%)	99 (2%)	2 (0%)	100	100
1	B	4198/4967 (84%)	4099 (98%)	97 (2%)	2 (0%)	100	100
1	C	4198/4967 (84%)	4099 (98%)	97 (2%)	2 (0%)	100	100
1	D	4198/4967 (84%)	4098 (98%)	98 (2%)	2 (0%)	100	100
2	E	105/108 (97%)	102 (97%)	3 (3%)	0	100	100
2	F	105/108 (97%)	102 (97%)	3 (3%)	0	100	100
2	G	105/108 (97%)	103 (98%)	2 (2%)	0	100	100
2	H	105/108 (97%)	102 (97%)	3 (3%)	0	100	100
All	All	17212/20300 (85%)	16802 (98%)	402 (2%)	8 (0%)	100	100

All (8) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	A	2988	ARG
1	A	4641	PRO
1	B	2988	ARG
1	B	4641	PRO
1	C	2988	ARG
1	C	4641	PRO
1	D	2988	ARG

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	D	4641	PRO

5.3.2 Protein sidechains ⓘ

In the following table, the Percentiles column shows the percent sidechain outliers of the chain as a percentile score with respect to all PDB entries followed by that with respect to all EM entries.

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	
1	A	3708/4358 (85%)	3688 (100%)	20 (0%)	81	82
1	B	3708/4358 (85%)	3688 (100%)	20 (0%)	81	82
1	C	3708/4358 (85%)	3688 (100%)	20 (0%)	81	82
1	D	3708/4358 (85%)	3688 (100%)	20 (0%)	81	82
2	E	88/89 (99%)	86 (98%)	2 (2%)	44	66
2	F	88/89 (99%)	86 (98%)	2 (2%)	44	66
2	G	88/89 (99%)	86 (98%)	2 (2%)	44	66
2	H	88/89 (99%)	86 (98%)	2 (2%)	44	66
All	All	15184/17788 (85%)	15096 (99%)	88 (1%)	76	80

All (88) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	317	MET
1	A	995	MET
1	A	999	LEU
1	A	1300	MET
1	A	1564	MET
1	A	2172	MET
1	A	2681	MET
1	A	2695	MET
1	A	2772	ARG
1	A	2884	LYS
1	A	3057	LEU
1	A	3072	MET
1	A	3123	LEU
1	A	3190	ARG

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	A	3215	MET
1	A	3241	MET
1	A	3316	LYS
1	A	4002	MET
1	A	4256	MET
1	A	4504	MET
2	E	18	LYS
2	E	45	LYS
1	B	317	MET
1	B	995	MET
1	B	999	LEU
1	B	1300	MET
1	B	1564	MET
1	B	2172	MET
1	B	2681	MET
1	B	2695	MET
1	B	2772	ARG
1	B	2884	LYS
1	B	3057	LEU
1	B	3072	MET
1	B	3123	LEU
1	B	3190	ARG
1	B	3215	MET
1	B	3241	MET
1	B	3316	LYS
1	B	4002	MET
1	B	4256	MET
1	B	4504	MET
1	C	317	MET
1	C	995	MET
1	C	999	LEU
1	C	1300	MET
1	C	1564	MET
1	C	2172	MET
1	C	2681	MET
1	C	2695	MET
1	C	2772	ARG
1	C	2884	LYS
1	C	3057	LEU
1	C	3072	MET
1	C	3123	LEU
1	C	3190	ARG

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	C	3215	MET
1	C	3241	MET
1	C	3316	LYS
1	C	4002	MET
1	C	4256	MET
1	C	4504	MET
1	D	317	MET
1	D	995	MET
1	D	999	LEU
1	D	1300	MET
1	D	1564	MET
1	D	2172	MET
1	D	2681	MET
1	D	2695	MET
1	D	2772	ARG
1	D	2884	LYS
1	D	3057	LEU
1	D	3072	MET
1	D	3123	LEU
1	D	3190	ARG
1	D	3215	MET
1	D	3241	MET
1	D	3316	LYS
1	D	4002	MET
1	D	4256	MET
1	D	4504	MET
2	F	18	LYS
2	F	45	LYS
2	G	18	LYS
2	G	45	LYS
2	H	18	LYS
2	H	45	LYS

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (208) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	29	HIS
1	A	57	ASN
1	A	71	GLN
1	A	150	GLN
1	A	193	HIS
1	A	313	ASN

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	A	604	HIS
1	A	635	ASN
1	A	658	ASN
1	A	716	ASN
1	A	731	HIS
1	A	808	HIS
1	A	903	GLN
1	A	1021	GLN
1	A	1233	GLN
1	A	1498	GLN
1	A	1551	ASN
1	A	1554	GLN
1	A	1577	ASN
1	A	1581	GLN
1	A	1710	HIS
1	A	1905	GLN
1	A	1917	GLN
1	A	1920	HIS
1	A	1974	ASN
1	A	2251	ASN
1	A	2802	ASN
1	A	2830	ASN
1	A	2973	GLN
1	A	3031	ASN
1	A	3111	HIS
1	A	3114	GLN
1	A	3178	HIS
1	A	3179	ASN
1	A	3256	ASN
1	A	3274	ASN
1	A	3622	GLN
1	A	3774	GLN
1	A	3775	GLN
1	A	3831	GLN
1	A	3869	ASN
1	A	4055	HIS
1	A	4097	ASN
1	A	4178	ASN
1	A	4201	GLN
1	A	4762	HIS
1	A	4766	GLN
1	A	4787	ASN

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	A	4879	GLN
2	E	26	HIS
2	E	33	ASN
2	E	106	ASN
1	B	29	HIS
1	B	57	ASN
1	B	71	GLN
1	B	150	GLN
1	B	193	HIS
1	B	313	ASN
1	B	658	ASN
1	B	716	ASN
1	B	731	HIS
1	B	781	ASN
1	B	808	HIS
1	B	903	GLN
1	B	1022	GLN
1	B	1233	GLN
1	B	1551	ASN
1	B	1554	GLN
1	B	1577	ASN
1	B	1581	GLN
1	B	1710	HIS
1	B	1905	GLN
1	B	1917	GLN
1	B	1920	HIS
1	B	1974	ASN
1	B	1978	ASN
1	B	2251	ASN
1	B	2802	ASN
1	B	2973	GLN
1	B	3031	ASN
1	B	3111	HIS
1	B	3114	GLN
1	B	3178	HIS
1	B	3179	ASN
1	B	3256	ASN
1	B	3274	ASN
1	B	3622	GLN
1	B	3774	GLN
1	B	3775	GLN
1	B	3831	GLN

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	B	3844	GLN
1	B	3869	ASN
1	B	3915	GLN
1	B	3953	HIS
1	B	4055	HIS
1	B	4097	ASN
1	B	4178	ASN
1	B	4201	GLN
1	B	4762	HIS
1	B	4787	ASN
1	B	4879	GLN
1	C	29	HIS
1	C	57	ASN
1	C	71	GLN
1	C	150	GLN
1	C	193	HIS
1	C	604	HIS
1	C	635	ASN
1	C	658	ASN
1	C	716	ASN
1	C	731	HIS
1	C	808	HIS
1	C	903	GLN
1	C	1021	GLN
1	C	1233	GLN
1	C	1294	ASN
1	C	1498	GLN
1	C	1554	GLN
1	C	1577	ASN
1	C	1581	GLN
1	C	1710	HIS
1	C	1905	GLN
1	C	1917	GLN
1	C	1920	HIS
1	C	1974	ASN
1	C	2251	ASN
1	C	2639	HIS
1	C	2802	ASN
1	C	2973	GLN
1	C	3031	ASN
1	C	3111	HIS
1	C	3114	GLN

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	C	3178	HIS
1	C	3179	ASN
1	C	3256	ASN
1	C	3274	ASN
1	C	3622	GLN
1	C	3774	GLN
1	C	3775	GLN
1	C	3831	GLN
1	C	3869	ASN
1	C	4055	HIS
1	C	4097	ASN
1	C	4178	ASN
1	C	4201	GLN
1	C	4558	HIS
1	C	4736	ASN
1	C	4762	HIS
1	C	4766	GLN
1	C	4787	ASN
1	C	4879	GLN
1	D	29	HIS
1	D	57	ASN
1	D	71	GLN
1	D	150	GLN
1	D	193	HIS
1	D	313	ASN
1	D	394	HIS
1	D	604	HIS
1	D	658	ASN
1	D	716	ASN
1	D	731	HIS
1	D	781	ASN
1	D	903	GLN
1	D	1233	GLN
1	D	1294	ASN
1	D	1498	GLN
1	D	1551	ASN
1	D	1554	GLN
1	D	1577	ASN
1	D	1581	GLN
1	D	1710	HIS
1	D	1905	GLN
1	D	1917	GLN

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	D	1920	HIS
1	D	1974	ASN
1	D	2251	ASN
1	D	2802	ASN
1	D	2973	GLN
1	D	3031	ASN
1	D	3114	GLN
1	D	3178	HIS
1	D	3179	ASN
1	D	3256	ASN
1	D	3274	ASN
1	D	3622	GLN
1	D	3774	GLN
1	D	3775	GLN
1	D	3831	GLN
1	D	3869	ASN
1	D	4055	HIS
1	D	4097	ASN
1	D	4178	ASN
1	D	4201	GLN
1	D	4558	HIS
1	D	4762	HIS
1	D	4766	GLN
1	D	4787	ASN
1	D	4879	GLN
2	F	26	HIS
2	F	33	ASN
2	F	106	ASN
2	G	26	HIS
2	G	33	ASN
2	G	106	ASN
2	H	26	HIS
2	H	33	ASN
2	H	106	ASN

5.3.3 RNA ⓘ

There are no RNA molecules in this entry.

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

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 12 ligands modelled in this entry, 4 are monoatomic - leaving 8 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	$\# Z > 2$
4	ATP	D	5002	-	32,33,33	0.34	0	48,52,52	0.36	0
4	ATP	C	5003	-	32,33,33	0.26	0	48,52,52	0.27	0
4	ATP	B	5002	-	32,33,33	0.33	0	48,52,52	0.36	0
4	ATP	D	5003	-	32,33,33	0.26	0	48,52,52	0.28	0
4	ATP	A	5003	-	32,33,33	0.26	0	48,52,52	0.27	0
4	ATP	B	5003	-	32,33,33	0.26	0	48,52,52	0.27	0
4	ATP	C	5002	-	32,33,33	0.34	0	48,52,52	0.36	0
4	ATP	A	5002	-	32,33,33	0.34	0	48,52,52	0.36	0

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
4	ATP	D	5002	-	-	10/22/38/38	0/3/3/3
4	ATP	C	5003	-	-	7/22/38/38	0/3/3/3
4	ATP	B	5002	-	-	10/22/38/38	0/3/3/3
4	ATP	D	5003	-	-	7/22/38/38	0/3/3/3
4	ATP	A	5003	-	-	7/22/38/38	0/3/3/3
4	ATP	B	5003	-	-	7/22/38/38	0/3/3/3
4	ATP	C	5002	-	-	10/22/38/38	0/3/3/3
4	ATP	A	5002	-	-	10/22/38/38	0/3/3/3

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

All (68) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
4	A	5002	ATP	C5'-O5'-PA-O1A
4	A	5002	ATP	C5'-O5'-PA-O3A
4	A	5003	ATP	C5'-O5'-PA-O1A
4	A	5003	ATP	C5'-O5'-PA-O2A
4	A	5003	ATP	C5'-O5'-PA-O3A
4	A	5003	ATP	O4'-C4'-C5'-O5'
4	B	5002	ATP	C5'-O5'-PA-O1A
4	B	5002	ATP	C5'-O5'-PA-O3A
4	B	5003	ATP	C5'-O5'-PA-O1A
4	B	5003	ATP	C5'-O5'-PA-O2A
4	B	5003	ATP	C5'-O5'-PA-O3A
4	B	5003	ATP	O4'-C4'-C5'-O5'
4	C	5002	ATP	C5'-O5'-PA-O1A
4	C	5002	ATP	C5'-O5'-PA-O3A
4	C	5003	ATP	C5'-O5'-PA-O1A
4	C	5003	ATP	C5'-O5'-PA-O2A
4	C	5003	ATP	C5'-O5'-PA-O3A
4	C	5003	ATP	O4'-C4'-C5'-O5'
4	D	5002	ATP	C5'-O5'-PA-O1A
4	D	5002	ATP	C5'-O5'-PA-O3A
4	D	5003	ATP	C5'-O5'-PA-O1A
4	D	5003	ATP	C5'-O5'-PA-O2A
4	D	5003	ATP	C5'-O5'-PA-O3A
4	D	5003	ATP	O4'-C4'-C5'-O5'
4	A	5003	ATP	C3'-C4'-C5'-O5'
4	B	5003	ATP	C3'-C4'-C5'-O5'
4	C	5003	ATP	C3'-C4'-C5'-O5'
4	D	5003	ATP	C3'-C4'-C5'-O5'
4	B	5003	ATP	PA-O3A-PB-O3B
4	D	5003	ATP	PA-O3A-PB-O3B
4	A	5002	ATP	PB-O3A-PA-O1A
4	B	5002	ATP	PB-O3A-PA-O1A
4	C	5002	ATP	PB-O3A-PA-O1A
4	D	5002	ATP	PB-O3A-PA-O1A
4	A	5003	ATP	PA-O3A-PB-O3B
4	C	5003	ATP	PA-O3A-PB-O3B
4	A	5002	ATP	PB-O3A-PA-O5'
4	A	5003	ATP	PB-O3A-PA-O5'
4	B	5002	ATP	PB-O3A-PA-O5'

Continued on next page...

Continued from previous page...

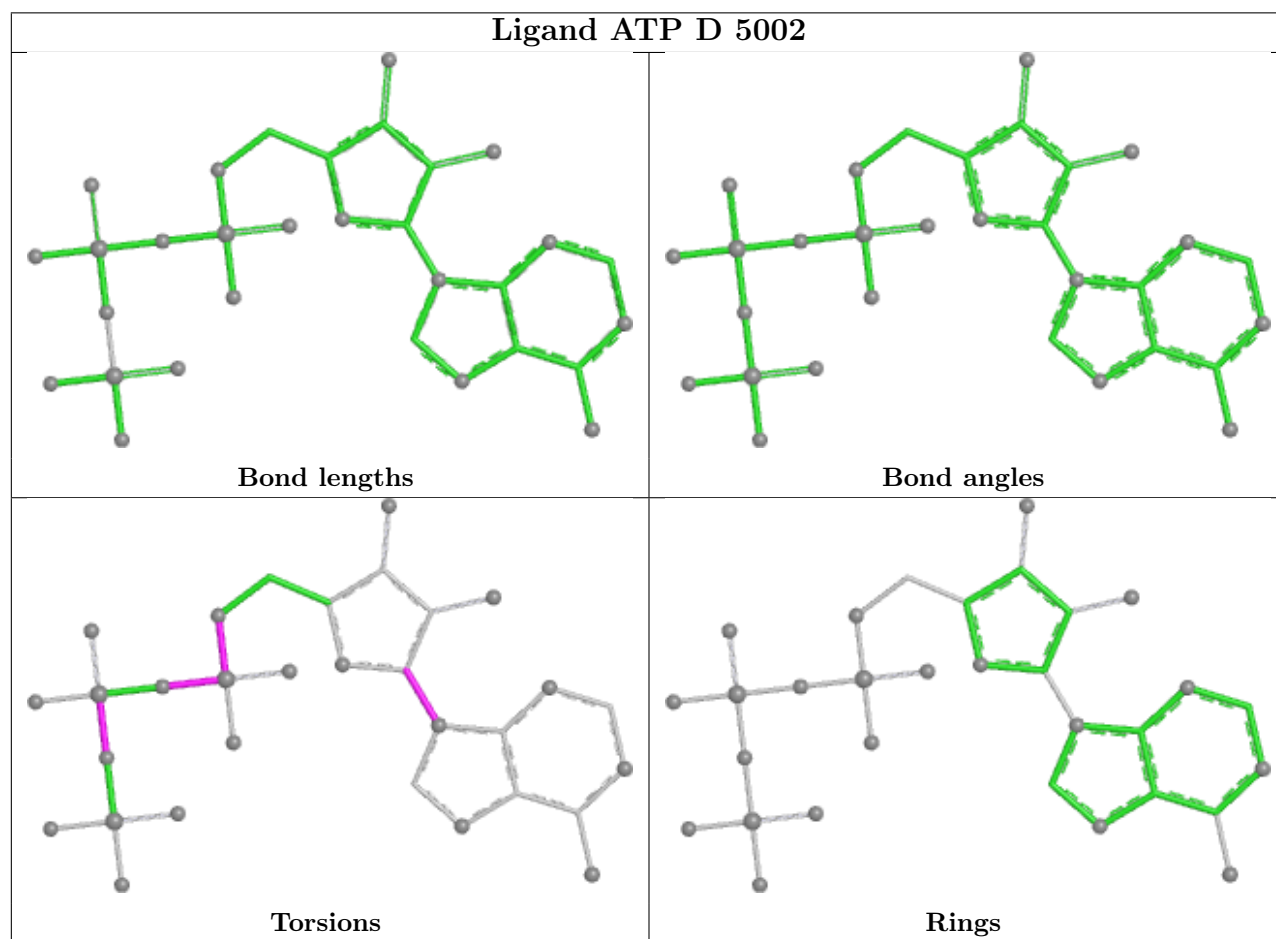
Mol	Chain	Res	Type	Atoms
4	B	5003	ATP	PB-O3A-PA-O5'
4	C	5002	ATP	PB-O3A-PA-O5'
4	C	5003	ATP	PB-O3A-PA-O5'
4	D	5002	ATP	PB-O3A-PA-O5'
4	D	5003	ATP	PB-O3A-PA-O5'
4	A	5002	ATP	C5'-O5'-PA-O2A
4	B	5002	ATP	C5'-O5'-PA-O2A
4	C	5002	ATP	C5'-O5'-PA-O2A
4	D	5002	ATP	C5'-O5'-PA-O2A
4	A	5002	ATP	O4'-C1'-N9-C8
4	B	5002	ATP	O4'-C1'-N9-C8
4	C	5002	ATP	O4'-C1'-N9-C8
4	D	5002	ATP	O4'-C1'-N9-C8
4	A	5002	ATP	O4'-C1'-N9-C4
4	B	5002	ATP	O4'-C1'-N9-C4
4	C	5002	ATP	O4'-C1'-N9-C4
4	D	5002	ATP	O4'-C1'-N9-C4
4	A	5002	ATP	PG-O3B-PB-O3A
4	B	5002	ATP	PG-O3B-PB-O3A
4	C	5002	ATP	PG-O3B-PB-O3A
4	D	5002	ATP	PG-O3B-PB-O3A
4	A	5002	ATP	PG-O3B-PB-O1B
4	A	5002	ATP	PG-O3B-PB-O2B
4	B	5002	ATP	PG-O3B-PB-O1B
4	B	5002	ATP	PG-O3B-PB-O2B
4	C	5002	ATP	PG-O3B-PB-O1B
4	C	5002	ATP	PG-O3B-PB-O2B
4	D	5002	ATP	PG-O3B-PB-O1B
4	D	5002	ATP	PG-O3B-PB-O2B

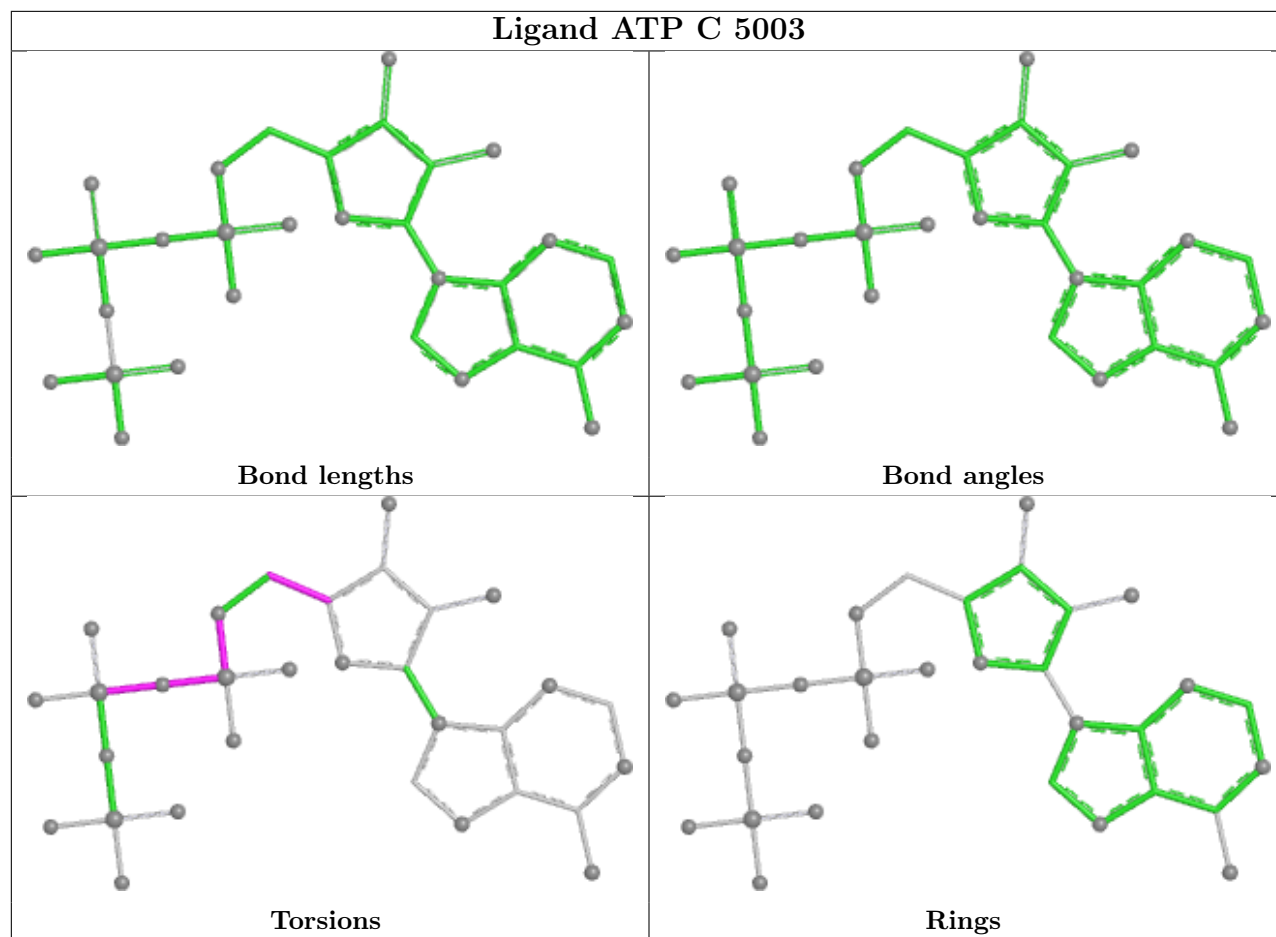
There are no ring outliers.

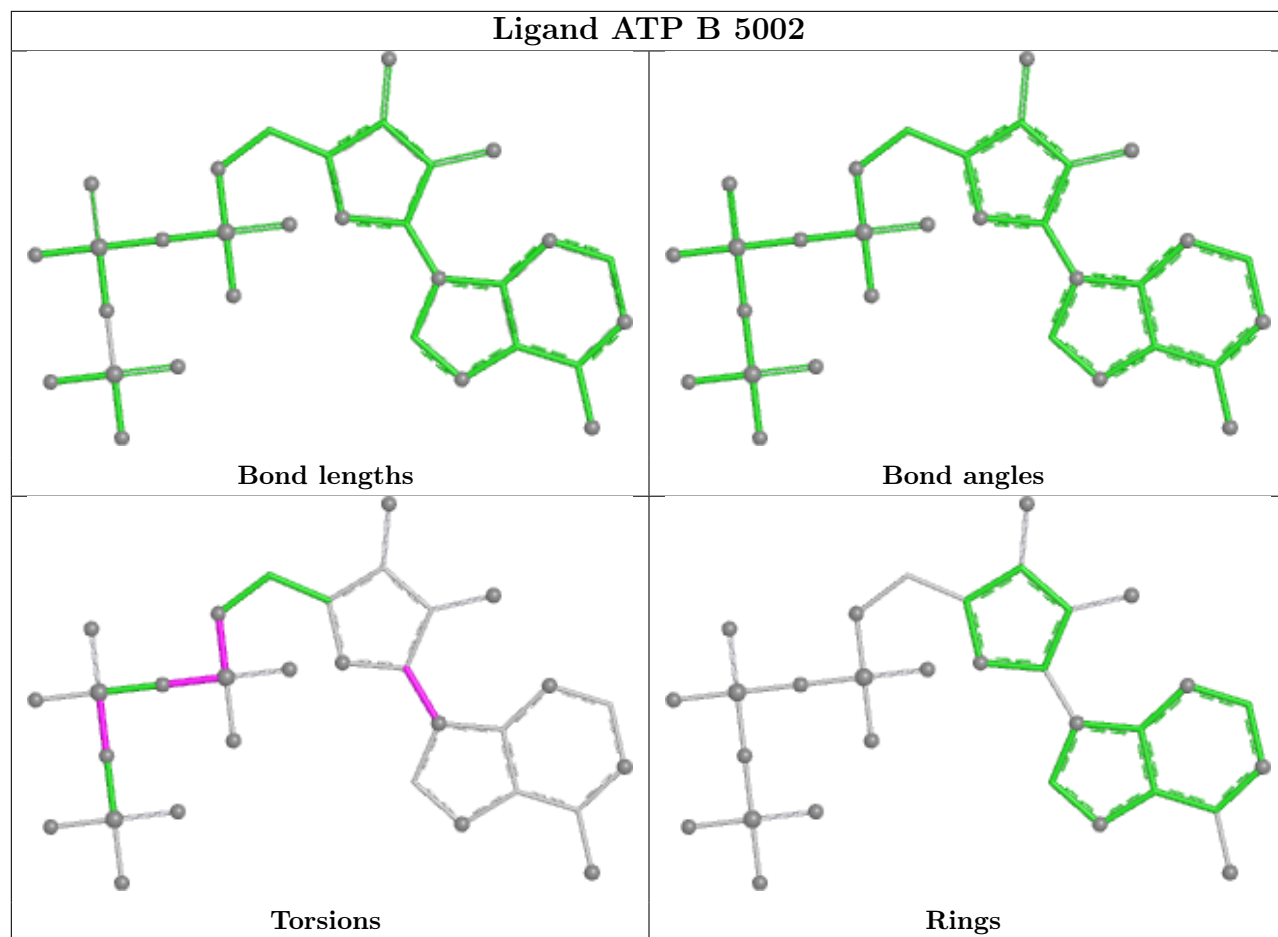
No monomer is involved in short contacts.

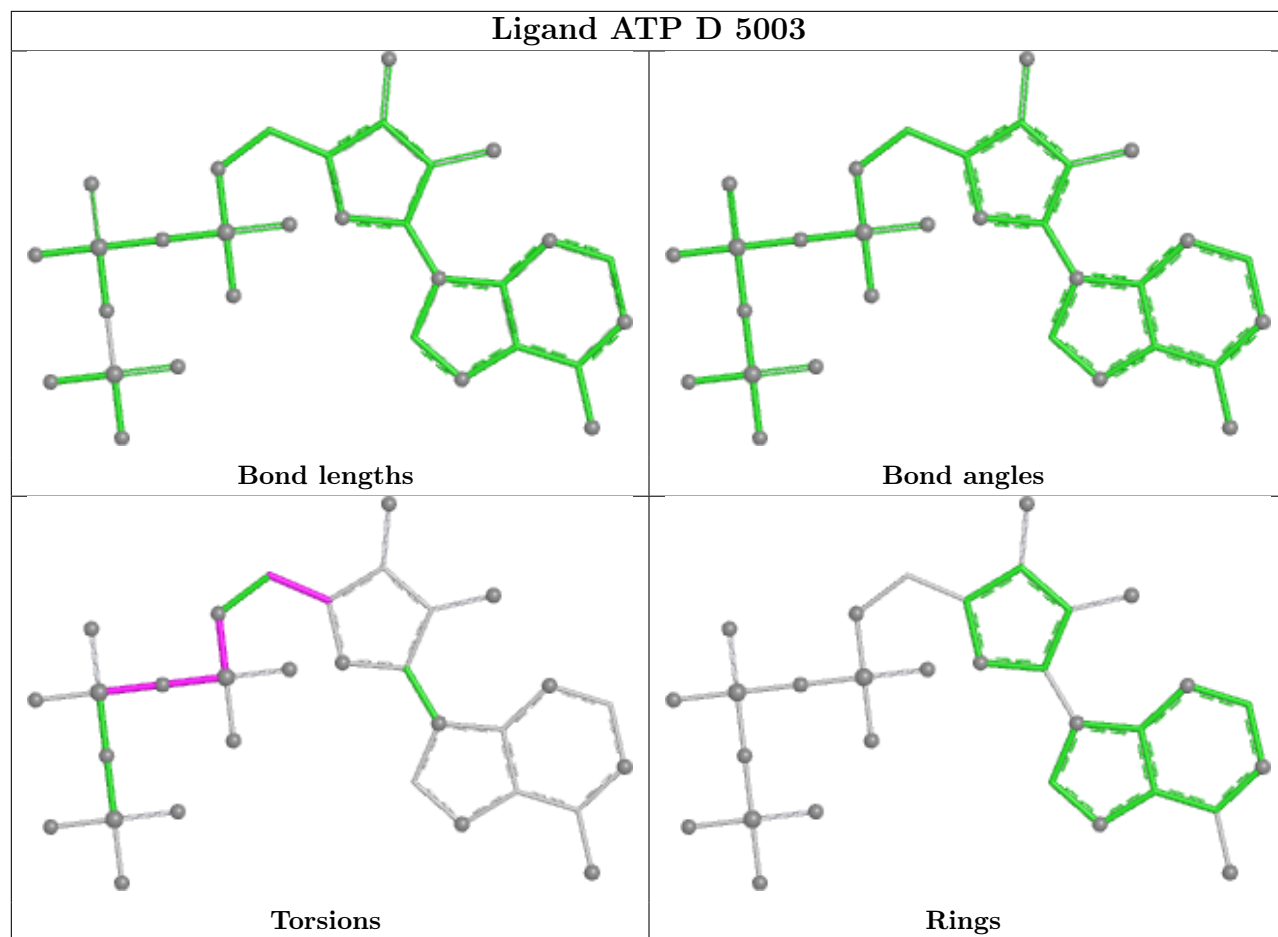
The following is a two-dimensional graphical depiction of Mogul quality analysis of bond lengths, bond angles, torsion angles, and ring geometry for all instances of the Ligand of Interest. In addition, ligands with molecular weight > 250 and outliers as shown on the validation Tables will also be included. For torsion angles, if less than 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.

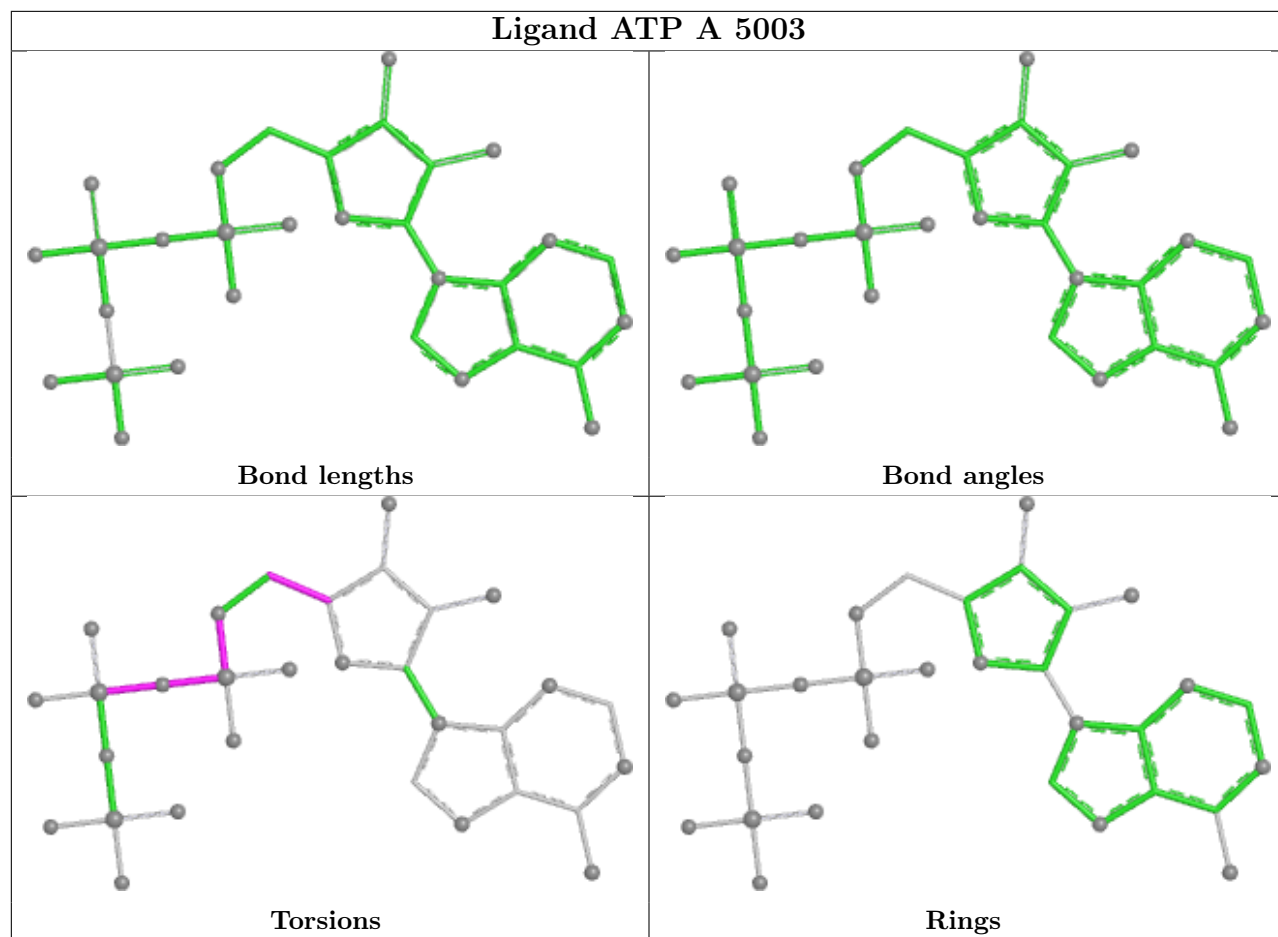
The outliers are highlighted in purple. The color gray indicates Mogul did not find sufficient equivalents in the CSD to analyse the geometry.

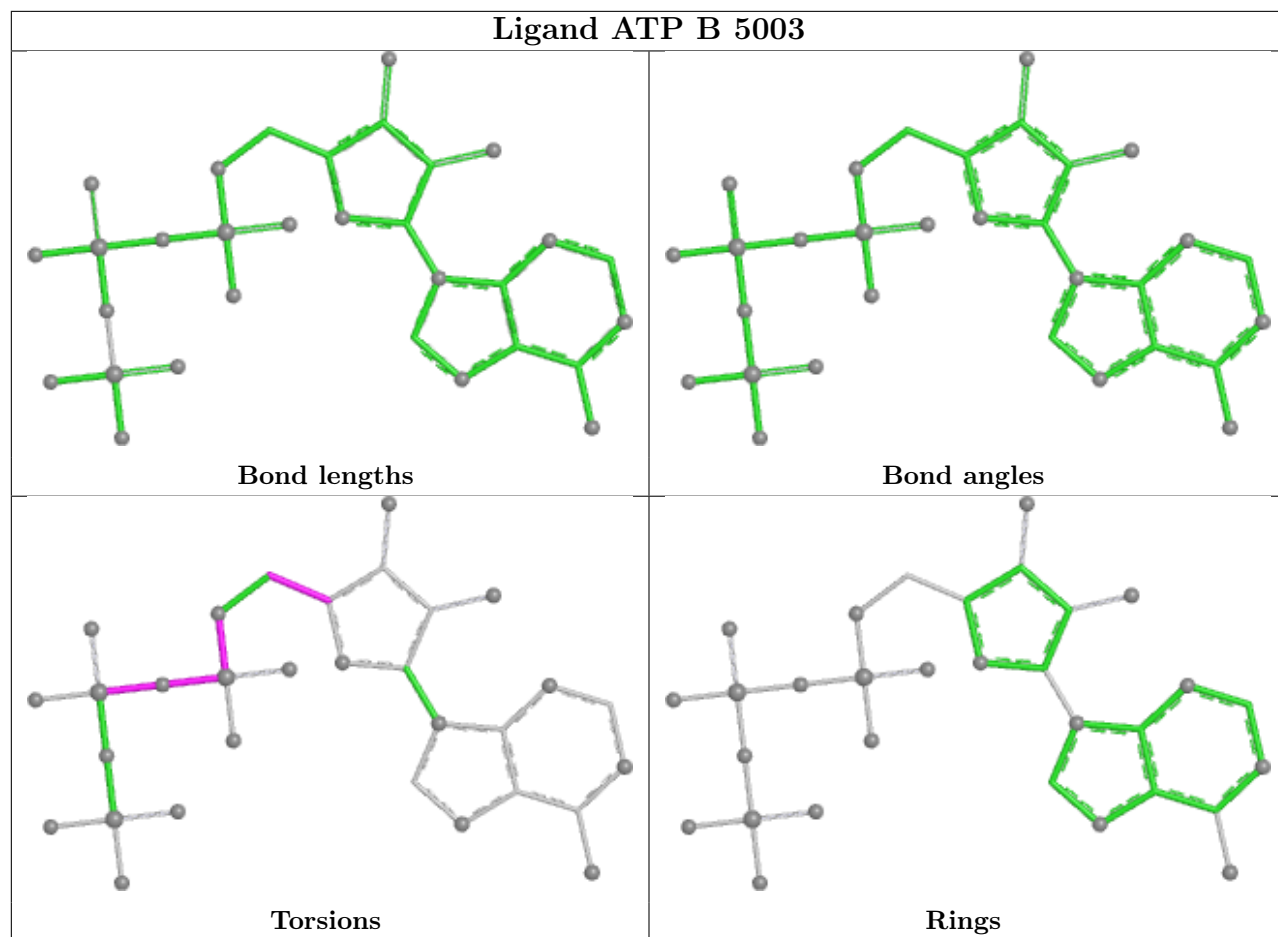


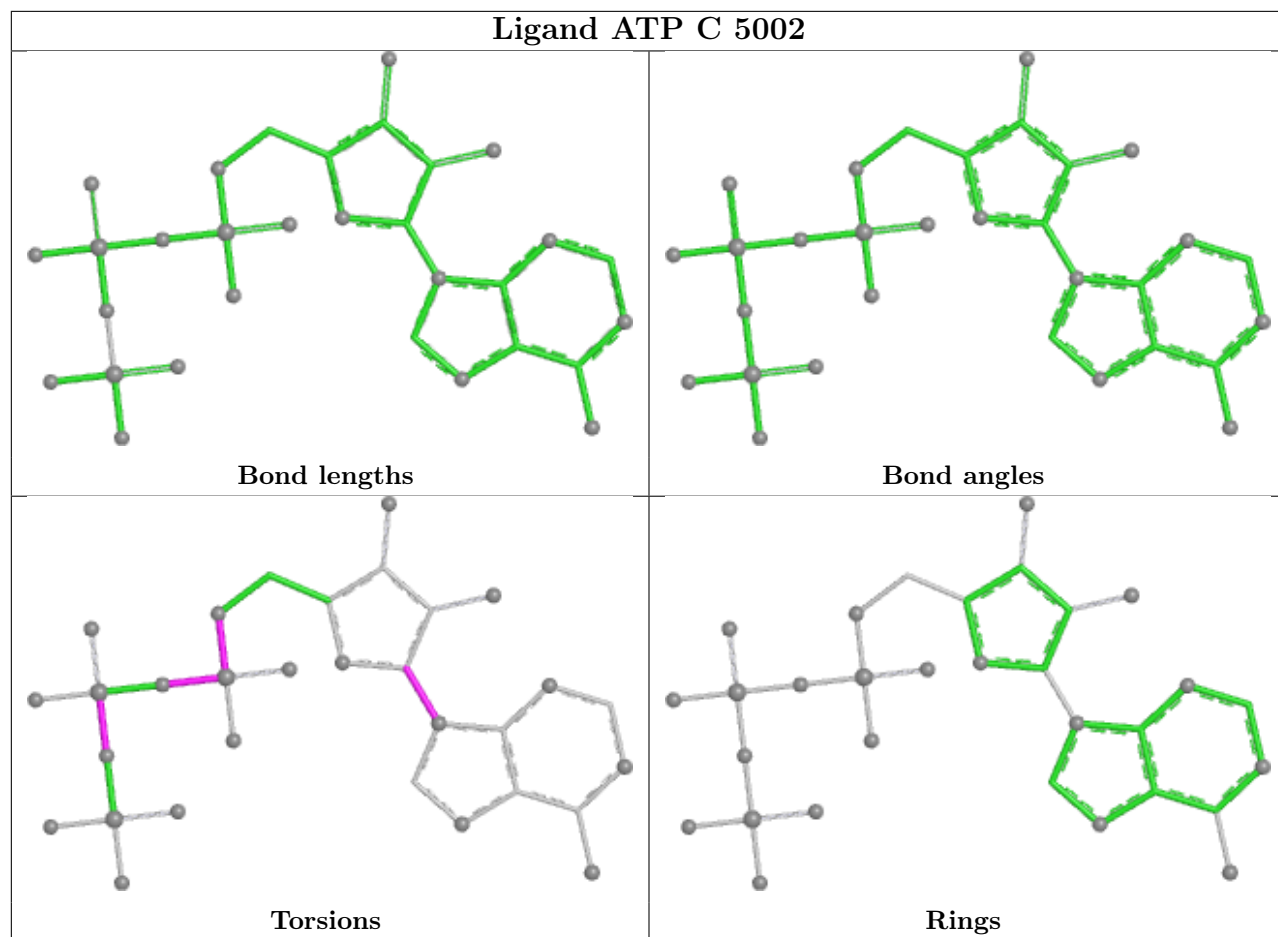


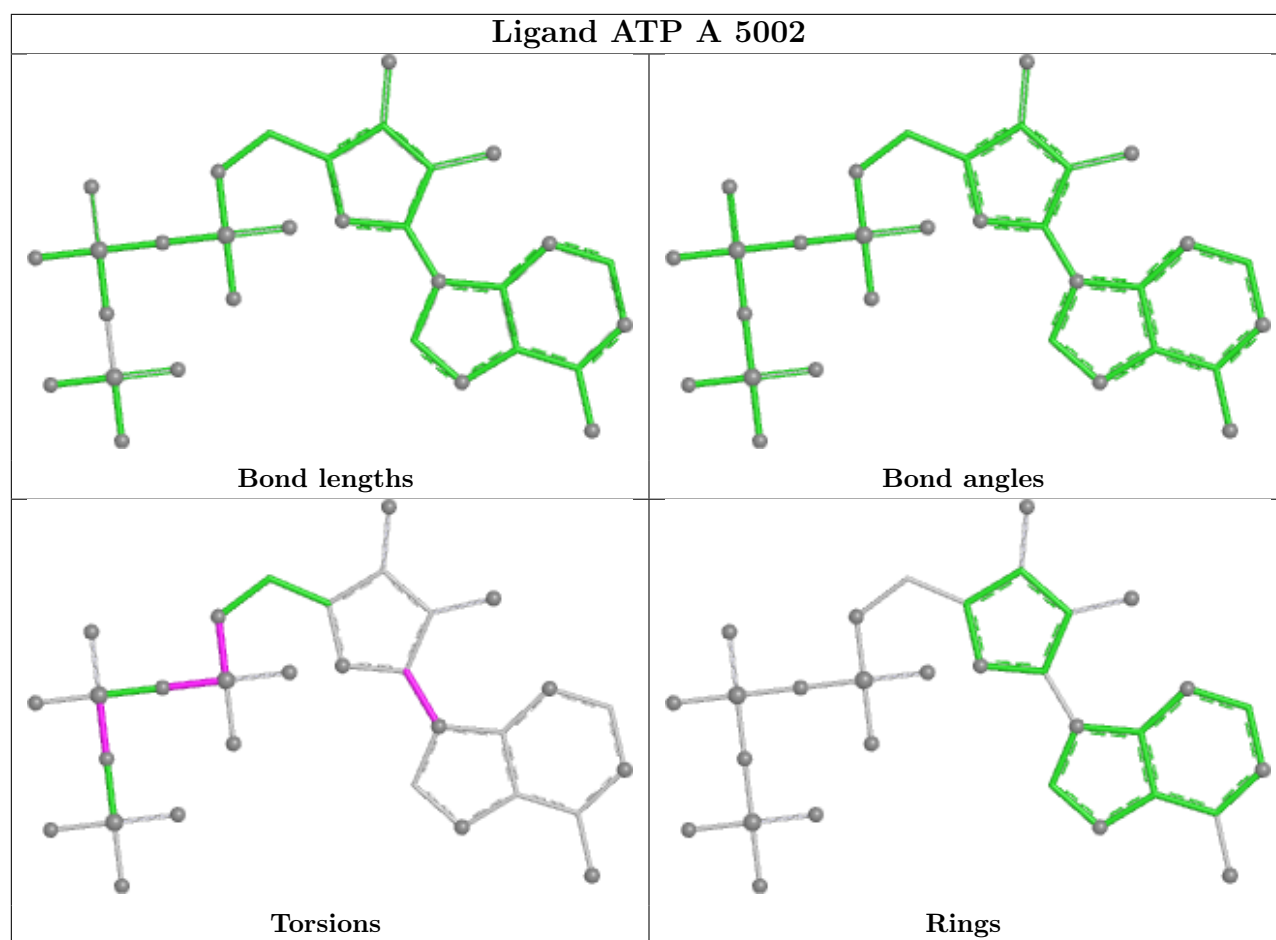












5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-42762. These allow visual inspection of the internal detail of the map and identification of artifacts.

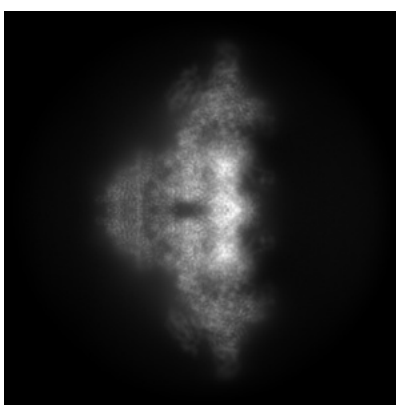
No raw map or half-maps were deposited for this entry and therefore no images, graphs, etc. pertaining to the raw map can be shown.

6.1 Orthogonal projections [i](#)

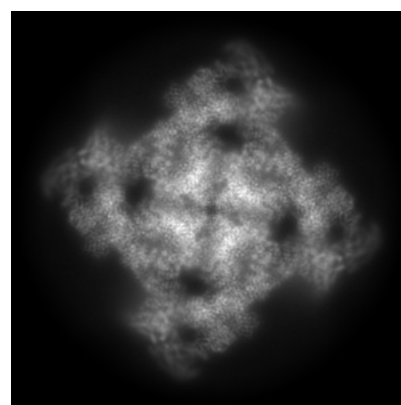
6.1.1 Primary map



X



Y

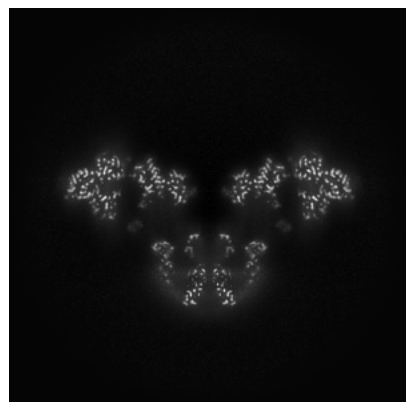


Z

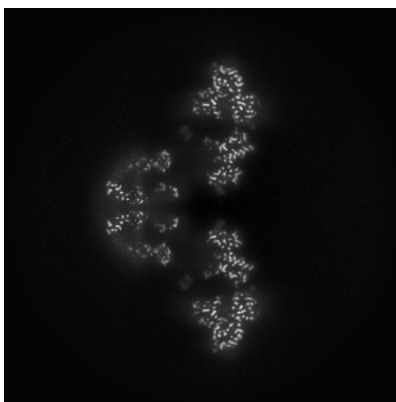
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

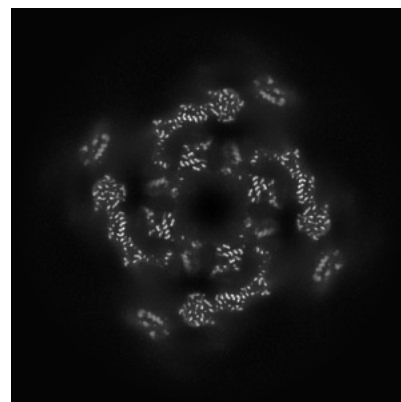
6.2.1 Primary map



X Index: 256



Y Index: 256



Z Index: 256

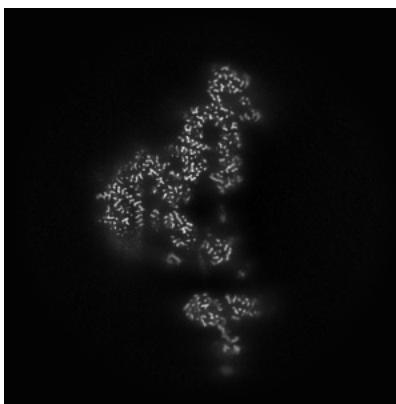
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

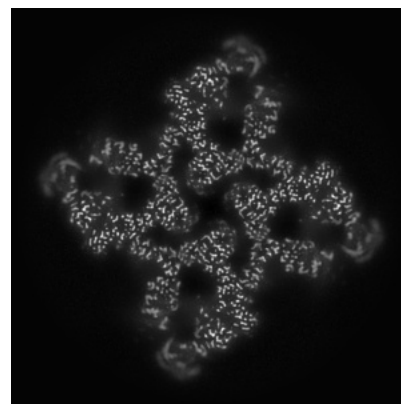
6.3.1 Primary map



X Index: 238



Y Index: 274

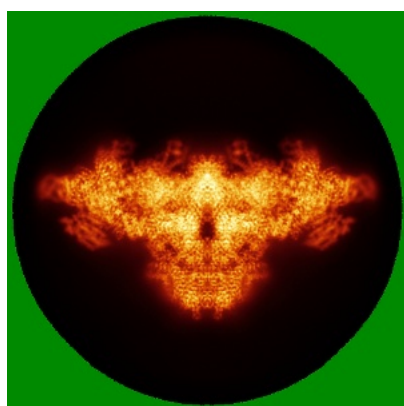


Z Index: 282

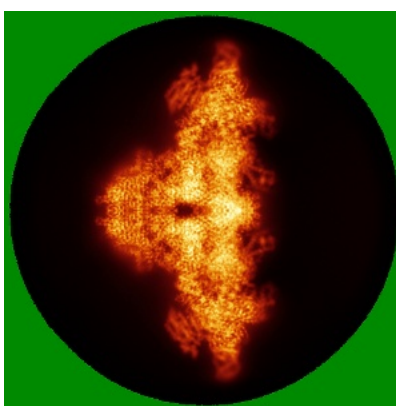
The images above show the largest variance slices of the map in three orthogonal directions.

6.4 Orthogonal standard-deviation projections (False-color) [i](#)

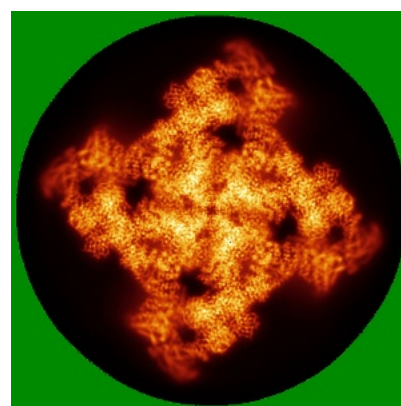
6.4.1 Primary map



X



Y

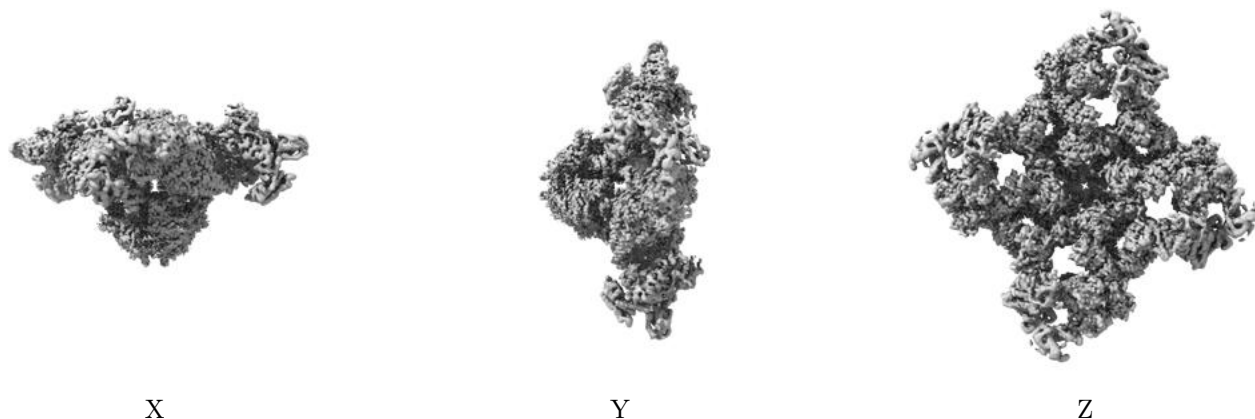


Z

The images above show the map standard deviation projections with false color in three orthogonal directions. Minimum values are shown in green, max in blue, and dark to light orange shades represent small to large values respectively.

6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.12. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

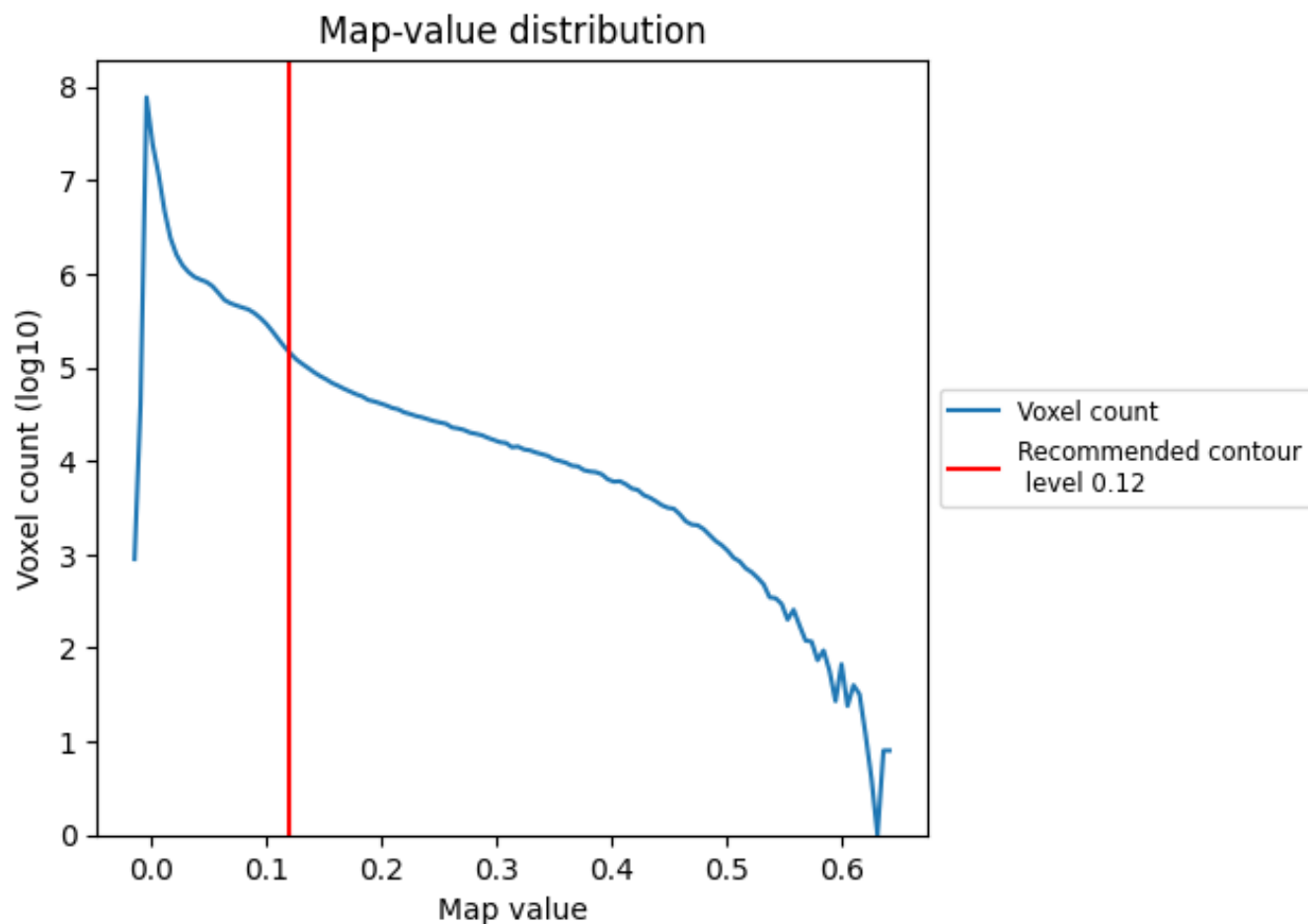
6.6 Mask visualisation [i](#)

This section was not generated. No masks/segmentation were deposited.

7 Map analysis [i](#)

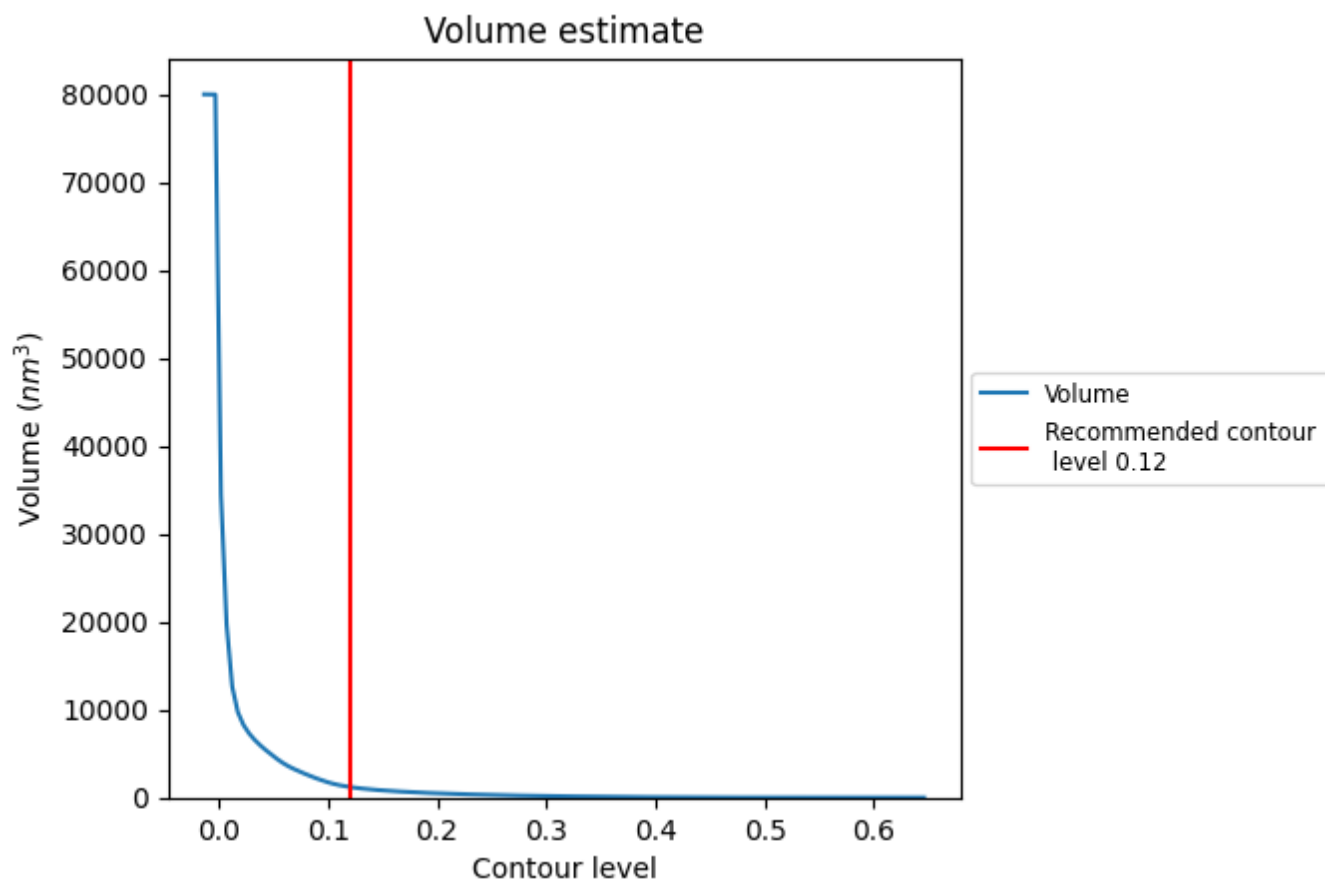
This section contains the results of statistical analysis of the map.

7.1 Map-value distribution [i](#)



The map-value distribution is plotted in 128 intervals along the x-axis. The y-axis is logarithmic. A spike in this graph at zero usually indicates that the volume has been masked.

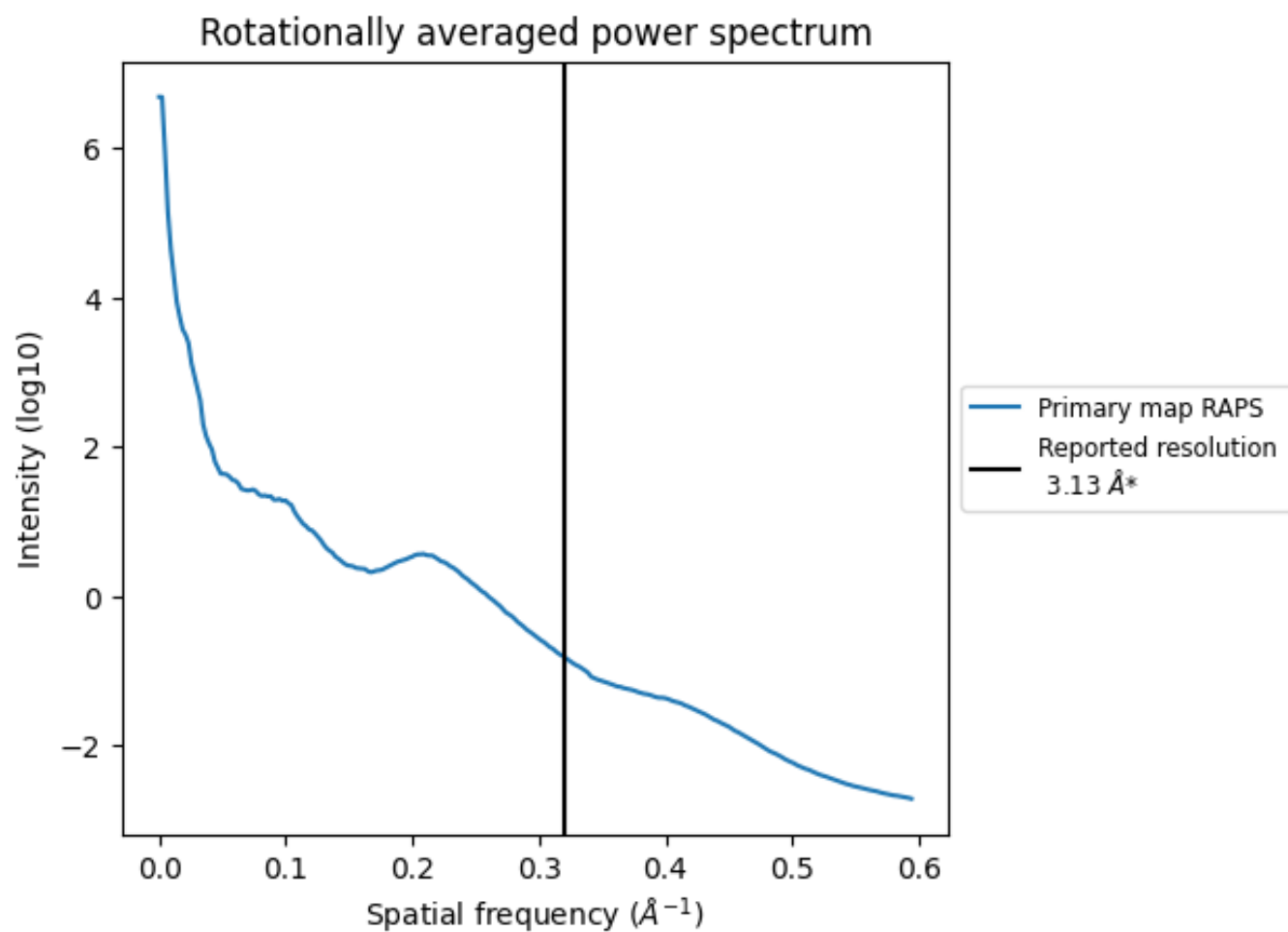
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 1200 nm³; this corresponds to an approximate mass of 1084 kDa.

The volume estimate graph shows how the enclosed volume varies with the contour level. The recommended contour level is shown as a vertical line and the intersection between the line and the curve gives the volume of the enclosed surface at the given level.

7.3 Rotationally averaged power spectrum ⓘ



*Reported resolution corresponds to spatial frequency of 0.319 Å⁻¹

8 Fourier-Shell correlation

This section was not generated. No FSC curve or half-maps provided.

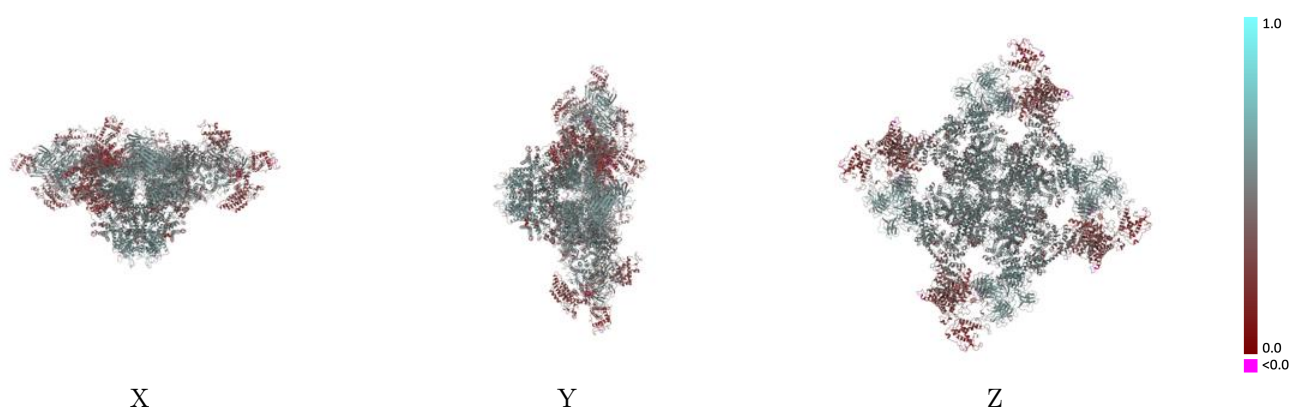
9 Map-model fit [i](#)

This section contains information regarding the fit between EMDB map EMD-42762 and PDB model 8UXF. Per-residue inclusion information can be found in section 3 on page 7.

9.1 Map-model overlay [i](#)

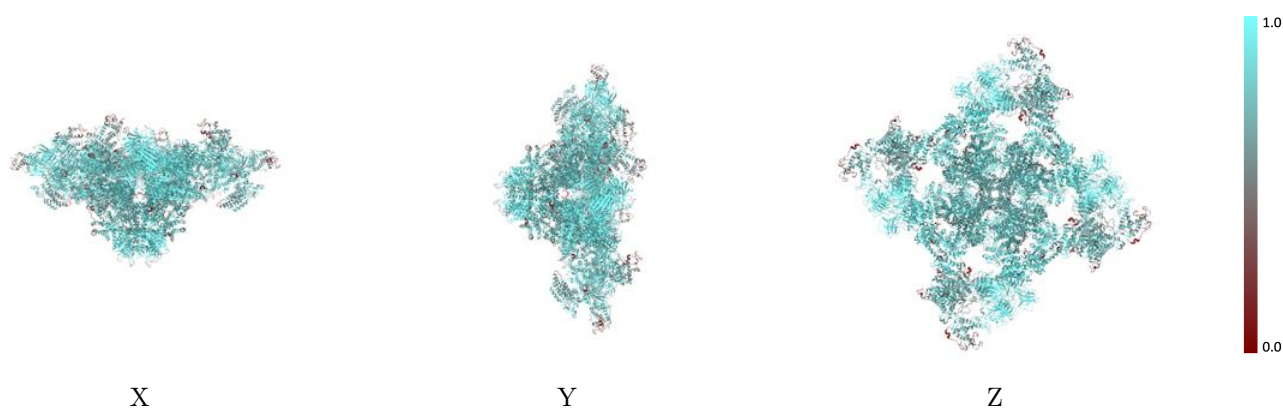
This section was not generated.

9.2 Q-score mapped to coordinate model [i](#)



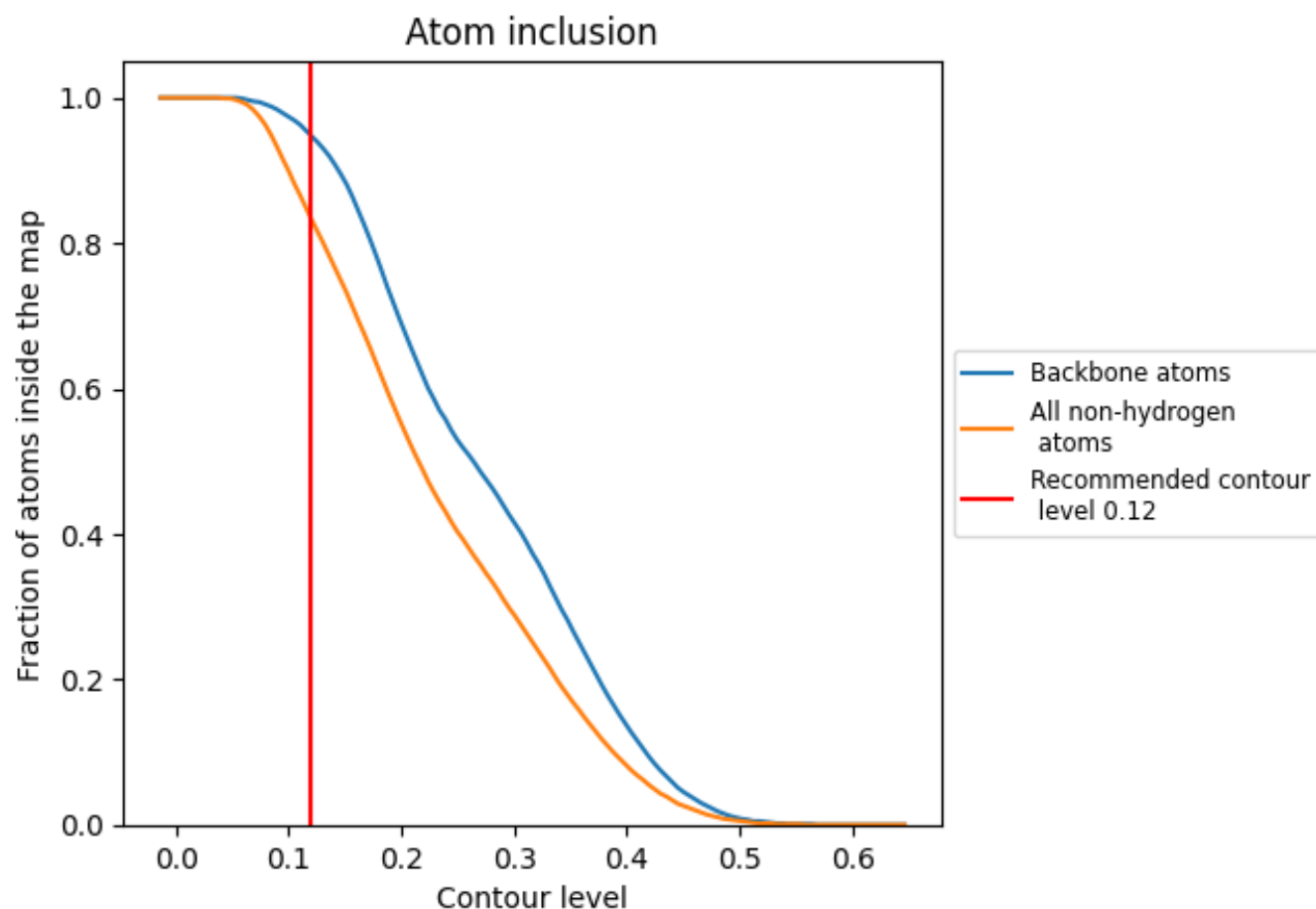
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions at which atoms can be resolved. Low Q-score values may therefore be expected for many entries.

9.3 Atom inclusion mapped to coordinate model [i](#)



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (0.12).

9.4 Atom inclusion [i](#)



At the recommended contour level, 95% of all backbone atoms, 83% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary ⓘ

The table lists the average atom inclusion at the recommended contour level (0.12) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	0.8330	0.4540
A	0.8300	0.4510
B	0.8300	0.4510
C	0.8310	0.4530
D	0.8310	0.4520
E	0.9430	0.5530
F	0.9420	0.5530
G	0.9430	0.5540
H	0.9440	0.5540

1.0

0.0

<0.0