



Full wwPDB X-ray Structure Validation Report ⓘ

Mar 6, 2026 – 01:29 PM UTC

PDB ID : 1QVG / pdb_00001qvg
Title : Structure of CCA oligonucleotide bound to the tRNA binding sites of the large ribosomal subunit of *Haloarcula marismortui*
Authors : Schmeing, T.M.; Moore, P.B.; Steitz, T.A.
Deposited on : 2003-08-27
Resolution : 2.90 Å(reported)

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

We welcome your comments at validation@mail.wwpdb.org

A user guide is available at

<https://www.wwpdb.org/validation/2017/XrayValidationReportHelp>

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

<http://www.wwpdb.org/validation/2017/FAQs#types>.

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

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

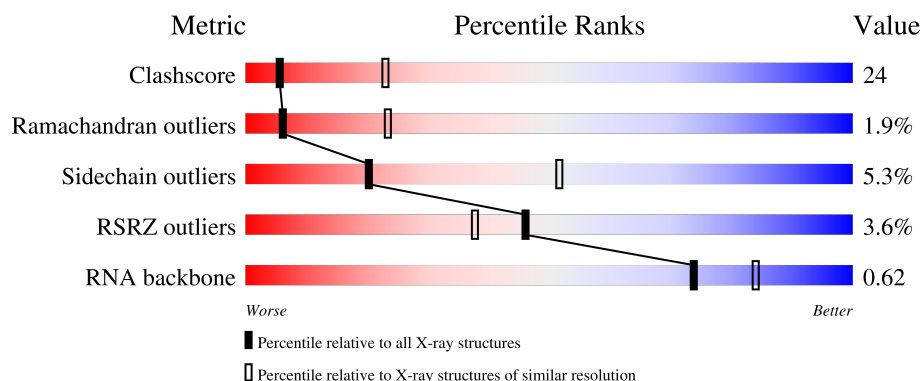
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 2.90 Å.

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



Metric	Whole archive (#Entries)	Similar resolution (#Entries, resolution range(Å))
Clashscore	190562	2690 (2.90-2.90)
Ramachandran outliers	187476	2623 (2.90-2.90)
Sidechain outliers	187428	2625 (2.90-2.90)
RSRZ outliers	180081	2481 (2.90-2.90)
RNA backbone	3983	1120 (3.10-2.70)

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

Mol	Chain	Length	Quality of chain
1	0	2922	45% 41% 7% 6%
2	9	122	4% 44% 39% 13%
3	3	3	33% 33% 67%
3	4	3	33% 33% 67%
3	5	3	67% 33%

Continued on next page...

Continued from previous page...

Mol	Chain	Length	Quality of chain
4	A	239	
5	B	337	
6	C	246	
7	D	176	
8	E	177	
9	F	119	
10	G	348	
11	H	167	
12	I	145	
13	J	132	
14	K	164	
15	L	194	
16	M	186	
17	N	115	
18	O	148	
19	P	95	
20	Q	154	
21	R	84	
22	S	119	
23	T	66	
24	U	70	
25	V	154	
26	W	91	
27	X	240	
28	Y	73	

Continued on next page...

Continued from previous page...

Mol	Chain	Length	Quality of chain
29	Z	56	
30	1	48	
31	2	92	

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

Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
34	NA	0	8329	-	-	-	X
34	NA	0	8384	-	-	-	X
34	NA	Q	8386	-	-	-	X
35	CL	I	8502	-	-	X	-

2 Entry composition

There are 37 unique types of molecules in this entry. The entry contains 98494 atoms, of which 0 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a RNA chain called 23S ribosomal rna.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
1	0	2754	Total	C	N	O	P	0	0	0
			59017	26346	10878	19048	2745			

- Molecule 2 is a RNA chain called 5S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
2	9	122	Total	C	N	O	P	0	0	0
			2600	1160	472	847	121			

- Molecule 3 is a RNA chain called Oligonucleotide CCA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
3	3	3	Total	C	N	O	P	0	0	0
			59	28	11	18	2			
3	4	3	Total	C	N	O	P	0	0	0
			59	28	11	18	2			
3	5	3	Total	C	N	O	P	0	0	0
			59	28	11	18	2			

- Molecule 4 is a protein called 50S ribosomal protein L2P.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
4	A	237	Total	C	N	O	S	0	0	0
			1754	1072	352	325	5			

- Molecule 5 is a protein called 50S ribosomal protein L3P.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
5	B	337	Total	C	N	O	S	0	0	0
			2624	1616	493	510	5			

There are 2 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
B	?	-	PRO	deletion	UNP P20279
B	310	ARG	PHE	conflict	UNP P20279

- Molecule 6 is a protein called 50S ribosomal protein L4E.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
6	C	246	Total	C	N	O	S	0	0	0
			1858	1131	344	382	1			

- Molecule 7 is a protein called 50S ribosomal protein L5P.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
7	D	140	Total	C	N	O	S	0	0	0
			1094	685	195	210	4			

- Molecule 8 is a protein called 50S ribosomal protein L6P.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
8	E	172	Total	C	N	O	S	0	0	0
			1357	840	224	289	4			

- Molecule 9 is a protein called 50S ribosomal protein L7Ae.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
9	F	119	Total	C	N	O	S	0	0	0
			885	552	141	191	1			

- Molecule 10 is a protein called 50S RIBOSOMAL PROTEIN L10E.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	G	29	Total	C	N	O	S	0	0	0
			240	149	39	51	1			

- Molecule 11 is a protein called L10 Ribosomal Protein.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
11	H	156	Total	C	N	O	S	0	0	0
			1215	766	233	212	4			

- Molecule 12 is a protein called 50S ribosomal protein L13P.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
12	I	142	Total	C	N	O	S	0	0	0
			1119	696	199	221	3			

- Molecule 13 is a protein called 50S ribosomal protein L14P.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
13	J	132	Total	C	N	O	S	0	0	0
			993	609	189	191	4			

- Molecule 14 is a protein called 50S ribosomal protein L15P.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
14	K	145	Total	C	N	O	S	0	0	0
			1114	668	222	224				

- Molecule 15 is a protein called L15 Ribosomal Protein.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
15	L	194	Total	C	N	O	S	0	0	0
			1605	988	346	266	5			

- Molecule 16 is a protein called 50S ribosomal protein L18P.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
16	M	186	Total	C	N	O	S	0	0	0
			1444	895	262	285	2			

- Molecule 17 is a protein called 50S ribosomal protein L18e.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
17	N	115	Total	C	N	O	S	0	0	0
			864	529	161	174				

- Molecule 18 is a protein called 50S ribosomal protein L19E.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
18	O	143	Total	C	N	O	S	0	0	0
			1133	680	230	223				

There is a discrepancy between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
O	71	LYS	TYR	conflict	UNP P14119

- Molecule 19 is a protein called 50S ribosomal protein L21e.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
19	P	95	Total	C	N	O	0	0	0
			734	450	141	143			

- Molecule 20 is a protein called 50S ribosomal protein L22P.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
20	Q	150	Total	C	N	O	S	0	0	0
			1149	713	209	223	4			

- Molecule 21 is a protein called 50S ribosomal protein L23P.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
21	R	81	Total	C	N	O	S	0	0	0
			641	389	111	138	3			

- Molecule 22 is a protein called 50S ribosomal protein L24P.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
22	S	119	Total	C	N	O	0	0	0
			949	568	180	201			

- Molecule 23 is a protein called 50S ribosomal protein L24E.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
23	T	53	Total	C	N	O	S	0	0	0
			410	244	75	86	5			

- Molecule 24 is a protein called 50S ribosomal protein L29P.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
24	U	65	Total	C	N	O	S	0	0	0
			499	304	94	100	1			

- Molecule 25 is a protein called 50S ribosomal protein L30P.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
25	V	154	Total	C	N	O	S	0	0	0
			1195	737	209	243	6			

- Molecule 26 is a protein called 50S ribosomal protein L31e.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	W	82	Total	C	N	O	S	0	0	0
			654	402	129	122	1			

- Molecule 27 is a protein called 50S ribosomal protein L32E.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
27	X	142	Total	C	N	O	S	0	0	0
			1130	686	228	216				

- Molecule 28 is a protein called L37Ae 50S ribosomal protein.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
28	Y	73	Total	C	N	O	S	0	0	0
			563	359	111	86	7			

- Molecule 29 is a protein called 50S ribosomal protein L37e.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
29	Z	56	Total	C	N	O	S	0	0	0
			430	258	86	82	4			

- Molecule 30 is a protein called 50S ribosomal protein L39e.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
30	1	46	Total	C	N	O	S	0	0	0
			393	238	86	68	1			

There is a discrepancy between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
1	?	-	ARG	deletion	UNP P22452

- Molecule 31 is a protein called 50S ribosomal protein L44E.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
31	2	92	Total	C	N	O	S	0	0	0
			755	458	153	137	7			

- Molecule 32 is MAGNESIUM ION (CCD ID: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
32	0	107	Total	Mg	0	0
			107	107		
32	9	1	Total	Mg	0	0
			1	1		
32	A	3	Total	Mg	0	0
			3	3		
32	B	1	Total	Mg	0	0
			1	1		
32	J	1	Total	Mg	0	0
			1	1		
32	S	1	Total	Mg	0	0
			1	1		
32	X	1	Total	Mg	0	0
			1	1		
32	2	1	Total	Mg	0	0
			1	1		

- Molecule 33 is POTASSIUM ION (CCD ID: K) (formula: K).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
33	0	2	Total	K	0	0
			2	2		

- Molecule 34 is SODIUM ION (CCD ID: NA) (formula: Na).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
34	0	73	Total	Na	0	0
			73	73		
34	9	2	Total	Na	0	0
			2	2		
34	A	1	Total	Na	0	0
			1	1		
34	C	1	Total	Na	0	0
			1	1		
34	H	1	Total	Na	0	0
			1	1		

Continued on next page...

Continued from previous page...

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
34	I	1	Total 1	Na 1	0	0
34	K	1	Total 1	Na 1	0	0
34	L	1	Total 1	Na 1	0	0
34	P	1	Total 1	Na 1	0	0
34	Q	3	Total 3	Na 3	0	0
34	R	1	Total 1	Na 1	0	0

- Molecule 35 is CHLORIDE ION (CCD ID: CL) (formula: Cl).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
35	0	9	Total 9	Cl 9	0	0
35	A	1	Total 1	Cl 1	0	0
35	B	1	Total 1	Cl 1	0	0
35	I	3	Total 3	Cl 3	0	0
35	J	1	Total 1	Cl 1	0	0
35	K	1	Total 1	Cl 1	0	0
35	L	1	Total 1	Cl 1	0	0
35	M	1	Total 1	Cl 1	0	0
35	N	1	Total 1	Cl 1	0	0
35	Q	1	Total 1	Cl 1	0	0
35	X	1	Total 1	Cl 1	0	0
35	2	1	Total 1	Cl 1	0	0

- Molecule 36 is CADMIUM ION (CCD ID: CD) (formula: Cd).

Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
36	N	1	Total Cd 1 1	0	0
36	T	1	Total Cd 1 1	0	0
36	Y	1	Total Cd 1 1	0	0
36	Z	1	Total Cd 1 1	0	0
36	2	1	Total Cd 1 1	0	0

- Molecule 37 is water.

Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
37	0	5766	Total O 5766 5766	0	0
37	9	148	Total O 148 148	0	0
37	4	1	Total O 1 1	0	0
37	5	2	Total O 2 2	0	0
37	A	115	Total O 115 115	0	0
37	B	146	Total O 146 146	0	0
37	C	166	Total O 166 166	0	0
37	D	48	Total O 48 48	0	0
37	E	43	Total O 43 43	0	0
37	F	25	Total O 25 25	0	0
37	G	20	Total O 20 20	0	0
37	H	77	Total O 77 77	0	0
37	I	56	Total O 56 56	0	0
37	J	56	Total O 56 56	0	0
37	K	80	Total O 80 80	0	0

Continued on next page...

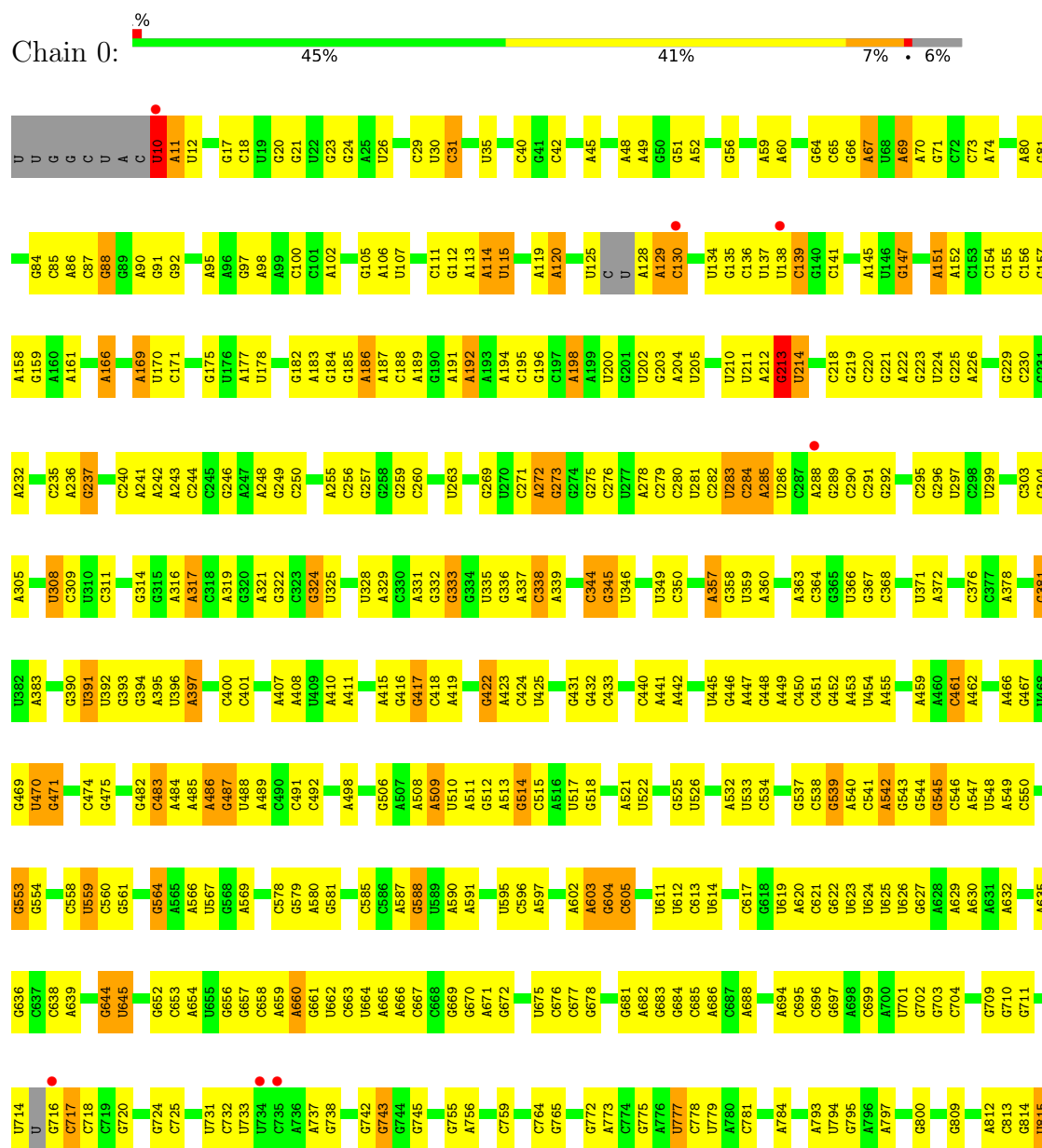
Continued from previous page...

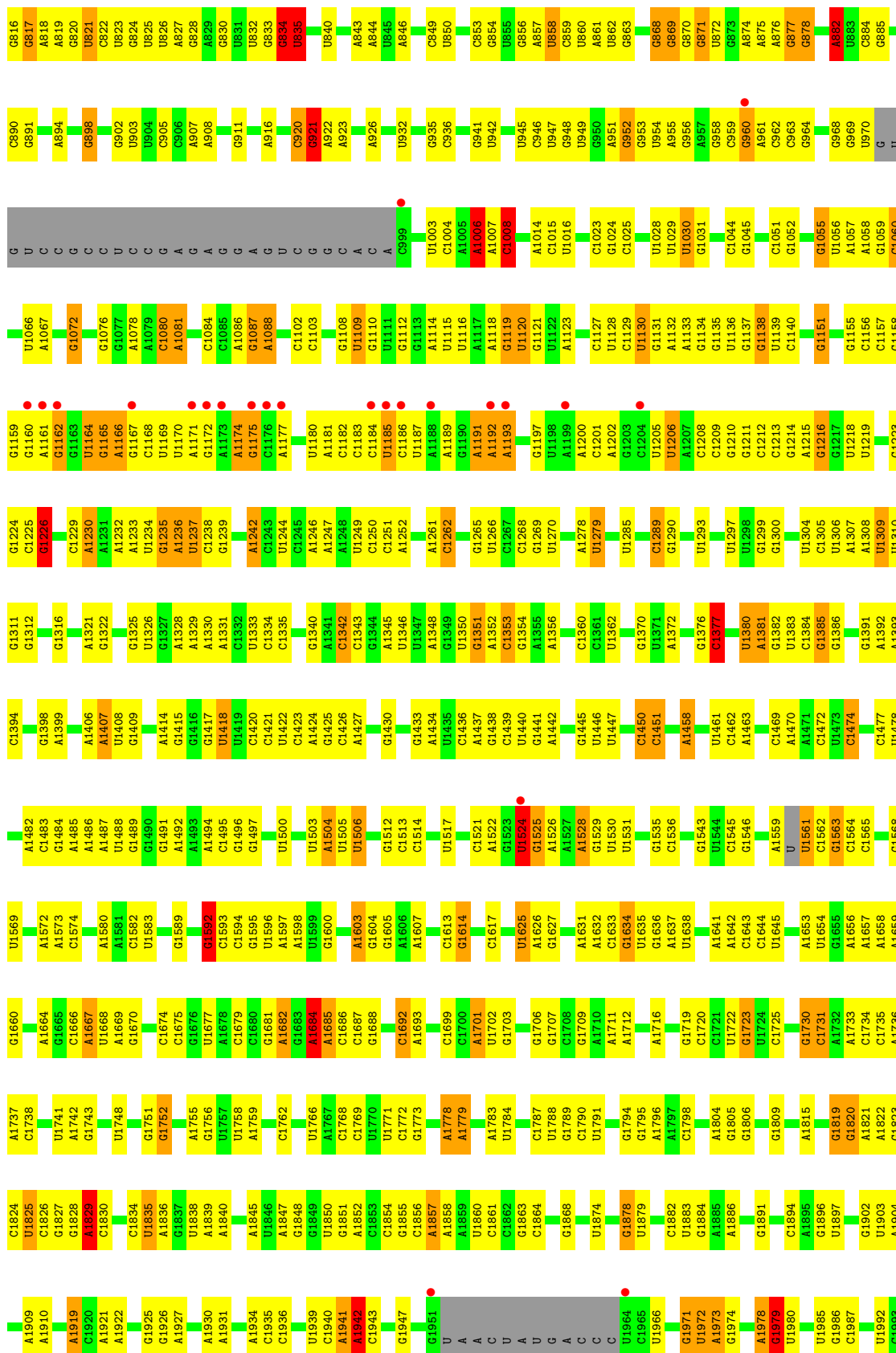
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
37	L	129	Total 129	O 129	0	0
37	M	56	Total 56	O 56	0	0
37	N	43	Total 43	O 43	0	0
37	O	58	Total 58	O 58	0	0
37	P	57	Total 57	O 57	0	0
37	Q	85	Total 85	O 85	0	0
37	R	31	Total 31	O 31	0	0
37	S	38	Total 38	O 38	0	0
37	T	30	Total 30	O 30	0	0
37	U	12	Total 12	O 12	0	0
37	V	69	Total 69	O 69	0	0
37	W	27	Total 27	O 27	0	0
37	X	97	Total 97	O 97	0	0
37	Y	35	Total 35	O 35	0	0
37	Z	54	Total 54	O 54	0	0
37	1	42	Total 42	O 42	0	0
37	2	56	Total 56	O 56	0	0

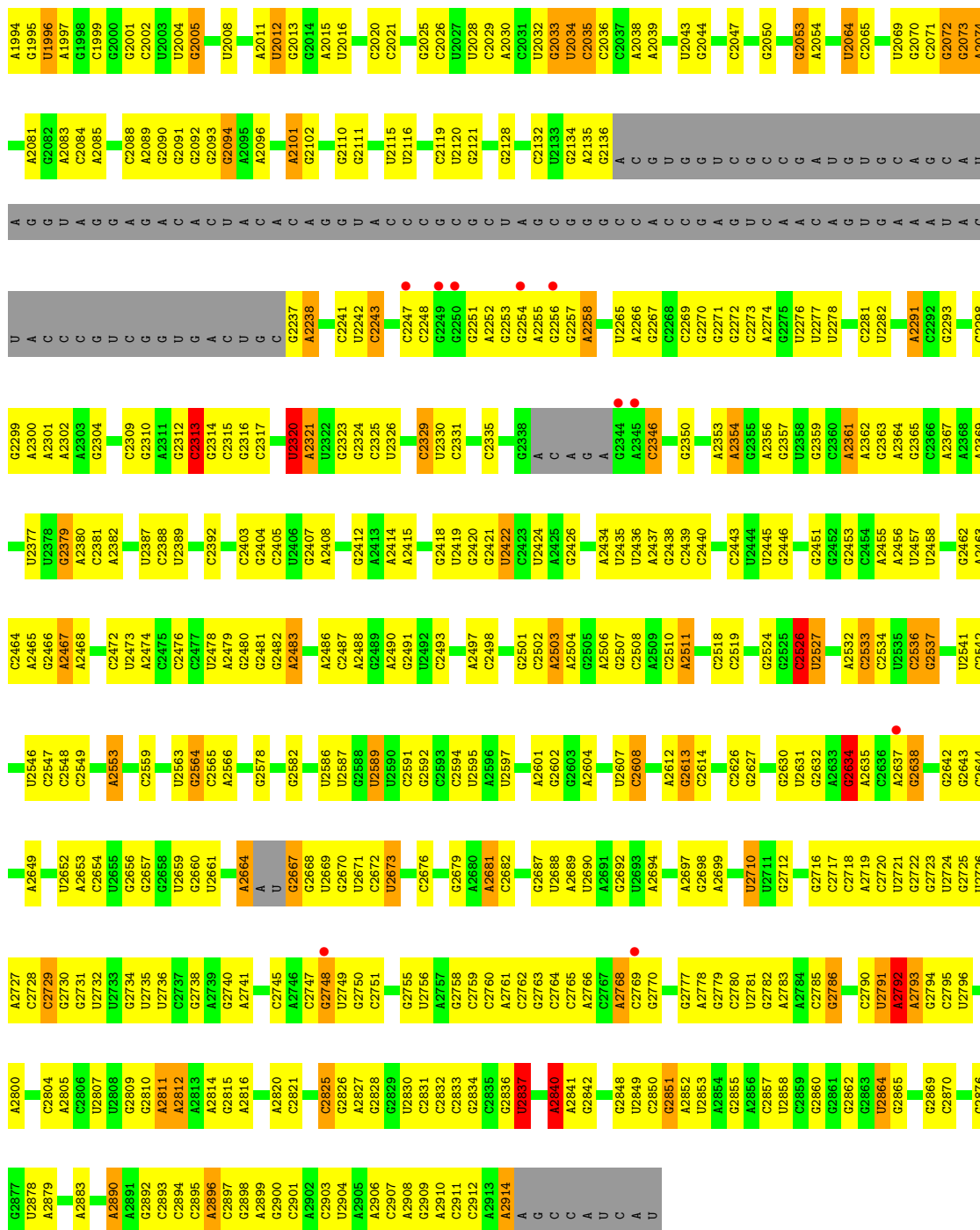
3 Residue-property plots

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 electron density. Residues are color-coded according to the number of geometric quality criteria for which they contain at least one outlier: green = 0, yellow = 1, orange = 2 and red = 3 or more. A red dot above a residue indicates a poor fit to the electron density ($RSRZ > 2$). Stretches of 2 or more consecutive residues without any outlier are shown as a green connector. Residues present in the sample, but not in the model, are shown in grey.

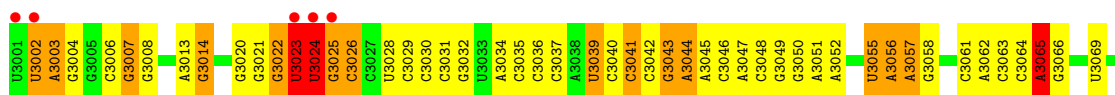
- Molecule 1: 23S ribosomal rna

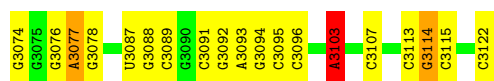






- Molecule 2: 5S ribosomal RNA





• Molecule 3: Oligonucleotide CCA



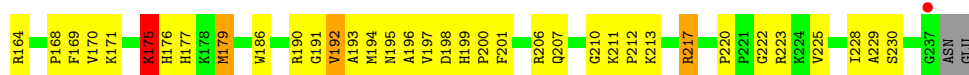
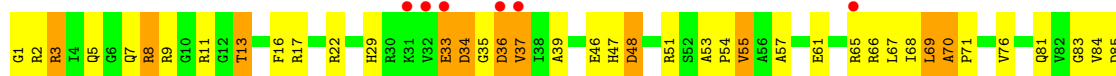
• Molecule 3: Oligonucleotide CCA



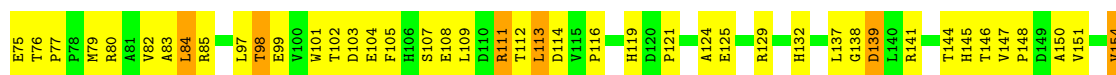
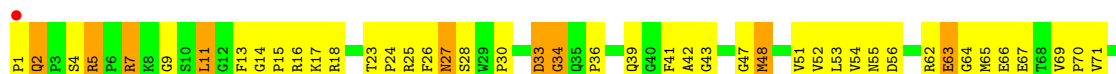
• Molecule 3: Oligonucleotide CCA



• Molecule 4: 50S ribosomal protein L2P

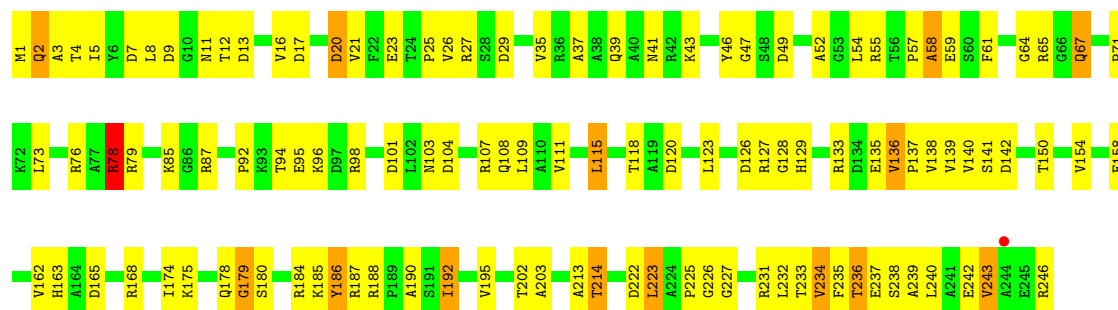


• Molecule 5: 50S ribosomal protein L3P

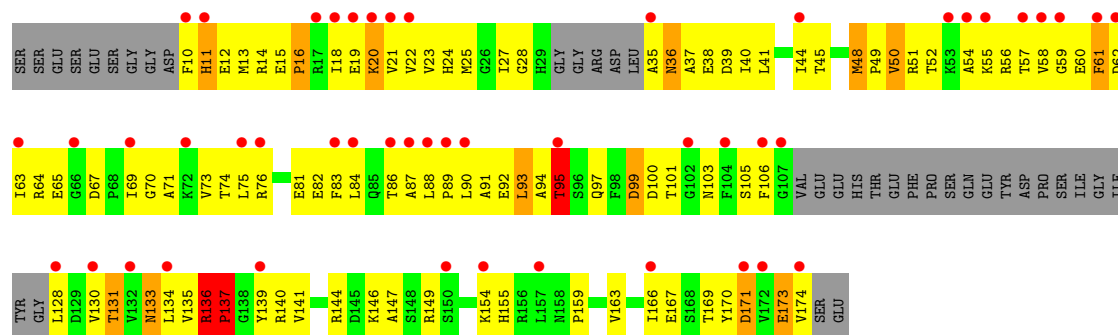




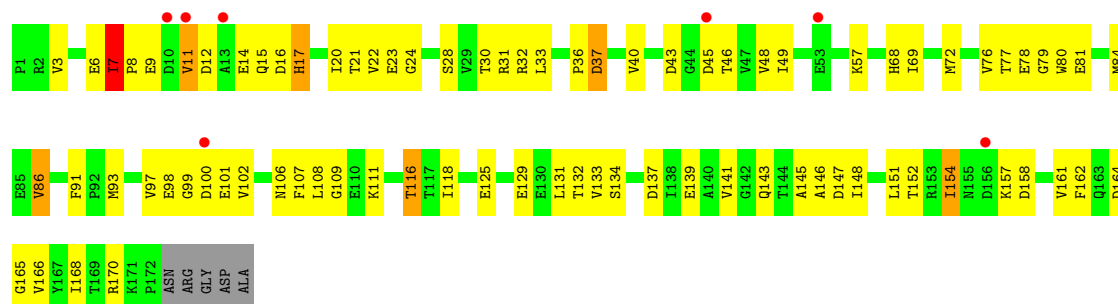
• Molecule 6: 50S ribosomal protein L4E



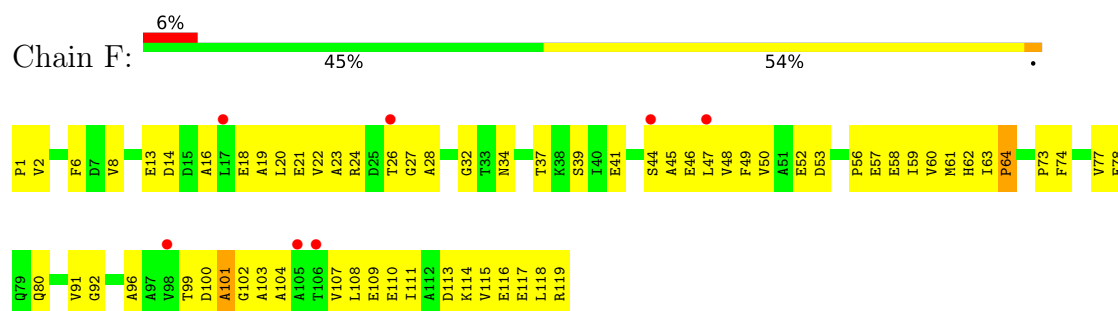
• Molecule 7: 50S ribosomal protein L5P



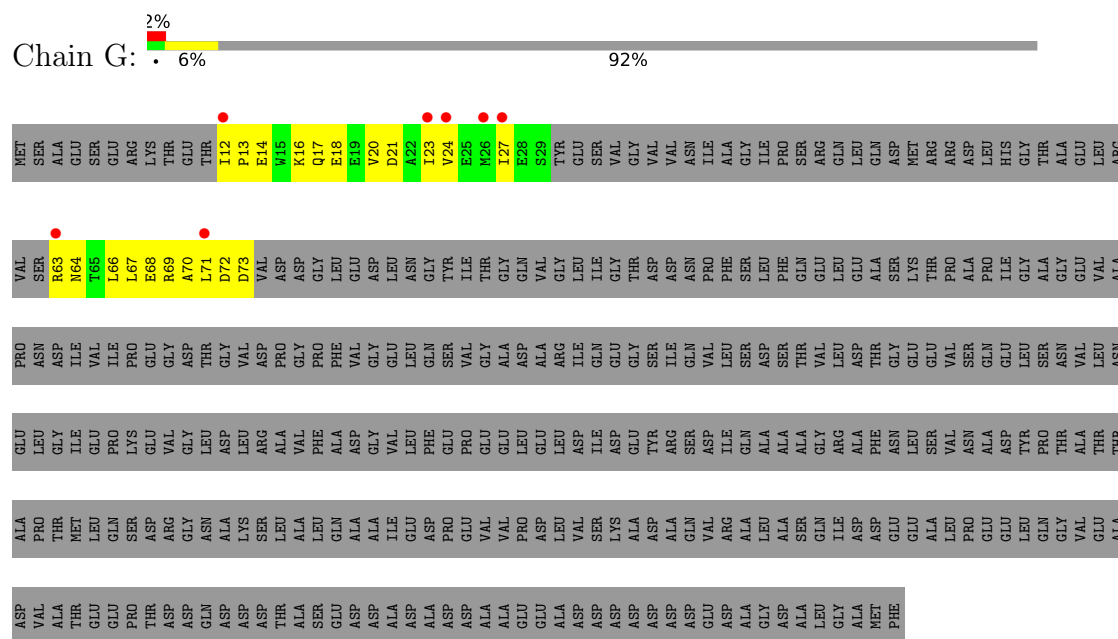
• Molecule 8: 50S ribosomal protein L6P



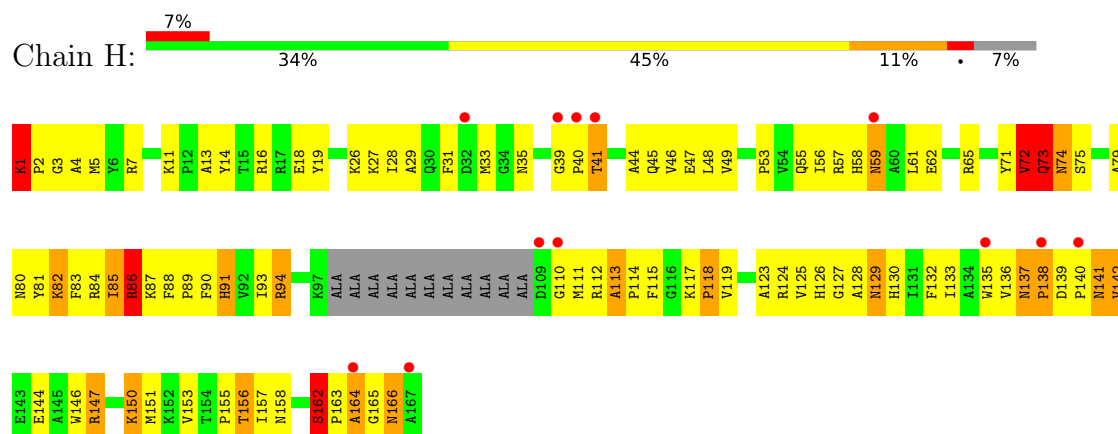
- Molecule 9: 50S ribosomal protein L7Ae



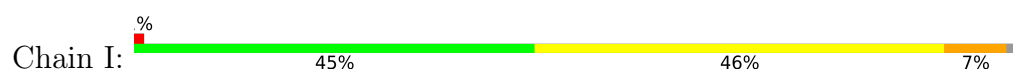
- Molecule 10: 50S RIBOSOMAL PROTEIN L10E

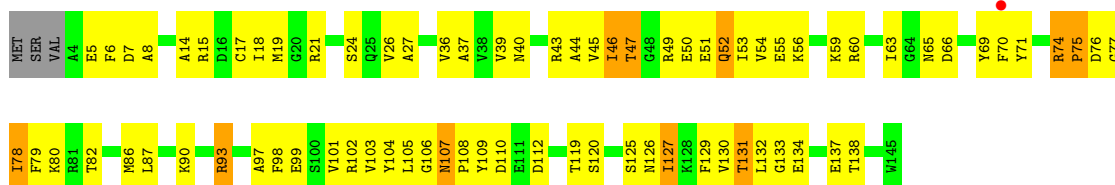


- Molecule 11: L10 Ribosomal Protein

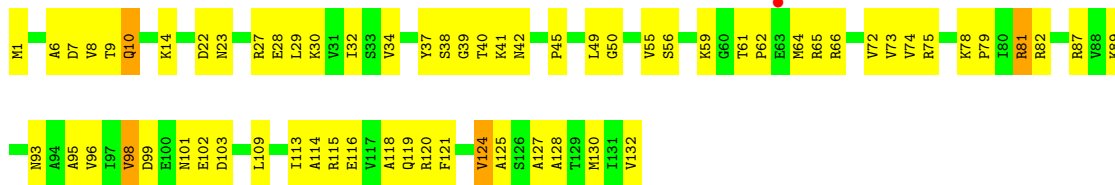


- Molecule 12: 50S ribosomal protein L13P

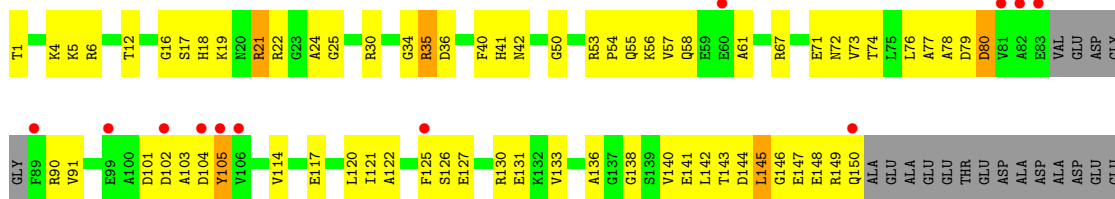




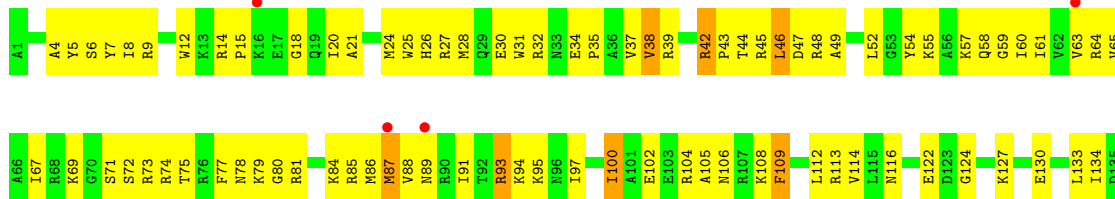
• Molecule 13: 50S ribosomal protein L14P



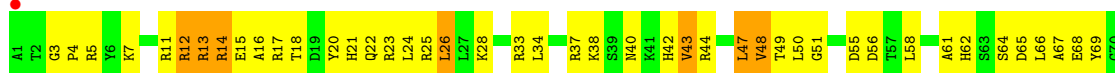
• Molecule 14: 50S ribosomal protein L15P

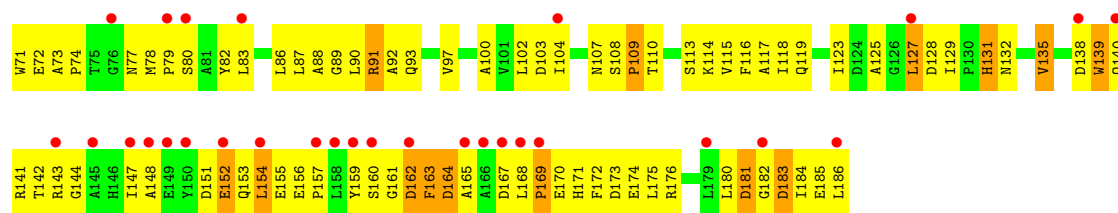


• Molecule 15: L15 Ribosomal Protein

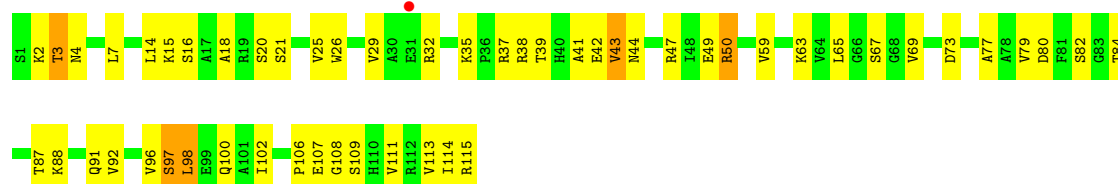


• Molecule 16: 50S ribosomal protein L18P

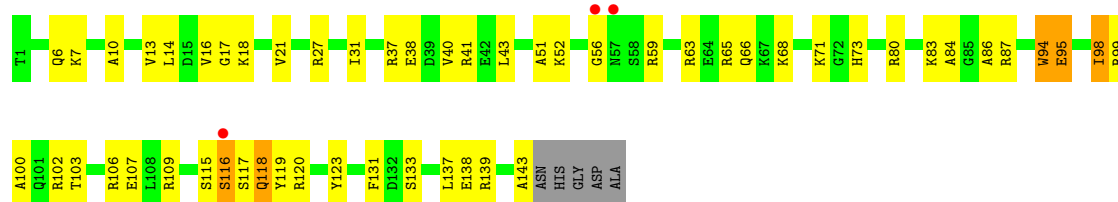




• Molecule 17: 50S ribosomal protein L18e



• Molecule 18: 50S ribosomal protein L19E



• Molecule 19: 50S ribosomal protein L21e

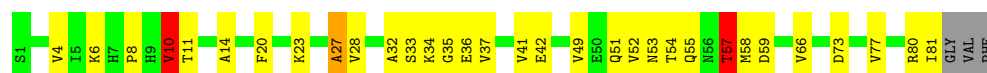


• Molecule 20: 50S ribosomal protein L22P



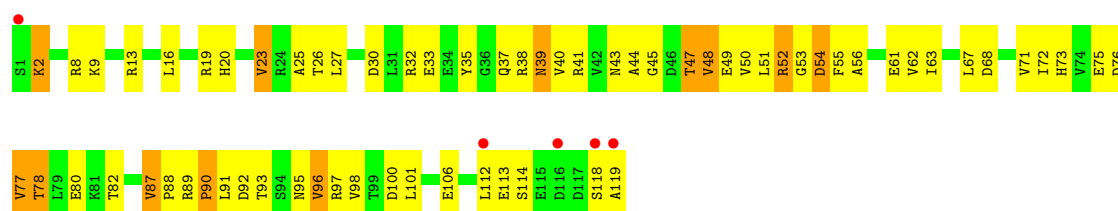
• Molecule 21: 50S ribosomal protein L23P

Chain R:  58% 35% ..



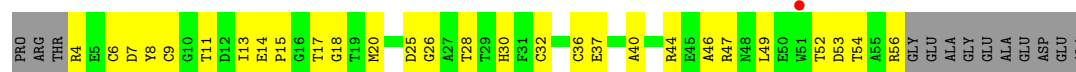
• Molecule 22: 50S ribosomal protein L24P

Chain S:  4% 45% 45% 10%



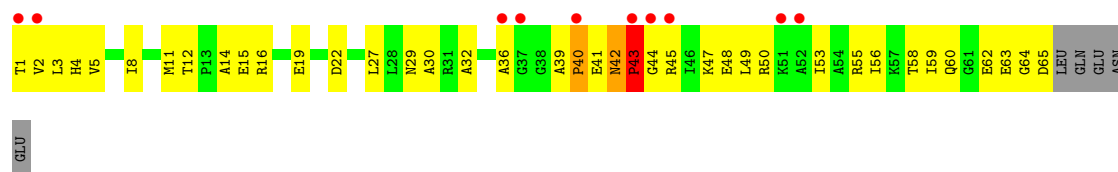
• Molecule 23: 50S ribosomal protein L24E

Chain T:  2% 38% 42% 20%

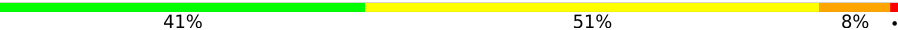


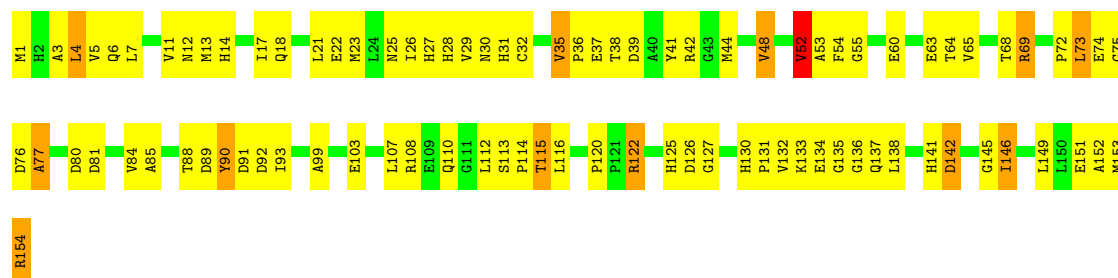
• Molecule 24: 50S ribosomal protein L29P

Chain U:  14% 37% 51% 7%



• Molecule 25: 50S ribosomal protein L30P

Chain V:  41% 51% 8%

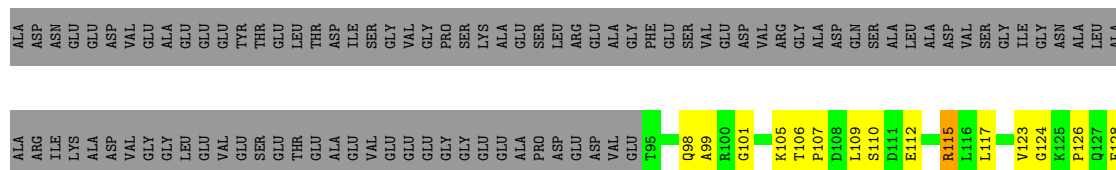
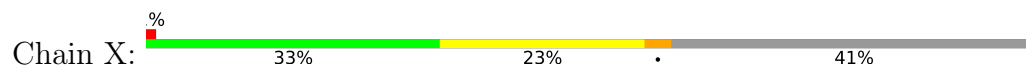


• Molecule 26: 50S ribosomal protein L31e

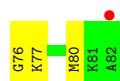
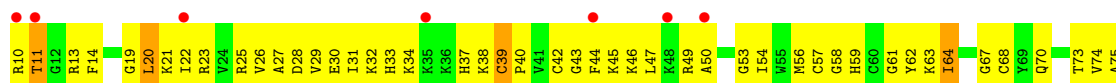
Chain W:  2% 38% 44% 7% 10%



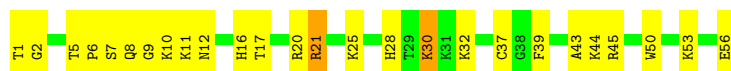
• Molecule 27: 50S ribosomal protein L32E



• Molecule 28: L37Ae 50S ribosomal protein



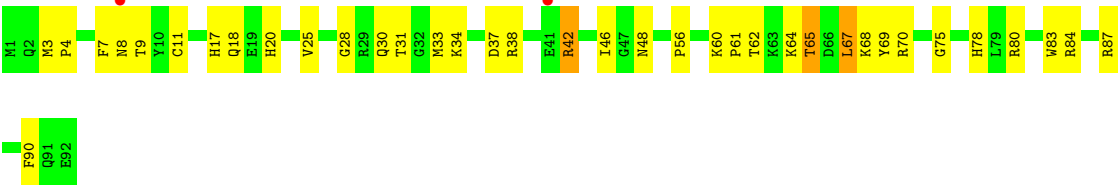
• Molecule 29: 50S ribosomal protein L37e



• Molecule 30: 50S ribosomal protein L39e



• Molecule 31: 50S ribosomal protein L44E



4 Data and refinement statistics

Property	Value	Source
Space group	C 2 2 21	Depositor
Cell constants a, b, c, α , β , γ	213.52Å 300.61Å 573.93Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	50.00 – 2.90 50.00 – 2.90	Depositor EDS
% Data completeness (in resolution range)	(Not available) (50.00-2.90) 89.6 (50.00-2.90)	Depositor EDS
R_{merge}	(Not available)	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.62 (at 2.81Å)	Xtriage
Refinement program	CNS	Depositor
R, R_{free}	0.201 , 0.258 0.199 , (Not available)	Depositor DCC
R_{free} test set	No test flags present.	wwPDB-VP
Wilson B-factor (Å ²)	51.8	Xtriage
Anisotropy	0.440	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.30 , 70.0	EDS
L-test for twinning ²	$\langle L \rangle = 0.46$, $\langle L^2 \rangle = 0.29$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
F_o, F_c correlation	0.93	EDS
Total number of atoms	98494	wwPDB-VP
Average B, all atoms (Å ²)	60.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.58% of the height of the origin peak. No significant pseudotranslation is detected.*

¹Intensities estimated from amplitudes.

²Theoretical values of $\langle |L| \rangle$, $\langle L^2 \rangle$ for acentric reflections are 0.5, 0.333 respectively for untwinned datasets, and 0.375, 0.2 for perfectly twinned datasets.

5 Model quality ⓘ

5.1 Standard geometry ⓘ

Bond lengths and bond angles in the following residue types are not validated in this section: NA, CD, MG, K, CL

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	0	0.55	0/66076	0.74	42/103052 (0.0%)
2	9	0.53	0/2905	0.76	7/4528 (0.2%)
3	3	0.54	0/65	0.84	0/99
3	4	0.73	0/65	0.77	0/99
3	5	0.47	0/65	0.79	0/99
4	A	0.68	0/1787	1.09	12/2409 (0.5%)
5	B	0.73	0/2689	1.13	24/3652 (0.7%)
6	C	0.80	0/1883	1.12	7/2551 (0.3%)
7	D	0.54	0/1111	0.98	5/1498 (0.3%)
8	E	0.72	0/1382	1.00	5/1880 (0.3%)
9	F	0.60	0/896	0.97	2/1219 (0.2%)
10	G	0.60	0/241	0.84	1/324 (0.3%)
11	H	0.80	1/1246 (0.1%)	1.40	22/1686 (1.3%)
12	I	0.79	0/1135	1.07	5/1530 (0.3%)
13	J	0.71	0/1003	1.12	4/1351 (0.3%)
14	K	0.62	0/1126	1.05	3/1504 (0.2%)
15	L	0.75	0/1633	1.13	10/2180 (0.5%)
16	M	0.59	0/1473	1.16	13/1999 (0.7%)
17	N	0.74	0/873	1.07	6/1181 (0.5%)
18	O	0.66	0/1143	0.98	3/1521 (0.2%)
19	P	0.74	0/748	1.18	6/1005 (0.6%)
20	Q	0.77	0/1172	1.07	5/1578 (0.3%)
21	R	0.57	0/648	0.91	3/875 (0.3%)
22	S	0.62	0/957	1.04	8/1289 (0.6%)
23	T	0.69	0/417	1.05	0/562
24	U	0.52	0/502	1.05	2/675 (0.3%)
25	V	0.83	0/1218	1.10	4/1655 (0.2%)
26	W	0.81	0/664	1.13	7/895 (0.8%)
27	X	0.74	0/1146	1.06	2/1536 (0.1%)
28	Y	0.66	0/575	1.04	1/763 (0.1%)
29	Z	0.70	0/437	1.04	3/578 (0.5%)
30	1	0.62	0/398	0.82	0/527

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
31	2	0.68	0/771	1.00	1/1024 (0.1%)
All	All	0.60	1/98450 (0.0%)	0.84	213/147324 (0.1%)

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

Mol	Chain	#Chirality outliers	#Planarity outliers
1	0	0	105
2	9	0	4
25	V	0	1
All	All	0	110

All (1) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
11	H	41	THR	CA-CB	5.49	1.60	1.53

All (213) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
11	H	74	ASN	N-CA-C	-15.73	91.57	114.39
1	0	1563	G	C2'-C3'-O3'	14.45	131.17	109.50
11	H	141	ASN	N-CA-C	-12.69	97.78	113.38
15	L	141	ILE	N-CA-C	-10.95	102.01	112.96
1	0	1979	G	C2'-C3'-O3'	10.69	125.53	109.50
11	H	156	THR	N-CA-C	-10.21	96.77	110.55
16	M	135	VAL	N-CA-C	-9.73	103.64	113.47
19	P	47	VAL	CA-C-N	9.06	128.40	118.97
19	P	47	VAL	C-N-CA	9.06	128.40	118.97
1	0	1524	U	C2'-C3'-O3'	9.04	123.06	109.50
8	E	11	VAL	N-CA-C	8.80	122.84	109.20
7	D	136	ARG	CA-C-N	8.63	130.63	119.84
7	D	136	ARG	C-N-CA	8.63	130.63	119.84
9	F	8	VAL	N-CA-C	8.13	114.96	107.56
1	0	213	G	C2'-C3'-O3'	8.05	121.58	109.50
19	P	17	LYS	N-CA-C	-7.94	99.74	109.83
11	H	1	LYS	CA-C-N	7.77	128.38	120.14
11	H	1	LYS	C-N-CA	7.77	128.38	120.14
7	D	100	ASP	N-CA-C	-7.74	103.86	113.38
16	M	13	ARG	N-CA-C	-7.43	104.34	113.41

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
25	V	69	ARG	N-CA-C	7.33	121.93	112.41
5	B	265	LEU	N-CA-C	7.31	121.30	110.48
5	B	157	LYS	N-CA-C	-7.31	104.97	114.04
11	H	137	ASN	N-CA-C	-7.30	100.58	110.36
4	A	222	GLY	N-CA-C	-7.22	105.86	113.58
6	C	64	GLY	N-CA-C	7.17	122.17	113.79
11	H	147	ARG	N-CA-C	-7.16	103.80	112.54
29	Z	21	ARG	N-CA-C	7.16	120.14	111.40
2	9	3024	U	C2'-C3'-O3'	7.06	120.09	109.50
29	Z	43	ALA	N-CA-C	-7.01	103.72	111.36
6	C	179	GLY	N-CA-C	-6.97	104.32	112.83
16	M	153	GLN	N-CA-C	-6.93	103.30	112.72
5	B	217	ARG	N-CA-C	6.91	118.81	111.28
6	C	175	LYS	N-CA-C	6.90	119.74	110.35
21	R	57	THR	N-CA-C	6.87	120.65	110.48
14	K	145	LEU	N-CA-C	-6.86	103.89	111.36
16	M	51	GLY	CA-C-N	6.78	127.33	119.47
16	M	51	GLY	C-N-CA	6.78	127.33	119.47
5	B	323	LEU	N-CA-C	6.77	119.04	110.24
16	M	169	PRO	N-CA-C	-6.71	105.77	114.03
6	C	78	ARG	N-CA-C	6.64	118.86	108.96
5	B	23	THR	CA-C-N	6.62	126.60	119.78
5	B	23	THR	C-N-CA	6.62	126.60	119.78
5	B	154	VAL	CA-C-N	6.58	126.09	119.24
5	B	154	VAL	C-N-CA	6.58	126.09	119.24
1	0	1942	A	C5'-C4'-C3'	6.58	125.87	116.00
22	S	52	ARG	N-CA-C	6.57	121.92	113.18
16	M	102	LEU	N-CA-C	6.56	118.84	108.67
5	B	222	LYS	N-CA-C	-6.55	100.27	110.42
24	U	42	ASN	CA-C-N	6.54	128.01	119.84
24	U	42	ASN	C-N-CA	6.54	128.01	119.84
2	9	3039	U	C4'-C3'-O3'	-6.51	103.24	113.00
15	L	139	PRO	N-CA-C	-6.49	102.05	111.33
20	Q	137	ASN	N-CA-C	6.47	119.97	110.59
1	0	2466	G	C4'-C3'-O3'	-6.46	103.31	113.00
1	0	1120	U	C5'-C4'-C3'	-6.46	106.31	116.00
19	P	68	GLY	N-CA-C	-6.42	97.97	113.18
17	N	43	VAL	N-CA-C	6.34	117.05	108.17
1	0	2291	A	N9-C1'-C2'	6.34	121.50	112.00
5	B	47	GLY	N-CA-C	6.34	118.69	110.45
5	B	80	ARG	N-CA-C	6.33	118.76	107.80
13	J	8	VAL	N-CA-C	6.33	117.70	108.53

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	0	874	A	C4'-C3'-O3'	-6.32	103.52	113.00
20	Q	13	THR	N-CA-C	6.31	119.79	109.06
17	N	50	ARG	N-CA-C	6.30	119.13	111.82
1	0	920	C	C2'-C3'-O3'	-6.28	104.28	113.70
20	Q	34	GLU	N-CA-C	6.28	118.13	111.28
4	A	103	VAL	N-CA-C	6.27	115.45	108.05
22	S	47	THR	N-CA-C	-6.27	99.98	109.95
1	0	1563	G	C4'-C3'-O3'	6.26	118.79	109.40
17	N	69	VAL	N-CA-C	6.23	117.45	108.48
11	H	86	ARG	N-CA-C	6.22	120.60	112.89
18	O	118	GLN	N-CA-C	-6.21	104.43	111.14
12	I	110	ASP	N-CA-C	-6.19	104.44	112.68
20	Q	122	GLN	N-CA-C	-6.14	99.16	108.67
26	W	79	GLU	N-CA-C	6.13	120.60	113.18
7	D	48	MET	N-CA-C	6.13	116.65	109.60
6	C	186	TYR	N-CA-C	6.04	119.01	110.14
8	E	7	ILE	CA-C-N	6.02	126.04	119.90
8	E	7	ILE	C-N-CA	6.02	126.04	119.90
15	L	89	ASN	N-CA-C	6.02	117.53	110.97
4	A	122	SER	N-CA-C	6.01	119.54	110.64
1	0	1261	A	N9-C1'-C2'	6.01	121.01	112.00
4	A	198	ASP	N-CA-C	6.00	120.67	113.23
26	W	68	SER	N-CA-C	5.99	119.69	112.38
15	L	109	PHE	CA-C-N	5.95	127.28	119.84
15	L	109	PHE	C-N-CA	5.95	127.28	119.84
11	H	113	ALA	CA-C-N	5.92	128.05	120.89
11	H	113	ALA	C-N-CA	5.92	128.05	120.89
29	Z	30	LYS	N-CA-C	-5.91	105.73	113.12
12	I	112	ASP	N-CA-C	-5.89	100.39	109.76
22	S	2	LYS	N-CA-C	-5.89	106.14	113.38
17	N	2	LYS	N-CA-C	-5.88	101.78	110.48
13	J	124	VAL	N-CA-C	-5.88	105.12	110.82
1	0	921	G	N9-C1'-C2'	5.87	120.81	112.00
1	0	2526	C	N1-C1'-C2'	5.84	120.76	112.00
6	C	192	ILE	N-CA-C	5.82	117.12	109.21
27	X	178	HIS	N-CA-C	-5.82	101.83	110.20
15	L	100	ILE	N-CA-C	-5.81	105.07	110.53
11	H	162	SER	CA-C-N	-5.80	112.59	119.84
11	H	162	SER	C-N-CA	-5.80	112.59	119.84
5	B	151	VAL	CA-C-N	5.78	125.25	119.24
5	B	151	VAL	C-N-CA	5.78	125.25	119.24
20	Q	90	ASP	N-CA-C	-5.78	104.42	112.45

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	0	1030	U	C5'-C4'-O4'	5.73	117.69	109.10
26	W	51	ASP	CA-C-N	5.69	126.95	119.84
26	W	51	ASP	C-N-CA	5.69	126.95	119.84
16	M	14	ARG	N-CA-C	-5.68	104.77	110.97
16	M	168	LEU	N-CA-C	5.68	121.48	108.81
2	9	3039	U	N1-C1'-C2'	5.67	120.50	112.00
12	I	75	PRO	N-CA-C	-5.65	107.08	114.03
15	L	142	LYS	N-CA-C	-5.65	105.27	111.82
1	0	2664	A	N9-C1'-C2'	5.64	122.47	114.00
2	9	3065	A	N9-C1'-C2'	5.62	120.43	112.00
15	L	100	ILE	CB-CA-C	-5.62	104.67	112.02
28	Y	39	CYS	N-CA-C	5.61	116.77	109.64
25	V	115	THR	N-CA-C	5.59	118.02	108.90
1	0	834	G	C2'-C3'-O3'	-5.58	105.33	113.70
5	B	48	MET	N-CA-C	5.58	118.31	109.50
11	H	7	ARG	N-CA-C	5.56	119.68	113.01
16	M	117	ALA	N-CA-C	-5.55	105.13	111.07
17	N	59	VAL	N-CA-C	5.55	115.58	107.37
4	A	135	VAL	N-CA-C	5.55	116.47	108.48
4	A	228	ILE	N-CA-C	5.54	116.16	108.84
16	M	91	ARG	N-CA-C	-5.54	105.34	111.71
7	D	137	PRO	N-CA-C	5.53	123.87	112.47
11	H	155	PRO	N-CA-C	5.53	118.97	111.22
25	V	25	ASN	N-CA-C	5.51	119.58	112.86
14	K	91	VAL	N-CA-C	5.50	116.03	108.12
15	L	42	ARG	CA-C-N	5.49	125.41	119.76
15	L	42	ARG	C-N-CA	5.49	125.41	119.76
1	0	1996	U	C3'-C2'-O2'	-5.47	106.39	114.60
19	P	57	ASP	N-CA-C	-5.47	101.75	109.96
1	0	1504	A	N9-C1'-C2'	5.46	120.18	112.00
1	0	2537	G	N9-C1'-C2'	5.43	120.15	112.00
1	0	1941	A	C2'-C3'-O3'	5.43	117.64	109.50
4	A	175	LYS	N-CA-C	-5.43	105.44	111.36
4	A	70	ALA	N-CA-C	5.42	116.78	109.24
1	0	1723	G	C4'-C3'-O3'	-5.41	104.89	113.00
1	0	129	A	C2'-C3'-O3'	5.39	121.78	113.70
9	F	62	HIS	N-CA-C	5.39	118.07	111.82
14	K	42	ASN	N-CA-C	5.38	119.01	111.74
25	V	52	VAL	N-CA-CB	-5.38	106.42	112.45
22	S	54	ASP	N-CA-C	-5.37	106.56	113.01
5	B	242	TRP	N-CA-C	-5.37	105.32	111.07
18	O	56	GLY	N-CA-C	-5.37	102.63	111.38

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
5	B	279	THR	N-CA-C	5.36	117.21	110.24
18	O	95	GLU	N-CA-C	-5.36	105.52	111.36
26	W	10	VAL	N-CA-C	5.35	116.44	108.46
13	J	81	ARG	N-CA-C	5.35	117.24	108.52
11	H	129	ASN	CA-C-N	-5.35	115.34	122.72
11	H	129	ASN	C-N-CA	-5.35	115.34	122.72
16	M	131	HIS	N-CA-C	5.34	115.33	108.34
11	H	91	HIS	N-CA-C	5.32	116.81	109.15
13	J	38	SER	N-CA-C	-5.32	100.36	109.24
2	9	3103	A	C4'-C3'-C2'	-5.32	97.28	102.60
17	N	82	SER	N-CA-C	-5.32	103.51	110.53
6	C	20	ASP	N-CA-C	5.31	117.07	111.28
1	0	835	U	C4'-C3'-O3'	-5.31	105.03	113.00
1	0	1356	A	C3'-C2'-O2'	-5.31	106.64	114.60
11	H	115	PHE	N-CA-C	-5.30	102.07	109.96
19	P	84	ILE	N-CA-C	5.29	115.52	108.11
1	0	1738	C	C5'-C4'-C3'	5.29	123.93	116.00
31	2	67	LEU	N-CA-C	5.26	117.86	109.81
22	S	78	THR	N-CA-C	5.25	117.39	109.41
1	0	2291	A	C4'-C3'-O3'	-5.25	105.13	113.00
5	B	124	ALA	N-CA-C	-5.24	105.26	110.97
2	9	3065	A	C4'-C3'-O3'	-5.24	105.15	113.00
5	B	282	GLY	N-CA-C	-5.23	108.47	114.69
1	0	2667	G	O5'-C5'-C4'	5.23	119.34	111.50
12	I	107	ASN	CA-C-N	5.23	126.16	120.83
12	I	107	ASN	C-N-CA	5.23	126.16	120.83
22	S	77	VAL	CB-CA-C	-5.22	107.37	113.22
1	0	1693	A	C3'-C2'-O2'	-5.22	106.77	114.60
1	0	2792	A	C3'-C2'-O2'	5.22	118.53	110.70
27	X	143	TRP	N-CA-C	5.21	118.05	109.76
26	W	80	GLU	N-CA-C	-5.20	106.89	114.12
11	H	73	GLN	N-CA-C	-5.19	104.11	110.65
1	0	10	U	O5'-C5'-C4'	5.17	119.25	111.50
5	B	114	ASP	N-CA-C	-5.15	98.92	108.24
1	0	1504	A	C4'-C3'-O3'	-5.13	105.30	113.00
1	0	1592	G	N9-C1'-C2'	5.13	119.70	112.00
1	0	1723	G	N9-C1'-C2'	5.13	119.70	112.00
21	R	27	ALA	N-CA-C	-5.13	99.42	108.20
4	A	8	ARG	N-CA-C	-5.13	105.77	111.36
5	B	175	LEU	N-CA-C	-5.12	105.59	111.07
8	E	37	ASP	N-CA-C	5.11	118.78	112.24
22	S	87	VAL	CA-C-N	5.11	125.02	119.76

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	S	87	VAL	C-N-CA	5.11	125.02	119.76
1	0	2536	C	O4'-C4'-C3'	-5.11	100.99	106.10
2	9	3065	A	C2'-C3'-O3'	5.10	121.36	113.70
5	B	154	VAL	N-CA-C	5.09	112.19	107.56
4	A	133	ARG	N-CA-C	-5.09	106.92	113.23
1	0	1006	A	N9-C1'-C2'	5.08	119.62	112.00
11	H	88	PHE	CA-C-N	5.07	126.18	119.84
11	H	88	PHE	C-N-CA	5.07	126.18	119.84
8	E	48	VAL	N-CA-C	5.07	115.21	108.11
1	0	1941	A	C4'-C3'-O3'	-5.07	101.80	109.40
1	0	2467	A	C1'-O4'-C4'	-5.06	104.64	109.70
4	A	48	ASP	CA-C-N	5.06	124.84	119.28
4	A	48	ASP	C-N-CA	5.06	124.84	119.28
1	0	2313	C	C4'-C3'-C2'	-5.05	97.55	102.60
5	B	111	ARG	N-CA-C	-5.05	107.07	113.18
21	R	10	VAL	N-CA-C	5.05	115.04	107.51
11	H	13	ALA	N-CA-C	-5.04	102.39	109.96
1	0	1504	A	C1'-O4'-C4'	-5.04	104.86	109.90
5	B	324	ASP	CA-C-N	5.04	124.80	119.76
5	B	324	ASP	C-N-CA	5.04	124.80	119.76
1	0	1561	U	O5'-C5'-C4'	5.03	119.05	111.50
1	0	1087	G	C3'-C2'-O2'	-5.03	107.06	114.60
26	W	29	ALA	N-CA-C	-5.02	105.49	110.97
10	G	14	GLU	N-CA-C	5.01	116.83	111.36
16	M	48	VAL	N-CA-C	5.01	116.17	108.71

There are no chirality outliers.

All (110) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	0	1008	C	Sidechain
1	0	1055	G	Sidechain
1	0	1078	A	Sidechain
1	0	1108	G	Sidechain
1	0	1191	A	Sidechain
1	0	1226	G	Sidechain
1	0	1230	A	Sidechain
1	0	1262	C	Sidechain
1	0	1300	G	Sidechain
1	0	1304	U	Sidechain
1	0	1309	U	Sidechain
1	0	1316	G	Sidechain

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
1	0	1340	G	Sidechain
1	0	1350	U	Sidechain
1	0	1351	G	Sidechain
1	0	1376	G	Sidechain
1	0	1377	C	Sidechain
1	0	1385	G	Sidechain
1	0	1417	G	Sidechain
1	0	1418	U	Sidechain
1	0	1430	G	Sidechain
1	0	1458	A	Sidechain
1	0	147	G	Sidechain
1	0	1487	A	Sidechain
1	0	1614	G	Sidechain
1	0	1684	A	Sidechain
1	0	171	C	Sidechain
1	0	1720	C	Sidechain
1	0	1758	U	Sidechain
1	0	1809	G	Sidechain
1	0	1825	U	Sidechain
1	0	1829	A	Sidechain
1	0	1835	U	Sidechain
1	0	1845	A	Sidechain
1	0	1848	G	Sidechain
1	0	1863	G	Sidechain
1	0	1878	G	Sidechain
1	0	1971	G	Sidechain
1	0	1972	U	Sidechain
1	0	1978	A	Sidechain
1	0	2034	U	Sidechain
1	0	2035	C	Sidechain
1	0	2053	G	Sidechain
1	0	2094	G	Sidechain
1	0	2101	A	Sidechain
1	0	213	G	Sidechain
1	0	2273	C	Sidechain
1	0	2312	G	Sidechain
1	0	2313	C	Sidechain
1	0	2320	U	Sidechain
1	0	246	G	Sidechain
1	0	2493	C	Sidechain
1	0	2526	C	Sidechain
1	0	2564	G	Sidechain

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
1	0	2597	U	Sidechain
1	0	26	U	Sidechain
1	0	2631	U	Sidechain
1	0	2632	G	Sidechain
1	0	2634	G	Sidechain
1	0	2673	U	Sidechain
1	0	2692	G	Sidechain
1	0	2710	U	Sidechain
1	0	2729	C	Sidechain
1	0	2793	A	Sidechain
1	0	2811	A	Sidechain
1	0	2837	U	Sidechain
1	0	2840	A	Sidechain
1	0	2842	G	Sidechain
1	0	2849	U	Sidechain
1	0	2853	U	Sidechain
1	0	2855	G	Sidechain
1	0	2864	U	Sidechain
1	0	324	G	Sidechain
1	0	333	G	Sidechain
1	0	344	C	Sidechain
1	0	391	U	Sidechain
1	0	395	A	Sidechain
1	0	397	A	Sidechain
1	0	422	G	Sidechain
1	0	469	G	Sidechain
1	0	470	U	Sidechain
1	0	471	G	Sidechain
1	0	48	A	Sidechain
1	0	483	C	Sidechain
1	0	486	A	Sidechain
1	0	49	A	Sidechain
1	0	517	U	Sidechain
1	0	518	G	Sidechain
1	0	548	U	Sidechain
1	0	554	G	Sidechain
1	0	564	G	Sidechain
1	0	619	U	Sidechain
1	0	662	U	Sidechain
1	0	742	G	Sidechain
1	0	743	G	Sidechain
1	0	781	C	Sidechain

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
1	0	795	G	Sidechain
1	0	815	U	Sidechain
1	0	817	G	Sidechain
1	0	868	G	Sidechain
1	0	882	A	Sidechain
1	0	898	G	Sidechain
1	0	903	U	Sidechain
1	0	916	A	Sidechain
1	0	952	G	Sidechain
2	9	3023	U	Sidechain
2	9	3065	A	Sidechain
2	9	3087	U	Sidechain
2	9	3094	G	Sidechain
25	V	90	TYR	Sidechain

5.2 Too-close contacts

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	0	59017	0	29802	1343	0
2	9	2600	0	1326	87	0
3	3	59	0	35	1	0
3	4	59	0	35	3	0
3	5	59	0	35	0	0
4	A	1754	0	1763	128	0
5	B	2624	0	2533	215	0
6	C	1858	0	1816	136	0
7	D	1094	0	1085	149	0
8	E	1357	0	1266	90	0
9	F	885	0	854	73	0
10	G	240	0	231	30	0
11	H	1215	0	1215	178	0
12	I	1119	0	1098	86	0
13	J	993	0	1027	80	0
14	K	1114	0	1072	73	0
15	L	1605	0	1676	176	0
16	M	1444	0	1401	166	0
17	N	864	0	873	47	0

Continued on next page...

Continued from previous page...

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
18	O	1133	0	1127	64	0
19	P	734	0	728	38	0
20	Q	1149	0	1122	81	0
21	R	641	0	605	33	0
22	S	949	0	923	80	0
23	T	410	0	364	41	0
24	U	499	0	511	39	0
25	V	1195	0	1137	118	0
26	W	654	0	653	55	0
27	X	1130	0	1133	80	0
28	Y	563	0	597	58	0
29	Z	430	0	426	35	0
30	1	393	0	406	44	0
31	2	755	0	728	37	0
32	0	107	0	0	0	0
32	2	1	0	0	0	0
32	9	1	0	0	0	0
32	A	3	0	0	0	0
32	B	1	0	0	0	0
32	J	1	0	0	0	0
32	S	1	0	0	0	0
32	X	1	0	0	0	0
33	0	2	0	0	0	0
34	0	73	0	0	0	0
34	9	2	0	0	0	0
34	A	1	0	0	0	0
34	C	1	0	0	0	0
34	H	1	0	0	0	0
34	I	1	0	0	0	0
34	K	1	0	0	0	0
34	L	1	0	0	0	0
34	P	1	0	0	0	0
34	Q	3	0	0	0	0
34	R	1	0	0	0	0
35	0	9	0	0	1	0
35	2	1	0	0	0	0
35	A	1	0	0	0	0
35	B	1	0	0	0	0
35	I	3	0	0	2	0
35	J	1	0	0	1	0
35	K	1	0	0	0	0
35	L	1	0	0	1	0

Continued on next page...

Continued from previous page...

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
35	M	1	0	0	1	0
35	N	1	0	0	0	0
35	Q	1	0	0	0	0
35	X	1	0	0	0	0
36	2	1	0	0	0	0
36	N	1	0	0	0	0
36	T	1	0	0	0	0
36	Y	1	0	0	0	0
36	Z	1	0	0	0	0
37	0	5766	0	0	297	0
37	1	42	0	0	4	0
37	2	56	0	0	10	0
37	4	1	0	0	0	0
37	5	2	0	0	0	0
37	9	148	0	0	15	0
37	A	115	0	0	23	0
37	B	146	0	0	27	0
37	C	166	0	0	38	0
37	D	48	0	0	24	0
37	E	43	0	0	15	0
37	F	25	0	0	10	0
37	G	20	0	0	7	0
37	H	77	0	0	28	0
37	I	56	0	0	11	0
37	J	56	0	0	12	0
37	K	80	0	0	18	0
37	L	129	0	0	22	0
37	M	56	0	0	27	0
37	N	43	0	0	13	0
37	O	58	0	0	5	0
37	P	57	0	0	8	0
37	Q	85	0	0	11	0
37	R	31	0	0	9	0
37	S	38	0	0	7	0
37	T	30	0	0	9	0
37	U	12	0	0	2	0
37	V	69	0	0	14	0
37	W	27	0	0	7	0
37	X	97	0	0	21	0
37	Y	35	0	0	7	0
37	Z	54	0	0	6	0
All	All	98494	0	59603	3529	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 24.

All (3529) 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)
11:H:86:ARG:NH1	11:H:133:ILE:HG13	1.55	1.20
20:Q:99:ALA:HB1	20:Q:109:MET:HE1	1.21	1.14
7:D:25:MET:HE2	7:D:41:LEU:HG	1.29	1.14
24:U:12:THR:HG22	24:U:15:GLU:HG3	1.32	1.11
12:I:19:MET:HE3	12:I:132:LEU:HD11	1.29	1.07
6:C:236:THR:HG22	6:C:239:ALA:H	1.00	1.07
6:C:127:ARG:NH2	6:C:225:PRO:HG2	1.70	1.06
11:H:45:GLN:HB3	11:H:163:PRO:HD2	1.31	1.06
11:H:86:ARG:HH11	11:H:133:ILE:CG1	1.68	1.06
6:C:115:LEU:HD13	6:C:223:LEU:HD21	1.36	1.05
1:O:1134:G:H4'	11:H:151:MET:HE1	1.39	1.05
1:O:1119:G:H2'	12:I:52:GLN:HE22	1.22	1.04
7:D:105:SER:HB2	7:D:131:THR:HG23	1.37	1.03
28:Y:58:GLY:HA3	37:Y:8436:HOH:O	1.58	1.03
2:9:3024:U:O2'	2:9:3025:G:H4'	1.57	1.03
15:L:164:THR:HG22	15:L:167:GLY:H	1.21	1.03
5:B:162:MET:HE1	5:B:308:LEU:HD21	1.37	1.02
5:B:238:ASN:HD22	5:B:240:GLY:H	1.07	1.02
22:S:71:VAL:HG11	22:S:90:PRO:HB3	1.40	1.01
2:9:3056:A:H2'	2:9:3057:A:H5''	1.41	1.01
1:O:156:C:H5''	15:L:171:ARG:HD3	1.38	1.01
15:L:35:PRO:CG	15:L:38:VAL:HG23	1.91	1.01
1:O:1160:G:H5'	1:O:1161:A:H5'	1.41	1.00
37:O:3213:HOH:O	15:L:157:LEU:HD11	1.60	1.00
1:O:2637:A:H4'	1:O:2638:G:O5'	1.55	1.00
6:C:5:ILE:HD11	6:C:16:VAL:HG23	1.40	1.00
1:O:1878:G:H1'	37:O:5422:HOH:O	1.61	0.99
1:O:200:U:H2'	37:O:9930:HOH:O	1.60	0.99
13:J:10:GLN:NE2	13:J:10:GLN:H	1.61	0.99
37:O:3984:HOH:O	15:L:94:LYS:HE3	1.60	0.99
15:L:87:MET:HB3	31:2:46:ILE:HD13	1.45	0.98
9:F:91:VAL:HG12	9:F:92:GLY:H	1.29	0.98
13:J:14:LYS:HB2	13:J:45:PRO:HG2	1.42	0.98
11:H:165:GLY:HA3	37:H:8384:HOH:O	1.64	0.97
25:V:149:LEU:HG	25:V:153:MET:HE2	1.47	0.97
1:O:871:G:H5'	1:O:871:G:H8	1.27	0.96
2:9:3076:G:H3'	2:9:3077:A:H5''	1.45	0.96

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:1242:A:H5'	12:I:82:THR:HG23	1.46	0.95
11:H:86:ARG:HH11	11:H:133:ILE:HG13	0.79	0.95
21:R:57:THR:HG22	21:R:59:ASP:H	1.32	0.95
28:Y:10:ARG:HA	37:Y:8414:HOH:O	1.67	0.95
1:O:871:G:H5'	1:O:871:G:C8	2.01	0.95
11:H:162:SER:HB2	11:H:163:PRO:HD3	1.46	0.94
1:O:870:G:H2'	1:O:871:G:H5''	1.47	0.94
20:Q:8:ALA:HB1	20:Q:13:THR:HG21	1.49	0.93
11:H:2:PRO:HB2	37:H:8351:HOH:O	1.67	0.93
15:L:106:ASN:HD22	15:L:114:VAL:HG23	1.32	0.92
11:H:139:ASP:HA	37:H:8355:HOH:O	1.69	0.91
13:J:62:PRO:HG3	13:J:65:ARG:HH21	1.34	0.91
15:L:87:MET:HB2	15:L:91:ILE:HD11	1.52	0.91
18:O:115:SER:H	18:O:118:GLN:HE21	0.96	0.91
7:D:134:LEU:HD11	7:D:166:ILE:HD11	1.49	0.91
27:X:187:VAL:HG23	27:X:192:ASP:HB2	1.50	0.91
5:B:62:ARG:HA	5:B:65:MET:HE3	1.50	0.91
1:O:1119:G:H2'	12:I:52:GLN:NE2	1.85	0.91
20:Q:39:THR:HG22	20:Q:42:GLU:H	1.34	0.91
6:C:242:GLU:HG3	37:C:8378:HOH:O	1.69	0.91
1:O:1751:G:H2'	1:O:1752:G:H5''	1.53	0.91
1:O:856:G:H2'	37:O:4865:HOH:O	1.70	0.91
1:O:960:G:H4'	37:O:6805:HOH:O	1.71	0.90
13:J:74:VAL:HG11	13:J:113:ILE:HG12	1.53	0.90
2:9:3006:C:H5''	16:M:37:ARG:NH1	1.87	0.90
11:H:55:GLN:HE21	11:H:124:ARG:HE	1.16	0.90
1:O:1116:U:O2'	1:O:1118:A:H2	1.53	0.90
18:O:143:ALA:HA	37:O:5521:HOH:O	1.72	0.90
15:L:52:LEU:HD11	37:L:8617:HOH:O	1.72	0.90
20:Q:18:LEU:HB2	20:Q:143:VAL:HG12	1.54	0.90
2:9:3023:U:H3'	37:9:8485:HOH:O	1.72	0.90
1:O:1474:C:H6	1:O:1474:C:H5'	1.37	0.89
1:O:2637:A:H5'	37:O:5481:HOH:O	1.72	0.89
7:D:146:LYS:NZ	16:M:107:ASN:HD21	1.70	0.89
6:C:236:THR:HG22	6:C:239:ALA:N	1.86	0.89
1:O:2812:A:H2	1:O:2814:A:H62	1.17	0.89
10:G:12:ILE:HA	37:G:4499:HOH:O	1.71	0.89
2:9:3023:U:H4'	2:9:3024:U:OP2	1.69	0.89
1:O:1184:C:H1'	37:O:6843:HOH:O	1.72	0.89
1:O:1559:A:H1'	37:O:5293:HOH:O	1.72	0.89
1:O:282:C:H1'	1:O:368:C:N4	1.88	0.88

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:1165:G:H3'	1:0:1165:G:OP1	1.73	0.88
15:L:102:GLU:OE1	15:L:164:THR:HG21	1.71	0.88
11:H:26:LYS:HD2	11:H:28:ILE:HD12	1.54	0.88
1:0:542:A:H5'	1:0:542:A:H8	1.39	0.88
29:Z:21:ARG:HD2	29:Z:39:PHE:HB2	1.55	0.88
1:0:1835:U:H5	1:0:1840:A:N7	1.70	0.88
1:0:711:G:H1'	37:0:6483:HOH:O	1.73	0.87
37:0:4667:HOH:O	13:J:39:GLY:HA2	1.74	0.87
7:D:27:ILE:HG22	7:D:28:GLY:H	1.38	0.87
14:K:79:ASP:HB3	37:K:8559:HOH:O	1.75	0.87
15:L:84:LYS:HE2	37:L:8576:HOH:O	1.74	0.87
16:M:47:LEU:HD11	16:M:127:LEU:HD21	1.55	0.87
4:A:223:ARG:HG3	37:A:8596:HOH:O	1.75	0.86
5:B:201:ASP:HB2	5:B:312:ARG:HD2	1.54	0.86
1:0:1116:U:H3	1:0:1246:A:H62	1.24	0.86
11:H:59:ASN:H	11:H:59:ASN:HD22	1.24	0.86
8:E:100:ASP:HB2	37:E:2789:HOH:O	1.74	0.86
15:L:164:THR:HG22	15:L:167:GLY:N	1.90	0.86
1:0:111:C:O2'	29:Z:20:ARG:HG2	1.76	0.86
1:0:541:C:H2'	1:0:542:A:H5''	1.56	0.86
14:K:67:ARG:O	14:K:71:GLU:HG3	1.75	0.86
21:R:51:GLN:HE21	21:R:53:ASN:HD21	1.20	0.86
5:B:190:MET:HE2	5:B:194:PHE:CD1	2.11	0.85
12:I:131:THR:HG22	12:I:134:GLU:H	1.41	0.85
13:J:29:LEU:HB3	13:J:55:VAL:HG11	1.57	0.85
16:M:86:LEU:HD12	16:M:125:ALA:HB2	1.59	0.85
25:V:68:THR:HG23	25:V:69:ARG:HG2	1.58	0.85
4:A:192:VAL:HB	37:A:8587:HOH:O	1.76	0.85
13:J:81:ARG:HB2	13:J:87:ARG:HH11	1.39	0.85
26:W:78:GLU:HG2	26:W:79:GLU:H	1.41	0.85
28:Y:38:LYS:HE2	28:Y:45:LYS:HE2	1.58	0.85
1:0:289:G:H22	1:0:363:A:H2	1.24	0.84
5:B:195:ARG:HG2	5:B:323:LEU:HD22	1.58	0.84
1:0:962:C:H1'	16:M:5:ARG:NH1	1.93	0.84
7:D:20:LYS:HA	7:D:75:LEU:O	1.78	0.84
37:0:3245:HOH:O	22:S:9:LYS:HD2	1.76	0.84
14:K:77:ALA:HB3	37:K:8531:HOH:O	1.76	0.84
17:N:32:ARG:HB2	37:N:4656:HOH:O	1.77	0.84
1:0:2751:C:H3'	37:0:6655:HOH:O	1.76	0.84
15:L:164:THR:HG23	15:L:165:SER:N	1.90	0.84
11:H:139:ASP:N	11:H:140:PRO:HD3	1.93	0.84

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:U:1:THR:HG23	24:U:2:VAL:H	1.39	0.84
25:V:5:VAL:HG11	25:V:153:MET:HE1	1.58	0.84
11:H:41:THR:HA	37:H:8381:HOH:O	1.76	0.84
1:O:1450:C:H4'	1:O:1451:C:OP2	1.77	0.83
1:O:2310:G:OP2	11:H:114:PRO:HD2	1.77	0.83
37:9:8479:HOH:O	16:M:23:ARG:HD3	1.77	0.83
5:B:321:PRO:HA	37:B:8657:HOH:O	1.78	0.83
1:O:1372:A:H3'	37:O:6577:HOH:O	1.77	0.83
4:A:88:ILE:HD13	4:A:100:PRO:HD3	1.57	0.83
28:Y:46:LYS:HD3	28:Y:59:HIS:HB2	1.59	0.83
30:1:41:HIS:H	30:1:45:ASN:HD22	1.26	0.83
1:O:545:G:H5'	1:O:545:G:H8	1.44	0.83
11:H:29:ALA:HB3	11:H:65:ARG:HH12	1.44	0.83
16:M:87:LEU:HD12	16:M:186:LEU:HD21	1.59	0.83
5:B:41:PHE:CD1	5:B:79:MET:HE2	2.13	0.83
5:B:202:VAL:HG11	5:B:301:VAL:HG13	1.61	0.82
18:O:103:THR:HA	18:O:106:ARG:NH1	1.93	0.82
1:O:962:C:H1'	16:M:5:ARG:HH12	1.44	0.82
37:O:3329:HOH:O	11:H:11:LYS:HE2	1.79	0.82
4:A:121:ALA:O	4:A:124:VAL:HG22	1.77	0.82
12:I:93:ARG:HB3	12:I:93:ARG:HH11	1.41	0.82
1:O:1164:U:H4'	1:O:1165:G:OP1	1.79	0.82
25:V:72:PRO:HG2	25:V:77:ALA:HB3	1.59	0.82
25:V:88:THR:HB	37:V:6679:HOH:O	1.78	0.82
1:O:1187:U:H2'	37:O:6289:HOH:O	1.79	0.82
1:O:2890:A:H1'	23:T:56:ARG:NH2	1.94	0.82
26:W:30:MET:HE1	26:W:58:ALA:HB3	1.61	0.82
18:O:115:SER:OG	18:O:118:GLN:HG3	1.79	0.82
11:H:27:LYS:H	11:H:58:HIS:HD2	1.25	0.82
1:O:877:G:H5'	1:O:878:G:OP1	1.80	0.82
8:E:107:PHE:CE2	8:E:108:LEU:HD13	2.15	0.81
9:F:1:PRO:HB2	37:F:5897:HOH:O	1.78	0.81
23:T:20:MET:HE2	23:T:30:HIS:NE2	1.95	0.81
1:O:1834:C:H2'	1:O:1840:A:N6	1.95	0.81
16:M:113:SER:HB2	37:M:6448:HOH:O	1.80	0.81
1:O:541:C:C2'	1:O:542:A:H5''	2.11	0.81
1:O:2094:G:H4'	5:B:245:SER:HB3	1.63	0.81
1:O:2274:A:N3	15:L:86:MET:HE1	1.95	0.81
8:E:97:VAL:HG12	37:E:4191:HOH:O	1.81	0.81
25:V:4:LEU:HD22	25:V:52:VAL:HG21	1.61	0.81
2:9:3024:U:H4'	2:9:3025:G:OP1	1.79	0.81

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:9:3014:G:H5'	2:9:3014:G:H8	1.44	0.81
8:E:20:ILE:HD11	8:E:40:VAL:HG11	1.61	0.81
1:0:1741:U:H5'	1:0:1742:A:OP1	1.80	0.80
22:S:32:ARG:NH1	22:S:38:ARG:HH12	1.78	0.80
37:0:4316:HOH:O	15:L:14:ARG:HG2	1.78	0.80
26:W:72:VAL:HG22	26:W:85:VAL:HG12	1.63	0.80
1:0:157:G:H4'	15:L:95:LYS:HE3	1.63	0.80
1:0:1667:A:H8	1:0:1667:A:H5'	1.46	0.80
9:F:91:VAL:HG12	9:F:92:GLY:N	1.95	0.80
26:W:37:LEU:HD13	26:W:85:VAL:HG21	1.62	0.80
20:Q:9:ASP:O	20:Q:13:THR:HB	1.81	0.80
20:Q:18:LEU:HB2	20:Q:143:VAL:CG1	2.12	0.80
24:U:12:THR:HG22	24:U:15:GLU:CG	2.10	0.80
5:B:162:MET:HE2	5:B:310:ARG:HD3	1.64	0.80
7:D:154:LYS:H	7:D:154:LYS:HD2	1.46	0.80
13:J:62:PRO:HG3	13:J:65:ARG:NH2	1.95	0.80
31:2:60:LYS:HG3	31:2:61:PRO:HD2	1.63	0.80
2:9:3056:A:C2'	2:9:3057:A:H5''	2.11	0.80
1:0:282:C:H1'	1:0:368:C:H42	1.47	0.79
8:E:23:GLU:HG2	8:E:28:SER:HB3	1.64	0.79
15:L:35:PRO:HG2	15:L:38:VAL:HG23	1.61	0.79
1:0:714:U:H3'	37:0:6334:HOH:O	1.82	0.79
1:0:1909:A:H2'	1:0:1910:A:C8	2.17	0.79
20:Q:99:ALA:HB1	20:Q:109:MET:CE	2.10	0.79
30:1:39:ARG:HG2	37:1:3143:HOH:O	1.82	0.79
1:0:2478:U:H2'	1:0:2479:A:H8	1.47	0.79
25:V:6:GLN:HB2	25:V:26:ILE:HD12	1.64	0.79
26:W:41:PHE:O	26:W:43:VAL:HG23	1.82	0.79
31:2:25:VAL:HG22	31:2:68:LYS:HG3	1.63	0.79
1:0:630:A:H5''	37:0:4210:HOH:O	1.82	0.79
4:A:35:GLY:O	4:A:36:ASP:HB3	1.82	0.79
2:9:3006:C:H5''	16:M:37:ARG:HH12	1.48	0.79
7:D:91:ALA:HB1	37:D:5198:HOH:O	1.81	0.79
11:H:140:PRO:HB3	37:H:8366:HOH:O	1.82	0.79
12:I:76:ASP:HA	37:I:5907:HOH:O	1.80	0.79
12:I:133:GLY:O	12:I:137:GLU:HG3	1.83	0.79
16:M:80:SER:HB2	37:M:4257:HOH:O	1.82	0.79
8:E:84:MET:HE1	8:E:148:ILE:HD13	1.64	0.79
1:0:1822:A:O2'	1:0:1823:G:H5'	1.82	0.78
27:X:99:ALA:HB2	27:X:233:TYR:CZ	2.18	0.78
6:C:236:THR:HG21	37:C:8370:HOH:O	1.83	0.78

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:462:A:C2	30:1:37:HIS:HB3	2.18	0.78
1:0:812:A:H1'	37:0:3438:HOH:O	1.82	0.78
1:0:188:C:H5''	15:L:163:LEU:HD21	1.64	0.78
1:0:559:U:H6	1:0:559:U:H5'	1.48	0.78
15:L:24:MET:HE3	15:L:28:MET:HE3	1.62	0.78
1:0:1118:A:C8	1:0:1118:A:H3'	2.18	0.78
15:L:72:SER:HB2	15:L:93:ARG:HG2	1.66	0.78
1:0:2637:A:H4'	1:0:2638:G:C5'	2.14	0.78
37:0:6832:HOH:O	5:B:211:THR:HG21	1.83	0.78
5:B:238:ASN:HD22	5:B:240:GLY:N	1.81	0.78
1:0:182:G:H5'	37:0:4607:HOH:O	1.84	0.78
9:F:96:ALA:HA	37:F:3111:HOH:O	1.83	0.78
15:L:37:VAL:HG21	15:L:108:LYS:HG3	1.65	0.78
1:0:1166:A:H1'	1:0:1192:A:C2	2.18	0.78
18:O:115:SER:H	18:O:118:GLN:NE2	1.78	0.78
26:W:71:ARG:HB3	26:W:88:GLU:OE1	1.84	0.78
12:I:19:MET:CE	12:I:132:LEU:HD11	2.13	0.78
1:0:814:G:H8	37:0:6598:HOH:O	1.66	0.77
27:X:200:THR:HG22	27:X:201:GLU:HG2	1.67	0.77
1:0:1701:A:H5'	37:0:5694:HOH:O	1.84	0.77
15:L:186:SER:O	15:L:189:VAL:HG12	1.82	0.77
1:0:871:G:H8	1:0:871:G:C5'	1.97	0.77
37:0:5654:HOH:O	4:A:5:GLN:HB3	1.83	0.77
5:B:195:ARG:HD2	5:B:324:ASP:OD1	1.84	0.77
18:O:115:SER:N	18:O:118:GLN:HE21	1.80	0.77
28:Y:38:LYS:HG2	28:Y:45:LYS:HG2	1.65	0.77
4:A:153:ARG:HB2	4:A:153:ARG:HH11	1.49	0.77
4:A:192:VAL:HG12	4:A:207:GLN:HB3	1.65	0.77
23:T:14:GLU:OE1	23:T:15:PRO:HD2	1.85	0.77
11:H:26:LYS:HG2	11:H:28:ILE:H	1.49	0.77
11:H:162:SER:HB2	11:H:163:PRO:CD	2.13	0.77
27:X:212:ARG:HD2	37:X:8606:HOH:O	1.83	0.77
1:0:2414:A:H2'	1:0:2415:A:C8	2.20	0.77
1:0:2586:U:H3	1:0:2592:G:H22	1.28	0.77
4:A:211:LYS:HB3	4:A:212:PRO:HD2	1.66	0.77
5:B:304:PRO:HD2	5:B:307:ARG:HD2	1.66	0.77
25:V:4:LEU:HD22	25:V:52:VAL:CG2	2.14	0.77
1:0:794:U:H3	1:0:819:A:H61	1.32	0.77
16:M:7:LYS:HE3	19:P:21:ARG:O	1.85	0.77
1:0:1206:U:H5'	1:0:1206:U:H6	1.49	0.77
4:A:36:ASP:OD2	4:A:85:ASP:HB2	1.85	0.77

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:V:88:THR:HG22	25:V:89:ASP:H	1.48	0.77
5:B:175:LEU:C	5:B:175:LEU:HD23	2.10	0.76
15:L:38:VAL:C	15:L:63:VAL:HG13	2.10	0.76
16:M:169:PRO:O	16:M:172:PHE:HB3	1.85	0.76
22:S:9:LYS:HE3	22:S:13:ARG:NH1	1.99	0.76
6:C:78:ARG:HH11	6:C:78:ARG:HG3	1.51	0.76
27:X:187:VAL:HG23	27:X:192:ASP:CB	2.15	0.76
1:O:1165:G:H4'	1:O:1174:A:O2'	1.85	0.76
1:O:2908:A:H2'	1:O:2909:G:O4'	1.86	0.76
8:E:166:VAL:HG12	37:E:3134:HOH:O	1.85	0.76
20:Q:106:GLY:HA2	20:Q:109:MET:HE3	1.67	0.76
5:B:156:LYS:HE3	37:B:8628:HOH:O	1.85	0.76
9:F:46:GLU:O	9:F:73:PRO:HD2	1.84	0.76
16:M:151:ASP:OD1	16:M:154:LEU:HD13	1.86	0.76
4:A:199:HIS:HD2	4:A:201:PHE:H	1.33	0.76
10:G:23:ILE:HD13	10:G:67:LEU:HD23	1.67	0.76
14:K:136:ALA:HB3	37:K:8573:HOH:O	1.86	0.76
16:M:49:THR:HG22	16:M:56:ASP:HB2	1.68	0.76
25:V:21:LEU:HD21	25:V:48:VAL:CG1	2.16	0.76
14:K:148:GLU:HB2	37:K:8587:HOH:O	1.84	0.76
15:L:164:THR:CG2	15:L:167:GLY:H	1.96	0.76
1:O:560:C:H42	1:O:597:A:H61	1.33	0.76
1:O:1625:U:H4'	37:O:4122:HOH:O	1.86	0.75
1:O:1679:C:H5'	37:O:8840:HOH:O	1.86	0.75
37:O:5706:HOH:O	7:D:99:ASP:HA	1.85	0.75
14:K:143:THR:HG21	37:K:8539:HOH:O	1.84	0.75
5:B:162:MET:CE	5:B:308:LEU:HD21	2.14	0.75
6:C:5:ILE:HD11	6:C:16:VAL:CG2	2.15	0.75
1:O:2533:C:H5'	1:O:2533:C:H6	1.51	0.75
37:O:3272:HOH:O	15:L:189:VAL:HG21	1.85	0.75
1:O:870:G:C2'	1:O:871:G:H5''	2.16	0.75
4:A:105:VAL:HG13	4:A:155:THR:O	1.85	0.75
1:O:1474:C:H5'	1:O:1474:C:C6	2.21	0.75
13:J:74:VAL:CG1	13:J:113:ILE:HG12	2.15	0.75
1:O:1164:U:H3	1:O:1192:A:H2	1.31	0.75
1:O:1160:G:C5'	1:O:1161:A:H5'	2.17	0.75
1:O:2637:A:H5''	1:O:2638:G:H5'	1.68	0.75
16:M:89:GLY:O	16:M:92:ALA:HB3	1.87	0.75
1:O:56:G:H5''	24:U:50:ARG:HH12	1.52	0.75
25:V:149:LEU:CG	25:V:153:MET:HE2	2.17	0.75
28:Y:37:HIS:HB2	28:Y:47:LEU:HB2	1.69	0.75

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:2851:G:O2'	1:0:2852:A:H5'	1.87	0.74
37:0:3171:HOH:O	15:L:79:LYS:HD3	1.87	0.74
23:T:14:GLU:O	23:T:17:THR:HB	1.87	0.74
1:0:381:G:H5''	37:0:3789:HOH:O	1.86	0.74
8:E:23:GLU:HG2	8:E:28:SER:CB	2.17	0.74
15:L:34:GLU:HB3	15:L:35:PRO:HD2	1.69	0.74
2:9:3025:G:H3'	2:9:3026:C:H5'	1.68	0.74
11:H:47:GLU:HB3	11:H:133:ILE:CD1	2.17	0.74
25:V:21:LEU:HD21	25:V:48:VAL:HG11	1.68	0.74
2:9:3025:G:H3'	2:9:3026:C:C5'	2.18	0.74
5:B:168:GLY:N	5:B:174:ARG:HD3	2.02	0.74
8:E:84:MET:HE1	8:E:148:ILE:CD1	2.18	0.74
1:0:2325:C:H1'	37:0:3624:HOH:O	1.88	0.74
5:B:27:ASN:HD22	5:B:27:ASN:H	1.35	0.74
5:B:253:GLN:HA	37:B:8620:HOH:O	1.87	0.74
15:L:139:PRO:O	15:L:140:ALA:HB3	1.86	0.74
21:R:57:THR:HG22	21:R:59:ASP:N	2.00	0.74
26:W:72:VAL:HG22	26:W:85:VAL:CG1	2.17	0.74
1:0:317:A:H5'	22:S:52:ARG:HD2	1.68	0.74
30:1:41:HIS:N	30:1:45:ASN:HD22	1.84	0.74
1:0:21:G:C5'	20:Q:2:ILE:HA	2.18	0.74
2:9:3048:C:H4'	16:M:141:ARG:HH21	1.51	0.74
5:B:18:ARG:HG3	5:B:256:GLN:HG3	1.68	0.74
7:D:64:ARG:HG2	7:D:67:ASP:HB3	1.68	0.74
11:H:4:ALA:HB3	37:H:8351:HOH:O	1.88	0.74
1:0:240:C:H4'	15:L:146:GLN:NE2	2.02	0.74
1:0:1383:U:H5''	37:0:6061:HOH:O	1.86	0.74
1:0:2291:A:C8	1:0:2309:C:H5'	2.22	0.74
13:J:10:GLN:H	13:J:10:GLN:HE21	1.35	0.73
17:N:47:ARG:HA	17:N:50:ARG:HH12	1.51	0.73
1:0:1328:A:OP1	27:X:169:ARG:HD2	1.88	0.73
12:I:107:ASN:ND2	12:I:109:TYR:H	1.87	0.73
1:0:2502:C:H4'	11:H:151:MET:HG2	1.71	0.73
7:D:105:SER:CB	7:D:131:THR:HG23	2.15	0.73
1:0:1134:G:C4'	11:H:151:MET:HE1	2.16	0.73
1:0:2004:U:H4'	37:0:4752:HOH:O	1.88	0.73
4:A:69:LEU:HD21	4:A:120:ARG:HB3	1.69	0.73
25:V:122:ARG:HG2	25:V:122:ARG:HH11	1.54	0.73
7:D:19:GLU:O	7:D:20:LYS:HG2	1.89	0.73
7:D:37:ALA:O	7:D:40:ILE:HG12	1.88	0.73
12:I:52:GLN:HG3	12:I:53:ILE:N	2.03	0.73

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:M:144:GLY:O	16:M:147:ILE:HG22	1.89	0.73
5:B:225:GLY:HA3	37:B:8568:HOH:O	1.88	0.73
11:H:28:ILE:HA	11:H:62:GLU:OE1	1.88	0.73
29:Z:25:LYS:HD2	30:1:49:GLU:N	2.04	0.73
28:Y:59:HIS:HA	37:Y:8438:HOH:O	1.87	0.73
1:O:2587:U:H2'	1:O:2589:U:H5''	1.71	0.72
1:O:1118:A:H3'	1:O:1118:A:H8	1.54	0.72
1:O:2256:G:H2'	1:O:2257:G:H5'	1.71	0.72
26:W:76:ARG:HH11	26:W:76:ARG:HG3	1.52	0.72
28:Y:40:PRO:HD3	28:Y:47:LEU:HD11	1.71	0.72
8:E:43:ASP:HA	37:E:5864:HOH:O	1.88	0.72
11:H:56:ILE:HG22	11:H:61:LEU:HD22	1.72	0.72
15:L:74:ARG:HH11	15:L:74:ARG:HG3	1.53	0.72
1:O:2637:A:C5'	37:O:5481:HOH:O	2.34	0.72
37:O:5220:HOH:O	15:L:170:CYS:SG	2.46	0.72
17:N:42:GLU:HB2	37:N:2176:HOH:O	1.89	0.72
17:N:47:ARG:HA	17:N:50:ARG:NH1	2.04	0.72
1:O:285:A:H2'	1:O:286:U:O4'	1.90	0.72
1:O:1028:U:H1'	37:O:3135:HOH:O	1.89	0.72
1:O:1234:U:N3	5:B:244:PRO:HB3	2.04	0.72
1:O:541:C:H2'	1:O:542:A:C5'	2.19	0.72
1:O:1666:C:O2'	1:O:1667:A:H5''	1.90	0.72
11:H:5:MET:HG3	37:H:8351:HOH:O	1.89	0.72
5:B:43:GLY:O	5:B:308:LEU:HD12	1.89	0.72
9:F:58:GLU:HA	9:F:61:MET:HG3	1.71	0.72
11:H:137:ASN:O	11:H:139:ASP:N	2.22	0.72
11:H:151:MET:HA	11:H:151:MET:HE3	1.72	0.72
25:V:13:MET:HE2	25:V:18:GLN:CA	2.20	0.72
1:O:1044:C:H5''	37:O:8544:HOH:O	1.88	0.72
1:O:1666:C:H2'	1:O:1667:A:H5'	1.71	0.72
7:D:36:ASN:HA	37:D:7500:HOH:O	1.88	0.72
1:O:232:A:H4'	37:O:5501:HOH:O	1.90	0.71
22:S:52:ARG:HB2	22:S:95:ASN:HB3	1.71	0.71
25:V:14:HIS:HB2	25:V:17:ILE:HG13	1.71	0.71
1:O:2426:G:H1'	37:O:5510:HOH:O	1.88	0.71
11:H:14:TYR:H	11:H:91:HIS:CE1	2.07	0.71
11:H:47:GLU:HB3	11:H:133:ILE:HD13	1.72	0.71
25:V:21:LEU:HD22	25:V:26:ILE:HD11	1.71	0.71
25:V:122:ARG:HH21	25:V:154:ARG:HD2	1.55	0.71
9:F:2:VAL:HG22	9:F:57:GLU:OE1	1.90	0.71
16:M:154:LEU:O	16:M:155:GLU:HB3	1.88	0.71

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:1119:G:N2	1:0:1246:A:C2	2.58	0.71
8:E:37:ASP:OD1	12:I:125:SER:HB3	1.90	0.71
16:M:163:PHE:HE1	16:M:171:HIS:HD1	1.38	0.71
29:Z:28:HIS:HD2	29:Z:30:LYS:H	1.35	0.71
1:0:2559:C:H4'	37:0:6644:HOH:O	1.89	0.71
1:0:2862:G:H4'	5:B:336:GLN:O	1.91	0.71
5:B:119:HIS:O	5:B:121:PRO:HD3	1.91	0.71
1:0:777:U:O2'	29:Z:11:LYS:HG2	1.91	0.71
1:0:2405:C:H5'	37:0:5993:HOH:O	1.90	0.71
25:V:154:ARG:C	37:V:4276:HOH:O	2.34	0.71
27:X:200:THR:HG22	27:X:201:GLU:CG	2.20	0.71
27:X:235:GLU:H	27:X:235:GLU:CD	1.97	0.71
1:0:2508:C:H2'	37:0:6145:HOH:O	1.89	0.71
7:D:135:VAL:HG22	7:D:136:ARG:H	1.55	0.71
9:F:104:ALA:HA	37:F:6617:HOH:O	1.90	0.71
13:J:74:VAL:HG13	13:J:113:ILE:HG23	1.72	0.71
16:M:151:ASP:O	16:M:154:LEU:HB2	1.91	0.71
16:M:183:ASP:OD2	16:M:186:LEU:HD12	1.89	0.71
1:0:1080:C:H4'	1:0:1081:A:OP1	1.90	0.71
1:0:2897:C:H2'	1:0:2898:G:H8	1.56	0.71
26:W:15:ARG:HB3	26:W:15:ARG:HH11	1.55	0.71
7:D:27:ILE:HG22	7:D:28:GLY:N	2.07	0.70
12:I:74:ARG:HD3	37:I:5061:HOH:O	1.91	0.70
1:0:1835:U:C5	1:0:1840:A:N7	2.56	0.70
1:0:21:G:H5'	20:Q:2:ILE:HA	1.74	0.70
1:0:2276:U:H2'	1:0:2277:U:C6	2.27	0.70
1:0:2780:C:H1'	8:E:143:GLN:HE21	1.57	0.70
37:0:4403:HOH:O	2:9:3103:A:H4'	1.91	0.70
16:M:33:ARG:NH1	16:M:103:ASP:OD2	2.22	0.70
7:D:38:GLU:OE2	7:D:51:ARG:CZ	2.40	0.70
7:D:39:ASP:HB2	37:D:5583:HOH:O	1.91	0.70
13:J:14:LYS:HD2	35:J:8512:CL:CL	2.28	0.70
18:O:59:ARG:NH2	18:O:66:GLN:HE22	1.89	0.70
1:0:1285:U:H4'	25:V:74:GLU:OE1	1.92	0.70
37:0:9204:HOH:O	5:B:254:GLN:HG3	1.90	0.70
7:D:55:LYS:HA	37:D:6752:HOH:O	1.92	0.70
1:0:1684:A:H1'	30:1:43:ARG:HH22	1.57	0.70
1:0:2478:U:H2'	1:0:2479:A:C8	2.26	0.70
13:J:81:ARG:HB2	13:J:87:ARG:NH1	2.06	0.70
7:D:88:LEU:HB2	7:D:89:PRO:HD3	1.73	0.70
1:0:56:G:H5''	24:U:50:ARG:NH1	2.07	0.69

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:1330:A:H5''	1:0:1331:A:OP2	1.92	0.69
2:9:3092:G:H2'	2:9:3093:A:C8	2.27	0.69
12:I:104:TYR:HA	37:I:2238:HOH:O	1.92	0.69
27:X:133:HIS:HD2	37:X:8583:HOH:O	1.74	0.69
1:0:2346:C:O2'	7:D:52:THR:HG21	1.92	0.69
6:C:236:THR:H	6:C:239:ALA:HB3	1.57	0.69
7:D:41:LEU:HA	7:D:44:ILE:HG22	1.74	0.69
11:H:26:LYS:HD2	11:H:28:ILE:CD1	2.22	0.69
1:0:1162:G:H2'	37:0:5981:HOH:O	1.92	0.69
37:0:6529:HOH:O	29:Z:1:THR:HB	1.91	0.69
15:L:153:THR:HB	37:L:8613:HOH:O	1.92	0.69
24:U:39:ALA:N	24:U:40:PRO:HD2	2.08	0.69
1:0:506:G:H3'	37:0:3260:HOH:O	1.91	0.69
1:0:214:U:H5'	37:0:5555:HOH:O	1.91	0.69
13:J:115:ARG:HG3	13:J:116:GLU:N	2.06	0.69
1:0:2301:A:H5''	1:0:2302:A:H5'	1.75	0.69
2:9:3035:C:H5''	37:9:8459:HOH:O	1.93	0.69
1:0:1120:U:H5''	1:0:1120:U:C6	2.28	0.69
8:E:11:VAL:HG12	8:E:12:ASP:N	2.06	0.69
10:G:12:ILE:N	10:G:13:PRO:HD3	2.06	0.69
16:M:143:ARG:HA	16:M:172:PHE:CD2	2.27	0.69
1:0:2256:G:C2'	1:0:2257:G:H5'	2.23	0.69
37:0:3582:HOH:O	5:B:158:LYS:HB2	1.91	0.69
7:D:101:THR:HG22	37:D:7400:HOH:O	1.92	0.69
11:H:59:ASN:HD22	11:H:59:ASN:N	1.90	0.69
16:M:164:ASP:CG	16:M:167:ASP:HA	2.18	0.69
23:T:13:ILE:HG12	23:T:32:CYS:HB3	1.75	0.69
25:V:88:THR:HG23	25:V:110:GLN:NE2	2.06	0.69
26:W:25:ARG:HD2	37:W:3861:HOH:O	1.92	0.69
1:0:656:G:H5'	17:N:3:THR:HB	1.74	0.69
1:0:2679:G:H2'	1:0:2681:A:OP2	1.92	0.69
9:F:99:THR:HA	37:F:3461:HOH:O	1.92	0.69
12:I:74:ARG:HB3	12:I:74:ARG:HH11	1.55	0.69
31:2:30:GLN:HB3	37:2:8544:HOH:O	1.91	0.69
1:0:236:A:H4'	1:0:237:G:H5'	1.75	0.69
1:0:2690:U:O2'	8:E:111:LYS:HE3	1.93	0.69
9:F:63:ILE:HB	9:F:64:PRO:HD3	1.75	0.69
10:G:12:ILE:HB	37:G:4714:HOH:O	1.92	0.69
24:U:49:LEU:O	24:U:53:ILE:HG13	1.93	0.69
25:V:4:LEU:HD23	25:V:54:PHE:HB3	1.75	0.69
37:0:5730:HOH:O	7:D:55:LYS:HB2	1.91	0.68

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:9:3014:G:H5'	2:9:3014:G:C8	2.28	0.68
8:E:9:GLU:HA	37:E:5240:HOH:O	1.93	0.68
23:T:6:CYS:HA	23:T:13:ILE:HD11	1.75	0.68
25:V:4:LEU:O	25:V:32:CYS:HA	1.93	0.68
1:0:1589:G:N2	1:0:1605:G:H1'	2.08	0.68
4:A:199:HIS:CD2	4:A:201:PHE:H	2.11	0.68
11:H:162:SER:CB	11:H:163:PRO:HD3	2.22	0.68
14:K:133:VAL:HA	37:K:8573:HOH:O	1.92	0.68
22:S:50:VAL:HG12	22:S:56:ALA:HA	1.75	0.68
1:0:821:U:H2'	1:0:822:C:H6	1.59	0.68
1:0:1116:U:HO2'	1:0:1118:A:H2	0.74	0.68
1:0:1191:A:H3'	1:0:1192:A:H5''	1.75	0.68
5:B:212:GLN:HB2	5:B:257:THR:HG21	1.75	0.68
16:M:48:VAL:CG1	16:M:55:ASP:HB3	2.23	0.68
17:N:87:THR:O	17:N:91:GLN:HG3	1.94	0.68
20:Q:40:ALA:O	20:Q:44:VAL:HG23	1.93	0.68
1:0:338:C:H5''	37:0:5264:HOH:O	1.92	0.68
1:0:2241:C:H2'	1:0:2242:U:C6	2.29	0.68
2:9:3039:U:H1'	2:9:3044:A:H61	1.58	0.68
4:A:101:GLU:OE2	4:A:131:HIS:HB2	1.92	0.68
11:H:139:ASP:H	11:H:140:PRO:HD3	1.58	0.68
23:T:9:CYS:HA	23:T:52:THR:HG23	1.75	0.68
1:0:1185:U:H2'	1:0:1186:C:C6	2.29	0.68
5:B:238:ASN:ND2	5:B:240:GLY:H	1.86	0.68
10:G:64:ASN:O	10:G:68:GLU:HG3	1.93	0.68
11:H:48:LEU:HG	11:H:157:ILE:HG21	1.76	0.68
4:A:210:GLY:HA3	37:A:8579:HOH:O	1.93	0.68
5:B:168:GLY:H	5:B:174:ARG:HD3	1.58	0.68
6:C:129:HIS:CE1	6:C:231:ARG:HA	2.28	0.68
7:D:49:PRO:HG3	37:D:5828:HOH:O	1.94	0.68
14:K:143:THR:HG22	14:K:144:ASP:N	2.07	0.68
1:0:454:U:C2	37:0:8548:HOH:O	2.44	0.68
16:M:11:ARG:HG3	16:M:14:ARG:NH1	2.08	0.68
22:S:9:LYS:HE3	22:S:13:ARG:HH11	1.58	0.68
28:Y:11:THR:OG1	28:Y:23:ARG:HB2	1.94	0.68
30:1:35:ARG:HB2	37:1:2691:HOH:O	1.93	0.68
1:0:2638:G:H5'	37:0:5481:HOH:O	1.94	0.68
5:B:175:LEU:HD23	5:B:175:LEU:O	1.94	0.68
7:D:57:THR:HG23	7:D:63:ILE:HG22	1.75	0.68
7:D:64:ARG:CG	7:D:67:ASP:HB3	2.24	0.68
27:X:186:ARG:HH11	27:X:186:ARG:HG2	1.57	0.68

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:Y:49:ARG:HD2	37:Y:8425:HOH:O	1.93	0.68
1:O:2890:A:H2'	37:O:4637:HOH:O	1.93	0.67
9:F:50:VAL:HG21	9:F:63:ILE:HG21	1.76	0.67
15:L:152:ARG:HB3	37:L:8639:HOH:O	1.93	0.67
1:O:154:C:H2'	1:O:155:C:H6	1.57	0.67
1:O:182:G:H4'	15:L:157:LEU:HD13	1.77	0.67
8:E:6:GLU:HA	8:E:46:THR:HG22	1.74	0.67
20:Q:39:THR:HB	20:Q:42:GLU:HG3	1.76	0.67
1:O:553:G:H5'	37:O:9985:HOH:O	1.95	0.67
11:H:45:GLN:HB3	11:H:163:PRO:CD	2.16	0.67
14:K:143:THR:HG22	14:K:145:LEU:H	1.58	0.67
25:V:13:MET:HE2	25:V:18:GLN:HA	1.77	0.67
26:W:21:PRO:HG2	26:W:24:LYS:HD3	1.77	0.67
20:Q:119:VAL:HG21	20:Q:142:ASP:CG	2.19	0.67
27:X:220:GLU:HG2	37:X:8550:HOH:O	1.93	0.67
1:O:2793:A:H5'	37:O:4014:HOH:O	1.94	0.67
6:C:237:GLU:HB2	37:C:8427:HOH:O	1.95	0.67
16:M:141:ARG:N	37:M:7307:HOH:O	2.27	0.67
13:J:55:VAL:HG12	13:J:56:SER:N	2.10	0.67
16:M:155:GLU:O	16:M:156:GLU:HG3	1.95	0.67
5:B:240:GLY:HA3	37:B:8529:HOH:O	1.94	0.67
4:A:88:ILE:O	4:A:88:ILE:HG22	1.94	0.67
5:B:36:PRO:HA	5:B:168:GLY:HA2	1.77	0.67
25:V:122:ARG:HH22	25:V:154:ARG:C	2.03	0.67
15:L:60:ILE:C	15:L:61:ILE:HD12	2.19	0.67
17:N:88:LYS:HB3	37:N:7061:HOH:O	1.94	0.67
20:Q:39:THR:HG23	20:Q:107:GLU:O	1.95	0.67
25:V:88:THR:HG23	25:V:110:GLN:HE21	1.60	0.67
1:O:338:C:H4'	6:C:174:ILE:CD1	2.24	0.66
1:O:2502:C:C4'	11:H:151:MET:HG2	2.25	0.66
8:E:81:GLU:HG2	8:E:134:SER:HB3	1.77	0.66
14:K:72:ASN:O	14:K:76:LEU:HG	1.95	0.66
30:1:44:ARG:HG2	37:1:5527:HOH:O	1.94	0.66
1:O:2827:A:H2'	1:O:2828:G:O4'	1.96	0.66
21:R:51:GLN:NE2	21:R:53:ASN:HD21	1.93	0.66
1:O:902:G:N7	14:K:18:HIS:HD2	1.92	0.66
1:O:1165:G:OP1	1:O:1165:G:C3'	2.44	0.66
1:O:2237:G:H1'	37:O:4307:HOH:O	1.94	0.66
37:9:8471:HOH:O	16:M:147:ILE:HD12	1.94	0.66
5:B:51:VAL:HG13	5:B:53:LEU:HD13	1.76	0.66
6:C:139:VAL:HG13	37:C:8444:HOH:O	1.94	0.66

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:U:42:ASN:HB3	37:U:7247:HOH:O	1.96	0.66
31:2:17:HIS:O	31:2:18:GLN:HG3	1.95	0.66
2:9:3064:C:H2'	2:9:3065:A:H5'	1.76	0.66
6:C:1:MET:HG2	6:C:2:GLN:H	1.60	0.66
26:W:15:ARG:HB3	26:W:15:ARG:NH1	2.11	0.66
1:0:2064:U:H4'	1:0:2653:A:OP1	1.95	0.66
37:0:8909:HOH:O	15:L:94:LYS:HE2	1.95	0.66
2:9:3006:C:OP1	16:M:37:ARG:NH1	2.28	0.66
5:B:104:GLU:HG3	37:B:8593:HOH:O	1.96	0.66
31:2:65:THR:HG23	31:2:67:LEU:HG	1.77	0.66
1:0:2256:G:H2'	1:0:2257:G:C5'	2.26	0.66
9:F:91:VAL:CG1	9:F:92:GLY:H	2.06	0.66
10:G:63:ARG:N	37:G:2569:HOH:O	2.28	0.66
12:I:90:LYS:HB2	35:I:8502:CL:CL	2.32	0.66
20:Q:14:ALA:HB3	20:Q:147:LEU:HB2	1.77	0.66
25:V:81:ASP:OD1	25:V:92:ASP:HB2	1.95	0.66
1:0:1160:G:H5'	1:0:1161:A:C5'	2.22	0.66
11:H:27:LYS:N	11:H:58:HIS:HD2	1.94	0.66
11:H:46:VAL:O	11:H:146:TRP:HH2	1.79	0.66
12:I:107:ASN:HD22	12:I:107:ASN:C	2.02	0.66
15:L:139:PRO:O	15:L:140:ALA:CB	2.43	0.66
15:L:173:LEU:HD23	15:L:183:VAL:HG12	1.77	0.66
1:0:1593:C:H5'	18:O:116:SER:O	1.95	0.66
1:0:1603:A:H5'	1:0:1605:G:O4'	1.94	0.66
13:J:34:VAL:HB	37:J:7169:HOH:O	1.96	0.66
24:U:5:VAL:HG23	37:U:2271:HOH:O	1.96	0.66
27:X:185:VAL:HA	37:X:8565:HOH:O	1.94	0.66
7:D:23:VAL:HG23	7:D:23:VAL:O	1.96	0.66
12:I:19:MET:HE2	12:I:79:PHE:HA	1.78	0.66
1:0:1159:G:H21	1:0:1189:A:H8	1.44	0.65
1:0:2094:G:C4'	5:B:245:SER:HB3	2.25	0.65
8:E:7:ILE:HD11	8:E:11:VAL:C	2.22	0.65
15:L:97:ILE:HA	15:L:100:ILE:HD12	1.77	0.65
16:M:24:LEU:HD13	19:P:26:PRO:HB3	1.77	0.65
1:0:289:G:N2	1:0:363:A:H2	1.92	0.65
1:0:869:G:OP1	15:L:79:LYS:HE2	1.96	0.65
5:B:62:ARG:CA	5:B:65:MET:HE3	2.25	0.65
14:K:114:VAL:HG11	37:K:8573:HOH:O	1.95	0.65
1:0:485:A:N3	1:0:487:G:H5''	2.11	0.65
6:C:107:ARG:NH1	6:C:107:ARG:HB3	2.12	0.65
8:E:31:ARG:NH1	37:E:5919:HOH:O	2.29	0.65

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:L:35:PRO:HG3	15:L:38:VAL:HG23	1.76	0.65
29:Z:10:LYS:HG3	37:Z:8432:HOH:O	1.95	0.65
1:0:2111:G:H1'	37:0:8566:HOH:O	1.96	0.65
1:0:2722:G:H4'	37:J:5029:HOH:O	1.96	0.65
24:U:44:GLY:O	24:U:48:GLU:HG2	1.95	0.65
27:X:187:VAL:HB	37:X:8571:HOH:O	1.96	0.65
2:9:3107:C:C5	37:9:8440:HOH:O	2.49	0.65
5:B:125:GLU:O	5:B:129:ARG:HG3	1.97	0.65
5:B:280:VAL:HG13	5:B:334:SER:HA	1.77	0.65
7:D:135:VAL:HG22	7:D:136:ARG:N	2.11	0.65
25:V:7:LEU:HD12	25:V:53:ALA:HB2	1.79	0.65
2:9:3024:U:O2'	2:9:3025:G:C4'	2.41	0.65
5:B:36:PRO:HA	5:B:168:GLY:CA	2.27	0.65
1:0:462:A:C8	37:0:4337:HOH:O	2.50	0.65
2:9:3028:U:H2'	2:9:3029:C:C6	2.32	0.65
5:B:154:VAL:HG12	5:B:156:LYS:HG2	1.79	0.65
12:I:26:VAL:HG13	12:I:36:VAL:HG11	1.79	0.65
13:J:81:ARG:HD3	13:J:87:ARG:NH1	2.12	0.65
1:0:1972:U:H2'	1:0:1973:A:H5'	1.79	0.65
1:0:2035:C:O2'	1:0:2036:C:H5'	1.97	0.65
5:B:54:VAL:HB	37:B:8611:HOH:O	1.95	0.65
7:D:146:LYS:HZ3	16:M:107:ASN:HD21	1.45	0.65
28:Y:30:GLU:HA	28:Y:33:HIS:HB3	1.79	0.65
5:B:179:LEU:O	5:B:183:GLU:HG2	1.95	0.65
6:C:16:VAL:HG12	6:C:17:ASP:N	2.11	0.65
6:C:162:VAL:HG12	6:C:192:ILE:HD11	1.79	0.65
11:H:35:ASN:ND2	11:H:80:ASN:HA	2.12	0.65
11:H:71:TYR:C	11:H:73:GLN:H	2.05	0.65
20:Q:82:GLU:O	20:Q:86:LYS:HG3	1.97	0.65
1:0:567:U:H5''	37:V:5817:HOH:O	1.97	0.65
1:0:694:A:H2'	1:0:695:C:H5'	1.79	0.65
5:B:314:ALA:HB3	5:B:317:PRO:HG3	1.79	0.65
26:W:71:ARG:HD3	37:W:7542:HOH:O	1.97	0.65
1:0:542:A:H5'	1:0:542:A:C8	2.28	0.64
1:0:544:G:H2'	1:0:545:G:H5''	1.79	0.64
1:0:1118:A:H62	1:0:1244:U:H3	1.45	0.64
1:0:2241:C:H2'	1:0:2242:U:H6	1.62	0.64
1:0:2837:U:H2'	37:0:6230:HOH:O	1.97	0.64
2:9:3003:A:H2'	37:9:8424:HOH:O	1.97	0.64
2:9:3040:C:N4	7:D:51:ARG:HB2	2.11	0.64
6:C:141:SER:HA	37:C:8376:HOH:O	1.97	0.64

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:Z:25:LYS:HD2	30:1:49:GLU:H	1.62	0.64
1:0:1015:C:H2'	1:0:1016:U:H6	1.61	0.64
7:D:99:ASP:HB3	7:D:103:ASN:H	1.60	0.64
11:H:84:ARG:NH2	11:H:135:TRP:HH2	1.95	0.64
13:J:10:GLN:HE21	13:J:10:GLN:N	1.95	0.64
14:K:104:ASP:O	14:K:105:TYR:HB3	1.97	0.64
27:X:107:PRO:HB3	27:X:182:PHE:CE2	2.32	0.64
27:X:149:GLN:NE2	37:X:8607:HOH:O	2.29	0.64
1:0:545:G:H5'	1:0:545:G:C8	2.31	0.64
1:0:1120:U:H5''	1:0:1120:U:H6	1.62	0.64
7:D:64:ARG:CD	7:D:67:ASP:HB3	2.27	0.64
7:D:146:LYS:NZ	16:M:107:ASN:ND2	2.45	0.64
13:J:10:GLN:NE2	13:J:10:GLN:N	2.41	0.64
14:K:17:SER:C	14:K:19:LYS:H	2.06	0.64
27:X:107:PRO:HB3	27:X:182:PHE:CD2	2.33	0.64
1:0:2326:U:H4'	1:0:2412:G:H4'	1.79	0.64
1:0:2719:A:C2	5:B:70:PRO:HG3	2.33	0.64
1:0:2768:A:H2'	1:0:2769:C:O4'	1.96	0.64
30:1:40:ARG:HG3	30:1:45:ASN:CB	2.28	0.64
1:0:138:U:H5''	1:0:139:C:OP2	1.98	0.64
1:0:1209:C:H4'	37:0:4724:HOH:O	1.96	0.64
5:B:55:ASN:HB3	5:B:63:GLU:HA	1.79	0.64
11:H:150:LYS:HE2	37:H:8368:HOH:O	1.96	0.64
25:V:110:GLN:NE2	25:V:110:GLN:HA	2.13	0.64
1:0:2594:C:O2'	1:0:2595:U:H5'	1.98	0.64
8:E:15:GLN:HG3	8:E:20:ILE:HG12	1.78	0.64
11:H:46:VAL:HG12	11:H:146:TRP:HZ3	1.63	0.64
14:K:55:GLN:HA	14:K:58:GLN:NE2	2.11	0.64
17:N:14:LEU:HD23	17:N:102:ILE:HD11	1.79	0.64
26:W:25:ARG:NH1	37:W:3861:HOH:O	2.30	0.64
10:G:16:LYS:O	10:G:20:VAL:HG23	1.97	0.64
11:H:26:LYS:HD2	11:H:28:ILE:HB	1.78	0.64
11:H:26:LYS:HD3	11:H:89:PRO:HG3	1.80	0.64
16:M:37:ARG:NE	37:M:3863:HOH:O	2.31	0.64
11:H:136:VAL:HG22	11:H:137:ASN:O	1.97	0.64
18:O:27:ARG:HA	37:O:3969:HOH:O	1.96	0.64
20:Q:99:ALA:CB	20:Q:109:MET:HE1	2.14	0.64
23:T:37:GLU:HB3	37:T:408:HOH:O	1.96	0.64
26:W:30:MET:HE3	26:W:55:ASN:OD1	1.97	0.64
27:X:141:THR:HG23	37:X:8591:HOH:O	1.98	0.64
1:0:272:A:H5'	1:0:273:G:OP2	1.98	0.64

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:1164:U:C4'	1:0:1165:G:OP1	2.46	0.64
1:0:1362:U:H5'	37:0:9756:HOH:O	1.98	0.64
1:0:2486:A:H1'	3:4:76:A:H2'	1.80	0.64
37:0:6799:HOH:O	22:S:9:LYS:HB2	1.96	0.64
11:H:44:ALA:HA	11:H:163:PRO:O	1.98	0.64
25:V:88:THR:HG22	25:V:89:ASP:N	2.13	0.64
1:0:419:A:H1'	1:0:1921:A:C2	2.32	0.64
5:B:264:GLU:HG2	5:B:267:LYS:HE2	1.79	0.64
17:N:47:ARG:HG3	17:N:47:ARG:HH11	1.62	0.64
31:2:75:GLY:HA2	37:2:8547:HOH:O	1.97	0.64
1:0:1008:C:H5''	11:H:16:ARG:HH12	1.63	0.63
15:L:164:THR:CG2	15:L:165:SER:N	2.57	0.63
1:0:660:A:H4'	1:0:661:G:O5'	1.98	0.63
1:0:814:G:H4'	37:0:9624:HOH:O	1.98	0.63
1:0:1130:U:H5'	37:0:7046:HOH:O	1.98	0.63
8:E:84:MET:HE2	8:E:133:VAL:CG2	2.28	0.63
20:Q:39:THR:HB	20:Q:42:GLU:CD	2.23	0.63
24:U:12:THR:CG2	24:U:15:GLU:HG3	2.19	0.63
25:V:84:VAL:HG12	37:V:6679:HOH:O	1.98	0.63
1:0:506:G:N2	1:0:508:A:H3'	2.13	0.63
1:0:1535:G:H2'	1:0:1536:C:C6	2.33	0.63
12:I:19:MET:HE1	12:I:78:ILE:HG22	1.81	0.63
16:M:22:GLN:HG2	16:M:26:LEU:HD22	1.79	0.63
26:W:85:VAL:HG12	26:W:86:GLU:N	2.12	0.63
30:1:40:ARG:HA	30:1:45:ASN:ND2	2.13	0.63
1:0:1213:C:O2'	1:0:1214:G:H5'	1.98	0.63
9:F:99:THR:HG23	9:F:99:THR:O	1.99	0.63
12:I:45:VAL:HG23	12:I:130:VAL:O	1.98	0.63
14:K:53:ARG:NH2	14:K:57:VAL:HG12	2.14	0.63
16:M:61:ALA:HB3	16:M:88:ALA:HB2	1.80	0.63
23:T:47:ARG:HG3	37:T:4381:HOH:O	1.97	0.63
25:V:5:VAL:HG22	25:V:32:CYS:HB2	1.78	0.63
26:W:78:GLU:HG2	26:W:79:GLU:N	2.12	0.63
1:0:67:A:H5''	1:0:69:A:C8	2.34	0.63
1:0:656:G:H1'	37:0:6667:HOH:O	1.98	0.63
1:0:962:C:H5''	37:0:4368:HOH:O	1.98	0.63
1:0:2361:A:H2'	1:0:2362:A:C8	2.33	0.63
11:H:33:MET:HB2	11:H:83:PHE:HB3	1.81	0.63
29:Z:12:ASN:HB3	37:Z:8449:HOH:O	1.97	0.63
1:0:1205:U:C2'	1:0:1206:U:H5''	2.29	0.63
11:H:55:GLN:NE2	11:H:124:ARG:HE	1.92	0.63

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:O:3245:HOH:O	22:S:9:LYS:HB2	1.99	0.63
2:9:3006:C:C5'	16:M:37:ARG:NH1	2.61	0.63
8:E:116:THR:HG22	8:E:151:LEU:HD22	1.80	0.63
12:I:93:ARG:HB3	12:I:93:ARG:NH1	2.12	0.63
17:N:32:ARG:O	17:N:32:ARG:HD3	1.97	0.63
1:O:2816:A:H3'	37:O:5166:HOH:O	1.99	0.63
4:A:170:VAL:HG22	28:Y:22:ILE:HG23	1.79	0.63
6:C:76:ARG:HD2	37:C:8430:HOH:O	1.99	0.63
15:L:104:ARG:O	15:L:108:LYS:HE2	1.99	0.63
18:O:13:VAL:HG11	18:O:40:VAL:HG11	1.81	0.63
20:Q:17:MET:HE1	20:Q:19:ARG:NH2	2.14	0.63
24:U:8:ILE:HA	24:U:11:MET:HE3	1.81	0.63
28:Y:42:CYS:SG	28:Y:44:PHE:HB2	2.38	0.63
30:1:40:ARG:HH11	30:1:40:ARG:HG2	1.64	0.63
1:O:684:G:H5''	37:O:3542:HOH:O	1.99	0.63
1:O:1205:U:H2'	1:O:1206:U:H5''	1.81	0.63
5:B:162:MET:HG3	5:B:310:ARG:HD3	1.80	0.63
10:G:71:LEU:C	10:G:73:ASP:H	2.07	0.63
15:L:64:ARG:HD2	37:L:8584:HOH:O	1.99	0.63
22:S:106:GLU:HG3	37:S:4913:HOH:O	1.98	0.63
1:O:513:A:N3	37:O:3150:HOH:O	2.31	0.62
1:O:1206:U:H5'	1:O:1206:U:C6	2.34	0.62
1:O:1874:U:H2'	4:A:120:ARG:HG3	1.78	0.62
37:O:4425:HOH:O	11:H:57:ARG:HG3	1.99	0.62
6:C:104:ASP:HA	6:C:107:ARG:HH12	1.63	0.62
7:D:95:THR:C	7:D:97:GLN:H	2.07	0.62
7:D:163:VAL:HA	37:D:6326:HOH:O	1.97	0.62
20:Q:132:ARG:HG2	20:Q:133:ALA:N	2.14	0.62
22:S:32:ARG:NH1	22:S:38:ARG:NH1	2.47	0.62
25:V:1:MET:HB2	25:V:103:GLU:HG2	1.81	0.62
1:O:2254:G:H1'	37:O:4974:HOH:O	1.98	0.62
5:B:145:HIS:HD2	5:B:146:THR:O	1.81	0.62
26:W:25:ARG:HD3	26:W:64:ALA:O	1.99	0.62
1:O:638:C:H2'	1:O:639:A:C8	2.34	0.62
1:O:2276:U:H2'	1:O:2277:U:H6	1.63	0.62
4:A:100:PRO:HG2	4:A:103:VAL:HG21	1.79	0.62
1:O:20:G:H21	20:Q:117:HIS:HD2	1.47	0.62
1:O:2547:C:OP2	5:B:5:ARG:NH1	2.32	0.62
7:D:25:MET:HE1	7:D:37:ALA:O	1.98	0.62
21:R:51:GLN:HE21	21:R:53:ASN:ND2	1.95	0.62
1:O:558:C:H2'	1:O:559:U:C5'	2.29	0.62

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:1596:U:H2'	1:O:1598:A:OP2	1.99	0.62
7:D:97:GLN:HG2	7:D:97:GLN:O	1.99	0.62
11:H:49:VAL:O	11:H:157:ILE:HG23	2.00	0.62
12:I:75:PRO:HG2	12:I:105:LEU:HD21	1.81	0.62
16:M:71:TRP:CE3	16:M:175:LEU:HD22	2.35	0.62
20:Q:29:LYS:HB3	37:Q:8534:HOH:O	2.00	0.62
22:S:71:VAL:HG11	22:S:90:PRO:CB	2.24	0.62
23:T:47:ARG:CG	37:T:4381:HOH:O	2.48	0.62
27:X:130:ARG:HB2	27:X:142:SER:O	2.00	0.62
29:Z:25:LYS:O	29:Z:25:LYS:HG2	1.99	0.62
1:O:213:G:H1'	1:O:214:U:OP2	1.99	0.62
1:O:2533:C:H5'	1:O:2533:C:C6	2.34	0.62
37:A:8606:HOH:O	28:Y:75:ALA:HB3	1.99	0.62
7:D:140:ARG:O	7:D:144:ARG:HG2	2.00	0.62
15:L:39:ARG:HA	15:L:63:VAL:HG22	1.82	0.62
15:L:74:ARG:HG3	15:L:74:ARG:NH1	2.11	0.62
1:O:1733:A:H4'	5:B:212:GLN:HA	1.80	0.62
5:B:82:VAL:O	5:B:82:VAL:HG12	1.98	0.62
6:C:233:THR:HG22	6:C:234:VAL:N	2.14	0.62
7:D:50:VAL:O	7:D:71:ALA:HA	2.00	0.62
18:O:38:GLU:HA	18:O:41:ARG:NH1	2.15	0.62
28:Y:61:GLY:HA3	37:Y:8423:HOH:O	1.99	0.62
1:O:871:G:C8	1:O:871:G:C5'	2.76	0.62
1:O:1187:U:O2'	1:O:1189:A:H2	1.83	0.62
37:O:3138:HOH:O	17:N:3:THR:HG21	2.00	0.62
8:E:101:GLU:HB2	8:E:116:THR:O	1.99	0.62
15:L:61:ILE:HD12	15:L:61:ILE:N	2.14	0.62
16:M:152:GLU:C	16:M:154:LEU:H	2.06	0.62
21:R:37:VAL:O	21:R:41:VAL:HG23	2.00	0.62
28:Y:57:CYS:SG	28:Y:59:HIS:HB3	2.40	0.62
1:O:2637:A:C5'	1:O:2638:G:H5'	2.29	0.62
6:C:115:LEU:O	6:C:118:THR:HB	2.00	0.62
9:F:58:GLU:HG3	9:F:61:MET:HE2	1.81	0.62
11:H:26:LYS:HG2	11:H:28:ILE:N	2.15	0.62
11:H:75:SER:O	11:H:79:ALA:HB2	2.00	0.62
16:M:180:LEU:O	16:M:181:ASP:HB3	2.00	0.62
16:M:184:ILE:HG22	16:M:185:GLU:HG3	1.80	0.62
31:2:11:CYS:HB2	31:2:20:HIS:CE1	2.35	0.62
2:9:3114:G:O6	16:M:11:ARG:HD3	2.00	0.62
12:I:19:MET:HE3	12:I:132:LEU:CD1	2.19	0.62
14:K:90:ARG:NH2	14:K:121:ILE:HD11	2.15	0.62

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:M:12:ARG:HD3	16:M:18:THR:OG1	1.99	0.62
1:0:189:A:OP1	15:L:171:ARG:NH2	2.33	0.61
1:0:1450:C:O2'	1:0:1494:A:H5'	1.99	0.61
1:0:1634:G:H3'	37:0:3377:HOH:O	1.97	0.61
1:0:1688:G:H4'	29:Z:8:GLN:HG3	1.81	0.61
6:C:25:PRO:HA	37:C:8356:HOH:O	2.00	0.61
6:C:129:HIS:HE1	6:C:231:ARG:HA	1.63	0.61
16:M:154:LEU:HD11	16:M:157:PRO:HA	1.81	0.61
17:N:49:GLU:HG2	37:N:5191:HOH:O	2.00	0.61
18:O:16:VAL:HG12	18:O:17:GLY:N	2.15	0.61
28:Y:62:TYR:CE2	28:Y:64:ILE:HG23	2.35	0.61
1:0:777:U:HO2'	29:Z:11:LYS:HG2	1.65	0.61
1:0:820:G:H5'	1:0:821:U:H5'	1.81	0.61
1:0:2001:G:O2'	1:0:2002:C:H5'	2.00	0.61
37:0:5867:HOH:O	5:B:27:ASN:HB3	1.99	0.61
16:M:164:ASP:OD2	16:M:167:ASP:HA	1.99	0.61
1:0:263:U:O4'	9:F:59:ILE:HD13	2.00	0.61
1:0:280:C:H2'	1:0:281:U:O4'	2.00	0.61
1:0:281:U:H2'	1:0:282:C:O4'	1.99	0.61
4:A:131:HIS:O	4:A:132:ASP:HB2	1.99	0.61
6:C:1:MET:HG2	6:C:2:GLN:NE2	2.15	0.61
7:D:99:ASP:CB	7:D:103:ASN:H	2.13	0.61
1:0:558:C:C2'	1:0:559:U:H5''	2.31	0.61
1:0:1657:A:H2'	1:0:1658:A:C8	2.34	0.61
1:0:2768:A:O2'	1:0:2769:C:H5'	2.01	0.61
2:9:3043:G:H5'	37:9:8410:HOH:O	2.00	0.61
25:V:13:MET:HE2	25:V:18:GLN:N	2.14	0.61
1:0:1636:G:O2'	1:0:1637:A:H5'	2.00	0.61
1:0:2320:U:H4'	1:0:2321:A:O4'	2.00	0.61
1:0:2578:G:H5'	1:0:2578:G:H8	1.65	0.61
37:0:5695:HOH:O	27:X:158:LYS:HD3	1.99	0.61
4:A:129:LEU:CD1	4:A:145:MET:HE1	2.29	0.61
13:J:14:LYS:HB2	13:J:45:PRO:CG	2.25	0.61
16:M:119:GLN:O	16:M:123:ILE:HG13	2.00	0.61
26:W:15:ARG:HH11	26:W:15:ARG:CB	2.13	0.61
1:0:514:G:O5'	1:0:514:G:H8	1.84	0.61
1:0:1418:U:OP1	30:1:42:TRP:HB3	2.00	0.61
1:0:1909:A:N1	1:0:2128:G:H1'	2.16	0.61
4:A:211:LYS:NZ	37:A:8565:HOH:O	2.33	0.61
6:C:163:HIS:HD2	37:C:8435:HOH:O	1.82	0.61
15:L:30:GLU:O	15:L:34:GLU:HG3	1.99	0.61

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:L:155:HIS:CE1	15:L:158:ARG:HE	2.17	0.61
25:V:65:VAL:HA	25:V:68:THR:HG22	1.82	0.61
1:O:1711:A:O2'	1:O:1712:A:H5'	2.00	0.61
37:O:4957:HOH:O	5:B:298:LYS:HD3	2.00	0.61
2:9:3029:C:H2'	2:9:3030:C:H5'	1.82	0.61
6:C:78:ARG:HG3	6:C:78:ARG:NH1	2.16	0.61
7:D:69:ILE:HG22	7:D:69:ILE:O	1.99	0.61
1:O:2415:A:N3	16:M:26:LEU:HD13	2.16	0.61
4:A:109:GLU:HG2	4:A:116:GLY:N	2.15	0.61
11:H:26:LYS:CG	11:H:28:ILE:H	2.14	0.61
15:L:87:MET:HB2	15:L:91:ILE:CD1	2.28	0.61
1:O:1751:G:C2'	1:O:1752:G:H5''	2.29	0.61
37:O:6164:HOH:O	16:M:4:PRO:HD2	2.00	0.61
15:L:37:VAL:HG21	15:L:108:LYS:CG	2.29	0.61
24:U:56:ILE:O	24:U:60:GLN:HG3	2.01	0.61
26:W:20:GLU:CD	26:W:21:PRO:HD2	2.26	0.61
27:X:189:ASN:HB2	37:X:8525:HOH:O	2.00	0.61
1:O:902:G:N7	14:K:18:HIS:CD2	2.69	0.61
11:H:150:LYS:HG2	37:H:8368:HOH:O	1.99	0.61
22:S:47:THR:HB	22:S:100:ASP:HB3	1.83	0.61
1:O:1299:G:O6	14:K:6:ARG:HD3	2.01	0.60
1:O:1641:A:H2'	1:O:1642:A:H5'	1.82	0.60
1:O:1654:U:H2'	4:A:47:HIS:HD2	1.66	0.60
1:O:2786:G:H2'	37:O:6575:HOH:O	2.00	0.60
7:D:22:VAL:HG22	7:D:74:THR:HG22	1.81	0.60
11:H:58:HIS:HA	11:H:61:LEU:HD23	1.83	0.60
26:W:43:VAL:HG12	26:W:44:ASP:N	2.15	0.60
1:O:539:G:H2'	1:O:540:A:C8	2.36	0.60
1:O:1759:A:N7	37:O:9064:HOH:O	2.31	0.60
18:O:27:ARG:O	18:O:31:ILE:HG13	2.00	0.60
23:T:52:THR:CG2	23:T:54:THR:HB	2.31	0.60
26:W:18:ARG:NH1	37:W:4132:HOH:O	2.26	0.60
27:X:189:ASN:ND2	27:X:192:ASP:H	1.99	0.60
1:O:1741:U:O2'	1:O:2723:G:H4'	2.01	0.60
2:9:3114:G:H2'	2:9:3115:C:C6	2.37	0.60
5:B:162:MET:HE2	5:B:310:ARG:CD	2.30	0.60
6:C:140:VAL:HB	37:C:8446:HOH:O	2.01	0.60
7:D:136:ARG:HD2	7:D:155:HIS:O	2.01	0.60
9:F:21:GLU:O	9:F:24:ARG:HG3	2.01	0.60
9:F:110:GLU:O	9:F:114:LYS:HG3	2.01	0.60
11:H:26:LYS:HD2	11:H:28:ILE:CG1	2.30	0.60

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:J:109:LEU:HD13	13:J:113:ILE:HD11	1.82	0.60
1:O:1441:G:O2'	1:O:1442:A:H5'	2.00	0.60
1:O:2346:C:H6	1:O:2346:C:O5'	1.85	0.60
5:B:146:THR:O	5:B:159:PRO:HB3	2.00	0.60
8:E:31:ARG:HH12	8:E:68:HIS:CE1	2.19	0.60
9:F:58:GLU:OE1	15:L:27:ARG:NH2	2.30	0.60
11:H:166:ASN:HD22	11:H:166:ASN:N	1.99	0.60
26:W:9:VAL:HG13	26:W:88:GLU:OE2	2.01	0.60
1:O:1268:C:H2'	1:O:1269:G:H8	1.67	0.60
1:O:1278:A:H4'	1:O:1279:U:C4	2.37	0.60
4:A:105:VAL:HG11	4:A:154:ALA:HB1	1.82	0.60
11:H:26:LYS:CD	11:H:28:ILE:HB	2.31	0.60
11:H:57:ARG:HG3	11:H:57:ARG:HH11	1.67	0.60
17:N:21:SER:OG	17:N:106:PRO:HB2	2.02	0.60
26:W:85:VAL:HG12	26:W:86:GLU:H	1.67	0.60
1:O:1353:C:P	37:O:4134:HOH:O	2.58	0.60
1:O:2755:G:H1'	37:O:4138:HOH:O	2.01	0.60
7:D:23:VAL:HG22	7:D:73:VAL:HB	1.82	0.60
7:D:51:ARG:HD3	37:D:7636:HOH:O	2.01	0.60
1:O:282:C:O2'	1:O:283:U:H5'	2.01	0.60
1:O:793:A:H5''	18:O:83:LYS:HG2	1.83	0.60
6:C:35:VAL:HG21	6:C:227:GLY:HA2	1.82	0.60
21:R:32:ALA:HA	21:R:36:GLU:OE1	2.01	0.60
1:O:2748:G:H5'	37:O:6698:HOH:O	2.00	0.60
13:J:32:ILE:HD11	13:J:56:SER:HB3	1.83	0.60
20:Q:18:LEU:HD12	20:Q:143:VAL:HG11	1.83	0.60
25:V:65:VAL:HG12	25:V:116:LEU:HD13	1.84	0.60
28:Y:19:GLY:O	28:Y:23:ARG:HG2	2.02	0.60
1:O:661:G:C5	1:O:686:A:C2	2.89	0.60
1:O:1130:U:H2'	1:O:1131:G:O4'	2.01	0.60
1:O:2304:G:H5'	37:O:9861:HOH:O	2.01	0.60
37:O:5059:HOH:O	5:B:2:GLN:HG3	2.02	0.60
4:A:16:PHE:HB3	37:A:8548:HOH:O	2.02	0.60
7:D:135:VAL:HG21	7:D:139:TYR:CD1	2.37	0.60
9:F:37:THR:O	9:F:41:GLU:HG3	2.02	0.60
27:X:189:ASN:HD22	27:X:189:ASN:C	2.10	0.60
1:O:1667:A:H5'	1:O:1667:A:C8	2.34	0.60
1:O:2769:C:O2'	1:O:2770:G:H5'	2.02	0.60
2:9:3039:U:H1'	2:9:3044:A:N6	2.16	0.60
7:D:146:LYS:HZ1	16:M:107:ASN:HD21	1.47	0.60
9:F:100:ASP:O	9:F:101:ALA:O	2.20	0.60

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:F:107:VAL:HG23	37:F:6617:HOH:O	2.01	0.60
17:N:47:ARG:HB2	37:N:6739:HOH:O	2.01	0.60
1:0:564:G:H1'	37:0:5720:HOH:O	2.01	0.59
1:0:2443:C:H3'	37:0:9959:HOH:O	2.00	0.59
4:A:95:PRO:HG2	4:A:98:GLU:HG2	1.84	0.59
16:M:34:LEU:HD13	16:M:47:LEU:HD21	1.83	0.59
16:M:83:LEU:HD13	16:M:175:LEU:HD23	1.84	0.59
25:V:141:HIS:HB2	25:V:146:ILE:HG12	1.84	0.59
27:X:99:ALA:HB2	27:X:233:TYR:CE2	2.37	0.59
28:Y:13:ARG:NH1	28:Y:14:PHE:CE2	2.70	0.59
4:A:94:LEU:HG	4:A:99:ILE:HD11	1.85	0.59
4:A:94:LEU:HD23	4:A:94:LEU:N	2.17	0.59
4:A:191:GLY:HA2	4:A:194:MET:HE2	1.83	0.59
22:S:48:VAL:CG2	22:S:98:VAL:HA	2.32	0.59
25:V:137:GLN:HE21	25:V:141:HIS:HE1	1.48	0.59
26:W:37:LEU:CD1	26:W:85:VAL:HG21	2.31	0.59
1:0:1118:A:C8	1:0:1118:A:C3'	2.84	0.59
1:0:1477:C:H5'	1:0:1868:G:H5''	1.82	0.59
6:C:127:ARG:HG2	6:C:127:ARG:HH11	1.66	0.59
12:I:17:CYS:HA	12:I:119:THR:O	2.01	0.59
17:N:44:ASN:OD1	17:N:65:LEU:HB2	2.03	0.59
21:R:53:ASN:ND2	37:R:2190:HOH:O	2.34	0.59
1:0:349:U:O2'	1:0:350:C:H5'	2.02	0.59
1:0:558:C:H2'	1:0:559:U:H5'	1.84	0.59
1:0:657:G:H2'	1:0:658:C:H6	1.65	0.59
1:0:1675:C:H3'	37:0:7180:HOH:O	2.03	0.59
22:S:38:ARG:HH11	22:S:38:ARG:HG3	1.67	0.59
2:9:3022:G:O2'	2:9:3024:U:H5'	2.02	0.59
11:H:86:ARG:NH1	11:H:130:HIS:CD2	2.70	0.59
12:I:80:LYS:HE2	12:I:98:PHE:CZ	2.37	0.59
1:0:1191:A:C3'	1:0:1192:A:H5''	2.32	0.59
5:B:24:PRO:O	5:B:25:ARG:HD3	2.02	0.59
15:L:138:HIS:ND1	15:L:139:PRO:O	2.29	0.59
24:U:64:GLY:O	24:U:65:ASP:HB2	2.02	0.59
26:W:14:LEU:HD12	26:W:67:PRO:O	2.02	0.59
29:Z:25:LYS:HE2	37:1:7213:HOH:O	2.02	0.59
1:0:88:G:H8	1:0:88:G:H5'	1.68	0.59
1:0:2694:A:H4'	8:E:91:PHE:CE1	2.37	0.59
5:B:258:GLY:H	5:B:260:HIS:CE1	2.20	0.59
7:D:95:THR:O	7:D:97:GLN:N	2.29	0.59
28:Y:73:THR:O	28:Y:74:VAL:C	2.46	0.59

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:328:U:O4'	6:C:202:THR:HG22	2.03	0.59
1:0:645:U:OP2	14:K:4:LYS:HE2	2.03	0.59
1:0:797:A:H4'	28:Y:10:ARG:N	2.17	0.59
1:0:2420:G:O2'	1:0:2421:G:H5'	2.03	0.59
4:A:194:MET:HE3	4:A:199:HIS:HB2	1.84	0.59
11:H:75:SER:C	11:H:79:ALA:HB2	2.28	0.59
16:M:93:GLN:HG2	37:M:6239:HOH:O	2.02	0.59
1:0:553:G:P	27:X:204:ARG:HH22	2.25	0.59
11:H:27:LYS:H	11:H:58:HIS:CD2	2.14	0.59
20:Q:47:LEU:O	20:Q:51:ILE:HG13	2.02	0.59
22:S:26:THR:HA	22:S:39:ASN:HB3	1.85	0.59
25:V:38:THR:O	25:V:42:ARG:HB2	2.02	0.59
1:0:657:G:H2'	1:0:658:C:C6	2.38	0.59
1:0:820:G:O2'	1:0:856:G:H4'	2.03	0.59
1:0:2783:A:H3'	37:0:4676:HOH:O	2.02	0.59
5:B:56:ASP:OD1	5:B:322:ARG:HB3	2.03	0.59
5:B:279:THR:OG1	5:B:290:VAL:HB	2.03	0.59
12:I:39:VAL:HG12	12:I:40:ASN:ND2	2.18	0.59
25:V:65:VAL:CG1	25:V:116:LEU:HD13	2.33	0.59
28:Y:53:GLY:HA2	28:Y:67:GLY:O	2.03	0.59
1:0:100:C:H4'	22:S:16:LEU:HB2	1.85	0.58
1:0:1015:C:H2'	1:0:1016:U:C6	2.38	0.58
1:0:2769:C:C2'	1:0:2770:G:H5'	2.33	0.58
6:C:214:THR:HG21	37:C:8399:HOH:O	2.03	0.58
12:I:52:GLN:HG3	12:I:53:ILE:H	1.67	0.58
16:M:115:VAL:HG22	37:M:5851:HOH:O	2.03	0.58
1:0:284:C:H4'	1:0:285:A:O5'	2.02	0.58
2:9:3078:G:N2	2:9:3103:A:OP2	2.33	0.58
6:C:95:GLU:HG3	37:C:8469:HOH:O	2.03	0.58
11:H:85:ILE:HB	11:H:132:PHE:CE2	2.38	0.58
12:I:47:THR:CB	37:I:3661:HOH:O	2.51	0.58
17:N:41:ALA:HA	37:N:5104:HOH:O	2.03	0.58
28:Y:28:ASP:O	28:Y:31:ILE:HG22	2.03	0.58
1:0:2363:G:O2'	19:P:11:ARG:HG3	2.03	0.58
37:0:4002:HOH:O	11:H:151:MET:HE2	2.03	0.58
4:A:36:ASP:HA	4:A:83:GLY:HA3	1.86	0.58
11:H:65:ARG:HB3	37:H:8370:HOH:O	2.01	0.58
11:H:123:ALA:HA	37:H:8310:HOH:O	2.02	0.58
15:L:52:LEU:HD13	15:L:116:ASN:HB3	1.85	0.58
24:U:39:ALA:C	24:U:41:GLU:H	2.11	0.58
1:0:157:G:H4'	15:L:95:LYS:CE	2.32	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:1393:A:H2'	1:0:1394:C:C6	2.38	0.58
1:0:1589:G:H22	1:0:1605:G:H1'	1.67	0.58
1:0:2435:U:H1'	37:0:4868:HOH:O	2.03	0.58
19:P:25:PRO:HB2	37:P:4350:HOH:O	2.02	0.58
21:R:6:LYS:HB2	21:R:27:ALA:O	2.02	0.58
22:S:63:ILE:HD11	22:S:75:GLU:HB2	1.84	0.58
28:Y:33:HIS:HE1	28:Y:49:ARG:NE	2.02	0.58
1:0:1003:U:O2	11:H:90:PHE:HZ	1.85	0.58
4:A:192:VAL:CG1	4:A:207:GLN:HB3	2.31	0.58
6:C:27:ARG:HG3	6:C:29:ASP:OD1	2.03	0.58
16:M:78:MET:HB2	16:M:79:PRO:HD3	1.85	0.58
25:V:80:ASP:O	25:V:84:VAL:HG23	2.02	0.58
28:Y:29:VAL:O	28:Y:33:HIS:HB2	2.04	0.58
1:0:346:U:H4'	37:0:6234:HOH:O	2.02	0.58
1:0:1348:A:H3'	37:0:3275:HOH:O	2.02	0.58
1:0:2241:C:O2'	1:0:2242:U:H5'	2.04	0.58
1:0:2468:A:H61	31:2:48:ASN:HD21	1.52	0.58
1:0:2672:C:H1'	37:B:8633:HOH:O	2.04	0.58
37:0:3945:HOH:O	18:O:37:ARG:HB2	2.03	0.58
10:G:64:ASN:HD22	10:G:64:ASN:N	2.02	0.58
15:L:52:LEU:HD21	37:L:8617:HOH:O	2.02	0.58
17:N:113:VAL:O	17:N:114:ILE:HD13	2.04	0.58
27:X:155:ARG:NH1	37:X:8558:HOH:O	2.37	0.58
28:Y:50:ALA:HB3	28:Y:54:ILE:HG22	1.86	0.58
1:0:542:A:H1'	37:0:4132:HOH:O	2.03	0.58
14:K:133:VAL:HB	37:K:8557:HOH:O	2.03	0.58
20:Q:39:THR:CG2	20:Q:42:GLU:HG3	2.34	0.58
23:T:6:CYS:C	23:T:8:TYR:H	2.12	0.58
25:V:88:THR:HG23	25:V:110:GLN:HB3	1.84	0.58
1:0:1003:U:O2	11:H:90:PHE:CZ	2.57	0.58
1:0:1766:U:O2	1:0:1778:A:H5'	2.04	0.58
11:H:139:ASP:HB2	37:H:8334:HOH:O	2.03	0.58
17:N:73:ASP:HA	17:N:92:VAL:O	2.03	0.58
20:Q:39:THR:HB	20:Q:42:GLU:CG	2.33	0.58
30:1:41:HIS:H	30:1:45:ASN:ND2	2.00	0.58
1:0:31:C:OP2	22:S:8:ARG:HD2	2.04	0.58
1:0:656:G:OP2	17:N:37:ARG:HD2	2.04	0.58
1:0:1667:A:H2'	1:0:1668:U:C6	2.38	0.58
1:0:2694:A:H4'	8:E:91:PHE:HE1	1.69	0.58
1:0:2904:U:H4'	26:W:8:ARG:NH1	2.19	0.58
8:E:86:VAL:CG1	8:E:129:GLU:HA	2.34	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:L:106:ASN:ND2	35:L:8518:CL:CL	2.74	0.58
19:P:93:ARG:HH11	19:P:93:ARG:HG3	1.68	0.58
25:V:6:GLN:HB2	25:V:26:ILE:CD1	2.31	0.58
1:O:1185:U:H5'	37:O:6843:HOH:O	2.03	0.58
4:A:109:GLU:HG2	4:A:116:GLY:H	1.68	0.58
5:B:175:LEU:C	5:B:175:LEU:CD2	2.77	0.58
6:C:214:THR:HG23	37:C:8434:HOH:O	2.04	0.58
8:E:32:ARG:O	8:E:33:LEU:HD23	2.03	0.58
11:H:150:LYS:HB2	11:H:157:ILE:HD12	1.86	0.58
15:L:137:ASP:C	15:L:142:LYS:HE3	2.29	0.58
16:M:71:TRP:HE3	16:M:175:LEU:HD22	1.68	0.58
23:T:17:THR:HG22	23:T:18:GLY:N	2.19	0.58
28:Y:46:LYS:O	28:Y:57:CYS:HA	2.04	0.58
1:O:1687:C:O2	29:Z:9:GLY:HA2	2.04	0.57
5:B:315:VAL:HG23	5:B:316:ARG:HG2	1.86	0.57
14:K:148:GLU:HG3	37:K:8552:HOH:O	2.03	0.57
16:M:24:LEU:O	16:M:28:LYS:HG2	2.04	0.57
1:O:1134:G:C5'	11:H:151:MET:HE1	2.34	0.57
1:O:1942:A:O2'	1:O:1943:C:H5'	2.05	0.57
1:O:2365:G:H4'	19:P:45:PRO:O	2.03	0.57
13:J:32:ILE:CD1	13:J:56:SER:HB3	2.34	0.57
19:P:26:PRO:HG3	37:P:2847:HOH:O	2.05	0.57
1:O:547:A:H3'	37:O:4398:HOH:O	2.05	0.57
1:O:2326:U:H4'	1:O:2412:G:C4'	2.35	0.57
1:O:2472:C:O2'	1:O:2634:G:H4'	2.04	0.57
37:O:6796:HOH:O	15:L:154:ARG:HD3	2.02	0.57
4:A:153:ARG:HH11	4:A:153:ARG:CB	2.17	0.57
15:L:133:LEU:O	15:L:134:ILE:HD13	2.03	0.57
28:Y:38:LYS:HE2	28:Y:45:LYS:CE	2.33	0.57
1:O:1377:C:H5'	1:O:1377:C:H6	1.68	0.57
1:O:1422:U:H2'	1:O:1423:C:C6	2.39	0.57
2:9:3004:G:O2'	16:M:44:ARG:NH2	2.38	0.57
2:9:3044:A:O4'	7:D:76:ARG:NE	2.38	0.57
7:D:170:TYR:O	7:D:171:ASP:HB3	2.04	0.57
11:H:147:ARG:HA	11:H:150:LYS:HZ2	1.70	0.57
17:N:4:ASN:HB3	17:N:7:LEU:HB3	1.86	0.57
22:S:61:GLU:HG3	37:S:3851:HOH:O	2.04	0.57
30:1:22:PRO:HG2	30:1:25:VAL:HG23	1.86	0.57
1:O:447:A:OP1	22:S:2:LYS:HG2	2.04	0.57
1:O:1127:C:H2'	1:O:1128:U:H5'	1.86	0.57
1:O:1919:A:H4'	37:O:4303:HOH:O	2.05	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:2501:G:H1'	37:0:4002:HOH:O	2.03	0.57
1:0:2769:C:H2'	1:0:2770:G:O4'	2.04	0.57
2:9:3069:U:OP1	16:M:4:PRO:HG3	2.05	0.57
4:A:105:VAL:CG1	4:A:154:ALA:HB1	2.35	0.57
4:A:135:VAL:HG21	4:A:147:ARG:NH1	2.18	0.57
8:E:93:MET:HE1	8:E:165:GLY:N	2.20	0.57
15:L:87:MET:HB3	31:2:46:ILE:HG21	1.86	0.57
1:0:775:G:H1'	37:0:8832:HOH:O	2.05	0.57
1:0:1058:A:H2'	1:0:1060:C:H5''	1.87	0.57
1:0:1132:A:N6	1:0:1229:C:H2'	2.20	0.57
1:0:1151:G:OP1	10:G:16:LYS:NZ	2.36	0.57
5:B:53:LEU:HD21	5:B:270:ILE:HD12	1.87	0.57
6:C:12:THR:HB	37:C:8437:HOH:O	2.05	0.57
6:C:236:THR:HA	37:C:8446:HOH:O	2.04	0.57
8:E:3:VAL:HG22	8:E:49:ILE:HB	1.86	0.57
12:I:107:ASN:HD21	12:I:109:TYR:HB2	1.70	0.57
15:L:48:ARG:NH2	37:L:8559:HOH:O	2.37	0.57
21:R:52:VAL:HG22	21:R:66:VAL:HG22	1.85	0.57
26:W:73:ARG:O	26:W:85:VAL:HG13	2.05	0.57
1:0:1995:G:O2'	1:0:1997:A:N7	2.38	0.57
1:0:2748:G:C5'	37:0:6698:HOH:O	2.52	0.57
1:0:2878:U:H2'	1:0:2879:A:O4'	2.04	0.57
5:B:75:GLU:C	5:B:77:PRO:HD3	2.30	0.57
5:B:280:VAL:CG1	5:B:334:SER:HA	2.35	0.57
6:C:2:GLN:HB3	37:C:8337:HOH:O	2.03	0.57
15:L:38:VAL:O	15:L:63:VAL:HG13	2.04	0.57
26:W:78:GLU:CG	26:W:79:GLU:H	2.16	0.57
1:0:512:G:O3'	1:0:513:A:H8	1.86	0.57
1:0:1407:A:O2'	1:0:1408:U:H3'	2.05	0.57
1:0:2032:U:O2'	1:0:2033:G:H5''	2.04	0.57
37:0:3045:HOH:O	15:L:152:ARG:HG3	2.04	0.57
4:A:190:ARG:NH2	4:A:207:GLN:OE1	2.37	0.57
6:C:236:THR:O	6:C:237:GLU:C	2.47	0.57
18:O:13:VAL:HG11	18:O:40:VAL:CG1	2.35	0.57
27:X:144:ARG:NH1	37:X:8577:HOH:O	2.36	0.57
1:0:136:C:H2'	1:0:137:U:O4'	2.04	0.57
1:0:1244:U:OP1	12:I:18:ILE:HD13	2.04	0.57
1:0:1653:A:N6	37:0:3737:HOH:O	2.36	0.57
1:0:1874:U:OP1	4:A:51:ARG:HD2	2.03	0.57
4:A:161:GLY:O	28:Y:68:CYS:SG	2.56	0.57
4:A:179:MET:HG2	4:A:186:TRP:CB	2.35	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:D:59:GLY:C	7:D:61:PHE:H	2.13	0.57
8:E:106:ASN:ND2	8:E:109:GLY:HA2	2.19	0.57
13:J:74:VAL:HG12	13:J:75:ARG:HG3	1.87	0.57
14:K:104:ASP:HB2	37:K:8576:HOH:O	2.04	0.57
14:K:122:ALA:HB3	14:K:125:PHE:CZ	2.40	0.57
14:K:149:ARG:O	14:K:150:GLN:HB2	2.05	0.57
1:O:182:G:O3'	15:L:157:LEU:CD1	2.53	0.57
1:O:1625:U:H5''	37:O:5443:HOH:O	2.05	0.57
37:O:6257:HOH:O	4:A:211:LYS:HD3	2.04	0.57
4:A:101:GLU:HG2	4:A:131:HIS:ND1	2.20	0.57
10:G:23:ILE:O	10:G:27:ILE:HG13	2.05	0.57
11:H:130:HIS:CD2	11:H:133:ILE:HD11	2.40	0.57
1:O:461:C:H2'	37:O:3478:HOH:O	2.04	0.56
1:O:629:A:C2	1:O:2074:A:C2	2.93	0.56
1:O:926:A:O2'	14:K:41:HIS:CD2	2.58	0.56
1:O:2269:C:C2'	1:O:2270:G:H5'	2.34	0.56
37:O:6266:HOH:O	15:L:178:LYS:HB2	2.05	0.56
37:O:6359:HOH:O	24:U:4:HIS:HB3	2.05	0.56
12:I:74:ARG:O	12:I:78:ILE:HG12	2.04	0.56
1:O:407:A:H2'	1:O:408:A:C8	2.40	0.56
2:9:3030:C:OP1	7:D:137:PRO:O	2.23	0.56
5:B:190:MET:HE2	5:B:194:PHE:HD1	1.66	0.56
17:N:79:VAL:HA	37:N:6810:HOH:O	2.05	0.56
1:O:625:U:H5'	37:O:9677:HOH:O	2.05	0.56
1:O:657:G:OP1	6:C:27:ARG:NH2	2.28	0.56
1:O:1477:C:O2'	1:O:1478:U:H5'	2.04	0.56
1:O:1666:C:C2'	1:O:1667:A:H5'	2.35	0.56
1:O:2612:A:H4'	37:O:3173:HOH:O	2.05	0.56
37:O:5090:HOH:O	22:S:68:ASP:HB2	2.05	0.56
5:B:42:ALA:HB1	5:B:308:LEU:HD11	1.85	0.56
5:B:305:ASP:O	5:B:306:LYS:HB2	2.05	0.56
10:G:12:ILE:N	10:G:13:PRO:CD	2.68	0.56
11:H:83:PHE:HZ	11:H:146:TRP:HE1	1.50	0.56
11:H:139:ASP:N	11:H:140:PRO:CD	2.67	0.56
18:O:94:TRP:CZ2	18:O:98:ILE:HG13	2.41	0.56
22:S:23:VAL:C	22:S:93:THR:HG21	2.31	0.56
1:O:450:C:OP1	6:C:184:ARG:NH2	2.36	0.56
1:O:2463:A:H4'	1:O:2464:C:OP2	2.06	0.56
6:C:236:THR:CG2	6:C:239:ALA:H	1.94	0.56
6:C:246:ARG:HB3	6:C:246:ARG:NH1	2.20	0.56
8:E:107:PHE:CD2	8:E:108:LEU:HD13	2.41	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:H:5:MET:N	37:H:8351:HOH:O	2.37	0.56
25:V:85:ALA:HB2	25:V:91:ASP:O	2.05	0.56
1:0:926:A:O2'	14:K:41:HIS:HD2	1.88	0.56
1:0:2415:A:C2	16:M:25:ARG:HB3	2.41	0.56
1:0:2419:U:H5''	1:0:2420:G:H5'	1.87	0.56
6:C:235:PHE:HE2	6:C:243:VAL:HG21	1.71	0.56
13:J:75:ARG:CZ	37:J:4172:HOH:O	2.53	0.56
15:L:12:TRP:O	15:L:15:PRO:HD3	2.06	0.56
1:0:825:U:H5''	1:0:826:U:OP1	2.06	0.56
1:0:1543:G:N1	1:0:1641:A:OP2	2.34	0.56
4:A:1:GLY:HA2	4:A:197:VAL:HG23	1.87	0.56
13:J:55:VAL:HG12	13:J:56:SER:H	1.70	0.56
1:0:820:G:C6	4:A:171:LYS:HB2	2.40	0.56
1:0:1834:C:H2'	1:0:1840:A:H62	1.70	0.56
1:0:2379:G:H4'	1:0:2380:A:H5''	1.88	0.56
4:A:37:VAL:HG22	37:A:8590:HOH:O	2.06	0.56
11:H:81:TYR:CD1	11:H:81:TYR:C	2.83	0.56
37:O:9108:HOH:O	15:L:165:SER:HB3	2.06	0.56
2:9:3048:C:H4'	16:M:141:ARG:NH2	2.20	0.56
5:B:36:PRO:HG3	5:B:168:GLY:HA3	1.88	0.56
5:B:320:GLN:HG3	5:B:321:PRO:HD2	1.88	0.56
7:D:65:GLU:HG3	37:D:6752:HOH:O	2.06	0.56
8:E:7:ILE:HD11	8:E:11:VAL:O	2.05	0.56
22:S:55:PHE:CD2	22:S:77:VAL:HG13	2.41	0.56
1:0:681:G:N3	1:0:681:G:H5'	2.21	0.56
7:D:21:VAL:HG13	7:D:131:THR:O	2.06	0.56
12:I:45:VAL:HG22	12:I:46:ILE:N	2.20	0.56
16:M:87:LEU:CD1	16:M:186:LEU:HD21	2.35	0.56
27:X:187:VAL:CG2	27:X:192:ASP:HB2	2.29	0.56
28:Y:77:LYS:HA	28:Y:80:MET:HE2	1.88	0.56
1:0:1659:A:H2'	1:0:1660:G:O4'	2.05	0.56
1:0:2256:G:O2'	1:0:2257:G:H5'	2.05	0.56
4:A:57:ALA:HA	4:A:67:LEU:HD23	1.87	0.56
11:H:142:VAL:HG13	37:H:8366:HOH:O	2.05	0.56
11:H:163:PRO:O	11:H:164:ALA:HB2	2.06	0.56
16:M:132:ASN:O	16:M:135:VAL:HG12	2.06	0.56
17:N:96:VAL:HG12	17:N:97:SER:O	2.06	0.56
25:V:60:GLU:O	25:V:63:GLU:HB2	2.06	0.56
1:0:125:U:H2'	37:O:3255:HOH:O	2.05	0.55
1:0:935:G:H1'	37:O:6862:HOH:O	2.06	0.55
1:0:1617:C:C4	1:0:1643:C:H4'	2.41	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:1942:A:H3'	37:0:6726:HOH:O	2.06	0.55
1:0:1979:G:H2'	37:0:9786:HOH:O	2.06	0.55
1:0:2083:A:N1	37:0:4588:HOH:O	2.33	0.55
7:D:11:HIS:O	7:D:12:GLU:HB3	2.06	0.55
11:H:47:GLU:HG2	11:H:133:ILE:HD12	1.86	0.55
16:M:48:VAL:HG11	16:M:55:ASP:HB3	1.88	0.55
1:0:256:C:H2'	1:0:257:G:O4'	2.06	0.55
1:0:1381:A:H4'	1:0:1382:G:O5'	2.06	0.55
1:0:1654:U:H2'	4:A:47:HIS:CD2	2.41	0.55
1:0:2734:G:H4'	37:0:9071:HOH:O	2.05	0.55
4:A:87:GLU:HB3	37:A:8614:HOH:O	2.06	0.55
4:A:217:ARG:HG2	4:A:229:ALA:HB2	1.88	0.55
9:F:53:ASP:OD1	9:F:80:GLN:HB2	2.05	0.55
1:0:558:C:O2'	1:0:559:U:H5''	2.07	0.55
1:0:669:G:O2'	1:0:670:G:H5'	2.07	0.55
1:0:1230:A:H5'	37:0:6420:HOH:O	2.06	0.55
1:0:2070:G:H5''	37:0:3269:HOH:O	2.06	0.55
1:0:2323:G:H5'	37:0:6412:HOH:O	2.06	0.55
4:A:39:ALA:HB3	4:A:61:GLU:OE2	2.06	0.55
5:B:71:VAL:HG11	5:B:296:LEU:HB3	1.88	0.55
9:F:21:GLU:HA	9:F:24:ARG:HE	1.71	0.55
11:H:72:VAL:HG11	11:H:81:TYR:CZ	2.41	0.55
11:H:127:GLY:O	11:H:128:ALA:HB3	2.06	0.55
12:I:107:ASN:ND2	12:I:107:ASN:C	2.63	0.55
1:0:1197:G:N2	37:0:5647:HOH:O	2.39	0.55
1:0:1790:C:H2'	1:0:1791:U:H6	1.71	0.55
5:B:66:GLU:OE1	5:B:328:ARG:HD2	2.05	0.55
14:K:35:ARG:O	14:K:40:PHE:HA	2.06	0.55
16:M:154:LEU:O	16:M:155:GLU:CB	2.55	0.55
21:R:57:THR:C	21:R:59:ASP:H	2.15	0.55
22:S:38:ARG:NH1	22:S:38:ARG:HG3	2.21	0.55
25:V:21:LEU:HD22	25:V:26:ILE:CD1	2.35	0.55
1:0:288:A:H61	1:0:364:C:H42	1.54	0.55
1:0:1857:A:N6	1:0:2247:C:H1'	2.22	0.55
1:0:2613:G:O2'	1:0:2614:C:H5'	2.06	0.55
14:K:17:SER:C	14:K:19:LYS:N	2.62	0.55
18:O:59:ARG:HH22	18:O:66:GLN:HE22	1.53	0.55
1:0:111:C:H2'	1:0:112:G:O4'	2.07	0.55
1:0:212:A:O4'	1:0:214:U:C6	2.59	0.55
1:0:1423:C:O2'	1:0:1424:A:H5'	2.06	0.55
1:0:1748:U:H4'	37:0:6898:HOH:O	2.06	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:2054:A:N3	20:Q:128:ARG:NH2	2.55	0.55
1:0:2502:C:C2'	1:0:2503:A:H5'	2.35	0.55
37:0:8596:HOH:O	5:B:214:PRO:HD2	2.05	0.55
16:M:161:GLY:O	16:M:162:ASP:C	2.49	0.55
25:V:90:TYR:N	25:V:90:TYR:CD1	2.74	0.55
1:0:396:U:H1'	37:0:7004:HOH:O	2.06	0.55
1:0:1174:A:C5	1:0:1201:C:H4'	2.41	0.55
2:9:3020:G:O2'	2:9:3021:G:H5'	2.07	0.55
2:9:3064:C:C2'	2:9:3065:A:H5'	2.37	0.55
5:B:52:VAL:O	5:B:53:LEU:HD12	2.07	0.55
7:D:10:PHE:CG	7:D:11:HIS:N	2.75	0.55
8:E:84:MET:HB2	8:E:131:LEU:HB2	1.88	0.55
14:K:145:LEU:HB2	37:K:8539:HOH:O	2.05	0.55
17:N:15:LYS:O	17:N:18:ALA:N	2.40	0.55
20:Q:23:MET:HE3	20:Q:28:SER:OG	2.06	0.55
30:1:40:ARG:HG3	30:1:45:ASN:HB3	1.87	0.55
1:0:303:C:O2'	1:0:304:G:H5'	2.07	0.55
1:0:338:C:H4'	6:C:174:ILE:HD12	1.89	0.55
1:0:797:A:C4'	28:Y:10:ARG:N	2.69	0.55
1:0:1119:G:H22	1:0:1246:A:H2	1.50	0.55
1:0:1164:U:N3	1:0:1192:A:H2	2.02	0.55
1:0:1333:U:H2'	1:0:1334:C:C6	2.42	0.55
1:0:1398:G:H2'	1:0:1399:A:C8	2.42	0.55
1:0:1669:A:H2'	1:0:1670:G:C8	2.42	0.55
1:0:2132:C:H2'	37:0:6933:HOH:O	2.07	0.55
1:0:2382:A:H5'	37:0:4200:HOH:O	2.07	0.55
1:0:2548:C:OP2	5:B:5:ARG:NH2	2.39	0.55
7:D:94:ALA:HB3	7:D:174:VAL:HA	1.89	0.55
7:D:166:ILE:HD12	37:D:6326:HOH:O	2.06	0.55
14:K:30:ARG:NH2	37:K:8522:HOH:O	2.38	0.55
18:O:38:GLU:HA	18:O:41:ARG:HH11	1.71	0.55
1:0:675:U:H2'	1:0:676:C:H5'	1.89	0.55
1:0:849:C:O2'	1:0:850:U:H5'	2.06	0.55
1:0:1139:U:H2'	1:0:1140:C:C6	2.41	0.55
1:0:2266:A:H2'	1:0:2267:G:C8	2.42	0.55
7:D:54:ALA:HB2	7:D:69:ILE:HD12	1.89	0.55
7:D:128:LEU:N	37:D:6007:HOH:O	2.39	0.55
14:K:61:ALA:HA	37:K:8564:HOH:O	2.07	0.55
14:K:148:GLU:HA	37:K:8572:HOH:O	2.05	0.55
15:L:9:ARG:HG3	37:L:8544:HOH:O	2.06	0.55
16:M:138:ASP:O	16:M:140:GLN:N	2.33	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:N:77:ALA:HB1	17:N:98:LEU:HD12	1.89	0.55
28:Y:25:ARG:O	28:Y:29:VAL:HG23	2.06	0.55
28:Y:40:PRO:HG2	28:Y:64:ILE:HD13	1.89	0.55
1:0:31:C:H4'	37:0:6799:HOH:O	2.07	0.55
1:0:159:G:H5''	15:L:74:ARG:HH22	1.71	0.55
1:0:559:U:H2'	1:0:560:C:O4'	2.07	0.55
1:0:954:U:O2'	1:0:955:A:H5'	2.07	0.55
7:D:35:ALA:N	37:D:5576:HOH:O	2.39	0.55
7:D:155:HIS:NE2	37:D:7597:HOH:O	2.33	0.55
13:J:62:PRO:CG	13:J:65:ARG:HH21	2.13	0.55
14:K:73:VAL:HG23	14:K:74:THR:N	2.22	0.55
15:L:65:VAL:HG21	15:L:105:ALA:HB2	1.89	0.55
1:0:2405:C:C5'	37:0:5993:HOH:O	2.52	0.54
2:9:3002:U:OP2	2:9:3003:A:H5'	2.07	0.54
6:C:133:ARG:HD2	37:C:8405:HOH:O	2.05	0.54
7:D:58:VAL:HG12	7:D:59:GLY:N	2.22	0.54
9:F:28:ALA:HB3	9:F:99:THR:O	2.08	0.54
19:P:64:GLU:HG3	19:P:74:ASP:OD2	2.06	0.54
1:0:2269:C:H2'	1:0:2270:G:H5'	1.87	0.54
6:C:109:LEU:HD12	6:C:109:LEU:O	2.06	0.54
11:H:31:PHE:CD2	11:H:85:ILE:HG23	2.43	0.54
12:I:103:VAL:HG12	37:I:5907:HOH:O	2.06	0.54
15:L:47:ASP:CG	15:L:48:ARG:N	2.65	0.54
15:L:157:LEU:HB3	15:L:160:PHE:HD1	1.72	0.54
18:O:98:ILE:HD12	18:O:102:ARG:NE	2.22	0.54
1:0:100:C:H5'	22:S:16:LEU:HD12	1.89	0.54
1:0:970:U:H2'	37:0:5738:HOH:O	2.07	0.54
1:0:1864:C:OP1	15:L:75:THR:HG23	2.07	0.54
1:0:2676:C:H4'	12:I:70:PHE:CE1	2.42	0.54
1:0:2896:A:H5''	37:0:5517:HOH:O	2.06	0.54
11:H:45:GLN:HG3	11:H:135:TRP:NE1	2.22	0.54
12:I:99:GLU:HA	37:I:7377:HOH:O	2.06	0.54
21:R:4:VAL:HG23	37:R:2334:HOH:O	2.07	0.54
29:Z:45:ARG:HB3	37:Z:8419:HOH:O	2.07	0.54
1:0:289:G:O2'	1:0:290:C:H5'	2.06	0.54
1:0:581:G:H5'	37:0:7057:HOH:O	2.06	0.54
4:A:88:ILE:HD13	4:A:100:PRO:CD	2.32	0.54
6:C:142:ASP:OD2	6:C:238:SER:OG	2.24	0.54
11:H:35:ASN:ND2	11:H:79:ALA:O	2.41	0.54
12:I:14:ALA:HB1	12:I:44:ALA:HB2	1.87	0.54
20:Q:18:LEU:HD11	20:Q:87:ALA:O	2.07	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:R:57:THR:HG22	21:R:59:ASP:HB2	1.88	0.54
30:1:48:ASP:O	30:1:49:GLU:HB2	2.08	0.54
2:9:3049:G:H5''	37:9:8471:HOH:O	2.07	0.54
7:D:86:THR:O	7:D:90:LEU:HG	2.07	0.54
8:E:132:THR:HG23	8:E:132:THR:O	2.07	0.54
15:L:12:TRP:CE2	15:L:20:ILE:HD11	2.43	0.54
23:T:49:LEU:HD11	37:T:3805:HOH:O	2.08	0.54
1:0:1947:G:N2	1:0:1966:U:C2	2.75	0.54
1:0:119:A:N3	37:0:8653:HOH:O	2.34	0.54
1:0:182:G:O3'	15:L:157:LEU:HD13	2.07	0.54
1:0:1118:A:H8	1:0:1119:G:H5''	1.73	0.54
1:0:1666:C:O2'	1:0:1667:A:C5'	2.56	0.54
1:0:1886:A:N3	37:0:4278:HOH:O	2.33	0.54
1:0:2072:G:C6	1:0:2533:C:H1'	2.43	0.54
37:0:5729:HOH:O	18:O:100:ALA:HA	2.07	0.54
37:0:9481:HOH:O	14:K:22:ARG:HG2	2.07	0.54
14:K:143:THR:CG2	14:K:144:ASP:N	2.70	0.54
23:T:11:THR:HG22	23:T:53:ASP:OD2	2.08	0.54
28:Y:56:MET:CE	28:Y:63:LYS:HE3	2.37	0.54
1:0:138:U:OP2	1:0:139:C:H5	1.90	0.54
1:0:2265:U:H2'	1:0:2266:A:C8	2.43	0.54
8:E:11:VAL:HG12	8:E:12:ASP:H	1.70	0.54
9:F:48:VAL:HG23	9:F:74:PHE:CB	2.38	0.54
13:J:37:TYR:HE2	13:J:45:PRO:HA	1.73	0.54
15:L:24:MET:HE3	15:L:28:MET:CE	2.35	0.54
22:S:48:VAL:HG23	22:S:98:VAL:HA	1.90	0.54
23:T:52:THR:HG22	23:T:54:THR:HB	1.89	0.54
1:0:241:A:C2	1:0:378:A:H4'	2.43	0.54
1:0:542:A:H2'	1:0:543:G:O4'	2.08	0.54
1:0:625:U:H5''	1:0:1044:C:N4	2.23	0.54
5:B:82:VAL:HG12	5:B:101:TRP:CE3	2.43	0.54
8:E:8:PRO:HB2	8:E:11:VAL:HG23	1.89	0.54
17:N:96:VAL:CG1	17:N:100:GLN:HB2	2.38	0.54
22:S:9:LYS:CE	22:S:13:ARG:NH1	2.70	0.54
22:S:80:GLU:HA	37:S:6653:HOH:O	2.07	0.54
1:0:371:U:H2'	1:0:372:A:H8	1.72	0.54
1:0:844:A:C6	1:0:882:A:C5	2.95	0.54
1:0:1561:U:H2'	37:0:3619:HOH:O	2.06	0.54
1:0:2825:C:H4'	1:0:2826:G:O5'	2.08	0.54
2:9:3013:A:O2'	2:9:3014:G:H5''	2.08	0.54
4:A:17:ARG:HD2	37:A:8537:HOH:O	2.07	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:A:81:GLN:HB2	4:A:92:ASN:ND2	2.22	0.54
5:B:48:MET:HB2	37:B:8561:HOH:O	2.08	0.54
5:B:177:HIS:O	5:B:181:ILE:HG13	2.07	0.54
5:B:205:VAL:O	5:B:307:ARG:NE	2.40	0.54
11:H:31:PHE:HD2	11:H:85:ILE:O	1.91	0.54
11:H:62:GLU:HA	37:H:8370:HOH:O	2.08	0.54
13:J:28:GLU:HB3	13:J:59:LYS:HB2	1.90	0.54
15:L:164:THR:HG23	15:L:165:SER:H	1.70	0.54
16:M:82:TYR:C	16:M:82:TYR:CD2	2.86	0.54
25:V:26:ILE:HG13	25:V:26:ILE:O	2.08	0.54
1:0:947:U:O2'	1:0:948:G:H5'	2.08	0.53
1:0:2377:U:H6	1:0:2377:U:O5'	1.91	0.53
2:9:3003:A:N6	2:9:3022:G:H1'	2.23	0.53
5:B:238:ASN:ND2	5:B:240:GLY:N	2.51	0.53
6:C:162:VAL:HG13	6:C:232:LEU:HD21	1.90	0.53
12:I:47:THR:HB	37:I:3661:HOH:O	2.08	0.53
15:L:35:PRO:C	37:L:8537:HOH:O	2.51	0.53
1:0:544:G:C2'	1:0:545:G:H5''	2.38	0.53
1:0:1209:C:H2'	1:0:1210:G:H8	1.73	0.53
1:0:1562:C:O2	1:0:1562:C:H2'	2.06	0.53
1:0:1787:C:H4'	1:0:2883:A:O4'	2.09	0.53
9:F:19:ALA:O	9:F:22:VAL:HG22	2.08	0.53
18:O:139:ARG:NH2	37:O:6072:HOH:O	2.41	0.53
20:Q:96:VAL:HG13	20:Q:106:GLY:HA3	1.89	0.53
1:0:821:U:H2'	1:0:822:C:C6	2.42	0.53
11:H:47:GLU:CB	11:H:133:ILE:HD13	2.38	0.53
13:J:49:LEU:HD21	13:J:74:VAL:O	2.07	0.53
16:M:159:TYR:HB3	16:M:162:ASP:HB2	1.91	0.53
24:U:39:ALA:N	24:U:40:PRO:CD	2.71	0.53
1:0:2064:U:H5'	1:0:2652:U:O3'	2.08	0.53
1:0:2760:C:OP1	5:B:209:LYS:NZ	2.40	0.53
5:B:141:ARG:HG2	5:B:165:ARG:HA	1.89	0.53
12:I:40:ASN:OD1	12:I:106:GLY:HA2	2.08	0.53
15:L:77:PHE:HD2	37:L:8528:HOH:O	1.91	0.53
16:M:5:ARG:HG3	19:P:18:PRO:HB3	1.89	0.53
22:S:55:PHE:HB2	37:S:6384:HOH:O	2.08	0.53
1:0:394:G:H1	15:L:181:GLU:CD	2.17	0.53
1:0:1120:U:H5'	1:0:1121:G:OP2	2.09	0.53
1:0:1804:A:H2'	1:0:1805:G:C8	2.43	0.53
5:B:132:HIS:HB2	5:B:137:LEU:HD22	1.89	0.53
5:B:221:GLN:HE22	13:J:42:ASN:HD22	1.56	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:I:130:VAL:HG12	12:I:131:THR:N	2.21	0.53
14:K:57:VAL:HG12	14:K:57:VAL:O	2.08	0.53
15:L:47:ASP:CG	15:L:48:ARG:H	2.17	0.53
15:L:63:VAL:HG21	15:L:109:PHE:CE1	2.44	0.53
27:X:112:GLU:OE1	27:X:112:GLU:HA	2.09	0.53
30:1:16:ASN:C	30:1:18:ASN:H	2.17	0.53
1:0:1342:C:C2'	1:0:1343:C:H5'	2.39	0.53
1:0:1477:C:H5'	1:0:1868:G:C5'	2.38	0.53
1:0:1595:G:O2'	1:0:1596:U:H5'	2.09	0.53
1:0:2359:G:N7	37:0:3190:HOH:O	2.34	0.53
1:0:2896:A:OP1	26:W:15:ARG:NH1	2.41	0.53
4:A:48:ASP:HB3	37:A:8602:HOH:O	2.07	0.53
4:A:168:PRO:O	4:A:170:VAL:HG23	2.07	0.53
5:B:13:PHE:O	5:B:16:ARG:HD2	2.08	0.53
7:D:44:ILE:HG23	7:D:45:THR:HG23	1.89	0.53
11:H:72:VAL:CG1	11:H:81:TYR:CZ	2.92	0.53
25:V:108:ARG:HE	25:V:114:PRO:HG3	1.73	0.53
31:2:31:THR:HB	31:2:33:MET:HE3	1.90	0.53
1:0:319:A:H4'	1:0:338:C:C5	2.44	0.53
1:0:1189:A:H3'	37:0:7055:HOH:O	2.08	0.53
1:0:1328:A:C8	27:X:169:ARG:HD3	2.44	0.53
1:0:1528:A:H2'	1:0:1529:G:O4'	2.08	0.53
1:0:1735:C:O2'	1:0:1736:A:H5'	2.08	0.53
1:0:1771:U:H4'	28:Y:20:LEU:HD21	1.90	0.53
1:0:2860:G:H1'	37:0:6193:HOH:O	2.07	0.53
1:0:2909:G:H2'	1:0:2910:A:H8	1.73	0.53
2:9:3061:C:H2'	2:9:3062:A:H8	1.74	0.53
11:H:26:LYS:HD2	11:H:28:ILE:CB	2.38	0.53
12:I:77:GLY:O	12:I:78:ILE:C	2.50	0.53
12:I:93:ARG:HH11	12:I:93:ARG:CB	2.15	0.53
13:J:118:ALA:HA	13:J:125:ALA:HB2	1.90	0.53
1:0:521:A:H2'	1:0:522:U:H5'	1.91	0.53
1:0:1783:A:C2'	1:0:1784:U:H5'	2.39	0.53
1:0:2898:G:H4'	5:B:288:GLY:HA2	1.90	0.53
37:0:5622:HOH:O	5:B:2:GLN:HA	2.08	0.53
5:B:141:ARG:HD2	5:B:163:GLU:OE2	2.08	0.53
7:D:54:ALA:CB	7:D:69:ILE:HD12	2.38	0.53
20:Q:25:PHE:CE2	20:Q:29:LYS:HE2	2.43	0.53
31:2:65:THR:HB	31:2:83:TRP:H	1.74	0.53
1:0:858:U:H2'	1:0:859:C:H6	1.72	0.53
1:0:2274:A:H1'	15:L:86:MET:SD	2.49	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:9:3076:G:C3'	2:9:3077:A:H5''	2.29	0.53
11:H:56:ILE:HG22	11:H:61:LEU:CD2	2.37	0.53
17:N:32:ARG:HE	17:N:35:LYS:HD2	1.73	0.53
1:0:175:G:H2'	15:L:192:ALA:HB3	1.91	0.53
1:0:702:G:O2'	1:0:703:G:H5'	2.09	0.53
1:0:1003:U:HO2'	11:H:90:PHE:HE1	1.56	0.53
1:0:1790:C:O2'	1:0:1791:U:H5'	2.08	0.53
2:9:3025:G:C3'	2:9:3026:C:H5'	2.36	0.53
5:B:148:PRO:HD2	37:B:8583:HOH:O	2.08	0.53
11:H:45:GLN:HE21	11:H:135:TRP:HE1	1.57	0.53
20:Q:132:ARG:CZ	37:Q:8585:HOH:O	2.57	0.53
22:S:89:ARG:HD2	22:S:89:ARG:C	2.34	0.53
25:V:122:ARG:HG2	25:V:152:ALA:O	2.08	0.53
25:V:125:HIS:HE1	37:V:3071:HOH:O	1.92	0.53
25:V:151:GLU:O	25:V:154:ARG:HB3	2.09	0.53
28:Y:56:MET:HA	28:Y:62:TYR:O	2.09	0.53
1:0:169:A:O2'	31:2:48:ASN:ND2	2.42	0.52
1:0:280:C:H5'	37:0:5735:HOH:O	2.08	0.52
1:0:714:U:H4'	37:0:5169:HOH:O	2.09	0.52
1:0:1878:G:O2'	1:0:1879:U:C6	2.61	0.52
1:0:2136:G:H2'	37:0:4843:HOH:O	2.10	0.52
1:0:2781:U:C2'	1:0:2782:G:H5'	2.39	0.52
4:A:13:THR:HB	37:A:8531:HOH:O	2.09	0.52
20:Q:33:ARG:NH1	37:Q:8546:HOH:O	2.41	0.52
31:2:69:TYR:HB2	31:2:78:HIS:CE1	2.43	0.52
1:0:1151:G:OP1	10:G:63:ARG:NH1	2.43	0.52
8:E:31:ARG:NH1	8:E:68:HIS:CG	2.77	0.52
8:E:49:ILE:HD11	8:E:69:ILE:HD12	1.90	0.52
15:L:25:TRP:HE3	15:L:26:HIS:HD2	1.57	0.52
16:M:73:ALA:N	37:M:6988:HOH:O	2.43	0.52
22:S:48:VAL:HG22	22:S:97:ARG:C	2.34	0.52
1:0:338:C:H4'	6:C:174:ILE:HD11	1.91	0.52
1:0:737:A:H2'	1:0:738:G:O4'	2.09	0.52
1:0:1189:A:O2'	1:0:1208:C:H2'	2.09	0.52
1:0:1783:A:O2'	1:0:1784:U:H5'	2.09	0.52
1:0:2716:G:H5''	5:B:206:THR:HG21	1.90	0.52
1:0:2729:C:H2'	1:0:2730:G:H8	1.74	0.52
37:0:3541:HOH:O	5:B:27:ASN:HB2	2.08	0.52
37:0:3701:HOH:O	30:1:38:LYS:HE3	2.08	0.52
2:9:3055:U:H4'	2:9:3056:A:C8	2.45	0.52
13:J:1:MET:HE1	37:J:6646:HOH:O	2.08	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:K:142:LEU:HG	14:K:146:GLY:HA3	1.91	0.52
29:Z:21:ARG:HD2	29:Z:37:CYS:SG	2.50	0.52
1:0:198:A:H2'	37:0:5321:HOH:O	2.09	0.52
1:0:1205:U:H2'	1:0:1206:U:C5'	2.40	0.52
1:0:1682:A:H5''	37:0:8966:HOH:O	2.08	0.52
1:0:1847:A:OP1	4:A:175:LYS:HG3	2.10	0.52
1:0:2314:G:C2'	1:0:2315:C:H5'	2.40	0.52
37:0:9947:HOH:O	12:I:46:ILE:HD12	2.09	0.52
4:A:125:ASN:HB3	4:A:158:VAL:HG12	1.91	0.52
5:B:48:MET:N	37:B:8561:HOH:O	2.41	0.52
6:C:16:VAL:HG12	6:C:17:ASP:H	1.73	0.52
8:E:107:PHE:CE1	8:E:152:THR:HB	2.45	0.52
22:S:49:GLU:OE2	22:S:97:ARG:HD2	2.10	0.52
1:0:316:A:H5'	22:S:54:ASP:OD2	2.08	0.52
1:0:396:U:OP2	31:2:38:ARG:HD2	2.09	0.52
4:A:33:GLU:O	4:A:34:ASP:HB2	2.08	0.52
10:G:67:LEU:O	10:G:71:LEU:HG	2.10	0.52
16:M:67:ALA:C	16:M:69:TYR:H	2.17	0.52
16:M:74:PRO:HG2	16:M:159:TYR:CZ	2.44	0.52
18:O:131:PHE:CD1	18:O:137:LEU:HD13	2.43	0.52
23:T:52:THR:HG22	23:T:54:THR:N	2.24	0.52
25:V:1:MET:N	25:V:37:GLU:HG3	2.24	0.52
1:0:120:A:H5'	29:Z:20:ARG:HH21	1.74	0.52
1:0:677:C:H4'	6:C:246:ARG:NH2	2.25	0.52
1:0:2047:C:H5'	37:0:9311:HOH:O	2.09	0.52
1:0:2756:U:H3	1:0:2896:A:H2	1.58	0.52
2:9:3107:C:H5	37:9:8440:HOH:O	1.89	0.52
6:C:47:GLY:HA2	6:C:92:PRO:HB2	1.92	0.52
8:E:20:ILE:CD1	8:E:40:VAL:HG11	2.34	0.52
8:E:80:TRP:O	8:E:134:SER:HA	2.08	0.52
11:H:48:LEU:CG	11:H:157:ILE:HG21	2.39	0.52
14:K:138:GLY:HA3	37:K:8554:HOH:O	2.08	0.52
1:0:229:G:O2'	1:0:230:C:H5'	2.10	0.52
2:9:3031:C:H2'	2:9:3032:G:O4'	2.09	0.52
5:B:207:LYS:HG2	5:B:304:PRO:HB3	1.89	0.52
6:C:180:SER:HB2	37:C:8441:HOH:O	2.08	0.52
9:F:56:PRO:HG2	15:L:44:THR:HA	1.91	0.52
15:L:149:TRP:O	15:L:152:ARG:HG2	2.10	0.52
22:S:35:TYR:CG	22:S:112:LEU:HD22	2.45	0.52
23:T:9:CYS:CA	23:T:52:THR:HG23	2.40	0.52
1:0:42:C:H1'	37:0:4131:HOH:O	2.08	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:1025:C:H5'	25:V:23:MET:O	2.10	0.52
1:0:2781:U:O2'	1:0:2782:G:H5'	2.10	0.52
6:C:118:THR:O	6:C:136:VAL:HG13	2.10	0.52
11:H:55:GLN:HE22	11:H:91:HIS:CD2	2.28	0.52
11:H:147:ARG:HA	11:H:150:LYS:NZ	2.24	0.52
12:I:46:ILE:HA	37:I:1123:HOH:O	2.08	0.52
15:L:67:ILE:HD11	15:L:104:ARG:HD2	1.92	0.52
15:L:81:ARG:O	15:L:86:MET:HE2	2.09	0.52
16:M:47:LEU:CD1	16:M:97:VAL:HG11	2.39	0.52
16:M:67:ALA:C	16:M:69:TYR:N	2.68	0.52
19:P:30:VAL:O	19:P:30:VAL:HG12	2.10	0.52
27:X:126:PRO:HG2	27:X:128:PHE:CE1	2.44	0.52
1:0:1926:G:H2'	1:0:1927:A:C8	2.45	0.52
1:0:2359:G:H3'	37:0:5121:HOH:O	2.09	0.52
1:0:2745:C:H5	37:0:5311:HOH:O	1.92	0.52
26:W:43:VAL:CG1	26:W:47:ALA:HB3	2.39	0.52
27:X:178:HIS:CG	27:X:179:PRO:HD2	2.45	0.52
27:X:186:ARG:HG2	27:X:186:ARG:NH1	2.24	0.52
1:0:92:G:H4'	24:U:44:GLY:HA3	1.90	0.52
1:0:359:U:H2'	1:0:360:A:H8	1.75	0.52
1:0:1138:G:H4'	37:0:5139:HOH:O	2.09	0.52
1:0:1234:U:C4	5:B:244:PRO:HB3	2.45	0.52
1:0:1236:A:H2'	1:0:1237:U:O4'	2.10	0.52
1:0:1450:C:C4'	1:0:1451:C:OP2	2.55	0.52
5:B:1:PRO:O	5:B:2:GLN:HB2	2.10	0.52
5:B:16:ARG:HB3	5:B:217:ARG:HH21	1.75	0.52
5:B:36:PRO:HG3	5:B:169:GLY:H	1.75	0.52
5:B:85:ARG:NH1	37:B:8633:HOH:O	2.43	0.52
6:C:168:ARG:NH2	6:C:190:ALA:O	2.43	0.52
6:C:195:VAL:HA	6:C:213:ALA:O	2.09	0.52
7:D:59:GLY:O	7:D:61:PHE:N	2.35	0.52
11:H:71:TYR:C	11:H:73:GLN:N	2.67	0.52
11:H:86:ARG:CZ	11:H:130:HIS:CD2	2.93	0.52
13:J:23:ASN:HA	37:J:7075:HOH:O	2.10	0.52
13:J:99:ASP:OD1	13:J:101:ASN:N	2.42	0.52
15:L:57:LYS:HE2	15:L:140:ALA:O	2.10	0.52
15:L:67:ILE:CD1	15:L:104:ARG:HD2	2.39	0.52
20:Q:133:ALA:HB3	37:Q:8588:HOH:O	2.09	0.52
25:V:7:LEU:CD1	25:V:53:ALA:HB2	2.40	0.52
27:X:205:ILE:O	27:X:206:ALA:C	2.53	0.52
27:X:213:LYS:O	27:X:217:ILE:HG13	2.10	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:113:A:OP2	1:0:114:A:H2'	2.10	0.51
1:0:1072:G:OP2	27:X:154:ARG:NH2	2.38	0.51
1:0:1819:G:H2'	1:0:1820:G:H4'	1.90	0.51
2:9:3051:A:H5'	16:M:160:SER:HB3	1.91	0.51
7:D:99:ASP:HB2	7:D:103:ASN:HB2	1.91	0.51
15:L:104:ARG:O	15:L:108:LYS:HG2	2.09	0.51
20:Q:4:TYR:N	37:Q:8550:HOH:O	2.42	0.51
26:W:41:PHE:CZ	26:W:74:ALA:HB3	2.45	0.51
30:1:40:ARG:HG2	30:1:40:ARG:NH1	2.26	0.51
1:0:652:G:H8	37:0:9504:HOH:O	1.93	0.51
1:0:784:A:O2'	1:0:1458:A:N3	2.41	0.51
1:0:2119:C:O2'	1:0:2120:U:H5'	2.10	0.51
4:A:105:VAL:HG12	4:A:106:CYS:N	2.25	0.51
5:B:13:PHE:CD1	5:B:13:PHE:N	2.78	0.51
7:D:11:HIS:C	7:D:13:MET:H	2.18	0.51
7:D:64:ARG:HB3	7:D:67:ASP:OD2	2.10	0.51
8:E:154:ILE:HD11	8:E:157:LYS:HB2	1.92	0.51
9:F:115:VAL:O	9:F:118:LEU:N	2.44	0.51
10:G:71:LEU:C	10:G:73:ASP:N	2.67	0.51
11:H:118:PRO:HD2	37:H:8326:HOH:O	2.09	0.51
12:I:75:PRO:HG2	12:I:105:LEU:CD2	2.39	0.51
15:L:59:GLY:HA3	15:L:141:ILE:CD1	2.39	0.51
16:M:115:VAL:O	16:M:118:ILE:HB	2.10	0.51
20:Q:119:VAL:O	20:Q:119:VAL:HG12	2.09	0.51
27:X:165:GLU:HB3	37:X:8598:HOH:O	2.10	0.51
28:Y:56:MET:HE2	28:Y:63:LYS:HE3	1.92	0.51
1:0:960:G:N3	1:0:960:G:H2'	2.25	0.51
1:0:2434:A:O3'	31:2:28:GLY:HA3	2.11	0.51
1:0:2503:A:H1'	37:0:5772:HOH:O	2.10	0.51
1:0:2503:A:OP1	11:H:147:ARG:NH2	2.40	0.51
7:D:25:MET:CE	7:D:37:ALA:HB1	2.40	0.51
9:F:117:GLU:C	9:F:119:ARG:H	2.19	0.51
12:I:54:VAL:HG11	12:I:138:THR:HG21	1.92	0.51
17:N:35:LYS:HD3	37:N:3360:HOH:O	2.09	0.51
20:Q:19:ARG:HA	20:Q:142:ASP:OD1	2.10	0.51
20:Q:44:VAL:O	20:Q:48:GLU:HG3	2.11	0.51
27:X:185:VAL:HG12	37:X:8571:HOH:O	2.08	0.51
31:2:3:MET:O	31:2:90:PHE:HA	2.09	0.51
1:0:86:A:C2	30:1:25:VAL:HG13	2.45	0.51
1:0:240:C:O2	1:0:240:C:H2'	2.11	0.51
1:0:558:C:C2'	1:0:559:U:C5'	2.88	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:B:258:GLY:HA2	37:B:8560:HOH:O	2.10	0.51
8:E:108:LEU:HD11	8:E:164:ASP:HB2	1.92	0.51
11:H:83:PHE:HE1	11:H:146:TRP:CZ2	2.28	0.51
14:K:17:SER:O	14:K:19:LYS:N	2.44	0.51
19:P:66:LYS:HB2	19:P:70:ALA:O	2.10	0.51
20:Q:17:MET:SD	37:Q:8550:HOH:O	2.59	0.51
20:Q:39:THR:CB	20:Q:42:GLU:HG3	2.41	0.51
25:V:28:HIS:HD2	25:V:31:HIS:CE1	2.28	0.51
30:1:36:ASN:HB3	30:1:39:ARG:NE	2.25	0.51
1:0:1268:C:H2'	1:0:1269:G:C8	2.45	0.51
1:0:1839:A:H5'	1:0:2643:G:H4'	1.92	0.51
1:0:2115:U:H2'	1:0:2116:U:C6	2.46	0.51
1:0:2524:G:H21	1:0:2526:C:N4	2.09	0.51
4:A:191:GLY:HA2	4:A:194:MET:CE	2.40	0.51
5:B:138:GLY:O	5:B:139:ASP:O	2.28	0.51
6:C:127:ARG:CZ	6:C:225:PRO:HG2	2.39	0.51
7:D:94:ALA:HB3	7:D:174:VAL:CA	2.40	0.51
15:L:5:TYR:O	15:L:7:TYR:N	2.44	0.51
15:L:184:ARG:HG3	15:L:185:PRO:HA	1.92	0.51
16:M:143:ARG:HH12	16:M:173:ASP:CG	2.16	0.51
25:V:122:ARG:NH2	25:V:154:ARG:HD2	2.23	0.51
1:0:184:G:H5''	15:L:153:THR:HG22	1.91	0.51
1:0:344:C:H2'	1:0:345:G:O4'	2.10	0.51
1:0:603:A:H5''	1:0:604:G:OP1	2.10	0.51
1:0:1181:A:H2'	1:0:1182:C:O4'	2.10	0.51
1:0:1187:U:C3'	37:0:6289:HOH:O	2.59	0.51
1:0:1470:A:OP1	15:L:93:ARG:HD2	2.11	0.51
1:0:2044:G:OP1	26:W:23:HIS:HE1	1.94	0.51
1:0:2491:G:H1'	37:0:6260:HOH:O	2.09	0.51
4:A:55:VAL:HG11	4:A:67:LEU:HD13	1.93	0.51
5:B:7:ARG:HD3	5:B:9:GLY:O	2.09	0.51
6:C:5:ILE:HG23	37:C:8427:HOH:O	2.09	0.51
11:H:45:GLN:HE21	11:H:135:TRP:NE1	2.09	0.51
17:N:47:ARG:HG3	17:N:47:ARG:NH1	2.25	0.51
25:V:108:ARG:NH2	37:V:2359:HOH:O	2.44	0.51
1:0:185:G:H4'	1:0:186:A:H4'	1.92	0.51
1:0:621:C:H5'	27:X:132:ASP:OD2	2.11	0.51
1:0:1285:U:H1'	37:0:6879:HOH:O	2.11	0.51
1:0:1293:U:O2'	27:X:149:GLN:NE2	2.38	0.51
1:0:2071:C:H5'	37:0:9034:HOH:O	2.11	0.51
2:9:3025:G:H8	37:M:2665:HOH:O	1.94	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:9:8479:HOH:O	16:M:23:ARG:NH1	2.40	0.51
9:F:46:GLU:N	37:F:3461:HOH:O	2.44	0.51
15:L:69:LYS:HG2	15:L:127:LYS:HG3	1.93	0.51
1:0:177:A:H2'	1:0:178:U:O4'	2.09	0.51
1:0:622:G:O2'	1:0:623:U:H5'	2.11	0.51
1:0:703:G:O2'	1:0:704:C:H5'	2.10	0.51
1:0:1503:U:H2'	1:0:1504:A:O4'	2.11	0.51
1:0:1603:A:H5'	1:0:1605:G:H5'	1.92	0.51
5:B:41:PHE:CD2	5:B:190:MET:HE3	2.46	0.51
5:B:260:HIS:HA	37:B:8624:HOH:O	2.10	0.51
7:D:95:THR:C	7:D:97:GLN:N	2.68	0.51
7:D:174:VAL:HG13	37:D:6555:HOH:O	2.11	0.51
13:J:27:ARG:HD2	37:J:4747:HOH:O	2.10	0.51
14:K:12:THR:HG21	14:K:16:GLY:O	2.11	0.51
16:M:22:GLN:HG2	16:M:26:LEU:CD2	2.39	0.51
17:N:14:LEU:CD2	17:N:102:ILE:HD11	2.41	0.51
19:P:50:GLY:HA3	19:P:87:THR:OG1	2.11	0.51
1:0:1119:G:N2	1:0:1246:A:H2	2.06	0.51
1:0:1307:A:H2'	1:0:1308:A:C8	2.46	0.51
5:B:248:ARG:O	5:B:251:VAL:CG1	2.59	0.51
7:D:23:VAL:HG21	7:D:45:THR:HG21	1.92	0.51
8:E:16:ASP:O	8:E:17:HIS:HB2	2.10	0.51
8:E:69:ILE:HA	8:E:72:MET:CE	2.40	0.51
13:J:113:ILE:HG22	13:J:114:ALA:O	2.10	0.51
18:O:115:SER:C	18:O:117:SER:H	2.19	0.51
22:S:41:ARG:HG2	22:S:41:ARG:HH11	1.75	0.51
26:W:76:ARG:HG3	26:W:76:ARG:NH1	2.21	0.51
27:X:172:THR:HG22	27:X:173:ALA:N	2.26	0.51
1:0:80:A:H3'	22:S:43:ASN:OD1	2.11	0.51
1:0:281:U:O2'	1:0:282:C:H5'	2.11	0.51
1:0:324:G:O2'	1:0:325:U:H5'	2.10	0.51
1:0:694:A:C2'	1:0:695:C:H5'	2.40	0.51
1:0:755:G:O2'	1:0:756:A:H5'	2.11	0.51
1:0:823:U:H2'	1:0:824:G:O4'	2.11	0.51
1:0:1183:C:N4	37:0:3869:HOH:O	2.36	0.51
1:0:1701:A:H4'	1:0:1702:U:O5'	2.11	0.51
1:0:1719:G:H1'	37:0:3215:HOH:O	2.11	0.51
1:0:1768:C:H2'	1:0:1769:C:O4'	2.10	0.51
1:0:1825:U:O4'	1:0:1999:C:H5''	2.11	0.51
1:0:2780:C:H2'	1:0:2781:U:C6	2.46	0.51
4:A:94:LEU:HG	4:A:99:ILE:CD1	2.40	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:B:62:ARG:HA	5:B:65:MET:CE	2.31	0.51
6:C:49:ASP:HB3	6:C:52:ALA:HB2	1.93	0.51
8:E:84:MET:HE2	8:E:133:VAL:HG23	1.93	0.51
9:F:28:ALA:CB	9:F:99:THR:HG23	2.41	0.51
13:J:29:LEU:HB3	13:J:55:VAL:CG1	2.36	0.51
15:L:37:VAL:CG2	15:L:108:LYS:HG3	2.37	0.51
18:O:87:ARG:HG2	37:O:5182:HOH:O	2.10	0.51
20:Q:113:HIS:HE1	20:Q:144:GLU:CD	2.19	0.51
21:R:33:SER:O	21:R:37:VAL:HG23	2.11	0.51
21:R:57:THR:CG2	21:R:59:ASP:HB2	2.41	0.51
1:O:130:C:H5'	37:O:4659:HOH:O	2.10	0.50
1:O:290:C:O2'	1:O:291:C:H5'	2.11	0.50
1:O:772:G:H2'	1:O:773:A:O4'	2.11	0.50
1:O:1051:C:H2'	1:O:1052:G:O4'	2.11	0.50
1:O:1306:U:OP1	6:C:184:ARG:HD2	2.11	0.50
1:O:1334:C:H2'	1:O:1335:C:H6	1.76	0.50
1:O:1370:G:C4	37:O:9632:HOH:O	2.64	0.50
1:O:1789:G:O6	18:O:73:HIS:HE1	1.94	0.50
1:O:2252:A:C5	1:O:2253:G:H1'	2.45	0.50
6:C:127:ARG:HG2	6:C:127:ARG:NH1	2.26	0.50
6:C:133:ARG:NH2	37:C:8423:HOH:O	2.43	0.50
16:M:61:ALA:CB	16:M:88:ALA:HB2	2.42	0.50
25:V:39:ASP:HB2	37:V:3580:HOH:O	2.10	0.50
27:X:169:ARG:NH2	37:X:8531:HOH:O	2.42	0.50
31:2:84:ARG:HD3	37:2:8541:HOH:O	2.10	0.50
1:O:659:A:N7	17:N:63:LYS:NZ	2.52	0.50
1:O:894:A:C2	6:C:87:ARG:NH2	2.79	0.50
1:O:1189:A:H1'	1:O:1209:C:C1'	2.42	0.50
1:O:1500:U:P	18:O:41:ARG:HH22	2.34	0.50
1:O:1593:C:OP1	18:O:117:SER:CB	2.60	0.50
1:O:1664:A:H8	1:O:1664:A:OP1	1.93	0.50
1:O:2361:A:H5''	37:O:8523:HOH:O	2.10	0.50
1:O:2670:G:N2	37:O:3113:HOH:O	2.32	0.50
4:A:8:ARG:NH1	37:A:8547:HOH:O	2.43	0.50
7:D:41:LEU:HA	7:D:44:ILE:CG2	2.41	0.50
11:H:59:ASN:H	11:H:59:ASN:ND2	1.99	0.50
25:V:149:LEU:HG	25:V:153:MET:CE	2.32	0.50
1:O:281:U:H3'	37:O:6595:HOH:O	2.11	0.50
1:O:820:G:H5'	1:O:821:U:C5'	2.42	0.50
1:O:1674:C:H2'	1:O:1675:C:H6	1.77	0.50
1:O:2906:A:H5'	1:O:2907:C:O4'	2.12	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:9:3042:C:H5'	2:9:3043:G:OP2	2.11	0.50
11:H:85:ILE:HG23	11:H:85:ILE:O	2.10	0.50
17:N:39:THR:O	17:N:115:ARG:NH2	2.44	0.50
20:Q:100:ASP:C	20:Q:102:GLN:H	2.19	0.50
22:S:75:GLU:O	22:S:76:ASP:HB2	2.10	0.50
23:T:17:THR:CG2	23:T:18:GLY:N	2.74	0.50
24:U:58:THR:O	24:U:62:GLU:HG3	2.11	0.50
28:Y:30:GLU:HA	28:Y:33:HIS:CB	2.41	0.50
1:0:816:G:H5'	1:0:1598:A:H4'	1.93	0.50
1:0:958:G:O2'	1:0:959:C:H5'	2.12	0.50
1:0:1087:G:H4'	1:0:1088:A:OP1	2.12	0.50
1:0:1730:G:H5'	1:0:1731:C:C6	2.46	0.50
8:E:11:VAL:CG1	8:E:12:ASP:N	2.73	0.50
9:F:78:GLU:CB	37:F:2750:HOH:O	2.59	0.50
23:T:4:ARG:N	37:T:5334:HOH:O	2.43	0.50
24:U:1:THR:C	24:U:3:LEU:N	2.69	0.50
1:0:128:A:C8	1:0:128:A:H3'	2.47	0.50
1:0:154:C:H2'	1:0:155:C:C6	2.44	0.50
1:0:832:U:H2'	1:0:833:G:C8	2.46	0.50
1:0:920:C:H5'	1:0:921:G:C4	2.47	0.50
1:0:1386:G:N3	37:0:9680:HOH:O	2.34	0.50
4:A:36:ASP:HB2	4:A:84:VAL:N	2.27	0.50
5:B:108:GLU:HB3	5:B:111:ARG:HD2	1.93	0.50
5:B:248:ARG:NH1	37:B:8614:HOH:O	2.45	0.50
9:F:26:THR:HG21	9:F:103:ALA:CB	2.41	0.50
20:Q:117:HIS:HA	37:Q:8537:HOH:O	2.10	0.50
25:V:110:GLN:HE21	25:V:110:GLN:HA	1.75	0.50
29:Z:25:LYS:CG	30:1:49:GLU:H	2.25	0.50
4:A:2:ARG:HB3	37:A:8523:HOH:O	2.11	0.50
5:B:14:GLY:HA2	5:B:15:PRO:C	2.35	0.50
5:B:51:VAL:CG2	5:B:327:VAL:HG13	2.41	0.50
13:J:118:ALA:O	13:J:120:ARG:N	2.44	0.50
15:L:172:GLY:O	15:L:183:VAL:HG11	2.12	0.50
19:P:32:GLU:HA	19:P:71:TYR:OH	2.11	0.50
22:S:71:VAL:HG13	22:S:91:LEU:O	2.11	0.50
25:V:13:MET:HA	37:V:4944:HOH:O	2.10	0.50
1:0:396:U:O2'	1:0:418:C:H4'	2.12	0.50
1:0:1211:G:O2'	1:0:1212:C:H5'	2.11	0.50
1:0:1883:U:O2'	1:0:1884:G:H5'	2.12	0.50
1:0:1930:A:H2'	1:0:1931:A:C8	2.47	0.50
1:0:2424:U:H5'	37:0:6673:HOH:O	2.11	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:2626:C:H2'	1:0:2627:G:C8	2.47	0.50
1:0:2672:C:P	5:B:25:ARG:HH11	2.35	0.50
1:0:2894:C:O2'	1:0:2895:C:H5'	2.11	0.50
5:B:7:ARG:NH1	5:B:11:LEU:HD22	2.27	0.50
6:C:85:LYS:NZ	37:C:8324:HOH:O	2.38	0.50
8:E:57:LYS:HG3	37:E:358:HOH:O	2.12	0.50
9:F:57:GLU:O	9:F:61:MET:HG3	2.11	0.50
16:M:47:LEU:HD13	16:M:97:VAL:HG11	1.93	0.50
25:V:14:HIS:HB2	25:V:17:ILE:CG1	2.40	0.50
29:Z:25:LYS:HD2	30:1:48:ASP:HA	1.93	0.50
1:0:1634:G:H2'	1:0:1635:U:H6	1.77	0.50
5:B:254:GLN:HG2	5:B:255:GLY:N	2.25	0.50
7:D:57:THR:HG23	7:D:63:ILE:CB	2.42	0.50
27:X:109:LEU:HA	37:X:8572:HOH:O	2.11	0.50
1:0:319:A:H4'	1:0:338:C:C4	2.47	0.50
1:0:558:C:H2'	1:0:559:U:H5''	1.92	0.50
1:0:603:A:H4'	1:0:604:G:O5'	2.12	0.50
1:0:1060:C:H6	1:0:1060:C:H5'	1.77	0.50
1:0:2053:G:OP1	20:Q:138:SER:OG	2.29	0.50
1:0:2748:G:C2'	37:0:6917:HOH:O	2.60	0.50
5:B:217:ARG:HG3	5:B:257:THR:CG2	2.42	0.50
8:E:31:ARG:HH12	8:E:68:HIS:CG	2.30	0.50
8:E:79:GLY:HA3	37:E:7046:HOH:O	2.12	0.50
37:E:2512:HOH:O	12:I:127:ILE:HD11	2.12	0.50
9:F:56:PRO:CG	15:L:44:THR:HA	2.42	0.50
14:K:78:ALA:HB2	37:K:8560:HOH:O	2.12	0.50
21:R:81:ILE:HG23	37:R:4527:HOH:O	2.12	0.50
23:T:44:ARG:CB	37:T:3805:HOH:O	2.60	0.50
27:X:117:LEU:HD23	37:X:8616:HOH:O	2.11	0.50
1:0:858:U:H2'	1:0:859:C:C6	2.46	0.49
1:0:1342:C:O2'	1:0:1343:C:H5'	2.12	0.49
1:0:1943:C:O4'	4:A:212:PRO:HA	2.12	0.49
1:0:2727:A:H2'	1:0:2728:C:H5'	1.93	0.49
1:0:2840:A:H3'	37:0:7023:HOH:O	2.12	0.49
37:0:4530:HOH:O	5:B:216:LYS:HA	2.12	0.49
5:B:132:HIS:CE1	5:B:171:VAL:HG21	2.46	0.49
5:B:195:ARG:NH1	5:B:324:ASP:OD1	2.45	0.49
6:C:55:ARG:NH2	29:Z:56:GLU:OE2	2.31	0.49
7:D:59:GLY:C	7:D:61:PHE:N	2.70	0.49
7:D:154:LYS:HD3	37:D:1796:HOH:O	2.11	0.49
9:F:107:VAL:O	9:F:111:ILE:HG13	2.11	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:J:37:TYR:CE2	13:J:45:PRO:HA	2.46	0.49
31:2:18:GLN:HB3	37:2:8514:HOH:O	2.11	0.49
1:0:105:G:O2'	1:0:106:A:H5'	2.12	0.49
1:0:945:U:H2'	1:0:946:C:C6	2.48	0.49
1:0:1380:U:O4	1:0:2043:U:H4'	2.12	0.49
1:0:1702:U:H5'	37:0:9913:HOH:O	2.12	0.49
1:0:2028:U:H2'	1:0:2029:C:C6	2.47	0.49
1:0:2064:U:H2'	1:0:2065:C:H6	1.78	0.49
37:0:8873:HOH:O	29:Z:1:THR:HA	2.11	0.49
4:A:200:PRO:HG2	4:A:225:VAL:HG21	1.94	0.49
7:D:57:THR:HG23	7:D:63:ILE:CG2	2.40	0.49
11:H:47:GLU:CB	11:H:133:ILE:CD1	2.88	0.49
15:L:27:ARG:O	15:L:30:GLU:N	2.45	0.49
15:L:74:ARG:HD3	15:L:91:ILE:HD12	1.94	0.49
15:L:80:GLY:O	15:L:81:ARG:HD3	2.12	0.49
18:O:80:ARG:HG2	18:O:87:ARG:CZ	2.43	0.49
19:P:46:SER:O	19:P:48:PRO:HD3	2.12	0.49
19:P:93:ARG:HG3	19:P:93:ARG:NH1	2.27	0.49
1:0:359:U:H2'	1:0:360:A:C8	2.47	0.49
1:0:366:U:H2'	1:0:367:G:O4'	2.12	0.49
1:0:392:U:C5'	15:L:193:LYS:HB3	2.43	0.49
1:0:827:A:H2'	1:0:828:G:O4'	2.12	0.49
1:0:2004:U:H2'	1:0:2005:G:OP1	2.12	0.49
1:0:2330:U:H4'	1:0:2331:C:OP1	2.11	0.49
4:A:179:MET:HG2	4:A:186:TRP:CG	2.47	0.49
11:H:117:LYS:O	11:H:119:VAL:HG13	2.13	0.49
13:J:40:THR:O	13:J:41:LYS:C	2.55	0.49
15:L:31:TRP:HA	15:L:34:GLU:HG3	1.94	0.49
15:L:61:ILE:HA	37:L:8626:HOH:O	2.11	0.49
1:0:204:A:H2'	1:0:205:U:H5'	1.93	0.49
1:0:222:A:H2'	1:0:223:G:O4'	2.12	0.49
1:0:682:A:H2'	1:0:683:G:O4'	2.13	0.49
1:0:861:A:H2'	1:0:862:U:C6	2.47	0.49
1:0:920:C:H5''	1:0:921:G:O5'	2.13	0.49
1:0:1116:U:H3	1:0:1246:A:N6	2.01	0.49
1:0:2672:C:OP2	5:B:25:ARG:NH1	2.46	0.49
1:0:2812:A:C2	1:0:2814:A:N6	2.73	0.49
5:B:30:PRO:HB2	5:B:39:GLN:NE2	2.28	0.49
6:C:107:ARG:NE	37:C:8452:HOH:O	2.43	0.49
6:C:246:ARG:NE	37:C:8421:HOH:O	2.44	0.49
9:F:39:SER:HB3	9:F:45:ALA:HB2	1.95	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:H:53:PRO:HG3	11:H:127:GLY:H	1.76	0.49
13:J:49:LEU:HD23	13:J:73:VAL:O	2.12	0.49
27:X:123:VAL:HG12	27:X:124:GLY:O	2.12	0.49
1:0:31:C:H2'	37:0:7063:HOH:O	2.12	0.49
1:0:120:A:H2'	1:0:120:A:N3	2.28	0.49
1:0:635:A:H2'	1:0:636:G:H5''	1.94	0.49
1:0:820:G:C5	4:A:171:LYS:HB2	2.47	0.49
1:0:2300:A:H4'	1:0:2301:A:O5'	2.13	0.49
1:0:2353:A:H4'	1:0:2354:A:O5'	2.12	0.49
1:0:2781:U:H2'	1:0:2782:G:H5'	1.94	0.49
37:0:9267:HOH:O	14:K:41:HIS:HE1	1.95	0.49
2:9:3041:C:O4'	7:D:50:VAL:HG23	2.13	0.49
2:9:3055:U:H4'	2:9:3056:A:H8	1.77	0.49
7:D:95:THR:HG21	7:D:174:VAL:HG22	1.95	0.49
8:E:77:THR:OG1	8:E:78:GLU:N	2.44	0.49
10:G:12:ILE:HG22	10:G:17:GLN:NE2	2.26	0.49
15:L:77:PHE:N	37:L:8528:HOH:O	2.40	0.49
16:M:91:ARG:HG3	16:M:186:LEU:HD23	1.95	0.49
21:R:42:GLU:HG2	21:R:49:VAL:HG23	1.93	0.49
1:0:259:G:O2'	1:0:260:C:H5'	2.13	0.49
1:0:545:G:H2'	1:0:546:C:O4'	2.12	0.49
1:0:1003:U:O2'	11:H:90:PHE:HE1	1.96	0.49
1:0:1496:G:H5'	1:0:1572:A:H1'	1.94	0.49
1:0:2050:G:H5''	20:Q:80:TYR:O	2.13	0.49
2:9:3023:U:C4'	2:9:3024:U:OP2	2.51	0.49
5:B:314:ALA:CB	5:B:317:PRO:HG3	2.42	0.49
6:C:39:GLN:O	6:C:43:LYS:HD3	2.12	0.49
8:E:69:ILE:HA	8:E:72:MET:HE3	1.93	0.49
11:H:136:VAL:HG23	37:H:8330:HOH:O	2.12	0.49
13:J:72:VAL:HG11	13:J:121:PHE:CD1	2.47	0.49
16:M:34:LEU:HA	16:M:47:LEU:HD23	1.93	0.49
16:M:143:ARG:NH1	16:M:173:ASP:OD2	2.37	0.49
17:N:25:VAL:HG23	17:N:26:TRP:N	2.28	0.49
17:N:38:ARG:NH1	37:N:7674:HOH:O	2.44	0.49
24:U:39:ALA:O	24:U:41:GLU:N	2.45	0.49
28:Y:13:ARG:NH1	28:Y:14:PHE:CZ	2.81	0.49
1:0:671:A:O2'	1:0:672:G:H2'	2.13	0.49
1:0:1289:C:O2'	1:0:1290:G:H5'	2.13	0.49
1:0:2526:C:O2'	1:0:2527:U:H5'	2.12	0.49
1:0:2785:C:H4'	1:0:2786:G:OP2	2.13	0.49
37:0:6810:HOH:O	6:C:163:HIS:HE1	1.95	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:A:3:ARG:HB3	4:A:7:GLN:HB2	1.95	0.49
13:J:101:ASN:O	13:J:102:GLU:HB2	2.13	0.49
14:K:24:ALA:HB2	14:K:30:ARG:HD2	1.94	0.49
14:K:145:LEU:O	14:K:148:GLU:HG3	2.13	0.49
15:L:78:ASN:C	15:L:79:LYS:HG2	2.38	0.49
20:Q:32:ALA:O	20:Q:33:ARG:C	2.54	0.49
21:R:33:SER:OG	21:R:36:GLU:HG3	2.12	0.49
26:W:43:VAL:CG1	26:W:44:ASP:N	2.75	0.49
1:O:255:A:H2'	1:O:256:C:C6	2.48	0.49
1:O:417:G:P	37:O:6794:HOH:O	2.71	0.49
1:O:911:G:H5'	1:O:932:U:OP1	2.12	0.49
1:O:935:G:O2'	1:O:936:C:H5'	2.13	0.49
1:O:1488:U:H4'	1:O:1489:G:OP1	2.11	0.49
2:9:3056:A:H1'	7:D:14:ARG:HG2	1.94	0.49
6:C:142:ASP:OD1	6:C:237:GLU:HB3	2.12	0.49
6:C:178:GLN:O	6:C:179:GLY:C	2.56	0.49
14:K:53:ARG:HH22	14:K:57:VAL:HG12	1.77	0.49
16:M:42:HIS:CG	16:M:62:HIS:HE1	2.31	0.49
16:M:139:TRP:CE3	16:M:139:TRP:HA	2.47	0.49
20:Q:29:LYS:CD	37:Q:8543:HOH:O	2.61	0.49
25:V:130:HIS:O	25:V:136:GLY:HA3	2.13	0.49
27:X:200:THR:HG22	27:X:201:GLU:HG3	1.94	0.49
1:O:305:A:C5	1:O:329:A:C2	3.01	0.49
1:O:371:U:H2'	1:O:372:A:C8	2.48	0.49
1:O:738:G:H3'	37:O:6439:HOH:O	2.11	0.49
1:O:1321:A:H2'	1:O:1322:G:C8	2.47	0.49
1:O:1909:A:H2'	1:O:1910:A:H8	1.75	0.49
37:O:5682:HOH:O	18:O:63:ARG:NH2	2.44	0.49
6:C:236:THR:C	37:C:8444:HOH:O	2.55	0.49
23:T:52:THR:HG22	23:T:54:THR:H	1.78	0.49
25:V:38:THR:HG22	25:V:39:ASP:N	2.28	0.49
1:O:1634:G:H2'	1:O:1635:U:C6	2.48	0.49
1:O:1936:C:H3'	37:O:6563:HOH:O	2.12	0.49
1:O:2436:U:H5'	31:2:68:LYS:HE2	1.94	0.49
1:O:2748:G:H2'	37:O:6917:HOH:O	2.13	0.49
4:A:65:ARG:C	4:A:66:ARG:HG3	2.38	0.49
5:B:82:VAL:O	5:B:83:ALA:HB2	2.13	0.49
5:B:211:THR:HA	5:B:255:GLY:O	2.13	0.49
7:D:95:THR:CG2	7:D:174:VAL:HG22	2.43	0.49
15:L:55:LYS:HB2	15:L:60:ILE:CD1	2.43	0.49
16:M:64:SER:C	16:M:66:LEU:H	2.19	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:M:139:TRP:HH2	16:M:176:ARG:HH11	1.59	0.49
16:M:147:ILE:HG23	16:M:148:ALA:N	2.28	0.49
1:0:459:A:H4'	37:0:8963:HOH:O	2.13	0.48
1:0:1189:A:H1'	1:0:1209:C:O4'	2.13	0.48
1:0:2238:A:H3'	37:0:6070:HOH:O	2.12	0.48
1:0:2293:G:C8	1:0:2464:C:C4	3.01	0.48
5:B:26:PHE:HA	37:B:8582:HOH:O	2.13	0.48
5:B:109:LEU:HG	5:B:113:LEU:HD12	1.94	0.48
5:B:248:ARG:O	5:B:251:VAL:HG12	2.13	0.48
7:D:159:PRO:O	7:D:163:VAL:HG23	2.12	0.48
16:M:115:VAL:HG23	16:M:116:PHE:H	1.78	0.48
19:P:40:HIS:CE1	19:P:94:GLN:HA	2.48	0.48
24:U:64:GLY:O	24:U:65:ASP:CB	2.60	0.48
25:V:38:THR:HG22	25:V:39:ASP:H	1.78	0.48
25:V:125:HIS:CD2	25:V:127:GLY:H	2.31	0.48
1:0:1134:G:OP2	11:H:156:THR:HG23	2.13	0.48
1:0:1166:A:H61	1:0:1180:U:H3	1.60	0.48
1:0:1406:A:H4'	1:0:1407:A:H5''	1.95	0.48
1:0:1829:A:H5''	37:0:9574:HOH:O	2.13	0.48
1:0:2091:G:O3'	5:B:235:ARG:HD3	2.12	0.48
1:0:2421:G:H3'	1:0:2422:U:H5''	1.94	0.48
1:0:2563:U:H2'	1:0:2565:C:O5'	2.12	0.48
5:B:7:ARG:HG2	5:B:7:ARG:HH11	1.78	0.48
7:D:84:LEU:C	7:D:86:THR:H	2.21	0.48
9:F:27:GLY:HA3	9:F:101:ALA:O	2.13	0.48
9:F:50:VAL:CG2	9:F:63:ILE:HG21	2.43	0.48
11:H:46:VAL:O	11:H:146:TRP:CH2	2.62	0.48
15:L:137:ASP:O	15:L:142:LYS:HE3	2.12	0.48
23:T:8:TYR:O	23:T:46:ALA:CB	2.61	0.48
25:V:13:MET:HE3	25:V:17:ILE:CG2	2.43	0.48
25:V:88:THR:CG2	25:V:110:GLN:NE2	2.76	0.48
1:0:224:U:H1'	37:0:9387:HOH:O	2.12	0.48
1:0:383:A:H4'	37:0:4771:HOH:O	2.13	0.48
1:0:1594:C:OP2	18:O:120:ARG:HD2	2.13	0.48
1:0:1825:U:O2'	1:0:1826:C:H5'	2.12	0.48
1:0:2630:G:O6	4:A:206:ARG:NH2	2.46	0.48
1:0:2712:G:H5'	37:0:4667:HOH:O	2.13	0.48
5:B:36:PRO:HD3	5:B:169:GLY:H	1.77	0.48
7:D:23:VAL:HG12	7:D:130:VAL:HG22	1.95	0.48
11:H:153:VAL:HA	37:H:8337:HOH:O	2.12	0.48
16:M:5:ARG:HG3	19:P:18:PRO:CB	2.43	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:N:97:SER:HB3	17:N:100:GLN:HE21	1.78	0.48
20:Q:39:THR:O	20:Q:40:ALA:C	2.55	0.48
23:T:36:CYS:O	23:T:37:GLU:C	2.56	0.48
27:X:189:ASN:HD22	27:X:191:ASP:N	2.11	0.48
1:0:962:C:C1'	16:M:5:ARG:NH1	2.73	0.48
1:0:1184:C:O2'	1:0:1185:U:P	2.72	0.48
1:0:1446:U:H4'	1:0:1447:U:OP2	2.13	0.48
1:0:1495:C:H1'	1:0:1573:A:H1'	1.95	0.48
1:0:1778:A:H2'	1:0:1779:A:H5'	1.95	0.48
1:0:2634:G:O2'	1:0:2635:A:H5'	2.14	0.48
1:0:2851:G:C2'	1:0:2852:A:H5'	2.43	0.48
2:9:3061:C:H2'	2:9:3062:A:C8	2.48	0.48
6:C:126:ASP:C	6:C:128:GLY:N	2.70	0.48
9:F:52:GLU:HG3	9:F:77:VAL:O	2.13	0.48
16:M:38:LYS:HD2	16:M:114:LYS:HE3	1.95	0.48
22:S:19:ARG:NH1	22:S:68:ASP:O	2.47	0.48
29:Z:2:GLY:O	29:Z:7:SER:OG	2.25	0.48
1:0:195:C:H2'	1:0:196:G:H5'	1.95	0.48
1:0:308:U:H5'	22:S:97:ARG:NH2	2.28	0.48
1:0:2364:A:H5''	19:P:15:LYS:HD3	1.95	0.48
1:0:2453:G:H4'	14:K:50:GLY:C	2.39	0.48
37:0:9562:HOH:O	20:Q:83:LYS:HB3	2.13	0.48
6:C:85:LYS:CE	37:C:8324:HOH:O	2.62	0.48
10:G:64:ASN:N	10:G:64:ASN:ND2	2.60	0.48
11:H:14:TYR:N	11:H:91:HIS:CE1	2.77	0.48
11:H:113:ALA:N	11:H:114:PRO:HD3	2.28	0.48
12:I:6:PHE:O	12:I:8:ALA:N	2.47	0.48
24:U:16:ARG:NH1	24:U:65:ASP:O	2.45	0.48
30:1:41:HIS:O	30:1:45:ASN:HB2	2.12	0.48
1:0:2015:A:H2'	1:0:2016:U:O4'	2.13	0.48
1:0:2135:A:O4'	1:0:2243:C:N4	2.47	0.48
1:0:2764:C:O2'	1:0:2765:C:H5'	2.13	0.48
1:0:2830:U:H4'	37:0:9584:HOH:O	2.13	0.48
5:B:185:GLY:HA2	37:B:8632:HOH:O	2.13	0.48
11:H:166:ASN:N	11:H:166:ASN:ND2	2.61	0.48
12:I:27:ALA:HB1	12:I:87:LEU:CD2	2.44	0.48
25:V:21:LEU:HD21	25:V:48:VAL:HG13	1.94	0.48
25:V:90:TYR:CE2	25:V:99:ALA:HB2	2.49	0.48
1:0:588:G:O6	25:V:154:ARG:NH1	2.46	0.48
1:0:1675:C:H5''	30:1:5:LYS:HD2	1.96	0.48
1:0:2270:G:O3'	4:A:223:ARG:NH1	2.47	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:2735:U:H2'	1:0:2736:U:C6	2.48	0.48
37:0:5336:HOH:O	15:L:189:VAL:HG23	2.13	0.48
37:0:5661:HOH:O	23:T:56:ARG:HD3	2.13	0.48
2:9:3008:G:O6	16:M:11:ARG:NH1	2.47	0.48
6:C:46:TYR:CE2	6:C:98:ARG:NH1	2.81	0.48
8:E:158:ASP:HA	37:E:2712:HOH:O	2.12	0.48
11:H:132:PHE:O	11:H:133:ILE:HD13	2.13	0.48
12:I:60:ARG:HD3	12:I:71:TYR:CE1	2.48	0.48
17:N:84:THR:O	17:N:88:LYS:HG3	2.13	0.48
21:R:80:ARG:HG2	37:R:4527:HOH:O	2.14	0.48
25:V:1:MET:N	25:V:103:GLU:OE2	2.43	0.48
1:0:35:U:H5'	6:C:47:GLY:O	2.14	0.48
1:0:134:U:C2	1:0:145:A:C2	3.02	0.48
1:0:907:A:H2'	1:0:908:A:C8	2.47	0.48
1:0:1123:A:C2	1:0:1129:C:H4'	2.49	0.48
1:0:1878:G:O2'	1:0:1879:U:O4'	2.31	0.48
5:B:227:HIS:CD2	37:B:8531:HOH:O	2.66	0.48
5:B:329:TYR:HE2	23:T:15:PRO:HG2	1.78	0.48
6:C:20:ASP:HB2	37:C:8392:HOH:O	2.13	0.48
6:C:129:HIS:CE1	6:C:232:LEU:H	2.32	0.48
8:E:137:ASP:O	8:E:141:VAL:HG23	2.14	0.48
15:L:59:GLY:HA3	15:L:141:ILE:HD12	1.96	0.48
15:L:134:ILE:HG23	15:L:141:ILE:HD13	1.96	0.48
16:M:139:TRP:CH2	16:M:176:ARG:NH1	2.82	0.48
18:O:84:ALA:C	18:O:86:ALA:H	2.20	0.48
27:X:115:ARG:NE	37:X:8556:HOH:O	2.47	0.48
31:2:34:LYS:HB2	31:2:37:ASP:OD2	2.14	0.48
1:0:1191:A:H3'	1:0:1192:A:C5'	2.43	0.48
1:0:1305:C:H5'	37:0:9330:HOH:O	2.13	0.48
1:0:1613:C:H2'	1:0:1614:G:O4'	2.13	0.48
1:0:1894:C:C2	1:0:1939:U:C4	3.01	0.48
5:B:195:ARG:N	5:B:198:GLU:OE1	2.42	0.48
6:C:238:SER:O	37:C:8378:HOH:O	2.20	0.48
7:D:144:ARG:NH2	37:D:3839:HOH:O	2.47	0.48
8:E:23:GLU:HG2	8:E:28:SER:HB2	1.95	0.48
16:M:37:ARG:NH2	37:M:3863:HOH:O	2.46	0.48
21:R:80:ARG:NH1	37:R:7263:HOH:O	2.46	0.48
23:T:44:ARG:HB3	37:T:3805:HOH:O	2.12	0.48
30:1:10:ARG:HD2	30:1:49:GLU:OE2	2.13	0.48
1:0:1250:C:O2'	1:0:1251:C:H5'	2.14	0.48
1:0:1600:G:OP2	1:0:1600:G:H8	1.97	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:A:169:PHE:O	4:A:170:VAL:HB	2.13	0.48
9:F:28:ALA:HB3	9:F:99:THR:HG23	1.95	0.48
11:H:31:PHE:HA	11:H:85:ILE:CG2	2.44	0.48
11:H:150:LYS:CD	37:H:8368:HOH:O	2.60	0.48
15:L:37:VAL:CG1	15:L:63:VAL:HG11	2.44	0.48
16:M:20:TYR:N	37:M:4363:HOH:O	2.33	0.48
16:M:77:ASN:OD1	16:M:80:SER:HB2	2.13	0.48
17:N:47:ARG:NH2	37:N:510:HOH:O	2.47	0.48
19:P:11:ARG:NH1	37:P:5620:HOH:O	2.46	0.48
29:Z:5:THR:HB	29:Z:6:PRO:CD	2.44	0.48
1:O:1215:A:O3'	1:O:1216:G:H4'	2.14	0.47
1:O:2073:G:C6	1:O:2607:U:C2	3.02	0.47
5:B:102:THR:HG22	37:B:8613:HOH:O	2.13	0.47
6:C:107:ARG:NH2	37:C:8452:HOH:O	2.44	0.47
11:H:126:HIS:O	11:H:127:GLY:C	2.55	0.47
21:R:81:ILE:HA	37:R:6969:HOH:O	2.14	0.47
1:O:335:U:H4'	22:S:92:ASP:OD2	2.14	0.47
1:O:474:C:O3'	6:C:73:LEU:HD21	2.14	0.47
1:O:941:G:C5	1:O:942:U:C4	3.03	0.47
1:O:1269:G:H2'	1:O:1270:U:C6	2.49	0.47
1:O:1439:C:H5''	30:1:41:HIS:CE1	2.49	0.47
1:O:1730:G:H5'	1:O:1731:C:C5	2.49	0.47
1:O:2039:A:H4'	1:O:2760:C:O2'	2.15	0.47
1:O:2254:G:C1'	37:O:4974:HOH:O	2.58	0.47
1:O:2717:C:O2'	1:O:2718:C:H5'	2.14	0.47
5:B:304:PRO:HD2	5:B:307:ARG:CD	2.40	0.47
6:C:37:ALA:O	6:C:41:ASN:ND2	2.47	0.47
7:D:23:VAL:O	7:D:23:VAL:CG2	2.62	0.47
9:F:20:LEU:O	9:F:23:ALA:HB3	2.13	0.47
13:J:37:TYR:HD2	37:J:7169:HOH:O	1.98	0.47
16:M:143:ARG:HA	16:M:172:PHE:CE2	2.48	0.47
19:P:28:ARG:HG2	37:P:4350:HOH:O	2.14	0.47
20:Q:33:ARG:NH2	37:Q:8534:HOH:O	2.46	0.47
22:S:20:HIS:HB3	22:S:41:ARG:HD2	1.96	0.47
25:V:26:ILE:O	25:V:26:ILE:CG1	2.61	0.47
26:W:31:ILE:O	26:W:35:GLU:HG3	2.14	0.47
26:W:75:ALA:O	26:W:83:ALA:HA	2.13	0.47
26:W:76:ARG:O	26:W:77:PHE:HB3	2.14	0.47
1:O:213:G:C1'	1:O:214:U:OP2	2.61	0.47
1:O:482:G:H4'	1:O:508:A:N1	2.29	0.47
1:O:485:A:H4'	1:O:486:A:OP1	2.14	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:1055:G:OP2	11:H:94:ARG:NH1	2.48	0.47
1:0:1351:G:OP1	6:C:96:LYS:NZ	2.42	0.47
1:0:1666:C:C2'	1:0:1667:A:C5'	2.92	0.47
1:0:1805:G:H2'	1:0:1806:G:H8	1.78	0.47
1:0:2831:C:H2'	1:0:2832:C:H5'	1.96	0.47
2:9:3049:G:O2'	2:9:3050:G:H5'	2.14	0.47
8:E:32:ARG:C	8:E:33:LEU:HD23	2.39	0.47
11:H:71:TYR:O	11:H:73:GLN:N	2.48	0.47
16:M:37:ARG:CZ	37:M:3863:HOH:O	2.62	0.47
16:M:165:ALA:HA	37:M:2052:HOH:O	2.14	0.47
20:Q:39:THR:HG22	20:Q:42:GLU:N	2.16	0.47
25:V:76:ASP:O	25:V:77:ALA:C	2.57	0.47
27:X:110:SER:HB2	37:X:8557:HOH:O	2.13	0.47
1:0:204:A:C2'	1:0:205:U:H5'	2.44	0.47
1:0:450:C:H4'	6:C:46:TYR:CE1	2.50	0.47
1:0:521:A:C2'	1:0:522:U:H5'	2.45	0.47
1:0:816:G:C6	1:0:817:G:N1	2.82	0.47
1:0:1421:C:O2'	1:0:1422:U:H5'	2.13	0.47
1:0:1855:G:O6	4:A:142:SER:HB3	2.14	0.47
1:0:2672:C:O2'	1:0:2673:U:H5'	2.14	0.47
5:B:16:ARG:HB3	5:B:217:ARG:NH2	2.29	0.47
5:B:329:TYR:CE2	23:T:15:PRO:HG2	2.49	0.47
6:C:3:ALA:HA	37:C:8448:HOH:O	2.13	0.47
8:E:146:ALA:O	8:E:147:ASP:C	2.57	0.47
10:G:12:ILE:HG22	10:G:12:ILE:O	2.14	0.47
15:L:122:GLU:OE2	15:L:127:LYS:HE2	2.15	0.47
15:L:134:ILE:O	15:L:136:PRO:HD3	2.14	0.47
16:M:62:HIS:O	16:M:65:ASP:OD1	2.32	0.47
16:M:182:GLY:N	37:M:7390:HOH:O	2.37	0.47
22:S:48:VAL:HG22	22:S:98:VAL:HA	1.97	0.47
22:S:113:GLU:O	22:S:114:SER:C	2.56	0.47
29:Z:25:LYS:CD	30:1:49:GLU:H	2.25	0.47
30:1:16:ASN:C	30:1:18:ASN:N	2.72	0.47
1:0:907:A:H2'	1:0:908:A:H8	1.80	0.47
1:0:1269:G:H2'	1:0:1270:U:H6	1.79	0.47
1:0:1297:U:H1'	37:0:9876:HOH:O	2.12	0.47
1:0:1524:U:O2'	1:0:1525:G:O5'	2.31	0.47
1:0:1699:C:OP2	37:0:5097:HOH:O	2.20	0.47
1:0:1790:C:H2'	1:0:1791:U:C6	2.48	0.47
1:0:2502:C:H2'	1:0:2503:A:H5'	1.96	0.47
1:0:2591:C:H2'	1:0:2592:G:O4'	2.13	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:2659:U:H4'	20:Q:76:ASP:HB3	1.96	0.47
5:B:55:ASN:HB3	5:B:64:GLY:H	1.80	0.47
5:B:275:GLY:O	5:B:291:ASP:HA	2.14	0.47
6:C:21:VAL:C	6:C:23:GLU:H	2.23	0.47
7:D:94:ALA:O	7:D:95:THR:O	2.33	0.47
12:I:51:GLU:O	12:I:55:GLU:HG3	2.15	0.47
15:L:139:PRO:C	15:L:141:ILE:H	2.22	0.47
18:O:7:LYS:HD3	18:O:21:VAL:CG2	2.44	0.47
25:V:13:MET:HE3	25:V:17:ILE:HG22	1.96	0.47
26:W:23:HIS:HB2	37:W:7830:HOH:O	2.14	0.47
1:0:166:A:N7	14:K:25:GLY:HA2	2.29	0.47
1:0:955:A:H2'	1:0:956:G:O4'	2.15	0.47
1:0:1086:A:C6	25:V:11:VAL:HG11	2.48	0.47
1:0:1707:G:N2	1:0:1709:G:H3'	2.30	0.47
1:0:1926:G:H2'	1:0:1927:A:H8	1.80	0.47
1:0:2451:G:O2'	31:2:38:ARG:NH2	2.47	0.47
1:0:2478:U:O2'	1:0:2479:A:H5'	2.15	0.47
1:0:2837:U:H1'	5:B:307:ARG:HH12	1.78	0.47
37:9:8526:HOH:O	16:M:107:ASN:HB3	2.15	0.47
5:B:33:ASP:O	5:B:34:GLY:O	2.33	0.47
6:C:79:ARG:O	6:C:87:ARG:HG2	2.15	0.47
11:H:48:LEU:HD13	11:H:146:TRP:HB3	1.96	0.47
11:H:74:ASN:ND2	11:H:141:ASN:OD1	2.47	0.47
12:I:74:ARG:NH1	12:I:76:ASP:HB2	2.30	0.47
12:I:97:ALA:O	12:I:101:VAL:HG23	2.15	0.47
13:J:125:ALA:C	13:J:127:ALA:H	2.22	0.47
14:K:120:LEU:HD12	14:K:133:VAL:HG21	1.96	0.47
15:L:49:ALA:C	15:L:54:TYR:HB3	2.40	0.47
15:L:172:GLY:C	15:L:183:VAL:HG11	2.39	0.47
16:M:151:ASP:OD2	16:M:165:ALA:O	2.32	0.47
20:Q:114:VAL:HG13	20:Q:114:VAL:O	2.13	0.47
26:W:70:ILE:HG23	26:W:70:ILE:O	2.15	0.47
1:0:275:G:C2	1:0:376:C:N3	2.82	0.47
1:0:299:U:H5'	37:0:6718:HOH:O	2.15	0.47
1:0:484:A:N1	1:0:506:G:H4'	2.30	0.47
1:0:653:C:H2'	1:0:654:A:C8	2.50	0.47
1:0:1116:U:H1'	37:0:3990:HOH:O	2.14	0.47
1:0:1175:G:H1'	1:0:1193:A:H2'	1.97	0.47
1:0:1482:A:H1'	37:0:8932:HOH:O	2.13	0.47
1:0:1743:G:N7	37:0:8775:HOH:O	2.36	0.47
1:0:2687:G:O2'	1:0:2688:U:H5'	2.14	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:2833:C:C2	1:0:2848:G:N2	2.83	0.47
37:0:9267:HOH:O	14:K:41:HIS:CE1	2.68	0.47
4:A:220:PRO:HD2	4:A:223:ARG:HD3	1.96	0.47
5:B:312:ARG:HD3	5:B:315:VAL:HG13	1.97	0.47
5:B:320:GLN:HG3	5:B:321:PRO:CD	2.44	0.47
6:C:54:LEU:HD23	6:C:79:ARG:HG3	1.97	0.47
6:C:95:GLU:H	6:C:95:GLU:CD	2.20	0.47
6:C:150:THR:HA	6:C:203:ALA:O	2.15	0.47
10:G:66:LEU:O	10:G:69:ARG:HB3	2.15	0.47
15:L:109:PHE:HB3	15:L:112:LEU:HD12	1.96	0.47
18:O:10:ALA:HA	18:O:13:VAL:HG12	1.97	0.47
22:S:51:LEU:HD11	22:S:97:ARG:HB2	1.97	0.47
25:V:72:PRO:HB2	25:V:74:GLU:O	2.14	0.47
1:0:21:G:H4'	20:Q:2:ILE:HG22	1.97	0.47
1:0:119:A:H2'	1:0:120:A:H5''	1.97	0.47
1:0:400:C:O2'	1:0:401:C:H5'	2.15	0.47
1:0:1246:A:O2'	1:0:1247:A:H3'	2.15	0.47
1:0:1249:U:H2'	1:0:1250:C:C6	2.50	0.47
1:0:1266:U:H4'	27:X:115:ARG:HH21	1.78	0.47
1:0:1517:U:C2	1:0:1670:G:N2	2.83	0.47
1:0:1681:G:H5''	1:0:1682:A:H5'	1.95	0.47
1:0:2661:U:H3	1:0:2812:A:H62	1.63	0.47
2:9:3047:A:C2	2:9:3048:C:C2	3.01	0.47
5:B:162:MET:HE2	5:B:310:ARG:CG	2.45	0.47
6:C:104:ASP:HA	6:C:107:ARG:NH1	2.26	0.47
7:D:146:LYS:CE	16:M:107:ASN:ND2	2.78	0.47
11:H:117:LYS:HB2	37:H:8326:HOH:O	2.14	0.47
16:M:182:GLY:O	16:M:183:ASP:O	2.33	0.47
27:X:197:ASP:OD1	27:X:199:ASP:HB2	2.15	0.47
28:Y:26:VAL:O	28:Y:30:GLU:HG3	2.14	0.47
28:Y:33:HIS:CE1	28:Y:49:ARG:NE	2.83	0.47
1:0:541:C:O2'	1:0:542:A:H5''	2.15	0.47
1:0:1189:A:H1'	1:0:1209:C:H1'	1.97	0.47
1:0:1593:C:OP1	18:O:117:SER:HB3	2.15	0.47
1:0:1743:G:H1'	37:0:4345:HOH:O	2.15	0.47
1:0:2054:A:C2	20:Q:128:ARG:NH2	2.83	0.47
1:0:2506:A:N6	1:0:2511:A:O2'	2.48	0.47
1:0:2676:C:H4'	12:I:70:PHE:HE1	1.80	0.47
4:A:164:ARG:HB2	28:Y:68:CYS:SG	2.54	0.47
5:B:4:SER:O	5:B:5:ARG:HB2	2.15	0.47
11:H:151:MET:HA	11:H:151:MET:CE	2.39	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:J:99:ASP:OD1	13:J:99:ASP:C	2.57	0.47
14:K:1:THR:HB	14:K:6:ARG:NH1	2.30	0.47
1:0:236:A:H8	1:0:236:A:OP1	1.98	0.47
1:0:935:G:H4'	17:N:38:ARG:HH12	1.80	0.47
1:0:1182:C:H1'	1:0:1192:A:H8	1.80	0.47
1:0:2001:G:C2'	1:0:2002:C:H5'	2.45	0.47
1:0:2093:G:H5''	37:B:8528:HOH:O	2.15	0.47
1:0:2455:A:H2'	1:0:2456:A:O4'	2.15	0.47
1:0:2761:A:C4	1:0:2763:G:C8	3.02	0.47
7:D:166:ILE:HB	37:D:6326:HOH:O	2.13	0.47
17:N:47:ARG:NH1	37:N:4564:HOH:O	2.47	0.47
1:0:515:C:H5	37:0:6485:HOH:O	1.98	0.46
1:0:566:A:H2'	1:0:567:U:O4'	2.15	0.46
1:0:890:C:O2'	1:0:891:G:H5'	2.15	0.46
1:0:894:A:H1'	37:0:4672:HOH:O	2.14	0.46
1:0:952:G:OP1	19:P:42:LYS:HE2	2.15	0.46
1:0:1333:U:H2'	1:0:1334:C:H6	1.80	0.46
1:0:1594:C:C5	18:O:120:ARG:NH1	2.83	0.46
1:0:1794:G:H3'	37:0:9997:HOH:O	2.14	0.46
1:0:1878:G:H5''	37:0:4614:HOH:O	2.14	0.46
1:0:2507:G:H2'	1:0:2510:C:H42	1.80	0.46
1:0:2779:G:H21	8:E:143:GLN:NE2	2.12	0.46
37:0:7213:HOH:O	4:A:190:ARG:HG3	2.14	0.46
5:B:144:THR:HG22	5:B:145:HIS:N	2.30	0.46
6:C:98:ARG:NH1	37:C:8357:HOH:O	2.47	0.46
7:D:25:MET:SD	7:D:40:ILE:HD11	2.55	0.46
9:F:50:VAL:HG13	9:F:60:VAL:HG11	1.97	0.46
12:I:37:ALA:HA	12:I:102:ARG:O	2.15	0.46
13:J:132:VAL:C	37:J:3160:HOH:O	2.57	0.46
15:L:46:LEU:HG	37:L:8624:HOH:O	2.15	0.46
16:M:180:LEU:O	16:M:181:ASP:CB	2.62	0.46
17:N:39:THR:HB	37:N:3360:HOH:O	2.14	0.46
27:X:106:THR:HG22	27:X:107:PRO:O	2.15	0.46
27:X:235:GLU:CD	27:X:235:GLU:N	2.67	0.46
1:0:95:A:H5''	1:0:97:G:O4'	2.15	0.46
1:0:135:G:OP1	15:L:39:ARG:NH1	2.47	0.46
1:0:626:U:C4	1:0:627:G:C6	3.02	0.46
1:0:745:G:N2	14:K:67:ARG:HD2	2.31	0.46
1:0:1574:C:O5'	1:0:1574:C:H6	1.98	0.46
1:0:1771:U:H4'	1:0:1772:C:OP2	2.15	0.46
1:0:2316:G:H4'	37:0:5510:HOH:O	2.15	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:2421:G:H3'	1:0:2422:U:C5'	2.44	0.46
1:0:2656:G:O2'	1:0:2657:G:H5'	2.16	0.46
1:0:2670:G:O2'	1:0:2671:U:H5'	2.14	0.46
1:0:2688:U:H2'	1:0:2689:A:C8	2.50	0.46
1:0:2769:C:H2'	1:0:2770:G:C5'	2.45	0.46
2:9:3036:C:C5	2:9:3037:C:C5	3.03	0.46
5:B:279:THR:HG22	5:B:280:VAL:N	2.31	0.46
7:D:19:GLU:O	7:D:133:ASN:HB3	2.15	0.46
7:D:41:LEU:CA	7:D:44:ILE:HG22	2.44	0.46
11:H:26:LYS:HD3	11:H:89:PRO:CG	2.45	0.46
12:I:107:ASN:HD22	12:I:108:PRO:N	2.13	0.46
13:J:9:THR:O	13:J:10:GLN:C	2.57	0.46
14:K:130:ARG:O	14:K:131:GLU:C	2.58	0.46
15:L:20:ILE:HG22	37:L:8591:HOH:O	2.15	0.46
15:L:71:SER:O	15:L:73:ARG:NH1	2.48	0.46
19:P:87:THR:HB	37:P:1295:HOH:O	2.14	0.46
20:Q:25:PHE:CE2	20:Q:29:LYS:CE	2.98	0.46
24:U:16:ARG:NH2	24:U:63:GLU:HG3	2.30	0.46
1:0:29:C:O2'	1:0:30:U:H5'	2.15	0.46
1:0:1118:A:C8	1:0:1119:G:H5''	2.50	0.46
1:0:1262:C:H1'	25:V:120:PRO:HG3	1.97	0.46
1:0:1308:A:O4'	6:C:226:GLY:HA3	2.15	0.46
8:E:84:MET:HG2	8:E:168:ILE:HD13	1.97	0.46
9:F:26:THR:HG21	9:F:103:ALA:HB2	1.97	0.46
13:J:124:VAL:HG23	37:J:2659:HOH:O	2.14	0.46
24:U:1:THR:C	24:U:3:LEU:H	2.22	0.46
25:V:26:ILE:HG22	37:V:5420:HOH:O	2.14	0.46
27:X:212:ARG:HB3	37:X:8535:HOH:O	2.14	0.46
29:Z:25:LYS:HG3	30:1:49:GLU:H	1.79	0.46
1:0:243:A:H61	1:0:269:G:H1'	1.80	0.46
1:0:391:U:H2'	1:0:392:U:C6	2.51	0.46
1:0:491:C:O2'	1:0:492:C:H5'	2.16	0.46
1:0:585:C:H6	37:0:5513:HOH:O	1.99	0.46
1:0:1462:C:H2'	1:0:1463:A:C8	2.50	0.46
1:0:1878:G:O2'	1:0:1879:U:H6	1.98	0.46
1:0:2324:G:H4'	1:0:2418:G:O2'	2.15	0.46
2:9:3091:C:H2'	2:9:3092:G:O4'	2.16	0.46
4:A:36:ASP:O	4:A:37:VAL:C	2.58	0.46
5:B:252:PRO:HA	37:B:8572:HOH:O	2.16	0.46
7:D:35:ALA:C	7:D:37:ALA:N	2.73	0.46
8:E:11:VAL:CG1	8:E:12:ASP:H	2.29	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:E:24:GLY:HA3	8:E:76:VAL:HB	1.97	0.46
8:E:125:GLU:O	8:E:132:THR:HG22	2.15	0.46
10:G:12:ILE:CD1	37:G:692:HOH:O	2.63	0.46
11:H:35:ASN:HD21	11:H:80:ASN:HA	1.79	0.46
14:K:21:ARG:N	37:K:8532:HOH:O	2.48	0.46
16:M:37:ARG:HA	16:M:37:ARG:HD3	1.83	0.46
25:V:14:HIS:HA	37:V:2978:HOH:O	2.16	0.46
26:W:71:ARG:HD3	37:W:2171:HOH:O	2.16	0.46
1:0:1076:G:H1'	37:0:3913:HOH:O	2.15	0.46
1:0:1414:A:H2'	1:0:1415:G:O4'	2.16	0.46
1:0:1787:C:O2'	1:0:1788:U:H5'	2.16	0.46
1:0:1878:G:O2'	1:0:1879:U:O5'	2.34	0.46
1:0:2064:U:H5'	1:0:2652:U:H4'	1.97	0.46
1:0:2064:U:H2'	1:0:2065:C:C6	2.49	0.46
1:0:2081:A:H4'	12:I:69:TYR:CE1	2.50	0.46
1:0:2270:G:H4'	4:A:223:ARG:HH12	1.81	0.46
2:9:3056:A:C3'	2:9:3057:A:H5''	2.45	0.46
5:B:279:THR:CG2	5:B:280:VAL:N	2.79	0.46
12:I:126:ASN:O	12:I:129:PHE:HE2	1.98	0.46
19:P:26:PRO:O	19:P:27:GLN:C	2.57	0.46
20:Q:39:THR:N	20:Q:42:GLU:OE1	2.41	0.46
22:S:23:VAL:CA	22:S:93:THR:HG21	2.46	0.46
24:U:1:THR:HG23	24:U:2:VAL:N	2.18	0.46
1:0:151:A:H2'	1:0:152:A:O4'	2.15	0.46
1:0:423:A:O2'	1:0:424:C:H5'	2.15	0.46
1:0:779:U:H5'	1:0:1836:A:C2	2.50	0.46
1:0:832:U:H2'	1:0:833:G:H8	1.79	0.46
1:0:947:U:H2'	1:0:948:G:C8	2.51	0.46
1:0:1804:A:H2'	1:0:1805:G:H8	1.78	0.46
1:0:2073:G:OP2	1:0:2490:A:H5'	2.15	0.46
1:0:2407:G:O2'	1:0:2408:A:H5'	2.16	0.46
1:0:2720:C:O2	13:J:87:ARG:NH2	2.49	0.46
37:0:6120:HOH:O	19:P:2:SER:HA	2.15	0.46
5:B:243:ASN:HA	5:B:244:PRO:C	2.41	0.46
6:C:1:MET:HG2	6:C:2:GLN:N	2.30	0.46
6:C:142:ASP:OD1	6:C:236:THR:HG23	2.16	0.46
7:D:86:THR:C	7:D:89:PRO:HD2	2.40	0.46
12:I:39:VAL:CG1	12:I:107:ASN:HB2	2.46	0.46
15:L:5:TYR:O	15:L:8:ILE:N	2.49	0.46
15:L:113:ARG:NH2	15:L:156:ARG:HG2	2.30	0.46
16:M:115:VAL:HG23	16:M:116:PHE:N	2.31	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:S:44:ALA:HA	22:S:62:VAL:HG12	1.98	0.46
22:S:101:LEU:HD13	22:S:112:LEU:HD11	1.97	0.46
25:V:4:LEU:HD22	25:V:52:VAL:HG22	1.95	0.46
25:V:68:THR:HG23	25:V:69:ARG:N	2.30	0.46
25:V:133:LYS:NZ	37:V:4031:HOH:O	2.48	0.46
1:0:849:C:C2'	1:0:850:U:H5'	2.46	0.46
1:0:2251:G:H2'	1:0:2252:A:C8	2.51	0.46
4:A:70:ALA:HA	4:A:71:PRO:HD3	1.77	0.46
6:C:7:ASP:O	6:C:9:ASP:N	2.49	0.46
9:F:47:LEU:HB2	9:F:108:LEU:HD11	1.98	0.46
11:H:75:SER:HB3	11:H:79:ALA:HB1	1.98	0.46
13:J:89:LYS:HA	37:J:7064:HOH:O	2.16	0.46
16:M:115:VAL:HG23	37:M:6448:HOH:O	2.16	0.46
17:N:25:VAL:O	17:N:29:VAL:HG23	2.15	0.46
17:N:26:TRP:CE3	17:N:26:TRP:HA	2.51	0.46
30:1:9:LYS:O	30:1:12:ALA:HB3	2.15	0.46
1:0:595:U:O2'	1:0:596:C:H5'	2.16	0.46
1:0:695:C:O2'	1:0:696:C:H5'	2.16	0.46
1:0:1192:A:O2'	1:0:1193:A:P	2.73	0.46
1:0:1311:G:C2	1:0:1312:G:C8	3.04	0.46
1:0:2587:U:C2	1:0:2589:U:H5'	2.51	0.46
37:0:6835:HOH:O	6:C:188:ARG:CD	2.64	0.46
37:0:9840:HOH:O	19:P:16:ASN:HB2	2.15	0.46
5:B:180:ASP:O	5:B:181:ILE:C	2.58	0.46
6:C:7:ASP:C	6:C:9:ASP:H	2.24	0.46
8:E:145:ALA:O	8:E:148:ILE:HB	2.16	0.46
9:F:13:GLU:O	9:F:16:ALA:HB3	2.16	0.46
12:I:131:THR:CG2	12:I:134:GLU:HG3	2.46	0.46
13:J:6:ALA:HB3	13:J:116:GLU:HG2	1.98	0.46
16:M:93:GLN:CG	37:M:6239:HOH:O	2.63	0.46
37:M:4624:HOH:O	19:P:19:ARG:HD2	2.15	0.46
20:Q:25:PHE:CZ	20:Q:29:LYS:HE2	2.51	0.46
27:X:112:GLU:CD	27:X:115:ARG:NH1	2.73	0.46
31:2:42:ARG:HD2	37:2:8507:HOH:O	2.15	0.46
1:0:159:G:H5''	15:L:74:ARG:NH2	2.31	0.46
1:0:449:A:N7	6:C:43:LYS:HG2	2.31	0.46
1:0:559:U:O2'	1:0:560:C:H5'	2.16	0.46
1:0:644:G:H5'	1:0:644:G:N3	2.31	0.46
1:0:800:G:H4'	37:0:6447:HOH:O	2.15	0.46
1:0:843:A:C2	1:0:846:A:C8	3.04	0.46
1:0:1641:A:C2'	1:0:1642:A:H5'	2.45	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:1766:U:O4'	1:0:1779:A:N6	2.48	0.46
1:0:1947:G:N2	1:0:1966:U:O2	2.49	0.46
1:0:2269:C:H2'	1:0:2270:G:C5'	2.46	0.46
37:0:9688:HOH:O	14:K:4:LYS:HG3	2.14	0.46
5:B:125:GLU:OE2	5:B:129:ARG:NH1	2.48	0.46
5:B:238:ASN:HA	37:B:8522:HOH:O	2.14	0.46
5:B:307:ARG:HH11	5:B:307:ARG:HB2	1.80	0.46
7:D:141:VAL:HG13	7:D:144:ARG:HH21	1.80	0.46
7:D:146:LYS:HE2	16:M:107:ASN:ND2	2.31	0.46
8:E:108:LEU:HD12	8:E:108:LEU:N	2.31	0.46
15:L:35:PRO:CD	15:L:38:VAL:HG23	2.46	0.46
15:L:81:ARG:HG3	15:L:85:ARG:HB2	1.97	0.46
15:L:162:GLY:HA2	37:L:8519:HOH:O	2.16	0.46
16:M:151:ASP:CG	16:M:165:ALA:O	2.59	0.46
16:M:171:HIS:CE1	37:M:6988:HOH:O	2.69	0.46
22:S:37:GLN:OE1	22:S:118:SER:HA	2.16	0.46
1:0:195:C:C5	1:0:196:G:C5	3.04	0.46
1:0:240:C:C4'	15:L:146:GLN:NE2	2.78	0.46
1:0:295:C:H2'	1:0:296:G:O4'	2.16	0.46
1:0:431:G:P	15:L:48:ARG:HH12	2.39	0.46
1:0:602:A:O2'	1:0:605:C:H4'	2.15	0.46
1:0:1116:U:O2'	1:0:1118:A:C2	2.41	0.46
1:0:1162:G:H2'	1:0:1162:G:N3	2.31	0.46
1:0:1329:A:H2	37:0:4139:HOH:O	1.98	0.46
1:0:1420:C:C2	1:0:1445:G:N2	2.84	0.46
1:0:1496:G:H4'	37:0:5092:HOH:O	2.15	0.46
1:0:1787:C:OP1	18:O:68:LYS:HE2	2.16	0.46
1:0:2748:G:C5'	37:0:6917:HOH:O	2.64	0.46
37:0:3245:HOH:O	22:S:9:LYS:CD	2.51	0.46
2:9:3041:C:H2'	2:9:3042:C:H6	1.81	0.46
5:B:54:VAL:O	5:B:55:ASN:C	2.58	0.46
5:B:154:VAL:CG1	5:B:156:LYS:HG2	2.46	0.46
5:B:294:TYR:HE2	37:B:8649:HOH:O	1.99	0.46
7:D:92:GLU:O	7:D:93:LEU:O	2.33	0.46
8:E:98:GLU:N	37:E:4191:HOH:O	2.49	0.46
8:E:137:ASP:C	8:E:137:ASP:OD1	2.58	0.46
11:H:82:LYS:HB2	11:H:82:LYS:NZ	2.32	0.46
12:I:6:PHE:HB3	12:I:109:TYR:OH	2.15	0.46
12:I:45:VAL:CG2	12:I:46:ILE:N	2.78	0.46
13:J:55:VAL:CG1	13:J:56:SER:N	2.78	0.46
13:J:66:ARG:HH11	13:J:66:ARG:HG2	1.81	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:J:72:VAL:O	13:J:95:ALA:HA	2.16	0.46
14:K:143:THR:HG22	14:K:144:ASP:H	1.81	0.46
15:L:32:ARG:HA	37:L:8647:HOH:O	2.15	0.46
16:M:90:LEU:HB2	16:M:186:LEU:HD22	1.97	0.46
16:M:110:THR:HG22	37:M:5537:HOH:O	2.14	0.46
22:S:25:ALA:O	22:S:39:ASN:HB2	2.15	0.46
22:S:25:ALA:O	22:S:39:ASN:CB	2.65	0.46
27:X:105:LYS:HE2	27:X:198:GLY:O	2.16	0.46
1:0:332:G:H4'	22:S:2:LYS:O	2.16	0.45
1:0:447:A:O2'	1:0:448:G:H5'	2.17	0.45
1:0:1850:U:H2'	1:0:1851:G:H8	1.80	0.45
1:0:2710:U:H6	1:0:2710:U:O5'	1.99	0.45
13:J:22:ASP:C	13:J:22:ASP:OD1	2.58	0.45
13:J:118:ALA:C	13:J:120:ARG:H	2.23	0.45
16:M:58:LEU:HD12	16:M:58:LEU:N	2.32	0.45
16:M:62:HIS:HB3	16:M:65:ASP:OD1	2.15	0.45
30:1:25:VAL:O	30:1:29:THR:HG23	2.16	0.45
1:0:64:G:H2'	1:0:65:C:O4'	2.16	0.45
1:0:275:G:N2	1:0:376:C:C2	2.84	0.45
1:0:1821:A:O2'	1:0:1822:A:H5'	2.16	0.45
1:0:1827:G:H2'	1:0:1828:G:C8	2.51	0.45
1:0:2270:G:H4'	4:A:223:ARG:NH1	2.31	0.45
1:0:2436:U:OP1	31:2:68:LYS:NZ	2.45	0.45
1:0:2488:A:H61	1:0:2534:C:H42	1.63	0.45
1:0:2834:G:OP1	26:W:39:LYS:HE2	2.16	0.45
2:9:3037:C:H4'	16:M:110:THR:HG23	1.96	0.45
2:9:3063:C:O2'	2:9:3064:C:H5'	2.16	0.45
4:A:53:ALA:HB1	4:A:54:PRO:HD2	1.97	0.45
7:D:15:GLU:O	7:D:16:PRO:O	2.33	0.45
12:I:47:THR:HG22	37:I:2409:HOH:O	2.16	0.45
16:M:151:ASP:CG	16:M:154:LEU:HD13	2.41	0.45
18:O:10:ALA:HA	18:O:13:VAL:CG1	2.47	0.45
22:S:19:ARG:HD3	22:S:67:LEU:O	2.17	0.45
25:V:21:LEU:CD2	25:V:48:VAL:HG11	2.43	0.45
1:0:213:G:H1'	1:0:225:G:N1	2.31	0.45
1:0:278:A:H2'	1:0:279:C:O4'	2.17	0.45
1:0:1603:A:H5'	1:0:1605:G:C5'	2.47	0.45
2:9:3088:G:OP1	25:V:130:HIS:NE2	2.39	0.45
5:B:36:PRO:HG3	5:B:169:GLY:N	2.32	0.45
5:B:307:ARG:HH11	5:B:307:ARG:CG	2.28	0.45
6:C:57:PRO:O	6:C:58:ALA:C	2.59	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:C:76:ARG:HG2	6:C:78:ARG:NH1	2.31	0.45
7:D:81:GLU:C	7:D:83:PHE:N	2.73	0.45
8:E:125:GLU:HB2	8:E:132:THR:CG2	2.47	0.45
11:H:150:LYS:CE	37:H:8368:HOH:O	2.61	0.45
18:O:83:LYS:O	18:O:86:ALA:HB3	2.17	0.45
25:V:132:VAL:HG23	25:V:138:LEU:O	2.16	0.45
26:W:30:MET:CE	26:W:58:ALA:HB3	2.41	0.45
31:2:64:LYS:HE2	37:2:8558:HOH:O	2.15	0.45
31:2:84:ARG:CD	37:2:8541:HOH:O	2.65	0.45
1:0:65:C:O2'	1:0:66:G:H5'	2.16	0.45
1:0:244:C:H6	1:0:244:C:O5'	2.00	0.45
1:0:249:G:O2'	1:0:250:C:H5'	2.17	0.45
1:0:724:G:O2'	1:0:725:C:H5'	2.17	0.45
1:0:1168:C:H2'	1:0:1169:U:O4'	2.15	0.45
1:0:1940:C:H4'	37:0:6726:HOH:O	2.15	0.45
37:0:6956:HOH:O	28:Y:31:ILE:HG13	2.15	0.45
8:E:11:VAL:HG11	8:E:22:VAL:HG13	1.97	0.45
8:E:166:VAL:HB	37:E:6341:HOH:O	2.16	0.45
11:H:150:LYS:CG	37:H:8368:HOH:O	2.62	0.45
12:I:27:ALA:HB1	12:I:87:LEU:HD21	1.96	0.45
13:J:82:ARG:NH2	13:J:115:ARG:HG2	2.32	0.45
18:O:109:ARG:NH1	18:O:119:TYR:CE2	2.85	0.45
23:T:14:GLU:OE1	23:T:15:PRO:CD	2.58	0.45
25:V:29:VAL:O	25:V:30:ASN:HB2	2.16	0.45
27:X:126:PRO:HG2	27:X:128:PHE:CZ	2.51	0.45
1:0:161:A:H3'	37:0:8848:HOH:O	2.17	0.45
1:0:240:C:C5'	15:L:146:GLN:NE2	2.79	0.45
1:0:814:G:N2	1:0:815:U:H1'	2.32	0.45
1:0:2379:G:N7	1:0:2408:A:N1	2.64	0.45
1:0:2445:U:H2'	1:0:2446:G:C8	2.51	0.45
1:0:2900:G:H2'	1:0:2901:C:O4'	2.17	0.45
5:B:36:PRO:CD	5:B:169:GLY:H	2.30	0.45
6:C:174:ILE:HG12	6:C:186:TYR:CE2	2.51	0.45
7:D:55:LYS:O	7:D:56:ARG:HB2	2.16	0.45
8:E:81:GLU:N	37:E:6931:HOH:O	2.40	0.45
9:F:13:GLU:OE2	9:F:78:GLU:HG2	2.17	0.45
9:F:48:VAL:HG23	9:F:74:PHE:HB3	1.98	0.45
11:H:86:ARG:H	11:H:86:ARG:HG2	1.38	0.45
11:H:158:ASN:ND2	37:H:8372:HOH:O	2.49	0.45
13:J:75:ARG:O	13:J:93:ASN:HA	2.16	0.45
15:L:84:LYS:O	15:L:87:MET:HG2	2.16	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:M:154:LEU:C	16:M:156:GLU:H	2.24	0.45
20:Q:31:ILE:O	20:Q:32:ALA:C	2.59	0.45
22:S:37:GLN:HB3	37:S:6711:HOH:O	2.16	0.45
26:W:26:ALA:O	26:W:27:ASP:C	2.59	0.45
27:X:151:SER:HB3	27:X:154:ARG:HB3	1.97	0.45
1:0:183:A:H5'	15:L:157:LEU:HD12	1.98	0.45
1:0:579:G:H2'	1:0:580:A:C8	2.51	0.45
1:0:665:A:C6	1:0:666:A:C6	3.05	0.45
1:0:1530:U:O2'	1:0:1531:U:H5'	2.17	0.45
1:0:1771:U:O2'	1:0:1773:G:N7	2.47	0.45
1:0:1829:A:H2'	1:0:1830:C:H5'	1.99	0.45
1:0:2134:G:C6	1:0:2258:A:C8	3.05	0.45
37:0:5608:HOH:O	15:L:174:ARG:HD3	2.15	0.45
2:9:3114:G:H2'	2:9:3115:C:H6	1.81	0.45
5:B:162:MET:HG3	5:B:310:ARG:CD	2.46	0.45
5:B:166:VAL:O	5:B:174:ARG:HD2	2.17	0.45
7:D:166:ILE:O	7:D:169:THR:N	2.50	0.45
9:F:48:VAL:CG2	9:F:74:PHE:HB3	2.45	0.45
11:H:65:ARG:CZ	37:H:8370:HOH:O	2.65	0.45
16:M:100:ALA:O	16:M:129:ILE:HG12	2.16	0.45
18:O:6:GLN:OE1	18:O:6:GLN:N	2.45	0.45
19:P:31:GLU:CD	19:P:93:ARG:HH12	2.24	0.45
20:Q:35:ILE:O	20:Q:38:LYS:HB2	2.17	0.45
20:Q:100:ASP:C	20:Q:102:GLN:N	2.74	0.45
25:V:11:VAL:O	25:V:12:ASN:HB2	2.17	0.45
1:0:489:A:C8	22:S:82:THR:HG22	2.52	0.45
1:0:1058:A:H2'	1:0:1060:C:C5'	2.47	0.45
1:0:1439:C:O5'	1:0:1439:C:H6	1.99	0.45
1:0:2481:G:C3'	1:0:2482:G:H5''	2.47	0.45
1:0:2667:G:H1'	1:0:2914:A:N3	2.31	0.45
1:0:2730:G:O2'	1:0:2731:G:H5'	2.17	0.45
1:0:2897:C:H2'	1:0:2898:G:C8	2.45	0.45
2:9:3058:G:H1'	37:D:3839:HOH:O	2.17	0.45
4:A:1:GLY:N	37:A:8604:HOH:O	2.49	0.45
5:B:217:ARG:HG3	5:B:257:THR:HG22	1.97	0.45
7:D:128:LEU:C	7:D:128:LEU:HD23	2.42	0.45
11:H:31:PHE:HE2	11:H:87:LYS:O	1.99	0.45
12:I:74:ARG:HH11	12:I:74:ARG:CB	2.28	0.45
15:L:87:MET:HE3	37:L:8592:HOH:O	2.16	0.45
25:V:122:ARG:CG	25:V:152:ALA:O	2.65	0.45
1:0:363:A:H8	1:0:363:A:O5'	1.99	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:1052:G:H2'	1:0:1052:G:N3	2.31	0.45
1:0:1477:C:C5'	1:0:1868:G:H5''	2.46	0.45
1:0:1504:A:H5'	37:0:3881:HOH:O	2.17	0.45
1:0:1942:A:O3'	4:A:213:LYS:HE2	2.17	0.45
1:0:2253:G:N2	37:0:4974:HOH:O	2.28	0.45
37:0:6737:HOH:O	4:A:177:HIS:HE1	1.99	0.45
2:9:3049:G:H2'	2:9:3050:G:O4'	2.16	0.45
4:A:105:VAL:HG11	4:A:154:ALA:CB	2.46	0.45
4:A:153:ARG:HB2	4:A:153:ARG:NH1	2.27	0.45
7:D:35:ALA:O	7:D:37:ALA:N	2.50	0.45
8:E:31:ARG:HH12	8:E:68:HIS:CD2	2.34	0.45
11:H:84:ARG:CZ	11:H:135:TRP:CH2	2.99	0.45
11:H:110:GLY:N	37:H:8382:HOH:O	2.49	0.45
11:H:144:GLU:OE1	11:H:144:GLU:HA	2.16	0.45
16:M:155:GLU:C	16:M:156:GLU:HG3	2.42	0.45
25:V:41:TYR:HA	25:V:44:MET:HE3	1.99	0.45
26:W:26:ALA:HB1	26:W:59:TRP:CE2	2.51	0.45
27:X:189:ASN:ND2	27:X:189:ASN:C	2.74	0.45
1:0:659:A:H5''	37:N:6799:HOH:O	2.16	0.45
1:0:1007:A:H2'	11:H:19:TYR:CZ	2.52	0.45
1:0:1114:A:H2'	1:0:1115:U:H6	1.82	0.45
1:0:1180:U:H2'	1:0:1181:A:O4'	2.17	0.45
1:0:2299:G:O6	19:P:1:PRO:HA	2.16	0.45
1:0:2697:A:H2'	1:0:2698:G:O4'	2.17	0.45
1:0:2869:G:H2'	1:0:2870:C:C6	2.51	0.45
4:A:211:LYS:NZ	37:A:8612:HOH:O	2.39	0.45
5:B:168:GLY:H	5:B:174:ARG:HH11	1.65	0.45
6:C:140:VAL:HG12	6:C:141:SER:N	2.32	0.45
6:C:178:GLN:C	6:C:180:SER:N	2.72	0.45
7:D:154:LYS:H	7:D:154:LYS:CD	2.22	0.45
11:H:157:ILE:HG22	11:H:158:ASN:N	2.32	0.45
12:I:45:VAL:HG21	12:I:129:PHE:CD1	2.52	0.45
15:L:147:LEU:O	15:L:149:TRP:N	2.50	0.45
16:M:72:GLU:H	16:M:171:HIS:CE1	2.34	0.45
18:O:14:LEU:HD13	18:O:51:ALA:HB2	1.98	0.45
18:O:16:VAL:CG1	18:O:17:GLY:N	2.80	0.45
18:O:120:ARG:NH2	18:O:123:TYR:CD2	2.84	0.45
19:P:16:ASN:HD22	19:P:16:ASN:HA	1.50	0.45
19:P:32:GLU:O	19:P:93:ARG:NH2	2.50	0.45
1:0:154:C:C2	1:0:155:C:C5	3.05	0.45
1:0:890:C:O2'	29:Z:50:TRP:O	2.33	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:1384:C:H5'	26:W:30:MET:HG2	1.98	0.45
1:0:2256:G:C2'	1:0:2257:G:C5'	2.92	0.45
4:A:88:ILE:CD1	4:A:100:PRO:HD3	2.39	0.45
5:B:41:PHE:HB3	5:B:190:MET:HE1	1.99	0.45
11:H:53:PRO:HA	11:H:125:VAL:O	2.17	0.45
11:H:84:ARG:CZ	11:H:135:TRP:HH2	2.30	0.45
14:K:146:GLY:C	14:K:148:GLU:H	2.25	0.45
22:S:89:ARG:O	22:S:90:PRO:C	2.59	0.45
23:T:11:THR:HG22	23:T:53:ASP:CG	2.42	0.45
24:U:19:GLU:O	24:U:22:ASP:HB2	2.17	0.45
27:X:101:GLY:HA3	37:X:8560:HOH:O	2.17	0.45
27:X:117:LEU:HD12	27:X:174:VAL:CG1	2.46	0.45
29:Z:44:LYS:NZ	37:Z:8454:HOH:O	2.50	0.45
1:0:242:A:H5'	37:0:5252:HOH:O	2.18	0.44
1:0:331:A:H1'	37:0:4253:HOH:O	2.16	0.44
1:0:357:A:N7	37:0:4583:HOH:O	2.36	0.44
1:0:462:A:N3	30:1:37:HIS:HB3	2.31	0.44
1:0:696:C:O2'	1:0:697:G:H5'	2.16	0.44
1:0:818:A:O2'	28:Y:13:ARG:HD3	2.16	0.44
1:0:1008:C:OP1	11:H:16:ARG:NH2	2.47	0.44
1:0:1677:U:OP2	30:1:8:LYS:NZ	2.49	0.44
1:0:1706:G:H1'	1:0:1712:A:H61	1.82	0.44
1:0:1755:A:H2'	1:0:1756:G:O4'	2.17	0.44
4:A:99:ILE:O	4:A:131:HIS:CE1	2.70	0.44
4:A:130:THR:C	37:A:8566:HOH:O	2.60	0.44
4:A:149:ASP:OD1	4:A:151:GLN:HB2	2.17	0.44
5:B:76:THR:N	5:B:77:PRO:HD3	2.32	0.44
5:B:275:GLY:C	37:B:8649:HOH:O	2.60	0.44
6:C:85:LYS:HE2	37:C:8324:HOH:O	2.17	0.44
10:G:18:GLU:O	10:G:21:ASP:HB2	2.17	0.44
13:J:78:LYS:HA	13:J:79:PRO:HD3	1.85	0.44
15:L:5:TYR:C	15:L:7:TYR:N	2.74	0.44
16:M:79:PRO:HG3	16:M:142:THR:O	2.17	0.44
18:O:31:ILE:HG12	18:O:43:LEU:HD13	1.99	0.44
20:Q:106:GLY:HA2	20:Q:109:MET:CE	2.41	0.44
25:V:13:MET:CE	25:V:17:ILE:HG22	2.47	0.44
1:0:213:G:C2'	1:0:214:U:OP2	2.65	0.44
1:0:303:C:H2'	1:0:304:G:O4'	2.17	0.44
1:0:308:U:C5'	22:S:97:ARG:NH2	2.80	0.44
1:0:963:C:O5'	1:0:963:C:H6	2.00	0.44
1:0:1497:G:H4'	1:0:1627:G:O2'	2.17	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:1565:C:O4'	1:0:2738:G:H1'	2.17	0.44
1:0:1794:G:P	18:O:133:SER:HB2	2.58	0.44
1:0:1820:G:C6	1:0:2030:A:C2	3.05	0.44
1:0:2353:A:O2'	16:M:7:LYS:HB3	2.17	0.44
1:0:2443:C:O3'	14:K:56:LYS:HE3	2.17	0.44
1:0:2483:A:H4'	37:0:6875:HOH:O	2.17	0.44
37:0:3180:HOH:O	8:E:143:GLN:HG2	2.16	0.44
2:9:3107:C:C6	37:9:8440:HOH:O	2.68	0.44
6:C:237:GLU:N	37:C:8444:HOH:O	2.49	0.44
37:C:8366:HOH:O	22:S:2:LYS:HE2	2.16	0.44
7:D:23:VAL:CG2	7:D:73:VAL:HB	2.47	0.44
8:E:162:PHE:N	8:E:162:PHE:CD1	2.84	0.44
9:F:104:ALA:O	9:F:108:LEU:HB3	2.17	0.44
11:H:46:VAL:CG1	11:H:146:TRP:HZ3	2.29	0.44
11:H:57:ARG:HG3	11:H:57:ARG:NH1	2.32	0.44
14:K:121:ILE:HG12	14:K:141:GLU:HB2	2.00	0.44
16:M:67:ALA:O	16:M:69:TYR:N	2.51	0.44
19:P:13:LYS:NZ	37:P:4956:HOH:O	2.50	0.44
21:R:34:LYS:HG2	21:R:54:THR:HG23	1.99	0.44
25:V:122:ARG:CZ	37:V:5817:HOH:O	2.65	0.44
1:0:814:G:C8	37:0:6598:HOH:O	2.54	0.44
1:0:1170:U:O2'	1:0:1172:G:N7	2.34	0.44
1:0:1855:G:H8	4:A:144:GLU:OE2	2.00	0.44
1:0:2038:A:O2'	1:0:2039:A:H5'	2.18	0.44
1:0:2437:A:H2'	1:0:2438:G:C8	2.52	0.44
1:0:2548:C:H2'	1:0:2549:C:H6	1.83	0.44
1:0:2637:A:H4'	1:0:2638:G:H5'	1.97	0.44
1:0:2807:U:OP2	5:B:28:SER:OG	2.35	0.44
1:0:2892:G:C6	1:0:2893:C:N3	2.86	0.44
2:9:3029:C:C2'	2:9:3030:C:H5'	2.48	0.44
5:B:41:PHE:HB3	5:B:190:MET:CE	2.47	0.44
6:C:246:ARG:HB3	6:C:246:ARG:HH11	1.82	0.44
7:D:35:ALA:HB1	37:D:3279:HOH:O	2.17	0.44
7:D:93:LEU:HB3	7:D:97:GLN:OE1	2.17	0.44
7:D:167:GLU:C	7:D:169:THR:H	2.25	0.44
7:D:170:TYR:CD1	7:D:170:TYR:N	2.85	0.44
10:G:12:ILE:O	10:G:13:PRO:C	2.59	0.44
10:G:71:LEU:O	10:G:73:ASP:N	2.50	0.44
13:J:50:GLY:O	13:J:120:ARG:NH1	2.49	0.44
15:L:37:VAL:HG11	15:L:108:LYS:HG3	1.99	0.44
16:M:33:ARG:HD2	16:M:103:ASP:OD2	2.17	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:O:84:ALA:C	18:O:86:ALA:N	2.75	0.44
19:P:26:PRO:O	19:P:30:VAL:HG23	2.18	0.44
24:U:42:ASN:O	24:U:44:GLY:N	2.49	0.44
25:V:154:ARG:HB3	25:V:154:ARG:HE	1.60	0.44
31:2:70:ARG:HG2	31:2:70:ARG:HH11	1.81	0.44
1:0:73:C:O2'	1:0:74:A:H5'	2.17	0.44
1:0:396:U:P	31:2:38:ARG:HH11	2.39	0.44
1:0:488:U:H2'	37:0:3484:HOH:O	2.17	0.44
1:0:532:A:N1	1:0:2660:G:O2'	2.41	0.44
1:0:860:U:H2'	1:0:861:A:C8	2.53	0.44
1:0:1235:G:C1'	12:I:63:ILE:HG23	2.48	0.44
1:0:1299:G:N2	37:0:4139:HOH:O	2.50	0.44
1:0:1603:A:C5'	1:0:1605:G:H5'	2.48	0.44
1:0:1819:G:H2'	1:0:1820:G:C5'	2.47	0.44
1:0:2497:A:H2'	1:0:2498:C:C6	2.52	0.44
1:0:2668:G:H2'	1:0:2669:U:C6	2.53	0.44
37:0:9949:HOH:O	2:9:3103:A:H1'	2.17	0.44
7:D:10:PHE:CD1	7:D:11:HIS:N	2.86	0.44
9:F:117:GLU:C	9:F:119:ARG:N	2.75	0.44
16:M:131:HIS:NE2	37:M:4678:HOH:O	2.36	0.44
17:N:107:GLU:O	17:N:108:GLY:C	2.58	0.44
19:P:42:LYS:NZ	19:P:43:ILE:O	2.46	0.44
23:T:25:ASP:OD2	23:T:26:GLY:N	2.50	0.44
25:V:6:GLN:CB	25:V:26:ILE:HD12	2.42	0.44
25:V:65:VAL:HA	25:V:68:THR:CG2	2.47	0.44
27:X:112:GLU:OE1	27:X:115:ARG:NH1	2.50	0.44
27:X:219:GLU:HG3	27:X:220:GLU:N	2.31	0.44
28:Y:73:THR:O	28:Y:76:GLY:N	2.50	0.44
1:0:812:A:H2'	1:0:813:C:C6	2.52	0.44
1:0:1181:A:O2'	1:0:1182:C:H5'	2.18	0.44
1:0:1192:A:O2'	1:0:1193:A:OP1	2.35	0.44
1:0:1667:A:H2'	1:0:1668:U:H6	1.83	0.44
1:0:1972:U:C2'	1:0:1973:A:H5'	2.45	0.44
1:0:2387:U:H2'	1:0:2388:C:C6	2.52	0.44
1:0:2518:C:H2'	1:0:2519:C:O4'	2.16	0.44
4:A:29:HIS:CE1	4:A:107:ASN:ND2	2.86	0.44
8:E:170:ARG:HE	8:E:170:ARG:HB2	1.63	0.44
9:F:32:GLY:N	37:F:3111:HOH:O	2.49	0.44
11:H:1:LYS:CA	37:H:8353:HOH:O	2.65	0.44
20:Q:46:TYR:CD2	20:Q:47:LEU:HD23	2.52	0.44
20:Q:75:TRP:CG	20:Q:76:ASP:H	2.35	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:Y:27:ALA:O	28:Y:28:ASP:C	2.59	0.44
1:0:440:C:O2'	1:0:441:A:H5'	2.18	0.44
1:0:1209:C:H2'	1:0:1210:G:C8	2.51	0.44
1:0:1226:G:N2	37:0:4048:HOH:O	2.44	0.44
1:0:1500:U:OP2	18:O:41:ARG:NH2	2.51	0.44
1:0:2542:C:H5''	1:0:2608:C:N4	2.32	0.44
4:A:53:ALA:HB3	37:A:8602:HOH:O	2.17	0.44
4:A:66:ARG:HH11	4:A:66:ARG:HB2	1.81	0.44
4:A:76:VAL:HG23	28:Y:63:LYS:HB3	1.99	0.44
7:D:52:THR:O	7:D:52:THR:HG22	2.17	0.44
7:D:174:VAL:CG1	37:D:6555:HOH:O	2.65	0.44
11:H:45:GLN:CB	11:H:163:PRO:HD2	2.22	0.44
13:J:98:VAL:HG13	13:J:99:ASP:N	2.32	0.44
16:M:176:ARG:O	16:M:180:LEU:HG	2.18	0.44
31:2:8:ASN:O	31:2:9:THR:HB	2.17	0.44
1:0:415:A:O2'	1:0:416:G:H5'	2.18	0.44
1:0:485:A:O2'	1:0:487:G:H5'	2.18	0.44
1:0:611:U:H2'	1:0:612:U:C6	2.53	0.44
1:0:716:G:H2'	1:0:717:C:O5'	2.18	0.44
1:0:876:A:H2'	1:0:876:A:N3	2.33	0.44
1:0:1925:G:O2'	1:0:1926:G:H5'	2.18	0.44
1:0:2032:U:H2'	1:0:2033:G:C5'	2.48	0.44
1:0:2356:A:H2'	1:0:2357:G:O4'	2.17	0.44
1:0:2457:U:H2'	1:0:2458:U:C6	2.53	0.44
1:0:2769:C:H2'	1:0:2770:G:H5'	2.00	0.44
1:0:2777:G:O2'	1:0:2778:A:H5'	2.17	0.44
1:0:2815:G:N7	12:I:80:LYS:NZ	2.65	0.44
7:D:35:ALA:C	7:D:37:ALA:H	2.25	0.44
11:H:144:GLU:HG3	37:H:8332:HOH:O	2.17	0.44
17:N:77:ALA:HA	17:N:96:VAL:O	2.18	0.44
1:0:794:U:H3	1:0:819:A:N6	2.10	0.44
1:0:812:A:H2'	1:0:813:C:O4'	2.17	0.44
1:0:1854:C:O2'	1:0:1858:A:N3	2.45	0.44
1:0:1882:C:O2'	1:0:2012:U:OP2	2.31	0.44
1:0:2248:C:H3'	37:0:4880:HOH:O	2.18	0.44
1:0:2488:A:H1'	37:0:8613:HOH:O	2.16	0.44
1:0:2721:U:H4'	13:J:87:ARG:HG3	2.00	0.44
1:0:2791:U:H1'	1:0:2792:A:H5''	2.00	0.44
7:D:25:MET:HE1	7:D:40:ILE:CG1	2.48	0.44
12:I:130:VAL:CG1	12:I:131:THR:N	2.81	0.44
25:V:65:VAL:CA	25:V:68:THR:HG22	2.47	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:W:43:VAL:HG22	26:W:76:ARG:NH1	2.32	0.44
27:X:177:LYS:HE2	27:X:183:GLU:OE2	2.18	0.44
31:2:3:MET:HG3	31:2:4:PRO:HD2	1.99	0.44
1:0:681:G:H1'	1:0:683:G:O6	2.18	0.44
1:0:1486:A:C5	30:1:2:LYS:HG3	2.52	0.44
1:0:1597:A:O4'	18:O:95:GLU:HG2	2.18	0.44
1:0:2255:A:C6	1:0:2256:G:C5	3.06	0.44
1:0:2911:C:H2'	1:0:2912:C:C6	2.52	0.44
4:A:149:ASP:OD1	4:A:151:GLN:CB	2.66	0.44
5:B:132:HIS:CE1	5:B:171:VAL:CG2	3.01	0.44
5:B:168:GLY:O	5:B:169:GLY:O	2.36	0.44
5:B:322:ARG:HB2	37:B:8606:HOH:O	2.17	0.44
6:C:111:VAL:HB	37:C:8321:HOH:O	2.17	0.44
6:C:162:VAL:HG12	6:C:162:VAL:O	2.18	0.44
8:E:21:THR:HG23	8:E:30:THR:OG1	2.18	0.44
11:H:39:GLY:O	11:H:41:THR:N	2.51	0.44
15:L:28:MET:HA	15:L:31:TRP:HB2	2.00	0.44
16:M:151:ASP:HB3	37:M:3251:HOH:O	2.17	0.44
22:S:45:GLY:C	37:S:3851:HOH:O	2.61	0.44
22:S:98:VAL:HG11	22:S:101:LEU:CD2	2.48	0.44
25:V:31:HIS:C	37:V:5420:HOH:O	2.60	0.44
1:0:283:U:H5''	1:0:284:C:P	2.58	0.43
1:0:1014:A:H2'	1:0:1015:C:H5'	2.00	0.43
1:0:1135:G:C6	1:0:1136:U:C4	3.07	0.43
1:0:1589:G:H4'	37:0:6248:HOH:O	2.17	0.43
1:0:1592:G:O2'	1:0:1593:C:O4'	2.35	0.43
1:0:1692:C:H1'	37:0:8974:HOH:O	2.16	0.43
1:0:2265:U:H2'	1:0:2266:A:H8	1.82	0.43
2:9:3002:U:H1'	37:9:8484:HOH:O	2.17	0.43
4:A:191:GLY:C	4:A:193:ALA:H	2.26	0.43
5:B:278:PRO:HD3	5:B:294:TYR:CZ	2.52	0.43
13:J:30:LYS:O	13:J:55:VAL:HG13	2.17	0.43
16:M:175:LEU:O	16:M:176:ARG:C	2.61	0.43
23:T:6:CYS:C	23:T:8:TYR:N	2.73	0.43
1:0:221:G:H2'	1:0:222:A:C8	2.53	0.43
1:0:226:A:H1'	1:0:393:G:C5	2.53	0.43
1:0:624:U:H5''	37:0:9026:HOH:O	2.17	0.43
1:0:1114:A:H2'	1:0:1115:U:C6	2.53	0.43
1:0:1215:A:O3'	1:0:1216:G:C4'	2.65	0.43
1:0:1838:U:O2'	1:0:2644:C:H5'	2.17	0.43
1:0:1934:A:C8	1:0:1935:C:C5	3.07	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:2782:G:O6	1:0:2790:C:H5''	2.18	0.43
1:0:2897:C:O2'	1:0:2898:G:H5'	2.19	0.43
2:9:3034:A:H2'	2:9:3035:C:O4'	2.18	0.43
5:B:145:HIS:CD2	5:B:146:THR:O	2.68	0.43
7:D:27:ILE:CG2	7:D:28:GLY:H	2.20	0.43
10:G:73:ASP:O	37:G:2218:HOH:O	2.21	0.43
14:K:101:ASP:C	14:K:103:ALA:H	2.26	0.43
14:K:143:THR:CG2	14:K:144:ASP:H	2.31	0.43
14:K:147:GLU:C	37:K:8552:HOH:O	2.61	0.43
16:M:86:LEU:O	16:M:90:LEU:HG	2.18	0.43
21:R:11:THR:H	21:R:14:ALA:HB3	1.83	0.43
22:S:40:VAL:HG23	22:S:119:ALA:C	2.44	0.43
28:Y:20:LEU:O	28:Y:21:LYS:C	2.61	0.43
28:Y:34:LYS:HE2	37:Y:8422:HOH:O	2.18	0.43
1:0:51:G:O2'	1:0:52:A:H5'	2.18	0.43
1:0:236:A:H4'	1:0:237:G:OP1	2.18	0.43
1:0:699:C:C2	1:0:743:G:N2	2.87	0.43
1:0:1155:G:H2'	1:0:1156:C:C6	2.52	0.43
1:0:1265:G:H1'	37:0:4454:HOH:O	2.17	0.43
1:0:1874:U:P	4:A:51:ARG:HD2	2.58	0.43
1:0:2546:U:H2'	1:0:2547:C:C6	2.54	0.43
4:A:194:MET:HE3	4:A:199:HIS:CB	2.47	0.43
5:B:102:THR:CG2	5:B:182:VAL:HG12	2.48	0.43
5:B:102:THR:O	5:B:105:PHE:CZ	2.71	0.43
9:F:108:LEU:HG	9:F:109:GLU:N	2.33	0.43
12:I:15:ARG:CZ	12:I:43:ARG:NH1	2.81	0.43
12:I:59:LYS:O	12:I:63:ILE:HG13	2.18	0.43
13:J:115:ARG:CG	13:J:116:GLU:N	2.77	0.43
15:L:45:ARG:CZ	15:L:48:ARG:HG3	2.48	0.43
16:M:14:ARG:C	16:M:16:ALA:H	2.26	0.43
17:N:26:TRP:HA	17:N:26:TRP:HE3	1.82	0.43
17:N:63:LYS:HG3	17:N:80:ASP:O	2.18	0.43
19:P:29:ALA:HB1	37:P:7270:HOH:O	2.18	0.43
20:Q:61:GLN:CD	37:Q:8543:HOH:O	2.60	0.43
25:V:107:LEU:O	25:V:112:LEU:HB2	2.18	0.43
27:X:189:ASN:HA	27:X:217:ILE:HD11	2.00	0.43
1:0:10:U:HO2'	1:0:11:A:P	2.42	0.43
1:0:218:C:C5	1:0:220:C:C4	3.06	0.43
1:0:445:U:H2'	1:0:446:G:H8	1.83	0.43
1:0:483:C:C4	1:0:484:A:C6	3.06	0.43
1:0:695:C:H2'	1:0:696:C:C6	2.54	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:732:C:H2'	1:0:733:U:C6	2.53	0.43
1:0:1545:C:H2'	1:0:1546:G:O4'	2.19	0.43
1:0:1701:A:H4'	1:0:1702:U:C5'	2.49	0.43
1:0:2335:C:C2	1:0:2350:G:C2	3.06	0.43
1:0:2367:A:H5''	37:M:4034:HOH:O	2.18	0.43
1:0:2758:G:H2'	1:0:2759:C:C6	2.54	0.43
1:0:2780:C:H1'	37:0:3180:HOH:O	2.17	0.43
1:0:2791:U:C1'	1:0:2792:A:H5''	2.49	0.43
1:0:2831:C:C2'	1:0:2832:C:H5'	2.49	0.43
1:0:2892:G:C6	1:0:2893:C:C4	3.06	0.43
37:0:6835:HOH:O	6:C:188:ARG:HD3	2.18	0.43
4:A:120:ARG:NH2	37:A:8577:HOH:O	2.50	0.43
4:A:125:ASN:CB	4:A:158:VAL:HG12	2.48	0.43
5:B:271:ASP:HB3	5:B:296:LEU:HD12	2.00	0.43
7:D:173:GLU:O	7:D:174:VAL:C	2.60	0.43
9:F:26:THR:HB	9:F:102:GLY:C	2.43	0.43
9:F:49:PHE:N	9:F:49:PHE:CD1	2.87	0.43
9:F:53:ASP:CG	9:F:80:GLN:HB2	2.43	0.43
9:F:113:ASP:O	9:F:117:GLU:HG3	2.18	0.43
12:I:39:VAL:HG11	12:I:107:ASN:HB2	2.00	0.43
14:K:55:GLN:HA	14:K:58:GLN:HE21	1.82	0.43
16:M:116:PHE:CG	37:M:6448:HOH:O	2.70	0.43
20:Q:8:ALA:CB	20:Q:13:THR:HG21	2.34	0.43
23:T:47:ARG:HG2	37:T:4381:HOH:O	2.15	0.43
26:W:9:VAL:HG13	26:W:88:GLU:CD	2.43	0.43
27:X:117:LEU:HD12	27:X:174:VAL:HG11	2.00	0.43
1:0:90:A:H2'	1:0:91:G:O4'	2.18	0.43
1:0:158:A:H2'	1:0:159:G:O4'	2.19	0.43
1:0:613:C:H2'	1:0:614:U:H6	1.83	0.43
1:0:870:G:C3'	1:0:871:G:H5''	2.48	0.43
1:0:1242:A:H5'	12:I:82:THR:CG2	2.33	0.43
1:0:1851:G:H1'	37:0:3899:HOH:O	2.18	0.43
1:0:2403:C:H2'	1:0:2404:G:O5'	2.18	0.43
1:0:2672:C:P	5:B:25:ARG:NH1	2.92	0.43
37:0:3877:HOH:O	4:A:11:ARG:CZ	2.66	0.43
2:9:3026:C:P	37:9:8445:HOH:O	2.77	0.43
9:F:22:VAL:HG21	9:F:104:ALA:HB2	2.00	0.43
12:I:49:ARG:O	12:I:50:GLU:C	2.61	0.43
15:L:153:THR:C	15:L:155:HIS:H	2.26	0.43
15:L:164:THR:HG22	15:L:167:GLY:CA	2.49	0.43
16:M:147:ILE:CG2	16:M:148:ALA:N	2.82	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:M:152:GLU:C	16:M:154:LEU:N	2.75	0.43
18:O:131:PHE:CE1	18:O:137:LEU:HD13	2.53	0.43
22:S:40:VAL:HG22	22:S:41:ARG:N	2.33	0.43
23:T:6:CYS:O	23:T:8:TYR:N	2.52	0.43
29:Z:17:THR:HA	30:1:49:GLU:HA	2.01	0.43
29:Z:45:ARG:NH1	37:Z:8435:HOH:O	2.52	0.43
1:0:814:G:H2'	1:0:815:U:O4'	2.19	0.43
1:0:853:C:H2'	1:0:854:G:O4'	2.17	0.43
1:0:951:A:C2'	1:0:952:G:H5'	2.48	0.43
1:0:1345:A:H2'	1:0:1346:U:C6	2.53	0.43
1:0:1345:A:H2'	1:0:1346:U:H6	1.84	0.43
1:0:1592:G:O2'	1:0:1593:C:O5'	2.36	0.43
1:0:1644:C:H2'	1:0:1645:U:H6	1.84	0.43
1:0:1992:U:H2'	1:0:1994:A:OP2	2.19	0.43
1:0:2039:A:OP2	5:B:234:ARG:NH2	2.51	0.43
1:0:2898:G:O2'	1:0:2899:A:H5'	2.17	0.43
4:A:36:ASP:CB	4:A:85:ASP:H	2.32	0.43
4:A:132:ASP:OD1	4:A:133:ARG:N	2.49	0.43
5:B:145:HIS:CD2	5:B:145:HIS:C	2.97	0.43
6:C:123:LEU:HA	6:C:123:LEU:HD23	1.85	0.43
7:D:15:GLU:HA	7:D:16:PRO:HD3	1.94	0.43
7:D:60:GLU:C	7:D:62:ASP:N	2.74	0.43
9:F:6:PHE:O	9:F:6:PHE:CD1	2.71	0.43
13:J:39:GLY:HA3	37:J:992:HOH:O	2.18	0.43
15:L:61:ILE:N	15:L:61:ILE:CD1	2.81	0.43
16:M:175:LEU:HD12	16:M:175:LEU:HA	1.84	0.43
20:Q:119:VAL:CG2	20:Q:142:ASP:HB2	2.49	0.43
21:R:73:ASP:O	21:R:77:VAL:HG23	2.18	0.43
22:S:53:GLY:HA3	37:S:6384:HOH:O	2.19	0.43
23:T:20:MET:HE1	37:T:7438:HOH:O	2.19	0.43
26:W:27:ASP:OD2	26:W:27:ASP:N	2.41	0.43
27:X:234:VAL:HG12	27:X:235:GLU:N	2.33	0.43
28:Y:39:CYS:HA	28:Y:40:PRO:HD3	1.89	0.43
1:0:1582:C:O2'	1:0:1583:U:H5'	2.19	0.43
1:0:2438:G:H2'	1:0:2439:C:O4'	2.19	0.43
1:0:2642:G:H2'	1:0:2643:G:O4'	2.18	0.43
1:0:2765:C:H2'	1:0:2766:A:C8	2.54	0.43
37:O:3339:HOH:O	11:H:90:PHE:HD2	2.01	0.43
6:C:26:VAL:N	37:C:8356:HOH:O	2.26	0.43
7:D:48:MET:HA	7:D:49:PRO:HD3	1.83	0.43
9:F:21:GLU:O	9:F:24:ARG:CG	2.65	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:H:82:LYS:HB2	11:H:82:LYS:HZ2	1.83	0.43
13:J:101:ASN:O	13:J:102:GLU:CB	2.67	0.43
18:O:103:THR:O	18:O:107:GLU:HG3	2.19	0.43
25:V:63:GLU:HG2	25:V:93:ILE:HG22	2.00	0.43
27:X:98:GLN:HA	37:X:8537:HOH:O	2.18	0.43
27:X:148:GLY:O	27:X:154:ARG:HD3	2.18	0.43
29:Z:53:LYS:HD3	29:Z:53:LYS:HA	1.88	0.43
1:O:128:A:C8	1:O:128:A:C3'	3.01	0.43
1:O:187:A:H3'	1:O:188:C:H6	1.82	0.43
1:O:1436:C:O2'	1:O:1437:A:H5'	2.19	0.43
1:O:2073:G:N2	37:O:4143:HOH:O	2.52	0.43
1:O:2120:U:H2'	1:O:2121:G:O4'	2.19	0.43
1:O:2809:G:H2'	1:O:2810:G:O4'	2.18	0.43
1:O:2894:C:H2'	1:O:2895:C:H6	1.84	0.43
2:9:3095:C:O2'	2:9:3096:C:H5'	2.19	0.43
6:C:120:ASP:OD1	6:C:120:ASP:C	2.62	0.43
7:D:166:ILE:O	7:D:167:GLU:C	2.59	0.43
9:F:111:ILE:O	9:F:115:VAL:HG23	2.19	0.43
20:Q:111:ILE:HG23	20:Q:145:LEU:HD11	2.00	0.43
25:V:4:LEU:HD23	25:V:4:LEU:HA	1.83	0.43
30:1:18:ASN:HD22	30:1:18:ASN:HA	1.57	0.43
31:2:70:ARG:HG2	31:2:70:ARG:NH1	2.33	0.43
1:O:11:A:H5'	1:O:12:U:OP2	2.18	0.43
1:O:248:A:H5'	1:O:249:G:OP2	2.19	0.43
1:O:283:U:H5	1:O:284:C:N4	2.17	0.43
1:O:401:C:C5'	37:O:5220:HOH:O	2.67	0.43
1:O:1056:U:H2'	1:O:1057:A:O4'	2.19	0.43
1:O:1268:C:O2'	1:O:1269:G:H5'	2.19	0.43
1:O:2135:A:O2'	1:O:2136:G:H5'	2.18	0.43
1:O:2795:C:O2'	1:O:2796:U:H5'	2.19	0.43
2:9:3041:C:H4'	7:D:48:MET:HB2	2.01	0.43
5:B:109:LEU:HD11	5:B:113:LEU:HD11	2.01	0.43
6:C:103:ASN:O	6:C:104:ASP:C	2.60	0.43
7:D:81:GLU:O	7:D:84:LEU:N	2.51	0.43
11:H:157:ILE:CG2	11:H:158:ASN:N	2.82	0.43
12:I:74:ARG:C	12:I:76:ASP:N	2.77	0.43
15:L:35:PRO:HD2	15:L:38:VAL:HG21	1.99	0.43
16:M:37:ARG:HD3	35:M:8507:CL:CL	2.56	0.43
22:S:78:THR:HB	22:S:87:VAL:O	2.18	0.43
25:V:122:ARG:HG2	25:V:122:ARG:NH1	2.28	0.43
25:V:131:PRO:HG2	25:V:134:GLU:HB2	2.00	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:Y:32:LYS:NZ	28:Y:70:GLN:NE2	2.66	0.43
28:Y:38:LYS:CG	28:Y:45:LYS:HG2	2.43	0.43
1:0:113:A:H3'	1:0:114:A:H5''	2.00	0.43
1:0:488:U:O2'	22:S:82:THR:HG21	2.18	0.43
1:0:525:G:H2'	1:0:526:U:O4'	2.18	0.43
1:0:1187:U:H3'	37:0:6289:HOH:O	2.17	0.43
1:0:1524:U:O2'	1:0:1525:G:P	2.77	0.43
1:0:1568:G:O2'	1:0:1569:U:H5'	2.18	0.43
1:0:2473:U:O3'	1:0:2474:A:H3'	2.19	0.43
5:B:41:PHE:HA	5:B:79:MET:CE	2.48	0.43
6:C:61:PHE:HB3	37:C:8440:HOH:O	2.19	0.43
8:E:106:ASN:HD21	8:E:109:GLY:HA2	1.83	0.43
12:I:56:LYS:NZ	37:I:2828:HOH:O	2.52	0.43
13:J:61:THR:O	13:J:64:MET:N	2.50	0.43
15:L:138:HIS:C	15:L:139:PRO:O	2.60	0.43
18:O:13:VAL:HG21	18:O:41:ARG:HG2	2.01	0.43
21:R:23:LYS:HE2	37:R:3430:HOH:O	2.19	0.43
21:R:49:VAL:HG13	21:R:66:VAL:HG13	2.01	0.43
23:T:52:THR:HG21	23:T:54:THR:HB	2.01	0.43
25:V:3:ALA:O	25:V:54:PHE:HA	2.18	0.43
27:X:117:LEU:HA	27:X:174:VAL:HG11	2.00	0.43
1:0:1109:U:O4	12:I:21:ARG:HA	2.19	0.42
1:0:1335:C:OP2	27:X:207:SER:HB2	2.19	0.42
1:0:1433:G:H2'	1:0:1434:A:O4'	2.19	0.42
1:0:1669:A:H2'	1:0:1670:G:H8	1.84	0.42
1:0:1823:G:O2'	1:0:1824:C:H5'	2.19	0.42
1:0:2637:A:C4'	1:0:2638:G:C5'	2.93	0.42
1:0:2794:G:N2	1:0:2795:C:H1'	2.33	0.42
1:0:2812:A:H2	1:0:2814:A:N6	1.99	0.42
6:C:118:THR:HG22	6:C:137:PRO:HB3	2.01	0.42
6:C:246:ARG:NH2	37:C:8421:HOH:O	2.52	0.42
11:H:1:LYS:HA	11:H:2:PRO:HD3	1.82	0.42
11:H:62:GLU:CD	37:H:8370:HOH:O	2.62	0.42
15:L:39:ARG:HG3	37:L:8601:HOH:O	2.19	0.42
22:S:23:VAL:O	22:S:93:THR:HG21	2.17	0.42
1:0:590:A:H2'	1:0:591:A:H5'	2.01	0.42
1:0:1218:U:H2'	1:0:1219:U:C6	2.53	0.42
1:0:1391:G:H2'	1:0:1392:A:H5'	2.01	0.42
1:0:2362:A:H2'	1:0:2363:G:C8	2.54	0.42
1:0:2381:C:H4'	31:2:80:ARG:NH2	2.34	0.42
1:0:2670:G:H4'	5:B:112:THR:HG22	2.01	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:B:82:VAL:O	5:B:82:VAL:CG1	2.66	0.42
5:B:162:MET:HG3	5:B:310:ARG:CZ	2.48	0.42
5:B:302:PRO:C	37:B:8652:HOH:O	2.62	0.42
6:C:138:VAL:O	6:C:234:VAL:HA	2.19	0.42
9:F:101:ALA:HB2	9:F:108:LEU:HD22	2.01	0.42
11:H:86:ARG:HD3	11:H:130:HIS:HD2	1.84	0.42
11:H:129:ASN:N	11:H:129:ASN:HD22	2.17	0.42
14:K:144:ASP:HA	14:K:147:GLU:HG3	2.00	0.42
16:M:11:ARG:O	16:M:15:GLU:HG3	2.19	0.42
16:M:71:TRP:N	37:M:4394:HOH:O	2.52	0.42
16:M:152:GLU:HA	16:M:152:GLU:OE1	2.18	0.42
1:O:192:A:N6	1:O:194:A:C2	2.88	0.42
1:O:297:U:H1'	37:O:3416:HOH:O	2.19	0.42
1:O:661:G:C4	1:O:686:A:C2	3.08	0.42
1:O:699:C:H5'	37:O:3489:HOH:O	2.19	0.42
1:O:1086:A:N6	25:V:11:VAL:HG11	2.34	0.42
1:O:1167:G:O2'	1:O:1168:C:H5'	2.18	0.42
1:O:1593:C:C6	18:O:120:ARG:HD3	2.54	0.42
1:O:2035:C:H2'	1:O:2036:C:H6	1.84	0.42
1:O:2456:A:H2'	1:O:2457:U:C6	2.54	0.42
1:O:2890:A:H1'	23:T:56:ARG:HH21	1.77	0.42
11:H:74:ASN:HD22	11:H:141:ASN:CG	2.27	0.42
11:H:82:LYS:NZ	11:H:82:LYS:CB	2.82	0.42
18:O:71:LYS:O	18:O:71:LYS:HG3	2.19	0.42
1:O:731:U:O2'	1:O:732:C:H5'	2.20	0.42
1:O:1186:C:N4	1:O:1187:U:C4	2.87	0.42
1:O:1308:A:C5	1:O:1309:U:C5	3.07	0.42
1:O:2314:G:H2'	1:O:2315:C:H5'	2.00	0.42
1:O:2392:C:N3	37:O:4305:HOH:O	2.36	0.42
1:O:2781:U:H2'	1:O:2782:G:C5'	2.48	0.42
1:O:2820:A:H2'	1:O:2821:C:O4'	2.19	0.42
2:9:3007:G:OP1	16:M:23:ARG:NE	2.52	0.42
2:9:3042:C:O2	7:D:76:ARG:NH1	2.51	0.42
4:A:211:LYS:HB3	4:A:212:PRO:CD	2.43	0.42
5:B:305:ASP:O	5:B:306:LYS:CB	2.66	0.42
7:D:95:THR:OG1	7:D:174:VAL:HG22	2.19	0.42
9:F:6:PHE:CD1	9:F:6:PHE:C	2.97	0.42
13:J:109:LEU:CD1	13:J:113:ILE:HD11	2.48	0.42
13:J:130:MET:SD	23:T:25:ASP:O	2.77	0.42
16:M:163:PHE:HA	37:M:2180:HOH:O	2.18	0.42
16:M:170:GLU:O	16:M:174:GLU:HG3	2.18	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:N:44:ASN:HB3	17:N:67:SER:O	2.19	0.42
20:Q:119:VAL:O	20:Q:119:VAL:CG1	2.67	0.42
21:R:8:PRO:HD2	24:U:32:ALA:HA	2.01	0.42
28:Y:11:THR:CG2	28:Y:23:ARG:HB2	2.49	0.42
1:0:17:G:H2'	1:0:18:C:C6	2.55	0.42
1:0:40:C:H4'	37:0:6394:HOH:O	2.18	0.42
1:0:317:A:OP1	22:S:52:ARG:O	2.37	0.42
1:0:453:A:H4'	1:0:455:A:N7	2.34	0.42
1:0:466:A:H2'	1:0:467:G:O4'	2.19	0.42
1:0:578:C:O2	1:0:1112:G:H4'	2.20	0.42
1:0:710:G:O2'	1:0:711:G:H5'	2.20	0.42
1:0:732:C:O2'	1:0:733:U:H5'	2.19	0.42
1:0:2089:A:O2'	1:0:2090:G:H5'	2.20	0.42
1:0:2748:G:C4'	37:0:6917:HOH:O	2.67	0.42
4:A:69:LEU:CD2	4:A:120:ARG:HB3	2.44	0.42
4:A:105:VAL:CG1	4:A:106:CYS:N	2.82	0.42
4:A:176:HIS:CD2	4:A:176:HIS:C	2.98	0.42
5:B:24:PRO:HG2	5:B:204:GLY:HA2	2.01	0.42
7:D:27:ILE:HD11	7:D:37:ALA:CB	2.50	0.42
14:K:105:TYR:C	14:K:105:TYR:CD1	2.98	0.42
14:K:126:SER:O	14:K:127:GLU:C	2.62	0.42
15:L:18:GLY:O	15:L:21:ALA:HB3	2.19	0.42
15:L:27:ARG:O	15:L:28:MET:C	2.63	0.42
15:L:35:PRO:HD2	15:L:38:VAL:CG2	2.49	0.42
15:L:42:ARG:HA	15:L:43:PRO:HD3	1.86	0.42
21:R:10:VAL:HG11	24:U:36:ALA:HA	2.01	0.42
25:V:6:GLN:HA	25:V:52:VAL:HG23	2.01	0.42
31:2:87:ARG:NH1	37:2:8520:HOH:O	2.53	0.42
1:0:424:C:H2'	1:0:425:U:C6	2.54	0.42
1:0:553:G:O4'	1:0:1325:G:H5'	2.19	0.42
1:0:1084:C:O5'	1:0:1084:C:H6	2.02	0.42
1:0:1133:A:H2'	1:0:1134:G:O4'	2.19	0.42
1:0:1224:G:H2'	1:0:1225:C:C6	2.54	0.42
1:0:1512:G:O2'	1:0:1513:C:H5'	2.18	0.42
1:0:2028:U:H2'	1:0:2029:C:H6	1.84	0.42
1:0:2807:U:H1'	37:0:9700:HOH:O	2.19	0.42
1:0:2857:C:O2'	1:0:2858:U:H5'	2.19	0.42
1:0:2864:U:H3'	1:0:2865:G:C8	2.55	0.42
4:A:35:GLY:O	4:A:36:ASP:CB	2.58	0.42
6:C:13:ASP:N	37:C:8437:HOH:O	2.52	0.42
6:C:154:VAL:O	6:C:158:GLU:HG3	2.19	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:H:58:HIS:CE1	11:H:59:ASN:ND2	2.88	0.42
16:M:3:GLY:HA3	37:M:853:HOH:O	2.19	0.42
20:Q:82:GLU:HG3	20:Q:83:LYS:N	2.34	0.42
1:0:23:G:C6	1:0:24:G:N1	2.88	0.42
1:0:102:A:H4'	37:0:6713:HOH:O	2.20	0.42
1:0:210:U:O2'	1:0:211:U:H5'	2.20	0.42
1:0:470:U:O2'	29:Z:16:HIS:CD2	2.73	0.42
1:0:818:A:H2	28:Y:13:ARG:HA	1.85	0.42
1:0:1406:A:H4'	1:0:1407:A:C5'	2.49	0.42
1:0:1504:A:H4'	1:0:1506:U:C5	2.54	0.42
1:0:1513:C:O2'	1:0:1514:C:H5'	2.19	0.42
1:0:2346:C:H4'	7:D:52:THR:HG22	2.02	0.42
1:0:2388:C:OP1	37:0:4061:HOH:O	2.21	0.42
1:0:2440:C:H5''	37:0:3304:HOH:O	2.19	0.42
1:0:2526:C:C6	1:0:2526:C:H5'	2.54	0.42
1:0:2547:C:H2'	1:0:2548:C:H6	1.84	0.42
1:0:2804:C:H2'	1:0:2805:A:O4'	2.20	0.42
37:0:9185:HOH:O	27:X:163:THR:HG23	2.18	0.42
5:B:165:ARG:HG2	5:B:165:ARG:HH11	1.85	0.42
5:B:223:ARG:HG3	5:B:232:TRP:O	2.19	0.42
16:M:143:ARG:NH1	16:M:173:ASP:CG	2.77	0.42
22:S:48:VAL:CG1	22:S:96:VAL:HG13	2.49	0.42
26:W:9:VAL:HG22	26:W:88:GLU:OE2	2.19	0.42
29:Z:45:ARG:NH2	37:Z:8427:HOH:O	2.46	0.42
30:1:22:PRO:HG2	30:1:25:VAL:CG2	2.49	0.42
1:0:115:U:H2'	37:0:8671:HOH:O	2.19	0.42
1:0:422:G:O2'	1:0:423:A:H5'	2.19	0.42
1:0:560:C:H2'	1:0:561:G:H8	1.84	0.42
1:0:716:G:C2'	1:0:717:C:O5'	2.68	0.42
1:0:834:G:H5''	1:0:835:U:O5'	2.20	0.42
1:0:963:C:H2'	1:0:964:G:C8	2.54	0.42
1:0:1252:A:O3'	37:0:7105:HOH:O	2.22	0.42
1:0:1594:C:O2'	1:0:1607:A:H4'	2.20	0.42
1:0:2301:A:H5''	1:0:2302:A:C5'	2.48	0.42
1:0:2740:G:H2'	1:0:2741:A:O4'	2.19	0.42
1:0:2831:C:H2'	1:0:2832:C:C5'	2.49	0.42
4:A:125:ASN:ND2	37:A:8532:HOH:O	2.52	0.42
6:C:59:GLU:C	6:C:71:PRO:HA	2.45	0.42
7:D:27:ILE:O	7:D:69:ILE:HG22	2.19	0.42
7:D:52:THR:N	7:D:70:GLY:O	2.53	0.42
7:D:128:LEU:N	37:D:5495:HOH:O	2.53	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:F:60:VAL:O	9:F:61:MET:C	2.63	0.42
11:H:150:LYS:HA	11:H:153:VAL:HG22	2.02	0.42
15:L:169:ARG:HD2	37:L:8589:HOH:O	2.20	0.42
16:M:73:ALA:HB1	16:M:74:PRO:CD	2.49	0.42
16:M:74:PRO:HB3	37:M:4713:HOH:O	2.20	0.42
16:M:78:MET:CB	16:M:79:PRO:HD3	2.50	0.42
22:S:27:LEU:HG	22:S:39:ASN:HA	2.02	0.42
1:O:151:A:C2	1:O:442:A:C8	3.08	0.42
1:O:332:G:O2'	1:O:333:G:H5'	2.19	0.42
1:O:661:G:C6	1:O:686:A:C2	3.08	0.42
1:O:862:U:H2'	1:O:863:G:C8	2.55	0.42
1:O:1127:C:C5	1:O:1128:U:C4	3.07	0.42
1:O:1860:U:H2'	1:O:1861:C:O4'	2.20	0.42
1:O:1979:G:P	37:O:5724:HOH:O	2.77	0.42
1:O:2084:C:H2'	1:O:2085:A:H8	1.85	0.42
1:O:2088:C:H1'	1:O:2841:A:N1	2.34	0.42
1:O:2381:C:H2'	1:O:2382:A:H8	1.85	0.42
4:A:95:PRO:O	4:A:99:ILE:HG12	2.20	0.42
5:B:63:GLU:O	5:B:63:GLU:HG3	2.19	0.42
5:B:147:VAL:O	5:B:150:ALA:HB3	2.20	0.42
7:D:23:VAL:HG23	7:D:41:LEU:HD22	2.01	0.42
7:D:81:GLU:O	7:D:83:PHE:N	2.53	0.42
11:H:57:ARG:C	11:H:59:ASN:N	2.76	0.42
11:H:111:MET:O	11:H:114:PRO:HD3	2.19	0.42
12:I:131:THR:HG22	12:I:134:GLU:N	2.21	0.42
13:J:10:GLN:H	13:J:10:GLN:CD	2.24	0.42
16:M:67:ALA:HA	16:M:71:TRP:HB3	2.02	0.42
16:M:139:TRP:HA	16:M:139:TRP:HE3	1.85	0.42
18:O:99:ARG:HA	37:O:7545:HOH:O	2.19	0.42
20:Q:129:ALA:O	20:Q:130:MET:HB2	2.19	0.42
22:S:73:HIS:CD2	22:S:88:PRO:HG3	2.54	0.42
25:V:55:GLY:CA	25:V:146:ILE:HG13	2.49	0.42
29:Z:28:HIS:O	29:Z:32:LYS:N	2.44	0.42
1:O:617:C:O2'	27:X:158:LYS:O	2.32	0.42
1:O:682:A:H3'	1:O:683:G:H8	1.85	0.42
1:O:778:C:C4	1:O:779:U:C4	3.08	0.42
1:O:949:U:O2'	19:P:40:HIS:HE1	2.02	0.42
1:O:1213:C:C2'	1:O:1214:G:H5'	2.50	0.42
1:O:1351:G:H1'	37:O:3528:HOH:O	2.20	0.42
1:O:2266:A:H2'	1:O:2267:G:H8	1.85	0.42
1:O:2727:A:C2'	1:O:2728:C:H5'	2.49	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:A:109:GLU:CD	4:A:113:GLY:H	2.28	0.42
6:C:65:ARG:HG3	6:C:67:GLN:HB2	2.02	0.42
6:C:185:LYS:HD3	6:C:186:TYR:CE1	2.55	0.42
9:F:34:ASN:HA	15:L:4:ALA:HB2	2.01	0.42
12:I:66:ASP:HA	37:I:6706:HOH:O	2.19	0.42
18:O:115:SER:C	18:O:117:SER:N	2.78	0.42
19:P:10:THR:O	19:P:11:ARG:C	2.63	0.42
19:P:88:ALA:N	37:P:1295:HOH:O	2.52	0.42
22:S:87:VAL:HB	22:S:88:PRO:HD2	2.01	0.42
25:V:126:ASP:HB3	25:V:135:GLY:O	2.20	0.42
28:Y:23:ARG:NH1	37:Y:8404:HOH:O	2.52	0.42
1:0:410:A:H5'	1:0:411:A:H2'	2.01	0.41
1:0:432:G:O2'	1:0:433:C:H5'	2.20	0.41
1:0:539:G:H2'	1:0:540:A:H8	1.84	0.41
1:0:549:A:O2'	1:0:550:C:H5'	2.20	0.41
1:0:663:C:H2'	1:0:664:U:O4'	2.20	0.41
1:0:830:G:N3	37:0:8779:HOH:O	2.37	0.41
1:0:1815:A:H4'	1:0:2751:C:O4'	2.20	0.41
1:0:1829:A:C2'	1:0:1830:C:H5'	2.49	0.41
9:F:26:THR:HB	9:F:102:GLY:HA3	2.02	0.41
16:M:108:SER:HA	16:M:109:PRO:HD3	1.84	0.41
20:Q:3:SER:O	20:Q:4:TYR:C	2.62	0.41
20:Q:114:VAL:HA	20:Q:144:GLU:O	2.19	0.41
1:0:470:U:H2'	1:0:471:G:O4'	2.20	0.41
1:0:533:U:H4'	1:0:534:C:O5'	2.20	0.41
1:0:684:G:H2'	1:0:685:C:C6	2.55	0.41
1:0:694:A:H2'	1:0:695:C:C5'	2.49	0.41
1:0:720:G:H5'	37:0:8867:HOH:O	2.20	0.41
1:0:1852:A:H4'	4:A:230:SER:HB2	2.01	0.41
1:0:1902:G:H2'	1:0:1903:U:O4'	2.20	0.41
1:0:2255:A:N1	1:0:2256:G:C4	2.88	0.41
1:0:2274:A:O2'	15:L:86:MET:HE2	2.20	0.41
1:0:2389:U:H4'	19:P:53:HIS:CD2	2.55	0.41
1:0:2439:C:H5'	37:0:4924:HOH:O	2.19	0.41
1:0:2547:C:H2'	1:0:2548:C:C6	2.55	0.41
1:0:2734:G:O2'	1:0:2735:U:H5'	2.20	0.41
4:A:9:ARG:HG2	4:A:16:PHE:CD2	2.55	0.41
5:B:55:ASN:HB3	5:B:64:GLY:N	2.35	0.41
6:C:104:ASP:O	6:C:108:GLN:HG3	2.21	0.41
6:C:162:VAL:HG12	6:C:192:ILE:CD1	2.48	0.41
7:D:58:VAL:CG1	7:D:59:GLY:N	2.82	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:D:169:THR:C	7:D:170:TYR:HD1	2.28	0.41
9:F:110:GLU:HG2	37:F:6926:HOH:O	2.20	0.41
10:G:12:ILE:HD12	37:G:692:HOH:O	2.19	0.41
15:L:169:ARG:HB3	37:L:8589:HOH:O	2.20	0.41
18:O:7:LYS:HD3	18:O:21:VAL:HG21	2.03	0.41
18:O:18:LYS:O	18:O:21:VAL:HG22	2.20	0.41
22:S:71:VAL:HG12	22:S:72:ILE:N	2.33	0.41
26:W:61:ARG:N	37:W:1449:HOH:O	2.50	0.41
27:X:133:HIS:CD2	37:X:8583:HOH:O	2.58	0.41
1:0:154:C:H3'	15:L:188:ARG:NH1	2.36	0.41
1:0:711:G:C2	1:0:718:C:C2	3.08	0.41
1:0:1200:A:H4'	37:0:6722:HOH:O	2.20	0.41
1:0:1223:G:O2'	1:0:1224:G:H5'	2.20	0.41
1:0:1706:G:OP1	18:O:65:ARG:HD2	2.20	0.41
1:0:1768:C:C2'	1:0:1769:C:H5'	2.50	0.41
1:0:2346:C:O3'	7:D:52:THR:HG23	2.20	0.41
1:0:2379:G:H4'	1:0:2380:A:C5'	2.50	0.41
1:0:2553:A:H2'	1:0:2553:A:N3	2.34	0.41
1:0:2604:A:H5'	37:0:5218:HOH:O	2.19	0.41
1:0:2637:A:H8	3:4:75:C:OP1	2.03	0.41
1:0:2729:C:H2'	1:0:2730:G:C8	2.54	0.41
1:0:2832:C:H5	37:0:6602:HOH:O	2.02	0.41
1:0:2904:U:H4'	26:W:8:ARG:HH12	1.84	0.41
4:A:1:GLY:HA3	37:A:8525:HOH:O	2.19	0.41
5:B:36:PRO:CG	5:B:169:GLY:H	2.32	0.41
5:B:109:LEU:CG	5:B:113:LEU:HD12	2.50	0.41
5:B:168:GLY:HA3	5:B:174:ARG:HB3	2.03	0.41
15:L:153:THR:C	15:L:155:HIS:N	2.79	0.41
18:O:137:LEU:O	18:O:138:GLU:C	2.63	0.41
22:S:30:ASP:O	22:S:33:GLU:HB3	2.20	0.41
1:0:81:G:N3	1:0:98:A:C2	2.88	0.41
1:0:314:G:N2	1:0:316:A:H3'	2.35	0.41
1:0:331:A:C6	1:0:332:G:C4	3.08	0.41
1:0:462:A:C4	30:1:37:HIS:CG	3.09	0.41
1:0:1325:G:O2'	1:0:1326:U:H5'	2.20	0.41
1:0:1329:A:N1	35:0:8513:CL:CL	2.90	0.41
1:0:1439:C:H5''	30:1:41:HIS:HE1	1.85	0.41
1:0:1469:C:N3	1:0:1472:C:OP2	2.53	0.41
1:0:1486:A:C4	30:1:2:LYS:HG3	2.54	0.41
1:0:1631:A:H2'	1:0:1632:A:C8	2.55	0.41
1:0:2025:G:C6	1:0:2026:C:C4	3.08	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:2688:U:H2'	1:0:2689:A:H8	1.85	0.41
1:0:2724:U:H2'	1:0:2725:G:O4'	2.20	0.41
2:9:3042:C:N4	2:9:3044:A:N1	2.68	0.41
7:D:140:ARG:HH11	7:D:140:ARG:HG3	1.84	0.41
9:F:110:GLU:HA	9:F:113:ASP:OD2	2.20	0.41
10:G:63:ARG:HB2	10:G:66:LEU:HG	2.02	0.41
14:K:80:ASP:HB2	14:K:90:ARG:O	2.20	0.41
15:L:5:TYR:C	15:L:7:TYR:H	2.28	0.41
16:M:170:GLU:HA	16:M:173:ASP:OD2	2.21	0.41
20:Q:52:GLU:CG	20:Q:54:ASP:OD2	2.68	0.41
27:X:189:ASN:HD21	27:X:192:ASP:H	1.69	0.41
1:0:84:G:O2'	1:0:85:C:H5'	2.21	0.41
1:0:202:U:H2'	1:0:203:G:O4'	2.21	0.41
1:0:475:G:H5'	6:C:73:LEU:HD23	2.01	0.41
1:0:1006:A:N3	1:0:2298:C:O2'	2.45	0.41
1:0:1165:G:H5'	37:0:5033:HOH:O	2.19	0.41
1:0:1425:G:O2'	1:0:1426:C:H5'	2.21	0.41
1:0:1583:U:H1'	37:0:9471:HOH:O	2.20	0.41
1:0:1795:G:H2'	1:0:1796:A:O4'	2.20	0.41
1:0:2132:C:H1'	15:L:124:GLY:HA3	2.03	0.41
1:0:2281:C:C5	1:0:2282:U:C4	3.08	0.41
1:0:2654:C:H5'	37:B:8661:HOH:O	2.20	0.41
1:0:2816:A:H4'	37:0:3386:HOH:O	2.21	0.41
1:0:2821:C:H4'	5:B:116:PRO:HG3	2.02	0.41
37:0:5144:HOH:O	13:J:87:ARG:CZ	2.68	0.41
6:C:4:THR:HB	6:C:135:GLU:OE1	2.20	0.41
6:C:7:ASP:OD1	6:C:11:ASN:O	2.38	0.41
12:I:52:GLN:CG	12:I:53:ILE:N	2.78	0.41
14:K:54:PRO:HG2	14:K:57:VAL:CG2	2.51	0.41
14:K:73:VAL:HG23	14:K:74:THR:H	1.84	0.41
21:R:81:ILE:HG22	24:U:29:ASN:OD1	2.20	0.41
22:S:43:ASN:C	22:S:45:GLY:H	2.28	0.41
24:U:42:ASN:N	24:U:43:PRO:HD3	2.34	0.41
24:U:55:ARG:O	24:U:59:ILE:HG12	2.19	0.41
25:V:7:LEU:HA	25:V:7:LEU:HD23	1.89	0.41
1:0:111:C:O2'	1:0:112:G:H5'	2.20	0.41
1:0:962:C:C2'	1:0:963:C:H5'	2.51	0.41
1:0:1446:U:H2'	21:R:55:GLN:NE2	2.36	0.41
1:0:1638:U:H5'	37:0:6658:HOH:O	2.21	0.41
1:0:1891:G:H1'	1:0:1972:U:O2	2.20	0.41
1:0:2054:A:H4'	20:Q:135:ALA:O	2.21	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:2090:G:H2'	1:0:2091:G:C8	2.55	0.41
1:0:2329:C:O2'	1:0:2330:U:H5'	2.20	0.41
1:0:2582:G:O3'	13:J:41:LYS:HA	2.20	0.41
37:0:4185:HOH:O	16:M:21:HIS:HD2	2.04	0.41
2:9:3088:G:N2	2:9:3089:C:C2	2.89	0.41
37:9:8493:HOH:O	25:V:133:LYS:HG3	2.19	0.41
4:A:46:GLU:O	4:A:55:VAL:N	2.46	0.41
5:B:83:ALA:HB2	5:B:101:TRP:CD2	2.55	0.41
5:B:182:VAL:O	5:B:183:GLU:C	2.63	0.41
5:B:224:LYS:HD3	5:B:224:LYS:HA	1.72	0.41
6:C:107:ARG:NH1	6:C:107:ARG:CB	2.82	0.41
8:E:14:GLU:HG2	8:E:15:GLN:N	2.35	0.41
8:E:133:VAL:HG12	8:E:141:VAL:HG13	2.03	0.41
12:I:24:SER:HA	12:I:86:MET:SD	2.61	0.41
16:M:13:ARG:NH1	16:M:13:ARG:O	2.53	0.41
24:U:27:LEU:O	24:U:30:ALA:HB3	2.20	0.41
25:V:22:GLU:HG2	25:V:27:HIS:CD2	2.55	0.41
1:0:569:A:H5''	1:0:587:A:N1	2.34	0.41
1:0:1310:U:C2'	1:0:1311:G:O5'	2.68	0.41
1:0:1385:G:O3'	26:W:49:ARG:NH1	2.53	0.41
1:0:1461:U:H2'	1:0:1462:C:C6	2.56	0.41
1:0:1666:C:H2'	1:0:1667:A:H8	1.85	0.41
1:0:1825:U:C4'	1:0:1999:C:H5''	2.51	0.41
1:0:1986:G:C6	1:0:1987:C:N4	2.89	0.41
1:0:2092:G:H5''	1:0:2613:G:OP1	2.21	0.41
1:0:2278:U:H3'	37:0:4546:HOH:O	2.20	0.41
1:0:2748:G:H5''	37:0:6698:HOH:O	2.20	0.41
2:9:3028:U:H5''	16:M:40:ASN:ND2	2.36	0.41
7:D:23:VAL:HG21	7:D:45:THR:CG2	2.50	0.41
7:D:55:LYS:C	37:D:4069:HOH:O	2.64	0.41
8:E:7:ILE:HG22	8:E:45:ASP:O	2.21	0.41
8:E:157:LYS:NZ	37:E:2401:HOH:O	2.54	0.41
11:H:136:VAL:HG22	11:H:137:ASN:N	2.36	0.41
12:I:36:VAL:HG12	12:I:37:ALA:N	2.35	0.41
13:J:22:ASP:HB3	13:J:96:VAL:HG13	2.02	0.41
16:M:17:ARG:NE	37:M:922:HOH:O	2.32	0.41
16:M:143:ARG:NH1	16:M:173:ASP:OD1	2.53	0.41
22:S:92:ASP:O	22:S:93:THR:C	2.63	0.41
23:T:37:GLU:O	23:T:40:ALA:HB3	2.20	0.41
1:0:45:A:N6	1:0:147:G:C4	2.89	0.41
1:0:235:C:O2'	1:0:236:A:H2'	2.21	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:597:A:H8	1:0:597:A:O5'	2.03	0.41
1:0:844:A:C6	1:0:882:A:C6	3.09	0.41
1:0:1066:U:H2'	1:0:1067:A:C8	2.55	0.41
1:0:1157:C:H2'	1:0:1158:G:H8	1.85	0.41
1:0:1206:U:H6	1:0:1206:U:C5'	2.24	0.41
1:0:1491:G:H4'	1:0:1492:A:OP2	2.20	0.41
1:0:1667:A:O2'	1:0:1668:U:H5'	2.21	0.41
1:0:1667:A:C2	1:0:1668:U:C2	3.09	0.41
1:0:1921:A:C6	1:0:1922:A:C2	3.08	0.41
1:0:2765:C:H2'	1:0:2766:A:H8	1.86	0.41
37:0:5996:HOH:O	14:K:102:ASP:HA	2.21	0.41
2:9:3050:G:C6	2:9:3051:A:C6	3.09	0.41
5:B:84:LEU:HD23	5:B:178:ALA:HB1	2.03	0.41
6:C:141:SER:HB3	37:C:8413:HOH:O	2.20	0.41
6:C:165:ASP:O	6:C:168:ARG:HB3	2.21	0.41
8:E:31:ARG:NH1	8:E:68:HIS:ND1	2.69	0.41
11:H:130:HIS:CG	11:H:133:ILE:HD11	2.55	0.41
11:H:137:ASN:O	11:H:138:PRO:C	2.63	0.41
37:J:1387:HOH:O	23:T:20:MET:HE3	2.20	0.41
16:M:154:LEU:HD12	16:M:156:GLU:O	2.21	0.41
18:O:37:ARG:O	18:O:40:VAL:HB	2.21	0.41
20:Q:46:TYR:HD2	20:Q:47:LEU:HD23	1.86	0.41
21:R:35:GLY:N	37:R:1504:HOH:O	2.37	0.41
22:S:55:PHE:O	22:S:56:ALA:C	2.63	0.41
25:V:137:GLN:HG3	25:V:137:GLN:O	2.21	0.41
28:Y:27:ALA:O	28:Y:31:ILE:N	2.50	0.41
1:0:21:G:H5''	20:Q:2:ILE:HA	1.97	0.41
1:0:240:C:O2	1:0:240:C:C2'	2.68	0.41
1:0:321:A:O2'	1:0:322:G:H5'	2.20	0.41
1:0:451:C:N4	1:0:452:G:C6	2.89	0.41
1:0:622:G:P	27:X:148:GLY:HA3	2.60	0.41
1:0:696:C:HO2'	1:0:697:G:H5'	1.86	0.41
1:0:1023:C:H2'	1:0:1024:G:O4'	2.21	0.41
1:0:1354:G:O6	14:K:5:LYS:HE3	2.21	0.41
1:0:1603:A:H5'	1:0:1605:G:C4'	2.50	0.41
1:0:1685:A:H4'	1:0:1686:C:OP2	2.21	0.41
1:0:1762:C:H4'	37:0:4111:HOH:O	2.21	0.41
1:0:1943:C:C4'	4:A:212:PRO:HA	2.50	0.41
1:0:2050:G:OP1	20:Q:79:ARG:HB3	2.21	0.41
1:0:2487:C:H5	37:0:4342:HOH:O	2.03	0.41
1:0:2502:C:H4'	11:H:151:MET:CG	2.45	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:2690:U:H4'	8:E:111:LYS:CE	2.50	0.41
1:0:2698:G:H2'	1:0:2699:A:C8	2.55	0.41
1:0:2748:G:C3'	37:0:6917:HOH:O	2.68	0.41
1:0:2810:G:OP1	5:B:17:LYS:NZ	2.41	0.41
37:0:5721:HOH:O	8:E:139:GLU:HA	2.21	0.41
5:B:84:LEU:HD13	5:B:84:LEU:C	2.46	0.41
7:D:87:ALA:O	7:D:88:LEU:C	2.63	0.41
8:E:11:VAL:HG13	8:E:23:GLU:O	2.20	0.41
9:F:115:VAL:O	9:F:116:GLU:C	2.64	0.41
11:H:47:GLU:CG	11:H:133:ILE:HD12	2.49	0.41
14:K:34:GLY:C	14:K:36:ASP:N	2.77	0.41
15:L:54:TYR:CG	15:L:55:LYS:N	2.89	0.41
16:M:43:VAL:HG13	16:M:118:ILE:HD11	2.02	0.41
16:M:69:TYR:HE2	16:M:183:ASP:OD2	2.03	0.41
18:O:7:LYS:CD	18:O:21:VAL:CG2	2.99	0.41
21:R:81:ILE:HG12	37:R:4527:HOH:O	2.21	0.41
24:U:39:ALA:C	24:U:41:GLU:N	2.79	0.41
25:V:48:VAL:CG1	25:V:48:VAL:O	2.68	0.41
26:W:20:GLU:CG	26:W:21:PRO:HD2	2.50	0.41
26:W:85:VAL:CG1	26:W:86:GLU:N	2.81	0.41
27:X:189:ASN:ND2	27:X:191:ASP:N	2.68	0.41
28:Y:42:CYS:SG	28:Y:43:GLY:N	2.94	0.41
29:Z:25:LYS:HD2	30:1:48:ASP:CA	2.51	0.41
1:0:276:C:H4'	37:0:6202:HOH:O	2.20	0.41
1:0:292:G:H1'	1:0:360:A:N6	2.36	0.41
1:0:922:A:N7	1:0:2281:C:H5'	2.36	0.41
1:0:1169:U:C5	1:0:1170:U:C4	3.09	0.41
1:0:1235:G:H1'	12:I:63:ILE:HG23	2.03	0.41
1:0:1310:U:H2'	1:0:1311:G:O5'	2.20	0.41
1:0:1427:A:H61	1:0:1440:U:H1'	1.85	0.41
1:0:1789:G:H2'	1:0:1790:C:O5'	2.21	0.41
1:0:1985:U:H5''	37:0:4645:HOH:O	2.20	0.41
1:0:2414:A:C2	1:0:2415:A:C6	3.09	0.41
1:0:2613:G:H2'	1:0:2614:C:C6	2.55	0.41
4:A:195:ASN:O	4:A:196:ALA:C	2.64	0.41
5:B:69:VAL:HA	5:B:70:PRO:HD3	1.93	0.41
6:C:107:ARG:CB	6:C:107:ARG:HH11	2.33	0.41
7:D:18:ILE:HD13	7:D:84:LEU:CD1	2.51	0.41
7:D:36:ASN:C	37:D:7500:HOH:O	2.64	0.41
7:D:91:ALA:HB2	7:D:106:PHE:CD2	2.56	0.41
9:F:56:PRO:HG2	15:L:43:PRO:O	2.20	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:G:20:VAL:O	10:G:24:VAL:HG23	2.21	0.41
11:H:112:ARG:O	11:H:113:ALA:C	2.64	0.41
12:I:90:LYS:N	35:I:8502:CL:CL	2.81	0.41
13:J:14:LYS:HD2	13:J:45:PRO:HG3	2.03	0.41
13:J:49:LEU:HA	13:J:73:VAL:CG1	2.51	0.41
13:J:101:ASN:HB2	13:J:103:ASP:OD2	2.21	0.41
20:Q:33:ARG:HG3	37:Q:8568:HOH:O	2.20	0.41
22:S:98:VAL:HG11	22:S:101:LEU:HD21	2.02	0.41
24:U:12:THR:HG23	24:U:14:ALA:H	1.86	0.41
1:0:106:A:H2'	1:0:107:U:O4'	2.21	0.40
1:0:275:G:H1'	37:0:3022:HOH:O	2.22	0.40
1:0:636:G:N2	37:0:8769:HOH:O	2.47	0.40
1:0:709:G:O2'	17:N:25:VAL:HG12	2.21	0.40
1:0:1102:C:O2'	1:0:1103:C:H5'	2.21	0.40
1:0:1735:C:OP2	5:B:234:ARG:HG3	2.21	0.40
1:0:1788:U:O2'	1:0:1789:G:H5'	2.21	0.40
1:0:2274:A:H1'	15:L:86:MET:CE	2.51	0.40
1:0:2731:G:H2'	1:0:2732:U:O4'	2.21	0.40
2:9:3045:A:H2'	2:9:3046:C:H6	1.85	0.40
4:A:135:VAL:N	37:A:8589:HOH:O	2.54	0.40
5:B:55:ASN:CB	5:B:63:GLU:HA	2.49	0.40
5:B:144:THR:CG2	5:B:145:HIS:N	2.84	0.40
8:E:84:MET:HE1	8:E:148:ILE:HD12	1.97	0.40
8:E:99:GLY:N	37:E:4191:HOH:O	2.53	0.40
11:H:84:ARG:NH2	11:H:135:TRP:CH2	2.84	0.40
11:H:113:ALA:N	11:H:114:PRO:CD	2.83	0.40
15:L:20:ILE:O	15:L:24:MET:HG2	2.21	0.40
15:L:32:ARG:NH2	37:L:8596:HOH:O	2.53	0.40
16:M:104:ILE:O	16:M:104:ILE:HG13	2.19	0.40
25:V:113:SER:C	25:V:115:THR:H	2.29	0.40
27:X:141:THR:O	27:X:142:SER:C	2.64	0.40
1:0:10:U:H5'	37:0:5458:HOH:O	2.20	0.40
1:0:533:U:H2'	1:0:2814:A:C6	2.57	0.40
1:0:968:G:O2'	1:0:969:G:H5'	2.21	0.40
1:0:1072:G:P	27:X:154:ARG:NH2	2.94	0.40
1:0:1139:U:H2'	1:0:1140:C:H6	1.86	0.40
1:0:2020:C:O2'	1:0:2021:C:H5'	2.20	0.40
1:0:2069:U:H5'	37:0:4221:HOH:O	2.20	0.40
1:0:2464:C:H5''	1:0:2465:A:OP1	2.22	0.40
1:0:2547:C:C2	1:0:2548:C:C5	3.10	0.40
4:A:22:ARG:NH1	37:A:8558:HOH:O	2.53	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:B:98:THR:HG22	5:B:99:GLU:H	1.87	0.40
7:D:57:THR:HG23	7:D:63:ILE:HA	2.03	0.40
8:E:81:GLU:HA	8:E:133:VAL:O	2.21	0.40
8:E:101:GLU:HA	8:E:118:ILE:HG13	2.02	0.40
11:H:27:LYS:HG3	11:H:58:HIS:CD2	2.56	0.40
11:H:57:ARG:O	11:H:61:LEU:HD22	2.21	0.40
12:I:129:PHE:CD1	12:I:129:PHE:C	2.99	0.40
13:J:113:ILE:HD12	13:J:128:ALA:HB2	2.04	0.40
20:Q:111:ILE:HG23	20:Q:145:LEU:CD1	2.51	0.40
21:R:20:PHE:N	21:R:20:PHE:CD2	2.87	0.40
23:T:46:ALA:HB1	23:T:52:THR:HG21	2.04	0.40
30:1:18:ASN:HD21	30:1:40:ARG:H	1.69	0.40
1:0:407:A:C2	1:0:408:A:C4	3.10	0.40
1:0:506:G:N1	1:0:509:A:OP2	2.53	0.40
1:0:876:A:N3	1:0:876:A:C2'	2.84	0.40
1:0:1004:C:H1'	37:0:4292:HOH:O	2.21	0.40
1:0:1201:C:H2'	1:0:1202:A:H5'	2.03	0.40
1:0:1562:C:O2	1:0:1562:C:C2'	2.70	0.40
1:0:1703:G:C2	1:0:1716:A:C4	3.09	0.40
3:3:75:C:H6	3:3:75:C:O5'	2.05	0.40
7:D:64:ARG:NE	7:D:67:ASP:HB3	2.37	0.40
7:D:170:TYR:O	7:D:171:ASP:CB	2.70	0.40
9:F:58:GLU:CD	15:L:27:ARG:HH22	2.22	0.40
10:G:12:ILE:HG22	10:G:17:GLN:HE22	1.86	0.40
15:L:37:VAL:CB	15:L:108:LYS:HG3	2.51	0.40
15:L:174:ARG:HG3	37:L:8521:HOH:O	2.20	0.40
25:V:75:GLY:HA3	37:V:5763:HOH:O	2.21	0.40
29:Z:2:GLY:O	29:Z:6:PRO:HG2	2.21	0.40
31:2:7:PHE:CD1	31:2:7:PHE:C	2.99	0.40
31:2:69:TYR:CB	31:2:78:HIS:CE1	3.04	0.40
1:0:212:A:H5'	1:0:214:U:O4'	2.22	0.40
1:0:311:C:O5'	1:0:311:C:H6	2.05	0.40
1:0:390:G:H5''	37:0:9056:HOH:O	2.20	0.40
1:0:508:A:C2'	1:0:509:A:H5'	2.52	0.40
1:0:731:U:H2'	1:0:732:C:C6	2.56	0.40
1:0:764:C:H2'	1:0:765:G:O4'	2.21	0.40
1:0:797:A:H5'	28:Y:10:ARG:HG2	2.02	0.40
1:0:962:C:H2'	1:0:963:C:H5'	2.03	0.40
1:0:1015:C:O5'	1:0:1015:C:H6	2.04	0.40
1:0:1116:U:C2'	1:0:1118:A:C2	3.05	0.40
1:0:1483:C:O2'	1:0:1484:G:H5'	2.22	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:1521:C:C4	1:0:1522:A:N7	2.90	0.40
1:0:1641:A:C8	1:0:1702:U:O4	2.75	0.40
1:0:1819:G:H5''	37:0:4167:HOH:O	2.20	0.40
1:0:1896:G:C6	1:0:1897:U:C4	3.09	0.40
1:0:2252:A:C6	1:0:2253:G:H1'	2.55	0.40
1:0:2408:A:H2	37:2:8515:HOH:O	2.04	0.40
2:9:3074:G:H1	2:9:3107:C:H42	1.68	0.40
4:A:51:ARG:NH2	37:A:8602:HOH:O	2.53	0.40
4:A:179:MET:HG2	4:A:186:TRP:HB2	2.02	0.40
5:B:55:ASN:ND2	5:B:67:GLU:OE2	2.54	0.40
5:B:175:LEU:O	5:B:178:ALA:HB3	2.22	0.40
5:B:202:VAL:CG1	5:B:301:VAL:HG13	2.41	0.40
6:C:21:VAL:C	6:C:23:GLU:N	2.77	0.40
7:D:40:ILE:HG23	37:D:5583:HOH:O	2.21	0.40
9:F:14:ASP:O	9:F:18:GLU:HG3	2.21	0.40
9:F:78:GLU:HB2	37:F:2750:HOH:O	2.22	0.40
11:H:3:GLY:HA2	11:H:57:ARG:NH1	2.36	0.40
13:J:98:VAL:HG22	13:J:102:GLU:C	2.47	0.40
15:L:60:ILE:HG12	15:L:134:ILE:HD12	2.03	0.40
18:O:115:SER:O	18:O:117:SER:N	2.54	0.40
24:U:45:ARG:C	24:U:47:LYS:N	2.77	0.40
25:V:64:THR:O	25:V:68:THR:HG22	2.20	0.40
25:V:142:ASP:HB3	25:V:145:GLY:H	1.85	0.40
27:X:182:PHE:HD2	27:X:200:THR:O	2.05	0.40
1:0:56:G:C5'	24:U:50:ARG:HH12	2.26	0.40
1:0:259:G:H21	15:L:58:GLN:NE2	2.20	0.40
1:0:666:A:H2'	1:0:667:C:O4'	2.22	0.40
1:0:678:G:OP2	6:C:107:ARG:NH2	2.54	0.40
1:0:1165:G:H4'	1:0:1174:A:HO2'	1.82	0.40
1:0:1166:A:N3	1:0:1166:A:H2'	2.36	0.40
1:0:1594:C:C5	18:O:120:ARG:CZ	3.05	0.40
1:0:1734:C:OP1	5:B:234:ARG:HD3	2.21	0.40
1:0:2381:C:H2'	1:0:2382:A:C8	2.57	0.40
1:0:2566:A:H4'	8:E:161:VAL:HG21	2.03	0.40
1:0:2659:U:C4'	20:Q:76:ASP:HB3	2.52	0.40
37:0:3666:HOH:O	27:X:186:ARG:HD2	2.21	0.40
3:4:75:C:H6	3:4:75:C:O5'	2.04	0.40
5:B:36:PRO:HA	5:B:168:GLY:HA3	2.00	0.40
10:G:23:ILE:HG22	10:G:70:ALA:CB	2.52	0.40
10:G:73:ASP:C	37:G:2218:HOH:O	2.63	0.40
14:K:40:PHE:CD1	14:K:40:PHE:C	3.00	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:L:59:GLY:C	15:L:141:ILE:HD11	2.45	0.40
15:L:87:MET:CB	31:2:46:ILE:HG21	2.50	0.40
16:M:72:GLU:HB3	16:M:171:HIS:HE1	1.87	0.40
24:U:27:LEU:HA	24:U:49:LEU:HD13	2.02	0.40
25:V:35:VAL:HA	25:V:36:PRO:HD3	1.75	0.40
25:V:73:LEU:CD1	25:V:112:LEU:C	2.95	0.40
31:2:62:THR:HB	37:2:8541:HOH:O	2.22	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 X-ray entries followed by that with respect to entries of similar resolution.

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
4	A	235/239 (98%)	204 (87%)	27 (12%)	4 (2%)	7	26
5	B	335/337 (99%)	295 (88%)	31 (9%)	9 (3%)	4	16
6	C	244/246 (99%)	221 (91%)	21 (9%)	2 (1%)	16	44
7	D	134/176 (76%)	93 (69%)	29 (22%)	12 (9%)	0	1
8	E	170/177 (96%)	157 (92%)	12 (7%)	1 (1%)	21	51
9	F	117/119 (98%)	103 (88%)	11 (9%)	3 (3%)	4	17
10	G	25/348 (7%)	21 (84%)	3 (12%)	1 (4%)	2	9
11	H	152/167 (91%)	131 (86%)	16 (10%)	5 (3%)	3	13
12	I	140/145 (97%)	128 (91%)	8 (6%)	4 (3%)	3	15
13	J	130/132 (98%)	111 (85%)	18 (14%)	1 (1%)	16	44
14	K	141/164 (86%)	118 (84%)	20 (14%)	3 (2%)	5	21
15	L	192/194 (99%)	165 (86%)	24 (12%)	3 (2%)	7	27
16	M	184/186 (99%)	161 (88%)	15 (8%)	8 (4%)	2	8
17	N	113/115 (98%)	104 (92%)	7 (6%)	2 (2%)	6	25

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
18	O	141/148 (95%)	132 (94%)	8 (6%)	1 (1%)	18	48
19	P	93/95 (98%)	88 (95%)	4 (4%)	1 (1%)	11	36
20	Q	148/154 (96%)	136 (92%)	12 (8%)	0	100	100
21	R	79/84 (94%)	75 (95%)	3 (4%)	1 (1%)	9	32
22	S	117/119 (98%)	99 (85%)	17 (14%)	1 (1%)	14	41
23	T	51/66 (77%)	45 (88%)	5 (10%)	1 (2%)	6	22
24	U	63/70 (90%)	56 (89%)	5 (8%)	2 (3%)	3	13
25	V	152/154 (99%)	139 (91%)	12 (8%)	1 (1%)	18	48
26	W	80/91 (88%)	72 (90%)	6 (8%)	2 (2%)	4	17
27	X	140/240 (58%)	134 (96%)	6 (4%)	0	100	100
28	Y	71/73 (97%)	57 (80%)	13 (18%)	1 (1%)	9	30
29	Z	54/56 (96%)	51 (94%)	3 (6%)	0	100	100
30	1	42/48 (88%)	40 (95%)	2 (5%)	0	100	100
31	2	90/92 (98%)	84 (93%)	6 (7%)	0	100	100
All	All	3633/4235 (86%)	3220 (89%)	344 (10%)	69 (2%)	6	23

All (69) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
5	B	139	ASP
7	D	16	PRO
7	D	93	LEU
7	D	95	THR
7	D	137	PRO
7	D	173	GLU
9	F	101	ALA
11	H	162	SER
11	H	164	ALA
16	M	154	LEU
16	M	162	ASP
16	M	164	ASP
16	M	181	ASP
16	M	183	ASP
24	U	43	PRO
5	B	34	GLY
5	B	169	GLY
5	B	184	ASP

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
6	C	8	LEU
7	D	20	LYS
7	D	61	PHE
7	D	147	ALA
7	D	171	ASP
12	I	5	GLU
13	J	119	GLN
14	K	80	ASP
15	L	6	SER
17	N	20	SER
18	O	116	SER
4	A	34	ASP
5	B	107	SER
7	D	11	HIS
7	D	36	ASN
11	H	72	VAL
11	H	138	PRO
12	I	7	ASP
12	I	78	ILE
14	K	21	ARG
16	M	139	TRP
19	P	23	THR
23	T	7	ASP
25	V	77	ALA
26	W	77	PHE
4	A	132	ASP
10	G	72	ASP
14	K	105	TYR
17	N	16	SER
5	B	185	GLY
5	B	206	THR
6	C	58	ALA
7	D	82	GLU
8	E	17	HIS
9	F	44	SER
9	F	64	PRO
11	H	40	PRO
15	L	148	SER
16	M	68	GLU
21	R	58	MET
26	W	70	ILE
28	Y	20	LEU

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
4	A	192	VAL
5	B	2	GLN
12	I	65	ASN
22	S	90	PRO
16	M	109	PRO
24	U	40	PRO
4	A	37	VAL
5	B	5	ARG
15	L	88	VAL

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 X-ray entries followed by that with respect to entries of similar resolution.

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
4	A	179/181 (99%)	166 (93%)	13 (7%)	13	38
5	B	282/282 (100%)	262 (93%)	20 (7%)	13	40
6	C	193/193 (100%)	178 (92%)	15 (8%)	11	35
7	D	117/147 (80%)	108 (92%)	9 (8%)	12	35
8	E	152/155 (98%)	146 (96%)	6 (4%)	28	63
9	F	92/92 (100%)	92 (100%)	0	100	100
10	G	27/283 (10%)	27 (100%)	0	100	100
11	H	122/122 (100%)	108 (88%)	14 (12%)	5	18
12	I	118/121 (98%)	110 (93%)	8 (7%)	14	42
13	J	106/106 (100%)	103 (97%)	3 (3%)	38	72
14	K	112/126 (89%)	109 (97%)	3 (3%)	39	73
15	L	166/166 (100%)	159 (96%)	7 (4%)	26	60
16	M	149/149 (100%)	140 (94%)	9 (6%)	17	47
17	N	93/93 (100%)	87 (94%)	6 (6%)	15	44
18	O	113/116 (97%)	110 (97%)	3 (3%)	39	73
19	P	79/79 (100%)	75 (95%)	4 (5%)	21	53
20	Q	117/121 (97%)	113 (97%)	4 (3%)	32	66

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
21	R	71/73 (97%)	68 (96%)	3 (4%)	26	60
22	S	105/105 (100%)	101 (96%)	4 (4%)	29	64
23	T	44/52 (85%)	43 (98%)	1 (2%)	44	76
24	U	51/56 (91%)	50 (98%)	1 (2%)	48	78
25	V	130/130 (100%)	121 (93%)	9 (7%)	14	41
26	W	66/73 (90%)	61 (92%)	5 (8%)	12	36
27	X	120/195 (62%)	112 (93%)	8 (7%)	15	43
28	Y	56/56 (100%)	54 (96%)	2 (4%)	31	65
29	Z	46/46 (100%)	46 (100%)	0	100	100
30	1	42/44 (96%)	41 (98%)	1 (2%)	43	75
31	2	79/79 (100%)	76 (96%)	3 (4%)	29	64
All	All	3027/3441 (88%)	2866 (95%)	161 (5%)	20	52

All (161) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
4	A	3	ARG
4	A	13	THR
4	A	33	GLU
4	A	36	ASP
4	A	55	VAL
4	A	68	ILE
4	A	69	LEU
4	A	94	LEU
4	A	120	ARG
4	A	153	ARG
4	A	175	LYS
4	A	179	MET
4	A	217	ARG
5	B	7	ARG
5	B	11	LEU
5	B	27	ASN
5	B	33	ASP
5	B	63	GLU
5	B	84	LEU
5	B	97	LEU
5	B	98	THR
5	B	103	ASP

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
5	B	113	LEU
5	B	190	MET
5	B	195	ARG
5	B	245	SER
5	B	251	VAL
5	B	254	GLN
5	B	256	GLN
5	B	257	THR
5	B	304	PRO
5	B	307	ARG
5	B	312	ARG
6	C	2	GLN
6	C	67	GLN
6	C	78	ARG
6	C	94	THR
6	C	101	ASP
6	C	115	LEU
6	C	136	VAL
6	C	187	ARG
6	C	214	THR
6	C	222	ASP
6	C	223	LEU
6	C	234	VAL
6	C	236	THR
6	C	240	LEU
6	C	243	VAL
7	D	24	HIS
7	D	50	VAL
7	D	95	THR
7	D	99	ASP
7	D	131	THR
7	D	133	ASN
7	D	136	ARG
7	D	137	PRO
7	D	149	ARG
8	E	7	ILE
8	E	36	PRO
8	E	86	VAL
8	E	102	VAL
8	E	116	THR
8	E	154	ILE
11	H	1	LYS

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
11	H	18	GLU
11	H	59	ASN
11	H	72	VAL
11	H	73	GLN
11	H	82	LYS
11	H	85	ILE
11	H	86	ARG
11	H	93	ILE
11	H	94	ARG
11	H	118	PRO
11	H	142	VAL
11	H	150	LYS
11	H	166	ASN
12	I	46	ILE
12	I	47	THR
12	I	52	GLN
12	I	74	ARG
12	I	93	ARG
12	I	120	SER
12	I	127	ILE
12	I	131	THR
13	J	7	ASP
13	J	10	GLN
13	J	98	VAL
14	K	35	ARG
14	K	117	GLU
14	K	140	VAL
15	L	38	VAL
15	L	46	LEU
15	L	87	MET
15	L	93	ARG
15	L	130	GLU
15	L	159	THR
15	L	164	THR
16	M	12	ARG
16	M	26	LEU
16	M	43	VAL
16	M	47	LEU
16	M	50	LEU
16	M	127	LEU
16	M	128	ASP
16	M	152	GLU

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
16	M	163	PHE
17	N	3	THR
17	N	43	VAL
17	N	97	SER
17	N	98	LEU
17	N	109	SER
17	N	111	VAL
18	O	52	LYS
18	O	94	TRP
18	O	98	ILE
19	P	11	ARG
19	P	16	ASN
19	P	57	ASP
19	P	95	GLU
20	Q	13	THR
20	Q	39	THR
20	Q	82	GLU
20	Q	132	ARG
21	R	10	VAL
21	R	28	VAL
21	R	57	THR
22	S	23	VAL
22	S	39	ASN
22	S	48	VAL
22	S	96	VAL
23	T	28	THR
24	U	43	PRO
25	V	4	LEU
25	V	35	VAL
25	V	48	VAL
25	V	52	VAL
25	V	73	LEU
25	V	122	ARG
25	V	142	ASP
25	V	146	ILE
25	V	154	ARG
26	W	15	ARG
26	W	27	ASP
26	W	49	ARG
26	W	72	VAL
26	W	79	GLU
27	X	115	ARG

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
27	X	154	ARG
27	X	163	THR
27	X	172	THR
27	X	189	ASN
27	X	200	THR
27	X	203	VAL
27	X	235	GLU
28	Y	11	THR
28	Y	64	ILE
30	1	18	ASN
31	2	42	ARG
31	2	56	PRO
31	2	65	THR

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (104) such sidechains are listed below:

Mol	Chain	Res	Type
4	A	29	HIS
4	A	47	HIS
4	A	127	GLN
4	A	199	HIS
4	A	218	ASN
5	B	27	ASN
5	B	145	HIS
5	B	221	GLN
5	B	238	ASN
5	B	260	HIS
5	B	332	ASN
6	C	2	GLN
6	C	39	GLN
6	C	67	GLN
6	C	129	HIS
6	C	163	HIS
7	D	103	ASN
7	D	133	ASN
8	E	106	ASN
8	E	143	GLN
9	F	80	GLN
10	G	17	GLN
10	G	64	ASN
11	H	35	ASN
11	H	45	GLN

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
11	H	55	GLN
11	H	58	HIS
11	H	59	ASN
11	H	74	ASN
11	H	80	ASN
11	H	91	HIS
11	H	129	ASN
11	H	130	HIS
11	H	137	ASN
11	H	166	ASN
12	I	25	GLN
12	I	52	GLN
12	I	107	ASN
13	J	10	GLN
13	J	67	GLN
14	K	18	HIS
14	K	41	HIS
14	K	43	HIS
14	K	58	GLN
14	K	116	HIS
15	L	26	HIS
15	L	58	GLN
15	L	78	ASN
15	L	176	GLN
16	M	107	ASN
17	N	53	GLN
17	N	100	GLN
18	O	50	GLN
18	O	66	GLN
18	O	73	HIS
18	O	118	GLN
19	P	16	ASN
19	P	40	HIS
19	P	59	GLN
19	P	67	GLN
20	Q	61	GLN
20	Q	98	ASN
20	Q	113	HIS
20	Q	117	HIS
20	Q	122	GLN
20	Q	123	GLN
20	Q	140	GLN

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
21	R	51	GLN
21	R	53	ASN
21	R	55	GLN
22	S	39	ASN
22	S	73	HIS
22	S	95	ASN
23	T	39	ASN
23	T	48	ASN
24	U	4	HIS
24	U	6	GLN
24	U	60	GLN
25	V	6	GLN
25	V	14	HIS
25	V	27	HIS
25	V	28	HIS
25	V	87	HIS
25	V	110	GLN
25	V	119	HIS
25	V	125	HIS
25	V	141	HIS
26	W	23	HIS
27	X	121	HIS
27	X	133	HIS
27	X	134	HIS
27	X	149	GLN
27	X	189	ASN
28	Y	33	HIS
28	Y	70	GLN
29	Z	8	GLN
29	Z	16	HIS
29	Z	28	HIS
30	1	16	ASN
30	1	18	ASN
30	1	45	ASN
31	2	30	GLN
31	2	39	GLN
31	2	48	ASN

5.3.3 RNA ⓘ

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	0	2746/2922 (93%)	277 (10%)	55 (2%)
2	9	121/122 (99%)	20 (16%)	7 (5%)
3	3	2/3 (66%)	1 (50%)	0
3	4	2/3 (66%)	0	0
3	5	2/3 (66%)	1 (50%)	0
All	All	2873/3053 (94%)	299 (10%)	62 (2%)

All (299) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	0	11	A
1	0	31	C
1	0	60	A
1	0	67	A
1	0	69	A
1	0	70	A
1	0	71	G
1	0	87	C
1	0	88	G
1	0	114	A
1	0	115	U
1	0	120	A
1	0	130	C
1	0	139	C
1	0	141	C
1	0	151	A
1	0	166	A
1	0	169	A
1	0	170	U
1	0	186	A
1	0	191	A
1	0	192	A
1	0	198	A
1	0	213	G
1	0	214	U
1	0	219	G
1	0	237	G
1	0	271	C
1	0	272	A
1	0	273	G
1	0	283	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	0	284	C
1	0	285	A
1	0	308	U
1	0	309	C
1	0	317	A
1	0	336	G
1	0	337	A
1	0	338	C
1	0	339	A
1	0	345	G
1	0	358	G
1	0	381	G
1	0	397	A
1	0	417	G
1	0	461	C
1	0	487	G
1	0	498	A
1	0	509	A
1	0	510	U
1	0	511	A
1	0	514	G
1	0	537	G
1	0	538	C
1	0	539	G
1	0	542	A
1	0	545	G
1	0	553	G
1	0	559	U
1	0	588	G
1	0	604	G
1	0	605	C
1	0	620	A
1	0	632	A
1	0	644	G
1	0	645	U
1	0	660	A
1	0	688	A
1	0	701	U
1	0	717	C
1	0	759	C
1	0	777	U
1	0	809	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	0	821	U
1	0	835	U
1	0	840	U
1	0	857	A
1	0	858	U
1	0	868	G
1	0	869	G
1	0	871	G
1	0	872	U
1	0	875	A
1	0	877	G
1	0	878	G
1	0	882	A
1	0	884	C
1	0	885	G
1	0	898	G
1	0	905	C
1	0	921	G
1	0	923	A
1	0	953	G
1	0	960	G
1	0	961	A
1	0	1006	A
1	0	1008	C
1	0	1029	U
1	0	1030	U
1	0	1031	G
1	0	1045	G
1	0	1059	G
1	0	1060	C
1	0	1072	G
1	0	1081	A
1	0	1088	A
1	0	1109	U
1	0	1110	G
1	0	1119	G
1	0	1130	U
1	0	1137	G
1	0	1138	G
1	0	1151	G
1	0	1162	G
1	0	1164	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	0	1165	G
1	0	1166	A
1	0	1171	A
1	0	1174	A
1	0	1175	G
1	0	1177	A
1	0	1185	U
1	0	1192	A
1	0	1193	A
1	0	1206	U
1	0	1216	G
1	0	1226	G
1	0	1232	A
1	0	1233	A
1	0	1235	G
1	0	1236	A
1	0	1237	U
1	0	1238	C
1	0	1239	G
1	0	1242	A
1	0	1279	U
1	0	1289	C
1	0	1342	C
1	0	1353	C
1	0	1360	C
1	0	1377	C
1	0	1380	U
1	0	1381	A
1	0	1407	A
1	0	1409	G
1	0	1438	G
1	0	1451	C
1	0	1474	C
1	0	1485	A
1	0	1505	U
1	0	1506	U
1	0	1524	U
1	0	1525	G
1	0	1526	A
1	0	1528	A
1	0	1563	G
1	0	1564	C

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	0	1580	A
1	0	1592	G
1	0	1603	A
1	0	1604	G
1	0	1625	U
1	0	1626	A
1	0	1633	C
1	0	1634	G
1	0	1656	A
1	0	1667	A
1	0	1682	A
1	0	1684	A
1	0	1685	A
1	0	1692	C
1	0	1701	A
1	0	1722	U
1	0	1723	G
1	0	1725	C
1	0	1730	G
1	0	1731	C
1	0	1737	A
1	0	1752	G
1	0	1778	A
1	0	1779	A
1	0	1798	C
1	0	1819	G
1	0	1820	G
1	0	1829	A
1	0	1856	C
1	0	1857	A
1	0	1904	A
1	0	1919	A
1	0	1941	A
1	0	1942	A
1	0	1971	G
1	0	1973	A
1	0	1974	G
1	0	1978	A
1	0	1980	U
1	0	1996	U
1	0	2005	G
1	0	2008	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	0	2011	A
1	0	2012	U
1	0	2013	G
1	0	2033	G
1	0	2034	U
1	0	2064	U
1	0	2072	G
1	0	2073	G
1	0	2074	A
1	0	2096	A
1	0	2101	A
1	0	2102	G
1	0	2110	G
1	0	2238	A
1	0	2243	C
1	0	2258	A
1	0	2271	G
1	0	2272	G
1	0	2317	C
1	0	2320	U
1	0	2321	A
1	0	2329	C
1	0	2346	C
1	0	2354	A
1	0	2361	A
1	0	2369	A
1	0	2379	G
1	0	2422	U
1	0	2462	G
1	0	2467	A
1	0	2476	C
1	0	2480	G
1	0	2483	A
1	0	2503	A
1	0	2504	A
1	0	2511	A
1	0	2526	C
1	0	2527	U
1	0	2532	A
1	0	2533	C
1	0	2537	G
1	0	2541	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	0	2553	A
1	0	2564	G
1	0	2589	U
1	0	2601	A
1	0	2602	G
1	0	2608	C
1	0	2613	G
1	0	2634	G
1	0	2638	G
1	0	2649	A
1	0	2664	A
1	0	2681	A
1	0	2682	C
1	0	2726	U
1	0	2747	C
1	0	2748	G
1	0	2749	U
1	0	2750	G
1	0	2762	C
1	0	2768	A
1	0	2786	G
1	0	2792	A
1	0	2800	A
1	0	2811	A
1	0	2812	A
1	0	2825	C
1	0	2836	G
1	0	2837	U
1	0	2840	A
1	0	2850	C
1	0	2851	G
1	0	2876	G
1	0	2890	A
1	0	2896	A
1	0	2903	C
1	0	2914	A
2	9	3002	U
2	9	3003	A
2	9	3007	G
2	9	3014	G
2	9	3022	G
2	9	3023	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
2	9	3024	U
2	9	3025	G
2	9	3026	C
2	9	3041	C
2	9	3043	G
2	9	3044	A
2	9	3052	A
2	9	3055	U
2	9	3056	A
2	9	3057	A
2	9	3066	G
2	9	3077	A
2	9	3114	G
2	9	3122	C
3	3	76	A
3	5	76	A

All (62) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
1	0	10	U
1	0	59	A
1	0	87	C
1	0	129	A
1	0	169	A
1	0	213	G
1	0	284	C
1	0	338	C
1	0	357	A
1	0	509	A
1	0	603	A
1	0	604	G
1	0	644	G
1	0	834	G
1	0	857	A
1	0	871	G
1	0	877	G
1	0	898	G
1	0	1030	U
1	0	1080	C
1	0	1137	G
1	0	1164	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	0	1192	A
1	0	1232	A
1	0	1235	G
1	0	1237	U
1	0	1352	A
1	0	1377	C
1	0	1380	U
1	0	1450	C
1	0	1524	U
1	0	1563	G
1	0	1603	A
1	0	1667	A
1	0	1730	G
1	0	1752	G
1	0	1856	C
1	0	1941	A
1	0	1979	G
1	0	2011	A
1	0	2102	G
1	0	2313	C
1	0	2320	U
1	0	2369	A
1	0	2379	G
1	0	2467	A
1	0	2503	A
1	0	2526	C
1	0	2536	C
1	0	2748	G
1	0	2749	U
1	0	2791	U
1	0	2836	G
1	0	2837	U
1	0	2850	C
2	9	3002	U
2	9	3023	U
2	9	3024	U
2	9	3055	U
2	9	3065	A
2	9	3103	A
2	9	3113	C

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

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

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

Of 231 ligands modelled in this entry, 231 are monoatomic - leaving 0 for Mogul analysis.

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

No monomer is involved in short contacts.

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 Fit of model and data ⓘ

6.1 Protein, DNA and RNA chains ⓘ

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

Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
1	0	2754/2922 (94%)	-0.27	40 (1%) 72 64	23, 53, 103, 166	0
2	9	122/122 (100%)	0.11	5 (4%) 41 33	36, 67, 104, 164	0
3	3	3/3 (100%)	1.03	1 (33%) 1 1	68, 68, 83, 120	0
3	4	3/3 (100%)	1.77	1 (33%) 1 1	88, 88, 89, 103	0
3	5	3/3 (100%)	0.14	0 100 100	57, 57, 58, 64	0
4	A	237/239 (99%)	0.33	7 (2%) 52 43	33, 66, 100, 126	0
5	B	337/337 (100%)	0.21	1 (0%) 90 87	27, 60, 86, 95	0
6	C	246/246 (100%)	0.16	1 (0%) 88 85	27, 57, 81, 88	0
7	D	140/176 (79%)	1.64	48 (34%) 1 1	65, 111, 129, 137	0
8	E	172/177 (97%)	0.43	7 (4%) 41 33	43, 69, 95, 101	0
9	F	119/119 (100%)	0.81	7 (5%) 28 22	65, 88, 112, 117	0
10	G	29/348 (8%)	1.19	7 (24%) 2 2	72, 95, 107, 111	0
11	H	156/167 (93%)	0.39	12 (7%) 19 16	31, 56, 83, 92	0
12	I	142/145 (97%)	0.06	1 (0%) 84 79	37, 50, 75, 87	0
13	J	132/132 (100%)	0.30	1 (0%) 82 77	41, 60, 83, 90	0
14	K	145/164 (88%)	0.70	12 (8%) 17 14	31, 77, 121, 127	0
15	L	194/194 (100%)	0.30	7 (3%) 46 38	37, 56, 80, 88	0
16	M	186/186 (100%)	1.03	30 (16%) 4 4	43, 78, 123, 135	0
17	N	115/115 (100%)	0.34	1 (0%) 81 75	47, 65, 83, 86	0
18	O	143/148 (96%)	0.58	3 (2%) 63 54	39, 65, 85, 94	0
19	P	95/95 (100%)	0.01	0 100 100	35, 49, 67, 86	0
20	Q	150/154 (97%)	-0.20	0 100 100	35, 47, 69, 80	0
21	R	81/84 (96%)	0.46	0 100 100	56, 80, 96, 103	0
22	S	119/119 (100%)	0.56	5 (4%) 40 32	52, 71, 101, 120	0

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
23	T	53/66 (80%)	0.29	1 (1%) 66 58	48, 62, 79, 91	0
24	U	65/70 (92%)	1.03	10 (15%) 5 4	67, 93, 127, 133	0
25	V	154/154 (100%)	-0.10	0 100 100	32, 49, 70, 80	0
26	W	82/91 (90%)	0.39	2 (2%) 59 50	46, 63, 84, 101	0
27	X	142/240 (59%)	0.01	2 (1%) 73 65	28, 52, 77, 93	0
28	Y	73/73 (100%)	0.85	8 (10%) 10 9	55, 72, 90, 96	0
29	Z	56/56 (100%)	-0.41	0 100 100	29, 42, 50, 53	0
30	1	46/48 (95%)	1.42	13 (28%) 1 1	42, 77, 136, 138	0
31	2	92/92 (100%)	0.55	2 (2%) 62 53	44, 67, 80, 91	0
All	All	6586/7288 (90%)	0.12	235 (3%) 46 38	23, 60, 108, 166	0

All (235) RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
24	U	1	THR	7.2
7	D	88	LEU	6.7
2	9	3001	U	5.6
30	1	42	TRP	5.3
7	D	10	PHE	5.1
5	B	1	PRO	4.9
30	1	48	ASP	4.8
14	K	83	GLU	4.7
30	1	37	HIS	4.5
16	M	148	ALA	4.5
2	9	3025	G	4.4
8	E	45	ASP	4.4
30	1	38	LYS	4.4
14	K	82	ALA	4.3
30	1	46	ASP	4.3
16	M	168	LEU	4.2
16	M	165	ALA	4.2
22	S	119	ALA	4.1
7	D	90	LEU	4.1
16	M	147	ILE	4.0
30	1	49	GLU	4.0
7	D	106	PHE	4.0
15	L	152	ARG	4.0
1	0	138	U	3.9
7	D	107	GLY	3.8

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	RSRZ
11	H	40	PRO	3.8
30	1	47	THR	3.8
30	1	44	ARG	3.8
16	M	162	ASP	3.7
10	G	27	ILE	3.6
7	D	174	VAL	3.6
24	U	2	VAL	3.5
1	0	1171	A	3.5
1	0	2637	A	3.5
7	D	104	PHE	3.4
1	0	1176	C	3.4
7	D	19	GLU	3.4
15	L	87	MET	3.4
7	D	17	ARG	3.4
22	S	118	SER	3.4
1	0	1175	G	3.3
2	9	3002	U	3.2
1	0	1199	A	3.2
7	D	55	LYS	3.2
2	9	3024	U	3.2
30	1	43	ARG	3.2
14	K	104	ASP	3.2
16	M	166	ALA	3.2
7	D	21	VAL	3.2
2	9	3023	U	3.1
9	F	106	THR	3.1
16	M	158	LEU	3.1
7	D	63	ILE	3.1
24	U	44	GLY	3.1
1	0	10	U	3.1
11	H	167	ALA	3.0
16	M	186	LEU	3.0
14	K	102	ASP	3.0
7	D	130	VAL	3.0
1	0	735	C	3.0
30	1	36	ASN	3.0
7	D	134	LEU	3.0
7	D	89	PRO	3.0
16	M	1	ALA	3.0
16	M	159	TYR	3.0
7	D	75	LEU	3.0
16	M	154	LEU	3.0

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	RSRZ
11	H	32	ASP	3.0
31	2	41	GLU	3.0
30	1	35	ARG	2.9
1	0	2254	G	2.9
24	U	36	ALA	2.9
3	4	76	A	2.9
7	D	171	ASP	2.9
8	E	156	ASP	2.9
16	M	80	SER	2.9
1	0	1177	A	2.9
30	1	41	HIS	2.8
9	F	44	SER	2.8
7	D	35	ALA	2.8
7	D	87	ALA	2.8
26	W	65	ASN	2.8
28	Y	11	THR	2.8
1	0	716	G	2.8
4	A	37	VAL	2.8
7	D	18	ILE	2.7
16	M	83	LEU	2.7
16	M	169	PRO	2.7
1	0	1184	C	2.7
1	0	960	G	2.7
16	M	149	GLU	2.7
27	X	235	GLU	2.7
28	Y	44	PHE	2.7
31	2	8	ASN	2.7
1	0	1524	U	2.7
7	D	20	LYS	2.6
7	D	61	PHE	2.6
16	M	143	ARG	2.6
1	0	2748	G	2.6
15	L	16	LYS	2.6
24	U	51	LYS	2.6
7	D	22	VAL	2.6
7	D	166	ILE	2.6
10	G	63	ARG	2.6
7	D	172	VAL	2.6
14	K	81	VAL	2.6
10	G	26	MET	2.6
1	0	2250	G	2.5
1	0	734	U	2.5

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	RSRZ
1	0	1185	U	2.5
24	U	40	PRO	2.5
24	U	43	PRO	2.5
7	D	84	LEU	2.5
10	G	71	LEU	2.5
10	G	12	ILE	2.5
28	Y	22	ILE	2.5
4	A	36	ASP	2.5
4	A	65	ARG	2.5
1	0	2345	A	2.5
4	A	31	LYS	2.5
4	A	237	GLY	2.5
16	M	145	ALA	2.5
28	Y	10	ARG	2.5
15	L	89	ASN	2.5
7	D	102	GLY	2.4
15	L	165	SER	2.4
23	T	51	TRP	2.4
7	D	154	LYS	2.4
1	0	1204	C	2.4
1	0	288	A	2.4
7	D	44	ILE	2.4
22	S	116	ASP	2.4
18	O	56	GLY	2.4
28	Y	48	LYS	2.4
16	M	179	LEU	2.4
22	S	112	LEU	2.4
7	D	59	GLY	2.4
16	M	167	ASP	2.4
14	K	99	GLU	2.4
16	M	160	SER	2.4
11	H	140	PRO	2.4
11	H	110	GLY	2.4
15	L	63	VAL	2.4
11	H	138	PRO	2.4
11	H	59	ASN	2.3
27	X	216	ARG	2.3
28	Y	35	LYS	2.3
11	H	39	GLY	2.3
8	E	53	GLU	2.3
1	0	1173	A	2.3
22	S	1	SER	2.3

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	RSRZ
16	M	104	ILE	2.3
12	I	70	PHE	2.3
8	E	100	ASP	2.3
16	M	138	ASP	2.3
3	3	74	C	2.3
1	0	1172	G	2.3
6	C	244	ALA	2.3
11	H	164	ALA	2.3
9	F	17	LEU	2.3
7	D	150	SER	2.3
1	0	1161	A	2.2
1	0	2344	G	2.2
28	Y	82	ALA	2.2
10	G	23	ILE	2.2
7	D	11	HIS	2.2
7	D	62	ASP	2.2
4	A	32	VAL	2.2
7	D	57	THR	2.2
7	D	58	VAL	2.2
7	D	95	THR	2.2
7	D	132	VAL	2.2
17	N	31	GLU	2.2
1	0	1964	U	2.2
14	K	150	GLN	2.2
1	0	1192	A	2.2
7	D	86	THR	2.2
7	D	53	LYS	2.2
7	D	69	ILE	2.2
7	D	76	ARG	2.2
1	0	1951	G	2.2
1	0	1193	A	2.2
16	M	157	PRO	2.2
1	0	999	C	2.2
1	0	2247	C	2.2
11	H	135	TRP	2.2
7	D	72	LYS	2.2
7	D	83	PHE	2.2
14	K	60	GLU	2.2
8	E	11	VAL	2.1
9	F	98	VAL	2.1
16	M	79	PRO	2.1
8	E	10	ASP	2.1

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	RSRZ
11	H	109	ASP	2.1
1	0	1162	G	2.1
1	0	1188	A	2.1
7	D	66	GLY	2.1
7	D	128	LEU	2.1
14	K	89	PHE	2.1
30	1	39	ARG	2.1
16	M	140	GLN	2.1
10	G	24	VAL	2.1
26	W	85	VAL	2.1
13	J	63	GLU	2.1
1	0	130	C	2.1
1	0	1186	C	2.1
1	0	2769	C	2.1
7	D	54	ALA	2.1
24	U	52	ALA	2.1
7	D	139	TYR	2.1
14	K	105	TYR	2.1
16	M	150	TYR	2.1
1	0	1167	G	2.1
16	M	127	LEU	2.1
14	K	125	PHE	2.1
24	U	45	ARG	2.1
4	A	33	GLU	2.1
18	O	116	SER	2.0
18	O	57	ASN	2.0
1	0	1160	G	2.0
24	U	37	GLY	2.0
9	F	26	THR	2.0
9	F	105	ALA	2.0
7	D	157	LEU	2.0
9	F	47	LEU	2.0
16	M	76	GLY	2.0
16	M	182	GLY	2.0
8	E	13	ALA	2.0
11	H	41	THR	2.0
15	L	192	ALA	2.0
28	Y	50	ALA	2.0
1	0	2249	G	2.0
1	0	2256	G	2.0
16	M	152	GLU	2.0
14	K	106	VAL	2.0

6.2 Non-standard residues in protein, DNA, RNA chains [i](#)

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

6.3 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

6.4 Ligands [i](#)

In the following table, the Atoms column lists the number of modelled atoms in the group and the number defined in the chemical component dictionary. The B-factors column lists the minimum, median, 95th percentile and maximum values of B factors of atoms in the group. The column labelled 'Q< 0.9' lists the number of atoms with occupancy less than 0.9.

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
34	NA	0	8329	1/1	0.50	0.41	135,135,135,135	0
34	NA	9	8351	1/1	0.53	0.24	89,89,89,89	0
34	NA	R	8312	1/1	0.61	0.17	60,60,60,60	0
32	MG	0	8092	1/1	0.77	0.17	109,109,109,109	0
34	NA	0	8384	1/1	0.77	0.43	132,132,132,132	0
34	NA	Q	8386	1/1	0.78	0.48	84,84,84,84	0
32	MG	9	8095	1/1	0.79	0.12	73,73,73,73	0
34	NA	0	8341	1/1	0.81	0.07	42,42,42,42	0
34	NA	0	8372	1/1	0.81	0.43	71,71,71,71	0
35	CL	N	8508	1/1	0.82	0.22	101,101,101,101	0
34	NA	0	8381	1/1	0.84	0.09	47,47,47,47	0
34	NA	0	8382	1/1	0.84	0.15	64,64,64,64	0
34	NA	9	8383	1/1	0.84	0.10	62,62,62,62	0
34	NA	0	8326	1/1	0.85	0.17	52,52,52,52	0
34	NA	0	8369	1/1	0.85	0.22	68,68,68,68	0
34	NA	Q	8337	1/1	0.85	0.08	43,43,43,43	0
34	NA	0	8302	1/1	0.86	0.07	40,40,40,40	0
34	NA	0	8324	1/1	0.86	0.17	62,62,62,62	0
34	NA	0	8350	1/1	0.86	0.23	61,61,61,61	0
34	NA	0	8352	1/1	0.86	0.30	61,61,61,61	0
35	CL	0	8505	1/1	0.86	0.18	87,87,87,87	0
32	MG	0	8049	1/1	0.86	0.10	64,64,64,64	0
34	NA	0	8365	1/1	0.87	0.15	58,58,58,58	0
34	NA	0	8368	1/1	0.87	0.12	51,51,51,51	0
32	MG	0	8104	1/1	0.87	0.09	58,58,58,58	0
32	MG	0	8087	1/1	0.87	0.08	83,83,83,83	0
32	MG	0	8011	1/1	0.87	0.25	1,1,1,1	0

Continued on next page...

Continued from previous page...

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
34	NA	0	8317	1/1	0.87	0.09	75,75,75,75	0
32	MG	0	8101	1/1	0.87	0.15	68,68,68,68	0
35	CL	0	8511	1/1	0.88	0.18	90,90,90,90	0
32	MG	J	8069	1/1	0.88	0.11	114,114,114,114	0
34	NA	0	8363	1/1	0.89	0.16	47,47,47,47	0
35	CL	0	8503	1/1	0.89	0.15	67,67,67,67	0
34	NA	0	8316	1/1	0.89	0.15	39,39,39,39	0
32	MG	0	8076	1/1	0.89	0.09	83,83,83,83	0
34	NA	0	8359	1/1	0.89	0.28	66,66,66,66	0
34	NA	0	8362	1/1	0.90	0.23	73,73,73,73	0
34	NA	0	8371	1/1	0.90	0.18	58,58,58,58	0
32	MG	0	8103	1/1	0.90	0.18	88,88,88,88	0
34	NA	0	8360	1/1	0.91	0.24	47,47,47,47	0
32	MG	0	8046	1/1	0.91	0.07	61,61,61,61	0
34	NA	0	8303	1/1	0.91	0.19	54,54,54,54	0
34	NA	0	8364	1/1	0.91	0.14	51,51,51,51	0
34	NA	0	8307	1/1	0.91	0.08	40,40,40,40	0
34	NA	0	8357	1/1	0.91	0.11	63,63,63,63	0
35	CL	M	8507	1/1	0.91	0.08	66,66,66,66	0
32	MG	0	8067	1/1	0.91	0.07	78,78,78,78	0
34	NA	P	8348	1/1	0.92	0.06	47,47,47,47	0
34	NA	0	8308	1/1	0.92	0.09	54,54,54,54	0
34	NA	0	8373	1/1	0.92	0.10	53,53,53,53	0
34	NA	0	8325	1/1	0.92	0.16	45,45,45,45	0
32	MG	0	8050	1/1	0.92	0.07	50,50,50,50	0
32	MG	0	8099	1/1	0.92	0.13	73,73,73,73	0
34	NA	0	8332	1/1	0.92	0.07	30,30,30,30	0
35	CL	0	8515	1/1	0.92	0.13	96,96,96,96	0
35	CL	0	8517	1/1	0.92	0.09	64,64,64,64	0
35	CL	K	8510	1/1	0.92	0.13	71,71,71,71	0
34	NA	0	8322	1/1	0.92	0.28	65,65,65,65	0
34	NA	K	8380	1/1	0.92	0.16	77,77,77,77	0
34	NA	0	8356	1/1	0.93	0.23	77,77,77,77	0
34	NA	0	8374	1/1	0.93	0.24	74,74,74,74	0
32	MG	0	8113	1/1	0.93	0.07	39,39,39,39	0
34	NA	0	8370	1/1	0.93	0.28	60,60,60,60	0
35	CL	I	8502	1/1	0.93	0.06	54,54,54,54	0
34	NA	0	8306	1/1	0.93	0.15	57,57,57,57	0
34	NA	0	8385	1/1	0.93	0.23	44,44,44,44	0
32	MG	0	8024	1/1	0.93	0.31	146,146,146,146	0
35	CL	X	8520	1/1	0.93	0.08	43,43,43,43	0
34	NA	0	8305	1/1	0.94	0.07	39,39,39,39	0

Continued on next page...

Continued from previous page...

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
32	MG	0	8082	1/1	0.94	0.10	77,77,77,77	0
35	CL	0	8522	1/1	0.94	0.23	68,68,68,68	0
35	CL	A	8509	1/1	0.94	0.14	78,78,78,78	0
35	CL	I	8501	1/1	0.94	0.08	64,64,64,64	0
32	MG	0	8044	1/1	0.94	0.11	54,54,54,54	0
32	MG	0	8047	1/1	0.94	0.08	48,48,48,48	0
34	NA	0	8333	1/1	0.94	0.07	34,34,34,34	0
34	NA	0	8339	1/1	0.94	0.08	34,34,34,34	0
34	NA	0	8379	1/1	0.94	0.08	34,34,34,34	0
34	NA	0	8310	1/1	0.95	0.14	20,20,20,20	0
34	NA	0	8314	1/1	0.95	0.11	26,26,26,26	0
32	MG	0	8051	1/1	0.95	0.07	86,86,86,86	0
32	MG	0	8084	1/1	0.95	0.13	101,101,101,101	0
34	NA	0	8358	1/1	0.95	0.18	102,102,102,102	0
34	NA	0	8320	1/1	0.95	0.07	44,44,44,44	0
34	NA	L	8347	1/1	0.95	0.06	38,38,38,38	0
32	MG	0	8085	1/1	0.95	0.07	82,82,82,82	0
34	NA	0	8361	1/1	0.95	0.14	47,47,47,47	0
32	MG	0	8057	1/1	0.95	0.08	51,51,51,51	0
32	MG	0	8090	1/1	0.95	0.11	65,65,65,65	0
32	MG	0	8091	1/1	0.95	0.07	73,73,73,73	0
34	NA	0	8328	1/1	0.95	0.10	51,51,51,51	0
34	NA	0	8367	1/1	0.95	0.08	54,54,54,54	0
35	CL	0	8513	1/1	0.95	0.07	62,62,62,62	0
35	CL	0	8514	1/1	0.95	0.12	80,80,80,80	0
32	MG	0	8059	1/1	0.95	0.08	84,84,84,84	0
34	NA	0	8330	1/1	0.95	0.06	45,45,45,45	0
32	MG	0	8097	1/1	0.95	0.17	37,37,37,37	0
32	MG	0	8041	1/1	0.95	0.08	53,53,53,53	0
35	CL	B	8519	1/1	0.95	0.17	58,58,58,58	0
34	NA	0	8334	1/1	0.95	0.09	36,36,36,36	0
32	MG	0	8100	1/1	0.95	0.10	74,74,74,74	0
32	MG	0	8045	1/1	0.95	0.07	67,67,67,67	0
34	NA	0	8378	1/1	0.95	0.38	45,45,45,45	0
34	NA	0	8342	1/1	0.95	0.07	36,36,36,36	0
34	NA	0	8343	1/1	0.95	0.05	33,33,33,33	0
34	NA	0	8335	1/1	0.96	0.17	54,54,54,54	0
34	NA	0	8375	1/1	0.96	0.13	55,55,55,55	0
34	NA	0	8323	1/1	0.96	0.12	46,46,46,46	0
34	NA	0	8340	1/1	0.96	0.10	36,36,36,36	0
32	MG	0	8116	1/1	0.96	0.04	40,40,40,40	0
34	NA	0	8311	1/1	0.96	0.10	44,44,44,44	0

Continued on next page...

Continued from previous page...

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
32	MG	0	8088	1/1	0.96	0.08	34,34,34,34	0
34	NA	0	8366	1/1	0.96	0.05	48,48,48,48	0
32	MG	A	8066	1/1	0.96	0.06	44,44,44,44	0
32	MG	A	8105	1/1	0.96	0.09	29,29,29,29	0
34	NA	A	8345	1/1	0.96	0.06	58,58,58,58	0
34	NA	C	8304	1/1	0.96	0.05	59,59,59,59	0
34	NA	0	8355	1/1	0.96	0.12	59,59,59,59	0
35	CL	I	8521	1/1	0.96	0.07	54,54,54,54	0
34	NA	0	8318	1/1	0.96	0.09	20,20,20,20	0
32	MG	0	8098	1/1	0.96	0.13	18,18,18,18	0
34	NA	0	8321	1/1	0.96	0.09	45,45,45,45	0
33	K	0	8201	1/1	0.96	0.17	74,74,74,74	0
35	CL	2	8504	1/1	0.96	0.08	77,77,77,77	0
36	CD	N	8405	1/1	0.96	0.05	90,90,90,90	0
32	MG	0	8108	1/1	0.97	0.07	61,61,61,61	0
34	NA	Q	8338	1/1	0.97	0.09	96,96,96,96	0
32	MG	0	8111	1/1	0.97	0.04	57,57,57,57	0
34	NA	0	8353	1/1	0.97	0.06	40,40,40,40	0
34	NA	0	8354	1/1	0.97	0.14	44,44,44,44	0
32	MG	0	8028	1/1	0.97	0.05	93,93,93,93	0
34	NA	0	8327	1/1	0.97	0.05	37,37,37,37	0
34	NA	0	8376	1/1	0.97	0.10	33,33,33,33	0
34	NA	0	8377	1/1	0.97	0.18	66,66,66,66	0
32	MG	0	8093	1/1	0.97	0.04	37,37,37,37	0
35	CL	0	8516	1/1	0.97	0.17	65,65,65,65	0
32	MG	0	8094	1/1	0.97	0.07	56,56,56,56	0
34	NA	0	8313	1/1	0.97	0.06	77,77,77,77	0
32	MG	0	8062	1/1	0.97	0.05	67,67,67,67	0
34	NA	0	8315	1/1	0.97	0.08	41,41,41,41	0
32	MG	0	8063	1/1	0.97	0.07	143,143,143,143	0
32	MG	0	8055	1/1	0.97	0.04	54,54,54,54	0
34	NA	0	8336	1/1	0.97	0.06	50,50,50,50	0
35	CL	J	8512	1/1	0.97	0.08	40,40,40,40	0
32	MG	0	8071	1/1	0.97	0.06	104,104,104,104	0
35	CL	L	8518	1/1	0.97	0.08	52,52,52,52	0
34	NA	0	8301	1/1	0.97	0.04	30,30,30,30	0
34	NA	H	8309	1/1	0.97	0.08	29,29,29,29	0
32	MG	0	8089	1/1	0.97	0.04	60,60,60,60	0
32	MG	0	8075	1/1	0.97	0.03	54,54,54,54	0
32	MG	0	8042	1/1	0.97	0.06	32,32,32,32	0
32	MG	0	8102	1/1	0.98	0.06	61,61,61,61	0
32	MG	0	8117	1/1	0.98	0.05	28,28,28,28	0

Continued on next page...

Continued from previous page...

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
32	MG	0	8048	1/1	0.98	0.04	35,35,35,35	0
34	NA	0	8344	1/1	0.98	0.04	21,21,21,21	0
34	NA	0	8349	1/1	0.98	0.10	51,51,51,51	0
32	MG	0	8074	1/1	0.98	0.04	29,29,29,29	0
32	MG	0	8106	1/1	0.98	0.04	58,58,58,58	0
32	MG	0	8054	1/1	0.98	0.09	17,17,17,17	0
32	MG	S	8073	1/1	0.98	0.04	64,64,64,64	0
32	MG	X	8109	1/1	0.98	0.04	34,34,34,34	0
34	NA	0	8331	1/1	0.98	0.16	50,50,50,50	0
32	MG	0	8110	1/1	0.98	0.05	34,34,34,34	0
33	K	0	8202	1/1	0.98	0.08	79,79,79,79	0
32	MG	0	8012	1/1	0.98	0.07	18,18,18,18	0
34	NA	0	8319	1/1	0.98	0.04	37,37,37,37	0
35	CL	Q	8506	1/1	0.98	0.06	52,52,52,52	0
32	MG	0	8112	1/1	0.98	0.08	52,52,52,52	0
32	MG	0	8039	1/1	0.98	0.04	39,39,39,39	0
32	MG	0	8115	1/1	0.98	0.06	27,27,27,27	0
32	MG	0	8008	1/1	0.99	0.04	39,39,39,39	0
32	MG	0	8025	1/1	0.99	0.03	24,24,24,24	0
32	MG	0	8027	1/1	0.99	0.05	69,69,69,69	0
32	MG	0	8009	1/1	0.99	0.03	53,53,53,53	0
32	MG	0	8052	1/1	0.99	0.08	42,42,42,42	0
32	MG	0	8053	1/1	0.99	0.04	47,47,47,47	0
34	NA	I	8346	1/1	0.99	0.02	39,39,39,39	0
32	MG	0	8029	1/1	0.99	0.03	38,38,38,38	0
32	MG	0	8096	1/1	0.99	0.04	39,39,39,39	0
32	MG	0	8030	1/1	0.99	0.04	25,25,25,25	0
32	MG	0	8032	1/1	0.99	0.06	26,26,26,26	0
32	MG	0	8058	1/1	0.99	0.04	63,63,63,63	0
32	MG	0	8034	1/1	0.99	0.06	16,16,16,16	0
32	MG	0	8060	1/1	0.99	0.03	34,34,34,34	0
32	MG	0	8061	1/1	0.99	0.02	23,23,23,23	0
32	MG	0	8035	1/1	0.99	0.04	50,50,50,50	0
32	MG	0	8037	1/1	0.99	0.03	32,32,32,32	0
32	MG	0	8064	1/1	0.99	0.12	22,22,22,22	0
32	MG	0	8107	1/1	0.99	0.03	52,52,52,52	0
32	MG	0	8004	1/1	0.99	0.03	44,44,44,44	0
32	MG	0	8068	1/1	0.99	0.03	52,52,52,52	0
32	MG	0	8070	1/1	0.99	0.07	41,41,41,41	0
32	MG	0	8040	1/1	0.99	0.03	53,53,53,53	0
32	MG	0	8072	1/1	0.99	0.07	38,38,38,38	0
32	MG	0	8006	1/1	0.99	0.03	46,46,46,46	0

Continued on next page...

Continued from previous page...

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
32	MG	0	8013	1/1	0.99	0.04	64,64,64,64	0
32	MG	0	8043	1/1	0.99	0.07	58,58,58,58	0
32	MG	0	8079	1/1	0.99	0.03	50,50,50,50	0
32	MG	0	8081	1/1	0.99	0.04	34,34,34,34	0
32	MG	0	8014	1/1	0.99	0.05	16,16,16,16	0
32	MG	B	8056	1/1	0.99	0.03	65,65,65,65	0
32	MG	0	8083	1/1	0.99	0.07	36,36,36,36	0
32	MG	0	8016	1/1	0.99	0.03	50,50,50,50	0
32	MG	0	8018	1/1	0.99	0.03	75,75,75,75	0
32	MG	2	8078	1/1	0.99	0.02	31,31,31,31	0
32	MG	0	8086	1/1	0.99	0.08	61,61,61,61	0
32	MG	0	8023	1/1	0.99	0.03	20,20,20,20	0
36	CD	2	8404	1/1	0.99	0.03	76,76,76,76	0
32	MG	A	8065	1/1	1.00	0.02	56,56,56,56	0
32	MG	0	8021	1/1	1.00	0.03	19,19,19,19	0
32	MG	0	8077	1/1	1.00	0.02	31,31,31,31	0
32	MG	0	8038	1/1	1.00	0.01	23,23,23,23	0
32	MG	0	8080	1/1	1.00	0.02	39,39,39,39	0
32	MG	0	8022	1/1	1.00	0.06	27,27,27,27	0
32	MG	0	8005	1/1	1.00	0.05	55,55,55,55	0
32	MG	0	8002	1/1	1.00	0.03	23,23,23,23	0
32	MG	0	8007	1/1	1.00	0.03	14,14,14,14	0
32	MG	0	8026	1/1	1.00	0.02	25,25,25,25	0
32	MG	0	8003	1/1	1.00	0.03	26,26,26,26	0
32	MG	0	8015	1/1	1.00	0.02	61,61,61,61	0
32	MG	0	8001	1/1	1.00	0.02	40,40,40,40	0
32	MG	0	8017	1/1	1.00	0.04	33,33,33,33	0
32	MG	0	8031	1/1	1.00	0.06	19,19,19,19	0
32	MG	0	8010	1/1	1.00	0.04	19,19,19,19	0
32	MG	0	8033	1/1	1.00	0.01	15,15,15,15	0
32	MG	0	8019	1/1	1.00	0.05	24,24,24,24	0
32	MG	0	8020	1/1	1.00	0.02	25,25,25,25	0
36	CD	T	8401	1/1	1.00	0.02	78,78,78,78	0
36	CD	Y	8403	1/1	1.00	0.02	82,82,82,82	0
36	CD	Z	8402	1/1	1.00	0.03	75,75,75,75	0
32	MG	0	8036	1/1	1.00	0.03	36,36,36,36	0

6.5 Other polymers

There are no such residues in this entry.