



Full wwPDB EM Validation Report ⓘ

Mar 27, 2026 – 06:40 PM UTC

PDB ID : 6QUL / pdb_00006qul
EMDB ID : EMD-4638
Title : Structure of a bacterial 50S ribosomal subunit in complex with the novel quinoxolidinone antibiotic cadazolid
Authors : Scaiola, A.; Leibundgut, M.; Boehringer, D.; Ritz, D.
Deposited on : 2019-02-27
Resolution : 3.00 Å(reported)

This is a Full wwPDB EM Validation Report for a publicly released PDB entry.

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

A user guide is available at

<https://www.wwpdb.org/validation/2017/EMValidationReportHelp>
with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

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

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

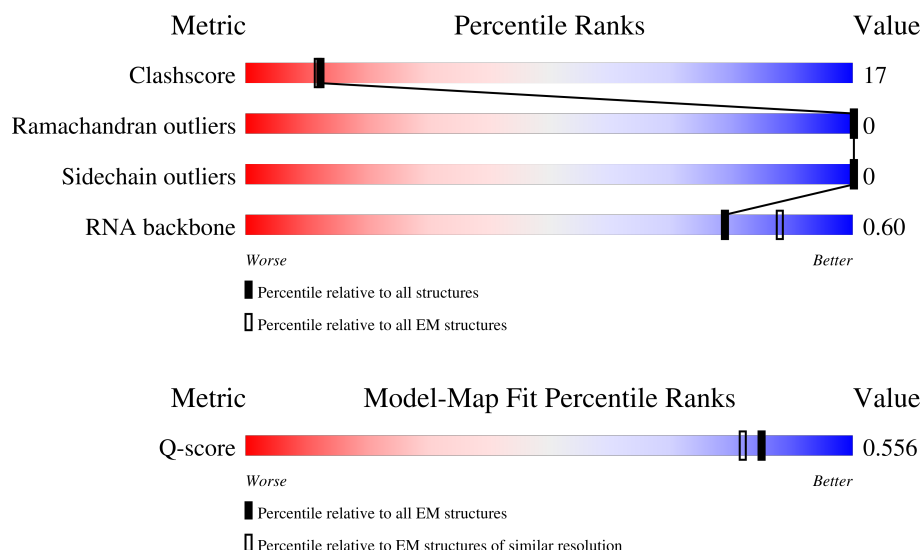
EMDB validation analysis : 0.0.1.dev132
Mogul : 2022.3.0, CSD as543be (2022)
MolProbity : 4-5-2 with Phenix2.0
Buster-report : wwPDB partial adaption of 1.1.7 (2018)
Percentile statistics : 20250101.v01 (using entries in the PDB archive January 1st 2025)
EM percentile statistics : 202505.v01 (Using data in the EMDB archive up until May 2025)
MapQ : 1.9.13
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.49

1 Overall quality at a glance

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

The reported resolution of this entry is 3.00 Å.

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



Metric	Whole archive (#Entries)	EM structures (#Entries)	Similar EM resolution (#Entries, resolution range(Å))
Clashscore	229148	23984	-
Ramachandran outliers	224038	23583	-
Sidechain outliers	223484	23102	-
RNA backbone	8273	3508	-
Q-score	-	25397	14081 (2.50 - 3.50)

The table below summarises the geometric issues observed across the polymeric chains and their fit to the map. The red, orange, yellow and green segments of the bar indicate the fraction of residues that contain outliers for ≥ 3 , 2, 1 and 0 types of geometric quality criteria respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions $\leq 5\%$. The upper red bar (where present) indicates the fraction of residues that have poor fit to the EM map (all-atom inclusion $< 40\%$). The numeric value is given above the bar.

Mol	Chain	Length	Quality of chain
1	1	4	
2	A	2795	
3	B	120	

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Mol	Chain	Length	Quality of chain
4	C	273	
5	D	209	
6	E	201	
7	F	179	
8	G	177	
9	H	149	
10	K	142	
11	L	123	
12	M	144	
13	N	136	
14	O	127	
15	P	117	
16	Q	115	
17	R	118	
18	S	103	
19	T	110	
20	U	100	
21	V	104	
22	W	94	
23	X	85	
24	Y	78	
25	Z	63	
26	a	59	
27	b	57	
28	c	55	

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Mol	Chain	Length	Quality of chain
29	d	46	
30	e	65	
31	f	38	

2 Entry composition

There are 36 unique types of molecules in this entry. The entry contains 87635 atoms, of which 28 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a RNA chain called P-site fMet-tRNA(fMet).

Mol	Chain	Residues	Atoms					AltConf	Trace
1	1	4	Total	C	N	O	P	0	0
			84	38	16	26	4		

- Molecule 2 is a RNA chain called 23S rRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	A	2795	Total	C	N	O	P	3	0
			60085	26803	11076	19408	2798		

- Molecule 3 is a RNA chain called 5S rRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
3	B	120	Total	C	N	O	P	0	0
			2569	1144	468	837	120		

- Molecule 4 is a protein called 50S ribosomal protein L2.

Mol	Chain	Residues	Atoms					AltConf	Trace
4	C	271	Total	C	N	O	S	0	0
			2082	1288	423	364	7		

- Molecule 5 is a protein called 50S ribosomal protein L3.

Mol	Chain	Residues	Atoms					AltConf	Trace
5	D	209	Total	C	N	O	S	0	0
			1564	979	288	293	4		

- Molecule 6 is a protein called 50S ribosomal protein L4.

Mol	Chain	Residues	Atoms					AltConf	Trace
6	E	201	Total	C	N	O	S	0	0
			1551	974	283	289	5		

- Molecule 7 is a protein called 50S ribosomal protein L5.

Mol	Chain	Residues	Atoms					AltConf	Trace
7	F	177	Total	C	N	O	S	0	0
			1410	899	249	256	6		

- Molecule 8 is a protein called 50S ribosomal protein L6.

Mol	Chain	Residues	Atoms					AltConf	Trace
8	G	176	Total	C	N	O	S	0	0
			1322	832	243	245	2		

- Molecule 9 is a protein called 50S ribosomal protein L9.

Mol	Chain	Residues	Atoms					AltConf	Trace
9	H	50	Total	C	N	O	S	0	0
			384	247	68	68	1		

- Molecule 10 is a protein called 50S ribosomal protein L13.

Mol	Chain	Residues	Atoms					AltConf	Trace
10	K	142	Total	C	N	O	S	0	0
			1128	714	212	198	4		

- Molecule 11 is a protein called 50S ribosomal protein L14.

Mol	Chain	Residues	Atoms					AltConf	Trace
11	L	123	Total	C	N	O	S	0	0
			946	593	181	166	6		

- Molecule 12 is a protein called 50S ribosomal protein L15.

Mol	Chain	Residues	Atoms					AltConf	Trace
12	M	144	Total	C	N	O	S	0	0
			1052	654	207	189	2		

- Molecule 13 is a protein called 50S ribosomal protein L16.

Mol	Chain	Residues	Atoms					AltConf	Trace
13	N	136	Total	C	N	O	S	0	0
			1073	686	205	176	6		

- Molecule 14 is a protein called 50S ribosomal protein L17.

Mol	Chain	Residues	Atoms					AltConf	Trace
14	O	125	Total	C	N	O	S	0	0
			993	613	202	173	5		

- Molecule 15 is a protein called 50S ribosomal protein L18.

Mol	Chain	Residues	Atoms					AltConf	Trace
15	P	117	Total	C	N	O	S	0	0
			899	557	179	162	1		

- Molecule 16 is a protein called 50S ribosomal protein L19.

Mol	Chain	Residues	Atoms					AltConf	Trace
16	Q	114	Total	C	N	O	S	0	0
			916	574	179	162	1		

- Molecule 17 is a protein called 50S ribosomal protein L20.

Mol	Chain	Residues	Atoms					AltConf	Trace
17	R	117	Total	C	N	O	S	0	0
			946	604	192	150			

- Molecule 18 is a protein called 50S ribosomal protein L21.

Mol	Chain	Residues	Atoms					AltConf	Trace
18	S	103	Total	C	N	O	S	0	0
			815	516	153	144	2		

- Molecule 19 is a protein called 50S ribosomal protein L22.

Mol	Chain	Residues	Atoms					AltConf	Trace
19	T	110	Total	C	N	O	S	0	0
			856	532	166	155	3		

- Molecule 20 is a protein called 50S ribosomal protein L23.

Mol	Chain	Residues	Atoms					AltConf	Trace
20	U	95	Total	C	N	O	S	0	0
			756	479	141	135	1		

- Molecule 21 is a protein called 50S ribosomal protein L24.

Mol	Chain	Residues	Atoms				AltConf	Trace
21	V	102	Total	C	N	O		
			779	492	146	141	0	0

- Molecule 22 is a protein called 50S ribosomal protein L25.

Mol	Chain	Residues	Atoms				AltConf	Trace
22	W	94	Total	C	N	O	S	
			752	479	137	133	3	0

- Molecule 23 is a protein called 50S ribosomal protein L27.

Mol	Chain	Residues	Atoms				AltConf	Trace
23	X	76	Total	C	N	O	S	
			579	359	117	102	1	0

- Molecule 24 is a protein called 50S ribosomal protein L28.

Mol	Chain	Residues	Atoms				AltConf	Trace
24	Y	77	Total	C	N	O	S	
			624	388	129	105	2	0

- Molecule 25 is a protein called 50S ribosomal protein L29.

Mol	Chain	Residues	Atoms				AltConf	Trace
25	Z	62	Total	C	N	O	S	
			500	308	98	93	1	0

- Molecule 26 is a protein called 50S ribosomal protein L30.

Mol	Chain	Residues	Atoms				AltConf	Trace
26	a	58	Total	C	N	O	S	
			448	281	87	78	2	0

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

Mol	Chain	Residues	Atoms				AltConf	Trace
27	b	56	Total	C	N	O	S	
			443	269	94	79	1	0

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

Mol	Chain	Residues	Atoms				AltConf	Trace
28	c	51	Total	C	N	O	0	0
			414	266	76	72		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
29	d	46	Total	C	N	O	S	0	0
			376	228	90	56	2		

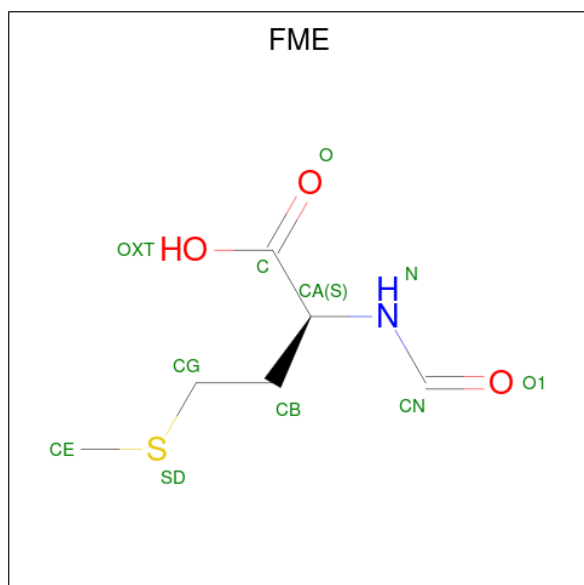
- Molecule 30 is a protein called 50S ribosomal protein L35.

Mol	Chain	Residues	Atoms					AltConf	Trace
30	e	64	Total	C	N	O	S	0	0
			503	323	105	73	2		

- Molecule 31 is a protein called 50S ribosomal protein L36.

Mol	Chain	Residues	Atoms					AltConf	Trace
31	f	38	Total	C	N	O	S	0	0
			301	185	65	47	4		

- Molecule 32 is N-FORMYLMETHIONINE (CCD ID: FME) (formula: $C_6H_{11}NO_3S$).

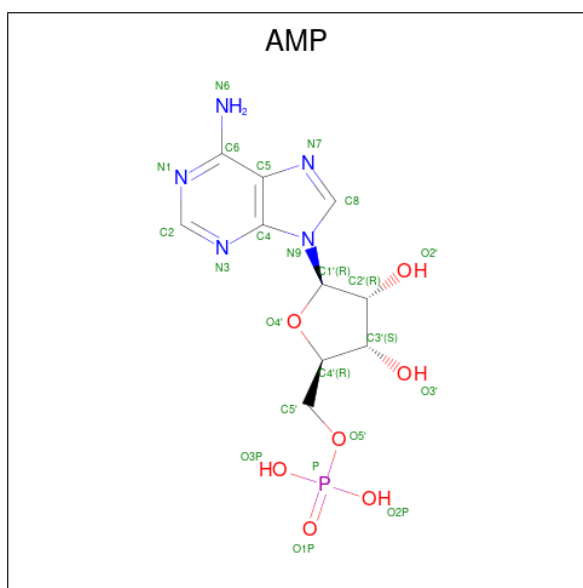


Mol	Chain	Residues	Atoms					AltConf
32	1	1	Total	C	N	O	S	0
			10	6	1	2	1	

- Molecule 33 is MAGNESIUM ION (CCD ID: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		AltConf
33	1	1	Total	Mg	0
			1	1	
33	A	369	Total	Mg	0
			369	369	
33	B	8	Total	Mg	0
			8	8	
33	C	1	Total	Mg	0
			1	1	
33	O	1	Total	Mg	0
			1	1	
33	R	1	Total	Mg	0
			1	1	
33	b	1	Total	Mg	0
			1	1	

- Molecule 34 is ADENOSINE MONOPHOSPHATE (CCD ID: AMP) (formula: C₁₀H₁₄N₅O₇P).



Mol	Chain	Residues	Atoms					AltConf
34	A	1	Total	C	N	O	P	0
			22	10	5	6	1	

- Molecule 35 is cadazolid (CCD ID: JJH) (formula: C₂₉H₂₉F₂N₃O₈) (labeled as "Ligand of Interest" by depositor).



Mol	Chain	Residues	Atoms						AltConf
35	A	1	Total 70	C 29	F 2	H 28	N 3	O 8	0

- Molecule 36 is ZINC ION (CCD ID: ZN) (formula: Zn).

Mol	Chain	Residues	Atoms	AltConf
36	f	1	Total Zn 1 1	0

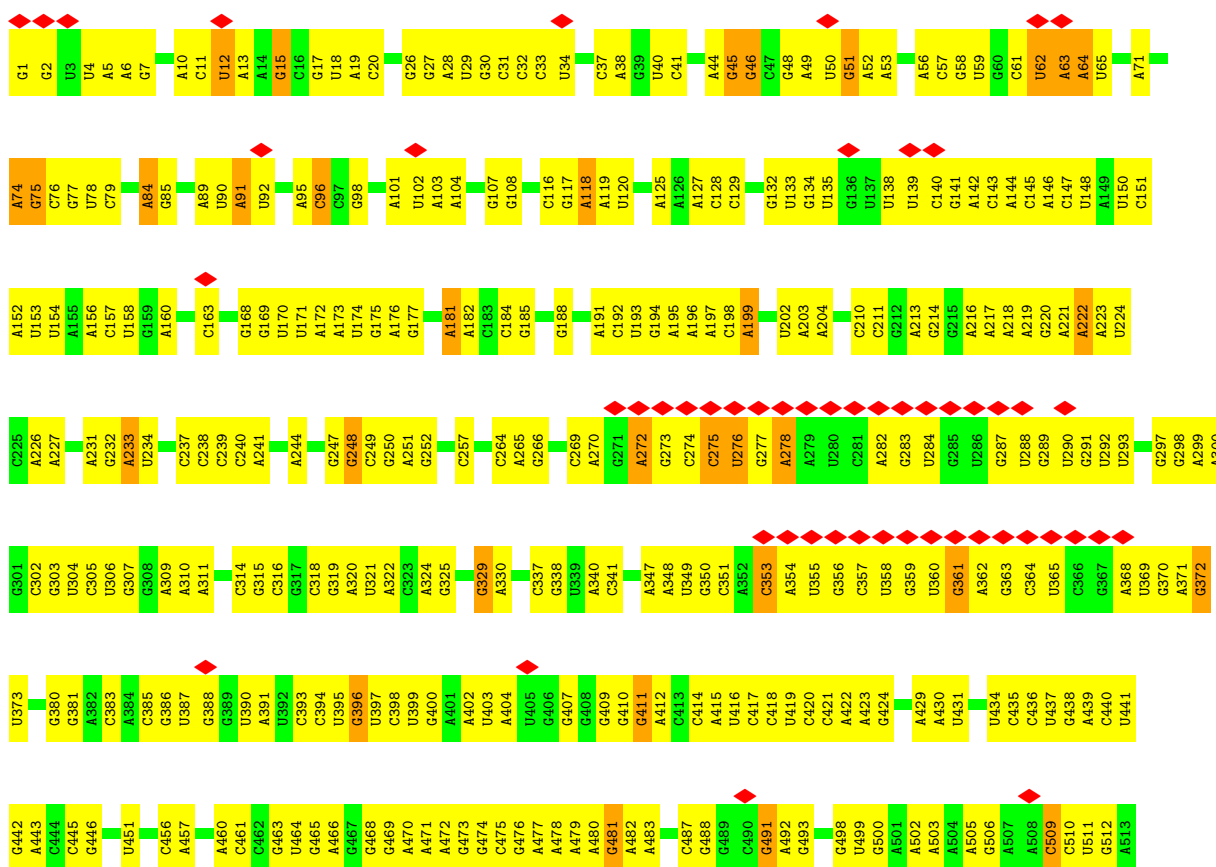
3 Residue-property plots

These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and atom inclusion in map density. Residues are color-coded according to the number of geometric quality criteria for which they contain at least one outlier: green = 0, yellow = 1, orange = 2 and red = 3 or more. A red diamond above a residue indicates a poor fit to the EM map for this residue (all-atom inclusion < 40%). Stretches of 2 or more consecutive residues without any outlier are shown as a green connector. Residues present in the sample, but not in the model, are shown in grey.

• Molecule 1: P-site fMet-tRNA(fMet)

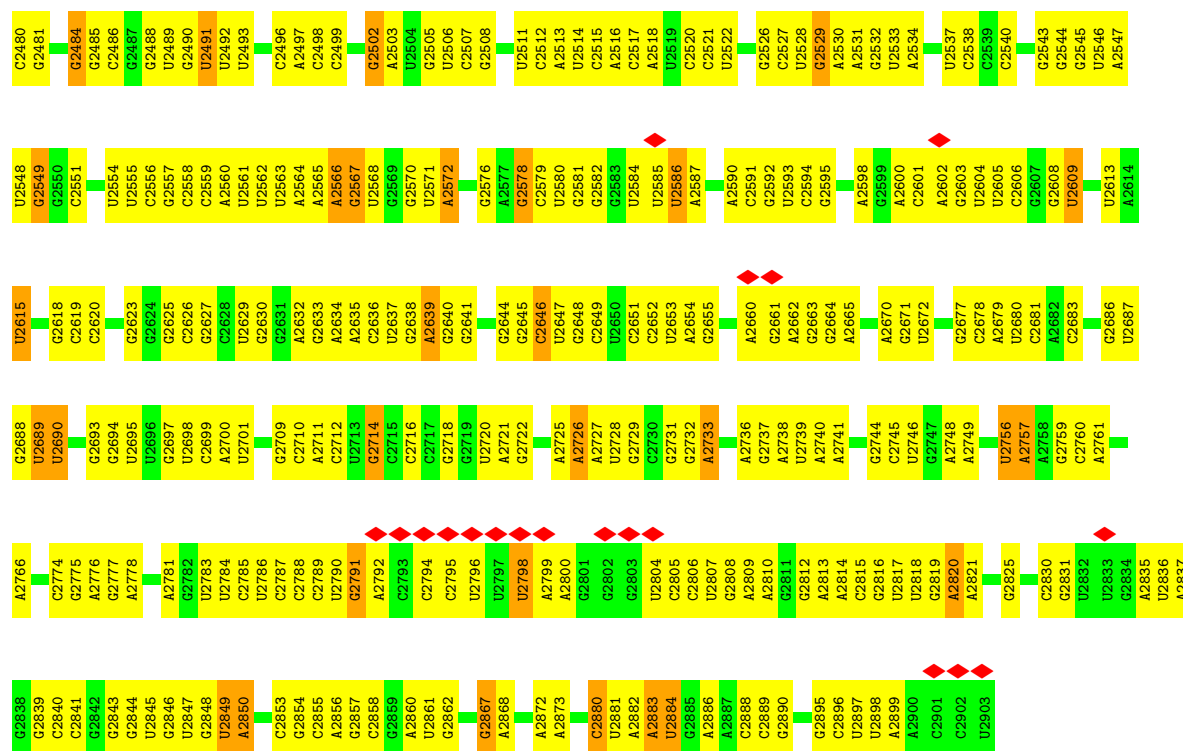


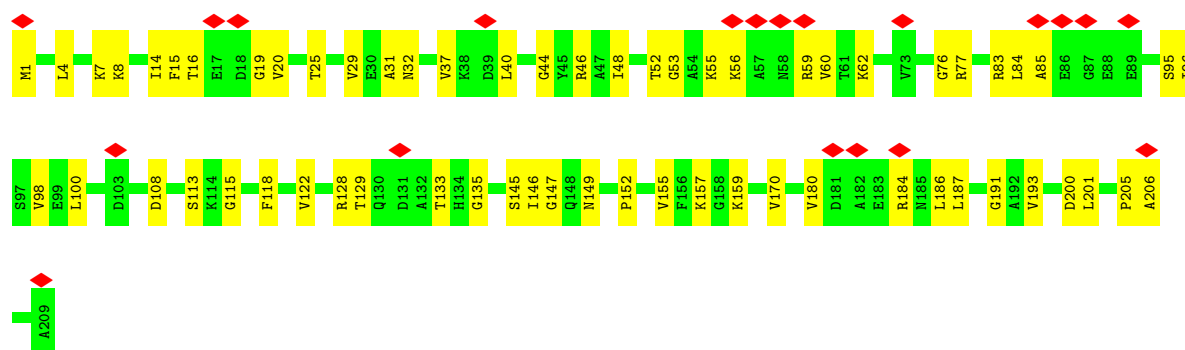
• Molecule 2: 23S rRNA



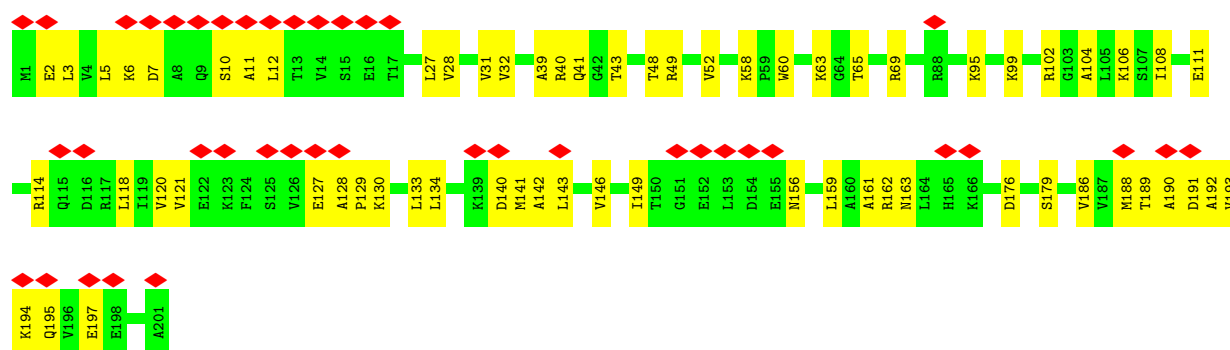




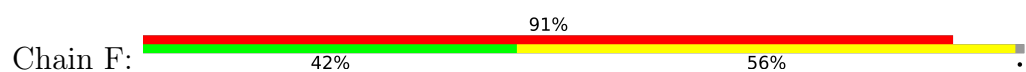




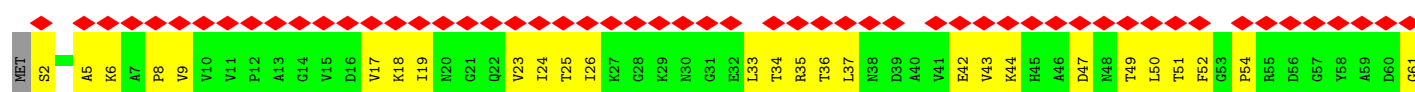
• Molecule 6: 50S ribosomal protein L4

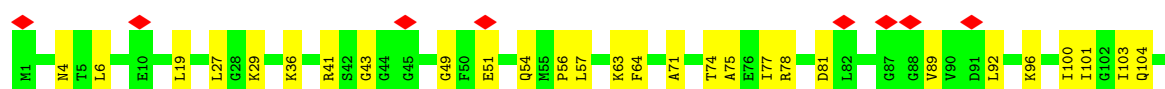


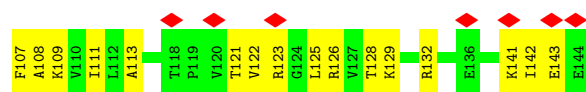
• Molecule 7: 50S ribosomal protein L5



• Molecule 8: 50S ribosomal protein L6







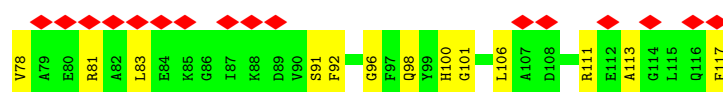
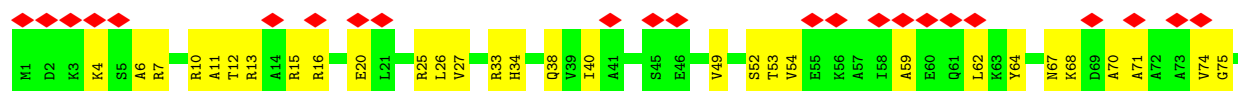
- Molecule 13: 50S ribosomal protein L16



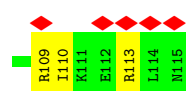
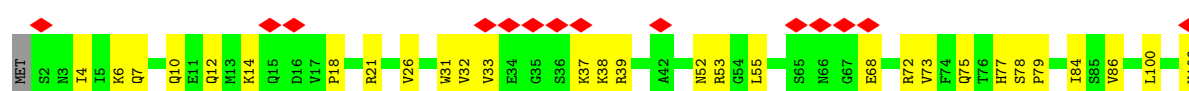
- Molecule 14: 50S ribosomal protein L17



- Molecule 15: 50S ribosomal protein L18

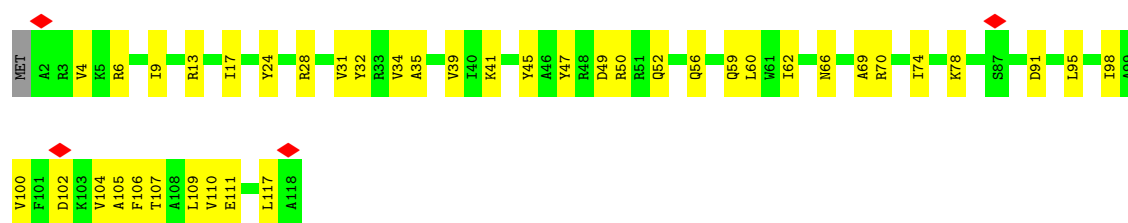


- Molecule 16: 50S ribosomal protein L19



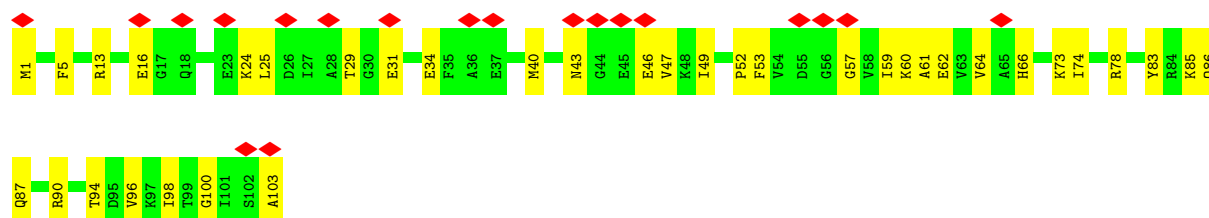
- Molecule 17: 50S ribosomal protein L20

Chain R:  65% 34%



- Molecule 18: 50S ribosomal protein L21

Chain S:  18% 65% 35%



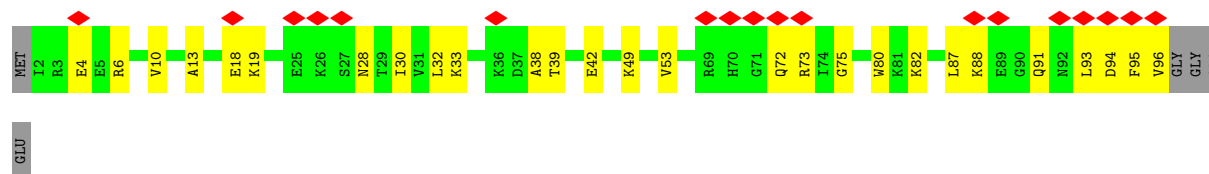
- Molecule 19: 50S ribosomal protein L22

Chain T:  12% 64% 36%



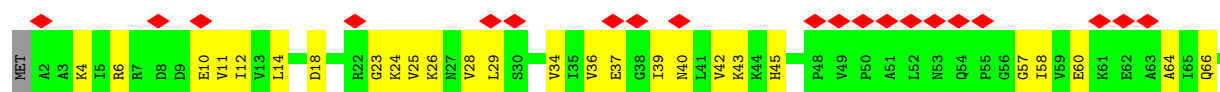
- Molecule 20: 50S ribosomal protein L23

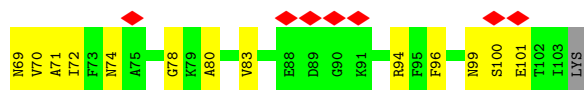
Chain U:  18% 68% 27% 5%



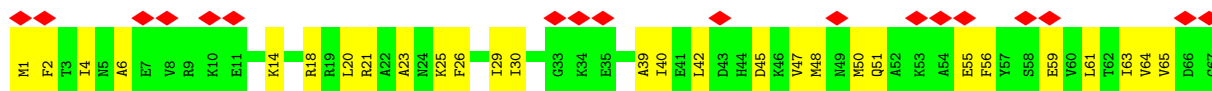
- Molecule 21: 50S ribosomal protein L24

Chain V:  26% 61% 38%





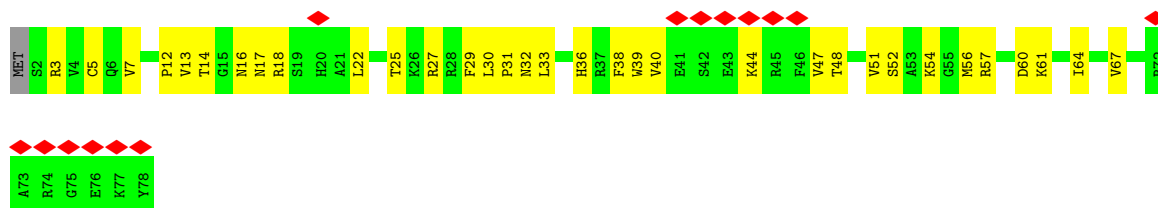
- Molecule 22: 50S ribosomal protein L25



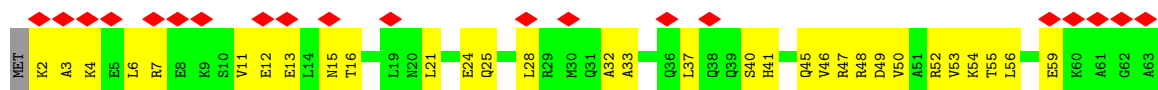
- Molecule 23: 50S ribosomal protein L27



- Molecule 24: 50S ribosomal protein L28



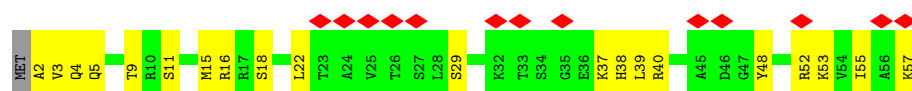
- Molecule 25: 50S ribosomal protein L29



- Molecule 26: 50S ribosomal protein L30



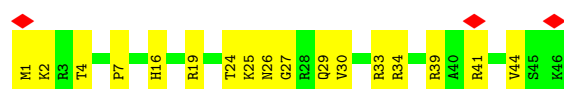
- Molecule 27: 50S ribosomal protein L32



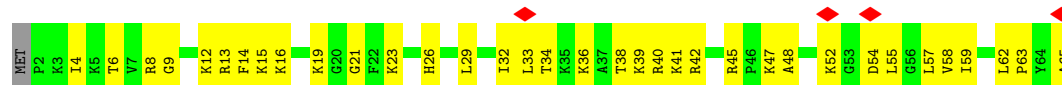
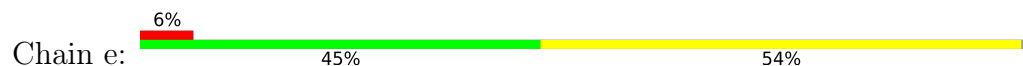
- Molecule 28: 50S ribosomal protein L33



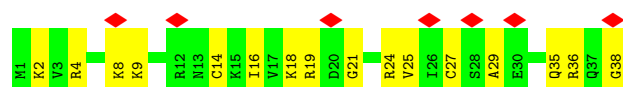
- Molecule 29: 50S ribosomal protein L34



- Molecule 30: 50S ribosomal protein L35



- Molecule 31: 50S ribosomal protein L36



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C1	Depositor
Number of particles used	59148	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	1.14	Depositor
Minimum defocus (nm)	Not provided	
Maximum defocus (nm)	Not provided	
Magnification	Not provided	
Image detector	FEI FALCON II (4k x 4k)	Depositor
Maximum map value	0.616	Depositor
Minimum map value	-0.384	Depositor
Average map value	0.002	Depositor
Map value standard deviation	0.026	Depositor
Recommended contour level	0.1	Depositor
Map size (Å)	312.91, 312.91, 302.12	wwPDB
Map dimensions	290, 290, 280	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	1.079, 1.079, 1.079	Depositor

5 Model quality [i](#)

5.1 Standard geometry [i](#)

Bond lengths and bond angles in the following residue types are not validated in this section: AMP, MG, JJH, FME, ZN

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	1	0.10	0/93	0.35	0/142
2	A	0.10	0/67296	0.20	0/104979
3	B	0.08	0/2872	0.16	0/4478
4	C	0.14	0/2121	0.36	0/2852
5	D	0.12	0/1585	0.34	0/2134
6	E	0.12	0/1570	0.32	0/2113
7	F	0.14	0/1434	0.39	0/1926
8	G	0.13	0/1342	0.35	0/1816
9	H	0.14	0/389	0.41	0/523
10	K	0.12	0/1151	0.30	0/1551
11	L	0.14	0/955	0.33	0/1279
12	M	0.15	0/1061	0.39	0/1413
13	N	0.13	0/1092	0.32	0/1460
14	O	0.13	0/1006	0.37	0/1345
15	P	0.11	0/909	0.31	0/1219
16	Q	0.13	0/928	0.34	0/1242
17	R	0.13	0/959	0.34	0/1278
18	S	0.12	0/828	0.30	0/1107
19	T	0.12	0/863	0.32	0/1156
20	U	0.10	0/763	0.30	0/1021
21	V	0.12	0/787	0.33	0/1051
22	W	0.13	0/765	0.33	0/1025
23	X	0.13	0/586	0.36	0/776
24	Y	0.12	0/634	0.35	0/848
25	Z	0.10	0/501	0.32	0/667
26	a	0.11	0/452	0.29	0/605
27	b	0.12	0/449	0.32	0/599
28	c	0.12	0/421	0.31	0/561
29	d	0.13	0/379	0.38	0/498
30	e	0.13	0/512	0.36	0/676
31	f	0.12	0/302	0.30	0/397
All	All	0.11	0/95005	0.24	0/142737

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

There are no planarity outliers.

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	1	84	0	44	2	0
2	A	60085	0	30220	1583	0
3	B	2569	0	1301	66	0
4	C	2082	0	2154	83	0
5	D	1564	0	1616	61	0
6	E	1551	0	1619	53	0
7	F	1410	0	1444	102	0
8	G	1322	0	1371	68	0
9	H	384	0	405	21	0
10	K	1128	0	1162	36	0
11	L	946	0	1023	50	0
12	M	1052	0	1129	43	0
13	N	1073	0	1157	35	0
14	O	993	0	1034	41	0
15	P	899	0	935	38	0
16	Q	916	0	962	31	0
17	R	946	0	1019	33	0
18	S	815	0	839	28	0
19	T	856	0	922	32	0
20	U	756	0	817	22	0
21	V	779	0	831	32	0
22	W	752	0	780	40	0
23	X	579	0	594	14	0
24	Y	624	0	652	34	0
25	Z	500	0	531	28	0
26	a	448	0	488	26	0
27	b	443	0	458	22	0
28	c	414	0	442	16	0
29	d	376	0	418	15	0
30	e	503	0	572	43	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
31	f	301	0	340	16	0
32	1	10	0	10	0	0
33	1	1	0	0	0	0
33	A	369	0	0	0	0
33	B	8	0	0	0	0
33	C	1	0	0	0	0
33	O	1	0	0	0	0
33	R	1	0	0	0	0
33	b	1	0	0	0	0
34	A	22	0	12	1	0
35	A	42	28	0	0	0
36	f	1	0	0	0	0
All	All	87607	28	57301	2423	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 17.

All (2423) 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)
2:A:775:G:H4'	2:A:776:G:H5'	1.45	0.97
13:N:77:PRO:HG2	13:N:80:VAL:HG21	1.47	0.96
20:U:53:VAL:HG11	20:U:87:LEU:HD13	1.48	0.94
7:F:108:VAL:HG11	7:F:176:PRO:HG2	1.47	0.93
8:G:164:TYR:HB2	8:G:167:GLU:HG2	1.50	0.91
2:A:1009:A:H5'	17:R:59:GLN:HG3	1.53	0.90
7:F:12:VAL:HG13	7:F:16:LEU:HD13	1.53	0.89
2:A:45:G:H5'	2:A:46:G:H5'	1.56	0.88
21:V:28:VAL:HG12	21:V:34:VAL:HG12	1.54	0.88
8:G:44:LYS:HB2	8:G:51:THR:HB	1.58	0.86
2:A:545:U:H3	2:A:548:G:H1	1.22	0.86
13:N:75:GLU:HB2	13:N:90:GLU:HG3	1.55	0.86
19:T:4:ILE:HG12	19:T:106:VAL:HG22	1.57	0.86
26:a:6:LYS:HG2	26:a:37:GLU:HG2	1.57	0.85
4:C:29:PRO:HG2	4:C:34:LEU:HD11	1.58	0.85
2:A:2508:G:H1	2:A:2580:U:H3	1.26	0.84
2:A:1597:A:H5''	2:A:1598:A:H5'	1.58	0.84
11:L:102:PRO:HB3	11:L:121:GLU:HB3	1.58	0.84
2:A:181:A:H1'	2:A:435:C:H5'	1.60	0.84
2:A:479:A:H4'	2:A:480:A:H5'	1.61	0.83
8:G:37:LEU:HD13	8:G:68:ALA:HB1	1.60	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:D:56:LYS:HB2	5:D:59:ARG:HB2	1.61	0.82
11:L:108:ARG:HG3	11:L:113:MET:HE1	1.61	0.82
7:F:69:LYS:HE3	7:F:84:PRO:HG3	1.61	0.82
7:F:142:ASP:HB3	7:F:145:LYS:HD3	1.59	0.82
6:E:5:LEU:HD11	6:E:12:LEU:HD23	1.60	0.81
13:N:20:LEU:HD13	22:W:81:PRO:HG2	1.62	0.81
16:Q:14:LYS:HE2	16:Q:77:HIS:HA	1.63	0.81
7:F:160:ALA:HB1	7:F:165:GLU:HB2	1.63	0.81
31:f:18:LYS:HE2	31:f:21:GLY:HA2	1.61	0.81
7:F:103:LEU:HA	7:F:107:ALA:HB3	1.64	0.80
2:A:1798:U:H5''	4:C:258:ARG:HB2	1.64	0.79
2:A:1645:G:H5''	2:A:1646:C:H5'	1.64	0.79
2:A:1664:A:H61	2:A:1996:C:H42	1.27	0.79
12:M:57:LEU:HD22	30:e:54:ASP:HB3	1.64	0.78
2:A:358:U:H2'	2:A:359:G:H8	1.49	0.77
25:Z:2:LYS:HE3	25:Z:6:LEU:HD11	1.67	0.76
6:E:149:ILE:HB	6:E:188:MET:HG2	1.68	0.76
2:A:820:A:H4'	2:A:836:G:H22	1.51	0.76
7:F:34:ILE:HG12	7:F:156:ILE:HG12	1.67	0.76
2:A:2343:U:HO2'	2:A:2373:G:HO2'	1.27	0.76
7:F:12:VAL:HG22	7:F:172:ALA:HB1	1.68	0.76
2:A:834:G:H5'	30:e:57:LEU:HD11	1.68	0.75
2:A:1125:G:OP2	2:A:1126:A:O2'	2.03	0.75
18:S:24:LYS:HA	18:S:94:THR:HG23	1.68	0.75
3:B:30:C:H1'	3:B:57:A:H61	1.51	0.74
9:H:40:THR:OG1	9:H:43:ASN:OD1	2.05	0.74
29:d:24:THR:HG23	29:d:27:GLY:H	1.52	0.74
2:A:1942:C:OP2	2:A:1943:U:O2'	2.05	0.74
22:W:55:GLU:HB2	22:W:59:GLU:HG2	1.67	0.74
2:A:1048:A:H1'	2:A:1112:G:H21	1.52	0.74
2:A:2511:U:OP1	5:D:128:ARG:NH1	2.21	0.74
6:E:41:GLN:HG2	6:E:43:THR:HG23	1.69	0.74
2:A:764:A:H5'	4:C:209:GLY:HA2	1.70	0.74
4:C:232:HIS:HA	4:C:242:LYS:HD2	1.68	0.74
7:F:103:LEU:HD12	7:F:107:ALA:HB3	1.69	0.73
2:A:2304:G:H4'	7:F:130:MET:HA	1.70	0.73
8:G:35:ARG:HG2	8:G:71:LEU:HD21	1.71	0.73
13:N:53:MET:HG3	13:N:120:ALA:HB2	1.70	0.73
2:A:992:C:OP1	17:R:47:TYR:OH	2.07	0.72
2:A:1068:G:N2	2:A:1095:A:O2'	2.22	0.72
2:A:1082:U:H3'	2:A:1083:U:H5''	1.71	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:Y:12:PRO:HB3	24:Y:30:LEU:HD23	1.71	0.72
2:A:1818:U:OP2	4:C:156:ARG:NH1	2.22	0.72
4:C:245:VAL:HG12	4:C:251:GLN:HA	1.69	0.72
2:A:219:A:N3	2:A:234:U:O2'	2.23	0.72
2:A:1664:A:H61	2:A:1996:C:N4	1.87	0.72
2:A:2683:C:O2	11:L:70:ARG:NH2	2.22	0.72
2:A:1044:C:H4'	2:A:1048:A:H5'	1.72	0.72
2:A:1826:G:O2'	2:A:1971:U:OP2	2.07	0.71
2:A:775:G:H5'	2:A:777:G:H1'	1.71	0.71
7:F:110:ARG:HB2	7:F:137:ILE:HG23	1.73	0.71
8:G:25:THR:HA	8:G:34:THR:HG22	1.73	0.70
8:G:86:LYS:HG3	8:G:165:ALA:HB2	1.72	0.70
14:O:121:LYS:HA	14:O:124:ALA:HB3	1.71	0.70
4:C:132:MET:HE2	4:C:188:CYS:HB2	1.74	0.70
17:R:105:ALA:HB1	18:S:40:MET:HE1	1.73	0.70
30:e:33:LEU:HA	30:e:36:LYS:HD2	1.74	0.70
2:A:2566:A:N1	11:L:28:SER:OG	2.23	0.70
5:D:14:ILE:HD13	16:Q:12:GLN:HE22	1.56	0.69
2:A:845:A:O2'	2:A:846:U:O4'	2.09	0.69
3:B:1:U:H2'	3:B:2:G:H8	1.56	0.69
7:F:125:ARG:O	7:F:127:ASN:ND2	2.24	0.69
10:K:114:LEU:HG	10:K:118:MET:HE3	1.74	0.69
2:A:1797:G:O2'	4:C:257:THR:OG1	2.10	0.69
8:G:17:VAL:HG12	8:G:26:ILE:HG13	1.73	0.69
2:A:276:U:O2'	2:A:278:A:N7	2.20	0.69
2:A:2506:U:OP2	2:A:2576:G:N1	2.25	0.69
2:A:224:U:O4	2:A:419:U:O2'	2.11	0.69
2:A:1654:A:O2'	5:D:118:PHE:O	2.11	0.69
2:A:1779:U:OP2	2:A:1784:A:N6	2.21	0.69
2:A:2285:C:OP2	28:c:6:ARG:NE	2.26	0.69
25:Z:21:LEU:HB3	25:Z:50:VAL:HG12	1.75	0.69
2:A:1800:C:H3'	4:C:146:MET:HE1	1.76	0.68
2:A:84:A:N1	2:A:98:G:O2'	2.27	0.68
2:A:1918:A:O2'	2:A:1920:C:N4	2.26	0.68
5:D:25:THR:OG1	5:D:191:GLY:O	2.11	0.68
8:G:33:LEU:HB3	8:G:75:MET:HE3	1.75	0.68
19:T:38:TYR:OH	27:b:37:LYS:O	2.12	0.68
2:A:698:C:O2'	2:A:734:A:N6	2.26	0.68
22:W:4:ILE:HB	22:W:63:ILE:HD13	1.76	0.68
2:A:320:A:N3	6:E:163:ASN:ND2	2.41	0.68
2:A:249:C:O2	30:e:12:LYS:NZ	2.26	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:1019:U:OP1	2:A:1035:U:O2'	2.10	0.68
2:A:210:C:OP1	29:d:29:GLN:NE2	2.27	0.68
2:A:414:C:O2	2:A:1864:U:O2'	2.12	0.68
2:A:411:G:OP2	2:A:2406:A:O2'	2.12	0.68
2:A:2898:U:H2'	2:A:2899:A:H8	1.59	0.68
2:A:2491:U:H5'	2:A:2570:G:H5''	1.74	0.67
21:V:40:ASN:ND2	21:V:64:ALA:O	2.27	0.67
23:X:25:ARG:HG3	23:X:31:VAL:HG11	1.76	0.67
2:A:577:G:O2'	2:A:1254:A:OP1	2.12	0.67
2:A:2303:G:O2'	7:F:121:SER:O	2.08	0.67
24:Y:36:HIS:HB2	24:Y:56:MET:HE1	1.74	0.67
2:A:2258:C:O2'	2:A:2427:C:OP2	2.11	0.67
12:M:19:LEU:HD23	12:M:27:LEU:HD13	1.76	0.67
19:T:86:MET:HE3	19:T:88:ARG:HD3	1.76	0.67
2:A:1132:U:H3'	2:A:1133:A:H5''	1.75	0.67
2:A:2328:A:H2'	2:A:2329:U:C6	2.30	0.67
11:L:114:LYS:HE3	11:L:118:LEU:HD11	1.75	0.67
2:A:1264:A:OP1	27:b:16:ARG:NH1	2.28	0.67
2:A:381:G:OP1	24:Y:18:ARG:NH2	2.28	0.67
2:A:993:G:OP1	17:R:50:ARG:NE	2.25	0.66
2:A:1250:G:H5''	17:R:6:ARG:HD3	1.76	0.66
2:A:2298:A:OP1	7:F:71:ARG:NH2	2.28	0.66
21:V:14:LEU:HD11	21:V:71:ALA:HB2	1.77	0.66
2:A:1866:A:N6	2:A:1875:G:O2'	2.28	0.66
12:M:132:ARG:HG3	12:M:142:ILE:HD12	1.76	0.66
12:M:141:LYS:NZ	12:M:143:GLU:OE1	2.26	0.66
9:H:41:LYS:HA	9:H:44:ILE:HG12	1.76	0.66
2:A:380:G:H5'	24:Y:18:ARG:HG2	1.78	0.66
2:A:1094:U:H1'	2:A:1097:U:H5	1.60	0.66
8:G:123:ALA:HB2	8:G:133:LEU:HG	1.77	0.66
2:A:1432:G:H2'	2:A:1433:A:C8	2.31	0.66
2:A:117:G:OP2	2:A:119:A:O2'	2.13	0.66
2:A:239:C:HO2'	2:A:622:G:HO2'	1.44	0.66
2:A:2602[B]:A:H3'	2:A:2603[B]:G:H5''	1.77	0.66
2:A:2756:U:OP2	31:f:19:ARG:NE	2.27	0.66
19:T:24:ILE:HD13	19:T:36:LEU:HD11	1.76	0.66
25:Z:50:VAL:HA	25:Z:53:VAL:HG12	1.78	0.66
2:A:743:A:O2'	2:A:1659:G:OP1	2.12	0.66
8:G:19:ILE:HG12	8:G:24:ILE:HG23	1.76	0.66
2:A:2848:G:O2'	2:A:2867:G:N2	2.28	0.65
5:D:29:VAL:HB	5:D:98:VAL:HG12	1.78	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:476:G:N1	2:A:479:A:OP2	2.29	0.65
2:A:1149:G:H2'	2:A:1150:C:C6	2.30	0.65
2:A:1340:U:OP2	20:U:82:LYS:NZ	2.30	0.65
7:F:101:GLU:O	7:F:105:THR:HG22	1.97	0.65
21:V:12:ILE:HG21	21:V:80:ALA:HB2	1.79	0.65
2:A:197:A:N6	2:A:2430:A:O2'	2.30	0.65
2:A:1392:A:N6	20:U:18:GLU:OE1	2.29	0.65
2:A:2798:U:H4'	2:A:2799:A:H5'	1.79	0.65
2:A:1270:C:H5''	2:A:1271:G:H5'	1.79	0.65
2:A:1564:C:H2'	2:A:1565:C:C6	2.32	0.65
2:A:1827:U:OP1	2:A:1971:U:O2'	2.15	0.65
2:A:1853:A:N6	2:A:1888:G:O2'	2.29	0.65
2:A:601:C:O2'	2:A:605:G:OP1	2.14	0.65
2:A:1715:G:O2'	2:A:1743:G:O6	2.12	0.65
2:A:1757:A:O2'	2:A:1758:U:OP1	2.12	0.65
6:E:48:THR:O	6:E:52:VAL:HG23	1.97	0.65
2:A:329:G:OP2	21:V:69:ASN:ND2	2.29	0.65
5:D:55:LYS:HD3	5:D:60:VAL:HG22	1.77	0.65
2:A:160:A:N3	2:A:2208:C:O2'	2.30	0.64
2:A:1485:U:H2'	2:A:1486:U:C6	2.32	0.64
3:B:66:A:OP2	3:B:108:A:N6	2.26	0.64
5:D:32:ASN:ND2	5:D:52:THR:OG1	2.29	0.64
2:A:188:G:H5'	24:Y:14:THR:HG21	1.79	0.64
2:A:2619:C:OP1	5:D:157:LYS:NZ	2.28	0.64
2:A:671:C:OP1	12:M:43:GLY:N	2.26	0.64
2:A:2291:U:H2'	2:A:2292:U:C6	2.33	0.64
2:A:2562:U:H4'	11:L:25:LEU:HD22	1.79	0.64
2:A:1387:A:H5'	2:A:1469:A:H1'	1.80	0.64
2:A:2086:U:H2'	2:A:2087:G:C8	2.33	0.64
6:E:159:LEU:HD22	6:E:162:ARG:HH11	1.62	0.64
11:L:69:VAL:HG21	11:L:104:THR:HG21	1.79	0.64
28:c:8:LYS:HE3	30:e:34:THR:HG21	1.78	0.64
2:A:1211:C:H3'	2:A:1212:G:H5'	1.79	0.64
12:M:78:ARG:HG2	12:M:113:ALA:HB3	1.80	0.64
22:W:2:PHE:HB2	22:W:61:LEU:HD22	1.80	0.64
2:A:2350:C:OP2	30:e:45:ARG:NH2	2.30	0.64
22:W:30:ILE:HD11	22:W:40:ILE:HD13	1.79	0.64
2:A:1746:A:H2'	2:A:1747:U:C6	2.33	0.64
8:G:107:LEU:HD13	8:G:152:ARG:HB2	1.80	0.64
2:A:1264:A:OP2	2:A:1265:A:O2'	2.11	0.64
5:D:25:THR:HG21	5:D:193:VAL:HG22	1.80	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:a:6:LYS:HB2	26:a:58:GLU:HG2	1.80	0.64
2:A:807:U:OP2	12:M:41:ARG:NH1	2.28	0.63
2:A:1057:A:N6	2:A:1087:G:OP2	2.31	0.63
4:C:205:LEU:HB3	4:C:210:ALA:HB3	1.79	0.63
11:L:76:VAL:HG12	16:Q:73:VAL:HB	1.80	0.63
19:T:17:VAL:HG12	19:T:76:VAL:HG21	1.80	0.63
2:A:703:U:H2'	2:A:704:G:O4'	1.98	0.63
2:A:729:G:O2'	2:A:763:G:H4'	1.98	0.63
2:A:878:A:H3'	2:A:879:G:H8	1.64	0.63
2:A:1301:A:O2'	2:A:1607:C:N3	2.27	0.63
7:F:101:GLU:HA	7:F:104:ILE:HG12	1.79	0.63
2:A:1028:A:OP2	2:A:1126:A:N6	2.26	0.63
2:A:2547:A:H2'	2:A:2548:U:C6	2.33	0.63
18:S:25:LEU:H	18:S:94:THR:HG21	1.62	0.63
2:A:1754:A:N1	2:A:2716:C:O2'	2.27	0.63
2:A:2639:A:H2'	2:A:2640:G:O4'	1.97	0.63
20:U:30:ILE:HD12	20:U:32:LEU:HD21	1.79	0.63
2:A:2530:A:N7	8:G:172:LYS:NZ	2.41	0.63
2:A:529:A:OP2	10:K:116:ARG:NH2	2.24	0.63
2:A:1386:C:H2'	2:A:1387:A:C8	2.34	0.63
2:A:2304:G:H22	2:A:2312:U:H3	1.46	0.63
7:F:126:GLY:HA3	7:F:160:ALA:HB3	1.81	0.63
21:V:36:VAL:HB	21:V:39:ILE:HB	1.81	0.63
2:A:1818:U:O2'	4:C:153:GLN:O	2.12	0.63
2:A:2291:U:OP1	2:A:2380:C:O2'	2.17	0.63
9:H:3:VAL:HB	9:H:36:ALA:HB1	1.80	0.63
2:A:605:G:OP1	6:E:99:LYS:NZ	2.30	0.62
21:V:11:VAL:CG1	21:V:70:VAL:HB	2.29	0.62
2:A:1816:C:N4	4:C:35:GLU:OE2	2.25	0.62
27:b:55:ILE:HG22	27:b:57:LYS:H	1.64	0.62
3:B:43:C:O2	7:F:92:ARG:NH2	2.32	0.62
7:F:94:GLU:O	7:F:98:GLU:HG3	2.00	0.62
19:T:5:ALA:HB3	19:T:54:ALA:HB2	1.81	0.62
2:A:306:U:H2'	2:A:307:G:O4'	2.00	0.62
2:A:589:U:H2'	2:A:590:A:H8	1.63	0.62
2:A:1048:A:H1'	2:A:1112:G:N2	2.15	0.62
2:A:240:C:OP2	2:A:241:A:O2'	2.10	0.62
2:A:414:C:H1'	2:A:1864:U:H1'	1.82	0.62
7:F:115:ARG:O	7:F:178:ARG:NH2	2.33	0.62
2:A:515:A:H1'	2:A:581:C:H1'	1.81	0.62
2:A:878:A:N6	2:A:899:A:O2'	2.32	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:976:G:HO2'	2:A:1155:A:HO2'	1.44	0.62
2:A:2474:U:OP2	2:A:2475:C:N4	2.27	0.62
2:A:2788:C:O2'	2:A:2809:A:N3	2.32	0.62
11:L:19:VAL:HB	11:L:41:ILE:HD12	1.82	0.62
2:A:2799:A:O2'	2:A:2800:A:H5''	2.00	0.62
10:K:1:MET:HE1	18:S:13:ARG:HB3	1.81	0.62
2:A:1869:G:N2	2:A:1871:A:O2'	2.32	0.62
2:A:2491:U:C5'	2:A:2570:G:H5''	2.30	0.62
25:Z:12:GLU:O	25:Z:16:THR:HG23	1.98	0.62
2:A:1296:G:OP1	2:A:2709:G:O2'	2.14	0.61
2:A:400:G:N7	24:Y:57:ARG:NH1	2.44	0.61
2:A:1567:G:OP2	4:C:83:TYR:OH	2.15	0.61
2:A:1727:C:H2'	2:A:1728:C:C6	2.36	0.61
3:B:77:U:OP1	22:W:21:ARG:NH1	2.31	0.61
10:K:13:ARG:NH1	10:K:49:ASP:O	2.31	0.61
13:N:41:LEU:HG	13:N:96:ILE:HG13	1.81	0.61
2:A:1433:A:H2'	2:A:1434:A:C8	2.35	0.61
2:A:2286:G:H4'	2:A:2287:A:O4'	2.00	0.61
13:N:66:ARG:NH1	13:N:104:GLU:OE2	2.33	0.61
2:A:675:A:OP1	6:E:58:LYS:NZ	2.27	0.61
2:A:910:A:H2'	2:A:911:A:C8	2.35	0.61
2:A:1581:G:H2'	2:A:1582:C:C6	2.34	0.61
2:A:2071:A:H2'	2:A:2072:C:C6	2.35	0.61
4:C:107:PRO:HG2	4:C:110:LEU:HD23	1.81	0.61
2:A:926:G:H2'	2:A:927:A:H8	1.66	0.61
2:A:1682:G:H2'	2:A:1683:U:C6	2.36	0.61
6:E:189:THR:OG1	6:E:191:ASP:OD1	2.14	0.61
7:F:26:MET:SD	7:F:30:ARG:NH1	2.73	0.61
13:N:17:ASN:O	13:N:38:ARG:NH1	2.33	0.61
2:A:1799:G:OP1	4:C:258:ARG:NH1	2.33	0.61
6:E:12:LEU:HD21	6:E:120:VAL:HG11	1.82	0.61
8:G:23:VAL:HA	8:G:36:THR:HA	1.81	0.61
2:A:125:A:OP2	29:d:19:ARG:NH2	2.31	0.61
2:A:1485:U:H2'	2:A:1486:U:H6	1.64	0.61
2:A:1515:A:H3'	2:A:1516:G:H8	1.66	0.61
2:A:1589:U:H2'	2:A:1590:A:H8	1.66	0.61
2:A:2070:A:H2'	2:A:2071:A:C8	2.35	0.61
24:Y:17:ASN:HD21	24:Y:27:ARG:HB3	1.66	0.61
2:A:45:G:C5'	2:A:46:G:H5'	2.30	0.61
2:A:548:G:H2'	2:A:549:G:C8	2.35	0.61
2:A:2898:U:H2'	2:A:2899:A:C8	2.36	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:M:81:ASP:HB3	12:M:100:ILE:HD13	1.83	0.61
15:P:68:LYS:HE2	15:P:101:GLY:HA2	1.83	0.61
2:A:576:U:H2'	2:A:577:G:C8	2.35	0.60
2:A:2215:C:H2'	2:A:2216:G:C8	2.37	0.60
2:A:1814:G:OP2	2:A:1815:A:O2'	2.09	0.60
2:A:2070:A:H2'	2:A:2071:A:H8	1.66	0.60
2:A:2727:A:O2'	2:A:2728:U:H5'	2.01	0.60
4:C:120:VAL:HG13	4:C:134:ASN:HD21	1.65	0.60
23:X:66:LYS:HG3	23:X:85:GLU:OE2	2.01	0.60
2:A:1437:C:HO2'	2:A:1516:G:HO2'	1.50	0.60
2:A:2014:A:H2'	2:A:2015:A:C8	2.36	0.60
25:Z:2:LYS:HE2	25:Z:52:ARG:HE	1.65	0.60
2:A:90:U:OP2	2:A:91:A:O2'	2.11	0.60
2:A:2532:G:N2	2:A:2663:G:O2'	2.35	0.60
2:A:2810:A:H4'	5:D:62:LYS:HD2	1.84	0.60
17:R:49:ASP:HA	17:R:52:GLN:HB2	1.82	0.60
2:A:829:A:N7	2:A:2247:A:O2'	2.34	0.60
12:M:51:GLU:OE2	30:e:57:LEU:HD13	2.00	0.60
2:A:807:U:O2'	2:A:2060:A:N1	2.32	0.60
2:A:1802:A:H2'	2:A:1803:A:C8	2.37	0.60
8:G:87:LEU:HD12	8:G:162:VAL:CG1	2.32	0.60
8:G:118:PRO:HD2	8:G:121:ILE:HG21	1.82	0.60
2:A:75:G:H4'	25:Z:48:ARG:NH2	2.17	0.60
2:A:720:U:H2'	2:A:721:A:C8	2.36	0.60
2:A:1287:A:O2'	2:A:1288:G:H5'	2.01	0.60
2:A:2025:C:H2'	2:A:2026:U:C6	2.37	0.60
2:A:2323:G:O2'	2:A:2324:U:H5'	2.02	0.60
2:A:832:U:H2'	2:A:833:A:C8	2.37	0.59
2:A:894:U:H2'	2:A:895:U:C6	2.37	0.59
2:A:2788:C:H2'	2:A:2789:C:C6	2.37	0.59
25:Z:11:VAL:O	25:Z:15:ASN:ND2	2.35	0.59
2:A:307:G:N1	2:A:310:A:OP2	2.29	0.59
2:A:1669:A:O2'	2:A:2549:G:OP1	2.14	0.59
2:A:2687:U:H2'	2:A:2688:G:O4'	2.02	0.59
14:O:29:VAL:HG11	14:O:75:ILE:HG23	1.83	0.59
2:A:685:A:N1	2:A:787:C:H1'	2.17	0.59
2:A:729:G:OP1	4:C:13:ARG:HG2	2.03	0.59
2:A:918:A:N3	3:B:80:U:O2'	2.33	0.59
2:A:1496:A:H2'	2:A:1498:C:C5	2.38	0.59
2:A:2306:C:OP2	2:A:2307:G:O2'	2.13	0.59
7:F:102:ARG:HA	7:F:106:ILE:CD1	2.33	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:1360:G:H2'	2:A:1361:G:H5'	1.84	0.59
6:E:111:GLU:OE1	6:E:114:ARG:NH1	2.35	0.59
7:F:102:ARG:HA	7:F:106:ILE:HD13	1.83	0.59
2:A:396:G:H1'	24:Y:29:PHE:HB3	1.83	0.59
2:A:463:G:N2	2:A:466:A:OP2	2.25	0.59
2:A:580:U:H2'	2:A:581:C:H6	1.67	0.59
2:A:1361:G:H2'	2:A:1362:C:H6	1.67	0.59
2:A:2290:G:H2'	2:A:2291:U:C6	2.38	0.59
2:A:2641:G:H5''	10:K:78:THR:HB	1.84	0.59
3:B:7:G:OP1	15:P:4:LYS:NZ	2.27	0.59
28:c:11:LEU:N	28:c:21:TYR:O	2.35	0.59
2:A:926:G:H2'	2:A:927:A:C8	2.37	0.59
2:A:1712:U:OP2	2:A:1713:A:O2'	2.13	0.59
2:A:2074:U:H2'	2:A:2075:U:C6	2.37	0.59
2:A:2637:U:H2'	2:A:2638:G:O4'	2.02	0.59
10:K:125:TYR:OH	10:K:132:HIS:NE2	2.27	0.59
2:A:859:G:O2'	2:A:916:G:O6	2.20	0.59
2:A:2557:G:H2'	2:A:2558:C:C6	2.38	0.59
3:B:8:C:OP1	15:P:15:ARG:NH2	2.31	0.59
2:A:570:G:H2'	2:A:2030:A:N7	2.18	0.59
2:A:1445:G:O6	2:A:1466:U:O2	2.20	0.59
7:F:117:LEU:HD11	7:F:130:MET:HE2	1.85	0.59
2:A:2644:G:O2'	2:A:2645:G:H5'	2.03	0.59
5:D:48:ILE:HG23	5:D:84:LEU:HD11	1.85	0.59
2:A:435:C:H2'	2:A:436:C:H5'	1.85	0.58
2:A:720:U:H2'	2:A:721:A:H8	1.67	0.58
2:A:2540:C:O2'	2:A:2740:A:N3	2.33	0.58
7:F:105:THR:HG23	7:F:106:ILE:HD12	1.84	0.58
2:A:156:A:H2'	2:A:157:C:C6	2.38	0.58
2:A:2215:C:H2'	2:A:2216:G:H8	1.68	0.58
2:A:2731:G:H2'	2:A:2732:G:C8	2.38	0.58
5:D:16:THR:HG22	5:D:20:VAL:O	2.02	0.58
7:F:40:VAL:HG11	7:F:43:ALA:HB2	1.85	0.58
2:A:2243:U:H2'	2:A:2244:U:C6	2.39	0.58
2:A:2529:G:H4'	8:G:175:LYS:HG3	1.84	0.58
2:A:2804:U:H2'	2:A:2805:C:C6	2.39	0.58
11:L:58:LEU:HA	11:L:89:ASN:HD22	1.67	0.58
2:A:582:A:H2'	2:A:583:G:H8	1.69	0.58
2:A:1026:G:OP1	2:A:1134:A:O2'	2.21	0.58
2:A:1808:A:H3'	2:A:1809:A:C8	2.39	0.58
2:A:287:G:H2'	2:A:288:U:C6	2.38	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:391:A:H1'	2:A:411:G:O4'	2.03	0.58
2:A:742:A:H2'	2:A:743:A:C8	2.38	0.58
2:A:2366:A:H4'	23:X:62:LYS:HE3	1.86	0.58
15:P:4:LYS:HD2	15:P:7:ARG:HH21	1.68	0.58
19:T:82:MET:CG	19:T:98:LYS:HB2	2.33	0.58
2:A:177:G:OP2	2:A:177:G:N2	2.21	0.58
2:A:1433:A:N1	2:A:1560:G:O6	2.37	0.58
2:A:2193:G:H2'	2:A:2194:U:C6	2.38	0.58
2:A:2567:G:H2'	2:A:2568:U:C6	2.38	0.58
2:A:2595:G:N2	2:A:2598:A:OP2	2.26	0.58
7:F:29:PRO:HB2	7:F:169:LEU:HD13	1.86	0.58
2:A:32:C:H2'	2:A:33:C:C6	2.38	0.58
2:A:833:A:H2'	2:A:834:G:C8	2.39	0.58
2:A:1454:C:C5	14:O:64:ARG:HG2	2.39	0.58
2:A:2365:G:N7	30:e:39:LYS:NZ	2.47	0.58
3:B:49:C:H2'	3:B:50:A:C8	2.38	0.58
2:A:1297:C:OP1	2:A:2710:C:H4'	2.03	0.58
2:A:1469:A:H2'	2:A:1470:A:C8	2.38	0.58
2:A:1495:A:H2	2:A:1578:U:H1'	1.69	0.58
2:A:2508:G:O6	2:A:2580:U:O4	2.21	0.58
17:R:24:TYR:HB3	17:R:28:ARG:HB2	1.86	0.58
2:A:832:U:H2'	2:A:833:A:H8	1.69	0.58
2:A:2652:C:H2'	2:A:2653:U:O4'	2.04	0.58
9:H:9:VAL:CG1	9:H:12:LEU:HB2	2.34	0.58
11:L:41:ILE:HD11	11:L:86:LEU:CD2	2.34	0.58
13:N:50:ARG:HD3	13:N:65:ILE:HD11	1.86	0.58
18:S:1:MET:HG3	18:S:43:ASN:OD1	2.03	0.58
22:W:70:ILE:HG22	22:W:72:VAL:HG13	1.86	0.58
2:A:611:C:H2'	2:A:612:G:O4'	2.04	0.57
2:A:1270:C:H5''	2:A:1271:G:C5'	2.34	0.57
2:A:1874:C:H2'	2:A:1875:G:O4'	2.03	0.57
8:G:33:LEU:HD12	8:G:79:VAL:HG13	1.85	0.57
10:K:7:LYS:O	10:K:11:VAL:HG13	2.03	0.57
12:M:19:LEU:HD23	12:M:27:LEU:HB3	1.86	0.57
17:R:9:ILE:O	17:R:13:ARG:HG3	2.04	0.57
2:A:227:A:C2	2:A:2407:A:H1'	2.38	0.57
2:A:778:G:H5''	4:C:48:ARG:HD2	1.87	0.57
2:A:912:C:O2'	2:A:913:U:H5'	2.03	0.57
2:A:1854:A:H4'	2:A:2233:U:H4'	1.84	0.57
2:A:2094:A:O2'	2:A:2095:A:H5'	2.05	0.57
2:A:2233:U:H2'	2:A:2234:G:C8	2.39	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:W:4:ILE:HB	22:W:63:ILE:CD1	2.34	0.57
22:W:50:MET:HE2	22:W:56:PHE:HE1	1.70	0.57
2:A:2375:G:N2	2:A:2378:A:OP2	2.35	0.57
3:B:37:C:O2	15:P:100:HIS:NE2	2.28	0.57
2:A:590:A:H2'	2:A:591:U:C6	2.38	0.57
2:A:1292:G:H2'	2:A:1293:C:C6	2.39	0.57
2:A:2271:G:OP1	23:X:18:ALA:HB1	2.05	0.57
13:N:36:VAL:HG12	22:W:82:TYR:HB2	1.86	0.57
13:N:53:MET:HE3	13:N:117:PHE:HE1	1.69	0.57
22:W:77:VAL:HG21	22:W:86:LEU:HD22	1.86	0.57
31:f:2:LYS:HG3	31:f:4:ARG:HG3	1.86	0.57
2:A:514:A:N3	2:A:581:C:O2'	2.35	0.57
2:A:1856:U:H2'	2:A:1857:G:O4'	2.04	0.57
2:A:2556:C:H2'	2:A:2557:G:O4'	2.03	0.57
3:B:38:C:H2'	3:B:39:A:C8	2.39	0.57
3:B:40:U:N3	3:B:44:G:OP2	2.28	0.57
2:A:222:A:N6	2:A:232:G:H1'	2.19	0.57
2:A:399:U:O4	24:Y:54:LYS:NZ	2.38	0.57
2:A:1086:A:O2'	2:A:1087:G:N7	2.37	0.57
2:A:1486:U:H2'	2:A:1487:U:H6	1.69	0.57
2:A:1794:A:H2'	2:A:1795:C:H6	1.70	0.57
2:A:2011:U:H2'	2:A:2012:G:O4'	2.04	0.57
2:A:2282:G:H4'	2:A:2389:G:O2'	2.03	0.57
2:A:2311:A:N3	7:F:85:ILE:HD11	2.18	0.57
2:A:2324:U:H3'	2:A:2325:G:H5''	1.86	0.57
3:B:104:A:H2'	3:B:105:G:O4'	2.05	0.57
5:D:31:ALA:HB1	5:D:95:SER:OG	2.05	0.57
21:V:18:ASP:OD2	21:V:40:ASN:N	2.33	0.57
2:A:305:C:H2'	2:A:306:U:C6	2.39	0.57
2:A:355:U:H2'	2:A:356:G:H8	1.70	0.57
2:A:1149:G:H2'	2:A:1150:C:H6	1.69	0.57
2:A:1638:C:H1'	2:A:2698:U:O2'	2.05	0.57
2:A:2020:A:H5'	27:b:9:THR:HG21	1.87	0.57
2:A:2066:C:O2'	2:A:2067:G:H5'	2.05	0.57
7:F:103:LEU:HD12	7:F:107:ALA:CB	2.35	0.57
30:e:6:THR:HG22	30:e:63:PRO:HD2	1.87	0.57
31:f:16:ILE:HG12	31:f:25:VAL:HG22	1.86	0.57
2:A:77:G:H2'	2:A:78:U:C6	2.40	0.57
2:A:355:U:H2'	2:A:356:G:C8	2.39	0.57
2:A:492:A:H2'	2:A:493:G:O4'	2.05	0.57
2:A:848:C:H2'	2:A:849:A:H8	1.68	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:1794:A:H2'	2:A:1795:C:C6	2.40	0.57
2:A:1954:G:O2'	2:A:1956:U:O4	2.08	0.57
2:A:2196:C:H2'	2:A:2197:U:C6	2.40	0.57
2:A:2305:U:C2	7:F:151:GLY:HA3	2.39	0.57
2:A:2314:A:OP1	7:F:88:LYS:NZ	2.38	0.57
2:A:2537:U:H2'	2:A:2538:C:C6	2.40	0.57
9:H:30:LEU:HB3	9:H:36:ALA:HB3	1.87	0.57
2:A:645:C:H2'	2:A:647:G:N7	2.20	0.57
2:A:1051:G:H1	2:A:1108:U:H3	1.53	0.57
2:A:1361:G:H2'	2:A:1362:C:C6	2.39	0.57
2:A:1394:U:H2'	2:A:1395:A:O4'	2.05	0.57
2:A:1505:A:H2'	2:A:1506:U:O4'	2.04	0.57
2:A:2329:U:H2'	2:A:2330:G:C8	2.39	0.57
2:A:357:C:H2'	2:A:358:U:C6	2.40	0.57
2:A:445:C:H2'	2:A:446:G:O4'	2.04	0.57
2:A:522:A:H2'	2:A:523:C:C6	2.40	0.57
2:A:1353:A:H2'	2:A:1354:A:H8	1.70	0.57
2:A:1437:C:O2'	2:A:1516:G:O2'	2.21	0.57
2:A:1684:G:H2'	2:A:1685:C:C6	2.40	0.57
2:A:2516:A:O2'	2:A:2517:C:H5'	2.05	0.57
2:A:372:G:H5''	24:Y:61:LYS:HD3	1.87	0.56
2:A:402:A:H2'	2:A:403:U:H5'	1.87	0.56
2:A:586:A:N1	2:A:809:G:O2'	2.36	0.56
2:A:1069:A:H4'	2:A:1070:A:H5''	1.87	0.56
2:A:1649:G:O2'	14:O:106:ASP:OD2	2.14	0.56
2:A:2464:G:H2'	2:A:2465:C:H6	1.69	0.56
2:A:2636:C:H2'	2:A:2637:U:C6	2.40	0.56
18:S:24:LYS:HA	18:S:94:THR:CG2	2.35	0.56
2:A:170:U:H2'	2:A:171:U:H6	1.70	0.56
2:A:687:C:H1'	29:d:4:THR:HG22	1.86	0.56
2:A:1932:A:H2'	2:A:1933:G:O4'	2.05	0.56
3:B:42:C:C5	7:F:66:LEU:HD22	2.39	0.56
24:Y:7:VAL:HG23	24:Y:51:VAL:HG12	1.87	0.56
26:a:48:ILE:HG21	26:a:57:VAL:HG21	1.86	0.56
2:A:471:A:H2'	2:A:472:A:O4'	2.05	0.56
2:A:851:C:OP1	26:a:19:LYS:HE2	2.06	0.56
2:A:880:G:H22	2:A:897:C:H42	1.52	0.56
2:A:1442:U:H2'	2:A:1443:U:C6	2.40	0.56
2:A:1447:C:H2'	2:A:1448:G:H8	1.70	0.56
2:A:1521:G:OP2	2:A:1522:A:O2'	2.11	0.56
2:A:1710:G:H4'	2:A:2858:C:O2	2.05	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:2038:G:H2'	2:A:2039:U:O4'	2.05	0.56
2:A:2295:C:O2'	2:A:2296:U:H5'	2.05	0.56
2:A:2419:U:H5''	28:c:22:THR:HG21	1.87	0.56
2:A:2581:G:OP2	2:A:2581:G:N2	2.23	0.56
2:A:2850:A:N7	2:A:2868:A:O2'	2.28	0.56
3:B:55:U:O3'	7:F:24:SER:OG	2.22	0.56
7:F:29:PRO:HB2	7:F:169:LEU:CD1	2.35	0.56
20:U:4:GLU:OE1	20:U:49:LYS:HE2	2.05	0.56
2:A:248:G:O5'	2:A:249:C:H5''	2.06	0.56
2:A:580:U:H2'	2:A:581:C:C6	2.40	0.56
2:A:1056:G:O2'	2:A:1086:A:H1'	2.06	0.56
2:A:1079:C:H2'	2:A:1080:A:C8	2.40	0.56
2:A:1710:G:H2'	2:A:1711:A:C8	2.40	0.56
2:A:2300:C:H2'	2:A:2301:C:C6	2.41	0.56
2:A:2618:G:H21	5:D:155:VAL:HG21	1.71	0.56
8:G:72:LEU:O	8:G:76:VAL:HG13	2.06	0.56
2:A:1353:A:H2'	2:A:1354:A:C8	2.40	0.56
7:F:38:MET:HE1	7:F:150:ARG:HD2	1.88	0.56
8:G:137:ASP:OD1	8:G:140:VAL:HG23	2.05	0.56
2:A:360:U:H2'	2:A:361:G:O4'	2.06	0.56
2:A:697:G:H2'	2:A:698:C:C6	2.40	0.56
3:B:49:C:H2'	3:B:50:A:H8	1.70	0.56
2:A:322:A:H5'	2:A:340:A:H1'	1.88	0.56
2:A:696:G:H1	2:A:766:U:H3	1.54	0.56
2:A:1319:C:O2'	2:A:1320:C:H5'	2.06	0.56
2:A:1747:U:H2'	2:A:1748:C:C6	2.41	0.56
2:A:1857:G:O2'	2:A:1884:G:N2	2.39	0.56
2:A:2533:U:H2'	2:A:2534:A:O4'	2.04	0.56
2:A:1037:G:O2'	2:A:1038:G:H5'	2.06	0.56
2:A:1281:G:H2'	2:A:1282:U:C6	2.40	0.56
2:A:1447:C:O2'	2:A:1544:A:N3	2.36	0.56
2:A:1664:A:N6	2:A:1996:C:H42	1.98	0.56
15:P:62:LEU:HD22	15:P:70:ALA:HA	1.87	0.56
16:Q:26:VAL:HG12	16:Q:86:VAL:HG22	1.88	0.56
2:A:1071:G:H2'	2:A:1072:C:C6	2.41	0.56
2:A:2257:U:O2'	2:A:2258:C:H5'	2.06	0.56
2:A:2489:U:H2'	2:A:2490:G:O4'	2.06	0.56
2:A:2845:U:H5''	16:Q:52:ASN:O	2.05	0.56
8:G:44:LYS:N	8:G:51:THR:O	2.27	0.56
10:K:4:PHE:CD2	17:R:100:VAL:HG11	2.41	0.56
14:O:30:ARG:NH2	14:O:75:ILE:HD11	2.20	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:1059:G:H2'	2:A:1060:U:C5	2.41	0.56
2:A:1625:C:O2'	2:A:1626:A:H5'	2.05	0.56
3:B:1:U:H2'	3:B:2:G:C8	2.40	0.56
2:A:12:U:O2	2:A:2626:C:H4'	2.06	0.55
2:A:760:G:H2'	2:A:761:A:O4'	2.05	0.55
2:A:833:A:H2'	2:A:834:G:H8	1.70	0.55
2:A:1286:A:H1'	2:A:1288:G:OP2	2.06	0.55
2:A:2328:A:H2'	2:A:2329:U:H6	1.69	0.55
2:A:2391:G:H3'	30:e:32:ILE:HD12	1.88	0.55
4:C:120:VAL:HG13	4:C:134:ASN:ND2	2.21	0.55
13:N:110:GLU:O	13:N:114:ARG:HG3	2.06	0.55
22:W:77:VAL:CG2	22:W:86:LEU:HD22	2.36	0.55
2:A:434:U:O2	2:A:435:C:N4	2.35	0.55
2:A:1469:A:H2'	2:A:1470:A:H8	1.71	0.55
2:A:1809:A:H2'	2:A:1810:A:C8	2.41	0.55
2:A:2795:C:H2'	2:A:2796:U:C6	2.41	0.55
15:P:34:HIS:ND1	15:P:53:THR:OG1	2.34	0.55
15:P:92:PHE:HB2	15:P:117:PHE:CE1	2.41	0.55
2:A:722:A:H2'	2:A:723:C:C6	2.42	0.55
2:A:848:C:H2'	2:A:849:A:C8	2.41	0.55
2:A:2671:G:H2'	2:A:2672:U:C6	2.42	0.55
8:G:94:TYR:HA	8:G:106:SER:O	2.06	0.55
2:A:534:U:O2'	17:R:49:ASP:OD2	2.15	0.55
2:A:590:A:H2'	2:A:591:U:H6	1.72	0.55
2:A:1443:U:H2'	2:A:1444:G:H8	1.72	0.55
2:A:1846:G:H2'	2:A:1847:A:N9	2.21	0.55
2:A:582:A:H2'	2:A:583:G:C8	2.41	0.55
2:A:654:A:O2'	2:A:655:A:H5'	2.07	0.55
2:A:2292:U:H2'	2:A:2293:G:C8	2.42	0.55
9:H:1:MET:HE3	9:H:23:ALA:HA	1.88	0.55
28:c:17:THR:HG21	28:c:43:VAL:HG23	1.89	0.55
2:A:64:A:H2'	2:A:65:U:C6	2.41	0.55
2:A:680:C:H2'	2:A:681:G:C8	2.41	0.55
2:A:1592:C:O2'	2:A:1593:A:H5'	2.07	0.55
2:A:2299:U:H2'	2:A:2300:C:C6	2.42	0.55
2:A:2830:C:O3'	5:D:56:LYS:HE3	2.07	0.55
2:A:2831:G:O2'	2:A:2884:U:OP1	2.24	0.55
5:D:52:THR:O	5:D:77:ARG:HG2	2.07	0.55
8:G:87:LEU:HD23	8:G:87:LEU:H	1.72	0.55
8:G:117:LEU:HB3	8:G:121:ILE:HG23	1.89	0.55
2:A:632:A:H2'	2:A:633:A:C8	2.41	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:1729:U:H3'	2:A:1730:C:O4'	2.07	0.55
7:F:8:TYR:HB2	7:F:173:PHE:CZ	2.42	0.55
7:F:40:VAL:HG21	7:F:50:LEU:HD23	1.89	0.55
15:P:52:SER:OG	15:P:54:VAL:HG22	2.06	0.55
2:A:356:G:H2'	2:A:357:C:C6	2.42	0.55
2:A:709:U:H2'	2:A:710:U:C6	2.41	0.55
2:A:856:G:H2'	2:A:857:G:C8	2.42	0.55
2:A:973:A:H5'	2:A:1188:U:H1'	1.88	0.55
2:A:2580:U:O2'	2:A:2581:G:H5'	2.07	0.55
8:G:99:LYS:N	8:G:102:VAL:O	2.29	0.55
20:U:13:ALA:HB1	25:Z:33:ALA:HB1	1.88	0.55
2:A:934:U:H2'	2:A:935:C:C6	2.41	0.55
2:A:2050:C:H2'	2:A:2051:A:O4'	2.07	0.55
2:A:2317:A:H2'	2:A:2318:G:O4'	2.07	0.55
2:A:2646:C:H2'	2:A:2647:U:O4'	2.06	0.55
4:C:230:HIS:CD2	4:C:247:PRO:HB3	2.42	0.55
16:Q:7:GLN:HA	16:Q:10:GLN:HE21	1.70	0.55
18:S:29:THR:OG1	18:S:64:VAL:O	2.24	0.55
30:e:33:LEU:HA	30:e:36:LYS:CD	2.37	0.55
2:A:364:C:H2'	2:A:365:U:C6	2.42	0.55
2:A:593:U:H2'	2:A:594:U:C6	2.42	0.55
2:A:947:A:H2'	2:A:948:C:C6	2.41	0.55
2:A:1652:A:H2'	2:A:1653:G:O4'	2.06	0.55
3:B:66:A:H61	3:B:107:G:H2'	1.72	0.55
7:F:33:LYS:HD3	7:F:92:ARG:NH1	2.21	0.55
7:F:119:ALA:O	7:F:167:ARG:NE	2.38	0.55
10:K:69:ARG:O	10:K:89:PHE:HB3	2.06	0.55
2:A:589:U:H2'	2:A:590:A:C8	2.41	0.54
2:A:634:C:H2'	2:A:635:C:C6	2.41	0.54
2:A:680:C:H2'	2:A:681:G:H8	1.72	0.54
2:A:813:U:H2'	2:A:814:C:H6	1.71	0.54
2:A:2364:C:H2'	2:A:2365:G:O4'	2.07	0.54
9:H:31:VAL:HB	9:H:32:PRO:HD3	1.90	0.54
25:Z:55:THR:O	25:Z:59:GLU:HG3	2.06	0.54
2:A:1858:A:H2'	2:A:1859:U:O4'	2.07	0.54
4:C:132:MET:HE1	4:C:184:VAL:CG1	2.37	0.54
2:A:959:A:H2'	2:A:960:A:C8	2.43	0.54
2:A:1901:A:OP2	4:C:253:LYS:NZ	2.33	0.54
2:A:2547:A:H2'	2:A:2548:U:H6	1.71	0.54
2:A:2709:G:OP1	14:O:18:GLN:NE2	2.40	0.54
4:C:71:LYS:HG2	4:C:74:ILE:HD12	1.88	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:F:5:HIS:HB2	7:F:97:TRP:CD1	2.43	0.54
7:F:122:PHE:CZ	7:F:170:LEU:HD12	2.42	0.54
8:G:107:LEU:HD13	8:G:152:ARG:CB	2.38	0.54
2:A:592:A:C2	30:e:4:ILE:HD11	2.42	0.54
2:A:629:G:N3	2:A:639:U:O2'	2.40	0.54
2:A:1887:C:H2'	2:A:1888:G:O4'	2.07	0.54
2:A:1890:A:H2'	2:A:1891:G:O4'	2.07	0.54
8:G:155:GLU:OE1	8:G:157:TYR:HB2	2.08	0.54
25:Z:24:GLU:O	25:Z:28:LEU:HG	2.07	0.54
2:A:29:U:H2'	2:A:30:G:C8	2.43	0.54
2:A:538:A:H2'	2:A:539:G:O4'	2.08	0.54
2:A:721:A:H2'	2:A:722:A:C8	2.43	0.54
2:A:814:C:H1'	2:A:1225:G:H21	1.72	0.54
2:A:1211:C:H3'	2:A:1212:G:C5'	2.37	0.54
2:A:1383:A:H2'	2:A:1384:A:C8	2.42	0.54
2:A:1703:G:H2'	2:A:1704:C:C6	2.42	0.54
8:G:170:ARG:HH22	31:f:29:ALA:HA	1.72	0.54
9:H:27:ARG:NH2	24:Y:60:ASP:OD2	2.39	0.54
14:O:38:LEU:HB3	14:O:39:PRO:HD3	1.89	0.54
2:A:231:A:H2'	2:A:232:G:O4'	2.07	0.54
2:A:825:A:H2'	2:A:826:U:C6	2.42	0.54
2:A:1687:G:H2'	2:A:1688:U:C6	2.43	0.54
2:A:2335:A:OP1	15:P:13:ARG:HD2	2.06	0.54
5:D:29:VAL:HB	5:D:98:VAL:CG1	2.38	0.54
8:G:92:VAL:HG12	8:G:160:LYS:HZ3	1.73	0.54
22:W:64:VAL:HA	22:W:68:LYS:O	2.08	0.54
26:a:4:THR:CG2	26:a:37:GLU:HB3	2.38	0.54
26:a:4:THR:HG21	26:a:37:GLU:HB3	1.89	0.54
2:A:542:C:H2'	2:A:543:G:C8	2.42	0.54
2:A:969:G:H2'	2:A:970:U:C6	2.42	0.54
2:A:1024:G:OP2	2:A:1025:G:O2'	2.26	0.54
2:A:1486:U:H2'	2:A:1487:U:C6	2.43	0.54
2:A:1556:C:H2'	2:A:1557:C:C6	2.43	0.54
2:A:1720:U:H2'	2:A:1721:G:O4'	2.08	0.54
5:D:1:MET:HG2	5:D:205:PRO:HG2	1.89	0.54
18:S:5:PHE:HB3	18:S:59:ILE:HD12	1.90	0.54
24:Y:17:ASN:OD1	24:Y:27:ARG:HD2	2.07	0.54
24:Y:36:HIS:CB	24:Y:56:MET:HE1	2.37	0.54
2:A:5:A:H2'	2:A:6:A:H8	1.73	0.54
2:A:145:C:H2'	2:A:146:A:H8	1.73	0.54
2:A:636:G:OP1	12:M:129:LYS:HG3	2.07	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:819:A:H5''	2:A:973:A:N1	2.23	0.54
2:A:1761:C:H2'	2:A:1762:A:O4'	2.07	0.54
2:A:1879:C:H2'	2:A:1880:U:O4'	2.08	0.54
4:C:165:VAL:HG21	4:C:181:MET:HE1	1.89	0.54
9:H:1:MET:N	9:H:21:VAL:O	2.28	0.54
12:M:89:VAL:HG21	12:M:123:ARG:CZ	2.38	0.54
27:b:38:HIS:ND1	27:b:39:LEU:O	2.40	0.54
2:A:297:G:H2'	2:A:298:G:O4'	2.08	0.54
2:A:1282:U:H2'	2:A:1283:G:O4'	2.08	0.54
2:A:2020:A:H5'	27:b:9:THR:CG2	2.38	0.54
2:A:2095:A:H2'	2:A:2096:C:C6	2.42	0.54
2:A:2324:U:H3'	2:A:2325:G:C5'	2.38	0.54
2:A:2461:A:H2'	2:A:2462:C:C6	2.43	0.54
2:A:2774:C:H2'	2:A:2775:G:O4'	2.08	0.54
2:A:2819:G:O2'	2:A:2820:A:H5''	2.08	0.54
7:F:123:ASP:OD1	7:F:127:ASN:HB2	2.07	0.54
30:e:14:PHE:C	30:e:15:LYS:HD2	2.33	0.54
2:A:5:A:H2'	2:A:6:A:C8	2.43	0.54
2:A:1419:A:O2'	2:A:1421:G:N7	2.35	0.54
2:A:1440:U:H2'	2:A:1441:G:C8	2.42	0.54
2:A:1652:A:OP1	14:O:8:ARG:HD3	2.08	0.54
2:A:2457:U:O2'	2:A:2458:G:H5'	2.08	0.54
2:A:2625:G:H2'	2:A:2626:C:C6	2.43	0.54
2:A:2646:C:OP2	2:A:2732:G:O2'	2.25	0.54
2:A:2677:G:H2'	2:A:2678:C:C6	2.43	0.54
2:A:2749:A:H5''	8:G:2:SER:HB3	1.90	0.54
10:K:73:VAL:CG1	10:K:86:GLN:HB2	2.37	0.54
11:L:63:VAL:HA	11:L:107:LEU:HD11	1.89	0.54
2:A:1292:G:H2'	2:A:1293:C:H6	1.72	0.53
2:A:1571:A:H2'	2:A:1572:A:C8	2.43	0.53
2:A:2502:G:H5''	2:A:2503:A:H5''	1.89	0.53
26:a:36:VAL:HG12	26:a:38:ARG:HG2	1.89	0.53
2:A:192:C:H2'	2:A:193:U:H5'	1.90	0.53
2:A:2332:C:OP1	23:X:77:ARG:NH2	2.41	0.53
2:A:2443:C:OP1	6:E:63:LYS:HD3	2.08	0.53
4:C:161:TYR:HB3	4:C:194:GLU:HG2	1.89	0.53
16:Q:18:PRO:HG3	16:Q:84:ILE:O	2.08	0.53
21:V:66:GLN:HB2	21:V:69:ASN:ND2	2.23	0.53
2:A:732:C:H2'	2:A:733:G:O4'	2.08	0.53
2:A:814:C:H1'	2:A:1225:G:N2	2.23	0.53
2:A:1831:G:H2'	2:A:1832:C:C6	2.44	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:22:U:H2'	3:B:23:G:C8	2.43	0.53
5:D:7:LYS:HE3	5:D:77:ARG:NH1	2.24	0.53
5:D:108:ASP:OD1	5:D:206:ALA:HA	2.08	0.53
5:D:186:LEU:HD21	16:Q:4:ILE:CG2	2.38	0.53
2:A:127:A:H5''	2:A:128:C:O4'	2.07	0.53
2:A:324:A:H2'	2:A:325:G:O4'	2.09	0.53
2:A:579:G:H2'	2:A:580:U:C6	2.43	0.53
2:A:813:U:H2'	2:A:814:C:C6	2.43	0.53
2:A:1409:U:H2'	2:A:1410:G:H8	1.73	0.53
2:A:2412:A:H2'	2:A:2413:G:O4'	2.08	0.53
2:A:2637:U:H5''	5:D:83:ARG:NH2	2.23	0.53
13:N:53:MET:HE1	13:N:103:TYR:HB3	1.91	0.53
16:Q:100:LEU:HD11	16:Q:110:ILE:HD11	1.91	0.53
20:U:53:VAL:HG13	20:U:91:GLN:HB3	1.89	0.53
27:b:29:SER:OG	27:b:40:ARG:HG2	2.07	0.53
2:A:368:A:O2'	2:A:369:U:H5'	2.09	0.53
2:A:542:C:H2'	2:A:543:G:H8	1.73	0.53
2:A:2236:U:H2'	2:A:2237:G:O4'	2.09	0.53
3:B:50:A:OP2	15:P:67:ASN:HA	2.09	0.53
3:B:53:A:H2'	3:B:54:G:O4'	2.08	0.53
4:C:144:VAL:HG21	4:C:162:VAL:HG21	1.91	0.53
16:Q:37:LYS:NZ	16:Q:39:ARG:HD3	2.23	0.53
22:W:48:MET:O	22:W:51:GLN:HG3	2.08	0.53
31:f:9:LYS:HE3	31:f:14:CYS:O	2.08	0.53
2:A:156:A:H2'	2:A:157:C:H6	1.73	0.53
2:A:479:A:C4'	2:A:480:A:H5'	2.36	0.53
2:A:713:G:H2'	2:A:714:U:C6	2.43	0.53
2:A:796:C:H2'	2:A:797:G:C8	2.42	0.53
2:A:876:C:H2'	2:A:877:A:C1'	2.39	0.53
2:A:975:A:H1'	2:A:990:A:C2	2.44	0.53
3:B:30:C:H2'	3:B:31:C:H5'	1.91	0.53
4:C:62:TYR:HA	4:C:86:ASN:HD21	1.74	0.53
5:D:56:LYS:HB2	5:D:59:ARG:CB	2.37	0.53
7:F:61:SER:OG	7:F:91:LEU:HD21	2.08	0.53
8:G:118:PRO:O	8:G:121:ILE:HG22	2.08	0.53
17:R:105:ALA:HB1	18:S:40:MET:CE	2.38	0.53
25:Z:46:VAL:O	25:Z:50:VAL:HG13	2.08	0.53
2:A:581:C:H2'	2:A:582:A:H8	1.73	0.53
2:A:1311:G:OP2	2:A:1311:G:N2	2.24	0.53
2:A:1386:C:H2'	2:A:1387:A:H8	1.73	0.53
2:A:2095:A:H2'	2:A:2096:C:H6	1.74	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:2329:U:H2'	2:A:2330:G:H8	1.74	0.53
3:B:7:G:H1'	15:P:38:GLN:HE22	1.74	0.53
10:K:8:PRO:O	10:K:11:VAL:HG22	2.09	0.53
14:O:33:ILE:HG13	14:O:114:GLU:HB3	1.91	0.53
23:X:47:ALA:HB1	23:X:51:VAL:O	2.09	0.53
25:Z:6:LEU:HB2	25:Z:56:LEU:HD12	1.91	0.53
2:A:559:G:N2	17:R:49:ASP:OD1	2.41	0.53
2:A:609:A:H2'	2:A:610:C:O4'	2.09	0.53
2:A:955:U:OP1	13:N:86:LYS:NZ	2.40	0.53
2:A:1094:U:H1'	2:A:1097:U:C5	2.42	0.53
2:A:1440:U:H2'	2:A:1441:G:H8	1.74	0.53
2:A:1625:C:H2'	2:A:1626:A:O4'	2.08	0.53
2:A:2468:A:OP2	2:A:2476:A:N6	2.41	0.53
9:H:2:GLN:NE2	9:H:20:ASN:OD1	2.41	0.53
30:e:42:ARG:HG3	30:e:45:ARG:NH2	2.24	0.53
31:f:2:LYS:HE3	31:f:4:ARG:HG2	1.91	0.53
2:A:482:A:H4'	21:V:45:HIS:ND1	2.24	0.53
2:A:572:A:H5''	2:A:573:U:OP2	2.09	0.53
2:A:1070:A:H2'	2:A:1097:U:OP1	2.09	0.53
2:A:1212:G:H1'	2:A:1237:A:N6	2.23	0.53
2:A:1482:G:H1'	2:A:1509:A:N1	2.23	0.53
2:A:1589:U:H2'	2:A:1590:A:C8	2.44	0.53
2:A:1914:C:H2'	2:A:1915:U:C6	2.44	0.53
2:A:1958:C:O2'	2:A:1959:G:H5'	2.09	0.53
2:A:2586:U:H2'	2:A:2587:A:O4'	2.09	0.53
2:A:244:A:OP2	30:e:8:ARG:NH2	2.41	0.53
2:A:1084:A:N3	2:A:1105:U:O2'	2.40	0.53
2:A:1447:C:H2'	2:A:1448:G:C8	2.44	0.53
2:A:1483:G:H2'	2:A:1484:U:C6	2.43	0.53
2:A:1684:G:H2'	2:A:1685:C:H6	1.73	0.53
2:A:1870:C:H2'	2:A:1871:A:C5	2.44	0.53
2:A:2199:A:N1	2:A:2226:C:N4	2.56	0.53
7:F:50:LEU:CD1	7:F:84:PRO:HB2	2.38	0.53
2:A:191:A:H2'	2:A:192:C:C6	2.44	0.52
2:A:283:G:H2'	2:A:284:U:O4'	2.09	0.52
2:A:483:A:O4'	21:V:45:HIS:HB3	2.08	0.52
2:A:682:G:H5'	29:d:26:ASN:CG	2.34	0.52
2:A:1526:C:H2'	2:A:1527:G:O4'	2.09	0.52
2:A:1645:G:H5''	2:A:1646:C:C5'	2.35	0.52
2:A:2081:U:H2'	2:A:2082:A:H8	1.74	0.52
7:F:110:ARG:HB2	7:F:137:ILE:CG2	2.37	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:P:49:VAL:HG11	15:P:81:ARG:HB2	1.90	0.52
2:A:742:A:H2'	2:A:743:A:H8	1.73	0.52
2:A:2246:G:H2'	2:A:2247:A:C8	2.45	0.52
2:A:2281:A:O2'	2:A:2282:G:H5'	2.07	0.52
2:A:2395:C:H2'	2:A:2396:G:O4'	2.09	0.52
4:C:72:ASP:OD1	4:C:189:ARG:NH1	2.42	0.52
28:c:5:ILE:O	28:c:28:ARG:NH2	2.29	0.52
2:A:668:A:H2'	2:A:670:A:H62	1.74	0.52
2:A:871:U:H2'	2:A:872:U:C6	2.44	0.52
2:A:905:A:H2'	2:A:906:U:H5'	1.90	0.52
2:A:1370:C:H2'	2:A:1371:G:O4'	2.09	0.52
2:A:1570:A:H2'	2:A:1571:A:C8	2.43	0.52
2:A:2098:U:H2'	2:A:2099:U:O4'	2.09	0.52
2:A:2655:G:O2'	2:A:2664:G:O6	2.25	0.52
6:E:6:LYS:HE2	6:E:141:MET:HB3	1.90	0.52
11:L:107:LEU:O	11:L:116:ILE:HD11	2.09	0.52
2:A:773:U:O2'	4:C:48:ARG:HD3	2.09	0.52
2:A:1275:A:N1	2:A:1295:C:O2'	2.36	0.52
2:A:1790:C:O2'	4:C:208:ALA:HB2	2.08	0.52
2:A:2305:U:H2'	2:A:2306:C:C6	2.44	0.52
2:A:2464:G:H2'	2:A:2465:C:C6	2.44	0.52
5:D:19:GLY:HA2	16:Q:79:PRO:HG2	1.91	0.52
10:K:6:ALA:O	10:K:48:VAL:HG21	2.09	0.52
20:U:72:GLN:HG3	20:U:73:ARG:NH1	2.23	0.52
2:A:2377:A:H2'	2:A:2378:A:C8	2.44	0.52
2:A:2853:C:H2'	2:A:2854:G:H8	1.75	0.52
3:B:48:U:H2'	3:B:49:C:H6	1.75	0.52
8:G:89:LEU:HD23	8:G:94:TYR:HB3	1.90	0.52
20:U:38:ALA:HA	20:U:42:GLU:OE2	2.08	0.52
2:A:597:G:H2'	2:A:598:U:C6	2.45	0.52
2:A:704:G:H1'	2:A:727:A:N6	2.24	0.52
2:A:2074:U:H2'	2:A:2075:U:H6	1.74	0.52
2:A:2078:C:O2'	2:A:2079:U:H5'	2.10	0.52
6:E:60:TRP:CD1	6:E:65:THR:HG21	2.45	0.52
8:G:52:PHE:CZ	8:G:72:LEU:HD12	2.45	0.52
2:A:639:U:H2'	2:A:640:C:C6	2.44	0.52
2:A:741:U:H2'	2:A:742:A:C8	2.45	0.52
2:A:784:G:H5'	2:A:785:G:OP1	2.09	0.52
2:A:902:C:O2'	2:A:903:C:H5'	2.10	0.52
2:A:1000:A:H2'	2:A:1001:A:C8	2.44	0.52
2:A:1046:A:H3'	2:A:1047:G:C5'	2.39	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:51:G:H5''	15:P:64:TYR:CD2	2.45	0.52
7:F:36:LEU:HB2	7:F:89:VAL:HG12	1.92	0.52
8:G:17:VAL:HG12	8:G:26:ILE:CG1	2.40	0.52
19:T:3:THR:OG1	19:T:62:ASP:OD2	2.28	0.52
2:A:64:A:O2'	20:U:75:GLY:HA3	2.09	0.52
2:A:272:A:H2'	2:A:273:G:C8	2.45	0.52
2:A:457:A:N1	2:A:470:A:H5''	2.24	0.52
2:A:554:U:O2'	2:A:555:G:H5'	2.10	0.52
2:A:755:U:H2'	2:A:756:A:C8	2.45	0.52
2:A:1166:G:H2'	2:A:1167:C:C6	2.45	0.52
2:A:2241:A:H2'	2:A:2242:G:C8	2.45	0.52
2:A:2427:C:C5'	2:A:2429:G:H5'	2.40	0.52
2:A:2632:A:H2'	2:A:2633:G:C8	2.45	0.52
8:G:140:VAL:O	8:G:144:VAL:HG23	2.10	0.52
24:Y:17:ASN:ND2	24:Y:27:ARG:HB3	2.25	0.52
2:A:633:A:O2'	2:A:2404:U:OP1	2.25	0.52
2:A:2362:C:OP1	30:e:40:ARG:NE	2.43	0.52
4:C:35:GLU:HG3	4:C:64:ILE:HD11	1.92	0.52
7:F:56:ASP:OD2	7:F:135:GLN:HG3	2.10	0.52
19:T:82:MET:HG3	19:T:98:LYS:HB2	1.92	0.52
2:A:350:G:H2'	2:A:351:C:O4'	2.10	0.52
2:A:1829:A:H3'	2:A:1830:C:H6	1.75	0.52
2:A:2291:U:O2'	2:A:2374:C:O2	2.28	0.52
2:A:2405:G:O2'	2:A:2411:A:N6	2.43	0.52
17:R:109:LEU:HD23	18:S:49:ILE:HD13	1.92	0.52
22:W:14:LYS:O	22:W:18:ARG:HG3	2.10	0.52
26:a:24:LEU:HD11	26:a:54:MET:SD	2.50	0.52
2:A:414:C:H2'	2:A:415:A:C8	2.45	0.51
2:A:511:U:H2'	2:A:512:G:H5'	1.91	0.51
2:A:551:G:O2'	2:A:552:U:H5'	2.11	0.51
2:A:818:G:N1	2:A:1188:U:OP2	2.28	0.51
2:A:1340:U:OP1	20:U:19:LYS:NZ	2.38	0.51
2:A:1571:A:H2'	2:A:1572:A:H8	1.73	0.51
2:A:2321:U:H3'	2:A:2322:A:C5'	2.41	0.51
2:A:2364:C:H4'	23:X:56:ASP:OD1	2.10	0.51
2:A:479:A:H1'	2:A:481:G:H5''	1.92	0.51
2:A:581:C:H2'	2:A:582:A:C8	2.44	0.51
2:A:924:G:H2'	2:A:925:A:H8	1.75	0.51
2:A:967:U:H2'	2:A:968:C:C6	2.45	0.51
2:A:1542:U:H2'	2:A:1543:G:O4'	2.10	0.51
2:A:2064:C:H2'	2:A:2065:C:H6	1.75	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:L:116:ILE:HG23	11:L:122:VAL:HG21	1.92	0.51
2:A:1081:U:H2'	2:A:1082:U:C6	2.46	0.51
2:A:1125:G:H5'	31:f:38:GLY:HA2	1.91	0.51
2:A:1316:U:H2'	2:A:1317:G:C8	2.45	0.51
2:A:1431:A:O2'	2:A:1432:G:H5'	2.11	0.51
2:A:1465:G:H2'	2:A:1466:U:O4'	2.10	0.51
2:A:1854:A:C4'	2:A:2233:U:H4'	2.40	0.51
2:A:2398:U:H2'	2:A:2399:G:H8	1.75	0.51
2:A:2400:G:O2'	2:A:2401:U:H5'	2.11	0.51
5:D:180:VAL:HG22	5:D:187:LEU:CD1	2.40	0.51
7:F:105:THR:CG2	7:F:106:ILE:HD12	2.39	0.51
11:L:19:VAL:CB	11:L:41:ILE:HD12	2.41	0.51
2:A:477:A:H2'	2:A:478:A:C8	2.46	0.51
2:A:532:A:H4'	2:A:533:G:C8	2.45	0.51
2:A:1482:G:H2'	2:A:1483:G:H8	1.75	0.51
2:A:1871:A:H1'	2:A:1872:A:C2	2.46	0.51
2:A:2210:U:H4'	2:A:2211:A:H5'	1.91	0.51
2:A:2290:G:H2'	2:A:2291:U:H6	1.74	0.51
2:A:2897:U:H2'	2:A:2898:U:C6	2.46	0.51
7:F:12:VAL:HG13	7:F:16:LEU:CD1	2.35	0.51
8:G:87:LEU:HD12	8:G:162:VAL:HG11	1.91	0.51
2:A:1736:U:O2'	2:A:1737:G:H5'	2.10	0.51
2:A:1922:G:H2'	2:A:1923:U:C6	2.46	0.51
2:A:2228:G:H2'	2:A:2229:U:C6	2.45	0.51
2:A:2467:C:H2'	2:A:2468:A:O4'	2.10	0.51
2:A:2619:C:O2'	2:A:2620:C:H5'	2.11	0.51
2:A:2787:C:O2'	2:A:2788:C:H5'	2.10	0.51
7:F:37:ASN:O	7:F:153:ASP:HB2	2.10	0.51
22:W:6:ALA:HB2	22:W:42:LEU:HD23	1.93	0.51
2:A:248:G:H5'	2:A:250:G:N7	2.26	0.51
2:A:638:G:H2'	2:A:639:U:C6	2.46	0.51
2:A:1018:U:O2'	2:A:1120:G:N2	2.41	0.51
2:A:2197:U:O2'	2:A:2198:A:H2'	2.11	0.51
2:A:2514:U:H2'	2:A:2515:C:C6	2.46	0.51
2:A:2819:G:H2'	2:A:2821:A:N7	2.25	0.51
3:B:28:C:H2'	3:B:29:A:C8	2.45	0.51
8:G:52:PHE:CE1	8:G:72:LEU:HD12	2.44	0.51
17:R:107:THR:O	17:R:111:GLU:HG2	2.09	0.51
2:A:468:G:N7	29:d:39:ARG:NH2	2.55	0.51
2:A:613:A:H2'	2:A:614:A:O4'	2.11	0.51
2:A:741:U:H2'	2:A:742:A:H8	1.75	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:1090:A:C2	2:A:1102:C:H1'	2.45	0.51
2:A:1845:G:O2'	2:A:1846:G:H5'	2.11	0.51
2:A:2334:U:H5'	15:P:12:THR:HB	1.92	0.51
2:A:2576:G:O2'	2:A:2579:C:OP2	2.28	0.51
2:A:2836:U:H2'	2:A:2837:A:C8	2.45	0.51
3:B:109:A:O2'	3:B:110:C:H5'	2.10	0.51
2:A:19:A:H2'	2:A:20:C:H6	1.76	0.51
2:A:644:A:H2'	2:A:645:C:O4'	2.11	0.51
2:A:1077:A:H2'	2:A:1078:U:H5'	1.93	0.51
2:A:1405:U:H2'	2:A:1406:U:H6	1.76	0.51
2:A:2467:C:O2'	2:A:2468:A:H5'	2.11	0.51
2:A:2804:U:H2'	2:A:2805:C:H6	1.73	0.51
12:M:49:GLY:O	12:M:56:PRO:HB3	2.11	0.51
2:A:19:A:H2'	2:A:20:C:C6	2.46	0.51
2:A:1520:U:H2'	2:A:1521:G:O4'	2.10	0.51
2:A:2648:G:H2'	2:A:2649:C:C6	2.46	0.51
4:C:205:LEU:HB3	4:C:210:ALA:CB	2.40	0.51
10:K:36:LEU:O	10:K:51:GLY:HA3	2.10	0.51
18:S:40:MET:HE3	18:S:47:VAL:CG1	2.41	0.51
2:A:145:C:H2'	2:A:146:A:C8	2.45	0.51
2:A:150:U:H2'	2:A:151:C:C6	2.46	0.51
2:A:347:A:H2'	2:A:348:A:C8	2.45	0.51
2:A:544:C:H2'	2:A:545:U:C6	2.46	0.51
2:A:1601:G:O2'	2:A:1602:U:H5'	2.11	0.51
2:A:2537:U:H2'	2:A:2538:C:H6	1.75	0.51
2:A:2578:G:N7	5:D:145:SER:HB3	2.26	0.51
3:B:79:G:N7	22:W:14:LYS:NZ	2.59	0.51
3:B:106:G:H2'	3:B:107:G:O4'	2.11	0.51
2:A:147:C:O2'	2:A:148:U:H5'	2.11	0.50
2:A:1036:G:O2'	2:A:1037:G:H5'	2.11	0.50
2:A:1038:G:H2'	2:A:1039:A:C8	2.46	0.50
2:A:1141:U:H4'	2:A:1142:A:O4'	2.11	0.50
2:A:1147:A:O2'	2:A:1148:U:H5'	2.10	0.50
2:A:1280:G:O2'	2:A:1281:G:H5'	2.11	0.50
2:A:1821:A:H2'	2:A:1822:C:C6	2.46	0.50
2:A:2419:U:P	30:e:33:LEU:HD13	2.51	0.50
2:A:2430:A:C3'	2:A:2431:U:H5'	2.41	0.50
12:M:109:LYS:HE2	12:M:128:THR:HG22	1.91	0.50
19:T:17:VAL:HA	19:T:43:ALA:HB1	1.92	0.50
2:A:552:U:H2'	2:A:553:G:H8	1.76	0.50
2:A:764:A:O2'	2:A:765:C:H5'	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:807:U:OP1	12:M:36:LYS:HD3	2.11	0.50
2:A:877:A:H5'	2:A:878:A:OP1	2.11	0.50
2:A:1145:C:O2'	2:A:1146:C:H5'	2.10	0.50
2:A:1243:C:H1'	12:M:4:ASN:O	2.11	0.50
2:A:1515:A:H2'	2:A:1516:G:O4'	2.11	0.50
2:A:1897:G:H2'	2:A:1898:U:C6	2.46	0.50
2:A:2209:G:OP2	2:A:2210:U:O2'	2.28	0.50
24:Y:13:VAL:CG2	24:Y:29:PHE:HB2	2.41	0.50
2:A:282:A:H2'	2:A:283:G:C8	2.46	0.50
2:A:314:C:O2'	2:A:315:G:H5'	2.10	0.50
2:A:1069:A:C2	2:A:1096:A:H5''	2.46	0.50
2:A:1283:G:H1'	2:A:1329:U:O2	2.10	0.50
2:A:1425:G:H2'	2:A:1426:G:C8	2.46	0.50
2:A:1812:U:H2'	2:A:1813:G:C8	2.46	0.50
2:A:2321:U:H5'	2:A:2322:A:OP2	2.11	0.50
2:A:2376:A:H2'	2:A:2377:A:O4'	2.11	0.50
2:A:2419:U:OP2	30:e:33:LEU:HD13	2.10	0.50
2:A:2520:C:O2'	2:A:2521:C:H5'	2.11	0.50
2:A:2627:G:O2'	2:A:2781:A:N1	2.39	0.50
2:A:2737:G:H2'	2:A:2738:A:C8	2.47	0.50
2:A:315:G:H2'	2:A:316:C:C6	2.46	0.50
2:A:387:U:H4'	2:A:388:G:O4'	2.11	0.50
2:A:678:C:H2'	2:A:679:C:C6	2.46	0.50
2:A:933:A:H5'	2:A:934:U:OP2	2.11	0.50
2:A:1019:U:H2'	2:A:1020:A:C8	2.46	0.50
2:A:1661:G:O2'	2:A:1662:U:H5'	2.12	0.50
2:A:1812:U:H2'	2:A:1813:G:H8	1.76	0.50
2:A:2034:U:O2'	2:A:2035:G:H5'	2.11	0.50
2:A:2394:C:N3	34:A:3001:AMP:O2'	2.43	0.50
2:A:2467:C:O2	13:N:123:LYS:NZ	2.43	0.50
11:L:108:ARG:HA	11:L:116:ILE:CD1	2.42	0.50
12:M:51:GLU:HG3	12:M:56:PRO:HA	1.93	0.50
14:O:35:LYS:HB2	14:O:112:TYR:CE1	2.46	0.50
15:P:26:LEU:HD12	15:P:38:GLN:O	2.11	0.50
22:W:50:MET:HE2	22:W:56:PHE:CE1	2.46	0.50
2:A:363:G:O2'	2:A:364:C:H5'	2.11	0.50
2:A:764:A:H5'	4:C:209:GLY:CA	2.40	0.50
2:A:1853:A:O2'	2:A:2234:G:H5'	2.12	0.50
2:A:2073:C:O2'	2:A:2074:U:H5'	2.11	0.50
2:A:2680:U:O2'	2:A:2681:C:H5'	2.11	0.50
2:A:2889:C:H2'	2:A:2890:G:O4'	2.12	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:L:103:VAL:HB	11:L:107:LEU:HD12	1.94	0.50
19:T:51:LEU:O	19:T:55:ILE:HG13	2.12	0.50
2:A:937:C:OP1	30:e:52:LYS:HE2	2.11	0.50
2:A:2480:C:H2'	2:A:2481:G:O4'	2.11	0.50
7:F:127:ASN:OD1	7:F:157:THR:HA	2.12	0.50
15:P:11:ALA:HB2	15:P:96:GLY:N	2.26	0.50
20:U:30:ILE:HD11	20:U:32:LEU:HD11	1.94	0.50
21:V:72:ILE:HG21	21:V:83:VAL:CG2	2.41	0.50
25:Z:2:LYS:HE3	25:Z:6:LEU:CD1	2.39	0.50
28:c:39:PHE:CZ	28:c:44:ARG:HA	2.47	0.50
2:A:84:A:H4'	2:A:85:G:O5'	2.12	0.50
2:A:107:G:O2'	2:A:108:G:H5'	2.11	0.50
2:A:1827:U:H5'	2:A:1971:U:H4'	1.92	0.50
2:A:1948:G:O2'	2:A:1949:G:H5'	2.12	0.50
6:E:5:LEU:HD11	6:E:12:LEU:CD2	2.38	0.50
13:N:35:ALA:HB1	13:N:126:ILE:HD12	1.92	0.50
15:P:92:PHE:HB2	15:P:117:PHE:HE1	1.77	0.50
18:S:83:TYR:OH	18:S:85:LYS:HE3	2.12	0.50
2:A:358:U:H2'	2:A:359:G:C8	2.38	0.50
2:A:445:C:O2'	2:A:446:G:H5'	2.12	0.50
2:A:665:U:O2'	2:A:666:A:H5'	2.11	0.50
2:A:987:C:H2'	2:A:988:A:O4'	2.11	0.50
2:A:1521:G:H5''	2:A:1522:A:OP2	2.12	0.50
2:A:2250:G:O2'	2:A:2496:C:OP1	2.17	0.50
6:E:191:ASP:O	6:E:195:GLN:HG3	2.11	0.50
14:O:12:ARG:O	14:O:17:ARG:NH2	2.45	0.50
15:P:6:ALA:O	15:P:10:ARG:HG3	2.12	0.50
20:U:6:ARG:O	20:U:10:VAL:HG13	2.10	0.50
1:1:75:C:H5''	1:1:76:A:OP2	2.11	0.50
2:A:796:C:H2'	2:A:797:G:H8	1.76	0.50
2:A:960:A:H5''	2:A:961:C:OP1	2.12	0.50
2:A:2086:U:H2'	2:A:2087:G:H8	1.73	0.50
2:A:2313:C:H5''	7:F:88:LYS:HD3	1.94	0.50
2:A:2476:A:H2'	2:A:2477:U:O4'	2.12	0.50
2:A:2686:G:H2'	2:A:2687:U:C6	2.47	0.50
2:A:2812:G:H2'	2:A:2813:A:O4'	2.12	0.50
11:L:14:SER:HB2	11:L:86:LEU:HD12	1.93	0.50
15:P:70:ALA:O	15:P:74:VAL:HG23	2.12	0.50
21:V:83:VAL:HG13	21:V:94:ARG:HB3	1.94	0.50
27:b:48:TYR:CE2	27:b:53:LYS:HB2	2.47	0.50
2:A:948:C:H2'	2:A:949:G:H8	1.77	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:1075:C:H2'	2:A:1076:C:C6	2.46	0.49
2:A:1316:U:H2'	2:A:1317:G:H8	1.77	0.49
2:A:1405:U:H2'	2:A:1406:U:C6	2.46	0.49
2:A:2064:C:H2'	2:A:2065:C:C6	2.46	0.49
2:A:2229:U:H2'	2:A:2230:G:H8	1.77	0.49
2:A:2466:C:O2'	2:A:2467:C:H5'	2.12	0.49
2:A:2880:C:H1'	14:O:92:GLY:O	2.11	0.49
3:B:28:C:H2'	3:B:29:A:H8	1.77	0.49
3:B:66:A:N6	3:B:107:G:H2'	2.26	0.49
7:F:132:VAL:N	7:F:152:LEU:O	2.45	0.49
22:W:1:MET:HA	22:W:1:MET:HE3	1.93	0.49
2:A:170:U:H2'	2:A:171:U:C6	2.46	0.49
2:A:364:C:O2'	2:A:365:U:H5'	2.11	0.49
2:A:1672:A:C2	2:A:2582:G:H5'	2.48	0.49
2:A:1797:G:O2'	2:A:1798:U:H5'	2.12	0.49
2:A:1825:U:H2'	2:A:1826:G:C8	2.46	0.49
22:W:2:PHE:HD2	22:W:50:MET:HE3	1.77	0.49
27:b:55:ILE:HG22	27:b:57:LYS:N	2.25	0.49
2:A:52:A:H2'	2:A:53:A:C8	2.47	0.49
2:A:417:C:H2'	2:A:418:C:C6	2.48	0.49
2:A:580:U:O3'	17:R:31:VAL:HG13	2.12	0.49
2:A:678:C:H2'	2:A:679:C:H6	1.76	0.49
2:A:1190:G:H2'	2:A:1191:G:H8	1.76	0.49
2:A:1495:A:H2'	2:A:1496:A:C8	2.47	0.49
2:A:1628:G:O2'	2:A:1629:U:H5'	2.12	0.49
2:A:1796:U:H2'	2:A:1797:G:C8	2.48	0.49
2:A:1800:C:C3'	4:C:146:MET:HE1	2.41	0.49
2:A:2474:U:H3'	2:A:2475:C:C5	2.47	0.49
8:G:9:VAL:CG2	8:G:69:ARG:HG2	2.42	0.49
26:a:51:VAL:O	26:a:55:VAL:HG22	2.11	0.49
2:A:600:G:N2	2:A:605:G:O3'	2.44	0.49
2:A:604:G:O2'	2:A:605:G:H5'	2.13	0.49
2:A:629:G:H1'	2:A:639:U:H1'	1.94	0.49
2:A:772:C:O2'	2:A:773:U:H5'	2.13	0.49
2:A:1009:A:H5'	17:R:59:GLN:CG	2.35	0.49
2:A:1018:U:O2'	2:A:1019:U:H5'	2.13	0.49
2:A:1567:G:OP1	4:C:60:GLN:NE2	2.36	0.49
2:A:2055:C:H5'	2:A:2056:G:O5'	2.12	0.49
2:A:2292:U:H2'	2:A:2293:G:H8	1.76	0.49
2:A:2333:A:H5'	2:A:2335:A:H1'	1.94	0.49
2:A:2653:U:OP2	2:A:2654:A:O2'	2.25	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:E:41:GLN:HG2	6:E:43:THR:CG2	2.39	0.49
7:F:43:ALA:O	7:F:47:LYS:HD2	2.12	0.49
12:M:92:LEU:HG	12:M:96:LYS:HE3	1.93	0.49
2:A:291:G:O2'	2:A:292:U:H5'	2.13	0.49
2:A:749:A:H4'	2:A:1271:G:N3	2.28	0.49
2:A:858:G:N3	2:A:2268:A:H2'	2.27	0.49
2:A:1239:G:H2'	2:A:1240:U:O4'	2.13	0.49
2:A:1588:G:O2'	2:A:1589:U:H5'	2.13	0.49
2:A:2266:A:H4'	2:A:2267:A:N3	2.28	0.49
14:O:117:ASP:OD2	14:O:121:LYS:NZ	2.45	0.49
26:a:9:GLN:HB2	26:a:29:LEU:HD13	1.95	0.49
2:A:726:G:H5''	2:A:1432:G:H1'	1.94	0.49
2:A:1596:A:O2'	2:A:1597:A:H5'	2.12	0.49
2:A:2470:G:O2'	2:A:2471:A:H5'	2.11	0.49
27:b:52:ARG:HB2	27:b:52:ARG:NH2	2.28	0.49
2:A:6:A:O2'	2:A:7:G:H5'	2.13	0.49
2:A:566:U:O2'	2:A:567:U:H5'	2.12	0.49
2:A:711:G:H1	2:A:720:U:H3	1.61	0.49
2:A:1118:C:H2'	2:A:1119:U:O4'	2.13	0.49
2:A:1730:C:O2	2:A:1731:G:N1	2.46	0.49
2:A:2473:U:O2	2:A:2473:U:H2'	2.13	0.49
5:D:4:LEU:HD21	5:D:96:ILE:CG2	2.43	0.49
6:E:127:GLU:HG2	6:E:133:LEU:HD13	1.93	0.49
7:F:4:LEU:HG	7:F:101:GLU:HB3	1.93	0.49
7:F:68:THR:N	7:F:86:GLY:O	2.43	0.49
24:Y:12:PRO:HB3	24:Y:30:LEU:CD2	2.41	0.49
24:Y:64:ILE:HD12	24:Y:67:VAL:CG2	2.43	0.49
31:f:2:LYS:HE3	31:f:4:ARG:CG	2.43	0.49
2:A:222:A:H61	2:A:232:G:H1'	1.77	0.49
2:A:223:A:N1	2:A:407:G:O2'	2.38	0.49
2:A:418:C:H2'	2:A:419:U:C6	2.47	0.49
2:A:420:C:O2'	2:A:421:C:H5'	2.12	0.49
2:A:949:G:O2'	2:A:950:G:H5'	2.13	0.49
2:A:1442:U:H2'	2:A:1443:U:H6	1.78	0.49
2:A:1710:G:H2'	2:A:1711:A:H8	1.76	0.49
2:A:1854:A:N6	2:A:1888:G:H1'	2.28	0.49
2:A:2467:C:OP2	31:f:4:ARG:NH2	2.46	0.49
2:A:2661:G:H2'	2:A:2662:A:O4'	2.13	0.49
6:E:118:LEU:HD12	6:E:186:VAL:HB	1.94	0.49
14:O:72:ASP:O	14:O:76:VAL:HG23	2.13	0.49
2:A:30:G:H2'	2:A:31:C:C6	2.48	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:172:A:H2'	2:A:173:A:C8	2.47	0.49
2:A:754:U:H2'	2:A:755:U:C6	2.48	0.49
2:A:1637:A:H4'	2:A:2711:A:O2'	2.12	0.49
2:A:1773:A:N7	2:A:1829:A:H1'	2.28	0.49
2:A:2334:U:C2	15:P:16:ARG:HG3	2.48	0.49
2:A:2348:U:P	30:e:38:THR:HG21	2.52	0.49
7:F:69:LYS:HA	7:F:84:PRO:HA	1.95	0.49
11:L:122:VAL:O	11:L:123:LEU:HD23	2.13	0.49
21:V:11:VAL:HG11	21:V:70:VAL:HB	1.95	0.49
2:A:729:G:P	4:C:13:ARG:HG2	2.53	0.49
2:A:1007:C:H5''	10:K:37:ARG:NH2	2.28	0.49
2:A:1946:U:O2'	2:A:1947:C:H5'	2.13	0.49
2:A:2071:A:H2'	2:A:2072:C:H6	1.78	0.49
2:A:2461:A:H2'	2:A:2462:C:H6	1.77	0.49
2:A:2471:A:H2'	2:A:2472:G:O4'	2.13	0.49
2:A:2557:G:H2'	2:A:2558:C:H6	1.78	0.49
2:A:2844:G:H2'	2:A:2845:U:O4'	2.13	0.49
2:A:2895:G:H2'	2:A:2896:C:C6	2.47	0.49
4:C:25:HIS:CG	4:C:80:ARG:HD2	2.48	0.49
4:C:115:GLN:O	4:C:125:LYS:NZ	2.38	0.49
8:G:43:VAL:O	8:G:44:LYS:HE2	2.13	0.49
19:T:73:LYS:HB2	19:T:106:VAL:HB	1.95	0.49
2:A:198:C:O2'	2:A:199:A:H5'	2.13	0.48
2:A:371:A:H1'	24:Y:61:LYS:NZ	2.28	0.48
2:A:419:U:H2'	2:A:420:C:C6	2.48	0.48
2:A:1383:A:H1'	2:A:1405:U:O2'	2.12	0.48
2:A:1680:U:H2'	2:A:1681:G:O4'	2.13	0.48
2:A:1877:A:H2'	2:A:1878:G:O4'	2.12	0.48
2:A:2376:A:O2'	15:P:111:ARG:NH2	2.43	0.48
4:C:107:PRO:HG2	4:C:110:LEU:CD2	2.43	0.48
15:P:64:TYR:HB3	15:P:67:ASN:ND2	2.28	0.48
20:U:39:THR:OG1	20:U:42:GLU:HG2	2.13	0.48
20:U:94:ASP:OD2	20:U:96:VAL:HG13	2.13	0.48
30:e:33:LEU:O	30:e:41:LYS:HD3	2.12	0.48
2:A:373:U:H4'	2:A:423:A:O2'	2.13	0.48
2:A:394:C:O2'	2:A:395:U:H5'	2.13	0.48
2:A:721:A:H2'	2:A:722:A:H8	1.77	0.48
2:A:2040:G:H2'	2:A:2041:U:O4'	2.13	0.48
2:A:2847:U:H2'	2:A:2848:G:O4'	2.13	0.48
2:A:2855:C:O2'	2:A:2856:A:H5'	2.13	0.48
5:D:113:SER:HB3	5:D:170:VAL:HG21	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:F:77:PHE:HB2	7:F:79:ILE:HG12	1.96	0.48
7:F:102:ARG:HG2	7:F:106:ILE:HD13	1.94	0.48
8:G:5:ALA:HB2	8:G:66:GLY:N	2.27	0.48
8:G:75:MET:O	8:G:79:VAL:HG22	2.13	0.48
16:Q:4:ILE:O	16:Q:7:GLN:HG3	2.13	0.48
20:U:33:LYS:HG2	20:U:80:TRP:CE3	2.47	0.48
2:A:74:A:H4'	2:A:75:G:O5'	2.14	0.48
2:A:402:A:C2'	2:A:403:U:H5'	2.42	0.48
2:A:637:A:H4'	2:A:638:G:O5'	2.13	0.48
2:A:1853:A:H2'	2:A:1854:A:H8	1.78	0.48
2:A:2305:U:O4	7:F:38:MET:HA	2.13	0.48
2:A:2489:U:O2'	2:A:2490:G:H5'	2.13	0.48
2:A:2641:G:OP1	10:K:78:THR:HG22	2.13	0.48
2:A:2816:G:O2'	2:A:2817:U:H5'	2.14	0.48
2:A:2818:U:H4'	2:A:2837:A:H4'	1.95	0.48
2:A:2853:C:H2'	2:A:2854:G:C8	2.48	0.48
4:C:140:THR:HG23	4:C:161:TYR:HB2	1.95	0.48
5:D:46:ARG:NH1	5:D:85:ALA:O	2.46	0.48
5:D:146:ILE:HA	5:D:159:LYS:NZ	2.28	0.48
7:F:120:LYS:O	7:F:120:LYS:HD3	2.13	0.48
9:H:15:LEU:HD12	9:H:15:LEU:O	2.13	0.48
11:L:66:LYS:HA	11:L:79:PHE:O	2.12	0.48
12:M:96:LYS:HD3	12:M:103:ILE:HD12	1.96	0.48
15:P:27:VAL:HG21	15:P:40:ILE:HD13	1.95	0.48
30:e:54:ASP:O	30:e:58:VAL:HG23	2.13	0.48
2:A:150:U:H2'	2:A:151:C:H6	1.79	0.48
2:A:282:A:H2'	2:A:283:G:H8	1.79	0.48
2:A:640:C:O2'	2:A:641:U:H5'	2.13	0.48
2:A:1435:G:H2'	2:A:1436:G:C8	2.48	0.48
2:A:2016:U:H1'	27:b:3:VAL:HG21	1.95	0.48
2:A:2559:C:O2'	2:A:2560:A:H5'	2.13	0.48
7:F:69:LYS:CE	7:F:84:PRO:HG3	2.39	0.48
10:K:76:HIS:O	10:K:84:ILE:HD12	2.13	0.48
16:Q:31:TRP:CE3	16:Q:38:LYS:HE2	2.49	0.48
2:A:348:A:H2'	2:A:349:U:O4'	2.14	0.48
2:A:794:A:H2'	2:A:795:C:C6	2.48	0.48
2:A:2418:A:OP1	30:e:45:ARG:HD3	2.13	0.48
2:A:2759:G:H1'	8:G:35:ARG:NH2	2.28	0.48
7:F:108:VAL:HG11	7:F:176:PRO:CG	2.32	0.48
7:F:108:VAL:HB	7:F:109:PRO:HD3	1.96	0.48
7:F:122:PHE:CE2	7:F:128:TYR:HB2	2.49	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:G:9:VAL:HB	8:G:50:LEU:HB2	1.95	0.48
22:W:48:MET:HE2	22:W:84:PRO:O	2.14	0.48
25:Z:47:ARG:O	25:Z:50:VAL:HG22	2.13	0.48
2:A:599:A:O2'	2:A:600:G:H5'	2.13	0.48
2:A:696:G:O2'	2:A:697:G:H5'	2.13	0.48
2:A:831:G:H2'	2:A:832:U:O4'	2.13	0.48
2:A:843:G:H2'	2:A:844:A:C8	2.49	0.48
2:A:963:U:O2'	2:A:964:C:H5'	2.14	0.48
2:A:1625:C:C2'	2:A:1626:A:H5'	2.43	0.48
2:A:2430:A:N3	2:A:2430:A:H2'	2.28	0.48
2:A:2547:A:H4'	11:L:29:HIS:NE2	2.29	0.48
4:C:43:ARG:HA	4:C:48:ARG:O	2.13	0.48
6:E:194:LYS:O	6:E:197:GLU:HG2	2.14	0.48
12:M:64:PHE:CZ	30:e:15:LYS:HD3	2.49	0.48
14:O:9:GLN:O	14:O:17:ARG:HD3	2.13	0.48
25:Z:50:VAL:O	25:Z:54:LYS:HG3	2.14	0.48
30:e:6:THR:HG22	30:e:63:PRO:HG2	1.95	0.48
2:A:615:U:O4	6:E:39:ALA:HB2	2.14	0.48
2:A:1153:C:H2'	2:A:1154:G:O4'	2.14	0.48
2:A:1321:A:H5'	2:A:1322:A:OP2	2.14	0.48
2:A:1548:A:H2'	2:A:1549:A:C8	2.48	0.48
2:A:1591:A:H2'	2:A:1592:C:C6	2.49	0.48
2:A:2312:U:H5'	7:F:85:ILE:HG21	1.95	0.48
3:B:11:C:OP1	23:X:72:LYS:HE3	2.14	0.48
5:D:122:VAL:HG21	5:D:129:THR:HG22	1.95	0.48
13:N:47:GLU:OE2	13:N:50:ARG:NH1	2.32	0.48
18:S:61:ALA:HB1	18:S:96:VAL:HB	1.95	0.48
2:A:133:U:O2'	2:A:134:G:H5'	2.13	0.48
2:A:143:C:H2'	2:A:144:A:H8	1.78	0.48
2:A:275:C:H3'	2:A:276:U:H5''	1.95	0.48
2:A:287:G:H2'	2:A:288:U:H6	1.77	0.48
2:A:523:C:O2'	2:A:524:G:H5'	2.14	0.48
2:A:672:C:O2'	2:A:673:C:H5'	2.13	0.48
2:A:784:G:H4'	2:A:785:G:O5'	2.14	0.48
2:A:918:A:H5''	3:B:97:C:O2'	2.14	0.48
2:A:1668:A:O2'	2:A:1674:G:N7	2.39	0.48
2:A:1745:A:O2'	2:A:1746:A:H5'	2.14	0.48
2:A:2241:A:H2'	2:A:2242:G:H8	1.79	0.48
2:A:2615:U:O4	27:b:2:ALA:HB1	2.13	0.48
2:A:2632:A:H2'	2:A:2633:G:H8	1.77	0.48
2:A:2857:G:N2	2:A:2860:A:OP2	2.28	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:2888:C:H2'	2:A:2889:C:C6	2.48	0.48
3:B:85:G:O2'	3:B:86:G:H5'	2.13	0.48
7:F:122:PHE:HZ	7:F:170:LEU:HD12	1.78	0.48
14:O:72:ASP:HB3	14:O:75:ILE:HD13	1.95	0.48
22:W:20:LEU:HB3	22:W:25:LYS:O	2.13	0.48
2:A:77:G:H2'	2:A:78:U:H6	1.78	0.48
2:A:775:G:C4'	2:A:776:G:H5'	2.30	0.48
2:A:876:C:H2'	2:A:877:A:H1'	1.95	0.48
2:A:921:C:O2'	2:A:922:C:H5'	2.13	0.48
2:A:1009:A:O2'	2:A:1010:A:O4'	2.29	0.48
2:A:1697:G:OP2	2:A:1698:A:O2'	2.22	0.48
2:A:2051:A:H4'	5:D:146:ILE:HG12	1.96	0.48
2:A:2427:C:H5'	2:A:2429:G:H5'	1.94	0.48
2:A:2570:G:H2'	2:A:2571:U:O4'	2.14	0.48
2:A:2602[B]:A:H3'	2:A:2603[B]:G:C5'	2.44	0.48
3:B:13:G:O2'	3:B:15:A:H5''	2.14	0.48
7:F:38:MET:CE	7:F:150:ARG:HD2	2.43	0.48
11:L:41:ILE:HD11	11:L:86:LEU:HD21	1.94	0.48
11:L:79:PHE:HB3	16:Q:68:GLU:OE2	2.14	0.48
18:S:74:ILE:HB	18:S:87:GLN:HG2	1.95	0.48
21:V:83:VAL:CG1	21:V:94:ARG:HB3	2.44	0.48
2:A:132:G:O2'	2:A:133:U:H5'	2.14	0.48
2:A:729:G:C8	4:C:207:LYS:HD2	2.49	0.48
2:A:857:G:H2'	2:A:858:G:O4'	2.14	0.48
2:A:1028:A:N6	2:A:1125:G:H2'	2.29	0.48
2:A:1993:U:H4'	5:D:133:THR:OG1	2.14	0.48
2:A:2221:G:O2'	2:A:2222:C:H5'	2.13	0.48
2:A:2299:U:H2'	2:A:2300:C:H6	1.77	0.48
2:A:2521:C:O2'	2:A:2522:U:H5'	2.14	0.48
2:A:2563:U:H2'	2:A:2565:A:OP2	2.14	0.48
6:E:31:VAL:HG21	6:E:104:ALA:CB	2.44	0.48
11:L:34:GLY:O	11:L:62:VAL:HB	2.14	0.48
17:R:106:PHE:O	17:R:110:VAL:HG23	2.14	0.48
19:T:1:MET:HE3	19:T:62:ASP:HB3	1.96	0.48
19:T:7:HIS:HB2	19:T:50:VAL:HG22	1.95	0.48
21:V:26:LYS:HD2	21:V:37:GLU:HG3	1.94	0.48
23:X:36:ILE:HD13	23:X:39:ARG:HH11	1.77	0.48
2:A:152:A:H2'	2:A:153:U:C6	2.48	0.47
2:A:221:A:C4	2:A:233:A:H1'	2.49	0.47
2:A:292:U:O2'	2:A:293:U:H5'	2.14	0.47
2:A:657:U:H2'	2:A:658:U:C6	2.49	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:727:A:O2'	2:A:728:G:H5'	2.13	0.47
2:A:1160:G:O2'	2:A:1161:C:H5'	2.14	0.47
2:A:1199:U:H1'	17:R:4:VAL:HG22	1.95	0.47
2:A:1255:U:O5'	2:A:1256:G:H5''	2.13	0.47
2:A:1409:U:H2'	2:A:1410:G:C8	2.49	0.47
2:A:1777:U:O2'	2:A:1778:U:H5'	2.13	0.47
2:A:1798:U:OP2	4:C:271:ARG:NH1	2.36	0.47
2:A:2283:C:O5'	2:A:2389:G:O2'	2.32	0.47
6:E:7:ASP:OD2	6:E:7:ASP:N	2.44	0.47
2:A:27:G:O2'	2:A:28:A:OP2	2.25	0.47
2:A:194:G:H2'	2:A:195:A:O4'	2.13	0.47
2:A:793:A:OP2	2:A:2071:A:O2'	2.30	0.47
2:A:812:C:H5''	2:A:1250:G:O2'	2.13	0.47
2:A:828:U:H4'	2:A:831:G:N1	2.29	0.47
2:A:1198:U:H2'	2:A:1199:U:C6	2.49	0.47
2:A:1441:G:H2'	2:A:1442:U:C6	2.49	0.47
2:A:1443:U:H2'	2:A:1444:G:C8	2.48	0.47
2:A:1753:G:N2	2:A:1755:A:H3'	2.29	0.47
2:A:1881:C:H2'	2:A:1882:U:O4'	2.14	0.47
2:A:2352:A:H2'	2:A:2353:G:H5'	1.96	0.47
2:A:2760:C:O2'	2:A:2761:A:H5'	2.13	0.47
3:B:114:C:H2'	3:B:115:A:H8	1.79	0.47
4:C:66:ASP:OD2	4:C:102:ARG:HD3	2.13	0.47
11:L:112:PHE:O	11:L:116:ILE:HD12	2.14	0.47
15:P:62:LEU:HD13	15:P:64:TYR:O	2.13	0.47
17:R:78:LYS:O	17:R:117:LEU:HD11	2.14	0.47
2:A:78:U:H2'	2:A:79:C:H6	1.79	0.47
2:A:498:G:O2'	2:A:499:U:H5'	2.14	0.47
2:A:521:U:H2'	2:A:522:A:H8	1.79	0.47
2:A:662:G:O2'	2:A:663:G:H5'	2.14	0.47
2:A:745:G:O2'	2:A:748:G:H1'	2.14	0.47
2:A:987:C:O2'	2:A:1000:A:N3	2.40	0.47
2:A:1474:U:H2'	2:A:1475:G:H5'	1.96	0.47
2:A:2016:U:H1'	27:b:3:VAL:CG2	2.44	0.47
2:A:2648:G:H2'	2:A:2649:C:H6	1.79	0.47
2:A:2698:U:H2'	2:A:2699:C:C6	2.50	0.47
2:A:2785:C:H2'	2:A:2786:U:C6	2.49	0.47
3:B:5:U:O2'	3:B:6:G:H5'	2.14	0.47
22:W:4:ILE:CG2	22:W:42:LEU:HD22	2.44	0.47
25:Z:4:LYS:HD3	25:Z:7:ARG:NH1	2.29	0.47
25:Z:41:HIS:O	25:Z:45:GLN:HG3	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:275:C:H3'	2:A:276:U:C5'	2.45	0.47
2:A:1164:C:H2'	2:A:1165:A:H8	1.79	0.47
2:A:1187:G:H5''	18:S:83:TYR:CE1	2.50	0.47
2:A:1939:U:OP1	2:A:2604:U:O2'	2.32	0.47
2:A:2300:C:H2'	2:A:2301:C:H6	1.77	0.47
2:A:2700:A:H2'	2:A:2701:U:H6	1.79	0.47
6:E:134:LEU:CD2	6:E:161:ALA:HB2	2.45	0.47
8:G:5:ALA:HB2	8:G:66:GLY:CA	2.44	0.47
10:K:76:HIS:CE1	10:K:85:LYS:HB2	2.49	0.47
11:L:110:GLU:OE2	11:L:111:LYS:HG3	2.14	0.47
13:N:26:VAL:HA	13:N:104:GLU:OE2	2.14	0.47
16:Q:52:ASN:O	16:Q:53:ARG:HD3	2.14	0.47
2:A:58:G:O2'	2:A:59:U:H5'	2.15	0.47
2:A:78:U:H2'	2:A:79:C:C6	2.49	0.47
2:A:929:U:O2	26:a:26:GLY:HA2	2.14	0.47
2:A:1268:A:H2'	2:A:1269:A:O4'	2.14	0.47
2:A:1399:C:O2'	2:A:1400:U:H5'	2.14	0.47
2:A:1969:A:H2'	2:A:1972:G:H21	1.78	0.47
2:A:2398:U:H2'	2:A:2399:G:C8	2.50	0.47
2:A:2508:G:HO2'	2:A:2554:U:HO2'	1.63	0.47
2:A:2679:A:H5'	5:D:170:VAL:HG11	1.97	0.47
2:A:2840:C:O2'	2:A:2841:C:H5'	2.14	0.47
2:A:2846:G:H2'	2:A:2847:U:C6	2.49	0.47
4:C:53:HIS:CE1	4:C:219:THR:HG23	2.49	0.47
4:C:72:ASP:HA	4:C:118:SER:O	2.14	0.47
7:F:56:ASP:O	7:F:60:ILE:HG13	2.14	0.47
13:N:24:THR:HG22	13:N:99:GLY:O	2.15	0.47
18:S:43:ASN:O	18:S:46:GLU:HB3	2.14	0.47
31:f:8:LYS:O	31:f:35:GLN:NE2	2.47	0.47
2:A:290:U:H2'	2:A:291:G:H8	1.80	0.47
2:A:603:A:H4'	2:A:604:G:O5'	2.15	0.47
2:A:729:G:C5	4:C:207:LYS:HB2	2.50	0.47
2:A:905:A:C2'	2:A:906:U:H5'	2.45	0.47
2:A:1591:A:H2'	2:A:1592:C:H6	1.79	0.47
2:A:2000:C:OP1	14:O:5:LYS:NZ	2.37	0.47
2:A:2099:U:H2'	2:A:2100:G:H8	1.79	0.47
2:A:2273:A:H2'	2:A:2274:A:C8	2.50	0.47
6:E:12:LEU:HD12	6:E:12:LEU:O	2.15	0.47
7:F:117:LEU:HD12	7:F:130:MET:HG3	1.95	0.47
9:H:37:VAL:CG2	9:H:38:PRO:HD2	2.44	0.47
11:L:10:VAL:HG21	11:L:16:ALA:HB3	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:V:10:GLU:O	21:V:72:ILE:HD12	2.15	0.47
2:A:289:G:H2'	2:A:290:U:C6	2.50	0.47
2:A:347:A:H2'	2:A:348:A:H8	1.80	0.47
2:A:372:G:O2'	2:A:400:G:O6	2.28	0.47
2:A:388:G:O6	2:A:390:U:O2'	2.22	0.47
2:A:479:A:H4'	2:A:480:A:C5'	2.39	0.47
2:A:896:A:H5'	2:A:897:C:OP2	2.15	0.47
2:A:995:C:O2	10:K:3:THR:OG1	2.28	0.47
2:A:1231:U:H2'	2:A:1232:G:H8	1.78	0.47
2:A:1355:G:O2'	2:A:1356:G:H5'	2.15	0.47
2:A:1435:G:H2'	2:A:1436:G:H8	1.80	0.47
2:A:2700:A:H2'	2:A:2701:U:C6	2.49	0.47
2:A:2739:U:O2'	2:A:2740:A:H5'	2.14	0.47
2:A:2821:A:OP2	5:D:115:GLY:N	2.40	0.47
7:F:8:TYR:CD1	7:F:12:VAL:HB	2.50	0.47
8:G:2:SER:O	8:G:6:LYS:HG3	2.15	0.47
8:G:47:ASP:N	8:G:47:ASP:OD1	2.48	0.47
10:K:46:PRO:HD3	17:R:60:LEU:HD13	1.97	0.47
11:L:64:ARG:HD3	11:L:102:PRO:O	2.14	0.47
22:W:4:ILE:HG21	22:W:42:LEU:HD22	1.97	0.47
23:X:36:ILE:HD13	23:X:39:ARG:NH1	2.30	0.47
24:Y:38:PHE:N	24:Y:47:VAL:O	2.29	0.47
2:A:48:G:N2	2:A:177:G:OP2	2.48	0.47
2:A:61:C:O2'	2:A:62:U:H5'	2.15	0.47
2:A:203:A:OP2	2:A:204:A:O2'	2.17	0.47
2:A:219:A:H2'	2:A:220:G:O4'	2.15	0.47
2:A:499:U:H2'	2:A:500:G:O4'	2.14	0.47
2:A:625:G:H2'	2:A:626:A:C8	2.49	0.47
2:A:971:G:H2'	2:A:972:A:O4'	2.14	0.47
2:A:2812:G:H2'	2:A:2813:A:C8	2.50	0.47
8:G:105:LEU:HD21	8:G:131:ILE:HD11	1.97	0.47
17:R:91:ASP:O	17:R:95:LEU:HG	2.14	0.47
18:S:16:GLU:OE1	18:S:100:GLY:HA2	2.15	0.47
22:W:6:ALA:HB1	22:W:40:ILE:HB	1.96	0.47
2:A:176:A:O2'	2:A:177:G:H5'	2.14	0.47
2:A:430:A:H5''	2:A:431:U:OP2	2.15	0.47
2:A:789:A:H3'	2:A:790:U:H5'	1.97	0.47
2:A:1025:G:H4'	2:A:1026:G:OP2	2.15	0.47
2:A:1213:A:H2'	2:A:1214:A:C8	2.50	0.47
2:A:1433:A:O2'	2:A:1434:A:H5'	2.15	0.47
2:A:1863:G:H2'	2:A:1864:U:C6	2.49	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:2663:G:H2'	2:A:2664:G:O4'	2.14	0.47
3:B:100:G:H2'	3:B:101:A:O4'	2.14	0.47
4:C:240:PHE:O	4:C:242:LYS:HG2	2.15	0.47
5:D:15:PHE:CD2	16:Q:55:LEU:HD11	2.49	0.47
5:D:56:LYS:O	5:D:60:VAL:HG23	2.15	0.47
12:M:51:GLU:O	12:M:54:GLN:HG2	2.15	0.47
12:M:77:ILE:HD11	12:M:101:ILE:CG2	2.45	0.47
13:N:112:LEU:O	13:N:115:GLU:HG2	2.15	0.47
2:A:15:G:H4'	27:b:18:SER:HB3	1.96	0.47
2:A:579:G:H2'	2:A:580:U:H6	1.79	0.47
2:A:744:U:H2'	2:A:745:G:O4'	2.14	0.47
7:F:102:ARG:HD2	7:F:140:GLU:OE2	2.15	0.47
8:G:96:ALA:N	8:G:128:GLN:O	2.48	0.47
14:O:24:MET:HE1	14:O:40:LYS:HD3	1.96	0.47
18:S:57:GLY:HA2	18:S:103:ALA:O	2.15	0.47
19:T:41:LYS:HE2	27:b:22:LEU:HD11	1.95	0.47
22:W:29:ILE:HD12	22:W:39:ALA:HA	1.96	0.47
25:Z:2:LYS:HG2	25:Z:3:ALA:H	1.80	0.47
2:A:61:C:O4'	25:Z:40:SER:HB3	2.15	0.46
2:A:479:A:H1'	2:A:481:G:C5'	2.45	0.46
2:A:500:G:N2	2:A:502:A:H3'	2.30	0.46
2:A:598:U:H2'	2:A:599:A:H8	1.80	0.46
2:A:671:C:H2'	2:A:672:C:C6	2.50	0.46
2:A:1916:A:H2'	2:A:1917:U:O4'	2.15	0.46
2:A:1928:A:H2'	2:A:1929:G:O4'	2.16	0.46
2:A:2233:U:H2'	2:A:2234:G:H8	1.76	0.46
2:A:2849:U:H4'	2:A:2868:A:C2	2.50	0.46
3:B:33:G:H2'	3:B:34:A:O4'	2.15	0.46
8:G:9:VAL:HG23	8:G:69:ARG:HG2	1.97	0.46
19:T:23:LEU:HD21	27:b:22:LEU:O	2.15	0.46
21:V:99:ASN:ND2	21:V:101:GLU:OE2	2.47	0.46
26:a:44:ILE:HD13	26:a:47:MET:HE2	1.97	0.46
2:A:653:U:H4'	2:A:654:A:O5'	2.15	0.46
2:A:706:A:H61	2:A:725:G:H1'	1.79	0.46
2:A:1457:U:H5''	2:A:1458:U:OP1	2.15	0.46
2:A:1501:G:O2'	2:A:1502:A:H5'	2.16	0.46
2:A:2246:G:H2'	2:A:2247:A:H8	1.79	0.46
2:A:2651:C:O2'	2:A:2652:C:H5'	2.15	0.46
17:R:52:GLN:O	17:R:56:GLN:HG3	2.15	0.46
2:A:29:U:H2'	2:A:30:G:H8	1.81	0.46
2:A:118:A:OP2	2:A:119:A:H2'	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:197:A:H4'	2:A:2069:G:OP2	2.15	0.46
2:A:1404:C:O2'	2:A:1405:U:H5'	2.15	0.46
2:A:1786:A:H1'	2:A:1938:A:N6	2.29	0.46
2:A:1817:G:H2'	2:A:1818:U:H5'	1.98	0.46
2:A:2290:G:O2'	2:A:2291:U:H5'	2.14	0.46
8:G:33:LEU:HB3	8:G:75:MET:CE	2.44	0.46
11:L:1:MET:SD	11:L:67:LYS:HE3	2.56	0.46
27:b:11:SER:O	27:b:15:MET:HG3	2.15	0.46
2:A:144:A:H2'	2:A:145:C:C6	2.50	0.46
2:A:319:G:H2'	2:A:320:A:O4'	2.15	0.46
2:A:319:G:O2'	2:A:320:A:H5'	2.15	0.46
2:A:755:U:H2'	2:A:756:A:H8	1.79	0.46
2:A:807:U:O2	6:E:69:ARG:NH2	2.45	0.46
2:A:1056:G:HO2'	2:A:1086:A:H1'	1.81	0.46
2:A:1562:U:H2'	2:A:1563:U:O4'	2.15	0.46
2:A:1863:G:H4'	2:A:2411:A:H4'	1.98	0.46
2:A:2262:U:O2'	2:A:2263:C:H5'	2.15	0.46
2:A:2507:C:H2'	2:A:2508:G:O4'	2.15	0.46
2:A:2880:C:HO2'	14:O:90:ARG:HD3	1.79	0.46
6:E:31:VAL:HG21	6:E:104:ALA:HB2	1.98	0.46
8:G:89:LEU:HG	8:G:162:VAL:HG22	1.97	0.46
2:A:414:C:H2'	2:A:415:A:H8	1.79	0.46
2:A:645:C:H2'	2:A:647:G:C8	2.50	0.46
2:A:722:A:H2'	2:A:723:C:H6	1.80	0.46
2:A:1749:A:O2'	2:A:1750:G:H5'	2.15	0.46
2:A:1965:C:H5''	2:A:1966:A:H2'	1.97	0.46
2:A:2636:C:H2'	2:A:2637:U:H6	1.81	0.46
2:A:2836:U:H2'	2:A:2837:A:H8	1.81	0.46
4:C:119:GLY:HA2	4:C:189:ARG:NH1	2.29	0.46
5:D:146:ILE:HA	5:D:159:LYS:HZ2	1.80	0.46
16:Q:106:LYS:HA	16:Q:109:ARG:HD2	1.97	0.46
22:W:6:ALA:HB2	22:W:42:LEU:CD2	2.45	0.46
2:A:173:A:H2'	2:A:174:U:C6	2.50	0.46
2:A:213:A:O2'	2:A:214:G:H5'	2.15	0.46
2:A:416:U:H2'	2:A:417:C:O4'	2.16	0.46
2:A:948:C:H1'	2:A:984:A:C8	2.50	0.46
2:A:1075:C:H2'	2:A:1076:C:H6	1.81	0.46
2:A:1401:G:H2'	2:A:1402:U:C6	2.51	0.46
2:A:1585:C:H2'	2:A:1586:A:O4'	2.15	0.46
2:A:1990:C:H2'	2:A:1991:U:C1'	2.46	0.46
2:A:2295:C:OP1	15:P:10:ARG:NH2	2.48	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:2820:A:H4'	14:O:3:HIS:CG	2.50	0.46
3:B:105:G:O2'	3:B:106:G:H5'	2.15	0.46
4:C:16:VAL:HG22	4:C:206:GLY:HA3	1.98	0.46
6:E:149:ILE:HG22	6:E:192:ALA:HB1	1.97	0.46
8:G:52:PHE:CE2	8:G:69:ARG:HA	2.50	0.46
20:U:33:LYS:HE2	20:U:80:TRP:CD2	2.50	0.46
26:a:12:SER:HB2	26:a:32:ILE:HD11	1.98	0.46
2:A:289:G:H2'	2:A:290:U:H6	1.81	0.46
2:A:843:G:O2'	2:A:844:A:H5'	2.16	0.46
2:A:928:A:O2'	2:A:929:U:H5'	2.16	0.46
2:A:1365:A:OP1	24:Y:3:ARG:NE	2.40	0.46
2:A:2585:U:H4'	2:A:2586:U:H5''	1.96	0.46
4:C:43:ARG:NH2	4:C:49:ILE:HD11	2.30	0.46
4:C:133:ARG:HB3	4:C:186:ALA:HB1	1.96	0.46
5:D:1:MET:HE3	5:D:100:LEU:HG	1.98	0.46
7:F:136:ILE:CG2	7:F:146:VAL:HG11	2.46	0.46
22:W:6:ALA:CB	22:W:40:ILE:HD12	2.46	0.46
29:d:30:VAL:HG22	29:d:33:ARG:NH1	2.30	0.46
2:A:4:U:H2'	2:A:5:A:C8	2.51	0.46
2:A:521:U:H2'	2:A:522:A:C8	2.51	0.46
2:A:1636:U:H2'	2:A:1637:A:C8	2.51	0.46
2:A:1651:G:H4'	14:O:39:PRO:HG2	1.98	0.46
2:A:1827:U:OP2	4:C:221:ARG:HD2	2.16	0.46
2:A:1853:A:H1'	2:A:2234:G:O4'	2.16	0.46
2:A:1870:C:H2'	2:A:1871:A:N7	2.31	0.46
2:A:2420:C:P	30:e:34:THR:HB	2.56	0.46
2:A:2528:U:O2'	2:A:2530:A:OP1	2.19	0.46
2:A:2806:C:H2'	2:A:2807:U:O4'	2.16	0.46
2:A:2899:A:H5'	10:K:136:GLN:OE1	2.15	0.46
3:B:48:U:H2'	3:B:49:C:C6	2.50	0.46
6:E:5:LEU:HD13	6:E:10:SER:O	2.16	0.46
11:L:71:ARG:NH2	11:L:123:LEU:O	2.48	0.46
14:O:100:CYS:SG	14:O:101:GLY:N	2.87	0.46
30:e:9:GLY:O	30:e:13:ARG:HD3	2.16	0.46
2:A:396:G:O3'	24:Y:31:PRO:HA	2.16	0.46
2:A:476:G:H4'	2:A:502:A:N1	2.31	0.46
2:A:487:C:H2'	2:A:488:G:O4'	2.16	0.46
2:A:878:A:H3'	2:A:879:G:C8	2.48	0.46
2:A:962:G:H4'	2:A:2496:C:O2'	2.16	0.46
2:A:2327:A:H2'	2:A:2328:A:C8	2.50	0.46
2:A:2341:G:O2'	2:A:2342:C:H5'	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:2580:U:H5''	5:D:135:GLY:O	2.16	0.46
5:D:31:ALA:HA	5:D:96:ILE:O	2.16	0.46
13:N:54:THR:HG22	13:N:59:ARG:HG3	1.98	0.46
14:O:114:GLU:HG3	14:O:118:ARG:HE	1.81	0.46
18:S:52:PRO:HG2	18:S:53:PHE:CD2	2.51	0.46
26:a:44:ILE:O	26:a:48:ILE:HG13	2.16	0.46
28:c:9:ILE:HD12	28:c:51:GLU:HG3	1.96	0.46
2:A:659:G:H4'	6:E:95:LYS:HB3	1.97	0.46
2:A:1109:C:H2'	2:A:1110:G:N9	2.31	0.46
2:A:1310:G:C2'	2:A:1311:G:H5'	2.46	0.46
2:A:1430:G:H2'	2:A:1431:A:C8	2.51	0.46
2:A:1597:A:C5'	2:A:1598:A:H5'	2.39	0.46
2:A:1854:A:H61	2:A:1888:G:H1'	1.80	0.46
2:A:2745:C:H2'	2:A:2746:U:C6	2.50	0.46
2:A:2896:C:O2'	2:A:2897:U:H5'	2.14	0.46
7:F:6:ASP:OD1	7:F:6:ASP:N	2.49	0.46
7:F:134:GLU:HA	7:F:150:ARG:O	2.16	0.46
12:M:89:VAL:HG21	12:M:123:ARG:NH2	2.30	0.46
13:N:75:GLU:HB2	13:N:90:GLU:CG	2.35	0.46
21:V:6:ARG:O	21:V:25:VAL:HG21	2.16	0.46
2:A:152:A:H2'	2:A:153:U:H6	1.80	0.45
2:A:226:A:O2'	2:A:227:A:H5'	2.16	0.45
2:A:322:A:H5'	2:A:340:A:C1'	2.46	0.45
2:A:628:G:O2'	2:A:629:G:H5'	2.15	0.45
2:A:687:C:H5''	29:d:2:LYS:HE2	1.98	0.45
2:A:1165:A:O2'	2:A:1166:G:H5'	2.15	0.45
2:A:1407:G:O2'	2:A:1408:G:H5'	2.16	0.45
2:A:2366:A:H2'	2:A:2367:G:O4'	2.16	0.45
2:A:2809:A:H2'	2:A:2810:A:C8	2.51	0.45
3:B:11:C:O2'	3:B:12:C:H5'	2.15	0.45
3:B:113:C:H2'	3:B:114:C:C6	2.51	0.45
4:C:37:ASN:HB2	4:C:62:TYR:HB2	1.97	0.45
13:N:20:LEU:HD13	22:W:81:PRO:CG	2.40	0.45
2:A:89:A:O2'	2:A:90:U:H5'	2.16	0.45
2:A:656:G:H2'	2:A:657:U:C6	2.50	0.45
2:A:1400:U:H2'	2:A:1401:G:C8	2.51	0.45
2:A:2272:U:H5''	2:A:2273:A:OP1	2.16	0.45
2:A:2677:G:H2'	2:A:2678:C:H6	1.79	0.45
2:A:2785:C:H2'	2:A:2786:U:H6	1.81	0.45
3:B:50:A:P	15:P:67:ASN:HA	2.56	0.45
16:Q:75:GLN:HB2	16:Q:78:SER:HB2	1.96	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:W:30:ILE:HA	22:W:91:PHE:O	2.16	0.45
2:A:40:U:O2'	2:A:41:C:H5'	2.15	0.45
2:A:269:C:O2'	2:A:270:A:H5'	2.17	0.45
2:A:597:G:O2'	2:A:598:U:H5'	2.16	0.45
2:A:752:A:H3'	29:d:1:MET:CE	2.46	0.45
2:A:864:G:H2'	2:A:865:C:C6	2.51	0.45
2:A:1039:A:O2'	22:W:45:ASP:OD2	2.14	0.45
2:A:1076:C:H2'	2:A:1077:A:C8	2.52	0.45
2:A:1085:A:H1'	2:A:1105:U:H1'	1.97	0.45
2:A:1278:C:OP1	14:O:36:THR:HG22	2.16	0.45
2:A:2191:A:O2'	2:A:2192:U:H5'	2.16	0.45
2:A:2392:A:OP1	30:e:32:ILE:HD12	2.16	0.45
2:A:2580:U:H3'	2:A:2581:G:C2	2.51	0.45
2:A:2886:A:N1	27:b:40:ARG:HD2	2.30	0.45
3:B:21:G:O2'	3:B:22:U:H5'	2.17	0.45
7:F:60:ILE:HG22	7:F:99:PHE:HE2	1.81	0.45
8:G:98:VAL:HG12	8:G:103:ILE:HG12	1.98	0.45
13:N:58:LYS:O	13:N:60:GLN:NE2	2.49	0.45
20:U:30:ILE:CD1	20:U:32:LEU:HD21	2.47	0.45
23:X:62:LYS:HB3	23:X:62:LYS:HE2	1.80	0.45
2:A:95:A:H2'	2:A:96:C:O4'	2.15	0.45
2:A:1083:U:H2'	2:A:1085:A:OP2	2.15	0.45
2:A:1259:G:O2'	2:A:1260:A:H5'	2.15	0.45
2:A:1376:C:H2'	2:A:1377:G:O4'	2.16	0.45
2:A:1446:C:H2'	2:A:1447:C:C6	2.51	0.45
2:A:1614:A:C2	19:T:93:ALA:HB2	2.52	0.45
2:A:1831:G:O2'	2:A:1832:C:H5'	2.16	0.45
2:A:1842:G:H1'	4:C:243:HIS:CD2	2.51	0.45
2:A:1910:G:H2'	2:A:1911:U:O4'	2.16	0.45
2:A:2794:C:H2'	2:A:2795:C:C6	2.52	0.45
3:B:27:C:O2'	3:B:28:C:H5'	2.17	0.45
7:F:102:ARG:O	7:F:106:ILE:HB	2.17	0.45
21:V:72:ILE:HG21	21:V:83:VAL:HG23	1.98	0.45
2:A:318:C:O2'	2:A:319:G:H5'	2.15	0.45
2:A:538:A:H4'	10:K:7:LYS:HG2	1.98	0.45
2:A:566:U:OP1	12:M:29:LYS:HE2	2.17	0.45
2:A:1429:G:H2'	2:A:1430:G:H8	1.82	0.45
2:A:1657:U:O2'	2:A:1658:C:H5'	2.16	0.45
2:A:2030:A:N3	2:A:2499:C:H5''	2.32	0.45
2:A:2052:A:O4'	5:D:147:GLY:HA3	2.17	0.45
2:A:2240:U:O2'	2:A:2241:A:H5'	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:2289:G:O2'	2:A:2290:G:H5'	2.17	0.45
2:A:2531:A:OP1	8:G:175:LYS:HB3	2.16	0.45
2:A:2555:U:H2'	2:A:2556:C:H5'	1.98	0.45
2:A:2602[B]:A:H2'	2:A:2602[B]:A:N3	2.32	0.45
2:A:2880:C:O2'	14:O:90:ARG:HD3	2.17	0.45
15:P:25:ARG:HA	15:P:91:SER:O	2.16	0.45
19:T:77:ASP:HB2	19:T:102:HIS:HB2	1.99	0.45
2:A:169:G:O2'	2:A:170:U:H5'	2.16	0.45
2:A:191:A:H2'	2:A:192:C:H6	1.81	0.45
2:A:353:C:O2	2:A:353:C:H2'	2.17	0.45
2:A:476:G:N2	2:A:478:A:H3'	2.32	0.45
2:A:565:C:H2'	2:A:566:U:C6	2.52	0.45
2:A:671:C:H2'	2:A:672:C:H6	1.82	0.45
2:A:687:C:O4'	29:d:4:THR:HA	2.16	0.45
2:A:729:G:C6	4:C:207:LYS:HB2	2.52	0.45
2:A:747:U:O2'	19:T:92:ARG:NH2	2.50	0.45
2:A:925:A:O2'	2:A:926:G:H5'	2.16	0.45
2:A:1082:U:H3'	2:A:1083:U:C5'	2.45	0.45
2:A:1558:C:H4'	2:A:1559:U:O5'	2.16	0.45
2:A:1773:A:H2'	2:A:1774:C:H5'	1.99	0.45
2:A:2210:U:H4'	2:A:2211:A:C5'	2.46	0.45
2:A:2484:G:OP1	13:N:44:ARG:NH2	2.38	0.45
2:A:2818:U:H4'	2:A:2837:A:C4'	2.46	0.45
4:C:180:GLU:OE2	4:C:267:ILE:HG23	2.17	0.45
5:D:16:THR:CG2	5:D:20:VAL:HB	2.46	0.45
7:F:103:LEU:O	7:F:108:VAL:HG23	2.17	0.45
19:T:83:LYS:HD2	19:T:95:ARG:HH11	1.81	0.45
2:A:96:C:H4'	25:Z:41:HIS:CE1	2.52	0.45
2:A:1922:G:H2'	2:A:1923:U:H6	1.80	0.45
2:A:2592:G:H2'	2:A:2593:U:C6	2.51	0.45
2:A:2808:G:O2'	2:A:2890:G:O6	2.20	0.45
5:D:40:LEU:HD23	5:D:44:GLY:C	2.41	0.45
6:E:190:ALA:O	6:E:193:VAL:HG12	2.17	0.45
7:F:30:ARG:O	7:F:159:THR:HG23	2.17	0.45
11:L:14:SER:HA	11:L:51:LYS:HD2	1.99	0.45
18:S:66:HIS:ND1	18:S:94:THR:HG22	2.32	0.45
21:V:45:HIS:CD2	21:V:58:ILE:HG12	2.50	0.45
31:f:25:VAL:HB	31:f:35:GLN:HG2	1.99	0.45
2:A:644:A:H2'	2:A:645:C:C4'	2.47	0.45
2:A:1088:A:N3	2:A:1088:A:H2'	2.32	0.45
2:A:1484:U:H2'	2:A:1485:U:H5''	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:2031:A:C6	2:A:2498:C:H1'	2.52	0.45
2:A:2352:A:C2'	2:A:2353:G:H5'	2.47	0.45
3:B:106:G:O2'	3:B:107:G:H5'	2.17	0.45
8:G:95:ARG:HB3	8:G:128:GLN:OE1	2.17	0.45
14:O:95:THR:HG22	14:O:115:LEU:HD23	1.98	0.45
16:Q:106:LYS:HA	16:Q:109:ARG:CD	2.47	0.45
19:T:1:MET:HE3	19:T:62:ASP:CG	2.42	0.45
26:a:24:LEU:HD11	26:a:54:MET:CE	2.47	0.45
30:e:55:LEU:HG	30:e:59:ILE:HD11	1.98	0.45
2:A:17:G:H2'	2:A:18:U:C6	2.51	0.45
2:A:597:G:H2'	2:A:598:U:H6	1.82	0.45
2:A:605:G:H1'	2:A:657:U:H1'	1.99	0.45
2:A:634:C:H2'	2:A:635:C:H6	1.79	0.45
2:A:708:G:H2'	2:A:709:U:C6	2.52	0.45
2:A:1153:C:H5''	17:R:62:ILE:HD13	1.98	0.45
2:A:1213:A:H2'	2:A:1214:A:H8	1.82	0.45
2:A:1491:G:H5'	4:C:98:ASP:OD2	2.17	0.45
2:A:1495:A:C2	2:A:1578:U:H1'	2.50	0.45
2:A:1831:G:H2'	2:A:1832:C:H6	1.82	0.45
2:A:1862:G:O2'	2:A:1863:G:H5'	2.17	0.45
2:A:2000:C:O2'	2:A:2001:C:H5'	2.17	0.45
2:A:2513:A:H2'	2:A:2514:U:C6	2.52	0.45
2:A:2783:U:H2'	2:A:2784:U:C6	2.52	0.45
6:E:28:VAL:HG12	12:M:6:LEU:HD13	1.98	0.45
6:E:140:ASP:OD1	6:E:141:MET:HG2	2.16	0.45
7:F:50:LEU:HD23	7:F:50:LEU:HA	1.86	0.45
12:M:109:LYS:HA	12:M:126:ARG:O	2.17	0.45
24:Y:32:ASN:O	24:Y:52:SER:HA	2.16	0.45
25:Z:2:LYS:CE	25:Z:6:LEU:HD11	2.44	0.45
2:A:443:A:N7	6:E:40:ARG:HG2	2.32	0.45
2:A:527:C:H41	2:A:2777:G:HO2'	1.62	0.45
2:A:839:U:H2'	2:A:840:C:C6	2.52	0.45
2:A:975:A:H5'	18:S:78:ARG:HH22	1.82	0.45
2:A:1220:G:O2'	2:A:1221:C:H5'	2.16	0.45
2:A:1309:G:H4'	29:d:7:PRO:HB2	1.98	0.45
2:A:1844:C:O2'	2:A:1845:G:H5'	2.16	0.45
2:A:2298:A:H2'	2:A:2299:U:O4'	2.16	0.45
2:A:2365:G:O6	30:e:39:LYS:HE3	2.17	0.45
2:A:2637:U:C2'	2:A:2638:G:H5'	2.47	0.45
2:A:2690:U:O2'	2:A:2872:A:H1'	2.17	0.45
2:A:2699:C:H2'	2:A:2700:A:H8	1.82	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:D:15:PHE:HD2	16:Q:55:LEU:HD11	1.82	0.45
7:F:73:SER:OG	7:F:80:ARG:HD3	2.16	0.45
13:N:8:LYS:HD2	13:N:9:PHE:CE2	2.51	0.45
24:Y:36:HIS:O	24:Y:48:THR:HA	2.17	0.45
2:A:863:A:O2'	2:A:864:G:H5'	2.16	0.44
2:A:981:A:OP2	2:A:982:C:N4	2.38	0.44
2:A:1829:A:H3'	2:A:1830:C:C6	2.52	0.44
2:A:1893:C:H2'	2:A:1894:C:H5'	1.98	0.44
2:A:2346:A:H3'	2:A:2347:C:H5''	1.99	0.44
3:B:51:G:O2'	3:B:52:A:H5'	2.16	0.44
5:D:53:GLY:O	5:D:76:GLY:HA2	2.17	0.44
6:E:176:ASP:OD1	6:E:179:SER:OG	2.27	0.44
7:F:66:LEU:O	7:F:87:CYS:HA	2.17	0.44
8:G:117:LEU:HB3	8:G:121:ILE:CG2	2.47	0.44
10:K:12:LYS:O	10:K:41:LYS:NZ	2.51	0.44
21:V:42:VAL:O	21:V:60:GLU:HA	2.16	0.44
2:A:806:C:OP2	12:M:41:ARG:NH1	2.42	0.44
2:A:1361:G:O2'	2:A:1362:C:H5'	2.18	0.44
2:A:1601:G:H2'	2:A:1602:U:O4'	2.17	0.44
2:A:2331:G:O2'	2:A:2336:A:N1	2.37	0.44
4:C:31:ALA:HB3	4:C:32:PRO:HD3	2.00	0.44
5:D:7:LYS:NZ	5:D:200:ASP:HB3	2.32	0.44
24:Y:3:ARG:O	24:Y:12:PRO:HD3	2.17	0.44
2:A:12:U:H2'	2:A:13:A:H5'	1.98	0.44
2:A:153:U:H2'	2:A:154:U:C6	2.52	0.44
2:A:397:U:O2'	2:A:398:C:H5'	2.18	0.44
2:A:631:A:OP1	30:e:47:LYS:NZ	2.28	0.44
2:A:664:G:O2'	2:A:940:G:OP1	2.22	0.44
2:A:1006:C:O2'	2:A:1007:C:H5'	2.17	0.44
2:A:1637:A:H5'	2:A:1760:C:O2'	2.17	0.44
2:A:1853:A:H2'	2:A:1854:A:C8	2.52	0.44
2:A:2281:A:C2'	2:A:2282:G:H5'	2.46	0.44
2:A:2600:A:H2'	2:A:2601[A]:C:C6	2.52	0.44
2:A:574:A:N6	2:A:2034:U:OP1	2.51	0.44
2:A:651:G:OP1	30:e:19:LYS:HB2	2.18	0.44
2:A:768:G:O2'	2:A:769:U:H5'	2.18	0.44
2:A:1913:A:H4'	2:A:1914:C:OP2	2.17	0.44
2:A:2634:A:O2'	2:A:2635:A:H5'	2.17	0.44
4:C:246:THR:HB	4:C:247:PRO:HD2	2.00	0.44
6:E:3:LEU:O	6:E:11:ALA:HA	2.17	0.44
7:F:122:PHE:HA	7:F:127:ASN:O	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:H:2:GLN:HB2	9:H:39:ALA:HB3	1.99	0.44
9:H:9:VAL:HG13	9:H:12:LEU:HB2	1.99	0.44
11:L:51:LYS:HZ3	11:L:96:GLY:HA2	1.81	0.44
19:T:1:MET:HE3	19:T:62:ASP:CB	2.47	0.44
21:V:96:PHE:O	21:V:100:SER:HA	2.17	0.44
31:f:14:CYS:SG	31:f:27:CYS:HB2	2.57	0.44
2:A:418:C:H2'	2:A:419:U:H6	1.83	0.44
2:A:443:A:C6	6:E:40:ARG:HD2	2.52	0.44
2:A:569:U:H2'	2:A:570:G:O4'	2.17	0.44
2:A:727:A:H2'	2:A:728:G:O4'	2.17	0.44
2:A:755:U:O2'	2:A:756:A:H5'	2.18	0.44
2:A:896:A:H3'	2:A:897:C:H5'	1.99	0.44
2:A:984:A:H2'	2:A:984:A:N3	2.33	0.44
2:A:1057:A:N7	2:A:1086:A:H2'	2.33	0.44
2:A:1118:C:H3'	2:A:1119:U:H5''	2.00	0.44
2:A:1750:G:O2'	2:A:1751:U:H5'	2.17	0.44
2:A:1824:G:O3'	4:C:247:PRO:HD3	2.17	0.44
2:A:1890:A:C2	2:A:1891:G:H1'	2.52	0.44
2:A:2028:U:H2'	2:A:2029:G:C8	2.53	0.44
2:A:2394:C:H5''	12:M:63:LYS:HE2	1.98	0.44
2:A:2514:U:H2'	2:A:2515:C:H6	1.82	0.44
2:A:2790:U:H4'	2:A:2791:G:OP1	2.18	0.44
3:B:119:A:H2'	3:B:120:U:O4'	2.17	0.44
4:C:30:PHE:CE2	4:C:32:PRO:HG2	2.53	0.44
4:C:141:VAL:HG23	4:C:162:VAL:CG2	2.48	0.44
14:O:6:SER:HB3	14:O:46:ARG:HH22	1.82	0.44
16:Q:6:LYS:O	16:Q:10:GLN:HG2	2.18	0.44
25:Z:25:GLN:HB2	25:Z:46:VAL:HG11	1.99	0.44
2:A:128:C:H2'	2:A:129:C:C6	2.53	0.44
2:A:156:A:O2'	2:A:157:C:H5'	2.18	0.44
2:A:188:G:H5'	24:Y:14:THR:CG2	2.47	0.44
2:A:397:U:H5''	24:Y:32:ASN:HB2	1.99	0.44
2:A:554:U:H2'	2:A:555:G:O4'	2.18	0.44
2:A:620:G:H4'	2:A:621:A:O5'	2.18	0.44
2:A:784:G:O2'	2:A:785:G:H5''	2.17	0.44
2:A:810:U:C5	12:M:29:LYS:HG2	2.53	0.44
2:A:1185:G:H5''	2:A:1186:G:OP1	2.17	0.44
2:A:1288:G:OP2	2:A:1288:G:N2	2.23	0.44
2:A:1604:C:H2'	2:A:1605:C:C6	2.53	0.44
2:A:1627:G:O2'	2:A:1628:G:H5'	2.18	0.44
2:A:1693:U:H4'	2:A:1694:C:OP2	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:2007:U:O2'	2:A:2008:C:H5'	2.17	0.44
2:A:2056:G:H4'	27:b:5:GLN:NE2	2.33	0.44
2:A:2437:G:O2'	2:A:2438:U:H5'	2.18	0.44
12:M:78:ARG:HG2	12:M:113:ALA:CB	2.46	0.44
19:T:31:GLN:O	19:T:35:ILE:HG13	2.17	0.44
22:W:30:ILE:HD11	22:W:40:ILE:CD1	2.47	0.44
29:d:34:ARG:NH1	29:d:41:ARG:O	2.51	0.44
2:A:1:G:H2'	2:A:2:G:H8	1.82	0.44
2:A:475:C:O2	2:A:479:A:N6	2.51	0.44
2:A:1003:G:O2'	2:A:1010:A:N1	2.44	0.44
2:A:1164:C:H2'	2:A:1165:A:C8	2.52	0.44
2:A:1275:A:H4'	2:A:1276:A:O5'	2.18	0.44
2:A:1301:A:H4'	2:A:1302:A:OP1	2.18	0.44
2:A:1432:G:H2'	2:A:1433:A:H8	1.82	0.44
2:A:1824:G:O2'	2:A:1825:U:H5'	2.18	0.44
2:A:2586:U:OP2	2:A:2608:G:N2	2.44	0.44
2:A:2670:A:O2'	2:A:2671:G:H5'	2.17	0.44
2:A:2710:C:H2'	2:A:2711:A:C8	2.53	0.44
3:B:4:C:O2'	3:B:5:U:H5'	2.18	0.44
3:B:33:G:O2'	3:B:34:A:H5'	2.18	0.44
3:B:78:A:H2'	3:B:79:G:O4'	2.17	0.44
4:C:161:TYR:CB	4:C:194:GLU:HG2	2.47	0.44
10:K:101:ILE:O	10:K:105:VAL:HG23	2.18	0.44
21:V:24:LYS:HB3	21:V:37:GLU:OE1	2.18	0.44
28:c:5:ILE:HB	28:c:28:ARG:NH2	2.33	0.44
2:A:103:A:H2'	2:A:104:A:O4'	2.18	0.44
2:A:250:G:H2'	2:A:251:A:C8	2.52	0.44
2:A:422:A:H2'	2:A:423:A:O4'	2.18	0.44
2:A:532:A:N7	2:A:2021:C:O2'	2.44	0.44
2:A:706:A:N6	2:A:725:G:H1'	2.32	0.44
2:A:717:C:H2'	2:A:718:A:H5'	2.00	0.44
2:A:740:C:O2'	2:A:741:U:H5'	2.18	0.44
2:A:804:A:H2'	2:A:806:C:C4	2.53	0.44
2:A:849:A:H2'	2:A:850:U:C6	2.52	0.44
2:A:2028:U:H2'	2:A:2029:G:O4'	2.18	0.44
2:A:2399:G:O2'	2:A:2400:G:H5'	2.18	0.44
2:A:2515:C:H2'	2:A:2516:A:H8	1.82	0.44
2:A:2605:U:H2'	2:A:2606:C:C6	2.53	0.44
3:B:27:C:OP1	15:P:34:HIS:NE2	2.42	0.44
4:C:79:GLU:HB2	4:C:93:LEU:O	2.18	0.44
7:F:34:ILE:O	7:F:91:LEU:HB2	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:F:135:GLN:OE1	7:F:135:GLN:N	2.49	0.44
12:M:51:GLU:CG	12:M:56:PRO:HA	2.48	0.44
14:O:32:GLU:O	14:O:114:GLU:HB2	2.17	0.44
24:Y:16:ASN:HA	24:Y:25:THR:O	2.17	0.44
26:a:27:LEU:CD2	26:a:47:MET:HE3	2.48	0.44
2:A:729:G:OP2	4:C:13:ARG:HD3	2.17	0.44
2:A:780:G:H2'	2:A:782:A:N7	2.33	0.44
2:A:1029:A:H2'	2:A:1030:C:O4'	2.18	0.44
2:A:1270:C:H5''	2:A:1271:G:O5'	2.18	0.44
2:A:1474:U:C2'	2:A:1475:G:H5'	2.48	0.44
2:A:1772:A:H2'	2:A:1773:A:O3'	2.17	0.44
2:A:2193:G:H2'	2:A:2194:U:H6	1.83	0.44
4:C:161:TYR:CG	4:C:194:GLU:HG2	2.53	0.44
4:C:164:ILE:HG23	4:C:172:VAL:CG1	2.48	0.44
4:C:175:ARG:HG3	4:C:181:MET:SD	2.58	0.44
13:N:53:MET:HE3	13:N:117:PHE:CE1	2.51	0.44
23:X:34:GLY:N	23:X:61:ALA:O	2.40	0.44
2:A:30:G:H2'	2:A:31:C:H6	1.83	0.43
2:A:184:C:H2'	2:A:185:G:H8	1.83	0.43
2:A:393:C:O2'	2:A:394:C:H5'	2.17	0.43
2:A:435:C:C2'	2:A:436:C:H5'	2.47	0.43
2:A:438:G:H2'	2:A:439:A:C8	2.53	0.43
2:A:536:G:H2'	2:A:537:G:O4'	2.18	0.43
2:A:553:G:H2'	2:A:554:U:H6	1.83	0.43
2:A:636:G:OP1	12:M:129:LYS:NZ	2.40	0.43
2:A:673:C:O2'	2:A:674:G:H5'	2.18	0.43
2:A:1100:C:H2'	2:A:1101:U:O4'	2.18	0.43
2:A:1431:A:H2'	2:A:1432:G:H8	1.83	0.43
2:A:2345:G:N3	2:A:2381:A:H2'	2.33	0.43
7:F:111:ILE:HD12	7:F:114:PHE:CE1	2.52	0.43
10:K:19:ASP:OD1	10:K:21:THR:HG22	2.18	0.43
15:P:33:ARG:HG2	15:P:34:HIS:CD2	2.53	0.43
22:W:77:VAL:HG23	22:W:89:ILE:HG12	2.00	0.43
26:a:27:LEU:HD23	26:a:47:MET:HE3	2.00	0.43
28:c:14:SER:OG	28:c:48:ILE:HG23	2.17	0.43
2:A:4:U:H2'	2:A:5:A:H8	1.83	0.43
2:A:103:A:C2	2:A:104:A:H1'	2.54	0.43
2:A:370:G:O2'	2:A:423:A:H3'	2.18	0.43
2:A:1323:C:O2'	2:A:1324:G:H5'	2.18	0.43
2:A:1363:C:O2'	2:A:1809:A:H1'	2.18	0.43
2:A:1389:G:O2'	2:A:1390:U:H5'	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:1591:A:O2'	2:A:1592:C:H5'	2.18	0.43
2:A:1682:G:H2'	2:A:1683:U:H6	1.80	0.43
2:A:1838:C:H4'	2:A:1839:G:C8	2.53	0.43
2:A:1973:G:H2'	2:A:1974:C:C6	2.52	0.43
2:A:2019:A:H4'	17:R:34:VAL:HG21	1.99	0.43
5:D:7:LYS:HE3	5:D:77:ARG:CZ	2.49	0.43
7:F:117:LEU:CD1	7:F:130:MET:HG3	2.47	0.43
11:L:105:ARG:NH2	16:Q:32:VAL:HB	2.33	0.43
12:M:75:ALA:O	12:M:108:ALA:HA	2.17	0.43
13:N:34:LYS:HE2	13:N:99:GLY:O	2.19	0.43
13:N:42:THR:O	13:N:46:ILE:HG13	2.19	0.43
15:P:27:VAL:CG2	15:P:40:ILE:HD13	2.48	0.43
17:R:66:ASN:O	17:R:70:ARG:HG3	2.18	0.43
25:Z:3:ALA:O	25:Z:7:ARG:NH1	2.51	0.43
30:e:23:LYS:HA	30:e:48:ALA:O	2.18	0.43
2:A:27:G:N2	2:A:512:G:H1'	2.33	0.43
2:A:419:U:H2'	2:A:420:C:H6	1.82	0.43
2:A:1130:U:C2	2:A:2025:C:H5''	2.54	0.43
2:A:1495:A:N3	2:A:1578:U:O2'	2.38	0.43
2:A:1651:G:H5'	14:O:39:PRO:HG2	2.00	0.43
2:A:1652:A:O2'	2:A:1653:G:H5'	2.18	0.43
2:A:1757:A:O3'	2:A:1758:U:H4'	2.17	0.43
2:A:1791:A:H4'	4:C:205:LEU:HB2	2.01	0.43
2:A:1796:U:H2'	2:A:1797:G:H8	1.82	0.43
2:A:2087:G:H2'	2:A:2088:A:H8	1.83	0.43
3:B:54:G:O2'	3:B:55:U:H5'	2.18	0.43
5:D:7:LYS:HZ1	5:D:200:ASP:HB3	1.82	0.43
7:F:25:VAL:O	7:F:28:VAL:HG22	2.18	0.43
10:K:78:THR:HG21	10:K:85:LYS:NZ	2.34	0.43
10:K:98:GLU:O	10:K:102:GLU:HG3	2.17	0.43
15:P:53:THR:O	15:P:59:ALA:HB2	2.17	0.43
17:R:102:ASP:OD1	17:R:104:VAL:HG22	2.18	0.43
2:A:337:C:H2'	2:A:338:G:O4'	2.19	0.43
2:A:364:C:H2'	2:A:365:U:H6	1.84	0.43
2:A:440:C:O2'	2:A:441:U:H5'	2.19	0.43
2:A:1432:G:O2'	2:A:1433:A:H5'	2.19	0.43
2:A:1530:G:O2'	2:A:1531:C:H5'	2.19	0.43
2:A:1681:G:P	2:A:1757:A:H61	2.41	0.43
2:A:1897:G:H2'	2:A:1898:U:H6	1.83	0.43
2:A:1955:U:H2'	2:A:2551:C:O2'	2.18	0.43
2:A:2359:C:O2'	2:A:2360:G:H5'	2.17	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:2547:A:H4'	11:L:29:HIS:CD2	2.53	0.43
3:B:51:G:H2'	3:B:52:A:C8	2.53	0.43
8:G:54:PRO:HB3	8:G:61:GLY:HA3	2.00	0.43
17:R:35:ALA:O	17:R:39:VAL:HG23	2.18	0.43
17:R:47:TYR:HD2	17:R:50:ARG:HH21	1.67	0.43
26:a:5:ILE:O	26:a:37:GLU:HA	2.18	0.43
2:A:576:U:H2'	2:A:577:G:H8	1.80	0.43
2:A:1038:G:O2'	2:A:1039:A:H5'	2.18	0.43
2:A:1410:G:H2'	2:A:1411:U:C6	2.54	0.43
3:B:106:G:H2'	3:B:107:G:C8	2.54	0.43
5:D:8:LYS:HB2	5:D:201:LEU:HD11	2.00	0.43
5:D:16:THR:O	16:Q:79:PRO:HG3	2.19	0.43
6:E:102:ARG:O	6:E:106:LYS:HG3	2.18	0.43
10:K:49:ASP:OD1	10:K:121:LYS:NZ	2.50	0.43
15:P:75:GLY:HA2	15:P:78:VAL:HG12	2.00	0.43
19:T:24:ILE:HB	19:T:32:ALA:HB1	1.98	0.43
20:U:53:VAL:CG1	20:U:91:GLN:HB3	2.49	0.43
22:W:23:ALA:O	22:W:25:LYS:HG3	2.18	0.43
22:W:63:ILE:HG22	22:W:65:VAL:HG23	1.99	0.43
25:Z:2:LYS:NZ	25:Z:49:ASP:OD1	2.23	0.43
2:A:49:A:O5'	2:A:51:G:H5'	2.18	0.43
2:A:202:U:H2'	2:A:203:A:O4'	2.18	0.43
2:A:251:A:H2'	2:A:252:G:O4'	2.19	0.43
2:A:500:G:N1	2:A:503:A:OP2	2.48	0.43
2:A:693:A:O2'	2:A:694:U:H5'	2.18	0.43
2:A:745:G:H2'	2:A:746:U:H5'	2.00	0.43
2:A:786:C:O2'	2:A:787:C:H5'	2.17	0.43
2:A:825:A:H2'	2:A:826:U:H6	1.82	0.43
2:A:1412:U:H2'	2:A:1413:A:C8	2.53	0.43
2:A:1427:A:H4'	2:A:1428:C:O5'	2.18	0.43
2:A:1694:C:OP1	4:C:8:PRO:HG3	2.18	0.43
2:A:1757:A:HO2'	2:A:1758:U:P	2.39	0.43
2:A:1771:C:H2'	2:A:1772:A:C8	2.53	0.43
2:A:1911:U:H2'	2:A:1918:A:C2	2.54	0.43
2:A:2030:A:C2	2:A:2499:C:H5''	2.54	0.43
2:A:2355:G:O2'	2:A:2356:U:H5'	2.17	0.43
2:A:2592:G:H2'	2:A:2593:U:H6	1.83	0.43
2:A:2626:C:O2'	2:A:2627:G:H5'	2.19	0.43
9:H:1:MET:HE3	9:H:23:ALA:CA	2.49	0.43
11:L:105:ARG:HG2	11:L:123:LEU:C	2.44	0.43
13:N:62:LYS:HD3	13:N:64:TRP:CZ2	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:O:25:ALA:O	14:O:29:VAL:HG23	2.19	0.43
17:R:17:ILE:HG13	17:R:32:TYR:HE1	1.84	0.43
17:R:69:ALA:HB1	17:R:74:ILE:HG23	2.00	0.43
26:a:7:ILE:HD11	26:a:48:ILE:HD11	2.00	0.43
2:A:64:A:H2'	2:A:65:U:H6	1.81	0.43
2:A:237:C:O2'	2:A:238:C:H5'	2.17	0.43
2:A:303:G:H2'	2:A:304:U:C6	2.54	0.43
2:A:594:U:H2'	2:A:595:C:C6	2.54	0.43
2:A:624:C:O2'	2:A:657:U:H5''	2.19	0.43
2:A:638:G:H2'	2:A:639:U:H6	1.84	0.43
2:A:645:C:O2'	2:A:646:U:H5''	2.18	0.43
2:A:1014:A:O2'	2:A:1015:U:H5'	2.17	0.43
2:A:1264:A:O5'	2:A:1265:A:H2'	2.19	0.43
4:C:107:PRO:HG2	4:C:110:LEU:CG	2.48	0.43
7:F:4:LEU:HD12	7:F:100:PHE:HB3	2.00	0.43
7:F:100:PHE:O	7:F:104:ILE:HG23	2.18	0.43
7:F:130:MET:O	7:F:154:ILE:N	2.41	0.43
11:L:22:ILE:HG12	11:L:40:LYS:O	2.19	0.43
12:M:63:LYS:HG2	30:e:13:ARG:HD2	1.99	0.43
13:N:33:LEU:HD13	13:N:117:PHE:HB3	2.00	0.43
22:W:51:GLN:HG2	22:W:86:LEU:HD11	1.99	0.43
2:A:172:A:H2'	2:A:173:A:H8	1.84	0.43
2:A:175:G:O2'	2:A:176:A:H5'	2.18	0.43
2:A:217:A:H2'	2:A:218:A:O4'	2.18	0.43
2:A:264:C:O2'	2:A:265:A:H2'	2.18	0.43
2:A:659:G:O2'	6:E:95:LYS:O	2.31	0.43
2:A:671:C:O2'	2:A:672:C:H5'	2.19	0.43
2:A:673:C:OP1	6:E:49:ARG:NH2	2.48	0.43
2:A:775:G:H5'	2:A:777:G:C1'	2.45	0.43
2:A:1006:C:O2'	10:K:108:MET:O	2.32	0.43
2:A:1148:U:O2'	2:A:1149:G:H5'	2.18	0.43
2:A:1446:C:O2'	2:A:1447:C:H5'	2.18	0.43
2:A:1456:G:O2'	2:A:1457:U:H5'	2.19	0.43
2:A:1502:A:H2'	2:A:1503:A:C8	2.54	0.43
2:A:1683:U:H2'	2:A:1684:G:C8	2.53	0.43
2:A:1827:U:O2'	2:A:1828:G:H5'	2.18	0.43
2:A:1882:U:O2'	2:A:1883:U:H5'	2.19	0.43
2:A:2321:U:H3'	2:A:2322:A:H5''	2.01	0.43
2:A:2585:U:H4'	2:A:2586:U:C5'	2.49	0.43
2:A:2615:U:C2	27:b:4:GLN:HA	2.53	0.43
2:A:2689:U:H4'	2:A:2690:U:OP2	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:2725:A:O2'	2:A:2726:A:H2'	2.19	0.43
9:H:2:GLN:HB2	9:H:39:ALA:CB	2.49	0.43
11:L:77:ILE:HD11	16:Q:72:ARG:CZ	2.48	0.43
16:Q:33:VAL:CG2	16:Q:38:LYS:HG2	2.49	0.43
17:R:98:ILE:HG23	17:R:105:ALA:HB3	2.00	0.43
24:Y:5:CYS:HB2	24:Y:33:LEU:HD21	2.00	0.43
26:a:6:LYS:HG2	26:a:37:GLU:CG	2.38	0.43
2:A:475:C:H4'	2:A:510:C:H5'	2.01	0.43
2:A:528:A:H5''	10:K:113:PRO:HG3	2.01	0.43
2:A:558:U:OP1	10:K:113:PRO:HD2	2.19	0.43
2:A:656:G:O2'	2:A:657:U:H5'	2.18	0.43
2:A:1130:U:N3	2:A:2025:C:OP1	2.42	0.43
2:A:1953:A:O2'	2:A:2559:C:O2	2.35	0.43
2:A:1975:G:H2'	2:A:1976:U:O4'	2.19	0.43
2:A:2491:U:H5'	2:A:2570:G:C5'	2.48	0.43
2:A:2721:A:H2'	2:A:2722:G:O4'	2.19	0.43
3:B:111:U:H2'	3:B:112:G:H8	1.84	0.43
7:F:106:ILE:HG21	7:F:139:PRO:HG2	2.00	0.43
7:F:129:SER:HA	7:F:155:THR:HA	2.01	0.43
10:K:37:ARG:HG3	10:K:118:MET:SD	2.59	0.43
10:K:97:PRO:HG2	10:K:126:ALA:HB2	2.00	0.43
13:N:47:GLU:O	13:N:51:ARG:HG3	2.18	0.43
14:O:59:SER:OG	14:O:62:ASN:HB2	2.19	0.43
22:W:26:PHE:HZ	22:W:47:VAL:HG11	1.84	0.43
2:A:473:G:C2'	2:A:474:G:H5'	2.49	0.43
2:A:506:G:H5''	2:A:509:C:H1'	2.01	0.43
2:A:541:A:O2'	2:A:542:C:H5'	2.19	0.43
2:A:594:U:H2'	2:A:595:C:H6	1.83	0.43
2:A:738:G:O2'	2:A:739:A:H5'	2.19	0.43
2:A:929:U:O2'	2:A:930:G:H5'	2.19	0.43
2:A:1046:A:H3'	2:A:1047:G:H5'	2.00	0.43
2:A:1446:C:H2'	2:A:1447:C:H6	1.84	0.43
2:A:2766:A:N3	2:A:2766:A:H2'	2.33	0.43
2:A:2814:A:H2'	2:A:2815:C:H6	1.83	0.43
2:A:2855:C:H2'	2:A:2856:A:H8	1.84	0.43
4:C:174:LEU:O	4:C:181:MET:HA	2.19	0.43
6:E:129:PRO:HB3	6:E:156:ASN:HA	2.00	0.43
7:F:57:LEU:HD12	7:F:87:CYS:SG	2.58	0.43
12:M:71:ALA:HA	12:M:74:THR:HG22	2.01	0.43
22:W:4:ILE:CG1	22:W:50:MET:HE1	2.49	0.43
25:Z:32:ALA:HB2	25:Z:37:LEU:HD23	2.01	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:171:U:O2'	2:A:172:A:H5'	2.19	0.42
2:A:533:G:H2'	2:A:534:U:C6	2.53	0.42
2:A:1450:G:O2'	2:A:1451:C:H5'	2.19	0.42
2:A:1722:A:N6	2:A:1738:G:H1'	2.33	0.42
2:A:2305:U:O2	7:F:151:GLY:HA3	2.19	0.42
4:C:110:LEU:HD21	4:C:126:PRO:O	2.19	0.42
6:E:5:LEU:HD22	6:E:10:SER:OG	2.19	0.42
16:Q:21:ARG:HD3	16:Q:113:ARG:CZ	2.49	0.42
22:W:2:PHE:CB	22:W:61:LEU:HD22	2.48	0.42
24:Y:13:VAL:HG23	24:Y:29:PHE:HB2	2.01	0.42
2:A:78:U:O2'	2:A:79:C:H5'	2.19	0.42
2:A:429:A:H2'	2:A:430:A:C8	2.55	0.42
2:A:631:A:OP2	30:e:23:LYS:NZ	2.38	0.42
2:A:675:A:N3	2:A:2443:C:O2'	2.45	0.42
2:A:878:A:H5'	2:A:879:G:OP2	2.18	0.42
2:A:1323:C:C2'	2:A:1324:G:H5'	2.49	0.42
2:A:1482:G:H2'	2:A:1483:G:C8	2.54	0.42
2:A:1515:A:H1'	2:A:1557:C:H1'	2.01	0.42
2:A:1759:A:O2'	2:A:2714:G:H1'	2.19	0.42
2:A:1856:U:O2'	2:A:1857:G:H5'	2.19	0.42
2:A:2572:A:H2'	5:D:149:ASN:ND2	2.34	0.42
2:A:2590:A:H2'	2:A:2591:C:H6	1.84	0.42
2:A:2816:G:H1'	2:A:2883:A:O2'	2.18	0.42
2:A:2839:G:O2'	14:O:92:GLY:HA2	2.19	0.42
4:C:10:SER:HB2	4:C:11:PRO:HD2	2.01	0.42
4:C:93:LEU:HD13	4:C:103:TYR:CE1	2.53	0.42
5:D:186:LEU:HD21	16:Q:4:ILE:HG23	2.01	0.42
8:G:9:VAL:C	8:G:49:THR:HG23	2.44	0.42
9:H:4:ILE:HG13	9:H:18:GLN:HB3	2.01	0.42
13:N:82:MET:HE2	13:N:82:MET:HA	2.01	0.42
17:R:109:LEU:CD2	18:S:49:ILE:HD13	2.48	0.42
18:S:31:GLU:O	18:S:62:GLU:HG3	2.20	0.42
2:A:151:C:H2'	2:A:152:A:H8	1.85	0.42
2:A:153:U:H2'	2:A:154:U:H6	1.84	0.42
2:A:210:C:O2'	2:A:211:C:H5'	2.19	0.42
2:A:1021:A:H3'	2:A:1021:A:N3	2.34	0.42
2:A:1456:G:H2'	2:A:1457:U:C6	2.54	0.42
2:A:1642:G:O2'	2:A:1643:G:H5'	2.19	0.42
2:A:1681:G:N2	2:A:1763:G:OP2	2.39	0.42
2:A:1736:U:H2'	2:A:1737:G:O4'	2.19	0.42
2:A:2027:G:O2'	2:A:2028:U:H5'	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:2898:U:O2'	10:K:134:ALA:O	2.26	0.42
20:U:93:LEU:HD23	20:U:95:PHE:CE2	2.54	0.42
2:A:31:C:O2'	2:A:32:C:H5'	2.20	0.42
2:A:37:C:O2'	2:A:38:A:H5'	2.20	0.42
2:A:396:G:C1'	24:Y:29:PHE:HB3	2.48	0.42
2:A:997:G:O2'	2:A:998:C:H5'	2.20	0.42
2:A:1187:G:N2	2:A:1188:U:O4	2.52	0.42
2:A:1212:G:H1'	2:A:1237:A:H61	1.84	0.42
2:A:1383:A:O2'	2:A:1384:A:O4'	2.22	0.42
2:A:1637:A:O2'	2:A:1638:C:H5'	2.20	0.42
2:A:1910:G:O2'	2:A:1911:U:H5'	2.20	0.42
2:A:1979:U:O2'	2:A:1980:G:H5'	2.19	0.42
2:A:2693:G:O2'	2:A:2694:G:H5'	2.20	0.42
3:B:14:U:O3'	3:B:107:G:O2'	2.37	0.42
6:E:28:VAL:O	6:E:32:VAL:HG13	2.19	0.42
7:F:98:GLU:O	7:F:101:GLU:HG2	2.19	0.42
7:F:152:LEU:HD23	7:F:154:ILE:HD11	2.02	0.42
10:K:15:TRP:O	10:K:137:PRO:HA	2.19	0.42
19:T:86:MET:HE2	19:T:86:MET:HB3	1.90	0.42
23:X:65:GLY:HA3	23:X:82:ILE:HG23	2.02	0.42
2:A:98:G:H1'	2:A:103:A:H1'	2.00	0.42
2:A:181:A:H2'	2:A:182:A:C8	2.55	0.42
2:A:257:C:O2	12:M:104:GLN:NE2	2.53	0.42
2:A:273:G:O2'	2:A:274:C:H5'	2.19	0.42
2:A:321:U:OP2	6:E:130:LYS:HA	2.19	0.42
2:A:851:C:O4'	26:a:47:MET:HG2	2.20	0.42
2:A:898:C:H2'	2:A:899:A:O4'	2.20	0.42
2:A:966:G:H2'	2:A:967:U:C6	2.55	0.42
2:A:1019:U:O2'	2:A:1021:A:N7	2.40	0.42
2:A:1222:U:H2'	2:A:1223:G:C8	2.54	0.42
2:A:1545:A:H2'	2:A:1546:G:O4'	2.20	0.42
2:A:1951:U:H2'	2:A:1953:A:OP2	2.19	0.42
2:A:2039:U:H2'	2:A:2040:G:C8	2.54	0.42
2:A:2081:U:H2'	2:A:2082:A:C8	2.54	0.42
2:A:2759:G:H1'	8:G:35:ARG:HH21	1.85	0.42
3:B:23:G:H5''	3:B:24:G:OP1	2.20	0.42
5:D:56:LYS:HD2	5:D:59:ARG:HG3	2.00	0.42
7:F:4:LEU:HD23	7:F:4:LEU:HA	1.90	0.42
11:L:2:ILE:HG21	11:L:8:LEU:HD11	2.01	0.42
11:L:51:LYS:NZ	11:L:96:GLY:HA2	2.35	0.42
12:M:19:LEU:CD2	12:M:27:LEU:HD13	2.48	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:M:122:VAL:HG12	12:M:125:LEU:HB2	2.02	0.42
17:R:41:LYS:HG3	17:R:45:TYR:CE2	2.54	0.42
18:S:34:GLU:OE2	18:S:60:LYS:NZ	2.53	0.42
23:X:26:PHE:O	23:X:67:VAL:HB	2.19	0.42
2:A:91:A:H4'	2:A:92:U:O4'	2.20	0.42
2:A:282:A:O2'	2:A:283:G:H5'	2.20	0.42
2:A:299:A:H2'	2:A:300:A:C8	2.54	0.42
2:A:468:G:H2'	2:A:469:G:O4'	2.19	0.42
2:A:543:G:H2'	2:A:544:C:C6	2.55	0.42
2:A:988:A:P	26:a:12:SER:HB3	2.60	0.42
2:A:1437:C:H2'	2:A:1438:U:C6	2.55	0.42
2:A:1561:C:H2'	2:A:1562:U:C6	2.55	0.42
2:A:1799:G:OP1	4:C:258:ARG:HD2	2.18	0.42
2:A:1883:U:C2'	2:A:1884:G:H5'	2.50	0.42
2:A:1998:A:H2'	2:A:1999:C:C6	2.54	0.42
2:A:2561:U:O2	11:L:23:LYS:HE3	2.18	0.42
6:E:27:LEU:O	6:E:31:VAL:HG23	2.20	0.42
6:E:142:ALA:C	6:E:143:LEU:HD12	2.44	0.42
18:S:61:ALA:HB2	18:S:98:ILE:HD13	2.02	0.42
24:Y:39:TRP:CZ2	24:Y:44:LYS:HD3	2.55	0.42
28:c:10:LYS:HD3	28:c:54:ILE:HA	2.00	0.42
2:A:11:C:H2'	2:A:12:U:H5'	2.02	0.42
2:A:116:C:H2'	2:A:117:G:O4'	2.20	0.42
2:A:239:C:H1'	2:A:621:A:H2	1.84	0.42
2:A:736:C:H2'	2:A:737:C:C6	2.54	0.42
2:A:909:A:H2'	2:A:912:C:H5	1.84	0.42
2:A:1086:A:H3'	2:A:1086:A:N3	2.35	0.42
2:A:1283:G:N1	2:A:1286:A:OP2	2.51	0.42
2:A:1378:A:O2'	2:A:1380:G:OP2	2.38	0.42
2:A:1528:A:OP2	2:A:1543:G:N2	2.53	0.42
2:A:1995:U:OP2	2:A:1996:C:O2'	2.32	0.42
2:A:2193:G:O2'	2:A:2194:U:H5'	2.19	0.42
2:A:2313:C:H4'	7:F:88:LYS:HD3	2.02	0.42
2:A:2369:A:O2'	2:A:2370:G:H5'	2.20	0.42
6:E:2:GLU:HB3	6:E:11:ALA:HB1	2.02	0.42
8:G:37:LEU:HD13	8:G:68:ALA:CB	2.40	0.42
8:G:76:VAL:HA	8:G:79:VAL:HG22	2.02	0.42
12:M:92:LEU:O	12:M:96:LYS:HG3	2.20	0.42
31:f:14:CYS:HB3	31:f:25:VAL:CG1	2.50	0.42
2:A:1080:A:O2'	2:A:1081:U:H5'	2.19	0.42
2:A:1431:A:H2'	2:A:1432:G:C8	2.55	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:1480:C:H2'	2:A:1481:U:O4'	2.20	0.42
2:A:1705:A:O2'	2:A:1706:C:H5'	2.19	0.42
2:A:1735:A:H2'	2:A:1736:U:O4'	2.20	0.42
2:A:1864:U:H5''	2:A:2410:G:O2'	2.19	0.42
2:A:2084:C:H2'	2:A:2085:U:O4'	2.19	0.42
2:A:2386:A:O2'	2:A:2387:U:H5'	2.19	0.42
2:A:2450:A:OP1	2:A:2497:A:O2'	2.25	0.42
2:A:2484:G:H1'	13:N:123:LYS:HD2	2.01	0.42
7:F:57:LEU:HD23	7:F:57:LEU:HA	1.93	0.42
9:H:29:PHE:O	9:H:33:GLN:HB2	2.19	0.42
15:P:83:LEU:HD11	15:P:113:ALA:O	2.19	0.42
18:S:34:GLU:HG2	18:S:60:LYS:HG2	2.01	0.42
28:c:36:LEU:C	28:c:48:ILE:HD12	2.45	0.42
30:e:14:PHE:O	30:e:15:LYS:HD2	2.20	0.42
2:A:62:U:H3'	2:A:63:A:C8	2.55	0.42
2:A:354:A:H2'	2:A:355:U:C6	2.55	0.42
2:A:460:A:H2'	2:A:461:C:O4'	2.20	0.42
2:A:627:A:C5	12:M:111:ILE:HD12	2.54	0.42
2:A:709:U:H2'	2:A:710:U:H6	1.83	0.42
2:A:754:U:H2'	2:A:755:U:H6	1.85	0.42
2:A:1009:A:H2'	2:A:1010:A:C8	2.55	0.42
2:A:1943:U:O4'	2:A:1945:G:H5'	2.19	0.42
2:A:2485:G:O3'	13:N:125:PRO:HB3	2.20	0.42
2:A:2697:G:H2'	2:A:2698:U:O4'	2.19	0.42
2:A:2720:U:H4'	2:A:2845:U:O2'	2.20	0.42
2:A:2736:A:O2'	2:A:2737:G:H5'	2.20	0.42
2:A:2740:A:H2'	2:A:2741:A:C8	2.55	0.42
2:A:2813:A:H2'	2:A:2814:A:H8	1.84	0.42
3:B:113:C:H2'	3:B:114:C:H6	1.84	0.42
4:C:84:ASP:OD2	4:C:87:ARG:NE	2.40	0.42
9:H:8:LYS:HE2	9:H:14:SER:HB3	2.02	0.42
18:S:73:LYS:HE3	18:S:86:GLN:HB3	2.02	0.42
19:T:17:VAL:HA	19:T:43:ALA:CB	2.49	0.42
19:T:65:ASP:O	19:T:69:LEU:HG	2.19	0.42
20:U:72:GLN:HG3	20:U:73:ARG:HD2	2.02	0.42
24:Y:22:LEU:HD23	24:Y:22:LEU:HA	1.91	0.42
26:a:21:LYS:HE3	26:a:21:LYS:HB2	1.78	0.42
2:A:464:U:H2'	2:A:465:G:O4'	2.19	0.42
2:A:635:C:O2'	2:A:639:U:OP1	2.38	0.42
2:A:659:G:O5'	6:E:95:LYS:HD3	2.20	0.42
2:A:843:G:H2'	2:A:844:A:H8	1.84	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:1056:G:H4'	2:A:1086:A:C8	2.55	0.42
2:A:1464:G:O2'	2:A:1465:G:H5'	2.20	0.42
2:A:1638:C:O2	2:A:2698:U:O2'	2.35	0.42
2:A:1683:U:H2'	2:A:1684:G:H8	1.85	0.42
2:A:2265:U:OP2	2:A:2266:A:O2'	2.16	0.42
2:A:2572:A:OP2	5:D:149:ASN:HB3	2.20	0.42
2:A:2591:C:H2'	2:A:2592:G:H8	1.85	0.42
7:F:69:LYS:HG3	7:F:82:GLY:O	2.20	0.42
8:G:18:LYS:HB3	8:G:25:THR:HB	2.01	0.42
11:L:87:LEU:HD11	11:L:112:PHE:HE1	1.85	0.42
21:V:23:GLY:HA3	21:V:36:VAL:CG1	2.50	0.42
2:A:289:G:O2'	2:A:290:U:H5'	2.20	0.41
2:A:689:A:H2'	2:A:690:G:C8	2.54	0.41
2:A:703:U:C2'	2:A:704:G:H5'	2.50	0.41
2:A:820:A:H4'	2:A:836:G:N2	2.27	0.41
2:A:851:C:O2'	26:a:43:ALA:O	2.37	0.41
2:A:1498:C:H2'	2:A:1499:C:C6	2.55	0.41
2:A:1833:C:H2'	2:A:1834:U:O4'	2.19	0.41
2:A:1886:U:O2'	2:A:1887:C:H5'	2.20	0.41
2:A:1952:A:C6	11:L:22:ILE:HB	2.55	0.41
2:A:2205:A:H2'	2:A:2206:C:C6	2.55	0.41
2:A:2526:G:O2'	2:A:2527:C:H5'	2.20	0.41
2:A:2593:U:O2'	2:A:2594:C:H5'	2.20	0.41
2:A:2712:C:OP1	2:A:2714:G:H4'	2.20	0.41
3:B:73:A:H2'	3:B:73:A:N3	2.35	0.41
11:L:66:LYS:HE2	11:L:66:LYS:HB3	1.84	0.41
27:b:29:SER:O	27:b:37:LYS:HA	2.19	0.41
2:A:76:C:O3'	25:Z:52:ARG:HG2	2.21	0.41
2:A:676:A:H62	2:A:802:A:H61	1.68	0.41
2:A:1782:U:O2'	2:A:2609:U:H5''	2.19	0.41
2:A:2564:A:C6	2:A:2647:U:H4'	2.55	0.41
4:C:25:HIS:HB3	4:C:82:GLU:HG2	2.01	0.41
2:A:1418:G:N1	2:A:1579:A:OP2	2.36	0.41
2:A:1485:U:O2'	2:A:1486:U:H5'	2.20	0.41
2:A:1771:C:O2'	2:A:1786:A:O4'	2.33	0.41
2:A:2017:U:H4'	27:b:5:GLN:O	2.20	0.41
2:A:2037:A:H2'	2:A:2038:G:C8	2.55	0.41
2:A:2404:U:O2'	2:A:2405:G:H5'	2.21	0.41
2:A:2548:U:O2	11:L:23:LYS:NZ	2.53	0.41
2:A:2694:G:H2'	2:A:2695:U:O4'	2.20	0.41
3:B:70:C:H2'	3:B:71:C:H6	1.85	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:E:128:ALA:HB1	6:E:129:PRO:HD2	2.02	0.41
8:G:91:GLY:HA3	8:G:94:TYR:CD2	2.55	0.41
8:G:168:VAL:O	8:G:168:VAL:HG13	2.20	0.41
16:Q:33:VAL:HB	16:Q:38:LYS:HG2	2.03	0.41
19:T:70:LYS:O	19:T:107:VAL:HA	2.20	0.41
30:e:29:LEU:O	30:e:33:LEU:HD23	2.20	0.41
1:1:74:C:H4'	2:A:2602[B]:A:OP1	2.20	0.41
2:A:56:A:H2'	2:A:57:C:O4'	2.20	0.41
2:A:371:A:H3'	2:A:371:A:OP1	2.20	0.41
2:A:441:U:O2'	2:A:442:G:H5'	2.19	0.41
2:A:483:A:O2'	21:V:57:GLY:N	2.40	0.41
2:A:917:A:H5''	2:A:2268:A:N1	2.35	0.41
2:A:1312:U:H4'	2:A:1313:U:O5'	2.21	0.41
2:A:1391:U:H2'	2:A:1393:A:OP2	2.20	0.41
2:A:2323:G:C2'	2:A:2324:U:H5'	2.50	0.41
2:A:2861:U:H2'	2:A:2862:G:C8	2.56	0.41
2:A:2881:U:O2'	2:A:2882:A:H5'	2.19	0.41
7:F:3:LYS:O	7:F:3:LYS:HD3	2.20	0.41
19:T:72:THR:HG21	19:T:108:SER:HB3	2.02	0.41
2:A:26:G:O2'	2:A:27:G:O4'	2.36	0.41
2:A:355:U:O2'	2:A:356:G:H5'	2.20	0.41
2:A:924:G:H2'	2:A:925:A:C8	2.54	0.41
2:A:1028:A:N3	2:A:2486:C:O2'	2.41	0.41
2:A:1130:U:C2	5:D:152:PRO:HB3	2.55	0.41
2:A:1178:C:H2'	2:A:1179:G:H8	1.86	0.41
2:A:1401:G:H2'	2:A:1402:U:O4'	2.21	0.41
2:A:1837:C:O2'	2:A:1927:A:H1'	2.20	0.41
2:A:2050:C:O2'	2:A:2051:A:H5'	2.20	0.41
2:A:2197:U:H1'	2:A:2198:A:C8	2.55	0.41
2:A:2545:G:H2'	2:A:2546:U:O4'	2.20	0.41
2:A:2662:A:H2'	2:A:2663:G:O4'	2.20	0.41
3:B:7:G:O2'	3:B:8:C:H5'	2.20	0.41
10:K:60:ASP:HB3	10:K:97:PRO:CB	2.50	0.41
21:V:12:ILE:CG2	21:V:80:ALA:HB2	2.49	0.41
25:Z:6:LEU:HB2	25:Z:56:LEU:CD1	2.49	0.41
28:c:35:GLU:HA	28:c:49:TYR:O	2.19	0.41
2:A:45:G:C4'	2:A:46:G:H5'	2.51	0.41
2:A:191:A:O2'	2:A:678:C:O2'	2.31	0.41
2:A:231:A:O2'	2:A:232:G:H5'	2.20	0.41
2:A:409:G:H2'	2:A:410:G:C8	2.56	0.41
2:A:491:G:O6	19:T:49:LYS:NZ	2.48	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:1068:G:O2'	2:A:1096:A:H1'	2.21	0.41
2:A:1825:U:H2'	2:A:1826:G:H8	1.86	0.41
2:A:2584:U:H2'	2:A:2585:U:C6	2.55	0.41
2:A:2623:G:H4'	2:A:2825:G:H8	1.86	0.41
2:A:2627:G:N2	2:A:2777:G:OP2	2.52	0.41
2:A:2776:A:H4'	2:A:2777:G:O5'	2.20	0.41
11:L:108:ARG:HA	11:L:116:ILE:HD11	2.02	0.41
12:M:121:THR:HG23	12:M:141:LYS:O	2.21	0.41
25:Z:12:GLU:HG2	25:Z:13:GLU:N	2.36	0.41
2:A:5:A:O2'	2:A:6:A:H5'	2.20	0.41
2:A:134:G:H2'	2:A:135:U:C6	2.55	0.41
2:A:1007:C:OP2	2:A:1008:A:O2'	2.35	0.41
2:A:1296:G:O2'	2:A:1297:C:H5'	2.21	0.41
2:A:1406:U:H2'	2:A:1407:G:C8	2.56	0.41
2:A:1632:A:H2'	2:A:1633:G:O4'	2.21	0.41
2:A:1636:U:H2'	2:A:1637:A:H8	1.85	0.41
2:A:1663:G:O2'	2:A:2686:G:H4'	2.21	0.41
2:A:1789:A:H2'	2:A:1790:C:O4'	2.19	0.41
2:A:2034:U:H2'	2:A:2035:G:C8	2.55	0.41
2:A:2295:C:OP2	15:P:10:ARG:HD3	2.20	0.41
2:A:2391:G:OP2	30:e:32:ILE:HG21	2.19	0.41
8:G:37:LEU:HD11	8:G:72:LEU:HD11	2.01	0.41
8:G:170:ARG:NH2	31:f:29:ALA:HA	2.35	0.41
16:Q:32:VAL:O	16:Q:32:VAL:HG23	2.20	0.41
19:T:5:ALA:HB2	19:T:54:ALA:HA	2.03	0.41
29:d:16:HIS:HB2	29:d:44:VAL:HG21	2.01	0.41
30:e:62:LEU:HB3	30:e:65:ALA:CB	2.50	0.41
2:A:158:U:H3	2:A:168:G:H1	1.67	0.41
2:A:774:G:O2'	2:A:775:G:OP2	2.36	0.41
2:A:1260:A:H2'	2:A:1261:C:O4'	2.20	0.41
2:A:1410:G:H2'	2:A:1411:U:H6	1.86	0.41
2:A:1420:A:O2'	2:A:2211:A:N7	2.45	0.41
2:A:1466:U:O3'	2:A:1546:G:O2'	2.38	0.41
2:A:1989:G:O2'	2:A:1990:C:H5'	2.21	0.41
2:A:2065:C:H2'	2:A:2066:C:H6	1.86	0.41
2:A:2308:G:H2'	2:A:2308:G:N3	2.35	0.41
2:A:2654:A:N1	2:A:2665:A:H5''	2.35	0.41
5:D:37:VAL:HG22	5:D:48:ILE:HG22	2.02	0.41
6:E:127:GLU:CG	6:E:133:LEU:HD13	2.51	0.41
9:H:4:ILE:HG13	9:H:18:GLN:CB	2.51	0.41
10:K:88:THR:OG1	10:K:89:PHE:N	2.51	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:L:24:VAL:CG1	11:L:30:ARG:HG2	2.51	0.41
11:L:43:ILE:CD1	11:L:52:VAL:HB	2.51	0.41
15:P:71:ALA:HB1	15:P:106:LEU:HB2	2.02	0.41
22:W:86:LEU:HD13	22:W:89:ILE:HD11	2.01	0.41
2:A:50:U:H3'	2:A:51:G:C5'	2.51	0.41
2:A:184:C:H2'	2:A:185:G:C8	2.56	0.41
2:A:247:G:OP2	2:A:249:C:N4	2.54	0.41
2:A:307:G:N2	2:A:309:A:H3'	2.36	0.41
2:A:340:A:H2'	2:A:341:C:O4'	2.21	0.41
2:A:383:C:H5''	2:A:385:C:OP2	2.20	0.41
2:A:680:C:O2'	2:A:681:G:H5'	2.20	0.41
2:A:841:G:O2'	2:A:842:U:H5'	2.21	0.41
2:A:967:U:H2'	2:A:968:C:H6	1.85	0.41
2:A:1112:G:H2'	2:A:1113:U:O4'	2.21	0.41
2:A:1114:C:H2'	2:A:1115:G:H8	1.85	0.41
2:A:1310:G:O2'	2:A:1311:G:H5'	2.21	0.41
2:A:1408:G:H2'	2:A:1409:U:H6	1.86	0.41
2:A:1412:U:H2'	2:A:1413:A:H8	1.86	0.41
2:A:1450:G:H2'	2:A:1451:C:C6	2.56	0.41
2:A:1485:U:H6	2:A:1485:U:H5'	1.86	0.41
2:A:1631:G:H1'	2:A:1635:A:H61	1.85	0.41
2:A:1746:A:H2'	2:A:1747:U:H6	1.81	0.41
2:A:1833:C:H2'	2:A:1834:U:C6	2.56	0.41
2:A:2347:C:O2'	28:c:21:TYR:OH	2.36	0.41
2:A:2350:C:H2'	2:A:2351:G:O4'	2.20	0.41
2:A:2846:G:OP1	16:Q:53:ARG:NH1	2.54	0.41
2:A:2888:C:H2'	2:A:2889:C:H6	1.85	0.41
3:B:40:U:H1'	3:B:45:A:N6	2.35	0.41
5:D:184:ARG:HB3	5:D:186:LEU:HD13	2.02	0.41
8:G:42:GLU:HG2	8:G:44:LYS:HE3	2.02	0.41
11:L:71:ARG:HD3	11:L:71:ARG:HA	1.84	0.41
13:N:21:ALA:CB	13:N:100:LYS:HB2	2.51	0.41
14:O:49:GLU:HB2	14:O:50:PRO:HD3	2.02	0.41
14:O:86:ARG:HD3	14:O:117:ASP:HB3	2.03	0.41
16:Q:21:ARG:HD3	16:Q:113:ARG:NH2	2.36	0.41
21:V:28:VAL:C	21:V:29:LEU:HD23	2.45	0.41
28:c:34:LEU:N	28:c:51:GLU:OE1	2.49	0.41
30:e:4:ILE:CG2	30:e:63:PRO:HG3	2.50	0.41
2:A:436:C:O2'	2:A:437:U:H5'	2.20	0.41
2:A:473:G:O2'	2:A:474:G:H5'	2.21	0.41
2:A:554:U:C2'	2:A:555:G:H5'	2.50	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:939:G:O2'	2:A:940:G:H5'	2.21	0.41
2:A:1347:A:H2'	2:A:1348:C:H5'	2.02	0.41
2:A:1430:G:H2'	2:A:1431:A:H8	1.85	0.41
2:A:1624:U:O2'	2:A:1625:C:H5'	2.21	0.41
2:A:2060:A:O4'	2:A:2502:G:H1'	2.20	0.41
2:A:2196:C:O2'	2:A:2197:U:H5'	2.20	0.41
2:A:2266:A:H4'	2:A:2267:A:O5'	2.21	0.41
2:A:2322:A:H2'	2:A:2323:G:C8	2.56	0.41
2:A:2694:G:H2'	2:A:2695:U:C6	2.56	0.41
3:B:96:G:O2'	3:B:97:C:H5'	2.21	0.41
9:H:33:GLN:HE21	9:H:33:GLN:HB3	1.68	0.41
14:O:20:MET:HE2	14:O:20:MET:HB3	1.99	0.41
14:O:24:MET:HG2	14:O:44:LEU:HD22	2.02	0.41
14:O:79:LEU:HD23	14:O:83:LEU:HD12	2.03	0.41
2:A:26:G:H1'	2:A:514:A:H61	1.86	0.40
2:A:272:A:H2'	2:A:273:G:H8	1.84	0.40
2:A:516:C:O2'	2:A:517:C:H5'	2.21	0.40
2:A:588:U:H2'	2:A:589:U:C6	2.56	0.40
2:A:598:U:H2'	2:A:599:A:C8	2.57	0.40
2:A:745:G:C2'	2:A:746:U:H5'	2.51	0.40
2:A:850:U:O2'	26:a:23:THR:HA	2.21	0.40
2:A:1468:U:H2'	2:A:1522:A:N6	2.36	0.40
2:A:2099:U:H2'	2:A:2100:G:C8	2.56	0.40
2:A:2258:C:O2'	2:A:2426:A:H4'	2.20	0.40
2:A:2294:G:H5'	15:P:98:GLN:HE22	1.87	0.40
2:A:2472:G:O6	2:A:2475:C:H2'	2.21	0.40
2:A:2660:A:H2'	2:A:2661:G:C8	2.56	0.40
2:A:2783:U:O2'	2:A:2784:U:H5'	2.21	0.40
3:B:30:C:C2'	3:B:31:C:H5'	2.51	0.40
8:G:73:ASN:O	8:G:76:VAL:HG22	2.21	0.40
19:T:83:LYS:HD3	19:T:95:ARG:HE	1.86	0.40
21:V:74:ASN:O	21:V:78:GLY:N	2.45	0.40
24:Y:40:VAL:O	24:Y:44:LYS:N	2.53	0.40
2:A:429:A:H2'	2:A:430:A:N9	2.36	0.40
2:A:511:U:H4'	2:A:1235:G:H4'	2.02	0.40
2:A:595:C:H2'	2:A:596:U:C6	2.57	0.40
2:A:644:A:C2	2:A:2369:A:H1'	2.57	0.40
2:A:1072:C:H2'	2:A:1093:G:O6	2.22	0.40
2:A:1182:G:H2'	2:A:1183:U:O4'	2.21	0.40
2:A:1190:G:H2'	2:A:1191:G:C8	2.56	0.40
2:A:1223:G:OP2	18:S:90:ARG:NH1	2.51	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:2430:A:O2'	2:A:2431:U:H5'	2.21	0.40
2:A:2492:U:H2'	2:A:2493:U:C6	2.56	0.40
2:A:2508:G:O3'	2:A:2555:U:H5'	2.21	0.40
5:D:184:ARG:CB	5:D:186:LEU:HD13	2.51	0.40
6:E:104:ALA:O	6:E:108:ILE:HG13	2.22	0.40
6:E:143:LEU:HD23	6:E:146:VAL:CG2	2.52	0.40
11:L:38:ILE:HD11	11:L:112:PHE:HZ	1.86	0.40
14:O:35:LYS:HD2	14:O:112:TYR:CZ	2.56	0.40
15:P:16:ARG:O	15:P:20:GLU:HG2	2.21	0.40
20:U:28:ASN:ND2	20:U:88:LYS:O	2.54	0.40
26:a:12:SER:OG	26:a:14:ILE:HG13	2.22	0.40
2:A:44:A:H2'	2:A:45:G:O4'	2.21	0.40
2:A:587:C:H4'	2:A:588:U:C6	2.56	0.40
2:A:682:G:H5'	29:d:26:ASN:ND2	2.36	0.40
2:A:704:G:H1'	2:A:727:A:H61	1.86	0.40
2:A:864:G:O2'	2:A:865:C:H5'	2.21	0.40
2:A:966:G:O4'	2:A:2267:A:N6	2.54	0.40
2:A:1079:C:H2'	2:A:1080:A:H8	1.84	0.40
2:A:1426:G:OP2	2:A:1427:A:O2'	2.17	0.40
2:A:1651:G:C5'	14:O:39:PRO:HG2	2.51	0.40
2:A:1666:G:C2'	2:A:1667:G:H5'	2.50	0.40
2:A:1942:C:P	2:A:1943:U:HO2'	2.35	0.40
2:A:2366:A:C2	2:A:2367:G:H1'	2.57	0.40
2:A:2512:C:H2'	2:A:2513:A:O4'	2.22	0.40
2:A:2757:A:H2'	2:A:2757:A:N3	2.36	0.40
5:D:4:LEU:HD21	5:D:96:ILE:HG21	2.02	0.40
12:M:89:VAL:HG21	12:M:123:ARG:NH1	2.37	0.40
21:V:96:PHE:HB2	21:V:99:ASN:OD1	2.21	0.40
31:f:24:ARG:HG2	31:f:36:ARG:HB2	2.03	0.40
2:A:48:G:H22	2:A:177:G:P	2.45	0.40
2:A:439:A:O2'	2:A:440:C:H5'	2.20	0.40
2:A:499:U:H5''	21:V:43:LYS:HE3	2.03	0.40
2:A:1196:C:H2'	2:A:1197:G:C8	2.56	0.40
2:A:1483:G:H2'	2:A:1484:U:H6	1.83	0.40
2:A:1851:U:O2'	2:A:1852:U:H5'	2.21	0.40
2:A:2420:C:OP1	30:e:34:THR:HB	2.21	0.40
2:A:2488:G:O2'	2:A:2489:U:H5'	2.21	0.40
2:A:2756:U:H4'	2:A:2757:A:OP1	2.19	0.40
8:G:8:PRO:HB2	8:G:49:THR:HG21	2.02	0.40
11:L:92:GLU:O	11:L:92:GLU:HG3	2.21	0.40
12:M:74:THR:HA	12:M:107:PHE:O	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:O:103:ARG:HD3	14:O:110:MET:SD	2.62	0.40
21:V:4:LYS:O	21:V:94:ARG:NH2	2.51	0.40
28:c:11:LEU:HD23	28:c:51:GLU:HA	2.03	0.40
30:e:16:LYS:HA	30:e:21:GLY:O	2.21	0.40
30:e:26:HIS:NE2	30:e:48:ALA:HB2	2.37	0.40
2:A:103:A:H3'	2:A:104:A:H8	1.87	0.40
2:A:736:C:H2'	2:A:737:C:H6	1.87	0.40
2:A:858:G:H3'	2:A:859:G:C8	2.56	0.40
2:A:1449:G:O2'	2:A:1450:G:H5'	2.21	0.40
2:A:1491:G:H4'	4:C:100:GLU:OE1	2.20	0.40
2:A:1604:C:O2'	2:A:1610:A:N1	2.55	0.40
2:A:1668:A:H4'	2:A:1669:A:O5'	2.22	0.40
2:A:1800:C:C2'	4:C:146:MET:HE1	2.51	0.40
2:A:2090:A:O2'	2:A:2091:C:H5'	2.22	0.40
2:A:2244:U:H2'	2:A:2245:U:O4'	2.21	0.40
2:A:2373:G:H2'	2:A:2374:C:C6	2.57	0.40
2:A:2543:G:H2'	2:A:2544:G:C8	2.56	0.40
2:A:2732:G:H3'	2:A:2733:A:O4'	2.21	0.40
2:A:2861:U:H2'	2:A:2862:G:H8	1.87	0.40
4:C:81:LEU:HD22	4:C:92:ALA:HB2	2.03	0.40
4:C:118:SER:HB2	4:C:129:THR:HB	2.04	0.40
6:E:121:VAL:O	6:E:189:THR:HA	2.20	0.40
14:O:29:VAL:CG1	14:O:75:ILE:HG23	2.49	0.40
29:d:25:LYS:HD3	29:d:29:GLN:NE2	2.36	0.40

There are no symmetry-related clashes.

5.3 Torsion angles [i](#)

5.3.1 Protein backbone [i](#)

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles
4	C	269/273 (98%)	258 (96%)	11 (4%)	0	100 100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
5	D	207/209 (99%)	203 (98%)	4 (2%)	0	100	100
6	E	199/201 (99%)	197 (99%)	2 (1%)	0	100	100
7	F	175/179 (98%)	172 (98%)	3 (2%)	0	100	100
8	G	174/177 (98%)	170 (98%)	4 (2%)	0	100	100
9	H	48/149 (32%)	45 (94%)	3 (6%)	0	100	100
10	K	140/142 (99%)	137 (98%)	3 (2%)	0	100	100
11	L	121/123 (98%)	118 (98%)	3 (2%)	0	100	100
12	M	142/144 (99%)	135 (95%)	7 (5%)	0	100	100
13	N	134/136 (98%)	134 (100%)	0	0	100	100
14	O	123/127 (97%)	119 (97%)	4 (3%)	0	100	100
15	P	115/117 (98%)	115 (100%)	0	0	100	100
16	Q	112/115 (97%)	110 (98%)	2 (2%)	0	100	100
17	R	115/118 (98%)	114 (99%)	1 (1%)	0	100	100
18	S	101/103 (98%)	99 (98%)	2 (2%)	0	100	100
19	T	108/110 (98%)	106 (98%)	2 (2%)	0	100	100
20	U	93/100 (93%)	93 (100%)	0	0	100	100
21	V	100/104 (96%)	95 (95%)	5 (5%)	0	100	100
22	W	92/94 (98%)	91 (99%)	1 (1%)	0	100	100
23	X	74/85 (87%)	73 (99%)	1 (1%)	0	100	100
24	Y	75/78 (96%)	75 (100%)	0	0	100	100
25	Z	60/63 (95%)	60 (100%)	0	0	100	100
26	a	56/59 (95%)	55 (98%)	1 (2%)	0	100	100
27	b	54/57 (95%)	54 (100%)	0	0	100	100
28	c	49/55 (89%)	49 (100%)	0	0	100	100
29	d	44/46 (96%)	42 (96%)	2 (4%)	0	100	100
30	e	62/65 (95%)	56 (90%)	6 (10%)	0	100	100
31	f	36/38 (95%)	35 (97%)	1 (3%)	0	100	100
All	All	3078/3267 (94%)	3010 (98%)	68 (2%)	0	100	100

There are no Ramachandran outliers to report.

5.3.2 Protein sidechains ⓘ

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
4	C	216/218 (99%)	216 (100%)	0	100	100
5	D	164/164 (100%)	164 (100%)	0	100	100
6	E	165/165 (100%)	165 (100%)	0	100	100
7	F	148/150 (99%)	148 (100%)	0	100	100
8	G	137/138 (99%)	137 (100%)	0	100	100
9	H	40/114 (35%)	40 (100%)	0	100	100
10	K	116/116 (100%)	116 (100%)	0	100	100
11	L	104/104 (100%)	104 (100%)	0	100	100
12	M	103/103 (100%)	103 (100%)	0	100	100
13	N	109/109 (100%)	109 (100%)	0	100	100
14	O	102/103 (99%)	102 (100%)	0	100	100
15	P	87/87 (100%)	87 (100%)	0	100	100
16	Q	99/100 (99%)	99 (100%)	0	100	100
17	R	89/90 (99%)	89 (100%)	0	100	100
18	S	84/84 (100%)	84 (100%)	0	100	100
19	T	93/93 (100%)	93 (100%)	0	100	100
20	U	82/84 (98%)	82 (100%)	0	100	100
21	V	83/85 (98%)	83 (100%)	0	100	100
22	W	78/78 (100%)	78 (100%)	0	100	100
23	X	57/63 (90%)	57 (100%)	0	100	100
24	Y	67/68 (98%)	67 (100%)	0	100	100
25	Z	54/55 (98%)	54 (100%)	0	100	100
26	a	48/49 (98%)	48 (100%)	0	100	100
27	b	47/48 (98%)	47 (100%)	0	100	100
28	c	45/49 (92%)	45 (100%)	0	100	100
29	d	38/38 (100%)	38 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
30	e	51/52 (98%)	51 (100%)	0	100	100
31	f	34/34 (100%)	34 (100%)	0	100	100
All	All	2540/2641 (96%)	2540 (100%)	0	100	100

There are no protein residues with a non-rotameric sidechain to report.

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (38) such sidechains are listed below:

Mol	Chain	Res	Type
4	C	15	HIS
4	C	86	ASN
4	C	134	ASN
5	D	32	ASN
5	D	42	ASN
5	D	94	GLN
5	D	173	GLN
6	E	62	GLN
6	E	115	GLN
7	F	52	ASN
7	F	81	GLN
8	G	22	GLN
8	G	45	HIS
8	G	115	HIS
9	H	2	GLN
9	H	11	ASN
9	H	20	ASN
9	H	33	GLN
10	K	47	HIS
10	K	128	ASN
10	K	138	GLN
11	L	89	ASN
11	L	90	ASN
11	L	93	GLN
12	M	104	GLN
13	N	3	GLN
15	P	38	GLN
16	Q	10	GLN
16	Q	12	GLN
16	Q	56	HIS
16	Q	66	ASN
17	R	52	GLN

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Mol	Chain	Res	Type
22	W	78	GLN
23	X	76	ASN
24	Y	20	HIS
24	Y	34	HIS
25	Z	39	GLN
29	d	26	ASN

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	1	3/4 (75%)	1 (33%)	0
2	A	2786/2795 (99%)	327 (11%)	8 (0%)
3	B	119/120 (99%)	9 (7%)	0
All	All	2908/2919 (99%)	337 (11%)	8 (0%)

All (337) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	1	76	A
2	A	10	A
2	A	12	U
2	A	15	G
2	A	34	U
2	A	45	G
2	A	46	G
2	A	51	G
2	A	63	A
2	A	64	A
2	A	71	A
2	A	74	A
2	A	75	G
2	A	84	A
2	A	91	A
2	A	96	C
2	A	101	A
2	A	102	U
2	A	118	A
2	A	120	U
2	A	138	U
2	A	139	U
2	A	140	C

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Mol	Chain	Res	Type
2	A	141	G
2	A	142	A
2	A	163	C
2	A	181	A
2	A	196	A
2	A	199	A
2	A	216	A
2	A	222	A
2	A	233	A
2	A	248	G
2	A	266	G
2	A	272	A
2	A	275	C
2	A	276	U
2	A	277	G
2	A	278	A
2	A	302	C
2	A	311	A
2	A	329	G
2	A	330	A
2	A	353	C
2	A	361	G
2	A	362	A
2	A	372	G
2	A	386	G
2	A	396	G
2	A	404	A
2	A	411	G
2	A	412	A
2	A	424	G
2	A	451	U
2	A	456	C
2	A	481	G
2	A	491	G
2	A	505	A
2	A	509	C
2	A	529	A
2	A	530	G
2	A	532	A
2	A	543	G
2	A	549	G
2	A	563	A

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Mol	Chain	Res	Type
2	A	573	U
2	A	575	A
2	A	603	A
2	A	614	A
2	A	615	U
2	A	616	A
2	A	621	A
2	A	627	A
2	A	637	A
2	A	645	C
2	A	646	U
2	A	647	G
2	A	654	A
2	A	655	A
2	A	686	U
2	A	704	G
2	A	717	C
2	A	726	G
2	A	729	G
2	A	730	A
2	A	739	A
2	A	747	U
2	A	764	A
2	A	765	C
2	A	782	A
2	A	784	G
2	A	785	G
2	A	789	A
2	A	790	U
2	A	805	G
2	A	812	C
2	A	827	U
2	A	828	U
2	A	846	U
2	A	858	G
2	A	877	A
2	A	878	A
2	A	896	A
2	A	897	C
2	A	901	C
2	A	902	C
2	A	910	A

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Mol	Chain	Res	Type
2	A	914	G
2	A	931	U
2	A	946	C
2	A	961	C
2	A	974	G
2	A	983	A
2	A	996	A
2	A	1009	A
2	A	1012	U
2	A	1013	C
2	A	1026	G
2	A	1033	U
2	A	1046	A
2	A	1047	G
2	A	1070	A
2	A	1071	G
2	A	1083	U
2	A	1088	A
2	A	1112	G
2	A	1119	U
2	A	1132	U
2	A	1133	A
2	A	1135	C
2	A	1142	A
2	A	1143	A
2	A	1157	G
2	A	1167	C
2	A	1173	U
2	A	1204	A
2	A	1206	G
2	A	1211	C
2	A	1212	G
2	A	1236	G
2	A	1238	G
2	A	1250	G
2	A	1253	A
2	A	1256	G
2	A	1271	G
2	A	1272	A
2	A	1276	A
2	A	1300	G
2	A	1301	A

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Mol	Chain	Res	Type
2	A	1321	A
2	A	1325	U
2	A	1329	U
2	A	1352	U
2	A	1359	A
2	A	1365	A
2	A	1378	A
2	A	1379	U
2	A	1383	A
2	A	1395	A
2	A	1416	G
2	A	1417	C
2	A	1420	A
2	A	1421	G
2	A	1427	A
2	A	1428	C
2	A	1435	G
2	A	1437	C
2	A	1452	G
2	A	1453	A
2	A	1455	G
2	A	1482	G
2	A	1485	U
2	A	1487	U
2	A	1488	C
2	A	1491	G
2	A	1494	A
2	A	1497	U
2	A	1504	A
2	A	1507	C
2	A	1508	A
2	A	1509	A
2	A	1515	A
2	A	1522	A
2	A	1524	G
2	A	1569	A
2	A	1584	U
2	A	1585	C
2	A	1594	U
2	A	1607	C
2	A	1608	A
2	A	1626	A

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Mol	Chain	Res	Type
2	A	1634	A
2	A	1646	C
2	A	1647	U
2	A	1648	U
2	A	1674	G
2	A	1715	G
2	A	1729	U
2	A	1730	C
2	A	1737	G
2	A	1738	G
2	A	1756	G
2	A	1758	U
2	A	1764	C
2	A	1773	A
2	A	1784	A
2	A	1800	C
2	A	1801	A
2	A	1808	A
2	A	1816	C
2	A	1829	A
2	A	1847	A
2	A	1866	A
2	A	1870	C
2	A	1906	G
2	A	1913	A
2	A	1914	C
2	A	1916	A
2	A	1929	G
2	A	1936	A
2	A	1937	A
2	A	1938	A
2	A	1955	U
2	A	1965	C
2	A	1967	C
2	A	1970	A
2	A	1971	U
2	A	1972	G
2	A	1991	U
2	A	1992	G
2	A	1993	U
2	A	1997	C
2	A	2021	C

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Mol	Chain	Res	Type
2	A	2023	C
2	A	2030	A
2	A	2031	A
2	A	2033	A
2	A	2036	C
2	A	2043	C
2	A	2055	C
2	A	2056	G
2	A	2060	A
2	A	2061	G
2	A	2069	G
2	A	2077	A
2	A	2092	U
2	A	2093	G
2	A	2097	A
2	A	2198	A
2	A	2199	A
2	A	2204	G
2	A	2211	A
2	A	2212	A
2	A	2225	A
2	A	2238	G
2	A	2239	G
2	A	2279	G
2	A	2283	C
2	A	2287	A
2	A	2297	A
2	A	2305	U
2	A	2308	G
2	A	2322	A
2	A	2323	G
2	A	2325	G
2	A	2333	A
2	A	2347	C
2	A	2350	C
2	A	2383	G
2	A	2385	C
2	A	2396	G
2	A	2402	U
2	A	2406	A
2	A	2425	A
2	A	2428	G

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Mol	Chain	Res	Type
2	A	2429	G
2	A	2430	A
2	A	2431	U
2	A	2435	A
2	A	2441	U
2	A	2447	G
2	A	2448	A
2	A	2473	U
2	A	2475	C
2	A	2476	A
2	A	2484	G
2	A	2491	U
2	A	2502	G
2	A	2505	G
2	A	2518	A
2	A	2529	G
2	A	2549	G
2	A	2566	A
2	A	2567	G
2	A	2572	A
2	A	2578	G
2	A	2586	U
2	A	2609	U
2	A	2613	U
2	A	2615	U
2	A	2629	U
2	A	2630	G
2	A	2639	A
2	A	2646	C
2	A	2689	U
2	A	2690	U
2	A	2714	G
2	A	2718	G
2	A	2726	A
2	A	2729	G
2	A	2733	A
2	A	2744	G
2	A	2748	A
2	A	2757	A
2	A	2778	A
2	A	2791	G
2	A	2792	A

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Mol	Chain	Res	Type
2	A	2798	U
2	A	2820	A
2	A	2835	A
2	A	2843	G
2	A	2849	U
2	A	2850	A
2	A	2867	G
2	A	2873	A
2	A	2880	C
2	A	2883	A
2	A	2884	U
3	B	13	G
3	B	15	A
3	B	24	G
3	B	35	C
3	B	41	G
3	B	56	G
3	B	89	U
3	B	90	C
3	B	109	A

All (8) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
2	A	62	U
2	A	784	G
2	A	1275	A
2	A	1358	G
2	A	1757	A
2	A	2211	A
2	A	2474	U
2	A	2756	U

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

Of 386 ligands modelled in this entry, 383 are monoatomic - leaving 3 for Mogul analysis.

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

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
35	JJH	A	3370	-	46,47,47	1.93	9 (19%)	66,71,71	2.59	20 (30%)
34	AMP	A	3001	-	21,24,25	0.24	0	30,35,38	0.32	0
32	FME	1	101	1	8,9,10	0.96	0	8,9,11	0.95	1 (12%)

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

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
35	JJH	A	3370	-	-	11/24/50/50	0/6/6/6
34	AMP	A	3001	-	-	0/7/25/26	0/3/3/3
32	FME	1	101	1	-	2/7/9/11	-

All (9) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	A	3370	JJH	CBB-CBF	-5.47	1.40	1.48
35	A	3370	JJH	CBJ-NAM	-4.97	1.33	1.43
35	A	3370	JJH	CBA-CBD	-4.47	1.39	1.48
35	A	3370	JJH	CAT-NAK	-3.92	1.35	1.40
35	A	3370	JJH	CAX-NAK	3.54	1.40	1.34
35	A	3370	JJH	CAW-NAL	-3.16	1.34	1.41
35	A	3370	JJH	CBK-CBL	-2.77	1.48	1.52
35	A	3370	JJH	CAP-CAN	2.31	1.53	1.48
35	A	3370	JJH	CBB-CBD	-2.03	1.39	1.44

All (21) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	A	3370	JJH	CBL-OAH-CBO	-11.45	100.96	110.10
35	A	3370	JJH	CBK-NAM-CBO	-8.28	105.05	111.17
35	A	3370	JJH	OAJ-CBO-NAM	-5.72	123.50	128.80
35	A	3370	JJH	OAH-CBL-CBK	-5.41	99.26	104.50
35	A	3370	JJH	CBL-CBK-NAM	-3.99	98.08	101.85
35	A	3370	JJH	CBE-CBC-CAW	-3.97	119.84	123.35
35	A	3370	JJH	CAY-OAD-CBG	-3.88	108.01	117.63
35	A	3370	JJH	CBK-NAM-CBJ	3.18	126.88	121.36
35	A	3370	JJH	CAZ-CAW-CBC	3.12	119.79	116.53
35	A	3370	JJH	CAV-NAL-CAU	3.09	118.51	111.57
35	A	3370	JJH	CBA-CBD-CBB	2.75	119.11	115.64
35	A	3370	JJH	CAV-CAS-CAQ	-2.69	108.14	112.02
35	A	3370	JJH	OAH-CBO-OAJ	2.68	125.55	122.40
35	A	3370	JJH	CAU-CAR-CAQ	-2.65	108.20	112.02
35	A	3370	JJH	CBB-CAX-NAK	-2.61	121.30	124.42
35	A	3370	JJH	CAV-NAL-CAW	2.57	122.29	116.19
35	A	3370	JJH	CAZ-CAW-NAL	-2.30	119.24	122.59
35	A	3370	JJH	CBM-CBH-CBG	-2.27	119.80	122.86
35	A	3370	JJH	FAA-CBC-CAW	2.26	120.52	118.37
32	1	101	FME	C-CA-N	2.04	113.44	109.50
35	A	3370	JJH	OAG-CBF-OAF	-2.04	119.06	123.90

There are no chirality outliers.

All (13) torsion outliers are listed below:

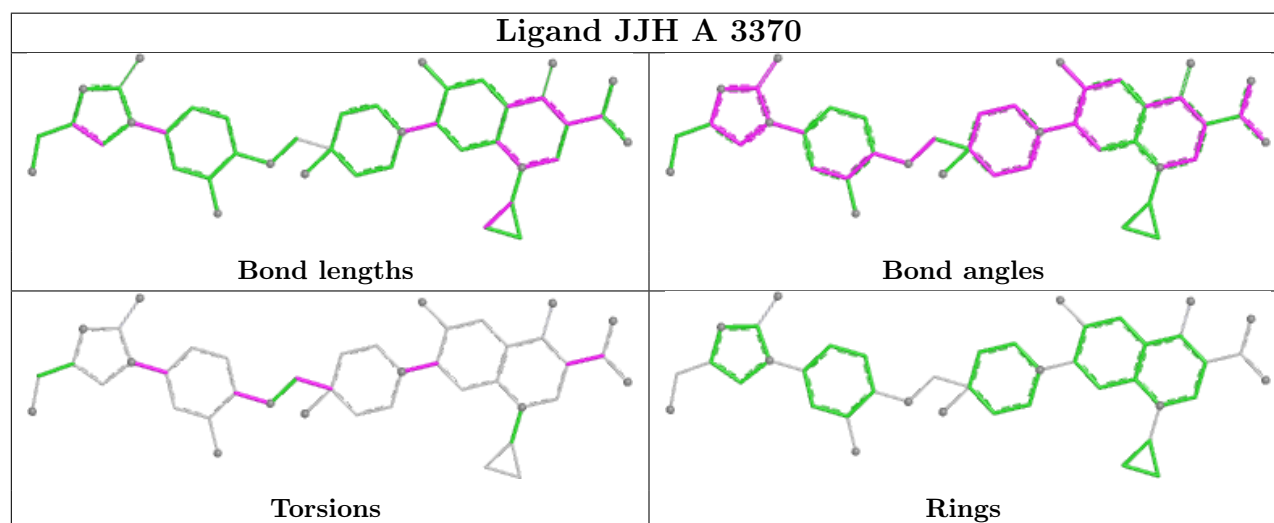
Mol	Chain	Res	Type	Atoms
32	1	101	FME	O1-CN-N-CA
32	1	101	FME	CB-CA-N-CN
35	A	3370	JJH	CAR-CAQ-CAY-OAD
35	A	3370	JJH	CAS-CAQ-CAY-OAD
35	A	3370	JJH	OAC-CAQ-CAY-OAD
35	A	3370	JJH	CBH-CBG-OAD-CAY
35	A	3370	JJH	CBC-CAW-NAL-CAU
35	A	3370	JJH	CBM-CBJ-NAM-CBO
35	A	3370	JJH	CBN-CBJ-NAM-CBO
35	A	3370	JJH	CBI-CBG-OAD-CAY
35	A	3370	JJH	CAZ-CAW-NAL-CAU
35	A	3370	JJH	CBD-CBB-CBF-OAF
35	A	3370	JJH	CBD-CBB-CBF-OAG

There are no ring outliers.

1 monomer is involved in 1 short contact:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
34	A	3001	AMP	1	0

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



5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

The following chains have linkage breaks:

Mol	Chain	Number of breaks
2	A	5

All chain breaks are listed below:

Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	A	1531:C	O3'	1540:G	P	18.12
1	A	883:G	O3'	893:C	P	17.33
1	A	2101:A	O3'	2188:U	P	16.18
1	A	545:U	O3'	548:G	P	15.10
1	A	1173:U	O3'	1177:G	P	12.82

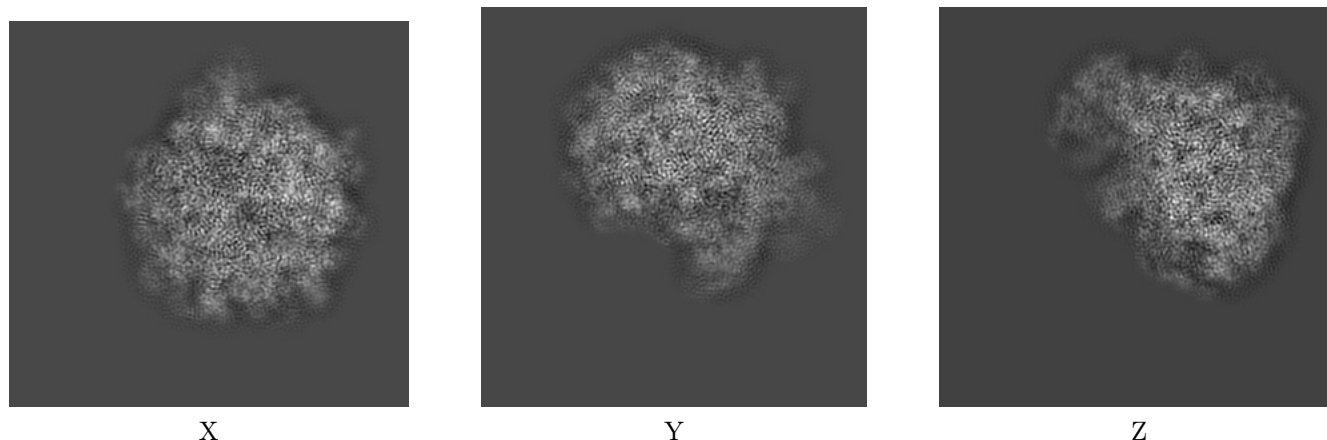
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-4638. These allow visual inspection of the internal detail of the map and identification of artifacts.

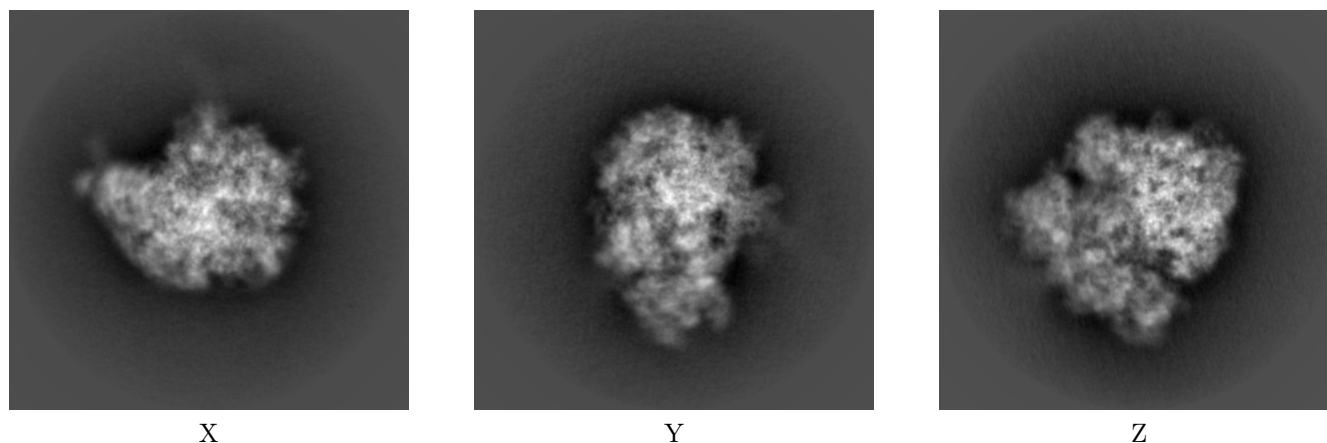
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections [i](#)

6.1.1 Primary map



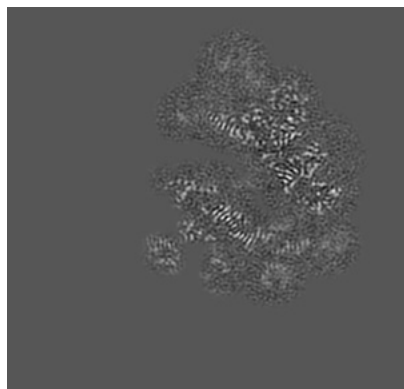
6.1.2 Raw map



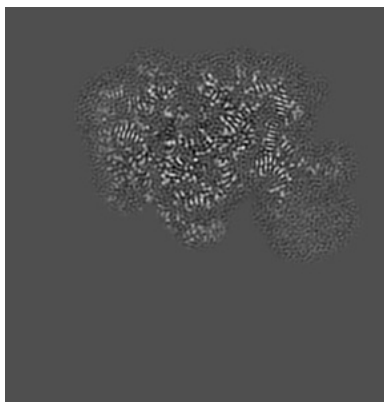
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

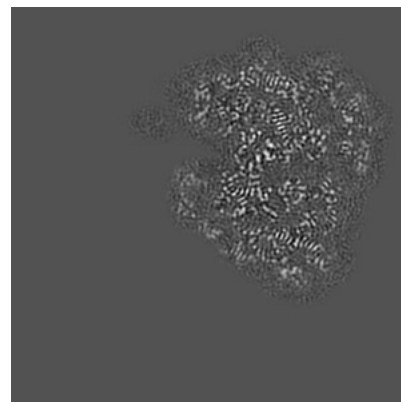
6.2.1 Primary map



X Index: 145

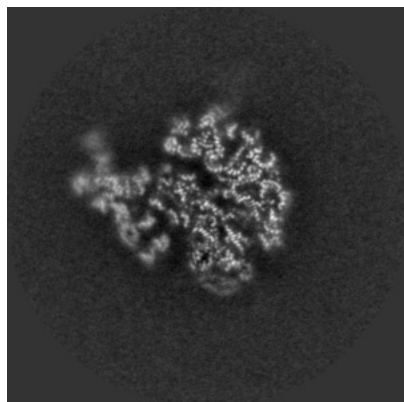


Y Index: 145

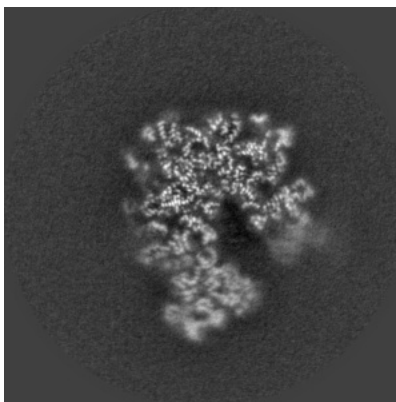


Z Index: 140

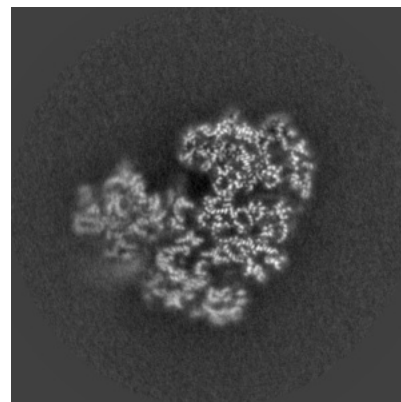
6.2.2 Raw map



X Index: 200



Y Index: 200

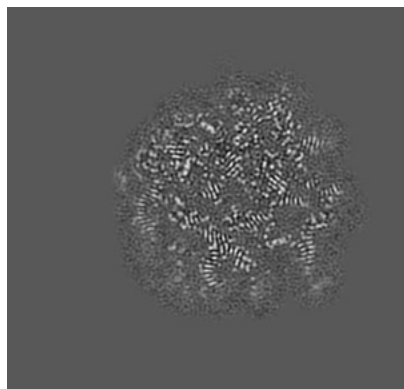


Z Index: 200

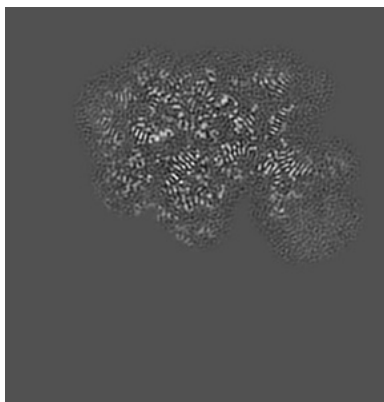
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

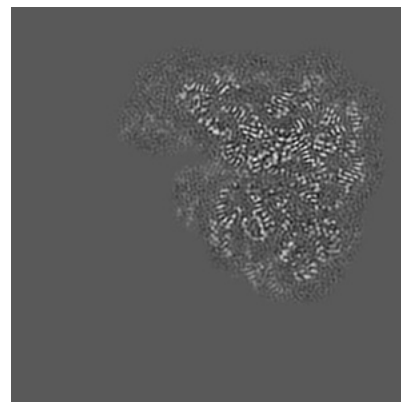
6.3.1 Primary map



X Index: 196

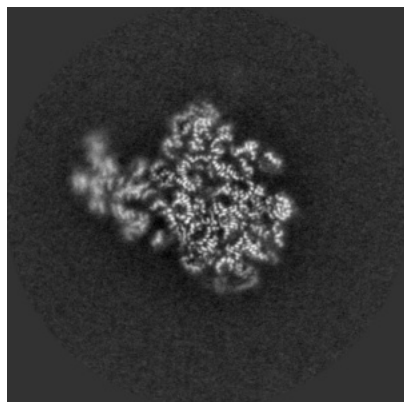


Y Index: 150

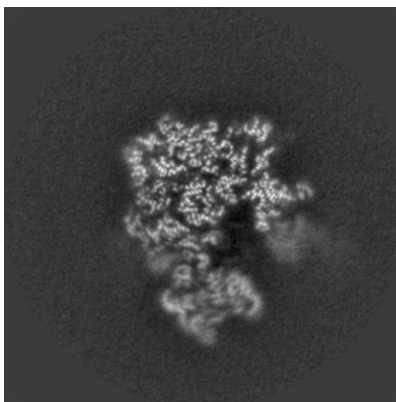


Z Index: 153

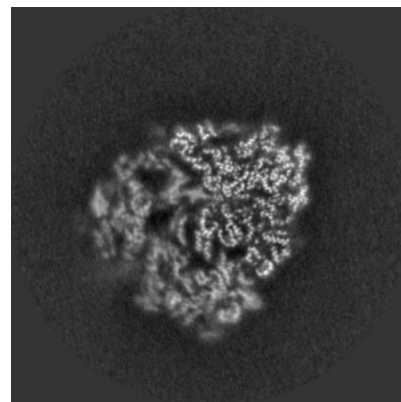
6.3.2 Raw map



X Index: 208



Y Index: 190

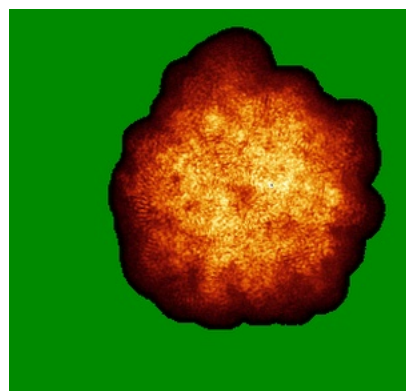


Z Index: 213

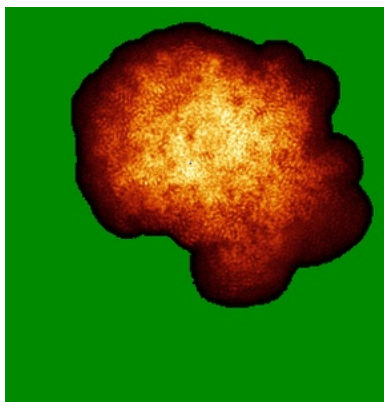
The images above show the largest variance slices of the map in three orthogonal directions.

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

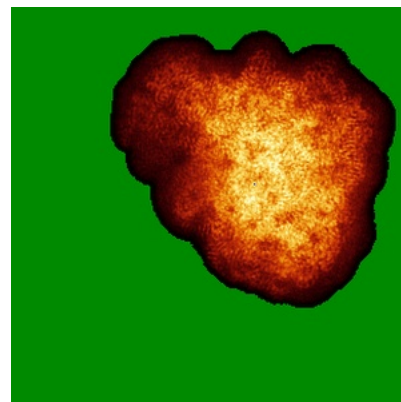
6.4.1 Primary map



X

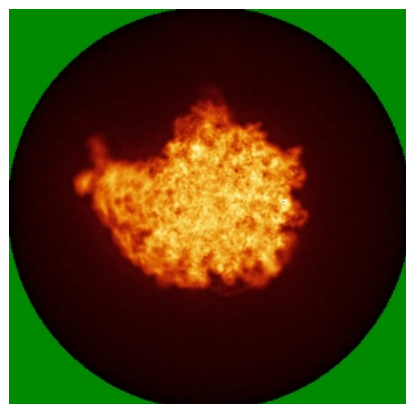


Y

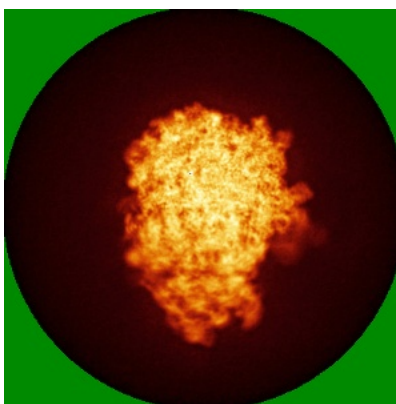


Z

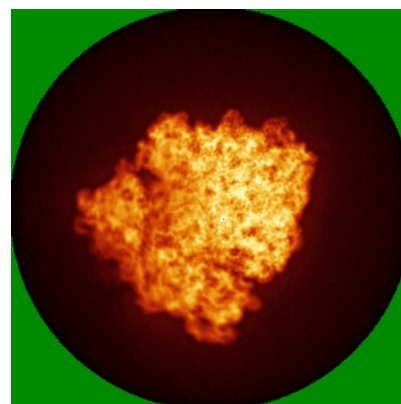
6.4.2 Raw map



X



Y

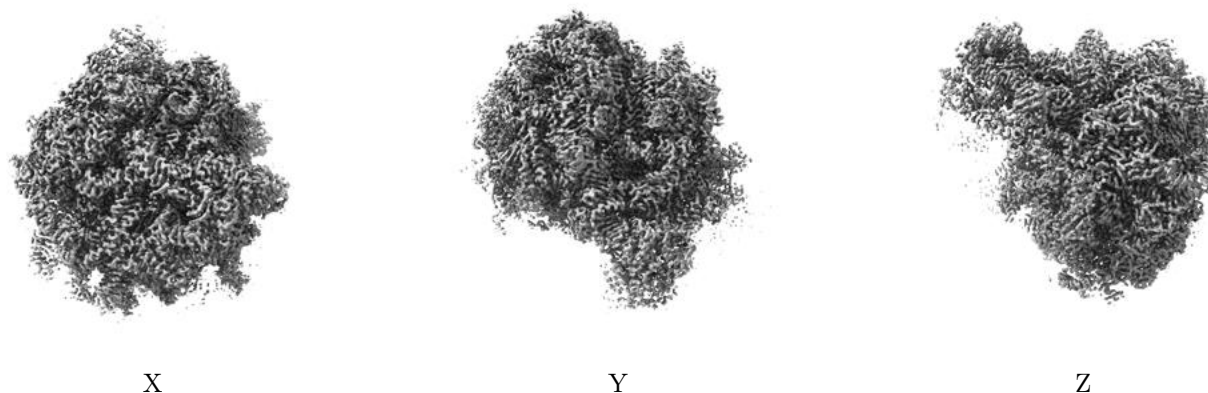


Z

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

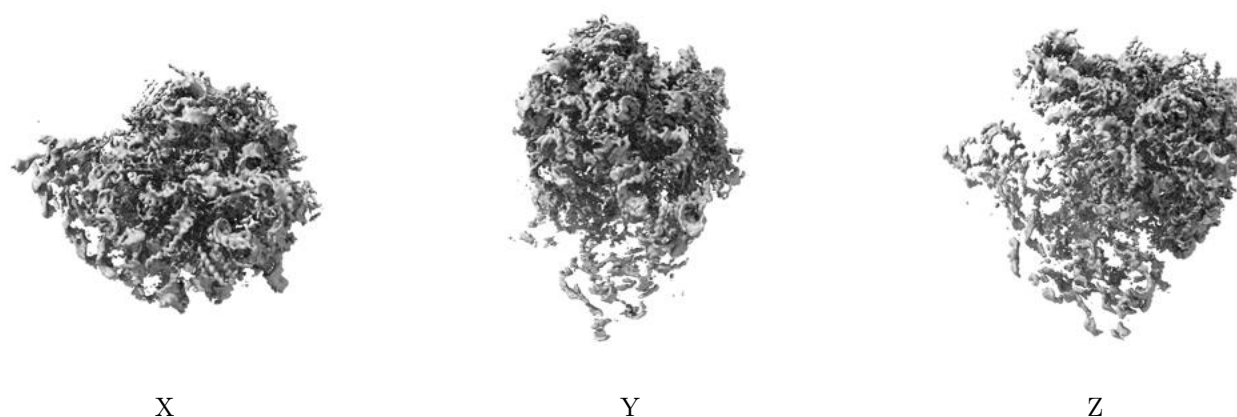
6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.1. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

6.5.2 Raw map



These images show the 3D surface of the raw map. The raw map's contour level was selected so that its surface encloses the same volume as the primary map does at its recommended contour level.

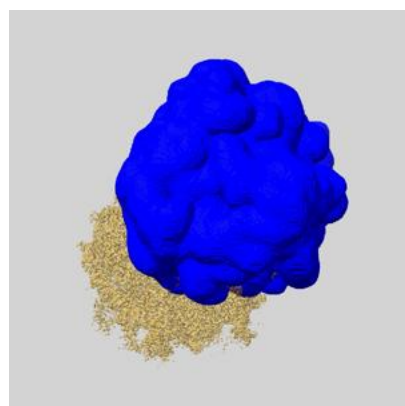
6.6 Mask visualisation [i](#)

This section shows the 3D surface view of the primary map at 50% transparency overlaid with the specified mask at 0% transparency

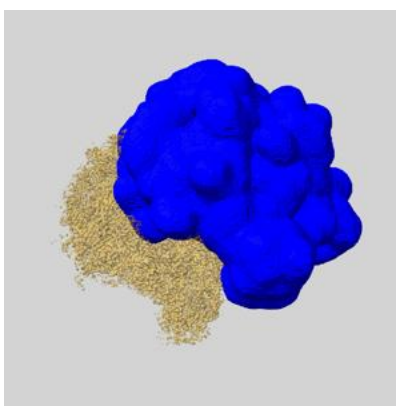
A mask typically either:

- Encompasses the whole structure
- Separates out a domain, a functional unit, a monomer or an area of interest from a larger structure

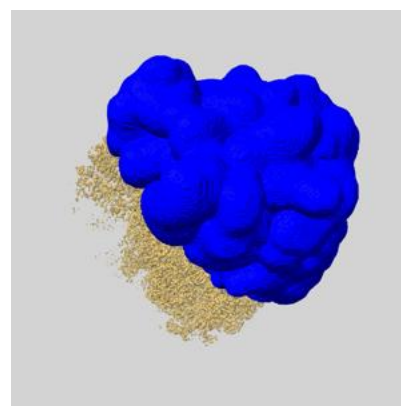
6.6.1 emd_4638_msk_1.map [i](#)



X



Y

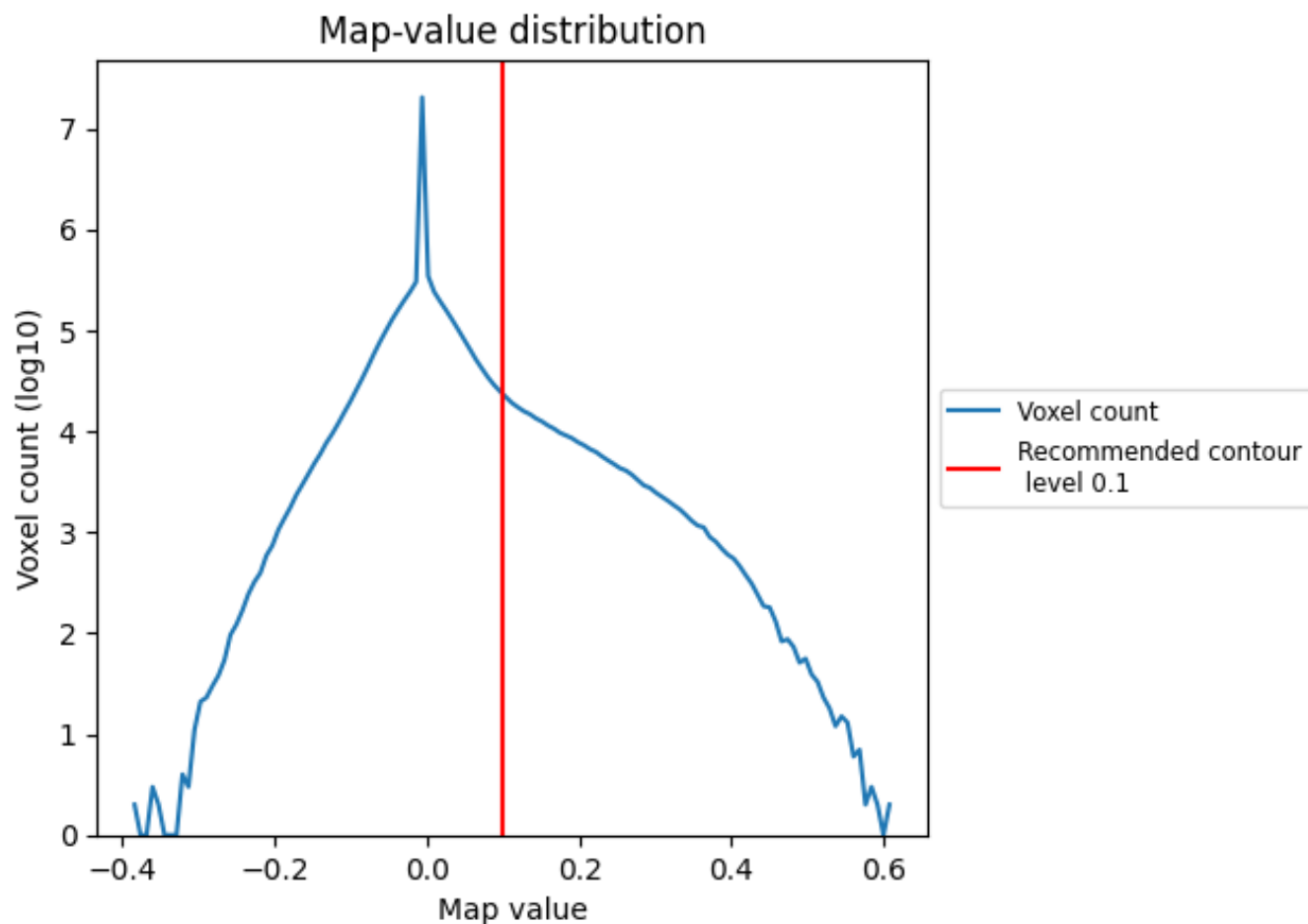


Z

7 Map analysis [i](#)

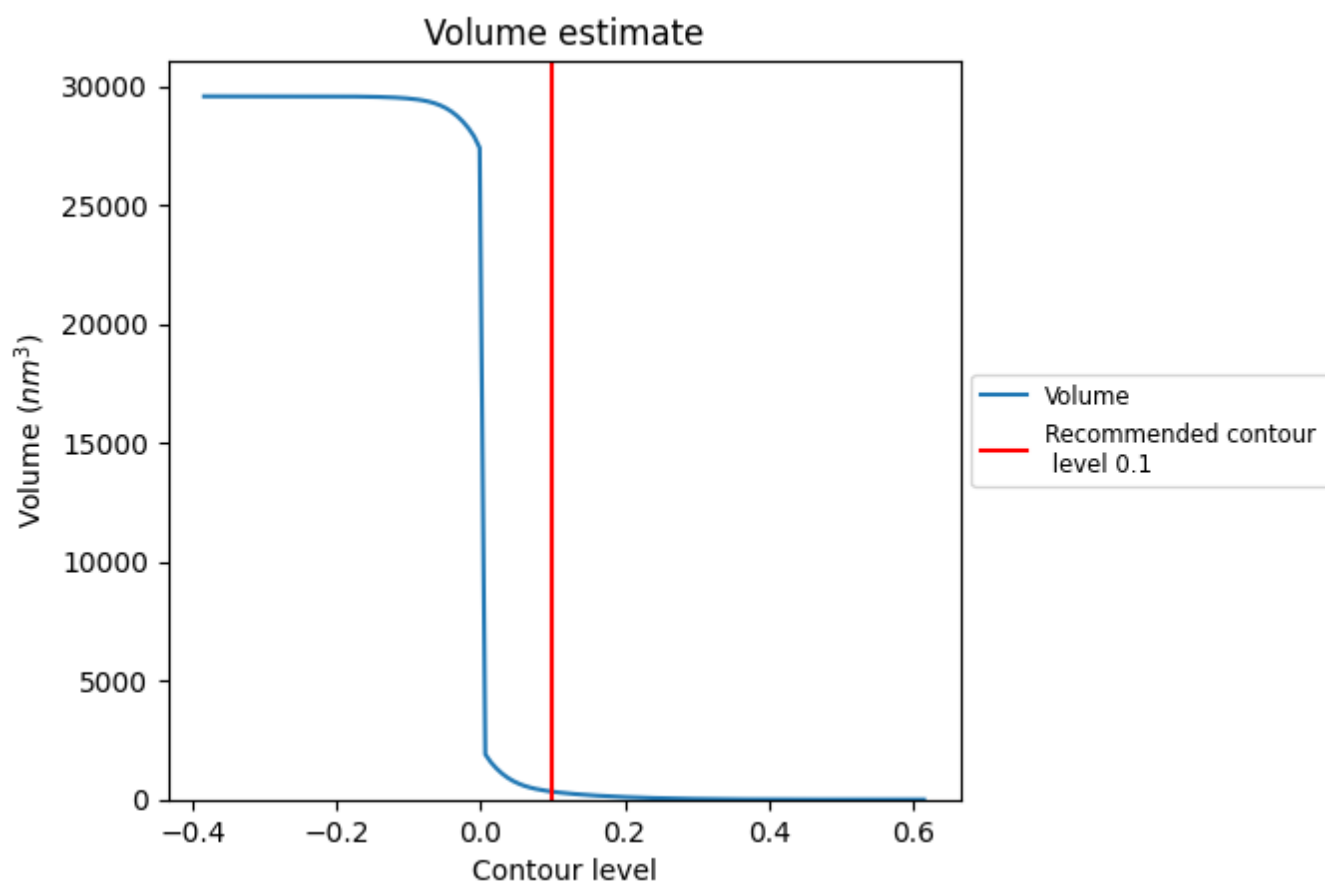
This section contains the results of statistical analysis of the map.

7.1 Map-value distribution [i](#)



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

7.2 Volume estimate [i](#)



The volume at the recommended contour level is 329 nm³; this corresponds to an approximate mass of 297 kDa.

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

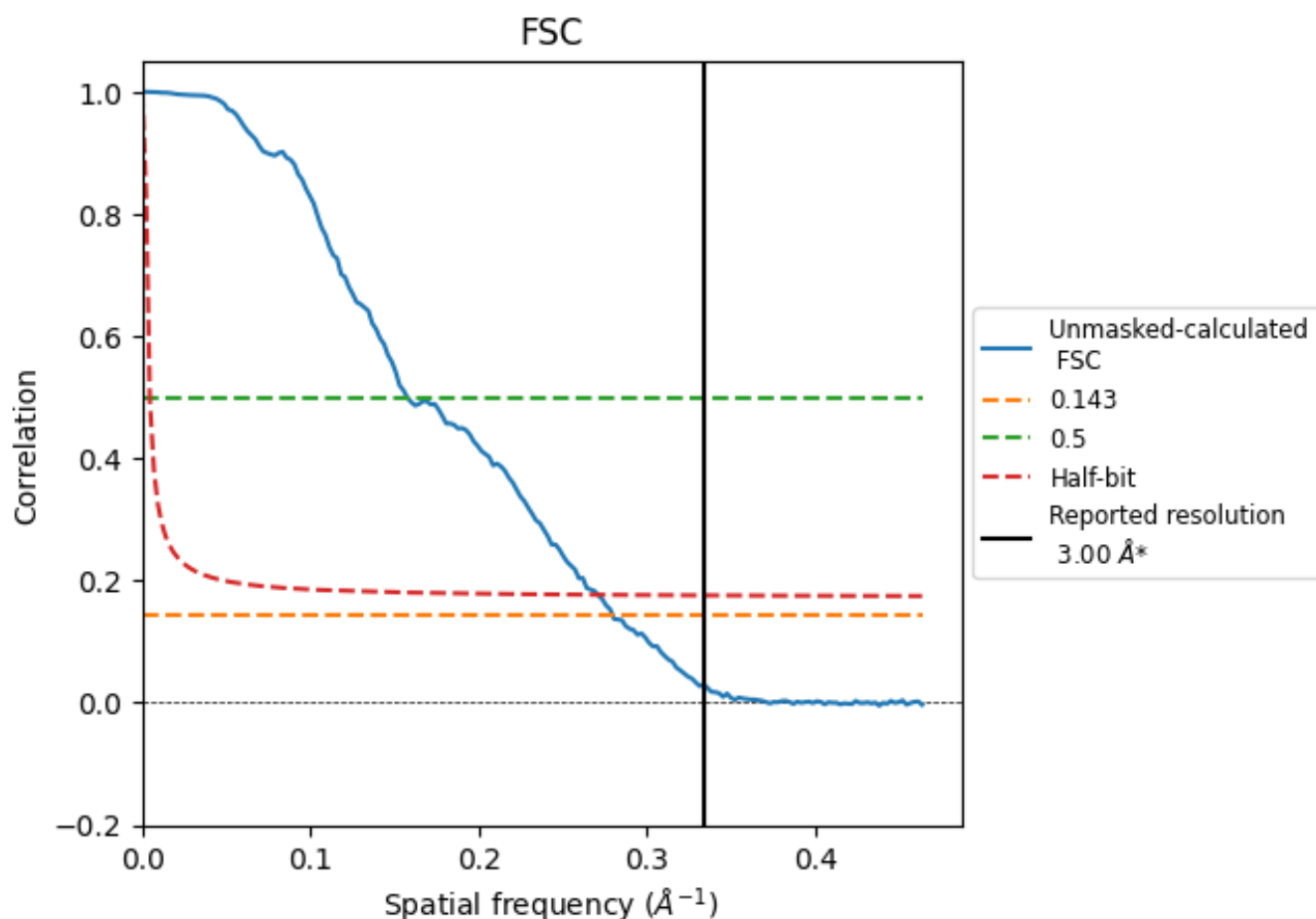
7.3 Rotationally averaged power spectrum [i](#)

This section was not generated. The rotationally averaged power spectrum is only generated for cubic maps.

8 Fourier-Shell correlation [i](#)

Fourier-Shell Correlation (FSC) is the most commonly used method to estimate the resolution of single-particle and subtomogram-averaged maps. The shape of the curve depends on the imposed symmetry, mask and whether or not the two 3D reconstructions used were processed from a common reference. The reported resolution is shown as a black line. A curve is displayed for the half-bit criterion in addition to lines showing the 0.143 gold standard cut-off and 0.5 cut-off.

8.1 FSC [i](#)



*Reported resolution corresponds to spatial frequency of 0.333 $\text{\AA}^{-1}$

8.2 Resolution estimates [i](#)

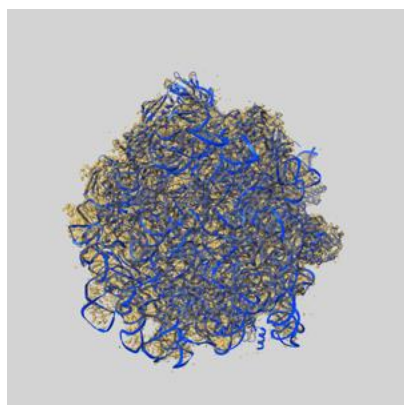
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	3.00	-	-
Author-provided FSC curve	-	-	-
Unmasked-calculated*	3.58	6.35	3.69

*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps. The value from deposited half-maps intersecting FSC 0.143 CUT-OFF 3.58 differs from the reported value 3.0 by more than 10 %

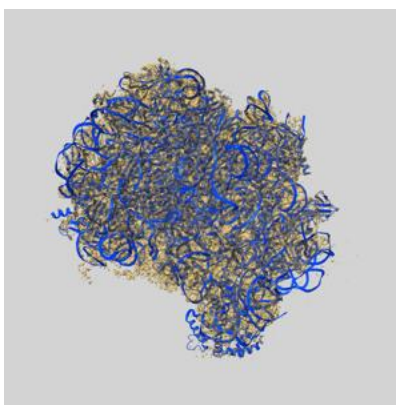
9 Map-model fit [i](#)

This section contains information regarding the fit between EMDB map EMD-4638 and PDB model 6QUL. Per-residue inclusion information can be found in section 3 on page 12.

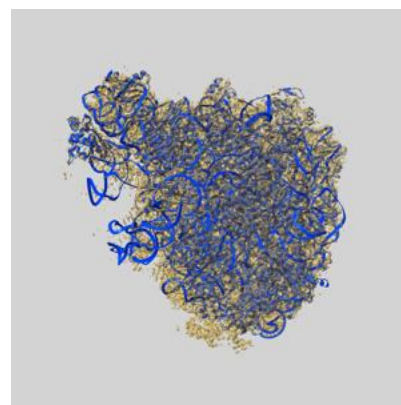
9.1 Map-model overlay [i](#)



X



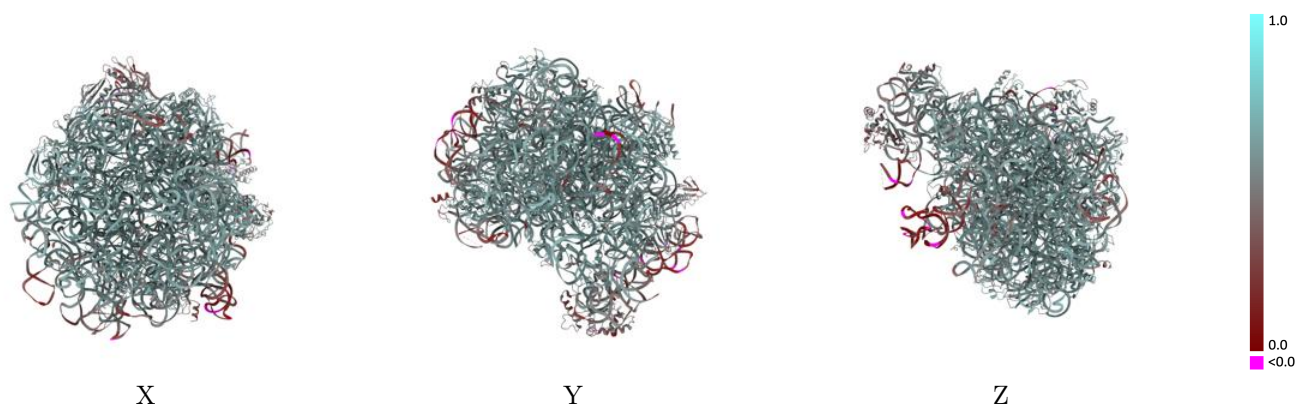
Y



Z

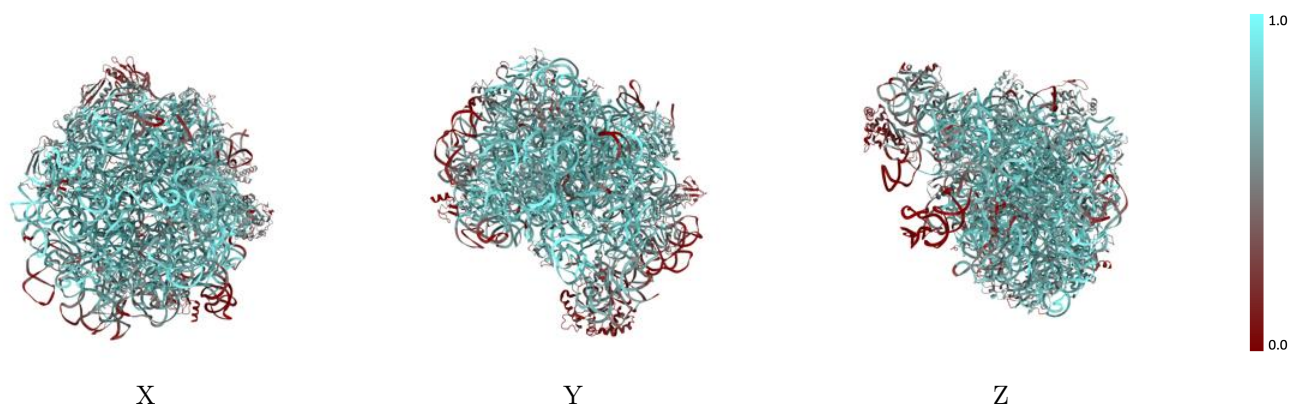
The images above show the 3D surface view of the map at the recommended contour level 0.1 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



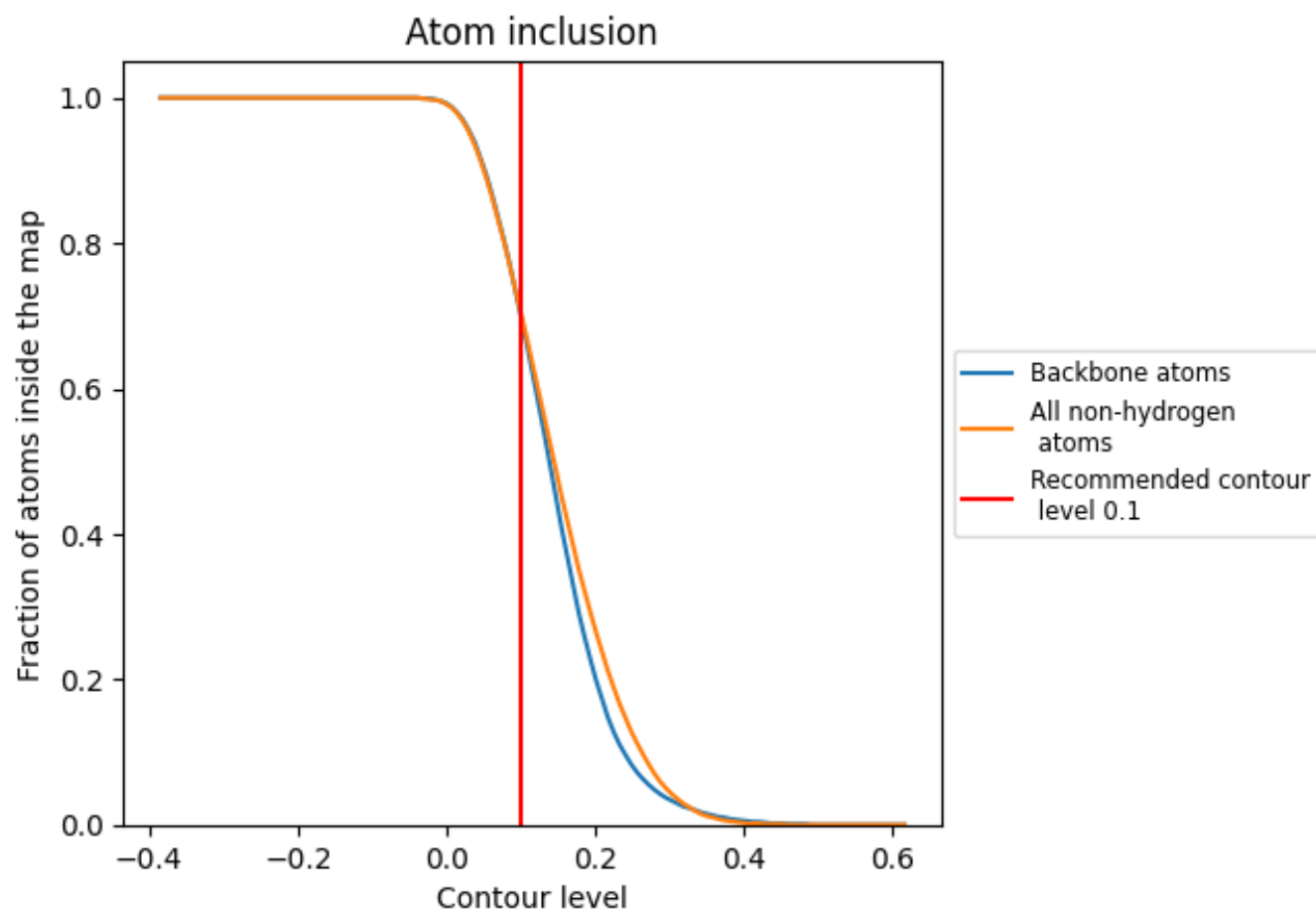
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions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

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



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (0.1).

9.4 Atom inclusion [i](#)



At the recommended contour level, 70% of all backbone atoms, 70% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary ⓘ

The table lists the average atom inclusion at the recommended contour level (0.1) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.7050	 0.5560
1	 0.3900	 0.4680
A	 0.7570	 0.5590
B	 0.6120	 0.5120
C	 0.7030	 0.5940
D	 0.6840	 0.5880
E	 0.5980	 0.5610
F	 0.1780	 0.4100
G	 0.3240	 0.4600
H	 0.2240	 0.4210
K	 0.7030	 0.5900
L	 0.6440	 0.5840
M	 0.6620	 0.5720
N	 0.6600	 0.5830
O	 0.7020	 0.5810
P	 0.5070	 0.5260
Q	 0.6080	 0.5710
R	 0.7380	 0.5940
S	 0.6490	 0.5750
T	 0.6570	 0.5800
U	 0.5650	 0.5500
V	 0.5350	 0.5300
W	 0.5480	 0.5390
X	 0.6980	 0.5910
Y	 0.6230	 0.5760
Z	 0.5080	 0.5210
a	 0.6650	 0.5750
b	 0.6450	 0.5610
c	 0.5860	 0.5550
d	 0.7510	 0.6080
e	 0.7310	 0.5940
f	 0.6030	 0.5590

