



Full wwPDB EM Validation Report ⓘ

Mar 9, 2026 – 11:40 AM UTC

PDB ID : 3J0O / pdb_00003j0o
EMDB ID : EMD-5327
Title : Core of mammalian 80S pre-ribosome in complex with tRNAs fitted to a 9A cryo-EM map: classic PRE state 2
Authors : Budkevich, T.; Giesebrecht, J.; Altman, R.; Munro, J.; Mielke, T.; Nierhaus, K.; Blanchard, S.; Spahn, C.M.
Deposited on : 2011-10-05
Resolution : 9.00 Å(reported)
Based on initial models : 2WDK, 3O58, 2XZM

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
MolProbity : 4-5-2 with Phenix2.0
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

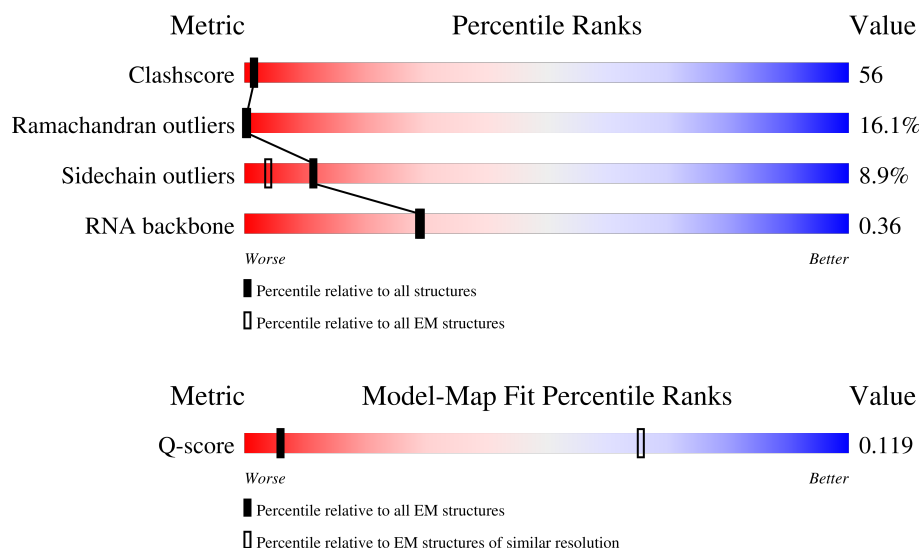
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 9.00 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	257 (8.50 - 9.50)

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	a	48	
2	b	12	
3	c	17	

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Mol	Chain	Length	Quality of chain
4	d	7	
5	e	4	
6	E	5	
7	f	21	
8	g	31	
9	G	13	
10	h	111	
11	T	192	
12	K	140	
13	L	141	
14	X	68	
15	S	125	
16	2	112	
17	3	12	
18	4	14	
19	5	6	
20	6	19	
21	7	50	
22	8	20	
23	B	213	
24	F	95	
25	V	76	
25	W	76	
25	Y	76	
26	v	3	

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Mol	Chain	Length	Quality of chain
26	y	3	33% 67%
27	w	2	100%

2 Entry composition

There are 27 unique types of molecules in this entry. The entry contains 22450 atoms, of which 0 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a RNA chain called 40S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	a	48	Total	C	N	O	P	0	0
			1029	459	190	332	48		

- Molecule 2 is a RNA chain called 40S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	b	12	Total	C	N	O	P	0	0
			260	116	49	83	12		

- Molecule 3 is a RNA chain called 40S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
3	c	17	Total	C	N	O	P	0	0
			362	162	66	117	17		

- Molecule 4 is a RNA chain called 40S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
4	d	7	Total	C	N	O	P	0	0
			155	69	33	46	7		

- Molecule 5 is a RNA chain called 40S ribosomal RNA fragment.

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

- Molecule 6 is a RNA chain called 40S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
6	E	5	Total	C	N	O	P	0	0
			100	45	13	37	5		

- Molecule 7 is a RNA chain called 40S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
7	f	21	Total	C	N	O	P	0	0
			452	200	79	152	21		

- Molecule 8 is a RNA chain called 40S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
8	g	31	Total	C	N	O	P	0	0
			660	295	118	216	31		

- Molecule 9 is a RNA chain called 40S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
9	G	13	Total	C	N	O	P	0	0
			276	123	49	91	13		

- Molecule 10 is a RNA chain called 40S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
10	h	111	Total	C	N	O	P	0	0
			2368	1060	431	766	111		

- Molecule 11 is a protein called Ribosomal protein S5.

Mol	Chain	Residues	Atoms					AltConf	Trace
11	T	192	Total	C	N	O	S	0	0
			1520	961	281	270	8		

- Molecule 12 is a protein called Ribosomal protein S14.

Mol	Chain	Residues	Atoms					AltConf	Trace
12	K	140	Total	C	N	O	S	0	0
			1063	654	206	197	6		

- Molecule 13 is a protein called Ribosomal protein S23.

Mol	Chain	Residues	Atoms					AltConf	Trace
13	L	141	Total	C	N	O	S	0	0
			1097	691	221	180	5		

- Molecule 14 is a protein called Ribosomal protein S30.

Mol	Chain	Residues	Atoms					AltConf	Trace
14	X	68	Total	C	N	O	S	0	0
			554	350	113	90	1		

- Molecule 15 is a protein called Ribosomal protein S15.

Mol	Chain	Residues	Atoms					AltConf	Trace
15	S	125	Total	C	N	O	S	0	0
			985	632	173	176	4		

- Molecule 16 is a RNA chain called 60S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
16	2	112	Total	C	N	O	P	0	0
			2392	1070	435	775	112		

- Molecule 17 is a RNA chain called 60S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
17	3	12	Total	C	N	O	P	0	0
			259	116	50	81	12		

- Molecule 18 is a RNA chain called 60S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
18	4	14	Total	C	N	O	P	0	0
			306	135	59	98	14		

- Molecule 19 is a RNA chain called 60S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
19	5	6	Total	C	N	O	P	0	0
			127	57	23	41	6		

- Molecule 20 is a RNA chain called 60S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
20	6	19	Total	C	N	O	P	0	0
			417	187	88	123	19		

- Molecule 21 is a RNA chain called 60S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
21	7	50	Total	C	N	O	P	0	0
			1054	471	173	360	50		

- Molecule 22 is a RNA chain called 60S ribosomal RNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
22	8	20	Total	C	N	O	P	0	0
			431	192	80	139	20		

- Molecule 23 is a protein called Ribosomal protein L10a.

Mol	Chain	Residues	Atoms					AltConf	Trace
23	B	213	Total	C	N	O		0	0
			1055	629	213	213			

- Molecule 24 is a protein called Ribosomal protein L36a.

Mol	Chain	Residues	Atoms					AltConf	Trace
24	F	95	Total	C	N	O		0	0
			467	277	95	95			

- Molecule 25 is a RNA chain called tRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
25	Y	75	Total	C	N	O	P	0	0
			1597	713	285	525	74		
25	V	75	Total	C	N	O	P	0	0
			1597	713	285	525	74		
25	W	76	Total	C	N	O	P	0	0
			1619	723	290	531	75		

- Molecule 26 is a RNA chain called mRNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
26	y	3	Total	C	N	O	P	0	0
			60	27	7	23	3		
26	v	3	Total	C	N	O	P	0	0
			60	27	7	23	3		

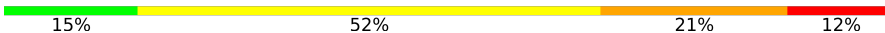
- Molecule 27 is a RNA chain called mRNA fragment.

Mol	Chain	Residues	Atoms					AltConf	Trace
27	w	2	Total	C	N	O	P	0	0
			44	20	10	12	2		

3 Residue-property plots

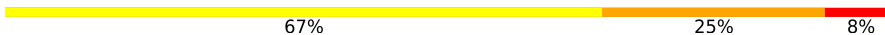
These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and atom inclusion 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: 40S ribosomal RNA fragment

Chain a: 



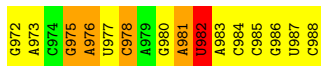
- Molecule 2: 40S ribosomal RNA fragment

Chain b: 

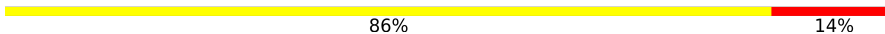


- Molecule 3: 40S ribosomal RNA fragment

Chain c: 



- Molecule 4: 40S ribosomal RNA fragment

Chain d: 



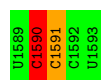
- Molecule 5: 40S ribosomal RNA fragment

Chain e: 



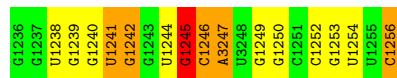
- Molecule 6: 40S ribosomal RNA fragment

Chain E: 



- Molecule 7: 40S ribosomal RNA fragment

Chain f: 29% 43% 24% 5%



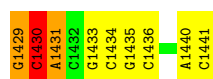
- Molecule 8: 40S ribosomal RNA fragment

Chain g: 6% 68% 19% 6%



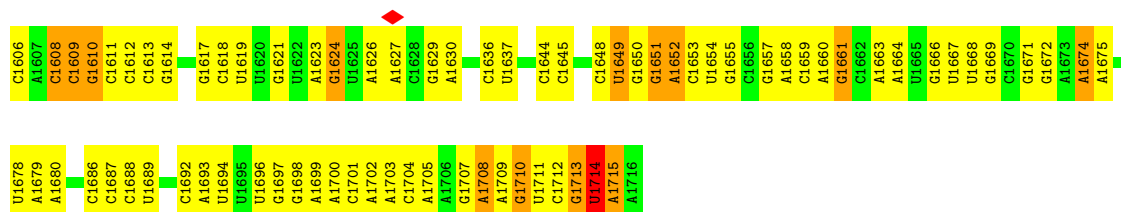
- Molecule 9: 40S ribosomal RNA fragment

Chain G: 31% 46% 15% 8%



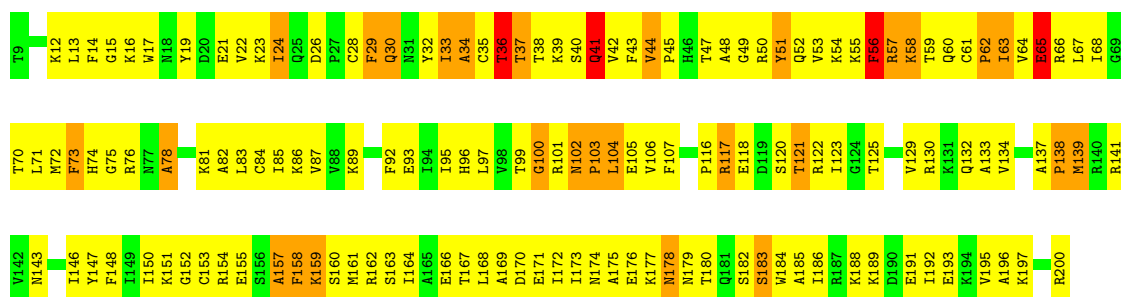
- Molecule 10: 40S ribosomal RNA fragment

Chain h: 33% 54% 12%

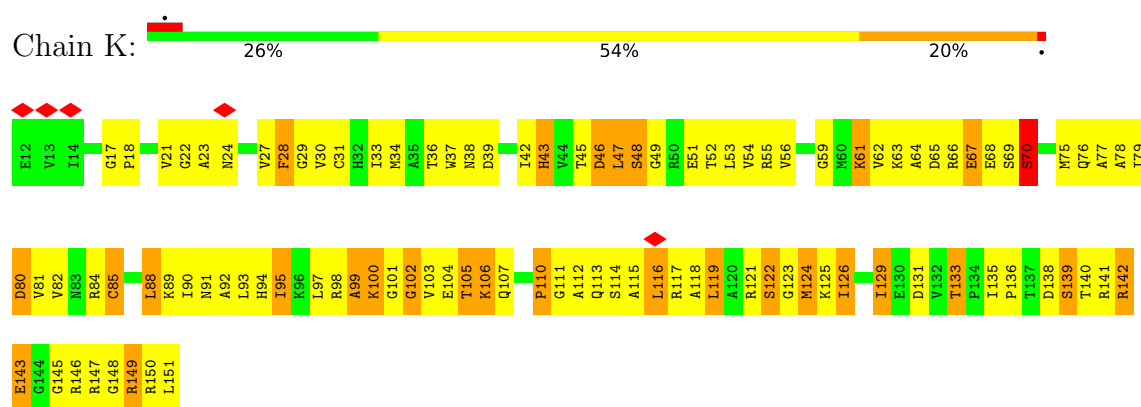


- Molecule 11: Ribosomal protein S5

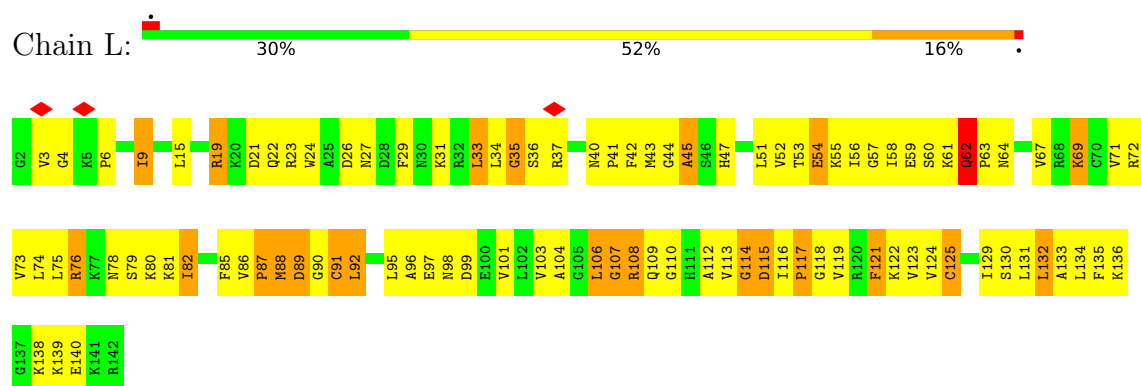
Chain T: 24% 59% 14%



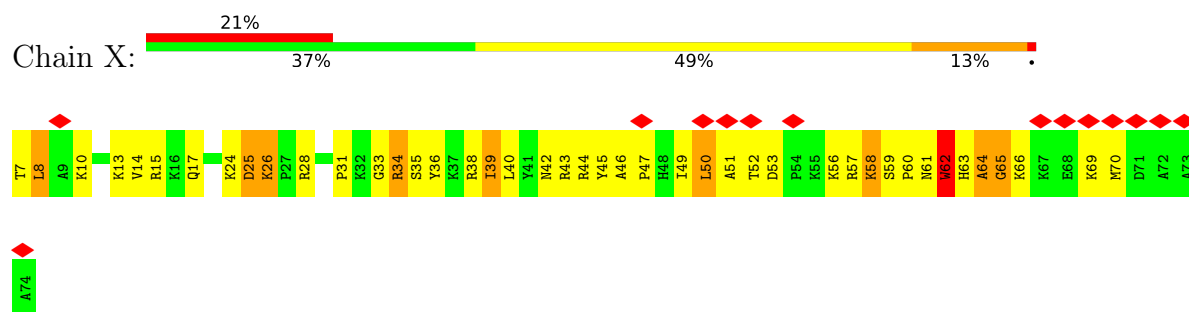
- Molecule 12: Ribosomal protein S14



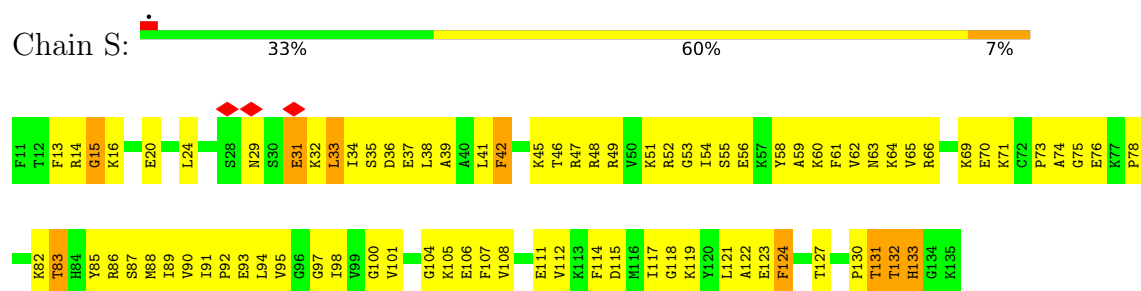
- Molecule 13: Ribosomal protein S23



- Molecule 14: Ribosomal protein S30



- Molecule 15: Ribosomal protein S15



- Molecule 16: 60S ribosomal RNA fragment





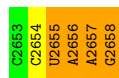
- Molecule 17: 60S ribosomal RNA fragment



- Molecule 18: 60S ribosomal RNA fragment



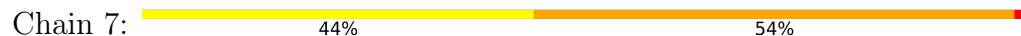
- Molecule 19: 60S ribosomal RNA fragment



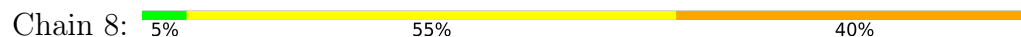
- Molecule 20: 60S ribosomal RNA fragment



- Molecule 21: 60S ribosomal RNA fragment

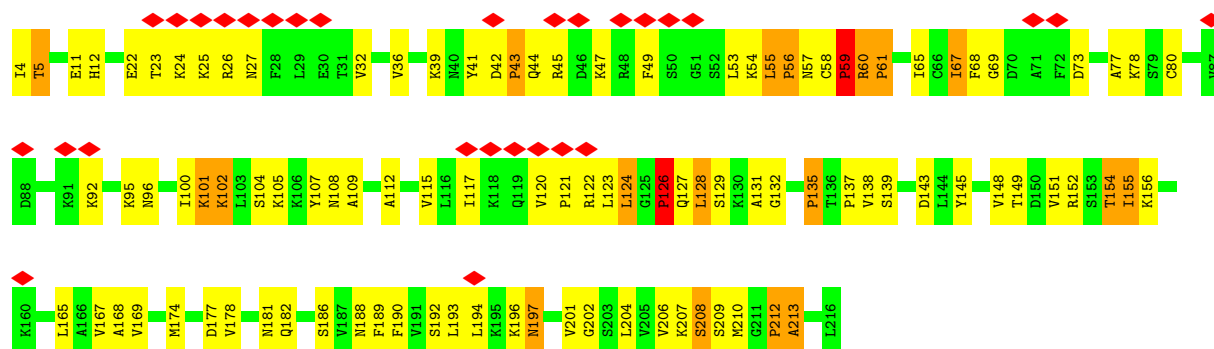


- Molecule 22: 60S ribosomal RNA fragment



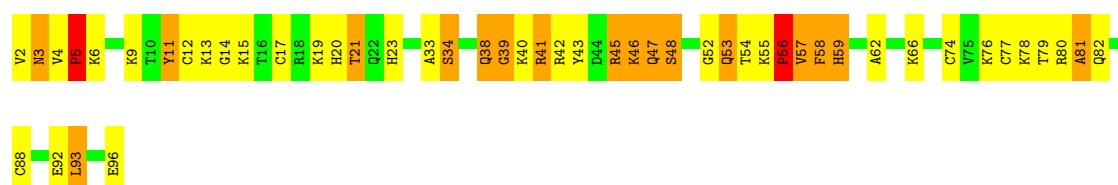
- Molecule 23: Ribosomal protein L10a





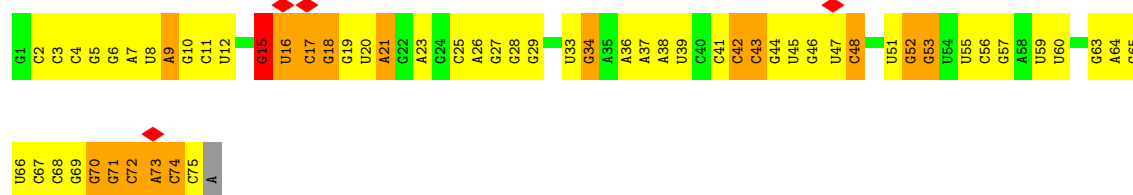
• Molecule 24: Ribosomal protein L36a

Chain F: 47% 33% 18% .



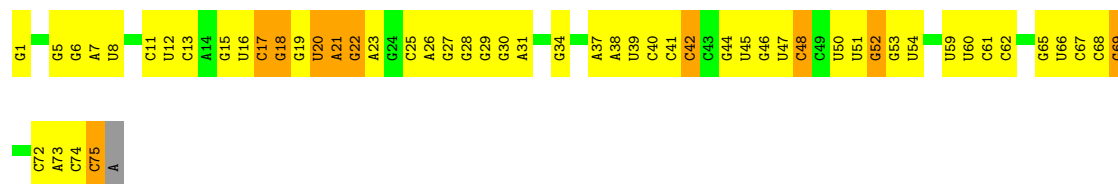
• Molecule 25: tRNA

Chain Y: 5% 21% 55% 21% ..



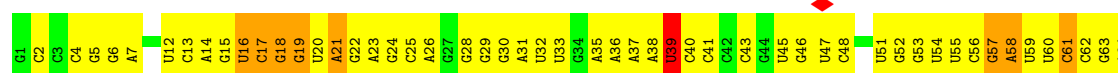
• Molecule 25: tRNA

Chain V: 28% 58% 13% .



• Molecule 25: tRNA

Chain W: 20% 66% 13% .





- Molecule 26: mRNA fragment

Chain y: 33% 67%



- Molecule 26: mRNA fragment

Chain v: 67% 33%



- Molecule 27: mRNA fragment

Chain w: 100%

There are no outlier residues recorded for this chain.

4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C1	Depositor
Number of particles used	81946	Depositor
Resolution determination method	Not provided	
CTF correction method	CTF CORRECTION OF EACH DEFOCUS GROUP VOLUME PRIOR TO BACK PROJECTION	Depositor
Microscope	FEI POLARA 300	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	20	Depositor
Minimum defocus (nm)	2000	Depositor
Maximum defocus (nm)	4000	Depositor
Magnification	65520	Depositor
Image detector	KODAK SO-163 FILM	Depositor
Maximum map value	21309.771	Depositor
Minimum map value	-5197.505	Depositor
Average map value	340.467	Depositor
Map value standard deviation	1535.379	Depositor
Recommended contour level	1000	Depositor
Map size ($\text{\AA}$)	453.6, 453.6, 453.6	wwPDB
Map dimensions	180, 180, 180	wwPDB
Map angles ($^\circ$)	90.0, 90.0, 90.0	wwPDB
Pixel spacing ($\text{\AA}$)	2.52, 2.52, 2.52	Depositor

5 Model quality ⓘ

5.1 Standard geometry ⓘ

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

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	$\# Z > 5$	RMSZ	$\# Z > 5$
1	a	0.55	0/1151	0.98	6/1793 (0.3%)
2	b	0.41	0/291	0.83	1/452 (0.2%)
3	c	0.46	0/404	0.93	3/627 (0.5%)
4	d	0.42	0/174	0.94	1/270 (0.4%)
5	e	0.41	0/93	0.72	0/142
6	E	0.41	0/109	1.04	1/166 (0.6%)
7	f	0.47	0/504	0.91	3/785 (0.4%)
8	g	0.50	0/737	0.95	4/1146 (0.3%)
9	G	0.44	0/307	0.93	2/476 (0.4%)
10	h	0.38	0/2650	0.70	2/4127 (0.0%)
11	T	0.54	0/1546	1.15	12/2079 (0.6%)
12	K	0.53	0/1078	1.09	9/1452 (0.6%)
13	L	0.53	0/1114	1.06	4/1485 (0.3%)
14	X	0.47	0/566	1.18	5/753 (0.7%)
15	S	0.45	0/1003	1.03	6/1342 (0.4%)
16	2	0.58	0/2677	0.64	0/4170
17	3	0.11	0/290	0.24	0/450
18	4	0.38	0/342	0.51	0/533
19	5	0.39	0/141	0.54	0/217
20	6	0.70	0/470	0.80	0/732
21	7	0.58	0/1174	0.74	3/1825 (0.2%)
22	8	0.76	0/482	0.64	0/750
23	B	0.51	0/1054	1.05	12/1468 (0.8%)
24	F	0.66	0/466	1.04	3/646 (0.5%)
25	V	0.40	0/1784	0.68	0/2780
25	W	0.44	0/1809	0.61	0/2819
25	Y	0.50	0/1784	0.68	2/2780 (0.1%)
26	v	0.40	0/65	0.70	0/98
26	y	0.44	0/65	0.54	0/98
27	w	0.42	0/49	0.62	0/74
All	All	0.50	0/24379	0.84	79/36535 (0.2%)

Chiral center outliers are detected by calculating the chiral volume of a chiral center and verifying if the center is modelled as a planar moiety or with the opposite hand. A planarity outlier is detected

by checking planarity of atoms in a peptide group, atoms in a mainchain group or atoms of a sidechain that are expected to be planar.

Mol	Chain	#Chirality outliers	#Planarity outliers
1	a	0	4
4	d	0	1
6	E	0	1
8	g	0	2
9	G	0	1
10	h	0	2
25	W	0	2
All	All	0	13

There are no bond length outliers.

All (79) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
11	T	36	THR	N-CA-C	10.62	123.93	111.71
15	S	131	THR	N-CA-C	9.81	125.24	109.24
10	h	1714	U	N1-C1'-C2'	9.06	125.60	112.00
11	T	73	PHE	N-CA-C	8.95	124.26	113.16
14	X	58	LYS	N-CA-C	8.92	122.12	110.43
23	B	121	PRO	N-CA-CB	8.52	110.49	101.88
24	F	5	PRO	N-CA-CB	7.93	111.58	103.25
15	S	132	THR	N-CA-C	-7.89	98.13	110.36
21	7	2845	A	N1-C6-N6	7.87	142.21	118.60
21	7	2845	A	C6-N1-C2	7.72	141.75	118.60
14	X	50	LEU	N-CA-C	-7.62	103.96	113.18
6	E	1590	C	C2'-C3'-O3'	7.59	120.88	109.50
14	X	62	TRP	N-CA-C	-7.46	103.44	112.54
23	B	56	PRO	N-CA-CB	7.35	110.97	103.25
1	a	559	C	N1-C1'-C2'	7.33	123.00	112.00
4	d	1545	A	C2'-C3'-O3'	-7.18	98.73	109.50
11	T	78	ALA	N-CA-C	7.10	120.89	110.17
15	S	133	HIS	N-CA-C	7.02	118.85	108.60
23	B	61	PRO	N-CA-CB	6.97	110.56	103.25
23	B	135	PRO	N-CA-CB	6.95	110.55	103.25
23	B	149	THR	N-CA-C	-6.92	104.10	112.54
12	K	133	THR	N-CA-C	-6.85	99.29	109.42
23	B	126	PRO	N-CA-CB	6.83	110.42	103.25
8	g	1172	G	C2'-C3'-O3'	6.80	123.90	113.70
24	F	56	PRO	N-CA-CB	6.80	110.39	103.25
23	B	43	PRO	N-CA-CB	6.78	110.36	103.25
1	a	553	A	C2'-C3'-O3'	6.76	119.64	109.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
12	K	88	LEU	N-CA-C	-6.76	105.00	113.18
23	B	137	PRO	N-CA-CB	6.74	110.33	103.25
23	B	59	PRO	N-CA-CB	6.72	110.30	103.25
23	B	212	PRO	N-CA-CB	6.69	110.27	103.25
24	F	38	GLN	N-CA-C	-6.64	103.14	111.11
8	g	1172	G	N9-C1'-C2'	6.53	121.79	112.00
11	T	104	LEU	N-CA-C	-6.43	104.48	112.90
11	T	162	ARG	N-CA-C	-6.40	100.80	110.28
7	f	1245	G	C2'-C3'-O3'	6.35	119.02	109.50
1	a	588	A	N9-C1'-C2'	6.33	123.50	114.00
11	T	65	GLU	N-CA-C	-6.25	104.47	111.28
13	L	115	ASP	N-CA-C	6.18	117.81	111.14
7	f	1241	U	C2'-C3'-O3'	6.17	118.76	109.50
23	B	23	THR	N-CA-C	6.13	113.81	108.78
15	S	115	ASP	N-CA-C	6.09	118.71	111.71
1	a	552	C	N1-C1'-C2'	6.07	121.11	112.00
12	K	48	SER	N-CA-C	-6.03	105.74	114.12
3	c	981	A	N9-C1'-C2'	5.98	122.97	114.00
12	K	143	GLU	N-CA-C	-5.97	102.65	110.53
13	L	62	GLN	N-CA-C	5.91	122.87	109.81
13	L	121	PHE	N-CA-C	5.91	118.67	109.52
1	a	547	C	C4'-C3'-O3'	5.90	118.26	109.40
3	c	982	U	C2'-C3'-O3'	-5.81	104.98	113.70
12	K	126	ILE	N-CA-C	5.77	116.17	107.80
11	T	34	ALA	N-CA-C	5.69	120.88	113.88
21	7	2845	A	C5-C6-N1	-5.67	100.69	117.70
14	X	65	GLY	N-CA-C	-5.66	107.18	115.72
15	S	61	PHE	N-CA-C	-5.65	105.20	111.36
10	h	1708	A	C2'-C3'-O3'	-5.61	105.29	113.70
11	T	29	PHE	N-CA-C	5.58	119.71	113.01
8	g	1162	C	C4'-C3'-O3'	-5.55	104.67	113.00
12	K	61	LYS	N-CA-C	5.55	120.33	113.50
11	T	37	THR	N-CA-C	5.54	118.71	110.52
8	g	1162	C	N1-C1'-C2'	5.42	120.14	112.00
14	X	39	ILE	CB-CA-C	-5.40	104.97	111.88
2	b	887	U	N1-C1'-C2'	5.33	120.00	112.00
11	T	73	PHE	CA-C-N	-5.31	115.13	122.30
11	T	73	PHE	C-N-CA	-5.31	115.13	122.30
15	S	42	PHE	N-CA-C	5.30	117.70	110.55
3	c	981	A	C2'-C3'-O3'	5.26	117.40	109.50
11	T	161	MET	N-CA-C	-5.26	106.82	113.18
25	Y	15	G	C2'-C3'-O3'	5.24	117.36	109.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	a	574	A	N9-C1'-C2'	5.24	121.86	114.00
12	K	70	SER	N-CA-C	5.22	121.35	109.81
7	f	1256	C	N1-C1'-C2'	5.19	119.78	112.00
9	G	1430	C	C2'-C3'-O3'	5.12	117.19	109.50
12	K	131	ASP	N-CA-C	-5.11	100.09	108.41
12	K	85	CYS	N-CA-C	5.05	116.86	111.36
25	Y	15	G	C4'-C3'-O3'	5.04	116.95	109.40
9	G	1429	G	C2'-C3'-O3'	-5.03	106.15	113.70
23	B	11	GLU	N-CA-C	-5.03	107.27	113.41
13	L	125	CYS	N-CA-C	5.02	117.59	109.06

There are no chirality outliers.

All (13) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
6	E	1590	C	Sidechain
9	G	1430	C	Sidechain
25	W	39	U	Sidechain
25	W	5	G	Sidechain
1	a	547	C	Sidechain
1	a	558	G	Sidechain
1	a	559	C	Sidechain
1	a	574	A	Sidechain
4	d	1545	A	Sidechain
8	g	1157	U	Sidechain
8	g	1158	U	Sidechain
10	h	1610	G	Sidechain
10	h	1714	U	Sidechain

5.2 Too-close contacts ⓘ

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	a	1029	0	521	73	0
2	b	260	0	130	31	0
3	c	362	0	185	23	0
4	d	155	0	78	6	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
5	e	84	0	45	2	0
6	E	100	0	54	7	0
7	f	452	0	226	27	0
8	g	660	0	335	49	0
9	G	276	0	141	18	0
10	h	2368	0	1197	162	0
11	T	1520	0	1569	198	0
12	K	1063	0	1088	130	0
13	L	1097	0	1169	118	0
14	X	554	0	604	58	0
15	S	985	0	1026	103	0
16	2	2392	0	1208	448	0
17	3	259	0	131	13	0
18	4	306	0	154	63	0
19	5	127	0	66	25	0
20	6	417	0	209	78	0
21	7	1054	0	532	175	0
22	8	431	0	216	50	0
23	B	1055	0	453	34	0
24	F	467	0	208	33	0
25	V	1597	0	807	94	0
25	W	1619	0	821	120	0
25	Y	1597	0	811	95	0
26	v	60	0	31	23	0
26	y	60	0	31	14	0
27	w	44	0	23	0	0
All	All	22450	0	14069	2044	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 56.

All (2044) 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)
10:h:1609:C:C6	25:V:34:G:C6	1.84	1.50
19:5:2657:A:C2	20:6:2694:A:N3	1.73	1.45
10:h:1609:C:C1'	26:v:18:C:N4	1.76	1.44
10:h:1709:A:OP1	26:y:20:U:C5'	1.64	1.41
18:4:2619:G:OP1	21:7:2866:U:C6	1.79	1.35
18:4:2619:G:OP1	21:7:2866:U:C4	1.80	1.34
10:h:1609:C:C5	25:V:34:G:C6	2.17	1.33

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:4:2618:G:H2'	21:7:2865:U:P	1.69	1.33
10:h:1609:C:O4'	26:v:18:C:N4	1.64	1.29
18:4:2618:G:C2'	21:7:2865:U:P	2.22	1.28
10:h:1609:C:C4'	26:v:18:C:N4	1.96	1.27
16:2:2286:U:OP1	25:Y:71:G:H5'	1.33	1.27
18:4:2619:G:P	21:7:2866:U:C5	2.27	1.26
10:h:1609:C:C4	25:V:34:G:C4	2.17	1.26
10:h:1609:C:C5	25:V:34:G:C5	2.23	1.25
16:2:2286:U:OP1	25:Y:71:G:C5'	1.83	1.24
10:h:1609:C:C6	25:V:34:G:O6	1.88	1.23
25:V:40:C:H5''	25:W:35:A:O3'	1.37	1.21
10:h:1608:C:H4'	10:h:1609:C:H5''	1.25	1.19
18:4:2618:G:H2'	21:7:2865:U:OP1	1.38	1.19
13:L:9:ILE:H	13:L:9:ILE:HD12	1.02	1.19
18:4:2618:G:H1'	21:7:2866:U:OP2	1.37	1.17
18:4:2619:G:OP1	21:7:2866:U:C5	0.79	1.14
10:h:1709:A:OP1	26:y:20:U:H5''	1.26	1.14
11:T:57:ARG:HA	11:T:57:ARG:HH11	1.02	1.13
16:2:2285:C:O2'	25:Y:71:G:H4'	1.45	1.13
18:4:2618:G:O2'	21:7:2865:U:P	2.07	1.12
16:2:2210:G:N2	25:W:71:G:O3'	1.83	1.11
16:2:2250:G:N2	16:2:2267:C:O2	1.82	1.11
16:2:2285:C:O3'	25:Y:72:C:P	2.08	1.10
20:6:2690:G:H2'	20:6:2690:G:N3	1.62	1.10
16:2:2213:A:H2'	16:2:2214:A:H8	1.16	1.10
16:2:2262:A:H3'	16:2:2263:C:H5''	1.31	1.10
16:2:2270:A:H5''	16:2:2270:A:C8	1.87	1.10
10:h:1709:A:OP1	26:y:20:U:H5'	1.38	1.09
25:V:20:U:H3'	25:V:21:A:H5'	1.34	1.08
16:2:2271:A:C2'	16:2:2272:G:H5''	1.84	1.07
10:h:1609:C:C4	25:V:34:G:C5	2.42	1.07
16:2:2271:A:H2'	16:2:2272:G:H5''	1.30	1.06
17:3:2484:A:H1'	25:W:19:G:N2	1.71	1.06
2:b:886:U:H3'	2:b:887:U:H5''	1.11	1.06
16:2:2213:A:H2'	16:2:2214:A:C8	1.89	1.06
10:h:1609:C:H4'	26:v:18:C:H41	1.18	1.05
16:2:2287:C:H5	25:Y:73:A:N7	1.53	1.05
7:f:1245:G:H4'	7:f:1246:C:H5''	1.38	1.05
16:2:2285:C:O3'	25:Y:72:C:OP1	1.73	1.04
10:h:1709:A:N6	25:Y:36:A:C2	2.26	1.04
20:6:2698:G:H2'	20:6:2699:G:H5''	1.40	1.04

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:h:1711:U:C4'	16:2:2256:A:OP2	2.06	1.03
13:L:19:ARG:NH1	13:L:19:ARG:HA	1.73	1.02
13:L:19:ARG:HA	13:L:19:ARG:HH11	1.20	1.02
12:K:21:VAL:HG12	12:K:22:GLY:H	1.24	1.02
7:f:1246:C:H42	25:Y:34:G:H1'	1.24	1.01
13:L:116:ILE:HG21	13:L:119:VAL:HB	1.41	1.00
19:5:2656:A:H4'	19:5:2657:A:OP1	1.55	1.00
8:g:1163:U:O2	25:V:34:G:H5'	1.62	0.99
10:h:1609:C:N4	25:V:34:G:C4	2.29	0.99
25:V:20:U:C3'	25:V:21:A:H5'	1.91	0.99
10:h:1609:C:O3'	26:v:18:C:C5	2.15	0.99
10:h:1711:U:H4'	16:2:2256:A:OP2	1.63	0.98
11:T:54:LYS:HG2	11:T:55:LYS:H	1.28	0.98
16:2:2223:A:O2'	16:2:2224:A:H8	1.45	0.98
10:h:1709:A:N6	25:Y:36:A:H2	1.62	0.98
25:W:38:A:H3'	25:W:39:U:H5''	1.43	0.98
16:2:2304:C:O2'	16:2:2305:G:H5'	1.64	0.98
10:h:1609:C:C1'	26:v:18:C:H42	1.62	0.97
11:T:37:THR:HG22	11:T:39:LYS:H	1.30	0.97
16:2:2285:C:H5	16:2:2286:U:C2	1.83	0.97
18:4:2618:G:O2'	21:7:2865:U:OP2	1.81	0.97
16:2:2287:C:C5	25:Y:73:A:N7	2.32	0.96
25:V:68:C:H2'	25:V:69:G:H5''	1.46	0.96
1:a:561:A:H1'	14:X:14:VAL:HG23	1.46	0.96
2:b:881:U:H2'	2:b:882:G:H5''	1.47	0.96
16:2:2285:C:H5''	25:Y:72:C:OP1	1.65	0.95
10:h:1609:C:H4'	26:v:18:C:N4	1.74	0.95
12:K:65:ASP:O	12:K:68:GLU:HG2	1.64	0.95
18:4:2616:C:H2'	18:4:2617:U:H5'	1.46	0.95
16:2:2270:A:H5''	16:2:2270:A:H8	1.25	0.95
16:2:2276:G:C5	16:2:2277:C:C5	2.55	0.95
16:2:2282:U:OP1	22:8:2973:G:O2'	1.84	0.94
20:6:2699:G:C8	20:6:2699:G:H5'	2.02	0.94
16:2:2281:A:N3	22:8:2974:U:C1'	2.29	0.94
25:W:69:G:H2'	25:W:70:G:H5''	1.49	0.94
10:h:1609:C:C4'	26:v:18:C:H41	1.69	0.94
16:2:2252:A:H2'	16:2:2253:G:H8	1.33	0.94
16:2:2286:U:OP1	25:Y:71:G:H5''	1.64	0.94
20:6:2689:A:N3	20:6:2702:A:C2	2.36	0.94
11:T:117:ARG:HD2	11:T:193:GLU:OE2	1.67	0.93
20:6:2698:G:C2'	20:6:2699:G:H5''	1.98	0.93

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:V:17:C:H3'	25:V:18:G:H5''	1.50	0.93
25:Y:28:G:H1	25:Y:42:C:H42	1.03	0.93
16:2:2251:G:H1'	16:2:2252:A:OP1	1.68	0.93
25:Y:41:C:H3'	25:Y:42:C:H5''	1.46	0.93
16:2:2268:U:H2'	16:2:2269:U:H5	1.33	0.93
20:6:2703:A:H5''	20:6:2704:A:H5'	1.46	0.93
2:b:886:U:H3'	2:b:887:U:C5'	1.98	0.92
21:7:2862:U:O2'	21:7:2863:G:H5'	1.69	0.92
18:4:2618:G:C2'	21:7:2865:U:OP1	2.16	0.92
10:h:1608:C:H4'	10:h:1609:C:C5'	1.99	0.91
21:7:2850:G:OP1	21:7:2850:G:H4'	1.68	0.91
13:L:9:ILE:HD12	13:L:9:ILE:N	1.84	0.91
23:B:67:ILE:HA	23:B:112:ALA:HB2	1.52	0.91
10:h:1613:C:O4'	10:h:1715:A:H2	1.53	0.91
12:K:54:VAL:HG11	12:K:81:VAL:HG13	1.50	0.91
1:a:552:C:H4'	14:X:66:LYS:HE2	1.53	0.91
14:X:7:THR:HG21	14:X:10:LYS:HB2	1.53	0.91
25:W:16:U:H3'	25:W:17:C:H5'	1.52	0.90
10:h:1709:A:O5'	26:y:19:U:O2'	1.87	0.90
11:T:57:ARG:HA	11:T:57:ARG:NH1	1.86	0.90
21:7:2869:U:H5''	21:7:2870:C:OP2	1.71	0.90
16:2:2299:A:C5	16:2:2300:G:C8	2.60	0.90
18:4:2619:G:C8	21:7:2866:U:C5	2.59	0.90
7:f:1244:U:H5'	7:f:1245:G:OP2	1.72	0.90
16:2:2251:G:O2'	16:2:2252:A:H5''	1.72	0.90
11:T:72:MET:HE1	11:T:81:LYS:HB2	1.52	0.90
16:2:2253:G:O6	16:2:2263:C:N4	2.04	0.90
25:V:40:C:C5'	25:W:35:A:O3'	2.19	0.90
16:2:2254:U:O2	16:2:2261:G:N2	2.05	0.89
21:7:2826:U:O2'	21:7:2827:U:H5'	1.72	0.89
12:K:95:ILE:HD12	12:K:129:ILE:HG23	1.54	0.89
20:6:2699:G:H5'	20:6:2699:G:H8	1.35	0.89
13:L:69:LYS:HZ2	13:L:69:LYS:HB3	1.36	0.89
16:2:2270:A:H2'	16:2:2271:A:C8	2.07	0.89
21:7:2847:A:H5'	21:7:2848:G:OP2	1.72	0.89
10:h:1609:C:C4'	26:v:18:C:C4	2.51	0.89
19:5:2657:A:C6	20:6:2694:A:O2'	2.26	0.89
1:a:564:A:N1	13:L:115:ASP:HB2	1.88	0.88
11:T:123:ILE:HD12	11:T:132:GLN:OE1	1.72	0.88
22:8:2966:G:O2'	22:8:2967:A:H5'	1.73	0.88
11:T:34:ALA:HB3	11:T:63:ILE:HG13	1.55	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:7:2834:G:O2'	21:7:2835:U:H6	1.56	0.88
20:6:2703:A:H5''	20:6:2704:A:C5'	2.03	0.88
16:2:2252:A:H2'	16:2:2253:G:C8	2.08	0.88
18:4:2619:G:OP2	21:7:2866:U:OP2	1.90	0.88
18:4:2619:G:P	21:7:2866:U:H5	1.77	0.87
16:2:2215:A:H2'	16:2:2215:A:N3	1.87	0.87
20:6:2689:A:H4'	20:6:2690:G:H5'	1.56	0.87
23:B:178:VAL:O	23:B:182:GLN:CB	2.23	0.87
19:5:2655:U:H4'	19:5:2656:A:C5'	2.04	0.87
11:T:92:PHE:HD1	11:T:103:PRO:HB2	1.38	0.86
25:Y:25:C:H2'	25:Y:26:A:H8	1.40	0.86
3:c:986:G:OP1	12:K:149:ARG:NH2	2.08	0.86
16:2:2262:A:H3'	16:2:2263:C:C5'	2.03	0.86
25:W:57:G:H2'	25:W:58:A:H5'	1.54	0.86
2:b:881:U:O2	2:b:883:A:H5''	1.76	0.86
20:6:2702:A:C8	20:6:2704:A:N6	2.43	0.86
21:7:2833:A:H2'	21:7:2834:G:H5'	1.55	0.86
10:h:1709:A:O5'	26:y:19:U:O3'	1.93	0.86
16:2:2293:C:H5	16:2:2294:U:H5	1.20	0.86
19:5:2657:A:H2	20:6:2694:A:N3	1.09	0.86
23:B:154:THR:HA	23:B:155:ILE:O	1.76	0.86
19:5:2657:A:N1	20:6:2694:A:O2'	2.07	0.85
25:W:20:U:H2'	25:W:21:A:H4'	1.58	0.85
25:W:69:G:C2'	25:W:70:G:H5''	2.06	0.85
16:2:2287:C:O2	16:2:2298:U:O4'	1.95	0.85
16:2:2286:U:OP2	25:Y:72:C:OP2	1.95	0.85
16:2:2259:A:H3'	16:2:2260:U:C6	2.11	0.85
16:2:2285:C:C4'	25:Y:72:C:OP1	2.24	0.85
8:g:1163:U:C2	25:V:34:G:C4'	2.60	0.85
11:T:95:ILE:CG2	11:T:103:PRO:HB3	2.06	0.85
13:L:58:ILE:HG21	13:L:117:PRO:HG3	1.57	0.85
8:g:1163:U:C2	25:V:34:G:H4'	2.12	0.85
10:h:1609:C:C4	25:V:34:G:C6	2.60	0.85
10:h:1613:C:O4'	10:h:1715:A:C2	2.30	0.85
13:L:71:VAL:HG21	13:L:95:LEU:HD13	1.57	0.85
16:2:2205:U:H2'	16:2:2206:G:H5'	1.58	0.84
12:K:28:PHE:O	12:K:47:LEU:HD21	1.76	0.84
16:2:2272:G:N3	16:2:2272:G:H5'	1.91	0.84
17:3:2484:A:H1'	25:W:19:G:C2	2.12	0.84
17:3:2485:A:OP1	25:W:56:C:O2	1.96	0.84
18:4:2618:G:O2'	21:7:2865:U:O5'	1.95	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:561:A:H1'	14:X:14:VAL:CG2	2.08	0.84
16:2:2285:C:C5'	25:Y:72:C:OP1	2.25	0.84
8:g:1158:U:O4	8:g:1172:G:N2	2.10	0.83
22:8:2971:A:OP1	22:8:2971:A:H3'	1.78	0.83
16:2:2281:A:N3	22:8:2974:U:H1'	1.87	0.83
1:a:572:U:H4'	1:a:573:A:C5'	2.08	0.83
16:2:2232:A:H2'	16:2:2233:A:O4'	1.79	0.83
21:7:2851:A:H2'	21:7:2852:C:O4'	1.77	0.83
25:V:40:C:H5'	25:W:35:A:H4'	1.58	0.83
11:T:197:LYS:HD3	11:T:200:ARG:NH2	1.92	0.83
25:W:16:U:C5	25:W:18:G:H3'	2.14	0.83
15:S:105:LYS:HG3	15:S:106:GLU:HG2	1.61	0.83
12:K:62:VAL:HG12	12:K:63:LYS:H	1.43	0.83
25:W:38:A:C3'	25:W:39:U:H5''	2.08	0.83
13:L:40:ASN:HB2	13:L:41:PRO:HD2	1.61	0.83
16:2:2293:C:C5	16:2:2294:U:H5	1.97	0.83
17:3:2484:A:N6	25:W:19:G:H2'	1.92	0.83
20:6:2689:A:N3	20:6:2702:A:H2	1.74	0.83
13:L:9:ILE:H	13:L:9:ILE:CD1	1.76	0.82
16:2:2210:G:N2	25:W:72:C:P	2.52	0.82
16:2:2253:G:N7	16:2:2254:U:C4	2.47	0.82
18:4:2618:G:C4	21:7:2865:U:OP1	2.32	0.82
16:2:2278:C:N4	16:2:2305:G:N7	2.27	0.82
25:Y:16:U:H3'	25:Y:17:C:H5'	1.61	0.82
25:W:56:C:H2'	25:W:57:G:O4'	1.79	0.82
13:L:62:GLN:HB3	13:L:63:PRO:HD3	1.62	0.82
16:2:2268:U:H2'	16:2:2269:U:C5	2.14	0.82
25:Y:28:G:H1	25:Y:42:C:N4	1.76	0.82
15:S:101:VAL:HG21	15:S:121:LEU:HD11	1.62	0.82
22:8:2960:C:H5''	22:8:2961:G:OP2	1.78	0.82
11:T:92:PHE:CD1	11:T:103:PRO:HB2	2.14	0.81
11:T:13:LEU:HD23	11:T:104:LEU:HD21	1.61	0.81
18:4:2618:G:C1'	21:7:2866:U:OP2	2.25	0.81
10:h:1661:G:H3'	10:h:1661:G:N3	1.95	0.81
16:2:2285:C:C5	16:2:2286:U:C2	2.68	0.81
25:V:41:C:C3'	25:V:42:C:H5''	2.09	0.81
16:2:2223:A:O2'	16:2:2224:A:C8	2.29	0.81
16:2:2271:A:H2'	16:2:2272:G:C5'	2.11	0.81
10:h:1613:C:C1'	10:h:1715:A:C2	2.63	0.81
16:2:2224:A:H2'	16:2:2225:U:O4'	1.79	0.81
11:T:121:THR:HG22	11:T:123:ILE:HG13	1.62	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:2:2253:G:H3'	16:2:2254:U:C5	2.16	0.80
10:h:1609:C:O2	26:v:18:C:C2	2.35	0.80
7:f:1245:G:C4'	7:f:1246:C:H5''	2.12	0.80
21:7:2842:U:O2	21:7:2842:U:H2'	1.82	0.80
13:L:62:GLN:HB3	13:L:63:PRO:CD	2.12	0.80
14:X:34:ARG:NH1	14:X:34:ARG:HB3	1.97	0.80
20:6:2702:A:H8	20:6:2704:A:C6	2.00	0.80
13:L:24:TRP:CH2	13:L:33:LEU:HD13	2.16	0.80
16:2:2285:C:C3'	25:Y:72:C:OP1	2.29	0.79
16:2:2272:G:H4'	16:2:2273:G:OP1	1.81	0.79
25:W:2:C:H42	25:W:71:G:H1	1.30	0.79
7:f:1246:C:N4	25:Y:34:G:H1'	1.97	0.79
16:2:2222:A:HO2'	16:2:2223:A:H8	1.28	0.79
23:B:177:ASP:O	23:B:181:ASN:CB	2.31	0.79
21:7:2825:C:H5'	21:7:2826:U:OP2	1.83	0.79
24:F:47:GLN:N	24:F:48:SER:HA	1.98	0.78
11:T:37:THR:HG22	11:T:39:LYS:N	1.97	0.78
16:2:2293:C:C5	16:2:2294:U:C5	2.72	0.78
16:2:2245:C:H2'	16:2:2246:G:O4'	1.83	0.78
16:2:2286:U:C4	16:2:2288:G:H1'	2.18	0.78
18:4:2619:G:H8	21:7:2866:U:C5	1.98	0.78
1:a:558:G:O2'	1:a:571:G:H4'	1.83	0.78
11:T:14:PHE:CE2	11:T:89:LYS:HA	2.19	0.78
3:c:975:G:H5'	3:c:976:A:OP2	1.82	0.78
20:6:2696:A:C8	20:6:2697:A:N7	2.51	0.78
16:2:2269:U:H2'	16:2:2269:U:O2	1.84	0.78
18:4:2619:G:H8	21:7:2866:U:H5	1.30	0.78
21:7:2837:A:H2'	21:7:2845:A:N1	1.98	0.78
10:h:1668:U:H2'	10:h:1669:G:H8	1.48	0.77
11:T:152:GLY:O	11:T:155:GLU:HB2	1.85	0.77
13:L:92:LEU:HD12	13:L:92:LEU:O	1.84	0.77
16:2:2251:G:C1'	16:2:2252:A:OP1	2.32	0.77
8:g:1163:U:O2	25:V:34:G:C5'	2.32	0.77
11:T:122:ARG:HE	11:T:129:VAL:HG13	1.50	0.77
21:7:2834:G:HO2'	21:7:2835:U:H6	0.77	0.77
16:2:2215:A:C2	16:2:2216:G:C8	2.73	0.77
21:7:2838:A:C2	21:7:2851:A:C4	2.72	0.77
25:V:68:C:C2'	25:V:69:G:H5''	2.14	0.77
8:g:1153:U:N3	8:g:1157:U:H5	1.81	0.77
25:Y:51:U:H3	25:Y:63:G:H1	1.30	0.77
13:L:3:VAL:HG12	13:L:4:GLY:H	1.48	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:2:2253:G:C6	16:2:2254:U:C2	2.73	0.77
21:7:2858:U:H2'	21:7:2859:U:C5	2.20	0.77
24:F:59:HIS:CB	25:W:76:A:N7	2.47	0.77
1:a:570:G:H4'	1:a:574:A:N3	2.00	0.76
25:V:51:U:H2'	25:V:52:G:C8	2.19	0.76
13:L:75:LEU:HD21	13:L:82:ILE:HD12	1.65	0.76
13:L:96:ALA:HB3	13:L:99:ASP:OD2	1.85	0.76
16:2:2280:A:C5	16:2:2282:U:H5	2.02	0.76
2:b:881:U:C2'	2:b:882:G:H5''	2.14	0.76
8:g:1159:U:O2'	8:g:1160:G:H5'	1.84	0.76
11:T:67:LEU:HD12	11:T:150:ILE:HD11	1.66	0.76
12:K:103:VAL:O	12:K:142:ARG:HD2	1.86	0.76
21:7:2834:G:C4	21:7:2835:U:C5	2.74	0.76
15:S:55:SER:HB2	15:S:58:TYR:CD2	2.20	0.76
10:h:1609:C:C3'	26:v:18:C:C5	2.57	0.76
25:Y:63:G:H2'	25:Y:64:A:C8	2.21	0.76
10:h:1609:C:H1'	26:v:18:C:H42	1.30	0.76
25:V:41:C:H3'	25:V:42:C:H5''	1.67	0.76
25:W:56:C:H3'	25:W:57:G:H5''	1.68	0.76
16:2:2253:G:H3'	16:2:2254:U:C6	2.20	0.76
10:h:1609:C:O2	26:v:18:C:N3	2.19	0.76
12:K:103:VAL:HG12	12:K:142:ARG:HG2	1.68	0.76
14:X:63:HIS:C	14:X:65:GLY:H	1.93	0.76
25:W:18:G:N1	25:W:55:U:H1'	1.99	0.76
3:c:983:A:H2'	3:c:984:C:C6	2.21	0.76
7:f:1245:G:O2'	7:f:1246:C:OP2	2.02	0.76
25:Y:5:G:H2'	25:Y:6:G:C8	2.20	0.76
20:6:2699:G:H8	20:6:2699:G:C5'	1.98	0.75
12:K:62:VAL:HG12	12:K:63:LYS:N	2.01	0.75
19:5:2656:A:OP1	24:F:96:GLU:O	2.04	0.75
20:6:2690:G:N3	20:6:2690:G:C2'	2.45	0.75
25:V:20:U:H3'	25:V:21:A:C5'	2.14	0.75
25:W:18:G:H1	25:W:55:U:H1'	1.48	0.75
7:f:1246:C:H42	25:Y:34:G:C1'	1.97	0.75
12:K:36:THR:HG22	12:K:37:TRP:N	2.02	0.75
15:S:46:THR:HG23	15:S:89:ILE:HD13	1.69	0.75
16:2:2279:A:H4'	16:2:2280:A:H5'	1.68	0.75
15:S:38:LEU:HB3	15:S:42:PHE:HE1	1.51	0.75
16:2:2278:C:N4	16:2:2305:G:C8	2.55	0.75
16:2:2260:U:H2'	16:2:2261:G:N7	2.02	0.75
25:W:69:G:C3'	25:W:70:G:H5''	2.16	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:h:1710:G:O2'	16:2:2256:A:C5'	2.35	0.74
16:2:2287:C:C2	16:2:2298:U:O2	2.40	0.74
22:8:2970:C:H3'	22:8:2971:A:C5'	2.17	0.74
8:g:1153:U:H3	8:g:1157:U:H5	1.34	0.74
16:2:2270:A:C2	16:2:2271:A:C4	2.76	0.74
21:7:2829:U:H2'	21:7:2830:G:H5'	1.67	0.74
11:T:99:THR:HG22	11:T:101:ARG:HG3	1.69	0.74
20:6:2702:A:H8	20:6:2704:A:N6	1.83	0.74
25:Y:41:C:C3'	25:Y:42:C:H5''	2.18	0.74
16:2:2280:A:C5	16:2:2282:U:C5	2.76	0.74
25:Y:27:G:H21	25:Y:43:C:H5	1.36	0.74
20:6:2702:A:C8	20:6:2704:A:C6	2.76	0.74
22:8:2961:G:C6	22:8:2962:U:C4	2.76	0.74
1:a:584:C:OP1	14:X:44:ARG:NH2	2.20	0.74
3:c:986:G:OP1	12:K:149:ARG:CZ	2.35	0.74
16:2:2211:U:O2	25:W:72:C:H4'	1.87	0.74
18:4:2619:G:C8	21:7:2866:U:H5	2.02	0.74
8:g:1171:G:O2'	8:g:1172:G:OP2	2.05	0.73
14:X:50:LEU:HD23	14:X:57:ARG:HH12	1.52	0.73
15:S:16:LYS:HG3	15:S:16:LYS:O	1.88	0.73
16:2:2249:G:H2'	16:2:2250:G:C1'	2.17	0.73
16:2:2296:A:O2'	16:2:2297:U:H5'	1.87	0.73
22:8:2970:C:H3'	22:8:2971:A:H5''	1.69	0.73
11:T:54:LYS:HG2	11:T:55:LYS:N	2.01	0.73
16:2:2257:C:H3'	16:2:2258:U:H6	1.52	0.73
19:5:2655:U:OP2	24:F:2:VAL:O	2.06	0.73
10:h:1657:G:H1	10:h:1668:U:H3	1.35	0.73
10:h:1709:A:P	26:y:20:U:H5'	2.28	0.73
16:2:2266:U:N3	16:2:2267:C:N4	2.36	0.73
21:7:2872:A:C4'	21:7:2873:U:OP1	2.35	0.73
10:h:1609:C:O2	26:v:18:C:O2	2.07	0.73
11:T:34:ALA:HB3	11:T:63:ILE:H	1.51	0.73
15:S:100:GLY:HA2	15:S:108:VAL:O	1.87	0.73
21:7:2854:U:O2'	21:7:2855:U:H5'	1.89	0.73
12:K:47:LEU:HD23	12:K:47:LEU:N	2.02	0.73
17:3:2485:A:OP1	25:W:56:C:C2	2.42	0.73
25:W:61:C:H2'	25:W:62:C:C6	2.24	0.73
7:f:1246:C:N4	25:Y:34:G:C1'	2.50	0.73
12:K:55:ARG:HG2	12:K:55:ARG:O	1.88	0.73
16:2:2278:C:H2'	16:2:2279:A:H5''	1.70	0.73
16:2:2289:U:O2'	16:2:2290:C:H5'	1.88	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:K:64:ALA:HB3	12:K:67:GLU:HG2	1.70	0.72
25:W:30:G:H2'	25:W:31:A:H8	1.54	0.72
8:g:1143:A:H2'	8:g:1144:A:C8	2.24	0.72
25:Y:16:U:C3'	25:Y:17:C:H5'	2.19	0.72
25:W:16:U:C3'	25:W:17:C:H5'	2.18	0.72
12:K:75:MET:HG2	12:K:121:ARG:NH2	2.04	0.72
12:K:119:LEU:O	12:K:124:MET:HB2	1.89	0.72
16:2:2294:U:H5'	16:2:2295:A:OP2	1.90	0.72
1:a:561:A:N3	14:X:14:VAL:HG21	2.04	0.72
10:h:1714:U:O2'	10:h:1715:A:H5'	1.89	0.72
11:T:158:PHE:HD2	11:T:159:LYS:HG3	1.54	0.72
16:2:2293:C:H5	16:2:2294:U:C5	2.05	0.72
15:S:55:SER:HB2	15:S:58:TYR:HD2	1.53	0.72
16:2:2281:A:O2'	16:2:2282:U:H5''	1.90	0.72
21:7:2852:C:C5	21:7:2853:A:C4	2.78	0.72
14:X:63:HIS:O	14:X:65:GLY:N	2.22	0.72
16:2:2279:A:O5'	16:2:2280:A:H5''	1.88	0.72
13:L:69:LYS:HB3	13:L:69:LYS:NZ	2.04	0.72
16:2:2288:G:C4	16:2:2289:U:C5	2.78	0.72
18:4:2618:G:N9	21:7:2865:U:OP1	2.23	0.72
20:6:2697:A:H2'	20:6:2698:G:H8	1.53	0.72
20:6:2701:U:O2'	20:6:2705:A:C8	2.42	0.72
13:L:19:ARG:O	13:L:23:ARG:HB2	1.90	0.72
21:7:2834:G:O2'	21:7:2835:U:O5'	2.08	0.72
9:G:1430:C:O2'	9:G:1431:A:H5'	1.89	0.71
16:2:2267:C:H2'	16:2:2268:U:O4'	1.90	0.71
16:2:2271:A:C3'	16:2:2272:G:H5''	2.20	0.71
20:6:2699:G:C8	20:6:2699:G:C5'	2.73	0.71
20:6:2699:G:H2'	20:6:2700:G:O4'	1.90	0.71
10:h:1608:C:C4'	10:h:1609:C:H5''	2.12	0.71
11:T:12:LYS:HB3	11:T:17:TRP:H	1.54	0.71
20:6:2702:A:O4'	20:6:2704:A:C8	2.43	0.71
21:7:2847:A:H3'	21:7:2848:G:H8	1.55	0.71
25:W:45:U:O2'	25:W:46:G:H5'	1.89	0.71
13:L:19:ARG:HH12	13:L:22:GLN:HB2	1.53	0.71
16:2:2286:U:P	25:Y:71:G:H5''	2.30	0.71
20:6:2697:A:H2'	20:6:2698:G:C8	2.26	0.71
20:6:2703:A:H3'	20:6:2704:A:C5'	2.21	0.71
12:K:121:ARG:HG3	12:K:122:SER:N	2.04	0.71
15:S:83:THR:HG23	15:S:85:TYR:H	1.55	0.71
16:2:2269:U:C6	16:2:2272:G:O6	2.44	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:6:2696:A:H62	20:6:2697:A:N6	1.88	0.71
10:h:1711:U:C5'	16:2:2256:A:OP2	2.38	0.71
15:S:46:THR:HG23	15:S:89:ILE:CD1	2.21	0.71
21:7:2853:A:C6	21:7:2854:U:N3	2.58	0.71
1:a:585:A:H2'	1:a:586:A:C8	2.26	0.71
15:S:98:ILE:O	15:S:98:ILE:HG13	1.89	0.70
19:5:2655:U:H4'	19:5:2656:A:O5'	1.90	0.70
4:d:1548:A:O4'	25:V:41:C:O2'	2.04	0.70
16:2:2284:C:H2'	16:2:2285:C:O2	1.90	0.70
16:2:2294:U:O2	16:2:2296:A:H2'	1.90	0.70
12:K:148:GLY:O	12:K:150:ARG:HG3	1.91	0.70
13:L:101:VAL:CG1	13:L:123:VAL:HG13	2.21	0.70
21:7:2846:U:O2	21:7:2850:G:O6	2.09	0.70
25:Y:43:C:H2'	25:Y:44:G:O4'	1.91	0.70
25:W:6:G:O2'	25:W:7:A:H5'	1.91	0.70
10:h:1609:C:H1'	26:v:18:C:N4	1.47	0.70
22:8:2963:C:C2'	22:8:2964:G:H5'	2.21	0.70
2:b:886:U:C3'	2:b:887:U:H5''	2.06	0.70
8:g:1165:A:H4'	8:g:1166:A:OP2	1.91	0.70
16:2:2301:U:H2'	16:2:2302:G:H8	1.55	0.70
7:f:1241:U:O2'	7:f:1242:G:OP2	2.06	0.70
10:h:1668:U:H2'	10:h:1669:G:C8	2.27	0.70
11:T:35:CYS:H	11:T:63:ILE:HD11	1.56	0.70
21:7:2834:G:N3	21:7:2835:U:C6	2.59	0.70
21:7:2872:A:H4'	21:7:2873:U:OP1	1.91	0.70
16:2:2218:G:O2'	16:2:2219:A:H5'	1.92	0.69
20:6:2703:A:C3'	20:6:2704:A:H5'	2.21	0.69
12:K:138:ASP:OD1	12:K:139:SER:N	2.25	0.69
16:2:2287:C:N3	16:2:2298:U:O2	2.25	0.69
25:Y:67:C:H2'	25:Y:68:C:C6	2.27	0.69
15:S:58:TYR:CD1	15:S:88:MET:HE2	2.27	0.69
16:2:2256:A:O2'	16:2:2257:C:P	2.49	0.69
16:2:2295:A:N6	16:2:2296:A:N6	2.40	0.69
20:6:2703:A:C5'	20:6:2704:A:H5'	2.21	0.69
13:L:69:LYS:NZ	13:L:92:LEU:HD23	2.06	0.69
25:W:16:U:H3'	25:W:17:C:C5'	2.21	0.69
16:2:2207:A:H8	16:2:2237:C:O2	1.75	0.69
16:2:2252:A:C2	16:2:2253:G:C5	2.81	0.69
16:2:2253:G:C8	16:2:2254:U:C5	2.81	0.69
19:5:2657:A:H1'	20:6:2694:A:O4'	1.93	0.69
13:L:116:ILE:CG2	13:L:119:VAL:HB	2.20	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:2:2221:G:N2	16:2:2225:U:N3	2.41	0.69
24:F:39:GLY:O	24:F:43:TYR:CB	2.41	0.69
15:S:14:ARG:HB3	15:S:114:PHE:CD2	2.28	0.69
15:S:97:GLY:O	15:S:112:VAL:HG23	1.93	0.69
11:T:24:ILE:HD11	11:T:36:THR:OG1	1.92	0.69
11:T:45:PRO:CG	11:T:85:ILE:HG23	2.22	0.69
16:2:2196:C:N4	16:2:2242:A:N7	2.40	0.69
16:2:2285:C:H5	16:2:2286:U:N3	1.90	0.69
21:7:2826:U:C2'	21:7:2827:U:H5'	2.22	0.69
11:T:158:PHE:CD2	11:T:159:LYS:HG3	2.27	0.69
18:4:2618:G:C2'	21:7:2865:U:OP2	2.36	0.69
16:2:2222:A:O2'	16:2:2223:A:C8	2.46	0.68
25:Y:63:G:H2'	25:Y:64:A:H8	1.57	0.68
25:Y:8:U:H4'	25:Y:48:C:H4'	1.75	0.68
21:7:2837:A:C8	21:7:2845:A:C2	2.81	0.68
25:Y:41:C:H2'	25:Y:42:C:O4'	1.92	0.68
10:h:1709:A:C5'	26:y:19:U:O3'	2.41	0.68
11:T:57:ARG:HH11	11:T:57:ARG:CA	1.94	0.68
16:2:2239:G:O2'	16:2:2240:G:H5'	1.93	0.68
21:7:2841:G:C6	21:7:2844:C:C4	2.82	0.68
1:a:546:G:H2'	1:a:547:C:C5	2.29	0.68
11:T:44:VAL:HG12	11:T:45:PRO:HD2	1.74	0.68
14:X:50:LEU:HD23	14:X:57:ARG:NH1	2.08	0.68
15:S:33:LEU:HD11	15:S:59:ALA:HA	1.76	0.68
20:6:2696:A:N7	20:6:2697:A:C5	2.61	0.68
1:a:572:U:H4'	1:a:573:A:H5'	1.75	0.68
10:h:1709:A:N6	26:y:19:U:H3	1.86	0.68
15:S:91:ILE:HG23	15:S:92:PRO:HD2	1.74	0.68
11:T:72:MET:O	11:T:78:ALA:HB1	1.94	0.68
12:K:21:VAL:HG12	12:K:22:GLY:N	2.04	0.68
16:2:2294:U:O2	16:2:2294:U:H2'	1.92	0.68
25:W:67:C:H2'	25:W:68:C:C6	2.29	0.68
10:h:1660:A:C6	10:h:1661:G:N7	2.62	0.67
12:K:46:ASP:OD2	12:K:48:SER:OG	2.10	0.67
7:f:1244:U:C5'	7:f:1245:G:OP2	2.41	0.67
12:K:56:VAL:HG11	12:K:77:ALA:HA	1.77	0.67
12:K:103:VAL:CG1	12:K:142:ARG:HG2	2.23	0.67
16:2:2249:G:O4'	16:2:2272:G:H8	1.76	0.67
25:W:38:A:H3'	25:W:39:U:C5'	2.20	0.67
18:4:2619:G:P	18:4:2619:G:H8	2.16	0.67
25:V:27:G:H2'	25:V:28:G:C8	2.29	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:569:C:H5'	1:a:570:G:OP2	1.93	0.67
1:a:572:U:H5''	1:a:573:A:OP1	1.95	0.67
3:c:983:A:H2'	3:c:984:C:H6	1.59	0.67
13:L:52:VAL:HG13	13:L:71:VAL:CG1	2.25	0.67
13:L:58:ILE:HG21	13:L:117:PRO:CG	2.24	0.67
16:2:2249:G:O4'	16:2:2272:G:C8	2.48	0.67
11:T:45:PRO:HG3	11:T:85:ILE:HG23	1.77	0.67
13:L:97:GLU:O	13:L:98:ASN:HB2	1.95	0.67
24:F:47:GLN:H	24:F:48:SER:HA	1.57	0.67
25:W:6:G:C2'	25:W:7:A:H5'	2.24	0.67
16:2:2222:A:O2'	16:2:2223:A:H8	1.78	0.67
16:2:2251:G:N3	16:2:2252:A:C8	2.63	0.67
16:2:2280:A:C6	16:2:2282:U:C5	2.83	0.67
16:2:2295:A:C5	16:2:2296:A:N6	2.63	0.67
21:7:2836:C:H5	21:7:2853:A:N1	1.93	0.67
25:V:16:U:O3'	25:V:17:C:H6	1.77	0.67
16:2:2279:A:C8	16:2:2288:G:C6	2.83	0.67
25:Y:66:U:H2'	25:Y:67:C:C6	2.30	0.67
11:T:55:LYS:O	11:T:56:PHE:HD1	1.78	0.66
16:2:2259:A:H3'	16:2:2260:U:H6	1.57	0.66
19:5:2657:A:O3'	20:6:2694:A:OP2	2.13	0.66
25:W:30:G:H2'	25:W:31:A:C8	2.29	0.66
10:h:1613:C:H1'	10:h:1715:A:C2	2.30	0.66
13:L:34:LEU:O	13:L:36:SER:N	2.28	0.66
20:6:2694:A:H2'	20:6:2695:A:H1'	1.77	0.66
11:T:70:THR:HB	11:T:150:ILE:HD12	1.76	0.66
13:L:117:PRO:HG2	13:L:118:GLY:H	1.60	0.66
16:2:2269:U:C4	16:2:2272:G:N1	2.61	0.66
16:2:2295:A:C6	16:2:2296:A:N6	2.62	0.66
10:h:1693:A:H2'	10:h:1694:U:H6	1.59	0.66
11:T:137:ALA:O	11:T:141:ARG:HB2	1.95	0.66
12:K:45:THR:CG2	12:K:49:GLY:HA2	2.25	0.66
12:K:95:ILE:HD11	12:K:126:ILE:CG2	2.25	0.66
13:L:86:VAL:HG12	13:L:91:CYS:HB3	1.78	0.66
16:2:2276:G:C6	16:2:2277:C:C4	2.84	0.66
1:a:546:G:C2	1:a:547:C:N4	2.63	0.66
13:L:112:ALA:HB2	13:L:119:VAL:O	1.96	0.66
16:2:2221:G:C8	16:2:2221:G:H3'	2.31	0.66
8:g:1163:U:N1	25:V:34:G:H4'	2.10	0.66
10:h:1671:G:H2'	10:h:1672:G:C8	2.31	0.66
16:2:2299:A:C4	16:2:2300:G:C8	2.84	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:7:2858:U:H2'	21:7:2859:U:C6	2.30	0.66
21:7:2861:U:H2'	21:7:2862:U:H6	1.58	0.66
22:8:2968:G:H2'	22:8:2969:A:O5'	1.94	0.66
1:a:564:A:N1	13:L:115:ASP:CB	2.58	0.66
10:h:1693:A:H2'	10:h:1694:U:C6	2.30	0.66
18:4:2619:G:OP1	18:4:2619:G:H8	1.79	0.66
11:T:95:ILE:HG21	11:T:103:PRO:HB3	1.78	0.66
15:S:69:LYS:HZ1	15:S:94:LEU:C	2.04	0.66
16:2:2226:U:HO2'	16:2:2227:C:H6	1.42	0.66
16:2:2285:C:O2'	25:Y:71:G:C4'	2.33	0.66
23:B:53:LEU:CB	23:B:54:LYS:HA	2.26	0.66
11:T:13:LEU:HD12	11:T:14:PHE:H	1.61	0.66
11:T:83:LEU:HD23	11:T:84:CYS:N	2.11	0.66
16:2:2225:U:O2'	16:2:2226:U:H5'	1.95	0.66
18:4:2619:G:O4'	21:7:2866:U:C5	2.48	0.66
24:F:14:GLY:N	24:F:15:LYS:HA	2.10	0.66
16:2:2282:U:C6	16:2:2282:U:H5'	2.30	0.66
19:5:2655:U:H4'	19:5:2656:A:H5'	1.78	0.66
20:6:2702:A:C4'	20:6:2704:A:C8	2.79	0.66
21:7:2868:U:H2'	21:7:2869:U:H6	1.61	0.66
22:8:2971:A:OP2	25:V:75:C:OP1	2.13	0.66
25:W:33:U:H2'	25:W:35:A:OP2	1.95	0.66
11:T:32:TYR:CE2	11:T:138:PRO:HG2	2.31	0.65
16:2:2254:U:H2'	16:2:2261:G:H1	1.60	0.65
13:L:19:ARG:HH11	13:L:19:ARG:CA	2.02	0.65
7:f:1239:G:O2'	7:f:1240:G:H5'	1.96	0.65
16:2:2262:A:H5''	16:2:2263:C:H5'	1.76	0.65
3:c:984:C:O2	12:K:151:LEU:HD22	1.96	0.65
12:K:82:VAL:HG21	12:K:122:SER:OG	1.95	0.65
16:2:2286:U:P	25:Y:71:G:C5'	2.84	0.65
8:g:1158:U:H2'	8:g:1159:U:O4'	1.96	0.65
14:X:66:LYS:HD2	14:X:69:LYS:HD2	1.77	0.65
9:G:1429:G:OP2	9:G:1430:C:H5	1.80	0.65
16:2:2288:G:C6	16:2:2289:U:O4	2.50	0.65
20:6:2689:A:C2	20:6:2702:A:H2	2.13	0.65
11:T:22:VAL:HG22	11:T:105:GLU:HB2	1.78	0.65
16:2:2194:G:C2'	16:2:2195:C:H5'	2.27	0.65
25:V:68:C:H2'	25:V:69:G:C5'	2.24	0.65
10:h:1709:A:O5'	26:y:19:U:C3'	2.45	0.65
13:L:53:THR:O	13:L:54:GLU:HB3	1.96	0.65
16:2:2223:A:C2	16:2:2224:A:C4	2.84	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:6:2690:G:O2'	20:6:2691:A:OP1	2.14	0.65
25:Y:55:U:H2'	25:Y:56:C:H3'	1.79	0.65
2:b:883:A:H5'	12:K:66:ARG:HB3	1.79	0.65
4:d:1549:A:O2'	25:V:31:A:O2'	2.06	0.65
16:2:2254:U:O2	16:2:2261:G:C2	2.50	0.65
18:4:2619:G:C8	21:7:2866:U:C4	2.85	0.65
15:S:132:THR:HG22	15:S:133:HIS:H	1.60	0.64
12:K:84:ARG:HD2	12:K:84:ARG:O	1.97	0.64
13:L:54:GLU:OE2	13:L:56:ILE:HD11	1.96	0.64
15:S:73:PRO:C	15:S:75:GLY:H	2.04	0.64
16:2:2278:C:C2'	16:2:2279:A:H5''	2.27	0.64
18:4:2616:C:C2'	18:4:2617:U:H5'	2.23	0.64
25:W:40:C:H2'	25:W:41:C:C6	2.32	0.64
10:h:1645:C:H42	10:h:1680:A:H61	1.46	0.64
11:T:177:LYS:O	11:T:178:ASN:HB2	1.98	0.64
22:8:2971:A:H4'	22:8:2972:G:O5'	1.95	0.64
16:2:2277:C:O2'	16:2:2278:C:O5'	2.13	0.64
10:h:1609:C:N4	25:V:34:G:N3	2.36	0.64
10:h:1648:C:H2'	10:h:1649:U:O4'	1.96	0.64
10:h:1696:U:O2'	10:h:1697:G:H5'	1.97	0.64
16:2:2276:G:C6	16:2:2277:C:C5	2.86	0.64
25:V:20:U:C3'	25:V:21:A:C5'	2.73	0.64
16:2:2197:C:N4	16:2:2241:U:H2'	2.13	0.64
21:7:2864:A:C5	21:7:2865:U:C5	2.86	0.64
22:8:2961:G:H5''	22:8:2962:U:OP2	1.97	0.64
25:Y:72:C:C2	25:Y:73:A:H2	2.16	0.64
11:T:122:ARG:HE	11:T:129:VAL:CG1	2.10	0.64
23:B:101:LYS:O	23:B:105:LYS:N	2.31	0.64
25:W:40:C:H2'	25:W:41:C:H6	1.63	0.64
11:T:13:LEU:HD12	11:T:14:PHE:N	2.12	0.64
11:T:71:LEU:O	11:T:71:LEU:HD12	1.98	0.64
15:S:83:THR:HG23	15:S:85:TYR:N	2.12	0.64
16:2:2251:G:N2	16:2:2252:A:C4	2.66	0.64
10:h:1609:C:H4'	26:v:18:C:C4	2.27	0.63
16:2:2253:G:C5	16:2:2254:U:C4	2.86	0.63
25:V:30:G:H2'	25:V:31:A:H5'	1.80	0.63
16:2:2211:U:H2'	16:2:2212:C:O4'	1.99	0.63
16:2:2285:C:C5	16:2:2286:U:N3	2.66	0.63
21:7:2830:G:O2'	21:7:2831:G:H5'	1.98	0.63
11:T:95:ILE:HG22	11:T:103:PRO:HB3	1.80	0.63
16:2:2255:A:C2	16:2:2260:U:O4	2.51	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:2:2289:U:C2	16:2:2290:C:C5	2.86	0.63
21:7:2830:G:C4	21:7:2831:G:C8	2.86	0.63
25:V:20:U:C2'	25:V:21:A:H5'	2.29	0.63
16:2:2211:U:O2	25:W:72:C:C5'	2.47	0.63
16:2:2279:A:H2	16:2:2285:C:H41	1.46	0.63
18:4:2621:G:H2'	18:4:2622:C:C6	2.33	0.63
25:W:12:U:H3	25:W:23:A:H61	1.44	0.63
10:h:1606:C:H3'	10:h:1606:C:H6	1.63	0.63
16:2:2294:U:C5'	16:2:2295:A:OP2	2.47	0.63
23:B:188:ASN:C	23:B:190:PHE:H	2.05	0.63
25:Y:5:G:H2'	25:Y:6:G:H8	1.64	0.63
8:g:1162:C:O2'	8:g:1163:U:H5''	1.97	0.63
16:2:2259:A:H5''	16:2:2260:U:H5	1.64	0.63
2:b:880:G:O6	12:K:68:GLU:CD	2.41	0.63
11:T:133:ALA:HB2	11:T:200:ARG:HB2	1.80	0.63
12:K:38:ASN:O	12:K:68:GLU:HG3	1.99	0.63
16:2:2209:U:C6	16:2:2209:U:OP2	2.52	0.63
16:2:2301:U:H2'	16:2:2302:G:C8	2.32	0.63
19:5:2657:A:C1'	20:6:2694:A:O4'	2.47	0.63
11:T:99:THR:HG22	11:T:99:THR:O	1.99	0.63
15:S:31:GLU:HG3	15:S:32:LYS:H	1.64	0.63
11:T:106:VAL:HA	11:T:176:GLU:OE1	1.99	0.63
13:L:6:PRO:HG2	13:L:15:LEU:HD21	1.79	0.63
16:2:2262:A:N3	16:2:2263:C:H5''	2.13	0.63
18:4:2618:G:H2'	21:7:2865:U:OP2	1.97	0.63
22:8:2966:G:C2'	22:8:2967:A:H5'	2.28	0.62
16:2:2251:G:C6	16:2:2266:U:O2	2.52	0.62
16:2:2279:A:N1	16:2:2283:G:N2	2.46	0.62
21:7:2857:C:H2'	21:7:2858:U:H6	1.65	0.62
1:a:570:G:H4'	1:a:574:A:C2	2.35	0.62
13:L:133:ALA:O	13:L:139:LYS:HB2	1.99	0.62
16:2:2246:G:C2	16:2:2247:G:C8	2.86	0.62
16:2:2279:A:H2	16:2:2285:C:N4	1.98	0.62
25:Y:27:G:H2'	25:Y:28:G:C8	2.34	0.62
5:e:1133:C:H1'	6:E:1591:C:H42	1.64	0.62
10:h:1709:A:N6	25:Y:37:A:H1'	2.14	0.62
16:2:2249:G:C4	16:2:2272:G:N7	2.67	0.62
21:7:2839:G:C5	21:7:2850:G:N2	2.67	0.62
21:7:2869:U:O2	21:7:2869:U:H2'	2.00	0.62
23:B:95:LYS:HA	23:B:100:ILE:N	2.15	0.62
25:Y:9:A:H2	25:Y:45:U:H3	1.42	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:T:72:MET:O	11:T:72:MET:HG3	1.99	0.62
11:T:163:SER:N	11:T:166:GLU:OE1	2.28	0.62
13:L:62:GLN:CB	13:L:63:PRO:CD	2.77	0.62
13:L:101:VAL:HA	13:L:125:CYS:O	2.00	0.62
12:K:99:ALA:O	12:K:100:LYS:C	2.43	0.62
12:K:145:GLY:O	12:K:147:ARG:N	2.33	0.62
14:X:44:ARG:HG3	14:X:45:TYR:N	2.14	0.62
15:S:42:PHE:CE2	15:S:118:GLY:HA2	2.34	0.62
15:S:66:ARG:NH1	15:S:93:GLU:HB3	2.15	0.62
20:6:2702:A:O4'	20:6:2704:A:N7	2.32	0.62
13:L:47:HIS:ND1	13:L:104:ALA:HB2	2.15	0.62
15:S:74:ALA:C	15:S:76:GLU:H	2.07	0.62
22:8:2960:C:H3'	22:8:2961:G:H8	1.65	0.62
11:T:137:ALA:HB1	11:T:138:PRO:HD2	1.80	0.62
15:S:58:TYR:O	15:S:62:VAL:HG23	1.99	0.62
1:a:586:A:C2'	1:a:587:U:H5'	2.30	0.62
21:7:2847:A:C4	21:7:2848:G:C8	2.88	0.62
24:F:38:GLN:O	24:F:39:GLY:C	2.43	0.62
9:G:1430:C:O2'	9:G:1431:A:C5'	2.48	0.61
10:h:1609:C:H4'	26:v:18:C:C5	2.34	0.61
11:T:169:ALA:O	11:T:173:ILE:HG13	2.00	0.61
17:3:2484:A:N6	25:W:19:G:C2'	2.62	0.61
15:S:78:PRO:HB2	15:S:98:ILE:HG12	1.80	0.61
15:S:124:PHE:CD1	15:S:124:PHE:N	2.68	0.61
16:2:2214:A:H2'	16:2:2215:A:H8	1.64	0.61
16:2:2285:C:HO2'	25:Y:71:G:H4'	1.63	0.61
25:Y:20:U:H2'	25:Y:21:A:H4'	1.82	0.61
11:T:193:GLU:O	11:T:197:LYS:HG3	1.99	0.61
13:L:78:ASN:HD21	13:L:80:LYS:HG3	1.64	0.61
20:6:2694:A:H2'	20:6:2695:A:C1'	2.30	0.61
1:a:574:A:N6	1:a:577:C:C2	2.69	0.61
10:h:1709:A:C5'	26:y:19:U:C3'	2.78	0.61
11:T:61:CYS:SG	11:T:66:ARG:HG2	2.41	0.61
12:K:101:GLY:O	12:K:102:GLY:C	2.43	0.61
16:2:2289:U:C2'	16:2:2290:C:H5'	2.29	0.61
6:E:1591:C:O2	6:E:1591:C:H2'	1.99	0.61
7:f:1252:C:H2'	7:f:1253:G:H8	1.65	0.61
11:T:13:LEU:CD2	11:T:104:LEU:HD21	2.28	0.61
13:L:58:ILE:CG2	13:L:117:PRO:HG3	2.27	0.61
15:S:24:LEU:HG	15:S:37:GLU:HB3	1.82	0.61
10:h:1671:G:H2'	10:h:1672:G:H8	1.64	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:T:21:GLU:HG2	11:T:102:ASN:HB2	1.81	0.61
16:2:2197:C:H4'	16:2:2198:A:C8	2.35	0.61
16:2:2215:A:N3	16:2:2216:G:C8	2.68	0.61
10:h:1609:C:C4'	26:v:18:C:C5	2.83	0.61
11:T:37:THR:HG22	11:T:38:THR:N	2.15	0.61
20:6:2690:G:HO2'	20:6:2691:A:P	2.23	0.61
16:2:2258:U:H3'	16:2:2259:A:H8	1.66	0.61
22:8:2973:G:O2'	22:8:2974:U:H5'	2.00	0.61
1:a:547:C:H5''	1:a:548:A:OP1	2.00	0.61
2:b:882:G:N7	2:b:883:A:C4	2.69	0.61
4:d:1548:A:H1'	25:V:41:C:H1'	1.82	0.61
20:6:2702:A:H4'	20:6:2704:A:C8	2.36	0.61
20:6:2703:A:C3'	20:6:2704:A:C5'	2.78	0.61
23:B:54:LYS:HA	23:B:55:LEU:CB	2.31	0.61
12:K:36:THR:HG22	12:K:37:TRP:H	1.65	0.60
16:2:2270:A:C2	16:2:2271:A:N3	2.69	0.60
11:T:40:SER:O	11:T:42:VAL:HG23	2.01	0.60
14:X:31:PRO:HG3	14:X:39:ILE:HD11	1.83	0.60
16:2:2235:C:C4	16:2:2236:G:C6	2.89	0.60
20:6:2696:A:N7	20:6:2697:A:N7	2.49	0.60
1:a:565:G:C8	1:a:566:C:C5	2.89	0.60
10:h:1710:G:O2'	16:2:2256:A:H5''	1.99	0.60
16:2:2219:A:H2'	16:2:2220:A:H5'	1.83	0.60
21:7:2851:A:H3'	21:7:2852:C:H5''	1.82	0.60
22:8:2961:G:C5	22:8:2962:U:C5	2.90	0.60
25:V:27:G:H2'	25:V:28:G:H8	1.67	0.60
11:T:16:LYS:O	11:T:17:TRP:CG	2.54	0.60
11:T:26:ASP:OD1	11:T:28:CYS:HB3	2.01	0.60
1:a:559:C:O2	1:a:559:C:H2'	2.00	0.60
16:2:2219:A:C2'	16:2:2220:A:H5'	2.31	0.60
20:6:2702:A:H8	20:6:2704:A:C5	2.19	0.60
23:B:55:LEU:N	23:B:186:SER:HA	2.17	0.60
3:c:986:G:OP1	12:K:149:ARG:NE	2.34	0.60
21:7:2861:U:H2'	21:7:2862:U:O4'	2.02	0.60
21:7:2831:G:C6	21:7:2832:C:C4	2.90	0.60
21:7:2842:U:O2	21:7:2842:U:C2'	2.49	0.60
25:W:56:C:C3'	25:W:57:G:H5''	2.32	0.60
7:f:1246:C:N4	25:Y:34:G:O4'	2.34	0.60
13:L:34:LEU:O	13:L:35:GLY:C	2.44	0.60
22:8:2966:G:C6	22:8:2967:A:N6	2.70	0.60
22:8:2970:C:C3'	22:8:2971:A:H5''	2.31	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:X:38:ARG:HG2	14:X:42:ASN:HD21	1.67	0.60
16:2:2204:C:O5'	16:2:2204:C:H6	1.85	0.60
18:4:2618:G:C2'	21:7:2865:U:O5'	2.48	0.60
25:Y:42:C:H3'	25:Y:43:C:H5''	1.84	0.60
15:S:56:GLU:N	15:S:56:GLU:CD	2.60	0.60
16:2:2258:U:H2'	16:2:2258:U:O2	2.01	0.60
23:B:95:LYS:CB	23:B:124:LEU:HA	2.31	0.60
13:L:54:GLU:HG3	13:L:56:ILE:HG13	1.83	0.59
15:S:86:ARG:NH1	15:S:122:ALA:HB1	2.17	0.59
16:2:2202:C:O2	16:2:2202:C:H2'	2.02	0.59
16:2:2269:U:O2	16:2:2269:U:C2'	2.50	0.59
16:2:2280:A:H4'	16:2:2281:A:OP1	2.02	0.59
25:V:68:C:C3'	25:V:69:G:H5''	2.33	0.59
1:a:546:G:H2'	1:a:547:C:C6	2.37	0.59
8:g:1157:U:O2'	8:g:1158:U:P	2.60	0.59
13:L:60:SER:OG	13:L:64:ASN:HB2	2.01	0.59
14:X:34:ARG:HB3	14:X:34:ARG:HH11	1.64	0.59
15:S:66:ARG:HH12	15:S:93:GLU:CB	2.15	0.59
22:8:2968:G:C2'	22:8:2969:A:O5'	2.49	0.59
11:T:101:ARG:O	11:T:103:PRO:HD3	2.02	0.59
12:K:27:VAL:HG21	12:K:89:LYS:O	2.02	0.59
21:7:2830:G:H1	21:7:2858:U:H3	1.48	0.59
25:W:58:A:H1'	25:W:60:U:C5	2.37	0.59
10:h:1701:C:H2'	10:h:1702:A:H5'	1.84	0.59
16:2:2253:G:C8	16:2:2254:U:C4	2.91	0.59
18:4:2618:G:C4'	18:4:2619:G:OP2	2.50	0.59
9:G:1429:G:H2'	9:G:1429:G:N3	2.16	0.59
11:T:189:LYS:C	11:T:189:LYS:HD3	2.28	0.59
25:Y:18:G:N1	25:Y:57:G:N7	2.50	0.59
12:K:121:ARG:HG3	12:K:122:SER:H	1.66	0.59
16:2:2290:C:C2	16:2:2303:A:C2	2.90	0.59
22:8:2963:C:H2'	22:8:2964:G:H5'	1.82	0.59
21:7:2827:U:O2	21:7:2827:U:H2'	2.02	0.59
23:B:95:LYS:HA	23:B:100:ILE:H	1.68	0.59
25:V:67:C:H2'	25:V:68:C:C6	2.38	0.59
12:K:62:VAL:CG1	12:K:63:LYS:H	2.14	0.59
16:2:2211:U:H1'	25:W:72:C:O4'	2.02	0.59
21:7:2849:C:O2	21:7:2849:C:H2'	2.03	0.59
9:G:1434:C:OP1	25:V:30:G:H4'	2.02	0.59
12:K:27:VAL:O	12:K:92:ALA:HB3	2.03	0.59
15:S:36:ASP:O	15:S:39:ALA:HB3	2.03	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:6:2689:A:O2'	20:6:2702:A:C6	2.54	0.59
21:7:2836:C:C2'	21:7:2836:C:O2	2.51	0.59
25:W:38:A:C2'	25:W:39:U:H5''	2.32	0.59
10:h:1707:G:H2'	10:h:1708:A:C8	2.38	0.58
11:T:34:ALA:CB	11:T:62:PRO:HA	2.33	0.58
16:2:2226:U:O2'	16:2:2227:C:H6	1.86	0.58
16:2:2294:U:C6	16:2:2297:U:H5	2.20	0.58
24:F:47:GLN:N	24:F:48:SER:CA	2.66	0.58
1:a:552:C:H4'	14:X:66:LYS:CE	2.31	0.58
11:T:55:LYS:O	11:T:56:PHE:CD1	2.56	0.58
13:L:78:ASN:ND2	13:L:80:LYS:CG	2.65	0.58
16:2:2285:C:C5	16:2:2286:U:C4	2.91	0.58
16:2:2299:A:C5	16:2:2300:G:N7	2.71	0.58
19:5:2655:U:C2	19:5:2656:A:N1	2.71	0.58
7:f:1246:C:O2'	7:f:3247:A:H5''	2.04	0.58
20:6:2698:G:C3'	20:6:2699:G:H5''	2.33	0.58
25:Y:68:C:H2'	25:Y:69:G:H8	1.68	0.58
25:W:37:A:H2'	25:W:38:A:O4'	2.03	0.58
11:T:183:SER:OG	11:T:186:ILE:HB	2.03	0.58
15:S:69:LYS:HE2	15:S:97:GLY:N	2.18	0.58
15:S:132:THR:HG22	15:S:133:HIS:N	2.19	0.58
16:2:2211:U:H1'	25:W:72:C:C4'	2.32	0.58
25:Y:6:G:O2'	25:Y:7:A:H5'	2.03	0.58
2:b:879:G:OP2	12:K:39:ASP:HB2	2.04	0.58
10:h:1710:G:O2'	16:2:2256:A:O5'	2.22	0.58
11:T:14:PHE:HE2	11:T:89:LYS:HA	1.64	0.58
12:K:36:THR:CG2	12:K:37:TRP:N	2.65	0.58
15:S:91:ILE:O	15:S:94:LEU:HB2	2.03	0.58
16:2:2244:A:O2'	16:2:2245:C:H5'	2.04	0.58
21:7:2859:U:H4'	21:7:2860:U:OP1	2.00	0.58
25:V:25:C:O2'	25:V:26:A:H5'	2.03	0.58
10:h:1608:C:N3	10:h:1610:G:C2	2.72	0.58
12:K:45:THR:HG22	12:K:49:GLY:HA2	1.85	0.58
14:X:63:HIS:C	14:X:65:GLY:N	2.56	0.58
16:2:2199:G:C6	16:2:2200:U:C4	2.92	0.58
16:2:2210:G:H22	25:W:72:C:P	2.27	0.58
20:6:2689:A:H4'	20:6:2690:G:C5'	2.33	0.58
21:7:2836:C:O2	21:7:2837:A:C8	2.57	0.58
25:W:14:A:H3'	25:W:15:G:H8	1.69	0.58
10:h:1629:G:H1	10:h:1696:U:H3	1.51	0.58
10:h:1701:C:C2'	10:h:1702:A:H5'	2.33	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:T:170:ASP:HA	11:T:173:ILE:HD12	1.85	0.58
12:K:95:ILE:HD12	12:K:129:ILE:CG2	2.32	0.58
16:2:2196:C:O2'	16:2:2271:A:O4'	2.22	0.58
16:2:2248:C:C4'	16:2:2271:A:H2	2.17	0.58
10:h:1613:C:C1'	10:h:1715:A:H2	2.12	0.58
15:S:82:LYS:HB3	15:S:107:PHE:CE2	2.39	0.58
16:2:2269:U:N3	16:2:2272:G:N1	2.45	0.58
10:h:1715:A:C8	10:h:1715:A:H3'	2.38	0.58
13:L:56:ILE:HG22	13:L:57:GLY:N	2.18	0.58
11:T:51:TYR:CE2	11:T:61:CYS:HA	2.39	0.58
16:2:2287:C:O2	16:2:2298:U:O2	2.22	0.58
21:7:2856:G:H2'	21:7:2857:C:C6	2.39	0.58
25:Y:9:A:C8	25:Y:46:G:N2	2.71	0.58
12:K:110:PRO:C	12:K:112:ALA:H	2.12	0.57
15:S:58:TYR:HD1	15:S:88:MET:HE2	1.67	0.57
16:2:2210:G:N2	25:W:72:C:H5'	2.18	0.57
16:2:2260:U:C2'	16:2:2261:G:N7	2.65	0.57
16:2:2288:G:N3	16:2:2289:U:C5	2.72	0.57
13:L:37:ARG:HG2	13:L:44:GLY:HA2	1.84	0.57
16:2:2211:U:O2'	25:W:72:C:C1'	2.52	0.57
21:7:2853:A:N6	21:7:2854:U:H3	2.00	0.57
10:h:1608:C:C4	10:h:1610:G:N3	2.72	0.57
13:L:52:VAL:HG13	13:L:71:VAL:HG13	1.84	0.57
16:2:2285:C:H5	16:2:2286:U:N1	2.03	0.57
15:S:56:GLU:CD	15:S:56:GLU:H	2.13	0.57
16:2:2210:G:C6	16:2:2211:U:C4	2.92	0.57
16:2:2253:G:C5	16:2:2254:U:N3	2.72	0.57
16:2:2260:U:H3'	16:2:2261:G:C8	2.39	0.57
25:V:40:C:C5'	25:W:35:A:H4'	2.32	0.57
25:W:68:C:O2'	25:W:69:G:H5'	2.05	0.57
12:K:22:GLY:C	12:K:24:ASN:H	2.11	0.57
18:4:2621:G:H2'	18:4:2622:C:H6	1.69	0.57
14:X:58:LYS:HD3	14:X:62:TRP:CH2	2.39	0.57
16:2:2261:G:N2	16:2:2263:C:C4	2.72	0.57
10:h:1606:C:H3'	10:h:1606:C:C6	2.39	0.57
11:T:24:ILE:HD12	11:T:30:GLN:HA	1.87	0.57
11:T:163:SER:HB3	11:T:166:GLU:HG3	1.85	0.57
11:T:96:HIS:O	11:T:100:GLY:N	2.38	0.57
12:K:116:LEU:HD23	12:K:116:LEU:C	2.29	0.57
16:2:2194:G:H1'	16:2:2274:U:O2	2.04	0.57
20:6:2704:A:C2	20:6:2706:G:C8	2.92	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:T:26:ASP:OD1	11:T:28:CYS:CB	2.53	0.57
12:K:36:THR:CG2	12:K:37:TRP:H	2.18	0.57
12:K:105:THR:C	12:K:107:GLN:H	2.13	0.57
16:2:2253:G:H3'	16:2:2254:U:H5	1.69	0.57
16:2:2290:C:H2'	16:2:2291:A:H8	1.70	0.57
20:6:2689:A:O2'	20:6:2702:A:N6	2.38	0.57
12:K:98:ARG:NH2	12:K:100:LYS:HA	2.20	0.57
13:L:51:LEU:HD21	13:L:76:ARG:HE	1.70	0.57
13:L:132:LEU:HD21	13:L:136:LYS:HE3	1.87	0.57
16:2:2266:U:C4	16:2:2267:C:N4	2.72	0.57
11:T:116:PRO:HG3	11:T:189:LYS:HA	1.86	0.56
13:L:106:LEU:HD21	13:L:121:PHE:C	2.30	0.56
16:2:2265:C:N4	16:2:2266:U:N3	2.53	0.56
16:2:2288:G:H2'	16:2:2289:U:C6	2.40	0.56
16:2:2304:C:C4	16:2:2305:G:N2	2.72	0.56
17:3:2482:U:H2'	17:3:2483:G:O4'	2.05	0.56
1:a:562:G:OP1	13:L:69:LYS:HE2	2.05	0.56
10:h:1653:C:H2'	10:h:1654:U:O4'	2.05	0.56
16:2:2202:C:H5''	16:2:2203:U:OP2	2.04	0.56
16:2:2204:C:H2'	16:2:2205:U:H5''	1.86	0.56
16:2:2260:U:C3'	16:2:2261:G:C8	2.88	0.56
16:2:2288:G:C2	16:2:2289:U:C4	2.93	0.56
25:V:17:C:H3'	25:V:18:G:C5'	2.29	0.56
8:g:1163:U:C2	25:V:34:G:C5'	2.87	0.56
10:h:1657:G:H2'	10:h:1658:A:C8	2.40	0.56
12:K:116:LEU:C	12:K:116:LEU:CD2	2.79	0.56
16:2:2246:G:N3	16:2:2247:G:C8	2.73	0.56
16:2:2299:A:C6	16:2:2300:G:C8	2.93	0.56
8:g:1157:U:O2'	8:g:1158:U:OP2	2.20	0.56
13:L:101:VAL:HG13	13:L:123:VAL:HG13	1.86	0.56
16:2:2221:G:C8	16:2:2221:G:C3'	2.88	0.56
16:2:2283:G:H1'	16:2:2284:C:H5	1.69	0.56
1:a:586:A:H2'	1:a:587:U:H5'	1.86	0.56
10:h:1697:G:H2'	10:h:1698:G:C8	2.40	0.56
13:L:116:ILE:HG23	13:L:117:PRO:HD2	1.88	0.56
16:2:2210:G:C6	16:2:2236:G:N2	2.73	0.56
16:2:2270:A:C8	16:2:2270:A:C5'	2.76	0.56
21:7:2842:U:O5'	21:7:2842:U:C6	2.58	0.56
11:T:153:CYS:O	11:T:167:THR:HG21	2.05	0.56
12:K:78:ALA:HB1	12:K:119:LEU:HD13	1.87	0.56
16:2:2286:U:C4	16:2:2288:G:C1'	2.88	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:7:2833:A:N1	21:7:2834:G:C8	2.74	0.56
21:7:2834:G:O2'	21:7:2835:U:C6	2.41	0.56
10:h:1704:C:O2'	10:h:1705:A:H5'	2.06	0.56
15:S:86:ARG:O	15:S:122:ALA:HB2	2.06	0.56
15:S:95:VAL:HG21	15:S:117:ILE:HD11	1.86	0.56
25:W:2:C:N4	25:W:71:G:H1	2.02	0.56
10:h:1609:C:C1'	25:V:34:G:O6	2.53	0.56
11:T:122:ARG:HH21	11:T:129:VAL:HG21	1.70	0.56
22:8:2967:A:O2'	22:8:2968:G:OP1	2.24	0.56
1:a:546:G:C4	1:a:547:C:C5	2.93	0.56
16:2:2260:U:C2'	16:2:2261:G:C8	2.89	0.56
21:7:2847:A:C2	21:7:2848:G:H1'	2.41	0.56
25:W:52:G:H2'	25:W:53:G:H8	1.71	0.56
1:a:544:G:C6	1:a:576:U:C5	2.94	0.55
11:T:13:LEU:HB2	11:T:19:TYR:CE1	2.41	0.55
11:T:189:LYS:O	11:T:193:GLU:HG3	2.06	0.55
13:L:81:LYS:HD2	13:L:81:LYS:O	2.06	0.55
15:S:31:GLU:HG3	15:S:32:LYS:HG2	1.88	0.55
16:2:2254:U:C6	16:2:2254:U:OP2	2.58	0.55
10:h:1693:A:O2'	10:h:1694:U:H5'	2.06	0.55
12:K:66:ARG:HG3	12:K:67:GLU:OE1	2.05	0.55
21:7:2833:A:C6	21:7:2834:G:N7	2.74	0.55
23:B:4:ILE:O	23:B:5:THR:C	2.48	0.55
25:W:12:U:H3	25:W:23:A:N6	2.03	0.55
25:W:38:A:H2'	25:W:39:U:C4'	2.35	0.55
10:h:1611:C:H2'	10:h:1612:C:H5'	1.87	0.55
11:T:180:THR:O	11:T:180:THR:HG22	2.05	0.55
15:S:88:MET:SD	15:S:89:ILE:N	2.79	0.55
16:2:2194:G:C1'	16:2:2274:U:O2	2.54	0.55
16:2:2223:A:O2'	16:2:2224:A:O5'	2.25	0.55
16:2:2290:C:O2'	16:2:2291:A:O4'	2.20	0.55
1:a:570:G:C4'	1:a:574:A:C2	2.88	0.55
11:T:189:LYS:HD3	11:T:189:LYS:O	2.06	0.55
21:7:2862:U:C2'	21:7:2863:G:H5'	2.36	0.55
25:W:38:A:C3'	25:W:39:U:C5'	2.83	0.55
25:W:68:C:H2'	25:W:69:G:H8	1.71	0.55
7:f:1245:G:C3'	7:f:1246:C:H5''	2.37	0.55
12:K:77:ALA:O	12:K:81:VAL:HG23	2.06	0.55
13:L:71:VAL:HG23	13:L:86:VAL:CG2	2.37	0.55
16:2:2211:U:O2'	25:W:72:C:C4'	2.55	0.55
21:7:2868:U:C5	21:7:2869:U:H5	2.24	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:B:155:ILE:HA	23:B:156:LYS:CB	2.37	0.55
25:W:39:U:H4'	25:W:39:U:OP1	2.07	0.55
11:T:37:THR:CG2	11:T:38:THR:N	2.69	0.55
15:S:86:ARG:HB3	15:S:122:ALA:HB2	1.89	0.55
16:2:2215:A:C4	16:2:2216:G:C8	2.94	0.55
16:2:2280:A:C4	16:2:2282:U:H5	2.25	0.55
20:6:2689:A:C4	20:6:2702:A:C2	2.94	0.55
24:F:40:LYS:O	24:F:42:ARG:N	2.40	0.55
25:Y:16:U:H3'	25:Y:17:C:C5'	2.34	0.55
25:Y:25:C:H2'	25:Y:26:A:C8	2.31	0.55
10:h:1652:A:N1	10:h:1674:A:H1'	2.22	0.55
11:T:47:THR:C	11:T:49:GLY:N	2.60	0.55
11:T:120:SER:OG	11:T:196:ALA:HA	2.06	0.55
16:2:2196:C:C4	16:2:2242:A:N7	2.75	0.55
16:2:2247:G:C6	16:2:2248:C:C4	2.95	0.55
16:2:2260:U:H4'	16:2:2261:G:OP2	2.07	0.55
18:4:2626:A:H3'	18:4:2626:A:N3	2.21	0.55
10:h:1608:C:H42	10:h:1610:G:N2	2.05	0.55
16:2:2217:U:O2'	16:2:2218:G:H5'	2.07	0.55
16:2:2265:C:N4	16:2:2266:U:C4	2.75	0.55
16:2:2275:A:H3'	16:2:2276:G:O4'	2.06	0.55
18:4:2615:G:C6	18:4:2616:C:N4	2.75	0.55
18:4:2622:C:H2'	18:4:2623:G:O4'	2.07	0.55
20:6:2699:G:H2'	20:6:2700:G:C8	2.42	0.55
10:h:1661:G:N3	10:h:1661:G:C3'	2.69	0.55
10:h:1710:G:H5'	10:h:1711:U:OP2	2.07	0.55
12:K:118:ALA:HA	12:K:121:ARG:HG2	1.88	0.55
14:X:59:SER:C	14:X:61:ASN:H	2.15	0.55
23:B:192:SER:HA	23:B:197:ASN:O	2.06	0.55
2:b:883:A:N6	2:b:884:A:C2	2.74	0.55
11:T:34:ALA:H	11:T:63:ILE:CG1	2.20	0.55
11:T:99:THR:O	11:T:101:ARG:N	2.40	0.55
16:2:2270:A:N1	16:2:2271:A:C2	2.75	0.55
16:2:2282:U:O2'	16:2:2283:G:C5'	2.54	0.55
21:7:2833:A:C6	21:7:2834:G:C8	2.95	0.55
25:V:53:G:H2'	25:V:54:U:C6	2.41	0.55
1:a:553:A:O2'	1:a:554:U:OP2	2.19	0.54
2:b:884:A:OP1	12:K:66:ARG:HG2	2.07	0.54
5:e:1133:C:O2'	5:e:1134:C:H5'	2.07	0.54
8:g:1163:U:C2	25:V:34:G:H5'	2.40	0.54
11:T:68:ILE:HD13	11:T:85:ILE:HA	1.88	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:K:106:LYS:HZ1	12:K:136:PRO:HD2	1.73	0.54
16:2:2223:A:C2'	16:2:2224:A:C8	2.89	0.54
16:2:2290:C:C2	16:2:2291:A:C8	2.96	0.54
16:2:2295:A:H62	16:2:2296:A:N6	2.05	0.54
13:L:42:PHE:CZ	13:L:103:VAL:HG23	2.42	0.54
15:S:101:VAL:CG2	15:S:121:LEU:HD11	2.36	0.54
16:2:2257:C:H3'	16:2:2258:U:C6	2.40	0.54
25:Y:64:A:H2'	25:Y:65:G:H8	1.72	0.54
11:T:66:ARG:HH22	11:T:143:ASN:HD21	1.54	0.54
11:T:66:ARG:NH2	11:T:143:ASN:HD21	2.05	0.54
16:2:2221:G:N2	16:2:2225:U:C2	2.75	0.54
21:7:2837:A:C8	21:7:2845:A:N1	2.75	0.54
25:V:65:G:O2'	25:V:66:U:H5'	2.07	0.54
25:W:57:G:H2'	25:W:57:G:N3	2.21	0.54
11:T:68:ILE:HG23	11:T:84:CYS:HB3	1.87	0.54
12:K:95:ILE:HD11	12:K:126:ILE:HG21	1.89	0.54
12:K:103:VAL:HG12	12:K:142:ARG:CG	2.37	0.54
16:2:2240:G:H2'	16:2:2241:U:O5'	2.07	0.54
10:h:1679:A:H2'	10:h:1680:A:C8	2.41	0.54
16:2:2290:C:N3	16:2:2303:A:N1	2.56	0.54
18:4:2617:U:C5	18:4:2620:G:OP1	2.61	0.54
18:4:2619:G:C8	21:7:2866:U:O4	2.60	0.54
21:7:2836:C:O2	21:7:2836:C:H3'	2.08	0.54
24:F:77:CYS:C	24:F:79:THR:H	2.14	0.54
1:a:560:C:O2'	14:X:10:LYS:HE2	2.07	0.54
11:T:22:VAL:HG12	11:T:23:LYS:N	2.22	0.54
14:X:42:ASN:HA	14:X:46:ALA:HB2	1.90	0.54
16:2:2214:A:H2'	16:2:2215:A:C8	2.41	0.54
16:2:2269:U:C5	16:2:2272:G:O6	2.60	0.54
19:5:2656:A:C8	19:5:2658:G:C8	2.96	0.54
25:Y:15:G:H22	25:Y:59:U:H3	1.54	0.54
25:V:30:G:C2'	25:V:31:A:H5'	2.37	0.54
1:a:551:U:O4	14:X:62:TRP:HB2	2.08	0.54
11:T:183:SER:C	11:T:185:ALA:N	2.63	0.54
15:S:70:GLU:HG3	15:S:71:LYS:H	1.71	0.54
16:2:2210:G:N2	25:W:72:C:OP1	2.40	0.54
21:7:2861:U:C2	21:7:2862:U:C6	2.96	0.54
21:7:2868:U:H5''	21:7:2869:U:OP2	2.08	0.54
25:V:41:C:O2	25:V:41:C:H2'	2.06	0.54
1:a:553:A:H2'	1:a:553:A:N3	2.21	0.54
11:T:200:ARG:HG3	11:T:200:ARG:OXT	2.07	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:S:49:ARG:HH22	15:S:87:SER:HB2	1.73	0.54
16:2:2257:C:H5'	16:2:2258:U:H5	1.72	0.54
23:B:155:ILE:HA	23:B:156:LYS:C	2.32	0.54
16:2:2254:U:C5	16:2:2254:U:OP2	2.61	0.54
16:2:2259:A:H3'	16:2:2260:U:C5	2.42	0.54
16:2:2261:G:N2	16:2:2263:C:N3	2.56	0.54
18:4:2619:G:OP1	18:4:2619:G:C8	2.60	0.54
21:7:2838:A:C2	21:7:2851:A:N9	2.76	0.54
21:7:2838:A:N1	21:7:2851:A:C8	2.75	0.54
21:7:2872:A:H1'	21:7:2873:U:H5'	1.90	0.54
2:b:878:A:P	12:K:59:GLY:HA3	2.47	0.54
10:h:1712:C:H2'	10:h:1713:G:O4'	2.08	0.54
11:T:64:VAL:O	11:T:68:ILE:HG13	2.08	0.54
11:T:122:ARG:HH21	11:T:129:VAL:CG2	2.21	0.54
13:L:6:PRO:HG2	13:L:15:LEU:CD2	2.38	0.54
16:2:2269:U:H2'	16:2:2270:A:H5'	1.89	0.54
25:Y:64:A:H2'	25:Y:65:G:C8	2.43	0.54
10:h:1660:A:H5'	10:h:1661:G:OP2	2.07	0.53
1:a:574:A:H3'	1:a:575:U:C5'	2.39	0.53
11:T:50:ARG:HG2	11:T:53:VAL:HG21	1.90	0.53
14:X:36:TYR:CE2	14:X:40:LEU:HD11	2.44	0.53
14:X:50:LEU:CD2	14:X:57:ARG:HH12	2.19	0.53
25:V:74:C:H2'	25:V:75:C:H5'	1.89	0.53
16:2:2285:C:O3'	25:Y:71:G:O3'	2.27	0.53
20:6:2692:A:C6	20:6:2693:C:C4	2.96	0.53
20:6:2704:A:C4	20:6:2706:G:N7	2.76	0.53
21:7:2833:A:H2'	21:7:2833:A:N3	2.23	0.53
24:F:3:ASN:HA	24:F:93:LEU:O	2.09	0.53
25:W:16:U:C4	25:W:18:G:H3'	2.43	0.53
11:T:157:ALA:O	11:T:158:PHE:O	2.25	0.53
12:K:43:HIS:HE1	12:K:52:THR:HG23	1.73	0.53
12:K:62:VAL:CG1	12:K:63:LYS:N	2.71	0.53
21:7:2853:A:N6	21:7:2854:U:N3	2.56	0.53
21:7:2861:U:H2'	21:7:2862:U:C6	2.41	0.53
24:F:80:ARG:O	24:F:81:ALA:C	2.51	0.53
25:W:39:U:O2	25:W:39:U:H5'	2.07	0.53
6:E:1590:C:O2'	6:E:1591:C:H5''	2.08	0.53
8:g:1171:G:HO2'	8:g:1172:G:P	2.32	0.53
21:7:2833:A:C2	21:7:2834:G:C8	2.97	0.53
25:Y:74:C:H2'	25:Y:75:C:H5'	1.89	0.53
7:f:1249:G:H3'	7:f:1250:G:C8	2.44	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:h:1650:G:H2'	10:h:1651:G:O4'	2.09	0.53
10:h:1654:U:H2'	10:h:1655:G:C8	2.43	0.53
12:K:42:ILE:HD11	12:K:115:ALA:HB1	1.91	0.53
15:S:15:GLY:N	15:S:114:PHE:HE2	2.07	0.53
16:2:2252:A:C2	16:2:2253:G:C4	2.97	0.53
25:Y:33:U:C3'	25:Y:34:G:H5''	2.38	0.53
8:g:1171:G:O2'	8:g:1172:G:P	2.67	0.53
12:K:34:MET:HA	12:K:98:ARG:HG3	1.90	0.53
13:L:47:HIS:CE1	13:L:104:ALA:HB2	2.44	0.53
14:X:53:ASP:OD2	14:X:56:LYS:HD2	2.09	0.53
15:S:59:ALA:O	15:S:62:VAL:HB	2.09	0.53
16:2:2281:A:N3	22:8:2974:U:O4'	2.42	0.53
21:7:2834:G:N3	21:7:2835:U:C5	2.76	0.53
24:F:4:VAL:O	24:F:5:PRO:CB	2.56	0.53
24:F:11:TYR:HA	24:F:20:HIS:HA	1.90	0.53
24:F:76:LYS:C	24:F:78:LYS:N	2.66	0.53
25:V:53:G:H2'	25:V:54:U:H6	1.73	0.53
25:W:21:A:H5'	25:W:21:A:H8	1.73	0.53
10:h:1629:G:H2'	10:h:1630:A:C8	2.43	0.53
11:T:133:ALA:HB2	11:T:200:ARG:CB	2.39	0.53
13:L:107:GLY:O	13:L:108:ARG:O	2.27	0.53
15:S:48:ARG:O	15:S:52:ARG:HB2	2.08	0.53
16:2:2270:A:O2'	16:2:2271:A:H5'	2.08	0.53
20:6:2700:G:H5''	20:6:2701:U:OP2	2.07	0.53
20:6:2701:U:H1'	20:6:2705:A:C4	2.44	0.53
21:7:2841:G:C4	21:7:2844:C:N4	2.77	0.53
7:f:1245:G:O2'	7:f:1246:C:P	2.66	0.53
10:h:1692:C:H2'	10:h:1693:A:H8	1.74	0.53
13:L:131:LEU:O	13:L:132:LEU:C	2.51	0.53
16:2:2299:A:C6	16:2:2300:G:N7	2.76	0.53
21:7:2829:U:C2'	21:7:2830:G:H5'	2.36	0.53
21:7:2852:C:H5	21:7:2853:A:C5	2.27	0.53
11:T:17:TRP:CD1	11:T:103:PRO:HD2	2.44	0.53
11:T:83:LEU:HD23	11:T:83:LEU:C	2.34	0.53
12:K:75:MET:HE3	12:K:121:ARG:HH22	1.74	0.53
1:a:564:A:H61	13:L:115:ASP:HB3	1.74	0.52
16:2:2291:A:C2	16:2:2292:U:C2	2.97	0.52
21:7:2836:C:O2	21:7:2836:C:H2'	2.09	0.52
25:W:18:G:H1	25:W:55:U:C1'	2.20	0.52
8:g:1148:G:H1'	8:g:1167:C:O2'	2.09	0.52
10:h:1623:A:O2'	10:h:1624:G:H8	1.92	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:2:2300:G:C6	16:2:2301:U:C4	2.97	0.52
20:6:2702:A:N7	20:6:2704:A:N6	2.57	0.52
21:7:2841:G:H3'	21:7:2842:U:H5'	1.91	0.52
12:K:45:THR:HG23	12:K:51:GLU:O	2.09	0.52
15:S:38:LEU:O	15:S:42:PHE:CD1	2.62	0.52
16:2:2302:G:C6	16:2:2303:A:C5	2.98	0.52
18:4:2619:G:O4'	21:7:2866:U:H5	1.93	0.52
24:F:57:VAL:HA	24:F:58:PHE:CB	2.39	0.52
7:f:1249:G:H3'	7:f:1250:G:H8	1.75	0.52
11:T:197:LYS:HD3	11:T:200:ARG:CZ	2.39	0.52
15:S:45:LYS:O	15:S:48:ARG:HB3	2.10	0.52
16:2:2233:A:C8	16:2:2234:G:C8	2.97	0.52
16:2:2259:A:C5	16:2:2260:U:C2	2.98	0.52
21:7:2866:U:H1'	21:7:2867:C:H5	1.74	0.52
22:8:2959:C:C5'	22:8:2960:C:OP2	2.57	0.52
22:8:2967:A:C2'	22:8:2968:G:OP1	2.57	0.52
22:8:2974:U:H2'	22:8:2975:U:C6	2.45	0.52
25:Y:59:U:O2'	25:Y:60:U:H5'	2.10	0.52
11:T:125:THR:HB	11:T:130:ARG:NH2	2.25	0.52
15:S:38:LEU:O	15:S:42:PHE:HD1	1.92	0.52
15:S:92:PRO:O	15:S:95:VAL:HG23	2.09	0.52
15:S:127:THR:O	15:S:127:THR:HG22	2.09	0.52
8:g:1166:A:O2'	8:g:1167:C:H5'	2.09	0.52
10:h:1651:G:H1'	10:h:1674:A:N6	2.24	0.52
12:K:30:VAL:HG22	12:K:94:HIS:HB2	1.92	0.52
13:L:112:ALA:CB	13:L:119:VAL:O	2.57	0.52
16:2:2304:C:O2'	16:2:2305:G:C5'	2.48	0.52
18:4:2617:U:C6	18:4:2620:G:OP1	2.63	0.52
25:V:5:G:H1'	25:V:69:G:N2	2.25	0.52
25:V:41:C:H3'	25:V:42:C:C5'	2.38	0.52
25:V:72:C:H2'	25:V:73:A:H5'	1.91	0.52
25:W:38:A:H2'	25:W:39:U:H5''	1.90	0.52
10:h:1709:A:H5'	26:y:19:U:O3'	2.09	0.52
11:T:107:PHE:HA	11:T:172:ILE:CG2	2.39	0.52
12:K:45:THR:HG22	12:K:46:ASP:N	2.24	0.52
12:K:106:LYS:HZ3	12:K:135:ILE:HG23	1.75	0.52
16:2:2252:A:C4	16:2:2253:G:C8	2.98	0.52
25:W:20:U:H2'	25:W:21:A:C4'	2.36	0.52
1:a:561:A:OP1	13:L:67:VAL:HB	2.10	0.52
11:T:96:HIS:O	11:T:100:GLY:HA2	2.10	0.52
16:2:2262:A:C5'	16:2:2263:C:OP2	2.58	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:3:2487:U:H4'	17:3:2488:A:OP2	2.08	0.52
25:V:41:C:C2'	25:V:42:C:H5''	2.40	0.52
1:a:542:G:H2'	1:a:543:A:H8	1.74	0.52
10:h:1608:C:H4'	10:h:1609:C:OP1	2.10	0.52
16:2:2207:A:H8	16:2:2237:C:C2	2.28	0.52
16:2:2290:C:C2'	16:2:2291:A:O5'	2.58	0.52
21:7:2850:G:C8	21:7:2850:G:H5''	2.44	0.52
24:F:38:GLN:O	24:F:40:LYS:N	2.42	0.52
8:g:1151:G:H2'	8:g:1152:C:C6	2.45	0.52
15:S:33:LEU:HD22	15:S:62:VAL:HG21	1.92	0.52
10:h:1654:U:H3	10:h:1671:G:H1	1.57	0.51
10:h:1701:C:H2'	10:h:1702:A:O4'	2.10	0.51
11:T:33:ILE:HG23	11:T:34:ALA:N	2.24	0.51
12:K:82:VAL:HG13	12:K:124:MET:HE3	1.92	0.51
13:L:42:PHE:HZ	13:L:103:VAL:HG23	1.74	0.51
15:S:33:LEU:CD2	15:S:62:VAL:HG21	2.40	0.51
16:2:2253:G:C6	16:2:2254:U:N3	2.78	0.51
16:2:2270:A:H2'	16:2:2271:A:H8	1.71	0.51
25:V:52:G:H1	25:V:62:C:H42	1.57	0.51
10:h:1610:G:OP2	26:v:18:C:OP2	2.28	0.51
12:K:46:ASP:OD1	12:K:51:GLU:HB2	2.11	0.51
14:X:59:SER:O	14:X:61:ASN:N	2.43	0.51
16:2:2253:G:C5	16:2:2254:U:C2	2.97	0.51
16:2:2271:A:C3'	16:2:2272:G:C5'	2.86	0.51
16:2:2289:U:H2'	16:2:2290:C:H5'	1.93	0.51
25:Y:28:G:H2'	25:Y:29:G:H8	1.75	0.51
1:a:562:G:OP1	13:L:69:LYS:CE	2.59	0.51
1:a:572:U:C4'	1:a:573:A:H5'	2.40	0.51
8:g:1149:C:O2'	8:g:1150:G:H5'	2.10	0.51
10:h:1709:A:O5'	26:y:19:U:C2'	2.58	0.51
11:T:62:PRO:O	11:T:64:VAL:N	2.42	0.51
12:K:121:ARG:C	12:K:123:GLY:H	2.18	0.51
16:2:2215:A:N3	16:2:2215:A:C2'	2.65	0.51
16:2:2255:A:C4'	16:2:2256:A:OP1	2.56	0.51
16:2:2280:A:O2'	16:2:2281:A:P	2.68	0.51
16:2:2299:A:H2'	16:2:2300:G:O5'	2.11	0.51
22:8:2963:C:H2'	22:8:2964:G:C5'	2.40	0.51
22:8:2963:C:O2'	22:8:2964:G:H5'	2.11	0.51
25:W:25:C:O2'	25:W:26:A:H5'	2.10	0.51
11:T:81:LYS:O	11:T:82:ALA:C	2.52	0.51
12:K:31:CYS:HA	12:K:43:HIS:O	2.09	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:K:46:ASP:C	12:K:48:SER:H	2.18	0.51
12:K:123:GLY:O	12:K:124:MET:O	2.29	0.51
12:K:150:ARG:O	12:K:151:LEU:CB	2.58	0.51
13:L:78:ASN:O	13:L:79:SER:HB2	2.10	0.51
15:S:47:ARG:O	15:S:51:LYS:HB2	2.10	0.51
15:S:58:TYR:HB3	15:S:88:MET:CE	2.41	0.51
16:2:2285:C:H4'	25:Y:72:C:OP1	2.09	0.51
16:2:2297:U:C2	16:2:2299:A:C6	2.99	0.51
20:6:2696:A:N7	20:6:2697:A:C6	2.79	0.51
1:a:561:A:H2'	1:a:562:G:O4'	2.11	0.51
7:f:1245:G:HO2'	7:f:1246:C:P	2.30	0.51
12:K:75:MET:HE3	12:K:121:ARG:HH12	1.75	0.51
12:K:101:GLY:CA	12:K:133:THR:HG22	2.41	0.51
16:2:2279:A:N7	16:2:2288:G:C6	2.79	0.51
23:B:59:PRO:O	23:B:60:ARG:CB	2.59	0.51
1:a:562:G:P	13:L:69:LYS:HD2	2.51	0.51
10:h:1701:C:H2'	10:h:1702:A:C5'	2.40	0.51
14:X:34:ARG:O	14:X:34:ARG:HG2	2.06	0.51
16:2:2198:A:C4	16:2:2199:G:C8	2.99	0.51
16:2:2200:U:C2	16:2:2201:G:C8	2.98	0.51
16:2:2251:G:C2	16:2:2252:A:C8	2.98	0.51
25:V:20:U:H2'	25:V:21:A:H5'	1.93	0.51
1:a:584:C:H5''	14:X:44:ARG:HH22	1.76	0.51
10:h:1661:G:H1'	10:h:1663:A:H62	1.74	0.51
10:h:1711:U:H5'	16:2:2256:A:OP2	2.09	0.51
14:X:40:LEU:HA	14:X:43:ARG:HG2	1.92	0.51
16:2:2223:A:C2'	16:2:2224:A:H8	2.20	0.51
25:W:38:A:H2'	25:W:39:U:O4'	2.10	0.51
2:b:882:G:O6	2:b:883:A:C2	2.64	0.51
10:h:1715:A:C8	10:h:1715:A:C3'	2.93	0.51
11:T:197:LYS:C	11:T:200:ARG:HG2	2.36	0.51
12:K:45:THR:HG21	12:K:49:GLY:HA2	1.92	0.51
16:2:2280:A:HO2'	16:2:2281:A:P	2.34	0.51
25:W:14:A:H1'	25:W:22:G:N2	2.26	0.51
10:h:1606:C:C6	10:h:1606:C:C3'	2.94	0.51
11:T:32:TYR:O	11:T:33:ILE:HD12	2.10	0.51
11:T:101:ARG:O	11:T:103:PRO:CD	2.58	0.51
12:K:54:VAL:CG1	12:K:81:VAL:HG13	2.32	0.51
16:2:2206:G:H2'	16:2:2207:A:H5'	1.92	0.51
16:2:2297:U:N3	16:2:2299:A:C6	2.79	0.51
21:7:2836:C:C5	21:7:2853:A:C2	2.99	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:S:91:ILE:CG2	15:S:92:PRO:HD2	2.40	0.51
11:T:13:LEU:C	11:T:15:GLY:N	2.69	0.50
12:K:53:LEU:HD13	12:K:88:LEU:HD13	1.94	0.50
16:2:2204:C:C2'	16:2:2205:U:H5''	2.41	0.50
17:3:2484:A:C6	25:W:19:G:C2'	2.93	0.50
18:4:2618:G:H4'	18:4:2619:G:OP2	2.09	0.50
21:7:2868:U:C4	21:7:2869:U:H5	2.28	0.50
25:Y:55:U:H3'	25:Y:55:U:O2	2.11	0.50
16:2:2253:G:H3'	16:2:2254:U:H6	1.74	0.50
21:7:2833:A:C2'	21:7:2834:G:H5'	2.34	0.50
21:7:2852:C:H5	21:7:2853:A:C4	2.29	0.50
25:V:15:G:O2'	25:V:16:U:H5'	2.11	0.50
25:W:65:G:H2'	25:W:66:U:C6	2.46	0.50
11:T:197:LYS:O	11:T:200:ARG:HG2	2.11	0.50
13:L:101:VAL:HG11	13:L:123:VAL:HG13	1.91	0.50
16:2:2279:A:C2	16:2:2283:G:C2	2.99	0.50
16:2:2281:A:N3	22:8:2974:U:C4'	2.74	0.50
21:7:2847:A:N3	21:7:2847:A:H2'	2.25	0.50
15:S:60:LYS:HZ3	15:S:64:LYS:CE	2.25	0.50
19:5:2655:U:O4'	19:5:2656:A:C4	2.64	0.50
21:7:2851:A:C8	21:7:2851:A:H5''	2.47	0.50
22:8:2957:G:H5'	22:8:2957:G:H8	1.76	0.50
11:T:191:GLU:O	11:T:195:VAL:HG23	2.11	0.50
12:K:142:ARG:NH2	12:K:143:GLU:CD	2.70	0.50
16:2:2285:C:H5	16:2:2286:U:C4	2.27	0.50
18:4:2618:G:H21	18:4:2618:G:P	2.35	0.50
21:7:2841:G:C5	21:7:2844:C:C4	2.99	0.50
21:7:2864:A:C6	21:7:2865:U:C4	2.99	0.50
2:b:879:G:C6	2:b:880:G:N1	2.80	0.50
2:b:879:G:C6	2:b:880:G:C6	2.99	0.50
16:2:2209:U:H6	16:2:2209:U:H3'	1.76	0.50
18:4:2627:C:OP2	18:4:2627:C:H4'	2.11	0.50
21:7:2841:G:N1	21:7:2844:C:N3	2.60	0.50
23:B:54:LYS:CA	23:B:55:LEU:CB	2.89	0.50
11:T:117:ARG:HE	11:T:189:LYS:HZ3	1.60	0.50
16:2:2249:G:H2'	16:2:2250:G:H1'	1.92	0.50
21:7:2847:A:N3	21:7:2848:G:C8	2.79	0.50
21:7:2856:G:C2	21:7:2857:C:C4	3.00	0.50
24:F:38:GLN:C	24:F:40:LYS:N	2.68	0.50
1:a:562:G:O2'	1:a:563:C:H5'	2.12	0.50
10:h:1608:C:N4	10:h:1610:G:N2	2.59	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:K:76:GLN:O	12:K:80:ASP:HB2	2.11	0.50
17:3:2484:A:C1'	25:W:19:G:C2	2.89	0.50
21:7:2827:U:O2	21:7:2827:U:C2'	2.60	0.50
21:7:2858:U:C2'	21:7:2859:U:C5	2.94	0.50
23:B:101:LYS:O	23:B:102:LYS:C	2.53	0.50
24:F:52:GLY:O	24:F:53:GLN:CB	2.60	0.50
10:h:1674:A:C2	10:h:1675:A:C8	3.00	0.50
11:T:158:PHE:C	11:T:160:SER:H	2.20	0.50
14:X:7:THR:CG2	14:X:10:LYS:HB2	2.34	0.50
15:S:42:PHE:HE2	15:S:118:GLY:HA2	1.77	0.50
21:7:2837:A:H2'	21:7:2845:A:C2	2.46	0.50
25:Y:26:A:H2'	25:Y:27:G:H5'	1.92	0.50
7:f:1252:C:H2'	7:f:1253:G:C8	2.47	0.49
8:g:1163:U:H2'	8:g:1164:C:C6	2.47	0.49
12:K:45:THR:CG2	12:K:46:ASP:N	2.74	0.49
12:K:106:LYS:NZ	12:K:136:PRO:HD2	2.27	0.49
13:L:85:PHE:HE1	13:L:87:PRO:HA	1.77	0.49
16:2:2207:A:C8	16:2:2237:C:C2	3.00	0.49
16:2:2294:U:H3'	16:2:2295:A:H5''	1.94	0.49
10:h:1711:U:O4'	16:2:2256:A:OP2	2.30	0.49
16:2:2213:A:C2'	16:2:2214:A:C8	2.81	0.49
19:5:2654:C:H5'	19:5:2657:A:C6	2.46	0.49
22:8:2976:A:H8	22:8:2976:A:O5'	1.96	0.49
10:h:1703:A:H2'	10:h:1704:C:H6	1.77	0.49
11:T:14:PHE:CD2	11:T:89:LYS:HG3	2.46	0.49
13:L:42:PHE:HB3	13:L:45:ALA:HB3	1.93	0.49
13:L:55:LYS:HD3	13:L:55:LYS:H	1.76	0.49
13:L:78:ASN:HD21	13:L:80:LYS:CG	2.25	0.49
15:S:121:LEU:HD12	15:S:121:LEU:O	2.12	0.49
16:2:2198:A:O2'	16:2:2199:G:H5'	2.13	0.49
16:2:2221:G:H8	16:2:2221:G:H5''	1.76	0.49
16:2:2249:G:N9	16:2:2272:G:C8	2.79	0.49
16:2:2264:U:O2'	16:2:2265:C:H5'	2.13	0.49
25:Y:53:G:N3	25:Y:53:G:H2'	2.27	0.49
10:h:1660:A:N6	10:h:1661:G:O6	2.45	0.49
13:L:73:VAL:HG12	13:L:74:LEU:N	2.26	0.49
16:2:2270:A:C2'	16:2:2271:A:C8	2.89	0.49
24:F:76:LYS:C	24:F:78:LYS:H	2.20	0.49
25:W:70:G:H2'	25:W:71:G:O4'	2.12	0.49
10:h:1678:U:H2'	10:h:1679:A:C8	2.48	0.49
11:T:178:ASN:HA	11:T:186:ILE:HD13	1.95	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:2:2211:U:O2'	25:W:72:C:O4'	2.30	0.49
16:2:2222:A:H8	16:2:2222:A:O5'	1.95	0.49
16:2:2276:G:N7	16:2:2277:C:C5	2.80	0.49
25:V:5:G:H2'	25:V:6:G:O4'	2.13	0.49
13:L:56:ILE:CG2	13:L:57:GLY:N	2.76	0.49
18:4:2615:G:C4	18:4:2616:C:C5	3.01	0.49
4:d:1547:G:O2'	4:d:1548:A:H5'	2.13	0.49
9:G:1434:C:O2	9:G:1434:C:H2'	2.11	0.49
15:S:60:LYS:NZ	15:S:64:LYS:HE3	2.28	0.49
16:2:2247:G:H2'	16:2:2247:G:N3	2.27	0.49
20:6:2689:A:C2	20:6:2702:A:C2	2.93	0.49
25:Y:74:C:H2'	25:Y:75:C:C5'	2.43	0.49
25:V:50:U:H2'	25:V:51:U:O4'	2.12	0.49
1:a:550:G:OP1	14:X:59:SER:HB3	2.12	0.49
1:a:576:U:H4'	1:a:577:C:OP1	2.10	0.49
2:b:879:G:N1	2:b:880:G:C2	2.80	0.49
8:g:1164:C:H5''	8:g:1165:A:H3'	1.95	0.49
10:h:1610:G:C6	10:h:1611:C:C2	3.01	0.49
11:T:22:VAL:HG12	11:T:23:LYS:H	1.76	0.49
11:T:125:THR:HB	11:T:130:ARG:CZ	2.42	0.49
20:6:2689:A:C4	20:6:2702:A:H2	2.29	0.49
25:Y:16:U:C2'	25:Y:17:C:H5'	2.43	0.49
11:T:54:LYS:HB3	11:T:57:ARG:HG2	1.95	0.49
11:T:150:ILE:O	11:T:151:LYS:C	2.55	0.49
15:S:49:ARG:NH1	15:S:87:SER:O	2.46	0.49
16:2:2206:G:H2'	16:2:2206:G:N3	2.28	0.49
16:2:2255:A:N1	16:2:2258:U:H5''	2.28	0.49
16:2:2286:U:P	25:Y:72:C:P	3.11	0.49
23:B:53:LEU:CB	23:B:54:LYS:CA	2.91	0.49
8:g:1147:U:H3	9:G:1435:G:H1	1.59	0.49
10:h:1609:C:O3'	26:v:18:C:C6	2.62	0.49
10:h:1687:C:H2'	10:h:1688:C:C6	2.48	0.49
16:2:2206:G:N2	16:2:2207:A:C8	2.80	0.49
18:4:2625:C:O2	18:4:2627:C:C5	2.66	0.49
25:V:40:C:H2'	25:V:41:C:H6	1.76	0.49
11:T:99:THR:HG21	11:T:101:ARG:HE	1.78	0.48
16:2:2252:A:N3	16:2:2253:G:C8	2.81	0.48
10:h:1608:C:N3	10:h:1610:G:N3	2.60	0.48
10:h:1688:C:O2'	10:h:1689:U:H5'	2.13	0.48
11:T:47:THR:O	11:T:48:ALA:C	2.56	0.48
13:L:52:VAL:HG22	13:L:71:VAL:HG11	1.96	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:2:2264:U:O5'	16:2:2264:U:H6	1.96	0.48
16:2:2299:A:C4	16:2:2300:G:H8	2.31	0.48
25:Y:16:U:H2'	25:Y:17:C:H5'	1.94	0.48
25:W:53:G:O2'	25:W:54:U:H5'	2.13	0.48
3:c:976:A:N6	3:c:982:U:OP2	2.47	0.48
10:h:1613:C:H2'	10:h:1614:G:C8	2.48	0.48
13:L:26:ASP:HB3	13:L:29:PHE:HB3	1.95	0.48
13:L:87:PRO:O	13:L:88:MET:O	2.31	0.48
15:S:13:PHE:O	15:S:14:ARG:HD3	2.13	0.48
18:4:2614:G:C6	18:4:2615:G:C5	3.01	0.48
22:8:2964:G:C5	22:8:2966:G:OP2	2.66	0.48
25:V:28:G:O2'	25:V:29:G:H5'	2.14	0.48
8:g:1158:U:O4	8:g:1159:U:C4	2.66	0.48
11:T:19:TYR:CE2	11:T:41:GLN:HA	2.48	0.48
16:2:2211:U:O2	25:W:72:C:C4'	2.60	0.48
16:2:2276:G:O6	16:2:2277:C:C4	2.67	0.48
23:B:154:THR:HA	23:B:155:ILE:C	2.38	0.48
25:W:63:G:H2'	25:W:64:A:C8	2.48	0.48
10:h:1711:U:O2'	10:h:1712:C:H5'	2.13	0.48
13:L:29:PHE:O	13:L:33:LEU:HB2	2.14	0.48
14:X:7:THR:HG22	14:X:8:LEU:H	1.78	0.48
14:X:62:TRP:CH2	14:X:63:HIS:NE2	2.81	0.48
15:S:60:LYS:HD3	15:S:64:LYS:HE3	1.94	0.48
18:4:2618:G:C1'	21:7:2865:U:OP1	2.59	0.48
21:7:2860:U:O2	21:7:2860:U:H2'	2.13	0.48
4:d:1544:G:O5'	4:d:1544:G:H8	1.97	0.48
16:2:2211:U:O2'	25:W:72:C:H4'	2.14	0.48
16:2:2279:A:C8	16:2:2288:G:C5	3.01	0.48
18:4:2616:C:H2'	18:4:2617:U:C5'	2.31	0.48
21:7:2847:A:C2	21:7:2848:G:C1'	2.96	0.48
25:W:52:G:H2'	25:W:53:G:C8	2.49	0.48
2:b:888:C:H6	2:b:888:C:H5'	1.78	0.48
3:c:987:U:C4	3:c:988:C:C5	3.02	0.48
10:h:1610:G:P	26:v:18:C:C6	3.07	0.48
10:h:1621:G:C4	10:h:1705:A:C2	3.01	0.48
11:T:71:LEU:HG	11:T:84:CYS:SG	2.54	0.48
11:T:105:GLU:O	11:T:105:GLU:HG2	2.12	0.48
11:T:152:GLY:HA2	11:T:184:TRP:HD1	1.78	0.48
13:L:29:PHE:CZ	13:L:33:LEU:HD12	2.49	0.48
13:L:61:LYS:HD2	13:L:117:PRO:HA	1.96	0.48
14:X:66:LYS:O	14:X:70:MET:HB2	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:2:2279:A:H4'	16:2:2280:A:C5'	2.40	0.48
21:7:2857:C:H2'	21:7:2858:U:C6	2.45	0.48
26:y:19:U:C5'	26:v:18:C:H5''	2.44	0.48
10:h:1644:C:H2'	10:h:1645:C:C6	2.48	0.48
11:T:59:THR:C	11:T:61:CYS:H	2.22	0.48
16:2:2254:U:C2'	16:2:2255:A:OP1	2.62	0.48
20:6:2693:C:H5''	20:6:2694:A:H5'	1.95	0.48
20:6:2702:A:O4'	20:6:2704:A:C5	2.67	0.48
25:W:6:G:H2'	25:W:7:A:H5'	1.93	0.48
25:W:76:A:H3'	25:W:76:A:OP2	2.14	0.48
12:K:88:LEU:O	12:K:89:LYS:HB2	2.13	0.48
11:T:63:ILE:O	11:T:66:ARG:HB2	2.14	0.48
11:T:193:GLU:O	11:T:197:LYS:N	2.46	0.48
16:2:2248:C:C4'	16:2:2271:A:C2	2.96	0.48
21:7:2834:G:C6	21:7:2855:U:C4	3.01	0.48
21:7:2841:G:C5	21:7:2844:C:N4	2.82	0.48
21:7:2869:U:O2	21:7:2869:U:C2'	2.62	0.48
25:W:21:A:H5'	25:W:21:A:C8	2.49	0.48
2:b:884:A:H2'	2:b:885:A:O4'	2.14	0.47
10:h:1609:C:N1	25:V:34:G:O6	2.18	0.47
10:h:1611:C:C2'	10:h:1612:C:H5'	2.43	0.47
10:h:1657:G:H2'	10:h:1658:A:H8	1.78	0.47
16:2:2253:G:C4	16:2:2254:U:C6	3.01	0.47
16:2:2254:U:O2'	16:2:2255:A:P	2.71	0.47
16:2:2261:G:N2	16:2:2263:C:C2	2.82	0.47
20:6:2701:U:O4'	20:6:2705:A:C6	2.67	0.47
22:8:2957:G:C6	22:8:2976:A:C2	3.02	0.47
25:V:40:C:H5''	25:W:36:A:P	2.51	0.47
1:a:562:G:OP1	13:L:69:LYS:HD2	2.13	0.47
6:E:1590:C:O2'	6:E:1591:C:OP2	2.28	0.47
12:K:29:GLY:O	12:K:93:LEU:HA	2.14	0.47
21:7:2826:U:N3	21:7:2827:U:C5	2.81	0.47
21:7:2850:G:H5''	21:7:2850:G:H8	1.79	0.47
23:B:55:LEU:HA	23:B:167:VAL:HA	1.96	0.47
1:a:577:C:H2'	1:a:577:C:O2	2.14	0.47
16:2:2290:C:C4	16:2:2303:A:N1	2.82	0.47
25:Y:38:A:H2'	25:Y:39:U:O4'	2.14	0.47
25:Y:51:U:H3'	25:Y:52:G:H8	1.79	0.47
10:h:1654:U:O2	10:h:1672:G:C2	2.68	0.47
11:T:102:ASN:O	11:T:105:GLU:N	2.46	0.47
12:K:99:ALA:O	12:K:101:GLY:N	2.48	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:K:103:VAL:HG12	12:K:142:ARG:CD	2.45	0.47
13:L:75:LEU:HD12	13:L:78:ASN:HD22	1.80	0.47
14:X:47:PRO:O	14:X:51:ALA:HB2	2.14	0.47
14:X:65:GLY:HA2	14:X:70:MET:HE1	1.95	0.47
15:S:90:VAL:HB	15:S:117:ILE:HA	1.96	0.47
18:4:2618:G:N9	21:7:2865:U:P	2.87	0.47
25:W:59:U:H2'	25:W:60:U:H5'	1.96	0.47
2:b:877:G:N2	2:b:878:A:N3	2.62	0.47
8:g:1155:A:N6	8:g:1156:A:N6	2.63	0.47
10:h:1609:C:C2	26:v:18:C:N3	2.81	0.47
16:2:2194:G:H2'	16:2:2195:C:H5'	1.97	0.47
16:2:2249:G:C2	16:2:2272:G:O6	2.68	0.47
22:8:2960:C:C3'	22:8:2961:G:H8	2.27	0.47
22:8:2966:G:C6	22:8:2967:A:C6	3.03	0.47
24:F:40:LYS:O	24:F:41:ARG:C	2.57	0.47
11:T:107:PHE:HA	11:T:172:ILE:HG23	1.96	0.47
13:L:55:LYS:HB2	13:L:92:LEU:HD21	1.97	0.47
16:2:2211:U:O2	25:W:72:C:H5''	2.14	0.47
16:2:2215:A:C2	16:2:2216:G:N9	2.83	0.47
22:8:2969:A:C6	22:8:2970:C:N4	2.83	0.47
25:Y:42:C:C3'	25:Y:43:C:H5''	2.45	0.47
1:a:570:G:C4'	1:a:574:A:N3	2.73	0.47
10:h:1608:C:C2	10:h:1610:G:C4	3.03	0.47
11:T:45:PRO:HB2	11:T:68:ILE:HD12	1.97	0.47
11:T:104:LEU:C	11:T:106:VAL:H	2.21	0.47
11:T:152:GLY:O	11:T:155:GLU:N	2.48	0.47
11:T:158:PHE:O	11:T:160:SER:N	2.36	0.47
12:K:69:SER:O	12:K:70:SER:O	2.33	0.47
12:K:78:ALA:HB1	12:K:119:LEU:CD1	2.45	0.47
12:K:104:GLU:O	12:K:105:THR:C	2.58	0.47
13:L:69:LYS:HZ2	13:L:92:LEU:HD23	1.80	0.47
13:L:131:LEU:O	13:L:134:LEU:N	2.47	0.47
15:S:42:PHE:CD2	15:S:46:THR:HG21	2.49	0.47
15:S:85:TYR:H	15:S:85:TYR:HD1	1.63	0.47
16:2:2248:C:H4'	16:2:2271:A:C2	2.49	0.47
21:7:2852:C:C5	21:7:2853:A:C5	3.03	0.47
25:Y:11:C:H2'	25:Y:12:U:C6	2.49	0.47
25:V:74:C:C2'	25:V:75:C:H5'	2.44	0.47
25:W:31:A:N1	25:W:39:U:O4	2.47	0.47
8:g:1165:A:HO2'	8:g:1166:A:P	2.37	0.47
12:K:67:GLU:C	12:K:69:SER:H	2.23	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:L:75:LEU:HD11	13:L:82:ILE:CD1	2.45	0.47
17:3:2484:A:C4	25:W:19:G:C2	2.47	0.47
18:4:2618:G:C1'	21:7:2865:U:P	2.99	0.47
21:7:2825:C:H42	21:7:2864:A:N6	2.12	0.47
1:a:572:U:O5'	1:a:572:U:H6	1.98	0.47
8:g:1153:U:C2	8:g:1157:U:H5	2.33	0.47
11:T:16:LYS:HE3	11:T:93:GLU:OE1	2.15	0.47
11:T:122:ARG:NH2	25:W:32:U:O2'	2.47	0.47
13:L:69:LYS:NZ	13:L:92:LEU:HB3	2.29	0.47
16:2:2259:A:H5''	16:2:2260:U:C5	2.47	0.47
16:2:2295:A:C6	16:2:2296:A:C6	3.02	0.47
19:5:2655:U:H1'	19:5:2656:A:C2	2.49	0.47
25:V:20:U:H4'	25:V:20:U:OP1	2.15	0.47
11:T:13:LEU:HB2	11:T:19:TYR:HE1	1.80	0.47
11:T:174:ASN:CB	11:T:182:SER:O	2.63	0.47
13:L:3:VAL:HG12	13:L:4:GLY:N	2.24	0.47
15:S:58:TYR:HB3	15:S:88:MET:HE2	1.97	0.47
15:S:114:PHE:O	15:S:117:ILE:HG13	2.15	0.47
16:2:2196:C:N4	16:2:2242:A:C8	2.83	0.47
16:2:2197:C:H4'	16:2:2198:A:H8	1.79	0.47
16:2:2199:G:C4	16:2:2200:U:C5	3.03	0.47
16:2:2223:A:N3	16:2:2224:A:C4	2.83	0.47
19:5:2656:A:C4'	19:5:2657:A:OP1	2.45	0.47
20:6:2706:G:C4	20:6:2707:C:C5	3.03	0.47
21:7:2850:G:OP1	21:7:2850:G:C4'	2.53	0.47
25:W:39:U:C5'	25:W:39:U:O2	2.63	0.47
14:X:8:LEU:C	14:X:10:LYS:N	2.73	0.46
14:X:33:GLY:C	14:X:35:SER:N	2.72	0.46
14:X:62:TRP:C	14:X:64:ALA:H	2.23	0.46
25:Y:15:G:O2'	25:Y:16:U:OP1	2.25	0.46
1:a:559:C:O2	1:a:559:C:C2'	2.63	0.46
16:2:2250:G:H2'	16:2:2250:G:N3	2.31	0.46
16:2:2290:C:N3	16:2:2303:A:C2	2.84	0.46
18:4:2617:U:O2	18:4:2617:U:O4'	2.32	0.46
24:F:33:ALA:O	24:F:34:SER:C	2.58	0.46
1:a:570:G:O4'	1:a:574:A:H2	1.99	0.46
3:c:981:A:O2'	3:c:982:U:OP2	2.29	0.46
10:h:1687:C:H2'	10:h:1688:C:H6	1.79	0.46
13:L:21:ASP:C	13:L:23:ARG:H	2.23	0.46
14:X:34:ARG:HB3	14:X:34:ARG:CZ	2.44	0.46
16:2:2205:U:C2'	16:2:2206:G:H5'	2.37	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:2:2215:A:C2	16:2:2216:G:C1'	2.98	0.46
16:2:2294:U:C2	16:2:2296:A:OP2	2.69	0.46
18:4:2618:G:C1'	21:7:2865:U:O5'	2.63	0.46
13:L:106:LEU:CD2	13:L:122:LYS:HB2	2.45	0.46
15:S:42:PHE:CD2	15:S:118:GLY:HA2	2.49	0.46
16:2:2225:U:C2	16:2:2226:U:C5	3.03	0.46
16:2:2249:G:C1'	16:2:2272:G:C8	2.97	0.46
16:2:2288:G:C2	16:2:2289:U:C5	3.03	0.46
16:2:2289:U:N3	16:2:2290:C:C5	2.84	0.46
19:5:2655:U:C1'	19:5:2656:A:C2	2.98	0.46
6:E:1590:C:O2'	6:E:1591:C:P	2.74	0.46
10:h:1663:A:H2'	10:h:1664:A:H8	1.81	0.46
15:S:32:LYS:O	15:S:34:ILE:N	2.48	0.46
15:S:69:LYS:NZ	15:S:93:GLU:O	2.45	0.46
15:S:86:ARG:CZ	15:S:122:ALA:HB1	2.45	0.46
16:2:2234:G:C5	16:2:2235:C:C5	3.04	0.46
16:2:2252:A:C4	16:2:2253:G:N7	2.84	0.46
16:2:2266:U:H5''	16:2:2267:C:OP2	2.15	0.46
16:2:2291:A:N1	16:2:2292:U:C2	2.84	0.46
22:8:2960:C:H2'	22:8:2961:G:C8	2.50	0.46
23:B:128:LEU:O	23:B:131:ALA:HB3	2.15	0.46
25:V:16:U:H4'	25:V:17:C:H5	1.81	0.46
8:g:1159:U:H3	8:g:1170:G:H1	1.63	0.46
15:S:60:LYS:HZ3	15:S:64:LYS:HE3	1.81	0.46
25:V:66:U:O2'	25:V:67:C:H5'	2.16	0.46
2:b:880:G:C5	2:b:881:U:H5	2.33	0.46
2:b:882:G:N7	2:b:883:A:C5	2.83	0.46
11:T:41:GLN:H	11:T:41:GLN:HG2	1.56	0.46
11:T:57:ARG:O	11:T:59:THR:N	2.48	0.46
14:X:44:ARG:HA	14:X:49:ILE:HD12	1.98	0.46
16:2:2289:U:C2'	16:2:2290:C:C5'	2.94	0.46
18:4:2619:G:O4'	21:7:2866:U:C6	2.69	0.46
20:6:2698:G:H2'	20:6:2699:G:C5'	2.28	0.46
21:7:2834:G:O2'	21:7:2835:U:O4'	2.34	0.46
1:a:580:G:H2'	1:a:581:C:C6	2.50	0.46
3:c:981:A:C2'	3:c:982:U:OP2	2.63	0.46
8:g:1170:G:H3'	8:g:1171:G:H5'	1.98	0.46
9:G:1440:A:H2'	9:G:1441:C:C6	2.50	0.46
11:T:13:LEU:C	11:T:15:GLY:H	2.22	0.46
11:T:70:THR:HB	11:T:150:ILE:CD1	2.44	0.46
11:T:153:CYS:HA	11:T:167:THR:CG2	2.46	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:K:110:PRO:HG2	12:K:111:GLY:N	2.31	0.46
16:2:2257:C:C2	16:2:2258:U:C6	3.04	0.46
25:Y:52:G:N3	25:Y:52:G:H2'	2.30	0.46
25:W:4:C:C2	25:W:70:G:C2	3.04	0.46
11:T:95:ILE:HG21	11:T:103:PRO:CB	2.43	0.46
12:K:36:THR:HG22	12:K:38:ASN:H	1.81	0.46
13:L:81:LYS:HD2	13:L:81:LYS:C	2.41	0.46
14:X:35:SER:O	14:X:39:ILE:HG13	2.15	0.46
16:2:2211:U:HO2'	16:2:2212:C:H5'	1.79	0.46
16:2:2240:G:C2'	16:2:2241:U:O5'	2.64	0.46
16:2:2285:C:H5	16:2:2286:U:C6	2.33	0.46
9:G:1435:G:C2	9:G:1436:C:C5	3.03	0.46
13:L:113:VAL:O	13:L:114:GLY:C	2.58	0.46
16:2:2250:G:N2	16:2:2267:C:C2	2.60	0.46
25:Y:36:A:C2	26:y:20:U:N3	2.84	0.46
1:a:583:C:O2'	14:X:61:ASN:HB2	2.17	0.45
10:h:1703:A:H2'	10:h:1704:C:C6	2.51	0.45
13:L:27:ASN:O	13:L:31:LYS:HB2	2.16	0.45
15:S:66:ARG:HH12	15:S:93:GLU:HB3	1.77	0.45
16:2:2235:C:H2'	16:2:2236:G:C8	2.52	0.45
16:2:2300:G:C2	16:2:2301:U:C2	3.04	0.45
21:7:2839:G:C6	21:7:2850:G:C2	3.04	0.45
25:V:16:U:H4'	25:V:17:C:C5	2.50	0.45
25:W:59:U:C2'	25:W:60:U:H5'	2.46	0.45
21:7:2852:C:N4	21:7:2853:A:N3	2.64	0.45
25:V:37:A:H3'	25:V:38:A:H8	1.80	0.45
25:V:41:C:H2'	25:V:42:C:H5''	1.99	0.45
12:K:54:VAL:HG22	12:K:84:ARG:HG3	1.98	0.45
16:2:2222:A:O2'	16:2:2223:A:H5''	2.17	0.45
16:2:2249:G:H2'	16:2:2250:G:O4'	2.16	0.45
21:7:2840:C:N4	21:7:2841:G:C5	2.85	0.45
25:W:18:G:O2'	25:W:57:G:O6	2.30	0.45
1:a:542:G:O2'	1:a:543:A:H5'	2.16	0.45
10:h:1651:G:H1'	10:h:1674:A:H62	1.81	0.45
10:h:1667:U:H2'	10:h:1668:U:O4'	2.17	0.45
15:S:73:PRO:C	15:S:75:GLY:N	2.66	0.45
16:2:2301:U:O2'	16:2:2302:G:H5'	2.16	0.45
20:6:2690:G:OP2	20:6:2691:A:N7	2.50	0.45
8:g:1162:C:C2'	8:g:1163:U:OP2	2.64	0.45
10:h:1626:A:C2	10:h:1627:A:C4	3.04	0.45
13:L:78:ASN:ND2	13:L:80:LYS:HG2	2.32	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:L:129:ILE:O	13:L:130:SER:C	2.59	0.45
16:2:2295:A:H2'	16:2:2296:A:C8	2.52	0.45
25:W:38:A:H2'	25:W:39:U:C5'	2.47	0.45
11:T:14:PHE:HD2	11:T:17:TRP:HZ3	1.63	0.45
11:T:19:TYR:C	11:T:21:GLU:H	2.24	0.45
11:T:52:GLN:HE22	11:T:58:LYS:HE2	1.82	0.45
11:T:152:GLY:O	11:T:153:CYS:C	2.59	0.45
12:K:27:VAL:HG11	12:K:90:ILE:HG12	1.98	0.45
22:8:2965:U:H2'	22:8:2966:G:H5'	1.99	0.45
23:B:68:PHE:HA	23:B:73:ASP:CB	2.46	0.45
25:Y:8:U:C4'	25:Y:48:C:H4'	2.44	0.45
10:h:1659:C:O5'	10:h:1659:C:H6	1.99	0.45
11:T:62:PRO:O	11:T:63:ILE:C	2.59	0.45
12:K:30:VAL:O	12:K:30:VAL:HG12	2.14	0.45
16:2:2232:A:C8	16:2:2232:A:H3'	2.51	0.45
18:4:2615:G:H2'	18:4:2616:C:C6	2.52	0.45
21:7:2836:C:H5	21:7:2853:A:C2	2.33	0.45
24:F:23:HIS:HA	24:F:74:CYS:HA	1.98	0.45
24:F:92:GLU:O	24:F:93:LEU:O	2.35	0.45
25:Y:74:C:C2'	25:Y:75:C:H5'	2.47	0.45
25:W:23:A:H2'	25:W:24:G:C8	2.52	0.45
4:d:1543:C:O2'	9:G:1441:C:O2	2.35	0.45
13:L:40:ASN:CB	13:L:41:PRO:HD2	2.37	0.45
13:L:52:VAL:HB	13:L:98:ASN:H	1.82	0.45
16:2:2302:G:H2'	16:2:2303:A:O4'	2.16	0.45
23:B:188:ASN:C	23:B:190:PHE:N	2.72	0.45
7:f:1252:C:O2'	7:f:1253:G:H5'	2.17	0.45
8:g:1163:U:N3	25:V:34:G:O4'	2.50	0.45
10:h:1608:C:C4	10:h:1610:G:C2	3.05	0.45
11:T:146:ILE:O	11:T:147:TYR:C	2.60	0.45
12:K:21:VAL:CG1	12:K:22:GLY:H	2.08	0.45
13:L:75:LEU:CD2	13:L:82:ILE:HD12	2.42	0.45
14:X:25:ASP:O	14:X:26:LYS:C	2.59	0.45
16:2:2211:U:O2'	16:2:2212:C:H5'	2.17	0.45
16:2:2270:A:C6	16:2:2271:A:C6	3.05	0.45
16:2:2297:U:O2	16:2:2299:A:C5	2.70	0.45
22:8:2959:C:H5'	22:8:2960:C:OP2	2.16	0.45
11:T:34:ALA:HB1	11:T:62:PRO:HA	1.97	0.45
12:K:104:GLU:O	12:K:106:LYS:N	2.50	0.45
12:K:148:GLY:O	12:K:149:ARG:C	2.60	0.45
13:L:75:LEU:HD11	13:L:82:ILE:HD11	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:2:2247:G:H5''	16:2:2248:C:OP2	2.17	0.45
16:2:2279:A:C3'	16:2:2280:A:C5'	2.94	0.45
20:6:2694:A:C5'	20:6:2695:A:OP2	2.65	0.45
10:h:1617:G:N2	10:h:1710:G:C4	2.85	0.44
11:T:171:GLU:OE2	11:T:183:SER:HB2	2.17	0.44
14:X:8:LEU:C	14:X:10:LYS:H	2.25	0.44
15:S:85:TYR:N	15:S:85:TYR:CD1	2.85	0.44
16:2:2209:U:OP2	16:2:2209:U:N1	2.50	0.44
16:2:2240:G:H2'	16:2:2241:U:C5'	2.47	0.44
16:2:2255:A:H4'	16:2:2256:A:OP1	2.12	0.44
16:2:2278:C:O2'	16:2:2279:A:H5''	2.17	0.44
16:2:2298:U:O4'	16:2:2298:U:O2	2.35	0.44
21:7:2847:A:C5'	21:7:2848:G:OP2	2.55	0.44
1:a:546:G:H2'	1:a:547:C:H5	1.76	0.44
11:T:34:ALA:H	11:T:63:ILE:HG12	1.82	0.44
12:K:47:LEU:N	12:K:47:LEU:CD2	2.71	0.44
13:L:21:ASP:C	13:L:23:ARG:N	2.74	0.44
16:2:2250:G:N1	16:2:2267:C:N3	2.54	0.44
16:2:2285:C:H41	16:2:2286:U:H3	1.65	0.44
16:2:2300:G:N3	16:2:2300:G:H2'	2.32	0.44
24:F:45:ARG:O	24:F:46:LYS:CB	2.65	0.44
8:g:1155:A:H5'	8:g:1156:A:OP2	2.17	0.44
8:g:1166:A:H2'	8:g:1167:C:C5'	2.47	0.44
9:G:1430:C:O2'	9:G:1431:A:P	2.75	0.44
10:h:1621:G:N3	10:h:1705:A:C2	2.85	0.44
10:h:1652:A:C2	10:h:1674:A:N3	2.86	0.44
10:h:1686:C:O2'	10:h:1687:C:H5'	2.17	0.44
10:h:1699:A:O2'	10:h:1700:A:H8	2.01	0.44
11:T:96:HIS:O	11:T:100:GLY:CA	2.65	0.44
11:T:184:TRP:CD1	11:T:188:LYS:HZ1	2.35	0.44
12:K:43:HIS:CE1	12:K:52:THR:HG23	2.51	0.44
12:K:95:ILE:HD11	12:K:126:ILE:HG23	1.99	0.44
12:K:138:ASP:CG	12:K:139:SER:H	2.17	0.44
16:2:2252:A:N1	16:2:2253:G:C6	2.86	0.44
16:2:2253:G:C3'	16:2:2254:U:C6	2.97	0.44
21:7:2872:A:O4'	21:7:2873:U:OP1	2.35	0.44
3:c:976:A:C2'	3:c:977:U:H5'	2.48	0.44
13:L:139:LYS:O	13:L:140:GLU:HG3	2.18	0.44
15:S:98:ILE:HG22	15:S:111:GLU:HG2	1.99	0.44
16:2:2287:C:O2	16:2:2298:U:H5''	2.17	0.44
16:2:2289:U:O2'	16:2:2290:C:C5'	2.62	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:6:2703:A:C4'	20:6:2704:A:H5'	2.47	0.44
21:7:2835:U:O2	21:7:2835:U:H2'	2.17	0.44
23:B:32:VAL:HA	23:B:208:SER:HA	2.00	0.44
24:F:47:GLN:H	24:F:48:SER:CA	2.26	0.44
25:Y:33:U:H3'	25:Y:34:G:H5''	1.99	0.44
25:W:16:U:N3	25:W:19:G:OP2	2.51	0.44
1:a:580:G:H2'	1:a:581:C:O4'	2.18	0.44
6:E:1591:C:O2	6:E:1591:C:C2'	2.64	0.44
11:T:83:LEU:O	11:T:86:LYS:N	2.49	0.44
12:K:91:ASN:HD22	12:K:125:LYS:HD2	1.83	0.44
21:7:2836:C:O2	21:7:2836:C:C3'	2.65	0.44
14:X:7:THR:O	14:X:8:LEU:HB2	2.17	0.44
16:2:2197:C:N3	16:2:2242:A:C2	2.86	0.44
16:2:2279:A:C2	16:2:2283:G:N2	2.86	0.44
22:8:2966:G:H8	22:8:2966:G:O5'	2.00	0.44
23:B:68:PHE:H	23:B:112:ALA:HB2	1.83	0.44
25:W:19:G:H5'	25:W:20:U:H5	1.82	0.44
8:g:1154:U:C3'	8:g:1155:A:H5''	2.47	0.44
8:g:1166:A:C2'	8:g:1167:C:H5'	2.48	0.44
10:h:1609:C:H3'	10:h:1610:G:H5'	1.99	0.44
12:K:61:LYS:HE3	12:K:76:GLN:HB3	1.99	0.44
12:K:150:ARG:O	12:K:151:LEU:HB3	2.17	0.44
15:S:46:THR:HG23	15:S:89:ILE:HD12	1.99	0.44
15:S:55:SER:HB2	15:S:58:TYR:CE2	2.52	0.44
16:2:2246:G:C4	16:2:2247:G:C8	3.06	0.44
8:g:1144:A:H2'	8:g:1145:C:C6	2.53	0.44
11:T:153:CYS:HA	11:T:167:THR:HG22	1.99	0.44
15:S:38:LEU:HA	15:S:41:LEU:HD12	2.00	0.44
15:S:46:THR:CG2	15:S:89:ILE:HD13	2.42	0.44
16:2:2217:U:H2'	16:2:2218:G:H8	1.83	0.44
16:2:2276:G:C4	16:2:2277:C:C5	3.05	0.44
16:2:2279:A:C4'	16:2:2280:A:H5'	2.43	0.44
25:W:12:U:H2'	25:W:13:C:O4'	2.18	0.44
2:b:882:G:O6	2:b:883:A:N1	2.51	0.44
2:b:883:A:C8	2:b:883:A:H3'	2.53	0.44
10:h:1644:C:O5'	10:h:1644:C:H6	2.01	0.44
17:3:2484:A:N7	25:W:19:G:O2'	2.43	0.44
18:4:2620:G:H2'	18:4:2621:G:H5'	2.00	0.44
21:7:2834:G:C4	21:7:2835:U:H5	2.35	0.44
21:7:2868:U:C4	21:7:2869:U:C5	3.06	0.44
22:8:2961:G:H5''	22:8:2962:U:P	2.58	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:Y:27:G:N2	25:Y:43:C:H5	2.10	0.44
25:V:68:C:H2'	25:V:69:G:O4'	2.17	0.44
11:T:57:ARG:O	11:T:58:LYS:C	2.62	0.43
16:2:2223:A:C2	16:2:2224:A:C5	3.06	0.43
16:2:2278:C:C4	16:2:2305:G:C5	3.06	0.43
11:T:24:ILE:CD1	11:T:30:GLN:HA	2.48	0.43
11:T:47:THR:O	11:T:47:THR:HG22	2.17	0.43
13:L:135:PHE:CE1	14:X:15:ARG:NH1	2.86	0.43
15:S:74:ALA:C	15:S:76:GLU:N	2.75	0.43
16:2:2255:A:C2	16:2:2258:U:OP2	2.71	0.43
16:2:2281:A:H1'	16:2:2282:U:OP1	2.18	0.43
19:5:2657:A:H1'	20:6:2694:A:C8	2.53	0.43
21:7:2831:G:C2	21:7:2832:C:C2	3.06	0.43
22:8:2968:G:O2'	22:8:2969:A:H5'	2.18	0.43
25:V:22:G:H2'	25:V:23:A:C8	2.54	0.43
8:g:1157:U:H4'	8:g:1158:U:OP2	2.17	0.43
9:G:1430:C:O2'	9:G:1431:A:O5'	2.36	0.43
10:h:1618:C:H2'	10:h:1619:U:C6	2.53	0.43
13:L:90:GLY:O	13:L:91:CYS:C	2.61	0.43
14:X:13:LYS:O	14:X:17:GLN:HG3	2.18	0.43
14:X:52:THR:O	14:X:53:ASP:C	2.61	0.43
16:2:2218:G:C4	16:2:2228:A:C2	3.06	0.43
17:3:2488:A:H2'	17:3:2488:A:N3	2.33	0.43
1:a:554:U:H2'	1:a:555:G:H8	1.83	0.43
3:c:977:U:H2'	3:c:978:C:O4'	2.19	0.43
3:c:981:A:O2'	3:c:982:U:C5'	2.67	0.43
3:c:986:G:O2'	3:c:987:U:H5'	2.18	0.43
12:K:43:HIS:HE1	12:K:52:THR:CG2	2.30	0.43
16:2:2197:C:C2	16:2:2242:A:C2	3.06	0.43
16:2:2256:A:O2'	16:2:2257:C:OP2	2.36	0.43
19:5:2655:U:O4'	19:5:2656:A:N3	2.51	0.43
2:b:880:G:C4	2:b:881:U:H5	2.36	0.43
8:g:1155:A:C6	8:g:1156:A:C6	3.07	0.43
10:h:1644:C:H2'	10:h:1645:C:H6	1.84	0.43
10:h:1692:C:O2'	10:h:1693:A:H5'	2.18	0.43
13:L:124:VAL:O	13:L:124:VAL:HG12	2.18	0.43
15:S:101:VAL:HG21	15:S:121:LEU:CD1	2.40	0.43
16:2:2199:G:C2	16:2:2200:U:C6	3.06	0.43
16:2:2257:C:O2	16:2:2257:C:O4'	2.36	0.43
21:7:2846:U:H2'	21:7:2847:A:OP2	2.19	0.43
21:7:2857:C:C2	21:7:2858:U:C6	3.06	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:Y:45:U:O5'	25:Y:45:U:H6	2.02	0.43
7:f:1241:U:HO2'	7:f:1242:G:P	2.35	0.43
11:T:62:PRO:O	11:T:65:GLU:N	2.44	0.43
14:X:31:PRO:CG	14:X:39:ILE:HD11	2.46	0.43
16:2:2210:G:H21	25:W:72:C:P	2.22	0.43
16:2:2224:A:H3'	16:2:2225:U:H6	1.83	0.43
16:2:2249:G:N9	16:2:2272:G:N7	2.66	0.43
16:2:2260:U:O2'	16:2:2261:G:C8	2.71	0.43
21:7:2830:G:H2'	21:7:2831:G:O5'	2.18	0.43
25:V:42:C:H6	25:V:42:C:H5'	1.84	0.43
25:V:72:C:C2'	25:V:73:A:H5'	2.47	0.43
3:c:986:G:C2	3:c:987:U:C5	3.07	0.43
13:L:106:LEU:O	13:L:107:GLY:C	2.61	0.43
15:S:35:SER:C	15:S:37:GLU:N	2.77	0.43
15:S:76:GLU:C	15:S:78:PRO:HD3	2.44	0.43
16:2:2218:G:C5	16:2:2228:A:C2	3.06	0.43
16:2:2249:G:C4	16:2:2272:G:C5	3.07	0.43
16:2:2254:U:H2'	16:2:2255:A:OP1	2.18	0.43
21:7:2864:A:C5	21:7:2865:U:C4	3.07	0.43
1:a:561:A:C1'	14:X:14:VAL:HG23	2.33	0.43
1:a:579:A:O2'	1:a:580:G:H5'	2.19	0.43
7:f:1253:G:O2'	7:f:1254:U:H5'	2.19	0.43
11:T:148:PHE:O	11:T:151:LYS:HB3	2.19	0.43
12:K:110:PRO:HG2	12:K:111:GLY:H	1.83	0.43
15:S:60:LYS:NZ	15:S:64:LYS:CE	2.82	0.43
15:S:95:VAL:CG2	15:S:117:ILE:HD11	2.46	0.43
16:2:2201:G:C4	16:2:2202:C:C6	3.07	0.43
21:7:2837:A:H2'	21:7:2845:A:C6	2.52	0.43
25:W:62:C:H2'	25:W:63:G:C8	2.53	0.43
1:a:570:G:O6	13:L:64:ASN:HA	2.19	0.43
8:g:1163:U:C2	25:V:34:G:O4'	2.72	0.43
11:T:68:ILE:CG2	11:T:84:CYS:HB3	2.49	0.43
11:T:182:SER:O	11:T:183:SER:HB3	2.19	0.43
12:K:53:LEU:HB2	12:K:54:VAL:H	1.70	0.43
15:S:101:VAL:O	15:S:108:VAL:N	2.46	0.43
16:2:2209:U:C6	16:2:2209:U:C3'	3.01	0.43
20:6:2702:A:C8	20:6:2704:A:C5	3.03	0.43
21:7:2828:G:H2'	21:7:2829:U:O5'	2.18	0.43
21:7:2830:G:H2'	21:7:2831:G:H8	1.83	0.43
21:7:2834:G:C2	21:7:2835:U:C6	3.07	0.43
23:B:101:LYS:O	23:B:104:SER:N	2.52	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:B:167:VAL:O	23:B:168:ALA:HB2	2.19	0.43
25:V:6:G:C6	25:V:7:A:C6	3.07	0.43
25:W:58:A:H1'	25:W:60:U:H5	1.80	0.43
11:T:154:ARG:HA	11:T:157:ALA:HB3	2.00	0.43
11:T:183:SER:O	11:T:184:TRP:C	2.60	0.43
12:K:22:GLY:C	12:K:24:ASN:N	2.76	0.43
13:L:42:PHE:O	13:L:43:MET:HB2	2.19	0.43
14:X:7:THR:CG2	14:X:8:LEU:H	2.28	0.43
16:2:2218:G:C6	16:2:2228:A:C6	3.07	0.43
16:2:2255:A:N1	16:2:2260:U:O4	2.52	0.43
20:6:2689:A:H1'	20:6:2702:A:H61	1.84	0.43
21:7:2856:G:N2	21:7:2857:C:C2	2.87	0.43
25:V:28:G:H1	25:V:42:C:H42	1.67	0.43
25:W:28:G:H2'	25:W:29:G:H8	1.84	0.43
3:c:972:G:C2'	3:c:973:A:H5'	2.48	0.42
11:T:26:ASP:OD1	11:T:28:CYS:N	2.52	0.42
12:K:17:GLY:O	12:K:18:PRO:C	2.62	0.42
16:2:2252:A:C2	16:2:2265:C:O2	2.72	0.42
16:2:2284:C:C5'	16:2:2285:C:OP2	2.66	0.42
16:2:2285:C:C3'	16:2:2285:C:C6	3.02	0.42
22:8:2960:C:H3'	22:8:2961:G:C8	2.49	0.42
23:B:126:PRO:O	23:B:128:LEU:N	2.52	0.42
25:V:67:C:H2'	25:V:68:C:H6	1.81	0.42
1:a:572:U:H4'	1:a:573:A:H5''	1.93	0.42
3:c:972:G:O2'	3:c:973:A:H5'	2.20	0.42
11:T:147:TYR:C	11:T:147:TYR:CD1	2.94	0.42
11:T:175:ALA:O	11:T:186:ILE:HD11	2.19	0.42
12:K:110:PRO:CG	12:K:111:GLY:N	2.82	0.42
21:7:2824:G:H5''	21:7:2825:C:P	2.59	0.42
24:F:77:CYS:C	24:F:79:THR:N	2.77	0.42
25:W:23:A:C8	25:W:24:G:N7	2.87	0.42
2:b:879:G:N2	2:b:887:U:O4	2.52	0.42
9:G:1436:C:O2	9:G:1436:C:H2'	2.19	0.42
10:h:1612:C:H6	10:h:1612:C:O5'	2.02	0.42
15:S:15:GLY:CA	15:S:114:PHE:HE2	2.31	0.42
15:S:60:LYS:CD	15:S:64:LYS:HE3	2.49	0.42
16:2:2229:A:H2'	16:2:2230:C:N1	2.34	0.42
16:2:2285:C:C5	16:2:2286:U:C6	3.07	0.42
18:4:2619:G:N3	18:4:2619:G:H2'	2.34	0.42
19:5:2654:C:H4'	19:5:2657:A:C5	2.54	0.42
21:7:2834:G:C2	21:7:2835:U:C5	3.07	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:Y:9:A:O2'	25:Y:10:G:C8	2.72	0.42
25:V:12:U:H2'	25:V:13:C:O4'	2.19	0.42
3:c:980:G:N1	3:c:981:A:N6	2.67	0.42
9:G:1430:C:O4'	15:S:131:THR:OG1	2.36	0.42
10:h:1623:A:C2'	10:h:1624:G:OP2	2.67	0.42
10:h:1660:A:N6	10:h:1666:G:C6	2.88	0.42
11:T:54:LYS:CG	11:T:55:LYS:N	2.76	0.42
16:2:2212:C:O2'	16:2:2233:A:N6	2.51	0.42
16:2:2247:G:C6	16:2:2248:C:N4	2.87	0.42
16:2:2262:A:C2	16:2:2263:C:O4'	2.72	0.42
22:8:2971:A:H3'	22:8:2971:A:P	2.59	0.42
22:8:2972:G:C2'	22:8:2973:G:O5'	2.67	0.42
25:V:11:C:H2'	25:V:12:U:C6	2.54	0.42
11:T:102:ASN:O	11:T:104:LEU:N	2.52	0.42
16:2:2288:G:C6	16:2:2289:U:C4	3.08	0.42
25:Y:3:C:H2'	25:Y:4:C:H6	1.85	0.42
7:f:1241:U:H4'	7:f:1242:G:C5'	2.49	0.42
9:G:1433:G:C2	9:G:1434:C:C6	3.08	0.42
10:h:1617:G:H22	10:h:1708:A:H2	1.67	0.42
11:T:61:CYS:SG	11:T:65:GLU:HB3	2.59	0.42
11:T:87:VAL:HG11	11:T:164:ILE:CG2	2.49	0.42
11:T:99:THR:C	11:T:101:ARG:N	2.77	0.42
11:T:117:ARG:HE	11:T:189:LYS:NZ	2.17	0.42
12:K:27:VAL:CG1	12:K:90:ILE:HG12	2.49	0.42
14:X:28:ARG:HA	14:X:28:ARG:HD3	1.57	0.42
15:S:20:GLU:O	15:S:24:LEU:HB2	2.20	0.42
16:2:2211:U:H2'	16:2:2212:C:C6	2.54	0.42
16:2:2214:A:C2'	16:2:2215:A:H8	2.32	0.42
16:2:2273:G:H1'	16:2:2274:U:H5	1.85	0.42
16:2:2280:A:O2'	16:2:2281:A:O5'	2.26	0.42
20:6:2690:G:O2'	20:6:2691:A:P	2.74	0.42
21:7:2840:C:C4	21:7:2841:G:C5	3.08	0.42
21:7:2843:U:P	21:7:2843:U:H3'	2.59	0.42
22:8:2969:A:C4	22:8:2970:C:C5	3.07	0.42
3:c:981:A:H2'	3:c:983:A:N7	2.35	0.42
3:c:985:C:C2	3:c:986:G:C8	3.07	0.42
8:g:1142:G:C2	8:g:1143:A:C8	3.07	0.42
10:h:1699:A:HO2'	10:h:1700:A:H8	1.64	0.42
10:h:1710:G:H2'	16:2:2256:A:O5'	2.20	0.42
11:T:45:PRO:HG2	11:T:85:ILE:HG23	2.00	0.42
11:T:76:ARG:H	11:T:76:ARG:HG2	1.68	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:X:39:ILE:HA	14:X:42:ASN:HD22	1.83	0.42
15:S:29:ASN:O	15:S:31:GLU:HG2	2.19	0.42
16:2:2213:A:N6	16:2:2232:A:C4	2.88	0.42
16:2:2276:G:H2'	16:2:2277:C:H5'	2.02	0.42
16:2:2290:C:O2'	16:2:2291:A:O5'	2.38	0.42
25:W:66:U:H2'	25:W:67:C:C6	2.55	0.42
8:g:1170:G:C3'	8:g:1171:G:H5'	2.49	0.42
10:h:1692:C:H2'	10:h:1693:A:C8	2.55	0.42
11:T:63:ILE:HG13	11:T:63:ILE:H	1.60	0.42
12:K:78:ALA:CB	12:K:118:ALA:HB3	2.49	0.42
16:2:2222:A:C2	16:2:2223:A:C6	3.08	0.42
16:2:2259:A:C6	16:2:2260:U:C2	3.08	0.42
16:2:2279:A:C4'	16:2:2280:A:C5'	2.97	0.42
16:2:2286:U:O4	16:2:2288:G:H1'	2.18	0.42
16:2:2300:G:N2	16:2:2301:U:H1'	2.34	0.42
23:B:145:TYR:HA	23:B:148:VAL:CB	2.50	0.42
8:g:1165:A:O2'	8:g:1166:A:OP1	2.37	0.42
10:h:1609:C:O4'	25:V:34:G:O6	2.38	0.42
12:K:33:ILE:O	12:K:97:LEU:HD12	2.19	0.42
16:2:2196:C:C2	16:2:2242:A:N6	2.87	0.42
16:2:2282:U:H5'	16:2:2282:U:H6	1.80	0.42
18:4:2625:C:C2	18:4:2627:C:C5	3.08	0.42
21:7:2853:A:C6	21:7:2854:U:C2	3.07	0.42
25:Y:43:C:C2'	25:Y:44:G:H5'	2.50	0.42
25:V:15:G:N2	25:V:21:A:H1'	2.35	0.42
25:V:21:A:C3'	25:V:22:G:H5''	2.50	0.42
1:a:565:G:H3'	1:a:565:G:H8	1.85	0.42
1:a:576:U:O2'	1:a:577:C:H5'	2.19	0.42
8:g:1163:U:C1'	25:V:34:G:H4'	2.49	0.42
12:K:67:GLU:C	12:K:69:SER:N	2.78	0.42
12:K:112:ALA:C	12:K:114:SER:N	2.78	0.42
14:X:35:SER:O	14:X:38:ARG:HB3	2.20	0.42
16:2:2258:U:H3'	16:2:2259:A:C8	2.50	0.42
20:6:2700:G:C6	20:6:2701:U:C4	3.08	0.42
25:Y:48:C:N3	25:Y:59:U:C2	2.87	0.42
2:b:887:U:C5'	2:b:887:U:C6	3.02	0.41
11:T:45:PRO:HB2	11:T:68:ILE:CD1	2.50	0.41
11:T:164:ILE:O	11:T:168:LEU:HB3	2.20	0.41
12:K:84:ARG:CZ	12:K:88:LEU:HD21	2.50	0.41
16:2:2212:C:H5'	25:W:72:C:HO2'	1.85	0.41
25:Y:4:C:H42	25:Y:69:G:H1	1.68	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:570:G:O4'	1:a:574:A:C2	2.73	0.41
2:b:888:C:H5'	2:b:888:C:C6	2.55	0.41
8:g:1148:G:C6	8:g:1149:C:C4	3.08	0.41
10:h:1636:C:O2'	10:h:1637:U:H5'	2.19	0.41
11:T:35:CYS:H	11:T:63:ILE:CD1	2.29	0.41
11:T:59:THR:O	11:T:61:CYS:N	2.53	0.41
11:T:96:HIS:HD2	11:T:97:LEU:HD23	1.85	0.41
11:T:184:TRP:CE2	11:T:188:LYS:HE3	2.55	0.41
13:L:44:GLY:O	13:L:45:ALA:O	2.38	0.41
15:S:117:ILE:C	15:S:119:LYS:H	2.27	0.41
24:F:79:THR:HA	24:F:80:ARG:HA	1.79	0.41
25:V:44:G:O2'	25:V:45:U:H5'	2.20	0.41
25:W:20:U:O4	25:W:59:U:O4	2.38	0.41
1:a:551:U:HO2'	1:a:552:C:P	2.43	0.41
1:a:561:A:C2	1:a:577:C:H1'	2.55	0.41
10:h:1608:C:C4'	10:h:1609:C:C5'	2.84	0.41
11:T:192:ILE:O	11:T:196:ALA:CB	2.68	0.41
12:K:61:LYS:HD3	12:K:61:LYS:HA	1.80	0.41
15:S:31:GLU:HG2	15:S:31:GLU:H	1.57	0.41
16:2:2262:A:H5''	16:2:2263:C:OP2	2.20	0.41
16:2:2273:G:O2'	16:2:2274:U:P	2.78	0.41
20:6:2706:G:C6	20:6:2707:C:N4	2.88	0.41
21:7:2831:G:C5	21:7:2832:C:C5	3.08	0.41
24:F:80:ARG:O	24:F:82:GLN:N	2.54	0.41
25:W:56:C:C2	25:W:57:G:H8	2.39	0.41
1:a:554:U:H2'	1:a:555:G:C8	2.55	0.41
11:T:52:GLN:HE22	11:T:58:LYS:CE	2.33	0.41
16:2:2196:C:C5'	16:2:2197:C:OP2	2.69	0.41
16:2:2249:G:C8	16:2:2272:G:C8	3.09	0.41
18:4:2619:G:OP1	21:7:2866:U:H5	0.76	0.41
21:7:2828:G:C2'	21:7:2829:U:O5'	2.68	0.41
21:7:2856:G:C6	21:7:2857:C:N4	2.89	0.41
25:Y:71:G:H2'	25:Y:71:G:N3	2.36	0.41
25:W:66:U:O2'	25:W:67:C:H5'	2.20	0.41
13:L:97:GLU:O	13:L:98:ASN:CB	2.67	0.41
15:S:32:LYS:O	15:S:33:LEU:C	2.64	0.41
15:S:63:ASN:HA	15:S:66:ARG:HB2	2.02	0.41
16:2:2299:A:C2'	16:2:2300:G:O5'	2.68	0.41
25:V:39:U:O2'	25:V:40:C:H5'	2.20	0.41
1:a:551:U:C2'	1:a:552:C:O5'	2.68	0.41
7:f:1246:C:HO2'	7:f:3247:A:H5''	1.85	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:h:1652:A:H5'	10:h:1653:C:OP2	2.20	0.41
12:K:105:THR:C	12:K:107:GLN:N	2.79	0.41
12:K:110:PRO:C	12:K:112:ALA:N	2.77	0.41
12:K:142:ARG:HB3	12:K:143:GLU:H	1.67	0.41
13:L:116:ILE:N	13:L:116:ILE:HD12	2.35	0.41
16:2:2213:A:C6	16:2:2214:A:N6	2.89	0.41
16:2:2294:U:C6	16:2:2297:U:C5	3.07	0.41
19:5:2654:C:H5'	19:5:2657:A:N6	2.34	0.41
21:7:2840:C:H5''	21:7:2840:C:H6	1.86	0.41
21:7:2856:G:N1	21:7:2857:C:C4	2.89	0.41
23:B:78:LYS:C	23:B:80:CYS:H	2.28	0.41
25:W:57:G:H21	25:W:58:A:H4'	1.85	0.41
2:b:883:A:N6	3:c:976:A:O2'	2.41	0.41
11:T:177:LYS:O	11:T:178:ASN:CB	2.67	0.41
12:K:61:LYS:CE	12:K:76:GLN:HB3	2.51	0.41
13:L:117:PRO:HG2	13:L:118:GLY:N	2.32	0.41
14:X:44:ARG:CG	14:X:45:TYR:N	2.83	0.41
22:8:2971:A:H4'	22:8:2972:G:C5'	2.50	0.41
22:8:2973:G:C2'	22:8:2974:U:H5'	2.51	0.41
25:V:1:G:O6	25:V:72:C:N4	2.52	0.41
2:b:878:A:C5'	12:K:39:ASP:OD2	2.69	0.41
7:f:1245:G:C2'	7:f:1246:C:OP2	2.68	0.41
9:G:1433:G:H4'	25:V:29:G:O3'	2.21	0.41
13:L:34:LEU:C	13:L:36:SER:N	2.79	0.41
15:S:83:THR:HG1	15:S:85:TYR:HD1	1.57	0.41
16:2:2210:G:C5	16:2:2211:U:C5	3.08	0.41
16:2:2273:G:O2'	16:2:2274:U:OP2	2.37	0.41
16:2:2276:G:N7	16:2:2277:C:H5	2.19	0.41
16:2:2280:A:H8	16:2:2280:A:H3'	1.85	0.41
16:2:2286:U:P	25:Y:72:C:OP2	2.79	0.41
21:7:2826:U:C4	21:7:2827:U:C4	3.08	0.41
21:7:2864:A:C6	21:7:2865:U:N3	2.89	0.41
25:V:48:C:H2'	25:V:59:U:C1'	2.51	0.41
10:h:1674:A:C6	10:h:1675:A:N7	2.88	0.41
10:h:1710:G:C2'	16:2:2256:A:O5'	2.69	0.41
11:T:72:MET:HB3	11:T:84:CYS:SG	2.61	0.41
11:T:118:GLU:OE1	11:T:197:LYS:HE2	2.20	0.41
11:T:138:PRO:HG2	11:T:139:MET:H	1.86	0.41
12:K:42:ILE:HD11	12:K:115:ALA:CB	2.51	0.41
16:2:2236:G:H2'	16:2:2237:C:O5'	2.21	0.41
16:2:2260:U:OP1	16:2:2260:U:C4'	2.68	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:2:2282:U:P	22:8:2973:G:O2'	2.77	0.41
21:7:2868:U:C6	21:7:2869:U:H5	2.38	0.41
23:B:12:HIS:CB	23:B:213:ALA:HB1	2.50	0.41
24:F:11:TYR:O	24:F:20:HIS:HA	2.20	0.41
24:F:54:THR:O	24:F:56:PRO:N	2.54	0.41
25:Y:28:G:N2	25:Y:43:C:C6	2.89	0.41
25:Y:43:C:O2'	25:Y:44:G:H5'	2.21	0.41
25:W:51:U:H3	25:W:63:G:H1	1.69	0.41
25:W:67:C:C4	25:W:68:C:N4	2.89	0.41
1:a:549:A:N3	1:a:584:C:H1'	2.36	0.41
10:h:1645:C:N4	10:h:1680:A:H61	2.16	0.41
11:T:42:VAL:HG12	11:T:43:PHE:N	2.36	0.41
12:K:116:LEU:HD23	12:K:116:LEU:O	2.21	0.41
14:X:59:SER:C	14:X:61:ASN:N	2.79	0.41
15:S:42:PHE:HB3	15:S:46:THR:HB	2.03	0.41
16:2:2194:G:N9	16:2:2274:U:O2	2.53	0.41
20:6:2696:A:C8	20:6:2697:A:C8	3.08	0.41
21:7:2872:A:H1'	21:7:2873:U:C5'	2.50	0.41
6:E:1590:C:O2'	6:E:1591:C:C5'	2.69	0.40
8:g:1157:U:C2	9:G:1429:G:C8	3.09	0.40
11:T:70:THR:HG21	11:T:147:TYR:HA	2.03	0.40
13:L:53:THR:HB	13:L:72:ARG:O	2.21	0.40
13:L:69:LYS:NZ	13:L:69:LYS:CB	2.74	0.40
20:6:2694:A:H5'	20:6:2695:A:OP2	2.21	0.40
20:6:2704:A:O2'	20:6:2705:A:H2'	2.21	0.40
25:Y:69:G:N2	25:Y:70:G:C6	2.89	0.40
1:a:551:U:H2'	1:a:552:C:O5'	2.21	0.40
1:a:564:A:H8	1:a:564:A:O5'	2.03	0.40
2:b:881:U:O2	2:b:883:A:OP2	2.39	0.40
7:f:1238:U:O2	7:f:1238:U:H2'	2.21	0.40
10:h:1654:U:H2'	10:h:1655:G:H8	1.86	0.40
11:T:45:PRO:O	11:T:65:GLU:OE1	2.40	0.40
11:T:99:THR:O	11:T:99:THR:CG2	2.65	0.40
12:K:110:PRO:CG	12:K:111:GLY:H	2.34	0.40
13:L:29:PHE:CZ	13:L:33:LEU:CD1	3.04	0.40
16:2:2285:C:C5	16:2:2286:U:N1	2.87	0.40
18:4:2620:G:C2'	18:4:2621:G:H5'	2.52	0.40
21:7:2843:U:H3'	21:7:2843:U:OP2	2.21	0.40
25:Y:28:G:N2	25:Y:43:C:C5	2.90	0.40
25:W:14:A:H3'	25:W:15:G:C8	2.54	0.40
25:W:14:A:H2'	25:W:14:A:N3	2.37	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:W:57:G:H2'	25:W:58:A:C5'	2.38	0.40
9:G:1429:G:OP2	9:G:1430:C:C5	2.68	0.40
11:T:67:LEU:HD22	11:T:107:PHE:CE1	2.56	0.40
15:S:31:GLU:CG	15:S:32:LYS:H	2.29	0.40
16:2:2213:A:C6	16:2:2214:A:C6	3.09	0.40
16:2:2267:C:N3	16:2:2268:U:O2	2.55	0.40
18:4:2625:C:O5'	18:4:2625:C:H6	2.04	0.40
19:5:2655:U:C4'	19:5:2656:A:H5'	2.49	0.40
25:V:59:U:O2'	25:V:60:U:H5'	2.21	0.40
11:T:29:PHE:CD1	11:T:29:PHE:N	2.90	0.40
11:T:99:THR:CG2	11:T:101:ARG:HE	2.33	0.40
16:2:2270:A:H2'	16:2:2271:A:O4'	2.20	0.40
18:4:2620:G:N1	25:V:75:C:C2	2.90	0.40
18:4:2626:A:H5'	18:4:2627:C:C5'	2.51	0.40
25:Y:42:C:H3'	25:Y:43:C:C5'	2.50	0.40
1:a:543:A:C2	1:a:550:G:C5	3.09	0.40
1:a:564:A:C2	1:a:568:G:O6	2.75	0.40
3:c:986:G:N1	3:c:987:U:C4	2.89	0.40
11:T:183:SER:C	11:T:185:ALA:H	2.30	0.40
13:L:130:SER:O	13:L:133:ALA:HB3	2.21	0.40
16:2:2223:A:O2'	16:2:2224:A:C5'	2.70	0.40
16:2:2229:A:H2'	16:2:2230:C:C1'	2.52	0.40
21:7:2830:G:H5'	21:7:2830:G:H8	1.86	0.40
21:7:2864:A:C4	21:7:2865:U:C6	3.10	0.40
23:B:138:VAL:O	23:B:139:SER:C	2.63	0.40
24:F:20:HIS:O	24:F:21:THR:CB	2.68	0.40
25:W:15:G:N2	25:W:48:C:C2	2.88	0.40
25:W:69:G:H3'	25:W:70:G:H5''	2.01	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	
11	T	190/192 (99%)	141 (74%)	33 (17%)	16 (8%)	0	9
12	K	138/140 (99%)	96 (70%)	28 (20%)	14 (10%)	0	7
13	L	139/141 (99%)	106 (76%)	17 (12%)	16 (12%)	0	5
14	X	66/68 (97%)	48 (73%)	12 (18%)	6 (9%)	0	8
15	S	123/125 (98%)	91 (74%)	25 (20%)	7 (6%)	1	14
23	B	211/213 (99%)	76 (36%)	67 (32%)	68 (32%)	0	0
24	F	93/95 (98%)	37 (40%)	28 (30%)	28 (30%)	0	0
All	All	960/974 (99%)	595 (62%)	210 (22%)	155 (16%)	0	2

All (155) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
11	T	56	PHE
11	T	60	GLN
11	T	63	ILE
11	T	75	GLY
11	T	158	PHE
12	K	70	SER
12	K	100	LYS
12	K	124	MET
12	K	142	ARG
13	L	45	ALA
13	L	62	GLN
13	L	87	PRO
13	L	88	MET
13	L	89	ASP
13	L	107	GLY
13	L	108	ARG
13	L	117	PRO
14	X	8	LEU
14	X	64	ALA
15	S	15	GLY
23	B	24	LYS
23	B	36	VAL
23	B	39	LYS
23	B	42	ASP
23	B	43	PRO
23	B	55	LEU
23	B	56	PRO
23	B	58	CYS
23	B	59	PRO

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Mol	Chain	Res	Type
23	B	60	ARG
23	B	61	PRO
23	B	109	ALA
23	B	115	VAL
23	B	120	VAL
23	B	122	ARG
23	B	124	LEU
23	B	126	PRO
23	B	127	GLN
23	B	128	LEU
23	B	129	SER
23	B	135	PRO
23	B	143	ASP
23	B	151	VAL
23	B	174	MET
23	B	201	VAL
23	B	207	LYS
23	B	208	SER
23	B	209	SER
23	B	210	MET
23	B	212	PRO
24	F	5	PRO
24	F	13	LYS
24	F	34	SER
24	F	47	GLN
24	F	53	GLN
24	F	66	LYS
24	F	93	LEU
11	T	58	LYS
11	T	100	GLY
11	T	157	ALA
11	T	159	LYS
11	T	183	SER
12	K	102	GLY
12	K	105	THR
12	K	110	PRO
12	K	139	SER
13	L	35	GLY
13	L	54	GLU
13	L	76	ARG
13	L	138	LYS
14	X	24	LYS

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Mol	Chain	Res	Type
15	S	31	GLU
15	S	33	LEU
23	B	25	LYS
23	B	26	ARG
23	B	41	TYR
23	B	44	GLN
23	B	47	LYS
23	B	67	ILE
23	B	77	ALA
23	B	96	ASN
23	B	117	ILE
23	B	123	LEU
23	B	169	VAL
23	B	193	LEU
23	B	194	LEU
23	B	196	LYS
23	B	197	ASN
23	B	204	LEU
23	B	206	VAL
24	F	6	LYS
24	F	9	LYS
24	F	11	TYR
24	F	21	THR
24	F	41	ARG
24	F	46	LYS
24	F	57	VAL
24	F	58	PHE
24	F	81	ALA
24	F	88	CYS
11	T	41	GLN
11	T	103	PRO
11	T	179	ASN
12	K	23	ALA
12	K	99	ALA
12	K	106	LYS
12	K	122	SER
12	K	141	ARG
13	L	109	GLN
14	X	60	PRO
15	S	130	PRO
23	B	22	GLU
23	B	49	PHE

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Mol	Chain	Res	Type
23	B	57	ASN
23	B	101	LYS
23	B	102	LYS
23	B	152	ARG
23	B	202	GLY
24	F	17	CYS
24	F	19	LYS
24	F	56	PRO
12	K	146	ARG
13	L	91	CYS
14	X	25	ASP
23	B	27	ASN
23	B	107	TYR
23	B	132	GLY
23	B	155	ILE
23	B	165	LEU
23	B	189	PHE
23	B	213	ALA
24	F	12	CYS
24	F	59	HIS
11	T	62	PRO
14	X	26	LYS
15	S	123	GLU
23	B	5	THR
23	B	45	ARG
23	B	92	LYS
23	B	108	ASN
23	B	154	THR
24	F	3	ASN
24	F	39	GLY
24	F	45	ARG
24	F	55	LYS
24	F	62	ALA
24	F	48	SER
11	T	102	ASN
11	T	138	PRO
15	S	104	GLY
23	B	69	GLY
13	L	110	GLY
23	B	65	ILE
13	L	114	GLY
15	S	53	GLY

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	
11	T	163/163 (100%)	146 (90%)	17 (10%)	7	22
12	K	112/112 (100%)	96 (86%)	16 (14%)	3	13
13	L	113/113 (100%)	103 (91%)	10 (9%)	9	28
14	X	57/57 (100%)	55 (96%)	2 (4%)	32	53
15	S	105/105 (100%)	101 (96%)	4 (4%)	29	50
All	All	550/550 (100%)	501 (91%)	49 (9%)	11	28

All (49) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
11	T	24	ILE
11	T	30	GLN
11	T	33	ILE
11	T	36	THR
11	T	41	GLN
11	T	44	VAL
11	T	51	TYR
11	T	56	PHE
11	T	57	ARG
11	T	65	GLU
11	T	73	PHE
11	T	74	HIS
11	T	117	ARG
11	T	121	THR
11	T	134	VAL
11	T	139	MET
11	T	178	ASN
12	K	28	PHE
12	K	43	HIS
12	K	46	ASP
12	K	47	LEU
12	K	67	GLU
12	K	79	ILE

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Mol	Chain	Res	Type
12	K	80	ASP
12	K	85	CYS
12	K	95	ILE
12	K	113	GLN
12	K	116	LEU
12	K	117	ARG
12	K	119	LEU
12	K	129	ILE
12	K	140	THR
12	K	149	ARG
13	L	9	ILE
13	L	19	ARG
13	L	33	LEU
13	L	59	GLU
13	L	69	LYS
13	L	82	ILE
13	L	89	ASP
13	L	92	LEU
13	L	106	LEU
13	L	132	LEU
14	X	34	ARG
14	X	62	TRP
15	S	54	ILE
15	S	65	VAL
15	S	83	THR
15	S	124	PHE

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (16) such sidechains are listed below:

Mol	Chain	Res	Type
11	T	10	GLN
11	T	41	GLN
11	T	46	HIS
11	T	52	GLN
11	T	96	HIS
11	T	102	ASN
11	T	143	ASN
12	K	43	HIS
13	L	47	HIS
13	L	64	ASN
13	L	78	ASN
13	L	98	ASN

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Mol	Chain	Res	Type
14	X	17	GLN
14	X	42	ASN
15	S	29	ASN
15	S	84	HIS

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	a	47/48 (97%)	16 (34%)	0
10	h	110/111 (99%)	12 (10%)	0
16	2	111/112 (99%)	55 (49%)	9 (8%)
17	3	11/12 (91%)	6 (54%)	1 (9%)
18	4	13/14 (92%)	5 (38%)	1 (7%)
19	5	5/6 (83%)	4 (80%)	1 (20%)
2	b	11/12 (91%)	4 (36%)	0
20	6	18/19 (94%)	13 (72%)	1 (5%)
21	7	49/50 (98%)	29 (59%)	5 (10%)
22	8	19/20 (95%)	8 (42%)	0
25	V	74/76 (97%)	15 (20%)	0
25	W	75/76 (98%)	13 (17%)	0
25	Y	74/76 (97%)	20 (27%)	1 (1%)
26	v	2/3 (66%)	0	0
26	y	2/3 (66%)	0	0
27	w	1/2 (50%)	0	0
3	c	16/17 (94%)	4 (25%)	0
4	d	6/7 (85%)	2 (33%)	0
5	e	3/4 (75%)	0	0
6	E	4/5 (80%)	1 (25%)	1 (25%)
7	f	20/21 (95%)	5 (25%)	0
8	g	30/31 (96%)	8 (26%)	0
9	G	12/13 (92%)	2 (16%)	1 (8%)
All	All	713/738 (96%)	222 (31%)	21 (2%)

All (222) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	a	544	G
1	a	547	C
1	a	548	A
1	a	552	C
1	a	553	A

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Mol	Chain	Res	Type
1	a	554	U
1	a	558	G
1	a	559	C
1	a	560	C
1	a	568	G
1	a	573	A
1	a	574	A
1	a	575	U
1	a	576	U
1	a	577	C
1	a	588	A
2	b	882	G
2	b	886	U
2	b	887	U
2	b	888	C
3	c	975	G
3	c	976	A
3	c	978	C
3	c	982	U
4	d	1545	A
4	d	1546	G
6	E	1591	C
7	f	1242	G
7	f	1245	G
7	f	1246	C
7	f	3247	A
7	f	1256	C
8	g	1155	A
8	g	1158	U
8	g	1163	U
8	g	1165	A
8	g	1166	A
8	g	1168	A
8	g	1169	C
8	g	1172	G
9	G	1430	C
9	G	1431	A
10	h	1608	C
10	h	1609	C
10	h	1624	G
10	h	1649	U
10	h	1651	G

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Mol	Chain	Res	Type
10	h	1652	A
10	h	1661	G
10	h	1674	A
10	h	1710	G
10	h	1713	G
10	h	1714	U
10	h	1715	A
16	2	2195	C
16	2	2201	G
16	2	2205	U
16	2	2206	G
16	2	2207	A
16	2	2208	A
16	2	2209	U
16	2	2210	G
16	2	2211	U
16	2	2215	A
16	2	2222	A
16	2	2223	A
16	2	2224	A
16	2	2227	C
16	2	2237	C
16	2	2244	A
16	2	2247	G
16	2	2248	C
16	2	2249	G
16	2	2250	G
16	2	2252	A
16	2	2254	U
16	2	2255	A
16	2	2256	A
16	2	2257	C
16	2	2259	A
16	2	2260	U
16	2	2261	G
16	2	2262	A
16	2	2263	C
16	2	2264	U
16	2	2266	U
16	2	2270	A
16	2	2271	A
16	2	2272	G

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Mol	Chain	Res	Type
16	2	2273	G
16	2	2275	A
16	2	2276	G
16	2	2277	C
16	2	2280	A
16	2	2281	A
16	2	2282	U
16	2	2283	G
16	2	2286	U
16	2	2288	G
16	2	2290	C
16	2	2291	A
16	2	2292	U
16	2	2295	A
16	2	2296	A
16	2	2297	U
16	2	2298	U
16	2	2299	A
16	2	2300	G
16	2	2303	A
17	3	2479	C
17	3	2484	A
17	3	2485	A
17	3	2486	A
17	3	2487	U
17	3	2488	A
18	4	2618	G
18	4	2619	G
18	4	2624	G
18	4	2626	A
18	4	2627	C
19	5	2655	U
19	5	2656	A
19	5	2657	A
19	5	2658	G
20	6	2690	G
20	6	2691	A
20	6	2693	C
20	6	2694	A
20	6	2695	A
20	6	2696	A
20	6	2697	A

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Mol	Chain	Res	Type
20	6	2699	G
20	6	2700	G
20	6	2702	A
20	6	2703	A
20	6	2704	A
20	6	2707	C
21	7	2825	C
21	7	2826	U
21	7	2828	G
21	7	2829	U
21	7	2830	G
21	7	2833	A
21	7	2834	G
21	7	2835	U
21	7	2836	C
21	7	2838	A
21	7	2839	G
21	7	2840	C
21	7	2842	U
21	7	2843	U
21	7	2844	C
21	7	2845	A
21	7	2846	U
21	7	2847	A
21	7	2849	C
21	7	2850	G
21	7	2851	A
21	7	2852	C
21	7	2859	U
21	7	2860	U
21	7	2867	C
21	7	2870	C
21	7	2871	G
21	7	2872	A
21	7	2873	U
22	8	2960	C
22	8	2961	G
22	8	2962	U
22	8	2965	U
22	8	2968	G
22	8	2971	A
22	8	2972	G

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Mol	Chain	Res	Type
22	8	2973	G
25	Y	2	C
25	Y	9	A
25	Y	16	U
25	Y	17	C
25	Y	18	G
25	Y	19	G
25	Y	21	A
25	Y	23	A
25	Y	34	G
25	Y	42	C
25	Y	43	C
25	Y	47	U
25	Y	48	C
25	Y	52	G
25	Y	53	G
25	Y	70	G
25	Y	71	G
25	Y	72	C
25	Y	73	A
25	Y	74	C
25	V	8	U
25	V	17	C
25	V	18	G
25	V	19	G
25	V	20	U
25	V	21	A
25	V	22	G
25	V	42	C
25	V	46	G
25	V	47	U
25	V	48	C
25	V	52	G
25	V	61	C
25	V	69	G
25	V	75	C
25	W	16	U
25	W	17	C
25	W	18	G
25	W	19	G
25	W	21	A
25	W	39	U

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Mol	Chain	Res	Type
25	W	43	C
25	W	47	U
25	W	57	G
25	W	58	A
25	W	61	C
25	W	70	G
25	W	76	A

All (21) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
6	E	1590	C
9	G	1430	C
16	2	2211	U
16	2	2251	G
16	2	2254	U
16	2	2255	A
16	2	2270	A
16	2	2280	A
16	2	2281	A
16	2	2290	C
16	2	2297	U
17	3	2487	U
18	4	2618	G
19	5	2656	A
20	6	2699	G
21	7	2834	G
21	7	2850	G
21	7	2851	A
21	7	2859	U
21	7	2872	A
25	Y	15	G

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

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

5.5 Carbohydrates ⓘ

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

There are no ligands in this entry.

5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

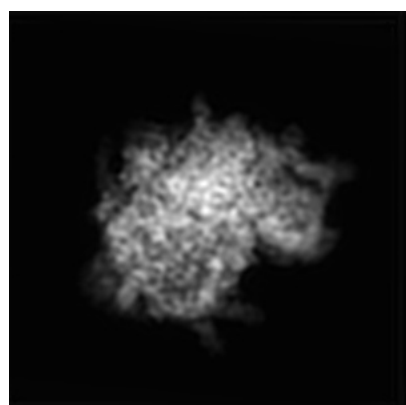
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-5327. These allow visual inspection of the internal detail of the map and identification of artifacts.

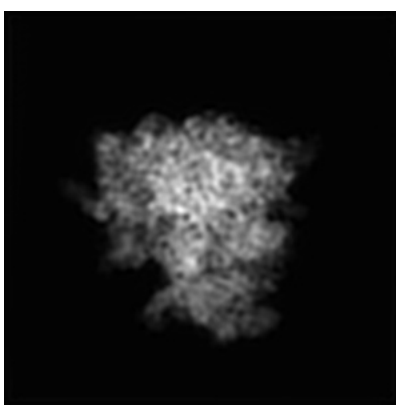
No raw map or half-maps were deposited for this entry and therefore no images, graphs, etc. pertaining to the raw map can be shown.

6.1 Orthogonal projections [i](#)

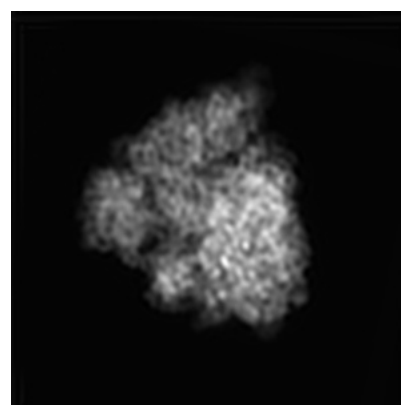
6.1.1 Primary map



X



Y

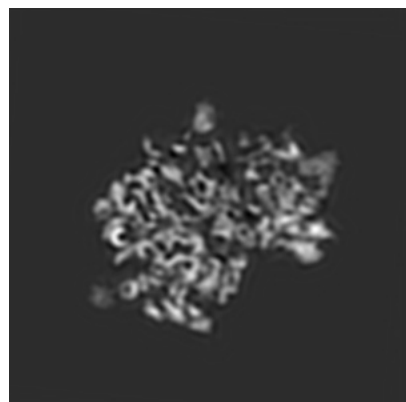


Z

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

6.2 Central slices [i](#)

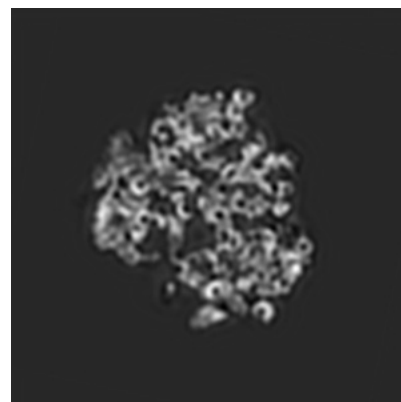
6.2.1 Primary map



X Index: 90



Y Index: 90

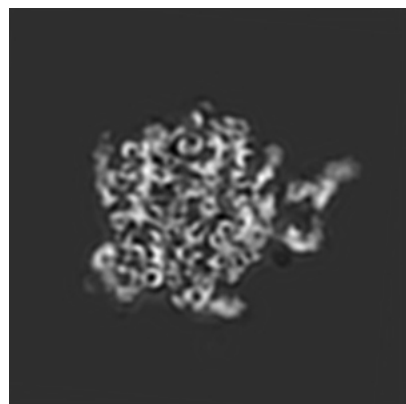


Z Index: 90

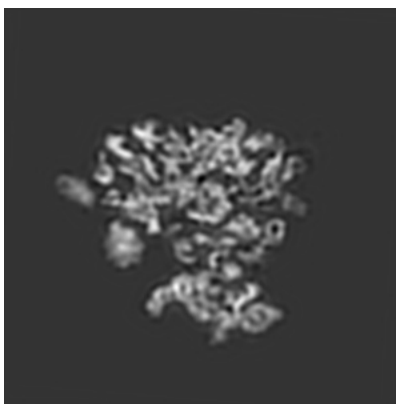
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

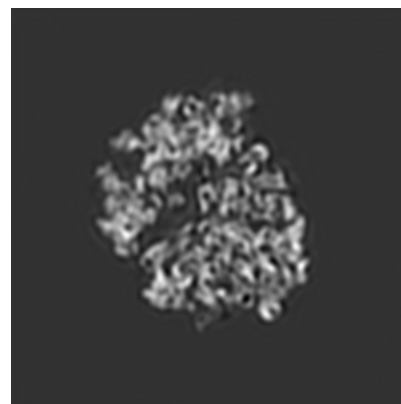
6.3.1 Primary map



X Index: 108



Y Index: 91

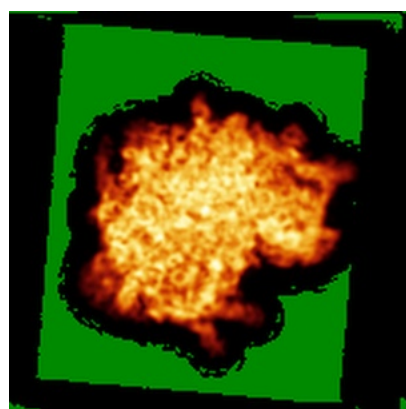


Z Index: 84

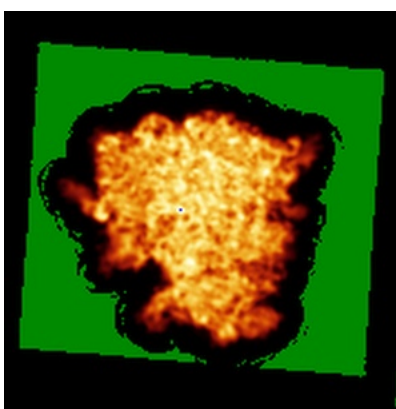
The images above show the largest variance slices of the map in three orthogonal directions.

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

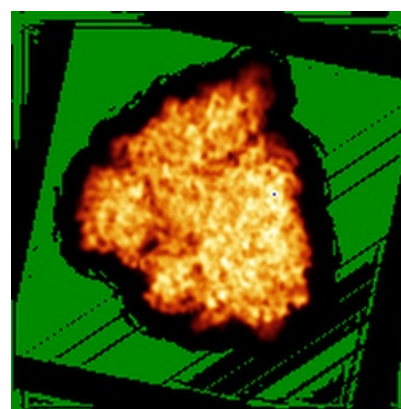
6.4.1 Primary map



X



Y



Z

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

6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 1000.0. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

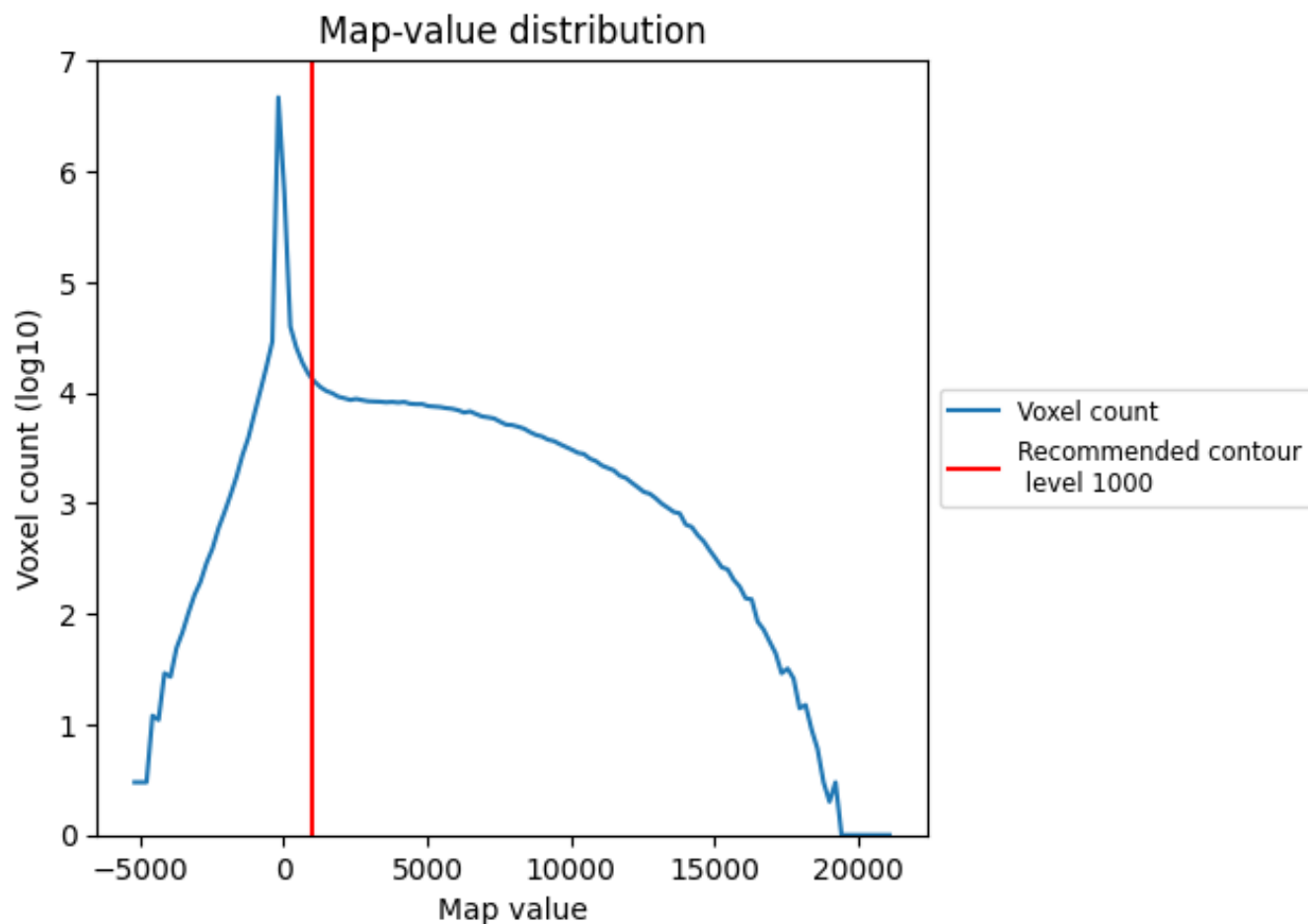
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

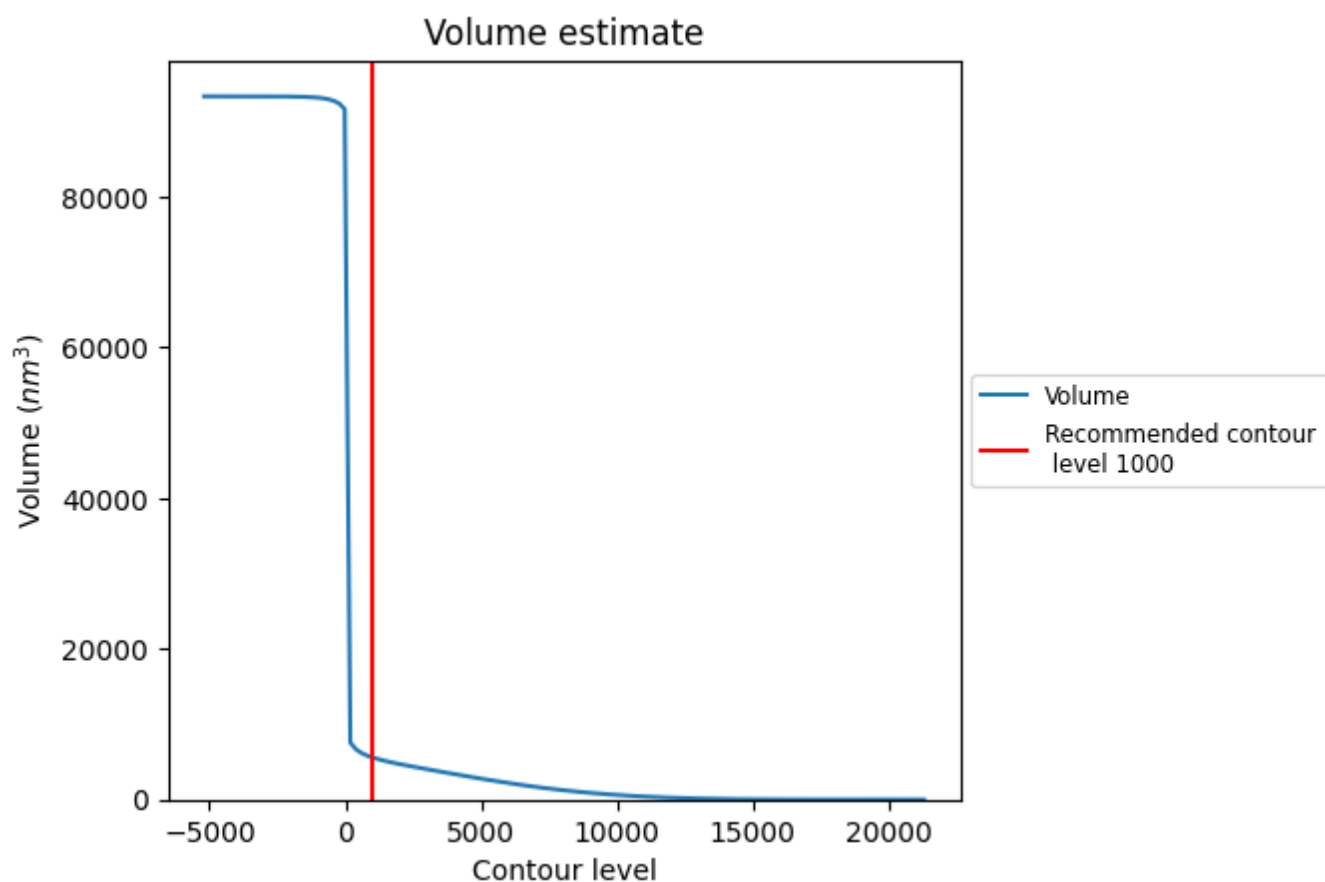
This section contains the results of statistical analysis of the map.

7.1 Map-value distribution [i](#)



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

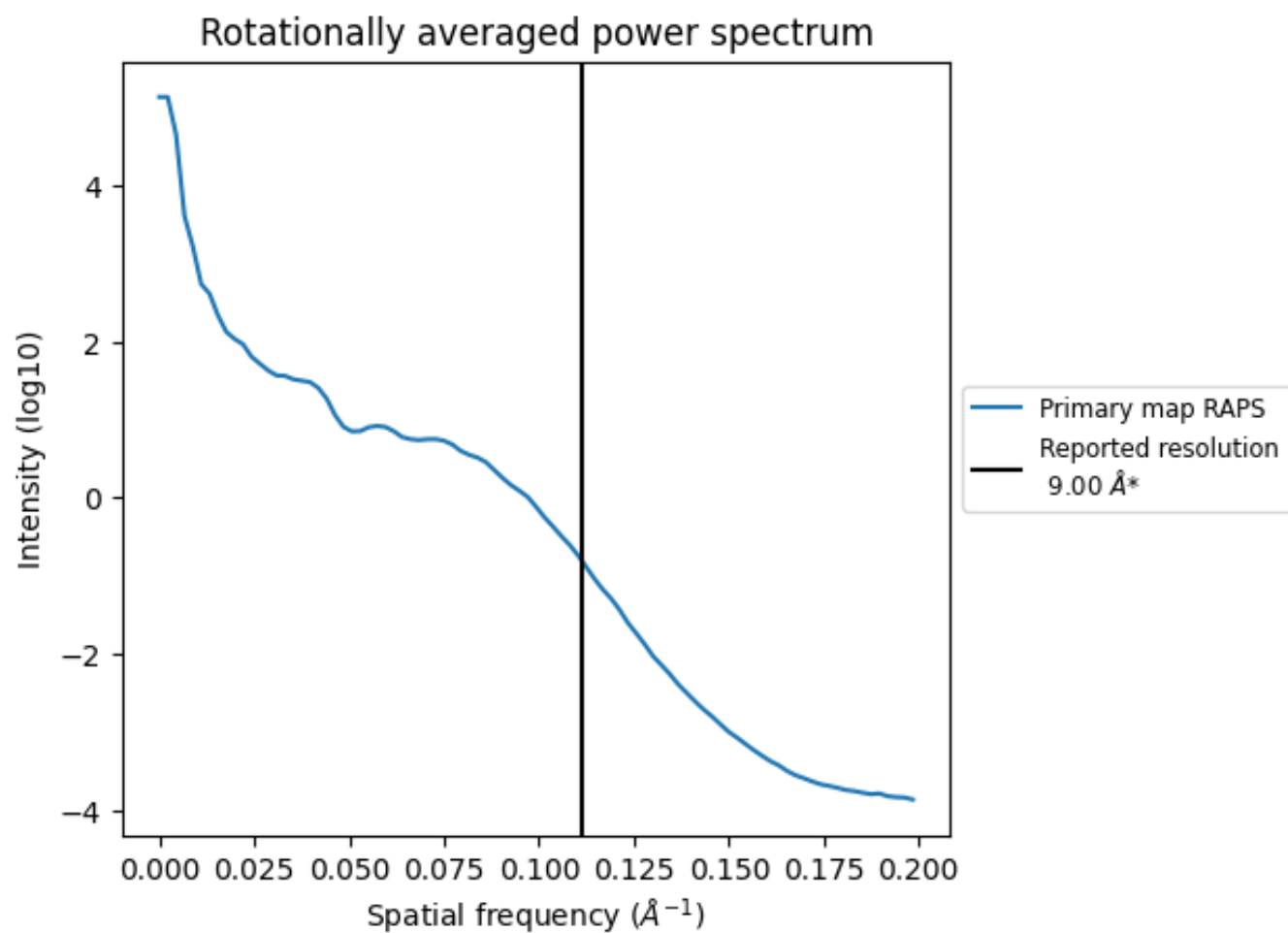
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 5581 nm^3 ; this corresponds to an approximate mass of 5042 kDa.

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

7.3 Rotationally averaged power spectrum ⓘ



*Reported resolution corresponds to spatial frequency of 0.111 Å⁻¹

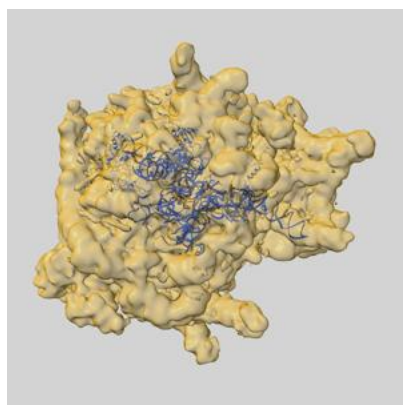
8 Fourier-Shell correlation

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

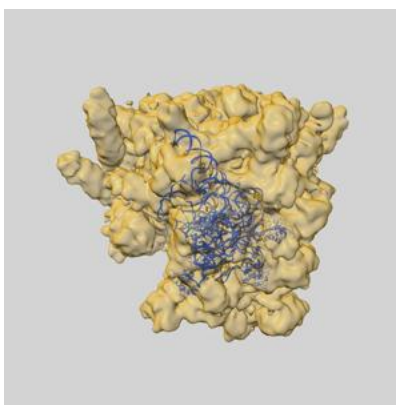
9 Map-model fit [i](#)

This section contains information regarding the fit between EMDB map EMD-5327 and PDB model 3J0O. Per-residue inclusion information can be found in section [3](#) on page [10](#).

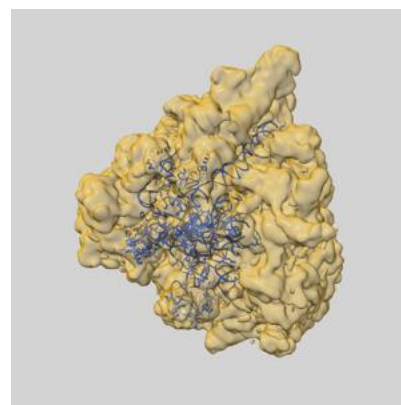
9.1 Map-model overlay [i](#)



X



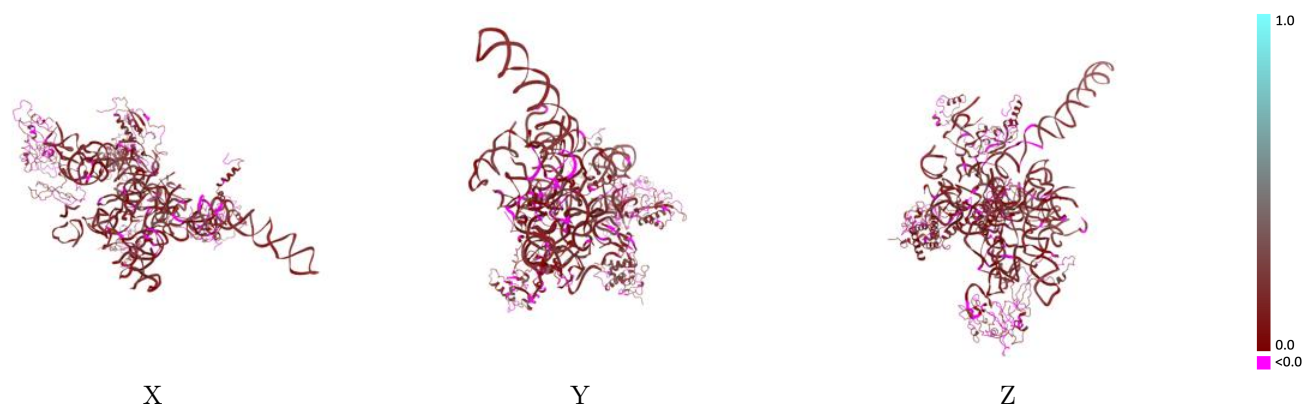
Y



Z

The images above show the 3D surface view of the map at the recommended contour level 1000.0 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](#)

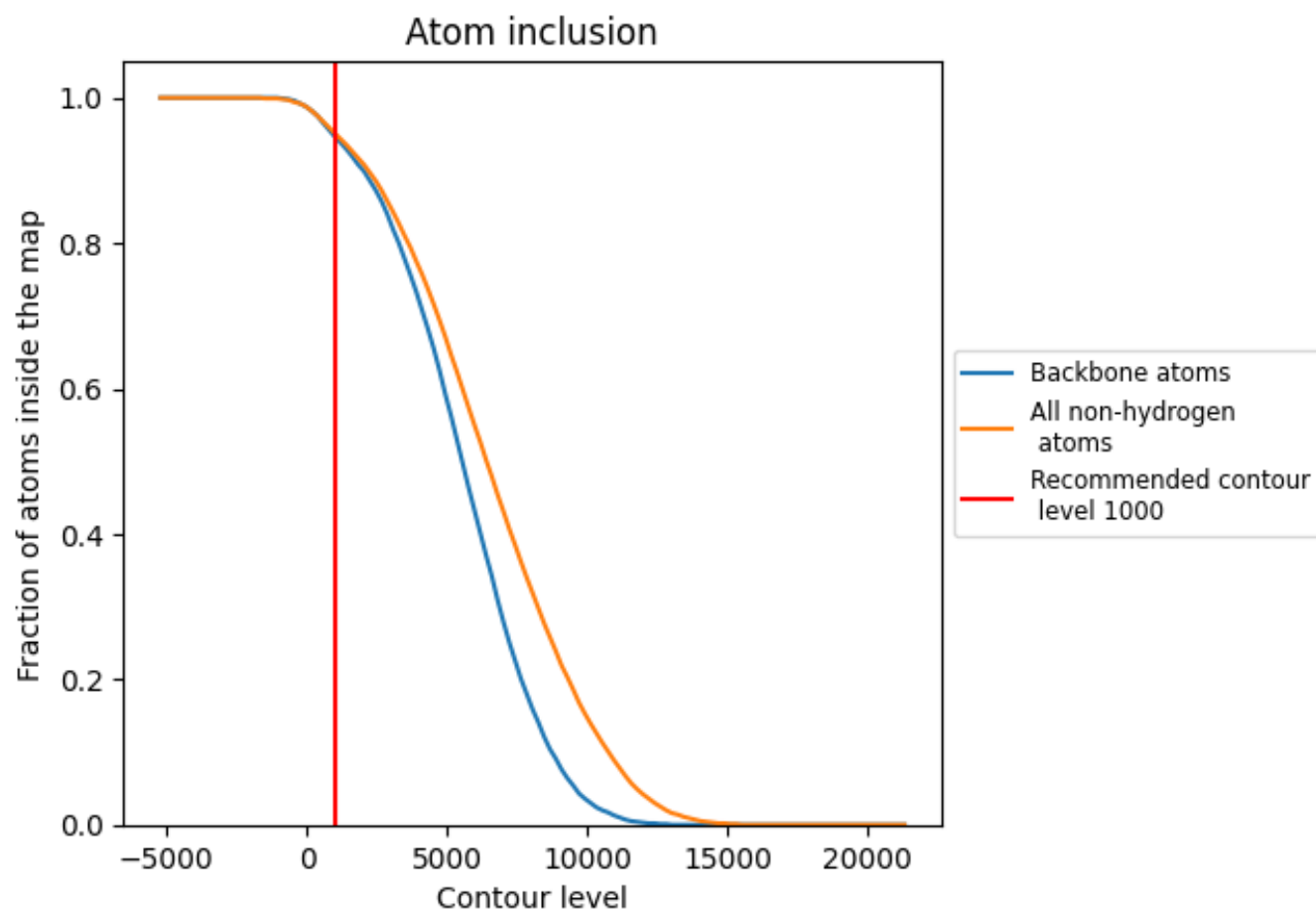


The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions 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](#)

This section was not generated.

9.4 Atom inclusion [i](#)



At the recommended contour level, 95% of all backbone atoms, 95% 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 (1000) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.9520	 0.1190
2	 0.9840	 0.1500
3	 1.0000	 0.1050
4	 0.9970	 0.1530
5	 1.0000	 0.1680
6	 1.0000	 0.1770
7	 0.9970	 0.1520
8	 0.9930	 0.1590
B	 0.8500	 0.0350
E	 1.0000	 0.1290
F	 0.9910	 0.1210
G	 1.0000	 0.1720
K	 0.9240	 0.0840
L	 0.9110	 0.0570
S	 0.9530	 0.0910
T	 0.9760	 0.1050
V	 0.9630	 0.1400
W	 0.9490	 0.1190
X	 0.7440	 0.0440
Y	 0.8770	 0.1200
a	 0.9960	 0.1350
b	 0.9960	 0.1440
c	 0.9860	 0.1360
d	 1.0000	 0.1320
e	 1.0000	 0.1310
f	 0.9980	 0.1380
g	 1.0000	 0.1320
h	 0.9450	 0.1310
v	 1.0000	 0.1860
w	 0.8640	 0.0260
y	 1.0000	 0.0860

