



Full wwPDB EM Validation Report ⓘ

Mar 28, 2026 – 06:42 PM UTC

PDB ID : 9V26 / pdb_00009v26
EMDB ID : EMD-64718
Title : Cryo- EM structure of 75S ribosome with A/P- & P/E- tRNAs from Entamoeba histolytica bound to antibiotic paromomycin
Authors : Sharma, S.; Mishra, S.; Gourinath, S.; Kaushal, P.S.
Deposited on : 2025-05-19
Resolution : 3.10 Å(reported)
Based on initial model : .

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

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

A user guide is available at

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

The types of validation reports are described at

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

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

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

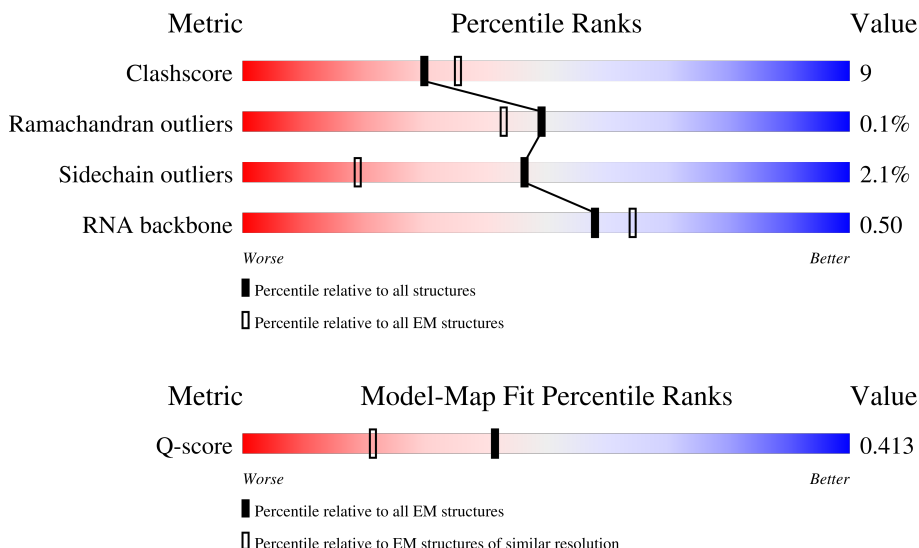
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 3.10 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	14724 (2.60 - 3.60)

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.

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1	1A	3503	
2	1B	155	
3	1C	117	

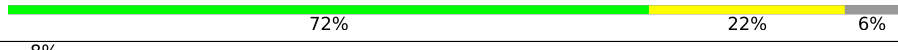
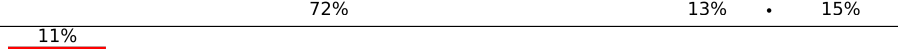
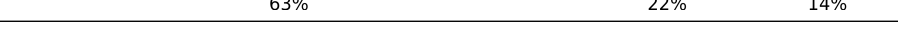
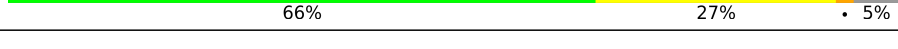
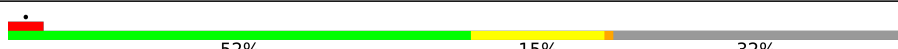
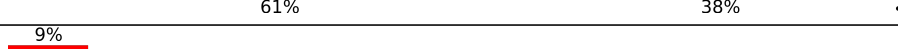
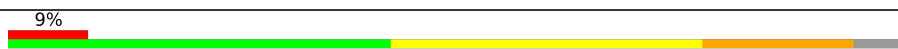
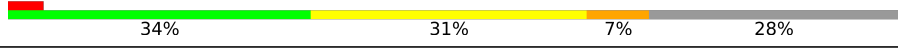
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Mol	Chain	Length	Quality of chain
4	ID	257	
5	IE	402	
6	IF	431	
7	IG	286	
8	IH	204	
9	II	230	
10	IJ	286	
11	IK	197	
12	IL	210	
13	IM	174	
14	IN	291	
15	IO	205	
16	IP	135	
17	IQ	205	
18	IR	179	
19	IS	168	
20	IT	173	
21	IU	198	
22	IV	166	
23	IW	137	
24	IX	140	
25	IY	121	
26	IZ	163	
27	la	213	
28	lb	139	

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Mol	Chain	Length	Quality of chain
29	lc	149	
30	ld	64	
31	le	109	
32	lf	150	
33	lg	134	
34	lh	137	
35	li	122	
36	lj	108	
37	lk	104	
38	ll	77	
39	lm	93	
40	ln	77	
41	lo	51	
42	lp	56	
43	lq	98	
44	sA	137	
45	sB	144	
46	sC	84	
47	sD	69	
48	sE	56	
49	sI	76	
50	sJ	77	
51	sK	10	
52	sa	1947	
53	sc	255	

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Mol	Chain	Length	Quality of chain
54	sd	244	
55	se	256	
56	sf	326	
57	sg	206	
58	sh	266	
59	si	201	
60	sj	237	
61	sk	185	
62	sl	127	
63	sm	156	
64	so	151	
65	sp	146	
66	sq	144	
67	sr	130	
68	ss	158	
69	st	117	
70	su	155	
71	sv	155	
72	sw	118	
73	sx	86	
74	sy	141	

2 Entry composition

There are 75 unique types of molecules in this entry. The entry contains 178778 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 25S rRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	1A	3142	Total	C	N	O	P	0	0
			67153	30106	12192	21713	3142		

- Molecule 2 is a RNA chain called 5.8S rRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	1B	145	Total	C	N	O	P	0	0
			3097	1390	560	1002	145		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
3	1C	117	Total	C	N	O	P	0	0
			2477	1108	425	827	117		

- Molecule 4 is a protein called Large ribosomal subunit protein uL2.

Mol	Chain	Residues	Atoms					AltConf	Trace
4	1D	246	Total	C	N	O	S	0	0
			1881	1165	382	326	8		

- Molecule 5 is a protein called 60S ribosomal protein L3, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
5	1E	387	Total	C	N	O	S	0	0
			3076	1956	578	527	15		

- Molecule 6 is a protein called 60S ribosomal protein L4, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
6	1F	425	Total	C	N	O	S	0	0
			3281	2091	624	552	14		

- Molecule 7 is a protein called 60S ribosomal protein L5, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
7	lG	278	Total	C	N	O	S	0	0
			2209	1412	399	390	8		

- Molecule 8 is a protein called Large ribosomal subunit protein eL6.

Mol	Chain	Residues	Atoms					AltConf	Trace
8	lH	203	Total	C	N	O	S	0	0
			1608	1054	272	278	4		

- Molecule 9 is a protein called 60S ribosomal protein L7, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
9	lI	210	Total	C	N	O	S	0	0
			1658	1067	301	282	8		

- Molecule 10 is a protein called 60S ribosomal protein L7a.

Mol	Chain	Residues	Atoms					AltConf	Trace
10	lJ	201	Total	C	N	O	S	0	0
			1640	1058	302	275	5		

- Molecule 11 is a protein called 60S ribosomal protein L9, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
11	lK	193	Total	C	N	O	S	0	0
			1538	974	279	279	6		

- Molecule 12 is a protein called Ribosomal protein L10, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
12	lL	200	Total	C	N	O	S	0	0
			1597	1017	302	264	14		

- Molecule 13 is a protein called 60S ribosomal protein L11, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
13	lM	170	Total	C	N	O	S	0	0
			1350	857	243	245	5		

- Molecule 14 is a protein called 60S ribosomal protein L13, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
14	lN	267	Total	C	N	O	S	0	0
			2130	1358	412	352	8		

- Molecule 15 is a protein called 60S ribosomal protein L13, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
15	lO	204	Total	C	N	O	S	0	0
			1616	1030	302	275	9		

- Molecule 16 is a protein called 60S ribosomal protein L14, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
16	lP	130	Total	C	N	O	S	0	0
			1021	654	188	175	4		

- Molecule 17 is a protein called Ribosomal protein L15.

Mol	Chain	Residues	Atoms					AltConf	Trace
17	lQ	204	Total	C	N	O	S	0	0
			1676	1051	356	264	5		

- Molecule 18 is a protein called 60S ribosomal protein L17, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
18	lR	158	Total	C	N	O	S	0	0
			1232	779	238	210	5		

- Molecule 19 is a protein called 60S ribosomal protein L18, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
19	lS	167	Total	C	N	O	S	0	0
			1321	835	258	219	9		

- Molecule 20 is a protein called 60S ribosomal protein L18a.

Mol	Chain	Residues	Atoms					AltConf	Trace
20	lT	173	Total	C	N	O	S	0	0
			1413	910	259	235	9		

- Molecule 21 is a protein called Ribosomal protein L19.

Mol	Chain	Residues	Atoms					AltConf	Trace
21	IU	150	Total	C	N	O	S	0	0
			1235	787	246	197	5		

- Molecule 22 is a protein called 60S ribosomal protein L21, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
22	IV	165	Total	C	N	O	S	0	0
			1320	846	254	217	3		

- Molecule 23 is a protein called Large ribosomal subunit protein eL22.

Mol	Chain	Residues	Atoms					AltConf	Trace
23	IW	93	Total	C	N	O	S	0	0
			763	493	132	133	5		

- Molecule 24 is a protein called 60S ribosomal protein L23, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
24	IX	133	Total	C	N	O	S	0	0
			1015	629	196	182	8		

- Molecule 25 is a protein called Ribosomal protein L23A, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
25	IY	116	Total	C	N	O	S	0	0
			926	597	166	159	4		

- Molecule 26 is a protein called 60S ribosomal protein L24, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
26	IZ	57	Total	C	N	O	S	0	0
			481	318	88	73	2		

- Molecule 27 is a protein called 60S ribosomal protein L26, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
27	la	210	Total	C	N	O	S	0	0
			1651	1055	304	285	7		

- Molecule 28 is a protein called 60S ribosomal protein L27, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
28	lb	137	Total	C	N	O	S	0	0
			1094	707	196	187	4		

- Molecule 29 is a protein called Large ribosomal subunit protein uL15A.

Mol	Chain	Residues	Atoms					AltConf	Trace
29	lc	148	Total	C	N	O	S	0	0
			1192	757	236	194	5		

- Molecule 30 is a protein called 60S ribosomal protein L29.

Mol	Chain	Residues	Atoms					AltConf	Trace
30	ld	60	Total	C	N	O	S	0	0
			478	297	97	82	2		

- Molecule 31 is a protein called 60S ribosomal protein L30, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
31	le	93	Total	C	N	O	S	0	0
			693	438	118	135	2		

- Molecule 32 is a protein called 60S ribosomal protein L31, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
32	lf	126	Total	C	N	O	S	0	0
			1032	662	191	174	5		

- Molecule 33 is a protein called 60S ribosomal protein L32, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
33	lg	123	Total	C	N	O	S	0	0
			1010	643	200	162	5		

- Molecule 34 is a protein called 60S ribosomal protein L34, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
34	lh	102	Total	C	N	O	S	0	0
			805	503	166	130	6		

- Molecule 35 is a protein called uL29.

Mol	Chain	Residues	Atoms					AltConf	Trace
35	li	122	Total	C	N	O	S	0	0
			974	620	188	162	4		

- Molecule 36 is a protein called 60S ribosomal protein L35a, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
36	lj	106	Total	C	N	O	S	0	0
			841	545	158	135	3		

- Molecule 37 is a protein called 60S ribosomal protein L36, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
37	lk	89	Total	C	N	O	S	0	0
			712	447	144	116	5		

- Molecule 38 is a protein called 60S ribosomal protein L37-A, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
38	ll	72	Total	C	N	O	S	0	0
			591	361	132	91	7		

- Molecule 39 is a protein called 60S ribosomal protein L37a, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
39	lm	90	Total	C	N	O	S	0	0
			693	432	136	119	6		

- Molecule 40 is a protein called 60S ribosomal protein L38 putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
40	ln	73	Total	C	N	O	S	0	0
			584	378	104	100	2		

- Molecule 41 is a protein called Ribosomal protein L39, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
41	lo	50	Total	C	N	O	S	0	0
			432	275	91	63	3		

- Molecule 42 is a protein called 60S ribosomal protein L40, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
42	lp	53	Total	C	N	O	S	0	0
			420	259	86	69	6		

- Molecule 43 is a protein called 60S ribosomal protein L44, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
43	lq	92	Total	C	N	O	S	0	0
			756	480	148	122	6		

- Molecule 44 is a protein called 40S ribosomal protein S25.

Mol	Chain	Residues	Atoms					AltConf	Trace
44	sA	10	Total	C	N	O		0	0
			91	62	15	14			

- Molecule 45 is a protein called 40S ribosomal protein S26.

Mol	Chain	Residues	Atoms					AltConf	Trace
45	sB	98	Total	C	N	O	S	0	0
			787	478	169	134	6		

- Molecule 46 is a protein called Small ribosomal subunit protein eS27.

Mol	Chain	Residues	Atoms					AltConf	Trace
46	sC	32	Total	C	N	O	S	0	0
			249	161	42	45	1		

- Molecule 47 is a protein called 40S ribosomal protein S28, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
47	sD	60	Total	C	N	O	S	0	0
			468	289	93	84	2		

- Molecule 48 is a protein called Ribosomal protein S29, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
48	sE	55	Total	C	N	O	S	0	0
			442	273	90	75	4		

- Molecule 49 is a RNA chain called A/P-tRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
49	sI	73	Total	C	N	O	P	0	0
			1556	695	283	506	72		

- Molecule 50 is a RNA chain called P/E-tRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
50	sJ	73	Total	C	N	O	P	0	0
			1549	691	271	514	73		

- Molecule 51 is a RNA chain called mRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
51	sK	10	Total	C	N	O	P	0	0
			215	97	41	67	10		

- Molecule 52 is a RNA chain called 17S rRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
52	sa	1401	Total	C	N	O	P	0	0
			29968	13412	5443	9712	1401		

- Molecule 53 is a protein called Small ribosomal subunit protein uS5.

Mol	Chain	Residues	Atoms					AltConf	Trace
53	sc	196	Total	C	N	O	S	0	0
			1499	960	267	265	7		

- Molecule 54 is a protein called 40S ribosomal protein S3.

Mol	Chain	Residues	Atoms					AltConf	Trace
54	sd	188	Total	C	N	O	S	0	0
			1440	912	257	260	11		

- Molecule 55 is a protein called Small ribosomal subunit protein eS1.

Mol	Chain	Residues	Atoms					AltConf	Trace
55	se	144	Total	C	N	O	S	0	0
			1166	746	208	205	7		

- Molecule 56 is a protein called 40S ribosomal protein S4, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
56	sf	256	Total	C	N	O	S	0	0
			2031	1297	378	345	11		

- Molecule 57 is a protein called Small ribosomal subunit protein uS7.

Mol	Chain	Residues	Atoms					AltConf	Trace
57	sg	179	Total	C	N	O	S	0	0
			1424	897	258	258	11		

- Molecule 58 is a protein called 40S ribosomal protein S6.

Mol	Chain	Residues	Atoms					AltConf	Trace
58	sh	52	Total	C	N	O	S	0	0
			400	243	83	70	4		

- Molecule 59 is a protein called 40S ribosomal protein S7.

Mol	Chain	Residues	Atoms					AltConf	Trace
59	si	57	Total	C	N	O	S	0	0
			445	285	82	77	1		

- Molecule 60 is a protein called 40S ribosomal protein S8.

Mol	Chain	Residues	Atoms					AltConf	Trace
60	sj	129	Total	C	N	O	S	0	0
			1009	623	203	179	4		

- Molecule 61 is a protein called Small ribosomal subunit protein uS4.

Mol	Chain	Residues	Atoms					AltConf	Trace
61	sk	82	Total	C	N	O	S	0	0
			677	430	132	110	5		

- Molecule 62 is a protein called 40S ribosomal protein S10, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
62	sl	66	Total	C	N	O	S	0	0
			536	352	90	85	9		

- Molecule 63 is a protein called 40S ribosomal protein S11, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
63	sm	141	Total	C	N	O	S	0	0
			1161	735	224	196	6		

- Molecule 64 is a protein called 40S ribosomal protein S13, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
64	so	76	Total	C	N	O	S	0	0
			644	410	123	108	3		

- Molecule 65 is a protein called Ribosomal protein S14, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
65	sp	128	Total	C	N	O	S	0	0
			964	592	187	179	6		

- Molecule 66 is a protein called 40S ribosomal protein S15, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
66	sq	105	Total	C	N	O	S	0	0
			842	543	150	144	5		

- Molecule 67 is a protein called 40S ribosomal protein S15a, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
67	sr	129	Total	C	N	O	S	0	0
			1022	650	186	181	5		

- Molecule 68 is a protein called 40S ribosomal protein S16, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
68	ss	137	Total	C	N	O	S	0	0
			1076	695	193	184	4		

- Molecule 69 is a protein called 40S ribosomal protein S17, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
69	st	81	Total	C	N	O	S	0	0
			676	424	135	116	1		

- Molecule 70 is a protein called Small ribosomal subunit protein uS13.

Mol	Chain	Residues	Atoms					AltConf	Trace
70	su	124	Total	C	N	O	S	0	0
			1006	625	207	170	4		

- Molecule 71 is a protein called Small ribosomal subunit protein eS19.

Mol	Chain	Residues	Atoms					AltConf	Trace
71	sv	127	Total	C	N	O	S	0	0
			1015	646	185	178	6		

- Molecule 72 is a protein called 40S ribosomal protein S20, putative.

Mol	Chain	Residues	Atoms					AltConf	Trace
72	sw	46	Total	C	N	O	S	0	0
			376	238	70	65	3		

- Molecule 73 is a protein called 40S ribosomal protein S21.

Mol	Chain	Residues	Atoms					AltConf	Trace
73	sx	63	Total	C	N	O	S	0	0
			494	313	90	88	3		

- Molecule 74 is a protein called 40S ribosomal protein S23, putative.

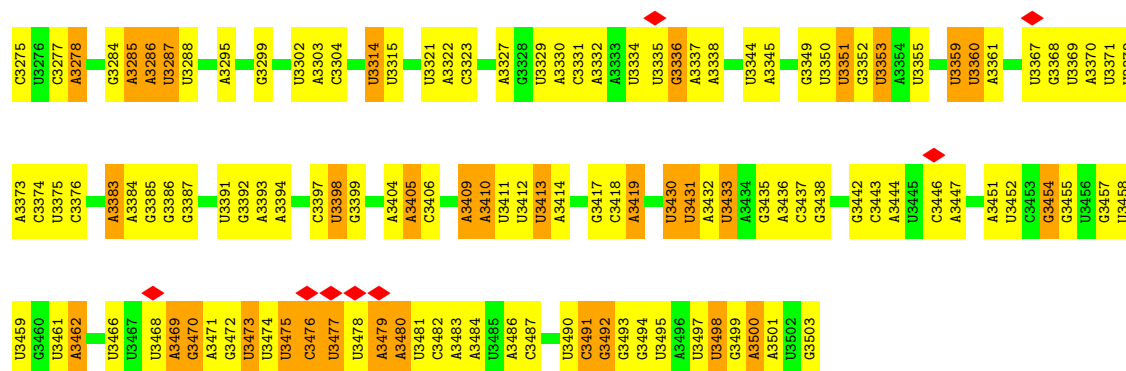
Mol	Chain	Residues	Atoms					AltConf	Trace
74	sy	106	Total	C	N	O	S	0	0
			836	522	169	142	3		

- Molecule 75 is PAROMOMYCIN (CCD ID: PAR) (formula: $C_{23}H_{45}N_5O_{14}$) (labeled as "Ligand of Interest" by depositor).



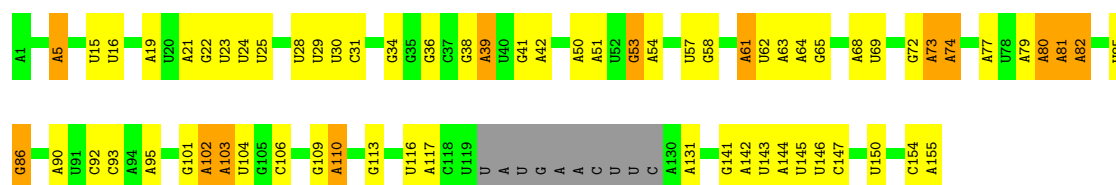


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G3163	U2970	U3074	U2877	A2896	U	A	G2467	G2379	A2274	A2189	A2083	U
G3166	U2971	A3076	U2878	U2699	U	C	G2468	A2379	U2281	A2190	G2084	A
A3170	U2972	A3077	U2790	U2702	C	U	G2470	G2381	U2285	G2193	A2089	C
A3171	U2973	A3078	U2791	G2705	A	G	A2473	G2382	G2286	U2194	G2095	A
A3175	A2981	U3084	C2795	A2706	U	C	A2478	G2383	G2297	U2195		U
G3176	U2884	U3085	C2796	A2707	U	C	G2479	G2384	A2298	A2196		A
	C2985		G2798	A2708	U	A	A2480	G2385	A2299	A2197		A
C3179	U2986	U3088	U2799	A2709	A	C	G2481	G2386	A2300	G2198		A
G3185	U2987	A3089	U2802	A2710	U	C	C2482	G2387	U2301	G2199		A
	A2988	U3090	U2803	U2711	U	G	G2485	G2388	U2302	G2200		A
U3196	U2989	U3091	U2804	A2712	U	G	U2486	G2389	A2303	G2201		A
C3197	U2990	U3092	G2815	A2713	U	G	U2487	G2390	A2304	G2202		A
	G2991	U3093	C2816	C2714	U	A	G2488	A2391	U2305	G2203		A
G3200	C2992	U3094	A2818	A2715	U	A	G2489	G2392	U2306	A2111		A
	G2993	U3095	U2819	A2716	U	A	U2490	G2393	A2307	A2112		U
A3203	U2994	A3096	U2820	U2717	U	C	C2491	G2394	U2308	A2113		U
C3204	G2995	A3097	A2821	G2721	U	C	G2492	G2395	A2309	A2114		A
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A3211	G3006	G3104	A2823	A2723	A	A	U2495	G2397	A2311	A2116		A
U3212	A3007	U3105	C2824	A2724	U	A	A2500	G2398	A2312	G2117		A
U3213	U3008	G3106	U2825	G2725	U	C	U2501	G2399	A2313	A2118		A
C3214	C3009	G3107	A2826	G2726	U	C	U2502	G2400	A2314	A2119		A
A3215	U3010	A3110	U2827	A2727	U	C	U2503	G2401	A2315	A2120		A
A3216	U3011	U3111	U2828	G2728	U	C	U2504	G2402	A2316	A2121		A
U3220	C3012	G3112	A2829	A2729	U	C	U2505	G2403	A2317	A2122		A
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G3232	U3017	U3117	U2652	G2734	U	C	U2511	G2408	A2322	A2127		A
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C3243	G3024	G3132	G2656	G2738	U	C	U2515	G2412	A2326	A2131		A
	U3025	U3133	U2657	A2739	U	C	U2516	G2413	A2327	A2132		A
A3246	A3030	U3134	A2658	G2740	U	C	U2517	G2414	A2328	A2133		A
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A3270	U3048	A3153	U2671	A2753	U	C	U2530	G2427	A2341	A2146		A
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			A2676	G2758	U	C	U2535	G2432	A2346	A2151		A
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			A2680	G2762	U	C	U2539	A2436	A2350	A2155		A
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			A2687	A2769	U	C	U2546	A2443	A2357	A2162		A
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			A2872	G2772	U	C	U2549	A2446	A2360	A2165		A
			C2873	A2773	U	C	U2550	A2447	A2361	A2166		A
				G2774	U	C	U2551	A2448	A2362	A2167		A
					U	C	U2552	A2449	A2363	A2168		A
					G	C	U2553	A2450	A2364	A2169		A
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					A	C	U2557	A2454	A2368	A2173		A
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					G	C	U2559	A2456	A2370	A2175		A
					A	C	U2560	A2457	A2371	A2176		A
					U	C	U2561	A2458	A2372	A2177		A
					G	C	U2562	A2459	A2373	A2178		A
					A	C	U2563	A2460	A2374	A2179		A
					U	C	U2564	A2461	A2375	A2180		A
					G	C	U2565	A2462	A2376	A2181		A
					A	C	U2566	A2463	A2377	A2182		A
					U	C	U2567	A2464	A2378	A2183		A
					G	C	U2568	A2465	A2379	A2184		A



• Molecule 2: 5.8S rRNA

Chain IB: 50% 35% 8% 6%



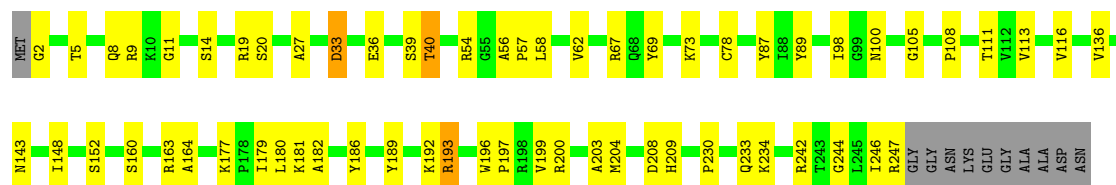
• Molecule 3: 5S rRNA

Chain IC: 41% 51% 8%



• Molecule 4: Large ribosomal subunit protein uL2

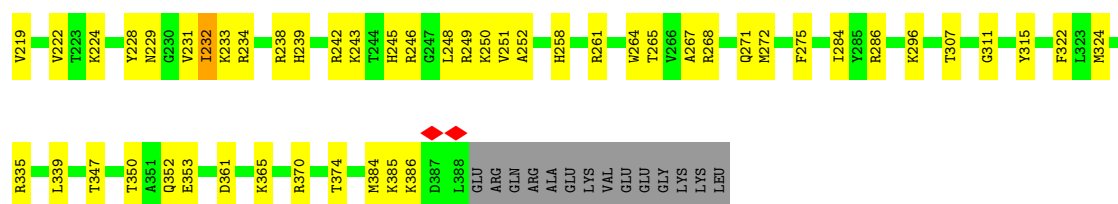
Chain ID: 72% 23%



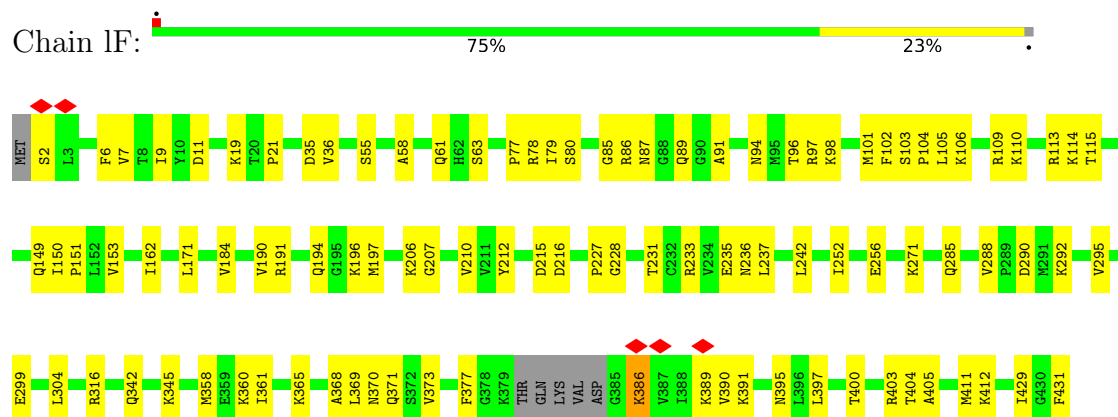
• Molecule 5: 60S ribosomal protein L3, putative

Chain IE: 76% 19%

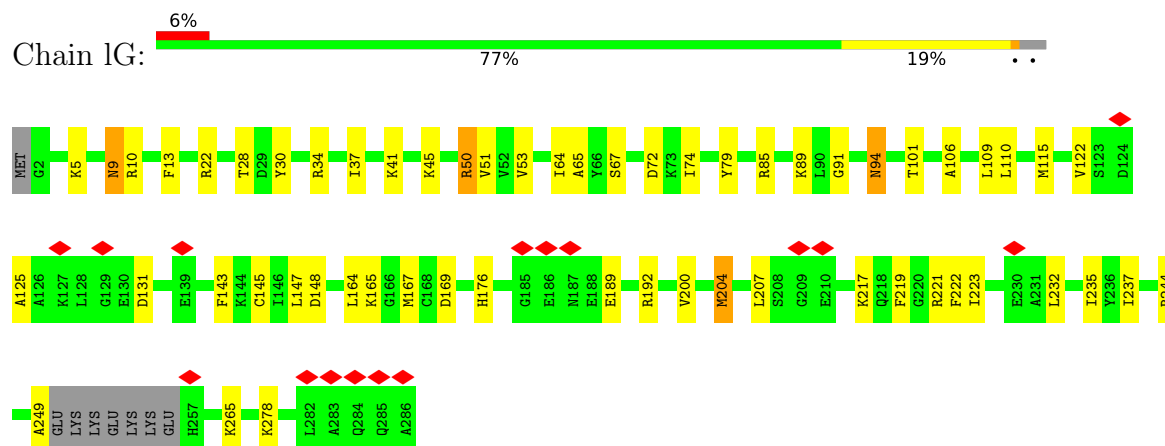




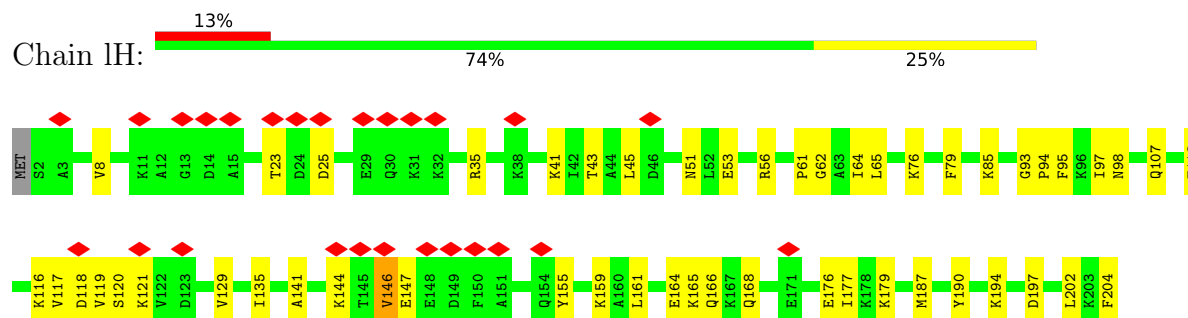
- Molecule 6: 60S ribosomal protein L4, putative



- Molecule 7: 60S ribosomal protein L5, putative

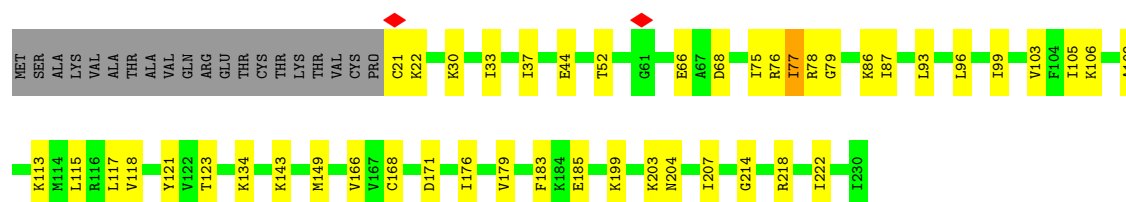


- Molecule 8: Large ribosomal subunit protein eL6

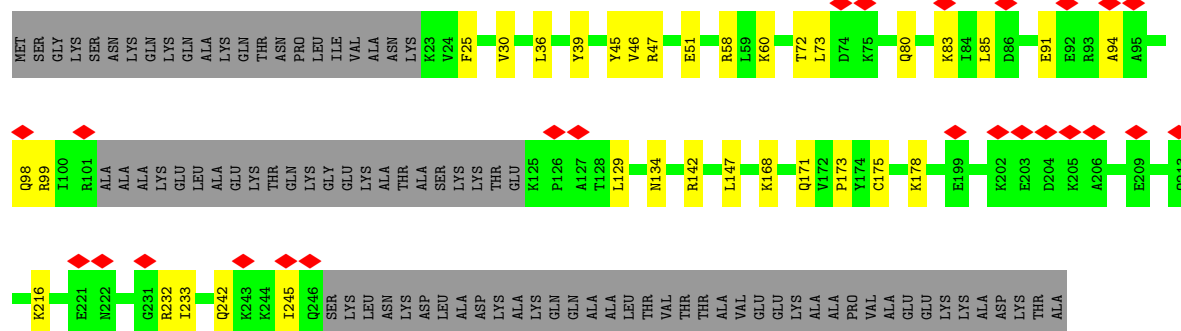


- Molecule 9: 60S ribosomal protein L7, putative

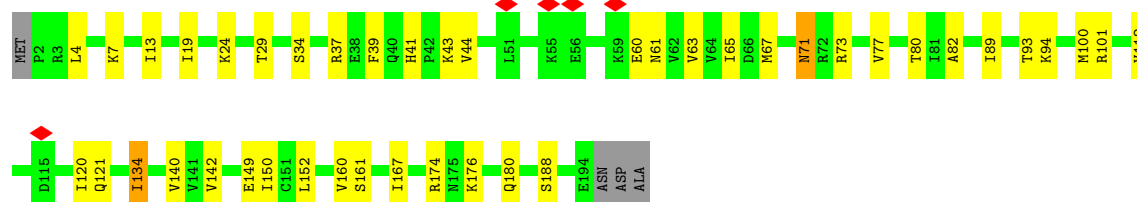
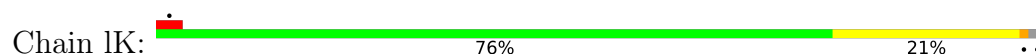




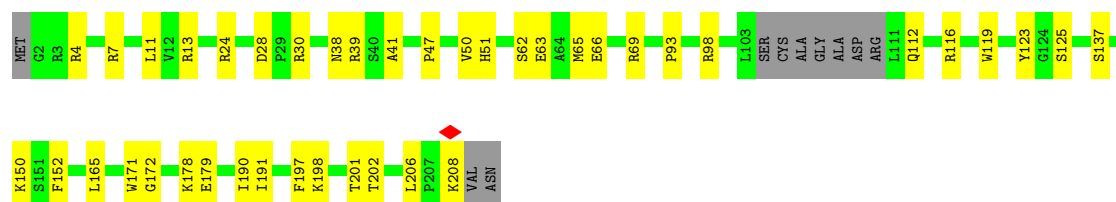
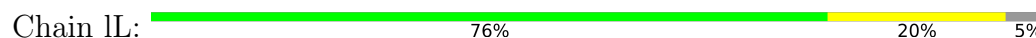
• Molecule 10: 60S ribosomal protein L7a



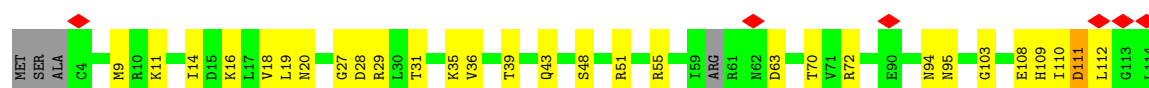
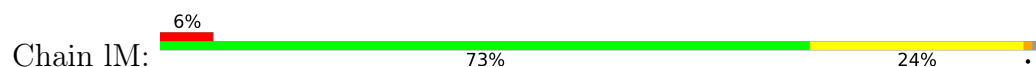
• Molecule 11: 60S ribosomal protein L9, putative



• Molecule 12: Ribosomal protein L10, putative

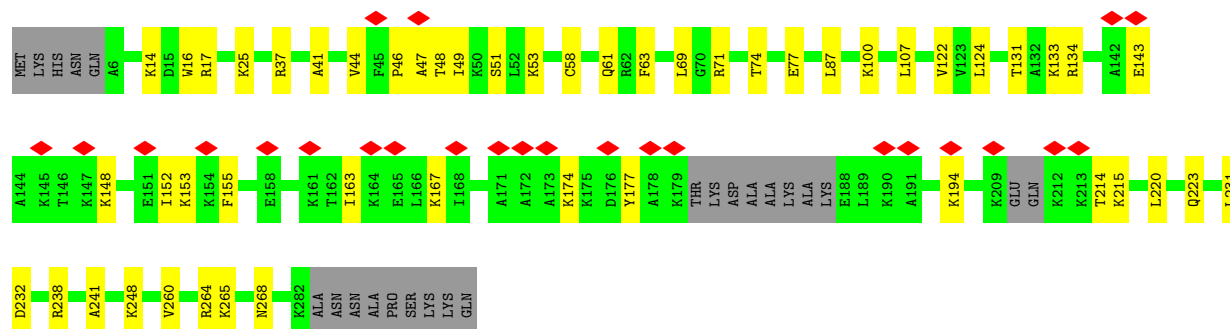
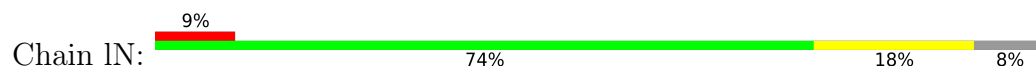


• Molecule 13: 60S ribosomal protein L11, putative

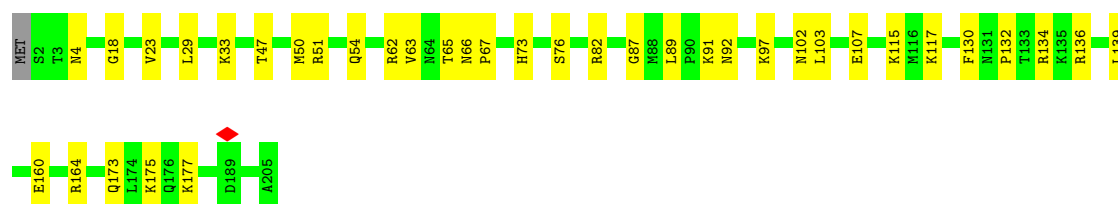
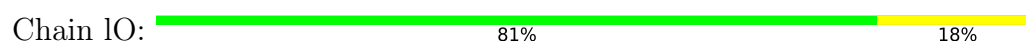




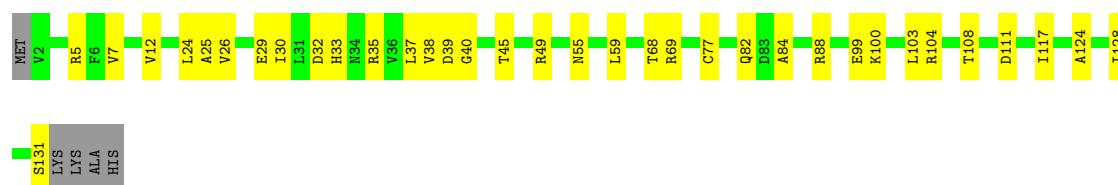
- Molecule 14: 60S ribosomal protein L13, putative



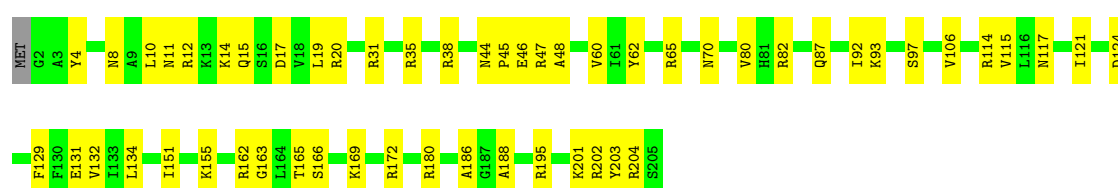
- Molecule 15: 60S ribosomal protein L13, putative



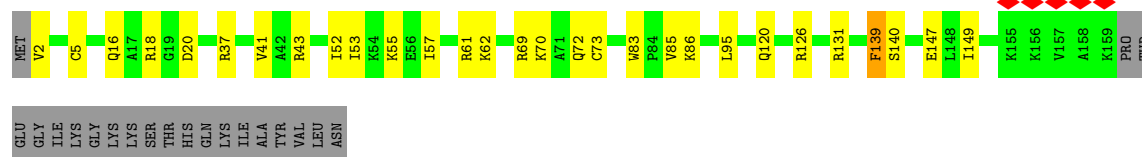
- Molecule 16: 60S ribosomal protein L14, putative



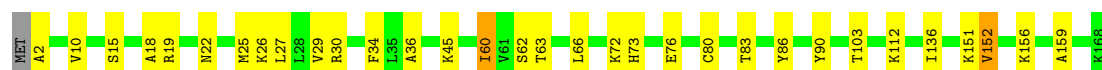
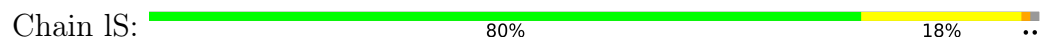
- Molecule 17: Ribosomal protein L15



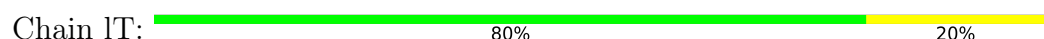
- Molecule 18: 60S ribosomal protein L17, putative



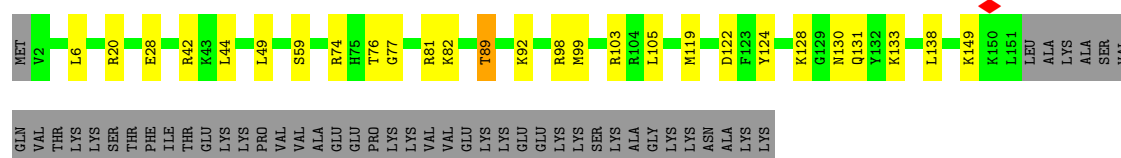
- Molecule 19: 60S ribosomal protein L18, putative



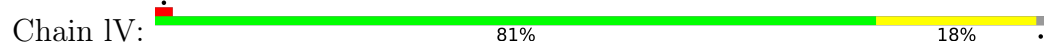
- Molecule 20: 60S ribosomal protein L18a



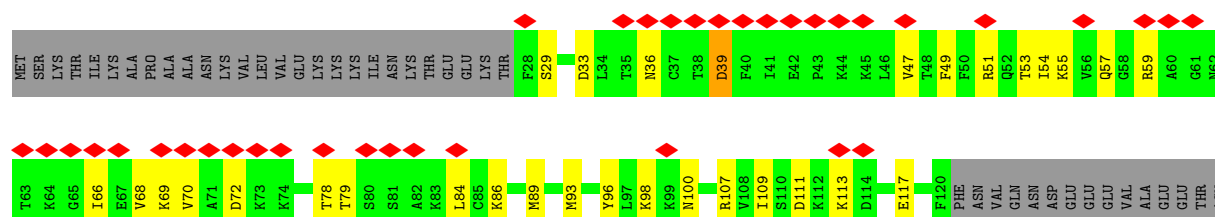
- Molecule 21: Ribosomal protein L19

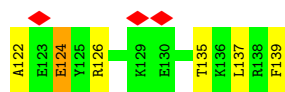


- Molecule 22: 60S ribosomal protein L21, putative

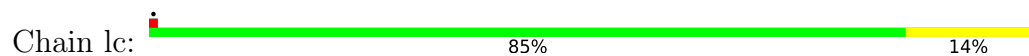


- Molecule 23: Large ribosomal subunit protein eL22





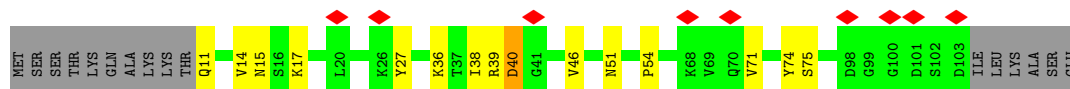
- Molecule 29: Large ribosomal subunit protein uL15A



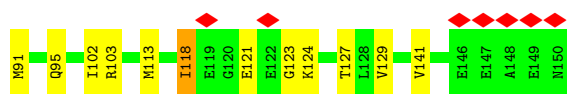
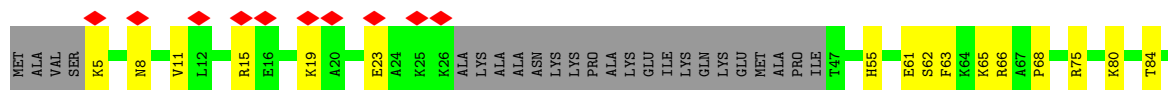
- Molecule 30: 60S ribosomal protein L29



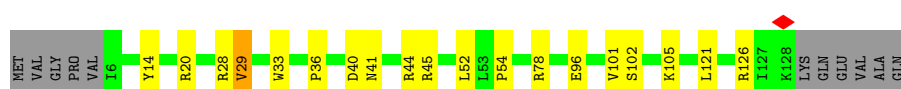
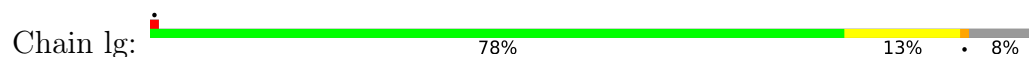
- Molecule 31: 60S ribosomal protein L30, putative



- Molecule 32: 60S ribosomal protein L31, putative

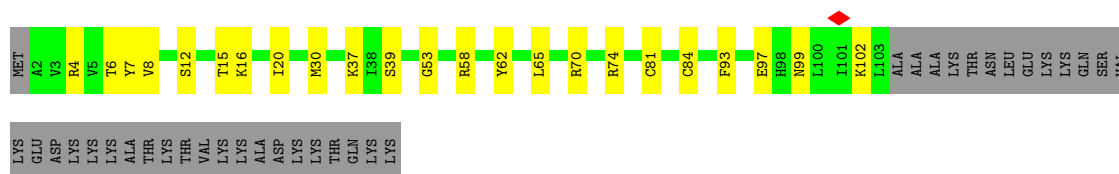


- Molecule 33: 60S ribosomal protein L32, putative

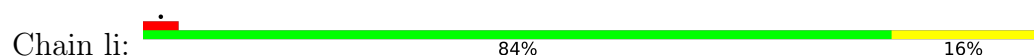


- Molecule 34: 60S ribosomal protein L34, putative

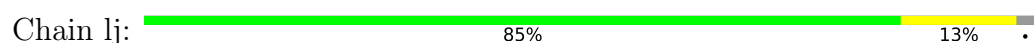




• Molecule 35: uL29



• Molecule 36: 60S ribosomal protein L35a, putative



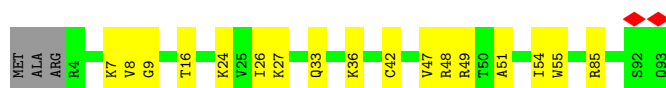
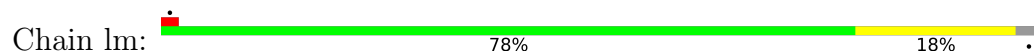
• Molecule 37: 60S ribosomal protein L36, putative



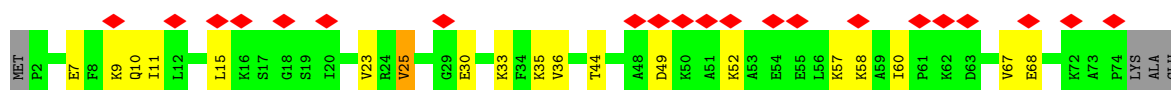
• Molecule 38: 60S ribosomal protein L37-A, putative

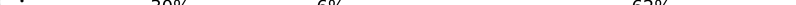


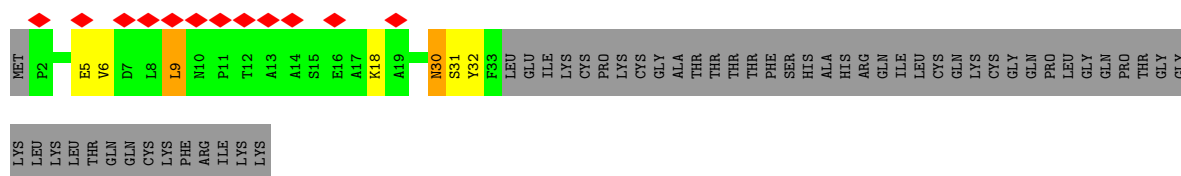
• Molecule 39: 60S ribosomal protein L37a, putative



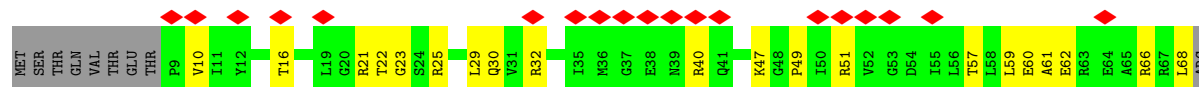
• Molecule 40: 60S ribosomal protein L38 putative



- Chain sC: 



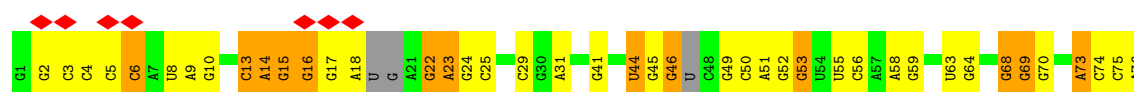
- Molecule 47: 40S ribosomal protein S28, putative



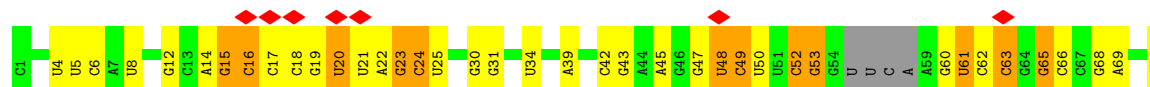
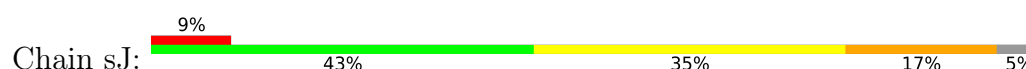
- Molecule 48: Ribosomal protein S29, putative



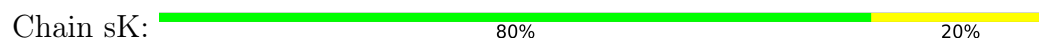
- Molecule 49: A/P-tRNA



- Molecule 50: P/E-tRNA



- Molecule 51: mRNA

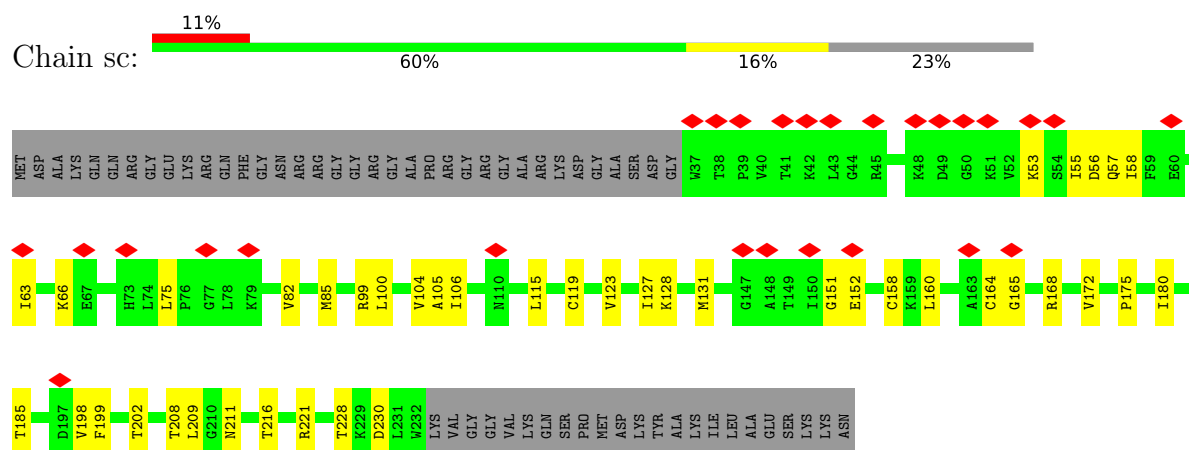


- Molecule 52: 17S rRNA

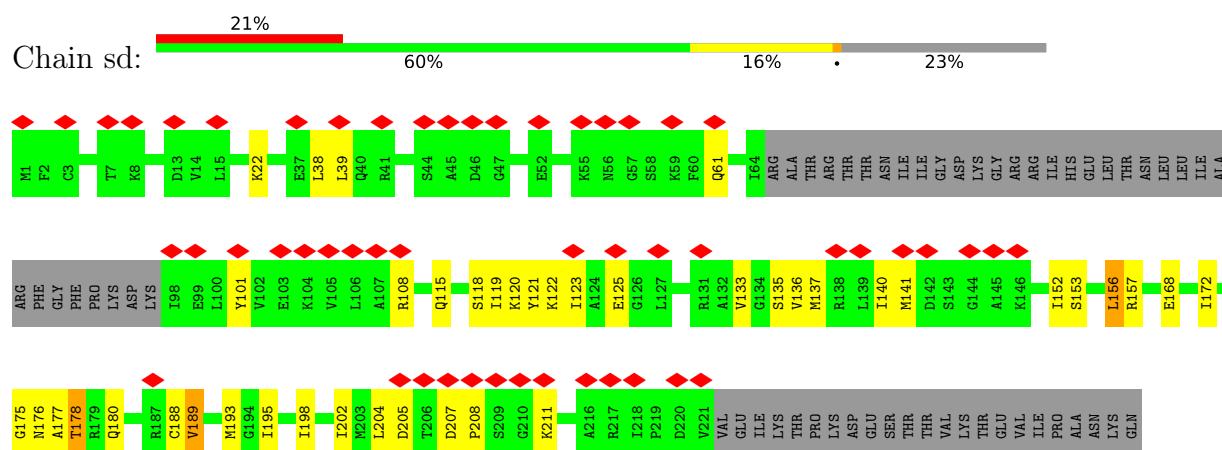




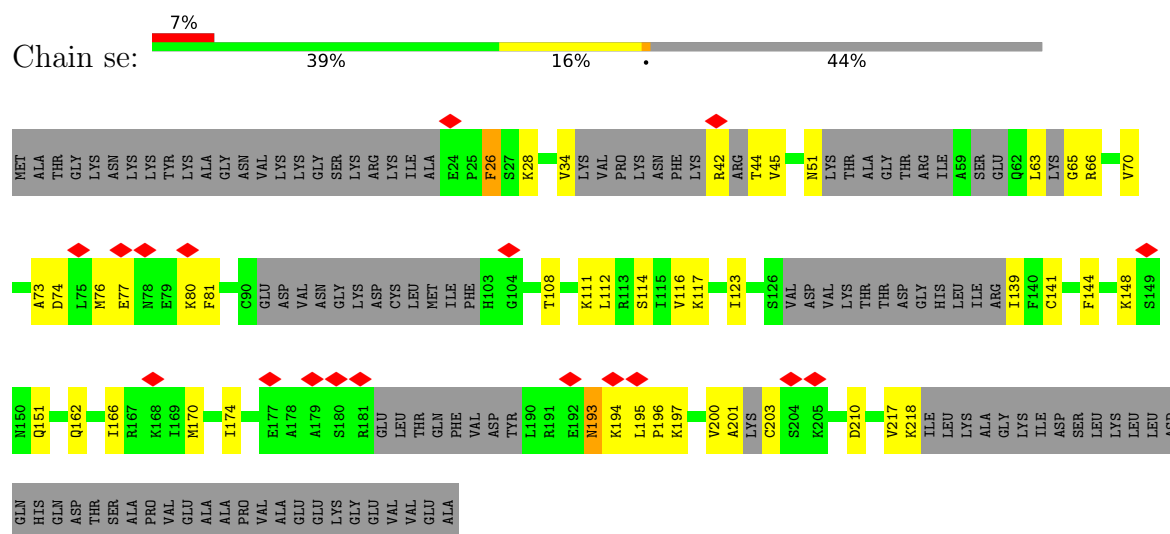




- Molecule 54: 40S ribosomal protein S3



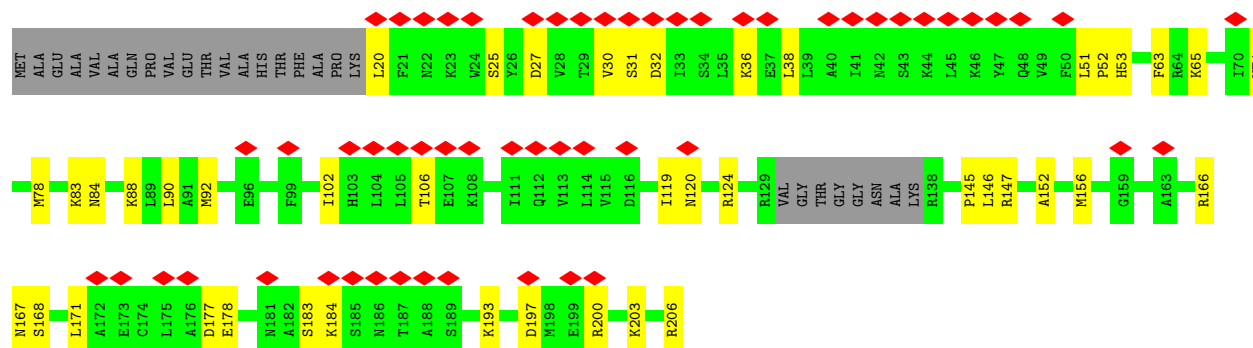
- Molecule 55: Small ribosomal subunit protein eS1



- Molecule 56: 40S ribosomal protein S4, putative

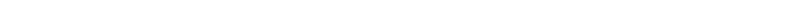


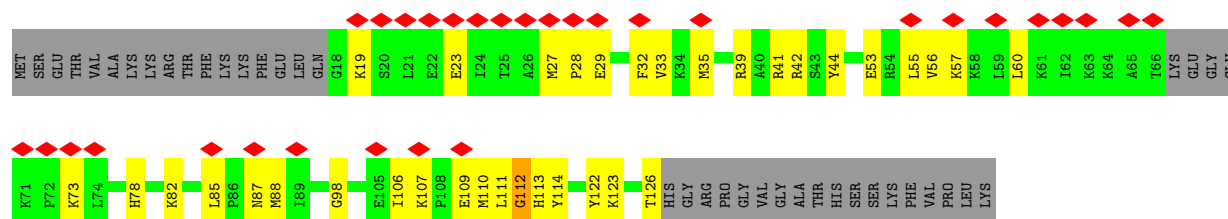
- Molecule 57: Small ribosomal subunit protein uS7



- Molecule 58: 40S ribosomal protein S6



- Chain sq: 



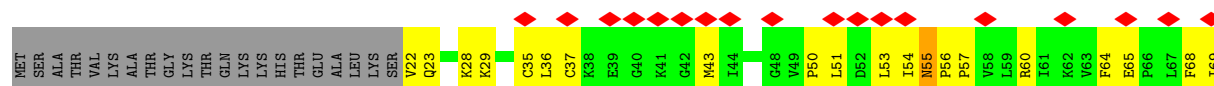
- Molecule 67: 40S ribosomal protein S15a, putative

Chain sr: 74% 25% ..



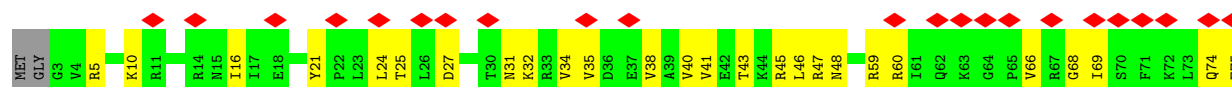
- Molecule 68: 40S ribosomal protein S16, putative

Chain ss: 29% 63% 23% 13%



- Molecule 69: 40S ribosomal protein S17, putative

Chain st: 26% 44% 25% 31%



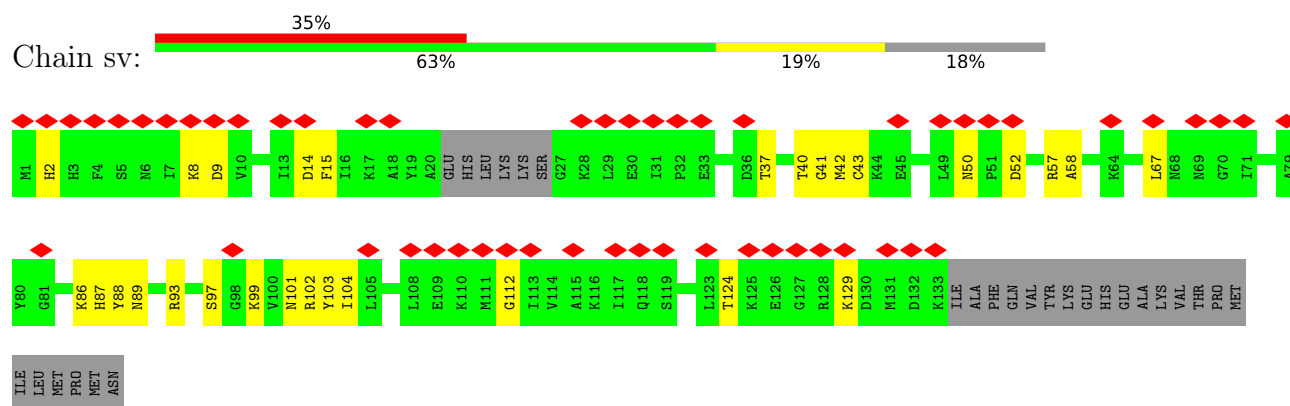
- Molecule 70: Small ribosomal subunit protein uS13

Chain su: 17% 56% 24% 20%

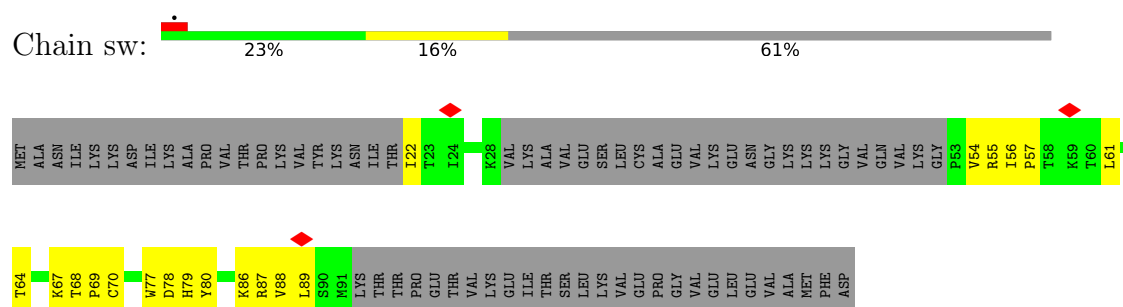


ALA

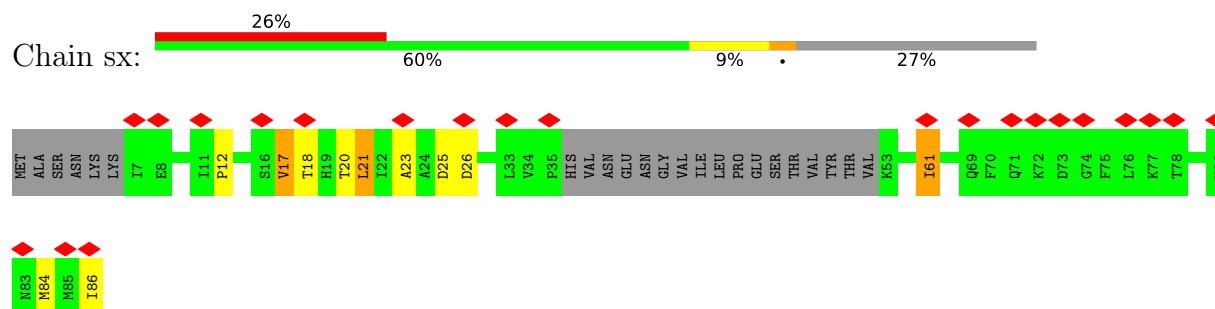
• Molecule 71: Small ribosomal subunit protein eS19



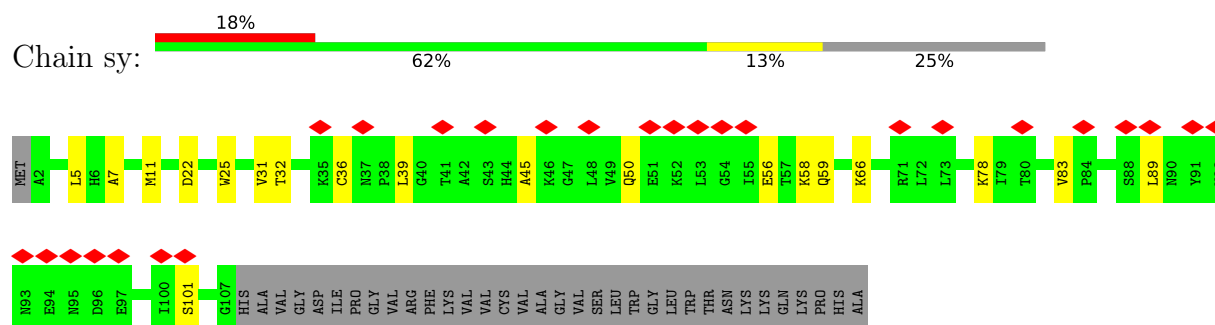
• Molecule 72: 40S ribosomal protein S20, putative



• Molecule 73: 40S ribosomal protein S21



• Molecule 74: 40S ribosomal protein S23, putative



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, Not provided	
Number of particles used	54889	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	TFS KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	1.09	Depositor
Minimum defocus (nm)	1800	Depositor
Maximum defocus (nm)	3000	Depositor
Magnification	75000	Depositor
Image detector	FEI FALCON III (4k x 4k)	Depositor
Maximum map value	23.075	Depositor
Minimum map value	-9.423	Depositor
Average map value	-0.000	Depositor
Map value standard deviation	1.000	Depositor
Recommended contour level	2.8	Depositor
Map size ($\text{\AA}$)	428.00003, 428.00003, 428.00003	wwPDB
Map dimensions	400, 400, 400	wwPDB
Map angles ($^\circ$)	90.0, 90.0, 90.0	wwPDB
Pixel spacing ($\text{\AA}$)	1.07, 1.07, 1.07	Depositor

5 Model quality

5.1 Standard geometry

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

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	1A	0.25	0/75250	0.33	2/117222 (0.0%)
2	1B	0.24	0/3470	0.32	0/5401
3	1C	0.25	0/2765	0.38	2/4303 (0.0%)
4	1D	0.25	0/1920	0.32	0/2582
5	1E	0.23	0/3140	0.30	0/4216
6	1F	0.22	0/3339	0.32	0/4479
7	1G	0.19	0/2248	0.28	0/3013
8	1H	0.17	0/1640	0.33	0/2204
9	1I	0.22	0/1680	0.29	0/2252
10	1J	0.17	0/1670	0.29	0/2243
11	1K	0.19	0/1562	0.27	0/2103
12	1L	0.20	0/1633	0.30	0/2184
13	1M	0.15	0/1369	0.24	0/1834
14	1N	0.19	0/2158	0.28	0/2875
15	1O	0.23	0/1646	0.30	0/2209
16	1P	0.21	0/1033	0.28	0/1389
17	1Q	0.26	0/1707	0.29	0/2276
18	1R	0.25	0/1251	0.29	0/1675
19	1S	0.24	0/1342	0.32	0/1796
20	1T	0.22	0/1445	0.29	0/1946
21	1U	0.21	0/1253	0.28	0/1666
22	1V	0.22	0/1351	0.30	0/1819
23	1W	0.14	0/774	0.37	0/1031
24	1X	0.23	0/1030	0.33	0/1384
25	1Y	0.19	0/941	0.25	0/1262
26	1Z	0.20	0/492	0.28	0/656
27	1a	0.18	0/1673	0.26	0/2236
28	1b	0.17	0/1112	0.26	0/1489
29	1c	0.26	0/1223	0.29	0/1636
30	1d	0.22	0/485	0.25	0/639
31	1e	0.20	0/701	0.29	0/945
32	1f	0.22	0/1050	0.27	0/1402

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
33	lg	0.24	0/1027	0.28	0/1370
34	lh	0.22	0/818	0.29	0/1094
35	li	0.17	0/984	0.22	0/1310
36	lj	0.27	0/862	0.30	0/1163
37	lk	0.17	0/721	0.24	0/955
38	ll	0.27	0/602	0.35	0/797
39	lm	0.22	0/701	0.34	0/934
40	ln	0.18	0/592	0.29	0/789
41	lo	0.26	0/444	0.29	0/587
42	lp	0.23	0/425	0.48	1/563 (0.2%)
43	lq	0.23	0/770	0.27	0/1019
44	sA	0.14	0/94	0.44	0/128
45	sB	0.14	0/797	0.25	0/1062
46	sC	0.13	0/255	0.29	0/346
47	sD	0.15	0/470	0.30	0/630
48	sE	0.14	0/449	0.31	0/595
49	sI	0.16	0/1737	0.38	0/2702
50	sJ	0.17	0/1727	0.40	0/2684
51	sK	0.16	0/241	0.27	0/373
52	sa	0.15	0/33554	0.28	0/52257
53	sc	0.13	0/1528	0.24	0/2064
54	sd	0.11	0/1456	0.27	0/1949
55	se	0.14	0/1174	0.31	0/1555
56	sf	0.11	0/2072	0.28	0/2792
57	sg	0.11	0/1443	0.26	0/1938
58	sh	0.11	0/402	0.26	0/535
59	si	0.11	0/453	0.28	0/610
60	sj	0.12	0/1021	0.27	0/1366
61	sk	0.09	0/688	0.25	0/916
62	sl	0.11	0/549	0.26	0/740
63	sm	0.12	0/1187	0.25	0/1586
64	so	0.13	0/654	0.22	0/866
65	sp	0.14	0/977	0.28	0/1311
66	sq	0.12	0/856	0.33	0/1143
67	sr	0.15	0/1040	0.31	0/1404
68	ss	0.11	0/1093	0.25	0/1466
69	st	0.11	0/684	0.24	0/914
70	su	0.13	0/1020	0.33	0/1364
71	sv	0.11	0/1034	0.23	0/1388
72	sw	0.12	0/382	0.24	0/512
73	sx	0.13	0/502	0.28	0/676
74	sy	0.12	0/848	0.30	0/1131
All	All	0.21	0/192686	0.31	5/283951 (0.0%)

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

Mol	Chain	#Chirality outliers	#Planarity outliers
6	lF	0	1
38	ll	0	1
48	sE	0	1
67	sr	0	1
All	All	0	4

There are no bond length outliers.

All (5) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	lC	62	U	OP2-P-O3'	-8.61	82.16	108.00
3	lC	62	U	OP1-P-O3'	-8.53	82.42	108.00
42	lp	24	CYS	CA-CB-SG	7.16	130.87	114.40
1	lA	2468	C	P-O3'-C3'	-5.13	112.50	120.20
1	lA	2468	C	O3'-P-O5'	5.10	111.66	104.00

There are no chirality outliers.

All (4) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
6	lF	368	ALA	Peptide
38	ll	39	TYR	Peptide
48	sE	15	GLY	Peptide
67	sr	76	SER	Peptide

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	lA	67153	0	33687	1002	0
2	lB	3097	0	1552	49	0
3	lC	2477	0	1252	71	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
4	ID	1881	0	1928	53	0
5	IE	3076	0	3209	63	0
6	IF	3281	0	3513	75	0
7	IG	2209	0	2287	43	0
8	IH	1608	0	1728	33	0
9	II	1658	0	1802	31	0
10	IJ	1640	0	1749	20	0
11	IK	1538	0	1598	28	0
12	IL	1597	0	1654	29	0
13	IM	1350	0	1390	34	0
14	IN	2130	0	2338	41	0
15	IO	1616	0	1700	25	0
16	IP	1021	0	1107	25	0
17	IQ	1676	0	1777	45	0
18	IR	1232	0	1307	22	0
19	IS	1321	0	1427	25	0
20	IT	1413	0	1479	26	0
21	IU	1235	0	1369	25	0
22	IV	1320	0	1406	23	0
23	IW	763	0	818	20	0
24	IX	1015	0	1054	23	0
25	IY	926	0	997	16	0
26	IZ	481	0	518	11	0
27	la	1651	0	1822	30	0
28	lb	1094	0	1174	27	0
29	lc	1192	0	1205	18	0
30	ld	478	0	507	13	0
31	le	693	0	721	9	0
32	lf	1032	0	1095	20	0
33	lg	1010	0	1091	16	0
34	lh	805	0	849	19	0
35	li	974	0	1093	13	0
36	lj	841	0	878	7	0
37	lk	712	0	755	21	0
38	ll	591	0	617	16	0
39	lm	693	0	738	10	0
40	ln	584	0	643	13	0
41	lo	432	0	444	10	0
42	lp	420	0	450	16	0
43	lq	756	0	821	16	0
44	sA	91	0	88	2	0
45	sB	787	0	833	18	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
46	sC	249	0	259	6	0
47	sD	468	0	500	15	0
48	sE	442	0	444	15	0
49	sI	1556	0	795	18	0
50	sJ	1549	0	789	20	0
51	sK	215	0	108	2	0
52	sa	29968	0	15064	504	0
53	sc	1499	0	1565	23	0
54	sd	1440	0	1522	29	0
55	se	1166	0	1227	29	0
56	sf	2031	0	2145	63	0
57	sg	1424	0	1485	38	0
58	sh	400	0	415	12	0
59	si	445	0	466	10	0
60	sj	1009	0	1055	40	0
61	sk	677	0	724	14	0
62	sl	536	0	559	19	0
63	sm	1161	0	1181	28	0
64	so	644	0	664	15	0
65	sp	964	0	988	27	0
66	sq	842	0	913	30	0
67	sr	1022	0	1051	28	0
68	ss	1076	0	1151	29	0
69	st	676	0	718	19	0
70	su	1006	0	1049	28	0
71	sv	1015	0	1043	20	0
72	sw	376	0	405	16	0
73	sx	494	0	508	8	0
74	sy	836	0	885	14	0
75	sa	42	0	45	1	0
All	All	178778	0	130193	2752	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 9.

All (2752) close contacts within the same asymmetric unit are listed below, sorted by their clash magnitude.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:lA:805:G:H21	1:lA:1496:A:N6	1.45	1.13
1:lA:869:G:N2	1:lA:872:A:H62	1.48	1.11
52:sa:139:G:H1	52:sa:166:A:N6	1.48	1.11

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:lA:869:G:H21	1:lA:872:A:N6	1.52	1.07
3:lC:79:G:N1	3:lC:95:A:C2	2.24	1.03
48:sE:2:GLY:N	48:sE:5:SER:HG	1.57	1.01
3:lC:27:G:H1	3:lC:50:A:H61	1.04	1.00
3:lC:27:G:H1	3:lC:50:A:N6	1.60	0.99
1:lA:869:G:N2	1:lA:872:A:N6	2.08	0.99
1:lA:805:G:N2	1:lA:1496:A:H61	1.61	0.98
52:sa:211:U:H3	52:sa:247:G:H1	1.15	0.94
3:lC:79:G:N1	3:lC:95:A:H2	1.61	0.91
19:lS:73:HIS:HB3	19:lS:76:GLU:HG3	1.51	0.91
1:lA:3454:G:H1	1:lA:3501:A:H2	0.91	0.89
1:lA:2269:U:H5'	1:lA:2270:G:H5'	1.54	0.89
1:lA:805:G:H21	1:lA:1496:A:H61	0.94	0.88
1:lA:3454:G:N1	1:lA:3501:A:C2	2.41	0.88
1:lA:457:G:N2	18:lR:5:CYS:SG	2.47	0.87
1:lA:2214:A:HO2'	38:ll:2:THR:N	1.72	0.87
1:lA:805:G:N2	1:lA:1496:A:N6	2.19	0.87
6:lF:101:MET:HE3	6:lF:104:PRO:HA	1.57	0.87
1:lA:206:U:H3	1:lA:303:G:H1	0.88	0.87
1:lA:2366:C:O2	52:sa:1892:G:N2	2.07	0.86
3:lC:79:G:H1	3:lC:95:A:H2	0.88	0.85
1:lA:3454:G:N1	1:lA:3501:A:H2	1.74	0.85
3:lC:52:G:N2	3:lC:54:U:O4	2.09	0.85
1:lA:873:U:H2'	1:lA:874:A:H8	1.42	0.84
52:sa:1821:A:N6	52:sa:1892:G:O6	2.10	0.83
58:sh:63:MET:HE1	58:sh:106:ILE:HD12	1.62	0.81
1:lA:2473:A:N6	1:lA:3012:U:O2'	2.14	0.81
13:lM:18:VAL:HG22	13:lM:70:THR:HG22	1.61	0.81
1:lA:774:U:OP1	29:lc:21:ARG:NH2	2.13	0.81
3:lC:79:G:O6	3:lC:95:A:N1	2.13	0.81
1:lA:1605:A:H2	1:lA:3430:U:H3	1.27	0.81
50:sJ:15:G:H21	50:sJ:16:C:H42	1.29	0.80
1:lA:1434:G:HO2'	1:lA:2456:U:HO2'	1.25	0.80
36:lj:5:ARG:NH1	36:lj:7:HIS:O	2.15	0.80
1:lA:3479:A:H2'	1:lA:3480:A:O4'	1.82	0.79
1:lA:2416:U:O2'	1:lA:3246:A:N1	2.14	0.79
1:lA:409:G:OP2	38:ll:56:ARG:NH2	2.16	0.79
22:IV:80:VAL:HG12	22:IV:81:ASN:H	1.47	0.79
57:sg:53:HIS:HB2	57:sg:92:MET:HE1	1.63	0.79
73:sx:18:THR:HG23	73:sx:20:THR:HG23	1.64	0.79
1:lA:225:U:H3	1:lA:282:G:H1	1.28	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:3454:G:O6	1:1A:3501:A:N1	2.15	0.79
1:1A:394:A:N3	1:1A:398:A:O2'	2.16	0.78
52:sa:389:U:H3	52:sa:396:A:H62	1.32	0.78
3:1C:26:U:H3	3:1C:51:G:H22	1.31	0.77
3:1C:27:G:N2	3:1C:50:A:N1	2.32	0.77
1:1A:500:G:N2	1:1A:572:U:O2	2.18	0.76
52:sa:1245:C:H2'	52:sa:1246:A:H8	1.49	0.76
52:sa:734:A:N1	52:sa:803:G:O6	2.19	0.76
62:sl:31:HIS:HD2	62:sl:41:VAL:HG21	1.50	0.76
50:sJ:53:G:H1	50:sJ:63:C:H42	1.34	0.76
27:la:21:ALA:O	27:la:26:ARG:NH1	2.19	0.76
7:1G:165:LYS:NZ	7:1G:169:ASP:OD2	2.19	0.75
52:sa:43:U:OP2	52:sa:432:A:N6	2.19	0.75
1:1A:2340:U:O2'	1:1A:2341:C:O5'	2.03	0.75
1:1A:2414:C:OP1	5:1E:238:ARG:NH1	2.20	0.75
1:1A:2688:G:O2'	1:1A:3008:U:OP1	2.04	0.75
1:1A:3263:G:OP2	42:lp:37:LYS:NZ	2.18	0.75
1:1A:3040:A:H5''	42:lp:50:LYS:HG3	1.68	0.75
3:1C:18:U:H3	3:1C:58:A:H61	1.35	0.75
1:1A:1251:G:H21	12:1L:116:ARG:HH22	1.34	0.74
1:1A:1177:A:H2'	1:1A:1178:A:C8	2.22	0.74
1:1A:1818:C:OP2	34:lh:74:ARG:NH2	2.20	0.74
1:1A:1030:C:OP2	4:1D:9:ARG:NH1	2.20	0.74
8:1H:76:LYS:O	8:1H:98:ASN:ND2	2.21	0.74
1:1A:2075:A:OP2	21:1U:20:ARG:NH1	2.20	0.74
3:1C:10:A:N6	7:1G:13:PHE:O	2.21	0.74
17:1Q:31:ARG:NH1	17:1Q:124:ASP:OD2	2.21	0.74
1:1A:1512:G:OP2	6:1F:191:ARG:NH1	2.21	0.73
1:1A:2705:A:H4'	1:1A:2706:C:H5'	1.68	0.73
1:1A:1033:A:N1	4:1D:204:MET:HE2	2.03	0.73
1:1A:1282:G:OP1	9:1I:78:ARG:NH1	2.22	0.73
1:1A:875:C:H2'	1:1A:876:A:H8	1.53	0.73
62:sl:3:ILE:HG12	62:sl:40:GLN:HB3	1.70	0.73
1:1A:3437:C:N3	18:1R:69:ARG:NH2	2.35	0.73
1:1A:3479:A:H3'	1:1A:3479:A:N3	2.03	0.73
34:lh:39:SER:OG	34:lh:58:ARG:NH1	2.21	0.73
52:sa:887:A:H2'	52:sa:888:A:H8	1.53	0.73
1:1A:737:A:H3'	1:1A:738:G:H21	1.53	0.72
61:sk:7:ASN:O	61:sk:8:HIS:ND1	2.22	0.72
35:li:76:THR:O	35:li:81:ARG:NH1	2.22	0.72
60:sj:67:TRP:HE1	60:sj:69:SER:HB3	1.53	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:lA:2629:A:H8	31:le:54:PRO:HB3	1.52	0.72
52:sa:783:A:N7	52:sa:784:U:O2'	2.20	0.72
59:si:199:VAL:HG23	59:si:200:VAL:HG23	1.71	0.72
52:sa:139:G:N2	52:sa:166:A:N1	2.34	0.72
1:lA:919:G:OP1	14:lN:17:ARG:NH2	2.22	0.72
7:lG:34:ARG:NH1	22:lV:26:ASN:O	2.22	0.72
52:sa:1350:A:H2'	52:sa:1351:A:H8	1.53	0.72
52:sa:1839:A:OP1	58:sh:94:ARG:NH2	2.23	0.71
37:lk:2:ALA:O	37:lk:4:GLY:N	2.22	0.71
72:sw:68:THR:OG1	72:sw:70:CYS:O	2.07	0.71
1:lA:500:G:H1	1:lA:572:U:H3	0.75	0.71
1:lA:873:U:H2'	1:lA:874:A:C8	2.24	0.71
1:lA:1086:G:H2'	1:lA:1087:G:H8	1.56	0.71
1:lA:1086:G:H1	1:lA:1240:C:H5	1.38	0.71
52:sa:103:U:OP2	52:sa:305:A:N6	2.24	0.71
52:sa:1580:U:H1'	57:sg:90:LEU:HD21	1.73	0.71
8:lH:56:ARG:NH2	8:lH:107:GLN:O	2.23	0.71
52:sa:630:A:H5''	67:sr:31:SER:HB2	1.71	0.71
1:lA:542:U:H2'	1:lA:543:A:H8	1.55	0.71
1:lA:1937:A:OP2	40:ln:33:LYS:NZ	2.20	0.71
3:lC:68:C:H42	3:lC:105:A:H61	1.39	0.71
52:sa:1716:A:OP1	66:sq:41:ARG:NH1	2.24	0.71
21:lU:99:MET:SD	21:lU:103:ARG:NH2	2.56	0.71
74:sy:66:LYS:HB3	74:sy:89:LEU:HD22	1.73	0.71
3:lC:69:U:H3	3:lC:104:G:H1	1.37	0.70
55:se:28:LYS:NZ	55:se:51:ASN:OD1	2.23	0.70
70:su:36:GLY:HA3	70:su:100:ILE:HA	1.73	0.70
1:lA:869:G:H21	1:lA:872:A:H62	1.11	0.70
1:lA:1555:U:H2'	1:lA:1556:G:H8	1.56	0.70
1:lA:1634:G:O6	41:lo:2:THR:N	2.24	0.70
52:sa:1395:A:H2'	52:sa:1396:A:C8	2.25	0.70
68:ss:35:CYS:SG	68:ss:36:LEU:N	2.64	0.70
1:lA:869:G:N2	1:lA:872:A:C6	2.58	0.70
1:lA:3491:C:O2'	5:lE:384:MET:SD	2.50	0.70
1:lA:3212:U:OP1	32:lf:55:HIS:NE2	2.25	0.70
5:lE:58:ARG:NH1	5:lE:353:GLU:OE2	2.25	0.70
1:lA:3455:G:H22	1:lA:3500:A:H2	1.38	0.70
4:lD:54:ARG:HG3	4:lD:56:ALA:H	1.56	0.70
52:sa:13:C:H4'	52:sa:1323:U:H3	1.57	0.70
52:sa:998:U:OP1	64:so:107:LYS:NZ	2.25	0.70
1:lA:871:A:H2'	1:lA:872:A:C8	2.27	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:IW:93:MET:HE2	23:IW:93:MET:HA	1.72	0.69
52:sa:1585:U:OP1	71:sv:40:THR:OG1	2.09	0.69
1:lA:1944:C:OP1	23:IW:107:ARG:NH2	2.24	0.69
48:sE:17:GLY:HA2	48:sE:27:ARG:HD2	1.75	0.69
54:sd:137:MET:HE2	54:sd:168:GLU:HG3	1.74	0.69
17:lQ:155:LYS:O	17:lQ:162:ARG:NH2	2.26	0.69
55:se:148:LYS:HB2	55:se:151:GLN:HB2	1.74	0.69
1:lA:639:G:H22	1:lA:650:A:H2	1.39	0.69
1:lA:816:G:N2	1:lA:829:G:OP2	2.23	0.69
1:lA:3329:U:HO2'	1:lA:3411:U:HO2'	1.33	0.69
1:lA:36:A:O2'	1:lA:1056:G:O6	2.10	0.69
1:lA:1892:C:H2'	1:lA:1893:A:H8	1.58	0.69
17:lQ:163:GLY:O	17:lQ:172:ARG:NH1	2.25	0.69
71:sv:97:SER:O	71:sv:101:ASN:ND2	2.26	0.69
30:ld:7:ARG:NH2	30:ld:9:SER:OG	2.25	0.69
16:lP:25:ALA:HA	16:lP:40:GLY:HA3	1.75	0.68
52:sa:1281:A:H4'	52:sa:1282:U:H5'	1.75	0.68
1:lA:3458:U:H3	1:lA:3494:G:H1	1.41	0.68
20:IT:78:VAL:HG22	20:IT:83:ILE:HG12	1.75	0.68
1:lA:1883:A:H62	1:lA:1931:U:H3	1.42	0.68
1:lA:1740:G:OP2	34:lh:37:LYS:NZ	2.24	0.68
1:lA:1841:G:N2	1:lA:1975:G:H22	1.92	0.68
1:lA:3147:U:OP1	5:lE:141:LYS:NZ	2.23	0.68
50:sJ:34:U:OP2	68:ss:158:ARG:NH2	2.27	0.68
1:lA:626:A:OP1	1:lA:664:A:N6	2.26	0.68
49:sI:16:G:H5''	49:sI:17:G:H5'	1.76	0.68
1:lA:869:G:N2	1:lA:872:A:C5	2.62	0.68
70:su:81:PRO:HG2	70:su:84:PHE:HB2	1.75	0.68
52:sa:286:U:H2'	52:sa:287:A:H8	1.58	0.68
1:lA:112:G:O2'	1:lA:311:A:N3	2.27	0.67
1:lA:1841:G:H22	1:lA:1975:G:H22	1.43	0.67
13:lM:116:TYR:OH	70:su:109:ARG:NH1	2.27	0.67
52:sa:1712:A:OP1	70:su:122:ARG:NH1	2.26	0.67
1:lA:1442:A:N6	15:lO:18:GLY:O	2.26	0.67
1:lA:2450:C:O2	5:lE:258:HIS:NE2	2.27	0.67
5:lE:219:VAL:HG12	5:lE:339:LEU:HD23	1.75	0.67
6:lF:101:MET:HE1	6:lF:105:LEU:HG	1.75	0.67
20:IT:5:TYR:HB2	20:IT:60:ILE:HD11	1.76	0.67
3:lC:79:G:C6	3:lC:95:A:N1	2.63	0.67
6:lF:377:PHE:HE1	16:lP:82:GLN:HG2	1.60	0.67
29:lc:59:ARG:NH2	43:lq:36:GLN:OE1	2.28	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:sg:52:PRO:HG2	57:sg:92:MET:HE2	1.76	0.67
1:lA:494:G:N2	1:lA:578:A:OP2	2.28	0.67
21:lU:99:MET:HE1	21:lU:128:LYS:HA	1.77	0.67
52:sa:751:A:N1	52:sa:785:A:N6	2.42	0.67
52:sa:783:A:H3'	52:sa:784:U:H4'	1.76	0.67
68:ss:54:ILE:O	68:ss:60:ARG:NH1	2.24	0.67
1:lA:95:A:OP1	17:lQ:195:ARG:NH1	2.25	0.67
1:lA:450:U:H4'	1:lA:1550:G:H4'	1.76	0.67
1:lA:893:U:H3	1:lA:897:A:H2	1.41	0.67
49:sI:63:U:H2'	49:sI:64:G:H8	1.58	0.67
52:sa:115:U:O2'	52:sa:328:C:O2	2.13	0.67
1:lA:3211:A:H2	1:lA:3239:G:H21	1.43	0.67
48:sE:41:ARG:O	52:sa:1540:G:N2	2.28	0.67
52:sa:356:C:H5'	52:sa:373:A:H5'	1.77	0.67
56:sf:104:LYS:O	56:sf:106:ARG:NH2	2.28	0.67
17:lQ:106:VAL:HG21	17:lQ:132:VAL:HG21	1.76	0.66
52:sa:1591:G:H21	52:sa:1772:A:H1'	1.61	0.66
56:sf:33:PRO:HD2	56:sf:81:PRO:HG2	1.77	0.66
66:sq:111:LEU:O	66:sq:113:HIS:N	2.28	0.66
1:lA:213:A:H5'	14:lN:133:LYS:H	1.60	0.66
1:lA:1664:G:OP2	1:lA:1752:U:O2'	2.13	0.66
2:lB:141:G:H2'	2:lB:142:A:H8	1.60	0.66
52:sa:63:U:O2'	52:sa:163:A:N3	2.25	0.66
23:lW:107:ARG:HE	23:lW:109:ILE:HD11	1.60	0.66
1:lA:925:A:N3	1:lA:2955:C:O2'	2.28	0.66
52:sa:90:G:OP1	52:sa:392:A:N6	2.29	0.66
52:sa:875:G:H21	65:sp:47:THR:HG21	1.60	0.66
52:sa:956:G:H1	52:sa:1003:A:HO2'	1.43	0.66
66:sq:27:MET:HG2	66:sq:28:PRO:HD2	1.76	0.66
1:lA:37:G:N2	1:lA:2946:A:N7	2.42	0.66
1:lA:2415:C:OP1	24:IX:54:ARG:NH2	2.28	0.66
1:lA:3243:C:OP1	5:lE:224:LYS:NZ	2.22	0.66
10:lJ:168:LYS:O	10:lJ:171:GLN:NE2	2.29	0.66
52:sa:107:A:H2'	52:sa:108:G:C8	2.31	0.66
52:sa:1717:U:OP2	66:sq:42:ARG:NH1	2.28	0.66
13:lM:43:GLN:NE2	13:lM:70:THR:O	2.28	0.66
1:lA:626:A:N6	1:lA:663:G:O2'	2.29	0.66
27:la:53:ASP:OD2	27:la:113:ARG:NH2	2.28	0.66
52:sa:394:A:OP1	60:sj:49:ARG:NH2	2.28	0.65
49:sI:13:C:O2	49:sI:14:A:H8	1.80	0.65
1:lA:631:G:N2	1:lA:632:G:N3	2.45	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:lA:1841:G:H22	1:lA:1975:G:H1	1.43	0.65
1:lA:1313:U:O2'	1:lA:1315:A:N7	2.27	0.65
1:lA:1523:U:OP1	33:lg:105:LYS:NZ	2.27	0.65
6:lF:55:SER:HB3	6:lF:58:ALA:HB2	1.78	0.65
6:lF:110:LYS:O	6:lF:113:ARG:NH1	2.28	0.65
52:sa:626:U:H4'	74:sy:7:ALA:HB2	1.79	0.65
1:lA:2462:G:H2'	1:lA:2463:A:H8	1.61	0.65
24:lX:83:ARG:NH2	24:lX:118:THR:O	2.28	0.65
28:lb:5:LEU:HD11	31:le:39:ARG:HD2	1.79	0.65
52:sa:1581:G:H2'	52:sa:1582:G:H8	1.59	0.65
1:lA:2173:A:H2'	1:lA:2174:U:H6	1.60	0.65
52:sa:251:C:O2'	60:sj:64:ASN:ND2	2.29	0.65
52:sa:1578:A:OP1	57:sg:166:ARG:NH1	2.29	0.65
1:lA:2337:G:O2'	1:lA:2339:C:N4	2.29	0.65
1:lA:2865:A:O2'	1:lA:2866:C:O4'	2.13	0.65
52:sa:213:A:H62	52:sa:243:U:H4'	1.62	0.65
38:ll:18:LEU:HD11	41:lo:51:MET:HE3	1.78	0.65
39:lm:85:ARG:NH2	52:sa:904:A:OP1	2.29	0.65
52:sa:584:C:H2'	52:sa:585:A:C8	2.31	0.65
55:se:63:LEU:O	55:se:65:GLY:N	2.30	0.65
65:sp:25:VAL:HG12	65:sp:89:HIS:HB2	1.79	0.65
66:sq:107:LYS:HG3	66:sq:109:GLU:H	1.61	0.65
56:sf:180:MET:HB2	56:sf:190:VAL:HG12	1.79	0.65
1:lA:1117:G:N3	1:lA:1120:A:N6	2.46	0.64
9:lI:109:ALA:HB2	22:IV:139:LYS:HB3	1.77	0.64
11:lK:100:MET:HB3	11:lK:150:ILE:HG13	1.79	0.64
16:lP:39:ASP:OD2	16:lP:49:ARG:NH2	2.30	0.64
52:sa:1816:U:H2'	52:sa:1817:A:C8	2.32	0.64
1:lA:59:A:N3	1:lA:74:U:O2'	2.24	0.64
3:lC:22:C:H2'	3:lC:23:A:H8	1.62	0.64
52:sa:213:A:N6	52:sa:244:U:OP2	2.31	0.64
7:lG:64:ILE:HD13	7:lG:109:LEU:HD22	1.79	0.64
52:sa:972:A:O2'	52:sa:1932:U:O2	2.14	0.64
1:lA:1157:A:H4'	7:lG:5:LYS:H	1.62	0.64
1:lA:2514:A:H61	1:lA:2583:U:H3	1.43	0.64
57:sg:63:PHE:HE1	57:sg:146:LEU:HD22	1.62	0.64
1:lA:1524:A:O2'	1:lA:1552:A:N1	2.29	0.64
53:sc:85:MET:HE2	53:sc:123:VAL:HG23	1.77	0.64
52:sa:1594:A:H2'	52:sa:1595:G:H8	1.63	0.64
56:sf:139:THR:OG1	56:sf:141:ASP:OD1	2.16	0.64
69:st:5:ARG:HB2	69:st:10:LYS:HE3	1.79	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
69:st:74:GLN:NE2	69:st:75:GLU:OE2	2.30	0.64
1:lA:1670:A:OP2	1:lA:1740:G:N2	2.30	0.64
52:sa:389:U:O4	52:sa:396:A:N7	2.30	0.64
52:sa:872:A:H2'	52:sa:873:A:C8	2.33	0.64
57:sg:120:ASN:ND2	57:sg:183:SER:O	2.30	0.64
5:lE:222:VAL:O	5:lE:335:ARG:NH1	2.31	0.64
45:sB:32:LYS:O	45:sB:37:LYS:NZ	2.31	0.64
1:lA:3262:U:OP1	42:lp:39:LYS:NZ	2.29	0.63
55:se:42:ARG:O	55:se:44:THR:N	2.31	0.63
1:lA:619:A:O2'	1:lA:620:U:O5'	2.16	0.63
1:lA:875:C:H2'	1:lA:876:A:C8	2.32	0.63
67:sr:36:GLU:HB3	67:sr:110:ILE:HG21	1.80	0.63
1:lA:926:A:H62	1:lA:1053:G:H1	1.46	0.63
1:lA:1327:A:N3	1:lA:2998:U:O2'	2.28	0.63
1:lA:1524:A:OP1	33:lg:102:SER:OG	2.13	0.63
3:lC:104:G:H2'	3:lC:105:A:H8	1.62	0.63
70:su:96:ASP:OD2	70:su:98:GLN:NE2	2.31	0.63
1:lA:612:A:H62	1:lA:715:A:H2'	1.64	0.63
55:se:174:ILE:HG12	55:se:195:LEU:HD11	1.79	0.63
5:lE:286:ARG:NH2	5:lE:296:LYS:O	2.32	0.63
52:sa:99:A:H61	52:sa:380:A:H1'	1.63	0.63
52:sa:852:A:H61	52:sa:935:C:H42	1.46	0.63
1:lA:64:A:OP2	1:lA:347:G:N2	2.29	0.63
1:lA:195:A:H61	1:lA:312:A:H2	1.47	0.63
1:lA:2233:G:H4'	1:lA:2234:A:H5'	1.80	0.63
38:ll:39:TYR:O	38:ll:41:ASP:N	2.31	0.63
4:lD:136:VAL:HA	4:lD:148:ILE:HG22	1.79	0.63
12:IL:171:TRP:CD1	12:IL:172:GLY:H	2.16	0.63
48:sE:14:TYR:HB3	52:sa:1763:A:C8	2.34	0.63
63:sm:57:CYS:HB3	63:sm:60:THR:HG22	1.79	0.63
1:lA:3367:U:H2'	1:lA:3368:G:C8	2.34	0.63
52:sa:865:A:H2'	52:sa:866:G:C8	2.34	0.63
1:lA:2114:U:N3	1:lA:2198:G:OP2	2.27	0.63
1:lA:2173:A:H2'	1:lA:2174:U:C6	2.33	0.63
20:IT:92:ASP:OD2	20:IT:101:GLN:NE2	2.31	0.63
52:sa:467:C:O2'	52:sa:760:A:N3	2.32	0.63
1:lA:43:C:OP2	1:lA:44:A:O2'	2.11	0.62
1:lA:628:C:O2'	20:IT:141:HIS:NE2	2.32	0.62
1:lA:2257:G:OP1	4:lD:193:ARG:NH2	2.30	0.62
1:lA:3355:U:O2	15:lO:4:ASN:ND2	2.31	0.62
52:sa:918:G:N2	52:sa:921:A:OP2	2.30	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:sa:1322:G:N2	52:sa:1325:A:OP2	2.31	0.62
1:1A:2705:A:C4'	1:1A:2706:C:H5'	2.28	0.62
50:sJ:23:G:H2'	50:sJ:24:C:C6	2.34	0.62
53:sc:164:CYS:SG	53:sc:165:GLY:N	2.73	0.62
55:se:74:ASP:OD1	65:sp:123:ARG:NH2	2.32	0.62
5:1E:84:MET:HG2	5:1E:166:THR:HA	1.82	0.62
1:1A:2990:A:OP1	42:lp:22:ARG:NH1	2.33	0.62
19:1S:60:ILE:HG22	19:1S:83:THR:HB	1.82	0.62
19:1S:152:VAL:HA	19:1S:159:ALA:H	1.63	0.62
52:sa:107:A:H2'	52:sa:108:G:H8	1.63	0.62
52:sa:1441:C:OP1	69:st:5:ARG:NH2	2.32	0.62
54:sd:137:MET:CE	54:sd:168:GLU:HG3	2.30	0.62
12:1L:51:HIS:ND1	12:1L:137:SER:OG	2.22	0.62
55:se:117:LYS:NZ	55:se:210:ASP:OD2	2.33	0.62
1:1A:1072:U:OP1	30:ld:12:GLN:NE2	2.30	0.62
1:1A:3349:G:OP2	16:1P:5:ARG:NH2	2.32	0.62
3:1C:26:U:O2	3:1C:52:G:N2	2.23	0.62
12:1L:112:GLN:OE1	49:sI:73:A:O2'	2.18	0.62
49:sI:63:U:H2'	49:sI:64:G:C8	2.34	0.62
60:sj:36:THR:O	60:sj:96:LEU:N	2.33	0.62
12:1L:66:GLU:OE1	12:1L:69:ARG:NH2	2.32	0.62
52:sa:258:A:OP2	52:sa:285:G:N2	2.31	0.62
3:1C:33:A:N1	3:1C:45:U:O2'	2.29	0.62
9:1I:76:ARG:NH2	9:1I:79:GLY:O	2.30	0.62
40:ln:60:ILE:H	40:ln:60:ILE:HD12	1.65	0.62
52:sa:938:U:O2'	52:sa:940:U:OP2	2.16	0.62
56:sf:41:PRO:HD2	56:sf:44:LEU:HD12	1.80	0.62
8:1H:64:ILE:O	8:1H:113:THR:OG1	2.17	0.62
52:sa:42:A:N1	52:sa:373:A:N6	2.48	0.62
55:se:73:ALA:HB3	65:sp:123:ARG:HH22	1.65	0.62
61:sk:78:ARG:O	61:sk:82:ASN:ND2	2.33	0.62
7:1G:65:ALA:HB2	7:1G:74:ILE:HD13	1.82	0.62
52:sa:401:U:H2'	52:sa:402:A:C8	2.34	0.62
3:1C:59:G:H5'	7:1G:265:LYS:HD3	1.82	0.61
68:ss:109:GLN:OE1	68:ss:117:LYS:NZ	2.32	0.61
1:1A:65:U:HO2'	1:1A:97:A:HO2'	1.47	0.61
1:1A:964:G:N2	1:1A:967:A:OP2	2.33	0.61
16:1P:29:GLU:OE2	20:1T:91:ARG:NH2	2.34	0.61
20:1T:79:SER:OG	20:1T:84:HIS:NE2	2.25	0.61
52:sa:840:U:H2'	52:sa:841:U:C6	2.35	0.61
52:sa:1718:U:O2'	52:sa:1763:A:N3	2.30	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1B:93:C:OP1	38:1L:73:ARG:NH1	2.33	0.61
25:1Y:78:VAL:HG21	25:1Y:103:VAL:HG21	1.81	0.61
38:1L:60:GLY:HA2	38:1L:64:MET:HE2	1.82	0.61
52:sa:145:U:O4	52:sa:159:A:N6	2.33	0.61
53:sc:58:ILE:HG23	53:sc:63:ILE:HB	1.82	0.61
58:sh:57:ASP:OD2	58:sh:98:ARG:NH1	2.24	0.61
1:1A:502:U:H2'	1:1A:503:A:C8	2.36	0.61
1:1A:2257:G:H5''	4:1D:193:ARG:HH12	1.65	0.61
1:1A:200:A:H62	1:1A:309:G:H21	1.47	0.61
69:st:77:GLU:OE1	69:st:80:ARG:NH1	2.33	0.61
1:1A:3336:G:O2'	15:1O:115:LYS:O	2.18	0.61
8:1H:165:LYS:NZ	8:1H:166:GLN:OE1	2.33	0.61
52:sa:239:U:N3	52:sa:242:G:OP2	2.29	0.61
52:sa:1556:U:H2'	52:sa:1557:C:H6	1.65	0.61
69:st:43:THR:HG22	69:st:46:LEU:H	1.65	0.61
1:1A:450:U:H2'	1:1A:451:G:H5'	1.83	0.61
1:1A:1634:G:N2	1:1A:1634:G:OP2	2.34	0.61
1:1A:2499:U:H2'	1:1A:2500:A:H8	1.66	0.61
3:1C:75:G:H1	3:1C:99:U:H5	1.48	0.61
6:1F:237:LEU:HD22	6:1F:242:LEU:HD21	1.82	0.61
20:1T:40:MET:HE3	22:1V:157:VAL:HG12	1.83	0.61
52:sa:797:G:OP2	52:sa:797:G:N2	2.33	0.61
1:1A:1217:U:H4'	22:1V:123:LYS:HE3	1.81	0.61
1:1A:3372:U:O2'	1:1A:3374:C:OP2	2.15	0.61
3:1C:3:G:N2	7:1G:79:TYR:OH	2.34	0.61
43:1q:72:CYS:SG	43:1q:75:CYS:N	2.73	0.61
5:1E:232:ILE:HG13	5:1E:249:ARG:HB3	1.82	0.61
22:1V:88:ARG:NH2	30:1d:28:ALA:O	2.34	0.61
28:1b:10:VAL:O	28:1b:83:THR:OG1	2.18	0.61
50:sJ:24:C:H2'	50:sJ:25:U:H6	1.65	0.61
68:ss:98:GLN:HE22	68:ss:134:ALA:HA	1.65	0.61
1:1A:1303:A:N3	1:1A:1453:U:O2'	2.31	0.60
52:sa:1670:A:H2'	52:sa:1671:A:C8	2.36	0.60
56:sf:209:LYS:NZ	56:sf:213:GLY:O	2.33	0.60
1:1A:1252:G:OP2	12:1L:98:ARG:NH1	2.33	0.60
1:1A:2468:C:O2	5:1E:268:ARG:NH2	2.34	0.60
3:1C:10:A:N1	3:1C:65:U:O2'	2.30	0.60
32:1f:63:PHE:HB3	32:1f:103:ARG:HG2	1.82	0.60
62:sl:21:ILE:HD13	62:sl:45:MET:HE1	1.83	0.60
1:1A:1561:U:OP2	29:1c:4:ARG:NH2	2.30	0.60
15:1O:82:ARG:NH2	15:1O:103:LEU:O	2.33	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:1266:C:O2'	1:1A:1278:A:N3	2.34	0.60
1:1A:2485:G:H3'	1:1A:2486:U:H4'	1.83	0.60
34:lh:81:CYS:SG	34:lh:84:CYS:N	2.74	0.60
48:sE:21:CYS:HB3	48:sE:25:GLY:H	1.66	0.60
52:sa:584:C:H2'	52:sa:585:A:H8	1.65	0.60
56:sf:98:ARG:HB3	56:sf:110:VAL:HG13	1.84	0.60
1:1A:3383:A:OP2	5:1E:148:ARG:NH2	2.30	0.60
52:sa:1907:G:H1'	52:sa:1928:A:H2	1.66	0.60
1:1A:1209:U:OP1	1:1A:1211:A:O2'	2.20	0.60
1:1A:1261:A:OP1	30:ld:7:ARG:NH1	2.35	0.60
2:1B:154:C:H2'	2:1B:155:A:C8	2.36	0.60
65:sp:56:ARG:NH2	65:sp:75:ASP:OD2	2.35	0.60
1:1A:1414:A:H2'	1:1A:1415:A:H8	1.66	0.60
1:1A:1476:G:O2'	1:1A:1477:A:H5'	2.01	0.60
1:1A:1868:U:OP2	23:1W:55:LYS:NZ	2.32	0.60
1:1A:2272:C:O2'	1:1A:2346:A:N3	2.31	0.60
1:1A:2368:U:H5''	52:sa:1824:G:H21	1.67	0.60
2:1B:25:U:OP2	27:la:15:ARG:NH1	2.34	0.60
6:1F:9:ILE:HD13	6:1F:256:GLU:HG3	1.82	0.60
10:1J:60:LYS:HG3	10:1J:232:ARG:HE	1.67	0.60
11:1K:101:ARG:NH1	11:1K:149:GLU:OE1	2.34	0.60
57:sg:197:ASP:OD1	57:sg:200:ARG:NH2	2.34	0.60
65:sp:93:ARG:HB3	65:sp:127:VAL:HG23	1.82	0.60
70:su:34:ILE:HG22	70:su:36:GLY:H	1.67	0.60
1:1A:2971:G:OP2	12:1L:7:ARG:NH2	2.35	0.60
52:sa:864:C:H2'	52:sa:865:A:C8	2.37	0.60
61:sk:48:MET:HE2	61:sk:48:MET:HA	1.83	0.60
1:1A:2634:A:OP1	4:1D:69:TYR:OH	2.20	0.60
1:1A:2798:G:OP1	19:1S:151:LYS:NZ	2.35	0.60
1:1A:3247:C:O2	24:1X:86:ARG:NH2	2.35	0.60
1:1A:3454:G:C6	1:1A:3501:A:N1	2.69	0.60
11:1K:101:ARG:HG3	11:1K:188:SER:HB3	1.84	0.60
48:sE:16:ASN:ND2	52:sa:1231:U:O2	2.35	0.60
52:sa:456:G:H2'	52:sa:457:G:H8	1.66	0.60
1:1A:543:A:H2'	1:1A:544:U:H6	1.67	0.60
1:1A:2414:C:O2	24:1X:21:ASN:ND2	2.26	0.60
52:sa:440:A:N6	52:sa:457:G:N3	2.43	0.60
52:sa:1759:A:H2'	52:sa:1760:G:H8	1.67	0.60
52:sa:1823:U:OP1	52:sa:1824:G:O2'	2.18	0.60
54:sd:108:ARG:NH1	54:sd:115:GLN:OE1	2.35	0.60
1:1A:2432:A:H2	18:1R:131:ARG:HH22	1.50	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:IC:5:A:OP2	7:IG:22:ARG:NH2	2.35	0.59
3:IC:73:G:O2'	20:IT:46:LYS:NZ	2.32	0.59
10:IJ:173:PRO:HG3	10:IJ:216:LYS:HE3	1.82	0.59
47:sD:21:ARG:HG2	47:sD:29:LEU:HD12	1.83	0.59
1:lA:718:A:H2'	1:lA:719:U:C6	2.37	0.59
1:lA:3176:G:O2'	1:lA:3185:G:O6	2.19	0.59
25:IY:85:LYS:HG2	25:IY:106:ALA:HB2	1.83	0.59
35:li:9:LYS:NZ	35:li:12:GLU:OE2	2.28	0.59
52:sa:1668:G:OP2	71:sv:102:ARG:NH1	2.35	0.59
52:sa:1760:G:OP2	52:sa:1762:C:N4	2.35	0.59
62:sl:31:HIS:CD2	62:sl:41:VAL:HG21	2.36	0.59
1:lA:1115:A:O2'	3:IC:76:U:O2	2.19	0.59
1:lA:1174:A:N3	3:IC:78:C:O2'	2.35	0.59
1:lA:3045:A:OP1	11:IK:176:LYS:NZ	2.29	0.59
10:IJ:94:ALA:O	10:IJ:98:GLN:NE2	2.30	0.59
56:sf:240:LYS:HD2	56:sf:240:LYS:O	2.02	0.59
21:IU:44:LEU:HD22	21:IU:49:LEU:HD12	1.84	0.59
52:sa:1245:C:H2'	52:sa:1246:A:C8	2.36	0.59
53:sc:99:ARG:NH2	53:sc:119:CYS:SG	2.75	0.59
1:lA:457:G:OP1	18:IR:62:LYS:NZ	2.34	0.59
1:lA:779:A:N6	1:lA:3012:U:OP1	2.35	0.59
1:lA:1177:A:H2'	1:lA:1178:A:H8	1.68	0.59
1:lA:2856:G:H2'	1:lA:2903:A:H61	1.66	0.59
7:IG:122:VAL:O	7:IG:244:ARG:NH2	2.35	0.59
38:ll:21:ARG:NH2	38:ll:38:GLY:O	2.35	0.59
52:sa:602:U:O5'	52:sa:604:G:N2	2.35	0.59
60:sj:87:ASP:OD1	60:sj:89:ASP:N	2.34	0.59
64:so:99:ARG:O	64:so:101:HIS:N	2.35	0.59
1:lA:1086:G:H2'	1:lA:1087:G:C8	2.38	0.59
1:lA:1494:G:OP2	29:lc:10:ARG:NH2	2.35	0.59
1:lA:1744:A:H2'	1:lA:1745:A:C8	2.37	0.59
1:lA:1892:C:H2'	1:lA:1893:A:C8	2.38	0.59
6:lF:80:SER:O	6:lF:87:ASN:ND2	2.35	0.59
32:lf:62:SER:HB2	32:lf:65:LYS:HD3	1.85	0.59
52:sa:1750:G:N2	52:sa:1776:G:O2'	2.35	0.59
60:sj:44:HIS:N	60:sj:56:ARG:O	2.35	0.59
3:IC:17:G:N2	3:IC:60:C:O2	2.36	0.59
6:lF:77:PRO:HB2	6:lF:91:ALA:HB3	1.83	0.59
24:IX:9:ARG:NH2	24:IX:11:GLY:O	2.32	0.59
1:lA:231:U:H4'	27:la:123:SER:HB3	1.84	0.59
1:lA:1112:G:O2'	1:lA:1172:G:O6	2.21	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:lA:1259:G:OP2	30:ld:5:LYS:NZ	2.24	0.59
24:IX:57:CYS:HG	24:IX:122:SER:HG	1.51	0.59
28:lb:104:GLU:HG2	28:lb:107:LYS:HD3	1.83	0.59
52:sa:139:G:H1	52:sa:166:A:H61	0.72	0.59
1:lA:1022:U:HO2'	1:lA:1677:A:HO2'	1.47	0.59
1:lA:1181:A:H5''	1:lA:1182:A:H5'	1.85	0.59
5:lE:370:ARG:HG2	26:lZ:34:ILE:HD11	1.85	0.59
1:lA:3150:A:H2'	1:lA:3151:A:H8	1.66	0.58
5:lE:361:ASP:OD2	5:lE:365:LYS:NZ	2.36	0.58
6:lF:360:LYS:HB3	6:lF:400:THR:HG21	1.85	0.58
1:lA:688:A:H2'	1:lA:689:A:H8	1.67	0.58
1:lA:862:G:O6	1:lA:879:U:O2	2.21	0.58
6:lF:61:GLN:OE1	38:ll:55:ARG:NH2	2.28	0.58
52:sa:1932:U:H2'	52:sa:1933:G:H8	1.68	0.58
70:su:25:ARG:NH1	70:su:29:TYR:O	2.37	0.58
1:lA:3038:C:O2'	42:lp:25:TYR:O	2.15	0.58
1:lA:3474:U:H3	1:lA:3480:A:H62	1.49	0.58
52:sa:802:U:N3	52:sa:803:G:O6	2.36	0.58
52:sa:1350:A:H2'	52:sa:1351:A:C8	2.37	0.58
63:sm:34:HIS:NE2	63:sm:47:ALA:O	2.35	0.58
71:sv:57:ARG:NH1	71:sv:101:ASN:OD1	2.37	0.58
1:lA:232:G:N1	1:lA:274:U:OP2	2.28	0.58
1:lA:779:A:H5'	1:lA:2448:A:H62	1.68	0.58
21:lU:89:THR:O	21:lU:89:THR:OG1	2.22	0.58
52:sa:160:G:H2'	52:sa:161:A:O4'	2.03	0.58
1:lA:1293:U:O4	1:lA:1454:G:O2'	2.19	0.58
1:lA:2225:A:O2'	4:lD:179:ILE:O	2.15	0.58
1:lA:2366:C:H1'	52:sa:1892:G:H21	1.68	0.58
52:sa:1440:A:N6	52:sa:1519:G:O2'	2.37	0.58
1:lA:2628:A:N7	4:lD:100:ASN:ND2	2.52	0.58
1:lA:3170:A:H2'	1:lA:3171:A:H8	1.68	0.58
1:lA:3315:U:O2'	1:lA:3418:C:OP2	2.19	0.58
3:lC:54:U:H4'	13:lM:152:HIS:HB2	1.86	0.58
28:lb:91:GLN:NE2	28:lb:124:GLU:OE2	2.37	0.58
37:lk:29:LYS:O	37:lk:32:HIS:ND1	2.37	0.58
52:sa:1581:G:H2'	52:sa:1582:G:C8	2.37	0.58
52:sa:1886:U:O4	52:sa:1887:A:N6	2.37	0.58
1:lA:1314:A:N7	11:lK:73:ARG:NH2	2.51	0.58
1:lA:2702:G:O6	1:lA:2717:A:N6	2.37	0.58
9:ll:203:LYS:HB2	9:ll:207:ILE:HD11	1.86	0.58
12:lL:190:ILE:HG23	12:lL:197:PHE:HB2	1.85	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:sI:22:G:H2'	49:sI:23:A:C8	2.38	0.58
52:sa:1007:A:OP1	52:sa:1936:G:O2'	2.22	0.58
52:sa:1126:G:H1'	67:sr:76:SER:HB3	1.86	0.58
54:sd:22:LYS:HG2	72:sw:61:LEU:HD11	1.86	0.58
1:lA:2597:U:H3	4:lD:67:ARG:HD2	1.68	0.58
2:lB:5:A:O2'	18:lR:61:ARG:NH1	2.36	0.58
4:lD:33:ASP:N	4:lD:33:ASP:OD1	2.34	0.58
4:lD:39:SER:OG	4:lD:40:THR:N	2.34	0.58
11:lK:19:ILE:HD11	11:lK:89:ILE:HD11	1.85	0.58
49:sI:68:G:H2'	49:sI:69:G:C8	2.38	0.58
1:lA:3107:G:N2	1:lA:3110:A:OP2	2.36	0.58
51:sK:8:A:O2'	74:sy:59:GLN:NE2	2.35	0.58
52:sa:1544:U:H2'	52:sa:1545:G:H8	1.69	0.58
57:sg:102:ILE:O	57:sg:106:THR:OG1	2.19	0.58
60:sj:56:ARG:NH1	60:sj:210:GLY:O	2.34	0.58
1:lA:772:C:N4	1:lA:773:G:O6	2.37	0.58
52:sa:1172:G:H2'	52:sa:1173:A:C8	2.38	0.58
52:sa:1341:C:HO2'	52:sa:1508:A:HO2'	1.51	0.58
71:sv:8:LYS:HE3	71:sv:67:LEU:HD22	1.86	0.58
1:lA:838:U:O2'	1:lA:907:U:OP1	2.21	0.57
1:lA:3091:C:OP1	5:lE:246:ARG:NH1	2.32	0.57
1:lA:3154:A:H2'	1:lA:3155:U:C6	2.38	0.57
28:lb:33:THR:OG1	28:lb:35:ASP:OD1	2.23	0.57
60:sj:67:TRP:NE1	60:sj:69:SER:HB3	2.19	0.57
1:lA:764:U:H2'	1:lA:765:A:H8	1.69	0.57
1:lA:2297:G:N1	1:lA:2300:A:OP2	2.33	0.57
1:lA:3350:U:OP2	1:lA:3351:U:O2'	2.16	0.57
7:lG:125:ALA:O	7:lG:192:ARG:NH1	2.37	0.57
15:lO:47:THR:H	15:lO:50:MET:HE3	1.69	0.57
22:lV:68:ASN:OD1	22:lV:69:PRO:HD2	2.02	0.57
31:le:40:ASP:N	31:le:40:ASP:OD1	2.35	0.57
68:ss:64:PHE:HB3	68:ss:68:PHE:HE2	1.69	0.57
73:sx:23:ALA:HB3	73:sx:26:ASP:HB2	1.86	0.57
1:lA:547:A:H2'	1:lA:548:A:H8	1.68	0.57
7:lG:85:ARG:NH2	7:lG:249:ALA:O	2.37	0.57
23:lW:70:VAL:HG12	23:lW:72:ASP:H	1.70	0.57
26:lZ:8:CYS:SG	26:lZ:11:SER:OG	2.62	0.57
52:sa:1396:A:H2'	52:sa:1425:A:C8	2.39	0.57
1:lA:311:A:H5'	37:lk:30:ARG:HB2	1.87	0.57
1:lA:1639:U:H2'	1:lA:1640:A:H8	1.70	0.57
1:lA:2590:A:N1	1:lA:2664:A:N6	2.52	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:2705:A:O2'	22:1V:10:ARG:NH2	2.36	0.57
4:1D:242:ARG:HH21	4:1D:246:ILE:HA	1.68	0.57
52:sa:15:U:H2'	52:sa:16:G:O4'	2.05	0.57
52:sa:841:U:O2'	52:sa:842:A:O4'	2.21	0.57
68:ss:65:GLU:OE1	68:ss:97:ARG:NH2	2.37	0.57
1:1A:844:G:N2	1:1A:847:A:OP2	2.36	0.57
1:1A:2815:G:N2	1:1A:2818:A:OP2	2.34	0.57
1:1A:2864:A:O2'	1:1A:2865:A:O5'	2.21	0.57
52:sa:1734:G:H4'	52:sa:1735:A:O5'	2.04	0.57
52:sa:1891:A:O3'	52:sa:1892:G:H8	1.86	0.57
55:se:66:ARG:NH1	65:sp:46:GLU:OE2	2.38	0.57
1:1A:361:C:OP1	14:1N:100:LYS:NZ	2.36	0.57
1:1A:2668:G:H2'	1:1A:2669:A:H8	1.68	0.57
1:1A:3359:U:OP1	8:1H:194:LYS:NZ	2.36	0.57
3:1C:42:U:OP1	13:1M:137:ARG:NH1	2.38	0.57
16:1P:69:ARG:NH2	20:1T:127:ASP:OD2	2.35	0.57
52:sa:615:A:O2'	52:sa:1132:C:O2'	2.22	0.57
1:1A:13:C:P	25:1Y:27:ARG:HH22	2.28	0.57
1:1A:1555:U:O2	33:lg:78:ARG:NH2	2.34	0.57
1:1A:2865:A:O2'	1:1A:2866:C:OP1	2.20	0.57
52:sa:335:U:H2'	52:sa:336:A:C8	2.39	0.57
52:sa:1141:U:H2'	52:sa:1142:A:H8	1.68	0.57
52:sa:1182:C:H2'	52:sa:1183:A:H8	1.70	0.57
52:sa:1356:A:OP1	69:st:45:ARG:NH2	2.37	0.57
52:sa:1595:G:O2'	52:sa:1661:C:O2	2.22	0.57
52:sa:1661:C:H2'	52:sa:1662:G:C8	2.39	0.57
56:sf:181:MET:HE3	56:sf:208:ILE:HD13	1.86	0.57
67:sr:23:ARG:NH2	73:sx:17:VAL:O	2.36	0.57
1:1A:815:A:H4'	1:1A:817:A:OP1	2.03	0.57
1:1A:1236:G:H2'	1:1A:1237:A:H8	1.69	0.57
1:1A:1269:U:OP2	33:lg:44:ARG:NH2	2.31	0.57
1:1A:2991:G:OP1	42:lp:25:TYR:OH	2.23	0.57
1:1A:3023:U:O2	5:1E:252:ALA:HB3	2.04	0.57
52:sa:782:A:C8	52:sa:784:U:H1'	2.40	0.57
54:sd:122:LYS:NZ	54:sd:135:SER:OG	2.38	0.57
57:sg:177:ASP:OD1	57:sg:178:GLU:N	2.37	0.57
1:1A:315:A:N7	17:1Q:12:ARG:NH1	2.53	0.57
9:1I:214:GLY:O	9:1I:218:ARG:NH2	2.30	0.57
31:1e:15:ASN:OD1	31:1e:74:TYR:OH	2.22	0.57
52:sa:1308:U:OP2	52:sa:1309:C:O2'	2.19	0.57
52:sa:1748:U:N3	52:sa:1780:A:N7	2.53	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:lA:1206:A:H2'	1:lA:1207:C:C6	2.40	0.57
1:lA:3397:C:OP2	1:lA:3398:U:O2'	2.23	0.57
52:sa:1556:U:H2'	52:sa:1557:C:C6	2.40	0.57
65:sp:35:THR:O	65:sp:53:GLY:N	2.35	0.57
69:st:25:THR:HG23	69:st:27:ASP:H	1.70	0.57
69:st:41:VAL:HG22	69:st:43:THR:H	1.70	0.57
1:lA:502:U:H2'	1:lA:503:A:H8	1.70	0.56
1:lA:2175:C:H2'	1:lA:2176:G:C8	2.38	0.56
57:sg:25:SER:OG	57:sg:27:ASP:OD1	2.23	0.56
57:sg:63:PHE:CE1	57:sg:146:LEU:HD22	2.40	0.56
60:sj:87:ASP:OD1	60:sj:88:ASN:N	2.37	0.56
67:sr:102:LEU:N	67:sr:113:ASN:OD1	2.35	0.56
5:lE:10:ARG:HH12	5:lE:265:THR:HB	1.70	0.56
1:lA:542:U:H2'	1:lA:543:A:C8	2.39	0.56
1:lA:1031:G:N1	4:lD:208:ASP:OD2	2.33	0.56
1:lA:1335:U:O2	20:lT:109:GLN:NE2	2.37	0.56
1:lA:3405:A:N6	8:lH:164:GLU:OE1	2.38	0.56
1:lA:3500:A:H2'	1:lA:3501:A:C8	2.39	0.56
8:lH:65:LEU:HD11	8:lH:79:PHE:HB2	1.86	0.56
16:lP:33:HIS:ND1	16:lP:33:HIS:O	2.39	0.56
44:sA:100:VAL:N	44:sA:108:ILE:O	2.38	0.56
52:sa:442:U:O2'	56:sf:25:PHE:O	2.23	0.56
52:sa:1722:A:OP1	66:sq:114:TYR:OH	2.19	0.56
70:su:97:SER:OG	70:su:98:GLN:N	2.37	0.56
1:lA:58:G:H21	17:lQ:186:ALA:HB1	1.71	0.56
1:lA:1128:G:OP1	12:lL:39:ARG:NH2	2.38	0.56
1:lA:1186:G:H21	1:lA:1220:A:N6	2.03	0.56
1:lA:2468:C:O2'	5:lE:268:ARG:NH1	2.39	0.56
13:lM:63:ASP:OD1	13:lM:63:ASP:N	2.38	0.56
16:lP:108:THR:OG1	16:lP:111:ASP:OD2	2.22	0.56
45:sB:5:ARG:NH1	52:sa:1943:C:OP2	2.37	0.56
52:sa:866:G:H2'	52:sa:867:A:H8	1.70	0.56
57:sg:38:LEU:HD12	57:sg:145:PRO:HB2	1.87	0.56
1:lA:701:A:H5''	1:lA:703:G:H21	1.69	0.56
10:lJ:80:GLN:HA	10:lJ:83:LYS:HE2	1.87	0.56
52:sa:1221:A:OP2	52:sa:1571:G:N2	2.24	0.56
61:sk:31:VAL:HA	61:sk:36:LEU:HD12	1.88	0.56
68:ss:75:TYR:OH	68:ss:100:ILE:O	2.21	0.56
71:sv:14:ASP:OD1	71:sv:15:PHE:N	2.38	0.56
1:lA:789:C:H2'	1:lA:790:A:H8	1.69	0.56
3:lC:21:C:N4	3:lC:52:G:O2'	2.39	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:lk:39:ARG:NH1	37:lk:43:GLY:O	2.35	0.56
45:sB:22:ARG:NH1	65:sp:140:GLY:O	2.38	0.56
49:sI:8:U:H3	49:sI:15:G:H1	1.52	0.56
52:sa:1748:U:O2	52:sa:1780:A:N6	2.34	0.56
1:lA:1252:G:N2	1:lA:1255:A:OP2	2.38	0.56
1:lA:2711:U:OP2	22:IV:9:ARG:NH2	2.34	0.56
52:sa:1002:U:O2'	52:sa:1151:A:N1	2.38	0.56
56:sf:69:LYS:HG2	56:sf:74:ILE:HB	1.86	0.56
62:sl:25:LYS:HG2	62:sl:61:ARG:HH21	1.69	0.56
1:lA:1555:U:H2'	1:lA:1556:G:C8	2.39	0.56
52:sa:440:A:H5''	56:sf:57:ARG:HH12	1.68	0.56
52:sa:1561:G:C6	52:sa:1562:G:N2	2.74	0.56
54:sd:133:VAL:HG21	54:sd:152:ILE:HD11	1.88	0.56
1:lA:871:A:O2'	1:lA:872:A:O5'	2.23	0.56
1:lA:2890:A:H2'	1:lA:2891:G:C8	2.41	0.56
14:lN:264:ARG:O	14:lN:268:ASN:ND2	2.39	0.56
46:sC:18:LYS:O	52:sa:1095:U:O2'	2.23	0.56
48:sE:22:ARG:NH1	48:sE:36:LEU:O	2.39	0.56
52:sa:919:A:OP2	64:so:6:ASN:ND2	2.39	0.56
52:sa:1820:G:H5''	52:sa:1821:A:H5'	1.86	0.56
52:sa:111:A:O2'	63:sm:66:ARG:NH1	2.39	0.56
55:se:201:ALA:O	55:se:203:CYS:N	2.39	0.56
56:sf:62:ILE:HG22	56:sf:68:ILE:HD11	1.87	0.56
1:lA:50:G:HO2'	1:lA:1688:A:HO2'	1.54	0.55
1:lA:760:A:H2'	1:lA:761:A:H8	1.70	0.55
1:lA:993:U:OP2	1:lA:2109:C:O2'	2.23	0.55
1:lA:1498:C:O3'	19:IS:26:LYS:NZ	2.39	0.55
1:lA:1622:A:C2	34:lh:4:ARG:HD3	2.41	0.55
1:lA:2795:C:N4	1:lA:2802:A:N1	2.54	0.55
1:lA:3267:G:O6	1:lA:3275:C:H5''	2.06	0.55
4:ID:116:VAL:HG13	4:ID:164:ALA:HB2	1.88	0.55
16:lP:29:GLU:HG2	20:lT:68:VAL:HG21	1.88	0.55
52:sa:1661:C:H2'	52:sa:1662:G:H8	1.71	0.55
52:sa:1776:G:H5''	57:sg:88:LYS:HD2	1.88	0.55
56:sf:67:THR:HG23	56:sf:92:ARG:HG3	1.87	0.55
1:lA:405:U:OP1	38:ll:11:ARG:NH2	2.37	0.55
1:lA:1649:U:HO2'	1:lA:2429:G:HO2'	1.53	0.55
1:lA:1937:A:OP1	40:ln:35:LYS:NZ	2.33	0.55
11:lK:112:VAL:HB	11:lK:121:GLN:HB3	1.87	0.55
32:lf:61:GLU:HB3	32:lf:66:ARG:HE	1.71	0.55
52:sa:1340:U:OP1	52:sa:1353:G:N2	2.33	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:sh:78:ASP:OD1	58:sh:78:ASP:N	2.38	0.55
62:sl:30:LEU:HA	62:sl:37:PRO:HA	1.88	0.55
1:lA:336:A:H5''	17:lQ:97:SER:HB3	1.88	0.55
1:lA:672:G:O6	6:lF:412:LYS:NZ	2.39	0.55
1:lA:2425:U:O2'	1:lA:3436:A:N3	2.40	0.55
1:lA:3476:C:H4'	1:lA:3476:C:OP1	2.05	0.55
2:lB:110:A:N6	41:lo:51:MET:O	2.39	0.55
4:lD:192:LYS:HD3	4:lD:193:ARG:HH21	1.71	0.55
9:II:176:ILE:HD13	9:II:183:PHE:HE1	1.71	0.55
52:sa:1827:U:H2'	52:sa:1828:A:H8	1.71	0.55
54:sd:211:LYS:NZ	69:st:40:VAL:O	2.38	0.55
68:ss:35:CYS:HB2	68:ss:82:ILE:HG12	1.88	0.55
1:lA:820:A:OP1	14:lN:37:ARG:NH2	2.37	0.55
6:lF:389:LYS:C	6:lF:391:LYS:H	2.14	0.55
65:sp:18:SER:OG	65:sp:19:THR:N	2.38	0.55
1:lA:211:U:OP1	14:lN:134:ARG:NH1	2.40	0.55
1:lA:1648:U:OP2	1:lA:2081:A:O2'	2.25	0.55
8:lH:176:GLU:HA	8:lH:179:LYS:HE2	1.88	0.55
1:lA:26:C:OP1	17:lQ:188:ALA:N	2.37	0.55
1:lA:1098:A:HO2'	1:lA:1183:G:HO2'	1.55	0.55
1:lA:2466:G:N2	5:lE:228:TYR:OH	2.33	0.55
1:lA:2469:G:OP1	1:lA:2469:G:C8	2.60	0.55
6:lF:11:ASP:OD1	27:la:146:TYR:OH	2.23	0.55
50:sJ:16:C:H2'	50:sJ:61:U:H3	1.72	0.55
52:sa:366:G:N2	52:sa:606:U:O2	2.40	0.55
52:sa:397:U:OP1	56:sf:1:ARG:NH2	2.39	0.55
52:sa:932:C:H2'	52:sa:933:A:C8	2.41	0.55
52:sa:1115:C:N4	52:sa:1116:U:O4	2.39	0.55
52:sa:1754:U:O4	52:sa:1755:A:N6	2.40	0.55
52:sa:1923:A:H2'	52:sa:1924:G:C8	2.41	0.55
67:sr:45:GLN:O	67:sr:68:ARG:NH1	2.40	0.55
70:su:70:ASP:OD1	70:su:70:ASP:N	2.37	0.55
29:lc:4:ARG:HA	29:lc:9:ARG:HG3	1.87	0.55
56:sf:127:ARG:HH11	56:sf:135:PRO:HB2	1.71	0.55
64:so:92:MET:O	64:so:94:ARG:N	2.39	0.55
1:lA:1320:U:O2'	3:lC:83:G:N2	2.40	0.55
1:lA:2416:U:OP2	5:lE:239:HIS:ND1	2.40	0.55
1:lA:2905:U:H5''	14:lN:265:LYS:HG2	1.88	0.55
43:lq:72:CYS:HB3	43:lq:77:TYR:H	1.72	0.55
45:sB:6:ARG:HB3	52:sa:1943:C:H5	1.72	0.55
52:sa:1284:U:O4	52:sa:1285:A:N6	2.40	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:729:U:O4	1:1A:730:A:N6	2.40	0.55
1:1A:1610:A:O2'	1:1A:1651:C:O2	2.25	0.55
1:1A:1744:A:H2'	1:1A:1745:A:H8	1.72	0.55
1:1A:1763:C:H2'	1:1A:1764:A:H8	1.72	0.55
1:1A:3214:C:O3'	1:1A:3493:G:N2	2.40	0.55
11:1K:93:THR:HG22	11:1K:94:LYS:HG3	1.88	0.55
43:lq:73:THR:O	43:lq:76:LYS:NZ	2.33	0.55
56:sf:239:ASP:OD1	56:sf:239:ASP:N	2.37	0.55
1:1A:591:C:H2'	1:1A:592:A:H8	1.72	0.55
1:1A:3008:U:O2'	1:1A:3009:C:OP1	2.22	0.55
13:1M:95:ASN:ND2	13:1M:103:GLY:O	2.40	0.55
52:sa:1196:A:O2'	52:sa:1736:A:N3	2.37	0.55
52:sa:1371:U:C4	52:sa:1373:A:N7	2.75	0.55
1:1A:1071:A:H4'	1:1A:1087:G:N2	2.22	0.54
18:1R:53:ILE:HG13	18:1R:55:LYS:HG2	1.88	0.54
1:1A:446:A:H5'	1:1A:447:G:H2'	1.89	0.54
1:1A:1251:G:N2	12:1L:116:ARG:HH22	2.04	0.54
1:1A:2848:A:N1	1:1A:2911:U:H5	2.05	0.54
5:1E:84:MET:HE1	5:1E:183:ILE:HD11	1.89	0.54
25:1Y:78:VAL:HG11	25:1Y:103:VAL:HG11	1.90	0.54
52:sa:1097:C:H4'	64:so:11:ILE:HD11	1.89	0.54
52:sa:1253:G:O2'	52:sa:1254:G:OP1	2.23	0.54
61:sk:59:MET:HA	61:sk:62:LYS:HE3	1.89	0.54
1:1A:106:G:OP2	14:1N:71:ARG:NH2	2.41	0.54
47:sD:60:GLU:OE1	57:sg:206:ARG:NH1	2.40	0.54
52:sa:887:A:H2'	52:sa:888:A:C8	2.38	0.54
54:sd:115:GLN:HE21	54:sd:136:VAL:HG13	1.71	0.54
1:1A:2037:C:O2'	1:1A:2043:A:N1	2.38	0.54
1:1A:3302:U:H2'	1:1A:3303:A:H8	1.72	0.54
10:1J:72:THR:HG21	10:1J:178:LYS:HG3	1.88	0.54
14:1N:77:GLU:HG3	14:1N:107:LEU:HD12	1.89	0.54
27:la:75:LYS:HB2	27:la:77:VAL:HG22	1.88	0.54
52:sa:346:U:O4	63:sm:101:LYS:NZ	2.29	0.54
52:sa:941:C:OP2	64:so:14:ARG:NH1	2.39	0.54
52:sa:1533:C:H3'	52:sa:1534:A:H4'	1.88	0.54
65:sp:73:ALA:HB1	65:sp:114:LEU:HG	1.89	0.54
1:1A:387:G:OP2	6:1F:194:GLN:NE2	2.29	0.54
1:1A:505:A:H2'	1:1A:506:A:C8	2.43	0.54
1:1A:2233:G:N7	4:1D:152:SER:OG	2.37	0.54
1:1A:3043:A:N3	1:1A:3179:C:O2'	2.35	0.54
3:1C:6:G:OP2	7:1G:28:THR:OG1	2.24	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:IG:50:ARG:NH1	7:IG:72:ASP:OD2	2.40	0.54
13:IM:116:TYR:O	70:su:102:ASN:ND2	2.41	0.54
27:la:47:ILE:HD11	27:la:107:LEU:HD21	1.88	0.54
31:le:14:VAL:HA	31:le:17:LYS:HD2	1.89	0.54
43:lq:63:THR:O	43:lq:85:ARG:NH1	2.36	0.54
52:sa:151:A:H2'	52:sa:152:A:O4'	2.07	0.54
52:sa:1880:U:H2'	52:sa:1881:C:H6	1.73	0.54
54:sd:188:CYS:HB2	54:sd:195:ILE:HD11	1.89	0.54
1:lA:1554:C:OP2	6:lF:196:LYS:NZ	2.41	0.54
3:lC:27:G:H5''	13:lM:137:ARG:HD3	1.89	0.54
6:lF:85:GLY:O	6:lF:89:GLN:NE2	2.40	0.54
13:lM:94:ASN:ND2	13:lM:94:ASN:O	2.41	0.54
19:lS:36:ALA:HB1	19:lS:45:LYS:HG2	1.88	0.54
34:lh:15:THR:HG22	34:lh:16:LYS:H	1.73	0.54
50:sJ:24:C:H2'	50:sJ:25:U:C6	2.42	0.54
52:sa:1828:A:H2'	52:sa:1829:A:H8	1.72	0.54
54:sd:141:MET:HE2	54:sd:141:MET:HA	1.90	0.54
1:lA:189:G:N7	10:lJ:134:ASN:HB2	2.23	0.54
1:lA:662:U:H2'	1:lA:663:G:O4'	2.08	0.54
1:lA:1639:U:H2'	1:lA:1640:A:C8	2.43	0.54
2:lB:95:A:OP1	35:li:59:ARG:NH2	2.41	0.54
8:lH:113:THR:HG22	8:lH:204:PHE:HB3	1.88	0.54
10:lJ:168:LYS:NZ	37:lk:40:GLU:OE2	2.38	0.54
50:sJ:4:U:H2'	50:sJ:5:U:O4'	2.07	0.54
52:sa:752:A:P	61:sk:51:HIS:HE2	2.30	0.54
52:sa:1838:A:H2'	52:sa:1839:A:C8	2.43	0.54
56:sf:34:HIS:HB2	56:sf:39:CYS:SG	2.48	0.54
63:sm:108:VAL:HG12	63:sm:137:PHE:HB2	1.90	0.54
1:lA:1553:A:OP1	2:lB:22:G:O2'	2.26	0.54
1:lA:3432:A:O2'	1:lA:3433:U:OP1	2.26	0.54
26:lZ:49:ARG:HG2	26:lZ:60:LEU:HD11	1.89	0.54
52:sa:967:G:N2	52:sa:993:A:OP1	2.41	0.54
54:sd:204:LEU:HD13	54:sd:208:PRO:HG2	1.90	0.54
14:lN:214:THR:OG1	14:lN:215:LYS:N	2.39	0.54
17:lQ:80:VAL:HB	17:lQ:82:ARG:HD2	1.90	0.54
32:lf:84:THR:OG1	32:lf:127:THR:OG1	2.26	0.54
52:sa:401:U:H2'	52:sa:402:A:H8	1.73	0.54
1:lA:3196:U:H2'	1:lA:3197:C:H6	1.73	0.54
9:II:66:GLU:OE2	9:II:66:GLU:N	2.38	0.54
15:lO:160:GLU:O	15:lO:164:ARG:NH1	2.41	0.54
23:lW:33:ASP:HB3	23:lW:117:GLU:HA	1.90	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:la:22:ASN:OD1	27:la:22:ASN:N	2.40	0.54
52:sa:599:A:OP2	52:sa:600:A:O2'	2.25	0.54
52:sa:1573:G:O2'	52:sa:1768:C:OP1	2.26	0.54
56:sf:123:LYS:HB2	56:sf:224:PHE:CE1	2.42	0.54
60:sj:10:LYS:HZ3	63:sm:133:LYS:HD2	1.73	0.54
74:sy:22:ASP:OD2	74:sy:25:TRP:HB3	2.08	0.54
1:lA:968:C:H2'	1:lA:969:U:C6	2.44	0.53
1:lA:1474:A:OP2	19:lS:30:ARG:NH2	2.40	0.53
1:lA:3285:A:C6	1:lA:3287:U:H1'	2.43	0.53
1:lA:3443:C:HO2'	1:lA:3447:A:HO2'	1.51	0.53
1:lA:3500:A:H2'	1:lA:3501:A:H8	1.73	0.53
17:lQ:114:ARG:HD3	17:lQ:151:ILE:HG12	1.91	0.53
27:la:47:ILE:HG22	27:la:120:LYS:HE3	1.89	0.53
50:sJ:62:C:H2'	50:sJ:63:C:C6	2.43	0.53
52:sa:144:C:H2'	52:sa:145:U:O4'	2.08	0.53
52:sa:585:A:H2'	52:sa:586:A:C8	2.44	0.53
52:sa:1196:A:H2'	52:sa:1197:G:C8	2.43	0.53
1:lA:3476:C:H5''	1:lA:3477:U:C2	2.44	0.53
6:lF:63:SER:O	6:lF:63:SER:OG	2.24	0.53
14:lN:74:THR:OG1	14:lN:77:GLU:OE1	2.24	0.53
58:sh:89:ASN:N	58:sh:89:ASN:OD1	2.42	0.53
63:sm:132:SER:OG	63:sm:133:LYS:N	2.38	0.53
67:sr:11:LEU:HD22	67:sr:72:CYS:HB2	1.90	0.53
67:sr:102:LEU:HB2	67:sr:113:ASN:HB3	1.90	0.53
1:lA:87:G:N2	1:lA:91:A:OP2	2.39	0.53
1:lA:572:U:H2'	1:lA:573:A:C8	2.43	0.53
1:lA:724:A:O2'	1:lA:725:A:H5''	2.09	0.53
1:lA:1626:G:N2	34:lh:4:ARG:HD2	2.24	0.53
1:lA:2884:U:H3'	1:lA:2885:G:N2	2.22	0.53
4:lD:19:ARG:NH1	4:lD:189:TYR:O	2.42	0.53
11:lK:13:ILE:HD11	11:lK:63:VAL:HG23	1.91	0.53
52:sa:247:G:OP1	56:sf:132:GLY:N	2.28	0.53
1:lA:206:U:O4	1:lA:303:G:O6	2.27	0.53
1:lA:1749:U:OP2	21:lU:42:ARG:NH2	2.41	0.53
1:lA:1813:G:N7	28:lb:17:ARG:NH2	2.56	0.53
1:lA:3444:A:H4'	1:lA:3447:A:H1'	1.89	0.53
1:lA:3497:U:H4'	1:lA:3498:U:H5'	1.91	0.53
3:lC:38:A:H2	13:lM:70:THR:HG23	1.72	0.53
7:lG:45:LYS:HG3	22:lV:33:THR:HG21	1.90	0.53
8:lH:135:ILE:HA	8:lH:159:LYS:HD2	1.89	0.53
52:sa:13:C:O2'	52:sa:1323:U:O4	2.23	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:sa:1822:U:H5'	52:sa:1823:U:H5''	1.90	0.53
56:sf:105:ARG:NH2	56:sf:185:ASN:OD1	2.41	0.53
1:lA:3492:G:H5'	5:lE:385:LYS:HD2	1.89	0.53
2:lB:57:U:C2	2:lB:58:G:C8	2.97	0.53
3:lC:26:U:H3	3:lC:51:G:N2	2.04	0.53
3:lC:104:G:H2'	3:lC:105:A:C8	2.44	0.53
52:sa:134:A:N1	52:sa:172:G:N2	2.56	0.53
52:sa:1248:U:H2'	52:sa:1249:A:H8	1.74	0.53
52:sa:1594:A:H2'	52:sa:1595:G:C8	2.43	0.53
56:sf:86:ASP:HA	56:sf:120:LYS:HG3	1.91	0.53
1:lA:220:A:H2'	1:lA:221:U:H6	1.74	0.53
1:lA:829:G:N2	6:lF:236:ASN:OD1	2.29	0.53
1:lA:836:C:H2'	1:lA:837:A:H8	1.74	0.53
1:lA:1475:A:HO2'	27:la:185:ARG:HH22	1.51	0.53
1:lA:2318:U:H5''	4:lD:244:GLY:HA3	1.91	0.53
1:lA:2348:G:N2	1:lA:2348:G:OP2	2.40	0.53
1:lA:2427:U:H2'	1:lA:2428:A:H8	1.73	0.53
5:lE:122:TRP:CE2	5:lE:127:LYS:HG2	2.43	0.53
7:lG:131:ASP:CG	7:lG:176:HIS:HE2	2.16	0.53
52:sa:540:U:H2'	52:sa:541:U:C6	2.44	0.53
52:sa:586:A:O2'	52:sa:590:U:OP1	2.26	0.53
52:sa:1680:G:O2'	52:sa:1682:A:N3	2.42	0.53
1:lA:897:A:O2'	1:lA:899:A:N1	2.42	0.53
1:lA:2213:U:OP2	1:lA:2218:A:N6	2.38	0.53
1:lA:2491:C:OP1	4:lD:2:GLY:N	2.41	0.53
1:lA:2499:U:H2'	1:lA:2500:A:C8	2.43	0.53
1:lA:3412:U:H5''	1:lA:3413:U:OP2	2.09	0.53
12:lL:30:ARG:HH21	12:lL:66:GLU:HG2	1.73	0.53
1:lA:2225:A:C2	1:lA:2264:A:H4'	2.44	0.53
1:lA:2851:A:H2'	1:lA:2852:G:H8	1.74	0.53
5:lE:307:THR:HG23	5:lE:311:GLY:HA2	1.90	0.53
52:sa:1396:A:C2	52:sa:1425:A:C5	2.96	0.53
1:lA:200:A:N6	1:lA:309:G:H21	2.06	0.53
1:lA:243:G:N2	1:lA:418:A:N7	2.57	0.53
1:lA:699:A:H5''	1:lA:746:A:H62	1.74	0.53
1:lA:2878:A:N7	37:lk:86:ARG:NH2	2.56	0.53
1:lA:3393:A:H2'	1:lA:3394:A:C8	2.43	0.53
23:IW:84:LEU:HD12	23:IW:89:MET:HE1	1.89	0.53
42:lp:29:PRO:HG2	42:lp:32:ALA:HB2	1.91	0.53
52:sa:1033:A:H61	52:sa:1091:C:N4	2.07	0.53
56:sf:201:GLY:HA2	63:sm:39:MET:SD	2.48	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
72:sw:64:THR:HG22	72:sw:77:TRP:HE3	1.74	0.53
1:lA:440:G:N1	1:lA:443:A:OP2	2.34	0.53
1:lA:466:G:O2'	1:lA:2441:U:OP2	2.24	0.53
1:lA:721:A:H2'	1:lA:722:A:H8	1.73	0.53
1:lA:2331:A:C8	52:sa:1905:U:H1'	2.44	0.53
1:lA:3142:G:H21	1:lA:3143:A:H62	1.57	0.53
32:lf:61:GLU:H	32:lf:66:ARG:HH21	1.56	0.53
41:lo:24:PRO:HG2	41:lo:27:PHE:HB2	1.90	0.53
52:sa:251:C:H2'	52:sa:252:C:C6	2.44	0.53
52:sa:1776:G:OP1	57:sg:53:HIS:NE2	2.39	0.53
54:sd:115:GLN:NE2	54:sd:136:VAL:HG13	2.24	0.53
54:sd:175:GLY:O	54:sd:178:THR:OG1	2.25	0.53
1:lA:998:U:H2'	18:IR:131:ARG:HH21	1.74	0.52
1:lA:2101:G:O2'	1:lA:2410:U:O4	2.24	0.52
1:lA:3446:C:O2'	1:lA:3447:A:H5'	2.09	0.52
24:IX:107:ASN:HD21	24:IX:111:GLU:HB2	1.73	0.52
41:lo:9:TYR:HE1	41:lo:51:MET:HE2	1.73	0.52
52:sa:882:G:OP2	65:sp:33:ASN:ND2	2.36	0.52
52:sa:888:A:C2	52:sa:989:U:H1'	2.44	0.52
52:sa:1396:A:H2'	52:sa:1425:A:H8	1.72	0.52
52:sa:1699:C:H4'	52:sa:1705:A:N6	2.24	0.52
56:sf:17:MET:HE2	56:sf:106:ARG:HD2	1.91	0.52
57:sg:30:VAL:HG13	57:sg:119:ILE:HD11	1.90	0.52
59:si:132:LEU:HD21	59:si:141:VAL:HG11	1.90	0.52
62:sl:11:ILE:HG13	62:sl:34:PHE:HE2	1.73	0.52
1:lA:365:G:OP1	17:IQ:47:ARG:NH2	2.31	0.52
1:lA:2509:U:O4	17:IQ:20:ARG:NH1	2.42	0.52
20:IT:106:LEU:HD13	20:IT:110:TYR:HD2	1.74	0.52
29:lc:64:LYS:HB2	29:lc:67:LYS:HE3	1.90	0.52
37:lk:2:ALA:HB1	37:lk:13:TYR:HE1	1.74	0.52
52:sa:164:A:O2'	52:sa:166:A:OP2	2.26	0.52
52:sa:846:G:OP1	64:so:2:GLY:N	2.42	0.52
52:sa:872:A:H2'	52:sa:873:A:H8	1.71	0.52
61:sk:33:GLN:HG2	61:sk:34:PHE:HD1	1.74	0.52
63:sm:62:ASP:OD1	63:sm:62:ASP:N	2.41	0.52
1:lA:228:A:H2'	1:lA:229:A:C8	2.44	0.52
1:lA:412:A:H5''	6:IF:97:ARG:HH22	1.73	0.52
1:lA:1220:A:H2'	1:lA:1220:A:N3	2.24	0.52
1:lA:1308:A:OP1	16:IP:55:ASN:ND2	2.37	0.52
1:lA:3088:G:O2'	1:lA:3091:C:OP2	2.25	0.52
9:II:106:LYS:HE2	9:II:183:PHE:HD2	1.74	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:lN:46:PRO:HB3	35:li:108:PHE:CZ	2.44	0.52
46:sC:30:ASN:N	46:sC:30:ASN:OD1	2.42	0.52
52:sa:389:U:H3	52:sa:396:A:N6	2.02	0.52
52:sa:864:C:H2'	52:sa:865:A:H8	1.72	0.52
70:su:119:ARG:HD2	70:su:119:ARG:O	2.09	0.52
1:lA:191:A:H2'	1:lA:192:A:H8	1.73	0.52
1:lA:500:G:O6	1:lA:572:U:O4	2.27	0.52
1:lA:1030:C:OP1	4:lD:14:SER:OG	2.22	0.52
3:lC:53:C:HO2'	13:lM:152:HIS:HD1	1.55	0.52
5:lE:386:LYS:N	5:lE:386:LYS:HD3	2.25	0.52
33:lg:96:GLU:HB2	33:lg:121:LEU:HD12	1.90	0.52
34:lh:65:LEU:O	34:lh:70:ARG:NH1	2.42	0.52
47:sD:51:ARG:HD2	57:sg:63:PHE:HD2	1.74	0.52
52:sa:169:U:H2'	52:sa:170:G:C8	2.44	0.52
52:sa:465:G:C2	52:sa:466:A:C8	2.98	0.52
52:sa:746:A:H2'	52:sa:747:A:C8	2.44	0.52
52:sa:1014:C:OP1	64:so:9:ARG:NH2	2.42	0.52
56:sf:159:LYS:HB3	56:sf:169:ASP:H	1.74	0.52
60:sj:4:THR:HG21	60:sj:24:LYS:HG2	1.92	0.52
1:lA:1938:G:O6	40:ln:35:LYS:NZ	2.42	0.52
1:lA:3162:A:OP1	15:lO:73:HIS:ND1	2.41	0.52
23:lW:66:ILE:HG13	23:lW:79:THR:HG21	1.92	0.52
25:lY:37:ASP:N	25:lY:37:ASP:OD1	2.41	0.52
57:sg:203:LYS:HG3	57:sg:206:ARG:HH21	1.74	0.52
1:lA:2079:A:H2'	1:lA:2080:A:O4'	2.10	0.52
28:lb:45:GLY:O	28:lb:70:SER:OG	2.21	0.52
45:sB:4:LYS:NZ	52:sa:1795:G:OP1	2.35	0.52
52:sa:799:C:O2	67:sr:124:ARG:NH1	2.40	0.52
52:sa:866:G:H2'	52:sa:867:A:C8	2.45	0.52
52:sa:1690:A:N1	52:sa:1774:A:H1'	2.25	0.52
52:sa:1701:U:H4'	52:sa:1702:G:O5'	2.09	0.52
57:sg:88:LYS:O	57:sg:92:MET:HG2	2.09	0.52
60:sj:65:PHE:HZ	60:sj:78:ILE:HG23	1.75	0.52
67:sr:77:PRO:HB2	74:sy:5:LEU:HB2	1.90	0.52
1:lA:378:A:N6	1:lA:379:G:O6	2.43	0.52
1:lA:597:A:H2'	1:lA:598:A:H8	1.75	0.52
1:lA:720:U:H2'	1:lA:721:A:H8	1.74	0.52
1:lA:1528:A:N3	1:lA:1553:A:O2'	2.41	0.52
1:lA:2752:C:O2'	13:lM:18:VAL:HG11	2.10	0.52
1:lA:3204:C:O2	24:lX:89:ARG:NH1	2.39	0.52
2:lB:5:A:HO2'	18:lR:61:ARG:HH11	1.57	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:IV:52:LEU:HD12	22:IV:53:PRO:HD2	1.91	0.52
33:lg:41:ASN:O	33:lg:45:ARG:HG3	2.09	0.52
52:sa:1382:C:H42	52:sa:1390:A:H61	1.57	0.52
53:sc:165:GLY:O	53:sc:168:ARG:NH1	2.41	0.52
54:sd:176:ASN:OD1	54:sd:180:GLN:NE2	2.43	0.52
1:lA:1250:U:OP2	12:lL:13:ARG:NH2	2.43	0.52
1:lA:1437:A:H2'	1:lA:1438:U:H6	1.74	0.52
1:lA:1498:C:N4	1:lA:1502:A:O3'	2.43	0.52
1:lA:2064:G:N1	1:lA:2067:A:OP2	2.43	0.52
32:lf:121:GLU:HB3	32:lf:124:LYS:HE2	1.92	0.52
52:sa:632:U:OP2	67:sr:32:LYS:NZ	2.42	0.52
72:sw:22:ILE:N	72:sw:89:LEU:O	2.43	0.52
74:sy:32:THR:HG23	74:sy:36:CYS:HB2	1.90	0.52
1:lA:2439:A:H2'	1:lA:2440:G:C8	2.45	0.52
1:lA:3021:G:H2'	1:lA:3022:C:H6	1.74	0.52
1:lA:3196:U:H2'	1:lA:3197:C:C6	2.45	0.52
8:lH:141:ALA:HB3	8:lH:144:LYS:HD2	1.91	0.52
16:lP:24:LEU:HB2	16:lP:45:THR:HG21	1.92	0.52
17:lQ:11:ASN:ND2	17:lQ:44:ASN:OD1	2.43	0.52
68:ss:55:ASN:O	68:ss:57:PRO:HD3	2.10	0.52
1:lA:836:C:H2'	1:lA:837:A:C8	2.45	0.52
1:lA:910:A:H5''	6:lF:113:ARG:HD2	1.90	0.52
1:lA:1006:A:H2'	1:lA:1007:C:C6	2.45	0.52
1:lA:2411:G:N2	1:lA:2415:C:O2'	2.39	0.52
14:lN:122:VAL:HG13	14:lN:223:GLN:HG3	1.92	0.52
50:sJ:52:C:H2'	50:sJ:53:G:O4'	2.10	0.52
52:sa:1441:C:O2'	52:sa:1442:A:O5'	2.28	0.52
52:sa:1442:A:H2'	52:sa:1443:U:H6	1.74	0.52
1:lA:2331:A:H8	52:sa:1904:G:H21	1.55	0.51
6:lF:184:VAL:HG11	6:lF:228:GLY:HA3	1.92	0.51
48:sE:34:TYR:OH	52:sa:1594:A:OP1	2.20	0.51
52:sa:327:U:OP1	60:sj:56:ARG:NH2	2.44	0.51
52:sa:1158:A:H2'	52:sa:1159:A:H8	1.75	0.51
1:lA:250:G:H2'	1:lA:251:A:C8	2.46	0.51
1:lA:341:G:OP2	17:lQ:15:GLN:NE2	2.42	0.51
1:lA:1841:G:H22	1:lA:1975:G:N2	2.06	0.51
1:lA:2331:A:H2	1:lA:2336:U:H3	1.59	0.51
1:lA:2950:U:O2'	1:lA:2952:C:OP1	2.28	0.51
3:lC:35:A:H8	3:lC:35:A:O5'	1.93	0.51
6:lF:7:VAL:HG13	6:lF:149:GLN:HE22	1.75	0.51
52:sa:1301:U:H2'	52:sa:1302:G:H8	1.75	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:lA:1208:U:H4'	30:ld:43:THR:HG23	1.91	0.51
1:lA:1219:A:H4'	1:lA:1220:A:H5'	1.92	0.51
1:lA:3154:A:H2'	1:lA:3155:U:H6	1.75	0.51
52:sa:327:U:H5''	60:sj:31:ARG:HH11	1.74	0.51
52:sa:802:U:C4	52:sa:803:G:O6	2.63	0.51
67:sr:31:SER:OG	67:sr:32:LYS:N	2.43	0.51
1:lA:2109:C:O2	5:lE:242:ARG:NH2	2.40	0.51
1:lA:3163:G:O2'	5:lE:14:LEU:O	2.27	0.51
2:lB:74:A:OP2	27:la:51:LYS:HB2	2.10	0.51
18:lR:18:ARG:NH1	18:lR:20:ASP:OD1	2.39	0.51
28:lb:36:ARG:HH12	28:lb:74:VAL:HG11	1.76	0.51
52:sa:1566:C:H5'	70:su:130:LEU:HD22	1.93	0.51
53:sc:208:THR:HG22	53:sc:211:ASN:H	1.76	0.51
65:sp:19:THR:OG1	65:sp:20:LYS:N	2.42	0.51
23:IW:39:ASP:OD1	23:IW:39:ASP:N	2.39	0.51
52:sa:319:U:OP1	63:sm:133:LYS:NZ	2.31	0.51
1:lA:250:G:H2'	1:lA:251:A:H8	1.76	0.51
1:lA:1847:A:H2'	1:lA:1848:A:C8	2.46	0.51
1:lA:1921:A:H2'	1:lA:1922:U:O2	2.09	0.51
1:lA:2379:A:O2'	52:sa:1894:G:O2'	2.03	0.51
31:le:46:VAL:HB	31:le:71:VAL:HG12	1.93	0.51
32:lf:113:MET:HG2	32:lf:129:VAL:HG22	1.91	0.51
52:sa:1816:U:H2'	52:sa:1817:A:H8	1.72	0.51
56:sf:171:VAL:HG23	56:sf:228:LYS:HB2	1.91	0.51
63:sm:4:GLN:NE2	63:sm:10:GLN:O	2.44	0.51
1:lA:114:U:P	10:lJ:142:ARG:HH22	2.33	0.51
1:lA:213:A:H62	14:lN:131:THR:HG1	1.57	0.51
1:lA:598:A:H2'	1:lA:599:U:C6	2.46	0.51
1:lA:625:A:O2'	1:lA:626:A:OP1	2.28	0.51
1:lA:1068:U:O2'	1:lA:1090:A:OP1	2.23	0.51
1:lA:2220:A:H1'	1:lA:2357:A:N6	2.26	0.51
1:lA:3090:G:N3	5:lE:252:ALA:HB1	2.25	0.51
52:sa:1546:C:H2'	52:sa:1547:C:H6	1.76	0.51
57:sg:183:SER:OG	57:sg:184:LYS:N	2.43	0.51
1:lA:765:A:H2'	1:lA:766:U:C6	2.45	0.51
1:lA:2743:G:N2	13:lM:20:ASN:HD21	2.09	0.51
5:lE:261:ARG:HB2	15:lO:65:THR:HG21	1.93	0.51
12:lL:93:PRO:HB2	12:lL:125:SER:HB3	1.91	0.51
23:IW:29:SER:HA	23:IW:78:THR:HA	1.93	0.51
52:sa:865:A:OP1	55:se:218:LYS:NZ	2.36	0.51
62:sl:15:LEU:HD13	62:sl:21:ILE:HG22	1.93	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:1090:A:H2'	1:1A:1091:A:H8	1.76	0.51
1:1A:2747:G:N2	1:1A:2750:A:OP2	2.44	0.51
8:1H:23:THR:OG1	8:1H:25:ASP:OD1	2.29	0.51
27:1a:87:LYS:O	27:1a:89:ASN:N	2.43	0.51
27:1a:179:MET:HE1	27:1a:187:TYR:CD2	2.46	0.51
37:1k:59:LEU:HD23	37:1k:62:LYS:HE2	1.93	0.51
50:sJ:23:G:H2'	50:sJ:24:C:C5	2.46	0.51
52:sa:325:U:H2'	52:sa:326:A:C8	2.46	0.51
1:1A:1476:G:H2'	1:1A:1476:G:N3	2.26	0.51
13:1M:27:GLY:O	13:1M:29:ARG:N	2.44	0.51
46:sC:31:SER:O	52:sa:939:G:N2	2.43	0.51
52:sa:1693:A:H2'	52:sa:1694:A:H8	1.74	0.51
55:se:197:LYS:HD3	55:se:197:LYS:C	2.35	0.51
56:sf:123:LYS:HB2	56:sf:224:PHE:HE1	1.75	0.51
57:sg:65:LYS:HD2	57:sg:146:LEU:HD21	1.93	0.51
1:1A:940:U:H2'	1:1A:941:A:C8	2.46	0.50
1:1A:2699:U:O2'	1:1A:2828:A:O2'	2.29	0.50
1:1A:3033:C:O2'	1:1A:3076:A:N3	2.31	0.50
1:1A:3093:G:OP2	1:1A:3093:G:N2	2.39	0.50
13:1M:16:LYS:HE2	13:1M:72:ARG:HH22	1.75	0.50
18:1R:57:ILE:HD12	18:1R:83:TRP:NE1	2.27	0.50
30:ld:2:SER:O	30:ld:2:SER:OG	2.26	0.50
52:sa:839:A:H8	59:si:119:PRO:HA	1.77	0.50
53:sc:105:ALA:HB2	53:sc:115:LEU:HD12	1.93	0.50
69:st:34:VAL:O	69:st:38:VAL:HG22	2.11	0.50
1:1A:495:A:H1'	1:1A:496:A:C8	2.45	0.50
1:1A:564:A:H4'	1:1A:565:U:O5'	2.12	0.50
1:1A:1184:A:OP2	30:ld:26:LYS:NZ	2.44	0.50
1:1A:1342:U:H2'	1:1A:1343:A:H4'	1.91	0.50
9:1I:87:ILE:HG21	9:1I:121:TYR:CE2	2.45	0.50
16:1P:99:GLU:O	16:1P:103:LEU:HD13	2.11	0.50
17:1Q:201:LYS:O	17:1Q:204:ARG:NH1	2.44	0.50
19:1S:63:THR:HA	19:1S:66:LEU:HD12	1.93	0.50
22:1V:80:VAL:HG12	22:1V:81:ASN:N	2.21	0.50
52:sa:252:C:H2'	52:sa:253:A:C8	2.46	0.50
52:sa:1823:U:H3'	52:sa:1824:G:H4'	1.93	0.50
1:1A:940:U:H2'	1:1A:941:A:H8	1.76	0.50
1:1A:1239:U:OP1	29:1c:23:GLY:N	2.39	0.50
1:1A:1254:A:H1'	1:1A:2969:U:O2'	2.11	0.50
3:1C:10:A:H4'	3:1C:12:C:C6	2.46	0.50
5:1E:27:ALA:HB2	5:1E:222:VAL:HG23	1.92	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:sa:1808:G:N3	52:sa:1928:A:O2'	2.43	0.50
53:sc:175:PRO:HG3	61:sk:57:LYS:HD3	1.93	0.50
61:sk:48:MET:O	61:sk:52:MET:HE3	2.09	0.50
71:sv:99:LYS:NZ	71:sv:103:TYR:HB2	2.26	0.50
1:lA:2179:U:H4'	21:lU:89:THR:HG22	1.93	0.50
1:lA:2223:A:H2'	1:lA:2224:U:O4'	2.12	0.50
1:lA:3274:U:H4'	42:lp:29:PRO:HG3	1.93	0.50
11:lK:4:LEU:HD23	11:lK:7:LYS:HB2	1.94	0.50
41:lo:49:MET:HE3	41:lo:51:MET:HG2	1.94	0.50
52:sa:1685:C:H2'	52:sa:1686:U:O4'	2.11	0.50
55:se:45:VAL:HG21	55:se:70:VAL:HG21	1.94	0.50
56:sf:112:LEU:HD22	56:sf:116:GLN:HG3	1.94	0.50
68:ss:22:VAL:HG13	68:ss:37:CYS:HB3	1.93	0.50
1:lA:720:U:O2'	1:lA:721:A:OP1	2.23	0.50
1:lA:1403:C:H2'	1:lA:1404:C:H6	1.75	0.50
3:lC:17:G:H1	3:lC:59:G:H1	1.59	0.50
3:lC:103:U:H2'	3:lC:104:G:C8	2.47	0.50
6:lF:370:ASN:H	6:lF:373:VAL:HG22	1.76	0.50
12:lL:30:ARG:HG3	12:lL:63:GLU:HG3	1.93	0.50
14:lN:46:PRO:HA	14:lN:49:ILE:HB	1.93	0.50
27:la:56:LYS:NZ	27:la:63:LYS:O	2.42	0.50
42:lp:24:CYS:HA	42:lp:40:CYS:HB3	1.94	0.50
52:sa:1719:G:N2	52:sa:1721:A:H3'	2.26	0.50
55:se:108:THR:HG23	55:se:111:LYS:H	1.77	0.50
1:lA:688:A:H2'	1:lA:689:A:C8	2.46	0.50
1:lA:1601:G:N2	1:lA:1608:A:OP2	2.37	0.50
14:lN:260:VAL:HG11	29:lc:132:GLU:HG2	1.93	0.50
16:lP:124:ALA:O	16:lP:128:ILE:HG12	2.11	0.50
52:sa:32:G:H22	52:sa:463:A:H5'	1.75	0.50
1:lA:219:G:H2'	1:lA:220:A:H8	1.77	0.50
12:lL:152:PHE:HB3	12:lL:165:LEU:HD21	1.93	0.50
23:lW:57:GLN:HB2	23:lW:59:ARG:NH2	2.27	0.50
46:sC:6:VAL:HG12	73:sx:84:MET:HG2	1.94	0.50
52:sa:1173:A:O2'	52:sa:1802:C:OP2	2.28	0.50
56:sf:198:VAL:HA	56:sf:204:GLU:HG3	1.93	0.50
67:sr:78:ARG:HB3	67:sr:124:ARG:HB3	1.94	0.50
1:lA:1325:C:H42	1:lA:3000:C:H5''	1.77	0.50
1:lA:1327:A:OP2	1:lA:1425:A:N6	2.43	0.50
1:lA:2637:A:H61	1:lA:2649:G:H2'	1.77	0.50
1:lA:3150:A:H2'	1:lA:3151:A:C8	2.46	0.50
52:sa:468:A:H2'	52:sa:469:C:O4'	2.12	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:sa:574:A:O2'	52:sa:576:U:OP1	2.29	0.50
52:sa:1442:A:OP2	69:st:59:ARG:NE	2.44	0.50
1:lA:619:A:H2'	1:lA:619:A:N3	2.26	0.50
1:lA:817:A:H2'	1:lA:818:A:C8	2.47	0.50
1:lA:1979:U:HO2'	1:lA:1981:A:H8	1.58	0.50
1:lA:2864:A:C4	1:lA:2865:A:C2	2.99	0.50
11:lK:41:HIS:ND1	11:lK:77:VAL:HG11	2.26	0.50
14:lN:58:CYS:HB2	14:lN:63:PHE:O	2.12	0.50
22:IV:40:VAL:HA	22:IV:99:SER:H	1.77	0.50
22:IV:122:LEU:HD22	22:IV:127:LYS:HD2	1.93	0.50
45:sB:59:LEU:HD11	65:sp:123:ARG:NE	2.27	0.50
52:sa:14:C:H2'	52:sa:15:U:C6	2.47	0.50
52:sa:936:U:OP1	52:sa:1097:C:O2'	2.30	0.50
52:sa:1794:U:H2'	52:sa:1795:G:C8	2.47	0.50
1:lA:220:A:H2'	1:lA:221:U:C6	2.46	0.49
1:lA:360:U:O4	37:lk:22:LYS:NZ	2.42	0.49
1:lA:713:G:H3'	1:lA:714:U:C5'	2.42	0.49
1:lA:2839:A:H5''	43:lq:78:LYS:HD2	1.94	0.49
5:lE:229:ASN:ND2	5:lE:272:MET:SD	2.76	0.49
24:IX:24:SER:O	24:IX:24:SER:OG	2.30	0.49
27:la:153:ILE:HA	27:la:156:LYS:HE2	1.94	0.49
38:ll:19:CYS:SG	38:ll:20:LYS:N	2.85	0.49
63:sm:115:ARG:HH12	63:sm:154:ALA:HB3	1.77	0.49
68:ss:57:PRO:HA	68:ss:60:ARG:HH11	1.77	0.49
1:lA:5:U:H2'	1:lA:6:U:C6	2.47	0.49
1:lA:589:A:OP1	8:lH:35:ARG:NH1	2.45	0.49
1:lA:678:A:H2'	1:lA:679:U:C6	2.47	0.49
1:lA:904:A:OP2	19:lS:62:SER:OG	2.25	0.49
1:lA:2867:A:N6	1:lA:2887:G:H1	2.10	0.49
3:lC:42:U:C2	3:lC:43:A:C8	2.99	0.49
6:lF:299:GLU:HG2	6:lF:304:LEU:HD12	1.94	0.49
9:lI:68:ASP:HB3	22:IV:141:PRO:HB3	1.93	0.49
12:lL:47:PRO:HB2	12:lL:178:LYS:HE2	1.93	0.49
48:sE:33:LYS:O	48:sE:36:LEU:HD12	2.12	0.49
52:sa:114:G:OP1	63:sm:66:ARG:NH1	2.44	0.49
52:sa:764:C:H4'	52:sa:765:A:H5'	1.93	0.49
52:sa:1834:G:H1	52:sa:1880:U:H3	1.60	0.49
55:se:34:VAL:HG21	55:se:45:VAL:HG22	1.94	0.49
59:si:132:LEU:O	59:si:136:LEU:HD23	2.12	0.49
60:sj:207:SER:OG	60:sj:208:ARG:N	2.45	0.49
1:lA:3089:A:H5''	1:lA:3090:G:H5'	1.95	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:lA:3155:U:H2'	1:lA:3156:U:C6	2.46	0.49
1:lA:3430:U:O5'	1:lA:3431:U:H5''	2.12	0.49
21:lU:105:LEU:HD23	21:lU:138:LEU:HD23	1.94	0.49
24:lX:82:ILE:HD12	24:lX:121:VAL:HG22	1.94	0.49
26:lZ:6:GLN:HB3	26:lZ:15:ILE:HD12	1.93	0.49
28:lb:83:THR:HG22	34:lh:93:PHE:CZ	2.48	0.49
41:lo:9:TYR:CE1	41:lo:51:MET:HE2	2.47	0.49
1:lA:518:A:H2'	1:lA:519:A:O4'	2.13	0.49
1:lA:705:U:H2'	1:lA:706:A:H8	1.77	0.49
1:lA:773:G:OP1	33:lg:41:ASN:ND2	2.44	0.49
1:lA:3222:U:OP1	21:lU:59:SER:N	2.38	0.49
5:lE:41:PRO:HA	5:lE:187:GLY:HA3	1.94	0.49
50:sJ:42:C:H2'	50:sJ:43:G:C8	2.47	0.49
52:sa:399:G:H2'	52:sa:400:C:H6	1.78	0.49
52:sa:1596:U:O2'	52:sa:1681:A:N6	2.45	0.49
60:sj:35:ASN:OD1	60:sj:35:ASN:N	2.45	0.49
60:sj:67:TRP:CZ2	60:sj:70:GLN:HG3	2.47	0.49
66:sq:33:VAL:HG21	66:sq:44:TYR:CD2	2.47	0.49
1:lA:721:A:H2'	1:lA:722:A:C8	2.48	0.49
1:lA:903:G:H8	19:lS:90:TYR:CD2	2.30	0.49
1:lA:968:C:H2'	1:lA:969:U:H6	1.77	0.49
1:lA:1090:A:H2'	1:lA:1091:A:C8	2.47	0.49
1:lA:1896:G:OP2	31:le:36:LYS:NZ	2.45	0.49
1:lA:2732:A:H2'	1:lA:2733:A:H8	1.78	0.49
1:lA:2863:A:N1	37:lk:16:HIS:HB3	2.26	0.49
3:lC:6:G:H5''	7:lG:22:ARG:HD3	1.94	0.49
3:lC:89:G:H4'	12:lL:11:LEU:HD22	1.95	0.49
3:lC:103:U:H2'	3:lC:104:G:H8	1.76	0.49
4:lD:20:SER:O	4:lD:20:SER:OG	2.30	0.49
15:lO:29:LEU:HD21	15:lO:89:LEU:HD22	1.95	0.49
15:lO:76:SER:OG	15:lO:107:GLU:OE2	2.30	0.49
25:lY:44:PHE:CE1	25:lY:64:LEU:HD12	2.47	0.49
26:lZ:11:SER:HA	26:lZ:54:THR:HG23	1.94	0.49
60:sj:78:ILE:HG22	60:sj:104:ILE:HG22	1.95	0.49
63:sm:37:ILE:HG13	63:sm:38:GLY:H	1.76	0.49
63:sm:58:PRO:HB3	63:sm:65:ILE:HD11	1.93	0.49
67:sr:20:GLN:OE1	67:sr:22:LYS:NZ	2.46	0.49
1:lA:191:A:H2'	1:lA:192:A:C8	2.47	0.49
1:lA:510:A:N1	1:lA:561:G:O2'	2.42	0.49
1:lA:708:U:H2'	1:lA:709:U:C5	2.47	0.49
1:lA:2030:G:H5''	1:lA:2031:C:H5'	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1B:62:U:O4	35:1i:51:ASN:ND2	2.43	0.49
11:1K:120:ILE:HG12	11:1K:121:GLN:H	1.77	0.49
49:sI:14:A:N3	49:sI:14:A:H2'	2.28	0.49
52:sa:592:U:H2'	52:sa:593:A:C8	2.47	0.49
52:sa:751:A:H2'	52:sa:752:A:O4'	2.12	0.49
52:sa:781:G:C8	56:sf:17:MET:HE1	2.47	0.49
52:sa:845:A:OP1	67:sr:28:ARG:NH2	2.45	0.49
52:sa:1162:U:O3'	74:sy:58:LYS:NZ	2.45	0.49
56:sf:9:ARG:HH21	56:sf:18:LEU:HD22	1.78	0.49
1:1A:228:A:H2'	1:1A:229:A:H8	1.78	0.49
1:1A:3383:A:H5''	5:1E:156:ARG:HE	1.78	0.49
3:1C:28:U:H3'	3:1C:29:G:H21	1.78	0.49
28:1b:95:ILE:HD11	28:1b:121:PHE:CE1	2.48	0.49
52:sa:1266:A:C8	66:sq:78:HIS:CE1	3.01	0.49
52:sa:1923:A:H2'	52:sa:1924:G:H8	1.77	0.49
56:sf:173:PHE:HE1	56:sf:223:VAL:HG11	1.77	0.49
1:1A:811:A:N1	1:1A:839:G:O2'	2.41	0.49
1:1A:874:A:H2'	1:1A:875:C:H6	1.78	0.49
1:1A:1095:A:H2	1:1A:1099:A:H2	1.59	0.49
1:1A:1465:G:H2'	1:1A:1466:U:C6	2.48	0.49
1:1A:1633:G:OP2	38:1l:14:LYS:NZ	2.41	0.49
1:1A:1864:C:OP1	23:1W:98:LYS:NZ	2.41	0.49
9:1I:204:ASN:OD1	9:1I:204:ASN:N	2.45	0.49
13:1M:109:HIS:HE1	13:1M:122:ILE:HA	1.78	0.49
16:1P:100:LYS:O	16:1P:104:ARG:HG3	2.12	0.49
18:1R:126:ARG:HG3	18:1R:140:SER:HB3	1.95	0.49
47:sD:62:GLU:O	65:sp:112:ARG:NH2	2.43	0.49
52:sa:545:G:OP2	52:sa:575:U:O2'	2.24	0.49
52:sa:932:C:H2'	52:sa:933:A:H8	1.77	0.49
52:sa:1371:U:C5	52:sa:1373:A:C8	3.01	0.49
65:sp:31:THR:OG1	65:sp:32:PHE:N	2.43	0.49
70:su:97:SER:O	70:su:99:LEU:HD23	2.12	0.49
1:1A:333:U:O2'	17:1Q:180:ARG:O	2.30	0.49
1:1A:660:U:H2'	1:1A:661:G:H8	1.78	0.49
1:1A:779:A:H1'	1:1A:781:A:OP2	2.13	0.49
1:1A:868:A:H2'	1:1A:869:G:C8	2.48	0.49
1:1A:2340:U:HO2'	1:1A:2341:C:P	2.33	0.49
1:1A:2489:A:H2'	1:1A:2490:G:H8	1.78	0.49
1:1A:2702:G:C6	1:1A:2717:A:C6	3.01	0.49
2:1B:28:U:H2'	2:1B:29:U:C6	2.47	0.49
2:1B:29:U:H2'	2:1B:30:U:H6	1.78	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1B:141:G:H2'	2:1B:142:A:C8	2.44	0.49
2:1B:144:A:H2'	2:1B:145:U:C6	2.48	0.49
6:1F:369:LEU:C	6:1F:371:GLN:H	2.20	0.49
7:1G:115:MET:HE1	7:1G:143:PHE:HB2	1.95	0.49
21:1U:98:ARG:NH1	21:1U:130:ASN:OD1	2.46	0.49
52:sa:611:U:H5'	52:sa:1011:U:O4'	2.13	0.49
60:sj:25:LYS:O	60:sj:28:THR:OG1	2.31	0.49
66:sq:122:TYR:OH	70:su:125:ARG:NH1	2.46	0.49
1:1A:710:A:O2'	1:1A:736:G:N2	2.23	0.49
1:1A:1027:G:H4'	1:1A:1028:G:O5'	2.11	0.49
1:1A:1098:A:H2	1:1A:1184:A:H5''	1.78	0.49
1:1A:1686:G:N2	1:1A:1689:A:OP2	2.45	0.49
1:1A:2904:U:H2'	1:1A:2905:U:C6	2.46	0.49
52:sa:443:U:H2'	52:sa:444:U:C6	2.48	0.49
52:sa:1332:A:H2'	52:sa:1333:G:O4'	2.13	0.49
52:sa:1917:U:H2'	52:sa:1918:U:C6	2.48	0.49
53:sc:228:THR:OG1	53:sc:230:ASP:OD1	2.30	0.49
55:se:26:PHE:CD2	65:sp:83:LEU:HD21	2.48	0.49
1:1A:249:U:H2'	1:1A:250:G:H8	1.78	0.48
1:1A:797:C:H2'	1:1A:798:U:C6	2.48	0.48
1:1A:3211:A:O2'	32:lf:55:HIS:NE2	2.46	0.48
1:1A:3480:A:C5	1:1A:3481:U:C5	3.01	0.48
2:1B:24:U:C5	6:1F:197:MET:HE1	2.47	0.48
16:1P:32:ASP:OD1	16:1P:35:ARG:NH2	2.31	0.48
45:sB:34:LYS:NZ	52:sa:1940:G:N7	2.61	0.48
47:sD:51:ARG:HG3	57:sg:63:PHE:HE2	1.77	0.48
47:sD:66:ARG:NE	47:sD:66:ARG:O	2.46	0.48
52:sa:161:A:H2'	52:sa:162:U:O4'	2.12	0.48
52:sa:622:G:OP1	64:so:120:SER:OG	2.31	0.48
52:sa:1243:A:N1	52:sa:1551:A:O2'	2.42	0.48
55:se:174:ILE:HG23	55:se:195:LEU:HD21	1.94	0.48
1:1A:103:A:O2'	1:1A:370:A:N3	2.44	0.48
1:1A:118:A:N6	1:1A:189:G:O2'	2.46	0.48
1:1A:546:U:H2'	1:1A:547:A:H8	1.78	0.48
1:1A:712:G:O2'	1:1A:713:G:H5'	2.14	0.48
1:1A:1968:U:H2'	1:1A:1969:U:C6	2.48	0.48
1:1A:2378:G:H4'	52:sa:1821:A:H2'	1.95	0.48
2:1B:145:U:H2'	2:1B:146:U:C6	2.47	0.48
11:1K:134:ILE:HG21	11:1K:167:ILE:HG12	1.94	0.48
22:1V:164:GLU:OE2	22:1V:164:GLU:N	2.32	0.48
52:sa:170:G:H1'	52:sa:259:A:N6	2.28	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:sa:1241:C:O2	52:sa:1551:A:N6	2.46	0.48
52:sa:1371:U:C4	52:sa:1373:A:C8	3.01	0.48
52:sa:1772:A:H2'	52:sa:1773:G:C8	2.47	0.48
58:sh:70:ASN:OD1	58:sh:70:ASN:N	2.46	0.48
63:sm:68:ARG:HB3	63:sm:127:GLN:HB2	1.94	0.48
63:sm:98:ARG:HG2	74:sy:5:LEU:HA	1.93	0.48
1:lA:943:U:C2	1:lA:1021:G:N2	2.82	0.48
1:lA:1640:A:H2'	1:lA:1641:A:C8	2.49	0.48
1:lA:1745:A:H2'	1:lA:1746:G:H8	1.78	0.48
1:lA:1847:A:H2'	1:lA:1848:A:H8	1.76	0.48
1:lA:3023:U:OP1	24:IX:50:ASN:ND2	2.46	0.48
1:lA:3043:A:O2'	11:lK:180:GLN:OE1	2.14	0.48
6:lF:389:LYS:HA	6:lF:389:LYS:HD3	1.61	0.48
11:lK:29:THR:HG22	11:lK:34:SER:HB2	1.95	0.48
52:sa:787:U:H4'	52:sa:788:C:H5	1.79	0.48
52:sa:1440:A:H1'	52:sa:1441:C:H5	1.79	0.48
66:sq:106:ILE:HD13	66:sq:110:MET:HE3	1.95	0.48
1:lA:2469:G:P	1:lA:2469:G:O4'	2.71	0.48
1:lA:3336:G:H5'	1:lA:3373:A:H61	1.79	0.48
1:lA:3352:G:C5	20:lT:155:ALA:HB2	2.49	0.48
2:lB:50:A:N7	2:lB:53:G:N1	2.62	0.48
5:lE:251:VAL:HG21	5:lE:267:ALA:HB3	1.95	0.48
6:lF:210:VAL:HB	6:lF:231:THR:HG22	1.95	0.48
11:lK:142:VAL:HG22	11:lK:152:LEU:HG	1.95	0.48
22:IV:97:LYS:HG2	22:IV:98:PRO:HD2	1.95	0.48
28:lb:95:ILE:HD11	28:lb:121:PHE:HE1	1.78	0.48
43:lq:31:ASP:OD1	43:lq:31:ASP:N	2.46	0.48
52:sa:1693:A:H2'	52:sa:1694:A:C8	2.48	0.48
57:sg:167:ASN:OD1	57:sg:168:SER:N	2.47	0.48
1:lA:793:G:H2'	1:lA:794:A:N7	2.28	0.48
1:lA:975:G:OP1	21:lU:92:LYS:NZ	2.34	0.48
1:lA:1729:G:O2'	1:lA:1730:G:O4'	2.26	0.48
1:lA:1902:A:H4'	1:lA:1915:A:H4'	1.95	0.48
1:lA:2856:G:H2'	1:lA:2903:A:N6	2.28	0.48
1:lA:3314:U:O2'	1:lA:3417:G:OP2	2.19	0.48
5:lE:62:ARG:NH1	5:lE:350:THR:OG1	2.47	0.48
5:lE:234:ARG:HG3	5:lE:272:MET:HB3	1.95	0.48
17:lQ:121:ILE:HG13	17:lQ:131:GLU:HG3	1.94	0.48
20:lT:7:THR:HG22	20:lT:22:ARG:HA	1.94	0.48
34:lh:62:TYR:O	34:lh:70:ARG:NH1	2.45	0.48
40:ln:49:ASP:HB3	40:ln:52:LYS:HG3	1.96	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
45:sB:37:LYS:HG3	45:sB:72:GLU:HG2	1.95	0.48
49:sI:4:C:H2'	49:sI:5:C:C6	2.48	0.48
52:sa:26:U:H2'	52:sa:27:A:C8	2.49	0.48
68:ss:50:PRO:HD2	68:ss:53:LEU:HD12	1.95	0.48
1:lA:1103:U:H2'	1:lA:1104:A:H8	1.78	0.48
13:lM:151:ASP:N	13:lM:151:ASP:OD1	2.34	0.48
18:lR:70:LYS:HD3	18:lR:70:LYS:HA	1.66	0.48
23:lW:49:PHE:O	23:lW:53:THR:OG1	2.22	0.48
55:se:141:CYS:SG	55:se:170:MET:HE3	2.53	0.48
1:lA:452:A:C2	2:lB:19:A:H1'	2.49	0.48
1:lA:500:G:HO2'	1:lA:501:U:H6	1.60	0.48
1:lA:2685:G:H2'	1:lA:2686:C:H6	1.77	0.48
7:lG:30:TYR:O	7:lG:34:ARG:HG3	2.13	0.48
11:lK:43:LYS:HE3	15:lO:132:PRO:HG2	1.95	0.48
14:lN:87:LEU:HD22	14:lN:220:LEU:HD11	1.96	0.48
33:lg:101:VAL:O	33:lg:126:ARG:NH1	2.46	0.48
43:lq:3:VAL:HA	43:lq:90:GLU:O	2.14	0.48
50:sJ:20:U:O2	50:sJ:20:U:H2'	2.13	0.48
52:sa:1757:C:H2'	52:sa:1758:G:H8	1.78	0.48
53:sc:82:VAL:HG13	53:sc:104:VAL:HG12	1.96	0.48
60:sj:37:ARG:HH22	60:sj:59:ARG:HH11	1.62	0.48
1:lA:219:G:H2'	1:lA:220:A:C8	2.49	0.48
1:lA:278:U:H2'	1:lA:279:U:H6	1.79	0.48
1:lA:495:A:N7	1:lA:577:G:O2'	2.46	0.48
1:lA:1664:G:C2	1:lA:1756:U:H4'	2.49	0.48
1:lA:2062:A:O2'	1:lA:3220:U:OP1	2.32	0.48
1:lA:2240:C:OP1	4:lD:8:GLN:NE2	2.47	0.48
1:lA:2884:U:H3'	1:lA:2885:G:H21	1.79	0.48
1:lA:3323:C:O2'	1:lA:3419:A:N1	2.43	0.48
12:lL:206:LEU:O	12:lL:208:LYS:NZ	2.47	0.48
21:lU:28:GLU:HG3	21:lU:49:LEU:HD22	1.94	0.48
35:li:70:LYS:HA	35:li:73:ARG:HE	1.79	0.48
52:sa:109:U:H2'	52:sa:110:A:C8	2.49	0.48
52:sa:1383:C:H5	52:sa:1389:G:H1	1.62	0.48
52:sa:1880:U:H2'	52:sa:1881:C:C6	2.48	0.48
1:lA:2256:A:N6	1:lA:2257:G:O6	2.47	0.48
1:lA:2462:G:H2'	1:lA:2463:A:C8	2.44	0.48
1:lA:2482:C:O2'	1:lA:2689:G:N2	2.47	0.48
1:lA:3404:A:H2'	8:lH:95:PHE:CZ	2.49	0.48
2:lB:28:U:H2'	2:lB:29:U:H6	1.79	0.48
2:lB:116:U:H2'	2:lB:117:A:H8	1.79	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:II:168:CYS:SG	9:II:171:ASP:HB2	2.54	0.48
47:sD:22:THR:OG1	47:sD:23:GLY:N	2.46	0.48
52:sa:61:A:O2'	52:sa:261:U:O2	2.25	0.48
52:sa:93:C:O2'	56:sf:5:HIS:HB2	2.14	0.48
52:sa:587:U:H4'	52:sa:589:G:H4'	1.96	0.48
52:sa:1557:C:H2'	52:sa:1558:U:H6	1.79	0.48
52:sa:1586:U:OP1	71:sv:57:ARG:NE	2.47	0.48
52:sa:1745:U:H2'	52:sa:1746:C:C6	2.48	0.48
55:se:139:ILE:HG22	55:se:217:VAL:HG22	1.95	0.48
56:sf:104:LYS:NZ	56:sf:249:GLN:OE1	2.47	0.48
63:sm:83:ILE:HG13	63:sm:108:VAL:HG23	1.95	0.48
1:lA:735:C:H2'	1:lA:736:G:O4'	2.13	0.48
1:lA:1689:A:C5	1:lA:1690:A:H1'	2.49	0.48
1:lA:1837:U:C5	1:lA:1984:A:H5''	2.49	0.48
1:lA:1849:A:H5'	34:lh:30:MET:HE1	1.96	0.48
1:lA:2236:A:H2'	1:lA:2237:A:H8	1.78	0.48
1:lA:2304:A:H2'	1:lA:2305:C:C6	2.49	0.48
1:lA:2628:A:N6	4:ID:98:ILE:O	2.47	0.48
1:lA:2737:A:OP1	13:IM:51:ARG:NH2	2.47	0.48
4:ID:113:VAL:HB	4:ID:164:ALA:HB1	1.96	0.48
10:IJ:73:LEU:HD22	10:IJ:175:CYS:HB2	1.94	0.48
46:sC:9:LEU:HD21	67:sr:51:THR:HG21	1.96	0.48
50:sJ:30:G:H2'	50:sJ:31:G:O4'	2.13	0.48
52:sa:258:A:P	52:sa:285:G:H22	2.36	0.48
56:sf:50:LEU:HB3	56:sf:52:TYR:CD1	2.49	0.48
58:sh:57:ASP:OD2	58:sh:98:ARG:HD2	2.14	0.48
68:ss:69:ILE:HG22	68:ss:70:VAL:HG13	1.95	0.48
1:lA:857:U:C2	1:lA:887:G:N2	2.82	0.47
1:lA:1019:G:H1'	1:lA:1737:A:N6	2.29	0.47
1:lA:1883:A:N6	1:lA:1931:U:H3	2.11	0.47
1:lA:2144:U:OP2	21:IU:74:ARG:NE	2.36	0.47
1:lA:2688:G:OP1	1:lA:2714:C:O2'	2.20	0.47
1:lA:3038:C:H5'	42:lp:27:ARG:HH21	1.78	0.47
1:lA:3265:C:O2'	11:IK:161:SER:OG	2.32	0.47
6:IF:233:ARG:NE	6:IF:235:GLU:OE1	2.32	0.47
6:IF:389:LYS:O	6:IF:391:LYS:N	2.47	0.47
13:IM:20:ASN:OD1	13:IM:126:ASP:HB2	2.14	0.47
16:IP:25:ALA:HB1	16:IP:38:VAL:HB	1.94	0.47
35:li:-9:ALA:HA	35:li:38:VAL:HG13	1.96	0.47
52:sa:251:C:HO2'	60:sj:64:ASN:HD21	1.56	0.47
52:sa:795:G:N2	67:sr:108:TYR:OH	2.47	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:sa:1438:A:N6	52:sa:1518:A:N7	2.61	0.47
52:sa:1439:C:H2'	52:sa:1440:A:H4'	1.96	0.47
52:sa:1717:U:O4	66:sq:39:ARG:NH2	2.46	0.47
52:sa:1872:G:H2'	52:sa:1873:A:H8	1.79	0.47
1:lA:2781:U:O2	1:lA:2814:U:H4'	2.14	0.47
2:lB:141:G:H5''	17:lQ:60:VAL:HG11	1.95	0.47
27:la:52:ASP:HA	27:la:67:LYS:HE3	1.96	0.47
40:ln:7:GLU:CD	40:ln:10:GLN:HE22	2.22	0.47
47:sD:40:ARG:NH2	47:sD:61:ALA:O	2.47	0.47
52:sa:4:C:OP1	53:sc:202:THR:OG1	2.29	0.47
52:sa:242:G:O2'	52:sa:243:U:OP1	2.32	0.47
52:sa:327:U:H5''	60:sj:31:ARG:NH1	2.29	0.47
55:se:76:MET:O	55:se:77:GLU:HG2	2.14	0.47
1:lA:255:A:H4'	1:lA:257:A:N7	2.28	0.47
1:lA:1033:A:C2	4:lD:204:MET:HE2	2.49	0.47
1:lA:1057:C:C5	29:lc:26:ARG:HD2	2.50	0.47
1:lA:1105:U:OP1	9:lI:113:LYS:NZ	2.48	0.47
1:lA:1639:U:H5''	21:lU:6:LEU:HD22	1.96	0.47
1:lA:2224:U:H2'	1:lA:2225:A:C8	2.49	0.47
2:lB:145:U:H2'	2:lB:146:U:H6	1.79	0.47
52:sa:328:C:C5	60:sj:49:ARG:HD2	2.50	0.47
52:sa:412:A:H5'	52:sa:413:G:C5	2.49	0.47
52:sa:1757:C:O3'	71:sv:93:ARG:NH2	2.47	0.47
52:sa:1828:A:H2'	52:sa:1829:A:C8	2.49	0.47
62:sl:45:MET:HB2	62:sl:48:LEU:HD22	1.96	0.47
1:lA:244:A:N7	1:lA:265:A:N6	2.62	0.47
1:lA:526:A:H5''	1:lA:538:A:N1	2.29	0.47
1:lA:676:A:N7	6:lF:358:MET:HG2	2.29	0.47
1:lA:802:A:H2'	1:lA:803:U:C6	2.49	0.47
1:lA:905:A:C8	19:lS:60:ILE:HD11	2.49	0.47
1:lA:926:A:H2	1:lA:2487:U:H1'	1.79	0.47
1:lA:2721:G:H5''	1:lA:2722:U:O4'	2.14	0.47
1:lA:3142:G:N2	1:lA:3143:A:H62	2.12	0.47
6:lF:19:LYS:HD3	6:lF:19:LYS:N	2.28	0.47
13:lM:109:HIS:CE1	13:lM:123:TYR:H	2.33	0.47
26:lZ:9:ALA:HB2	26:lZ:31:PHE:HB3	1.96	0.47
42:lp:35:CYS:O	42:lp:42:HIS:HA	2.14	0.47
52:sa:375:A:H2'	52:sa:375:A:N3	2.30	0.47
52:sa:972:A:C2	52:sa:973:A:C8	3.02	0.47
66:sq:32:PHE:HZ	66:sq:111:LEU:HD13	1.78	0.47
68:ss:97:ARG:NH1	68:ss:128:ASP:OD2	2.45	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
70:su:115:LEU:HA	70:su:118:MET:HE3	1.97	0.47
1:lA:851:A:H5'	29:lc:114:GLY:O	2.15	0.47
1:lA:2639:A:OP1	28:lb:55:ARG:HG3	2.14	0.47
1:lA:3452:U:H3	1:lA:3503:G:H1	1.61	0.47
1:lA:3455:G:N2	1:lA:3500:A:H2	2.09	0.47
3:lC:53:C:O2'	13:lM:152:HIS:ND1	2.48	0.47
43:lq:14:ARG:HB2	43:lq:77:TYR:CD2	2.50	0.47
52:sa:1319:G:H21	52:sa:1346:A:H8	1.61	0.47
52:sa:1693:A:O2'	52:sa:1694:A:OP1	2.29	0.47
70:su:24:ARG:O	70:su:56:ARG:NH1	2.34	0.47
1:lA:226:U:H2'	1:lA:227:A:H8	1.79	0.47
1:lA:249:U:C2	1:lA:250:G:C8	3.03	0.47
1:lA:343:A:H62	17:lQ:12:ARG:HH12	1.62	0.47
1:lA:907:U:H2'	1:lA:908:A:C8	2.49	0.47
1:lA:1033:A:O2'	4:lD:203:ALA:O	2.32	0.47
1:lA:1255:A:H2	1:lA:3007:A:N3	2.13	0.47
1:lA:1928:G:N2	1:lA:1929:G:O6	2.40	0.47
1:lA:2260:U:OP1	4:lD:209:HIS:NE2	2.40	0.47
1:lA:2632:A:OP2	4:lD:87:TYR:OH	2.21	0.47
14:lN:238:ARG:HE	14:lN:241:ALA:HA	1.80	0.47
20:lT:73:ILE:HD12	20:lT:90:TYR:HD2	1.79	0.47
24:lX:57:CYS:SG	24:lX:122:SER:OG	2.59	0.47
49:sI:14:A:N7	49:sI:22:G:C2	2.82	0.47
52:sa:325:U:H2'	52:sa:326:A:H8	1.79	0.47
52:sa:1567:A:OP2	70:su:145:ARG:NH2	2.48	0.47
56:sf:214:ASN:OD1	56:sf:214:ASN:N	2.48	0.47
63:sm:37:ILE:HG21	63:sm:41:PHE:HB2	1.96	0.47
66:sq:82:LYS:HE3	66:sq:88:MET:HE1	1.96	0.47
1:lA:322:U:H2'	1:lA:323:U:C6	2.48	0.47
1:lA:332:G:C6	43:lq:43:ARG:HD3	2.50	0.47
1:lA:547:A:H2'	1:lA:548:A:C8	2.49	0.47
1:lA:859:U:H2'	1:lA:860:G:O4'	2.13	0.47
1:lA:1111:G:N3	1:lA:2707:A:H2'	2.29	0.47
1:lA:1565:G:N7	29:lc:9:ARG:NH2	2.62	0.47
1:lA:2219:A:O2'	1:lA:2220:A:H2'	2.15	0.47
3:lC:2:G:O2'	3:lC:24:U:O2	2.32	0.47
5:lE:217:ILE:HD13	5:lE:284:ILE:HD11	1.96	0.47
7:lG:51:VAL:HG23	7:lG:145:CYS:HB3	1.97	0.47
10:lJ:85:LEU:HD22	10:lJ:129:LEU:HD21	1.97	0.47
22:lV:34:TYR:HB2	22:lV:72:ILE:HD12	1.97	0.47
27:la:2:LYS:HD2	27:la:7:VAL:HG23	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:lm:51:ALA:HB3	39:lm:54:ILE:HD12	1.96	0.47
43:lq:23:VAL:HG22	43:lq:70:ILE:HG12	1.96	0.47
49:sI:15:G:O2'	49:sI:59:G:N1	2.43	0.47
52:sa:345:A:H5''	52:sa:347:A:H5'	1.96	0.47
52:sa:1441:C:HO2'	52:sa:1442:A:H8	1.61	0.47
53:sc:56:ASP:OD1	53:sc:57:GLN:N	2.47	0.47
69:st:16:ILE:HG22	69:st:24:LEU:HD11	1.97	0.47
1:lA:65:U:O2'	1:lA:97:A:O2'	2.19	0.47
1:lA:543:A:H2'	1:lA:544:U:C6	2.49	0.47
1:lA:1225:A:H1'	9:ll:93:LEU:HD22	1.97	0.47
1:lA:1629:G:C2	1:lA:1630:A:C8	3.03	0.47
1:lA:1632:A:H5'	38:ll:12:HIS:O	2.14	0.47
1:lA:2236:A:H2'	1:lA:2237:A:C8	2.50	0.47
1:lA:3352:G:H4'	1:lA:3353:U:H5''	1.96	0.47
5:lE:250:LYS:HB3	5:lE:250:LYS:HE2	1.74	0.47
10:lJ:242:GLN:HA	10:lJ:245:ILE:HG22	1.96	0.47
32:lf:19:LYS:HE3	32:lf:23:GLU:HG2	1.97	0.47
52:sa:388:C:H2'	52:sa:389:U:H6	1.80	0.47
52:sa:1185:A:H2'	52:sa:1186:C:C6	2.49	0.47
52:sa:1317:G:H2'	52:sa:1318:U:C6	2.49	0.47
53:sc:53:LYS:HB3	53:sc:53:LYS:HE3	1.76	0.47
56:sf:44:LEU:HD23	56:sf:47:ARG:HH21	1.79	0.47
60:sj:85:ALA:HA	63:sm:7:ARG:HD3	1.97	0.47
66:sq:73:LYS:HA	66:sq:73:LYS:HD3	1.70	0.47
1:lA:1741:A:N3	1:lA:1763:C:O2'	2.46	0.47
1:lA:1846:A:H2'	1:lA:1847:A:C8	2.50	0.47
1:lA:2705:A:C3'	1:lA:2706:C:H5'	2.45	0.47
1:lA:3329:U:H4'	1:lA:3411:U:H4'	1.97	0.47
1:lA:3455:G:H1	1:lA:3500:A:H2	1.60	0.47
3:lC:48:U:OP1	7:lG:221:ARG:HB2	2.15	0.47
4:lD:36:GLU:OE2	4:lD:163:ARG:HD3	2.15	0.47
7:lG:37:ILE:HD12	7:lG:67:SER:HB2	1.97	0.47
7:lG:51:VAL:HG12	7:lG:53:VAL:HG23	1.97	0.47
10:lJ:147:LEU:HD12	10:lJ:173:PRO:HB2	1.97	0.47
15:lO:177:LYS:HD3	15:lO:177:LYS:HA	1.75	0.47
19:lS:156:LYS:HD2	19:lS:156:LYS:HA	1.60	0.47
24:lX:112:MET:HE1	24:lX:117:ILE:HG13	1.97	0.47
43:lq:19:THR:O	43:lq:21:HIS:HD2	1.98	0.47
52:sa:1126:G:H8	67:sr:76:SER:OG	1.98	0.47
52:sa:1751:U:H5''	68:ss:149:ALA:HB3	1.97	0.47
56:sf:40:LEU:HD12	56:sf:41:PRO:HD2	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:2261:G:O2'	1:1A:2390:U:OP2	2.32	0.47
3:1C:22:C:H2'	3:1C:23:A:C8	2.46	0.47
3:1C:94:U:H2'	3:1C:95:A:C8	2.50	0.47
6:1F:365:LYS:HA	6:1F:397:LEU:HD11	1.96	0.47
7:1G:219:PHE:HB3	7:1G:222:PHE:HB2	1.96	0.47
9:1I:143:LYS:HB2	9:1I:143:LYS:HE3	1.61	0.47
26:1Z:61:ASN:N	26:1Z:61:ASN:OD1	2.47	0.47
27:1a:109:LEU:HA	27:1a:113:ARG:HG3	1.97	0.47
28:1b:53:ILE:HD12	28:1b:62:GLN:HB3	1.97	0.47
28:1b:126:ARG:H	28:1b:126:ARG:HG2	1.50	0.47
48:sE:41:ARG:HG2	72:sw:69:PRO:HA	1.95	0.47
52:sa:96:C:H2'	52:sa:97:U:C6	2.50	0.47
52:sa:383:G:OP2	52:sa:418:G:O2'	2.33	0.47
52:sa:541:U:H2'	52:sa:542:G:C8	2.50	0.47
52:sa:628:G:N1	52:sa:945:U:OP2	2.46	0.47
67:sr:41:MET:HE2	67:sr:41:MET:HB2	1.77	0.47
1:1A:82:G:O2'	1:1A:94:G:O6	2.29	0.46
1:1A:187:G:H5''	1:1A:188:U:H3'	1.98	0.46
1:1A:210:U:OP1	14:1N:215:LYS:NZ	2.42	0.46
1:1A:310:A:N1	1:1A:2873:G:O2'	2.45	0.46
1:1A:877:U:H2'	1:1A:878:U:C6	2.50	0.46
1:1A:1816:A:H1'	1:1A:1892:C:O2'	2.15	0.46
1:1A:3039:A:H4'	42:lp:20:ILE:HG21	1.96	0.46
1:1A:3090:G:C2	5:1E:252:ALA:HB1	2.51	0.46
2:1B:63:A:OP1	35:li:41:LYS:NZ	2.48	0.46
52:sa:619:C:H2'	52:sa:620:U:C6	2.50	0.46
52:sa:939:G:C5	64:so:17:PRO:HG3	2.50	0.46
69:st:31:ASN:O	69:st:35:VAL:HG23	2.15	0.46
1:1A:200:A:H2'	1:1A:201:A:C8	2.49	0.46
1:1A:591:C:H2'	1:1A:592:A:C8	2.48	0.46
1:1A:1120:A:OP2	7:1G:10:ARG:NH1	2.37	0.46
1:1A:1483:U:O2'	9:1I:149:MET:SD	2.73	0.46
1:1A:3091:C:H5''	5:1E:245:HIS:HB3	1.96	0.46
1:1A:3370:A:H2'	1:1A:3371:U:C6	2.50	0.46
3:1C:104:G:C2	3:1C:105:A:C5	3.03	0.46
4:1D:62:VAL:HG22	4:1D:73:LYS:HG2	1.96	0.46
4:1D:177:LYS:NZ	39:lm:33:GLN:OE1	2.48	0.46
11:1K:24:LYS:HB2	11:1K:24:LYS:HE2	1.76	0.46
52:sa:604:G:HO2'	52:sa:607:G:HO2'	1.61	0.46
52:sa:905:G:C2	52:sa:968:A:H1'	2.50	0.46
52:sa:1674:G:H5'	52:sa:1717:U:H5'	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:sa:1756:U:H2'	52:sa:1757:C:H6	1.80	0.46
56:sf:123:LYS:HB3	56:sf:140:HIS:HB2	1.97	0.46
68:ss:51:LEU:HB3	68:ss:64:PHE:HE1	1.80	0.46
1:lA:579:A:C4	1:lA:580:A:C8	3.03	0.46
1:lA:723:U:O2'	1:lA:724:A:H5'	2.15	0.46
1:lA:779:A:O2'	1:lA:783:A:N6	2.49	0.46
1:lA:789:C:H2'	1:lA:790:A:C8	2.48	0.46
1:lA:1310:A:H2'	1:lA:1311:C:C6	2.50	0.46
1:lA:1930:C:H2'	1:lA:1931:U:H6	1.81	0.46
1:lA:2851:A:H2'	1:lA:2852:G:C8	2.50	0.46
1:lA:3204:C:O2'	26:lZ:18:ALA:O	2.33	0.46
5:lE:153:ILE:O	5:lE:157:CYS:HB2	2.16	0.46
8:lH:76:LYS:NZ	8:lH:190:TYR:O	2.49	0.46
9:lI:21:CYS:SG	9:lI:22:LYS:N	2.86	0.46
10:lJ:58:ARG:HD2	10:lJ:233:ILE:O	2.14	0.46
20:lT:19:GLU:OE2	22:IV:155:THR:HG21	2.16	0.46
43:lq:12:CYS:HB2	43:lq:21:HIS:CE1	2.50	0.46
44:sA:106:ILE:HD13	44:sA:108:ILE:HD12	1.97	0.46
49:sI:51:A:H2'	49:sI:52:G:O4'	2.14	0.46
59:si:154:LYS:HB2	59:si:154:LYS:HE3	1.65	0.46
72:sw:22:ILE:HB	72:sw:89:LEU:HB2	1.97	0.46
1:lA:560:U:H2'	1:lA:561:G:C8	2.50	0.46
1:lA:832:A:O2'	1:lA:833:U:OP1	2.29	0.46
1:lA:1078:C:O2	1:lA:2684:G:O2'	2.33	0.46
1:lA:1095:A:H2	1:lA:1099:A:C2	2.33	0.46
1:lA:2373:U:C2	1:lA:2375:A:C6	3.03	0.46
1:lA:3214:C:H1'	1:lA:3459:U:H1'	1.98	0.46
1:lA:3278:A:N1	11:lK:80:THR:OG1	2.49	0.46
49:sI:44:U:H2'	49:sI:45:G:O4'	2.16	0.46
52:sa:158:A:C8	52:sa:159:A:C8	3.04	0.46
52:sa:840:U:H2'	52:sa:841:U:H6	1.79	0.46
52:sa:1206:U:H4'	66:sq:126:THR:HG23	1.98	0.46
60:sj:98:LYS:HB2	60:sj:207:SER:O	2.15	0.46
71:sv:41:GLY:HA2	71:sv:97:SER:H	1.81	0.46
74:sy:83:VAL:HG21	74:sy:89:LEU:HD13	1.98	0.46
1:lA:564:A:H1'	1:lA:565:U:OP2	2.15	0.46
1:lA:1033:A:N7	4:lD:199:VAL:HG21	2.31	0.46
1:lA:1311:C:O3'	1:lA:1419:G:N2	2.48	0.46
1:lA:1763:C:H2'	1:lA:1764:A:C8	2.50	0.46
1:lA:2239:U:O2'	4:lD:11:GLY:HA3	2.16	0.46
1:lA:2867:A:H62	1:lA:2887:G:N2	2.13	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1B:57:U:N3	2:1B:58:G:N7	2.64	0.46
2:1B:85:U:H3'	2:1B:86:G:H5'	1.97	0.46
6:1F:242:LEU:HD22	6:1F:252:ILE:HD11	1.97	0.46
9:1I:96:LEU:HD21	9:1I:103:VAL:HG22	1.97	0.46
14:1N:148:LYS:HE3	14:1N:152:ILE:HD11	1.98	0.46
24:1X:88:TRP:HH2	24:1X:98:PHE:HE1	1.62	0.46
50:sJ:39:A:O2'	52:sa:981:A:OP1	2.25	0.46
52:sa:395:G:H1	60:sj:29:MET:HE1	1.80	0.46
52:sa:746:A:H2'	52:sa:747:A:H8	1.80	0.46
52:sa:982:G:O6	52:sa:983:A:N6	2.48	0.46
52:sa:1033:A:H61	52:sa:1091:C:H42	1.62	0.46
52:sa:1439:C:H5'	69:st:48:ASN:HB3	1.98	0.46
52:sa:1822:U:O4	52:sa:1891:A:H1'	2.16	0.46
1:1A:59:A:H8	1:1A:59:A:O5'	1.99	0.46
1:1A:1765:A:H2'	1:1A:1766:G:O4'	2.15	0.46
1:1A:2230:U:H2'	1:1A:2231:C:C6	2.50	0.46
1:1A:3410:A:H1'	1:1A:3411:U:H5	1.81	0.46
2:1B:102:A:C8	2:1B:103:A:C8	3.03	0.46
6:1F:150:ILE:HD12	6:1F:150:ILE:HA	1.78	0.46
9:1I:185:GLU:OE1	9:1I:185:GLU:N	2.46	0.46
52:sa:150:G:N1	52:sa:152:A:H5''	2.30	0.46
52:sa:597:U:H2'	52:sa:598:A:H8	1.81	0.46
52:sa:1224:G:OP1	52:sa:1225:G:H8	1.98	0.46
52:sa:1234:C:H42	52:sa:1562:G:H22	1.64	0.46
52:sa:1671:A:N3	52:sa:1729:G:O2'	2.34	0.46
52:sa:1776:G:H5''	57:sg:88:LYS:HB2	1.97	0.46
54:sd:119:ILE:HG22	54:sd:198:ILE:HD13	1.97	0.46
56:sf:196:ARG:HB3	56:sf:206:VAL:HG23	1.98	0.46
62:sl:1:MET:HE1	62:sl:39:LEU:HG	1.96	0.46
65:sp:48:ILE:HG22	65:sp:49:ILE:HG22	1.97	0.46
1:1A:527:G:H1	1:1A:545:U:H3	1.64	0.46
1:1A:718:A:H2'	1:1A:719:U:H6	1.78	0.46
1:1A:1675:A:OP2	1:1A:1734:G:N1	2.32	0.46
1:1A:3473:U:H2'	1:1A:3474:U:H6	1.80	0.46
5:1E:48:GLY:HA3	5:1E:81:THR:HG22	1.97	0.46
7:1G:189:GLU:HA	7:1G:192:ARG:HB3	1.97	0.46
8:1H:187:MET:HE3	8:1H:187:MET:HB3	1.76	0.46
8:1H:197:ASP:HB3	8:1H:202:LEU:HD21	1.97	0.46
17:1Q:17:ASP:N	17:1Q:17:ASP:OD1	2.48	0.46
45:sB:43:ASN:OD1	45:sB:43:ASN:N	2.49	0.46
48:sE:8:ASN:ND2	62:sl:28:VAL:HG12	2.31	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:sa:251:C:H2'	52:sa:252:C:H6	1.80	0.46
54:sd:38:LEU:HD13	62:sl:64:TYR:HE2	1.81	0.46
62:sl:39:LEU:O	62:sl:43:MET:HB2	2.16	0.46
1:lA:1581:G:N2	1:lA:2432:A:OP2	2.38	0.46
4:lD:204:MET:SD	4:lD:209:HIS:HB2	2.56	0.46
6:lF:135:ALA:HB3	27:la:175:ALA:HB1	1.98	0.46
6:lF:403:ARG:HE	6:lF:411:MET:HE1	1.81	0.46
8:lH:120:SER:O	8:lH:121:LYS:HG2	2.16	0.46
11:lK:71:ASN:OD1	11:lK:71:ASN:N	2.47	0.46
15:lO:92:ASN:HA	15:lO:97:LYS:HZ1	1.81	0.46
25:lY:55:ILE:HD11	25:lY:121:PHE:CE2	2.51	0.46
52:sa:1311:U:O2'	52:sa:1312:A:O4'	2.34	0.46
52:sa:1843:U:H2'	52:sa:1844:A:H8	1.80	0.46
53:sc:151:GLY:HA2	53:sc:152:GLU:HA	1.58	0.46
56:sf:87:VAL:HG13	56:sf:120:LYS:HB2	1.97	0.46
1:lA:2224:U:O2'	4:lD:182:ALA:HB2	2.15	0.46
1:lA:2395:C:HO2'	1:lA:2396:A:H8	1.63	0.46
13:lM:55:ARG:H	13:lM:55:ARG:HD3	1.81	0.46
39:lm:24:LYS:HA	39:lm:24:LYS:HD3	1.69	0.46
52:sa:1277:C:H2'	52:sa:1278:A:H8	1.80	0.46
52:sa:1678:A:H2'	52:sa:1679:G:O4'	2.16	0.46
1:lA:828:A:OP1	17:lQ:202:ARG:NH2	2.48	0.46
1:lA:2301:U:C2	1:lA:2302:U:C6	3.04	0.46
1:lA:3103:C:H2'	1:lA:3104:G:H8	1.80	0.46
1:lA:3104:G:H2'	1:lA:3105:U:C6	2.51	0.46
1:lA:3369:U:H2'	1:lA:3370:A:C8	2.51	0.46
1:lA:3435:G:OP1	5:lE:271:GLN:NE2	2.49	0.46
17:lQ:165:THR:O	17:lQ:169:LYS:HB2	2.16	0.46
19:lS:112:LYS:HE3	19:lS:112:LYS:HB2	1.61	0.46
33:lg:33:TRP:CZ2	33:lg:54:PRO:HD3	2.51	0.46
52:sa:338:C:H2'	52:sa:339:A:C8	2.51	0.46
52:sa:400:C:H5''	58:sh:93:ARG:HH21	1.81	0.46
52:sa:1590:A:H2	52:sa:1773:G:H1'	1.80	0.46
53:sc:172:VAL:HB	53:sc:199:PHE:HB2	1.98	0.46
70:su:27:VAL:HG23	70:su:55:LYS:O	2.16	0.46
1:lA:657:A:O2'	1:lA:658:A:OP2	2.26	0.45
1:lA:1485:A:H2'	1:lA:1486:C:O4'	2.16	0.45
1:lA:1520:A:H4'	1:lA:1521:A:O5'	2.15	0.45
1:lA:1533:U:O2'	1:lA:1534:U:OP1	2.29	0.45
1:lA:3474:U:H2'	1:lA:3475:U:O4'	2.16	0.45
10:lJ:47:ARG:O	10:lJ:51:GLU:HG2	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:1U:122:ASP:C	21:1U:122:ASP:OD1	2.59	0.45
22:1V:97:LYS:HE2	22:1V:97:LYS:HB3	1.78	0.45
32:1f:68:PRO:HD3	32:1f:102:ILE:HG13	1.98	0.45
52:sa:5:U:H2'	52:sa:6:G:C8	2.51	0.45
52:sa:96:C:H2'	52:sa:97:U:H6	1.80	0.45
52:sa:750:U:O4	52:sa:751:A:N6	2.49	0.45
52:sa:882:G:H1	65:sp:60:ASP:CG	2.24	0.45
67:sr:28:ARG:HB3	67:sr:29:PRO:HD3	1.98	0.45
1:1A:424:A:O2'	1:1A:425:C:OP2	2.26	0.45
1:1A:739:A:H2'	1:1A:740:A:H8	1.82	0.45
1:1A:1048:A:H2'	1:1A:1049:U:C6	2.51	0.45
1:1A:1305:A:H1'	1:1A:1306:A:C8	2.50	0.45
1:1A:2346:A:H2'	1:1A:2347:A:C8	2.51	0.45
1:1A:2972:U:C4	1:1A:2973:G:N7	2.84	0.45
6:1F:114:LYS:HG3	17:1Q:203:TYR:HB3	1.98	0.45
15:1O:33:LYS:HA	15:1O:102:ASN:HB3	1.98	0.45
15:1O:130:PHE:CE1	15:1O:136:ARG:HD2	2.52	0.45
52:sa:153:U:H1'	52:sa:415:A:O2'	2.17	0.45
52:sa:412:A:H4'	52:sa:413:G:O5'	2.15	0.45
52:sa:1591:G:H2'	52:sa:1592:A:C8	2.51	0.45
56:sf:85:MET:HB3	56:sf:100:MET:SD	2.57	0.45
59:si:137:TYR:CD1	59:si:138:PRO:HA	2.51	0.45
60:sj:22:GLN:HG3	60:sj:25:LYS:HE3	1.98	0.45
1:1A:353:G:O2'	1:1A:2299:A:H1'	2.17	0.45
1:1A:709:U:O2'	1:1A:710:A:OP1	2.29	0.45
11:1K:140:VAL:HG22	11:1K:160:VAL:HG22	1.96	0.45
52:sa:209:A:N6	52:sa:250:G:O6	2.50	0.45
52:sa:1272:U:H2'	52:sa:1273:C:C6	2.52	0.45
52:sa:1895:G:H2'	52:sa:1896:A:C8	2.51	0.45
56:sf:255:LYS:HE2	56:sf:255:LYS:HB2	1.70	0.45
68:ss:60:ARG:HH11	68:ss:60:ARG:HG3	1.81	0.45
73:sx:12:PRO:HB2	73:sx:21:LEU:HD21	1.99	0.45
1:1A:720:U:C2	1:1A:721:A:N7	2.84	0.45
1:1A:3469:A:H5''	1:1A:3470:G:N7	2.30	0.45
2:1B:24:U:H5	6:1F:197:MET:HE1	1.81	0.45
4:1D:230:PRO:HD2	4:1D:233:GLN:HG2	1.98	0.45
7:1G:278:LYS:HB2	7:1G:278:LYS:HE2	1.73	0.45
32:1f:121:GLU:CD	32:1f:123:GLY:H	2.23	0.45
52:sa:208:C:H2'	52:sa:209:A:C8	2.51	0.45
53:sc:185:THR:HG21	53:sc:209:LEU:HD12	1.99	0.45
56:sf:50:LEU:HB3	56:sf:52:TYR:HD1	1.81	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
62:sl:1:MET:HE2	62:sl:40:GLN:HA	1.97	0.45
71:sv:112:GLY:O	71:sv:124:THR:OG1	2.33	0.45
1:lA:285:A:H4'	1:lA:286:U:C5'	2.47	0.45
1:lA:2449:A:N3	1:lA:2967:G:O2'	2.45	0.45
1:lA:3370:A:C2	16:lP:128:ILE:HD12	2.52	0.45
1:lA:3386:G:H2'	1:lA:3387:G:H8	1.81	0.45
1:lA:3461:U:H4'	1:lA:3462:A:H5'	1.99	0.45
18:lR:57:ILE:HB	18:lR:72:GLN:HB2	1.97	0.45
32:lf:11:VAL:O	32:lf:15:ARG:HG3	2.16	0.45
34:lh:7:TYR:HE1	34:lh:20:ILE:HG12	1.81	0.45
52:sa:323:A:H2'	52:sa:324:G:C8	2.52	0.45
52:sa:1442:A:H2'	52:sa:1443:U:C6	2.51	0.45
52:sa:1565:G:H2'	52:sa:1565:G:N3	2.32	0.45
52:sa:1808:G:O2'	52:sa:1928:A:O3'	2.34	0.45
1:lA:265:A:H2'	1:lA:265:A:N3	2.32	0.45
1:lA:612:A:O4'	1:lA:716:A:N6	2.50	0.45
1:lA:846:A:H2'	1:lA:847:A:C8	2.52	0.45
1:lA:979:G:C5	4:lD:181:LYS:HB3	2.52	0.45
1:lA:1992:A:H2'	1:lA:1993:G:H8	1.82	0.45
1:lA:2744:A:C6	13:lM:124:GLY:HA3	2.51	0.45
1:lA:2863:A:C2	37:lk:16:HIS:HB3	2.52	0.45
6:lF:102:PHE:O	6:lF:103:SER:C	2.59	0.45
7:lG:232:LEU:HA	7:lG:235:ILE:HG22	1.99	0.45
14:lN:41:ALA:HB2	14:lN:48:THR:HG21	1.97	0.45
15:lO:175:LYS:HD3	15:lO:175:LYS:HA	1.72	0.45
27:la:83:LEU:HD23	27:la:97:ILE:HD11	1.98	0.45
52:sa:841:U:H2'	52:sa:842:A:C5	2.52	0.45
54:sd:115:GLN:O	54:sd:118:SER:OG	2.29	0.45
56:sf:157:THR:HB	56:sf:171:VAL:HG13	1.98	0.45
58:sh:62:PRO:HD2	58:sh:97:VAL:HG12	1.99	0.45
63:sm:52:TYR:CD2	63:sm:112:PRO:HG2	2.51	0.45
64:so:102:LEU:HA	64:so:102:LEU:HD23	1.81	0.45
66:sq:107:LYS:HG3	66:sq:109:GLU:N	2.31	0.45
1:lA:504:U:H2'	1:lA:505:A:H8	1.82	0.45
1:lA:1444:U:C4	1:lA:1445:U:O4	2.70	0.45
1:lA:2089:A:C2	1:lA:2467:G:H4'	2.52	0.45
1:lA:3148:C:H2'	1:lA:3149:A:C8	2.52	0.45
2:lB:15:U:H5'	18:lR:2:VAL:HG13	1.99	0.45
3:lC:69:U:H2'	3:lC:70:U:H6	1.82	0.45
7:lG:9:ASN:OD1	7:lG:9:ASN:N	2.50	0.45
14:lN:44:VAL:HG23	14:lN:47:ALA:HB2	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:lO:63:VAL:HG12	15:lO:66:ASN:H	1.82	0.45
52:sa:1110:G:N2	52:sa:1113:A:OP2	2.43	0.45
56:sf:57:ARG:O	56:sf:60:THR:OG1	2.34	0.45
58:sh:74:ARG:HG2	58:sh:96:THR:HB	1.98	0.45
64:so:3:ARG:HB2	64:so:6:ASN:O	2.17	0.45
71:sv:50:ASN:ND2	71:sv:52:ASP:OD1	2.49	0.45
1:lA:10:U:H2'	1:lA:11:A:H8	1.81	0.45
1:lA:11:A:H2'	1:lA:12:U:C6	2.52	0.45
1:lA:1314:A:C2	15:lO:134:ARG:HD3	2.52	0.45
1:lA:1430:G:O2'	1:lA:1431:G:H5''	2.16	0.45
1:lA:1904:U:OP2	21:lU:124:TYR:OH	2.29	0.45
1:lA:2861:U:O4'	1:lA:2892:A:N6	2.49	0.45
15:lO:62:ARG:HD2	15:lO:67:PRO:HB3	1.98	0.45
18:lR:52:ILE:HA	18:lR:85:VAL:HG22	1.99	0.45
22:lV:80:VAL:CG1	22:lV:81:ASN:H	2.25	0.45
28:lb:92:PHE:HA	28:lb:120:GLN:NE2	2.32	0.45
33:lg:36:PRO:HG2	33:lg:44:ARG:HB3	1.99	0.45
52:sa:429:G:N1	52:sa:432:A:OP2	2.49	0.45
52:sa:734:A:C2	52:sa:803:G:N1	2.78	0.45
52:sa:1595:G:H3'	52:sa:1682:A:H61	1.82	0.45
54:sd:205:ASP:O	54:sd:208:PRO:HD2	2.17	0.45
60:sj:106:PRO:HB2	60:sj:110:ARG:HG3	1.99	0.45
65:sp:34:ASP:OD1	65:sp:35:THR:N	2.48	0.45
66:sq:56:VAL:O	66:sq:60:LEU:HD13	2.17	0.45
1:lA:861:A:N6	1:lA:881:G:H1'	2.31	0.45
1:lA:1347:A:N7	1:lA:1410:A:C6	2.85	0.45
1:lA:1844:U:H2'	1:lA:1845:G:H8	1.82	0.45
1:lA:2112:C:H2'	1:lA:2113:A:H8	1.81	0.45
1:lA:2180:C:H2'	1:lA:2181:A:H8	1.82	0.45
1:lA:3161:U:H2'	1:lA:3162:A:C8	2.52	0.45
3:lC:48:U:H3'	7:lG:221:ARG:HG3	1.98	0.45
5:lE:242:ARG:HA	5:lE:248:LEU:HD21	1.99	0.45
17:lQ:38:ARG:HB2	17:lQ:62:TYR:CZ	2.52	0.45
27:la:62:GLN:HB3	27:la:64:VAL:HG13	1.98	0.45
28:lb:57:MET:HE2	28:lb:57:MET:HB3	1.85	0.45
43:lq:51:GLN:NE2	43:lq:53:LYS:O	2.30	0.45
52:sa:341:G:H5'	63:sm:78:HIS:HB2	1.98	0.45
52:sa:1196:A:H2'	52:sa:1197:G:H8	1.82	0.45
52:sa:1209:A:N1	52:sa:1562:G:H5'	2.32	0.45
52:sa:1363:C:H1'	52:sa:1517:A:C5	2.52	0.45
52:sa:1709:A:H5'	71:sv:88:TYR:CE2	2.52	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:se:123:ILE:HD13	55:se:166:ILE:HG21	1.99	0.45
56:sf:28:LYS:NZ	56:sf:36:MET:SD	2.89	0.45
70:su:27:VAL:O	70:su:31:LEU:HG	2.17	0.45
1:lA:760:A:H2'	1:lA:761:A:C8	2.51	0.45
1:lA:1615:A:O2'	32:lf:95:GLN:OE1	2.30	0.45
1:lA:2230:U:H5''	4:ID:242:ARG:O	2.16	0.45
1:lA:2485:G:O2'	1:lA:2487:U:H5''	2.17	0.45
3:lC:109:G:C6	3:lC:110:A:N1	2.85	0.45
12:lL:62:SER:HA	12:lL:65:MET:SD	2.56	0.45
19:lS:18:ALA:HB2	19:lS:29:VAL:HG21	1.99	0.45
24:lX:88:TRP:HH2	24:lX:98:PHE:CE1	2.35	0.45
52:sa:734:A:N1	52:sa:803:G:C6	2.85	0.45
52:sa:959:A:H4'	52:sa:1933:G:N2	2.31	0.45
53:sc:180:ILE:HG23	53:sc:198:VAL:HG12	1.99	0.45
59:si:129:GLU:OE2	64:so:20:ARG:NH1	2.50	0.45
74:sy:39:LEU:HD22	74:sy:45:ALA:HB2	1.98	0.45
1:lA:353:G:H2'	1:lA:354:U:C6	2.52	0.44
1:lA:914:G:H2'	1:lA:915:U:C6	2.52	0.44
1:lA:2150:G:C2	1:lA:2176:G:C2	3.05	0.44
9:lI:75:ILE:HD12	9:lI:222:ILE:HD11	1.99	0.44
25:lY:46:LEU:HG	25:lY:48:THR:HG23	1.98	0.44
40:ln:25:VAL:HG13	40:ln:68:GLU:HA	1.99	0.44
45:sB:19:GLN:NE2	52:sa:924:A:N7	2.64	0.44
49:sI:14:A:N6	49:sI:46:G:H1	2.15	0.44
52:sa:1689:A:H2	52:sa:1756:U:H1'	1.82	0.44
52:sa:1690:A:O2'	52:sa:1691:A:H8	2.00	0.44
60:sj:107:ALA:HB3	60:sj:108:PRO:HD3	1.97	0.44
67:sr:102:LEU:HD23	67:sr:128:PHE:HB3	1.98	0.44
1:lA:43:C:H5''	14:lN:14:LYS:HG2	1.98	0.44
1:lA:226:U:H2'	1:lA:227:A:C8	2.52	0.44
1:lA:857:U:H2'	1:lA:858:A:H8	1.83	0.44
1:lA:1593:A:OP1	32:lf:75:ARG:NH2	2.51	0.44
1:lA:1627:G:H21	34:lh:6:THR:HG22	1.82	0.44
1:lA:1736:A:C6	41:lo:4:HIS:HB3	2.52	0.44
1:lA:1745:A:H2'	1:lA:1746:G:C8	2.52	0.44
1:lA:1749:U:P	21:lU:42:ARG:HH22	2.40	0.44
1:lA:2445:G:H2'	1:lA:2446:G:C8	2.53	0.44
17:lQ:60:VAL:HG23	17:lQ:134:LEU:HB2	1.99	0.44
17:lQ:115:VAL:HG12	17:lQ:165:THR:HG21	1.97	0.44
25:lY:41:ILE:HG22	25:lY:42:ILE:HG13	1.99	0.44
28:lb:91:GLN:NE2	28:lb:91:GLN:O	2.50	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:ll:2:THR:O	38:ll:2:THR:OG1	2.31	0.44
52:sa:137:A:H2'	52:sa:138:G:H8	1.81	0.44
52:sa:620:U:H2'	52:sa:621:C:H6	1.82	0.44
54:sd:140:ILE:HG21	54:sd:202:ILE:HG12	1.99	0.44
55:se:111:LYS:HD3	55:se:111:LYS:HA	1.72	0.44
56:sf:127:ARG:NH1	56:sf:135:PRO:HB2	2.32	0.44
66:sq:85:LEU:H	66:sq:88:MET:HE3	1.82	0.44
74:sy:101:SER:O	74:sy:101:SER:OG	2.32	0.44
1:lA:592:A:H1'	1:lA:756:A:H2	1.82	0.44
1:lA:960:G:O2'	21:lU:131:GLN:OE1	2.34	0.44
1:lA:1544:U:O2'	33:lg:96:GLU:OE1	2.36	0.44
1:lA:2217:A:N1	1:lA:3099:A:H2'	2.33	0.44
3:lC:29:G:O2'	7:lG:217:LYS:NZ	2.36	0.44
6:lF:397:LEU:HD12	6:lF:397:LEU:HA	1.70	0.44
7:lG:41:LYS:HA	7:lG:41:LYS:HD2	1.67	0.44
17:lQ:10:LEU:HB2	37:lk:38:ILE:HD12	1.98	0.44
37:lk:68:LYS:HD3	37:lk:74:HIS:HA	2.00	0.44
49:sI:24:G:H2'	49:sI:25:C:C6	2.52	0.44
50:sJ:42:C:H2'	50:sJ:43:G:H8	1.81	0.44
52:sa:1431:U:H4'	72:sw:57:PRO:HG3	1.99	0.44
52:sa:1901:G:O2'	74:sy:56:GLU:OE2	2.31	0.44
70:su:27:VAL:HG11	70:su:51:ILE:HD13	1.98	0.44
1:lA:324:A:C6	1:lA:336:A:N1	2.85	0.44
1:lA:746:A:H5'	8:lH:41:LYS:HD3	1.99	0.44
1:lA:1089:C:O5'	1:lA:1089:C:H6	2.00	0.44
1:lA:2767:A:H2'	1:lA:2768:G:C8	2.52	0.44
1:lA:3149:A:H2'	1:lA:3150:A:C8	2.53	0.44
1:lA:3360:U:C4	16:lP:117:ILE:HD11	2.52	0.44
1:lA:3375:U:H2'	1:lA:3376:C:C6	2.52	0.44
12:lL:202:THR:O	12:lL:202:THR:OG1	2.34	0.44
27:la:10:ASN:HB3	27:la:13:LYS:HB2	1.99	0.44
48:sE:21:CYS:SG	48:sE:39:CYS:N	2.77	0.44
50:sJ:65:G:H2'	50:sJ:65:G:N3	2.33	0.44
52:sa:213:A:N6	52:sa:243:U:H4'	2.32	0.44
52:sa:373:A:H2'	52:sa:374:C:O4'	2.17	0.44
52:sa:380:A:H2'	52:sa:381:G:C8	2.53	0.44
52:sa:1021:G:H2'	52:sa:1022:G:C8	2.52	0.44
66:sq:19:LYS:HD3	66:sq:23:GLU:OE1	2.17	0.44
1:lA:424:A:OP1	1:lA:437:A:H2	2.00	0.44
1:lA:429:G:N2	1:lA:432:A:OP2	2.42	0.44
1:lA:857:U:H2'	1:lA:858:A:C8	2.53	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:1407:U:H2'	1:1A:1408:C:O4'	2.18	0.44
1:1A:1414:A:H2'	1:1A:1415:A:C8	2.48	0.44
1:1A:2358:U:OP1	1:1A:3116:G:O2'	2.27	0.44
1:1A:3103:C:H2'	1:1A:3104:G:C8	2.53	0.44
2:1B:90:A:H5'	27:1a:22:ASN:HD22	1.83	0.44
5:1E:80:GLU:OE1	5:1E:315:TYR:OH	2.35	0.44
19:1S:22:ASN:HB3	19:1S:25:MET:HB2	1.98	0.44
23:1W:68:VAL:C	23:1W:69:LYS:HG2	2.43	0.44
42:1p:21:CYS:HB3	42:1p:24:CYS:HB2	2.00	0.44
52:sa:959:A:N3	52:sa:1922:U:O2'	2.49	0.44
52:sa:1261:A:N6	52:sa:1262:G:O6	2.50	0.44
52:sa:1694:A:H5'	68:ss:57:PRO:HG2	1.97	0.44
52:sa:1756:U:H2'	52:sa:1757:C:C6	2.53	0.44
1:1A:199:G:N1	1:1A:311:A:OP2	2.49	0.44
1:1A:324:A:H2'	1:1A:325:U:C6	2.53	0.44
1:1A:597:A:H4'	8:1H:53:GLU:HG2	1.99	0.44
1:1A:709:U:HO2'	1:1A:710:A:P	2.40	0.44
1:1A:720:U:HO2'	1:1A:721:A:P	2.41	0.44
1:1A:2252:U:OP1	4:1D:54:ARG:NH2	2.44	0.44
1:1A:3147:U:N3	5:1E:144:SER:OG	2.49	0.44
1:1A:3369:U:H2'	1:1A:3370:A:H8	1.83	0.44
4:1D:143:ASN:OD1	4:1D:143:ASN:N	2.50	0.44
6:1F:207:GLY:HA3	6:1F:228:GLY:O	2.17	0.44
8:1H:93:GLY:O	8:1H:94:PRO:C	2.61	0.44
9:1I:87:ILE:HG23	9:1I:118:VAL:HG12	1.99	0.44
11:1K:65:ILE:HD13	11:1K:82:ALA:HB2	2.00	0.44
52:sa:170:G:C5	52:sa:171:A:H2	2.36	0.44
52:sa:323:A:H2'	52:sa:324:G:H8	1.82	0.44
52:sa:379:G:H2'	52:sa:380:A:C8	2.52	0.44
52:sa:592:U:H2'	52:sa:593:A:H8	1.81	0.44
52:sa:1566:C:OP1	70:su:125:ARG:NH2	2.41	0.44
52:sa:1771:G:OP1	72:sw:79:HIS:NE2	2.46	0.44
1:1A:258:G:C8	6:1F:227:PRO:HG3	2.52	0.44
1:1A:737:A:O5'	1:1A:738:G:N2	2.50	0.44
1:1A:913:U:C2	1:1A:914:G:C8	3.05	0.44
1:1A:1311:C:O2	20:1T:111:ARG:NH2	2.50	0.44
1:1A:2418:U:OP1	5:1E:233:LYS:NZ	2.47	0.44
36:1j:42:PHE:HE1	36:1j:108:ILE:HD11	1.82	0.44
38:1l:44:MET:HE3	38:1l:44:MET:HB3	1.71	0.44
40:1n:10:GLN:H	40:1n:10:GLN:CD	2.26	0.44
52:sa:587:U:H5''	61:sk:40:LYS:HE2	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:sa:799:C:O2	67:sr:124:ARG:HD2	2.17	0.44
52:sa:1250:G:H22	52:sa:1283:U:H3	1.66	0.44
54:sd:207:ASP:HB3	54:sd:208:PRO:HD3	1.98	0.44
1:lA:637:U:H1'	1:lA:638:A:C8	2.52	0.44
1:lA:996:C:HO2'	1:lA:999:G:HO2'	1.62	0.44
1:lA:1207:C:H2'	1:lA:1208:U:O4'	2.18	0.44
1:lA:1426:A:N7	1:lA:3000:C:O2'	2.50	0.44
1:lA:1733:U:C4	1:lA:1734:G:N7	2.86	0.44
1:lA:1877:A:H2'	1:lA:1878:A:H8	1.82	0.44
1:lA:2865:A:H2'	1:lA:2866:C:C2	2.52	0.44
1:lA:2868:U:H3	1:lA:2886:U:H3	1.66	0.44
1:lA:3134:U:O2'	1:lA:3438:G:N7	2.51	0.44
1:lA:3476:C:H5''	1:lA:3477:U:N1	2.33	0.44
4:lD:196:TRP:HB3	4:lD:197:PRO:HD3	2.00	0.44
6:lF:162:ILE:HD11	6:lF:171:LEU:HD22	2.00	0.44
17:lQ:65:ARG:HB2	17:lQ:129:PHE:HD1	1.82	0.44
19:lS:66:LEU:HD23	19:lS:136:ILE:HD12	2.00	0.44
21:lU:119:MET:HE3	21:lU:119:MET:HB2	1.78	0.44
22:lV:148:LYS:HD3	22:lV:148:LYS:HA	1.60	0.44
25:lY:112:MET:HE1	25:lY:121:PHE:CD2	2.52	0.44
33:lg:20:ARG:HD3	33:lg:29:VAL:HG12	1.98	0.44
36:lj:94:ALA:HA	36:lj:97:GLN:HG3	2.00	0.44
52:sa:150:G:C2	52:sa:152:A:H5''	2.53	0.44
52:sa:1579:A:H4'	52:sa:1580:U:H5'	1.99	0.44
52:sa:1752:A:H2'	52:sa:1753:A:O4'	2.18	0.44
56:sf:200:PRO:HB2	63:sm:39:MET:HG2	2.00	0.44
60:sj:6:ASP:HB3	60:sj:28:THR:HG21	1.99	0.44
1:lA:474:A:H2'	1:lA:475:A:H8	1.83	0.44
1:lA:496:A:H2'	1:lA:497:U:C6	2.53	0.44
1:lA:619:A:P	1:lA:619:A:H3'	2.58	0.44
1:lA:686:A:H5'	9:lI:134:LYS:HD3	2.00	0.44
1:lA:1293:U:OP1	9:lI:199:LYS:HG2	2.18	0.44
1:lA:2195:U:H2'	1:lA:2196:A:O4'	2.18	0.44
3:lC:7:A:OP2	7:lG:30:TYR:OH	2.28	0.44
4:lD:57:PRO:HG2	4:lD:78:CYS:HB3	1.99	0.44
12:lL:191:ILE:O	12:lL:198:LYS:N	2.51	0.44
16:lP:29:GLU:HG3	16:lP:30:ILE:N	2.32	0.44
17:lQ:8:ASN:HB2	17:lQ:46:GLU:HG3	2.00	0.44
21:lU:76:THR:O	21:lU:76:THR:OG1	2.31	0.44
25:lY:94:ARG:HH21	25:lY:98:LEU:HD12	1.83	0.44
39:lm:36:LYS:HE2	39:lm:48:ARG:HH11	1.82	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:sk:4:A:OP2	52:sa:1176:G:N1	2.36	0.44
52:sa:1107:A:N6	52:sa:1108:G:O6	2.51	0.44
52:sa:1383:C:H2'	52:sa:1384:U:C6	2.52	0.44
52:sa:1733:U:O4	52:sa:1734:G:N1	2.51	0.44
61:sk:57:LYS:HE3	61:sk:57:LYS:HB2	1.72	0.44
71:sv:37:THR:O	71:sv:37:THR:OG1	2.34	0.44
1:lA:247:A:H5'	1:lA:266:A:O2'	2.18	0.43
1:lA:661:G:H2'	1:lA:662:U:C6	2.53	0.43
1:lA:1222:U:O2'	22:IV:136:ARG:O	2.25	0.43
1:lA:1269:U:C4	1:lA:1488:A:H2	2.36	0.43
1:lA:1471:A:H2'	1:lA:1472:U:C6	2.53	0.43
1:lA:1519:A:H5''	1:lA:1521:A:N6	2.33	0.43
1:lA:2214:A:O2'	38:II:2:THR:N	2.44	0.43
1:lA:2381:G:N2	1:lA:2381:G:OP2	2.51	0.43
1:lA:2883:A:H2'	1:lA:2884:U:C6	2.53	0.43
1:lA:3391:U:H2'	1:lA:3392:G:C8	2.53	0.43
9:II:166:VAL:HG13	9:II:171:ASP:HB3	1.99	0.43
17:IQ:46:GLU:OE1	17:IQ:47:ARG:N	2.51	0.43
35:II:1:MET:HB3	35:II:5:ASP:HB3	2.00	0.43
52:sa:136:A:H4'	52:sa:137:A:H5'	1.99	0.43
52:sa:256:A:N6	52:sa:257:C:H41	2.16	0.43
56:sf:9:ARG:HD2	56:sf:9:ARG:HA	1.87	0.43
1:lA:213:A:N6	14:IN:131:THR:OG1	2.40	0.43
1:lA:1461:G:H2'	1:lA:1462:A:H8	1.83	0.43
1:lA:2034:G:O2'	41:IO:3:GLY:O	2.36	0.43
1:lA:2431:G:H4'	18:IR:139:PHE:CE1	2.53	0.43
2:IB:61:A:C4	2:IB:63:A:N6	2.86	0.43
5:IE:322:PHE:HE2	5:IE:324:MET:HE3	1.83	0.43
6:IF:295:VAL:HG22	19:IS:34:PHE:CE1	2.53	0.43
14:IN:174:LYS:HE2	14:IN:174:LYS:HB2	1.65	0.43
24:IX:140:VAL:HG11	26:IZ:24:VAL:HG21	2.00	0.43
52:sa:57:U:H3	52:sa:88:G:H1	1.66	0.43
52:sa:379:G:H2'	52:sa:380:A:H8	1.83	0.43
52:sa:1510:A:H2'	52:sa:1511:A:H8	1.83	0.43
56:sf:235:ASN:OD1	56:sf:235:ASN:N	2.44	0.43
71:sv:89:ASN:N	71:sv:89:ASN:OD1	2.50	0.43
1:lA:330:A:P	43:IQ:43:ARG:HH22	2.40	0.43
1:lA:504:U:H2'	1:lA:505:A:C8	2.53	0.43
1:lA:965:A:H2'	1:lA:966:A:O4'	2.17	0.43
1:lA:1022:U:OP2	38:II:30:GLN:NE2	2.51	0.43
1:lA:1622:A:H2'	1:lA:1622:A:N3	2.33	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:2220:A:H1'	1:1A:2357:A:H61	1.82	0.43
1:1A:2678:G:OP1	4:1D:2:GLY:HA3	2.19	0.43
1:1A:3442:G:OP1	5:1E:177:LYS:HD3	2.17	0.43
5:1E:217:ILE:HD12	5:1E:339:LEU:HD22	2.01	0.43
6:1F:292:LYS:HB2	6:1F:292:LYS:HE3	1.85	0.43
7:1G:89:LYS:HB2	7:1G:89:LYS:HE2	1.71	0.43
14:1N:49:ILE:HA	14:1N:231:LEU:O	2.18	0.43
15:1O:139:LEU:HD23	15:1O:139:LEU:HA	1.78	0.43
17:1Q:117:ASN:OD1	17:1Q:166:SER:HB3	2.18	0.43
18:1R:18:ARG:NH2	18:1R:147:GLU:HG2	2.33	0.43
24:1X:123:LYS:HG3	24:1X:139:ILE:HG22	1.98	0.43
45:sB:4:LYS:HD3	45:sB:5:ARG:NH2	2.33	0.43
52:sa:1302:G:H5''	54:sd:153:SER:HB2	2.01	0.43
52:sa:1305:C:O2'	72:sw:68:THR:HG22	2.18	0.43
55:se:114:SER:O	55:se:114:SER:OG	2.34	0.43
66:sq:53:GLU:O	66:sq:57:LYS:HG2	2.19	0.43
1:1A:213:A:C5'	14:1N:133:LYS:HB2	2.48	0.43
1:1A:619:A:N6	16:1P:77:CYS:HB3	2.34	0.43
1:1A:928:G:N1	1:1A:1051:U:O4	2.51	0.43
1:1A:1236:G:H2'	1:1A:1237:A:C8	2.52	0.43
1:1A:1275:A:H2	1:1A:2455:U:O2	2.01	0.43
1:1A:1289:G:C6	1:1A:1461:G:N1	2.87	0.43
1:1A:1502:A:N6	19:1S:15:SER:OG	2.40	0.43
1:1A:3287:U:H2'	1:1A:3288:U:C6	2.54	0.43
5:1E:20:LYS:HE3	5:1E:20:LYS:HB3	1.85	0.43
6:1F:35:ASP:OD1	6:1F:36:VAL:N	2.51	0.43
32:1f:118:ILE:HG22	32:1f:121:GLU:O	2.17	0.43
52:sa:1508:A:OP1	69:st:60:ARG:NH1	2.52	0.43
52:sa:1545:G:H2'	52:sa:1546:C:C6	2.53	0.43
54:sd:172:ILE:HD11	54:sd:177:ALA:HB3	2.00	0.43
71:sv:86:LYS:O	71:sv:87:HIS:ND1	2.51	0.43
71:sv:129:LYS:HZ2	71:sv:129:LYS:HG2	1.63	0.43
1:1A:215:A:H2'	1:1A:216:A:O4'	2.18	0.43
1:1A:227:A:H2'	1:1A:228:A:C8	2.53	0.43
1:1A:322:U:H2'	1:1A:323:U:H6	1.82	0.43
1:1A:343:A:H62	17:1Q:12:ARG:NH1	2.17	0.43
1:1A:467:A:C2	1:1A:2439:A:H4'	2.53	0.43
1:1A:551:A:H2'	1:1A:552:G:O4'	2.18	0.43
1:1A:1093:G:OP1	19:1S:19:ARG:NH2	2.45	0.43
1:1A:2469:G:OP1	1:1A:2469:G:O4'	2.37	0.43
1:1A:3203:A:H4'	5:1E:365:LYS:HD2	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:lA:3432:A:HO2'	1:lA:3433:U:P	2.41	0.43
1:lA:3482:C:H2'	1:lA:3483:A:O4'	2.18	0.43
2:lB:116:U:H2'	2:lB:117:A:C8	2.53	0.43
3:lC:2:G:H2'	3:lC:3:G:H8	1.84	0.43
6:lF:431:PHE:CZ	20:lT:24:ARG:HG3	2.53	0.43
8:lH:61:PRO:O	8:lH:116:LYS:NZ	2.52	0.43
8:lH:176:GLU:OE1	8:lH:176:GLU:N	2.39	0.43
12:lL:30:ARG:NH2	12:lL:66:GLU:HG2	2.33	0.43
14:lN:124:LEU:HD21	35:li:109:PHE:HE1	1.82	0.43
42:lp:20:ILE:HD11	42:lp:49:LYS:HD2	1.99	0.43
47:sD:10:VAL:HG11	57:sg:124:ARG:HH12	1.84	0.43
52:sa:456:G:H2'	52:sa:457:G:C8	2.49	0.43
52:sa:1264:U:H2'	52:sa:1265:U:O4'	2.19	0.43
52:sa:1266:A:H8	66:sq:78:HIS:CG	2.36	0.43
52:sa:1548:C:H2'	52:sa:1549:U:C6	2.54	0.43
52:sa:1584:G:H2'	52:sa:1585:U:C6	2.53	0.43
57:sg:51:LEU:HD23	57:sg:51:LEU:HA	1.90	0.43
60:sj:105:ASP:OD1	60:sj:105:ASP:N	2.43	0.43
69:st:32:LYS:O	69:st:47:ARG:NH1	2.51	0.43
1:lA:206:U:O2	1:lA:303:G:N2	2.35	0.43
1:lA:213:A:H5'	14:lN:133:LYS:HB2	2.00	0.43
1:lA:265:A:H1'	1:lA:266:A:H2	1.83	0.43
1:lA:720:U:H2'	1:lA:721:A:C8	2.53	0.43
1:lA:1064:C:O2'	1:lA:1540:A:H1'	2.18	0.43
1:lA:1430:G:O6	1:lA:2442:C:O2'	2.32	0.43
1:lA:2909:A:C6	1:lA:2910:G:N7	2.87	0.43
2:lB:68:A:H2'	2:lB:69:U:C6	2.53	0.43
2:lB:80:A:O2'	2:lB:81:A:OP1	2.33	0.43
4:lD:234:LYS:HE3	4:lD:234:LYS:HB2	1.78	0.43
5:lE:122:TRP:CZ2	5:lE:127:LYS:HG2	2.53	0.43
5:lE:347:THR:O	5:lE:352:GLN:NE2	2.52	0.43
17:lQ:4:TYR:HE2	17:lQ:46:GLU:HA	1.84	0.43
17:lQ:121:ILE:HD13	17:lQ:121:ILE:HA	1.77	0.43
25:lY:38:GLU:OE1	25:lY:38:GLU:N	2.47	0.43
32:lf:11:VAL:O	32:lf:15:ARG:N	2.52	0.43
33:lg:41:ASN:HB3	33:lg:44:ARG:HG2	2.00	0.43
34:lh:99:ASN:HA	34:lh:102:LYS:HB3	2.00	0.43
37:lk:48:GLU:HG2	37:lk:84:ILE:HD11	1.99	0.43
50:sJ:48:U:H5'	50:sJ:49:C:OP1	2.19	0.43
52:sa:1143:U:C2	52:sa:1144:G:C8	3.07	0.43
75:sa:5101:PAR:O44	75:sa:5101:PAR:N64	2.51	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:sc:221:ARG:NE	73:sx:25:ASP:OD2	2.51	0.43
56:sf:177:ASN:HA	56:sf:229:GLY:H	1.82	0.43
72:sw:67:LYS:HE3	72:sw:78:ASP:HB3	2.01	0.43
1:lA:443:A:H4'	1:lA:444:A:OP1	2.19	0.43
1:lA:503:A:H2'	1:lA:504:U:C6	2.54	0.43
1:lA:549:U:O2'	1:lA:550:A:H8	2.02	0.43
1:lA:687:U:C2	1:lA:688:A:C8	3.06	0.43
1:lA:1562:A:H2'	29:lc:3:THR:HG21	2.01	0.43
1:lA:1751:A:N3	1:lA:1751:A:H2'	2.34	0.43
1:lA:2726:A:N7	1:lA:2728:G:C8	2.87	0.43
5:lE:34:LYS:HE3	5:lE:34:LYS:HB2	1.85	0.43
13:lM:164:LYS:HD3	13:lM:171:VAL:HG22	2.01	0.43
15:lO:173:GLN:OE1	15:lO:173:GLN:HA	2.18	0.43
21:lU:133:LYS:HB2	21:lU:133:LYS:HE3	1.82	0.43
25:lY:29:LEU:O	35:li:54:ARG:NH2	2.42	0.43
52:sa:547:G:OP2	52:sa:548:C:O2'	2.30	0.43
52:sa:1170:U:C2	52:sa:1171:U:C5	3.07	0.43
52:sa:1256:A:O2'	62:sl:2:ARG:NH2	2.45	0.43
52:sa:1694:A:OP1	68:ss:57:PRO:HD2	2.19	0.43
52:sa:1932:U:OP2	65:sp:142:ARG:NH2	2.52	0.43
53:sc:128:LYS:HB3	53:sc:128:LYS:HE2	1.68	0.43
56:sf:63:VAL:HG11	56:sf:78:THR:HA	2.01	0.43
57:sg:83:LYS:HE3	57:sg:83:LYS:HB2	1.76	0.43
1:lA:747:C:H1'	8:lH:51:ASN:HD22	1.84	0.43
1:lA:814:G:H2'	1:lA:815:A:O4'	2.19	0.43
1:lA:817:A:O2'	1:lA:818:A:O4'	2.36	0.43
1:lA:822:C:H2'	1:lA:823:A:C8	2.54	0.43
1:lA:1444:U:O4	1:lA:1445:U:O4	2.36	0.43
1:lA:1856:G:H2'	1:lA:1857:A:H8	1.84	0.43
1:lA:1857:A:OP1	23:lW:86:LYS:HB2	2.19	0.43
1:lA:1889:U:H2'	1:lA:1890:U:C6	2.53	0.43
1:lA:3331:C:O2	32:lf:8:ASN:ND2	2.52	0.43
1:lA:3483:A:H2'	1:lA:3484:A:C2	2.54	0.43
1:lA:3486:A:H8	1:lA:3486:A:O5'	2.01	0.43
3:lC:27:G:H2'	3:lC:28:U:H5'	2.01	0.43
7:lG:91:GLY:O	7:lG:94:ASN:ND2	2.48	0.43
12:lL:39:ARG:H	12:lL:39:ARG:HG3	1.67	0.43
32:lf:80:LYS:HD2	32:lf:80:LYS:HA	1.78	0.43
48:sE:47:ALA:HB1	48:sE:52:PHE:HB2	2.01	0.43
52:sa:338:C:H2'	52:sa:339:A:H8	1.82	0.43
52:sa:1282:U:H5	62:sl:5:THR:HG22	1.84	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:sa:1356:A:H61	54:sd:175:GLY:HA2	1.83	0.43
58:sh:102:VAL:HG13	58:sh:106:ILE:HG13	2.00	0.43
1:lA:233:A:N3	1:lA:254:C:O2'	2.41	0.43
1:lA:865:U:H2'	1:lA:866:G:C8	2.54	0.43
1:lA:1403:C:H2'	1:lA:1404:C:C6	2.53	0.43
1:lA:1558:C:H2'	1:lA:1559:C:C6	2.54	0.43
1:lA:1636:U:H5	1:lA:2036:A:N1	2.16	0.43
1:lA:2429:G:H5''	18:lR:86:LYS:HB2	2.00	0.43
2:lB:30:U:C2	2:lB:31:C:C5	3.07	0.43
5:lE:19:ARG:HB3	5:lE:275:PHE:CZ	2.54	0.43
6:lF:162:ILE:CD1	6:lF:171:LEU:HD22	2.48	0.43
6:lF:288:VAL:O	6:lF:290:ASP:N	2.51	0.43
6:lF:358:MET:HA	6:lF:361:ILE:HD12	2.00	0.43
6:lF:404:THR:OG1	6:lF:405:ALA:N	2.51	0.43
7:lG:50:ARG:NH2	7:lG:148:ASP:OD2	2.30	0.43
8:lH:85:LYS:HA	8:lH:85:LYS:HD2	1.87	0.43
11:lK:44:VAL:HG12	11:lK:67:MET:HB2	2.01	0.43
14:lN:153:LYS:HA	14:lN:153:LYS:HD3	1.84	0.43
29:lc:103:ASP:OD1	29:lc:106:LYS:HG2	2.19	0.43
45:sB:23:CYS:HB3	45:sB:28:ARG:H	1.84	0.43
52:sa:866:G:C2'	52:sa:867:A:H8	2.31	0.43
52:sa:925:U:H2'	52:sa:926:U:C6	2.54	0.43
52:sa:1195:G:C2	52:sa:1196:A:C8	3.07	0.43
52:sa:1735:A:H2'	52:sa:1736:A:H8	1.84	0.43
52:sa:1843:U:H2'	52:sa:1844:A:C8	2.54	0.43
54:sd:193:MET:HE2	54:sd:193:MET:HA	2.01	0.43
64:so:98:VAL:HG11	64:so:115:LEU:HB2	2.01	0.43
66:sq:106:ILE:HA	66:sq:110:MET:HE2	2.00	0.43
1:lA:421:A:H2	1:lA:439:U:O2	2.02	0.43
1:lA:424:A:OP2	1:lA:438:G:H1'	2.18	0.43
1:lA:2059:A:O2'	1:lA:2060:A:OP2	2.34	0.43
6:lF:150:ILE:HG22	6:lF:151:PRO:HD3	2.01	0.43
16:lP:29:GLU:HB3	16:lP:37:LEU:HB3	2.01	0.43
25:lY:32:ASN:OD1	25:lY:32:ASN:N	2.52	0.43
28:lb:4:ILE:HD11	31:le:38:ILE:HG22	2.01	0.43
39:lm:33:GLN:HE21	39:lm:49:ARG:NH2	2.16	0.43
45:sB:87:ARG:NH2	45:sB:94:ASN:O	2.52	0.43
47:sD:49:PRO:HB3	52:sa:1781:C:OP1	2.19	0.43
52:sa:553:C:N4	52:sa:554:U:O4	2.52	0.43
52:sa:889:U:H2'	52:sa:890:C:C6	2.54	0.43
52:sa:1677:A:H5'	52:sa:1678:A:OP1	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:sc:66:LYS:HD2	53:sc:66:LYS:HA	1.67	0.43
1:lA:411:A:H4'	6:lF:86:ARG:HD3	2.01	0.42
1:lA:1945:A:N6	1:lA:1956:G:O6	2.52	0.42
1:lA:2111:A:H2'	1:lA:2112:C:H6	1.84	0.42
1:lA:2184:A:H2	1:lA:3471:A:N3	2.16	0.42
1:lA:2427:U:H2'	1:lA:2428:A:C8	2.53	0.42
1:lA:2461:G:O2'	1:lA:2462:G:P	2.77	0.42
1:lA:2732:A:H2'	1:lA:2733:A:C8	2.54	0.42
1:lA:2937:G:N2	50:sJ:77:A:O2'	2.52	0.42
6:lF:271:LYS:HE3	6:lF:271:LYS:HB2	1.81	0.42
14:lN:143:GLU:OE1	14:lN:143:GLU:N	2.43	0.42
18:lR:16:GLN:HG2	18:lR:149:ILE:HG12	2.01	0.42
20:lT:73:ILE:HG21	20:lT:102:MET:SD	2.59	0.42
23:lW:36:ASN:OD1	23:lW:36:ASN:N	2.51	0.42
24:lX:14:PHE:HE2	24:lX:91:LYS:HG2	1.83	0.42
26:lZ:24:VAL:HG22	26:lZ:30:SER:HB2	2.01	0.42
27:la:83:LEU:HD12	27:la:83:LEU:HA	1.88	0.42
28:lb:12:ILE:HD13	28:lb:137:LEU:HD22	2.01	0.42
28:lb:122:ALA:HB1	28:lb:126:ARG:HH21	1.84	0.42
32:lf:5:LYS:HD2	32:lf:5:LYS:HA	1.82	0.42
52:sa:122:G:O2'	56:sf:144:THR:N	2.46	0.42
52:sa:122:G:C6	52:sa:291:A:C6	3.07	0.42
52:sa:1745:U:H1'	68:ss:154:GLN:HG2	2.01	0.42
1:lA:11:A:H2'	1:lA:12:U:H6	1.84	0.42
1:lA:221:U:H2'	1:lA:222:A:H8	1.84	0.42
1:lA:463:U:O2'	1:lA:464:G:H5'	2.19	0.42
1:lA:560:U:O2'	1:lA:561:G:H5'	2.19	0.42
1:lA:686:A:H2'	1:lA:687:U:H5''	2.02	0.42
1:lA:699:A:H5''	1:lA:746:A:N6	2.33	0.42
1:lA:715:A:C8	1:lA:727:U:H5''	2.54	0.42
1:lA:1285:C:P	19:lS:2:ALA:HB2	2.58	0.42
1:lA:1572:U:H1'	6:lF:96:THR:HA	2.01	0.42
2:lB:23:U:O2'	2:lB:24:U:H5'	2.19	0.42
2:lB:31:C:H5''	14:lN:25:LYS:HG2	2.01	0.42
4:lD:27:ALA:HB2	4:lD:58:LEU:HD21	2.02	0.42
20:lT:73:ILE:HG12	20:lT:122:ILE:HD12	2.02	0.42
52:sa:29:G:H22	52:sa:541:U:H4'	1.84	0.42
52:sa:1586:U:H2'	52:sa:1587:A:C8	2.54	0.42
52:sa:1716:A:H2'	52:sa:1717:U:C6	2.54	0.42
52:sa:1724:U:OP1	70:su:121:HIS:ND1	2.43	0.42
56:sf:196:ARG:HA	56:sf:206:VAL:HA	2.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:sg:20:LEU:HD21	57:sg:71:VAL:HG21	2.02	0.42
60:sj:11:ARG:O	63:sm:133:LYS:NZ	2.49	0.42
66:sq:29:GLU:O	66:sq:33:VAL:HG13	2.18	0.42
66:sq:111:LEU:HB3	66:sq:112:GLY:H	1.73	0.42
1:lA:517:G:C6	1:lA:556:A:C6	3.07	0.42
1:lA:1195:G:N3	30:ld:46:GLN:HB2	2.34	0.42
1:lA:1562:A:OP2	29:lc:3:THR:HG22	2.19	0.42
1:lA:1886:A:H2'	1:lA:1887:A:C8	2.55	0.42
1:lA:2079:A:H3'	1:lA:2079:A:N3	2.34	0.42
1:lA:2382:C:H4'	1:lA:2383:G:H5'	2.01	0.42
1:lA:3321:U:H2'	1:lA:3322:A:O4'	2.20	0.42
2:lB:81:A:O2'	2:lB:82:A:OP1	2.37	0.42
6:lF:106:LYS:O	6:lF:109:ARG:HG3	2.19	0.42
11:lK:174:ARG:HA	11:lK:174:ARG:HD3	1.83	0.42
12:lL:4:ARG:N	12:lL:123:TYR:OH	2.44	0.42
13:lM:35:LYS:HA	13:lM:35:LYS:HD2	1.66	0.42
14:lN:51:SER:O	14:lN:53:LYS:HE3	2.19	0.42
20:lT:77:TYR:HD2	20:lT:112:THR:HG21	1.84	0.42
39:lm:7:LYS:O	39:lm:27:LYS:NZ	2.42	0.42
52:sa:555:G:H2'	52:sa:556:G:C8	2.55	0.42
52:sa:903:A:H2'	52:sa:904:A:C8	2.54	0.42
52:sa:919:A:H2'	52:sa:920:A:C8	2.54	0.42
52:sa:1311:U:O2'	52:sa:1312:A:O5'	2.37	0.42
55:se:80:LYS:HE2	55:se:81:PHE:CE1	2.54	0.42
60:sj:44:HIS:ND1	60:sj:58:LEU:HD11	2.34	0.42
68:ss:54:ILE:HG22	68:ss:60:ARG:HG3	2.01	0.42
71:sv:9:ASP:N	71:sv:9:ASP:OD1	2.52	0.42
1:lA:38:C:O2'	1:lA:39:A:H5'	2.19	0.42
1:lA:317:U:O2'	1:lA:364:A:H1'	2.19	0.42
1:lA:458:U:H5''	18:lR:37:ARG:NH2	2.35	0.42
1:lA:745:A:O2'	1:lA:746:A:N3	2.48	0.42
1:lA:1121:A:N1	1:lA:1168:C:O2'	2.41	0.42
1:lA:1620:U:H2'	1:lA:1621:G:H5'	2.02	0.42
6:lF:2:SER:O	6:lF:2:SER:OG	2.33	0.42
6:lF:358:MET:HA	6:lF:358:MET:HE2	2.01	0.42
7:lG:219:PHE:O	7:lG:223:ILE:HG13	2.19	0.42
12:lL:24:ARG:HB2	12:lL:24:ARG:HH11	1.84	0.42
17:lQ:45:PRO:HA	17:lQ:48:ALA:HB3	2.00	0.42
47:sD:47:LYS:HD3	57:sg:147:ARG:HE	1.85	0.42
52:sa:155:A:H2'	52:sa:156:U:O4'	2.19	0.42
52:sa:1111:A:H2'	52:sa:1112:A:C8	2.55	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:sa:1542:G:H4'	52:sa:1543:A:H5'	2.00	0.42
53:sc:127:ILE:O	53:sc:131:MET:HG2	2.19	0.42
60:sj:38:LEU:HD23	60:sj:96:LEU:HD21	2.01	0.42
68:ss:53:LEU:HA	68:ss:60:ARG:NH2	2.35	0.42
72:sw:22:ILE:HG22	72:sw:89:LEU:HD23	2.02	0.42
74:sy:50:GLN:HE22	74:sy:78:LYS:NZ	2.17	0.42
1:lA:721:A:C2	1:lA:722:A:C5	3.07	0.42
1:lA:721:A:O2'	1:lA:722:A:OP1	2.37	0.42
1:lA:765:A:C2	36:lj:92:PRO:HG2	2.54	0.42
1:lA:915:U:O2'	1:lA:916:U:H5'	2.18	0.42
1:lA:1961:G:H2'	1:lA:1962:U:O4'	2.19	0.42
1:lA:2649:G:OP2	1:lA:2650:A:O2'	2.30	0.42
1:lA:2872:A:N6	1:lA:2882:G:O6	2.52	0.42
1:lA:3129:U:C4	1:lA:3130:A:N7	2.88	0.42
5:lE:238:ARG:HE	5:lE:238:ARG:HB2	1.67	0.42
36:lj:86:PHE:CE1	36:lj:90:LEU:HD11	2.54	0.42
52:sa:1693:A:HO2'	52:sa:1694:A:P	2.41	0.42
60:sj:42:ARG:HD3	60:sj:59:ARG:NH2	2.35	0.42
65:sp:81:LYS:HA	65:sp:81:LYS:HD2	1.83	0.42
68:ss:28:LYS:HD2	68:ss:28:LYS:HA	1.88	0.42
70:su:126:HIS:CE1	70:su:132:VAL:HG21	2.55	0.42
1:lA:285:A:H4'	1:lA:286:U:H5'	2.01	0.42
1:lA:871:A:H4'	1:lA:872:A:OP1	2.19	0.42
1:lA:1239:U:H5''	29:lc:22:ILE:HD12	2.01	0.42
1:lA:1816:A:N3	1:lA:1892:C:O2'	2.48	0.42
1:lA:1858:A:HO2'	1:lA:1943:U:HO2'	1.59	0.42
1:lA:1977:C:N4	1:lA:1978:U:O4	2.53	0.42
1:lA:2632:A:O2'	28:lb:139:PHE:O	2.35	0.42
3:lC:18:U:H3	3:lC:58:A:N6	2.11	0.42
3:lC:29:G:O6	3:lC:46:G:O6	2.38	0.42
3:lC:84:G:N2	3:lC:91:G:N2	2.68	0.42
6:lF:7:VAL:HG13	6:lF:149:GLN:NE2	2.33	0.42
8:lH:120:SER:C	8:lH:121:LYS:HG2	2.44	0.42
13:lM:108:GLU:HG2	13:lM:122:ILE:HD12	2.02	0.42
23:lW:49:PHE:CE2	23:lW:96:TYR:HB2	2.54	0.42
37:lk:39:ARG:HD3	37:lk:39:ARG:HA	1.75	0.42
42:lp:24:CYS:CA	42:lp:40:CYS:HB3	2.50	0.42
47:sD:49:PRO:HG2	57:sg:146:LEU:HB3	2.00	0.42
52:sa:781:G:H5''	52:sa:782:A:O3'	2.20	0.42
52:sa:959:A:H4'	52:sa:1933:G:H21	1.84	0.42
52:sa:1378:U:H2'	52:sa:1379:C:H6	1.84	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:sd:61:GLN:HG2	54:sd:101:TYR:HE2	1.85	0.42
70:su:85:LEU:HD13	70:su:97:SER:H	1.84	0.42
1:lA:200:A:C4	1:lA:311:A:C6	3.08	0.42
1:lA:223:A:H2'	1:lA:224:U:C6	2.54	0.42
1:lA:605:C:H2'	1:lA:606:A:H8	1.84	0.42
1:lA:775:C:OP1	29:lc:21:ARG:HB3	2.20	0.42
1:lA:868:A:H2'	1:lA:869:G:H8	1.85	0.42
1:lA:1231:A:H2'	1:lA:1232:C:O4'	2.19	0.42
1:lA:1259:G:O2'	1:lA:2712:A:N3	2.44	0.42
1:lA:1904:U:O2'	1:lA:1906:A:N7	2.46	0.42
1:lA:1920:A:H2'	1:lA:1921:A:C8	2.55	0.42
42:lp:7:LEU:HD12	42:lp:7:LEU:HA	1.82	0.42
52:sa:744:A:H2'	52:sa:745:U:O4'	2.19	0.42
52:sa:1335:U:H2'	52:sa:1336:U:C6	2.55	0.42
59:si:118:LYS:HA	59:si:118:LYS:HD2	1.69	0.42
60:sj:22:GLN:HG2	60:sj:23:LYS:O	2.19	0.42
72:sw:55:ARG:HG2	72:sw:87:ARG:NH1	2.35	0.42
1:lA:235:U:C5	1:lA:252:G:N1	2.88	0.42
1:lA:338:A:H2'	1:lA:339:U:C6	2.54	0.42
1:lA:570:A:C4	1:lA:571:A:C8	3.07	0.42
1:lA:761:A:H2'	1:lA:762:A:C8	2.54	0.42
1:lA:1476:G:O2'	1:lA:1477:A:N3	2.48	0.42
1:lA:1975:G:H8	1:lA:1975:G:OP2	2.03	0.42
1:lA:2083:A:H2'	1:lA:2084:G:H8	1.85	0.42
1:lA:2205:C:H2'	1:lA:2206:G:H8	1.85	0.42
1:lA:2360:C:N4	1:lA:2384:C:O4'	2.53	0.42
1:lA:2734:C:O2'	1:lA:2735:U:H5'	2.20	0.42
1:lA:2867:A:H62	1:lA:2887:G:H1	1.67	0.42
1:lA:2867:A:H62	1:lA:2887:G:H22	1.67	0.42
1:lA:3095:G:H2'	1:lA:3096:U:O4'	2.20	0.42
3:lC:14:G:C2	3:lC:64:U:C4	3.08	0.42
5:lE:14:LEU:HD22	5:lE:264:TRP:CZ3	2.55	0.42
11:lK:60:GLU:HG3	11:lK:61:ASN:N	2.34	0.42
13:lM:35:LYS:O	13:lM:39:THR:HG23	2.20	0.42
28:lb:22:LYS:NZ	28:lb:135:THR:O	2.43	0.42
52:sa:291:A:H2'	52:sa:292:A:H8	1.84	0.42
52:sa:327:U:OP1	60:sj:31:ARG:HD2	2.20	0.42
52:sa:415:A:O2'	52:sa:416:A:H5'	2.20	0.42
67:sr:46:TYR:HB3	67:sr:69:LEU:HD13	2.02	0.42
1:lA:625:A:C6	1:lA:663:G:C5	3.08	0.42
1:lA:812:A:C5	1:lA:839:G:N2	2.88	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:948:C:O2'	1:1A:949:G:N7	2.50	0.42
1:1A:2116:G:O2'	21:1U:82:LYS:O	2.36	0.42
1:1A:3469:A:H5''	1:1A:3470:G:C8	2.55	0.42
3:1C:107:A:O2'	3:1C:108:U:H5'	2.20	0.42
4:1D:180:LEU:HA	4:1D:180:LEU:HD23	1.80	0.42
11:1K:37:ARG:HD3	11:1K:39:PHE:CE1	2.55	0.42
19:1S:27:LEU:HD12	19:1S:27:LEU:HA	1.86	0.42
27:1a:150:ASN:O	27:1a:153:ILE:HG12	2.20	0.42
27:1a:165:LYS:HE2	27:1a:165:LYS:HB2	1.75	0.42
52:sa:906:A:H2'	52:sa:907:U:O4'	2.20	0.42
52:sa:1093:U:H2'	52:sa:1094:G:H8	1.85	0.42
52:sa:1141:U:H2'	52:sa:1142:A:C8	2.51	0.42
55:se:116:VAL:HG22	55:se:144:PHE:HZ	1.85	0.42
60:sj:38:LEU:HD11	60:sj:80:ASN:HA	2.01	0.42
63:sm:92:TYR:HB2	63:sm:99:TYR:CE1	2.54	0.42
65:sp:30:ALA:HB2	65:sp:35:THR:HG23	2.01	0.42
66:sq:123:LYS:HB3	66:sq:123:LYS:HE2	1.78	0.42
1:1A:227:A:H2'	1:1A:228:A:H8	1.85	0.42
1:1A:474:A:H2'	1:1A:475:A:C8	2.54	0.42
1:1A:731:A:H2'	1:1A:732:U:C6	2.55	0.42
1:1A:1479:A:H4'	1:1A:1480:U:O5'	2.18	0.42
1:1A:1922:U:H2'	1:1A:1923:U:C6	2.55	0.42
1:1A:2062:A:O2'	1:1A:3220:U:H5''	2.20	0.42
1:1A:2116:G:P	1:1A:2133:G:H22	2.43	0.42
1:1A:2324:C:O2'	1:1A:2348:G:H1'	2.19	0.42
1:1A:2665:A:H2'	1:1A:2666:G:H8	1.85	0.42
1:1A:3479:A:C6	1:1A:3480:A:C6	3.07	0.42
4:1D:247:ARG:HD2	4:1D:247:ARG:HA	1.72	0.42
6:1F:386:LYS:H	6:1F:386:LYS:HG3	1.66	0.42
6:1F:429:ILE:O	20:1T:24:ARG:NH1	2.53	0.42
8:1H:62:GLY:O	8:1H:117:VAL:N	2.51	0.42
17:1Q:70:ASN:ND2	17:1Q:92:ILE:O	2.29	0.42
24:1X:91:LYS:HD2	24:1X:91:LYS:HA	1.84	0.42
45:sB:75:VAL:HB	52:sa:1940:G:N1	2.35	0.42
52:sa:5:U:H2'	52:sa:6:G:H8	1.85	0.42
52:sa:330:A:O2'	63:sm:130:PRO:O	2.36	0.42
52:sa:540:U:H2'	52:sa:541:U:H6	1.83	0.42
52:sa:597:U:H2'	52:sa:598:A:C8	2.55	0.42
52:sa:783:A:N1	52:sa:785:A:H5'	2.34	0.42
52:sa:1185:A:H2'	52:sa:1186:C:H6	1.85	0.42
52:sa:1905:U:C4	52:sa:1906:C:N4	2.88	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:sd:120:LYS:HG3	54:sd:189:VAL:HB	2.01	0.42
55:se:193:ASN:OD1	55:se:194:LYS:N	2.53	0.42
56:sf:79:ARG:HE	56:sf:79:ARG:HB2	1.64	0.42
57:sg:78:MET:HE2	57:sg:171:LEU:HD21	2.02	0.42
1:lA:359:A:H2'	1:lA:360:U:C6	2.55	0.41
1:lA:672:G:OP2	6:lF:403:ARG:NH1	2.49	0.41
1:lA:2752:C:H2'	1:lA:2753:U:H6	1.84	0.41
1:lA:2846:U:H2'	1:lA:2847:A:C8	2.54	0.41
1:lA:3303:A:H2'	1:lA:3304:C:C6	2.56	0.41
5:lE:146:ARG:O	5:lE:150:ILE:HG12	2.19	0.41
8:lH:45:LEU:HD11	8:lH:129:VAL:HG12	2.02	0.41
9:lI:30:LYS:HD3	9:lI:30:LYS:HA	1.78	0.41
9:lI:76:ARG:HG2	9:lI:99:ILE:HA	2.02	0.41
9:lI:105:ILE:HG21	9:lI:115:LEU:HD11	2.01	0.41
14:lN:49:ILE:HG12	14:lN:49:ILE:O	2.19	0.41
15:lO:50:MET:O	15:lO:54:GLN:HG2	2.20	0.41
15:lO:87:GLY:O	15:lO:91:LYS:NZ	2.52	0.41
17:lQ:62:TYR:CD1	17:lQ:134:LEU:HD12	2.55	0.41
24:lX:14:PHE:CE2	24:lX:91:LYS:HG2	2.54	0.41
28:lb:16:GLY:O	34:lh:74:ARG:HG3	2.20	0.41
28:lb:93:ASN:H	28:lb:120:GLN:HE22	1.66	0.41
52:sa:583:C:HO2'	52:sa:584:C:H6	1.67	0.41
61:sk:43:GLN:H	61:sk:43:GLN:HG3	1.67	0.41
65:sp:26:ALA:HB2	65:sp:88:LEU:HD13	2.01	0.41
1:lA:610:A:N6	1:lA:687:U:O4	2.53	0.41
1:lA:723:U:O2'	1:lA:724:A:N3	2.45	0.41
1:lA:764:U:H2'	1:lA:765:A:C8	2.52	0.41
1:lA:2040:A:O2'	1:lA:2041:U:H5''	2.20	0.41
1:lA:2107:G:O6	5:lE:243:LYS:NZ	2.33	0.41
1:lA:2341:C:N4	1:lA:2342:U:O4	2.53	0.41
1:lA:2747:G:H2'	1:lA:2747:G:N3	2.35	0.41
4:lD:105:GLY:HA3	4:lD:160:SER:HB3	2.01	0.41
12:lL:179:GLU:OE2	12:lL:179:GLU:C	2.63	0.41
23:lW:111:ASP:OD2	23:lW:113:LYS:HB2	2.20	0.41
40:ln:9:LYS:HA	40:ln:9:LYS:HD3	1.82	0.41
41:lo:44:TRP:O	41:lo:48:LYS:NZ	2.35	0.41
47:sD:25:ARG:HG2	52:sa:1785:C:N3	2.35	0.41
52:sa:1170:U:H2'	52:sa:1171:U:C6	2.55	0.41
52:sa:1937:A:O2'	52:sa:1938:A:H5'	2.20	0.41
54:sd:38:LEU:HD13	62:sl:64:TYR:CE2	2.55	0.41
57:sg:36:LYS:HA	57:sg:36:LYS:HD2	1.80	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
63:sm:37:ILE:HD11	63:sm:65:ILE:HD12	2.01	0.41
68:ss:88:GLY:H	68:ss:91:ALA:HB3	1.85	0.41
1:lA:606:A:H2'	1:lA:607:U:C6	2.56	0.41
1:lA:780:A:OP2	33:lg:40:ASP:HB3	2.20	0.41
1:lA:1259:G:N3	1:lA:2712:A:H2	2.18	0.41
1:lA:2262:U:OP2	4:ID:200:ARG:HD2	2.20	0.41
1:lA:2304:A:H2'	1:lA:2305:C:H6	1.85	0.41
1:lA:3008:U:O2'	1:lA:3009:C:P	2.79	0.41
1:lA:3008:U:HO2'	1:lA:3009:C:P	2.40	0.41
1:lA:3162:A:N1	1:lA:3295:A:C6	2.88	0.41
2:lB:42:A:OP2	2:lB:101:G:N1	2.38	0.41
3:lC:84:G:C2	3:lC:85:G:C8	3.08	0.41
6:lF:206:LYS:HD2	6:lF:206:LYS:HA	1.83	0.41
15:lO:51:ARG:HA	15:lO:51:ARG:HD2	1.70	0.41
19:lS:72:LYS:H	19:lS:72:LYS:HG2	1.68	0.41
21:lU:77:GLY:O	21:lU:81:ARG:NE	2.54	0.41
27:la:81:ASP:OD2	27:la:82:LYS:N	2.53	0.41
49:sI:5:C:H2'	49:sI:6:C:C6	2.55	0.41
52:sa:607:G:OP1	52:sa:607:G:N2	2.45	0.41
62:sl:4:ALA:HB3	62:sl:7:ASP:HB2	2.01	0.41
67:sr:48:THR:OG1	67:sr:49:ASP:OD1	2.29	0.41
68:ss:54:ILE:HG23	68:ss:56:PRO:HD2	2.02	0.41
69:st:21:TYR:C	69:st:21:TYR:CD1	2.99	0.41
1:lA:60:A:OP2	17:lQ:169:LYS:NZ	2.49	0.41
1:lA:311:A:H5'	37:lk:30:ARG:CB	2.49	0.41
1:lA:918:G:O2'	14:lN:16:TRP:NE1	2.52	0.41
1:lA:1203:U:H3'	1:lA:1204:A:H2	1.85	0.41
1:lA:1641:A:H2'	1:lA:1642:A:O4'	2.20	0.41
1:lA:1920:A:H2'	1:lA:1921:A:H8	1.86	0.41
1:lA:2866:C:H4'	1:lA:2867:A:OP1	2.20	0.41
1:lA:3031:U:C2	1:lA:3054:A:N6	2.88	0.41
1:lA:3286:A:C2	1:lA:3287:U:H5'	2.55	0.41
1:lA:3457:G:H1	1:lA:3495:U:H3	1.68	0.41
4:ID:108:PRO:O	4:ID:111:THR:OG1	2.28	0.41
4:ID:186:TYR:HB2	4:ID:196:TRP:CZ3	2.56	0.41
5:lE:264:TRP:H	5:lE:264:TRP:CD1	2.39	0.41
6:lF:6:PHE:HD2	6:lF:21:PRO:HB2	1.85	0.41
7:lG:237:ILE:HD13	7:lG:237:ILE:HA	1.88	0.41
8:lH:97:ILE:HD13	8:lH:177:ILE:HG21	2.01	0.41
8:lH:146:VAL:HG13	8:lH:147:GLU:H	1.85	0.41
21:lU:74:ARG:HD3	21:lU:74:ARG:HA	1.84	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:lb:83:THR:HG22	34:lh:93:PHE:CE1	2.56	0.41
45:sB:85:ARG:H	52:sa:1944:A:H61	1.68	0.41
52:sa:958:A:O2'	52:sa:1934:C:O2	2.31	0.41
52:sa:972:A:H2'	52:sa:973:A:H5'	2.02	0.41
52:sa:1580:U:O2'	57:sg:84:ASN:OD1	2.38	0.41
52:sa:1893:A:H4'	52:sa:1893:A:OP1	2.20	0.41
53:sc:85:MET:HE3	53:sc:100:LEU:HB3	2.03	0.41
57:sg:31:SER:OG	57:sg:32:ASP:N	2.54	0.41
66:sq:55:LEU:HD23	66:sq:55:LEU:HA	1.86	0.41
67:sr:69:LEU:HD12	67:sr:69:LEU:HA	1.85	0.41
1:lA:790:A:H2'	1:lA:791:A:C8	2.55	0.41
1:lA:1219:A:H4'	1:lA:1220:A:OP2	2.20	0.41
1:lA:1533:U:HO2'	1:lA:1534:U:P	2.41	0.41
1:lA:2634:A:O2'	10:IJ:30:VAL:HG13	2.20	0.41
2:lB:144:A:H2'	2:lB:145:U:H6	1.84	0.41
6:lF:345:LYS:HD2	6:lF:345:LYS:HA	1.81	0.41
7:lG:147:LEU:HD22	7:lG:164:LEU:HD12	2.03	0.41
8:lH:179:LYS:HE2	8:lH:179:LYS:HB2	1.86	0.41
9:II:33:ILE:O	9:II:37:ILE:HG12	2.20	0.41
9:II:44:GLU:OE2	9:II:44:GLU:HA	2.21	0.41
10:IJ:45:TYR:CE1	10:IJ:46:VAL:HG23	2.55	0.41
49:sI:52:G:H2'	49:sI:53:G:C8	2.56	0.41
52:sa:247:G:P	56:sf:132:GLY:H	2.41	0.41
52:sa:878:A:C2	52:sa:895:A:C5	3.08	0.41
52:sa:1208:A:H61	66:sq:98:GLY:HA3	1.86	0.41
52:sa:1234:C:H42	52:sa:1562:G:N2	2.19	0.41
56:sf:39:CYS:HA	56:sf:83:GLY:HA2	2.02	0.41
57:sg:27:ASP:OD1	57:sg:27:ASP:N	2.50	0.41
57:sg:30:VAL:HB	57:sg:36:LYS:HD3	2.02	0.41
58:sh:77:LEU:HD23	58:sh:77:LEU:HA	1.82	0.41
60:sj:81:VAL:HA	60:sj:102:ILE:HG13	2.01	0.41
72:sw:56:ILE:HB	72:sw:86:LYS:HB3	2.02	0.41
1:lA:205:C:H3'	1:lA:206:U:H5''	2.02	0.41
1:lA:229:A:H2'	1:lA:230:U:C6	2.56	0.41
1:lA:693:A:N6	1:lA:694:A:H62	2.18	0.41
1:lA:790:A:H2'	1:lA:791:A:H8	1.85	0.41
1:lA:2273:C:H4'	1:lA:2274:A:O5'	2.20	0.41
1:lA:3432:A:O2'	1:lA:3433:U:P	2.78	0.41
3:lC:27:G:C2	3:lC:51:G:C2	3.08	0.41
19:IS:86:TYR:OH	29:lc:84:LYS:NZ	2.34	0.41
31:le:11:GLN:HG3	31:le:75:SER:OG	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:sE:34:TYR:CZ	72:sw:61:LEU:HG	2.55	0.41
52:sa:42:A:N1	52:sa:373:A:C6	2.89	0.41
52:sa:42:A:C2	52:sa:373:A:C6	3.09	0.41
52:sa:62:G:O2'	52:sa:165:U:OP1	2.39	0.41
52:sa:284:A:H2'	52:sa:284:A:N3	2.35	0.41
52:sa:1394:A:C5	52:sa:1395:A:C8	3.09	0.41
52:sa:1903:A:H5''	52:sa:1904:G:OP2	2.21	0.41
55:se:26:PHE:HD2	65:sp:83:LEU:HD21	1.84	0.41
61:sk:33:GLN:HG2	61:sk:34:PHE:CD1	2.55	0.41
64:so:103:GLU:HB2	64:so:104:LYS:NZ	2.35	0.41
65:sp:52:THR:O	65:sp:55:MET:HB2	2.21	0.41
1:lA:22:A:C5	1:lA:376:A:C8	3.09	0.41
1:lA:515:G:C6	1:lA:558:G:C5	3.08	0.41
1:lA:896:A:O2'	1:lA:2790:U:OP1	2.38	0.41
1:lA:985:C:N4	1:lA:1012:C:O2'	2.53	0.41
1:lA:1265:G:C6	1:lA:1266:C:N4	2.88	0.41
1:lA:1764:A:H2'	1:lA:1765:A:C8	2.56	0.41
1:lA:3371:U:H5''	16:lP:131:SER:HA	2.02	0.41
2:lB:143:U:H2'	2:lB:144:A:C8	2.55	0.41
3:lC:79:G:C2	3:lC:95:A:H2	2.33	0.41
13:lM:111:ASP:HB3	13:lM:112:LEU:H	1.52	0.41
16:lP:84:ALA:O	16:lP:88:ARG:HG3	2.20	0.41
23:lW:47:VAL:O	23:lW:51:ARG:HG2	2.21	0.41
30:ld:54:LYS:HE2	30:ld:54:LYS:HB2	1.72	0.41
36:lj:18:ALA:HB3	36:lj:21:ASN:HB2	2.03	0.41
52:sa:204:A:H3'	52:sa:205:A:H8	1.85	0.41
56:sf:121:LEU:HD12	56:sf:121:LEU:HA	1.93	0.41
72:sw:54:VAL:HB	72:sw:88:VAL:HB	2.02	0.41
73:sx:61:ILE:HD13	73:sx:61:ILE:HA	1.94	0.41
1:lA:249:U:N3	1:lA:250:G:N7	2.68	0.41
1:lA:408:A:HO2'	1:lA:1047:U:HO2'	1.68	0.41
1:lA:445:G:H5''	1:lA:448:C:O2'	2.20	0.41
1:lA:1686:G:O6	17:lQ:35:ARG:NH2	2.50	0.41
1:lA:2025:C:H2'	1:lA:2026:U:C6	2.55	0.41
1:lA:2349:G:O2'	1:lA:2387:G:O6	2.34	0.41
2:lB:15:U:C2	2:lB:16:U:C5	3.09	0.41
6:lF:78:ARG:HA	6:lF:89:GLN:O	2.21	0.41
6:lF:212:TYR:HB2	6:lF:216:ASP:HB2	2.02	0.41
13:lM:155:THR:H	13:lM:158:ASP:HB2	1.84	0.41
14:lN:194:LYS:HE2	14:lN:194:LYS:HB3	1.86	0.41
24:lX:113:LYS:HE2	24:lX:113:LYS:HB2	1.96	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:lb:115:LYS:HD3	28:lb:115:LYS:HA	1.66	0.41
37:lk:84:ILE:HD13	37:lk:84:ILE:HA	1.91	0.41
40:ln:57:LYS:C	40:ln:58:LYS:HD3	2.45	0.41
45:sB:5:ARG:NH2	52:sa:1942:U:OP2	2.50	0.41
47:sD:16:THR:OG1	47:sD:32:ARG:O	2.31	0.41
52:sa:102:A:H4'	52:sa:103:U:O5'	2.20	0.41
52:sa:1112:A:H2'	52:sa:1113:A:C8	2.55	0.41
52:sa:1780:A:H3'	52:sa:1780:A:N3	2.35	0.41
56:sf:141:ASP:OD1	56:sf:141:ASP:N	2.53	0.41
66:sq:85:LEU:HD23	66:sq:85:LEU:HA	1.82	0.41
72:sw:80:TYR:HD1	72:sw:80:TYR:HA	1.77	0.41
1:lA:22:A:H2'	1:lA:23:C:H6	1.85	0.41
1:lA:58:G:N2	17:lQ:186:ALA:HB1	2.35	0.41
1:lA:277:A:H2'	1:lA:278:U:H6	1.86	0.41
1:lA:602:U:O3'	6:lF:316:ARG:NH2	2.54	0.41
1:lA:739:A:H2'	1:lA:740:A:C8	2.55	0.41
1:lA:810:G:N2	1:lA:906:G:C5	2.89	0.41
1:lA:827:A:N1	2:lB:30:U:O2'	2.40	0.41
1:lA:1091:A:H4'	1:lA:1494:G:H4'	2.02	0.41
1:lA:1105:U:H2'	1:lA:1106:U:C6	2.55	0.41
1:lA:1234:U:H2'	1:lA:1235:U:C6	2.56	0.41
1:lA:1437:A:H2'	1:lA:1438:U:C6	2.54	0.41
1:lA:1945:A:C6	1:lA:1956:G:C6	3.09	0.41
1:lA:1956:G:C2	1:lA:1957:A:C8	3.08	0.41
1:lA:2439:A:O2'	1:lA:2440:G:H5'	2.21	0.41
1:lA:2679:A:N3	17:lQ:87:GLN:NE2	2.58	0.41
1:lA:2710:A:H2'	1:lA:2711:U:O4'	2.20	0.41
1:lA:2864:A:O2'	1:lA:2865:A:N3	2.41	0.41
1:lA:2887:G:H2'	1:lA:2888:U:O4'	2.20	0.41
1:lA:3368:G:H2'	1:lA:3369:U:C6	2.56	0.41
2:lB:39:A:C6	2:lB:102:A:C5	3.08	0.41
3:lC:17:G:H1	3:lC:59:G:H22	1.69	0.41
3:lC:27:G:N2	3:lC:50:A:C2	2.89	0.41
6:lF:285:GLN:HG2	19:lS:25:MET:HG2	2.03	0.41
7:lG:106:ALA:O	7:lG:110:LEU:HG	2.21	0.41
8:lH:168:GLN:H	8:lH:168:GLN:HG3	1.74	0.41
10:lJ:98:GLN:CD	10:lJ:98:GLN:N	2.79	0.41
13:lM:11:LYS:HA	13:lM:11:LYS:HD3	1.91	0.41
14:lN:248:LYS:NZ	29:lc:146:GLU:OE2	2.47	0.41
16:lP:12:VAL:HG22	16:lP:26:VAL:HG22	2.03	0.41
25:lY:43:LYS:HB2	25:lY:64:LEU:O	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:IY:84:VAL:HG12	25:IY:109:VAL:HG21	2.03	0.41
30:ld:18:ARG:HA	30:ld:18:ARG:HD2	1.89	0.41
35:li:70:LYS:HG3	35:li:73:ARG:HH21	1.85	0.41
36:lj:24:PRO:O	36:lj:89:ASN:ND2	2.54	0.41
37:lk:80:LYS:HA	37:lk:80:LYS:HD3	1.94	0.41
39:lm:47:VAL:HG13	39:lm:55:TRP:HB3	2.03	0.41
40:ln:11:ILE:O	40:ln:15:LEU:HD12	2.20	0.41
52:sa:93:C:H2'	52:sa:94:G:O4'	2.21	0.41
52:sa:889:U:H2'	52:sa:890:C:H6	1.86	0.41
52:sa:919:A:H2'	52:sa:920:A:H8	1.85	0.41
52:sa:1200:U:H2'	52:sa:1201:G:C8	2.56	0.41
52:sa:1266:A:H2'	52:sa:1267:A:C4	2.56	0.41
52:sa:1576:A:H2'	52:sa:1577:C:C6	2.56	0.41
52:sa:1756:U:O2	52:sa:1773:G:N2	2.54	0.41
52:sa:1906:C:H2'	52:sa:1907:G:C8	2.56	0.41
63:sm:79:MET:HG2	63:sm:82:THR:HB	2.03	0.41
65:sp:94:GLY:O	65:sp:96:GLY:N	2.54	0.41
68:ss:28:LYS:HG3	68:ss:29:LYS:H	1.86	0.41
74:sy:7:ALA:O	74:sy:11:MET:HG3	2.21	0.41
1:lA:366:U:H2'	1:lA:367:C:H6	1.86	0.41
1:lA:494:G:OP2	1:lA:494:G:H8	2.04	0.41
1:lA:670:A:H2	6:lF:397:LEU:HD23	1.86	0.41
1:lA:1308:A:O2'	1:lA:1309:A:H5'	2.21	0.41
1:lA:1629:G:H1'	34:lh:12:SER:HB3	2.01	0.41
1:lA:2468:C:HO2'	1:lA:2469:G:P	2.43	0.41
1:lA:2639:A:OP2	28:lb:56:LYS:NZ	2.50	0.41
1:lA:2746:A:H5'	1:lA:2747:G:C8	2.55	0.41
1:lA:3409:A:O2'	1:lA:3410:A:OP2	2.32	0.41
2:lB:38:G:C5	35:li:78:ARG:HD3	2.56	0.41
10:lJ:91:GLU:OE2	10:lJ:99:ARG:NH1	2.47	0.41
12:lL:28:ASP:OD2	12:lL:28:ASP:C	2.64	0.41
12:lL:38:ASN:OD1	12:lL:41:ALA:HB2	2.21	0.41
15:lO:117:LYS:HA	15:lO:117:LYS:HD2	1.92	0.41
16:lP:7:VAL:HG22	16:lP:59:LEU:HD21	2.02	0.41
17:lQ:38:ARG:NH2	17:lQ:60:VAL:HG12	2.35	0.41
40:ln:23:VAL:HG22	40:ln:36:VAL:HG12	2.03	0.41
46:sC:5:GLU:HG2	73:sx:86:ILE:HD12	2.03	0.41
52:sa:29:G:C5	52:sa:591:G:N1	2.89	0.41
52:sa:311:C:C2	52:sa:350:G:C2	3.09	0.41
52:sa:905:G:N2	52:sa:968:A:H1'	2.36	0.41
52:sa:934:A:H2'	52:sa:935:C:C6	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
67:sr:79:PHE:CD2	67:sr:89:TRP:HH2	2.39	0.41
1:lA:58:G:H22	1:lA:75:U:H4'	1.86	0.40
1:lA:661:G:H2'	1:lA:662:U:H6	1.86	0.40
1:lA:811:A:OP2	19:lS:103:THR:HG21	2.20	0.40
1:lA:1529:A:N3	2:lB:21:A:O2'	2.50	0.40
1:lA:1845:G:N2	1:lA:1924:U:O2'	2.54	0.40
1:lA:1921:A:O2'	34:lh:53:GLY:N	2.48	0.40
1:lA:2598:A:H2'	10:lJ:39:TYR:O	2.21	0.40
2:lB:101:G:OP2	2:lB:103:A:O2'	2.39	0.40
3:lC:84:G:N3	3:lC:84:G:H2'	2.36	0.40
7:lG:200:VAL:HG12	7:lG:204:MET:HE2	2.03	0.40
21:lU:99:MET:HB3	21:lU:103:ARG:HE	1.85	0.40
27:la:79:ASN:O	27:la:80:ILE:HD13	2.21	0.40
40:ln:30:GLU:N	40:ln:30:GLU:OE1	2.55	0.40
48:sE:21:CYS:HA	48:sE:30:LEU:HD21	2.02	0.40
50:sJ:23:G:O2'	50:sJ:24:C:P	2.79	0.40
52:sa:387:G:OP1	52:sa:1876:U:O2'	2.39	0.40
52:sa:1129:C:H2'	52:sa:1130:U:C6	2.56	0.40
52:sa:1428:U:N3	68:ss:23:GLN:OE1	2.53	0.40
56:sf:124:ILE:HD11	56:sf:158:ILE:HG23	2.02	0.40
59:si:158:SER:OG	59:si:194:LYS:NZ	2.35	0.40
66:sq:32:PHE:HD1	66:sq:35:MET:HE2	1.85	0.40
67:sr:76:SER:O	67:sr:77:PRO:C	2.63	0.40
69:st:66:VAL:HG12	69:st:68:GLY:H	1.86	0.40
70:su:25:ARG:H	70:su:25:ARG:HG3	1.72	0.40
71:sv:58:ALA:HA	71:sv:104:ILE:HG21	2.02	0.40
1:lA:51:G:O2'	1:lA:52:A:H5'	2.22	0.40
1:lA:229:A:H2'	1:lA:230:U:H6	1.87	0.40
1:lA:237:C:H2'	1:lA:238:A:C8	2.55	0.40
1:lA:338:A:H2'	1:lA:339:U:H6	1.86	0.40
1:lA:572:U:H2'	1:lA:573:A:H8	1.85	0.40
1:lA:586:A:N6	8:lH:43:THR:O	2.55	0.40
1:lA:720:U:C2	1:lA:721:A:C8	3.09	0.40
1:lA:808:G:C6	1:lA:908:A:N1	2.89	0.40
1:lA:955:A:O2'	39:lm:9:GLY:O	2.27	0.40
1:lA:1305:A:C5	1:lA:1451:A:C6	3.09	0.40
1:lA:1689:A:H61	1:lA:2242:A:N6	2.18	0.40
1:lA:2224:U:H2'	1:lA:2225:A:N7	2.36	0.40
1:lA:2865:A:H4'	37:lk:21:ALA:HA	2.02	0.40
1:lA:3130:A:H2'	1:lA:3131:C:C6	2.56	0.40
1:lA:3483:A:H2'	1:lA:3484:A:H2	1.85	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:IK:41:HIS:CE1	11:IK:77:VAL:HG11	2.56	0.40
18:IR:41:VAL:HG23	18:IR:95:LEU:HD22	2.02	0.40
20:IT:10:CYS:HG	20:IT:15:HIS:HD1	1.60	0.40
20:IT:148:VAL:HG23	20:IT:172:LEU:HD23	2.03	0.40
20:IT:148:VAL:HG22	20:IT:149:HIS:H	1.87	0.40
33:lg:14:TYR:OH	33:lg:52:LEU:HD21	2.22	0.40
37:lk:55:LEU:HD23	37:lk:55:LEU:HA	1.89	0.40
39:lm:16:THR:O	39:lm:16:THR:OG1	2.29	0.40
43:lq:57:LYS:HE2	43:lq:57:LYS:HB2	1.87	0.40
47:sD:25:ARG:HG2	52:sa:1785:C:C4	2.55	0.40
52:sa:1710:U:OP1	70:su:135:GLN:NE2	2.51	0.40
54:sd:156:LEU:HD12	54:sd:157:ARG:H	1.87	0.40
56:sf:174:ASP:H	56:sf:177:ASN:ND2	2.19	0.40
59:si:125:LYS:HE2	59:si:125:LYS:HB3	1.92	0.40
60:sj:57:ALA:HB1	60:sj:60:LEU:HD11	2.02	0.40
66:sq:87:ASN:OD1	66:sq:88:MET:HE2	2.21	0.40
69:st:74:GLN:O	69:st:78:ARG:N	2.49	0.40
70:su:107:ARG:NE	70:su:110:GLU:OE2	2.54	0.40
1:lA:336:A:H2	17:lQ:93:LYS:HD3	1.86	0.40
1:lA:732:U:O2'	1:lA:733:C:H6	2.04	0.40
1:lA:798:U:H2'	1:lA:799:A:C8	2.56	0.40
1:lA:1047:U:H2'	1:lA:1048:A:C8	2.56	0.40
1:lA:1542:U:H2'	1:lA:1543:U:H6	1.85	0.40
1:lA:1558:C:H2'	1:lA:1559:C:H6	1.86	0.40
1:lA:1684:A:C6	1:lA:1685:U:C4	3.10	0.40
6:lF:98:LYS:HE2	6:lF:98:LYS:HB2	1.85	0.40
9:II:86:LYS:HE3	9:II:117:LEU:HD13	2.02	0.40
12:IL:150:LYS:HE3	12:IL:150:LYS:HB3	1.82	0.40
14:IN:49:ILE:O	14:IN:232:ASP:HA	2.22	0.40
17:lQ:14:LYS:HA	17:lQ:19:LEU:HD12	2.03	0.40
18:IR:57:ILE:HG12	18:IR:73:CYS:SG	2.61	0.40
23:IW:47:VAL:HA	23:IW:68:VAL:HG11	2.02	0.40
24:IX:25:THR:OG1	24:IX:38:THR:HG22	2.20	0.40
45:sB:45:VAL:HG21	45:sB:64:LEU:HG	2.03	0.40
45:sB:97:PRO:HA	45:sB:98:PRO:HD3	2.01	0.40
52:sa:254:C:H2'	52:sa:255:G:O4'	2.21	0.40
52:sa:594:U:H2'	52:sa:595:A:H8	1.86	0.40
52:sa:853:U:H2'	52:sa:854:U:C6	2.57	0.40
52:sa:1259:A:O2'	52:sa:1260:C:H5'	2.21	0.40
52:sa:1758:G:H2'	52:sa:1759:A:C8	2.56	0.40
53:sc:55:ILE:HG21	53:sc:75:LEU:HD11	2.03	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:sd:121:TYR:O	54:sd:125:GLU:HG3	2.21	0.40
55:se:193:ASN:O	55:se:196:PRO:HD2	2.22	0.40
56:sf:189:ARG:HG2	56:sf:243:LYS:HB2	2.03	0.40
57:sg:152:ALA:O	57:sg:156:MET:HG2	2.21	0.40
61:sk:25:ASP:OD1	61:sk:25:ASP:N	2.53	0.40
62:sl:45:MET:HA	62:sl:48:LEU:HD13	2.04	0.40
1:lA:549:U:HO2'	1:lA:550:A:H8	1.67	0.40
1:lA:835:A:H2'	1:lA:836:C:O4'	2.20	0.40
1:lA:1242:G:O6	1:lA:1267:G:N2	2.55	0.40
1:lA:2188:U:H3	1:lA:3216:A:H4'	1.86	0.40
2:lB:73:A:H4'	2:lB:74:A:H5'	2.03	0.40
4:lD:89:TYR:N	4:lD:100:ASN:OD1	2.40	0.40
7:lG:5:LYS:HA	7:lG:5:LYS:HD3	1.65	0.40
9:II:77:ILE:HG12	9:II:123:THR:HG22	2.03	0.40
14:lN:53:LYS:HB2	14:lN:69:LEU:HB3	2.04	0.40
20:IT:10:CYS:HG	20:IT:15:HIS:CG	2.38	0.40
30:ld:36:ASN:HB3	30:ld:39:VAL:HB	2.03	0.40
34:lh:93:PHE:O	34:lh:97:GLU:HG2	2.22	0.40
37:lk:48:GLU:OE2	37:lk:80:LYS:NZ	2.27	0.40
52:sa:291:A:H2'	52:sa:292:A:C8	2.57	0.40
52:sa:426:C:H2'	52:sa:427:G:C8	2.56	0.40
52:sa:798:A:C8	67:sr:107:SER:HA	2.56	0.40
70:su:85:LEU:HD13	70:su:97:SER:HB3	2.02	0.40
1:lA:247:A:C5	1:lA:248:U:C5	3.09	0.40
1:lA:789:C:OP1	33:lg:28:ARG:N	2.54	0.40
1:lA:1070:A:OP2	1:lA:1489:G:N2	2.37	0.40
1:lA:2756:U:OP1	13:lM:147:LYS:HE3	2.22	0.40
1:lA:3132:C:H2'	1:lA:3133:G:O4'	2.21	0.40
3:lC:28:U:H4'	13:lM:9:MET:HE1	2.02	0.40
5:lE:37:ALA:HA	5:lE:187:GLY:O	2.22	0.40
6:lF:395:ASN:N	6:lF:395:ASN:OD1	2.54	0.40
7:lG:207:LEU:HD12	7:lG:207:LEU:HA	1.85	0.40
13:lM:137:ARG:O	13:lM:137:ARG:HG2	2.22	0.40
14:lN:163:ILE:O	14:lN:167:LYS:HB2	2.21	0.40
24:IX:100:ASP:OD1	26:lZ:26:LEU:HD11	2.21	0.40
27:la:104:ILE:HG21	27:la:107:LEU:HD23	2.04	0.40
27:la:164:LYS:HD2	27:la:164:LYS:HA	1.80	0.40
30:ld:7:ARG:HG2	30:ld:8:SER:N	2.35	0.40
52:sa:427:G:H2'	52:sa:428:C:C6	2.57	0.40
52:sa:1923:A:C6	52:sa:1933:G:O6	2.73	0.40
55:se:162:GLN:O	55:se:166:ILE:HG12	2.22	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:sg:193:LYS:HE2	57:sg:193:LYS:HB3	1.96	0.40
68:ss:43:MET:HE1	71:sv:2:HIS:HA	2.02	0.40

There are no symmetry-related clashes.

5.3 Torsion angles [i](#)

5.3.1 Protein backbone [i](#)

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
4	ID	244/257 (95%)	226 (93%)	18 (7%)	0	100	100
5	IE	385/402 (96%)	362 (94%)	23 (6%)	0	100	100
6	IF	421/431 (98%)	392 (93%)	28 (7%)	1 (0%)	43	73
7	IG	274/286 (96%)	255 (93%)	19 (7%)	0	100	100
8	IH	201/204 (98%)	185 (92%)	14 (7%)	2 (1%)	12	41
9	II	208/230 (90%)	199 (96%)	8 (4%)	1 (0%)	24	57
10	IJ	197/286 (69%)	188 (95%)	9 (5%)	0	100	100
11	IK	191/197 (97%)	184 (96%)	6 (3%)	1 (0%)	24	57
12	IL	196/210 (93%)	184 (94%)	12 (6%)	0	100	100
13	IM	166/174 (95%)	162 (98%)	3 (2%)	1 (1%)	21	52
14	IN	261/291 (90%)	247 (95%)	14 (5%)	0	100	100
15	IO	202/205 (98%)	192 (95%)	10 (5%)	0	100	100
16	IP	128/135 (95%)	124 (97%)	4 (3%)	0	100	100
17	IQ	202/205 (98%)	194 (96%)	8 (4%)	0	100	100
18	IR	156/179 (87%)	148 (95%)	8 (5%)	0	100	100
19	IS	165/168 (98%)	151 (92%)	14 (8%)	0	100	100
20	IT	171/173 (99%)	162 (95%)	9 (5%)	0	100	100
21	IU	148/198 (75%)	145 (98%)	3 (2%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
22	IV	163/166 (98%)	157 (96%)	6 (4%)	0	100	100
23	IW	91/137 (66%)	83 (91%)	8 (9%)	0	100	100
24	IX	131/140 (94%)	125 (95%)	6 (5%)	0	100	100
25	IY	114/121 (94%)	112 (98%)	2 (2%)	0	100	100
26	IZ	55/163 (34%)	54 (98%)	1 (2%)	0	100	100
27	la	208/213 (98%)	197 (95%)	11 (5%)	0	100	100
28	lb	135/139 (97%)	132 (98%)	3 (2%)	0	100	100
29	lc	146/149 (98%)	134 (92%)	12 (8%)	0	100	100
30	ld	58/64 (91%)	54 (93%)	4 (7%)	0	100	100
31	le	91/109 (84%)	81 (89%)	10 (11%)	0	100	100
32	lf	122/150 (81%)	115 (94%)	7 (6%)	0	100	100
33	lg	121/134 (90%)	118 (98%)	3 (2%)	0	100	100
34	lh	100/137 (73%)	98 (98%)	2 (2%)	0	100	100
35	li	120/122 (98%)	118 (98%)	1 (1%)	1 (1%)	16	47
36	lj	104/108 (96%)	98 (94%)	6 (6%)	0	100	100
37	lk	83/104 (80%)	81 (98%)	2 (2%)	0	100	100
38	ll	70/77 (91%)	63 (90%)	5 (7%)	2 (3%)	3	19
39	lm	88/93 (95%)	82 (93%)	6 (7%)	0	100	100
40	ln	71/77 (92%)	68 (96%)	3 (4%)	0	100	100
41	lo	48/51 (94%)	45 (94%)	3 (6%)	0	100	100
42	lp	51/56 (91%)	48 (94%)	3 (6%)	0	100	100
43	lq	90/98 (92%)	89 (99%)	1 (1%)	0	100	100
44	sA	8/137 (6%)	6 (75%)	2 (25%)	0	100	100
45	sB	96/144 (67%)	93 (97%)	3 (3%)	0	100	100
46	sC	30/84 (36%)	30 (100%)	0	0	100	100
47	sD	58/69 (84%)	52 (90%)	6 (10%)	0	100	100
48	sE	53/56 (95%)	50 (94%)	3 (6%)	0	100	100
53	sc	194/255 (76%)	184 (95%)	10 (5%)	0	100	100
54	sd	184/244 (75%)	174 (95%)	10 (5%)	0	100	100
55	se	126/256 (49%)	121 (96%)	4 (3%)	1 (1%)	16	47
56	sf	254/326 (78%)	231 (91%)	23 (9%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
57	sg	175/206 (85%)	166 (95%)	9 (5%)	0	100	100
58	sh	49/266 (18%)	48 (98%)	1 (2%)	0	100	100
59	si	53/201 (26%)	53 (100%)	0	0	100	100
60	sj	125/237 (53%)	120 (96%)	5 (4%)	0	100	100
61	sk	80/185 (43%)	79 (99%)	1 (1%)	0	100	100
62	sl	64/127 (50%)	63 (98%)	1 (2%)	0	100	100
63	sm	137/156 (88%)	127 (93%)	10 (7%)	0	100	100
64	so	68/151 (45%)	66 (97%)	2 (3%)	0	100	100
65	sp	126/146 (86%)	119 (94%)	7 (6%)	0	100	100
66	sq	101/144 (70%)	92 (91%)	8 (8%)	1 (1%)	12	41
67	sr	127/130 (98%)	117 (92%)	10 (8%)	0	100	100
68	ss	135/158 (85%)	129 (96%)	6 (4%)	0	100	100
69	st	79/117 (68%)	79 (100%)	0	0	100	100
70	su	120/155 (77%)	108 (90%)	12 (10%)	0	100	100
71	sv	123/155 (79%)	116 (94%)	7 (6%)	0	100	100
72	sw	42/118 (36%)	42 (100%)	0	0	100	100
73	sx	59/86 (69%)	57 (97%)	2 (3%)	0	100	100
74	sy	104/141 (74%)	97 (93%)	7 (7%)	0	100	100
All	All	8946/11289 (79%)	8471 (95%)	464 (5%)	11 (0%)	49	78

All (11) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
38	ll	40	PRO
55	se	193	ASN
6	lF	390	VAL
8	lH	119	VAL
13	lM	28	ASP
66	sq	112	GLY
35	li	-6	LYS
38	ll	39	TYR
11	lK	134	ILE
8	lH	118	ASP
9	lI	179	VAL

5.3.2 Protein sidechains ⓘ

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
4	ID	195/201 (97%)	191 (98%)	4 (2%)	47	71
5	IE	330/343 (96%)	325 (98%)	5 (2%)	57	75
6	IF	339/345 (98%)	331 (98%)	8 (2%)	43	69
7	IG	223/231 (96%)	217 (97%)	6 (3%)	39	67
8	IH	172/173 (99%)	168 (98%)	4 (2%)	44	70
9	II	178/195 (91%)	176 (99%)	2 (1%)	65	78
10	IJ	177/242 (73%)	175 (99%)	2 (1%)	65	78
11	IK	171/174 (98%)	170 (99%)	1 (1%)	78	83
12	IL	169/176 (96%)	166 (98%)	3 (2%)	51	73
13	IM	144/147 (98%)	135 (94%)	9 (6%)	16	45
14	IN	224/243 (92%)	221 (99%)	3 (1%)	61	77
15	IO	167/168 (99%)	166 (99%)	1 (1%)	78	83
16	IP	114/118 (97%)	113 (99%)	1 (1%)	70	80
17	IQ	171/172 (99%)	171 (100%)	0	100	100
18	IR	129/147 (88%)	126 (98%)	3 (2%)	44	70
19	IS	142/143 (99%)	138 (97%)	4 (3%)	38	66
20	IT	156/156 (100%)	156 (100%)	0	100	100
21	IU	132/174 (76%)	130 (98%)	2 (2%)	57	75
22	IV	144/145 (99%)	141 (98%)	3 (2%)	47	71
23	IW	86/125 (69%)	83 (96%)	3 (4%)	32	62
24	IX	109/113 (96%)	105 (96%)	4 (4%)	30	61
25	IY	99/102 (97%)	98 (99%)	1 (1%)	68	79
26	IZ	52/137 (38%)	51 (98%)	1 (2%)	50	73
27	la	177/179 (99%)	173 (98%)	4 (2%)	44	70
28	lb	121/123 (98%)	119 (98%)	2 (2%)	53	74
29	lc	120/121 (99%)	116 (97%)	4 (3%)	33	63

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
30	ld	50/54 (93%)	50 (100%)	0	100	100
31	le	78/92 (85%)	75 (96%)	3 (4%)	29	60
32	lf	110/128 (86%)	107 (97%)	3 (3%)	39	67
33	lg	107/116 (92%)	106 (99%)	1 (1%)	70	80
34	lh	86/116 (74%)	85 (99%)	1 (1%)	63	78
35	li	103/103 (100%)	100 (97%)	3 (3%)	37	66
36	lj	89/91 (98%)	88 (99%)	1 (1%)	65	78
37	lk	71/82 (87%)	70 (99%)	1 (1%)	59	76
38	ll	60/64 (94%)	60 (100%)	0	100	100
39	lm	73/75 (97%)	70 (96%)	3 (4%)	27	59
40	ln	63/66 (96%)	60 (95%)	3 (5%)	23	54
41	lo	44/45 (98%)	42 (96%)	2 (4%)	24	56
42	lp	45/48 (94%)	45 (100%)	0	100	100
43	lq	85/91 (93%)	84 (99%)	1 (1%)	63	78
44	sA	10/112 (9%)	9 (90%)	1 (10%)	7	29
45	sB	87/127 (68%)	83 (95%)	4 (5%)	24	55
46	sC	28/73 (38%)	25 (89%)	3 (11%)	6	25
47	sD	50/59 (85%)	46 (92%)	4 (8%)	11	37
48	sE	45/46 (98%)	43 (96%)	2 (4%)	25	56
53	sc	157/199 (79%)	153 (98%)	4 (2%)	42	69
54	sd	156/206 (76%)	151 (97%)	5 (3%)	34	64
55	se	130/225 (58%)	127 (98%)	3 (2%)	44	70
56	sf	223/283 (79%)	220 (99%)	3 (1%)	61	77
57	sg	160/178 (90%)	160 (100%)	0	100	100
58	sh	42/220 (19%)	39 (93%)	3 (7%)	13	41
59	si	48/167 (29%)	46 (96%)	2 (4%)	26	58
60	sj	108/205 (53%)	106 (98%)	2 (2%)	50	73
61	sk	72/164 (44%)	72 (100%)	0	100	100
62	sl	60/111 (54%)	60 (100%)	0	100	100
63	sm	126/138 (91%)	124 (98%)	2 (2%)	55	75
64	so	68/129 (53%)	67 (98%)	1 (2%)	57	75

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
65	sp	99/114 (87%)	96 (97%)	3 (3%)	36	65
66	sq	94/127 (74%)	94 (100%)	0	100	100
67	sr	112/113 (99%)	109 (97%)	3 (3%)	39	67
68	ss	111/128 (87%)	108 (97%)	3 (3%)	39	67
69	st	73/106 (69%)	72 (99%)	1 (1%)	59	76
70	su	104/130 (80%)	101 (97%)	3 (3%)	37	66
71	sv	106/132 (80%)	104 (98%)	2 (2%)	50	73
72	sw	44/107 (41%)	44 (100%)	0	100	100
73	sx	56/77 (73%)	53 (95%)	3 (5%)	20	50
74	sy	86/114 (75%)	85 (99%)	1 (1%)	63	78
All	All	7760/9554 (81%)	7600 (98%)	160 (2%)	46	71

All (160) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
4	ID	5	THR
4	ID	33	ASP
4	ID	40	THR
4	ID	193	ARG
5	IE	166	THR
5	IE	213	THR
5	IE	231	VAL
5	IE	232	ILE
5	IE	374	THR
6	IF	79	ILE
6	IF	94	ASN
6	IF	115	THR
6	IF	153	VAL
6	IF	190	VAL
6	IF	215	ASP
6	IF	342	GLN
6	IF	386	LYS
7	IG	9	ASN
7	IG	50	ARG
7	IG	94	ASN
7	IG	101	THR
7	IG	167	MET
7	IG	204	MET

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Mol	Chain	Res	Type
8	1H	8	VAL
8	1H	146	VAL
8	1H	155	TYR
8	1H	161	LEU
9	1I	52	THR
9	1I	77	ILE
10	1J	25	PHE
10	1J	36	LEU
11	1K	71	ASN
12	1L	50	VAL
12	1L	119	TRP
12	1L	201	THR
13	1M	14	ILE
13	1M	19	LEU
13	1M	31	THR
13	1M	36	VAL
13	1M	48	SER
13	1M	110	ILE
13	1M	111	ASP
13	1M	148	ILE
13	1M	155	THR
14	1N	61	GLN
14	1N	155	PHE
14	1N	177	TYR
15	1O	23	VAL
16	1P	68	THR
18	1R	43	ARG
18	1R	120	GLN
18	1R	139	PHE
19	1S	10	VAL
19	1S	60	ILE
19	1S	80	CYS
19	1S	152	VAL
21	1U	89	THR
21	1U	149	LYS
22	1V	89	LEU
22	1V	91	ILE
22	1V	129	LEU
23	1W	39	ASP
23	1W	54	ILE
23	1W	100	ASN
24	1X	17	THR

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Mol	Chain	Res	Type
24	lX	59	ASP
24	lX	123	LYS
24	lX	128	LEU
25	lY	32	ASN
26	lZ	59	GLN
27	la	22	ASN
27	la	69	THR
27	la	109	LEU
27	la	158	VAL
28	lb	63	GLU
28	lb	124	GLU
29	lc	14	HIS
29	lc	97	GLU
29	lc	117	PHE
29	lc	141	VAL
31	le	27	TYR
31	le	40	ASP
31	le	51	ASN
32	lf	91	MET
32	lf	118	ILE
32	lf	141	VAL
33	lg	29	VAL
34	lh	8	VAL
35	li	12	GLU
35	li	47	LEU
35	li	85	THR
36	lj	79	HIS
37	lk	11	ARG
39	lm	8	VAL
39	lm	26	ILE
39	lm	42	CYS
40	ln	25	VAL
40	ln	44	THR
40	ln	67	VAL
41	lo	2	THR
41	lo	43	HIS
43	lq	33	LEU
44	sA	109	TYR
45	sB	30	VAL
45	sB	33	ASP
45	sB	45	VAL
45	sB	64	LEU

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Mol	Chain	Res	Type
46	sC	9	LEU
46	sC	30	ASN
46	sC	32	TYR
47	sD	30	GLN
47	sD	57	THR
47	sD	59	LEU
47	sD	68	LEU
48	sE	20	THR
48	sE	55	LEU
53	sc	106	ILE
53	sc	158	CYS
53	sc	160	LEU
53	sc	216	THR
54	sd	39	LEU
54	sd	123	ILE
54	sd	156	LEU
54	sd	178	THR
54	sd	189	VAL
55	se	26	PHE
55	se	112	LEU
55	se	200	VAL
56	sf	110	VAL
56	sf	171	VAL
56	sf	214	ASN
58	sh	70	ASN
58	sh	78	ASP
58	sh	97	VAL
59	si	121	SER
59	si	160	THR
60	sj	58	LEU
60	sj	96	LEU
63	sm	13	VAL
63	sm	61	SER
64	so	87	ASP
65	sp	39	VAL
65	sp	81	LYS
65	sp	126	ASP
67	sr	55	THR
67	sr	65	LEU
67	sr	98	HIS
68	ss	55	ASN
68	ss	124	PHE

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Mol	Chain	Res	Type
68	ss	135	ASP
69	st	69	ILE
70	su	77	ASN
70	su	80	ILE
70	su	137	THR
71	sv	42	MET
71	sv	43	CYS
73	sx	17	VAL
73	sx	21	LEU
73	sx	61	ILE
74	sy	31	VAL

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (67) such sidechains are listed below:

Mol	Chain	Res	Type
4	lD	29	HIS
4	lD	140	ASN
5	lE	175	GLN
6	lF	50	GLN
6	lF	87	ASN
6	lF	149	GLN
6	lF	241	ASN
6	lF	322	ASN
6	lF	394	ASN
7	lG	118	GLN
7	lG	257	HIS
7	lG	270	GLN
8	lH	5	GLN
9	lI	187	ASN
10	lJ	77	ASN
10	lJ	171	GLN
11	lK	10	HIS
13	lM	119	GLN
13	lM	161	ASN
14	lN	57	HIS
14	lN	200	GLN
15	lO	43	GLN
17	lQ	32	HIS
17	lQ	156	GLN
18	lR	25	HIS
18	lR	63	HIS
18	lR	120	GLN

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Mol	Chain	Res	Type
19	lS	13	HIS
19	lS	21	ASN
19	lS	96	ASN
21	lU	58	HIS
22	lV	45	ASN
24	lX	12	ASN
25	lY	18	ASN
29	lc	7	GLN
29	lc	39	HIS
29	lc	40	HIS
30	ld	10	HIS
30	ld	14	HIS
33	lg	66	ASN
33	lg	99	HIS
34	lh	36	GLN
41	lo	4	HIS
45	sB	94	ASN
47	sD	30	GLN
47	sD	41	GLN
47	sD	45	ASN
48	sE	8	ASN
53	sc	112	HIS
54	sd	18	ASN
54	sd	61	GLN
54	sd	176	ASN
54	sd	180	GLN
56	sf	34	HIS
56	sf	51	ASN
56	sf	140	HIS
56	sf	248	GLN
61	sk	33	GLN
61	sk	82	ASN
62	sl	17	ASN
62	sl	31	HIS
63	sm	91	HIS
67	sr	120	HIS
70	su	102	ASN
71	sv	69	ASN
74	sy	59	GLN
74	sy	75	ASN

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	lA	3127/3503 (89%)	593 (18%)	0
2	lB	143/155 (92%)	31 (21%)	0
3	lC	116/117 (99%)	18 (15%)	0
49	sI	70/76 (92%)	30 (42%)	0
50	sJ	71/77 (92%)	30 (42%)	0
51	sK	9/10 (90%)	0	0
52	sa	1383/1947 (71%)	305 (22%)	0
All	All	4919/5885 (83%)	1007 (20%)	0

All (1007) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	lA	18	G
1	lA	22	A
1	lA	29	U
1	lA	30	A
1	lA	36	A
1	lA	39	A
1	lA	45	A
1	lA	53	A
1	lA	54	G
1	lA	55	G
1	lA	56	A
1	lA	62	A
1	lA	65	U
1	lA	70	A
1	lA	73	A
1	lA	88	G
1	lA	105	A
1	lA	106	G
1	lA	107	A
1	lA	108	C
1	lA	115	G
1	lA	116	A
1	lA	117	U
1	lA	185	G
1	lA	186	A
1	lA	189	G
1	lA	198	A
1	lA	199	G
1	lA	200	A
1	lA	206	U
1	lA	213	A

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Mol	Chain	Res	Type
1	lA	214	A
1	lA	221	U
1	lA	233	A
1	lA	237	C
1	lA	243	G
1	lA	246	U
1	lA	256	G
1	lA	257	A
1	lA	259	A
1	lA	264	G
1	lA	265	A
1	lA	266	A
1	lA	267	A
1	lA	277	A
1	lA	285	A
1	lA	286	U
1	lA	287	A
1	lA	289	C
1	lA	293	A
1	lA	294	A
1	lA	302	U
1	lA	303	G
1	lA	311	A
1	lA	316	G
1	lA	332	G
1	lA	342	A
1	lA	344	A
1	lA	345	A
1	lA	346	G
1	lA	350	A
1	lA	369	A
1	lA	370	A
1	lA	376	A
1	lA	385	C
1	lA	396	G
1	lA	397	A
1	lA	416	U
1	lA	422	G
1	lA	423	A
1	lA	424	A
1	lA	425	C
1	lA	444	A

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Mol	Chain	Res	Type
1	lA	447	G
1	lA	448	C
1	lA	449	U
1	lA	451	G
1	lA	467	A
1	lA	474	A
1	lA	483	G
1	lA	494	G
1	lA	495	A
1	lA	496	A
1	lA	500	G
1	lA	501	U
1	lA	510	A
1	lA	511	A
1	lA	512	G
1	lA	517	G
1	lA	535	A
1	lA	536	G
1	lA	538	A
1	lA	539	G
1	lA	550	A
1	lA	562	A
1	lA	564	A
1	lA	565	U
1	lA	577	G
1	lA	583	A
1	lA	586	A
1	lA	588	G
1	lA	589	A
1	lA	610	A
1	lA	612	A
1	lA	613	G
1	lA	614	U
1	lA	615	G
1	lA	619	A
1	lA	620	U
1	lA	625	A
1	lA	626	A
1	lA	627	A
1	lA	630	A
1	lA	631	G
1	lA	632	G

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Mol	Chain	Res	Type
1	lA	637	U
1	lA	644	A
1	lA	645	A
1	lA	646	U
1	lA	647	C
1	lA	648	U
1	lA	649	U
1	lA	652	U
1	lA	658	A
1	lA	663	G
1	lA	675	A
1	lA	685	A
1	lA	696	G
1	lA	699	A
1	lA	701	A
1	lA	702	A
1	lA	703	G
1	lA	708	U
1	lA	709	U
1	lA	710	A
1	lA	714	U
1	lA	715	A
1	lA	720	U
1	lA	721	A
1	lA	722	A
1	lA	724	A
1	lA	725	A
1	lA	726	A
1	lA	727	U
1	lA	728	U
1	lA	733	C
1	lA	734	C
1	lA	740	A
1	lA	744	A
1	lA	745	A
1	lA	746	A
1	lA	747	C
1	lA	755	G
1	lA	770	C
1	lA	783	A
1	lA	794	A
1	lA	811	A

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Mol	Chain	Res	Type
1	lA	816	G
1	lA	817	A
1	lA	818	A
1	lA	825	G
1	lA	833	U
1	lA	834	A
1	lA	835	A
1	lA	841	A
1	lA	842	G
1	lA	849	C
1	lA	851	A
1	lA	853	A
1	lA	854	A
1	lA	855	A
1	lA	862	G
1	lA	871	A
1	lA	872	A
1	lA	873	U
1	lA	874	A
1	lA	881	G
1	lA	882	C
1	lA	889	A
1	lA	898	U
1	lA	903	G
1	lA	904	A
1	lA	905	A
1	lA	911	U
1	lA	915	U
1	lA	918	G
1	lA	925	A
1	lA	927	A
1	lA	934	G
1	lA	936	A
1	lA	960	G
1	lA	964	G
1	lA	968	C
1	lA	976	G
1	lA	979	G
1	lA	980	A
1	lA	993	U
1	lA	998	U
1	lA	999	G

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Mol	Chain	Res	Type
1	lA	1013	A
1	lA	1026	G
1	lA	1027	G
1	lA	1028	G
1	lA	1033	A
1	lA	1035	G
1	lA	1036	A
1	lA	1043	G
1	lA	1056	G
1	lA	1063	C
1	lA	1073	A
1	lA	1078	C
1	lA	1092	A
1	lA	1096	U
1	lA	1097	U
1	lA	1098	A
1	lA	1099	A
1	lA	1112	G
1	lA	1119	U
1	lA	1120	A
1	lA	1127	A
1	lA	1166	A
1	lA	1168	C
1	lA	1182	A
1	lA	1183	G
1	lA	1184	A
1	lA	1197	G
1	lA	1210	C
1	lA	1211	A
1	lA	1219	A
1	lA	1220	A
1	lA	1221	A
1	lA	1222	U
1	lA	1242	G
1	lA	1248	U
1	lA	1256	G
1	lA	1268	A
1	lA	1275	A
1	lA	1280	C
1	lA	1284	A
1	lA	1292	A
1	lA	1306	A

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Mol	Chain	Res	Type
1	lA	1312	A
1	lA	1314	A
1	lA	1315	A
1	lA	1316	C
1	lA	1325	C
1	lA	1343	A
1	lA	1344	U
1	lA	1345	U
1	lA	1409	G
1	lA	1410	A
1	lA	1411	A
1	lA	1417	G
1	lA	1431	G
1	lA	1433	U
1	lA	1440	A
1	lA	1441	U
1	lA	1450	A
1	lA	1455	A
1	lA	1475	A
1	lA	1476	G
1	lA	1477	A
1	lA	1478	C
1	lA	1480	U
1	lA	1486	C
1	lA	1495	A
1	lA	1499	G
1	lA	1502	A
1	lA	1503	U
1	lA	1505	G
1	lA	1516	A
1	lA	1517	C
1	lA	1519	A
1	lA	1521	A
1	lA	1534	U
1	lA	1553	A
1	lA	1568	G
1	lA	1569	A
1	lA	1570	U
1	lA	1571	C
1	lA	1580	A
1	lA	1584	G
1	lA	1589	U

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Mol	Chain	Res	Type
1	lA	1603	A
1	lA	1604	A
1	lA	1606	A
1	lA	1622	A
1	lA	1625	A
1	lA	1649	U
1	lA	1652	U
1	lA	1664	G
1	lA	1674	U
1	lA	1677	A
1	lA	1690	A
1	lA	1698	G
1	lA	1729	G
1	lA	1735	A
1	lA	1737	A
1	lA	1738	G
1	lA	1744	A
1	lA	1753	A
1	lA	1754	G
1	lA	1818	C
1	lA	1821	A
1	lA	1822	U
1	lA	1825	U
1	lA	1838	G
1	lA	1864	C
1	lA	1865	U
1	lA	1897	A
1	lA	1898	A
1	lA	1907	A
1	lA	1913	G
1	lA	1926	U
1	lA	1928	G
1	lA	1936	A
1	lA	1937	A
1	lA	1938	G
1	lA	1958	A
1	lA	1959	U
1	lA	1961	G
1	lA	1963	G
1	lA	1965	U
1	lA	1975	G
1	lA	1983	A

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Mol	Chain	Res	Type
1	lA	1984	A
1	lA	2040	A
1	lA	2043	A
1	lA	2044	C
1	lA	2049	A
1	lA	2050	C
1	lA	2051	A
1	lA	2067	A
1	lA	2079	A
1	lA	2080	A
1	lA	2081	A
1	lA	2082	U
1	lA	2095	G
1	lA	2108	G
1	lA	2132	A
1	lA	2133	G
1	lA	2178	A
1	lA	2179	U
1	lA	2190	A
1	lA	2193	G
1	lA	2197	G
1	lA	2198	G
1	lA	2207	A
1	lA	2215	A
1	lA	2216	U
1	lA	2218	A
1	lA	2234	A
1	lA	2236	A
1	lA	2245	A
1	lA	2246	G
1	lA	2263	G
1	lA	2264	A
1	lA	2269	U
1	lA	2274	A
1	lA	2281	U
1	lA	2285	U
1	lA	2286	G
1	lA	2301	U
1	lA	2320	A
1	lA	2328	A
1	lA	2329	G
1	lA	2330	U

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Mol	Chain	Res	Type
1	lA	2331	A
1	lA	2332	A
1	lA	2333	C
1	lA	2338	A
1	lA	2341	C
1	lA	2343	C
1	lA	2345	U
1	lA	2346	A
1	lA	2349	G
1	lA	2355	A
1	lA	2366	C
1	lA	2367	A
1	lA	2374	U
1	lA	2382	C
1	lA	2383	G
1	lA	2384	C
1	lA	2389	A
1	lA	2390	U
1	lA	2391	G
1	lA	2394	U
1	lA	2395	C
1	lA	2396	A
1	lA	2411	G
1	lA	2412	U
1	lA	2432	A
1	lA	2439	A
1	lA	2441	U
1	lA	2449	A
1	lA	2450	C
1	lA	2451	G
1	lA	2459	U
1	lA	2461	G
1	lA	2462	G
1	lA	2469	G
1	lA	2470	G
1	lA	2473	A
1	lA	2478	A
1	lA	2479	G
1	lA	2480	A
1	lA	2487	U
1	lA	2494	G
1	lA	2509	U

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Mol	Chain	Res	Type
1	lA	2513	A
1	lA	2514	A
1	lA	2589	C
1	lA	2594	C
1	lA	2595	G
1	lA	2597	U
1	lA	2598	A
1	lA	2601	U
1	lA	2629	A
1	lA	2637	A
1	lA	2651	A
1	lA	2655	G
1	lA	2656	G
1	lA	2657	C
1	lA	2663	A
1	lA	2676	G
1	lA	2677	G
1	lA	2684	G
1	lA	2689	G
1	lA	2696	A
1	lA	2705	A
1	lA	2706	C
1	lA	2707	A
1	lA	2722	U
1	lA	2726	A
1	lA	2727	A
1	lA	2742	G
1	lA	2744	A
1	lA	2747	G
1	lA	2759	A
1	lA	2760	G
1	lA	2761	A
1	lA	2764	A
1	lA	2766	A
1	lA	2773	U
1	lA	2774	A
1	lA	2784	G
1	lA	2795	C
1	lA	2796	C
1	lA	2797	A
1	lA	2798	G
1	lA	2799	U

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Mol	Chain	Res	Type
1	lA	2819	U
1	lA	2822	U
1	lA	2823	G
1	lA	2825	C
1	lA	2832	A
1	lA	2842	U
1	lA	2843	U
1	lA	2855	A
1	lA	2858	A
1	lA	2859	U
1	lA	2860	U
1	lA	2864	A
1	lA	2865	A
1	lA	2866	C
1	lA	2867	A
1	lA	2868	U
1	lA	2869	A
1	lA	2876	U
1	lA	2877	U
1	lA	2882	G
1	lA	2889	G
1	lA	2890	A
1	lA	2900	G
1	lA	2902	G
1	lA	2909	A
1	lA	2920	U
1	lA	2925	A
1	lA	2926	U
1	lA	2939	G
1	lA	2943	G
1	lA	2944	A
1	lA	2946	A
1	lA	2953	C
1	lA	2959	G
1	lA	2960	A
1	lA	2977	G
1	lA	2981	A
1	lA	2985	C
1	lA	2987	C
1	lA	2988	A
1	lA	2993	G
1	lA	3006	G

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Mol	Chain	Res	Type
1	lA	3009	C
1	lA	3010	C
1	lA	3014	G
1	lA	3015	A
1	lA	3018	U
1	lA	3030	A
1	lA	3032	C
1	lA	3041	G
1	lA	3066	U
1	lA	3070	C
1	lA	3073	U
1	lA	3078	U
1	lA	3079	A
1	lA	3084	A
1	lA	3085	C
1	lA	3090	G
1	lA	3114	A
1	lA	3126	C
1	lA	3133	G
1	lA	3141	A
1	lA	3142	G
1	lA	3143	A
1	lA	3144	U
1	lA	3151	A
1	lA	3166	G
1	lA	3175	A
1	lA	3200	G
1	lA	3203	A
1	lA	3210	U
1	lA	3212	U
1	lA	3213	U
1	lA	3223	G
1	lA	3232	G
1	lA	3240	A
1	lA	3246	A
1	lA	3247	C
1	lA	3254	U
1	lA	3259	U
1	lA	3264	G
1	lA	3270	A
1	lA	3277	C
1	lA	3278	A

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Mol	Chain	Res	Type
1	lA	3284	G
1	lA	3285	A
1	lA	3286	A
1	lA	3287	U
1	lA	3299	G
1	lA	3314	U
1	lA	3327	A
1	lA	3330	A
1	lA	3332	A
1	lA	3334	U
1	lA	3335	U
1	lA	3336	G
1	lA	3337	A
1	lA	3338	A
1	lA	3344	U
1	lA	3345	A
1	lA	3351	U
1	lA	3353	U
1	lA	3359	U
1	lA	3360	U
1	lA	3361	A
1	lA	3383	A
1	lA	3384	A
1	lA	3385	G
1	lA	3398	U
1	lA	3399	G
1	lA	3405	A
1	lA	3406	C
1	lA	3409	A
1	lA	3410	A
1	lA	3413	U
1	lA	3414	A
1	lA	3419	A
1	lA	3430	U
1	lA	3431	U
1	lA	3433	U
1	lA	3451	A
1	lA	3454	G
1	lA	3462	A
1	lA	3466	U
1	lA	3468	U
1	lA	3469	A

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Mol	Chain	Res	Type
1	1A	3470	G
1	1A	3472	G
1	1A	3473	U
1	1A	3475	U
1	1A	3476	C
1	1A	3477	U
1	1A	3478	U
1	1A	3479	A
1	1A	3480	A
1	1A	3487	C
1	1A	3490	U
1	1A	3491	C
1	1A	3492	G
1	1A	3498	U
1	1A	3499	G
1	1A	3500	A
2	1B	5	A
2	1B	34	G
2	1B	36	G
2	1B	39	A
2	1B	41	G
2	1B	51	A
2	1B	53	G
2	1B	54	A
2	1B	61	A
2	1B	64	A
2	1B	65	G
2	1B	72	G
2	1B	73	A
2	1B	74	A
2	1B	77	A
2	1B	79	A
2	1B	80	A
2	1B	81	A
2	1B	82	A
2	1B	86	G
2	1B	92	C
2	1B	102	A
2	1B	103	A
2	1B	104	U
2	1B	106	C
2	1B	109	G

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Mol	Chain	Res	Type
2	lB	110	A
2	lB	113	G
2	lB	131	A
2	lB	147	C
2	lB	150	U
3	lC	6	G
3	lC	17	G
3	lC	19	U
3	lC	22	C
3	lC	32	A
3	lC	37	A
3	lC	49	C
3	lC	52	G
3	lC	53	C
3	lC	55	C
3	lC	62	U
3	lC	63	C
3	lC	72	C
3	lC	73	G
3	lC	90	G
3	lC	96	U
3	lC	99	U
3	lC	109	G
49	sI	2	G
49	sI	3	C
49	sI	6	C
49	sI	9	A
49	sI	10	G
49	sI	13	C
49	sI	14	A
49	sI	15	G
49	sI	16	G
49	sI	18	A
49	sI	22	G
49	sI	23	A
49	sI	29	C
49	sI	31	A
49	sI	41	G
49	sI	44	U
49	sI	46	G
49	sI	49	G
49	sI	50	C

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Mol	Chain	Res	Type
49	sI	53	G
49	sI	55	U
49	sI	56	C
49	sI	58	A
49	sI	68	G
49	sI	69	G
49	sI	70	G
49	sI	73	A
49	sI	74	C
49	sI	75	C
49	sI	76	A
50	sJ	6	C
50	sJ	8	U
50	sJ	12	G
50	sJ	14	A
50	sJ	15	G
50	sJ	16	C
50	sJ	17	C
50	sJ	18	C
50	sJ	19	G
50	sJ	20	U
50	sJ	21	U
50	sJ	22	A
50	sJ	23	G
50	sJ	24	C
50	sJ	45	A
50	sJ	47	G
50	sJ	48	U
50	sJ	49	C
50	sJ	50	U
50	sJ	52	C
50	sJ	53	G
50	sJ	60	G
50	sJ	61	U
50	sJ	63	C
50	sJ	65	G
50	sJ	66	C
50	sJ	68	G
50	sJ	69	A
50	sJ	76	C
50	sJ	77	A
52	sa	4	C

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Mol	Chain	Res	Type
52	sa	24	U
52	sa	26	U
52	sa	33	A
52	sa	41	G
52	sa	42	A
52	sa	44	U
52	sa	46	A
52	sa	56	G
52	sa	59	U
52	sa	61	A
52	sa	62	G
52	sa	64	A
52	sa	65	U
52	sa	103	U
52	sa	113	A
52	sa	114	G
52	sa	115	U
52	sa	123	G
52	sa	135	C
52	sa	136	A
52	sa	137	A
52	sa	139	G
52	sa	140	A
52	sa	141	U
52	sa	142	A
52	sa	146	U
52	sa	148	G
52	sa	150	G
52	sa	152	A
52	sa	153	U
52	sa	154	G
52	sa	160	G
52	sa	161	A
52	sa	163	A
52	sa	171	A
52	sa	173	A
52	sa	211	U
52	sa	213	A
52	sa	239	U
52	sa	243	U
52	sa	244	U
52	sa	246	G

Continued on next page...

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Mol	Chain	Res	Type
52	sa	249	U
52	sa	253	A
52	sa	257	C
52	sa	258	A
52	sa	259	A
52	sa	260	U
52	sa	262	G
52	sa	263	U
52	sa	285	G
52	sa	288	A
52	sa	289	C
52	sa	293	U
52	sa	294	G
52	sa	295	A
52	sa	296	G
52	sa	299	U
52	sa	310	U
52	sa	311	C
52	sa	313	A
52	sa	317	G
52	sa	332	G
52	sa	333	A
52	sa	341	G
52	sa	345	A
52	sa	347	A
52	sa	354	A
52	sa	355	A
52	sa	356	C
52	sa	385	G
52	sa	395	G
52	sa	396	A
52	sa	397	U
52	sa	399	G
52	sa	411	A
52	sa	413	G
52	sa	417	G
52	sa	419	C
52	sa	420	A
52	sa	421	G
52	sa	429	G
52	sa	432	A
52	sa	434	U

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Mol	Chain	Res	Type
52	sa	439	C
52	sa	440	A
52	sa	456	G
52	sa	463	A
52	sa	470	A
52	sa	545	G
52	sa	549	A
52	sa	551	G
52	sa	553	C
52	sa	555	G
52	sa	559	C
52	sa	561	A
52	sa	562	G
52	sa	568	G
52	sa	572	U
52	sa	576	U
52	sa	579	A
52	sa	584	C
52	sa	588	A
52	sa	589	G
52	sa	600	A
52	sa	602	U
52	sa	604	G
52	sa	613	A
52	sa	614	A
52	sa	616	A
52	sa	617	C
52	sa	618	G
52	sa	628	G
52	sa	742	G
52	sa	746	A
52	sa	752	A
52	sa	756	G
52	sa	757	U
52	sa	759	U
52	sa	762	A
52	sa	782	A
52	sa	783	A
52	sa	784	U
52	sa	785	A
52	sa	787	U
52	sa	790	A

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Mol	Chain	Res	Type
52	sa	800	A
52	sa	803	G
52	sa	804	C
52	sa	843	A
52	sa	844	A
52	sa	845	A
52	sa	856	G
52	sa	876	G
52	sa	877	G
52	sa	879	G
52	sa	894	G
52	sa	904	A
52	sa	913	A
52	sa	915	A
52	sa	922	G
52	sa	924	A
52	sa	925	U
52	sa	927	U
52	sa	939	G
52	sa	940	U
52	sa	946	A
52	sa	950	A
52	sa	968	A
52	sa	972	A
52	sa	973	A
52	sa	984	U
52	sa	985	A
52	sa	1005	A
52	sa	1008	C
52	sa	1009	G
52	sa	1012	G
52	sa	1019	A
52	sa	1091	C
52	sa	1106	A
52	sa	1118	G
52	sa	1122	U
52	sa	1123	U
52	sa	1135	G
52	sa	1140	G
52	sa	1164	A
52	sa	1169	A
52	sa	1176	G

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Mol	Chain	Res	Type
52	sa	1177	A
52	sa	1184	C
52	sa	1185	A
52	sa	1192	G
52	sa	1210	U
52	sa	1219	A
52	sa	1221	A
52	sa	1224	G
52	sa	1225	G
52	sa	1227	A
52	sa	1232	U
52	sa	1233	A
52	sa	1239	A
52	sa	1242	G
52	sa	1243	A
52	sa	1253	G
52	sa	1254	G
52	sa	1260	C
52	sa	1267	A
52	sa	1269	A
52	sa	1270	G
52	sa	1271	U
52	sa	1274	U
52	sa	1276	U
52	sa	1277	C
52	sa	1281	A
52	sa	1282	U
52	sa	1283	U
52	sa	1284	U
52	sa	1287	U
52	sa	1290	G
52	sa	1299	C
52	sa	1310	U
52	sa	1311	U
52	sa	1312	A
52	sa	1316	G
52	sa	1317	G
52	sa	1323	U
52	sa	1326	U
52	sa	1331	C
52	sa	1332	A
52	sa	1337	A

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Mol	Chain	Res	Type
52	sa	1339	U
52	sa	1344	G
52	sa	1346	A
52	sa	1365	U
52	sa	1369	A
52	sa	1379	C
52	sa	1395	A
52	sa	1396	A
52	sa	1428	U
52	sa	1438	A
52	sa	1439	C
52	sa	1440	A
52	sa	1442	A
52	sa	1520	C
52	sa	1522	U
52	sa	1534	A
52	sa	1543	A
52	sa	1551	A
52	sa	1552	G
52	sa	1553	A
52	sa	1565	G
52	sa	1566	C
52	sa	1567	A
52	sa	1578	A
52	sa	1580	U
52	sa	1581	G
52	sa	1585	U
52	sa	1588	C
52	sa	1667	U
52	sa	1668	G
52	sa	1671	A
52	sa	1676	U
52	sa	1678	A
52	sa	1681	A
52	sa	1683	A
52	sa	1689	A
52	sa	1690	A
52	sa	1691	A
52	sa	1694	A
52	sa	1697	U
52	sa	1700	A
52	sa	1702	G

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Mol	Chain	Res	Type
52	sa	1703	A
52	sa	1706	G
52	sa	1708	G
52	sa	1711	A
52	sa	1712	A
52	sa	1713	A
52	sa	1714	U
52	sa	1717	U
52	sa	1723	U
52	sa	1725	A
52	sa	1734	G
52	sa	1735	A
52	sa	1736	A
52	sa	1750	G
52	sa	1767	A
52	sa	1773	G
52	sa	1777	A
52	sa	1780	A
52	sa	1781	C
52	sa	1782	G
52	sa	1785	C
52	sa	1788	G
52	sa	1797	A
52	sa	1800	C
52	sa	1801	A
52	sa	1821	A
52	sa	1822	U
52	sa	1824	G
52	sa	1825	A
52	sa	1830	A
52	sa	1831	G
52	sa	1832	A
52	sa	1846	G
52	sa	1849	U
52	sa	1850	C
52	sa	1852	G
52	sa	1883	C
52	sa	1885	U
52	sa	1893	A
52	sa	1894	G
52	sa	1902	A
52	sa	1907	G

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Mol	Chain	Res	Type
52	sa	1909	A
52	sa	1913	A
52	sa	1916	U
52	sa	1927	G
52	sa	1929	A
52	sa	1930	C
52	sa	1939	G
52	sa	1940	G
52	sa	1941	A
52	sa	1943	C

There are no RNA pucker outliers to report.

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

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

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

1 ligand is modelled in this entry.

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

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
75	PAR	sa	5101	-	44,45,45	0.71	0	63,67,67	0.83	0

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

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
75	PAR	sa	5101	-	-	3/18/94/94	0/4/4/4

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

All (3) torsion outliers are listed below:

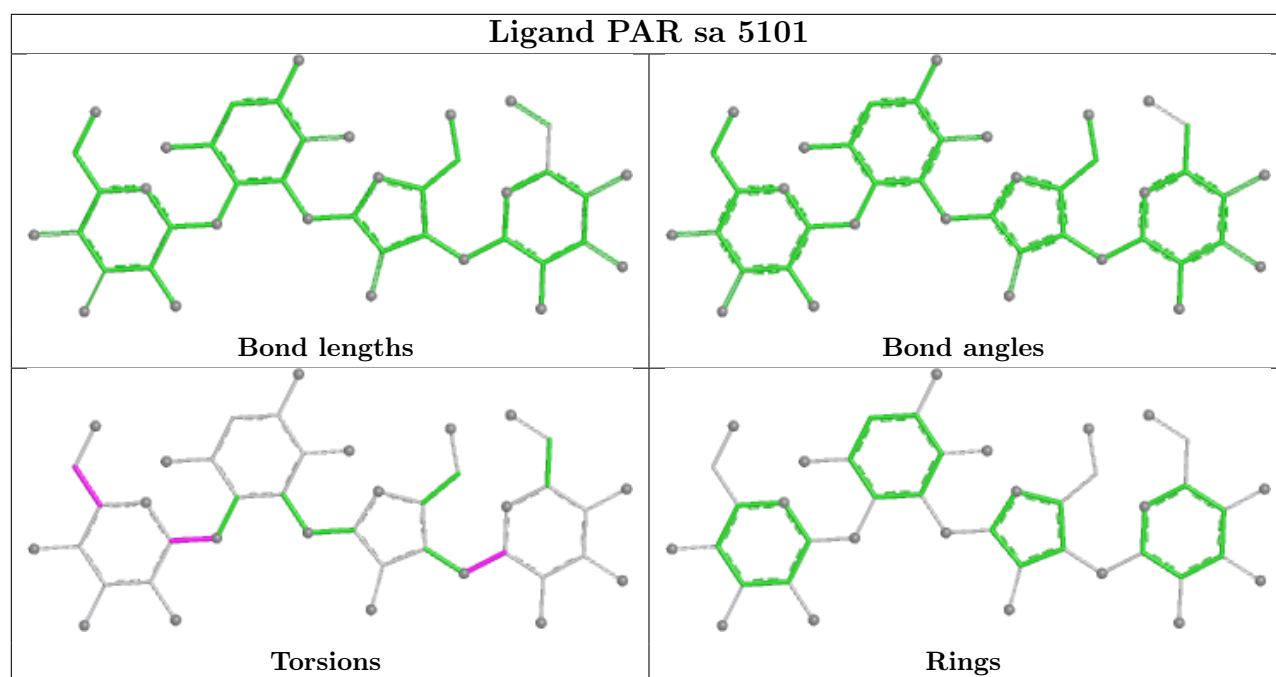
Mol	Chain	Res	Type	Atoms
75	sa	5101	PAR	O54-C14-O33-C33
75	sa	5101	PAR	O51-C51-C61-O61
75	sa	5101	PAR	O51-C11-O11-C42

There are no ring outliers.

1 monomer is involved in 1 short contact:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
75	sa	5101	PAR	1	0

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



5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

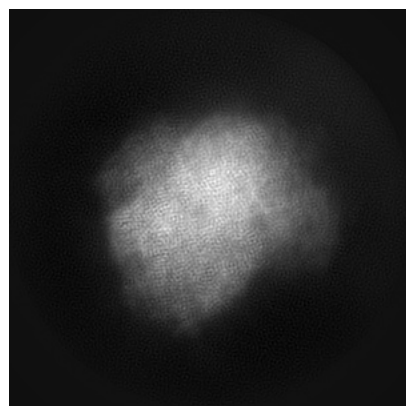
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-64718. These allow visual inspection of the internal detail of the map and identification of artifacts.

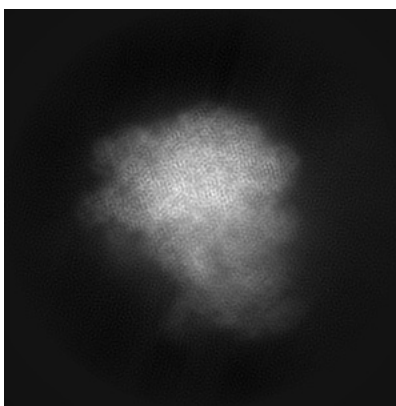
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections [i](#)

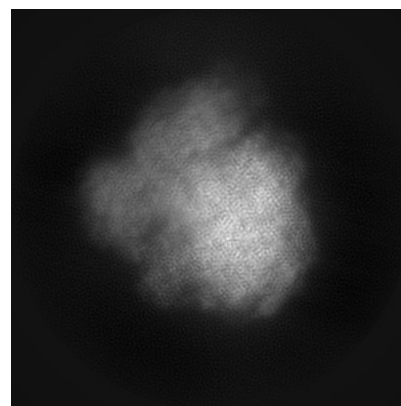
6.1.1 Primary map



X

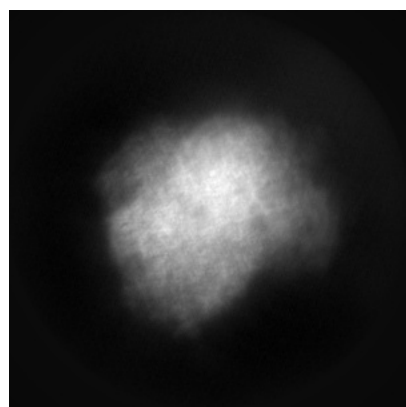


Y

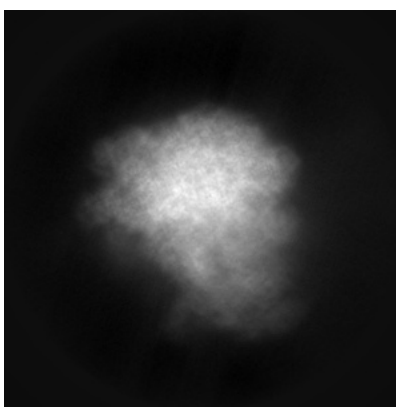


Z

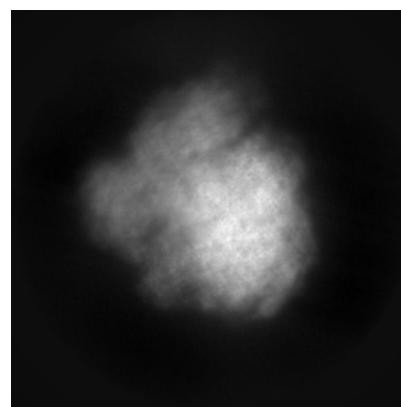
6.1.2 Raw map



X



Y

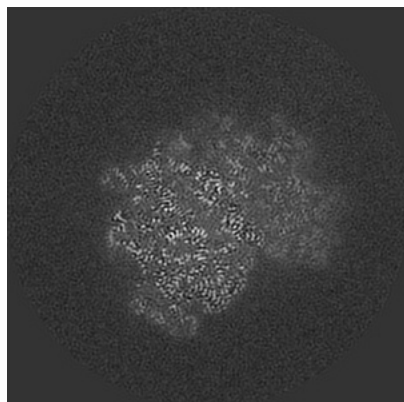


Z

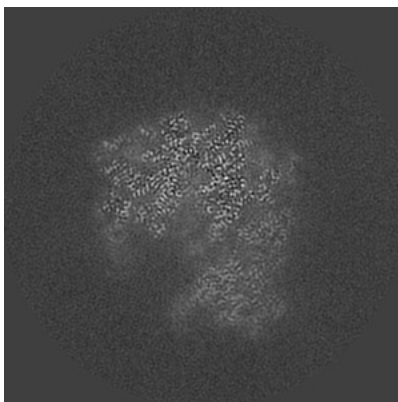
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

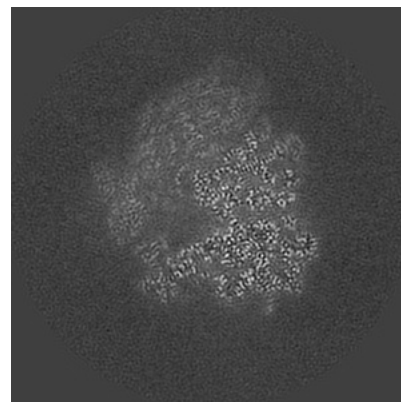
6.2.1 Primary map



X Index: 200

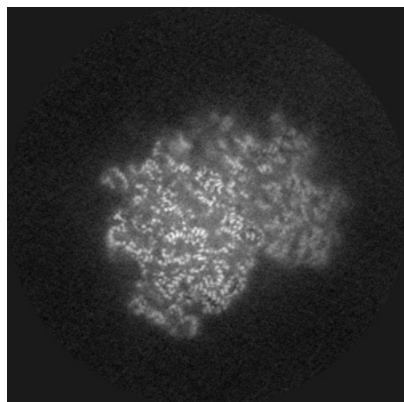


Y Index: 200

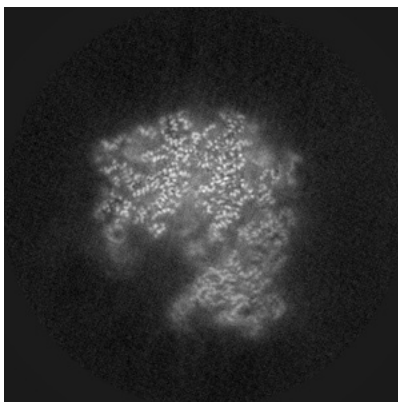


Z Index: 200

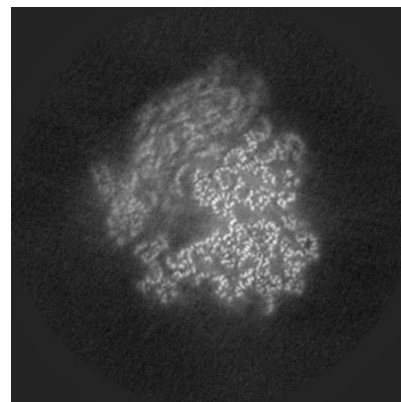
6.2.2 Raw map



X Index: 200



Y Index: 200

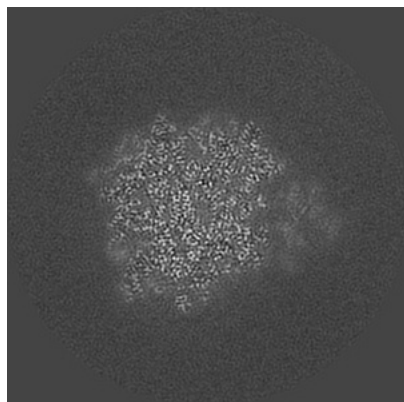


Z Index: 200

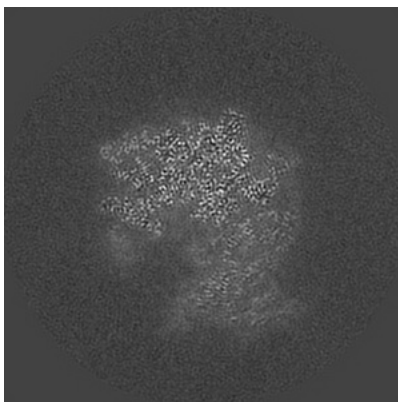
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

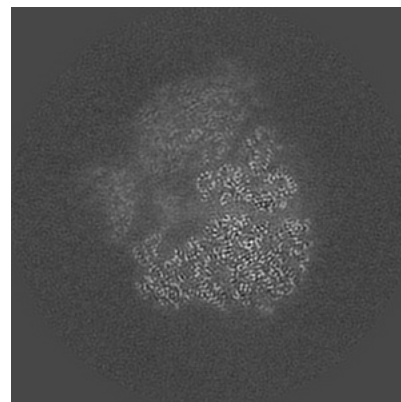
6.3.1 Primary map



X Index: 226

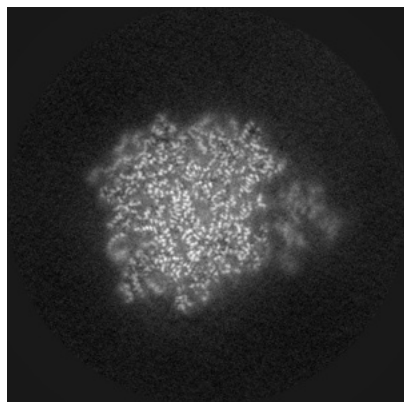


Y Index: 208

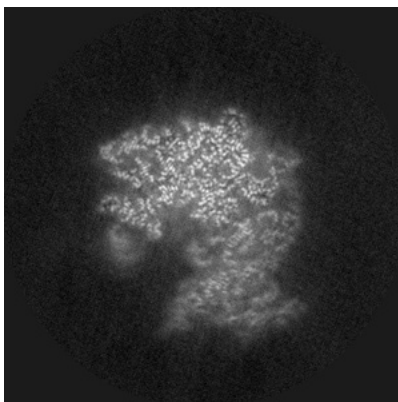


Z Index: 189

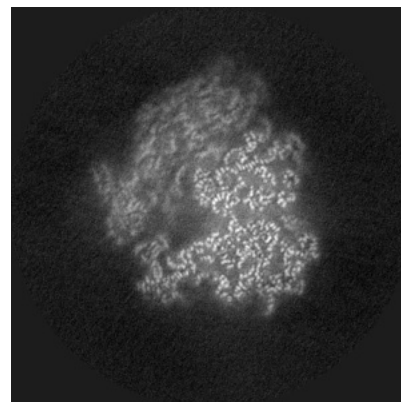
6.3.2 Raw map



X Index: 226



Y Index: 208

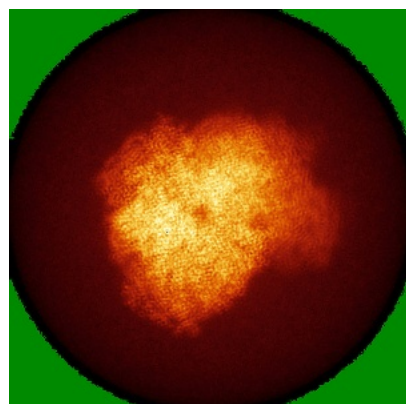


Z Index: 199

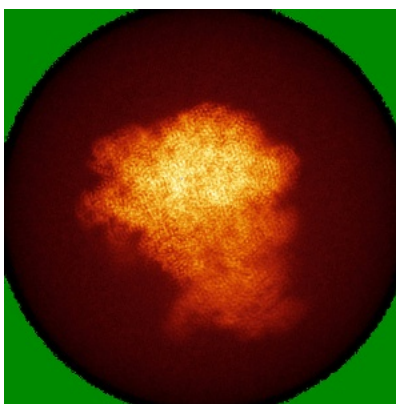
The images above show the largest variance slices of the map in three orthogonal directions.

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

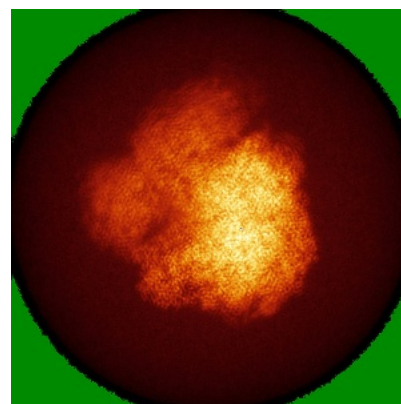
6.4.1 Primary map



X

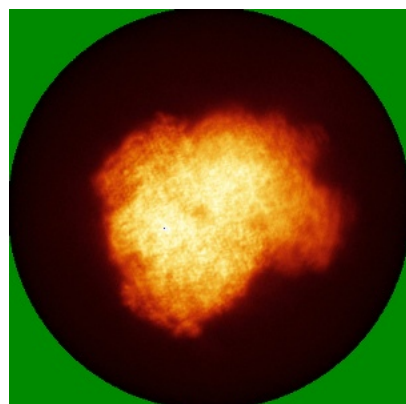


Y

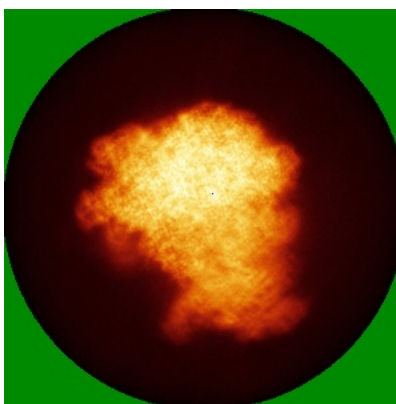


Z

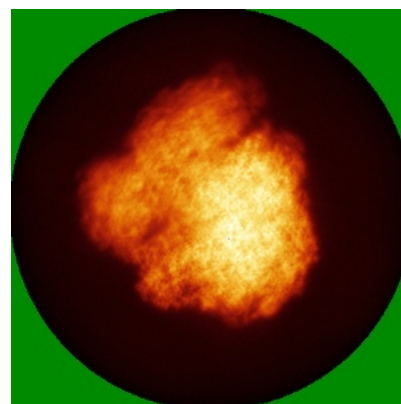
6.4.2 Raw map



X



Y



Z

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

6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 2.8. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

6.5.2 Raw map



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

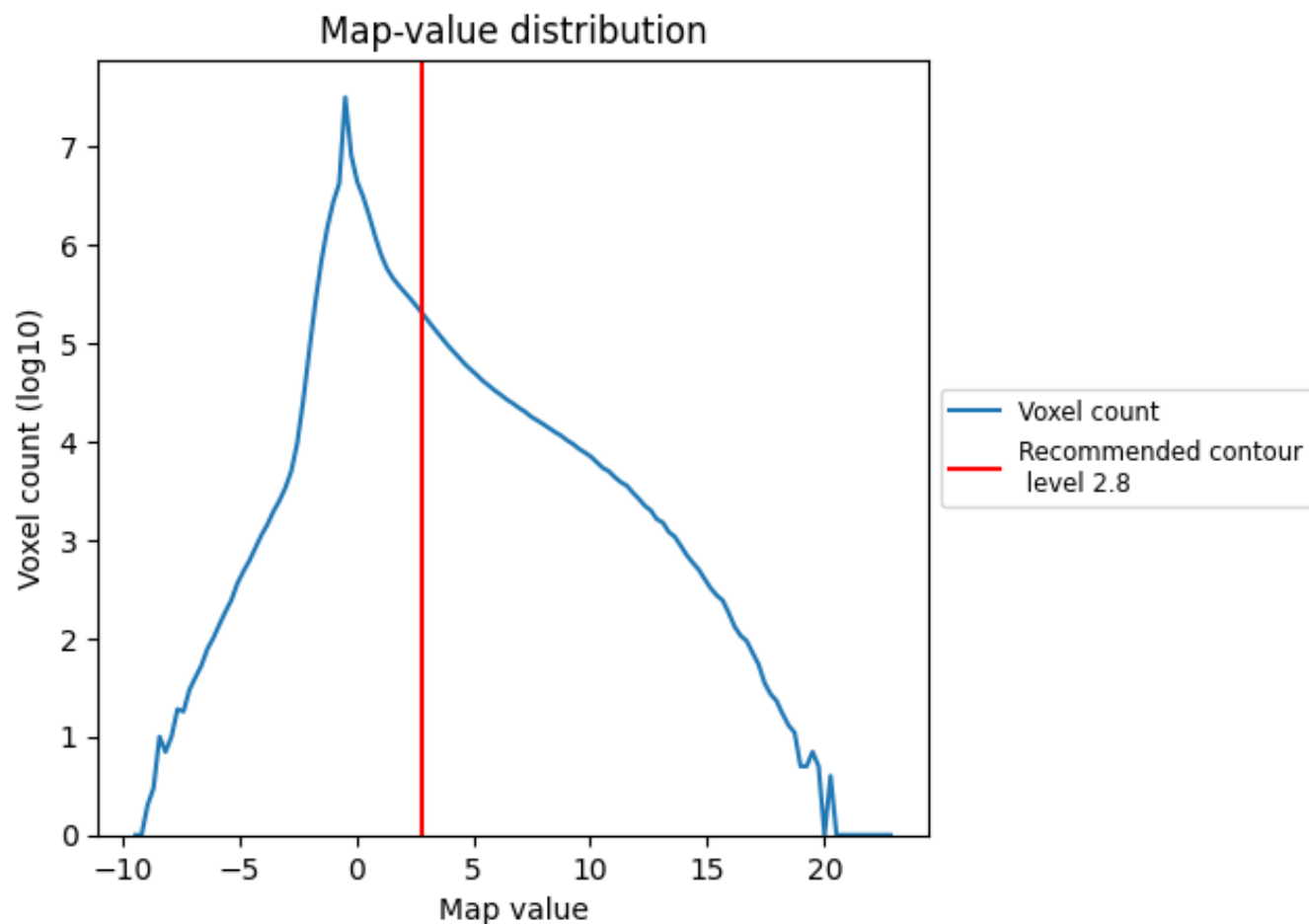
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

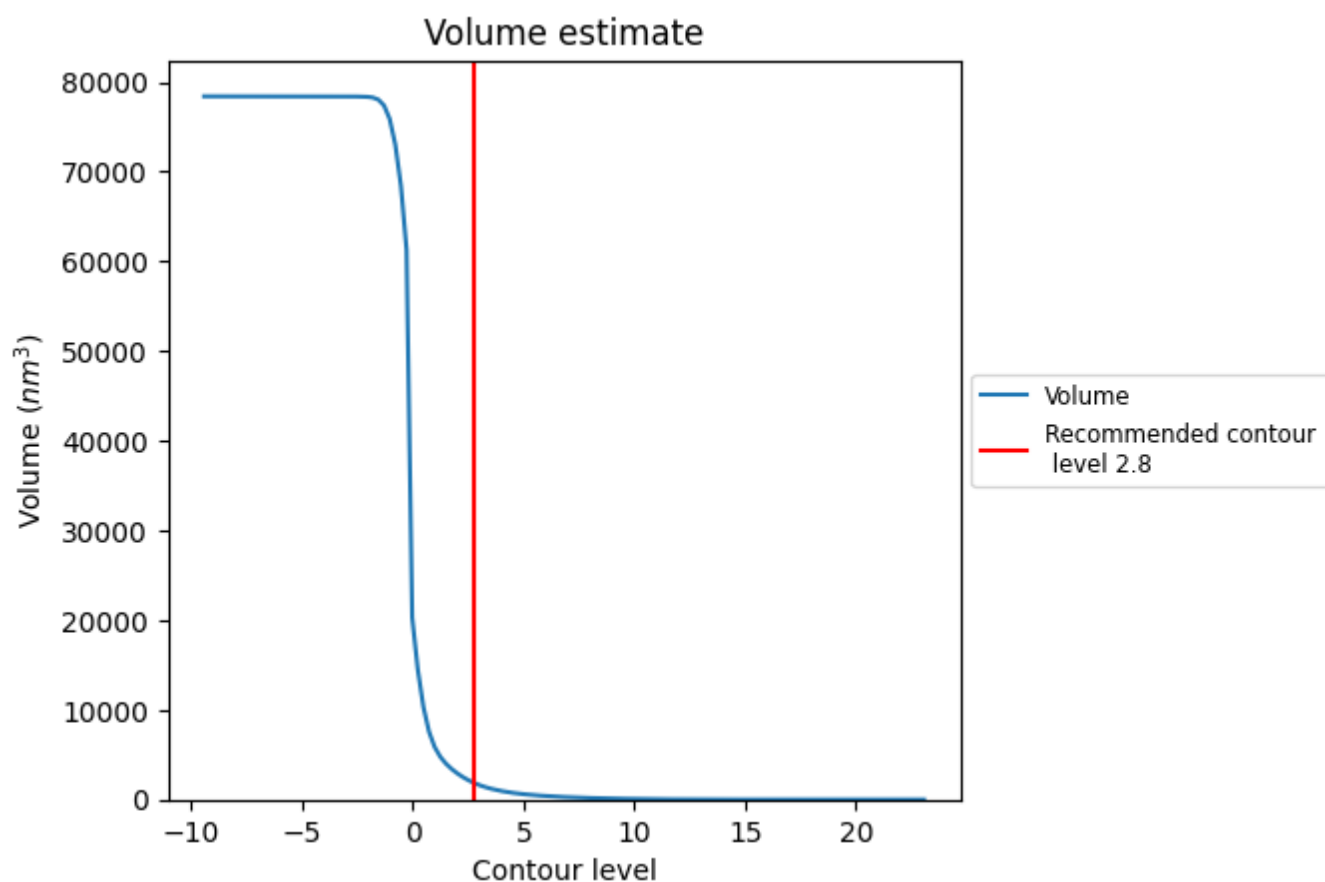
This section contains the results of statistical analysis of the map.

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.

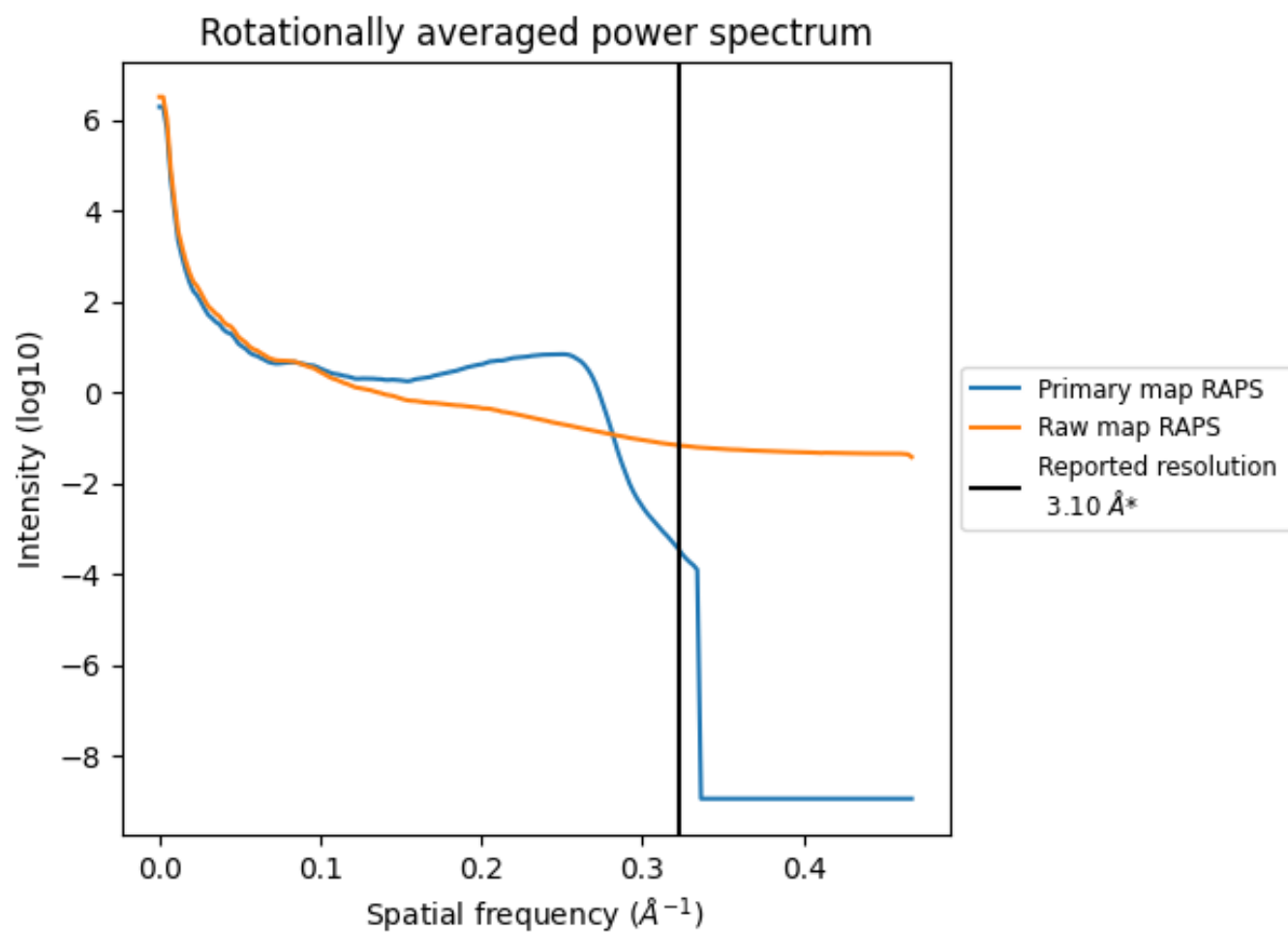
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 1820 nm^3 ; this corresponds to an approximate mass of 1644 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 ⓘ

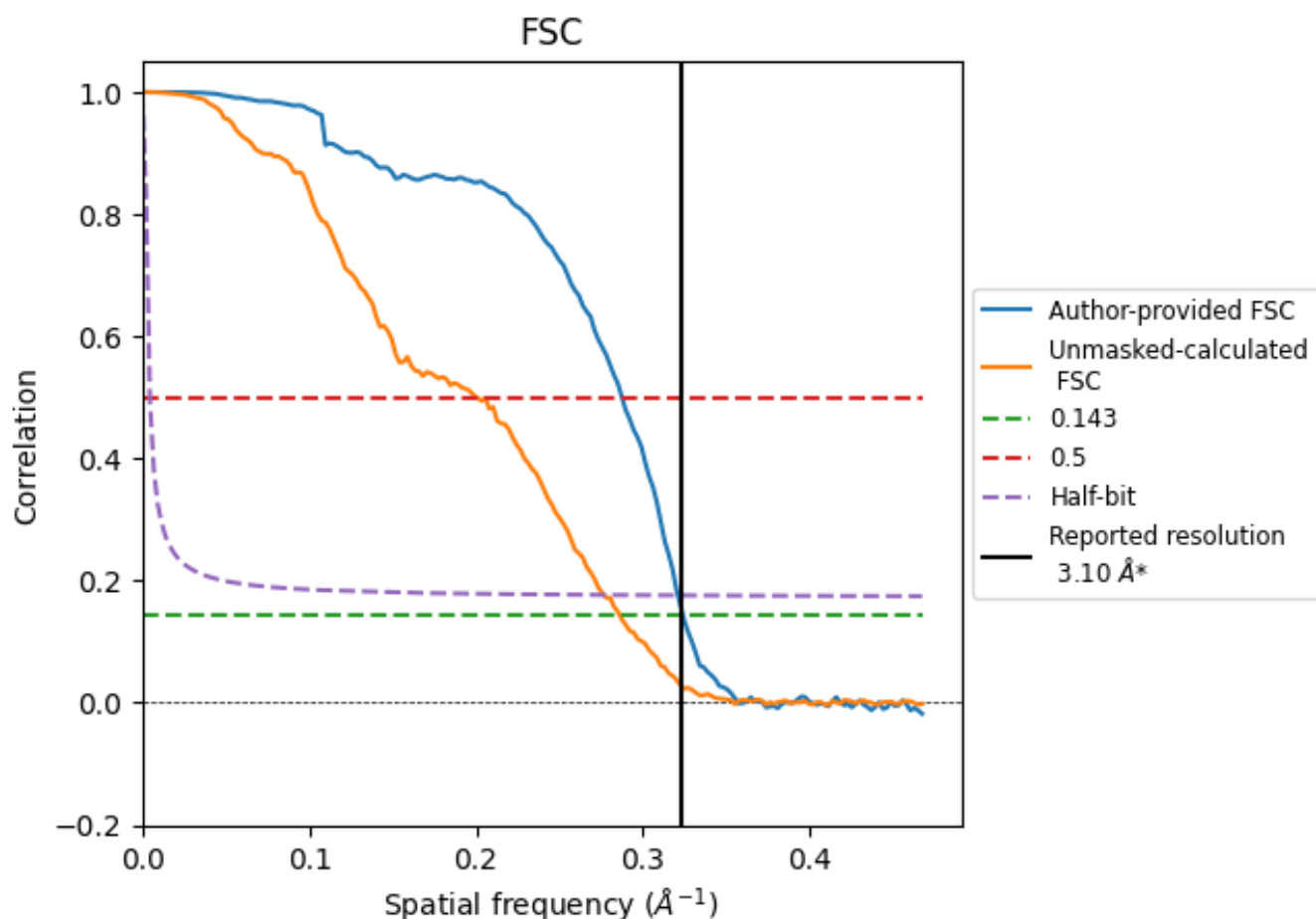


*Reported resolution corresponds to spatial frequency of 0.323 Å⁻¹

8 Fourier-Shell correlation [i](#)

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

8.1 FSC [i](#)



*Reported resolution corresponds to spatial frequency of 0.323 $\text{\AA}^{-1}$

8.2 Resolution estimates [i](#)

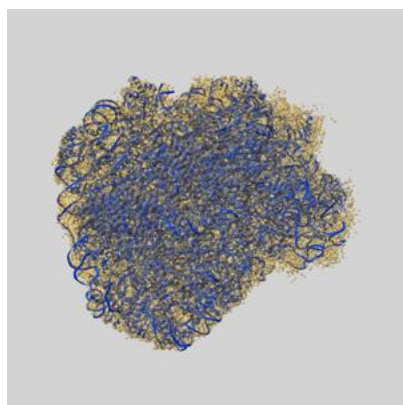
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	3.10	-	-
Author-provided FSC curve	3.09	3.48	3.12
Unmasked-calculated*	3.49	4.99	3.61

*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps. The value from deposited half-maps intersecting FSC 0.143 CUT-OFF 3.49 differs from the reported value 3.1 by more than 10 %

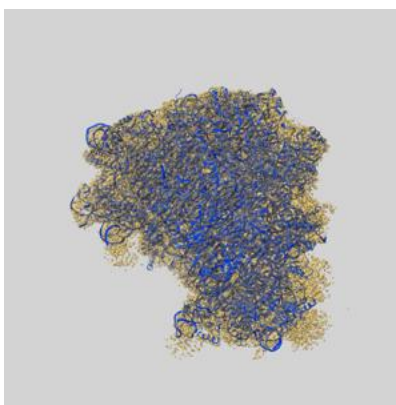
9 Map-model fit [i](#)

This section contains information regarding the fit between EMDB map EMD-64718 and PDB model 9V26. Per-residue inclusion information can be found in [section 3](#) on [page 18](#).

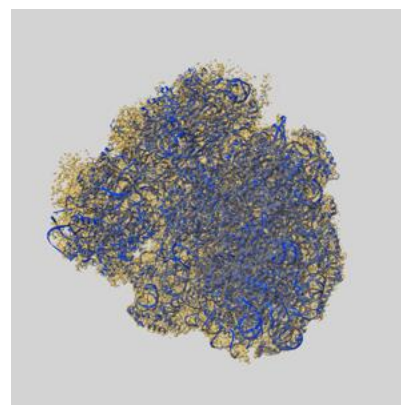
9.1 Map-model overlay [i](#)



X



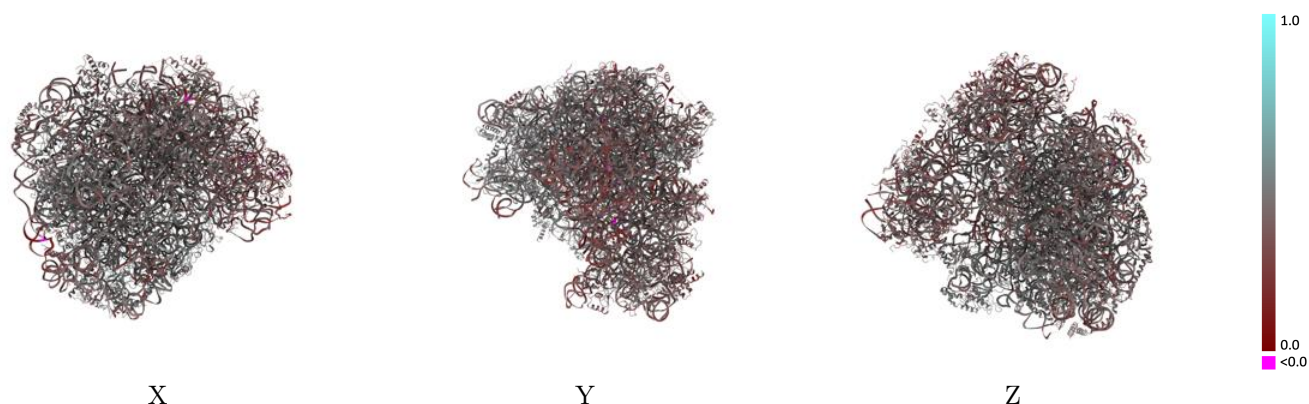
Y



Z

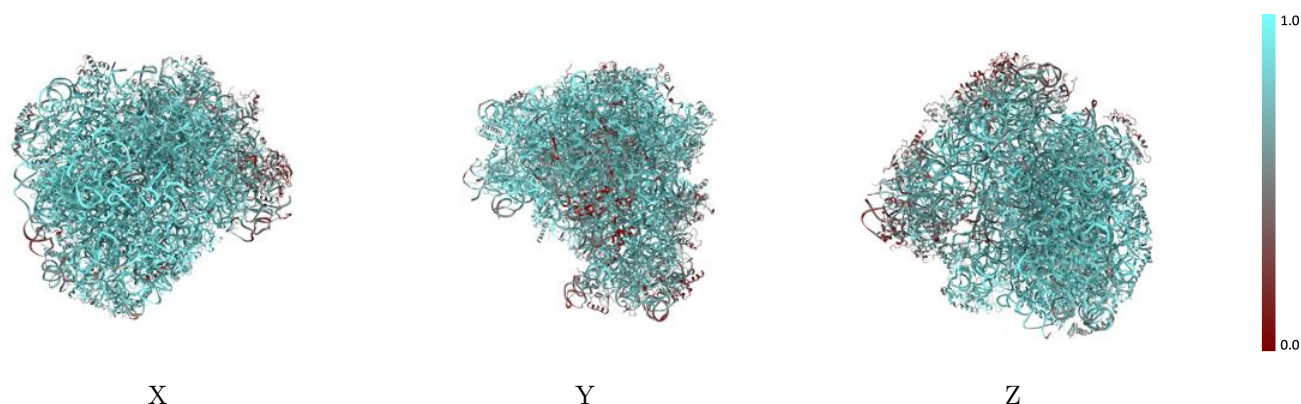
The images above show the 3D surface view of the map at the recommended contour level 2.8 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



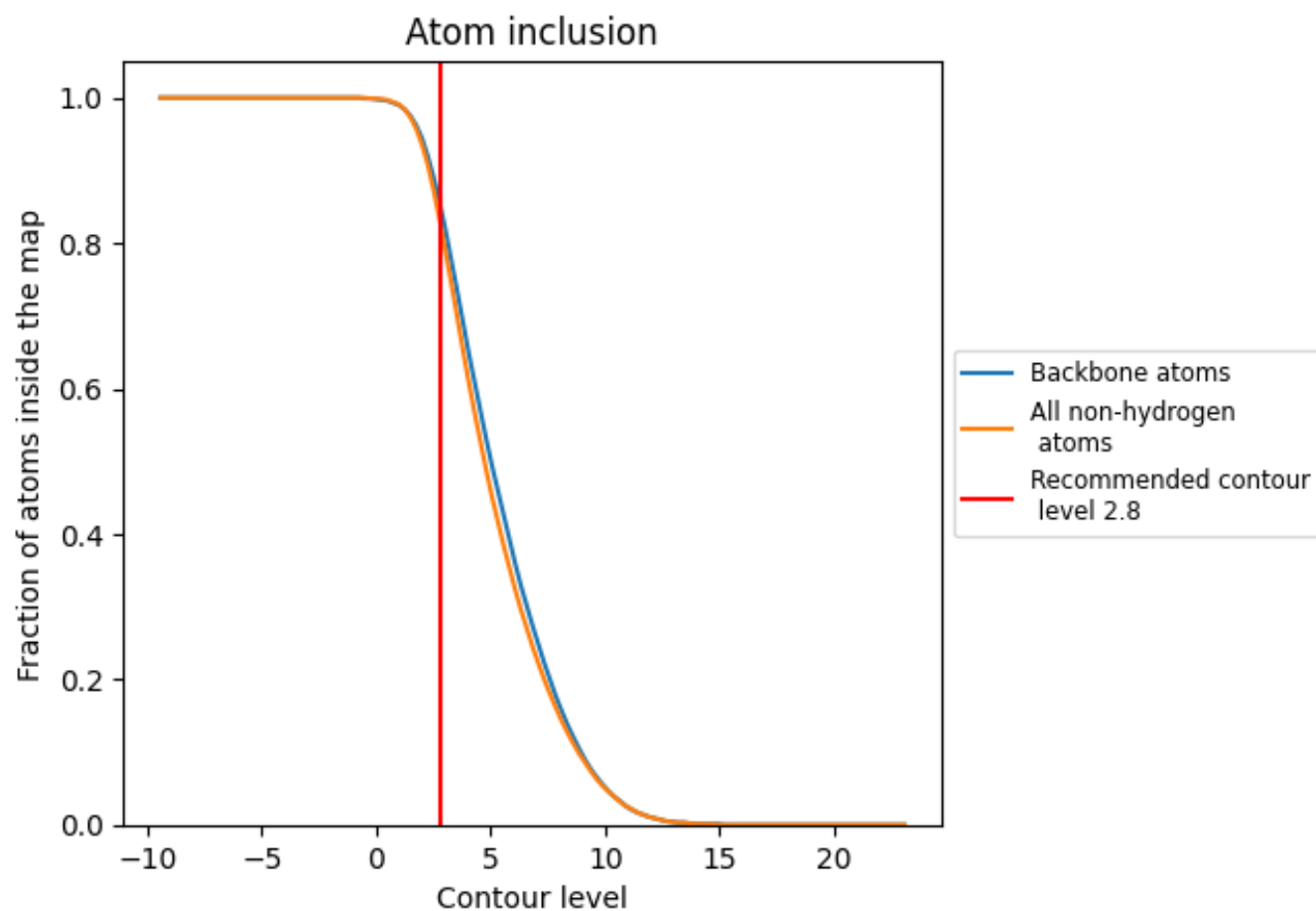
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions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

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



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (2.8).

9.4 Atom inclusion ⓘ



At the recommended contour level, 85% of all backbone atoms, 83% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary ⓘ

The table lists the average atom inclusion at the recommended contour level (2.8) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.8270	 0.4130
1A	 0.9070	 0.4200
1B	 0.9120	 0.4080
1C	 0.9100	 0.4230
1D	 0.9230	 0.4450
1E	 0.8900	 0.4610
1F	 0.8670	 0.4670
1G	 0.8250	 0.4750
1H	 0.6880	 0.4270
1I	 0.8840	 0.4840
1J	 0.7470	 0.4160
1K	 0.8330	 0.4650
1L	 0.8960	 0.4660
1M	 0.7560	 0.4500
1N	 0.7680	 0.4370
1O	 0.8830	 0.4700
1P	 0.8530	 0.4830
1Q	 0.9420	 0.4580
1R	 0.8720	 0.4480
1S	 0.9280	 0.4750
1T	 0.9250	 0.4890
1U	 0.8370	 0.4160
1V	 0.8980	 0.4880
1W	 0.4810	 0.3410
1X	 0.9260	 0.4520
1Y	 0.8150	 0.4240
1Z	 0.8800	 0.4470
1a	 0.7930	 0.4400
1b	 0.6850	 0.3910
1c	 0.9270	 0.4860
1d	 0.9320	 0.4920
1e	 0.7350	 0.4120
1f	 0.7430	 0.3900
1g	 0.9260	 0.4690
1h	 0.8770	 0.4200



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Chain	Atom inclusion	Q-score
li	 0.8150	 0.4280
lj	 0.9170	 0.4820
lk	 0.8470	 0.4520
ll	 0.9480	 0.4640
lm	 0.8700	 0.4310
ln	 0.5970	 0.3630
lo	 0.9350	 0.4390
lp	 0.8810	 0.4530
lq	 0.9050	 0.4670
sA	 0.4610	 0.3600
sB	 0.8250	 0.4330
sC	 0.5320	 0.3900
sD	 0.5410	 0.3950
sE	 0.8100	 0.4150
sI	 0.6790	 0.3060
sJ	 0.7450	 0.3500
sK	 0.9580	 0.4280
sa	 0.7950	 0.3650
sc	 0.6840	 0.4040
sd	 0.5750	 0.3790
se	 0.6700	 0.4030
sf	 0.3170	 0.3220
sg	 0.5430	 0.3810
sh	 0.6080	 0.3640
si	 0.3450	 0.3330
sj	 0.5530	 0.3520
sk	 0.4740	 0.3600
sl	 0.4260	 0.3510
sm	 0.6350	 0.3450
so	 0.7490	 0.3860
sp	 0.7710	 0.4280
sq	 0.5660	 0.3800
sr	 0.7420	 0.4050
ss	 0.5220	 0.3870
st	 0.4880	 0.3460
su	 0.6230	 0.4100
sv	 0.4890	 0.3920
sw	 0.7330	 0.4070
sx	 0.5400	 0.3940
sy	 0.6470	 0.3740