



Full wwPDB EM Validation Report ⓘ

May 18, 2026 – 07:18 pm BST

PDB ID : 9QZP / pdb_00009qzp
EMDB ID : EMD-53473
Title : Mouse Ribosome Classical Pre translocation state
Authors : Santo, P.E.; Astier, A.; Plisson-Chastang, C.
Deposited on : 2025-04-23
Resolution : 3.43 Å(reported)
Based on initial model : 7LS1

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 : 1.8.4, CSD as541be (2020)
MolProbity : 4-5-2 with Phenix2.0
Percentile statistics : 20250101.v01 (using entries in the PDB archive January 1st 2025)
EM percentile statistics : 202505.v01 (Using data in the EMDb archive up until May 2025)
MapQ : 1.9.13
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.49

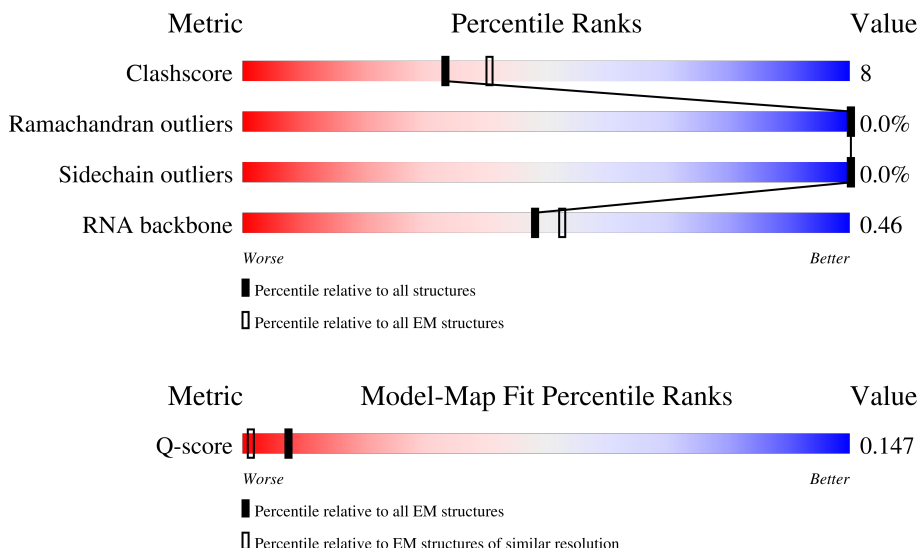
1 Overall quality at a glance i

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 3.43 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	13927 (2.93 - 3.93)

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

Mol	Chain	Length	Quality of chain
1	A1	270	
2	A2	3615	
3	A3	152	

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Mol	Chain	Length	Quality of chain
4	B1	266	
5	B2	121	
6	B3	145	
7	Bv	76	
7	Bz	76	
7	n2	76	
8	Bx	10	
9	By	22	
10	C1	192	
11	C2	156	
12	C3	119	
13	D1	214	
14	D2	257	
15	D3	83	
16	E1	178	
17	E2	403	
18	E3	143	
19	F1	211	
20	F2	419	
21	F3	115	
22	G1	217	
23	G2	297	
24	G3	69	
25	H1	204	
26	H2	296	

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Mol	Chain	Length	Quality of chain
27	H3	56	<div>93%</div> <div>75% 21%</div>
28	I2	203	<div>71%</div> <div>78% 21%</div>
29	I3	317	<div>99%</div> <div>70% 29%</div>
30	J2	184	<div>65%</div> <div>68% 15% 17%</div>
31	J3	293	<div>72%</div> <div>57% 18% 25%</div>
32	K2	188	<div>68%</div> <div>79% 20%</div>
33	K3	249	<div>89%</div> <div>72% 19% 9%</div>
34	L1	217	<div>74%</div> <div>55% 19% 26%</div>
35	L2	196	<div>79%</div> <div>80% 12% 9%</div>
36	L3	194	<div>93%</div> <div>75% 20% 5%</div>
37	M2	176	<div>73%</div> <div>85% 15%</div>
38	M3	132	<div>92%</div> <div>70% 23% 8%</div>
39	N2	160	<div>68%</div> <div>82% 18%</div>
40	N3	151	<div>96%</div> <div>85% 15%</div>
41	O2	128	<div>76%</div> <div>62% 16% 21%</div>
42	O3	151	<div>87%</div> <div>71% 17% 11%</div>
43	P2	140	<div>69%</div> <div>66% 26% 8%</div>
44	P3	130	<div>97%</div> <div>83% 16%</div>
45	Q2	157	<div>35%</div> <div>35% 61%</div>
46	Q3	133	<div>92%</div> <div>76% 16% 8%</div>
47	R2	156	<div>67%</div> <div>60% 15% 24%</div>
48	R3	125	<div>57%</div> <div>42% 15% 42%</div>
49	S2	145	<div>77%</div> <div>78% 14% 8%</div>
50	S3	84	<div>96%</div> <div>81% 18%</div>
51	T2	136	<div>87%</div> <div>80% 19%</div>

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Mol	Chain	Length	Quality of chain
52	T3	133	
53	U2	148	
54	U3	156	
55	V2	160	
56	W2	115	
57	X2	125	
58	Y2	135	
59	Z2	110	
60	a2	117	
61	b2	123	
62	c2	105	
63	d2	97	
64	e2	70	
65	f2	51	
66	g2	128	
67	h2	25	
68	i2	106	
69	j2	92	
70	k2	137	
71	o2	295	
72	p2	264	
73	q2	243	
74	r2	263	
75	s2	204	
76	t2	194	

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Mol	Chain	Length	Quality of chain
77	u2	208	<div><div></div><div>97%</div><div>76%</div><div>23%</div><div></div></div>
78	v2	165	<div><div></div><div>58%</div><div>39%</div><div>18%</div><div>42%</div><div></div></div>
79	w2	158	<div><div></div><div>91%</div><div>79%</div><div>16%</div><div>5%</div><div></div></div>
80	x2	145	<div><div></div><div>90%</div><div>58%</div><div>33%</div><div>9%</div><div></div></div>
81	y2	146	<div><div></div><div>96%</div><div>80%</div><div>17%</div><div></div><div></div></div>
82	z2	135	<div><div></div><div>50%</div><div>38%</div><div>13%</div><div>49%</div><div></div></div>
83	m2	1635	<div><div></div><div>86%</div><div>43%</div><div>47%</div><div>9%</div><div></div></div>

2 Entry composition

There are 86 unique types of molecules in this entry. The entry contains 214961 atoms, of which 0 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a protein called Large ribosomal subunit protein uL30.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	A1	222	Total	C	N	O	S	1	0
			1851	1190	356	297	8		

- Molecule 2 is a RNA chain called 28S ribosomal RNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	A2	3615	Total	C	N	O	P	0	0
			77547	34568	14148	25217	3614		

- Molecule 3 is a protein called Small ribosomal subunit protein uS13.

Mol	Chain	Residues	Atoms					AltConf	Trace
3	A3	140	Total	C	N	O	S	0	0
			1151	725	228	197	1		

- Molecule 4 is a protein called Large ribosomal subunit protein eL8.

Mol	Chain	Residues	Atoms					AltConf	Trace
4	B1	223	Total	C	N	O	S	1	0
			1812	1156	351	301	4		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
5	B2	119	Total	C	N	O	P	0	0
			2538	1132	454	834	118		

- Molecule 6 is a protein called Small ribosomal subunit protein eS19.

Mol	Chain	Residues	Atoms					AltConf	Trace
6	B3	141	Total	C	N	O	S	0	0
			1104	691	215	196	2		

- Molecule 7 is a RNA chain called transfer RNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
7	Bv	76	Total	C	N	O	P	0	0
			1623	723	290	534	76		
7	Bz	76	Total	C	N	O	P	0	0
			1623	723	290	534	76		
7	n2	76	Total	C	N	O	P	0	0
			1623	723	290	534	76		

- Molecule 8 is a RNA chain called messenger RNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
8	Bx	10	Total	C	N	O	P	0	0
			200	90	20	80	10		

- Molecule 9 is a protein called Nascent protein chain.

Mol	Chain	Residues	Atoms				AltConf	Trace
9	By	22	Total	C	N	O	0	0
			110	66	22	22		

- Molecule 10 is a protein called Large ribosomal subunit protein uL6.

Mol	Chain	Residues	Atoms					AltConf	Trace
10	C1	190	Total	C	N	O	S	0	0
			1519	956	284	273	6		

- Molecule 11 is a RNA chain called 5.8S ribosomal RNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
11	C2	156	Total	C	N	O	P	0	0
			3315	1481	585	1094	155		

- Molecule 12 is a protein called Small ribosomal subunit protein uS10.

Mol	Chain	Residues	Atoms					AltConf	Trace
12	C3	102	Total	C	N	O	S	0	0
			808	507	154	143	4		

- Molecule 13 is a protein called Large ribosomal subunit protein uL16.

Mol	Chain	Residues	Atoms					AltConf	Trace
13	D1	204	Total	C	N	O	S	0	0
			1656	1052	319	272	13		

- Molecule 14 is a protein called Large ribosomal subunit protein uL2.

Mol	Chain	Residues	Atoms					AltConf	Trace
14	D2	251	Total	C	N	O	S	0	0
			1921	1204	393	318	6		

- Molecule 15 is a protein called Small ribosomal subunit protein eS21.

Mol	Chain	Residues	Atoms					AltConf	Trace
15	D3	83	Total	C	N	O	S	0	0
			638	392	119	122	5		

- Molecule 16 is a protein called Large ribosomal subunit protein uL5.

Mol	Chain	Residues	Atoms					AltConf	Trace
16	E1	174	Total	C	N	O	S	0	0
			1397	880	260	251	6		

- Molecule 17 is a protein called Large ribosomal subunit protein uL3.

Mol	Chain	Residues	Atoms					AltConf	Trace
17	E2	402	Total	C	N	O	S	0	0
			3238	2060	609	555	14		

- Molecule 18 is a protein called Small ribosomal subunit protein uS12.

Mol	Chain	Residues	Atoms					AltConf	Trace
18	E3	139	Total	C	N	O	S	0	0
			1080	682	214	181	3		

- Molecule 19 is a protein called Large ribosomal subunit protein eL13.

Mol	Chain	Residues	Atoms					AltConf	Trace
19	F1	203	Total	C	N	O	S	0	0
			1643	1029	339	271	4		

- Molecule 20 is a protein called Large ribosomal subunit protein uL4.

Mol	Chain	Residues	Atoms					AltConf	Trace
20	F2	359	Total	C	N	O	S	0	0
			2867	1803	573	476	15		

- Molecule 21 is a protein called Small ribosomal subunit protein eS26.

Mol	Chain	Residues	Atoms					AltConf	Trace
21	F3	98	Total	C	N	O	S	1	0
			789	491	164	129	5		

- Molecule 22 is a protein called Large ribosomal subunit protein eL14.

Mol	Chain	Residues	Atoms					AltConf	Trace
22	G1	139	Total	C	N	O	S	0	0
			1143	732	221	183	7		

- Molecule 23 is a protein called Large ribosomal subunit protein uL18.

Mol	Chain	Residues	Atoms					AltConf	Trace
23	G2	293	Total	C	N	O	S	0	0
			2389	1509	441	425	14		

- Molecule 24 is a protein called Small ribosomal subunit protein eS28.

Mol	Chain	Residues	Atoms					AltConf	Trace
24	G3	62	Total	C	N	O	S	0	0
			488	297	97	92	2		

- Molecule 25 is a protein called Large ribosomal subunit protein eL15.

Mol	Chain	Residues	Atoms					AltConf	Trace
25	H1	203	Total	C	N	O	S	0	0
			1701	1072	359	266	4		

- Molecule 26 is a protein called Large ribosomal subunit protein eL6.

Mol	Chain	Residues	Atoms					AltConf	Trace
26	H2	221	Total	C	N	O	S	0	0
			1789	1145	342	298	4		

- Molecule 27 is a protein called Small ribosomal subunit protein uS14.

Mol	Chain	Residues	Atoms					AltConf	Trace
27	H3	54	Total	C	N	O	S	0	0
			455	284	93	73	5		

- Molecule 28 is a protein called Large ribosomal subunit protein uL13.

Mol	Chain	Residues	Atoms					AltConf	Trace
28	I2	201	Total	C	N	O	S	0	0
			1640	1055	320	259	6		

- Molecule 29 is a protein called Small ribosomal subunit protein RACK1.

Mol	Chain	Residues	Atoms					AltConf	Trace
29	I3	313	Total	C	N	O	S	0	0
			2436	1535	424	465	12		

- Molecule 30 is a protein called Large ribosomal subunit protein uL22.

Mol	Chain	Residues	Atoms					AltConf	Trace
30	J2	153	Total	C	N	O	S	0	0
			1242	777	241	215	9		

- Molecule 31 is a protein called Small ribosomal subunit protein uS5.

Mol	Chain	Residues	Atoms					AltConf	Trace
31	J3	219	Total	C	N	O	S	0	0
			1700	1101	292	298	9		

- Molecule 32 is a protein called Large ribosomal subunit protein eL18.

Mol	Chain	Residues	Atoms					AltConf	Trace
32	K2	186	Total	C	N	O	S	0	0
			1511	946	313	248	4		

- Molecule 33 is a protein called Small ribosomal subunit protein eS6.

Mol	Chain	Residues	Atoms					AltConf	Trace
33	K3	227	Total	C	N	O	S	0	0
			1840	1149	367	317	7		

- Molecule 34 is a protein called Large ribosomal subunit protein uL1.

Mol	Chain	Residues	Atoms					AltConf	Trace
34	L1	161	Total	C	N	O	S	0	0
			1300	833	230	231	6		

- Molecule 35 is a protein called Large ribosomal subunit protein eL19.

Mol	Chain	Residues	Atoms					AltConf	Trace
35	L2	179	Total	C	N	O	S	0	0
			1499	927	326	237	9		

- Molecule 36 is a protein called Small ribosomal subunit protein uS4.

Mol	Chain	Residues	Atoms					AltConf	Trace
36	L3	184	Total	C	N	O	S	0	0
			1518	964	305	247	2		

- Molecule 37 is a protein called Large ribosomal subunit protein eL20.

Mol	Chain	Residues	Atoms					AltConf	Trace
37	M2	175	Total	C	N	O	S	0	0
			1450	924	283	233	10		

- Molecule 38 is a protein called Small ribosomal subunit protein eS12.

Mol	Chain	Residues	Atoms					AltConf	Trace
38	M3	122	Total	C	N	O	S	0	0
			952	599	168	177	8		

There is a discrepancy between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
M3	69	LEU	CYS	variant	UNP P63323

- Molecule 39 is a protein called Large ribosomal subunit protein eL21.

Mol	Chain	Residues	Atoms					AltConf	Trace
39	N2	159	Total	C	N	O	S	0	0
			1299	824	252	217	6		

- Molecule 40 is a protein called Small ribosomal subunit protein uS15.

Mol	Chain	Residues	Atoms					AltConf	Trace
40	N3	150	Total	C	N	O	S	0	0
			1208	773	229	205	1		

- Molecule 41 is a protein called Large ribosomal subunit protein eL22.

Mol	Chain	Residues	Atoms					AltConf	Trace
41	O2	101	Total	C	N	O	S	0	0
			825	529	144	150	2		

- Molecule 42 is a protein called Small ribosomal subunit protein uS11.

Mol	Chain	Residues	Atoms					AltConf	Trace
42	O3	134	Total	C	N	O	S	0	0
			1002	612	197	187	6		

- Molecule 43 is a protein called Large ribosomal subunit protein uL14.

Mol	Chain	Residues	Atoms					AltConf	Trace
43	P2	129	Total	C	N	O	S	0	0
			969	613	182	169	5		

- Molecule 44 is a protein called Small ribosomal subunit protein uS8.

Mol	Chain	Residues	Atoms					AltConf	Trace
44	P3	129	Total	C	N	O	S	0	0
			1034	659	193	176	6		

- Molecule 45 is a protein called Large ribosomal subunit protein eL24.

Mol	Chain	Residues	Atoms					AltConf	Trace
45	Q2	62	Total	C	N	O	S	0	0
			519	332	101	83	3		

- Molecule 46 is a protein called Small ribosomal subunit protein eS24.

Mol	Chain	Residues	Atoms					AltConf	Trace
46	Q3	122	Total	C	N	O	S	0	0
			1002	635	196	166	5		

- Molecule 47 is a protein called Large ribosomal subunit protein uL23.

Mol	Chain	Residues	Atoms					AltConf	Trace
47	R2	118	Total	C	N	O	S	0	0
			967	618	181	167	1		

- Molecule 48 is a protein called Small ribosomal subunit protein eS25.

Mol	Chain	Residues	Atoms					AltConf	Trace
48	R3	72	Total	C	N	O	S	0	0
			570	366	104	99	1		

- Molecule 49 is a protein called Large ribosomal subunit protein uL24.

Mol	Chain	Residues	Atoms					AltConf	Trace
49	S2	134	Total	C	N	O	S	0	0
			1115	700	226	186	3		

- Molecule 50 is a protein called Small ribosomal subunit protein eS27.

Mol	Chain	Residues	Atoms					AltConf	Trace
50	S3	83	Total	C	N	O	S	0	0
			651	408	121	115	7		

- Molecule 51 is a protein called Large ribosomal subunit protein eL27.

Mol	Chain	Residues	Atoms					AltConf	Trace
51	T2	135	Total	C	N	O	S	0	0
			1107	714	208	182	3		

- Molecule 52 is a protein called Ubiquitin-like FUBI-ribosomal protein eS30 fusion protein.

Mol	Chain	Residues	Atoms					AltConf	Trace
52	T3	55	Total	C	N	O	S	0	0
			438	271	95	71	1		

- Molecule 53 is a protein called Large ribosomal subunit protein uL15.

Mol	Chain	Residues	Atoms					AltConf	Trace
53	U2	147	Total	C	N	O	S	0	0
			1164	736	239	185	4		

- Molecule 54 is a protein called Ubiquitin-ribosomal protein eS31 fusion protein.

Mol	Chain	Residues	Atoms					AltConf	Trace
54	U3	52	Total	C	N	O	S	0	0
			415	260	74	74	7		

- Molecule 55 is a protein called Large ribosomal subunit protein eL29.

Mol	Chain	Residues	Atoms					AltConf	Trace
55	V2	117	Total	C	N	O	S	0	0
			945	596	198	146	5		

- Molecule 56 is a protein called Large ribosomal subunit protein eL30.

Mol	Chain	Residues	Atoms					AltConf	Trace
56	W2	94	Total	C	N	O	S	0	0
			732	465	130	131	6		

- Molecule 57 is a protein called Large ribosomal subunit protein eL31.

Mol	Chain	Residues	Atoms					AltConf	Trace
57	X2	107	Total	C	N	O	S	0	0
			888	560	171	155	2		

- Molecule 58 is a protein called Large ribosomal subunit protein eL32.

Mol	Chain	Residues	Atoms					AltConf	Trace
58	Y2	128	Total	C	N	O	S	0	0
			1053	667	216	165	5		

- Molecule 59 is a protein called Large ribosomal subunit protein eL33.

Mol	Chain	Residues	Atoms					AltConf	Trace
59	Z2	109	Total	C	N	O	S	0	0
			876	555	174	143	4		

- Molecule 60 is a protein called Large ribosomal subunit protein eL34.

Mol	Chain	Residues	Atoms					AltConf	Trace
60	a2	114	Total	C	N	O	S	0	0
			906	565	187	148	6		

- Molecule 61 is a protein called Large ribosomal subunit protein uL29.

Mol	Chain	Residues	Atoms					AltConf	Trace
61	b2	120	Total	C	N	O	S	0	0
			1001	634	201	165	1		

- Molecule 62 is a protein called Large ribosomal subunit protein eL36.

Mol	Chain	Residues	Atoms					AltConf	Trace
62	c2	102	Total	C	N	O	S	0	0
			832	521	177	129	5		

- Molecule 63 is a protein called Large ribosomal subunit protein eL37.

Mol	Chain	Residues	Atoms					AltConf	Trace
63	d2	86	Total	C	N	O	S	0	0
			705	434	155	111	5		

- Molecule 64 is a protein called Large ribosomal subunit protein eL38.

Mol	Chain	Residues	Atoms					AltConf	Trace
64	e2	69	Total	C	N	O	S	0	0
			568	365	103	99	1		

- Molecule 65 is a protein called Large ribosomal subunit protein eL39.

Mol	Chain	Residues	Atoms					AltConf	Trace
65	f2	50	Total	C	N	O	S	0	0
			444	281	98	64	1		

- Molecule 66 is a protein called Ubiquitin-ribosomal protein eL40 fusion protein.

Mol	Chain	Residues	Atoms					AltConf	Trace
66	g2	52	Total	C	N	O	S	0	0
			429	266	90	67	6		

- Molecule 67 is a protein called 60S ribosomal protein L41.

Mol	Chain	Residues	Atoms					AltConf	Trace
67	h2	24	Total	C	N	O	S	0	0
			230	139	62	26	3		

- Molecule 68 is a protein called Large ribosomal subunit protein eL42.

Mol	Chain	Residues	Atoms					AltConf	Trace
68	i2	103	Total	C	N	O	S	0	0
			842	528	172	136	6		

- Molecule 69 is a protein called Large ribosomal subunit protein eL43.

Mol	Chain	Residues	Atoms					AltConf	Trace
69	j2	89	Total	C	N	O	S	0	0
			694	436	133	118	7		

- Molecule 70 is a protein called Large ribosomal subunit protein eL28.

Mol	Chain	Residues	Atoms					AltConf	Trace
70	k2	125	Total	C	N	O	S	0	0
			1001	621	207	168	5		

- Molecule 71 is a protein called Small ribosomal subunit protein uS2.

Mol	Chain	Residues	Atoms					AltConf	Trace
71	o2	214	Total	C	N	O	S	0	0
			1694	1077	297	312	8		

- Molecule 72 is a protein called 40S ribosomal protein S3a.

Mol	Chain	Residues	Atoms					AltConf	Trace
72	p2	212	Total	C	N	O	S	0	0
			1722	1093	308	307	14		

- Molecule 73 is a protein called Small ribosomal subunit protein uS3.

Mol	Chain	Residues	Atoms					AltConf	Trace
73	q2	220	Total	C	N	O	S	0	0
			1711	1092	308	304	7		

- Molecule 74 is a protein called Small ribosomal subunit protein eS4.

Mol	Chain	Residues	Atoms					AltConf	Trace
74	r2	262	Total	C	N	O	S	0	0
			2076	1324	386	358	8		

- Molecule 75 is a protein called Small ribosomal subunit protein uS7.

Mol	Chain	Residues	Atoms					AltConf	Trace
75	s2	183	Total	C	N	O	S	0	0
			1457	912	275	263	7		

- Molecule 76 is a protein called Small ribosomal subunit protein eS7.

Mol	Chain	Residues	Atoms					AltConf	Trace
76	t2	183	Total	C	N	O	S	0	0
			1278	822	243	213			

- Molecule 77 is a protein called Small ribosomal subunit protein eS8.

Mol	Chain	Residues	Atoms					AltConf	Trace
77	u2	206	Total	C	N	O	S	0	0
			1633	1025	322	281	5		

- Molecule 78 is a protein called Small ribosomal subunit protein eS10.

Mol	Chain	Residues	Atoms					AltConf	Trace
78	v2	95	Total	C	N	O	S	0	0
			800	522	142	131	5		

- Molecule 79 is a protein called Small ribosomal subunit protein uS17.

Mol	Chain	Residues	Atoms					AltConf	Trace
79	w2	150	Total	C	N	O	S	0	0
			1220	776	228	210	6		

- Molecule 80 is a protein called Small ribosomal subunit protein uS19.

Mol	Chain	Residues	Atoms					AltConf	Trace
80	x2	132	Total	C	N	O	S	0	0
			1081	688	203	183	7		

- Molecule 81 is a protein called Small ribosomal subunit protein uS9.

Mol	Chain	Residues	Atoms					AltConf	Trace
81	y2	142	Total	C	N	O	S	0	0
			1128	717	213	195	3		

- Molecule 82 is a protein called Small ribosomal subunit protein eS17.

Mol	Chain	Residues	Atoms					AltConf	Trace
82	z2	69	Total	C	N	O	S	0	0
			560	354	113	91	2		

- Molecule 83 is a RNA chain called 18S ribosomal RNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
83	m2	1635	Total	C	N	O	P	0	0
			34939	15614	6270	11420	1635		

- Molecule 84 is MAGNESIUM ION (CCD ID: MG) (formula: Mg) (labeled as "Ligand of Interest" by depositor).

Mol	Chain	Residues	Atoms		AltConf
84	A2	82	Total	Mg	0
			82	82	
84	Bv	2	Total	Mg	0
			2	2	
84	H1	1	Total	Mg	0
			1	1	
84	J2	1	Total	Mg	0
			1	1	
84	P2	1	Total	Mg	0
			1	1	
84	d2	1	Total	Mg	0
			1	1	
84	m2	30	Total	Mg	0
			30	30	

- Molecule 85 is ZINC ION (CCD ID: ZN) (formula: Zn) (labeled as "Ligand of Interest" by depositor).

Mol	Chain	Residues	Atoms		AltConf
85	F3	1	Total	Zn	0
			1	1	
85	H3	1	Total	Zn	0
			1	1	
85	d2	1	Total	Zn	0
			1	1	
85	g2	1	Total	Zn	0
			1	1	
85	i2	1	Total	Zn	0
			1	1	

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Mol	Chain	Residues	Atoms		AltConf
85	j2	1	Total	Zn	0
			1	1	

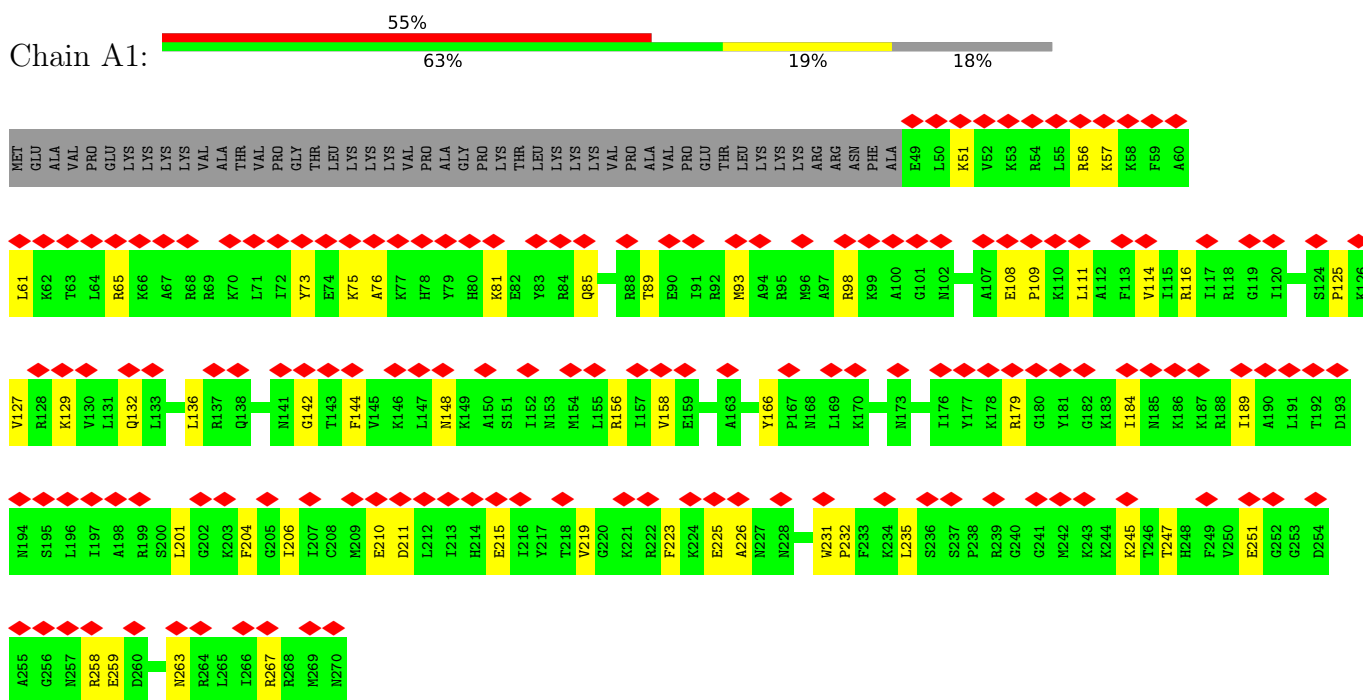
- Molecule 86 is water.

Mol	Chain	Residues	Atoms		AltConf
86	B1	1	Total	O	0
			1	1	

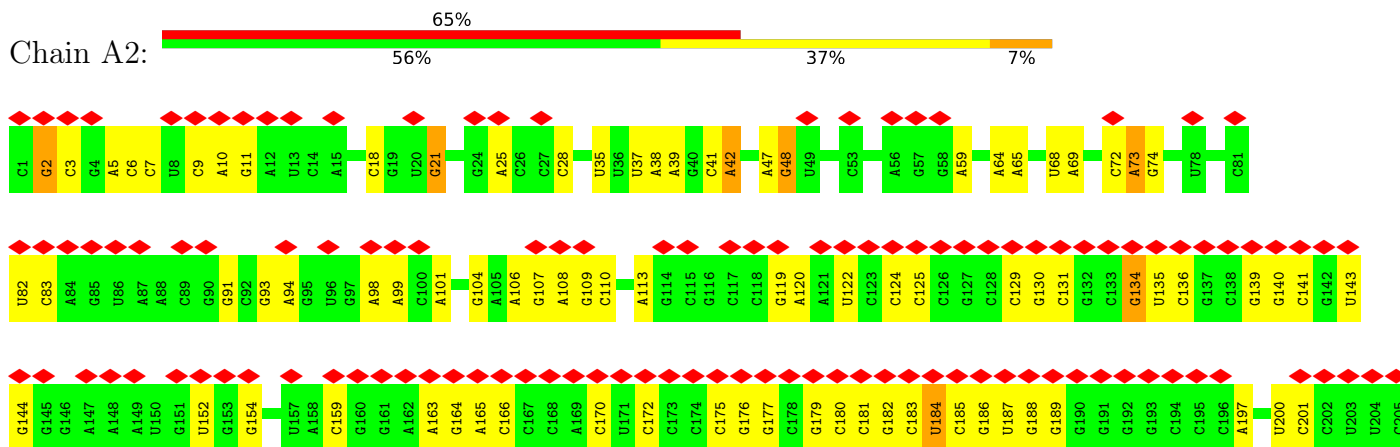
3 Residue-property plots

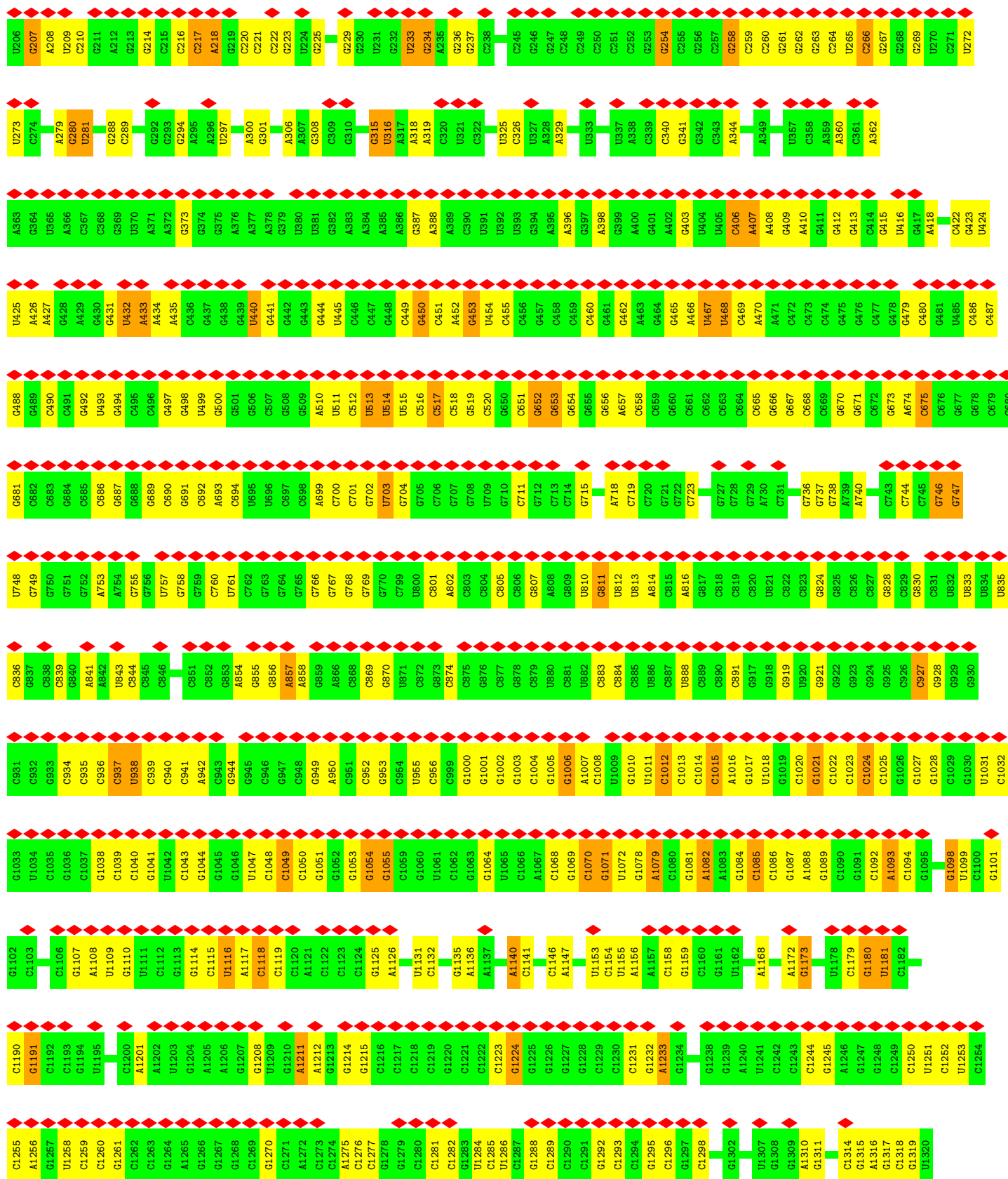
These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and atom inclusion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

- Molecule 1: Large ribosomal subunit protein uL30

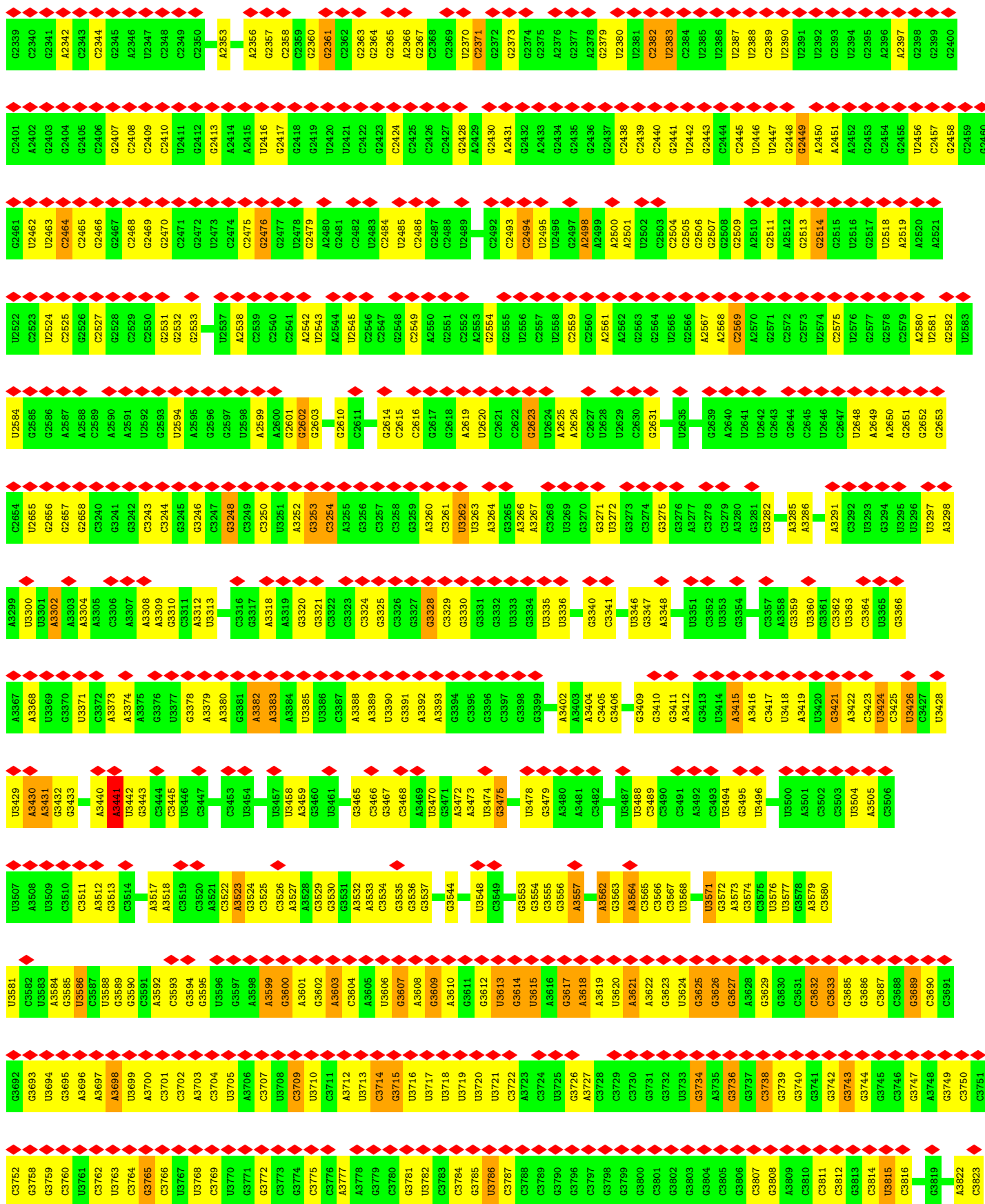


- Molecule 2: 28S ribosomal RNA

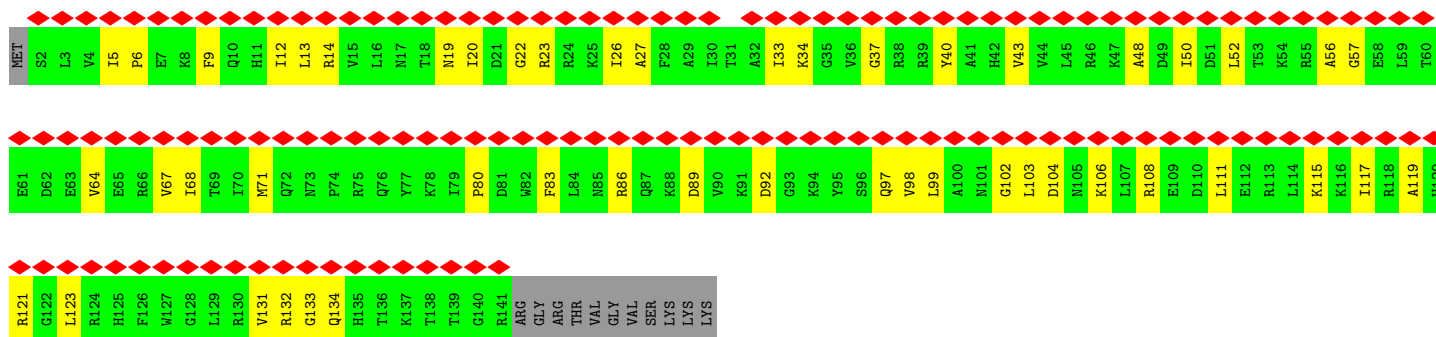




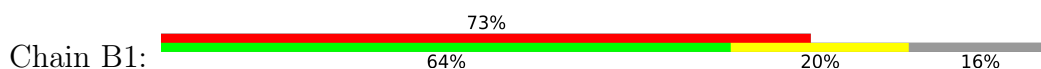




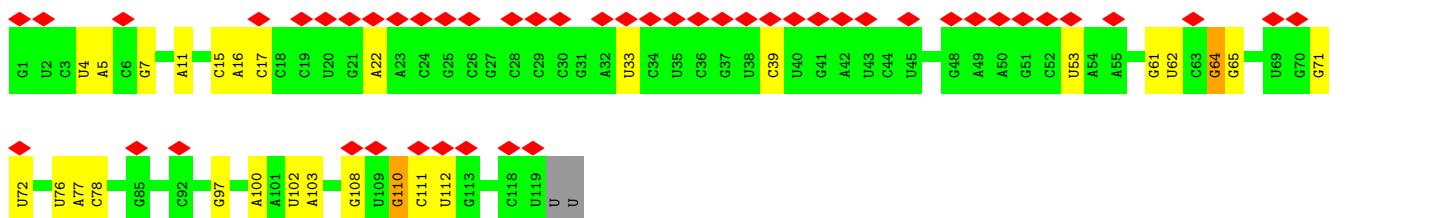
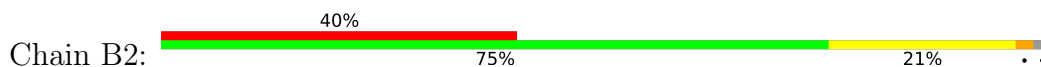




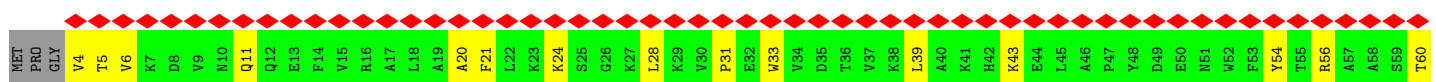
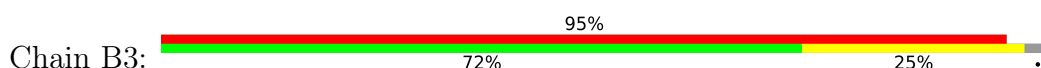
• Molecule 4: Large ribosomal subunit protein eL8

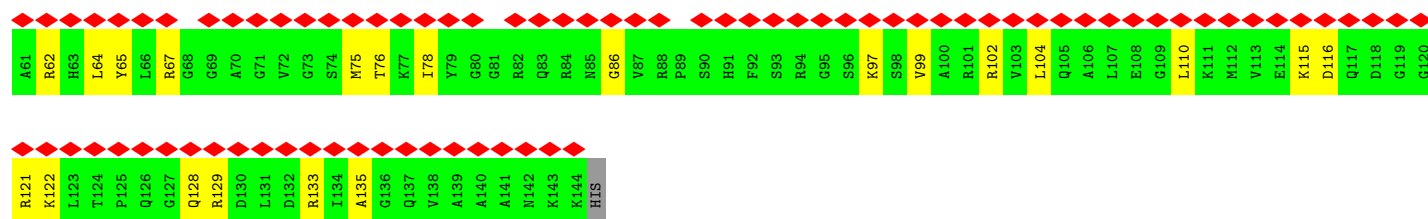


• Molecule 5: 5S ribosomal RNA



• Molecule 6: Small ribosomal subunit protein eS19





• Molecule 7: transfer RNA



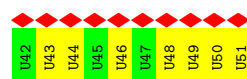
• Molecule 7: transfer RNA



• Molecule 7: transfer RNA

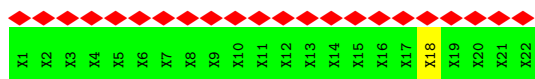


• Molecule 8: messenger RNA

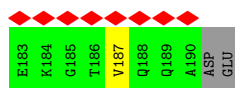
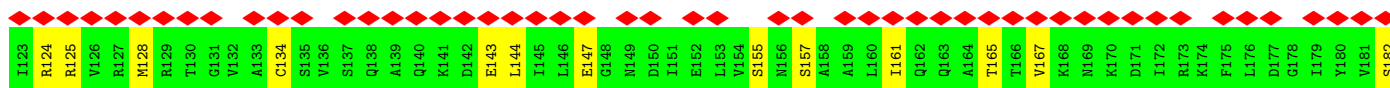
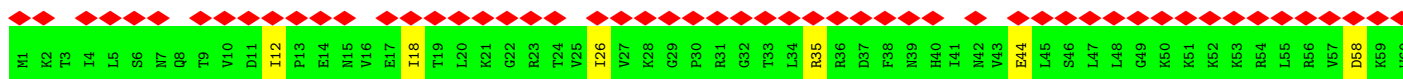
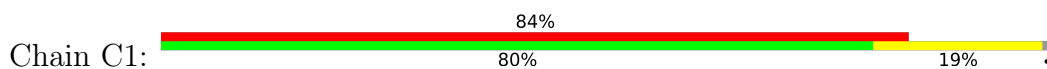


• Molecule 9: Nascent protein chain

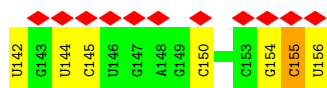
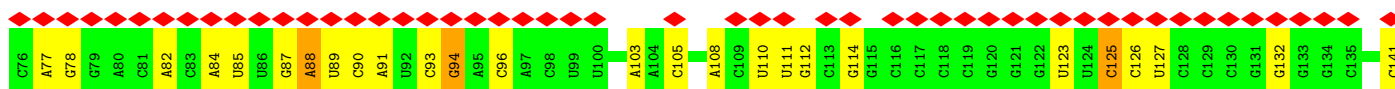
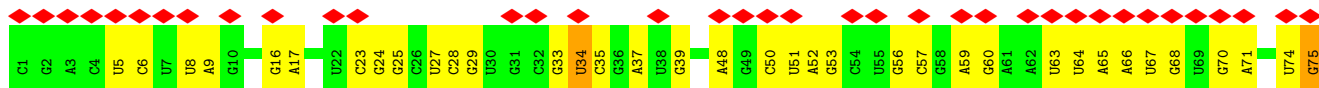




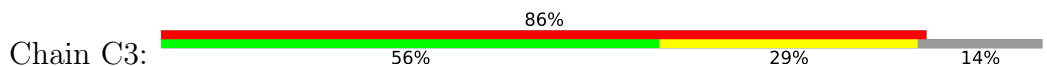
• Molecule 10: Large ribosomal subunit protein uL6



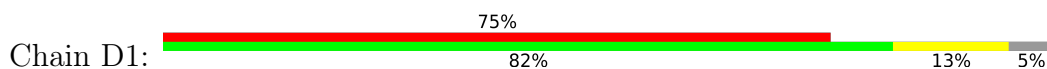
• Molecule 11: 5.8S ribosomal RNA

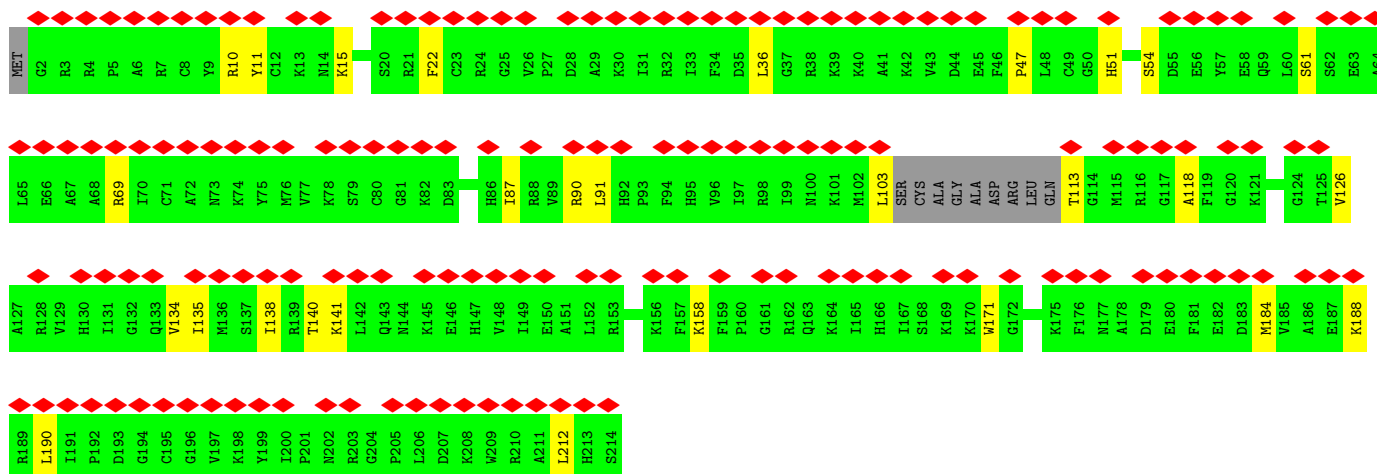


• Molecule 12: Small ribosomal subunit protein uS10

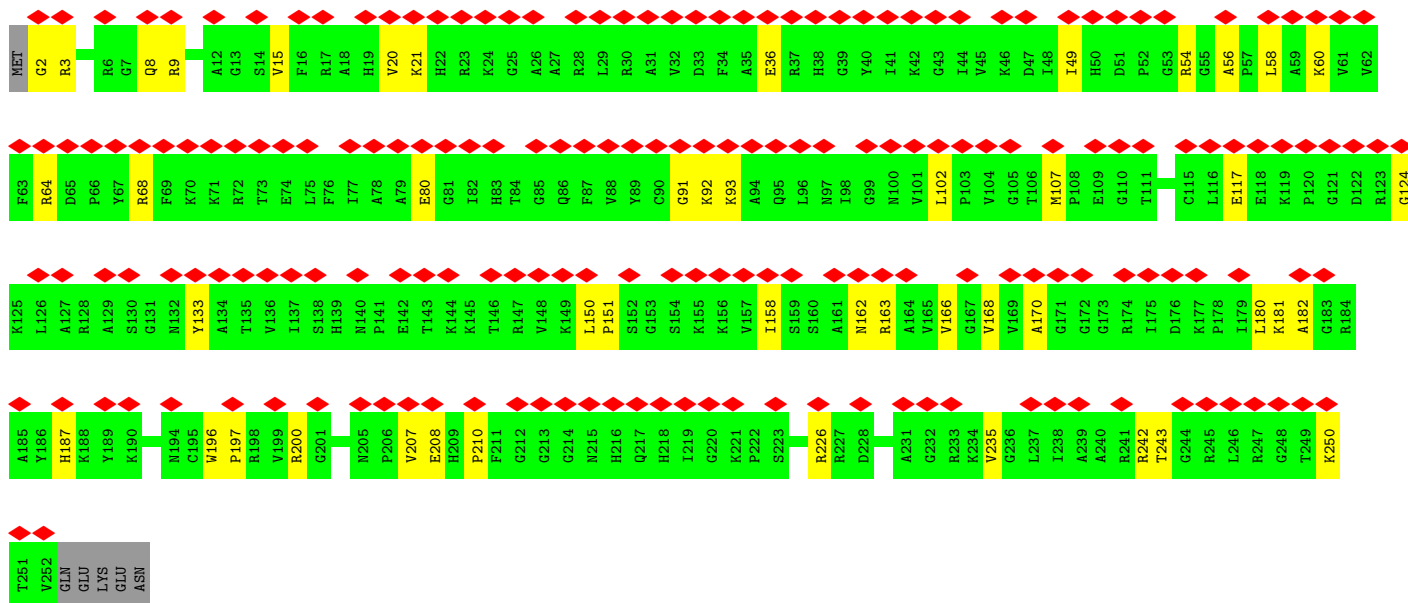
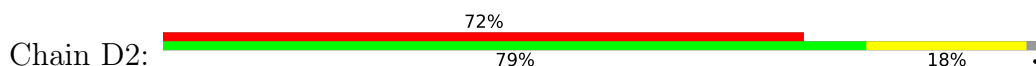


• Molecule 13: Large ribosomal subunit protein uL16

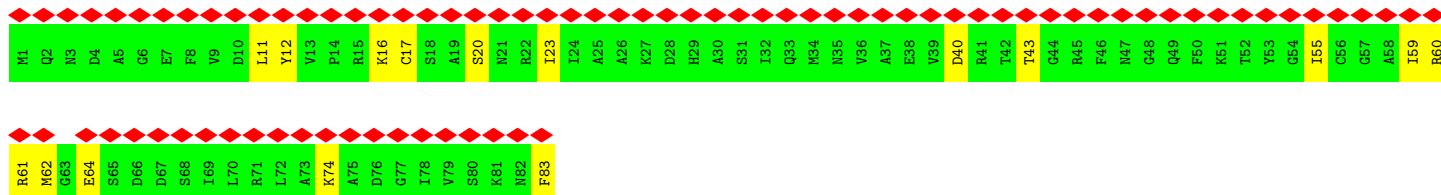
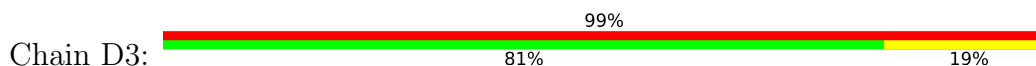




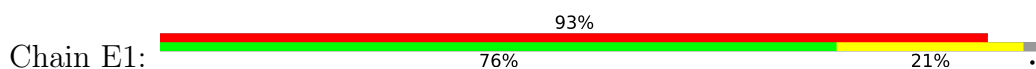
- Molecule 14: Large ribosomal subunit protein uL2

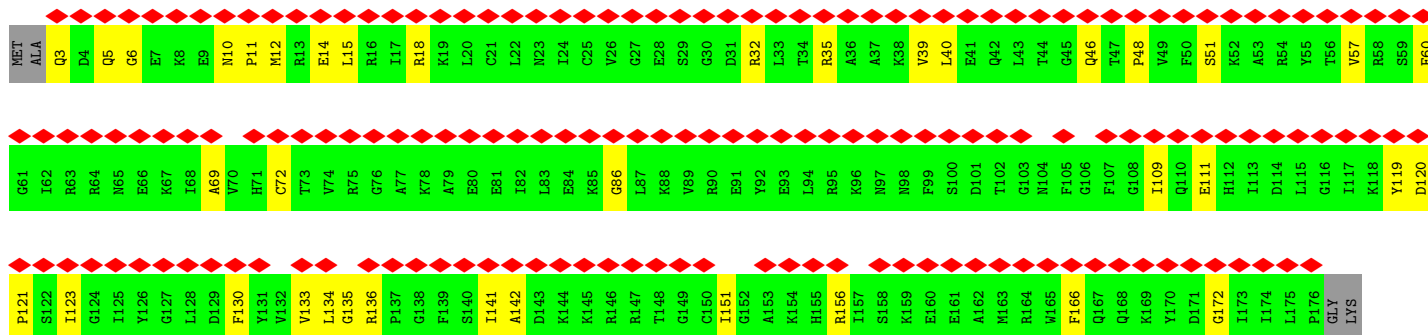


- Molecule 15: Small ribosomal subunit protein eS21

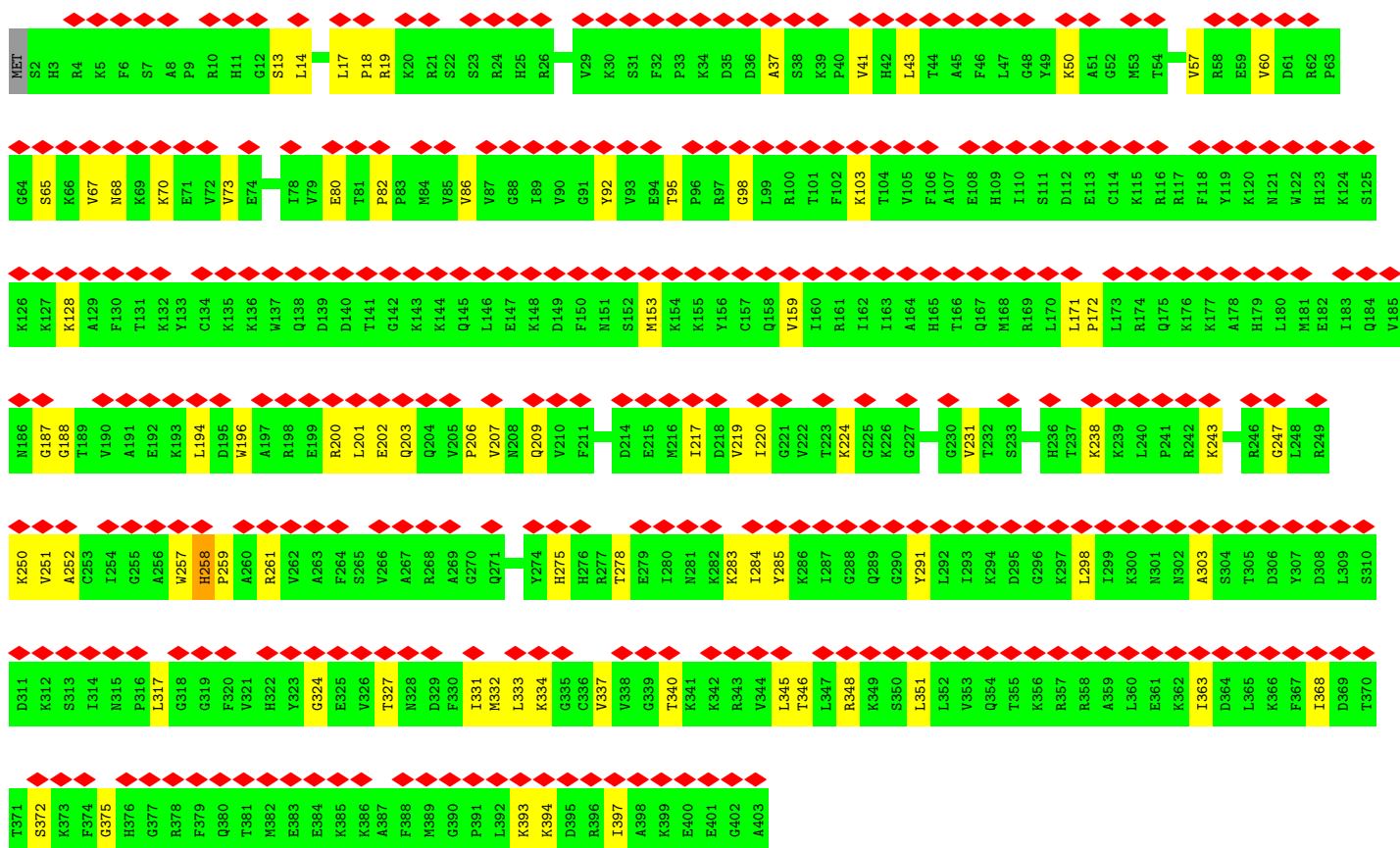
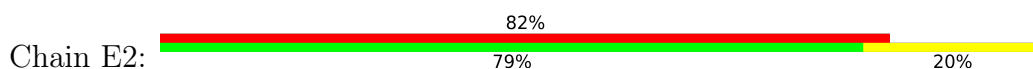


- Molecule 16: Large ribosomal subunit protein uL5

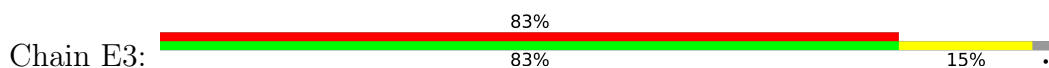




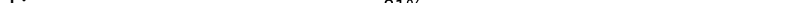
- Molecule 17: Large ribosomal subunit protein uL3

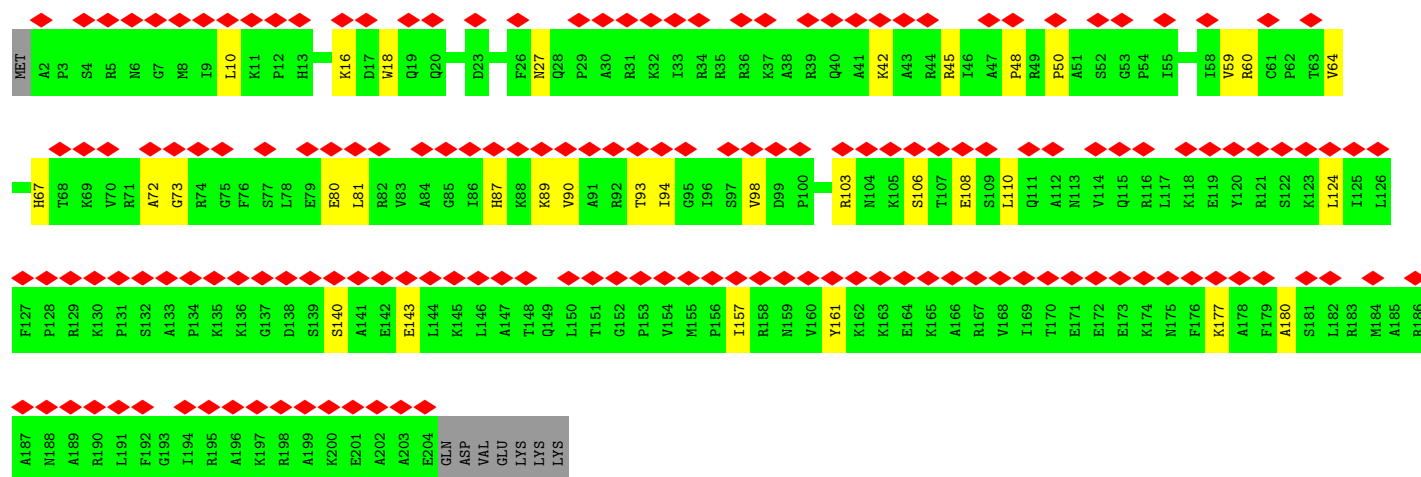


- Molecule 18: Small ribosomal subunit protein uS12

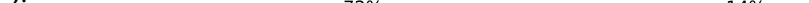


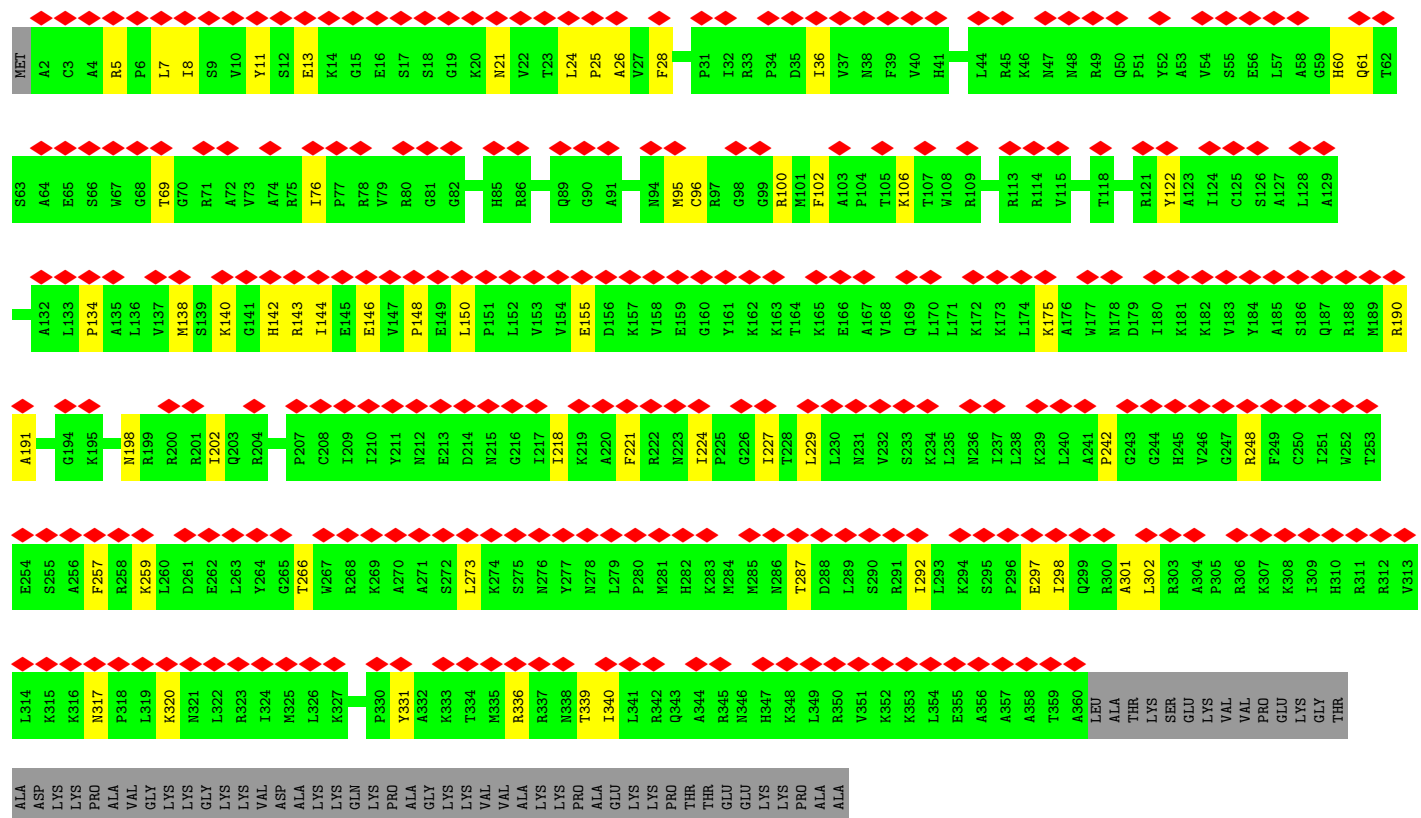
- Molecule 19: Large ribosomal subunit protein eL13

Chain F1: 



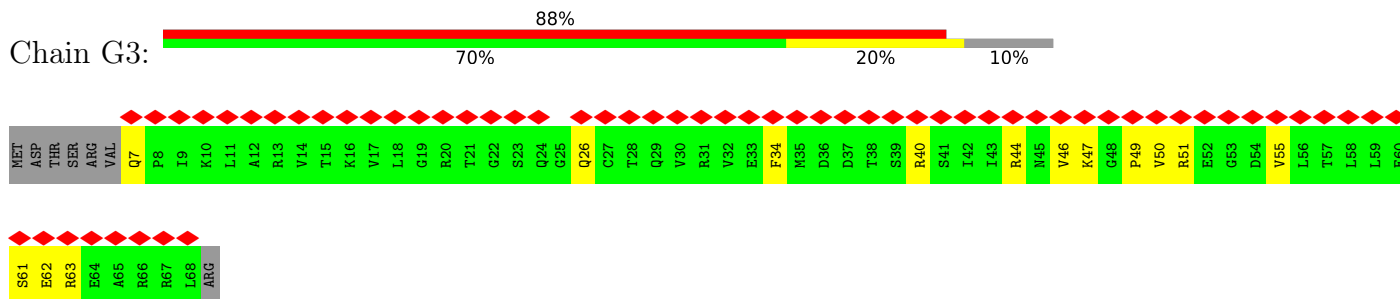
- Molecule 20: Large ribosomal subunit protein uL4

Chain F2: 



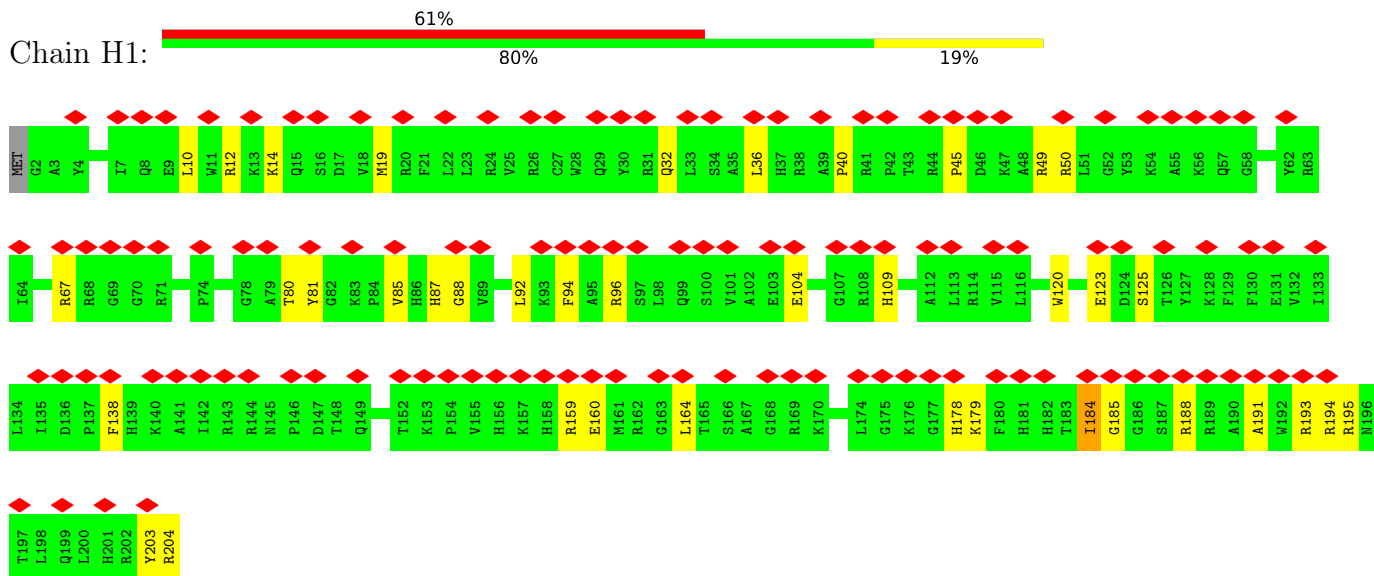
- Molecule 24: Small ribosomal subunit protein eS28

Chain G3:



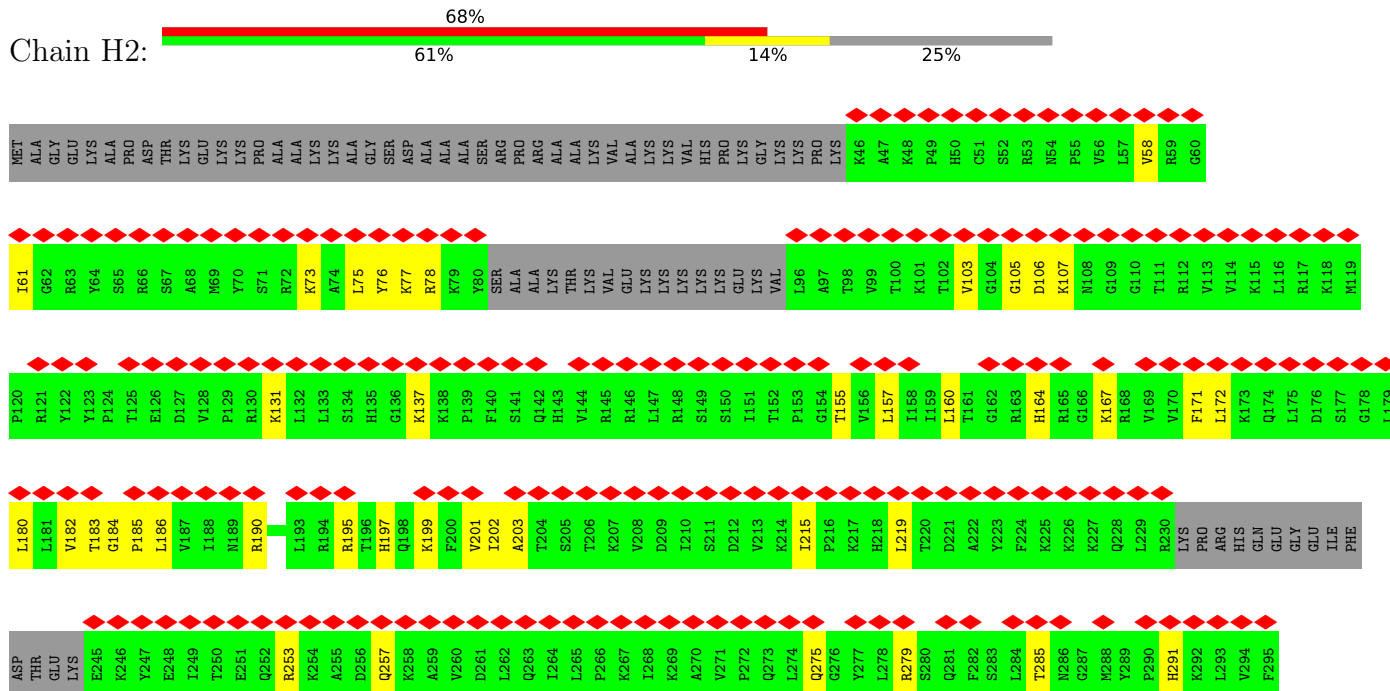
- Molecule 25: Large ribosomal subunit protein eL15

Chain H1:



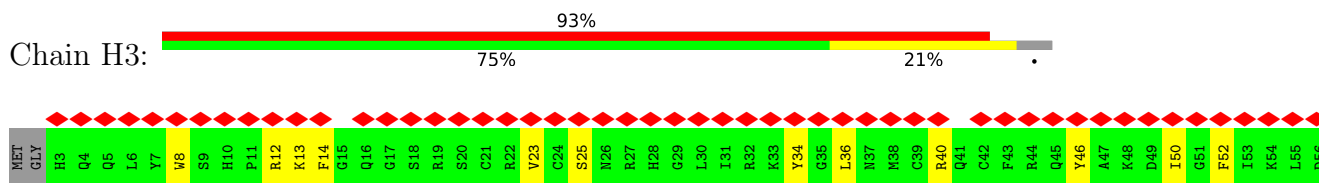
- Molecule 26: Large ribosomal subunit protein eL6

Chain H2:



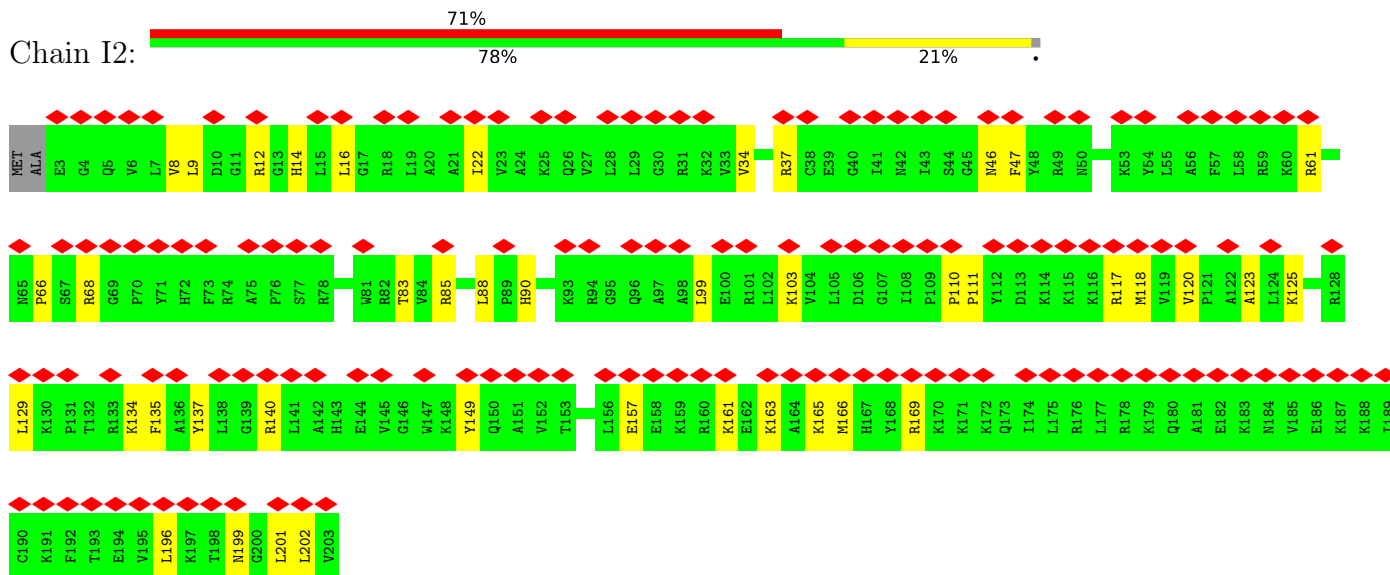
- Molecule 27: Small ribosomal subunit protein uS14

Chain H3:



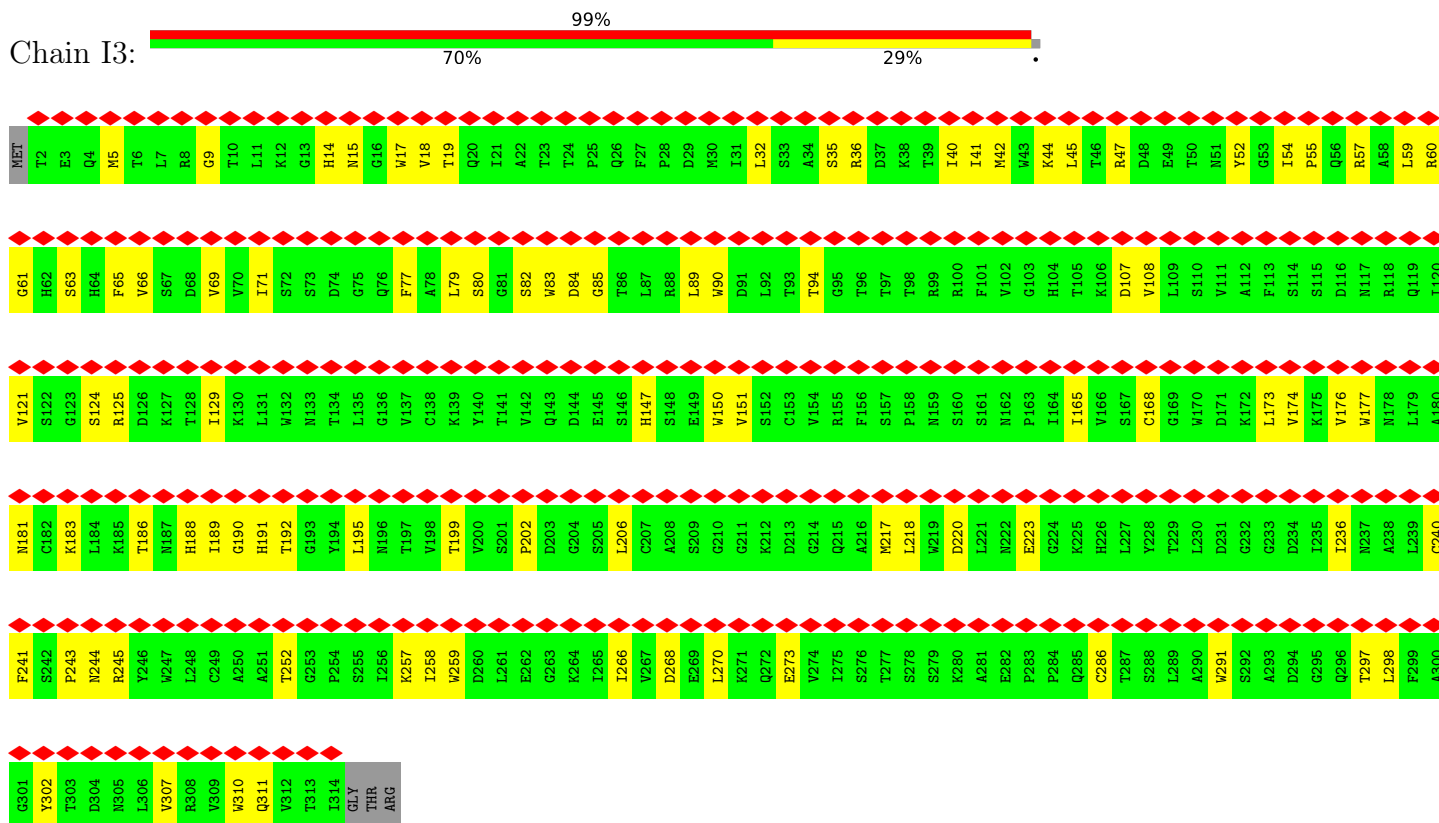
- Molecule 28: Large ribosomal subunit protein uL13

Chain I2:

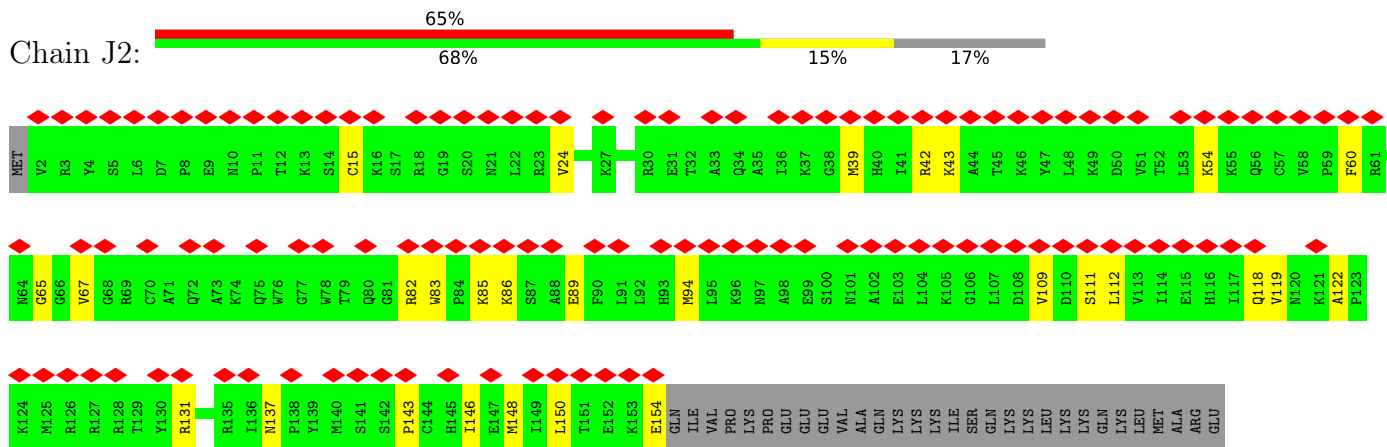


- Molecule 29: Small ribosomal subunit protein RACK1

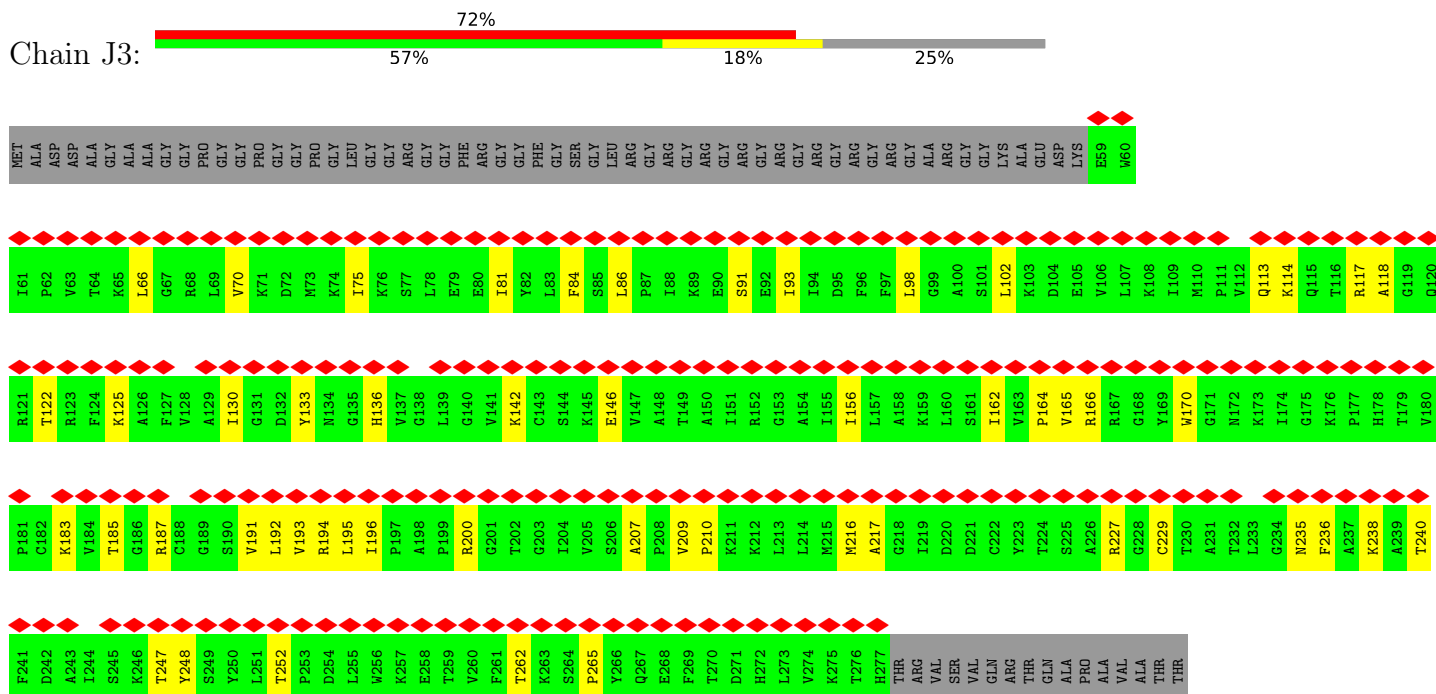
Chain I3:



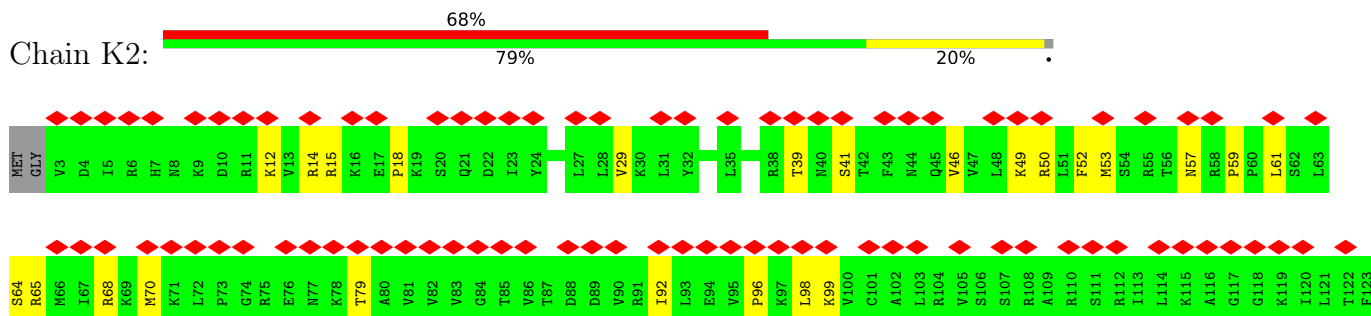
Chain J2:



Chain J3:

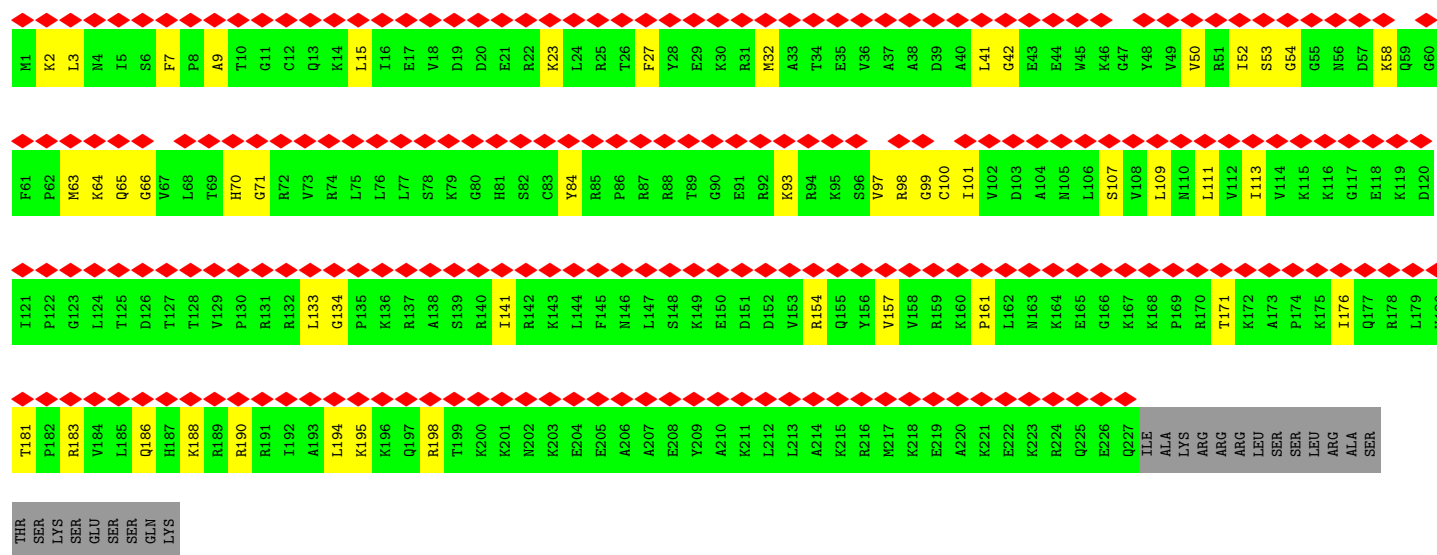
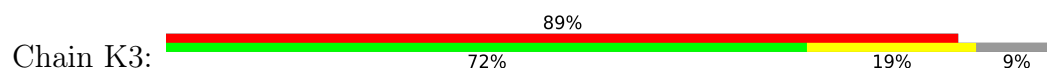


Chain K2:

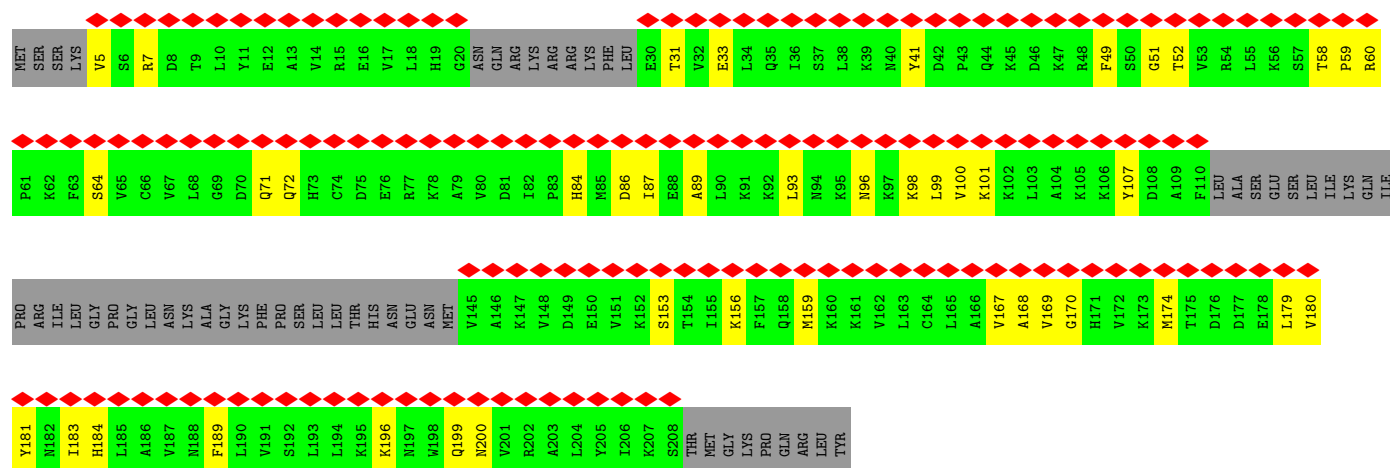
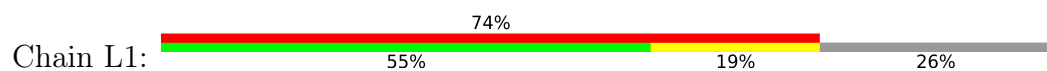




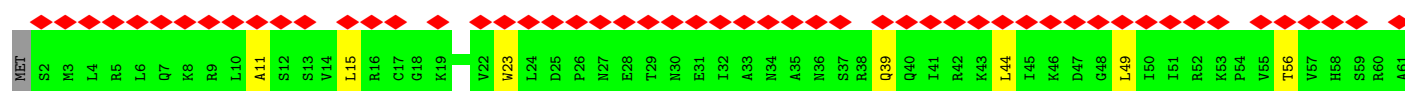
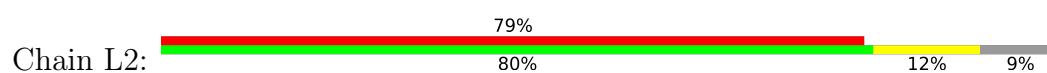
- Molecule 33: Small ribosomal subunit protein eS6

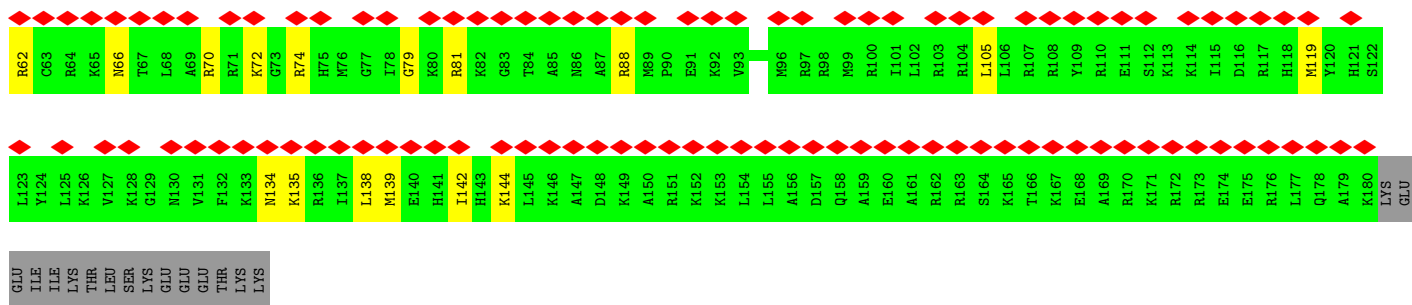


- Molecule 34: Large ribosomal subunit protein uL1

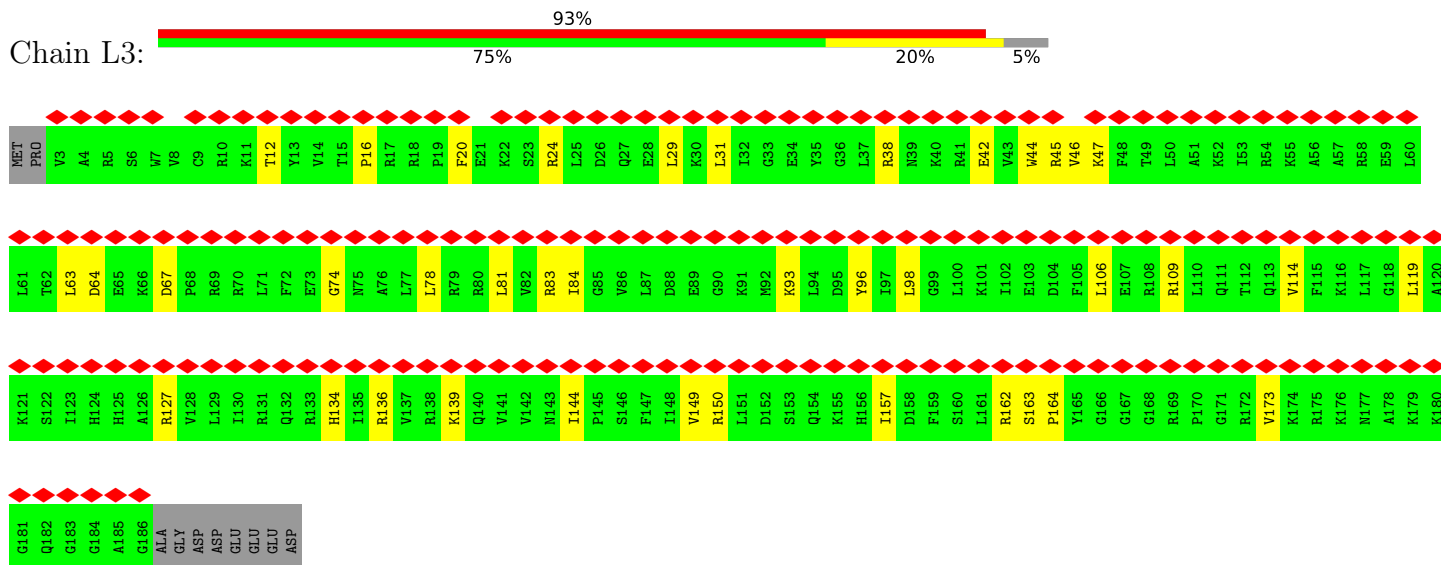


- Molecule 35: Large ribosomal subunit protein eL19

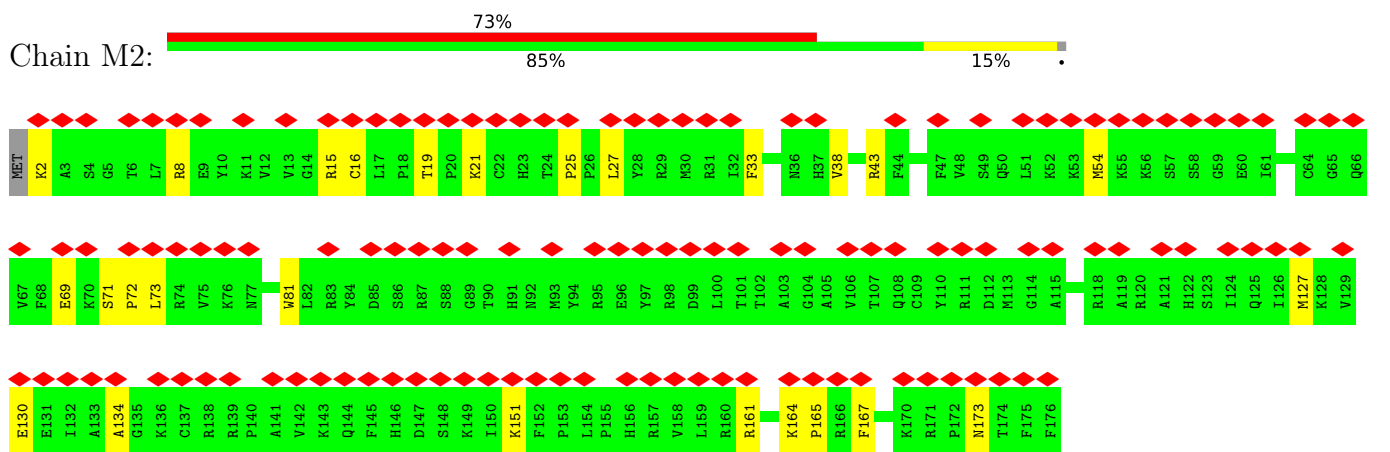




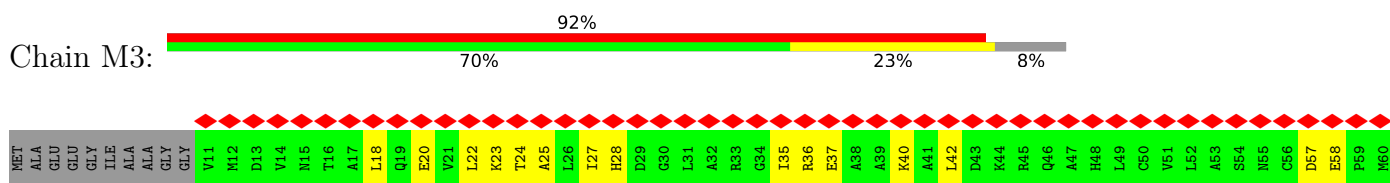
• Molecule 36: Small ribosomal subunit protein uS4



• Molecule 37: Large ribosomal subunit protein eL20

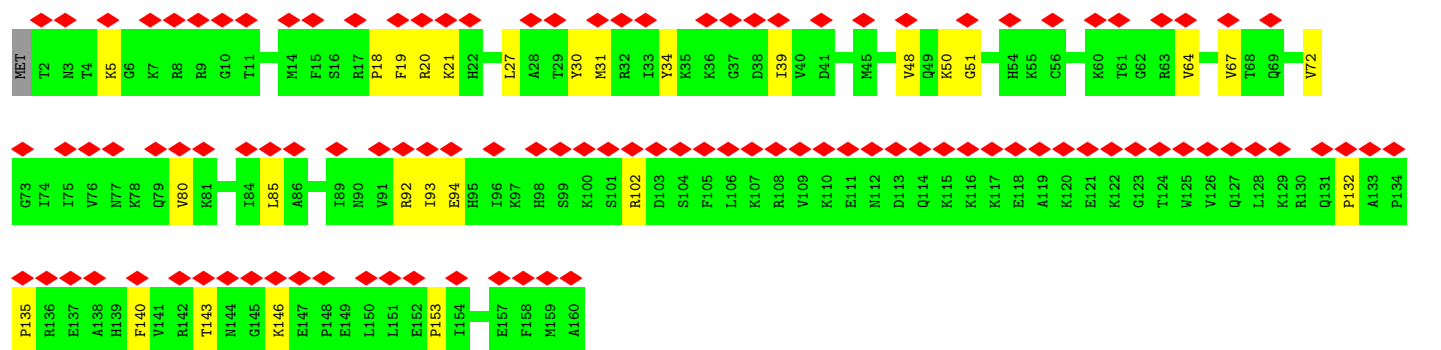
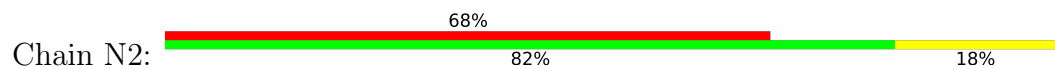


• Molecule 38: Small ribosomal subunit protein eS12

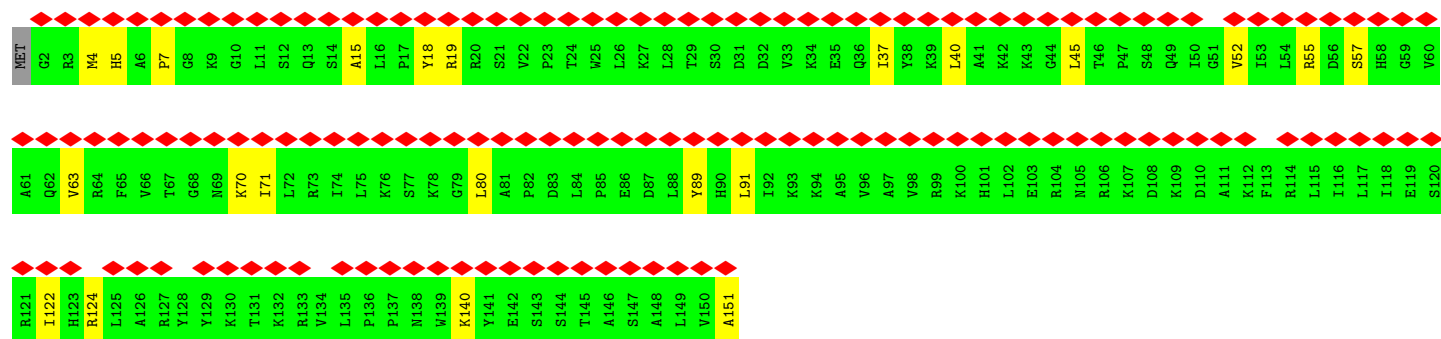
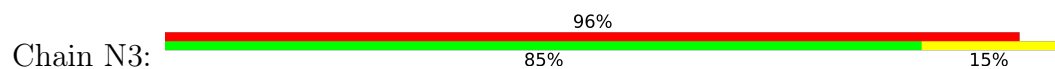




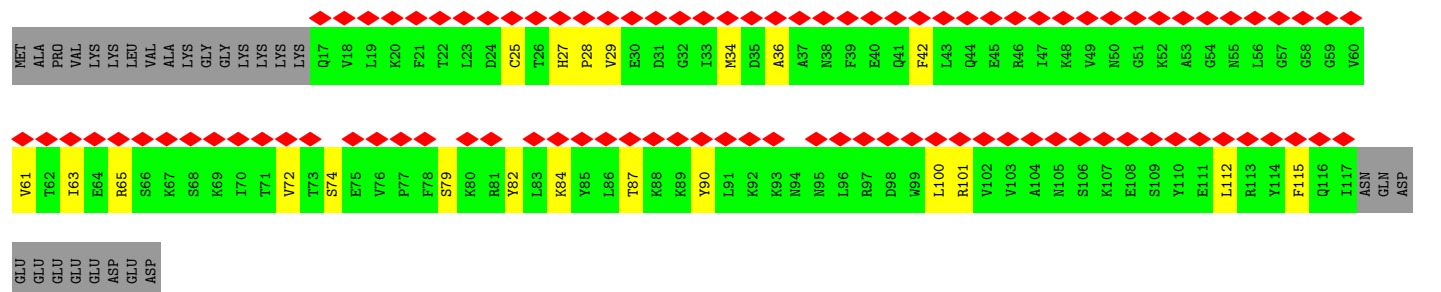
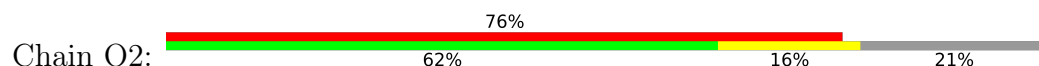
• Molecule 39: Large ribosomal subunit protein eL21



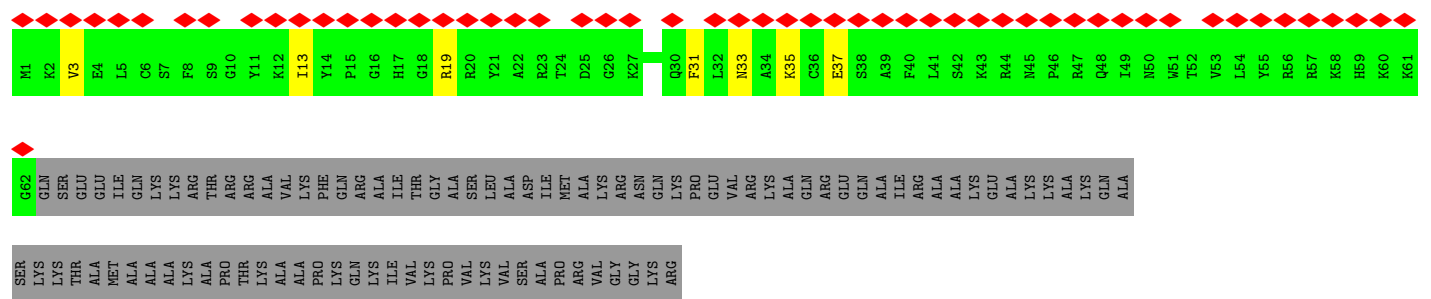
• Molecule 40: Small ribosomal subunit protein uS15



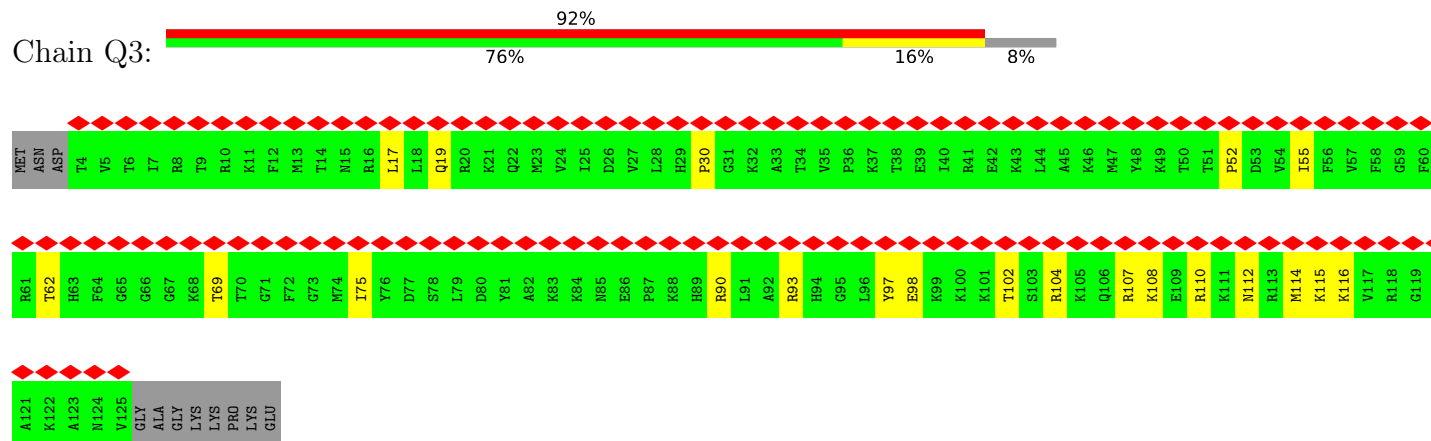
• Molecule 41: Large ribosomal subunit protein eL22



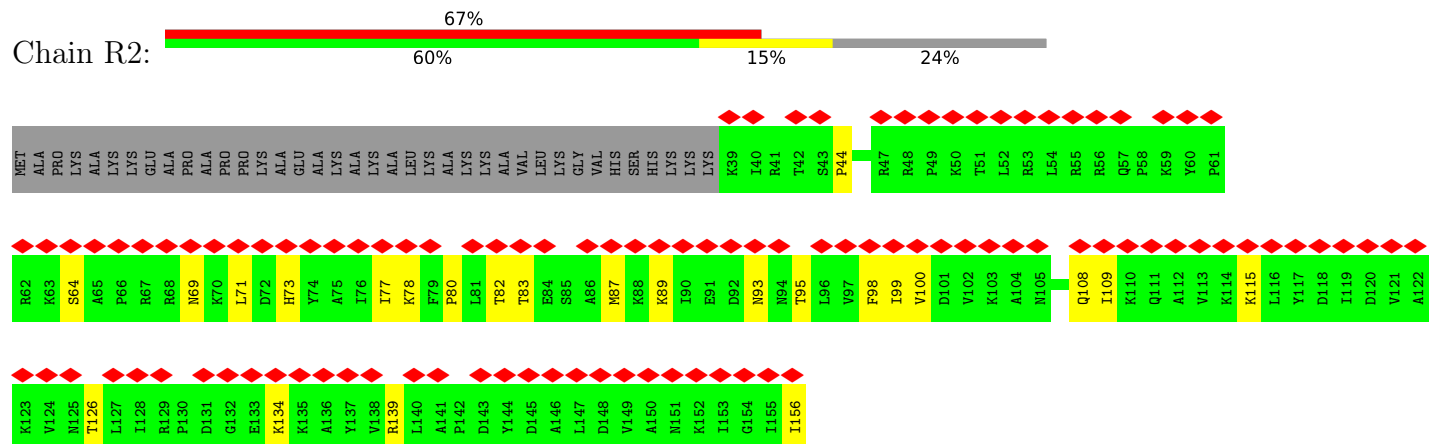




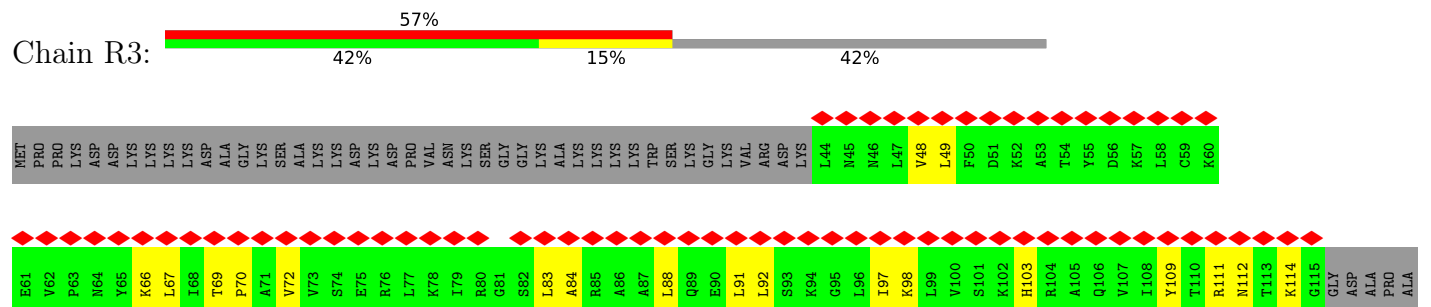
- Molecule 46: Small ribosomal subunit protein eS24



- Molecule 47: Large ribosomal subunit protein uL23

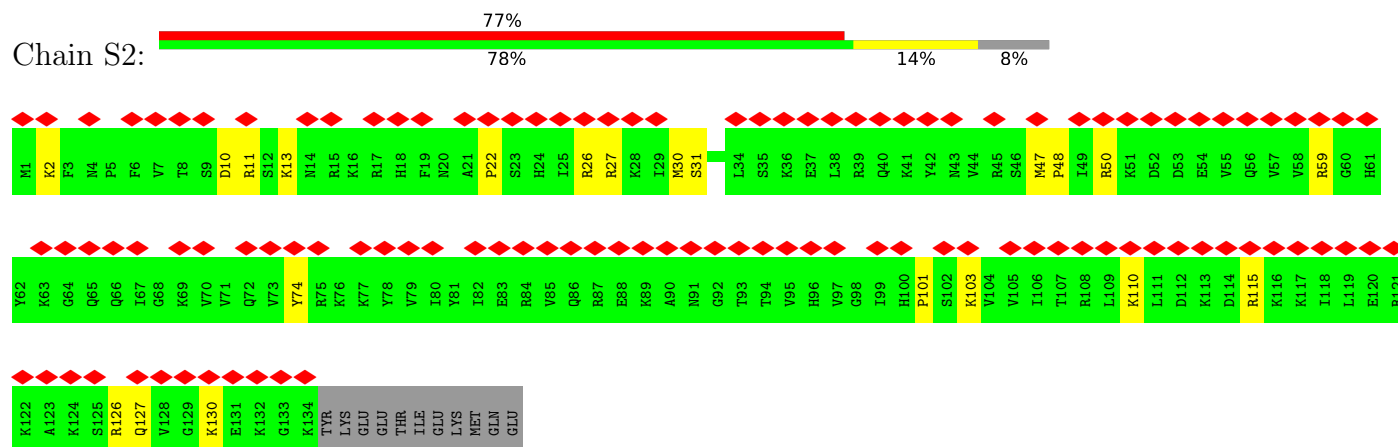


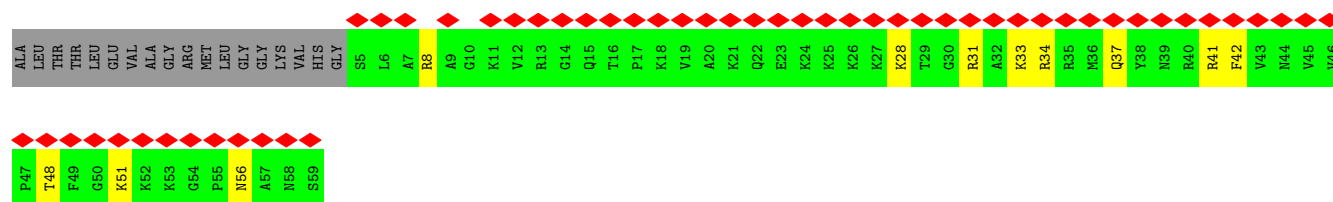
- Molecule 48: Small ribosomal subunit protein eS25



ALA
GLY
GLU
ASP
ALA

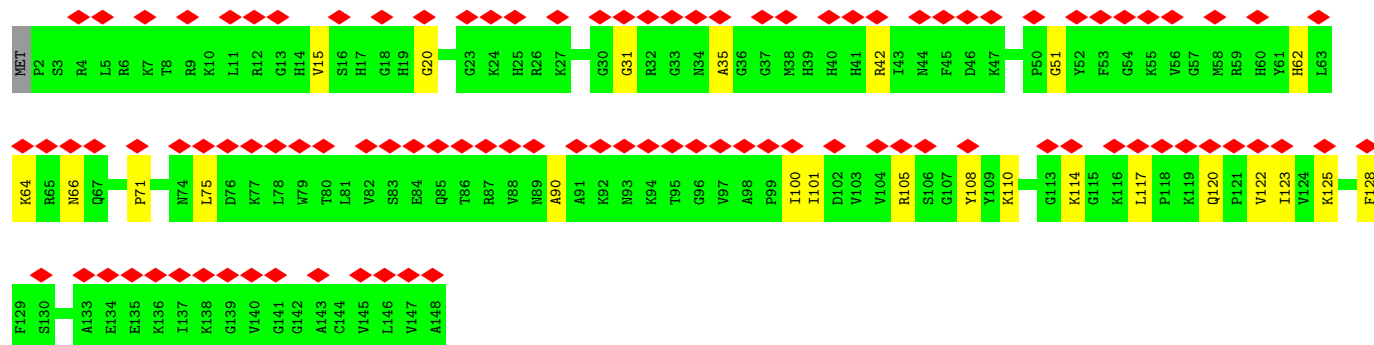
• Molecule 49: Large ribosomal subunit protein uL24





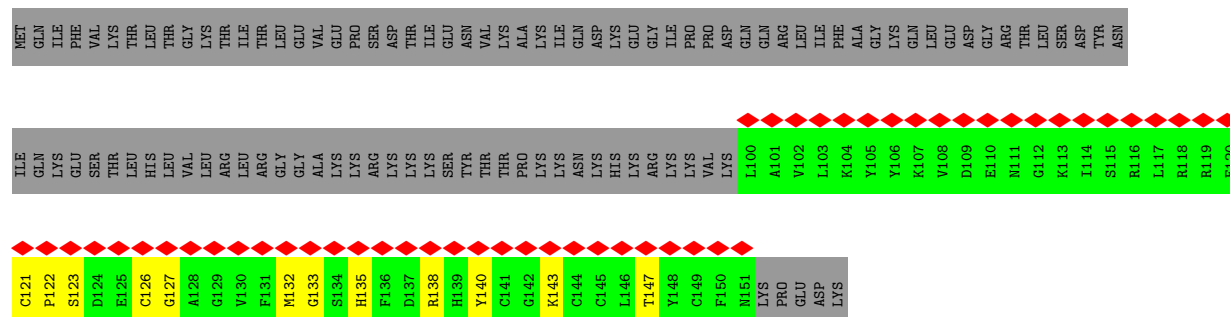
- Molecule 53: Large ribosomal subunit protein uL15

Chain U2: 68% 83% 16%



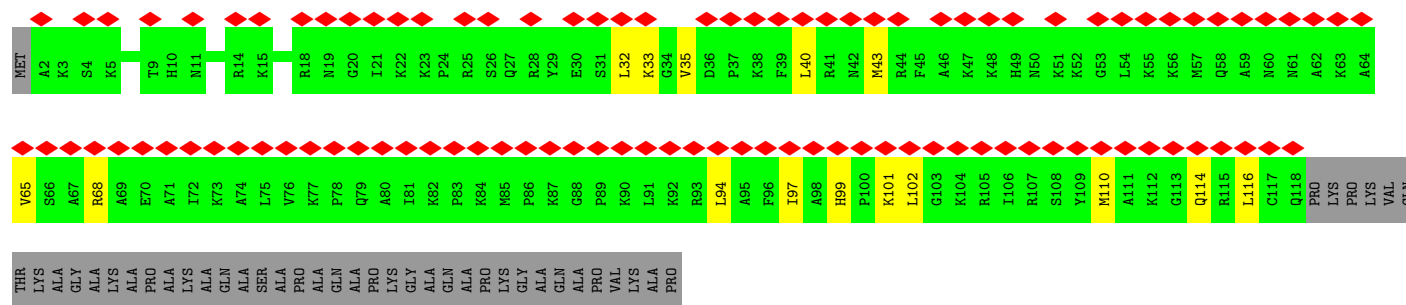
- Molecule 54: Ubiquitin-ribosomal protein eS31 fusion protein

Chain U3: 33% 26% 8% 67%

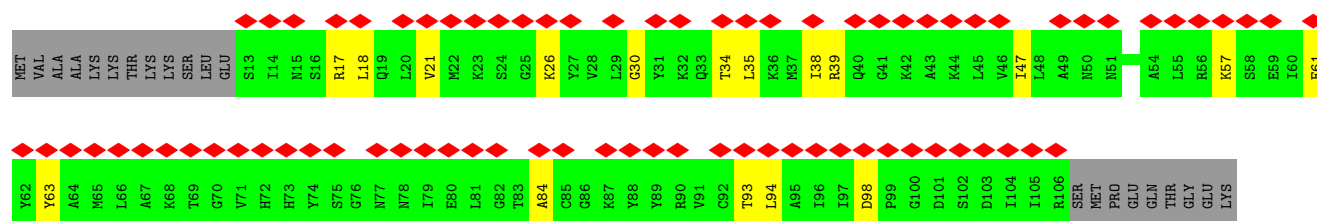


- Molecule 55: Large ribosomal subunit protein eL29

Chain V2: 62% 64% 9% 27%



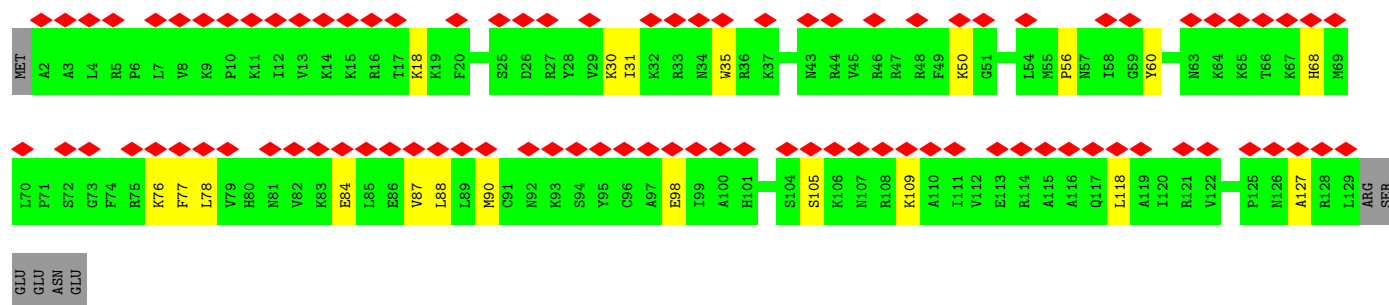
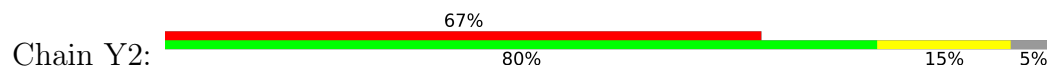
- Molecule 56: Large ribosomal subunit protein eL30



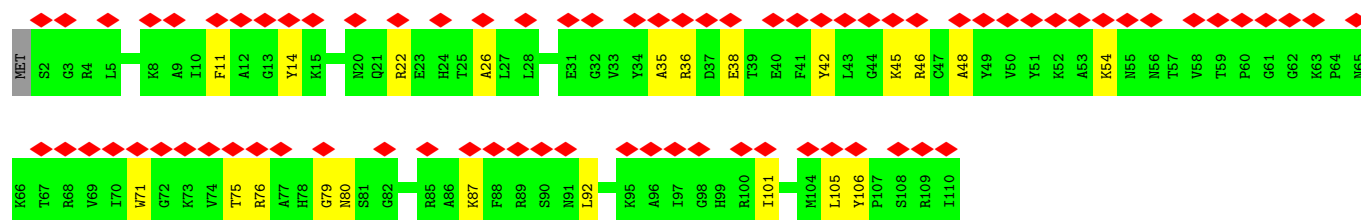
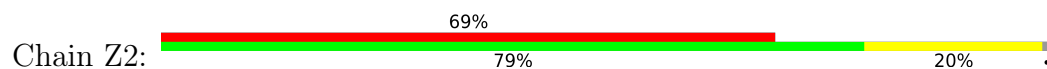
• Molecule 57: Large ribosomal subunit protein eL31




• Molecule 58: Large ribosomal subunit protein eL32

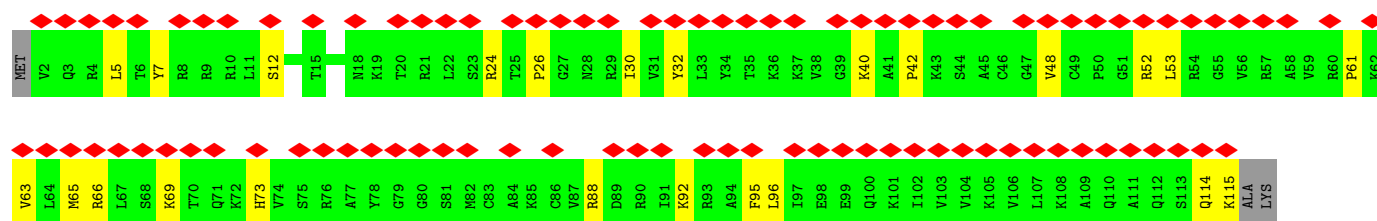


• Molecule 59: Large ribosomal subunit protein eL33




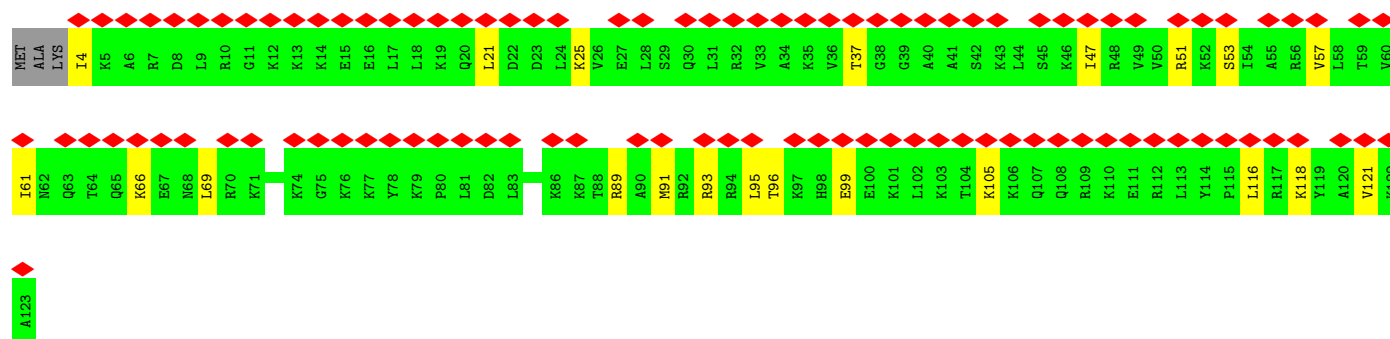
• Molecule 60: Large ribosomal subunit protein eL34

Chain a2: 




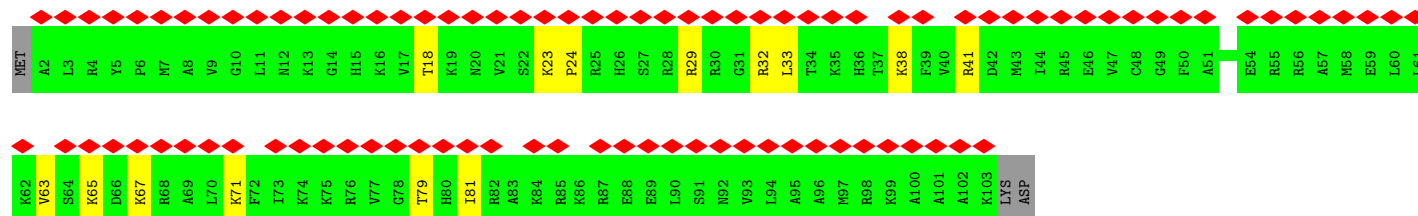
- Molecule 61: Large ribosomal subunit protein uL29

Chain b2: 




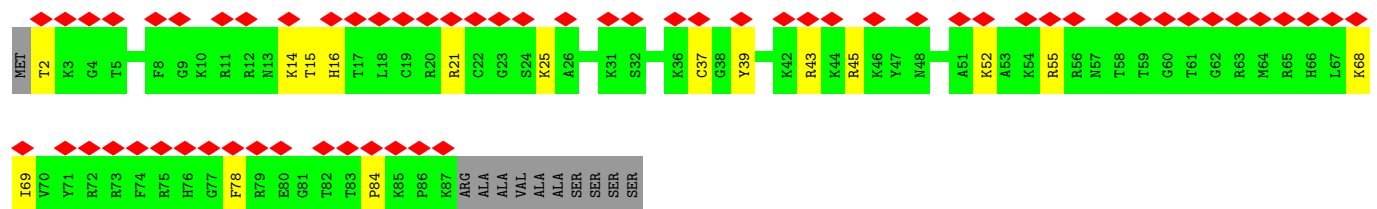
- Molecule 62: Large ribosomal subunit protein eL36

Chain c2: 



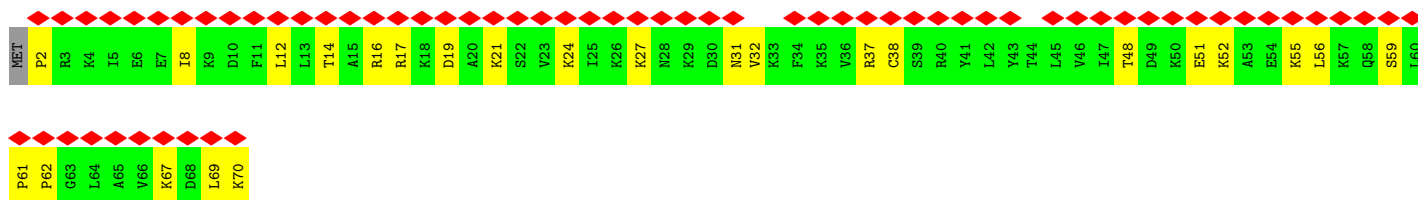
- Molecule 63: Large ribosomal subunit protein eL37

Chain d2: 

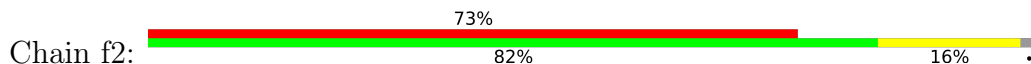


- Molecule 64: Large ribosomal subunit protein eL38

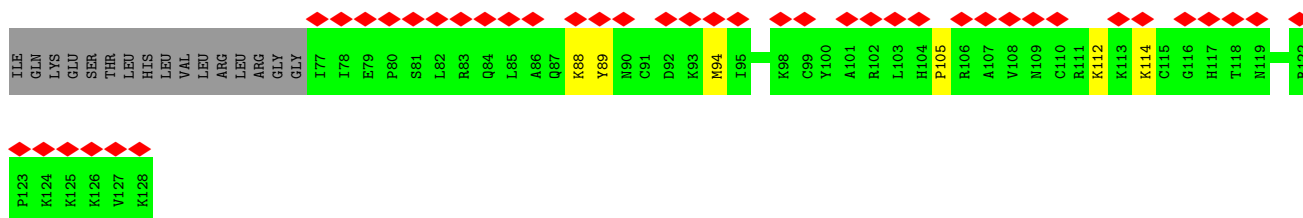
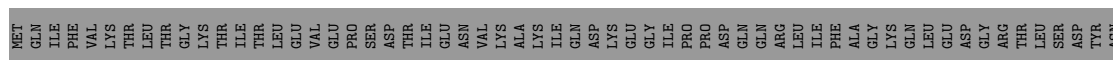
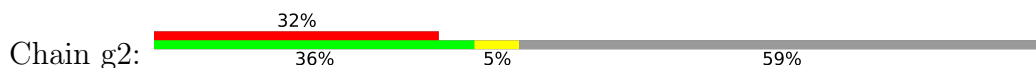
Chain e2: 



- Molecule 65: Large ribosomal subunit protein eL39



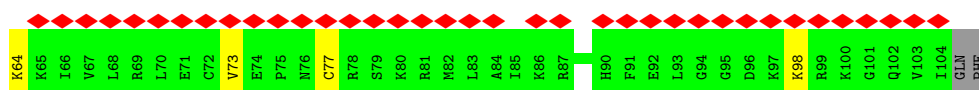
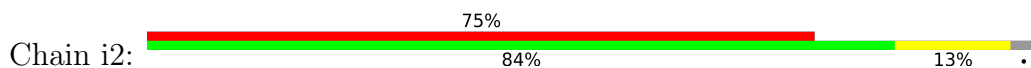
- Molecule 66: Ubiquitin-ribosomal protein eL40 fusion protein



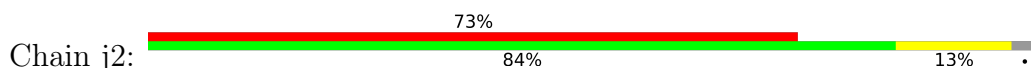
- Molecule 67: 60S ribosomal protein L41

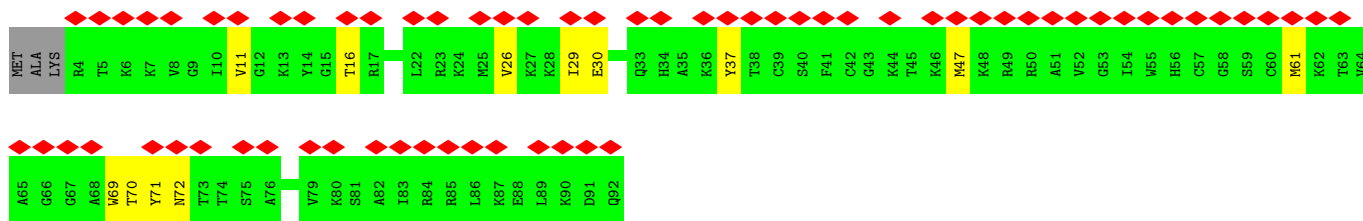


- Molecule 68: Large ribosomal subunit protein eL42

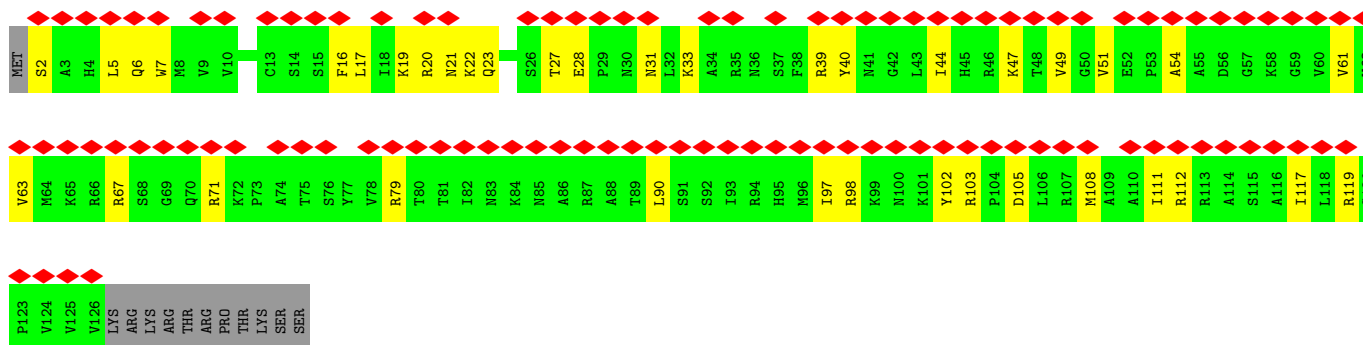
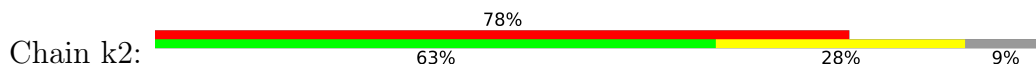


- Molecule 69: Large ribosomal subunit protein eL43

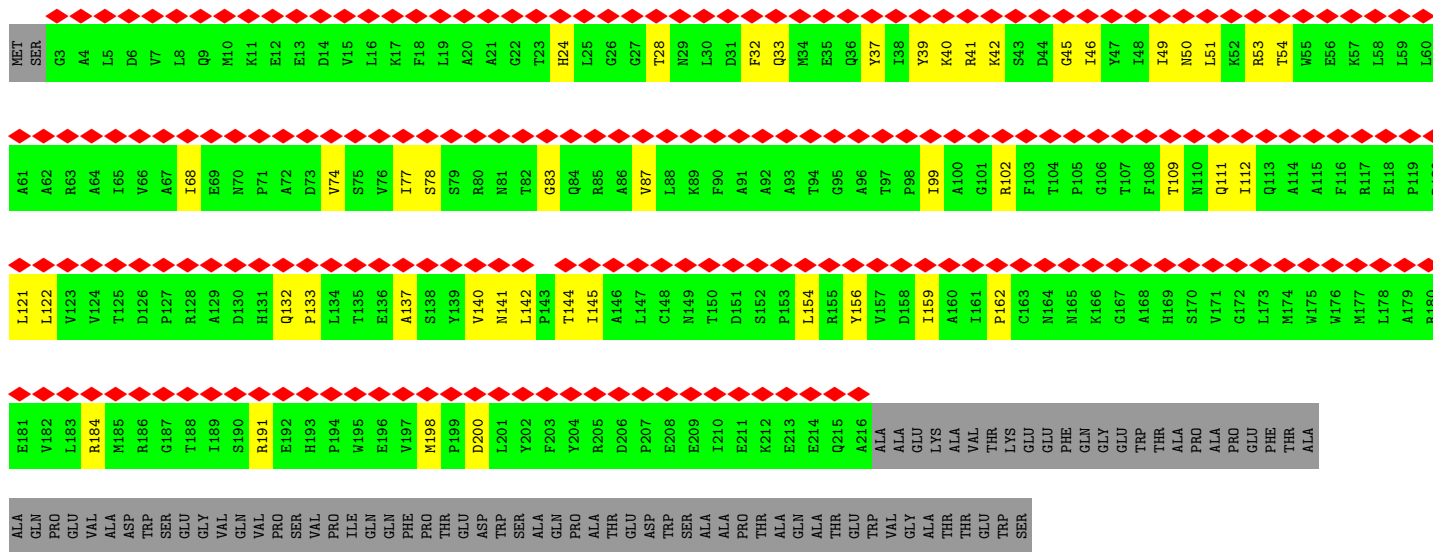
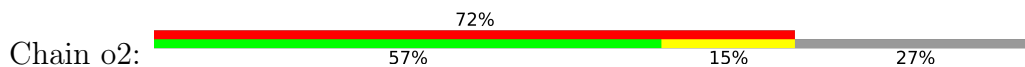




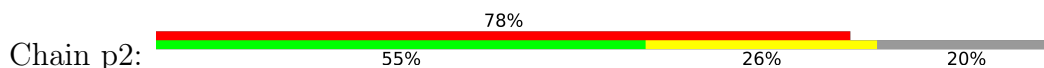
• Molecule 70: Large ribosomal subunit protein eL28

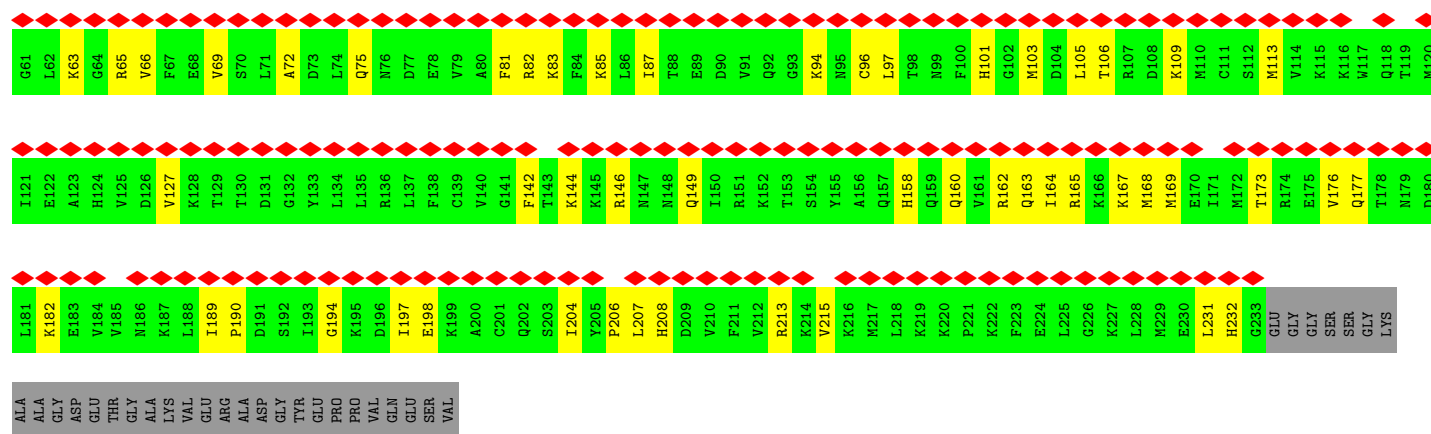


• Molecule 71: Small ribosomal subunit protein uS2

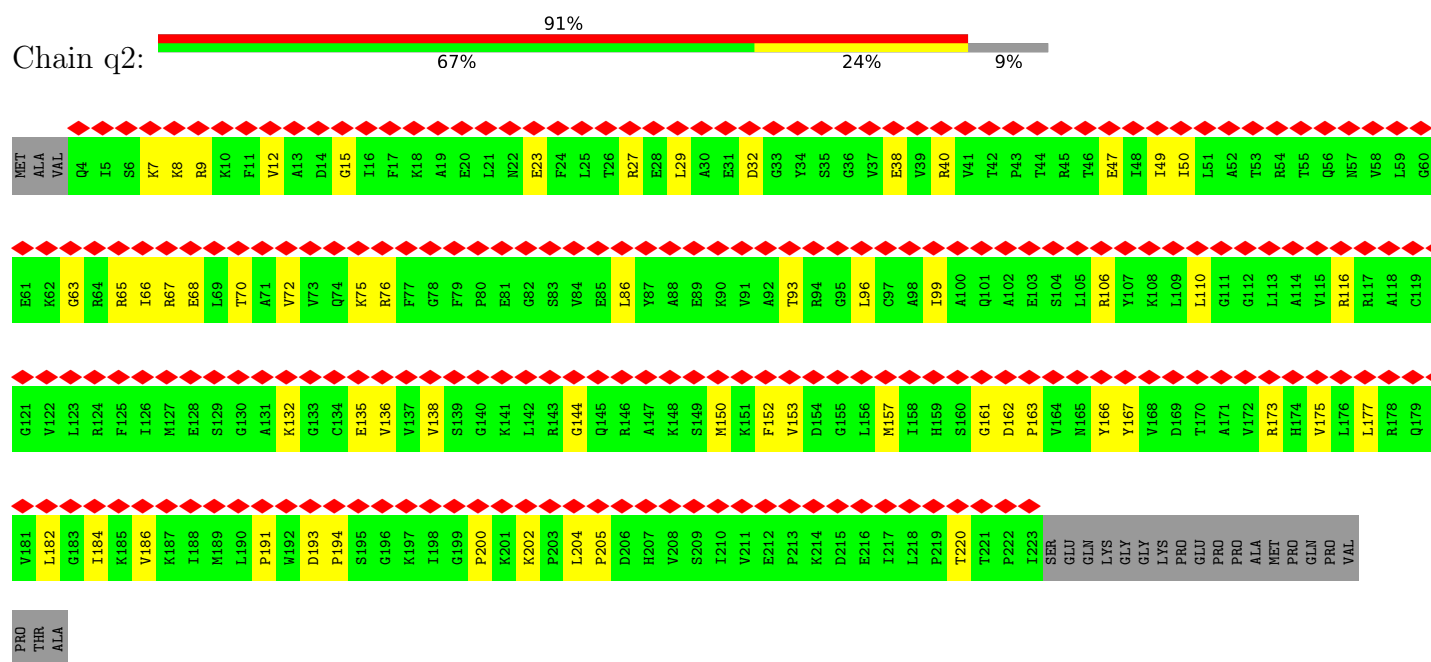


• Molecule 72: 40S ribosomal protein S3a

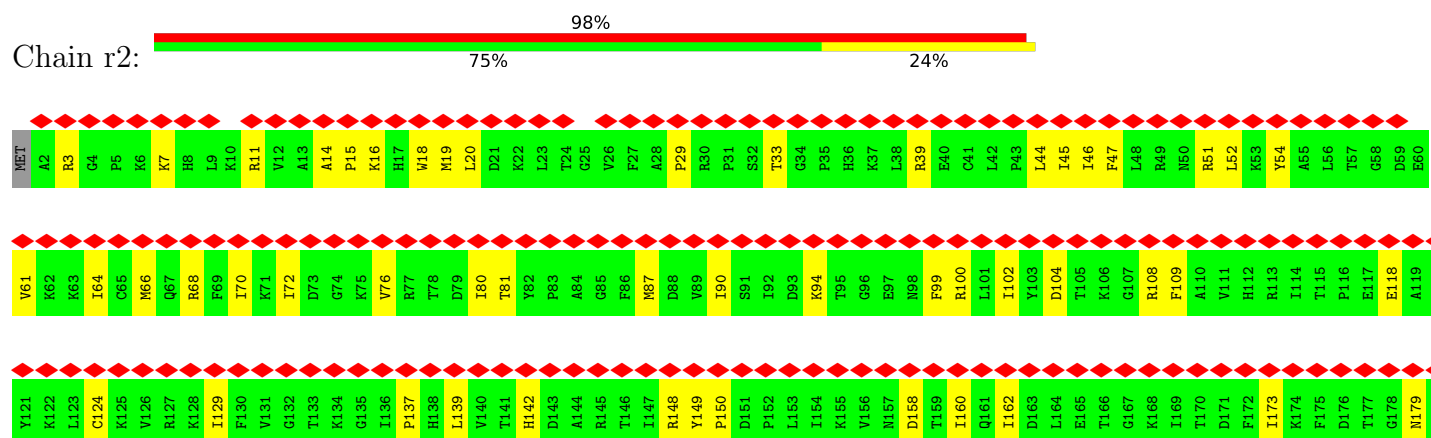




• Molecule 73: Small ribosomal subunit protein uS3

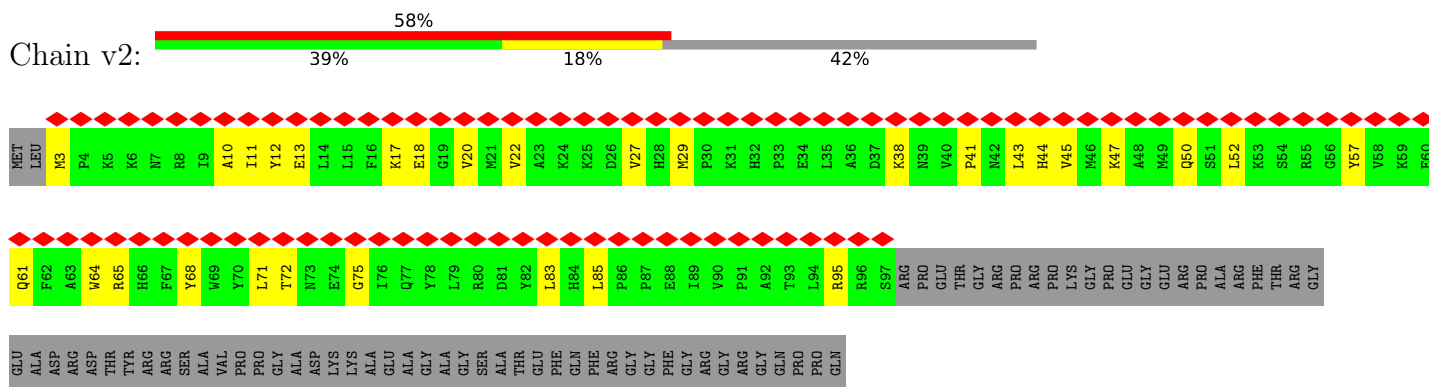


• Molecule 74: Small ribosomal subunit protein eS4

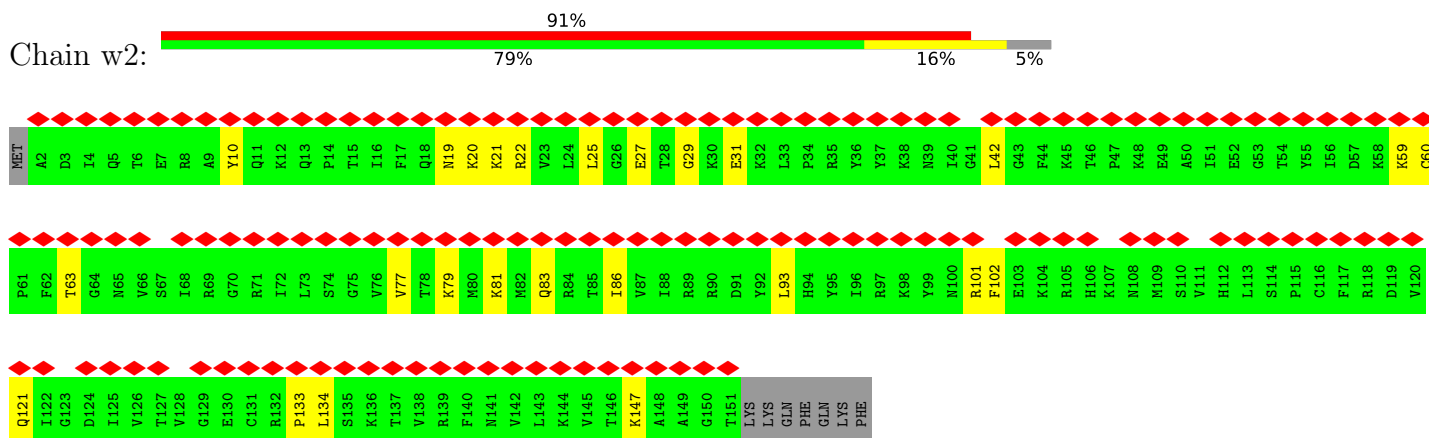




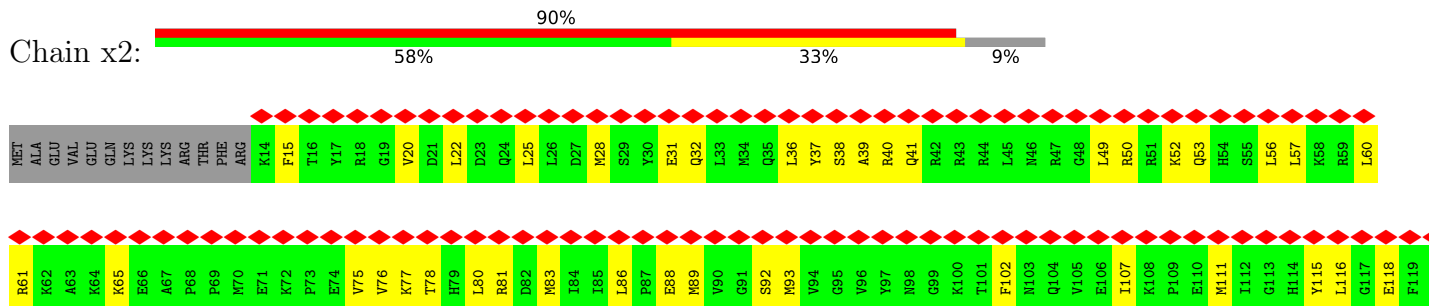
- Molecule 78: Small ribosomal subunit protein eS10

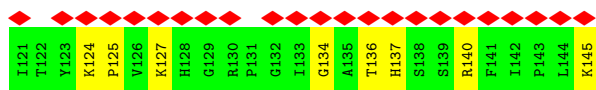


- Molecule 79: Small ribosomal subunit protein uS17

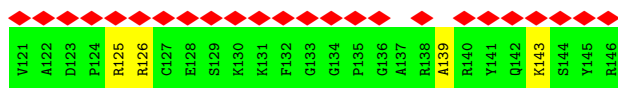
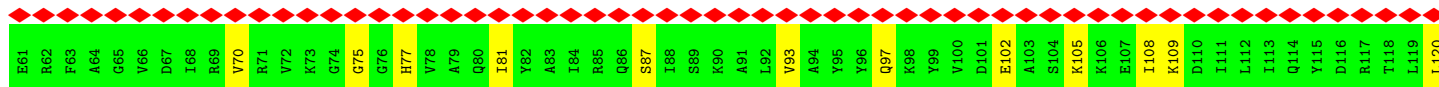
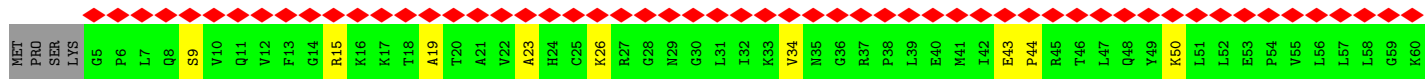
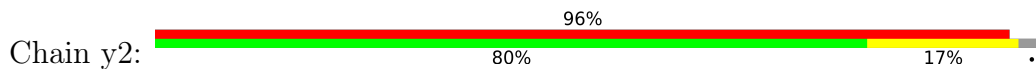


- Molecule 80: Small ribosomal subunit protein uS19

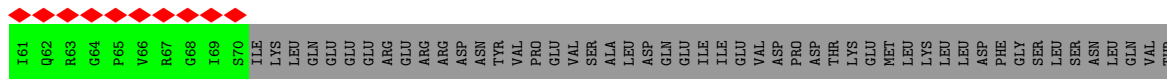
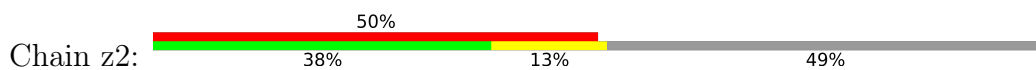




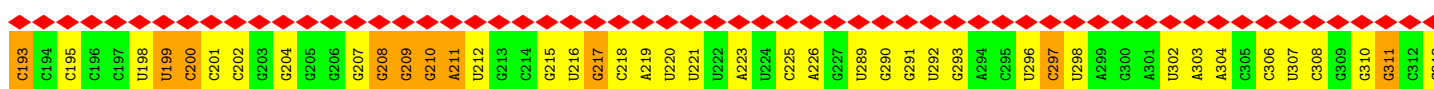
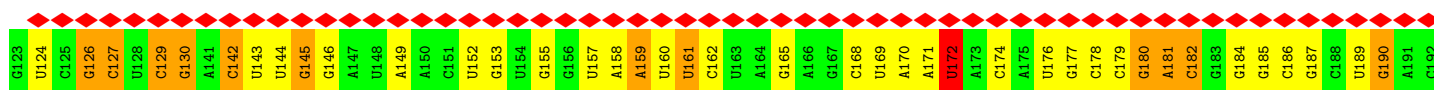
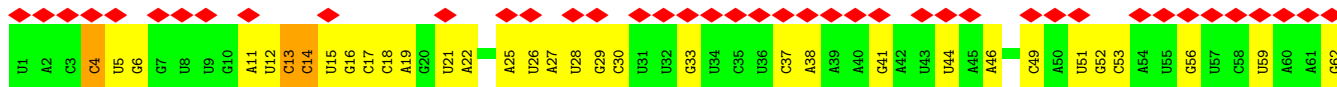
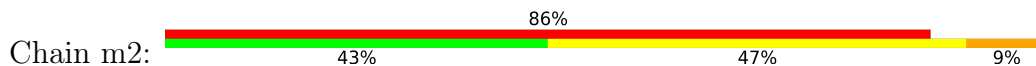
- Molecule 81: Small ribosomal subunit protein uS9



- Molecule 82: Small ribosomal subunit protein eS17



- Molecule 83: 18S ribosomal RNA



C1182	A1121	A992	G931	A871	A811	C640	U579	A502	G442	G382	G314
A1183	U1122	G993	C932	A872	A812	C641	C580	C503	C443	C383	A315
A1184	G1123	A994	C933	U873	A813	A642	C581	C504	C444	C384	U316
A1185	A1124	G997	G934	A874	A814	A643	U582	C505	G445	G385	C317
G1186	C1125	A998	G935	G875	A815	U644	U583	C506	G446	U386	G318
U1187	U1126	A999	G936	G876	U816	A645	U584	G507	A447	G387	C319
U1188	C1127	A1000	G937	A877	U817	U646	A585	G508	G448	C388	A320
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A1190	C1129	U1004	C939	C879	U819	U648	A587	A510	A450	U390	G322
A1191	U1130	U1005	A940	C880	A820	A650	C587	G511	A451	U391	C323
A1192	G1131	U1006	U941	C881	A821	U651	A589	G512	C452	A392	C324
C1193	U1132	G1007	U942	G882	U822	A652	G590	U513	G453	C393	G331
U1194	C1133	C1008	C943	G883	U823	U653	G591	A514	A454	A394	G332
U1195	U1134	C1009	G944	U884	U824	U654	A592	G515	C455	U395	C333
A1196	A1135	C1009	U945	U885	U825	A655	U593	U516	U456	G396	G334
A1197	G1136	A1010	A946	C886	C826	A656	C594	A518	A457	G397	G335
G1200	C1137	U1011	U947	U887	A827	U657	C595	C519	C458	U398	C336
A1201	U1138	G1012	U948	A888	A828	C658	A596	G520	A460	G399	G337
A1202	U1139	A1013	C949	U889	A829	U659	U597	A521	C461	A400	A338
U1203	C1140	U1015	C950	U890	G830	U660	U598	A522	A462	C401	C339
U1204	C1141	U1015	G951	U891	C831	G661	G599	A523	U463	C402	G340
G1205	G1142	U1083	C952	U892	A832	U662	G600	A524	C464	A403	A341
A1206	G1143	C953	C953	G893	G833	G664	A601	A525	C465	C404	C342
C1207	U1017	G954	G954	G894	G834	C665	G602	U526	A466	G405	C343
G1208	A1085	C955	C955	U895	C835	U668	G603	A527	A467	G406	C344
G1209	A1086	U956	U956	C896	C836	U669	G604	C528	G468	G407	C345
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A1211	C1021	G958	G958	U898	G838	A671	A607	A530	A471	G409	U347
G1212	U1022	A959	A959	U899	A839	A672	G608	A531	G472	C411	C348
G1213	A1023	C960	C960	U900	G840	A673	U609	U532	G473	G412	G349
U1214	U1024	G961	G961	U901	C941	A674	C610	A533	C474	G413	A350
C1215	A1025	U962	U962	C902	C942	C675	U611	C534	A475	G414	A351
A1216	A1026	U963	G963	C903	G843	U677	G612	A535	G476	G415	C352
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G1218	C1028	A965	A965	A905	C945	C679	U614	U554	U478	A417	U354
A1219	A1029	A966	A966	A906	C946	U680	G615	A555	G479	U418	C355
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A1222	C1033	C969	C969	G909	A849	U683	A618	U558	G482	G421	C358
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C1224	G1035	U971	U971	G911	A851	C685	G622	G560	G484	C423	A362
A1225	A1036	G972	G972	G912	C852	C686	C623	G561	C485	U424	U363
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G1228	A1039	C975	C975	A915	C855	C689	U626	U564	A488	G427	A366
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A1230	U1042	G979	G979	G917	G857	U801	G628	A566	U490	A429	U368
G1231	G1043	U980	U980	A918	C858	U802	U829	A567	A491	U430	U369
C1232	U1044	C981	C981	U919	U859	U803	A830	U568	C492	C431	U370
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G1235	U1047	A984	A984	A922	G862	U806	U633	A571	A495	C434	A373
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A1241	U1117	C990	C990	A928	U868	G869	U639	A577	A441	G440	G379
U1242	U1118	G991	G991	C929	G869	G870					C381

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C1857	C1279	C1339	U1399	U1459	G1519	G1579	A1639	C1699	G1759	C1857
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A1873	A1295	A1355	G1415	G1475	A1535	C1595	U1655	A1717	G1807	A1873
G1874	G1296	G1356	A1416	G1476	U1536	C1596	G1656	C1718	A1808	G1874
U1875	C1297	C1357	C1417	G1477	U1537	A1597	C1657	G1719	C1809	U1875
U1876	U1298	G1358	C1418	A1478	G1538	U1597	G1658	C1720	U1810	U1876
U1877	U1299	A1359	C1419	U1479	A1539	U1598	G1659	A1721	A1811	U1877
G1878	G1300	U1360	C1420	U1480	C1540	C1599	G1660	U1722	U1812	G1878
A1879	U1301	U1361	U1421	G1481	U1541	U1601	U1661		C1813	A1879
U1880	U1302	U1362				G1602	C1662			U1880

4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C1	Depositor
Number of particles used	9545	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	FEI TALOS ARCTICA	Depositor
Voltage (kV)	200	Depositor
Electron dose ($e^-/\text{\AA}^2$)	45	Depositor
Minimum defocus (nm)	800	Depositor
Maximum defocus (nm)	2900	Depositor
Magnification	100000	Depositor
Image detector	GATAN K2 QUANTUM (4k x 4k)	Depositor
Maximum map value	0.172	Depositor
Minimum map value	-0.084	Depositor
Average map value	0.002	Depositor
Map value standard deviation	0.010	Depositor
Recommended contour level	0.037	Depositor
Map size (Å)	333.3, 333.3, 333.3	wwPDB
Map dimensions	330, 330, 330	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	1.01, 1.01, 1.01	Depositor

5 Model quality ⓘ

5.1 Standard geometry ⓘ

Bond lengths and bond angles in the following residue types are not validated in this section: 1MA, 4AC, OMG, MG, 5MC, OMC, PSU, OMU, B8N, A2M, 2MG, B8T, ZN, UR3

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	A1	0.25	0/1888	0.40	0/2516
2	A2	0.23	1/84889 (0.0%)	0.32	0/132400
3	A3	0.15	0/1169	0.41	0/1568
4	B1	0.19	0/1847	0.37	0/2486
5	B2	0.20	0/2836	0.28	0/4421
6	B3	0.13	0/1122	0.33	0/1503
7	Bv	0.23	0/1813	0.39	0/2823
7	Bz	0.23	0/1813	0.41	0/2823
7	n2	0.14	0/1813	0.34	0/2823
8	Bx	0.41	0/219	0.53	0/336
10	C1	0.21	0/1537	0.39	0/2065
11	C2	0.22	0/3675	0.31	0/5725
12	C3	0.13	0/818	0.32	0/1099
13	D1	0.22	0/1694	0.38	0/2261
14	D2	0.25	0/1959	0.44	0/2627
15	D3	0.11	0/645	0.32	0/863
16	E1	0.20	0/1420	0.42	0/1899
17	E2	0.24	0/3305	0.45	0/4422
18	E3	0.13	0/1097	0.39	0/1464
19	F1	0.22	0/1674	0.41	0/2241
20	F2	0.24	0/2921	0.41	0/3921
21	F3	0.14	0/805	0.37	0/1079
22	G1	0.19	0/1165	0.37	0/1558
23	G2	0.20	0/2435	0.38	0/3260
24	G3	0.13	0/490	0.37	0/656
25	H1	0.27	0/1746	0.42	1/2338 (0.0%)
26	H2	0.19	0/1822	0.37	0/2443
27	H3	0.18	0/466	0.43	0/618
28	I2	0.25	0/1670	0.43	0/2232
29	I3	0.12	0/2493	0.36	0/3394
30	J2	0.24	0/1268	0.43	0/1700
31	J3	0.13	0/1737	0.35	0/2348

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
32	K2	0.23	0/1535	0.40	0/2048
33	K3	0.11	0/1863	0.32	0/2481
34	L1	0.13	0/1318	0.39	0/1767
35	L2	0.22	0/1515	0.36	0/2002
36	L3	0.10	0/1542	0.29	0/2058
37	M2	0.24	0/1490	0.41	0/2000
38	M3	0.10	0/962	0.27	0/1289
39	N2	0.22	0/1327	0.37	0/1771
40	N3	0.11	0/1232	0.31	0/1656
41	O2	0.17	0/839	0.38	0/1126
42	O3	0.12	0/1015	0.32	0/1361
43	P2	0.23	0/983	0.43	0/1319
44	P3	0.12	0/1051	0.31	0/1406
45	Q2	0.22	0/532	0.40	0/708
46	Q3	0.13	0/1019	0.36	0/1354
47	R2	0.21	0/984	0.39	0/1323
48	R3	0.13	0/576	0.35	0/774
49	S2	0.21	0/1132	0.41	0/1504
50	S3	0.14	0/665	0.40	0/891
51	T2	0.20	0/1130	0.39	0/1507
52	T3	0.12	0/443	0.35	0/582
53	U2	0.26	0/1193	0.39	0/1593
54	U3	0.12	0/424	0.32	0/566
55	V2	0.21	0/963	0.38	0/1275
56	W2	0.17	0/742	0.38	0/996
57	X2	0.21	0/903	0.38	0/1216
58	Y2	0.27	0/1071	0.40	0/1429
59	Z2	0.24	0/895	0.39	0/1198
60	a2	0.23	0/916	0.42	0/1221
61	b2	0.20	0/1009	0.39	0/1332
62	c2	0.19	0/843	0.37	0/1115
63	d2	0.26	0/720	0.46	0/952
64	e2	0.19	0/574	0.38	0/760
65	f2	0.24	0/454	0.36	0/599
66	g2	0.24	0/435	0.43	0/575
67	h2	0.15	0/231	0.37	0/294
68	i2	0.23	0/855	0.46	0/1128
69	j2	0.23	0/704	0.42	0/935
70	k2	0.23	0/1016	0.41	0/1363
71	o2	0.11	0/1731	0.28	0/2352
72	p2	0.15	0/1749	0.41	0/2340
73	q2	0.13	0/1739	0.34	0/2342
74	r2	0.14	0/2118	0.37	0/2849

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
75	s2	0.13	0/1477	0.33	0/1983
76	t2	0.13	0/1299	0.36	0/1767
77	u2	0.12	0/1662	0.36	0/2228
78	v2	0.14	0/824	0.33	0/1112
79	w2	0.13	0/1241	0.38	0/1662
80	x2	0.13	0/1103	0.35	0/1475
81	y2	0.15	0/1146	0.35	0/1534
82	z2	0.12	0/567	0.34	0/756
83	m2	0.42	3/38273 (0.0%)	0.34	12/59642 (0.0%)
All	All	0.26	4/228281 (0.0%)	0.35	13/335428 (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
17	E2	0	1

All (4) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
83	m2	1350	G	O3'-P	-75.31	0.48	1.61
2	A2	3441	A2M	O3'-P	5.15	1.61	1.56
83	m2	486	A2M	O3'-P	5.05	1.61	1.56
83	m2	670	A2M	O3'-P	5.03	1.61	1.56

All (13) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
83	m2	1350	G	P-O3'-C3'	21.54	152.50	120.20
83	m2	1350	G	OP2-P-O3'	-17.10	56.70	108.00
83	m2	1350	G	O3'-P-O5'	9.97	118.95	104.00
83	m2	1358	G	O3'-P-O5'	-8.61	91.09	104.00
83	m2	1357	C	C1'-C2'-O2'	-7.40	97.31	108.40
83	m2	1350	G	OP1-P-O3'	7.16	129.47	108.00
83	m2	1357	C	C4'-C3'-O3'	-6.74	102.89	113.00
83	m2	1357	C	O3'-P-O5'	-6.51	94.24	104.00
83	m2	1352	U	C4'-C3'-C2'	-5.89	96.71	102.60
83	m2	1351	G	C2'-C3'-O3'	-5.53	105.40	113.70
25	H1	184	ILE	N-CA-C	-5.29	107.20	111.91
83	m2	1359	A	P-O5'-C5'	-5.21	113.09	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
83	m2	1351	G	O4'-C1'-C2'	-5.04	102.56	107.60

There are no chirality outliers.

All (1) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
17	E2	258	HIS	Peptide

5.2 Too-close contacts [i](#)

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	A1	1851	0	1988	38	0
2	A2	77547	0	39261	880	0
3	A3	1151	0	1202	48	0
4	B1	1812	0	1947	33	0
5	B2	2538	0	1286	13	0
6	B3	1104	0	1139	29	0
7	Bv	1623	0	820	23	0
7	Bz	1623	0	821	31	0
7	n2	1623	0	821	34	0
8	Bx	200	0	101	5	0
9	By	110	0	31	1	0
10	C1	1519	0	1603	24	0
11	C2	3315	0	1685	34	0
12	C3	808	0	878	26	0
13	D1	1656	0	1706	18	0
14	D2	1921	0	2022	38	0
15	D3	638	0	635	13	0
16	E1	1397	0	1425	36	0
17	E2	3238	0	3380	58	0
18	E3	1080	0	1147	16	0
19	F1	1643	0	1750	29	0
20	F2	2867	0	3040	49	0
21	F3	789	0	841	21	0
22	G1	1143	0	1219	22	0
23	G2	2389	0	2420	32	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
24	G3	488	0	514	18	0
25	H1	1701	0	1749	29	0
26	H2	1789	0	1932	27	0
27	H3	455	0	445	17	0
28	I2	1640	0	1792	24	0
29	I3	2436	0	2393	62	0
30	J2	1242	0	1274	22	0
31	J3	1700	0	1786	44	0
32	K2	1511	0	1636	33	0
33	K3	1840	0	1989	38	0
34	L1	1300	0	1375	26	0
35	L2	1499	0	1651	19	0
36	L3	1518	0	1632	26	0
37	M2	1450	0	1488	19	0
38	M3	952	0	993	19	0
39	N2	1299	0	1368	20	0
40	N3	1208	0	1294	17	0
41	O2	825	0	850	11	0
42	O3	1002	0	1023	30	0
43	P2	969	0	1031	23	0
44	P3	1034	0	1080	20	0
45	Q2	519	0	533	4	0
46	Q3	1002	0	1075	16	0
47	R2	967	0	1040	17	0
48	R3	570	0	626	16	0
49	S2	1115	0	1205	18	0
50	S3	651	0	672	13	0
51	T2	1107	0	1182	22	0
52	T3	438	0	484	10	0
53	U2	1164	0	1213	17	0
54	U3	415	0	393	9	0
55	V2	945	0	1037	11	0
56	W2	732	0	769	10	0
57	X2	888	0	930	12	0
58	Y2	1053	0	1147	15	0
59	Z2	876	0	912	15	0
60	a2	906	0	997	20	0
61	b2	1001	0	1138	17	0
62	c2	832	0	917	9	0
63	d2	705	0	737	14	0
64	e2	568	0	635	17	0
65	f2	444	0	483	8	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
66	g2	429	0	465	3	0
67	h2	230	0	276	25	0
68	i2	842	0	912	13	0
69	j2	694	0	738	10	0
70	k2	1001	0	1066	28	0
71	o2	1694	0	1696	45	0
72	p2	1722	0	1794	47	0
73	q2	1711	0	1804	48	0
74	r2	2076	0	2177	52	0
75	s2	1457	0	1508	45	0
76	t2	1278	0	1207	26	0
77	u2	1633	0	1664	40	0
78	v2	800	0	818	25	0
79	w2	1220	0	1289	18	0
80	x2	1081	0	1134	42	0
81	y2	1128	0	1195	19	0
82	z2	560	0	614	13	0
83	m2	34939	0	17651	711	0
84	A2	82	0	0	0	0
84	Bv	2	0	0	0	0
84	H1	1	0	0	0	0
84	J2	1	0	0	0	0
84	P2	1	0	0	0	0
84	d2	1	0	0	0	0
84	m2	30	0	0	0	0
85	F3	1	0	0	0	0
85	H3	1	0	0	0	0
85	d2	1	0	0	0	0
85	g2	1	0	0	0	0
85	i2	1	0	0	0	0
85	j2	1	0	0	0	0
86	B1	1	0	0	0	0
All	All	214961	0	158596	3072	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 8.

All (3072) 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)
83:m2:1350:G:H2'	83:m2:1351:G:C5'	1.31	1.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
83:m2:1351:G:C2	83:m2:1383:G:N2	1.76	1.51
83:m2:1350:G:C2'	83:m2:1351:G:H5''	1.29	1.38
2:A2:1376:A:N6	83:m2:1030:A:N1	1.70	1.35
83:m2:1351:G:N1	83:m2:1383:G:C2	1.97	1.32
2:A2:4663:G:OP1	77:u2:92:ARG:HG3	1.28	1.25
83:m2:1350:G:O2'	83:m2:1351:G:H5''	1.36	1.25
83:m2:1351:G:C2	83:m2:1383:G:C2	2.25	1.24
83:m2:1351:G:N1	83:m2:1383:G:N2	1.84	1.24
83:m2:1350:G:C2'	83:m2:1351:G:C5'	1.89	1.22
3:A3:12:ILE:HD13	16:E1:121:PRO:CG	1.71	1.19
2:A2:3415:A:C2	83:m2:1828:G:N2	2.12	1.18
67:h2:16:LYS:NZ	83:m2:1820:A:H4'	1.57	1.18
83:m2:630:A:N6	83:m2:1502:G:N3	1.93	1.16
83:m2:1350:G:C8	83:m2:1351:G:C8	2.36	1.12
83:m2:1350:G:H2'	83:m2:1351:G:H5'	1.18	1.10
3:A3:12:ILE:HD13	16:E1:121:PRO:HG3	1.32	1.09
83:m2:1353:G:C2	83:m2:1381:A:C6	2.42	1.07
24:G3:62:GLU:HG2	42:O3:121:ARG:NE	1.75	1.01
24:G3:62:GLU:HG2	42:O3:121:ARG:CZ	1.90	0.99
2:A2:2623:G:OP1	83:m2:370:U:O4	1.79	0.99
83:m2:1351:G:C6	83:m2:1383:G:N2	2.31	0.99
83:m2:1351:G:H5'	83:m2:1351:G:H8	1.30	0.96
83:m2:1353:G:N1	83:m2:1381:A:N6	2.14	0.96
83:m2:1353:G:C6	83:m2:1381:A:N6	2.35	0.95
67:h2:3:ALA:HB3	83:m2:1844:4AC:OP1	1.67	0.94
83:m2:1290:U:H3	83:m2:1313:C:H42	1.15	0.94
2:A2:3415:A:N3	83:m2:1828:G:N2	2.14	0.94
3:A3:12:ILE:HD13	16:E1:121:PRO:CB	1.96	0.93
67:h2:11:ARG:HH22	83:m2:1846:U:P	1.90	0.93
83:m2:1351:G:N3	83:m2:1383:G:N2	2.16	0.93
2:A2:1376:A:N6	83:m2:1030:A:C2	2.35	0.93
2:A2:3348:A:H62	2:A2:3479:G:H21	1.14	0.93
67:h2:16:LYS:HZ3	83:m2:1820:A:H4'	1.26	0.92
71:o2:109:THR:HG22	83:m2:1353:G:H4'	1.52	0.91
67:h2:16:LYS:HZ1	83:m2:1820:A:H4'	1.35	0.91
2:A2:1259:C:H42	2:A2:1901:A:H61	1.15	0.90
2:A2:1374:G:N2	2:A2:1379:C:O2	2.04	0.90
83:m2:70:G:H21	83:m2:79:A:H62	1.10	0.89
50:S3:36:LYS:HD3	50:S3:43:ILE:HG13	1.54	0.89
7:Bv:30:G:OP1	80:x2:136:THR:O	1.91	0.89
74:r2:137:PRO:HB2	74:r2:150:PRO:HD2	1.54	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
83:m2:1290:U:H3	83:m2:1313:C:N4	1.72	0.88
83:m2:1351:G:C4	83:m2:1383:G:N2	2.41	0.88
31:J3:114:LYS:HG3	83:m2:1360:U:OP1	1.74	0.88
71:o2:109:THR:HG21	83:m2:1353:G:O3'	1.73	0.87
83:m2:1351:G:N2	83:m2:1383:G:N3	2.23	0.86
3:A3:12:ILE:CD1	16:E1:121:PRO:HG3	2.04	0.86
83:m2:1350:G:H2'	83:m2:1351:G:H8	1.41	0.86
2:A2:740:A:H62	2:A2:828:G:H21	1.25	0.85
2:A2:4663:G:OP1	77:u2:92:ARG:CG	2.22	0.84
46:Q3:90:ARG:HG2	46:Q3:93:ARG:HH21	1.40	0.84
83:m2:1351:G:C8	83:m2:1351:G:H5'	2.12	0.84
51:T2:99:ASP:HB3	51:T2:102:ARG:HH21	1.42	0.83
3:A3:12:ILE:HD13	16:E1:121:PRO:HB3	1.59	0.83
2:A2:3885:A:H4'	68:i2:98:LYS:HE3	1.59	0.82
2:A2:3593:C:H1'	25:H1:125:SER:HB3	1.61	0.82
3:A3:19:ASN:OD1	16:E1:119:TYR:CD2	2.33	0.81
32:K2:39:THR:HG22	32:K2:41:SER:H	1.44	0.81
67:h2:15:ARG:HG2	67:h2:18:ARG:HH21	1.43	0.81
71:o2:112:ILE:HD13	83:m2:1352:U:O2'	1.81	0.81
8:Bx:44:U:N3	7:n2:34:G:O6	2.13	0.81
55:V2:35:VAL:HG23	55:V2:40:LEU:HD21	1.63	0.80
78:v2:27:VAL:HB	78:v2:43:LEU:HD13	1.64	0.80
2:A2:181:C:H2'	2:A2:182:G:H8	1.47	0.80
36:L3:31:LEU:HD12	36:L3:106:LEU:HD12	1.64	0.80
43:P2:13:LYS:HD2	43:P2:128:LEU:HD11	1.61	0.80
83:m2:1398:A:H2	83:m2:1451:G:H1	1.27	0.79
29:I3:244:ASN:HD21	29:I3:245:ARG:HH21	1.29	0.79
67:h2:2:ARG:NE	83:m2:1844:4AC:OP2	2.15	0.79
2:A2:3415:A:H2	83:m2:1828:G:N2	1.78	0.79
83:m2:1353:G:N2	83:m2:1381:A:C5	2.51	0.78
71:o2:109:THR:CG2	83:m2:1353:G:O3'	2.30	0.78
2:A2:3422:A:O2'	83:m2:1830:C:H4'	1.83	0.78
28:I2:61:ARG:HD3	28:I2:66:PRO:HB3	1.64	0.78
50:S3:20:LYS:HG3	50:S3:21:LYS:HG2	1.66	0.78
83:m2:1273:C:H42	83:m2:1513:U:H3	1.31	0.78
2:A2:1505:C:H5''	32:K2:53:MET:HE3	1.65	0.78
2:A2:2279:U:H3	2:A2:2284:A:H2	1.31	0.77
42:O3:34:PHE:HB3	42:O3:41:PHE:HB2	1.67	0.77
57:X2:90:ARG:HD3	57:X2:102:LEU:HD23	1.65	0.77
80:x2:134:GLY:HA2	80:x2:140:ARG:HB3	1.67	0.76
23:G2:83:LEU:HB3	23:G2:88:VAL:HB	1.65	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:2276:G:H4'	60:a2:26:PRO:HD2	1.68	0.76
83:m2:1351:G:N1	83:m2:1383:G:N1	2.33	0.76
83:m2:145:G:H1	83:m2:174:OMC:H5	1.30	0.76
83:m2:1351:G:N2	83:m2:1383:G:C2	2.53	0.75
2:A2:4573:C:H4'	22:G1:121:ARG:HH12	1.49	0.75
49:S2:50:ARG:HH12	49:S2:110:LYS:HD3	1.51	0.75
12:C3:62:ARG:HG2	12:C3:81:GLN:HG2	1.68	0.75
43:P2:73:ARG:NH2	83:m2:1726:A:OP1	2.19	0.75
75:s2:179:ASN:HD21	75:s2:186:ASN:HB2	1.50	0.75
83:m2:1571:A:H8	83:m2:1615:G:H21	1.34	0.75
83:m2:153:G:H22	83:m2:165:G:H1	1.34	0.75
2:A2:1011:U:H3'	2:A2:1012:C:H5''	1.69	0.75
73:q2:106:ARG:HG3	73:q2:175:VAL:HG22	1.68	0.75
2:A2:2106:OMC:HM22	20:F2:95:MET:HG3	1.68	0.75
31:J3:136:HIS:HB3	31:J3:162:ILE:HD11	1.69	0.74
6:B3:129:ARG:HH12	6:B3:133:ARG:HB2	1.51	0.74
83:m2:70:G:N2	83:m2:79:A:H62	1.83	0.74
83:m2:1350:G:O2'	83:m2:1351:G:C5'	2.17	0.74
83:m2:1353:G:C2	83:m2:1381:A:C5	2.75	0.74
76:t2:43:LEU:HB3	76:t2:72:PHE:HE1	1.50	0.74
83:m2:1131:G:H3'	83:m2:1132:G:H21	1.53	0.74
51:T2:50:PRO:HD3	51:T2:68:ILE:HG12	1.69	0.74
17:E2:57:VAL:HG22	17:E2:73:VAL:HG22	1.70	0.74
82:z2:3:ARG:HH22	83:m2:1479:U:H5'	1.51	0.74
2:A2:3336:U:H5''	14:D2:54:ARG:HH21	1.51	0.74
7:Bz:69:G:H2'	7:Bz:70:G:H8	1.52	0.74
2:A2:3626:G:H4'	34:L1:168:ALA:HB2	1.70	0.73
2:A2:4555:A:H4'	17:E2:95:THR:HG22	1.69	0.73
27:H3:25:SER:HB2	78:v2:65:ARG:HD3	1.70	0.73
12:C3:61:LEU:HD22	27:H3:34:TYR:HE2	1.52	0.73
73:q2:135:GLU:HG2	73:q2:153:VAL:HG23	1.68	0.73
83:m2:602:G:H2'	83:m2:603:OMG:H8	1.53	0.73
83:m2:853:C:H5''	83:m2:854:G:H5'	1.71	0.73
7:n2:51:U:H3	7:n2:63:G:H1	1.35	0.73
83:m2:983:A:H2'	83:m2:984:G:C8	2.24	0.73
28:I2:125:LYS:HG3	28:I2:129:LEU:HD12	1.70	0.72
83:m2:1351:G:C5	83:m2:1383:G:N2	2.52	0.72
74:r2:20:LEU:HD21	74:r2:46:ILE:HD12	1.71	0.72
83:m2:930:G:H2'	83:m2:931:G:C8	2.25	0.72
12:C3:80:PHE:HB3	27:H3:52:PHE:HB3	1.69	0.72
29:I3:286:CYS:HA	29:I3:302:TYR:HA	1.69	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
83:m2:1398:A:N1	83:m2:1451:G:O6	2.23	0.72
2:A2:2366:A:H5'	2:A2:2443:G:H4'	1.70	0.71
2:A2:2302:G:H1	2:A2:2527:C:H5	1.38	0.71
2:A2:3625:G:H2'	2:A2:3626:G:C8	2.26	0.71
2:A2:1742:G:H22	2:A2:4086:C:H5''	1.55	0.71
24:G3:7:GLN:HG2	24:G3:61:SER:H	1.54	0.71
80:x2:37:TYR:HB3	80:x2:41:GLN:HB2	1.72	0.71
2:A2:2117:U:H2'	2:A2:2118:A2M:H8	1.73	0.71
83:m2:1351:G:C6	83:m2:1383:G:C2	2.78	0.71
75:s2:76:MET:HE2	75:s2:173:LEU:HD22	1.72	0.71
2:A2:4707:A:H4'	2:A2:4708:G:H5''	1.72	0.71
39:N2:48:VAL:HG21	39:N2:94:GLU:HG2	1.73	0.71
2:A2:1020:C:H2'	2:A2:1021:G:H8	1.55	0.71
2:A2:4391:C:H3'	2:A2:4393:A:H5''	1.73	0.71
3:A3:12:ILE:CD1	16:E1:121:PRO:CG	2.60	0.71
7:Bv:34:G:H1	8:Bx:48:U:H3	1.39	0.71
83:m2:1658:G:H1	83:m2:1670:U:H3	1.37	0.70
40:N3:19:ARG:HH22	50:S3:83:GLN:HG2	1.56	0.70
67:h2:16:LYS:HZ3	83:m2:1820:A:C4'	2.02	0.70
27:H3:8:TRP:HD1	27:H3:12:ARG:HH21	1.39	0.70
2:A2:1354:U:H3	2:A2:1431:G:H1	1.37	0.70
22:G1:104:MET:HE2	28:I2:199:ASN:HB3	1.72	0.70
7:n2:20:U:H1'	7:n2:21:A:H5''	1.74	0.70
2:A2:1314:C:H6	32:K2:68:ARG:HH21	1.39	0.70
31:J3:194:ARG:HD3	31:J3:196:ILE:HD11	1.72	0.70
2:A2:154:G:H4'	61:b2:105:LYS:HD2	1.73	0.70
2:A2:1158:C:H1'	19:F1:10:LEU:HD11	1.73	0.70
21:F3:22:ARG:HH21	42:O3:142:ARG:HB3	1.57	0.69
36:L3:162:ARG:HH21	46:Q3:30:PRO:HB3	1.57	0.69
77:u2:130:THR:HG23	77:u2:132:GLU:H	1.57	0.69
83:m2:510:A:H3'	83:m2:511:OMG:H8	1.55	0.69
2:A2:1068:C:H42	2:A2:1082:A:H61	1.40	0.69
4:B1:105:GLU:HB2	4:B1:109:GLU:HB2	1.74	0.69
31:J3:70:VAL:HG21	31:J3:93:ILE:HG23	1.72	0.69
83:m2:225:C:H2'	83:m2:226:A:H8	1.57	0.69
14:D2:102:LEU:HB2	14:D2:107:MET:HE3	1.75	0.69
29:I3:5:MET:HB3	29:I3:270:LEU:HD21	1.75	0.69
7:Bv:30:G:OP1	80:x2:136:THR:C	2.36	0.69
46:Q3:112:ASN:HA	46:Q3:115:LYS:HD3	1.73	0.69
56:W2:38:ILE:HG21	56:W2:63:TYR:HB3	1.74	0.69
72:p2:38:MET:HE2	72:p2:182:LYS:HA	1.75	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
79:w2:22:ARG:HG2	79:w2:27:GLU:HA	1.74	0.69
14:D2:54:ARG:HH11	14:D2:58:LEU:HD21	1.57	0.69
83:m2:529:C:H2'	83:m2:530:A:H8	1.57	0.69
12:C3:43:ALA:HB1	12:C3:50:VAL:HG21	1.75	0.69
81:y2:93:VAL:HG13	81:y2:105:LYS:HE3	1.75	0.68
31:J3:191:VAL:HG11	31:J3:236:PHE:HA	1.76	0.68
2:A2:3734:G:H1	14:D2:68:ARG:NH2	1.90	0.68
2:A2:3411:G:H1	2:A2:3424:U:H3	1.41	0.68
55:V2:32:LEU:HD13	55:V2:43:MET:HE1	1.75	0.68
2:A2:4543:G:H2'	2:A2:4544:G:C8	2.28	0.68
80:x2:83:MET:HB3	80:x2:116:LEU:HD13	1.76	0.68
83:m2:1350:G:C8	83:m2:1351:G:N7	2.61	0.68
83:m2:1351:G:C5'	83:m2:1351:G:H8	2.06	0.68
71:o2:41:ARG:HH21	71:o2:45:GLY:HA2	1.59	0.68
77:u2:98:LYS:HB3	83:m2:379:G:H5'	1.75	0.68
2:A2:1894:G:H4'	2:A2:1895:G:H4'	1.76	0.68
83:m2:65:C:H5'	83:m2:78:C:H41	1.58	0.68
3:A3:86:ARG:HD2	3:A3:106:LYS:HD3	1.75	0.68
29:I3:236:ILE:HG12	29:I3:252:THR:HG22	1.75	0.68
58:Y2:78:LEU:HB3	70:k2:20:ARG:HD3	1.76	0.68
72:p2:189:ILE:HG13	72:p2:190:PRO:HD3	1.75	0.68
83:m2:5:U:H2'	83:m2:6:G:H8	1.59	0.68
40:N3:89:TYR:HE1	40:N3:151:ALA:H	1.40	0.67
83:m2:973:G:HO2'	83:m2:974:A:H8	1.43	0.67
29:I3:40:ILE:HD11	29:I3:66:VAL:HG21	1.76	0.67
81:y2:93:VAL:HG21	81:y2:120:LEU:HD11	1.76	0.67
82:z2:26:ASN:HA	82:z2:58:MET:HE2	1.75	0.67
67:h2:16:LYS:NZ	83:m2:1820:A:C4'	2.48	0.67
83:m2:1230:A:H2'	83:m2:1231:G:C8	2.29	0.67
2:A2:746:G:HO2'	2:A2:747:G:H8	1.43	0.67
62:c2:67:LYS:HG3	62:c2:71:LYS:NZ	2.10	0.67
83:m2:1116:U:H3	83:m2:1121:A:H61	1.43	0.67
60:a2:5:LEU:HD21	60:a2:30:ILE:HG22	1.77	0.67
2:A2:928:G:H1	2:A2:1064:G:H22	1.41	0.67
73:q2:70:THR:HG22	73:q2:86:LEU:HD13	1.77	0.67
71:o2:77:ILE:HG21	71:o2:133:PRO:HG3	1.78	0.66
2:A2:1744:A:H2'	2:A2:1745:A:C8	2.30	0.66
31:J3:114:LYS:CG	83:m2:1360:U:H5'	2.25	0.66
33:K3:70:HIS:HB3	33:K3:101:ILE:HB	1.77	0.66
31:J3:113:GLN:HG2	31:J3:122:THR:HG22	1.78	0.66
2:A2:1604:A:H5''	2:A2:1605:G:H5'	1.78	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:A3:117:ILE:HD11	80:x2:111:MET:HE2	1.78	0.66
72:p2:144:LYS:HD2	72:p2:146:ARG:HH12	1.61	0.66
83:m2:1679:U:H2'	83:m2:1680:A2M:H8	1.76	0.66
62:c2:33:LEU:HD21	62:c2:38:LYS:HD2	1.78	0.66
22:G1:40:GLY:HA3	22:G1:45:VAL:HB	1.76	0.66
2:A2:181:C:H2'	2:A2:182:G:C8	2.29	0.66
74:r2:72:ILE:HG12	74:r2:90:ILE:HG12	1.76	0.66
79:w2:27:GLU:HG2	79:w2:29:GLY:H	1.60	0.66
80:x2:61:ARG:HG3	80:x2:89:MET:HE1	1.76	0.66
32:K2:53:MET:HE2	32:K2:57:ASN:HB3	1.77	0.66
71:o2:122:LEU:HD23	71:o2:144:THR:HG22	1.77	0.66
83:m2:876:G:H2'	83:m2:877:A:H8	1.59	0.66
19:F1:64:VAL:HA	19:F1:67:HIS:HD2	1.59	0.66
29:I3:61:GLY:HA3	29:I3:90:TRP:HH2	1.60	0.66
83:m2:641:C:H2'	83:m2:642:A:H8	1.60	0.66
83:m2:865:U:H2'	83:m2:866:A:H8	1.60	0.66
83:m2:950:C:H2'	83:m2:951:G:H8	1.59	0.66
3:A3:67:VAL:O	3:A3:71:MET:HG3	1.97	0.65
30:J2:131:ARG:HG3	30:J2:137:ASN:HD22	1.61	0.65
83:m2:1353:G:N1	83:m2:1381:A:C6	2.60	0.65
32:K2:178:ARG:HH21	53:U2:42:ARG:HH21	1.43	0.65
71:o2:102:ARG:CZ	83:m2:1379:U:C5	2.79	0.65
15:D3:60:ARG:HG2	71:o2:156:TYR:HD1	1.61	0.65
27:H3:8:TRP:CD1	27:H3:12:ARG:HH21	2.14	0.65
29:I3:63:SER:HB3	29:I3:84:ASP:HB3	1.77	0.65
83:m2:225:C:H2'	83:m2:226:A:C8	2.32	0.65
83:m2:1350:G:H2'	83:m2:1351:G:C8	2.30	0.65
83:m2:835:C:C4	83:m2:836:C:N4	2.64	0.65
2:A2:265:U:H5'	2:A2:266:C:H5'	1.77	0.65
2:A2:3621:A:H2'	2:A2:3698:A:H2'	1.78	0.65
31:J3:166:ARG:HB3	31:J3:247:THR:HB	1.77	0.65
25:H1:36:LEU:HD21	25:H1:109:HIS:CG	2.32	0.65
77:u2:48:VAL:HG11	77:u2:54:LYS:HE2	1.78	0.65
83:m2:1350:G:H2'	83:m2:1351:G:H5''	0.96	0.65
2:A2:74:G:H5'	19:F1:59:VAL:HG13	1.77	0.65
17:E2:257:TRP:HD1	17:E2:258:HIS:CD2	2.14	0.65
72:p2:49:VAL:HG22	72:p2:50:THR:H	1.62	0.65
74:r2:45:ILE:HA	74:r2:61:VAL:HG11	1.79	0.65
67:h2:1:MET:HG3	83:m2:1854:C:OP2	1.97	0.65
83:m2:118:C:H1'	83:m2:447:A:C5	2.31	0.65
2:A2:3775:C:H5'	4:B1:45:ILE:HD12	1.78	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:U3:121:CYS:HA	54:U3:132:MET:HE3	1.79	0.65
7:Bv:39:U:O2'	7:n2:35:A:O2'	2.14	0.65
2:A2:4663:G:P	77:u2:92:ARG:HG3	2.34	0.64
32:K2:18:PRO:HG3	32:K2:29:VAL:HG21	1.77	0.64
83:m2:1220:C:H1'	83:m2:1685:C:H42	1.59	0.64
2:A2:1672:C:H2'	2:A2:1673:A2M:H8	1.79	0.64
44:P3:69:LEU:HD21	44:P3:72:CYS:HB3	1.78	0.64
18:E3:52:LEU:HD11	18:E3:73:GLN:HB2	1.78	0.64
24:G3:55:VAL:HG13	75:s2:34:SER:HA	1.80	0.64
36:L3:109:ARG:HH22	36:L3:127:ARG:HH12	1.45	0.64
73:q2:63:GLY:O	73:q2:67:ARG:HG2	1.97	0.64
2:A2:229:G:H5''	49:S2:11:ARG:HG3	1.80	0.64
62:c2:23:LYS:HD2	62:c2:24:PRO:HD2	1.78	0.64
2:A2:453:G:H1	2:A2:1107:G:N2	1.96	0.64
17:E2:92:TYR:HB2	17:E2:159:VAL:HB	1.79	0.64
83:m2:1100:C:H2'	83:m2:1101:G:C8	2.33	0.64
2:A2:927:C:H2'	2:A2:928:G:C8	2.33	0.64
2:A2:938:U:H3	2:A2:1049:C:H42	1.46	0.64
42:O3:134:PRO:HB3	83:m2:946:A:H5''	1.80	0.64
73:q2:163:PRO:O	73:q2:167:TYR:HB2	1.98	0.64
2:A2:1894:G:H2'	2:A2:2017:G:C6	2.33	0.64
2:A2:3415:A:H2	83:m2:1828:G:H22	1.41	0.64
29:I3:258:ILE:HB	29:I3:268:ASP:HB3	1.78	0.64
2:A2:262:G:H2'	2:A2:263:G:H8	1.63	0.63
2:A2:2312:G:H1	2:A2:2325:U:H3	1.44	0.63
29:I3:176:VAL:HB	29:I3:186:THR:HG22	1.79	0.63
83:m2:102:A:H4'	83:m2:104:A:C8	2.33	0.63
83:m2:1223:G:H2'	83:m2:1224:G:H8	1.63	0.63
13:D1:188:LYS:HB3	13:D1:212:LEU:HD21	1.80	0.63
29:I3:32:LEU:HD22	29:I3:42:MET:HE2	1.80	0.63
43:P2:39:ILE:HG12	43:P2:61:VAL:HG11	1.79	0.63
2:A2:768:G:H2'	2:A2:769:G:C8	2.33	0.63
83:m2:1205:G:H2'	83:m2:1206:A:C8	2.34	0.63
12:C3:54:VAL:HB	12:C3:88:LEU:HB2	1.80	0.63
83:m2:5:U:H2'	83:m2:6:G:C8	2.33	0.63
29:I3:57:ARG:HD3	29:I3:94:THR:HA	1.80	0.63
29:I3:85:GLY:HA2	29:I3:107:ASP:HA	1.80	0.63
34:L1:52:THR:HG22	34:L1:156:LYS:HG3	1.80	0.63
2:A2:262:G:H2'	2:A2:263:G:C8	2.34	0.63
83:m2:1535:A:C8	83:m2:1606:G:H1'	2.34	0.63
7:n2:70:G:H2'	7:n2:71:G:C8	2.34	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:3312:A:H2'	2:A2:3313:U:H6	1.64	0.63
20:F2:134:PRO:HA	20:F2:150:LEU:HD22	1.80	0.63
83:m2:1130:C:H2'	83:m2:1131:G:H8	1.62	0.63
49:S2:30:MET:HB3	49:S2:101:PRO:HG3	1.81	0.62
74:r2:179:ASN:HA	74:r2:231:GLY:HA2	1.79	0.62
83:m2:971:U:H4'	83:m2:972:G:C8	2.34	0.62
83:m2:1191:A:H2'	83:m2:1192:A:H8	1.64	0.62
2:A2:1116:U:H4'	58:Y2:18:LYS:HA	1.81	0.62
17:E2:206:PRO:HG2	17:E2:209:GLN:HG3	1.79	0.62
19:F1:59:VAL:HG21	19:F1:73:GLY:HA3	1.81	0.62
81:y2:139:ALA:HB2	83:m2:1652:A:H5''	1.80	0.62
2:A2:467:U:H2'	2:A2:468:U:C6	2.35	0.62
2:A2:3566:C:H2'	2:A2:3567:C:C6	2.34	0.62
24:G3:62:GLU:HG2	42:O3:121:ARG:NH2	2.15	0.62
51:T2:124:THR:HG23	51:T2:126:LYS:H	1.64	0.62
1:A1:127:VAL:HG13	1:A1:158:VAL:HG12	1.81	0.62
2:A2:833:U:H5'	22:G1:44:ARG:HH21	1.64	0.62
2:A2:3348:A:H62	2:A2:3479:G:N2	1.94	0.62
30:J2:94:MET:HE2	30:J2:148:MET:HE3	1.81	0.62
33:K3:32:MET:HB2	33:K3:100:CYS:HB2	1.82	0.62
43:P2:16:ILE:HD11	43:P2:124:GLU:HG2	1.81	0.62
83:m2:302:U:H2'	83:m2:303:A:C8	2.35	0.62
83:m2:642:A:H2'	83:m2:643:A:C8	2.34	0.62
83:m2:1459:U:H2'	83:m2:1460:G:H8	1.63	0.62
2:A2:4506:G:H2'	2:A2:4507:G:H8	1.65	0.62
2:A2:4648:U:H4'	17:E2:375:GLY:HA3	1.81	0.62
6:B3:116:ASP:HB3	6:B3:122:LYS:HG2	1.82	0.62
6:B3:6:VAL:HG12	6:B3:135:ALA:HB2	1.82	0.62
13:D1:36:LEU:HD22	13:D1:69:ARG:HH11	1.65	0.62
16:E1:48:PRO:HB3	16:E1:72:CYS:HB2	1.82	0.62
25:H1:10:LEU:HG	25:H1:19:MET:HE3	1.82	0.62
29:I3:202:PRO:HG3	29:I3:243:PRO:HA	1.82	0.62
54:U3:140:TYR:HA	54:U3:147:THR:HG22	1.82	0.62
75:s2:23:TRP:HH2	75:s2:98:GLU:HG2	1.62	0.62
83:m2:302:U:H2'	83:m2:303:A:H8	1.64	0.62
2:A2:3713:U:H2'	2:A2:3715:G:H8	1.62	0.62
29:I3:15:ASN:HD21	83:m2:1474:C:H4'	1.65	0.62
1:A1:57:LYS:HZ2	1:A1:61:LEU:HD11	1.64	0.62
74:r2:198:ARG:HG3	74:r2:208:VAL:HG22	1.80	0.62
2:A2:1626:G:H4'	23:G2:44:TYR:CD2	2.34	0.62
81:y2:34:VAL:HG22	81:y2:70:VAL:HB	1.81	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:4394:G:H2'	2:A2:4395:G:O4'	1.99	0.61
7:Bv:37:A:H3'	7:Bv:38:A:H8	1.65	0.61
83:m2:1351:G:C6	83:m2:1383:G:N1	2.68	0.61
6:B3:76:THR:HG21	6:B3:97:LYS:HG2	1.81	0.61
7:n2:69:G:H2'	7:n2:70:G:H8	1.64	0.61
80:x2:15:PHE:HB3	80:x2:22:LEU:HD21	1.81	0.61
33:K3:181:THR:HG22	33:K3:183:ARG:H	1.65	0.61
2:A2:3336:U:H5''	14:D2:54:ARG:NH2	2.14	0.61
16:E1:151:ILE:HD11	16:E1:156:ARG:HG2	1.82	0.61
17:E2:217:ILE:HD11	17:E2:333:LEU:HD21	1.81	0.61
29:I3:173:LEU:HA	29:I3:188:HIS:O	2.00	0.61
48:R3:66:LYS:HD3	75:s2:102:LEU:HB3	1.81	0.61
77:u2:141:ARG:HB3	77:u2:144:LYS:HB2	1.82	0.61
83:m2:70:G:H21	83:m2:79:A:N6	1.91	0.61
55:V2:94:LEU:HD23	55:V2:97:ILE:HD12	1.82	0.61
2:A2:2132:C:O2	2:A2:2136:A:N6	2.33	0.61
46:Q3:52:PRO:HA	46:Q3:55:ILE:HD12	1.82	0.61
12:C3:50:VAL:HG22	12:C3:91:LEU:HG	1.82	0.61
21:F3:5:ARG:HH21	83:m2:1866:U:H3'	1.63	0.61
28:I2:163:LYS:HA	28:I2:166:MET:HE3	1.81	0.61
47:R2:89:LYS:HB3	47:R2:95:THR:OG1	2.00	0.61
67:h2:9:ARG:NH2	83:m2:1853:A:OP1	2.32	0.61
83:m2:1032:A:H2'	83:m2:1033:A2M:H8	1.82	0.61
83:m2:1538:G:H2'	83:m2:1539:A:C8	2.36	0.61
2:A2:2166:C:H2'	2:A2:2167:A:C8	2.36	0.61
33:K3:53:SER:HB3	83:m2:165:G:H4'	1.81	0.61
34:L1:170:GLY:HA2	34:L1:179:LEU:HD13	1.83	0.61
83:m2:959:A:H2'	83:m2:960:G:H8	1.65	0.61
83:m2:1350:G:N7	83:m2:1351:G:C5	2.69	0.61
2:A2:740:A:H62	2:A2:828:G:N2	1.95	0.61
2:A2:3586:U:H3	2:A2:3832:G:H1	1.48	0.61
2:A2:4395:G:H2'	2:A2:4396:A:C8	2.36	0.61
83:m2:1515:C:H2'	83:m2:1516:G:H8	1.66	0.61
2:A2:3415:A:N7	2:A2:3419:A:N1	2.49	0.61
2:A2:3513:G:H5''	30:J2:86:LYS:HB2	1.81	0.61
2:A2:4603:C:H2'	2:A2:4604:C:C6	2.36	0.61
29:I3:190:GLY:HA3	29:I3:217:MET:HE1	1.83	0.61
63:d2:21:ARG:HB2	63:d2:39:TYR:HD1	1.66	0.61
2:A2:3379:A:H2'	2:A2:3380:A2M:H8	1.83	0.60
13:D1:184:MET:HE2	13:D1:190:LEU:HG	1.82	0.60
72:p2:37:ALA:H	72:p2:232:HIS:HA	1.66	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A1:51:LYS:HE3	2:A2:1895:G:H1	1.66	0.60
3:A3:19:ASN:ND2	16:E1:119:TYR:CZ	2.69	0.60
7:Bv:51:U:H3	7:Bv:63:G:H22	1.47	0.60
18:E3:9:THR:HG22	83:m2:683:U:H4'	1.82	0.60
32:K2:59:PRO:HG3	32:K2:143:ARG:HA	1.82	0.60
34:L1:98:LYS:HA	34:L1:101:LYS:HE3	1.82	0.60
43:P2:35:LYS:HD2	43:P2:67:LYS:HG2	1.82	0.60
6:B3:129:ARG:NH1	6:B3:133:ARG:HB2	2.15	0.60
14:D2:54:ARG:HG2	14:D2:56:ALA:H	1.66	0.60
33:K3:195:LYS:HA	33:K3:198:ARG:HE	1.65	0.60
2:A2:492:G:H22	2:A2:670:G:H22	1.50	0.60
2:A2:1417:G:H2'	2:A2:1418:G:C8	2.36	0.60
14:D2:36:GLU:HG3	14:D2:91:GLY:HA2	1.83	0.60
57:X2:64:ILE:HG23	57:X2:68:LEU:HD23	1.82	0.60
83:m2:1607:G:H1'	83:m2:1636:A:H61	1.67	0.60
2:A2:1146:C:H2'	2:A2:1147:A:H8	1.63	0.60
31:J3:166:ARG:HH22	31:J3:252:THR:HG21	1.65	0.60
2:A2:3267:A:O2'	77:u2:92:ARG:CZ	2.49	0.60
6:B3:39:LEU:HD21	6:B3:56:ARG:HH21	1.66	0.60
20:F2:8:ILE:HD11	20:F2:257:PHE:CZ	2.35	0.60
34:L1:31:THR:HG21	34:L1:60:ARG:HD2	1.84	0.60
70:k2:47:LYS:HB2	70:k2:102:TYR:CZ	2.37	0.60
71:o2:198:MET:HG2	71:o2:200:ASP:H	1.66	0.60
83:m2:1009:C:H2'	83:m2:1010:A:C8	2.36	0.60
83:m2:1538:G:H2'	83:m2:1539:A:H8	1.66	0.60
2:A2:300:A:H2'	2:A2:301:G:H8	1.65	0.60
17:E2:50:LYS:HB2	17:E2:345:LEU:HD11	1.83	0.60
52:T3:48:THR:HG22	52:T3:51:LYS:HB2	1.84	0.60
83:m2:1856:U:H2'	83:m2:1857:G:H8	1.64	0.60
2:A2:1007:A:H2	2:A2:1017:G:H1	1.50	0.60
2:A2:1214:G:H2'	2:A2:1215:G:H8	1.66	0.60
14:D2:80:GLU:HB2	14:D2:170:ALA:HA	1.84	0.60
77:u2:141:ARG:HD2	77:u2:144:LYS:HD2	1.83	0.60
83:m2:1847:A:H2'	83:m2:1848:G:H8	1.66	0.60
2:A2:1668:G:H1	2:A2:4092:G:H21	1.49	0.60
2:A2:3517:A:H2'	2:A2:3518:A:H8	1.67	0.60
18:E3:46:HIS:CD2	18:E3:103:ALA:HB2	2.37	0.60
26:H2:186:LEU:HD12	26:H2:190:ARG:HA	1.83	0.60
2:A2:1377:A:OP1	40:N3:140:LYS:HD3	2.02	0.60
75:s2:17:ILE:HA	75:s2:48:TYR:HE1	1.67	0.60
2:A2:260:C:H2'	2:A2:261:G:H8	1.66	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:3325:G:H21	2:A2:3328:G:N2	2.00	0.59
2:A2:3602:G:H1	2:A2:3717:U:H3	1.50	0.59
3:A3:104:ASP:O	3:A3:108:ARG:HG3	2.03	0.59
60:a2:42:PRO:HB2	60:a2:53:LEU:HD12	1.83	0.59
1:A1:51:LYS:HE3	2:A2:1895:G:N1	2.18	0.59
38:M3:35:ILE:HD11	83:m2:1287:G:H5''	1.83	0.59
73:q2:50:ILE:HD11	73:q2:86:LEU:HD23	1.84	0.59
2:A2:1390:G:O2'	2:A2:1425:G:H4'	2.02	0.59
51:T2:79:HIS:HD2	60:a2:88:ARG:HH22	1.51	0.59
83:m2:181:A:C8	83:m2:182:C:H2'	2.37	0.59
2:A2:492:G:H22	2:A2:670:G:H1	1.50	0.59
19:F1:48:PRO:HG2	61:b2:118:LYS:HD2	1.83	0.59
37:M2:69:GLU:HG3	37:M2:71:SER:H	1.67	0.59
39:N2:64:VAL:HG22	39:N2:72:VAL:HG11	1.84	0.59
2:A2:4515:G:C6	22:G1:56:GLN:HG2	2.38	0.59
25:H1:185:GLY:HA3	25:H1:194:ARG:HH12	1.67	0.59
48:R3:98:LYS:HB3	48:R3:112:ASN:HB3	1.84	0.59
83:m2:1704:G:H2'	83:m2:1705:OMC:O4'	2.03	0.59
83:m2:1707:C:H2'	83:m2:1708:G:C8	2.38	0.59
2:A2:93:G:H2'	2:A2:94:A:C8	2.38	0.59
2:A2:2407:G:H22	69:j2:61:MET:HG2	1.68	0.59
2:A2:718:A:H2'	2:A2:719:C:C6	2.36	0.59
72:p2:163:GLN:HB3	72:p2:204:ILE:HD13	1.84	0.59
76:t2:146:VAL:HG22	76:t2:152:ARG:HG2	1.85	0.59
82:z2:29:HIS:HA	82:z2:32:LYS:HE2	1.85	0.59
1:A1:57:LYS:NZ	1:A1:61:LEU:HD11	2.18	0.59
32:K2:53:MET:HE1	32:K2:143:ARG:HH12	1.66	0.59
74:r2:182:MET:HE3	74:r2:192:ILE:HD11	1.84	0.59
80:x2:92:SER:HB2	80:x2:107:ILE:HD12	1.85	0.59
83:m2:180:G:H3'	83:m2:181:A:C8	2.37	0.59
2:A2:4526:U:H4'	2:A2:4527:U:H5	1.67	0.59
81:y2:15:ARG:HG2	81:y2:126:ARG:HH22	1.68	0.59
83:m2:1672:C:H2'	83:m2:1673:G:C8	2.38	0.59
2:A2:1088:A:H2'	2:A2:1089:G:H8	1.67	0.59
2:A2:2469:G:H2'	2:A2:2470:G:H8	1.68	0.59
11:C2:37:A:OP2	61:b2:89:ARG:HD2	2.03	0.59
77:u2:8:TRP:HA	77:u2:18:ARG:HD2	1.84	0.59
83:m2:1590:A:H2'	83:m2:1591:A:C8	2.38	0.59
2:A2:4400:U:H5''	59:Z2:54:LYS:HE3	1.85	0.58
30:J2:112:LEU:HD13	30:J2:150:LEU:HD23	1.84	0.58
83:m2:998:A:H2'	83:m2:999:A:C8	2.38	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
83:m2:1350:G:C2'	83:m2:1351:G:H5'	1.87	0.58
1:A1:259:GLU:HG2	37:M2:38:VAL:HG22	1.85	0.58
8:Bx:50:U:H1'	83:m2:1827:A:C8	2.38	0.58
7:Bz:18:G:H1	7:Bz:55:U:H1'	1.68	0.58
15:D3:11:LEU:HD23	15:D3:12:TYR:HD1	1.68	0.58
26:H2:157:LEU:HD13	26:H2:201:VAL:HG21	1.85	0.58
26:H2:160:LEU:HD11	26:H2:202:ILE:HG13	1.84	0.58
29:I3:44:LYS:HE2	29:I3:54:ILE:HD11	1.85	0.58
74:r2:51:ARG:HH21	74:r2:109:PHE:HB2	1.67	0.58
2:A2:2081:G:H5''	58:Y2:127:ALA:HB2	1.84	0.58
2:A2:3312:A:H2'	2:A2:3313:U:C6	2.37	0.58
5:B2:110:G:H2'	5:B2:111:C:C6	2.38	0.58
46:Q3:19:GLN:HE21	74:r2:94:LYS:NZ	2.02	0.58
56:W2:34:THR:HG21	56:W2:93:THR:HG22	1.86	0.58
62:c2:63:VAL:HG23	62:c2:65:LYS:HG3	1.84	0.58
3:A3:12:ILE:HB	16:E1:121:PRO:HG3	1.84	0.58
10:C1:113:GLU:HG2	10:C1:125:ARG:HG2	1.84	0.58
17:E2:332:MET:HE1	17:E2:368:ILE:HD13	1.84	0.58
21:F3:85[A]:ARG:HD2	83:m2:1211:A:O2'	2.04	0.58
21:F3:85[B]:ARG:HD2	83:m2:1211:A:O2'	2.04	0.58
83:m2:1350:G:N7	83:m2:1351:G:N7	2.52	0.58
83:m2:1847:A:H2'	83:m2:1848:G:C8	2.38	0.58
2:A2:1334:C:H5'	20:F2:95:MET:HE1	1.86	0.58
2:A2:2166:C:H2'	2:A2:2167:A:H8	1.68	0.58
2:A2:3923:A:H62	2:A2:3983:G:N2	2.02	0.58
72:p2:103:MET:HB3	72:p2:215:VAL:HG12	1.85	0.58
76:t2:43:LEU:HB3	76:t2:72:PHE:CE1	2.36	0.58
2:A2:3579:A:H2'	2:A2:3580:C:C6	2.38	0.58
7:Bz:63:G:H2'	7:Bz:64:A:C8	2.38	0.58
18:E3:36:LEU:HA	18:E3:39:ASN:HD21	1.67	0.58
31:J3:114:LYS:HG2	83:m2:1360:U:H5'	1.85	0.58
17:E2:348:ARG:HH12	17:E2:351:LEU:HG	1.67	0.58
28:I2:34:VAL:HG22	28:I2:103:LYS:HB3	1.86	0.58
71:o2:111:GLN:HG2	71:o2:112:ILE:HD12	1.85	0.58
83:m2:529:C:H2'	83:m2:530:A:C8	2.36	0.58
83:m2:984:G:H2'	83:m2:985:A:H8	1.69	0.58
8:Bx:50:U:H1'	83:m2:1827:A:H8	1.68	0.58
20:F2:7:LEU:HG	20:F2:21:ASN:HB3	1.85	0.58
29:I3:174:VAL:HB	29:I3:188:HIS:HB2	1.85	0.58
33:K3:2:LYS:HD3	33:K3:15:LEU:HD21	1.85	0.58
68:i2:22:LYS:HG3	68:i2:73:VAL:HG12	1.85	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
75:s2:68:ILE:HD12	75:s2:71:ARG:HD3	1.86	0.58
1:A1:108:GLU:HG3	39:N2:135:PRO:HB3	1.86	0.58
2:A2:1360:A:C2	2:A2:2568:A:H8	2.22	0.58
35:L2:72:LYS:HB3	35:L2:74:ARG:NH1	2.19	0.58
83:m2:27:A2M:HM'1	83:m2:485:C:H1'	1.86	0.58
83:m2:982:A:H2'	83:m2:983:A:C8	2.38	0.58
2:A2:433:A:C2	2:A2:3523:A:H4'	2.38	0.57
2:A2:3713:U:H2'	2:A2:3715:G:C8	2.39	0.57
3:A3:14:ARG:CB	16:E1:119:TYR:HB3	2.34	0.57
20:F2:144:ILE:O	20:F2:144:ILE:HG13	2.03	0.57
29:I3:83:TRP:HA	29:I3:107:ASP:HB2	1.86	0.57
36:L3:149:VAL:HG11	36:L3:157:ILE:HD11	1.86	0.57
8:Bx:44:U:C2	7:n2:34:G:O6	2.56	0.57
70:k2:63:VAL:HG22	70:k2:79:ARG:HG2	1.84	0.57
77:u2:3:ILE:H	77:u2:3:ILE:HD12	1.69	0.57
83:m2:909:G:H2'	83:m2:910:A:C8	2.39	0.57
83:m2:1846:U:H2'	83:m2:1847:A:C8	2.39	0.57
3:A3:97:GLN:HG3	3:A3:98:VAL:H	1.69	0.57
4:B1:207:VAL:HG21	4:B1:215:LEU:HD22	1.87	0.57
7:Bv:30:G:P	80:x2:136:THR:O	2.62	0.57
17:E2:317:LEU:HB2	17:E2:372:SER:HB2	1.86	0.57
40:N3:55:ARG:HD3	83:m2:1019:U:H5'	1.87	0.57
47:R2:87:MET:HE2	47:R2:156:ILE:HD12	1.84	0.57
71:o2:24:HIS:HB3	71:o2:51:LEU:HD11	1.86	0.57
42:O3:103:ASN:HD21	42:O3:142:ARG:HA	1.69	0.57
2:A2:4263:A:H2'	2:A2:4264:C:H6	1.70	0.57
71:o2:112:ILE:CD1	83:m2:1352:U:O2'	2.52	0.57
2:A2:424:U:H2'	2:A2:425:U:C6	2.39	0.57
2:A2:4500:G:H2'	2:A2:4501:G:H8	1.70	0.57
3:A3:5:ILE:HD12	3:A3:6:PRO:HD2	1.85	0.57
13:D1:54:SER:HB2	13:D1:135:ILE:HD11	1.86	0.57
24:G3:62:GLU:OE1	42:O3:121:ARG:HD3	2.05	0.57
24:G3:62:GLU:CG	42:O3:121:ARG:NE	2.62	0.57
71:o2:42:LYS:HD2	71:o2:46:ILE:HB	1.85	0.57
83:m2:589:A:H5'	83:m2:594:C:H41	1.70	0.57
2:A2:176:G:H2'	2:A2:177:G:H8	1.69	0.57
2:A2:2275:C:H2'	2:A2:2276:G:H8	1.70	0.57
2:A2:3566:C:H2'	2:A2:3567:C:H6	1.69	0.57
2:A2:4613:A:H2'	2:A2:4614:A:H8	1.69	0.57
4:B1:137[A]:ARG:HG2	4:B1:142:THR:HG21	1.86	0.57
23:G2:211:LEU:HD12	23:G2:223:PHE:HE2	1.70	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
83:m2:1800:C:H2'	83:m2:1801:G:O4'	2.05	0.57
33:K3:188:LYS:HD3	83:m2:142:C:H42	1.69	0.57
44:P3:105:THR:HG22	44:P3:110:ILE:HG12	1.86	0.57
2:A2:444:G:H2'	2:A2:445:U:H6	1.69	0.57
2:A2:2389:C:H2'	2:A2:2390:U:H6	1.70	0.57
2:A2:3815:U:H5'	2:A2:3816:C:H5''	1.87	0.57
19:F1:87:HIS:HE1	19:F1:89:LYS:HD2	1.70	0.57
29:I3:45:LEU:HD23	29:I3:47:ARG:HE	1.68	0.57
42:O3:26:ASN:C	42:O3:26:ASN:HD22	2.10	0.57
72:p2:144:LYS:HE2	72:p2:208:HIS:HB3	1.86	0.57
79:w2:59:LYS:HD3	79:w2:134:LEU:HD23	1.87	0.57
83:m2:971:U:H4'	83:m2:972:G:H8	1.70	0.57
83:m2:1103:U:H2'	83:m2:1104:G:H8	1.70	0.57
28:I2:165:LYS:HG3	28:I2:169:ARG:HH21	1.70	0.57
29:I3:17:TRP:HB2	29:I3:36:ARG:HG2	1.87	0.57
75:s2:81:ARG:HH21	83:m2:1538:G:H4'	1.69	0.57
83:m2:96:C:H2'	83:m2:97:U:C6	2.40	0.57
83:m2:1130:C:H2'	83:m2:1131:G:C8	2.40	0.57
83:m2:1201:A:H2'	83:m2:1202:A:C8	2.40	0.57
2:A2:139:G:H2'	2:A2:140:G:C8	2.40	0.56
51:T2:47:ASP:HB3	51:T2:69:LYS:HG2	1.85	0.56
70:k2:28:GLU:HG2	70:k2:31:ASN:HB2	1.87	0.56
2:A2:4505:G:H2'	2:A2:4506:G:C8	2.40	0.56
11:C2:29:G:H5''	19:F1:27:ASN:HB3	1.86	0.56
14:D2:117:GLU:HG2	14:D2:124:GLY:H	1.70	0.56
17:E2:224:LYS:HG2	17:E2:340:THR:HG22	1.85	0.56
50:S3:21:LYS:HE3	83:m2:923:G:H5'	1.85	0.56
7:n2:69:G:H2'	7:n2:70:G:C8	2.40	0.56
78:v2:12:TYR:HE1	78:v2:52:LEU:HD11	1.69	0.56
80:x2:77:LYS:HG2	80:x2:102:PHE:CZ	2.40	0.56
83:m2:15:U:H2'	83:m2:16:G:O4'	2.05	0.56
2:A2:223:G:H4'	2:A2:225:G:N7	2.20	0.56
2:A2:4613:A:H2'	2:A2:4614:A:C8	2.40	0.56
25:H1:120:TRP:HZ2	25:H1:123:GLU:HB2	1.70	0.56
29:I3:206:LEU:HD13	29:I3:218:LEU:HD12	1.87	0.56
31:J3:86:LEU:HD23	71:o2:141:ASN:HD22	1.68	0.56
31:J3:165:VAL:HG11	31:J3:217:ALA:HB1	1.88	0.56
41:O2:100:LEU:HD23	41:O2:112:LEU:HD23	1.87	0.56
50:S3:34:ASP:OD1	50:S3:43:ILE:HG22	2.06	0.56
76:t2:83:LEU:HD22	76:t2:92:VAL:HG11	1.87	0.56
76:t2:100:ILE:HG13	76:t2:125:VAL:HG21	1.87	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
77:u2:27:TYR:HB2	83:m2:383:C:N4	2.21	0.56
83:m2:375:G:N2	83:m2:394:A:H61	2.04	0.56
83:m2:390:U:H2'	83:m2:391:A:C8	2.40	0.56
12:C3:49:LYS:HB3	12:C3:92:HIS:HB3	1.86	0.56
20:F2:221:PHE:HB2	20:F2:229:LEU:HD21	1.86	0.56
83:m2:927:G:H1	83:m2:1019:U:H3	1.53	0.56
2:A2:4600:G:H2'	2:A2:4601:A:C8	2.41	0.56
35:L2:44:LEU:HD22	35:L2:49:LEU:HD12	1.88	0.56
51:T2:77:TYR:HB3	56:W2:39:ARG:HH11	1.71	0.56
67:h2:4:LYS:HD2	83:m2:1846:U:O4	2.06	0.56
71:o2:109:THR:HG22	83:m2:1353:G:C4'	2.31	0.56
73:q2:138:VAL:HG22	73:q2:184:ILE:HD12	1.86	0.56
2:A2:140:G:H2'	2:A2:141:C:C6	2.41	0.56
16:E1:51:SER:HB2	16:E1:69:ALA:HB3	1.86	0.56
28:I2:8:VAL:HG12	28:I2:117:ARG:HG3	1.87	0.56
76:t2:170:VAL:HG23	76:t2:187:PHE:HB2	1.88	0.56
83:m2:12:U:H2'	83:m2:13:C:C6	2.41	0.56
83:m2:435:A:H2'	83:m2:436:G:C8	2.40	0.56
83:m2:528:A:H2'	83:m2:529:C:C6	2.41	0.56
1:A1:201:LEU:HD21	1:A1:226:ALA:HA	1.88	0.56
2:A2:1054:G:H2'	2:A2:1055:G:C8	2.41	0.56
33:K3:154:ARG:HB3	83:m2:77:A:O5'	2.05	0.56
67:h2:11:ARG:NH2	83:m2:1846:U:OP2	2.37	0.56
71:o2:184:ARG:HD3	71:o2:191:ARG:HG3	1.88	0.56
83:m2:51:U:H2'	83:m2:52:G:H8	1.69	0.56
83:m2:107:A:H2'	83:m2:108:G:C8	2.41	0.56
83:m2:982:A:H2'	83:m2:983:A:H8	1.71	0.56
2:A2:1430:G:H1'	2:A2:2268:A:N6	2.20	0.56
2:A2:1541:G:H2'	2:A2:1542:C:C6	2.41	0.56
2:A2:3613:U:H1'	2:A2:3615:U:H5'	1.87	0.56
2:A2:4500:G:H2'	2:A2:4501:G:C8	2.39	0.56
28:I2:46:ASN:HB3	28:I2:134:LYS:HD3	1.86	0.56
83:m2:14:C:H2'	83:m2:15:U:C6	2.40	0.56
12:C3:40:ILE:HD11	12:C3:89:ILE:HD12	1.88	0.56
16:E1:120:ASP:HB3	16:E1:123:ILE:HG12	1.87	0.56
67:h2:5:TRP:HD1	83:m2:1854:C:OP2	1.89	0.56
72:p2:168:MET:HG2	72:p2:197:ILE:HG21	1.87	0.56
74:r2:148:ARG:NH1	83:m2:124:U:H5''	2.21	0.56
76:t2:64:VAL:HB	76:t2:72:PHE:CE2	2.41	0.56
2:A2:101:A:H1'	53:U2:66:ASN:ND2	2.20	0.56
2:A2:3298:A:H1'	63:d2:2:THR:HB	1.88	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:3565:C:H5	2:A2:4048:A:H61	1.54	0.56
2:A2:4582:U:H4'	2:A2:4583:U:OP1	2.05	0.56
29:I3:66:VAL:HA	29:I3:82:SER:HA	1.88	0.56
32:K2:61:LEU:HD11	32:K2:65:ARG:HG2	1.87	0.56
73:q2:9:ARG:HA	73:q2:12:VAL:HG22	1.87	0.56
83:m2:186:C:H2'	83:m2:187:G:C8	2.40	0.56
2:A2:6:C:H5''	4:B1:197:LYS:HB3	1.89	0.55
2:A2:72:C:H5	19:F1:67:HIS:HD1	1.54	0.55
2:A2:746:G:O2'	2:A2:747:G:H8	1.88	0.55
4:B1:110:LYS:HD2	4:B1:113:ARG:HH12	1.70	0.55
31:J3:114:LYS:CG	83:m2:1360:U:OP1	2.51	0.55
32:K2:96:PRO:HG2	32:K2:98:LEU:HD21	1.88	0.55
53:U2:75:LEU:HD22	53:U2:117:LEU:HD11	1.88	0.55
67:h2:9:ARG:NH2	83:m2:1853:A:P	2.79	0.55
75:s2:19:LEU:HG	75:s2:20:PHE:CD1	2.41	0.55
81:y2:97:GLN:HB2	81:y2:105:LYS:NZ	2.21	0.55
83:m2:1799:U:H2'	83:m2:1800:C:C6	2.39	0.55
2:A2:163:A:H2'	2:A2:164:G:C8	2.42	0.55
2:A2:3524:G:H22	2:A2:3556:G:H1'	1.71	0.55
2:A2:3624:U:H2'	2:A2:3625:G:C8	2.41	0.55
47:R2:64:SER:HB2	61:b2:69:LEU:HD13	1.88	0.55
82:z2:46:LEU:O	82:z2:50:ILE:HG12	2.05	0.55
83:m2:1191:A:H2'	83:m2:1192:A:C8	2.41	0.55
2:A2:939:C:H2'	2:A2:940:C:H6	1.70	0.55
29:I3:108:VAL:HG12	29:I3:124:SER:HB3	1.88	0.55
72:p2:87:ILE:HB	72:p2:101:HIS:HB2	1.88	0.55
73:q2:75:LYS:NZ	78:v2:18:GLU:HG2	2.21	0.55
74:r2:45:ILE:HD11	74:r2:80:ILE:HD12	1.87	0.55
83:m2:26:U:H2'	83:m2:27:A2M:H8	1.88	0.55
83:m2:1219:A:H2'	83:m2:1220:C:C6	2.42	0.55
1:A1:56:ARG:HH12	2:A2:1252:C:H4'	1.72	0.55
2:A2:1870:C:O2'	2:A2:1871:A:H5'	2.07	0.55
13:D1:51:HIS:HB3	13:D1:134:VAL:HG13	1.89	0.55
13:D1:90:ARG:O	13:D1:91:LEU:HD12	2.07	0.55
27:H3:13:LYS:HD2	83:m2:1558:A:H8	1.71	0.55
79:w2:133:PRO:HG2	83:m2:385:G:H21	1.70	0.55
2:A2:4253:U:H2'	2:A2:4254:A:H8	1.71	0.55
20:F2:301:ALA:HB1	32:K2:132:LYS:HE3	1.89	0.55
23:G2:216:GLU:HG3	23:G2:220:LYS:HE2	1.89	0.55
41:O2:84:LYS:HA	41:O2:87:THR:HG22	1.88	0.55
74:r2:45:ILE:HG23	74:r2:61:VAL:HG21	1.88	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
80:x2:49:LEU:HD22	80:x2:53:GLN:HG2	1.89	0.55
83:m2:444:C:H2'	83:m2:445:U:C6	2.41	0.55
83:m2:1397:C:H2'	83:m2:1398:A:C8	2.41	0.55
1:A1:129:LYS:HD2	1:A1:132:GLN:HE21	1.71	0.55
2:A2:3320:G:H2'	2:A2:3321:G:H8	1.71	0.55
2:A2:4023:G:H5'	7:n2:76:A:H62	1.71	0.55
4:B1:187:LYS:HD2	4:B1:198:THR:HG23	1.88	0.55
7:n2:21:A:H61	7:n2:46:G:H2'	1.71	0.55
83:m2:85:A:O2'	83:m2:149:A:H8	1.90	0.55
83:m2:842:C:H4'	83:m2:843:G:O5'	2.07	0.55
2:A2:444:G:H2'	2:A2:445:U:C6	2.42	0.55
2:A2:1153:U:H2'	2:A2:1154:OMC:C6	2.42	0.55
2:A2:3923:A:O2'	2:A2:3924:G:H5'	2.05	0.55
12:C3:22:ILE:HB	12:C3:89:ILE:HB	1.88	0.55
34:L1:181:TYR:HA	34:L1:184:HIS:CE1	2.41	0.55
74:r2:11:ARG:HH21	83:m2:497:U:H4'	1.71	0.55
83:m2:641:C:H2'	83:m2:642:A:C8	2.40	0.55
83:m2:944:G:H2'	83:m2:945:U:C6	2.42	0.55
83:m2:1146:A:H2'	83:m2:1147:A:C8	2.42	0.55
83:m2:1623:U:H5'	83:m2:1625:A:O2'	2.07	0.55
2:A2:955:U:H2'	2:A2:956:C:C6	2.41	0.55
30:J2:122:ALA:HB3	30:J2:143:PRO:HG2	1.88	0.55
71:o2:37:TYR:HA	71:o2:53:ARG:HG3	1.87	0.55
83:m2:472:G:H2'	83:m2:473:G:C8	2.41	0.55
83:m2:658:G:H21	83:m2:665:C:H5''	1.72	0.55
27:H3:50:ILE:HG23	73:q2:15:GLY:HA3	1.87	0.55
67:h2:9:ARG:HH21	83:m2:1853:A:P	2.29	0.55
72:p2:66:VAL:HB	72:p2:87:ILE:HD12	1.89	0.55
83:m2:369:U:H4'	83:m2:373:A:C8	2.41	0.55
83:m2:1162:U:H2'	83:m2:1163:U:H6	1.72	0.55
2:A2:744:C:H3'	2:A2:746:G:H5''	1.89	0.54
2:A2:1050:C:H2'	2:A2:1051:G:C8	2.41	0.54
2:A2:1068:C:N4	2:A2:1082:A:H61	2.05	0.54
17:E2:231:VAL:HG21	17:E2:251:VAL:HG23	1.88	0.54
36:L3:136:ARG:HD2	36:L3:139:LYS:HA	1.89	0.54
66:g2:94:MET:HG2	66:g2:105:PRO:HA	1.88	0.54
71:o2:132:GLN:NE2	83:m2:1380:A:C8	2.73	0.54
72:p2:146:ARG:NH1	72:p2:206:PRO:HB2	2.22	0.54
76:t2:64:VAL:HB	76:t2:72:PHE:HE2	1.72	0.54
1:A1:231:TRP:CD1	1:A1:232:PRO:HD2	2.42	0.54
4:B1:138:ALA:HB2	4:B1:194:VAL:HG11	1.89	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:Bz:69:G:H2'	7:Bz:70:G:C8	2.39	0.54
22:G1:13:ALA:HB1	22:G1:55:MET:HB2	1.89	0.54
24:G3:49:PRO:HB2	75:s2:61:PHE:CE1	2.42	0.54
32:K2:177:ALA:O	32:K2:184:ARG:HB2	2.07	0.54
33:K3:66:GLY:HA2	83:m2:1747:A:H1'	1.89	0.54
77:u2:106:SER:HB3	77:u2:171:LEU:HG	1.89	0.54
2:A2:1891:G:H1	2:A2:1897:A:H4'	1.73	0.54
2:A2:2651:G:H5''	35:L2:134:ASN:OD1	2.07	0.54
2:A2:4526:U:C4	22:G1:113:MET:HG2	2.42	0.54
83:m2:630:A:N6	83:m2:1502:G:C2	2.70	0.54
83:m2:931:G:H2'	83:m2:932:C:O4'	2.07	0.54
2:A2:3517:A:H2'	2:A2:3518:A:C8	2.43	0.54
4:B1:242:LEU:HB3	4:B1:246:SER:HB2	1.90	0.54
18:E3:93:PHE:HB3	18:E3:133:LEU:HD13	1.88	0.54
70:k2:47:LYS:HG3	70:k2:103:ARG:HD3	1.88	0.54
73:q2:65:ARG:O	73:q2:68:GLU:HG2	2.07	0.54
83:m2:126:G:C8	83:m2:181:A:H1'	2.42	0.54
83:m2:1860:G:H2'	83:m2:1861:A:H8	1.72	0.54
2:A2:1214:G:H2'	2:A2:1215:G:C8	2.43	0.54
2:A2:2167:A:H2'	2:A2:2168:U:C6	2.43	0.54
4:B1:37:LYS:HB2	4:B1:44:ASP:OD2	2.08	0.54
4:B1:171:PRO:HB3	4:B1:181:TYR:CE2	2.42	0.54
14:D2:133:TYR:HB3	14:D2:168:VAL:HG12	1.90	0.54
83:m2:456:U:H2'	83:m2:457:A:H8	1.71	0.54
83:m2:622:G:H2'	83:m2:622:G:N3	2.22	0.54
83:m2:1714:A:H2'	83:m2:1715:C:C6	2.43	0.54
83:m2:1841:U:H2'	83:m2:1842:U:C6	2.43	0.54
1:A1:111:LEU:HD11	1:A1:144:PHE:HB3	1.90	0.54
2:A2:3932:A:H5''	2:A2:3933:A:H5'	1.90	0.54
2:A2:4520:G:C2	28:I2:118:MET:HE1	2.42	0.54
15:D3:60:ARG:HH22	71:o2:144:THR:HG23	1.72	0.54
22:G1:11:ARG:HB3	22:G1:27:ILE:HD12	1.88	0.54
24:G3:40:ARG:CZ	24:G3:40:ARG:HA	2.37	0.54
34:L1:51:GLY:HA3	34:L1:189:PHE:HZ	1.73	0.54
52:T3:34:ARG:HA	52:T3:37:GLN:HE21	1.73	0.54
75:s2:187:SER:HB3	75:s2:190:ILE:HG22	1.89	0.54
83:m2:892:U:H2'	83:m2:893:G:H3'	1.90	0.54
83:m2:959:A:H2'	83:m2:960:G:C8	2.42	0.54
2:A2:175:C:H2'	2:A2:176:G:H8	1.73	0.54
2:A2:2320:A:H3'	2:A2:2321:G:C8	2.43	0.54
2:A2:3363:U:H2'	2:A2:3364:C:C6	2.43	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:H2:164:HIS:HB3	26:H2:167:LYS:HG3	1.88	0.54
31:J3:210:PRO:HD3	31:J3:236:PHE:HE2	1.72	0.54
36:L3:144:ILE:HG21	83:m2:826:C:C2	2.43	0.54
83:m2:12:U:H2'	83:m2:13:C:H6	1.73	0.54
2:A2:3263:U:H2'	2:A2:3264:A:H8	1.72	0.54
2:A2:4274:A:H4'	17:E2:13:SER:HB2	1.90	0.54
19:F1:60:ARG:HD2	19:F1:67:HIS:O	2.08	0.54
38:M3:20:GLU:HA	38:M3:23:LYS:HG2	1.89	0.54
64:e2:24:LYS:HD2	64:e2:69:LEU:HD21	1.89	0.54
72:p2:165:ARG:NH1	83:m2:1005:U:H5''	2.23	0.54
74:r2:66:MET:HG3	83:m2:504:C:O4'	2.07	0.54
77:u2:161:LEU:HD12	77:u2:199:LEU:HD12	1.89	0.54
2:A2:4433:G:HO2'	2:A2:4434:U:H6	1.56	0.54
29:I3:18:VAL:HG21	29:I3:307:VAL:HG22	1.90	0.54
48:R3:49:LEU:HD23	48:R3:83:LEU:HD21	1.89	0.54
69:j2:47:MET:HE2	69:j2:71:TYR:HE1	1.72	0.54
2:A2:1333:C:H5'	20:F2:102:PHE:HE1	1.73	0.54
2:A2:2456:U:H2'	2:A2:2457:C:C6	2.43	0.54
2:A2:2651:G:H5''	35:L2:134:ASN:ND2	2.23	0.54
18:E3:105:PHE:HD1	18:E3:121:LYS:HB3	1.72	0.54
36:L3:42:GLU:HG2	36:L3:45:ARG:HH21	1.73	0.54
53:U2:101:ILE:HD12	53:U2:122:VAL:HG21	1.89	0.54
60:a2:48:VAL:O	69:j2:61:MET:HG3	2.08	0.54
7:n2:15:G:N2	7:n2:21:A:H1'	2.23	0.54
80:x2:81:ARG:HD3	83:m2:1517:G:O3'	2.08	0.54
83:m2:1407:A:H2'	83:m2:1408:G:O4'	2.08	0.54
83:m2:1493:G:H2'	83:m2:1494:U:C6	2.43	0.54
83:m2:1738:G:H2'	83:m2:1739:G:H8	1.72	0.54
2:A2:2649:A:H2'	2:A2:2650:A:H8	1.73	0.53
22:G1:6:TYR:CG	37:M2:151:LYS:HE3	2.43	0.53
33:K3:183:ARG:HD3	33:K3:186:GLN:HE21	1.73	0.53
38:M3:57:ASP:HB2	83:m2:1287:G:H22	1.72	0.53
73:q2:136:VAL:HG22	73:q2:186:VAL:HG22	1.88	0.53
76:t2:66:VAL:HG11	83:m2:915:A:N3	2.24	0.53
81:y2:108:ILE:HG13	81:y2:109:LYS:N	2.23	0.53
83:m2:1094:G:H2'	83:m2:1095:A:H8	1.73	0.53
83:m2:1136:G:H2'	83:m2:1137:C:C6	2.43	0.53
83:m2:1742:C:H2'	83:m2:1743:U:C6	2.43	0.53
2:A2:3744:G:H22	2:A2:3765:G:N2	2.05	0.53
6:B3:65:TYR:HE2	6:B3:128:GLN:HG3	1.72	0.53
18:E3:105:PHE:CD2	18:E3:112:VAL:HB	2.43	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
71:o2:28:THR:HG22	71:o2:46:ILE:HG13	1.90	0.53
83:m2:75:G:H4'	83:m2:76:U:C5	2.43	0.53
2:A2:2320:A:H3'	2:A2:2321:G:H8	1.74	0.53
31:J3:193:VAL:HG11	31:J3:240:THR:HG22	1.91	0.53
51:T2:46:ILE:HG23	51:T2:68:ILE:HG23	1.90	0.53
55:V2:110:MET:O	55:V2:114:GLN:HG2	2.08	0.53
65:f2:44:TRP:CH2	65:f2:45:ARG:HD2	2.42	0.53
70:k2:51:VAL:HG12	70:k2:117:ILE:HD12	1.90	0.53
2:A2:4638:G:H2'	2:A2:4639:G:C8	2.43	0.53
6:B3:60:THR:HG23	6:B3:75:MET:HE2	1.89	0.53
22:G1:51:PRO:O	22:G1:55:MET:HE3	2.08	0.53
27:H3:34:TYR:O	27:H3:36:LEU:HG	2.09	0.53
53:U2:110:LYS:HG3	53:U2:128:PHE:HB2	1.90	0.53
56:W2:26:LYS:H	56:W2:98:ASP:HB3	1.73	0.53
77:u2:57:ALA:HB2	77:u2:183:GLY:HA2	1.90	0.53
78:v2:11:ILE:HG12	78:v2:45:VAL:HG22	1.90	0.53
83:m2:351:A:H2'	83:m2:352:C:C6	2.43	0.53
83:m2:636:A:H2'	83:m2:637:G:H8	1.73	0.53
83:m2:993:G:C6	83:m2:1136:G:H4'	2.43	0.53
2:A2:835:U:H5	22:G1:46:ARG:HH11	1.55	0.53
2:A2:2387:U:H2'	2:A2:2388:U:C6	2.44	0.53
2:A2:2475:C:H2'	2:A2:2476:G:O4'	2.08	0.53
16:E1:18:ARG:HG3	16:E1:135:GLY:HA3	1.91	0.53
74:r2:68:ARG:NH2	74:r2:76:VAL:HG11	2.24	0.53
77:u2:103:LEU:HG	77:u2:170:LYS:HB3	1.91	0.53
83:m2:1141:C:H2'	83:m2:1142:G:O4'	2.09	0.53
2:A2:418:A:C2	11:C2:17:A:H1'	2.43	0.53
2:A2:1027:G:H2'	2:A2:1028:G:H8	1.73	0.53
2:A2:1088:A:H2'	2:A2:1089:G:C8	2.43	0.53
2:A2:1146:C:H2'	2:A2:1147:A:C8	2.43	0.53
2:A2:4109:U:H1'	17:E2:252:ALA:HB3	1.89	0.53
22:G1:7:VAL:HG13	22:G1:27:ILE:HD13	1.89	0.53
31:J3:183:LYS:HA	31:J3:195:LEU:O	2.07	0.53
78:v2:41:PRO:HG2	78:v2:44:HIS:HD2	1.74	0.53
83:m2:172:OMU:HM22	83:m2:172:OMU:O2	2.09	0.53
83:m2:347:U:H2'	83:m2:348:C:C6	2.43	0.53
1:A1:98:ARG:HD2	2:A2:736:G:H5''	1.90	0.53
2:A2:3950:A:H4'	55:V2:33:LYS:HD3	1.91	0.53
64:e2:12:LEU:HD13	64:e2:16:ARG:HH12	1.73	0.53
76:t2:144:ILE:HG13	76:t2:154:ILE:HG13	1.90	0.53
83:m2:354:U:H2'	83:m2:355:OMC:C6	2.43	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:1527:U:H2'	2:A2:1528:U:H6	1.74	0.53
27:H3:14:PHE:HD1	83:m2:1663:A:C8	2.27	0.53
43:P2:32:THR:HG21	43:P2:105:ILE:HD12	1.89	0.53
57:X2:37:GLY:O	57:X2:41:ARG:HG3	2.08	0.53
67:h2:11:ARG:NH2	83:m2:1846:U:OP1	2.40	0.53
71:o2:54:THR:HG22	71:o2:162:PRO:HG2	1.91	0.53
83:m2:72:C:H3'	83:m2:74:G:H21	1.73	0.53
83:m2:390:U:H2'	83:m2:391:A:H8	1.74	0.53
1:A1:81:LYS:O	1:A1:85:GLN:HG3	2.09	0.53
2:A2:4067:1MA:H2'	2:A2:4068:G:O4'	2.09	0.53
2:A2:4351:U:H1'	2:A2:4352:A:H5''	1.90	0.53
17:E2:220:ILE:HB	17:E2:346:THR:HB	1.91	0.53
72:p2:59:SER:O	72:p2:63:LYS:HG2	2.09	0.53
72:p2:105:LEU:HD13	72:p2:213:ARG:HA	1.90	0.53
83:m2:25:A:O2'	83:m2:26:U:H5''	2.09	0.53
83:m2:96:C:H1'	83:m2:476:G:H5'	1.90	0.53
83:m2:578:A2M:HM'2	83:m2:579:U:H5'	1.91	0.53
83:m2:954:G:H2'	83:m2:955:C:C6	2.44	0.53
3:A3:12:ILE:CD1	16:E1:121:PRO:HB3	2.36	0.53
19:F1:87:HIS:HB3	19:F1:90:VAL:HG22	1.91	0.53
21:F3:88:SER:O	21:F3:92:ARG:HG3	2.09	0.53
29:I3:199:THR:HG21	29:I3:240:CYS:HA	1.89	0.53
83:m2:220:U:H2'	83:m2:221:U:C6	2.44	0.53
83:m2:434:G:H2'	83:m2:435:A:C8	2.44	0.53
2:A2:1760:A:H8	2:A2:1760:A:OP2	1.92	0.52
2:A2:3536:G:H2'	2:A2:3537:G:C8	2.43	0.52
19:F1:64:VAL:HA	19:F1:67:HIS:CD2	2.44	0.52
29:I3:52:TYR:HE2	29:I3:311:GLN:HE21	1.57	0.52
31:J3:102:LEU:HB3	31:J3:130:ILE:HD11	1.91	0.52
38:M3:24:THR:HA	38:M3:27:ILE:HG22	1.91	0.52
73:q2:162:ASP:N	73:q2:163:PRO:HD2	2.23	0.52
83:m2:208:G:H2'	83:m2:209:G:C8	2.43	0.52
83:m2:346:U:H2'	83:m2:347:U:H6	1.74	0.52
83:m2:1547:A:H2'	83:m2:1548:G:C8	2.43	0.52
2:A2:432:U:H4'	2:A2:433:A:H5''	1.91	0.52
2:A2:673:G:C8	2:A2:675:C:H2'	2.44	0.52
2:A2:2464:C:H5''	35:L2:39:GLN:HG3	1.91	0.52
2:A2:4200:A:C8	7:Bz:75:C:H5'	2.44	0.52
56:W2:35:LEU:HB3	56:W2:39:ARG:HH21	1.74	0.52
7:n2:20:U:H4'	7:n2:21:A:OP1	2.09	0.52
82:z2:20:TYR:O	82:z2:24:LEU:HD23	2.10	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
83:m2:457:A:H2'	83:m2:458:C:H6	1.74	0.52
83:m2:1096:C:H2'	83:m2:1097:U:C6	2.45	0.52
83:m2:1716:U:H2'	83:m2:1717:A:C8	2.43	0.52
2:A2:4264:C:C2	10:C1:120:GLU:HB2	2.44	0.52
3:A3:19:ASN:CG	16:E1:119:TYR:CE2	2.88	0.52
51:T2:41:ALA:HB2	51:T2:77:TYR:HE1	1.74	0.52
74:r2:19:MET:HE3	74:r2:51:ARG:CZ	2.39	0.52
83:m2:1595:C:H2'	83:m2:1596:A:C8	2.45	0.52
83:m2:1714:A:H2'	83:m2:1715:C:H6	1.74	0.52
2:A2:517:C:H2'	2:A2:518:C:C6	2.45	0.52
2:A2:1708:U:H2'	2:A2:1709:A:H8	1.75	0.52
2:A2:2500:A:H2'	2:A2:2501:A:C8	2.45	0.52
2:A2:2532:G:H5''	2:A2:2533:G:H5'	1.91	0.52
69:j2:37:TYR:HB2	69:j2:47:MET:HB3	1.92	0.52
83:m2:1200:G:H2'	83:m2:1201:A:H8	1.74	0.52
83:m2:1350:G:C5	83:m2:1351:G:N7	2.78	0.52
83:m2:1597:U:H2'	83:m2:1598:U:H6	1.74	0.52
2:A2:937:C:H2'	2:A2:938:U:H5'	1.90	0.52
2:A2:4506:G:H2'	2:A2:4507:G:C8	2.45	0.52
23:G2:166:ALA:HB1	23:G2:171:LEU:HD12	1.90	0.52
29:I3:32:LEU:HB3	29:I3:42:MET:HG2	1.91	0.52
40:N3:40:LEU:HB3	40:N3:45:LEU:HD12	1.92	0.52
47:R2:95:THR:HG22	47:R2:139:ARG:HA	1.91	0.52
52:T3:37:GLN:HB2	52:T3:41:ARG:HH12	1.75	0.52
64:e2:14:THR:HA	64:e2:17:ARG:HD3	1.92	0.52
74:r2:148:ARG:HH11	83:m2:124:U:H5''	1.74	0.52
75:s2:60:ARG:HD2	83:m2:1681:A:H2'	1.91	0.52
83:m2:1378:A:H2'	83:m2:1379:U:O4'	2.09	0.52
83:m2:1515:C:H2'	83:m2:1516:G:C8	2.44	0.52
2:A2:21:G:H5'	63:d2:43:ARG:HG2	1.90	0.52
52:T3:37:GLN:HB2	52:T3:41:ARG:NH1	2.24	0.52
67:h2:21:ARG:HD3	83:m2:1719:C:H4'	1.91	0.52
83:m2:346:U:H2'	83:m2:347:U:C6	2.45	0.52
83:m2:1162:U:H2'	83:m2:1163:U:C6	2.45	0.52
83:m2:1564:C:H2'	83:m2:1565:G:C8	2.45	0.52
2:A2:453:G:H1	2:A2:1107:G:H22	1.56	0.52
2:A2:3297:U:H5	2:A2:3302:A:N7	2.08	0.52
10:C1:103:VAL:HG11	10:C1:144:LEU:HD11	1.92	0.52
10:C1:118:LEU:HD11	10:C1:167:VAL:HG22	1.92	0.52
32:K2:79:THR:HG23	32:K2:99:LYS:HG2	1.91	0.52
42:O3:50:LYS:HD3	72:p2:65:ARG:HE	1.74	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
71:o2:32:PHE:CD1	83:m2:1099:G:H4'	2.44	0.52
71:o2:145:ILE:HG12	71:o2:159:ILE:HB	1.92	0.52
83:m2:528:A:H2'	83:m2:529:C:H6	1.75	0.52
83:m2:835:C:N4	83:m2:836:C:N4	2.58	0.52
83:m2:1339:B8T:O3'	83:m2:1340:G:H5'	2.09	0.52
83:m2:1669:U:H2'	83:m2:1670:U:C6	2.44	0.52
2:A2:261:G:H2'	2:A2:262:G:H8	1.74	0.52
2:A2:3825:G:H2'	2:A2:3826:U:C6	2.44	0.52
41:O2:27:HIS:HB2	41:O2:28:PRO:HD3	1.91	0.52
51:T2:74:VAL:HG23	51:T2:101:PHE:CE2	2.45	0.52
58:Y2:84:GLU:O	58:Y2:87:VAL:HG12	2.10	0.52
76:t2:95:ILE:HD11	76:t2:133:LEU:HD23	1.91	0.52
83:m2:434:G:H2'	83:m2:435:A:H8	1.74	0.52
83:m2:636:A:H2'	83:m2:637:G:C8	2.45	0.52
2:A2:233:U:H4'	2:A2:234:G:OP1	2.09	0.52
2:A2:325:U:H2'	2:A2:326:C:C6	2.44	0.52
2:A2:388:A:H1'	2:A2:403:G:N2	2.25	0.52
2:A2:440:U:H2'	2:A2:441:G:C8	2.45	0.52
2:A2:3415:A:C2	83:m2:1710:C:O2	2.63	0.52
2:A2:3912:U:H2'	2:A2:3913:C:C6	2.45	0.52
10:C1:93:ARG:HD2	10:C1:143:GLU:HB3	1.92	0.52
31:J3:75:ILE:HD11	31:J3:265:PRO:HB3	1.92	0.52
32:K2:167:VAL:HG22	32:K2:175:GLU:OE1	2.09	0.52
37:M2:15:ARG:HB3	37:M2:27:LEU:HD23	1.90	0.52
71:o2:68:ILE:HG21	71:o2:74:VAL:HG13	1.91	0.52
83:m2:470:A:H2'	83:m2:471:A:C8	2.45	0.52
83:m2:1707:C:H2'	83:m2:1708:G:H8	1.74	0.52
2:A2:325:U:H5''	19:F1:103:ARG:NH2	2.25	0.52
2:A2:1259:C:N4	2:A2:1901:A:H61	1.97	0.52
2:A2:1444:A:H2	14:D2:208:GLU:HG3	1.75	0.52
2:A2:1735:G:H2'	2:A2:1736:A:C8	2.44	0.52
51:T2:12:LEU:HD23	51:T2:22:LYS:HG2	1.91	0.52
63:d2:21:ARG:HB2	63:d2:39:TYR:CD1	2.44	0.52
83:m2:658:G:H5'	83:m2:664:G:N2	2.25	0.52
83:m2:806:U:H2'	83:m2:807:U:C6	2.45	0.52
83:m2:846:U:H2'	83:m2:847:G:C8	2.45	0.52
83:m2:1099:G:H2'	83:m2:1100:C:C6	2.45	0.52
2:A2:176:G:H2'	2:A2:177:G:C8	2.45	0.51
2:A2:3267:A:C2'	77:u2:92:ARG:CZ	2.84	0.51
6:B3:4:VAL:N	83:m2:1418:C:HO2'	2.08	0.51
83:m2:107:A:H2'	83:m2:108:G:H8	1.73	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
83:m2:534:C:H42	83:m2:555:A:H61	1.58	0.51
83:m2:1412:C:H2'	83:m2:1413:G:C8	2.44	0.51
83:m2:1738:G:H2'	83:m2:1739:G:C8	2.46	0.51
2:A2:1006:G:H2'	2:A2:1007:A:O4'	2.09	0.51
2:A2:1361:G:O2'	2:A2:2567:A:H8	1.92	0.51
2:A2:1626:G:H4'	23:G2:44:TYR:HD2	1.75	0.51
2:A2:3576:U:H2'	2:A2:3577:U:C6	2.46	0.51
2:A2:3762:C:H2'	2:A2:3763:U:H6	1.74	0.51
3:A3:111:LEU:O	3:A3:115:LYS:HG2	2.10	0.51
4:B1:200:THR:HG22	4:B1:201:THR:HG23	1.93	0.51
20:F2:138:MET:SD	20:F2:144:ILE:HG12	2.50	0.51
23:G2:196:ARG:HH12	23:G2:200:MET:HG3	1.74	0.51
41:O2:63:ILE:HG12	41:O2:72:VAL:HG22	1.92	0.51
74:r2:129:ILE:HD11	83:m2:291:G:H5'	1.92	0.51
83:m2:51:U:H2'	83:m2:52:G:C8	2.45	0.51
83:m2:630:A:C6	83:m2:1502:G:N3	2.75	0.51
2:A2:165:A:H2'	2:A2:166:C:C6	2.45	0.51
2:A2:487:C:H2'	2:A2:488:G:C8	2.46	0.51
2:A2:1154:OMC:HM21	20:F2:106:LYS:HB2	1.91	0.51
2:A2:2651:G:H5''	35:L2:134:ASN:HD21	1.75	0.51
23:G2:223:PHE:HB3	23:G2:226:TYR:HB2	1.93	0.51
30:J2:94:MET:HE1	30:J2:146:ILE:HG22	1.92	0.51
73:q2:132:LYS:HG3	73:q2:191:PRO:HA	1.92	0.51
2:A2:163:A:H2'	2:A2:164:G:H8	1.75	0.51
4:B1:44:ASP:N	14:D2:64:ARG:HH22	2.09	0.51
4:B1:209:SER:HA	4:B1:212:LYS:HG3	1.93	0.51
10:C1:103:VAL:HG11	10:C1:144:LEU:HD21	1.93	0.51
31:J3:229:CYS:HB2	83:m2:14:C:H5'	1.93	0.51
33:K3:70:HIS:HA	33:K3:98:ARG:HH12	1.75	0.51
38:M3:37:GLU:HA	38:M3:40:LYS:HG2	1.93	0.51
56:W2:17:ARG:O	56:W2:21:VAL:HG23	2.09	0.51
74:r2:52:LEU:HD22	74:r2:54:TYR:CZ	2.45	0.51
83:m2:1200:G:H2'	83:m2:1201:A:C8	2.45	0.51
83:m2:1541:U:H2'	83:m2:1542:G:C8	2.44	0.51
2:A2:1360:A:H2	2:A2:2568:A:H8	1.59	0.51
2:A2:3388:A:H2'	2:A2:3389:A:C8	2.44	0.51
2:A2:3617:G:N2	2:A2:3621:A:H62	2.08	0.51
2:A2:4543:G:H2'	2:A2:4544:G:H8	1.76	0.51
46:Q3:104:ARG:HD3	46:Q3:108:LYS:HE2	1.93	0.51
61:b2:89:ARG:O	61:b2:93:ARG:HG2	2.11	0.51
71:o2:74:VAL:HG12	71:o2:121:LEU:HB3	1.92	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
72:p2:160:GLN:O	72:p2:164:ILE:HD12	2.10	0.51
73:q2:150:MET:HG2	73:q2:152:PHE:CE1	2.45	0.51
75:s2:96:ALA:O	75:s2:100:ILE:HG12	2.11	0.51
83:m2:28:U:H2'	83:m2:29:G:H8	1.75	0.51
2:A2:1255:C:H2'	2:A2:1256:A:C8	2.46	0.51
2:A2:3390:U:H2'	2:A2:3391:G:O4'	2.11	0.51
3:A3:40:TYR:HD1	3:A3:83:PHE:HE2	1.59	0.51
29:I3:192:THR:HG21	73:q2:220:THR:HG22	1.92	0.51
33:K3:84:TYR:CE1	33:K3:93:LYS:HB3	2.46	0.51
70:k2:90:LEU:HD22	70:k2:111:ILE:HG23	1.92	0.51
30:J2:54:LYS:HA	30:J2:83:TRP:CD1	2.46	0.51
33:K3:133:LEU:HD22	83:m2:65:C:C2	2.46	0.51
74:r2:148:ARG:HH21	74:r2:149:TYR:HE2	1.58	0.51
83:m2:1350:G:C2'	83:m2:1351:G:H8	2.18	0.51
83:m2:1535:A:H2'	83:m2:1535:A:N3	2.26	0.51
83:m2:1608:G:H22	83:m2:1634:G:H2'	1.76	0.51
83:m2:1804:C:H2'	83:m2:1805:U:C6	2.46	0.51
2:A2:1244:C:H2'	2:A2:1245:G:O4'	2.11	0.51
2:A2:3942:U:OP1	68:i2:9:ARG:HD3	2.11	0.51
3:A3:99:LEU:HD12	3:A3:102:GLY:H	1.74	0.51
16:E1:3:GLN:HG3	16:E1:6:GLY:H	1.75	0.51
21:F3:95:ARG:HG2	83:m2:1868:A:H5'	1.92	0.51
24:G3:46:VAL:HG13	24:G3:50:VAL:HG21	1.91	0.51
26:H2:171:PHE:HA	26:H2:182:VAL:HG12	1.92	0.51
29:I3:19:THR:HG23	29:I3:35:SER:HA	1.93	0.51
83:m2:511:OMG:H2'	83:m2:512:G:H8	1.74	0.51
83:m2:611:U:H2'	83:m2:612:G:H8	1.75	0.51
83:m2:1203:U:H2'	83:m2:1204:U:C6	2.46	0.51
2:A2:1006:G:H1	2:A2:1018:U:H3	1.58	0.51
2:A2:1333:C:H5'	20:F2:102:PHE:CE1	2.46	0.51
2:A2:1441:C:H42	14:D2:3:ARG:HH11	1.56	0.51
2:A2:1626:G:H5'	23:G2:44:TYR:HD2	1.76	0.51
2:A2:2485:U:H2'	2:A2:2486:C:C6	2.46	0.51
2:A2:3926:A:H2'	2:A2:3927:G:C8	2.45	0.51
2:A2:4637:U:H2'	2:A2:4638:G:H8	1.75	0.51
3:A3:13:LEU:HD23	3:A3:20:ILE:HD11	1.91	0.51
11:C2:132:G:H4'	47:R2:69:ASN:ND2	2.26	0.51
17:E2:291:TYR:HB3	17:E2:298:LEU:HD11	1.93	0.51
20:F2:26:ALA:HB3	20:F2:266:THR:HA	1.93	0.51
29:I3:191:HIS:CD2	29:I3:195:LEU:HD21	2.46	0.51
73:q2:194:PRO:HD2	73:q2:200:PRO:O	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
79:w2:42:LEU:H	83:m2:293:G:H1'	1.75	0.51
81:y2:43:GLU:HB3	81:y2:44:PRO:HD3	1.92	0.51
83:m2:602:G:H2'	83:m2:603:OMG:C8	2.42	0.51
83:m2:933:C:H2'	83:m2:934:G:C8	2.45	0.51
83:m2:1114:U:H2'	83:m2:1115:A:C8	2.46	0.51
2:A2:492:G:H1	2:A2:670:G:H1	1.59	0.51
2:A2:1406:A:H5''	2:A2:2594:U:H5''	1.93	0.51
2:A2:1532:U:H2'	2:A2:1533:C:C6	2.47	0.51
2:A2:2409:C:H2'	2:A2:2410:C:H6	1.76	0.51
2:A2:3374:A2M:H2	2:A2:3590:G:O4'	2.10	0.51
2:A2:3423:C:O2'	2:A2:3424:U:H5'	2.10	0.51
2:A2:3717:U:H2'	2:A2:3718:U:C6	2.46	0.51
2:A2:4573:C:H4'	22:G1:121:ARG:NH1	2.24	0.51
3:A3:43:VAL:HG11	3:A3:83:PHE:CE2	2.46	0.51
6:B3:62:ARG:NH2	83:m2:1544:C:H5''	2.26	0.51
20:F2:61:GLN:O	63:d2:52:LYS:HE3	2.11	0.51
25:H1:159:ARG:HB3	25:H1:164:LEU:HB2	1.91	0.51
31:J3:235:ASN:ND2	83:m2:1357:C:O2	2.24	0.51
35:L2:105:LEU:HD23	35:L2:138:LEU:HD23	1.92	0.51
37:M2:19:THR:HG22	37:M2:21:LYS:H	1.76	0.51
58:Y2:76:LYS:HD2	58:Y2:98:GLU:OE1	2.11	0.51
65:f2:9:ILE:HD12	65:f2:51:LEU:HD21	1.92	0.51
72:p2:66:VAL:HG22	72:p2:85:LYS:HE2	1.93	0.51
72:p2:83:LYS:NZ	72:p2:106:THR:HA	2.26	0.51
73:q2:175:VAL:HB	73:q2:182:LEU:HB2	1.92	0.51
75:s2:35:LEU:HD22	75:s2:117:ILE:HD13	1.92	0.51
83:m2:118:C:H1'	83:m2:447:A:C4	2.46	0.51
83:m2:457:A:H2'	83:m2:458:C:C6	2.46	0.51
83:m2:519:OMC:H2'	83:m2:520:G:O4'	2.11	0.51
1:A1:166:TYR:CE2	1:A1:259:GLU:HB2	2.46	0.50
2:A2:718:A:OP1	26:H2:199:LYS:HD2	2.11	0.50
2:A2:2648:U:H2'	2:A2:2649:A:C8	2.46	0.50
6:B3:67:ARG:HD3	83:m2:1589:G:N7	2.26	0.50
11:C2:48:A:H5''	61:b2:51:ARG:HH12	1.76	0.50
22:G1:85:LYS:O	22:G1:89:THR:HG23	2.11	0.50
71:o2:137:ALA:HB1	71:o2:142:LEU:HB3	1.93	0.50
79:w2:121:GLN:HB2	79:w2:147:LYS:HE3	1.93	0.50
83:m2:1412:C:H2'	83:m2:1413:G:H8	1.76	0.50
83:m2:1535:A:H2	83:m2:1538:G:N3	2.09	0.50
2:A2:1317:G:H2'	2:A2:1318:C:C6	2.45	0.50
2:A2:1446:G:N2	14:D2:3:ARG:HH21	2.08	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:3720:U:H2'	2:A2:3721:U:C6	2.46	0.50
29:I3:14:HIS:NE2	29:I3:35:SER:HB3	2.26	0.50
39:N2:34:TYR:HE2	39:N2:93:ILE:HG23	1.76	0.50
83:m2:177:G:H1'	83:m2:315:A:H62	1.76	0.50
83:m2:947:U:H2'	83:m2:948:U:C6	2.47	0.50
1:A1:156:ARG:NH1	5:B2:97:G:H4'	2.27	0.50
2:A2:2220:C:H2'	2:A2:2221:G:O4'	2.12	0.50
2:A2:2275:C:H2'	2:A2:2276:G:C8	2.46	0.50
2:A2:3404:A:H2'	2:A2:3405:C:C6	2.47	0.50
2:A2:4564:G:O2'	2:A2:4565:C:H5'	2.11	0.50
3:A3:37:GLY:N	83:m2:1632:A:H5''	2.26	0.50
7:Bz:66:U:H2'	7:Bz:67:C:C6	2.46	0.50
34:L1:180:VAL:HA	34:L1:183:ILE:HG12	1.94	0.50
74:r2:14:ALA:HB1	74:r2:18:TRP:CE3	2.46	0.50
75:s2:72:LEU:HD23	75:s2:151:ILE:HG23	1.92	0.50
83:m2:839:A:H4'	83:m2:840:G:OP1	2.11	0.50
83:m2:1440:A:H2'	83:m2:1441:A:C8	2.45	0.50
1:A1:148:ASN:ND2	39:N2:132:PRO:HG2	2.26	0.50
2:A2:260:C:H2'	2:A2:261:G:C8	2.45	0.50
2:A2:280:G:H5''	25:H1:14:LYS:HE2	1.94	0.50
2:A2:702:G:H3'	2:A2:703:U:H4'	1.91	0.50
2:A2:3564:A:N1	9:By:18:UNK:HA	2.26	0.50
23:G2:44:TYR:OH	39:N2:67:VAL:HG21	2.11	0.50
25:H1:80:THR:OG1	25:H1:87:HIS:HA	2.11	0.50
33:K3:84:TYR:HE1	33:K3:93:LYS:HB3	1.75	0.50
46:Q3:110:ARG:HG2	46:Q3:114:MET:HE2	1.92	0.50
74:r2:87:MET:HE1	74:r2:236:ILE:HD13	1.94	0.50
77:u2:5:ARG:HD2	83:m2:381:C:O2	2.12	0.50
83:m2:106:C:H2'	83:m2:107:A:H8	1.76	0.50
83:m2:844:C:H2'	83:m2:845:C:C6	2.46	0.50
83:m2:1421:C:H4'	83:m2:1422:G:H5'	1.94	0.50
83:m2:1745:G:H21	83:m2:1793:A:H62	1.56	0.50
2:A2:1173:G:H4'	25:H1:203:TYR:HB2	1.94	0.50
2:A2:1438:OMG:HM23	25:H1:81:TYR:HE2	1.76	0.50
2:A2:2019:G:H2'	2:A2:2020:C:O4'	2.12	0.50
2:A2:2559:OMC:HM21	60:a2:32:TYR:HE2	1.77	0.50
2:A2:4044:OMG:N3	2:A2:4099:5MC:HM52	2.27	0.50
41:O2:61:VAL:HG23	41:O2:74:SER:HB3	1.92	0.50
83:m2:949:G:H2'	83:m2:950:C:C6	2.45	0.50
83:m2:1560:C:H2'	83:m2:1561:C:C6	2.47	0.50
83:m2:1564:C:H2'	83:m2:1565:G:H8	1.76	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
83:m2:1705:OMC:H2'	83:m2:1706:C:O4'	2.11	0.50
2:A2:4396:A:H2'	2:A2:4397:G:O4'	2.10	0.50
2:A2:4499:G:H2'	2:A2:4500:G:C8	2.47	0.50
10:C1:44:GLU:HG2	10:C1:58:ASP:HB2	1.93	0.50
41:O2:79:SER:HB3	41:O2:82:TYR:CE1	2.47	0.50
74:r2:100:ARG:HG2	74:r2:102:ILE:HG23	1.93	0.50
80:x2:75:VAL:HG12	80:x2:93:MET:HE2	1.94	0.50
83:m2:1012:G:H2'	83:m2:1013:A:H8	1.76	0.50
2:A2:129:C:H2'	2:A2:130:G:H8	1.77	0.50
2:A2:4343:A:H5'	10:C1:71:ARG:HE	1.76	0.50
2:A2:4413:G:H2'	2:A2:4414:A:C8	2.47	0.50
23:G2:64:ILE:HG13	23:G2:105:LEU:HD21	1.94	0.50
23:G2:181:PRO:HD2	23:G2:195:HIS:HD1	1.76	0.50
28:I2:12:ARG:HD3	28:I2:37:ARG:NH2	2.26	0.50
33:K3:157:VAL:HG21	33:K3:176:ILE:HD11	1.94	0.50
63:d2:25:LYS:HD3	65:f2:50:GLY:O	2.11	0.50
67:h2:12:ARG:HG3	67:h2:15:ARG:HH21	1.77	0.50
83:m2:17:C:H2'	83:m2:18:C:H6	1.76	0.50
2:A2:830:G:N2	2:A2:836:C:H2'	2.26	0.50
2:A2:928:G:H1	2:A2:1064:G:N2	2.07	0.50
2:A2:2250:U:H2'	2:A2:2251:G:H8	1.77	0.50
2:A2:3618:A:H3'	2:A2:3621:A:C2	2.47	0.50
2:A2:3720:U:H2'	2:A2:3721:U:H6	1.77	0.50
31:J3:66:LEU:HD21	31:J3:81:ILE:HD13	1.94	0.50
51:T2:102:ARG:HB2	51:T2:102:ARG:CZ	2.41	0.50
60:a2:63:VAL:HG12	60:a2:66:ARG:NH1	2.26	0.50
74:r2:124:CYS:HA	74:r2:142:HIS:CE1	2.47	0.50
83:m2:377:U:H2'	83:m2:378:A:C8	2.47	0.50
83:m2:1223:G:H2'	83:m2:1224:G:C8	2.45	0.50
2:A2:319:A:H1'	2:A2:3382:A:N3	2.27	0.50
2:A2:1040:C:H2'	2:A2:1041:G:H8	1.77	0.50
2:A2:1831:A:H2'	2:A2:1832:A:C8	2.46	0.50
2:A2:2102:A:C4	58:Y2:31:ILE:HD11	2.47	0.50
2:A2:3328:G:H4'	25:H1:67:ARG:HH12	1.75	0.50
70:k2:119:ARG:HA	70:k2:122:LYS:HE3	1.93	0.50
83:m2:335:G:H2'	83:m2:336:C:O4'	2.11	0.50
83:m2:865:U:H2'	83:m2:866:A:C8	2.45	0.50
1:A1:89:THR:O	1:A1:93:MET:HG2	2.12	0.49
2:A2:1836:G:H2'	2:A2:1837:C:C6	2.48	0.49
11:C2:67:U:H2'	11:C2:68:G:H8	1.77	0.49
17:E2:80:GLU:HG2	17:E2:82:PRO:HD3	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:U3:122:PRO:HA	54:U3:127:GLY:HA2	1.94	0.49
83:m2:118:C:H3'	83:m2:119:U:H6	1.77	0.49
83:m2:1459:U:H2'	83:m2:1460:G:C8	2.45	0.49
2:A2:217:C:H3'	2:A2:218:A:H4'	1.94	0.49
2:A2:854:A:H1'	2:A2:1878:G:H5''	1.94	0.49
2:A2:1574:C:H2'	2:A2:1575:U:C6	2.46	0.49
2:A2:1896:A:H2'	2:A2:1897:A:C4	2.47	0.49
2:A2:3472:A:OP1	2:A2:3474:OMU:H5	2.12	0.49
2:A2:4325:U:H2'	2:A2:4326:C:H6	1.76	0.49
2:A2:4431:C:H2'	2:A2:4432:C:C6	2.47	0.49
13:D1:47:PRO:HB3	13:D1:171:TRP:CZ2	2.47	0.49
15:D3:20:SER:HB3	15:D3:59:ILE:HD11	1.93	0.49
36:L3:134:HIS:CE1	36:L3:164:PRO:HD2	2.47	0.49
51:T2:12:LEU:HB2	51:T2:81:MET:HB3	1.92	0.49
51:T2:42:LEU:HA	51:T2:74:VAL:HG22	1.94	0.49
53:U2:100:ILE:HG12	53:U2:123:ILE:HB	1.94	0.49
59:Z2:11:PHE:HB2	59:Z2:101:ILE:HD12	1.94	0.49
83:m2:926:G:H1	83:m2:1020:U:H3	1.60	0.49
83:m2:955:C:H2'	83:m2:956:U:C6	2.47	0.49
83:m2:1038:A:H4'	83:m2:1857:G:N2	2.27	0.49
83:m2:1690:C:H2'	83:m2:1691:C:C6	2.47	0.49
83:m2:1846:U:H2'	83:m2:1847:A:H8	1.75	0.49
2:A2:2030:G:H2'	2:A2:2031:A:C8	2.47	0.49
2:A2:3762:C:H2'	2:A2:3763:U:C6	2.48	0.49
2:A2:4340:C:H2'	2:A2:4341:U:C6	2.48	0.49
3:A3:27:ALA:HB2	3:A3:52:LEU:HD23	1.94	0.49
7:Bv:27:G:H2'	7:Bv:28:G:H8	1.77	0.49
11:C2:91:A:H5'	49:S2:22:PRO:HB3	1.94	0.49
25:H1:191:ALA:O	25:H1:195:ARG:HG2	2.11	0.49
26:H2:106:ASP:OD2	26:H2:107:LYS:HG3	2.12	0.49
49:S2:31:SER:HA	49:S2:48:PRO:HA	1.93	0.49
7:n2:28:G:H2'	7:n2:29:G:C8	2.47	0.49
73:q2:177:LEU:HD12	73:q2:182:LEU:HD23	1.95	0.49
83:m2:16:G:H2'	83:m2:17:C:C6	2.47	0.49
83:m2:1351:G:C5'	83:m2:1351:G:C8	2.88	0.49
2:A2:318:A:H2'	2:A2:319:A:C8	2.47	0.49
2:A2:2649:A:H2'	2:A2:2650:A:C8	2.47	0.49
2:A2:3340:G:H2'	2:A2:3341:C:C6	2.47	0.49
2:A2:3880:OMG:H5''	2:A2:3881:U:O4'	2.12	0.49
2:A2:4637:U:H2'	2:A2:4638:G:C8	2.48	0.49
4:B1:162:ASP:HB2	4:B1:163:PRO:HD3	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:B3:21:PHE:HA	6:B3:24:LYS:NZ	2.28	0.49
17:E2:393:LYS:HE2	17:E2:397:ILE:HD11	1.93	0.49
21:F3:4:LYS:HE2	21:F3:92:ARG:NH1	2.28	0.49
35:L2:135:LYS:O	35:L2:139:MET:HG3	2.12	0.49
47:R2:100:VAL:HG21	47:R2:109:ILE:HD11	1.94	0.49
82:z2:7:LYS:HB2	83:m2:1375:C:OP1	2.12	0.49
2:A2:440:U:H2'	2:A2:441:G:H8	1.77	0.49
2:A2:811:G:H8	2:A2:811:G:OP2	1.95	0.49
2:A2:1482:A:H4'	2:A2:1498:G:N2	2.27	0.49
2:A2:3592:A:H2'	2:A2:3593:C:C6	2.47	0.49
2:A2:4550:C:H2'	2:A2:4551:C:C6	2.47	0.49
3:A3:132:ARG:HB2	3:A3:134:GLN:HE22	1.76	0.49
7:Bz:16:U:O3'	7:Bz:17:C:H2'	2.12	0.49
17:E2:19:ARG:HD3	17:E2:275:HIS:NE2	2.27	0.49
19:F1:81:LEU:HD11	19:F1:98:VAL:HG22	1.94	0.49
20:F2:36:ILE:HG21	20:F2:122:TYR:HD2	1.77	0.49
26:H2:215:ILE:HG23	26:H2:219:LEU:HD12	1.94	0.49
27:H3:34:TYR:HE1	83:m2:1552:G:P	2.35	0.49
31:J3:187:ARG:CZ	31:J3:192:LEU:HD12	2.43	0.49
33:K3:50:VAL:HG12	33:K3:113:ILE:HG12	1.95	0.49
34:L1:169:VAL:HG21	34:L1:183:ILE:HG22	1.94	0.49
48:R3:92:LEU:HD23	48:R3:97:ILE:HG13	1.94	0.49
51:T2:79:HIS:CD2	60:a2:88:ARG:HH22	2.29	0.49
70:k2:33:LYS:HD2	70:k2:40:TYR:CZ	2.47	0.49
7:n2:41:C:H5'	75:s2:198:ARG:NH1	2.27	0.49
73:q2:29:LEU:HB3	73:q2:32:ASP:HB2	1.92	0.49
76:t2:53:VAL:HG23	76:t2:175:GLY:HA3	1.93	0.49
83:m2:1713:U:H2'	83:m2:1714:A:H8	1.76	0.49
2:A2:1084:G:H1'	2:A2:1909:G:H21	1.77	0.49
2:A2:1532:U:H2'	2:A2:1533:C:H6	1.78	0.49
2:A2:2648:U:H2'	2:A2:2649:A:H8	1.77	0.49
2:A2:3266:A:H2'	2:A2:3267:A:H8	1.78	0.49
2:A2:4270:A2M:H5''	43:P2:15:ARG:HB2	1.94	0.49
2:A2:4382:C:H2'	2:A2:4383:G:O4'	2.13	0.49
17:E2:37:ALA:HA	17:E2:188:GLY:H	1.78	0.49
20:F2:146:GLU:HG3	20:F2:175:LYS:HB3	1.94	0.49
34:L1:71:GLN:HG3	34:L1:72:GLN:OE1	2.13	0.49
43:P2:62:MET:HE3	43:P2:76:VAL:HG12	1.93	0.49
71:o2:102:ARG:NE	83:m2:1379:U:C5	2.80	0.49
73:q2:38:GLU:HB3	73:q2:49:ILE:HG13	1.93	0.49
83:m2:29:G:H2'	83:m2:30:C:C6	2.48	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
83:m2:29:G:H2'	83:m2:30:C:H6	1.77	0.49
83:m2:96:C:H2'	83:m2:97:U:H6	1.76	0.49
2:A2:1277:C:H5''	55:V2:32:LEU:HD12	1.94	0.49
2:A2:1824:C:H2'	2:A2:1825:C:H6	1.77	0.49
2:A2:2599:A:O2'	2:A2:4283:G:H4'	2.13	0.49
2:A2:3263:U:H2'	2:A2:3264:A:C8	2.46	0.49
2:A2:3504:U:H2'	2:A2:3505:A:C8	2.47	0.49
7:Bz:28:G:H2'	7:Bz:29:G:H8	1.77	0.49
11:C2:70:G:H5''	49:S2:27:ARG:CZ	2.42	0.49
36:L3:93:LYS:HB3	36:L3:96:TYR:HD2	1.77	0.49
53:U2:100:ILE:HG21	53:U2:125:LYS:HE3	1.94	0.49
70:k2:119:ARG:O	70:k2:122:LYS:HG2	2.13	0.49
7:n2:59:U:H2'	7:n2:60:U:O4'	2.13	0.49
83:m2:945:U:C2	83:m2:946:A:C8	3.00	0.49
83:m2:1375:C:H2'	83:m2:1376:C:C6	2.48	0.49
2:A2:952:C:H2'	2:A2:953:G:H8	1.77	0.49
2:A2:1669:A:H2'	2:A2:1670:A:C8	2.48	0.49
2:A2:2203:G:H2'	2:A2:2204:A:C8	2.47	0.49
2:A2:2658:G:H5'	2:A2:3248:G:H21	1.77	0.49
19:F1:18:TRP:CD1	19:F1:18:TRP:H	2.30	0.49
20:F2:218:ILE:HA	20:F2:229:LEU:HG	1.95	0.49
21:F3:44:ILE:HD11	42:O3:97:LEU:HD23	1.93	0.49
22:G1:97:ALA:HA	22:G1:100:ARG:HG2	1.94	0.49
42:O3:26:ASN:O	42:O3:26:ASN:ND2	2.41	0.49
43:P2:25:VAL:HG22	43:P2:38:TYR:HD1	1.78	0.49
73:q2:27:ARG:HB3	78:v2:61:GLN:HE21	1.78	0.49
77:u2:38:ILE:HG12	77:u2:96:LEU:HD23	1.95	0.49
80:x2:107:ILE:HA	80:x2:111:MET:HE3	1.95	0.49
2:A2:259:C:H2'	2:A2:260:C:C6	2.48	0.49
2:A2:263:G:H2'	2:A2:264:C:C6	2.47	0.49
2:A2:2651:G:H5''	35:L2:134:ASN:CG	2.38	0.49
2:A2:3512:A:H5''	30:J2:83:TRP:O	2.13	0.49
16:E1:86:GLY:HA3	16:E1:130:PHE:HE2	1.77	0.49
61:b2:21:LEU:HG	61:b2:25:LYS:HE2	1.95	0.49
83:m2:1131:G:H2'	83:m2:1132:G:N3	2.28	0.49
2:A2:3253:G:H4'	2:A2:3254:C:OP1	2.13	0.49
14:D2:117:GLU:HG2	14:D2:124:GLY:N	2.28	0.49
83:m2:1229:G:C2	83:m2:1230:A:C8	3.01	0.49
83:m2:1327:G:H1'	83:m2:1512:G:H5''	1.95	0.49
83:m2:1595:C:H2'	83:m2:1596:A:H8	1.78	0.49
2:A2:129:C:H2'	2:A2:130:G:C8	2.47	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:1686:C:OP1	58:Y2:50:LYS:HG3	2.13	0.48
2:A2:2504:C:H2'	2:A2:2505:G:C8	2.48	0.48
2:A2:3588:U:H2'	2:A2:3589:G:C8	2.48	0.48
2:A2:3898:G:H2'	2:A2:3899:G:H8	1.78	0.48
16:E1:57:VAL:HG12	16:E1:60:PHE:H	1.78	0.48
29:I3:55:PRO:HB2	81:y2:102:GLU:OE2	2.13	0.48
33:K3:195:LYS:HE3	83:m2:126:G:C4	2.48	0.48
39:N2:64:VAL:HG13	39:N2:72:VAL:HG13	1.95	0.48
7:n2:15:G:H22	7:n2:21:A:H1'	1.78	0.48
72:p2:34:LYS:HB2	72:p2:97:LEU:HD13	1.95	0.48
72:p2:127:VAL:HG21	72:p2:176:VAL:HB	1.95	0.48
83:m2:1176:U:C2	83:m2:1177:G:C8	3.01	0.48
83:m2:1215:C:H2'	83:m2:1216:A:C8	2.48	0.48
83:m2:1607:G:H1'	83:m2:1636:A:N6	2.28	0.48
2:A2:152:U:P	25:H1:49:ARG:HH12	2.36	0.48
2:A2:406:C:H2'	2:A2:407:A:C8	2.48	0.48
2:A2:453:G:H22	2:A2:1107:G:H22	1.62	0.48
2:A2:2409:C:H2'	2:A2:2410:C:C6	2.48	0.48
2:A2:3425:C:H2'	2:A2:3426:U:C6	2.48	0.48
2:A2:4297:C:H3'	35:L2:62:ARG:HH22	1.78	0.48
4:B1:190:LEU:HB3	4:B1:199:CYS:HB3	1.94	0.48
12:C3:31:SER:HB3	12:C3:107:GLU:OE1	2.13	0.48
18:E3:36:LEU:HA	18:E3:39:ASN:ND2	2.28	0.48
29:I3:79:LEU:HD13	29:I3:89:LEU:HD12	1.95	0.48
69:j2:26:VAL:HG22	69:j2:30:GLU:HG3	1.93	0.48
80:x2:137:HIS:O	80:x2:140:ARG:HG2	2.14	0.48
2:A2:18:C:H4'	25:H1:138:PHE:CD1	2.48	0.48
2:A2:1250:C:H2'	2:A2:1251:U:C6	2.48	0.48
2:A2:2484:C:H2'	2:A2:2485:U:O4'	2.13	0.48
2:A2:3617:G:H21	2:A2:3621:A:H62	1.61	0.48
2:A2:3716:U:H2'	2:A2:3717:U:C6	2.48	0.48
2:A2:3721:U:H2'	2:A2:3722:C:H6	1.77	0.48
2:A2:4272:OMU:H5'	43:P2:51:ARG:NH1	2.28	0.48
5:B2:111:C:H2'	5:B2:112:U:O4'	2.14	0.48
7:Bz:17:C:O2'	7:Bz:18:G:H5'	2.13	0.48
15:D3:61:ARG:HH21	83:m2:1140:C:N4	2.11	0.48
17:E2:43:LEU:HG	17:E2:196:TRP:HH2	1.77	0.48
30:J2:85:LYS:HE2	30:J2:89:GLU:OE2	2.13	0.48
78:v2:13:GLU:O	78:v2:17:LYS:HG3	2.14	0.48
80:x2:115:TYR:HB2	80:x2:118:GLU:HG3	1.95	0.48
83:m2:178:C:H2'	83:m2:179:C:H6	1.78	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
83:m2:1205:G:H2'	83:m2:1206:A:H8	1.74	0.48
2:A2:1092:C:H2'	2:A2:1093:A:O4'	2.12	0.48
2:A2:1527:U:H2'	2:A2:1528:U:C6	2.49	0.48
2:A2:1680:G:H2'	2:A2:1681:C:C6	2.49	0.48
2:A2:4271:U:H2'	2:A2:4272:OMU:H6	1.94	0.48
31:J3:118:ALA:HB1	83:m2:1489:A:O4'	2.14	0.48
54:U3:135:HIS:CE1	54:U3:140:TYR:HB3	2.47	0.48
7:n2:35:A:H2'	7:n2:36:A:H8	1.78	0.48
81:y2:97:GLN:HB2	81:y2:105:LYS:HZ2	1.78	0.48
83:m2:1428:U:H2'	83:m2:1429:C:C6	2.48	0.48
83:m2:1530:G:H2'	83:m2:1531:C:C6	2.49	0.48
83:m2:1810:U:H2'	83:m2:1811:A:C8	2.48	0.48
2:A2:1223:C:H2'	2:A2:1224:G:O4'	2.14	0.48
2:A2:2299:G:H22	11:C2:125:C:H5'	1.79	0.48
2:A2:3346:U:H2'	2:A2:3347:G:O4'	2.14	0.48
3:A3:19:ASN:OD1	16:E1:119:TYR:CG	2.66	0.48
10:C1:93:ARG:HG2	10:C1:182:SER:HB3	1.96	0.48
10:C1:134:CYS:SG	10:C1:144:LEU:HD22	2.53	0.48
16:E1:18:ARG:HB2	16:E1:133:VAL:O	2.13	0.48
41:O2:101:ARG:HD2	41:O2:115:PHE:CE1	2.49	0.48
45:Q2:31:PHE:CD2	45:Q2:37:GLU:HG2	2.48	0.48
61:b2:91:MET:HE1	63:d2:78:PHE:CD2	2.49	0.48
71:o2:140:VAL:HG23	71:o2:142:LEU:HB2	1.96	0.48
76:t2:100:ILE:H	76:t2:100:ILE:HD12	1.79	0.48
77:u2:194:GLU:HG3	79:w2:10:TYR:CD2	2.48	0.48
78:v2:10:ALA:HB1	78:v2:38:LYS:HD3	1.94	0.48
79:w2:81:LYS:HB3	83:m2:396:G:H5'	1.96	0.48
80:x2:40:ARG:HG2	83:m2:1623:U:OP1	2.14	0.48
83:m2:1203:U:O2'	83:m2:1360:U:O2'	2.13	0.48
83:m2:1597:U:H2'	83:m2:1598:U:C6	2.48	0.48
1:A1:98:ARG:CD	2:A2:736:G:H5''	2.43	0.48
2:A2:1893:C:C4	2:A2:1896:A:H4'	2.49	0.48
2:A2:3709:C:H2'	2:A2:3710:U:C6	2.48	0.48
7:Bv:34:G:O4'	83:m2:1250:B8N:C32	2.62	0.48
17:E2:60:VAL:HG21	17:E2:67:VAL:HG23	1.95	0.48
24:G3:62:GLU:HG2	42:O3:121:ARG:HE	1.69	0.48
30:J2:131:ARG:HG3	30:J2:137:ASN:ND2	2.26	0.48
56:W2:30:GLY:O	56:W2:34:THR:HG23	2.12	0.48
74:r2:124:CYS:HA	74:r2:142:HIS:HE1	1.79	0.48
74:r2:203:GLY:HA2	83:m2:293:G:C2	2.48	0.48
83:m2:676:C:H2'	83:m2:677:U:C6	2.48	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:281:U:O2'	2:A2:329:A:H1'	2.14	0.48
2:A2:1023:C:H2'	2:A2:1024:C:C6	2.49	0.48
2:A2:1899:G:H5'	2:A2:1900:G:H4'	1.95	0.48
2:A2:2255:U:H2'	2:A2:2256:C:C6	2.48	0.48
2:A2:2389:C:H2'	2:A2:2390:U:C6	2.48	0.48
20:F2:224:ILE:HB	20:F2:227:ILE:HD12	1.96	0.48
23:G2:278:ASP:O	23:G2:282:GLN:HG3	2.14	0.48
33:K3:141:ILE:HD11	33:K3:157:VAL:HA	1.95	0.48
44:P3:57:ARG:HH21	83:m2:921:A:H5'	1.78	0.48
55:V2:65:VAL:HA	55:V2:68:ARG:HG2	1.95	0.48
70:k2:108:MET:O	70:k2:112:ARG:HG3	2.14	0.48
83:m2:71:G:H3'	83:m2:72:C:H5''	1.95	0.48
83:m2:439:G:H2'	83:m2:440:G:O4'	2.14	0.48
83:m2:486:A2M:H8	83:m2:486:A2M:O5'	2.14	0.48
2:A2:1910:G:H2'	2:A2:1911:G:O4'	2.12	0.48
2:A2:2602:G:OP1	43:P2:85:ARG:HD3	2.13	0.48
2:A2:3267:A:H2	2:A2:4662:A:H8	1.61	0.48
2:A2:3275:G:H4'	35:L2:79:GLY:O	2.13	0.48
2:A2:3419:A:C2	83:m2:1711:G:H4'	2.48	0.48
2:A2:3504:U:H2'	2:A2:3505:A:H8	1.79	0.48
2:A2:4649:U:H2'	2:A2:4650:C:C6	2.49	0.48
11:C2:144:U:H2'	11:C2:145:C:C6	2.49	0.48
31:J3:91:SER:HB3	31:J3:156:ILE:HG23	1.95	0.48
31:J3:117:ARG:NH1	83:m2:1690:C:OP1	2.46	0.48
74:r2:16:LYS:HD3	83:m2:814:A:H5''	1.96	0.48
83:m2:159:A:H2	83:m2:469:G:H21	1.62	0.48
1:A1:125:PRO:HD3	2:A2:1678:U:O3'	2.14	0.48
2:A2:3573:A:H2'	2:A2:3574:G:H8	1.79	0.48
2:A2:4104:U:H2'	2:A2:4174:G:O6	2.14	0.48
11:C2:74:U:C4	49:S2:74:TYR:HD1	2.32	0.48
12:C3:76:THR:HG21	83:m2:1340:G:H5''	1.95	0.48
29:I3:36:ARG:HD2	29:I3:65:PHE:CD2	2.49	0.48
30:J2:15:CYS:SG	30:J2:150:LEU:HB2	2.54	0.48
60:a2:88:ARG:HG2	60:a2:92:LYS:HE2	1.95	0.48
64:e2:12:LEU:HB3	64:e2:16:ARG:NH1	2.29	0.48
83:m2:14:C:H2'	83:m2:15:U:H6	1.78	0.48
83:m2:106:C:H5''	83:m2:433:G:O2'	2.13	0.48
83:m2:1225:A:H2'	83:m2:1226:G:O4'	2.13	0.48
2:A2:1668:G:H1	2:A2:4092:G:N2	2.11	0.48
2:A2:2464:C:H2'	2:A2:2465:C:C6	2.49	0.48
2:A2:3891:A:H2'	2:A2:3892:G:C8	2.49	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:4263:A:H2'	2:A2:4264:C:C6	2.49	0.48
17:E2:68:ASN:O	17:E2:70:LYS:HG3	2.14	0.48
23:G2:155:THR:HA	23:G2:179:ARG:HA	1.96	0.48
27:H3:46:TYR:O	27:H3:50:ILE:HD12	2.14	0.48
31:J3:210:PRO:HD3	31:J3:236:PHE:CE2	2.49	0.48
47:R2:126:THR:CG2	47:R2:134:LYS:HE2	2.43	0.48
77:u2:130:THR:OG1	77:u2:131:PRO:HD2	2.12	0.48
83:m2:1246:U:H2'	83:m2:1247:G:H8	1.78	0.48
2:A2:856:G:C2	26:H2:131:LYS:HD2	2.49	0.47
2:A2:1596:A:H5''	2:A2:3866:A:H61	1.79	0.47
2:A2:3618:A:C2	2:A2:3689:G:H5''	2.49	0.47
2:A2:3811:C:H2'	2:A2:3812:C:C6	2.49	0.47
2:A2:4581:G:C6	26:H2:190:ARG:HD3	2.49	0.47
7:Bz:36:A:H1'	83:m2:628:G:H21	1.78	0.47
7:Bz:43:C:H2'	7:Bz:44:G:H8	1.78	0.47
29:I3:18:VAL:HA	29:I3:35:SER:HB2	1.95	0.47
46:Q3:55:ILE:HG12	46:Q3:75:ILE:HD12	1.96	0.47
73:q2:175:VAL:HG12	73:q2:177:LEU:HG	1.94	0.47
83:m2:340:G:H2'	83:m2:341:A:C8	2.49	0.47
83:m2:680:U:H2'	83:m2:681:A:H8	1.79	0.47
83:m2:949:G:H2'	83:m2:950:C:H6	1.78	0.47
2:A2:2494:C:H5''	69:j2:69:TRP:CH2	2.49	0.47
2:A2:3926:A:H2'	2:A2:3927:G:H8	1.79	0.47
2:A2:4388:C:H2'	2:A2:4389:G:H8	1.79	0.47
3:A3:22:GLY:HA2	3:A3:56:ALA:HB3	1.95	0.47
11:C2:70:G:H5''	49:S2:27:ARG:NH2	2.28	0.47
11:C2:89:U:H2'	11:C2:90:C:C6	2.49	0.47
15:D3:16:LYS:HG2	15:D3:23:ILE:HD13	1.96	0.47
24:G3:34:PHE:CZ	24:G3:61:SER:HB3	2.48	0.47
29:I3:5:MET:HG2	29:I3:270:LEU:HD11	1.96	0.47
33:K3:195:LYS:C	33:K3:195:LYS:HD3	2.39	0.47
37:M2:165:PRO:C	37:M2:167:PHE:H	2.22	0.47
40:N3:15:ALA:HB2	50:S3:21:LYS:HD3	1.96	0.47
47:R2:78:LYS:HB2	47:R2:99:ILE:O	2.13	0.47
7:n2:21:A:N6	7:n2:46:G:H2'	2.29	0.47
72:p2:81:PHE:CE1	72:p2:82:ARG:HG3	2.50	0.47
75:s2:83:ASN:HD21	83:m2:1652:A:H2	1.62	0.47
83:m2:153:G:N2	83:m2:165:G:H22	2.12	0.47
83:m2:498:C:H2'	83:m2:499:C:H6	1.79	0.47
83:m2:1037:A:C4	83:m2:1038:A:C8	3.02	0.47
2:A2:2324:G:H2'	2:A2:2325:U:C6	2.49	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:Bz:63:G:H2'	7:Bz:64:A:H8	1.78	0.47
75:s2:77:MET:HB3	75:s2:83:ASN:HA	1.97	0.47
78:v2:83:LEU:HB2	78:v2:85:LEU:HG	1.95	0.47
83:m2:443:C:H2'	83:m2:444:C:C6	2.49	0.47
83:m2:1232:C:H2'	83:m2:1233:C:H6	1.79	0.47
2:A2:1155:U:H2'	2:A2:1156:A:C8	2.49	0.47
2:A2:1665:U:OP1	13:D1:15:LYS:HG3	2.14	0.47
10:C1:12:ILE:HG12	10:C1:18:ILE:HD12	1.96	0.47
15:D3:62:MET:HE3	15:D3:64:GLU:HG3	1.96	0.47
28:I2:16:LEU:HD21	28:I2:83:THR:HG21	1.95	0.47
29:I3:199:THR:HG22	29:I3:241:PHE:HD2	1.79	0.47
33:K3:161:PRO:HA	33:K3:171:THR:HG22	1.96	0.47
43:P2:106:VAL:HG12	43:P2:112:MET:HA	1.97	0.47
70:k2:2:SER:O	70:k2:6:GLN:HG3	2.14	0.47
80:x2:39:ALA:HB2	83:m2:1615:G:H3'	1.96	0.47
81:y2:23:ALA:HB2	81:y2:87:SER:OG	2.14	0.47
83:m2:534:C:N4	83:m2:555:A:H61	2.12	0.47
2:A2:1292:G:H2'	2:A2:1293:C:C6	2.49	0.47
2:A2:2297:G:H2'	2:A2:2298:A:C8	2.49	0.47
2:A2:3871:A:H2'	2:A2:3872:A:C8	2.48	0.47
11:C2:77:A:H2'	11:C2:78:G:C8	2.50	0.47
29:I3:77:PHE:CE2	29:I3:89:LEU:HD11	2.50	0.47
36:L3:74:GLY:O	36:L3:78:LEU:HD13	2.15	0.47
51:T2:83:THR:HG22	60:a2:95:PHE:CZ	2.50	0.47
55:V2:99:HIS:CE1	55:V2:101:LYS:HB3	2.49	0.47
56:W2:47:ILE:HD12	56:W2:94:LEU:HD11	1.95	0.47
81:y2:19:ALA:HB2	81:y2:75:GLY:HA3	1.96	0.47
83:m2:1137:C:H2'	83:m2:1138:U:C6	2.49	0.47
83:m2:1608:G:N2	83:m2:1634:G:H2'	2.30	0.47
1:A1:245:LYS:HD3	2:A2:1709:A:O2'	2.14	0.47
2:A2:1347:A2M:HM'3	2:A2:1450:A:N3	2.29	0.47
2:A2:3632:C:H3'	2:A2:3633:C:C6	2.50	0.47
2:A2:3826:U:H2'	2:A2:3827:G:H8	1.79	0.47
2:A2:3847:G:C8	2:A2:4094:PSU:H5'	2.49	0.47
2:A2:4024:U:OP2	68:i2:61:LYS:HG2	2.15	0.47
2:A2:4601:A:H2'	2:A2:4602:A:C8	2.50	0.47
2:A2:4675:C:H2'	2:A2:4676:U:H6	1.80	0.47
23:G2:156:GLY:HA2	23:G2:181:PRO:HG3	1.97	0.47
29:I3:291:TRP:CD1	29:I3:298:LEU:HD13	2.50	0.47
32:K2:12:LYS:HB2	32:K2:14:ARG:HE	1.80	0.47
33:K3:98:ARG:HD3	33:K3:99:GLY:O	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:N2:143:THR:O	39:N2:146:LYS:HG2	2.15	0.47
60:a2:114:GLN:HE22	60:a2:115:LYS:HE3	1.79	0.47
70:k2:49:VAL:HG11	70:k2:97:ILE:HD11	1.96	0.47
74:r2:29:PRO:HG2	74:r2:46:ILE:HD11	1.97	0.47
74:r2:68:ARG:HH21	74:r2:76:VAL:HG11	1.78	0.47
76:t2:62:ILE:HD11	76:t2:94:PHE:CE1	2.49	0.47
83:m2:177:G:H1'	83:m2:315:A:N6	2.30	0.47
83:m2:1131:G:H3'	83:m2:1132:G:N2	2.27	0.47
83:m2:1215:C:H2'	83:m2:1216:A:H8	1.79	0.47
2:A2:1155:U:H2'	2:A2:1156:A:H8	1.79	0.47
2:A2:1577:A:H2'	2:A2:1578:A:H8	1.79	0.47
2:A2:2353:A:N3	60:a2:61:PRO:HB3	2.30	0.47
2:A2:2360:G:O2'	2:A2:2361:G:H5'	2.15	0.47
2:A2:3511:C:H2'	2:A2:3512:A:H8	1.80	0.47
2:A2:3529:G:H2'	2:A2:3530:G:C8	2.49	0.47
2:A2:3607:G:H2'	2:A2:3608:A:H8	1.79	0.47
2:A2:4571:C:H2'	2:A2:4572:G:C8	2.49	0.47
6:B3:31:PRO:HG2	6:B3:33:TRP:O	2.14	0.47
6:B3:39:LEU:HD21	6:B3:56:ARG:NH2	2.29	0.47
11:C2:154:G:H2'	11:C2:155:C:C6	2.49	0.47
20:F2:25:PRO:HG2	20:F2:28:PHE:CE2	2.49	0.47
21:F3:28:ARG:HH12	42:O3:147:ARG:HG2	1.80	0.47
23:G2:279:ARG:O	23:G2:283:LYS:HG3	2.14	0.47
24:G3:26:GLN:HG2	83:m2:1221:C:O2'	2.15	0.47
28:I2:9:LEU:HD23	28:I2:118:MET:HB2	1.96	0.47
29:I3:14:HIS:HD1	29:I3:41:ILE:HB	1.79	0.47
34:L1:86:ASP:CG	34:L1:87:ILE:H	2.23	0.47
34:L1:181:TYR:HA	34:L1:184:HIS:HE1	1.78	0.47
36:L3:173:VAL:HG21	83:m2:563:A:N7	2.29	0.47
38:M3:42:LEU:HD11	38:M3:68:LEU:HB3	1.95	0.47
38:M3:58:GLU:O	38:M3:58:GLU:HG2	2.13	0.47
44:P3:35:VAL:O	44:P3:39:THR:HG23	2.14	0.47
59:Z2:48:ALA:HB2	59:Z2:71:TRP:CZ3	2.49	0.47
70:k2:16:PHE:HB3	70:k2:27:THR:H	1.79	0.47
80:x2:56:LEU:HD22	80:x2:80:LEU:HD12	1.97	0.47
83:m2:73:C:H2'	83:m2:76:U:O4	2.14	0.47
83:m2:129:C:H1'	83:m2:130:G:H5''	1.96	0.47
83:m2:199:U:H3'	83:m2:200:C:C5'	2.45	0.47
83:m2:498:C:H2'	83:m2:499:C:C6	2.49	0.47
83:m2:876:G:H2'	83:m2:877:A:C8	2.43	0.47
83:m2:1851:G:H2'	83:m2:1852:A:C8	2.49	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:281:U:HO2'	2:A2:329:A:HI1'	1.80	0.47
2:A2:512:C:H2'	2:A2:513:U:O4'	2.15	0.47
2:A2:835:U:H2'	2:A2:836:C:C6	2.49	0.47
2:A2:1337:A2M:H8	2:A2:3571:U:O2	2.15	0.47
2:A2:2445:C:H2'	2:A2:2446:U:C6	2.50	0.47
6:B3:28:LEU:HD11	6:B3:54:TYR:CE1	2.50	0.47
11:C2:50:C:H2'	65:f2:26:TRP:CZ3	2.49	0.47
13:D1:103:LEU:HD13	13:D1:113:THR:HG22	1.96	0.47
17:E2:86:VAL:HG12	17:E2:201:LEU:HD12	1.97	0.47
39:N2:39:ILE:HD13	39:N2:102:ARG:HD3	1.95	0.47
39:N2:51:GLY:HA3	39:N2:92:ARG:HG3	1.97	0.47
47:R2:73:HIS:CD2	47:R2:115:LYS:HG2	2.50	0.47
51:T2:25:ILE:HG23	51:T2:41:ALA:HB1	1.96	0.47
58:Y2:90:MET:HG3	70:k2:33:LYS:HA	1.97	0.47
64:e2:27:LYS:HG2	64:e2:32:VAL:HG22	1.97	0.47
68:i2:14:LYS:NZ	68:i2:77:CYS:HA	2.30	0.47
75:s2:73:THR:HG22	75:s2:93:VAL:HG21	1.96	0.47
83:m2:89:C:O2'	83:m2:501:G:H5''	2.14	0.47
83:m2:106:C:H2'	83:m2:107:A:C8	2.49	0.47
83:m2:190:G:N3	83:m2:190:G:H2'	2.30	0.47
2:A2:1508:U:H2'	2:A2:1509:C:C6	2.50	0.47
2:A2:2168:U:H2'	2:A2:2169:G:H8	1.80	0.47
2:A2:3285:A:O4'	83:m2:1723:U:C2	2.68	0.47
11:C2:88:A:H2'	11:C2:89:U:O4'	2.15	0.47
24:G3:62:GLU:OE1	42:O3:121:ARG:CD	2.63	0.47
31:J3:146:GLU:HG3	73:q2:116:ARG:CZ	2.44	0.47
38:M3:129:LYS:HD3	38:M3:132:LYS:HE2	1.96	0.47
83:m2:1206:A:H2'	83:m2:1207:C:C6	2.50	0.47
83:m2:1616:A:H2'	83:m2:1617:U:C6	2.49	0.47
2:A2:1085:C:H5	2:A2:1910:G:H21	1.63	0.47
2:A2:1697:G:H2'	2:A2:1698:A:O4'	2.14	0.47
2:A2:4176:G:N3	17:E2:252:ALA:HB1	2.29	0.47
2:A2:4351:U:H4'	2:A2:4352:A:OP1	2.15	0.47
2:A2:4663:G:P	77:u2:92:ARG:CG	3.00	0.47
19:F1:16:LYS:O	19:F1:16:LYS:HG2	2.14	0.47
19:F1:94:ILE:HG23	19:F1:124:LEU:HD22	1.97	0.47
19:F1:177:LYS:HB3	19:F1:180:ALA:HB3	1.97	0.47
33:K3:190:ARG:NH1	33:K3:194:LEU:HD22	2.29	0.47
47:R2:83:THR:O	47:R2:87:MET:HG2	2.15	0.47
53:U2:71:PRO:HG2	53:U2:108:TYR:HA	1.97	0.47
72:p2:44:ILE:HD11	72:p2:69:VAL:HB	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
83:m2:1015:U:H2'	83:m2:1016:G:H8	1.80	0.47
83:m2:1099:G:H2'	83:m2:1100:C:H6	1.79	0.47
83:m2:1103:U:H2'	83:m2:1104:G:C8	2.49	0.47
83:m2:1751:G:H2'	83:m2:1752:C:C6	2.50	0.47
2:A2:113:A:H1'	25:H1:50:ARG:HA	1.97	0.46
2:A2:689:G:H2'	2:A2:690:C:C6	2.50	0.46
2:A2:740:A:N6	2:A2:828:G:H21	2.04	0.46
2:A2:755:G:OP1	37:M2:134:ALA:HB3	2.14	0.46
2:A2:1233:A:H2'	2:A2:1233:A:N3	2.30	0.46
2:A2:1695:C:H1'	2:A2:1739:C:O2	2.14	0.46
2:A2:4133:U:H2'	2:A2:4134:U:C6	2.50	0.46
2:A2:4700:C:H4'	2:A2:4701:G:C8	2.50	0.46
3:A3:121:ARG:HD2	83:m2:1612:G:OP1	2.15	0.46
17:E2:238:LYS:HE2	17:E2:238:LYS:HB2	1.66	0.46
19:F1:140:SER:HA	19:F1:143:GLU:HB2	1.97	0.46
22:G1:41:PRO:HB3	22:G1:70:GLN:HG3	1.97	0.46
25:H1:178:HIS:NE2	25:H1:179:LYS:HE3	2.30	0.46
30:J2:24:VAL:HG12	30:J2:86:LYS:HG2	1.97	0.46
40:N3:18:TYR:HE2	76:t2:139:ILE:HG22	1.80	0.46
48:R3:66:LYS:HD2	48:R3:67:LEU:N	2.30	0.46
64:e2:31:ASN:HB3	64:e2:48:THR:HG22	1.97	0.46
7:n2:2:C:H2'	7:n2:3:C:C6	2.50	0.46
83:m2:13:C:H4'	83:m2:1357:C:O2	2.14	0.46
83:m2:617:C:H2'	83:m2:618:A:O4'	2.15	0.46
83:m2:942:U:H2'	83:m2:943:C:C6	2.50	0.46
1:A1:76:ALA:HB2	1:A1:210:GLU:HG3	1.97	0.46
2:A2:666:G:H2'	2:A2:667:G:H8	1.80	0.46
2:A2:1347:A2M:HM'3	2:A2:1450:A:C2	2.50	0.46
2:A2:1404:U:OP2	17:E2:243:LYS:HE2	2.15	0.46
2:A2:1707:U:P	59:Z2:76:ARG:HH12	2.38	0.46
2:A2:2498:A:H1'	14:D2:21:LYS:NZ	2.30	0.46
2:A2:4417:G:N2	37:M2:173:ASN:HD21	2.12	0.46
2:A2:4676:U:H2'	2:A2:4677:G:H8	1.79	0.46
15:D3:11:LEU:HD11	31:J3:136:HIS:CE1	2.50	0.46
15:D3:40:ASP:HB2	15:D3:43:THR:HB	1.96	0.46
17:E2:65:SER:OG	17:E2:67:VAL:HG22	2.15	0.46
17:E2:207:VAL:HG21	17:E2:331:ILE:HD11	1.97	0.46
25:H1:45:PRO:O	25:H1:49:ARG:HG3	2.14	0.46
31:J3:133:TYR:HD1	31:J3:216:MET:HA	1.80	0.46
36:L3:46:VAL:HG11	36:L3:106:LEU:HD21	1.96	0.46
42:O3:45:THR:HG22	42:O3:52:THR:HA	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
60:a2:65:MET:HB2	60:a2:65:MET:HE3	1.78	0.46
72:p2:38:MET:HE3	72:p2:231:LEU:HD21	1.97	0.46
72:p2:109:LYS:O	72:p2:113:MET:HE2	2.15	0.46
77:u2:81:VAL:HG22	77:u2:102:VAL:HG12	1.96	0.46
83:m2:377:U:H2'	83:m2:378:A:H8	1.80	0.46
83:m2:869:OMG:HM23	83:m2:869:OMG:H1'	1.70	0.46
83:m2:1449:G:H2'	83:m2:1450:A:C8	2.50	0.46
2:A2:416:U:H4'	2:A2:2085:G:H4'	1.97	0.46
2:A2:4406:G:N7	26:H2:285:THR:HB	2.30	0.46
4:B1:157:ILE:HB	4:B1:183:ILE:HD13	1.97	0.46
12:C3:32:LEU:HD21	12:C3:87:ARG:HG2	1.97	0.46
14:D2:49:ILE:HD13	14:D2:60:LYS:HE3	1.96	0.46
17:E2:200:ARG:HB2	17:E2:203:GLN:HB2	1.97	0.46
19:F1:42:LYS:HG3	19:F1:45:ARG:NH2	2.29	0.46
31:J3:238:LYS:NZ	83:m2:1357:C:O2'	2.35	0.46
46:Q3:108:LYS:HD3	83:m2:53:C:O3'	2.15	0.46
48:R3:111:ARG:HD2	48:R3:114:LYS:HB2	1.97	0.46
74:r2:7:LYS:HB2	83:m2:93:U:O2'	2.14	0.46
83:m2:186:C:H2'	83:m2:187:G:H8	1.79	0.46
83:m2:495:A:H1'	83:m2:576:A:O5'	2.15	0.46
83:m2:1012:G:H2'	83:m2:1013:A:C8	2.50	0.46
83:m2:1409:U:H2'	83:m2:1410:U:C6	2.51	0.46
2:A2:175:C:H2'	2:A2:176:G:C8	2.50	0.46
2:A2:3903:A:H2'	2:A2:3904:C:C6	2.51	0.46
3:A3:86:ARG:HE	3:A3:89:ASP:CG	2.24	0.46
21:F3:38:LYS:HG2	83:m2:1869:U:O2'	2.15	0.46
23:G2:232:THR:HB	23:G2:235:MET:HG2	1.98	0.46
33:K3:7:PHE:CE2	33:K3:9:ALA:HB3	2.50	0.46
44:P3:39:THR:O	44:P3:43:LYS:HG2	2.15	0.46
48:R3:91:LEU:HB3	48:R3:97:ILE:HG12	1.96	0.46
51:T2:33:THR:HG22	51:T2:35:ASP:H	1.80	0.46
57:X2:32:ARG:HB3	57:X2:48:GLU:HG3	1.98	0.46
62:c2:29:ARG:HG2	62:c2:32:ARG:HD3	1.97	0.46
72:p2:59:SER:HB3	72:p2:63:LYS:HE2	1.98	0.46
74:r2:47:PHE:O	74:r2:51:ARG:HB2	2.14	0.46
75:s2:39:ILE:HG22	75:s2:41:VAL:HG22	1.98	0.46
75:s2:143:PRO:O	75:s2:146:ARG:HG2	2.15	0.46
83:m2:890:U:H2'	83:m2:891:U:C6	2.50	0.46
83:m2:1590:A:H2'	83:m2:1591:A:H8	1.79	0.46
1:A1:109:PRO:HG3	1:A1:166:TYR:CD2	2.50	0.46
2:A2:740:A:O2'	37:M2:72:PRO:HD2	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:1118:C:H2'	2:A2:1119:C:C6	2.51	0.46
2:A2:2037:A:O3'	53:U2:20:GLY:HA2	2.16	0.46
2:A2:2256:C:C2'	2:A2:2257:G:H5'	2.45	0.46
2:A2:2365:G:H2'	2:A2:2366:A:H8	1.81	0.46
2:A2:2438:C:H2'	2:A2:2439:C:C6	2.51	0.46
2:A2:3363:U:H2'	2:A2:3364:C:H6	1.80	0.46
2:A2:4051:U:H2'	2:A2:4052:G:O4'	2.15	0.46
2:A2:4289:OMG:H1'	2:A2:4289:OMG:HM23	1.63	0.46
4:B1:171:PRO:HB3	4:B1:181:TYR:CZ	2.50	0.46
37:M2:127:MET:HA	39:N2:153:PRO:HD2	1.98	0.46
44:P3:3:ARG:HH12	44:P3:28:ARG:HE	1.64	0.46
56:W2:57:LYS:O	56:W2:61:GLU:HG3	2.15	0.46
59:Z2:36:ARG:HD2	59:Z2:79:GLY:O	2.16	0.46
59:Z2:46:ARG:HD3	59:Z2:106:TYR:HD2	1.80	0.46
64:e2:56:LEU:HD12	64:e2:59:SER:OG	2.16	0.46
73:q2:202:LYS:HB2	73:q2:202:LYS:HE2	1.75	0.46
74:r2:52:LEU:HD11	74:r2:99:PHE:HE2	1.80	0.46
83:m2:532:U:H2'	83:m2:533:A:C5	2.51	0.46
83:m2:1201:A:H2'	83:m2:1202:A:H8	1.81	0.46
83:m2:1246:U:H2'	83:m2:1247:G:C8	2.50	0.46
83:m2:1392:U:H2'	83:m2:1393:C:H6	1.80	0.46
83:m2:1752:C:H2'	83:m2:1753:C:O4'	2.16	0.46
2:A2:723:C:OP1	20:F2:317:ASN:HB2	2.16	0.46
2:A2:1475:C:H2'	2:A2:1476:C:C6	2.50	0.46
2:A2:1496:PSU:H2'	2:A2:1497:A:C8	2.51	0.46
2:A2:2247:C:H2'	2:A2:2248:G:H8	1.81	0.46
2:A2:3412:A:H61	2:A2:3423:C:H42	1.62	0.46
2:A2:3721:U:H2'	2:A2:3722:C:C6	2.50	0.46
2:A2:3844:A:H2'	2:A2:3845:C:H6	1.81	0.46
2:A2:3884:U:O2'	68:i2:5:PRO:HD3	2.15	0.46
3:A3:40:TYR:HD1	3:A3:83:PHE:CE2	2.32	0.46
4:B1:134:PRO:HB3	4:B1:206:GLN:HB2	1.96	0.46
18:E3:94:ILE:HG13	18:E3:125:VAL:HG11	1.97	0.46
42:O3:82:ALA:HA	42:O3:124:MET:HE3	1.98	0.46
59:Z2:14:TYR:HB3	59:Z2:22:ARG:O	2.15	0.46
7:n2:30:G:H2'	7:n2:31:A:O4'	2.15	0.46
79:w2:79:LYS:HB3	79:w2:81:LYS:HD3	1.98	0.46
80:x2:25:LEU:HD21	80:x2:36:LEU:HD22	1.98	0.46
83:m2:158:A:H1'	83:m2:159:A:H5'	1.97	0.46
83:m2:423:G:H4'	83:m2:662:C:N4	2.31	0.46
83:m2:1219:A:H2'	83:m2:1220:C:H6	1.81	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
83:m2:1715:C:H2'	83:m2:1716:U:C6	2.50	0.46
2:A2:518:C:H2'	2:A2:519:G:H8	1.81	0.46
2:A2:1180:G:H4'	2:A2:1181:U:H5	1.80	0.46
2:A2:1667:G:H5'	13:D1:118:ALA:O	2.16	0.46
2:A2:4544:G:H5''	2:A2:4545:U:OP2	2.16	0.46
3:A3:26:ILE:HD13	3:A3:56:ALA:HA	1.97	0.46
14:D2:180:LEU:HD23	14:D2:181:LYS:HG3	1.97	0.46
19:F1:161:TYR:CE2	53:U2:105:ARG:HG3	2.51	0.46
25:H1:94:PHE:CE2	25:H1:96:ARG:HB2	2.50	0.46
28:I2:110:PRO:N	28:I2:111:PRO:HD2	2.31	0.46
29:I3:121:VAL:HG11	29:I3:165:ILE:HD11	1.98	0.46
45:Q2:3:VAL:HG13	45:Q2:13:ILE:O	2.16	0.46
74:r2:158:ASP:HB3	74:r2:173:ILE:O	2.15	0.46
74:r2:188:ASN:HB3	74:r2:191:ARG:HD3	1.97	0.46
75:s2:94:LYS:O	75:s2:98:GLU:HG3	2.16	0.46
83:m2:1753:C:H1'	83:m2:1786:G:N1	2.30	0.46
2:A2:939:C:H2'	2:A2:940:C:C6	2.49	0.46
2:A2:1258:U:H2'	2:A2:1259:C:O4'	2.15	0.46
2:A2:3834:G:H5'	14:D2:226:ARG:HH12	1.81	0.46
3:A3:9:PHE:CE1	3:A3:57:GLY:HA3	2.50	0.46
7:Bz:36:A:H1'	83:m2:628:G:N2	2.30	0.46
16:E1:40:LEU:HD23	16:E1:40:LEU:HA	1.79	0.46
21:F3:10:ARG:HB2	21:F3:33:ASP:OD2	2.16	0.46
35:L2:119:MET:HE1	35:L2:142:ILE:HG23	1.98	0.46
53:U2:31:GLY:HA3	53:U2:35:ALA:HB3	1.97	0.46
71:o2:78:SER:HB3	71:o2:87:VAL:HG21	1.98	0.46
73:q2:93:THR:OG1	73:q2:96:LEU:HB2	2.16	0.46
80:x2:93:MET:HG3	80:x2:93:MET:O	2.16	0.46
82:z2:34:VAL:O	82:z2:38:ILE:HG12	2.16	0.46
83:m2:37:C:H2'	83:m2:38:A:O4'	2.16	0.46
83:m2:447:A:H2'	83:m2:448:G:C8	2.51	0.46
83:m2:895:U:H2'	83:m2:896:G:O4'	2.16	0.46
1:A1:204:PHE:CZ	1:A1:225:GLU:HG2	2.50	0.46
2:A2:3763:U:H2'	2:A2:3764:C:C6	2.50	0.46
7:Bz:8:U:H2'	7:Bz:46:G:H21	1.80	0.46
10:C1:90:TYR:CE2	10:C1:155:SER:HB3	2.51	0.46
10:C1:107:GLU:HB2	10:C1:110:SER:HB2	1.98	0.46
17:E2:291:TYR:CE2	17:E2:327:THR:HG22	2.51	0.46
21:F3:87:ARG:HD2	21:F3:92:ARG:HA	1.97	0.46
34:L1:196:LYS:HB3	34:L1:199:GLN:HB2	1.98	0.46
41:O2:42:PHE:CE2	41:O2:90:TYR:HB2	2.50	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
43:P2:43:LYS:HD3	43:P2:60:MET:SD	2.56	0.46
62:c2:33:LEU:HD11	62:c2:38:LYS:HB2	1.98	0.46
72:p2:146:ARG:HH11	72:p2:206:PRO:HB2	1.80	0.46
74:r2:15:PRO:HD2	74:r2:18:TRP:CZ3	2.50	0.46
75:s2:68:ILE:HD11	75:s2:151:ILE:HD11	1.97	0.46
76:t2:48:ALA:HA	76:t2:62:ILE:HA	1.97	0.46
77:u2:144:LYS:HD3	83:m2:193:C:H41	1.80	0.46
2:A2:3749:G:H2'	2:A2:3750:C:C6	2.51	0.46
15:D3:83:PHE:CZ	71:o2:53:ARG:HD2	2.51	0.46
20:F2:13:GLU:HB3	20:F2:155:GLU:OE2	2.16	0.46
23:G2:211:LEU:HD23	23:G2:211:LEU:HA	1.79	0.46
27:H3:34:TYR:CE2	73:q2:8:LYS:HD2	2.51	0.46
46:Q3:17:LEU:HD23	74:r2:64:ILE:HG21	1.98	0.46
52:T3:28:LYS:O	52:T3:33:LYS:HG3	2.16	0.46
57:X2:53:ALA:HA	57:X2:88:LEU:HD21	1.98	0.46
71:o2:112:ILE:CD1	83:m2:1352:U:H4'	2.46	0.46
75:s2:113:VAL:HA	75:s2:116:ILE:HG22	1.97	0.46
82:z2:9:VAL:HG13	82:z2:50:ILE:HD13	1.97	0.46
83:m2:115:U:H2'	83:m2:116:OMU:C6	2.46	0.46
83:m2:1283:G:H3'	83:m2:1284:A:H8	1.81	0.46
2:A2:10:A:H2'	2:A2:11:G:C8	2.52	0.45
2:A2:3912:U:H2'	2:A2:3913:C:H6	1.81	0.45
7:Bv:37:A:C4	7:Bv:38:A:C8	3.04	0.45
7:Bz:35:A:H2'	7:Bz:36:A:C8	2.51	0.45
10:C1:18:ILE:HD11	10:C1:81:ILE:HD11	1.97	0.45
21:F3:7:ASN:HB3	21:F3:10:ARG:O	2.16	0.45
22:G1:17:PHE:CE2	22:G1:54:CYS:HA	2.51	0.45
32:K2:178:ARG:N	53:U2:51:GLY:HA2	2.31	0.45
49:S2:126:ARG:HG2	49:S2:130:LYS:HE3	1.98	0.45
52:T3:8:ARG:HG2	83:m2:618:A:H5'	1.98	0.45
58:Y2:68:HIS:CE1	70:k2:22:LYS:HG3	2.51	0.45
72:p2:169:MET:O	72:p2:173:THR:HG23	2.17	0.45
72:p2:173:THR:O	72:p2:177:GLN:HG2	2.16	0.45
83:m2:200:C:O2	83:m2:200:C:H2'	2.16	0.45
83:m2:1716:U:H2'	83:m2:1717:A:H8	1.80	0.45
83:m2:1804:C:H2'	83:m2:1805:U:H6	1.80	0.45
2:A2:35:U:H4'	2:A2:1338:A:C2	2.51	0.45
2:A2:398:A2M:HM'3	2:A2:398:A2M:H1'	1.84	0.45
2:A2:1650:C:H2'	2:A2:1651:U:O4'	2.16	0.45
2:A2:3840:U:H2'	2:A2:3841:U:C6	2.52	0.45
2:A2:4230:G:H2'	2:A2:4231:U:C6	2.51	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:4395:G:H2'	2:A2:4396:A:H8	1.77	0.45
3:A3:12:ILE:HG21	16:E1:119:TYR:HE2	1.81	0.45
3:A3:89:ASP:HB3	3:A3:92:ASP:O	2.16	0.45
5:B2:39:C:O2'	16:E1:46:GLN:HB3	2.16	0.45
12:C3:55:ARG:HG2	12:C3:87:ARG:CZ	2.46	0.45
26:H2:73:LYS:HE2	26:H2:75:LEU:HD21	1.97	0.45
33:K3:63:MET:HE3	33:K3:100:CYS:HA	1.99	0.45
40:N3:7:PRO:HG3	83:m2:998:A:H5''	1.98	0.45
60:a2:24:ARG:HH11	60:a2:24:ARG:HG3	1.80	0.45
67:h2:16:LYS:HZ1	83:m2:1820:A:C4'	2.17	0.45
73:q2:76:ARG:HB2	78:v2:22:VAL:HG11	1.98	0.45
2:A2:294:G:O6	2:A2:315:G:H1'	2.16	0.45
2:A2:1285:C:H2'	2:A2:1286:U:H6	1.81	0.45
2:A2:1535:G:N3	2:A2:3866:A:H2'	2.32	0.45
6:B3:115:LYS:HE2	6:B3:121:ARG:NH1	2.30	0.45
17:E2:285:TYR:HA	17:E2:363:ILE:HD11	1.98	0.45
26:H2:76:TYR:CE1	26:H2:77:LYS:HG3	2.51	0.45
29:I3:32:LEU:HD21	29:I3:71:ILE:HG12	1.99	0.45
39:N2:18:PRO:HB2	39:N2:21:LYS:HD2	1.97	0.45
49:S2:47:MET:HE3	49:S2:47:MET:HB3	1.85	0.45
57:X2:87:ARG:NH1	57:X2:122:VAL:HG11	2.31	0.45
58:Y2:105:SER:O	58:Y2:109:LYS:HG3	2.16	0.45
7:n2:73:A:H2'	7:n2:74:C:H4'	1.99	0.45
83:m2:1353:G:C4	83:m2:1381:A:N1	2.84	0.45
83:m2:1381:A:H2'	83:m2:1382:C:C6	2.51	0.45
83:m2:1449:G:H2'	83:m2:1450:A:H8	1.81	0.45
83:m2:1856:U:H2'	83:m2:1857:G:C8	2.50	0.45
1:A1:75:LYS:HE2	1:A1:211:ASP:OD1	2.16	0.45
2:A2:1626:G:H2'	2:A2:1627:A:C8	2.50	0.45
2:A2:1702:C:H4'	58:Y2:60:TYR:CZ	2.51	0.45
2:A2:3443:G:H1'	2:A2:3445:C:N4	2.31	0.45
2:A2:4289:OMG:H4'	2:A2:4691:G:O2'	2.16	0.45
2:A2:4431:C:H2'	2:A2:4432:C:H6	1.81	0.45
7:Bv:34:G:C8	83:m2:1250:B8N:N34	2.84	0.45
14:D2:158:ILE:HB	14:D2:162:ASN:ND2	2.32	0.45
20:F2:76:ILE:HG12	20:F2:96:CYS:SG	2.57	0.45
23:G2:191:ASN:OD1	23:G2:194:VAL:HG23	2.16	0.45
32:K2:154:LYS:HD2	32:K2:158:THR:HG21	1.97	0.45
36:L3:63:LEU:HD13	36:L3:67:ASP:OD2	2.16	0.45
36:L3:114:VAL:HG13	36:L3:119:LEU:HB2	1.97	0.45
42:O3:26:ASN:C	42:O3:26:ASN:ND2	2.73	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:O3:103:ASN:ND2	42:O3:142:ARG:HA	2.30	0.45
58:Y2:118:LEU:C	70:k2:119:ARG:HH12	2.24	0.45
74:r2:100:ARG:HH21	74:r2:118:GLU:HG2	1.82	0.45
81:y2:77:HIS:O	81:y2:81:ILE:HG12	2.15	0.45
83:m2:210:G:C2'	83:m2:211:A:H5'	2.46	0.45
83:m2:380:U:H2'	83:m2:381:C:C6	2.52	0.45
83:m2:651:U:H2'	83:m2:652:A:C8	2.52	0.45
83:m2:806:U:H2'	83:m2:807:U:H6	1.82	0.45
2:A2:757:U:C2	2:A2:758:G:C8	3.04	0.45
2:A2:1473:U:OP1	53:U2:15:VAL:HA	2.16	0.45
2:A2:2048:U:H2'	2:A2:2049:G:C8	2.51	0.45
2:A2:2142:G:H5''	35:L2:23:TRP:CD1	2.51	0.45
2:A2:4027:C:H5'	2:A2:4029:G:H5'	1.98	0.45
2:A2:4352:A:O2'	10:C1:65:LYS:HD2	2.16	0.45
3:A3:48:ALA:O	3:A3:50:ILE:HG13	2.17	0.45
4:B1:100:HIS:CE1	4:B1:103:ARG:HH11	2.34	0.45
7:Bz:6:G:O2'	7:Bz:7:A:H5'	2.17	0.45
7:Bz:18:G:H5''	7:Bz:19:G:OP2	2.17	0.45
17:E2:103:LYS:O	17:E2:153:MET:HE2	2.17	0.45
17:E2:194:LEU:HD23	17:E2:194:LEU:HA	1.76	0.45
17:E2:196:TRP:CE2	17:E2:200:ARG:NH1	2.84	0.45
17:E2:284:ILE:HG23	17:E2:331:ILE:HG23	1.98	0.45
30:J2:111:SER:HB2	30:J2:154:GLU:HB2	1.98	0.45
36:L3:81:LEU:HA	36:L3:84:ILE:HG22	1.98	0.45
38:M3:111:VAL:C	38:M3:112:LYS:HZ2	2.24	0.45
44:P3:60:LYS:HB2	44:P3:60:LYS:HE3	1.73	0.45
79:w2:93:LEU:HD22	79:w2:102:PHE:HB3	1.97	0.45
83:m2:1310:U:H2'	83:m2:1311:C:C6	2.52	0.45
83:m2:1713:U:H2'	83:m2:1714:A:C8	2.50	0.45
83:m2:1753:C:H1'	83:m2:1786:G:H1	1.80	0.45
83:m2:1843:C:H2'	83:m2:1844:4AC:H6	1.99	0.45
2:A2:2357:G:H2'	2:A2:2358:C:C6	2.52	0.45
2:A2:2440:C:H2'	2:A2:2441:G:O4'	2.17	0.45
6:B3:104:LEU:HD13	6:B3:121:ARG:HH21	1.82	0.45
12:C3:55:ARG:HD2	83:m2:1448:A:N3	2.30	0.45
14:D2:107:MET:HE2	14:D2:166:VAL:HG22	1.99	0.45
20:F2:140:LYS:NZ	20:F2:242:PRO:HD2	2.32	0.45
20:F2:336:ARG:HD2	26:H2:61:ILE:HD11	1.98	0.45
21:F3:10:ARG:HH12	83:m2:1861:A:P	2.40	0.45
28:I2:47:PHE:CE1	28:I2:140:ARG:HG2	2.51	0.45
57:X2:38:PHE:HB3	57:X2:78:ARG:HG2	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
74:r2:45:ILE:HG13	74:r2:80:ILE:O	2.17	0.45
83:m2:72:C:H2'	83:m2:73:C:C6	2.51	0.45
83:m2:872:A:H62	83:m2:918:A:H5'	1.82	0.45
83:m2:1448:A:O2'	83:m2:1449:G:H5''	2.16	0.45
83:m2:1682:G:H2'	83:m2:1683:U:C6	2.51	0.45
83:m2:1819:G:H2'	83:m2:1820:A:C8	2.52	0.45
2:A2:1021:G:N3	2:A2:1021:G:H2'	2.31	0.45
2:A2:1190:C:H3'	2:A2:1191:G:C8	2.52	0.45
2:A2:2145:G:H22	2:A2:2580:A:H2	1.65	0.45
2:A2:3267:A:H2	2:A2:4662:A:C8	2.34	0.45
2:A2:4268:A:H2'	2:A2:4269:G:O4'	2.17	0.45
5:B2:4:U:H2'	5:B2:5:A:H8	1.80	0.45
12:C3:108:PRO:HB3	73:q2:40:ARG:HG2	1.98	0.45
29:I3:165:ILE:HG22	29:I3:177:TRP:HB2	1.98	0.45
46:Q3:62:THR:HA	46:Q3:69:THR:HG22	1.99	0.45
59:Z2:45:LYS:HD2	59:Z2:105:LEU:HA	1.99	0.45
66:g2:112:LYS:HB3	66:g2:114:LYS:HG2	1.99	0.45
72:p2:142:PHE:C	72:p2:207:LEU:HG	2.41	0.45
75:s2:167:LYS:HE3	75:s2:172:CYS:SG	2.57	0.45
83:m2:18:C:C2	83:m2:19:A:C8	3.04	0.45
83:m2:318:G:H2'	83:m2:319:C:C6	2.52	0.45
83:m2:1432:C:H2'	83:m2:1433:G:H8	1.80	0.45
83:m2:1696:U:H2'	83:m2:1698:C:H5	1.81	0.45
2:A2:37:U:H2'	2:A2:38:A:O4'	2.16	0.45
2:A2:308:G:H2'	62:c2:41:ARG:HH12	1.82	0.45
2:A2:883:C:H2'	2:A2:884:C:O4'	2.15	0.45
2:A2:1078:G:H3'	2:A2:1079:A:C8	2.52	0.45
2:A2:2333:G:O6	51:T2:17:ARG:HD2	2.16	0.45
2:A2:2447:U:O2	64:e2:2:PRO:HB2	2.17	0.45
2:A2:2625:A:H2'	2:A2:2626:A:C8	2.52	0.45
2:A2:4412:G:H2'	2:A2:4413:G:O4'	2.17	0.45
2:A2:4508:G:H2'	2:A2:4509:U:C6	2.52	0.45
3:A3:23:ARG:HA	48:R3:48:VAL:HG21	1.97	0.45
7:Bz:37:A:H3'	7:Bz:38:A:H8	1.82	0.45
11:C2:94:G:C5	63:d2:84:PRO:HG3	2.52	0.45
12:C3:37:ALA:O	12:C3:41:ARG:HG2	2.17	0.45
20:F2:11:TYR:OH	20:F2:148:PRO:HB2	2.16	0.45
22:G1:38:VAL:HG21	22:G1:50:MET:SD	2.57	0.45
25:H1:88:GLY:O	25:H1:92:LEU:HD11	2.16	0.45
33:K3:32:MET:HB3	33:K3:65:GLN:NE2	2.32	0.45
33:K3:195:LYS:HE3	83:m2:126:G:N9	2.32	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
63:d2:52:LYS:HG2	63:d2:55:ARG:HH22	1.80	0.45
71:o2:33:GLN:HB3	71:o2:154:LEU:HD12	1.99	0.45
71:o2:50:ASN:O	71:o2:54:THR:HG23	2.17	0.45
75:s2:20:PHE:CE2	75:s2:94:LYS:HB2	2.52	0.45
83:m2:94:G:C4	83:m2:95:G:C8	3.05	0.45
83:m2:961:G:H2'	83:m2:962:U:C6	2.52	0.45
83:m2:1675:U:H2'	83:m2:1676:G:O4'	2.17	0.45
83:m2:1799:U:H2'	83:m2:1800:C:H6	1.82	0.45
2:A2:362:A:C6	65:f2:35:ILE:HG23	2.52	0.45
2:A2:2170:U:H2'	2:A2:2171:G:O4'	2.17	0.45
2:A2:2382:C:H2'	2:A2:2383:U:C6	2.52	0.45
2:A2:3340:G:H2'	2:A2:3341:C:H6	1.82	0.45
2:A2:3996:U:H2'	2:A2:3997:C:C6	2.52	0.45
2:A2:4156:C:H2'	2:A2:4157:C:C6	2.51	0.45
2:A2:4700:C:H4'	2:A2:4701:G:H8	1.82	0.45
3:A3:64:VAL:O	3:A3:68:ILE:HG12	2.17	0.45
12:C3:48:LEU:HD13	12:C3:91:LEU:HD21	1.99	0.45
14:D2:150:LEU:HB3	14:D2:151:PRO:HD2	1.98	0.45
32:K2:15:ARG:HB3	32:K2:52:PHE:HB3	1.98	0.45
40:N3:124:ARG:HD2	83:m2:679:G:OP1	2.17	0.45
44:P3:14:ILE:HG13	44:P3:27:ILE:HD11	1.98	0.45
51:T2:74:VAL:HG23	51:T2:101:PHE:HE2	1.81	0.45
72:p2:52:THR:OG1	72:p2:57:ILE:HA	2.16	0.45
83:m2:350:A:H2'	83:m2:351:A:C8	2.52	0.45
83:m2:1036:A:C2	83:m2:1037:A:H1'	2.52	0.45
83:m2:1185:A:H2'	83:m2:1186:G:H8	1.81	0.45
2:A2:360:A:C4	11:C2:24:G:H1'	2.52	0.45
2:A2:426:A:H2'	2:A2:427:A:C8	2.52	0.45
2:A2:1068:C:H2'	2:A2:1069:G:C8	2.51	0.45
2:A2:1154:OMC:H1'	2:A2:1154:OMC:HM23	1.68	0.45
2:A2:1474:C:H2'	2:A2:1475:C:C6	2.53	0.45
2:A2:1668:G:C2	2:A2:4057:G:C8	3.05	0.45
2:A2:3898:G:H2'	2:A2:3899:G:C8	2.52	0.45
2:A2:4284:U:H2'	2:A2:4285:G:O4'	2.17	0.45
2:A2:4546:G:H2'	2:A2:4547:C:C6	2.52	0.45
4:B1:58:PRO:HG2	4:B1:61:ILE:HD12	1.99	0.45
5:B2:64:G:H2'	5:B2:65:G:H8	1.81	0.45
12:C3:63:ILE:HG22	12:C3:65:THR:HG23	1.99	0.45
14:D2:210:PRO:HG2	14:D2:235:VAL:HG21	1.99	0.45
16:E1:134:LEU:HD23	16:E1:134:LEU:HA	1.80	0.45
34:L1:84:HIS:NE2	34:L1:86:ASP:HB2	2.32	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:O3:137:SER:HB2	83:m2:944:G:H21	1.82	0.45
63:d2:37:CYS:HA	63:d2:45:ARG:HB3	1.99	0.45
68:i2:22:LYS:NZ	68:i2:73:VAL:HG12	2.31	0.45
76:t2:73:GLN:HB3	76:t2:135:PHE:CE1	2.52	0.45
77:u2:190:LEU:HD23	77:u2:190:LEU:HA	1.84	0.45
83:m2:129:C:H42	83:m2:181:A:H61	1.65	0.45
83:m2:530:A:H2'	83:m2:531:A:C8	2.51	0.45
83:m2:1558:A:H2	83:m2:1559:C:C4	2.35	0.45
1:A1:73:TYR:CZ	20:F2:331:TYR:HD1	2.35	0.44
1:A1:206:ILE:HD11	1:A1:215:GLU:HG2	1.99	0.44
2:A2:179:G:C6	2:A2:258:G:C6	3.05	0.44
2:A2:2233:C:H3'	2:A2:2234:G:H8	1.82	0.44
2:A2:3430:A:H2'	2:A2:3431:A:C8	2.52	0.44
2:A2:3567:C:H2'	2:A2:3568:U:H6	1.82	0.44
2:A2:3607:G:H2'	2:A2:3608:A:C8	2.52	0.44
2:A2:3627:G:H1	2:A2:3700:A:H61	1.64	0.44
2:A2:3738:C:H2'	2:A2:3739:G:C8	2.53	0.44
2:A2:3758:G:H2'	2:A2:3759:G:H8	1.82	0.44
4:B1:165:GLU:HA	4:B1:168:VAL:HG12	1.99	0.44
7:Bv:50:U:H2'	7:Bv:51:U:C6	2.52	0.44
11:C2:93:C:O2'	11:C2:94:G:H8	2.00	0.44
11:C2:96:C:H5''	61:b2:66:LYS:HG2	1.99	0.44
12:C3:94:PRO:HG2	12:C3:97:ILE:HD12	1.98	0.44
17:E2:247:GLY:HA3	17:E2:250:LYS:HE3	1.98	0.44
27:H3:23:VAL:HG22	78:v2:64:TRP:CD2	2.52	0.44
30:J2:39:MET:HE3	30:J2:43:LYS:HE3	1.99	0.44
54:U3:123:SER:HB3	54:U3:126:CYS:SG	2.57	0.44
61:b2:47:ILE:O	61:b2:51:ARG:HG2	2.17	0.44
71:o2:102:ARG:NH1	83:m2:1379:U:C6	2.85	0.44
73:q2:75:LYS:HZ1	78:v2:18:GLU:HG2	1.82	0.44
83:m2:1294:C:H2'	83:m2:1295:A:H8	1.82	0.44
2:A2:1098:G:H21	26:H2:137:LYS:HG3	1.82	0.44
6:B3:20:ALA:O	6:B3:24:LYS:HG3	2.18	0.44
6:B3:60:THR:O	6:B3:64:LEU:HG	2.17	0.44
7:Bz:39:U:H2'	7:Bz:40:C:C6	2.52	0.44
17:E2:172:PRO:HB2	17:E2:324:GLY:HA3	1.99	0.44
32:K2:168:ARG:HE	32:K2:168:ARG:HB3	1.65	0.44
35:L2:144:LYS:HE2	35:L2:144:LYS:HB3	1.83	0.44
48:R3:69:THR:H	48:R3:72:VAL:HG12	1.82	0.44
58:Y2:35:TRP:CZ2	58:Y2:56:PRO:HD2	2.52	0.44
69:j2:11:VAL:HG11	69:j2:26:VAL:HG13	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
70:k2:98:ARG:HG3	70:k2:98:ARG:NH1	2.31	0.44
71:o2:102:ARG:CZ	83:m2:1379:U:C6	3.00	0.44
80:x2:60:LEU:HD23	80:x2:76:VAL:HG21	2.00	0.44
80:x2:77:LYS:HG2	80:x2:102:PHE:HZ	1.83	0.44
83:m2:1680:A2M:O2'	83:m2:1681:A:H5'	2.17	0.44
2:A2:1684:U:C4	2:A2:2034:A:C2	3.04	0.44
2:A2:1847:G:O2'	2:A2:1848:G:H5''	2.17	0.44
2:A2:2259:C:H4'	2:A2:2260:C:OP2	2.16	0.44
2:A2:2365:G:H2'	2:A2:2366:A:C8	2.51	0.44
2:A2:2505:G:H2'	2:A2:2506:G:O4'	2.18	0.44
2:A2:3579:A:H2'	2:A2:3580:C:H6	1.79	0.44
2:A2:3584:A:H2'	2:A2:3585:G:O4'	2.17	0.44
2:A2:3954:U:H4'	39:N2:5:LYS:HD3	2.00	0.44
2:A2:4041:C:H2'	2:A2:4042:A:C8	2.53	0.44
2:A2:4224:U:H1'	2:A2:4372:C:C4	2.52	0.44
4:B1:87:LEU:HD13	4:B1:91:THR:HG22	1.99	0.44
11:C2:33:G:H4'	11:C2:34:U:C6	2.53	0.44
17:E2:14:LEU:O	17:E2:17:LEU:HB2	2.16	0.44
20:F2:5:ARG:HD2	20:F2:24:LEU:O	2.17	0.44
26:H2:275:GLN:O	26:H2:279:ARG:HG3	2.18	0.44
28:I2:22:ILE:HD13	28:I2:120:VAL:HG11	1.99	0.44
33:K3:64:LYS:HB2	33:K3:97:VAL:HG21	1.99	0.44
71:o2:49:ILE:HG21	71:o2:162:PRO:HB2	1.98	0.44
73:q2:23:GLU:HG2	78:v2:64:TRP:NE1	2.33	0.44
73:q2:40:ARG:HH22	73:q2:49:ILE:HG12	1.83	0.44
73:q2:135:GLU:CD	73:q2:157:MET:HG2	2.42	0.44
74:r2:139:LEU:HG	74:r2:150:PRO:HG3	2.00	0.44
83:m2:289:U:H2'	83:m2:290:G:C8	2.52	0.44
83:m2:833:G:H2'	83:m2:834:G:C8	2.52	0.44
83:m2:951:G:H2'	83:m2:952:C:C6	2.51	0.44
1:A1:263:ASN:O	1:A1:267:ARG:HG2	2.18	0.44
2:A2:279:A:N1	2:A2:306:A:H5''	2.33	0.44
2:A2:465:G:H2'	2:A2:466:A:C8	2.53	0.44
2:A2:518:C:H2'	2:A2:519:G:C8	2.52	0.44
2:A2:2140:U:H2'	2:A2:2141:U:H6	1.82	0.44
2:A2:4022:OMG:H5''	68:i2:64:LYS:HG3	2.00	0.44
23:G2:60:ILE:HB	23:G2:80:ALA:HB2	1.99	0.44
28:I2:85:ARG:HD3	28:I2:90:HIS:CE1	2.53	0.44
29:I3:259:TRP:CD1	29:I3:266:ILE:HG23	2.52	0.44
37:M2:161:ARG:NH2	37:M2:164:LYS:HE3	2.32	0.44
41:O2:36:ALA:HB1	41:O2:65:ARG:HD2	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:R2:126:THR:HG21	47:R2:134:LYS:HE2	2.00	0.44
54:U3:135:HIS:NE2	83:m2:1310:U:H1'	2.33	0.44
64:e2:51:GLU:O	64:e2:55:LYS:HG2	2.17	0.44
80:x2:53:GLN:O	80:x2:57:LEU:HD23	2.17	0.44
83:m2:603:OMG:H1'	83:m2:603:OMG:HM23	1.72	0.44
2:A2:1156:A:OP1	25:H1:204:ARG:HD2	2.17	0.44
2:A2:1288:G:H2'	2:A2:1289:C:C6	2.51	0.44
2:A2:3886:A:N7	2:A2:3923:A:N1	2.65	0.44
2:A2:3958:OMU:OP2	32:K2:159:PRO:HD2	2.17	0.44
2:A2:4138:C:H2'	2:A2:4139:A:O4'	2.16	0.44
2:A2:4325:U:H2'	2:A2:4326:C:C6	2.51	0.44
3:A3:102:GLY:O	3:A3:106:LYS:HG2	2.18	0.44
3:A3:121:ARG:HG3	3:A3:131:VAL:HG13	1.99	0.44
7:Bz:15:G:N2	7:Bz:48:C:H42	2.16	0.44
20:F2:190:ARG:HB2	20:F2:202:ILE:HG23	1.99	0.44
28:I2:135:PHE:HE2	28:I2:137:TYR:CZ	2.35	0.44
34:L1:96:ASN:HD21	34:L1:99:LEU:HD12	1.83	0.44
42:O3:136:PRO:HG3	42:O3:139:SER:HB2	1.99	0.44
49:S2:10:ASP:HB3	49:S2:13:LYS:HB2	2.00	0.44
60:a2:114:GLN:NE2	60:a2:115:LYS:HG3	2.33	0.44
64:e2:27:LYS:HD2	64:e2:70:LYS:NZ	2.33	0.44
74:r2:80:ILE:HG13	74:r2:81:THR:HG23	1.98	0.44
75:s2:72:LEU:HD12	75:s2:112:LEU:HD11	1.99	0.44
80:x2:77:LYS:HB3	83:m2:1300:G:H5'	1.98	0.44
83:m2:116:OMU:O5'	83:m2:116:OMU:H6	2.18	0.44
83:m2:331:G:H3'	83:m2:332:G:H8	1.81	0.44
83:m2:555:A:H2'	83:m2:556:A:C8	2.53	0.44
83:m2:651:U:H2'	83:m2:652:A:H8	1.82	0.44
83:m2:1398:A:N1	83:m2:1451:G:C6	2.85	0.44
83:m2:1408:G:H2'	83:m2:1409:U:C6	2.53	0.44
83:m2:1630:C:H2'	83:m2:1631:C:C6	2.53	0.44
1:A1:116:ARG:HB2	1:A1:136:LEU:HB3	2.00	0.44
1:A1:247:THR:HG21	1:A1:251:GLU:OE1	2.18	0.44
2:A2:106:A:H2'	2:A2:107:G:O4'	2.18	0.44
2:A2:2061:G:H1'	2:A2:2087:A:N6	2.32	0.44
2:A2:3588:U:H2'	2:A2:3589:G:H8	1.82	0.44
10:C1:147:GLU:HG2	10:C1:187:VAL:HG22	2.00	0.44
11:C2:5:U:H2'	11:C2:6:C:C6	2.52	0.44
12:C3:88:LEU:O	12:C3:89:ILE:HD13	2.18	0.44
13:D1:91:LEU:HD13	13:D1:135:ILE:HA	1.99	0.44
17:E2:220:ILE:HG13	17:E2:278:THR:HG23	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:F1:93:THR:HG22	61:b2:116:LEU:HD13	2.00	0.44
23:G2:233:PRO:HA	23:G2:236:MET:HG3	1.99	0.44
34:L1:89:ALA:O	34:L1:93:LEU:HG	2.17	0.44
40:N3:4:MET:HG2	40:N3:5:HIS:CE1	2.52	0.44
43:P2:89:ARG:HB2	43:P2:95:PHE:CE2	2.53	0.44
59:Z2:36:ARG:HB2	59:Z2:80:ASN:HA	1.99	0.44
7:n2:32:U:H2'	7:n2:33:U:C6	2.53	0.44
74:r2:15:PRO:HA	74:r2:39:ARG:NH2	2.32	0.44
78:v2:71:LEU:HD23	78:v2:71:LEU:HA	1.89	0.44
83:m2:25:A:HO2'	83:m2:26:U:H6	1.64	0.44
83:m2:404:C:C2	83:m2:405:G:C8	3.05	0.44
83:m2:850:U:H2'	83:m2:851:A:H8	1.82	0.44
83:m2:1328:U:H4'	83:m2:1329:G:H5''	1.99	0.44
83:m2:1364:U:H5''	83:m2:1365:C:H5	1.83	0.44
83:m2:1483:G:H2'	83:m2:1484:C:C6	2.53	0.44
83:m2:1484:C:H2'	83:m2:1485:A:O4'	2.18	0.44
83:m2:1653:A:H2'	83:m2:1654:G:H8	1.83	0.44
83:m2:1744:C:H2'	83:m2:1745:G:O4'	2.17	0.44
2:A2:700:C:H2'	2:A2:701:C:C6	2.52	0.44
2:A2:760:C:H2'	2:A2:761:U:C6	2.53	0.44
2:A2:1275:A:H2'	2:A2:1276:C:O4'	2.18	0.44
2:A2:1685:G:O6	2:A2:1698:A:H2	2.01	0.44
2:A2:2360:G:H2'	2:A2:2361:G:C8	2.53	0.44
2:A2:3958:OMU:OP2	32:K2:158:THR:HG23	2.18	0.44
2:A2:4042:A:H2'	2:A2:4043:G:O4'	2.17	0.44
2:A2:4236:A:H2'	2:A2:4237:U:O4'	2.17	0.44
16:E1:10:ASN:O	16:E1:14:GLU:HB2	2.18	0.44
23:G2:81:HIS:HA	23:G2:92:LEU:HD13	1.99	0.44
29:I3:259:TRP:CD1	29:I3:266:ILE:HG12	2.53	0.44
29:I3:297:THR:HA	29:I3:310:TRP:O	2.17	0.44
40:N3:91:LEU:HB3	40:N3:122:ILE:HG12	2.00	0.44
42:O3:128:ARG:HH21	72:p2:72:ALA:HB3	1.83	0.44
45:Q2:33:ASN:OD1	45:Q2:35:LYS:HB3	2.17	0.44
70:k2:17:LEU:HD21	70:k2:19:LYS:HE3	2.00	0.44
71:o2:50:ASN:HB3	71:o2:53:ARG:HG2	1.99	0.44
77:u2:12:ARG:HG2	77:u2:16:GLY:O	2.17	0.44
83:m2:456:U:H2'	83:m2:457:A:C8	2.52	0.44
83:m2:847:G:H2'	83:m2:848:G:C8	2.52	0.44
83:m2:1207:C:H2'	83:m2:1208:G:H8	1.82	0.44
83:m2:1220:C:H2'	83:m2:1221:C:H6	1.83	0.44
83:m2:1350:G:N7	83:m2:1351:G:C8	2.81	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A1:251:GLU:HA	37:M2:2:LYS:HA	2.00	0.44
2:A2:184:U:H5''	2:A2:254:G:C2	2.53	0.44
2:A2:927:C:H2'	2:A2:928:G:H8	1.78	0.44
2:A2:1360:A:HO2'	2:A2:2569:C:H6	1.66	0.44
2:A2:1363:G:H1	2:A2:1391:U:H3	1.65	0.44
2:A2:1551:A:H2'	2:A2:1552:G:C8	2.52	0.44
2:A2:3601:A:H2'	2:A2:3602:G:C8	2.53	0.44
2:A2:3625:G:H2'	2:A2:3626:G:H8	1.78	0.44
2:A2:3932:A:N1	23:G2:29:ASP:HA	2.32	0.44
2:A2:4414:A:H2'	2:A2:4415:U:O4'	2.18	0.44
6:B3:86:GLY:C	83:m2:1608:G:H5''	2.43	0.44
28:I2:14:HIS:HD2	28:I2:123:ALA:HB3	1.82	0.44
29:I3:9:GLY:HA3	29:I3:52:TYR:HB2	1.98	0.44
29:I3:52:TYR:OH	29:I3:297:THR:HG21	2.18	0.44
37:M2:43:ARG:HD2	37:M2:43:ARG:HA	1.66	0.44
39:N2:27:LEU:HA	39:N2:30:TYR:HD2	1.83	0.44
51:T2:135:ARG:HA	51:T2:135:ARG:HD2	1.84	0.44
71:o2:77:ILE:HD13	71:o2:99:ILE:HB	1.99	0.44
71:o2:112:ILE:HD13	83:m2:1352:U:H4'	1.99	0.44
72:p2:144:LYS:HD2	72:p2:146:ARG:NH1	2.30	0.44
73:q2:99:ILE:HG23	73:q2:173:ARG:HH21	1.83	0.44
74:r2:33:THR:O	83:m2:120:U:H1'	2.18	0.44
77:u2:51:GLY:HA3	83:m2:447:A:H5''	1.99	0.44
77:u2:155:ASN:HB3	79:w2:21:LYS:HE3	1.99	0.44
77:u2:188:TYR:HB3	83:m2:223:A:H4'	1.98	0.44
77:u2:192:GLY:HA3	79:w2:19:ASN:OD1	2.18	0.44
80:x2:57:LEU:HD13	80:x2:89:MET:HE3	1.99	0.44
83:m2:571:A:H2'	83:m2:572:C:O4'	2.17	0.44
83:m2:658:G:N2	83:m2:665:C:H5''	2.32	0.44
83:m2:1279:C:H2'	83:m2:1280:A:C8	2.53	0.44
83:m2:1322:G:H3'	83:m2:1323:G:C8	2.53	0.44
2:A2:952:C:H2'	2:A2:953:G:C8	2.52	0.44
2:A2:1330:2MG:H2'	2:A2:1331:A:N7	2.33	0.44
2:A2:2363:G:H2'	2:A2:2364:G:H8	1.82	0.44
2:A2:3423:C:C2'	2:A2:3424:U:H5'	2.47	0.44
2:A2:4515:G:C5	22:G1:56:GLN:HG2	2.51	0.44
6:B3:43:LYS:HB3	83:m2:1541:U:OP1	2.18	0.44
7:Bz:37:A:C4	7:Bz:38:A:C8	3.05	0.44
10:C1:128:MET:SD	10:C1:157:SER:HB3	2.58	0.44
17:E2:43:LEU:HG	17:E2:196:TRP:CH2	2.52	0.44
17:E2:283:LYS:HD3	17:E2:285:TYR:HE1	1.82	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:F2:242:PRO:HG3	20:F2:248:ARG:HD2	1.99	0.44
32:K2:64:SER:HB3	32:K2:92:ILE:HD11	1.99	0.44
42:O3:140:THR:HB	83:m2:1048:U:H1'	2.00	0.44
58:Y2:77:PHE:HD2	58:Y2:88:LEU:HD11	1.83	0.44
73:q2:163:PRO:HA	73:q2:166:TYR:CZ	2.52	0.44
75:s2:22:LYS:HB3	75:s2:22:LYS:HE2	1.69	0.44
78:v2:29:MET:HE3	78:v2:29:MET:HB3	1.82	0.44
83:m2:126:G:H4'	83:m2:127:C:OP2	2.15	0.44
83:m2:474:C:H4'	83:m2:476:G:OP1	2.18	0.44
83:m2:1432:C:H2'	83:m2:1433:G:C8	2.52	0.44
2:A2:221:C:H2'	2:A2:222:C:C6	2.52	0.43
2:A2:453:G:N2	2:A2:1107:G:H22	2.16	0.43
2:A2:656:G:H2'	2:A2:657:A:H8	1.82	0.43
2:A2:1146:C:OP2	58:Y2:30:LYS:HD3	2.18	0.43
2:A2:1590:A:H2'	13:D1:22:PHE:CZ	2.53	0.43
2:A2:1867:G:H2'	2:A2:1868:C:O4'	2.18	0.43
2:A2:2119:OMG:HM23	2:A2:2119:OMG:H1'	1.69	0.43
2:A2:2513:G:H2'	2:A2:2514:G:C5	2.53	0.43
2:A2:3830:A:H2'	2:A2:3831:G:C8	2.52	0.43
23:G2:67:ALA:HB1	39:N2:31:MET:HE2	2.00	0.43
28:I2:157:GLU:HG3	28:I2:161:LYS:NZ	2.33	0.43
31:J3:200:ARG:O	36:L3:98:LEU:HB3	2.18	0.43
31:J3:227:ARG:HD2	83:m2:1156:U:O4	2.18	0.43
44:P3:95:PRO:HG3	44:P3:130:PHE:CD1	2.53	0.43
50:S3:33:MET:HE1	50:S3:73:LEU:HD11	1.99	0.43
61:b2:4:ILE:HD11	61:b2:53:SER:HB3	2.00	0.43
65:f2:44:TRP:CZ3	65:f2:45:ARG:HD2	2.53	0.43
67:h2:11:ARG:NH2	83:m2:1846:U:P	2.73	0.43
73:q2:40:ARG:HB2	73:q2:47:GLU:HG3	1.99	0.43
80:x2:20:VAL:HG21	80:x2:36:LEU:HD21	1.99	0.43
83:m2:126:G:N7	83:m2:181:A:H1'	2.33	0.43
83:m2:129:C:H5'	83:m2:216:U:C4	2.53	0.43
2:A2:2259:C:H3'	2:A2:2261:G:C8	2.51	0.43
2:A2:2614:G:H2'	2:A2:2615:C:H6	1.82	0.43
2:A2:3426:U:H4'	7:Bv:13:C:H4'	2.00	0.43
2:A2:3685:G:P	2:A2:3685:G:H8	2.41	0.43
2:A2:4145:U:H2'	2:A2:4146:OMG:H8	1.83	0.43
2:A2:4560:G:H2'	2:A2:4561:G:C8	2.53	0.43
3:A3:34:LYS:O	3:A3:103:LEU:HD13	2.17	0.43
7:Bz:22:G:H2'	7:Bz:23:A:H8	1.82	0.43
12:C3:19:ARG:HG3	12:C3:92:HIS:ND1	2.32	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:F2:292:ILE:O	20:F2:298:ILE:HD12	2.17	0.43
30:J2:131:ARG:HG3	30:J2:137:ASN:HB2	1.99	0.43
32:K2:50:ARG:HA	32:K2:53:MET:SD	2.58	0.43
33:K3:3:LEU:HD22	33:K3:111:LEU:HD11	2.00	0.43
35:L2:11:ALA:O	35:L2:15:LEU:HB2	2.18	0.43
48:R3:92:LEU:HD11	48:R3:109:TYR:CZ	2.54	0.43
59:Z2:92:LEU:HD23	59:Z2:92:LEU:HA	1.86	0.43
63:d2:52:LYS:HG2	63:d2:55:ARG:NH2	2.33	0.43
67:h2:4:LYS:HE3	67:h2:5:TRP:CZ3	2.53	0.43
67:h2:15:ARG:HD3	83:m2:1185:A:P	2.58	0.43
7:n2:74:C:H5'	7:n2:75:C:OP2	2.18	0.43
80:x2:124:LYS:HD3	83:m2:1241:U:H4'	1.99	0.43
83:m2:503:C:H2'	83:m2:504:C:H5''	2.00	0.43
83:m2:561:G:O2'	83:m2:562:A:H8	2.01	0.43
83:m2:630:A:N6	83:m2:1334:A:C2	2.86	0.43
83:m2:1382:C:H2'	83:m2:1383:G:C8	2.52	0.43
83:m2:1392:U:H2'	83:m2:1393:C:C6	2.53	0.43
1:A1:184:ILE:HD12	1:A1:189:ILE:HB	2.00	0.43
2:A2:28:C:P	25:H1:193:ARG:HH22	2.41	0.43
2:A2:941:C:H2'	2:A2:942:A:H8	1.83	0.43
2:A2:1441:C:OP2	14:D2:9:ARG:HD2	2.17	0.43
2:A2:1744:A:N6	2:A2:1841:G:H2'	2.34	0.43
2:A2:1911:G:H2'	2:A2:1912:A:C8	2.53	0.43
2:A2:2321:G:H2'	2:A2:2322:G:H8	1.84	0.43
2:A2:3971:C:H2'	2:A2:3972:G:H8	1.83	0.43
7:Bv:20:U:H2'	7:Bv:21:A:H4'	2.00	0.43
19:F1:80:GLU:HB3	19:F1:110:LEU:HD12	1.99	0.43
36:L3:12:THR:HG23	83:m2:522:A:H5''	1.99	0.43
38:M3:117:GLU:HA	38:M3:120:ALA:HB3	2.00	0.43
40:N3:52:VAL:HG22	40:N3:55:ARG:NH2	2.33	0.43
70:k2:47:LYS:HB2	70:k2:102:TYR:CE2	2.53	0.43
7:n2:28:G:H2'	7:n2:29:G:H8	1.82	0.43
72:p2:149:GLN:HG3	83:m2:1125:C:O3'	2.17	0.43
74:r2:160:ILE:HD12	74:r2:162:ILE:HD11	1.99	0.43
83:m2:814:A:C4	83:m2:815:A:C8	3.06	0.43
83:m2:1330:OMG:HM23	83:m2:1330:OMG:H1'	1.88	0.43
1:A1:179:ARG:HD2	1:A1:235:LEU:HD12	2.00	0.43
2:A2:1474:C:H2'	2:A2:1475:C:H6	1.83	0.43
2:A2:1677:C:H2'	2:A2:1678:U:C6	2.53	0.43
2:A2:2448:G:H2'	2:A2:2449:G:N2	2.33	0.43
2:A2:3359:G:H2'	2:A2:3360:U:C6	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:3430:A:H8	2:A2:3430:A:OP1	2.01	0.43
2:A2:3602:G:H2'	2:A2:3603:A:H8	1.83	0.43
2:A2:3717:U:H2'	2:A2:3718:U:H6	1.81	0.43
2:A2:3744:G:H1	2:A2:3765:G:N2	2.17	0.43
2:A2:4652:U:H4'	2:A2:4653:A:H5'	1.99	0.43
6:B3:21:PHE:HA	6:B3:24:LYS:HZ3	1.83	0.43
7:Bz:19:G:H4'	7:Bz:20:U:OP2	2.19	0.43
10:C1:62:GLY:HA2	10:C1:66:GLU:OE2	2.19	0.43
20:F2:292:ILE:HG23	32:K2:128:LEU:HD21	2.01	0.43
22:G1:47:ARG:HG2	37:M2:73:LEU:HD22	2.00	0.43
33:K3:58:LYS:HA	33:K3:107:SER:HB2	1.99	0.43
47:R2:80:PRO:HA	47:R2:98:PHE:HA	2.00	0.43
49:S2:59:ARG:HB2	49:S2:103:LYS:HD2	2.00	0.43
52:T3:48:THR:HG22	52:T3:48:THR:O	2.18	0.43
75:s2:88:MET:O	75:s2:91:ARG:HG2	2.19	0.43
80:x2:53:GLN:HG3	80:x2:83:MET:SD	2.59	0.43
83:m2:572:C:H3'	83:m2:573:U:H6	1.84	0.43
83:m2:919:U:H2'	83:m2:920:U:C6	2.53	0.43
83:m2:1332:G:H4'	83:m2:1333:C:H3'	2.00	0.43
2:A2:99:A:H5''	25:H1:184:ILE:HD13	1.99	0.43
2:A2:949:G:H2'	2:A2:950:A:C8	2.54	0.43
2:A2:2264:C:H5'	11:C2:108:A:H4'	1.99	0.43
2:A2:3382:A:H2'	2:A2:3383:A:C8	2.53	0.43
2:A2:4369:A:O4'	17:E2:18:PRO:HG2	2.18	0.43
40:N3:57:SER:HB3	50:S3:54:VAL:HG13	2.01	0.43
40:N3:71:ILE:HD12	83:m2:1020:U:H5''	2.01	0.43
44:P3:69:LEU:HD11	44:P3:129:PHE:HB2	2.00	0.43
75:s2:143:PRO:O	75:s2:147:VAL:HG22	2.18	0.43
75:s2:173:LEU:O	75:s2:177:LEU:HG	2.18	0.43
77:u2:38:ILE:HD11	77:u2:81:VAL:HG23	1.99	0.43
83:m2:17:C:H2'	83:m2:18:C:C6	2.53	0.43
83:m2:931:G:N2	83:m2:1106:G:H4'	2.33	0.43
83:m2:1375:C:H2'	83:m2:1376:C:H6	1.82	0.43
2:A2:434:A:H2'	2:A2:435:A:O4'	2.19	0.43
2:A2:844:C:H5'	20:F2:336:ARG:NH1	2.33	0.43
2:A2:1884:G:H5''	32:K2:12:LYS:HG3	2.01	0.43
2:A2:2294:C:H2'	2:A2:2295:C:H6	1.83	0.43
2:A2:3402:A:H5'	14:D2:243:THR:HG23	2.00	0.43
3:A3:132:ARG:HD3	83:m2:1625:A:N7	2.34	0.43
7:Bz:22:G:N7	7:Bz:46:G:O6	2.51	0.43
14:D2:2:GLY:HA2	14:D2:207:VAL:HG23	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:I3:147:HIS:HE1	29:I3:168:CYS:HA	1.83	0.43
48:R3:70:PRO:HA	48:R3:84:ALA:HB1	2.00	0.43
60:a2:96:LEU:HD23	60:a2:96:LEU:HA	1.88	0.43
71:o2:122:LEU:HB2	71:o2:142:LEU:HD21	2.00	0.43
75:s2:19:LEU:HG	75:s2:20:PHE:HD1	1.80	0.43
79:w2:20:LYS:O	79:w2:22:ARG:HD2	2.18	0.43
79:w2:27:GLU:O	79:w2:31:GLU:HG3	2.19	0.43
83:m2:1104:G:H2'	83:m2:1105:C:C6	2.53	0.43
83:m2:1105:C:H2'	83:m2:1106:G:H8	1.82	0.43
83:m2:1362:U:H3'	83:m2:1363:G:C8	2.53	0.43
83:m2:1437:C:H2'	83:m2:1438:C:O4'	2.19	0.43
1:A1:219:VAL:HG23	1:A1:223:PHE:CG	2.54	0.43
2:A2:656:G:H2'	2:A2:657:A:C8	2.54	0.43
2:A2:1085:C:H5	2:A2:1910:G:N2	2.16	0.43
2:A2:1316:A:H4'	2:A2:1317:G:H5'	2.01	0.43
2:A2:1404:U:H5''	2:A2:4179:G:OP1	2.19	0.43
2:A2:1766:A:H1'	2:A2:4346:G:N2	2.34	0.43
2:A2:1766:A:H3'	2:A2:1767:G:H8	1.84	0.43
2:A2:2531:G:H2'	2:A2:2532:G:C8	2.54	0.43
2:A2:3324:C:H5'	14:D2:8:GLN:O	2.19	0.43
2:A2:3389:A:H2'	2:A2:3390:U:O4'	2.19	0.43
2:A2:3488:U:H2'	2:A2:3489:C:H6	1.84	0.43
2:A2:3933:A:O2'	2:A2:3934:A:H2'	2.18	0.43
2:A2:4531:C:H2'	2:A2:4532:U:C6	2.53	0.43
10:C1:44:GLU:CG	10:C1:58:ASP:HB2	2.48	0.43
11:C2:71:A:H3'	49:S2:50:ARG:HG3	2.01	0.43
20:F2:60:HIS:NE2	20:F2:100:ARG:HD3	2.34	0.43
31:J3:164:PRO:HG2	31:J3:248:TYR:HD2	1.84	0.43
31:J3:185:THR:HG22	31:J3:194:ARG:HG3	2.01	0.43
32:K2:70:MET:HE3	32:K2:70:MET:HB3	1.82	0.43
42:O3:31:CYS:HB2	42:O3:44:VAL:HG22	2.00	0.43
57:X2:87:ARG:HB3	57:X2:107:THR:OG1	2.19	0.43
60:a2:7:TYR:HD2	60:a2:12:SER:HA	1.83	0.43
63:d2:15:THR:HG23	63:d2:16:HIS:ND1	2.34	0.43
68:i2:32:SER:C	68:i2:34:TYR:H	2.27	0.43
72:p2:28:LYS:HD2	72:p2:50:THR:HA	2.00	0.43
83:m2:677:U:H2'	83:m2:678:C:H6	1.83	0.43
83:m2:951:G:H2'	83:m2:952:C:H6	1.84	0.43
83:m2:1136:G:H2'	83:m2:1137:C:H6	1.81	0.43
83:m2:1232:C:H2'	83:m2:1233:C:C6	2.54	0.43
83:m2:1388:A:H2'	83:m2:1389:G:O4'	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
83:m2:1522:G:H5''	83:m2:1523:C:OP2	2.19	0.43
83:m2:1803:A:H2'	83:m2:1804:C:C6	2.54	0.43
2:A2:318:A:H2'	2:A2:319:A:H8	1.82	0.43
2:A2:425:U:H2'	2:A2:426:A:H8	1.84	0.43
2:A2:1563:G:H2'	2:A2:1564:C:O4'	2.19	0.43
2:A2:1881:G:H2'	2:A2:1882:U:C6	2.54	0.43
2:A2:3266:A:H2'	2:A2:3267:A:C8	2.53	0.43
2:A2:4188:OMC:H2'	2:A2:4189:C:C6	2.53	0.43
10:C1:114:ILE:HB	10:C1:124:ARG:HB2	2.00	0.43
17:E2:41:VAL:HG22	17:E2:187:GLY:O	2.18	0.43
18:E3:105:PHE:CD1	18:E3:121:LYS:HB3	2.53	0.43
20:F2:140:LYS:HD2	20:F2:140:LYS:HA	1.89	0.43
20:F2:242:PRO:CG	20:F2:248:ARG:HD2	2.49	0.43
36:L3:134:HIS:ND1	36:L3:163:SER:HB2	2.33	0.43
60:a2:40:LYS:HD3	60:a2:52:ARG:NH2	2.34	0.43
72:p2:33:VAL:HG22	72:p2:96:CYS:SG	2.58	0.43
75:s2:40:ALA:HB3	75:s2:67:PRO:HA	2.00	0.43
77:u2:117:TYR:CD1	77:u2:152:ARG:HB3	2.53	0.43
80:x2:56:LEU:HD11	80:x2:78:THR:HB	2.01	0.43
83:m2:72:C:OP1	83:m2:72:C:H4'	2.18	0.43
83:m2:1027:U:H2'	83:m2:1028:C:O4'	2.19	0.43
83:m2:1094:G:H2'	83:m2:1095:A:C8	2.53	0.43
2:A2:207:G:H2'	2:A2:208:A:C8	2.54	0.43
2:A2:487:C:H2'	2:A2:488:G:H8	1.84	0.43
2:A2:690:C:H2'	2:A2:691:G:O4'	2.19	0.43
2:A2:1078:G:H3'	2:A2:1079:A:H8	1.83	0.43
2:A2:1285:C:H2'	2:A2:1286:U:C6	2.54	0.43
2:A2:1561:G:H2'	2:A2:1562:G:C8	2.54	0.43
2:A2:2652:G:H2'	2:A2:2653:G:H8	1.83	0.43
2:A2:3522:C:H2'	2:A2:3523:A:O4'	2.18	0.43
2:A2:3942:U:H3'	68:i2:9:ARG:O	2.18	0.43
2:A2:3952:U:H2'	2:A2:3953:U:C6	2.53	0.43
2:A2:4203:U:H2'	2:A2:4204:U:C6	2.54	0.43
2:A2:4218:U:H2'	2:A2:4219:G:O4'	2.19	0.43
2:A2:4584:G:H22	26:H2:257:GLN:HB2	1.83	0.43
11:C2:53:G:C2	65:f2:35:ILE:HD11	2.53	0.43
13:D1:140:THR:HG23	13:D1:141:LYS:O	2.19	0.43
17:E2:224:LYS:HG2	17:E2:340:THR:CG2	2.49	0.43
26:H2:155:THR:HG21	26:H2:203:ALA:HB1	2.00	0.43
29:I3:257:LYS:HB3	29:I3:259:TRP:NE1	2.33	0.43
33:K3:23:LYS:HD3	33:K3:42:GLY:H	1.84	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:V2:99:HIS:CE1	55:V2:102:LEU:HD13	2.54	0.43
73:q2:9:ARG:HH22	83:m2:1554:G:H4'	1.84	0.43
75:s2:78:MET:HB3	75:s2:159:ARG:HH11	1.83	0.43
76:t2:121:THR:HG23	76:t2:124:ALA:H	1.83	0.43
83:m2:21:U:H2'	83:m2:22:A:C8	2.54	0.43
83:m2:375:G:H2'	83:m2:376:G:H8	1.84	0.43
83:m2:1353:G:N3	83:m2:1381:A:N1	2.67	0.43
83:m2:1858:C:H2'	83:m2:1859:G:C8	2.53	0.43
2:A2:2:G:H2'	2:A2:3:C:C6	2.54	0.43
2:A2:9:C:OP1	25:H1:40:PRO:HD3	2.19	0.43
2:A2:1125:G:H2'	2:A2:1126:A:O4'	2.18	0.43
2:A2:3608:A:H2'	2:A2:3609:G:C8	2.54	0.43
2:A2:3933:A:O2'	2:A2:3934:A:C8	2.72	0.43
3:A3:13:LEU:HB3	3:A3:20:ILE:HG12	2.00	0.43
11:C2:75:OMG:HM23	11:C2:75:OMG:H1'	1.60	0.43
13:D1:61:SER:HA	13:D1:126:VAL:HG12	2.01	0.43
14:D2:196:TRP:HB3	14:D2:197:PRO:HD3	2.00	0.43
29:I3:60:ARG:HD2	81:y2:97:GLN:HE21	1.83	0.43
30:J2:60:PHE:CE1	30:J2:82:ARG:HB2	2.54	0.43
64:e2:21:LYS:HB2	64:e2:37:ARG:O	2.19	0.43
66:g2:88:LYS:HE3	66:g2:89:TYR:CZ	2.54	0.43
68:i2:53:LYS:HD3	7:n2:75:C:O2	2.19	0.43
70:k2:39:ARG:HH11	70:k2:105:ASP:CG	2.26	0.43
76:t2:63:PHE:HA	76:t2:95:ILE:O	2.18	0.43
79:w2:83:GLN:HE22	83:m2:394:A:N6	2.17	0.43
83:m2:682:G:H2'	83:m2:683:U:C6	2.54	0.43
83:m2:1034:C:H2'	83:m2:1035:G:O4'	2.19	0.43
2:A2:82:U:H2'	2:A2:83:C:O4'	2.19	0.42
2:A2:2379:G:H2'	2:A2:2380:U:H6	1.83	0.42
2:A2:3807:C:H2'	2:A2:3808:G:O4'	2.19	0.42
4:B1:58:PRO:HD2	4:B1:61:ILE:HD12	2.01	0.42
5:B2:108:G:P	23:G2:273:LEU:HD22	2.59	0.42
12:C3:55:ARG:HG2	12:C3:87:ARG:NE	2.34	0.42
14:D2:36:GLU:OE2	14:D2:163:ARG:HD2	2.19	0.42
25:H1:120:TRP:CZ2	25:H1:123:GLU:HB2	2.52	0.42
27:H3:14:PHE:O	83:m2:1663:A:H8	2.02	0.42
29:I3:129:ILE:HD11	29:I3:151:VAL:HG21	2.00	0.42
32:K2:46:VAL:O	32:K2:50:ARG:HG3	2.19	0.42
33:K3:32:MET:HB3	33:K3:65:GLN:HE22	1.84	0.42
34:L1:64:SER:HB3	34:L1:107:TYR:CD2	2.54	0.42
35:L2:66:ASN:HB3	35:L2:70:ARG:NH1	2.34	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:M3:120:ALA:HA	38:M3:123:VAL:HG12	2.00	0.42
46:Q3:116:LYS:HG2	83:m2:161:U:H5'	2.00	0.42
7:n2:14:A:H2'	7:n2:15:G:C8	2.54	0.42
74:r2:11:ARG:NH2	83:m2:497:U:H4'	2.33	0.42
75:s2:50:PRO:HB3	75:s2:69:VAL:HG12	2.00	0.42
77:u2:134:GLU:O	77:u2:138:ASN:HB2	2.20	0.42
78:v2:3:MET:HB3	78:v2:44:HIS:CE1	2.53	0.42
83:m2:11:A:N1	83:m2:1202:A:H2	2.17	0.42
83:m2:616:C:H5''	83:m2:617:C:C5	2.54	0.42
83:m2:898:U:H2'	83:m2:899:U:C6	2.54	0.42
83:m2:973:G:O2'	83:m2:974:A:H8	2.00	0.42
83:m2:1166:G:O2'	83:m2:1167:G:H5'	2.19	0.42
2:A2:184:U:H5''	2:A2:254:G:N1	2.34	0.42
2:A2:1115:C:H2'	2:A2:1117:A:C5	2.53	0.42
2:A2:1140:A2M:H2'	2:A2:1141:C:C6	2.54	0.42
2:A2:1318:C:H2'	2:A2:1319:G:C8	2.54	0.42
2:A2:2224:C:H5	2:A2:2226:G:H1	1.67	0.42
2:A2:2655:U:H2'	2:A2:2656:G:C8	2.54	0.42
2:A2:3906:G:H2'	2:A2:3906:G:N3	2.34	0.42
2:A2:4160:C:N3	2:A2:4164:U:H5	2.17	0.42
2:A2:4564:G:H2'	2:A2:4565:C:C6	2.54	0.42
4:B1:99:ALA:HB1	4:B1:136:LEU:HD11	2.00	0.42
11:C2:66:A:H2'	11:C2:67:U:C6	2.54	0.42
11:C2:67:U:H2'	11:C2:68:G:C8	2.54	0.42
20:F2:8:ILE:HG22	70:k2:71:ARG:HE	1.84	0.42
24:G3:47:LYS:HE2	75:s2:145:ARG:NH1	2.34	0.42
29:I3:181:ASN:HB3	29:I3:183:LYS:HG2	2.01	0.42
33:K3:54:GLY:HA2	33:K3:63:MET:HE2	2.02	0.42
39:N2:19:PHE:CD2	39:N2:20:ARG:HG3	2.54	0.42
43:P2:36:ASN:O	43:P2:37:LEU:HD23	2.20	0.42
71:o2:112:ILE:HD12	71:o2:112:ILE:H	1.84	0.42
72:p2:189:ILE:HG13	72:p2:190:PRO:CD	2.46	0.42
73:q2:110:LEU:HA	73:q2:177:LEU:HD21	2.00	0.42
83:m2:1175:A:H2'	83:m2:1176:U:C6	2.54	0.42
83:m2:1436:C:H4'	83:m2:1437:C:C6	2.54	0.42
83:m2:1440:A:H2'	83:m2:1441:A:H8	1.84	0.42
2:A2:1475:C:H2'	2:A2:1476:C:H6	1.85	0.42
2:A2:2287:C:H2'	2:A2:2288:C:C6	2.55	0.42
2:A2:2619:A:H2'	2:A2:2620:U:C6	2.54	0.42
2:A2:4082:G:H1'	13:D1:158:LYS:HD2	2.01	0.42
2:A2:4188:OMC:HM23	2:A2:4188:OMC:H1'	1.91	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:4285:G:O3'	2:A2:4286:U:H3'	2.18	0.42
29:I3:69:VAL:HG23	29:I3:80:SER:HB3	2.00	0.42
29:I3:273:GLU:H	29:I3:273:GLU:CD	2.28	0.42
33:K3:71:GLY:H	33:K3:98:ARG:NH1	2.18	0.42
36:L3:44:TRP:HA	36:L3:47:LYS:HG2	2.00	0.42
61:b2:95:LEU:HB3	61:b2:99:GLU:HG3	2.01	0.42
75:s2:41:VAL:HG23	75:s2:41:VAL:O	2.19	0.42
75:s2:136:ARG:O	75:s2:203:ASN:HB3	2.20	0.42
82:z2:26:ASN:CA	82:z2:58:MET:HE2	2.48	0.42
83:m2:184:G:H2'	83:m2:185:G:H8	1.84	0.42
83:m2:1458:G:H2'	83:m2:1459:U:C6	2.54	0.42
83:m2:1718:C:H2'	83:m2:1719:C:H6	1.84	0.42
2:A2:1070:C:H2'	2:A2:1071:G:O4'	2.20	0.42
2:A2:1559:U:C5	2:A2:3907:A:H5'	2.53	0.42
2:A2:2214:G:H2'	2:A2:2216:G:OP2	2.19	0.42
2:A2:3525:OMC:H1'	28:I2:68:ARG:NH2	2.34	0.42
3:A3:12:ILE:HG21	16:E1:119:TYR:CE2	2.54	0.42
12:C3:66:ARG:HB3	27:H3:40:ARG:NH1	2.35	0.42
17:E2:394:LYS:HE2	17:E2:394:LYS:HB3	1.76	0.42
35:L2:105:LEU:HD22	35:L2:135:LYS:HG3	2.01	0.42
36:L3:16:PRO:HG3	36:L3:24:ARG:NH1	2.34	0.42
37:M2:15:ARG:HD2	37:M2:25:PRO:HG2	2.00	0.42
64:e2:24:LYS:HA	64:e2:67:LYS:O	2.19	0.42
7:n2:8:U:H5'	7:n2:49:C:OP2	2.19	0.42
73:q2:66:ILE:O	73:q2:70:THR:HG23	2.20	0.42
78:v2:47:LYS:O	78:v2:50:GLN:HG2	2.20	0.42
82:z2:44:LYS:HB3	82:z2:44:LYS:HE3	1.79	0.42
83:m2:947:U:H2'	83:m2:948:U:H6	1.85	0.42
83:m2:1833:A:H1'	83:m2:1854:C:H5'	2.00	0.42
2:A2:1054:G:H2'	2:A2:1055:G:H8	1.84	0.42
2:A2:1281:C:H2'	2:A2:1282:C:H6	1.84	0.42
2:A2:2294:C:H2'	2:A2:2295:C:C6	2.55	0.42
2:A2:3383:A:H8	2:A2:3383:A:OP2	2.02	0.42
2:A2:3930:C:O2'	2:A2:3933:A:H1'	2.19	0.42
2:A2:4044:OMG:C4	2:A2:4099:5MC:HM52	2.54	0.42
2:A2:4160:C:H5'	43:P2:43:LYS:HG2	2.01	0.42
2:A2:4411:C:H2'	2:A2:4412:G:C8	2.54	0.42
6:B3:75:MET:HA	6:B3:78:ILE:HD12	2.01	0.42
12:C3:36:CYS:SG	12:C3:53:PRO:HB3	2.60	0.42
16:E1:166:PHE:CE2	16:E1:172:GLY:HA3	2.55	0.42
17:E2:128:LYS:HB3	17:E2:128:LYS:HE3	1.84	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:E3:123:VAL:HG12	18:E3:124:LYS:HG3	2.02	0.42
30:J2:112:LEU:HD23	30:J2:112:LEU:HA	1.88	0.42
34:L1:196:LYS:HA	34:L1:196:LYS:HD3	1.82	0.42
40:N3:80:LEU:HD23	40:N3:80:LEU:H	1.85	0.42
43:P2:111:GLU:HA	43:P2:131:ARG:HD2	2.01	0.42
44:P3:57:ARG:HD2	50:S3:26:GLN:NE2	2.35	0.42
60:a2:63:VAL:HG12	60:a2:66:ARG:HH12	1.84	0.42
64:e2:61:PRO:HA	64:e2:62:PRO:HD3	1.93	0.42
69:j2:29:ILE:HG23	69:j2:69:TRP:HB3	2.00	0.42
76:t2:100:ILE:HD11	76:t2:125:VAL:HG11	2.01	0.42
78:v2:72:THR:HG23	78:v2:75:GLY:H	1.84	0.42
81:y2:125:ARG:HE	81:y2:125:ARG:HB2	1.61	0.42
83:m2:29:G:C6	83:m2:648:G:C6	3.07	0.42
83:m2:961:G:O2'	83:m2:1067:G:H4'	2.19	0.42
83:m2:1056:G:C6	83:m2:1067:G:C6	3.08	0.42
83:m2:1456:A:C2	83:m2:1478:A:H1'	2.55	0.42
2:A2:261:G:H2'	2:A2:262:G:C8	2.53	0.42
2:A2:272:U:H2'	2:A2:273:U:C6	2.55	0.42
2:A2:1482:A:H4'	2:A2:1498:G:H22	1.85	0.42
2:A2:2240:U:H2'	2:A2:2241:G:C8	2.54	0.42
2:A2:2631:G:C8	69:j2:16:THR:HG22	2.55	0.42
20:F2:297:GLU:O	32:K2:131:PRO:HG3	2.20	0.42
23:G2:118:ILE:HG23	23:G2:135:ILE:HD12	2.00	0.42
26:H2:103:VAL:HG23	26:H2:105:GLY:H	1.84	0.42
27:H3:34:TYR:CZ	73:q2:8:LYS:HD2	2.54	0.42
35:L2:81:ARG:HG2	35:L2:88:ARG:CZ	2.49	0.42
42:O3:45:THR:OG1	42:O3:49:GLY:HA2	2.20	0.42
46:Q3:90:ARG:HG2	46:Q3:93:ARG:NH2	2.21	0.42
47:R2:82:THR:HG21	61:b2:37:THR:HG22	2.02	0.42
63:d2:68:LYS:HG3	63:d2:69:ILE:HD12	2.01	0.42
7:n2:35:A:H2'	7:n2:36:A:C8	2.54	0.42
77:u2:83:TYR:HB2	77:u2:202:ILE:HG22	2.01	0.42
78:v2:22:VAL:HG13	78:v2:68:TYR:HE1	1.84	0.42
79:w2:77:VAL:HG22	79:w2:86:ILE:HD12	2.02	0.42
83:m2:1549:C:H2'	83:m2:1550:G:O4'	2.19	0.42
1:A1:245:LYS:HD2	2:A2:1709:A:H4'	2.01	0.42
2:A2:41:C:O2'	2:A2:42:A:H5'	2.19	0.42
2:A2:492:G:N2	2:A2:670:G:H22	2.14	0.42
2:A2:1621:G:H2'	2:A2:1622:C:O4'	2.19	0.42
2:A2:2235:G:H2'	2:A2:2236:G:C8	2.55	0.42
2:A2:2495:U:O4'	14:D2:187:HIS:HB3	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:4322:C:O3'	2:A2:4323:C:H3'	2.20	0.42
2:A2:4499:G:H2'	2:A2:4500:G:H8	1.84	0.42
6:B3:33:TRP:CZ2	6:B3:102:ARG:HG3	2.55	0.42
7:Bv:7:A:H8	7:Bv:7:A:OP2	2.03	0.42
7:Bv:19:G:H4'	7:Bv:20:U:OP2	2.19	0.42
7:Bv:75:C:H2'	7:Bv:76:A:O4'	2.19	0.42
17:E2:82:PRO:HG3	17:E2:171:LEU:HD21	2.01	0.42
18:E3:77:ASN:ND2	18:E3:79:LYS:HE3	2.34	0.42
26:H2:184:GLY:O	26:H2:185:PRO:C	2.63	0.42
29:I3:42:MET:HB2	29:I3:57:ARG:HG2	2.02	0.42
29:I3:173:LEU:HD11	29:I3:189:ILE:HD12	2.01	0.42
38:M3:61:TYR:O	38:M3:64:LEU:HG	2.19	0.42
49:S2:47:MET:HE2	49:S2:115:ARG:HH21	1.84	0.42
70:k2:7:TRP:HB2	70:k2:44:ILE:HD12	2.01	0.42
70:k2:98:ARG:HG3	70:k2:98:ARG:HH11	1.83	0.42
77:u2:142:SER:HA	77:u2:146:GLN:HB3	2.02	0.42
83:m2:675:G:H2'	83:m2:676:C:C6	2.54	0.42
83:m2:845:C:H2'	83:m2:846:U:O4'	2.20	0.42
83:m2:878:C:H2'	83:m2:879:C:C6	2.55	0.42
2:A2:124:C:H2'	2:A2:125:C:H6	1.85	0.42
2:A2:700:C:H2'	2:A2:701:C:H6	1.85	0.42
2:A2:2061:G:H1'	2:A2:2087:A:H61	1.84	0.42
2:A2:2504:C:H2'	2:A2:2505:G:H8	1.81	0.42
2:A2:3261:C:H2'	2:A2:3262:U:O4'	2.19	0.42
2:A2:3602:G:H2'	2:A2:3603:A:C8	2.55	0.42
2:A2:3962:A:H2'	2:A2:3963:A:O4'	2.20	0.42
2:A2:4303:A:H2'	2:A2:4304:G:O4'	2.19	0.42
2:A2:4413:G:H2'	2:A2:4414:A:H8	1.84	0.42
2:A2:4427:C:H2'	2:A2:4428:C:C6	2.53	0.42
4:B1:70:LEU:HA	4:B1:70:LEU:HD23	1.80	0.42
5:B2:61:G:O2'	5:B2:62:U:H5'	2.19	0.42
7:Bv:48:C:C2	7:Bv:59:U:H1'	2.54	0.42
20:F2:143:ARG:O	20:F2:143:ARG:HG3	2.19	0.42
21:F3:82:LYS:HE3	83:m2:1211:A:H5''	2.02	0.42
23:G2:183:TYR:HA	23:G2:190:PHE:HA	2.02	0.42
34:L1:59:PRO:HD2	34:L1:153:SER:HB2	2.01	0.42
37:M2:81:TRP:HZ3	37:M2:130:GLU:HG2	1.85	0.42
43:P2:113:LYS:HE3	43:P2:113:LYS:HB2	1.89	0.42
44:P3:43:LYS:HG2	44:P3:43:LYS:H	1.71	0.42
50:S3:19:HIS:HB3	50:S3:22:LYS:HB2	2.02	0.42
60:a2:69:LYS:HG2	60:a2:73:HIS:CE1	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
64:e2:51:GLU:HG2	64:e2:52:LYS:N	2.35	0.42
69:j2:70:THR:HG23	69:j2:72:ASN:O	2.20	0.42
7:n2:16:U:H3	7:n2:61:C:H5'	1.84	0.42
72:p2:31:TYR:CD2	72:p2:94:LYS:HA	2.55	0.42
73:q2:72:VAL:HG13	78:v2:20:VAL:HG11	2.02	0.42
73:q2:193:ASP:N	73:q2:194:PRO:HD3	2.35	0.42
80:x2:125:PRO:HB2	80:x2:127:LYS:NZ	2.34	0.42
80:x2:145:LYS:NZ	83:m2:1828:G:H4'	2.35	0.42
83:m2:149:A:H2	83:m2:169:U:H3	1.66	0.42
83:m2:347:U:H2'	83:m2:348:C:H6	1.84	0.42
83:m2:1038:A:C4	83:m2:1039:G:C8	3.08	0.42
83:m2:1100:C:H2'	83:m2:1101:G:H8	1.83	0.42
83:m2:1279:C:H2'	83:m2:1280:A:H8	1.85	0.42
83:m2:1283:G:C6	83:m2:1284:A:C6	3.07	0.42
83:m2:1619:G:N2	83:m2:1621:A:H3'	2.34	0.42
2:A2:279:A:C4	25:H1:12:ARG:HG2	2.55	0.42
2:A2:1027:G:H2'	2:A2:1028:G:C8	2.52	0.42
2:A2:2283:G:H4'	2:A2:2538:A:H4'	2.02	0.42
2:A2:2500:A:H2'	2:A2:2501:A:H8	1.82	0.42
2:A2:3562:A:H2'	20:F2:69:THR:HG21	2.00	0.42
2:A2:4023:G:C5	7:n2:76:A:C2	3.08	0.42
4:B1:180:PRO:HG3	4:B1:219:VAL:HG13	2.02	0.42
20:F2:287:THR:HG21	70:k2:5:LEU:HD12	2.02	0.42
23:G2:211:LEU:HD12	23:G2:223:PHE:CE2	2.50	0.42
25:H1:85:VAL:HA	68:i2:49:GLY:HA2	2.02	0.42
31:J3:142:LYS:HB3	31:J3:142:LYS:HE3	1.77	0.42
31:J3:166:ARG:HB2	31:J3:248:TYR:CE1	2.54	0.42
34:L1:5:VAL:HG23	34:L1:7:ARG:H	1.85	0.42
48:R3:66:LYS:HZ3	75:s2:103:LEU:N	2.18	0.42
64:e2:14:THR:O	64:e2:17:ARG:HB2	2.20	0.42
67:h2:9:ARG:NH2	83:m2:1853:A:O5'	2.53	0.42
72:p2:66:VAL:HG13	72:p2:85:LYS:HE2	2.01	0.42
79:w2:60:CYS:SG	79:w2:63:THR:HG22	2.60	0.42
80:x2:50:ARG:CZ	83:m2:1620:C:H4'	2.50	0.42
83:m2:116:OMU:HM23	83:m2:116:OMU:H1'	1.77	0.42
83:m2:176:U:H2'	83:m2:177:G:O4'	2.19	0.42
83:m2:355:OMC:H1'	83:m2:355:OMC:HM23	1.80	0.42
83:m2:682:G:H2'	83:m2:683:U:H6	1.85	0.42
83:m2:1039:G:H4'	83:m2:1847:A:H4'	2.01	0.42
83:m2:1306:U:H2'	83:m2:1307:C:C6	2.55	0.42
2:A2:493:U:H2'	2:A2:494:G:C8	2.55	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:1490:PSU:H4'	2:A2:1493:G:C2	2.55	0.42
2:A2:1606:A:H4'	2:A2:1607:A:O5'	2.19	0.42
2:A2:1911:G:H2'	2:A2:1912:A:H8	1.85	0.42
2:A2:2141:U:C2	2:A2:2142:G:C8	3.08	0.42
2:A2:2194:G:H5'	2:A2:2533:G:OP2	2.20	0.42
2:A2:2284:A:N3	2:A2:2284:A:H2'	2.35	0.42
2:A2:2370:U:H2'	2:A2:2371:C:H6	1.85	0.42
2:A2:3253:G:H1'	2:A2:3254:C:O5'	2.19	0.42
2:A2:3335:U:C4	14:D2:20:VAL:HB	2.55	0.42
2:A2:3392:A:H2'	2:A2:3393:A:C8	2.55	0.42
2:A2:3526:C:H2'	2:A2:3527:A:H8	1.85	0.42
2:A2:3567:C:H2'	2:A2:3568:U:C6	2.55	0.42
2:A2:3713:U:H4'	2:A2:3714:C:H5	1.85	0.42
2:A2:3736:G:C8	47:R2:44:PRO:HG3	2.55	0.42
2:A2:3957:G:C6	39:N2:80:VAL:HG21	2.55	0.42
4:B1:51:LEU:O	4:B1:55:VAL:HG23	2.20	0.42
5:B2:102:U:H3'	5:B2:103:A:H8	1.84	0.42
7:Bv:6:G:O2'	7:Bv:7:A:H5'	2.20	0.42
12:C3:26:SER:HB3	12:C3:110:VAL:HA	2.02	0.42
16:E1:109:ILE:HG22	16:E1:111:GLU:H	1.83	0.42
17:E2:202:GLU:O	17:E2:202:GLU:HG2	2.20	0.42
19:F1:72:ALA:HB2	19:F1:157:ILE:HD12	2.01	0.42
24:G3:44:ARG:HH11	24:G3:63:ARG:HB3	1.85	0.42
25:H1:104:GLU:HA	25:H1:160:GLU:HG3	2.02	0.42
31:J3:84:PHE:HZ	31:J3:262:THR:HG23	1.84	0.42
31:J3:170:TRP:NE1	44:P3:97:ARG:HD2	2.35	0.42
43:P2:58:GLY:HA2	43:P2:125:CYS:HB3	2.02	0.42
48:R3:69:THR:HG23	48:R3:72:VAL:H	1.85	0.42
61:b2:57:VAL:O	61:b2:61:ILE:HG13	2.20	0.42
61:b2:96:THR:OG1	61:b2:99:GLU:HG2	2.19	0.42
64:e2:19:ASP:OD2	64:e2:38:CYS:HB3	2.20	0.42
7:n2:16:U:C2	7:n2:60:U:H2'	2.55	0.42
73:q2:161:GLY:HA3	83:m2:1390:A:H61	1.85	0.42
73:q2:163:PRO:HA	73:q2:166:TYR:CE1	2.55	0.42
1:A1:65:ARG:NH1	2:A2:874:C:H5''	2.35	0.41
2:A2:68:U:H2'	2:A2:69:A:O4'	2.19	0.41
2:A2:134:G:H1'	2:A2:135:U:C5	2.55	0.41
2:A2:686:C:H2'	2:A2:687:G:C8	2.55	0.41
2:A2:1078:G:H21	2:A2:1078:G:P	2.43	0.41
2:A2:1281:C:H2'	2:A2:1282:C:C6	2.55	0.41
2:A2:2235:G:H2'	2:A2:2236:G:H8	1.85	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:2388:U:H2'	2:A2:2389:C:C6	2.55	0.41
2:A2:2575:C:H5''	35:L2:56:THR:OG1	2.20	0.41
2:A2:3441:A2M:H2	2:A2:4203:U:O2	2.19	0.41
2:A2:3997:C:H2'	2:A2:3998:U:C6	2.55	0.41
2:A2:4119:A:O2'	2:A2:4162:A:H1'	2.20	0.41
2:A2:4417:G:H21	37:M2:173:ASN:HD21	1.68	0.41
2:A2:4648:U:H2'	2:A2:4649:U:H6	1.85	0.41
6:B3:110:LEU:HD23	6:B3:110:LEU:O	2.20	0.41
7:Bz:59:U:H2'	7:Bz:60:U:C6	2.55	0.41
15:D3:17:CYS:SG	15:D3:55:ILE:HA	2.60	0.41
32:K2:18:PRO:HD3	32:K2:52:PHE:CD1	2.55	0.41
34:L1:179:LEU:O	34:L1:183:ILE:HG23	2.20	0.41
36:L3:38:ARG:HG2	52:T3:31:ARG:HB2	2.02	0.41
39:N2:50:LYS:O	39:N2:92:ARG:HD3	2.20	0.41
43:P2:30:ASP:OD2	43:P2:104:VAL:HA	2.20	0.41
50:S3:73:LEU:HB3	50:S3:77:CYS:HB2	2.01	0.41
59:Z2:11:PHE:HB2	59:Z2:101:ILE:CD1	2.49	0.41
72:p2:83:LYS:O	72:p2:103:MET:HG2	2.19	0.41
74:r2:108:ARG:NH2	83:m2:848:G:H3'	2.35	0.41
77:u2:62:VAL:HA	77:u2:77:ARG:HA	2.02	0.41
80:x2:38:SER:OG	80:x2:41:GLN:HG2	2.20	0.41
81:y2:109:LYS:HE3	81:y2:109:LYS:HB2	1.90	0.41
83:m2:291:G:H2'	83:m2:292:U:C6	2.55	0.41
83:m2:612:G:H2'	83:m2:613:G:H8	1.85	0.41
83:m2:1033:A2M:HM'3	83:m2:1033:A2M:H1'	1.81	0.41
83:m2:1194:U:H2'	83:m2:1195:U:C6	2.54	0.41
2:A2:6:C:H2'	2:A2:7:C:H6	1.84	0.41
2:A2:686:C:H2'	2:A2:687:G:H8	1.85	0.41
2:A2:1114:G:H2'	2:A2:1115:C:C6	2.55	0.41
2:A2:3267:A:O2'	77:u2:92:ARG:NH1	2.53	0.41
2:A2:3348:A:N6	2:A2:3479:G:H21	1.97	0.41
2:A2:3421:G:H3'	2:A2:3422:A:C2	2.55	0.41
2:A2:3581:OMU:H1'	2:A2:3581:OMU:HM23	1.82	0.41
2:A2:4401:C:H2'	2:A2:4402:G:O4'	2.20	0.41
3:A3:80:PRO:HG2	3:A3:83:PHE:HB2	2.01	0.41
3:A3:119:ALA:O	3:A3:123:LEU:HG	2.20	0.41
10:C1:92:MET:SD	10:C1:161:ILE:HD11	2.60	0.41
11:C2:27:U:H2'	11:C2:28:C:C6	2.55	0.41
11:C2:141:C:H2'	11:C2:142:U:C6	2.54	0.41
13:D1:91:LEU:CD1	13:D1:135:ILE:HA	2.49	0.41
29:I3:220:ASP:OD2	29:I3:223:GLU:HB2	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:K2:53:MET:HE1	32:K2:143:ARG:NH1	2.33	0.41
54:U3:143:LYS:HG3	83:m2:1312:U:H4'	2.01	0.41
72:p2:113:MET:HE3	72:p2:113:MET:HB2	1.90	0.41
74:r2:3:ARG:HD3	83:m2:451:A:H4'	2.02	0.41
76:t2:178:LYS:HD3	76:t2:182:GLY:HA2	2.02	0.41
81:y2:9:SER:HB3	81:y2:26:LYS:HB3	2.02	0.41
83:m2:478:A:H2'	83:m2:479:G:O4'	2.19	0.41
83:m2:597:U:H2'	83:m2:598:U:C6	2.55	0.41
83:m2:662:C:H2'	83:m2:663:U:C6	2.55	0.41
83:m2:820:A:H2'	83:m2:821:G:C8	2.55	0.41
83:m2:888:A:H1'	83:m2:903:G:N2	2.36	0.41
83:m2:1206:A:H2'	83:m2:1207:C:H6	1.85	0.41
2:A2:1007:A:H2'	2:A2:1008:C:C6	2.56	0.41
2:A2:1135:G:H2'	2:A2:3532:A:N7	2.35	0.41
2:A2:2090:C:H2'	2:A2:2091:G:H8	1.85	0.41
2:A2:2241:G:C6	2:A2:2248:G:C6	3.08	0.41
2:A2:2290:G:H2'	2:A2:2291:A:C8	2.55	0.41
2:A2:3556:G:OP1	2:A2:3557:A:H4'	2.20	0.41
2:A2:4586:C:H1'	26:H2:253:ARG:HH22	1.85	0.41
11:C2:64:U:C2	11:C2:65:A:C8	3.08	0.41
15:D3:74:LYS:HE3	15:D3:74:LYS:HB3	1.91	0.41
17:E2:153:MET:HB2	17:E2:194:LEU:HD11	2.02	0.41
19:F1:108:GLU:OE2	62:c2:18:THR:HG22	2.20	0.41
30:J2:42:ARG:HH22	30:J2:109:VAL:HG23	1.85	0.41
32:K2:53:MET:CE	32:K2:143:ARG:HH12	2.30	0.41
59:Z2:75:THR:HG21	59:Z2:87:LYS:HG2	2.01	0.41
72:p2:32:ASP:HA	72:p2:46:LYS:HA	2.01	0.41
74:r2:201:HIS:NE2	83:m2:859:U:H4'	2.35	0.41
76:t2:10:LYS:HA	76:t2:45:ILE:O	2.20	0.41
1:A1:258:ARG:HG3	1:A1:258:ARG:O	2.21	0.41
2:A2:1553:A:H2'	2:A2:1554:G:C8	2.56	0.41
2:A2:1579:C:H2'	2:A2:1580:C:C6	2.55	0.41
2:A2:2118:A2M:HM'2	2:A2:2119:OMG:H5'	2.03	0.41
2:A2:2160:G:H5''	63:d2:14:LYS:HE2	2.02	0.41
2:A2:2167:A:H2'	2:A2:2168:U:H6	1.84	0.41
2:A2:2293:U:H2'	2:A2:2294:C:C6	2.55	0.41
2:A2:3404:A:H2'	2:A2:3405:C:H6	1.83	0.41
2:A2:3554:G:H2'	2:A2:3555:OMG:O4'	2.19	0.41
2:A2:3714:C:OP2	2:A2:3715:G:H5'	2.21	0.41
4:B1:100:HIS:HA	4:B1:103:ARG:HG3	2.02	0.41
5:B2:71:G:H2'	5:B2:72:U:O4'	2.19	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:E1:35:ARG:O	16:E1:39:VAL:HG23	2.20	0.41
29:I3:125:ARG:HG2	29:I3:150:TRP:CG	2.55	0.41
30:J2:119:VAL:HG12	30:J2:146:ILE:HG12	2.01	0.41
31:J3:209:VAL:HB	31:J3:210:PRO:HD3	2.02	0.41
32:K2:49:LYS:O	32:K2:53:MET:HG3	2.20	0.41
36:L3:29:LEU:HD23	52:T3:42:PHE:CE2	2.55	0.41
36:L3:134:HIS:HE1	36:L3:164:PRO:HD2	1.84	0.41
37:M2:8:ARG:O	37:M2:33:PHE:HA	2.20	0.41
40:N3:37:ILE:HD11	40:N3:63:VAL:HG11	2.01	0.41
46:Q3:102:THR:HG22	46:Q3:107:ARG:NH2	2.36	0.41
53:U2:90:ALA:HB3	53:U2:120:GLN:NE2	2.34	0.41
59:Z2:11:PHE:CE1	59:Z2:26:ALA:HB1	2.55	0.41
62:c2:79:THR:HG22	62:c2:81:ILE:HG22	2.03	0.41
83:m2:93:U:H2'	83:m2:94:G:O4'	2.20	0.41
83:m2:514:A2M:C2	83:m2:515:G:C8	3.03	0.41
83:m2:1174:U:H2'	83:m2:1175:A:H8	1.85	0.41
83:m2:1718:C:H2'	83:m2:1719:C:C6	2.55	0.41
2:A2:6:C:H2'	2:A2:7:C:C6	2.56	0.41
2:A2:1255:C:H2'	2:A2:1256:A:H8	1.84	0.41
2:A2:1352:G:O4'	14:D2:15:VAL:HG13	2.20	0.41
2:A2:1506:U:H2'	2:A2:1507:C:O4'	2.21	0.41
2:A2:1613:G:H2'	2:A2:1614:C:C6	2.55	0.41
2:A2:2140:U:H2'	2:A2:2141:U:C6	2.56	0.41
2:A2:2179:OMG:H1'	2:A2:2179:OMG:HM23	1.71	0.41
2:A2:3458:U:O2'	2:A2:3459:A:H5'	2.20	0.41
2:A2:3526:C:H2'	2:A2:3527:A:C8	2.56	0.41
2:A2:3618:A:H3'	2:A2:3621:A:H2	1.84	0.41
2:A2:3785:G:H4'	2:A2:3786:U:C5	2.56	0.41
2:A2:3834:G:C4'	14:D2:226:ARG:HH22	2.33	0.41
2:A2:4054:C:H42	2:A2:4093:A:H61	1.67	0.41
2:A2:4126:A:H2'	2:A2:4128:C:O5'	2.21	0.41
19:F1:80:GLU:HB3	19:F1:110:LEU:CD1	2.51	0.41
24:G3:26:GLN:OE1	75:s2:126:THR:HB	2.21	0.41
29:I3:32:LEU:HG	29:I3:71:ILE:HD11	2.02	0.41
31:J3:207:ALA:HB2	83:m2:4:C:H4'	2.02	0.41
33:K3:27:PHE:CE2	33:K3:41:LEU:HD11	2.56	0.41
33:K3:52:ILE:HG23	33:K3:109:LEU:HD11	2.03	0.41
38:M3:18:LEU:O	38:M3:22:LEU:HD23	2.20	0.41
40:N3:70:LYS:HD3	83:m2:1021:C:OP2	2.20	0.41
42:O3:98:ARG:HG3	42:O3:132:VAL:HG23	2.03	0.41
55:V2:65:VAL:HG13	55:V2:68:ARG:NH2	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:n2:24:G:H2'	7:n2:25:C:O4'	2.21	0.41
71:o2:83:GLY:O	71:o2:87:VAL:HG23	2.20	0.41
72:p2:35:ALA:HB3	72:p2:42:ARG:HA	2.01	0.41
82:z2:40:ILE:HG22	82:z2:40:ILE:O	2.20	0.41
83:m2:423:G:H4'	83:m2:662:C:H42	1.85	0.41
83:m2:1683:U:H2'	83:m2:1684:C:C6	2.56	0.41
2:A2:493:U:H2'	2:A2:494:G:H8	1.85	0.41
2:A2:1323:G:H2'	2:A2:1324:C:C6	2.55	0.41
2:A2:2089:C:C5	20:F2:191:ALA:HB2	2.56	0.41
2:A2:2283:G:H2'	2:A2:2284:A:O4'	2.21	0.41
2:A2:4090:U:H2'	2:A2:4091:U:O4'	2.21	0.41
2:A2:4541:G:H8	2:A2:4541:G:O5'	2.04	0.41
2:A2:4560:G:OP2	2:A2:4560:G:H8	2.04	0.41
2:A2:4676:U:H2'	2:A2:4677:G:C8	2.55	0.41
2:A2:4677:G:H2'	2:A2:4678:C:C6	2.55	0.41
10:C1:114:ILE:HD13	10:C1:165:THR:HG23	2.02	0.41
20:F2:198:ASN:OD1	49:S2:10:ASP:HA	2.21	0.41
20:F2:298:ILE:O	20:F2:302:LEU:HG	2.20	0.41
21:F3:22:ARG:HE	42:O3:142:ARG:HB3	1.86	0.41
23:G2:146:LEU:HD22	23:G2:163:LEU:HD22	2.02	0.41
23:G2:287:PHE:O	23:G2:291:GLN:HG2	2.20	0.41
34:L1:49:PHE:CZ	34:L1:159:MET:HB3	2.56	0.41
38:M3:36:ARG:HH22	83:m2:1312:U:H3'	1.86	0.41
48:R3:66:LYS:HE3	48:R3:66:LYS:HB3	1.94	0.41
56:W2:18:LEU:HD23	56:W2:84:ALA:HB1	2.02	0.41
59:Z2:35:ALA:HB3	59:Z2:38:GLU:HG3	2.02	0.41
78:v2:95:ARG:HH11	78:v2:95:ARG:HA	1.85	0.41
82:z2:50:ILE:O	82:z2:54:VAL:HG23	2.20	0.41
83:m2:1269:C:H2'	83:m2:1270:C:C6	2.55	0.41
83:m2:1553:U:H4'	83:m2:1554:G:C5	2.55	0.41
2:A2:431:G:H21	2:A2:3544:G:H3'	1.86	0.41
2:A2:513:U:O2'	2:A2:514:U:H5'	2.20	0.41
2:A2:1259:C:H42	2:A2:1901:A:N6	1.98	0.41
2:A2:1284:U:H2'	2:A2:1285:C:H6	1.85	0.41
2:A2:1577:A:H2'	2:A2:1578:A:C8	2.55	0.41
2:A2:2289:C:H2'	2:A2:2290:G:O4'	2.21	0.41
2:A2:3418:U:H3'	2:A2:3419:A:H8	1.85	0.41
2:A2:3848:OMG:HM23	2:A2:3848:OMG:H1'	1.66	0.41
2:A2:4063:G:H2'	2:A2:4064:C:H6	1.86	0.41
2:A2:4360:A:N3	2:A2:4361:U:H5'	2.35	0.41
2:A2:4569:C:O2	2:A2:4570:U:H1'	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:Bz:43:C:H2'	7:Bz:44:G:C8	2.56	0.41
10:C1:106:GLN:HB2	10:C1:111:LEU:HB3	2.03	0.41
17:E2:303:ALA:HA	17:E2:368:ILE:HD12	2.02	0.41
26:H2:172:LEU:HD11	26:H2:183:THR:HG22	2.03	0.41
33:K3:27:PHE:HE2	33:K3:41:LEU:HD11	1.85	0.41
36:L3:20:PHE:O	83:m2:605:C:H4'	2.21	0.41
38:M3:24:THR:OG1	38:M3:119:GLN:HG2	2.20	0.41
44:P3:57:ARG:HE	83:m2:921:A:H4'	1.85	0.41
48:R3:103:HIS:NE2	75:s2:91:ARG:HD2	2.36	0.41
49:S2:22:PRO:O	49:S2:26:ARG:HG3	2.19	0.41
57:X2:19:GLU:O	57:X2:90:ARG:HD2	2.21	0.41
72:p2:60:ASP:HA	72:p2:63:LYS:HG2	2.01	0.41
83:m2:152:U:H2'	83:m2:153:G:C8	2.55	0.41
83:m2:589:A:C8	83:m2:592:A:H2'	2.55	0.41
83:m2:919:U:H2'	83:m2:920:U:H6	1.85	0.41
2:A2:254:G:N3	2:A2:254:G:H2'	2.35	0.41
2:A2:261:G:C2	2:A2:262:G:C5	3.09	0.41
2:A2:857:A:H8	2:A2:857:A:OP2	2.03	0.41
2:A2:944:G:H5'	39:N2:140:PHE:CE1	2.56	0.41
2:A2:1211:A:C8	53:U2:114:LYS:HD2	2.55	0.41
2:A2:1458:C:H2'	2:A2:1459:A:C8	2.56	0.41
2:A2:2106:OMC:HM23	2:A2:2106:OMC:H1'	1.62	0.41
2:A2:3385:PSU:O5'	2:A2:3385:PSU:H6	2.04	0.41
2:A2:3764:C:H3'	2:A2:3765:G:C8	2.56	0.41
2:A2:3936:C:H2'	2:A2:3937:U:C6	2.56	0.41
2:A2:4226:U:H3'	2:A2:4227:G:H5''	2.01	0.41
5:B2:77:A:H2'	5:B2:78:C:O4'	2.21	0.41
7:Bz:18:G:H3'	7:Bz:57:G:H22	1.85	0.41
14:D2:92:LYS:HG3	14:D2:93:LYS:HG2	2.02	0.41
14:D2:242:ARG:HG2	14:D2:250:LYS:NZ	2.35	0.41
16:E1:3:GLN:NE2	16:E1:5:GLN:HB3	2.36	0.41
28:I2:88:LEU:HD12	28:I2:99:LEU:HD13	2.03	0.41
30:J2:67:VAL:HG13	30:J2:82:ARG:HG3	2.02	0.41
38:M3:36:ARG:NH2	83:m2:1312:U:H3'	2.35	0.41
43:P2:95:PHE:HB2	45:Q2:19:ARG:HG2	2.02	0.41
53:U2:62:HIS:NE2	53:U2:64:LYS:HD3	2.36	0.41
54:U3:135:HIS:HB2	54:U3:138:ARG:O	2.21	0.41
57:X2:40:LYS:C	57:X2:43:PRO:HD2	2.46	0.41
64:e2:8:ILE:HG23	64:e2:56:LEU:HD13	2.02	0.41
72:p2:163:GLN:O	72:p2:167:LYS:HG2	2.21	0.41
80:x2:28:MET:SD	80:x2:32:GLN:HB2	2.61	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
80:x2:145:LYS:HZ3	83:m2:1828:G:H4'	1.86	0.41
83:m2:112:U:H5''	83:m2:113:G:H2'	2.02	0.41
83:m2:220:U:H2'	83:m2:221:U:H6	1.83	0.41
83:m2:555:A:H2'	83:m2:556:A:H8	1.84	0.41
83:m2:630:A:H61	83:m2:1502:G:H1'	1.85	0.41
83:m2:668:U:H2'	83:m2:669:U:C6	2.56	0.41
2:A2:316:U:OP1	2:A2:316:U:H3'	2.20	0.41
2:A2:480:C:OP1	70:k2:67:ARG:HD2	2.20	0.41
2:A2:1015:C:H2'	2:A2:1016:A:C8	2.55	0.41
2:A2:1118:C:H2'	2:A2:1119:C:H6	1.86	0.41
2:A2:1125:G:C2	2:A2:1126:A:H1'	2.56	0.41
2:A2:1131:U:H2'	2:A2:1132:C:C6	2.56	0.41
2:A2:2256:C:H2'	2:A2:2257:G:H5'	2.02	0.41
2:A2:3362:C:H5''	14:D2:226:ARG:HB3	2.02	0.41
2:A2:3599:A:H5'	2:A2:3600:G:OP2	2.21	0.41
2:A2:3743:G:H2'	2:A2:3744:G:C8	2.55	0.41
2:A2:4111:U:H2'	2:A2:4112:U:C6	2.56	0.41
2:A2:4140:A:H4'	2:A2:4141:G:C8	2.56	0.41
2:A2:4170:A:N7	17:E2:257:TRP:CD1	2.89	0.41
2:A2:4538:A:H5''	22:G1:125:THR:HG21	2.03	0.41
4:B1:110:LYS:HD2	4:B1:113:ARG:NH1	2.32	0.41
5:B2:16:A:H2'	5:B2:17:C:C6	2.56	0.41
6:B3:33:TRP:HZ2	6:B3:102:ARG:HG3	1.86	0.41
11:C2:8:U:H2'	11:C2:9:A:C8	2.56	0.41
11:C2:56:G:H2'	11:C2:57:C:O4'	2.20	0.41
13:D1:10:ARG:HD3	13:D1:11:TYR:CE1	2.56	0.41
16:E1:15:LEU:HD23	16:E1:136:ARG:HG2	2.03	0.41
17:E2:219:VAL:HG11	17:E2:337:VAL:HB	2.03	0.41
20:F2:11:TYR:CZ	20:F2:148:PRO:HB2	2.55	0.41
20:F2:142:HIS:O	20:F2:143:ARG:C	2.63	0.41
21:F3:85[A]:ARG:NH1	83:m2:1212:G:H5''	2.36	0.41
21:F3:85[B]:ARG:NH1	83:m2:1212:G:H5''	2.36	0.41
31:J3:98:LEU:O	31:J3:102:LEU:HG	2.19	0.41
31:J3:207:ALA:HB3	31:J3:210:PRO:HD2	2.03	0.41
33:K3:63:MET:HG2	33:K3:99:GLY:O	2.21	0.41
34:L1:33:GLU:HB2	34:L1:167:VAL:O	2.21	0.41
34:L1:41:TYR:HD1	34:L1:200:ASN:ND2	2.19	0.41
41:O2:28:PRO:HB2	41:O2:34:MET:HB2	2.02	0.41
42:O3:54:CYS:SG	42:O3:84:ARG:HD2	2.61	0.41
44:P3:57:ARG:HD2	50:S3:26:GLN:HE21	1.85	0.41
47:R2:71:LEU:HD23	47:R2:108:GLN:HE22	1.86	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:T2:60:LYS:HB2	51:T2:60:LYS:HE3	1.87	0.41
57:X2:84:ILE:HD11	57:X2:108:TYR:CE1	2.56	0.41
7:n2:23:A:H2'	7:n2:24:G:C8	2.56	0.41
71:o2:32:PHE:CG	83:m2:1099:G:H4'	2.56	0.41
72:p2:194:GLY:O	72:p2:198:GLU:HG3	2.21	0.41
73:q2:204:LEU:HD12	73:q2:205:PRO:HD2	2.03	0.41
74:r2:52:LEU:HD21	74:r2:99:PHE:CE2	2.55	0.41
74:r2:104:ASP:O	74:r2:190:GLY:HA3	2.21	0.41
75:s2:76:MET:HB3	75:s2:89:THR:CG2	2.51	0.41
76:t2:120:ARG:HA	83:m2:915:A:N6	2.36	0.41
77:u2:123:ARG:NH1	77:u2:128:LYS:HE2	2.36	0.41
78:v2:57:TYR:O	78:v2:72:THR:HG22	2.21	0.41
83:m2:219:A:H1'	83:m2:311:G:N2	2.35	0.41
83:m2:297:C:H2'	83:m2:298:U:C6	2.56	0.41
83:m2:510:A:H5'	83:m2:511:OMG:OP2	2.21	0.41
83:m2:511:OMG:HM23	83:m2:511:OMG:H1'	1.87	0.41
83:m2:684:U:H2'	83:m2:685:OMG:O4'	2.21	0.41
83:m2:811:A:H2'	83:m2:812:A:O4'	2.20	0.41
83:m2:903:G:H8	83:m2:903:G:P	2.44	0.41
83:m2:972:G:H4'	83:m2:973:G:OP1	2.20	0.41
83:m2:1163:U:H2'	83:m2:1164:C:C6	2.56	0.41
83:m2:1830:C:H2'	83:m2:1831:G:C8	2.56	0.41
2:A2:47:A:H5''	19:F1:16:LYS:HG3	2.03	0.41
2:A2:73:A:OP1	19:F1:106:SER:HB3	2.21	0.41
2:A2:517:C:H2'	2:A2:518:C:H6	1.85	0.41
2:A2:810:U:H3'	2:A2:811:G:C8	2.56	0.41
2:A2:1723:C:C4	22:G1:17:PHE:HB3	2.56	0.41
2:A2:3626:G:H2'	2:A2:3627:G:C8	2.56	0.41
3:A3:33:ILE:H	3:A3:33:ILE:HG13	1.72	0.41
6:B3:33:TRP:CH2	6:B3:99:VAL:HG12	2.56	0.41
10:C1:26:ILE:HG13	10:C1:35:ARG:HG2	2.03	0.41
17:E2:217:ILE:HD13	17:E2:284:ILE:HD11	2.01	0.41
18:E3:5:ARG:HB3	44:P3:77:PRO:HG3	2.02	0.41
21:F3:53:ILE:HG23	42:O3:126:ILE:HD12	2.03	0.41
26:H2:78:ARG:HB2	55:V2:116:LEU:HD22	2.02	0.41
26:H2:180:LEU:O	26:H2:195:ARG:HA	2.21	0.41
29:I3:40:ILE:HB	29:I3:59:LEU:HB2	2.03	0.41
33:K3:134:GLY:O	83:m2:169:U:H4'	2.21	0.41
34:L1:58:THR:O	34:L1:174:MET:HE1	2.21	0.41
36:L3:64:ASP:OD1	36:L3:67:ASP:HB2	2.20	0.41
38:M3:91:LEU:HB2	38:M3:104:VAL:HG11	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:P3:53:ILE:HG13	76:t2:143:ARG:HD2	2.03	0.41
44:P3:105:THR:HB	83:m2:864:A:H8	1.86	0.41
47:R2:77:ILE:HD13	47:R2:100:VAL:HG12	2.02	0.41
49:S2:2:LYS:HE3	49:S2:2:LYS:HB3	1.82	0.41
57:X2:33:ILE:O	57:X2:36:VAL:HG22	2.21	0.41
65:f2:18:LYS:HB2	65:f2:18:LYS:HE3	1.83	0.41
67:h2:1:MET:HG3	83:m2:1854:C:P	2.60	0.41
74:r2:255:ARG:O	74:r2:259:LYS:HG3	2.21	0.41
76:t2:11:PRO:HD2	76:t2:45:ILE:O	2.21	0.41
80:x2:31:GLU:H	80:x2:31:GLU:CD	2.29	0.41
81:y2:143:LYS:NZ	83:m2:1251:C:H4'	2.36	0.41
83:m2:400:A:C8	83:m2:1029:A:H2	2.39	0.41
83:m2:486:A2M:HM'3	83:m2:486:A2M:H1'	1.90	0.41
83:m2:1654:G:H1	83:m2:1674:U:H3	1.68	0.41
2:A2:300:A:H2'	2:A2:301:G:C8	2.52	0.40
2:A2:422:C:H2'	2:A2:423:G:H8	1.86	0.40
2:A2:3346:U:O3'	2:A2:3475:G:H5''	2.22	0.40
2:A2:4080:A:H4'	7:Bz:63:G:O2'	2.20	0.40
2:A2:4332:G:H2'	2:A2:4333:A:C8	2.57	0.40
4:B1:148:GLU:HA	4:B1:177:MET:HE2	2.03	0.40
7:Bv:48:C:H2'	7:Bv:59:U:C4'	2.50	0.40
16:E1:141:ILE:HG13	16:E1:142:ALA:N	2.36	0.40
21:F3:4:LYS:HE2	21:F3:92:ARG:HH12	1.86	0.40
26:H2:197:HIS:HD2	26:H2:199:LYS:HB2	1.86	0.40
28:I2:196:LEU:HD23	28:I2:202:LEU:HD13	2.03	0.40
38:M3:25:ALA:HA	38:M3:28:HIS:CD2	2.56	0.40
38:M3:65:VAL:O	38:M3:69:LEU:HG	2.21	0.40
39:N2:85:LEU:HD23	39:N2:85:LEU:H	1.86	0.40
48:R3:88:LEU:HB3	48:R3:109:TYR:CE2	2.56	0.40
70:k2:21:ASN:O	70:k2:23:GLN:HG3	2.22	0.40
70:k2:54:ALA:HA	70:k2:61:VAL:HG23	2.03	0.40
7:n2:12:U:H2'	7:n2:13:C:O4'	2.21	0.40
73:q2:75:LYS:HE3	78:v2:20:VAL:O	2.22	0.40
80:x2:65:LYS:HE2	80:x2:65:LYS:HB2	1.90	0.40
80:x2:77:LYS:CB	83:m2:1300:G:H5'	2.50	0.40
83:m2:948:U:H2'	83:m2:949:G:H8	1.84	0.40
83:m2:1130:C:C2	83:m2:1131:G:C8	3.09	0.40
83:m2:1844:4AC:H2'	83:m2:1845:G:H8	1.86	0.40
2:A2:35:U:H4'	2:A2:1338:A:H2	1.87	0.40
2:A2:48:G:H3'	25:H1:188:ARG:HH12	1.85	0.40
2:A2:288:G:H2'	2:A2:289:C:C6	2.56	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:1069:G:H2'	2:A2:1070:C:H4'	2.03	0.40
2:A2:3614:G:H3'	2:A2:3615:U:C5'	2.51	0.40
2:A2:4509:U:H2'	2:A2:4510:C:H6	1.86	0.40
3:A3:133:GLY:HA3	83:m2:1625:A:H5''	2.02	0.40
7:Bz:15:G:H22	7:Bz:48:C:H42	1.68	0.40
16:E1:32:ARG:HD2	16:E1:35:ARG:HH21	1.86	0.40
18:E3:6:GLY:O	79:w2:101:ARG:HD3	2.22	0.40
18:E3:105:PHE:HD2	18:E3:112:VAL:HB	1.83	0.40
20:F2:317:ASN:ND2	20:F2:320:LYS:HG3	2.36	0.40
21:F3:94:ASP:OD1	21:F3:96:THR:HG22	2.20	0.40
23:G2:202:GLN:OE1	23:G2:237:GLU:HB2	2.21	0.40
28:I2:201:LEU:H	28:I2:201:LEU:HD12	1.85	0.40
31:J3:125:LYS:NZ	83:m2:1359:A:OP1	2.34	0.40
36:L3:83:ARG:HE	36:L3:150:ARG:HD3	1.85	0.40
43:P2:22:VAL:HG13	43:P2:40:ILE:HA	2.03	0.40
46:Q3:97:TYR:CG	46:Q3:98:GLU:N	2.90	0.40
51:T2:42:LEU:HD21	51:T2:96:VAL:HG12	2.03	0.40
77:u2:73:THR:O	77:u2:74:ARG:HD3	2.21	0.40
83:m2:458:C:H2'	83:m2:459:C:C6	2.56	0.40
83:m2:1803:A:H2'	83:m2:1804:C:H6	1.86	0.40
1:A1:114:VAL:O	1:A1:142:GLY:HA2	2.20	0.40
2:A2:768:G:H8	2:A2:768:G:O5'	2.05	0.40
2:A2:1284:U:H2'	2:A2:1285:C:C6	2.56	0.40
2:A2:1483:G:O2'	2:A2:1656:G:H5'	2.22	0.40
2:A2:1722:C:H3'	2:A2:1723:C:H5''	2.02	0.40
2:A2:2616:OMC:H1'	2:A2:2616:OMC:HM23	1.87	0.40
2:A2:3378:G:O2'	2:A2:3379:A:H5'	2.21	0.40
7:Bz:5:G:C2	7:Bz:69:G:C2	3.10	0.40
16:E1:11:PRO:C	16:E1:12:MET:HE2	2.47	0.40
17:E2:98:GLY:HA2	28:I2:149:TYR:CZ	2.57	0.40
19:F1:50:PRO:HG3	61:b2:121:VAL:HG23	2.03	0.40
23:G2:82:GLU:H	23:G2:82:GLU:HG2	1.71	0.40
27:H3:8:TRP:HZ3	83:m2:1514:C:H5'	1.86	0.40
54:U3:133:GLY:HA2	83:m2:1311:C:H5'	2.04	0.40
71:o2:39:TYR:CD2	71:o2:40:LYS:HG2	2.57	0.40
73:q2:144:GLY:HA2	83:m2:609:U:H3	1.86	0.40
77:u2:27:TYR:HB2	83:m2:383:C:H41	1.87	0.40
78:v2:41:PRO:HG2	78:v2:44:HIS:CD2	2.56	0.40
83:m2:84:A:H2'	83:m2:85:A:C8	2.56	0.40
83:m2:1163:U:H2'	83:m2:1164:C:H6	1.87	0.40
2:A2:180:C:H2'	2:A2:181:C:N1	2.37	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:221:C:H2'	2:A2:222:C:H6	1.87	0.40
2:A2:325:U:H2'	2:A2:326:C:H6	1.87	0.40
2:A2:450:G:H2'	2:A2:451:C:C6	2.56	0.40
2:A2:652:G:H3'	2:A2:653:G:H8	1.86	0.40
2:A2:665:C:H2'	2:A2:666:G:H8	1.86	0.40
2:A2:1885:C:H5''	2:A2:1886:C:OP2	2.21	0.40
2:A2:2117:U:H5'	30:J2:65:GLY:O	2.22	0.40
2:A2:3308:A:H2'	2:A2:3309:A:C8	2.56	0.40
2:A2:3346:U:OP2	14:D2:200:ARG:HD2	2.21	0.40
2:A2:3738:C:H2'	2:A2:3739:G:H8	1.85	0.40
2:A2:3958:OMU:H1'	2:A2:3958:OMU:HM23	1.87	0.40
2:A2:4230:G:H2'	2:A2:4231:U:H6	1.85	0.40
2:A2:4272:OMU:HM23	2:A2:4272:OMU:H1'	1.80	0.40
4:B1:147:VAL:HG13	4:B1:179:VAL:HG21	2.04	0.40
5:B2:15:C:H2'	5:B2:16:A:C8	2.56	0.40
6:B3:5:THR:HG22	6:B3:6:VAL:N	2.37	0.40
7:Bv:12:U:H2'	7:Bv:13:C:O4'	2.22	0.40
7:Bv:26:A:H2	7:Bv:44:G:H22	1.66	0.40
13:D1:87:ILE:HG12	13:D1:138:ILE:HG12	2.03	0.40
18:E3:107:ARG:HD3	18:E3:112:VAL:HG22	2.04	0.40
19:F1:90:VAL:O	19:F1:94:ILE:HG12	2.22	0.40
20:F2:340:ILE:HD13	26:H2:58:VAL:HG21	2.04	0.40
32:K2:178:ARG:H	53:U2:51:GLY:HA2	1.86	0.40
37:M2:16:CYS:SG	37:M2:54:MET:HE2	2.61	0.40
43:P2:42:VAL:HG22	43:P2:61:VAL:HG22	2.03	0.40
44:P3:51:GLU:CD	50:S3:8:LEU:HD21	2.46	0.40
46:Q3:19:GLN:HE21	74:r2:94:LYS:HZ2	1.70	0.40
49:S2:127:GLN:OE1	49:S2:130:LYS:HD2	2.21	0.40
52:T3:56:ASN:HB2	83:m2:640:C:O2'	2.22	0.40
72:p2:158:HIS:O	72:p2:162:ARG:HG3	2.21	0.40
75:s2:49:LEU:HB2	81:y2:50:LYS:HE2	2.04	0.40
75:s2:76:MET:HB3	75:s2:89:THR:HG23	2.03	0.40
76:t2:43:LEU:HD13	76:t2:72:PHE:CE1	2.56	0.40
83:m2:28:U:O2'	83:m2:29:G:H5'	2.21	0.40
83:m2:610:C:H2'	83:m2:611:U:C6	2.56	0.40
83:m2:1098:G:H2'	83:m2:1099:G:C8	2.56	0.40
83:m2:1708:G:H2'	83:m2:1709:U:H6	1.87	0.40
2:A2:418:A:H4'	2:A2:2066:C:H5'	2.04	0.40
2:A2:423:G:O2'	30:J2:118:GLN:HB2	2.21	0.40
2:A2:1260:C:C2	2:A2:1261:G:C8	3.10	0.40
2:A2:1373:A:H2'	2:A2:1374:G:O4'	2.22	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A2:1455:A:H2	2:A2:3574:G:H4'	1.86	0.40
2:A2:1901:A:C2'	2:A2:1902:A:H5'	2.52	0.40
2:A2:2032:C:H2'	2:A2:2033:G:C8	2.57	0.40
2:A2:2506:G:H2'	2:A2:2507:G:H8	1.86	0.40
2:A2:3308:A:O2'	14:D2:182:ALA:HB2	2.21	0.40
2:A2:3415:A:C2	83:m2:1828:G:H21	1.74	0.40
2:A2:3758:G:C4	2:A2:3759:G:C8	3.10	0.40
2:A2:3994:C:O3'	68:i2:37:GLY:HA3	2.22	0.40
2:A2:4123:U:H2'	2:A2:4124:G:C8	2.56	0.40
2:A2:4172:G:H2'	2:A2:4173:U:O4'	2.22	0.40
2:A2:4318:G:H2'	2:A2:4319:C:C6	2.57	0.40
2:A2:4389:G:H2'	2:A2:4390:C:H6	1.86	0.40
2:A2:4600:G:H2'	2:A2:4601:A:H8	1.84	0.40
4:B1:63:LEU:HD13	25:H1:32:GLN:HB2	2.03	0.40
6:B3:11:GLN:OE1	6:B3:11:GLN:HA	2.21	0.40
7:Bv:56:C:H2'	7:Bv:57:G:H8	1.87	0.40
7:Bz:25:C:H2'	7:Bz:26:A:H8	1.86	0.40
12:C3:25:THR:HG21	73:q2:7:LYS:HD2	2.04	0.40
15:D3:11:LEU:HD11	31:J3:136:HIS:HE1	1.86	0.40
17:E2:259:PRO:O	17:E2:261:ARG:N	2.54	0.40
17:E2:285:TYR:OH	17:E2:334:LYS:HD3	2.22	0.40
20:F2:259:LYS:HE2	20:F2:273:LEU:HD11	2.03	0.40
20:F2:336:ARG:HA	20:F2:339:THR:HG22	2.04	0.40
24:G3:51:ARG:NH2	75:s2:61:PHE:HB3	2.36	0.40
26:H2:291:HIS:HB3	59:Z2:42:TYR:CE1	2.57	0.40
34:L1:93:LEU:HB3	34:L1:100:VAL:HG22	2.03	0.40
41:O2:25:CYS:O	41:O2:29:VAL:HG13	2.21	0.40
44:P3:58:ALA:HA	83:m2:687:A:O2'	2.22	0.40
47:R2:93:ASN:O	47:R2:95:THR:HG23	2.22	0.40
74:r2:44:LEU:HD21	74:r2:70:ILE:HG21	2.03	0.40
80:x2:52:LYS:NZ	83:m2:1300:G:H21	2.20	0.40
80:x2:86:LEU:HB3	80:x2:88:GLU:OE1	2.21	0.40
82:z2:8:THR:HA	82:z2:11:LYS:HG2	2.04	0.40
83:m2:217:G:H2'	83:m2:218:C:C6	2.56	0.40
83:m2:820:A:H2'	83:m2:821:G:H8	1.86	0.40
83:m2:881:C:H2'	83:m2:882:G:H8	1.86	0.40
83:m2:972:G:H1'	83:m2:973:G:C1'	2.50	0.40

There are no symmetry-related clashes.

5.3 Torsion angles

5.3.1 Protein backbone

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	A1	221/270 (82%)	215 (97%)	6 (3%)	0	100	100
3	A3	138/152 (91%)	126 (91%)	12 (9%)	0	100	100
4	B1	220/266 (83%)	212 (96%)	8 (4%)	0	100	100
6	B3	139/145 (96%)	135 (97%)	4 (3%)	0	100	100
10	C1	188/192 (98%)	183 (97%)	5 (3%)	0	100	100
12	C3	100/119 (84%)	98 (98%)	2 (2%)	0	100	100
13	D1	200/214 (94%)	193 (96%)	7 (4%)	0	100	100
14	D2	249/257 (97%)	233 (94%)	16 (6%)	0	100	100
15	D3	81/83 (98%)	81 (100%)	0	0	100	100
16	E1	172/178 (97%)	160 (93%)	12 (7%)	0	100	100
17	E2	400/403 (99%)	382 (96%)	18 (4%)	0	100	100
18	E3	137/143 (96%)	133 (97%)	4 (3%)	0	100	100
19	F1	201/211 (95%)	193 (96%)	8 (4%)	0	100	100
20	F2	357/419 (85%)	345 (97%)	12 (3%)	0	100	100
21	F3	97/115 (84%)	95 (98%)	2 (2%)	0	100	100
22	G1	137/217 (63%)	133 (97%)	4 (3%)	0	100	100
23	G2	291/297 (98%)	288 (99%)	3 (1%)	0	100	100
24	G3	60/69 (87%)	57 (95%)	3 (5%)	0	100	100
25	H1	201/204 (98%)	192 (96%)	9 (4%)	0	100	100
26	H2	215/296 (73%)	207 (96%)	8 (4%)	0	100	100
27	H3	52/56 (93%)	45 (86%)	7 (14%)	0	100	100
28	I2	199/203 (98%)	194 (98%)	5 (2%)	0	100	100
29	I3	311/317 (98%)	295 (95%)	16 (5%)	0	100	100
30	J2	151/184 (82%)	146 (97%)	5 (3%)	0	100	100
31	J3	217/293 (74%)	211 (97%)	6 (3%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
32	K2	184/188 (98%)	176 (96%)	8 (4%)	0	100	100
33	K3	225/249 (90%)	222 (99%)	3 (1%)	0	100	100
34	L1	155/217 (71%)	146 (94%)	9 (6%)	0	100	100
35	L2	177/196 (90%)	176 (99%)	1 (1%)	0	100	100
36	L3	182/194 (94%)	177 (97%)	5 (3%)	0	100	100
37	M2	173/176 (98%)	162 (94%)	11 (6%)	0	100	100
38	M3	120/132 (91%)	117 (98%)	3 (2%)	0	100	100
39	N2	157/160 (98%)	154 (98%)	3 (2%)	0	100	100
40	N3	148/151 (98%)	147 (99%)	1 (1%)	0	100	100
41	O2	99/128 (77%)	95 (96%)	4 (4%)	0	100	100
42	O3	132/151 (87%)	128 (97%)	4 (3%)	0	100	100
43	P2	127/140 (91%)	121 (95%)	5 (4%)	1 (1%)	16	47
44	P3	127/130 (98%)	121 (95%)	6 (5%)	0	100	100
45	Q2	60/157 (38%)	57 (95%)	3 (5%)	0	100	100
46	Q3	120/133 (90%)	116 (97%)	4 (3%)	0	100	100
47	R2	116/156 (74%)	114 (98%)	2 (2%)	0	100	100
48	R3	70/125 (56%)	69 (99%)	1 (1%)	0	100	100
49	S2	132/145 (91%)	130 (98%)	2 (2%)	0	100	100
50	S3	81/84 (96%)	76 (94%)	4 (5%)	1 (1%)	10	38
51	T2	133/136 (98%)	127 (96%)	6 (4%)	0	100	100
52	T3	53/133 (40%)	52 (98%)	1 (2%)	0	100	100
53	U2	145/148 (98%)	137 (94%)	8 (6%)	0	100	100
54	U3	50/156 (32%)	45 (90%)	5 (10%)	0	100	100
55	V2	115/160 (72%)	112 (97%)	3 (3%)	0	100	100
56	W2	92/115 (80%)	89 (97%)	3 (3%)	0	100	100
57	X2	105/125 (84%)	101 (96%)	4 (4%)	0	100	100
58	Y2	126/135 (93%)	124 (98%)	2 (2%)	0	100	100
59	Z2	107/110 (97%)	102 (95%)	5 (5%)	0	100	100
60	a2	112/117 (96%)	108 (96%)	4 (4%)	0	100	100
61	b2	118/123 (96%)	117 (99%)	1 (1%)	0	100	100
62	c2	100/105 (95%)	99 (99%)	1 (1%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
63	d2	84/97 (87%)	83 (99%)	1 (1%)	0	100	100
64	e2	67/70 (96%)	67 (100%)	0	0	100	100
65	f2	48/51 (94%)	48 (100%)	0	0	100	100
66	g2	50/128 (39%)	48 (96%)	2 (4%)	0	100	100
67	h2	22/25 (88%)	22 (100%)	0	0	100	100
68	i2	101/106 (95%)	94 (93%)	7 (7%)	0	100	100
69	j2	87/92 (95%)	81 (93%)	6 (7%)	0	100	100
70	k2	123/137 (90%)	118 (96%)	5 (4%)	0	100	100
71	o2	212/295 (72%)	205 (97%)	7 (3%)	0	100	100
72	p2	210/264 (80%)	188 (90%)	22 (10%)	0	100	100
73	q2	218/243 (90%)	209 (96%)	9 (4%)	0	100	100
74	r2	260/263 (99%)	252 (97%)	8 (3%)	0	100	100
75	s2	179/204 (88%)	175 (98%)	4 (2%)	0	100	100
76	t2	179/194 (92%)	166 (93%)	13 (7%)	0	100	100
77	u2	204/208 (98%)	192 (94%)	12 (6%)	0	100	100
78	v2	93/165 (56%)	92 (99%)	1 (1%)	0	100	100
79	w2	148/158 (94%)	137 (93%)	10 (7%)	1 (1%)	18	50
80	x2	130/145 (90%)	127 (98%)	3 (2%)	0	100	100
81	y2	140/146 (96%)	132 (94%)	8 (6%)	0	100	100
82	z2	67/135 (50%)	65 (97%)	2 (3%)	0	100	100
All	All	11197/13004 (86%)	10770 (96%)	424 (4%)	3 (0%)	100	100

All (3) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
43	P2	92	ASP
50	S3	75	GLU
79	w2	25	LEU

5.3.2 Protein sidechains ⓘ

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	A1	194/234 (83%)	194 (100%)	0	100	100
3	A3	121/132 (92%)	121 (100%)	0	100	100
4	B1	193/223 (86%)	193 (100%)	0	100	100
6	B3	112/115 (97%)	112 (100%)	0	100	100
10	C1	169/171 (99%)	169 (100%)	0	100	100
12	C3	93/107 (87%)	93 (100%)	0	100	100
13	D1	174/181 (96%)	174 (100%)	0	100	100
14	D2	193/199 (97%)	193 (100%)	0	100	100
15	D3	67/67 (100%)	67 (100%)	0	100	100
16	E1	147/149 (99%)	147 (100%)	0	100	100
17	E2	347/348 (100%)	347 (100%)	0	100	100
18	E3	111/115 (96%)	111 (100%)	0	100	100
19	F1	170/178 (96%)	170 (100%)	0	100	100
20	F2	301/348 (86%)	301 (100%)	0	100	100
21	F3	86/98 (88%)	86 (100%)	0	100	100
22	G1	118/157 (75%)	118 (100%)	0	100	100
23	G2	246/249 (99%)	246 (100%)	0	100	100
24	G3	55/62 (89%)	55 (100%)	0	100	100
25	H1	171/172 (99%)	171 (100%)	0	100	100
26	H2	198/256 (77%)	198 (100%)	0	100	100
27	H3	48/49 (98%)	48 (100%)	0	100	100
28	I2	172/173 (99%)	172 (100%)	0	100	100
29	I3	272/275 (99%)	272 (100%)	0	100	100
30	J2	134/163 (82%)	134 (100%)	0	100	100
31	J3	185/224 (83%)	185 (100%)	0	100	100
32	K2	164/165 (99%)	164 (100%)	0	100	100
33	K3	198/218 (91%)	198 (100%)	0	100	100
34	L1	147/197 (75%)	147 (100%)	0	100	100
35	L2	158/175 (90%)	158 (100%)	0	100	100
36	L3	160/168 (95%)	160 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
37	M2	155/156 (99%)	155 (100%)	0	100	100
38	M3	104/108 (96%)	104 (100%)	0	100	100
39	N2	139/140 (99%)	139 (100%)	0	100	100
40	N3	130/131 (99%)	130 (100%)	0	100	100
41	O2	91/114 (80%)	91 (100%)	0	100	100
42	O3	104/119 (87%)	103 (99%)	1 (1%)	68	75
43	P2	100/107 (94%)	100 (100%)	0	100	100
44	P3	112/113 (99%)	112 (100%)	0	100	100
45	Q2	54/126 (43%)	54 (100%)	0	100	100
46	Q3	107/115 (93%)	107 (100%)	0	100	100
47	R2	106/133 (80%)	106 (100%)	0	100	100
48	R3	63/103 (61%)	63 (100%)	0	100	100
49	S2	124/135 (92%)	124 (100%)	0	100	100
50	S3	75/76 (99%)	75 (100%)	0	100	100
51	T2	117/118 (99%)	117 (100%)	0	100	100
52	T3	45/106 (42%)	45 (100%)	0	100	100
53	U2	120/121 (99%)	120 (100%)	0	100	100
54	U3	45/140 (32%)	45 (100%)	0	100	100
55	V2	98/124 (79%)	98 (100%)	0	100	100
56	W2	79/97 (81%)	79 (100%)	0	100	100
57	X2	98/110 (89%)	98 (100%)	0	100	100
58	Y2	114/121 (94%)	114 (100%)	0	100	100
59	Z2	88/89 (99%)	88 (100%)	0	100	100
60	a2	98/100 (98%)	98 (100%)	0	100	100
61	b2	108/110 (98%)	108 (100%)	0	100	100
62	c2	86/89 (97%)	86 (100%)	0	100	100
63	d2	73/80 (91%)	73 (100%)	0	100	100
64	e2	64/65 (98%)	64 (100%)	0	100	100
65	f2	47/48 (98%)	47 (100%)	0	100	100
66	g2	48/116 (41%)	48 (100%)	0	100	100
67	h2	23/24 (96%)	23 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
68	i2	91/94 (97%)	91 (100%)	0	100	100
69	j2	73/75 (97%)	73 (100%)	0	100	100
70	k2	109/121 (90%)	109 (100%)	0	100	100
71	o2	179/242 (74%)	179 (100%)	0	100	100
72	p2	193/229 (84%)	192 (100%)	1 (0%)	81	81
73	q2	184/202 (91%)	184 (100%)	0	100	100
74	r2	224/225 (100%)	224 (100%)	0	100	100
75	s2	156/170 (92%)	156 (100%)	0	100	100
76	t2	110/174 (63%)	110 (100%)	0	100	100
77	u2	165/180 (92%)	165 (100%)	0	100	100
78	v2	86/136 (63%)	86 (100%)	0	100	100
79	w2	134/142 (94%)	134 (100%)	0	100	100
80	x2	118/130 (91%)	118 (100%)	0	100	100
81	y2	117/121 (97%)	117 (100%)	0	100	100
82	z2	60/121 (50%)	60 (100%)	0	100	100
All	All	9718/11064 (88%)	9716 (100%)	2 (0%)	100	100

All (2) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
42	O3	26	ASN
72	p2	75	GLN

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (97) such sidechains are listed below:

Mol	Chain	Res	Type
1	A1	132	GLN
1	A1	214	HIS
1	A1	257	ASN
3	A3	10	GLN
4	B1	66	GLN
4	B1	112	GLN
4	B1	195	HIS
4	B1	227	ASN
6	B3	83	GLN
6	B3	126	GLN

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Mol	Chain	Res	Type
10	C1	98	HIS
13	D1	144	ASN
14	D2	38	HIS
16	E1	104	ASN
17	E2	138	GLN
17	E2	245	HIS
18	E3	23	HIS
18	E3	31	HIS
18	E3	39	ASN
18	E3	73	GLN
19	F1	149	GLN
20	F2	21	ASN
20	F2	50	GLN
20	F2	142	HIS
23	G2	131	ASN
23	G2	138	GLN
23	G2	291	GLN
25	H1	91	GLN
26	H2	50	HIS
26	H2	108	ASN
28	I2	199	ASN
29	I3	15	ASN
29	I3	76	GLN
29	I3	117	ASN
29	I3	178	ASN
29	I3	311	GLN
30	J2	56	GLN
30	J2	118	GLN
30	J2	137	ASN
31	J3	113	GLN
31	J3	277	HIS
33	K3	81	HIS
33	K3	186	GLN
36	L3	113	GLN
36	L3	154	GLN
37	M2	125	GLN
37	M2	173	ASN
39	N2	127	GLN
40	N3	36	GLN
40	N3	105	ASN
40	N3	123	HIS
41	O2	50	ASN

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Mol	Chain	Res	Type
43	P2	77	HIS
44	P3	15	ASN
46	Q3	19	GLN
49	S2	18	HIS
49	S2	72	GLN
52	T3	58	ASN
53	U2	60	HIS
54	U3	135	HIS
55	V2	6	ASN
55	V2	79	GLN
55	V2	118	GLN
57	X2	34	HIS
57	X2	69	ASN
58	Y2	80	HIS
59	Z2	20	ASN
59	Z2	21	GLN
60	a2	71	GLN
60	a2	112	GLN
60	a2	114	GLN
62	c2	15	HIS
63	d2	66	HIS
65	f2	19	GLN
66	g2	84	GLN
68	i2	25	GLN
68	i2	36	GLN
70	k2	21	ASN
70	k2	36	ASN
71	o2	141	ASN
72	p2	75	GLN
72	p2	118	GLN
74	r2	157	ASN
75	s2	36	GLN
75	s2	179	ASN
75	s2	186	ASN
76	t2	44	ASN
77	u2	35	ASN
77	u2	52	ASN
77	u2	146	GLN
77	u2	181	GLN
78	v2	28	HIS
78	v2	39	ASN
78	v2	42	ASN

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Mol	Chain	Res	Type
78	v2	44	HIS
79	w2	83	GLN
81	y2	35	ASN

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
11	C2	155/156 (99%)	31 (20%)	1 (0%)
2	A2	3593/3615 (99%)	764 (21%)	11 (0%)
5	B2	118/121 (97%)	9 (7%)	0
7	Bv	75/76 (98%)	20 (26%)	0
7	Bz	75/76 (98%)	23 (30%)	0
7	n2	75/76 (98%)	33 (44%)	0
8	Bx	9/10 (90%)	4 (44%)	0
83	m2	1627/1635 (99%)	400 (24%)	0
All	All	5727/5765 (99%)	1284 (22%)	12 (0%)

All (1284) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
2	A2	2	G
2	A2	5	A
2	A2	21	G
2	A2	25	A
2	A2	39	A
2	A2	42	A
2	A2	48	G
2	A2	59	A
2	A2	64	A
2	A2	65	A
2	A2	73	A
2	A2	91	G
2	A2	98	A
2	A2	104	G
2	A2	108	A
2	A2	109	G
2	A2	110	C
2	A2	119	G
2	A2	120	A
2	A2	122	U
2	A2	131	C

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Mol	Chain	Res	Type
2	A2	134	G
2	A2	136	C
2	A2	143	U
2	A2	144	G
2	A2	159	C
2	A2	170	C
2	A2	172	C
2	A2	183	C
2	A2	184	U
2	A2	185	C
2	A2	186	G
2	A2	187	U
2	A2	188	G
2	A2	189	G
2	A2	197	A
2	A2	200	U
2	A2	201	C
2	A2	207	G
2	A2	209	U
2	A2	210	C
2	A2	214	G
2	A2	216	C
2	A2	217	C
2	A2	218	A
2	A2	220	C
2	A2	233	U
2	A2	234	G
2	A2	237	G
2	A2	254	G
2	A2	258	G
2	A2	266	C
2	A2	267	G
2	A2	269	G
2	A2	280	G
2	A2	281	U
2	A2	297	U
2	A2	315	G
2	A2	316	U
2	A2	340	C
2	A2	341	G
2	A2	344	A
2	A2	373	G

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Mol	Chain	Res	Type
2	A2	387	G
2	A2	396	A
2	A2	407	A
2	A2	408	A
2	A2	409	G
2	A2	410	A
2	A2	412	G
2	A2	413	G
2	A2	415	G
2	A2	432	U
2	A2	433	A
2	A2	440	U
2	A2	449	C
2	A2	450	G
2	A2	452	A
2	A2	453	G
2	A2	454	U
2	A2	455	C
2	A2	460	C
2	A2	462	G
2	A2	467	U
2	A2	468	U
2	A2	469	C
2	A2	470	A
2	A2	479	G
2	A2	486	C
2	A2	490	C
2	A2	497	G
2	A2	498	G
2	A2	499	U
2	A2	500	G
2	A2	510	A
2	A2	511	U
2	A2	513	U
2	A2	514	U
2	A2	515	U
2	A2	516	C
2	A2	517	C
2	A2	520	C
2	A2	651	C
2	A2	652	G
2	A2	653	G

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Mol	Chain	Res	Type
2	A2	654	G
2	A2	658	C
2	A2	668	C
2	A2	671	G
2	A2	674	A
2	A2	675	C
2	A2	681	G
2	A2	692	C
2	A2	693	A
2	A2	694	C
2	A2	699	A
2	A2	703	U
2	A2	704	G
2	A2	711	C
2	A2	715	G
2	A2	737	G
2	A2	738	G
2	A2	746	G
2	A2	747	G
2	A2	748	U
2	A2	749	G
2	A2	753	A
2	A2	766	G
2	A2	767	G
2	A2	801	C
2	A2	802	A
2	A2	805	C
2	A2	807	G
2	A2	811	G
2	A2	812	U
2	A2	813	U
2	A2	814	A
2	A2	816	A
2	A2	824	G
2	A2	839	C
2	A2	841	A
2	A2	843	U
2	A2	855	G
2	A2	857	A
2	A2	858	A
2	A2	869	C
2	A2	870	G

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Mol	Chain	Res	Type
2	A2	888	U
2	A2	891	C
2	A2	919	G
2	A2	921	G
2	A2	927	C
2	A2	934	C
2	A2	935	C
2	A2	936	C
2	A2	937	C
2	A2	938	U
2	A2	1000	G
2	A2	1001	G
2	A2	1002	G
2	A2	1003	G
2	A2	1004	C
2	A2	1005	G
2	A2	1006	G
2	A2	1010	G
2	A2	1012	C
2	A2	1013	C
2	A2	1014	C
2	A2	1015	C
2	A2	1021	G
2	A2	1022	C
2	A2	1024	C
2	A2	1025	C
2	A2	1031	U
2	A2	1032	C
2	A2	1038	G
2	A2	1039	C
2	A2	1043	C
2	A2	1044	G
2	A2	1047	U
2	A2	1048	C
2	A2	1049	C
2	A2	1053	G
2	A2	1054	G
2	A2	1055	G
2	A2	1070	C
2	A2	1071	G
2	A2	1072	U
2	A2	1079	A

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Mol	Chain	Res	Type
2	A2	1081	G
2	A2	1082	A
2	A2	1085	C
2	A2	1086	C
2	A2	1087	G
2	A2	1093	A
2	A2	1094	C
2	A2	1098	G
2	A2	1099	U
2	A2	1101	G
2	A2	1108	A
2	A2	1109	U
2	A2	1110	G
2	A2	1116	U
2	A2	1118	C
2	A2	1136	A
2	A2	1140	A2M
2	A2	1159	G
2	A2	1168	A
2	A2	1172	A
2	A2	1173	G
2	A2	1179	C
2	A2	1180	G
2	A2	1181	U
2	A2	1191	G
2	A2	1201	A
2	A2	1208	G
2	A2	1211	A
2	A2	1212	A
2	A2	1224	G
2	A2	1231	C
2	A2	1232	G
2	A2	1233	A
2	A2	1253	U
2	A2	1270	G
2	A2	1295	G
2	A2	1296	C
2	A2	1298	C
2	A2	1310	A
2	A2	1311	G
2	A2	1315	G
2	A2	1329	G

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Mol	Chain	Res	Type
2	A2	1338	A
2	A2	1347	A2M
2	A2	1360	A
2	A2	1361	G
2	A2	1376	A
2	A2	1377	A
2	A2	1378	A
2	A2	1379	C
2	A2	1387	G
2	A2	1391	U
2	A2	1404	U
2	A2	1409	U
2	A2	1425	G
2	A2	1437	G
2	A2	1438	OMG
2	A2	1444	A
2	A2	1446	G
2	A2	1447	A
2	A2	1451	A
2	A2	1453	C
2	A2	1454	G
2	A2	1467	G
2	A2	1474	C
2	A2	1489	C
2	A2	1490	PSU
2	A2	1511	C
2	A2	1512	U
2	A2	1522	C
2	A2	1536	G
2	A2	1544	A
2	A2	1558	U
2	A2	1559	U
2	A2	1560	G
2	A2	1573	U
2	A2	1589	A
2	A2	1599	G
2	A2	1606	A
2	A2	1608	G
2	A2	1613	G
2	A2	1617	G
2	A2	1621	G
2	A2	1622	C

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Mol	Chain	Res	Type
2	A2	1623	G
2	A2	1624	U
2	A2	1626	G
2	A2	1636	U
2	A2	1638	G
2	A2	1639	A
2	A2	1644	G
2	A2	1645	A
2	A2	1657	G
2	A2	1671	G
2	A2	1684	U
2	A2	1690	A
2	A2	1699	A
2	A2	1718	G
2	A2	1719	A
2	A2	1721	G
2	A2	1722	C
2	A2	1723	C
2	A2	1724	G
2	A2	1727	G
2	A2	1733	C
2	A2	1734	A
2	A2	1738	C
2	A2	1750	G
2	A2	1753	G
2	A2	1757	G
2	A2	1760	A
2	A2	1762	A
2	A2	1763	G
2	A2	1764	A
2	A2	1766	A
2	A2	1821	C
2	A2	1822	U
2	A2	1825	C
2	A2	1827	A
2	A2	1828	A
2	A2	1846	U
2	A2	1848	G
2	A2	1850	U
2	A2	1857	G
2	A2	1858	G
2	A2	1871	A

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Mol	Chain	Res	Type
2	A2	1886	C
2	A2	1887	G
2	A2	1893	C
2	A2	1894	G
2	A2	1895	G
2	A2	1897	A
2	A2	1899	G
2	A2	1900	G
2	A2	1902	A
2	A2	1904	G
2	A2	1905	G
2	A2	1906	G
2	A2	1909	G
2	A2	1910	G
2	A2	1911	G
2	A2	2005	U
2	A2	2006	C
2	A2	2009	G
2	A2	2012	C
2	A2	2014	G
2	A2	2015	C
2	A2	2025	G
2	A2	2044	C
2	A2	2052	G
2	A2	2055	A
2	A2	2056	G
2	A2	2061	G
2	A2	2065	C
2	A2	2068	A
2	A2	2069	G
2	A2	2071	G
2	A2	2077	G
2	A2	2086	G
2	A2	2087	A
2	A2	2088	G
2	A2	2094	G
2	A2	2100	G
2	A2	2103	G
2	A2	2106	OMC
2	A2	2115	A
2	A2	2138	C
2	A2	2150	A

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Mol	Chain	Res	Type
2	A2	2152	G
2	A2	2153	U
2	A2	2176	G
2	A2	2177	OMC
2	A2	2180	U
2	A2	2196	C
2	A2	2205	G
2	A2	2208	A
2	A2	2224	C
2	A2	2225	C
2	A2	2226	G
2	A2	2229	G
2	A2	2233	C
2	A2	2235	G
2	A2	2239	A
2	A2	2245	U
2	A2	2249	U
2	A2	2257	G
2	A2	2258	G
2	A2	2259	C
2	A2	2260	C
2	A2	2261	G
2	A2	2268	A
2	A2	2273	G
2	A2	2275	C
2	A2	2276	G
2	A2	2284	A
2	A2	2285	U
2	A2	2301	G
2	A2	2302	G
2	A2	2310	G
2	A2	2311	G
2	A2	2314	G
2	A2	2321	G
2	A2	2322	G
2	A2	2328	A
2	A2	2342	A
2	A2	2344	C
2	A2	2356	A
2	A2	2361	G
2	A2	2367	G
2	A2	2371	C

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Mol	Chain	Res	Type
2	A2	2373	G
2	A2	2382	C
2	A2	2383	U
2	A2	2397	A
2	A2	2408	C
2	A2	2413	G
2	A2	2416	U
2	A2	2417	G
2	A2	2424	C
2	A2	2428	G
2	A2	2431	A
2	A2	2442	U
2	A2	2449	G
2	A2	2450	A
2	A2	2451	A
2	A2	2458	G
2	A2	2462	U
2	A2	2463	U
2	A2	2464	C
2	A2	2466	G
2	A2	2468	C
2	A2	2476	G
2	A2	2479	G
2	A2	2493	C
2	A2	2494	C
2	A2	2498	A
2	A2	2509	G
2	A2	2511	G
2	A2	2514	G
2	A2	2518	U
2	A2	2519	A
2	A2	2524	U
2	A2	2525	C
2	A2	2542	A2M
2	A2	2543	U
2	A2	2545	U
2	A2	2549	C
2	A2	2554	G
2	A2	2561	A
2	A2	2569	C
2	A2	2581	U
2	A2	2582	G

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Mol	Chain	Res	Type
2	A2	2584	U
2	A2	2601	G
2	A2	2602	G
2	A2	2603	G
2	A2	2610	G
2	A2	2623	G
2	A2	2657	G
2	A2	3243	C
2	A2	3244	C
2	A2	3246	G
2	A2	3248	G
2	A2	3250	C
2	A2	3252	A
2	A2	3253	G
2	A2	3254	C
2	A2	3260	A
2	A2	3262	U
2	A2	3271	G
2	A2	3272	U
2	A2	3282	G
2	A2	3286	A
2	A2	3291	A
2	A2	3300	U
2	A2	3302	A
2	A2	3304	A
2	A2	3310	G
2	A2	3318	A
2	A2	3328	G
2	A2	3329	C
2	A2	3330	G
2	A2	3366	G
2	A2	3368	A
2	A2	3373	A
2	A2	3382	A
2	A2	3383	A
2	A2	3406	G
2	A2	3409	G
2	A2	3410	G
2	A2	3415	A
2	A2	3416	A
2	A2	3417	C
2	A2	3421	G

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Mol	Chain	Res	Type
2	A2	3424	U
2	A2	3426	U
2	A2	3428	U
2	A2	3429	U
2	A2	3430	A
2	A2	3431	A
2	A2	3432	G
2	A2	3433	G
2	A2	3440	A
2	A2	3441	A2M
2	A2	3442	U
2	A2	3465	G
2	A2	3466	C
2	A2	3467	G
2	A2	3468	C
2	A2	3470	U
2	A2	3473	A
2	A2	3475	G
2	A2	3478	U
2	A2	3494	U
2	A2	3495	G
2	A2	3496	U
2	A2	3523	A
2	A2	3533	A
2	A2	3534	C
2	A2	3535	G
2	A2	3548	U
2	A2	3553	G
2	A2	3557	A
2	A2	3562	A
2	A2	3563	G
2	A2	3564	A
2	A2	3571	U
2	A2	3572	G
2	A2	3586	U
2	A2	3594	G
2	A2	3595	G
2	A2	3599	A
2	A2	3600	G
2	A2	3603	A
2	A2	3604	C
2	A2	3606	U

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Mol	Chain	Res	Type
2	A2	3607	G
2	A2	3609	G
2	A2	3610	A
2	A2	3612	G
2	A2	3613	U
2	A2	3614	G
2	A2	3615	U
2	A2	3617	G
2	A2	3618	A
2	A2	3619	A
2	A2	3620	U
2	A2	3621	A
2	A2	3622	A
2	A2	3623	G
2	A2	3625	G
2	A2	3626	G
2	A2	3627	G
2	A2	3629	G
2	A2	3632	C
2	A2	3633	C
2	A2	3686	G
2	A2	3687	C
2	A2	3689	G
2	A2	3690	C
2	A2	3693	G
2	A2	3694	U
2	A2	3695	G
2	A2	3696	A
2	A2	3697	A
2	A2	3698	A
2	A2	3699	U
2	A2	3701	C
2	A2	3702	C
2	A2	3703	A
2	A2	3704	C
2	A2	3705	U
2	A2	3707	C
2	A2	3709	C
2	A2	3712	A
2	A2	3714	C
2	A2	3715	G
2	A2	3719	U

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Mol	Chain	Res	Type
2	A2	3726	G
2	A2	3727	A
2	A2	3734	G
2	A2	3736	G
2	A2	3738	C
2	A2	3740	G
2	A2	3742	G
2	A2	3743	G
2	A2	3747	G
2	A2	3752	C
2	A2	3760	C
2	A2	3765	G
2	A2	3766	C
2	A2	3768	U
2	A2	3769	C
2	A2	3772	G
2	A2	3777	A
2	A2	3781	G
2	A2	3782	U
2	A2	3784	C
2	A2	3786	U
2	A2	3787	C
2	A2	3814	C
2	A2	3815	U
2	A2	3822	A
2	A2	3823	C
2	A2	3835	G
2	A2	3836	G
2	A2	3843	G
2	A2	3855	A
2	A2	3874	G
2	A2	3877	G
2	A2	3881	U
2	A2	3885	A
2	A2	3895	C
2	A2	3903	A
2	A2	3906	G
2	A2	3909	A
2	A2	3910	C
2	A2	3917	U
2	A2	3920	A
2	A2	3924	G

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Mol	Chain	Res	Type
2	A2	3925	A
2	A2	3933	A
2	A2	3943	G
2	A2	3947	U
2	A2	3957	G
2	A2	3958	OMU
2	A2	3966	C
2	A2	3981	G
2	A2	3982	G
2	A2	3990	G
2	A2	3991	A
2	A2	4002	C
2	A2	4004	U
2	A2	4006	U
2	A2	4023	G
2	A2	4025	G
2	A2	4029	G
2	A2	4030	A
2	A2	4031	A
2	A2	4033	A
2	A2	4039	C
2	A2	4043	G
2	A2	4046	A
2	A2	4071	U
2	A2	4072	U
2	A2	4073	C
2	A2	4074	A
2	A2	4078	C
2	A2	4096	C
2	A2	4100	G
2	A2	4101	A
2	A2	4104	U
2	A2	4105	C
2	A2	4116	A
2	A2	4118	C
2	A2	4127	G
2	A2	4151	OMG
2	A2	4152	PSU
2	A2	4162	A
2	A2	4164	U
2	A2	4165	A
2	A2	4170	A

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Mol	Chain	Res	Type
2	A2	4171	C
2	A2	4176	G
2	A2	4177	C
2	A2	4184	U
2	A2	4200	A
2	A2	4201	G
2	A2	4206	G
2	A2	4209	U
2	A2	4212	C
2	A2	4219	G
2	A2	4224	U
2	A2	4225	G
2	A2	4227	G
2	A2	4242	A
2	A2	4252	G
2	A2	4272	OMU
2	A2	4276	A
2	A2	4278	A
2	A2	4285	G
2	A2	4287	A
2	A2	4288	PSU
2	A2	4289	OMG
2	A2	4296	G
2	A2	4304	G
2	A2	4308	A
2	A2	4309	U
2	A2	4322	C
2	A2	4324	A
2	A2	4329	U
2	A2	4331	G
2	A2	4347	C
2	A2	4352	A
2	A2	4359	A
2	A2	4360	A
2	A2	4361	U
2	A2	4382	C
2	A2	4385	C
2	A2	4386	A
2	A2	4393	A
2	A2	4394	G
2	A2	4395	G
2	A2	4396	A

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Mol	Chain	Res	Type
2	A2	4397	G
2	A2	4403	G
2	A2	4406	G
2	A2	4409	C
2	A2	4411	C
2	A2	4413	G
2	A2	4417	G
2	A2	4422	G
2	A2	4433	G
2	A2	4434	U
2	A2	4435	C
2	A2	4497	U
2	A2	4498	C
2	A2	4499	G
2	A2	4503	C
2	A2	4504	C
2	A2	4505	G
2	A2	4508	G
2	A2	4511	C
2	A2	4515	G
2	A2	4516	C
2	A2	4520	G
2	A2	4521	A
2	A2	4522	G
2	A2	4525	G
2	A2	4527	U
2	A2	4528	C
2	A2	4540	C
2	A2	4541	G
2	A2	4545	U
2	A2	4546	G
2	A2	4547	C
2	A2	4548	G
2	A2	4552	G
2	A2	4553	G
2	A2	4555	A
2	A2	4557	G
2	A2	4559	G
2	A2	4560	G
2	A2	4565	C
2	A2	4569	C
2	A2	4570	U

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Mol	Chain	Res	Type
2	A2	4571	C
2	A2	4582	U
2	A2	4583	U
2	A2	4587	G
2	A2	4589	A
2	A2	4590	C
2	A2	4591	G
2	A2	4597	G
2	A2	4608	C
2	A2	4610	C
2	A2	4622	U
2	A2	4625	A
2	A2	4631	U
2	A2	4634	U
2	A2	4636	C
2	A2	4637	U
2	A2	4641	U
2	A2	4659	C
2	A2	4663	G
2	A2	4668	U
2	A2	4680	A
2	A2	4687	G
2	A2	4696	C
2	A2	4700	C
2	A2	4701	G
2	A2	4707	A
2	A2	4708	G
2	A2	4710	G
2	A2	4715	U
5	B2	7	G
5	B2	11	A
5	B2	22	A
5	B2	33	U
5	B2	53	U
5	B2	64	G
5	B2	76	U
5	B2	100	A
5	B2	110	G
7	Bv	7	A
7	Bv	8	U
7	Bv	14	A
7	Bv	16	U

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Mol	Chain	Res	Type
7	Bv	17	C
7	Bv	18	G
7	Bv	19	G
7	Bv	21	A
7	Bv	24	G
7	Bv	27	G
7	Bv	42	C
7	Bv	46	G
7	Bv	47	U
7	Bv	48	C
7	Bv	53	G
7	Bv	58	A
7	Bv	62	C
7	Bv	69	G
7	Bv	74	C
7	Bv	76	A
8	Bx	43	U
8	Bx	46	U
8	Bx	49	U
8	Bx	51	U
7	Bz	2	C
7	Bz	7	A
7	Bz	14	A
7	Bz	17	C
7	Bz	18	G
7	Bz	19	G
7	Bz	27	G
7	Bz	36	A
7	Bz	42	C
7	Bz	43	C
7	Bz	45	U
7	Bz	46	G
7	Bz	47	U
7	Bz	48	C
7	Bz	51	U
7	Bz	53	G
7	Bz	58	A
7	Bz	65	G
7	Bz	66	U
7	Bz	68	C
7	Bz	71	G
7	Bz	74	C

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Mol	Chain	Res	Type
7	Bz	76	A
11	C2	16	G
11	C2	23	C
11	C2	25	G
11	C2	34	U
11	C2	35	C
11	C2	39	G
11	C2	51	U
11	C2	52	A
11	C2	59	A
11	C2	60	G
11	C2	63	U
11	C2	75	OMG
11	C2	82	A
11	C2	84	A
11	C2	85	U
11	C2	87	G
11	C2	88	A
11	C2	94	G
11	C2	103	A
11	C2	105	C
11	C2	110	U
11	C2	111	U
11	C2	112	G
11	C2	114	G
11	C2	123	U
11	C2	125	C
11	C2	126	C
11	C2	127	U
11	C2	150	C
11	C2	155	C
11	C2	156	U
7	n2	2	C
7	n2	5	G
7	n2	9	A
7	n2	10	G
7	n2	13	C
7	n2	15	G
7	n2	16	U
7	n2	17	C
7	n2	19	G
7	n2	20	U

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Mol	Chain	Res	Type
7	n2	21	A
7	n2	23	A
7	n2	24	G
7	n2	27	G
7	n2	30	G
7	n2	32	U
7	n2	33	U
7	n2	40	C
7	n2	41	C
7	n2	44	G
7	n2	46	G
7	n2	47	U
7	n2	49	C
7	n2	52	G
7	n2	54	U
7	n2	55	U
7	n2	57	G
7	n2	58	A
7	n2	59	U
7	n2	66	U
7	n2	71	G
7	n2	75	C
7	n2	76	A
83	m2	4	C
83	m2	13	C
83	m2	14	C
83	m2	33	G
83	m2	41	G
83	m2	44	U
83	m2	46	A
83	m2	49	C
83	m2	56	G
83	m2	59	U
83	m2	62	G
83	m2	66	G
83	m2	67	C
83	m2	68	A
83	m2	71	G
83	m2	72	C
83	m2	74	G
83	m2	76	U
83	m2	77	A

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Mol	Chain	Res	Type
83	m2	78	C
83	m2	79	A
83	m2	103	A
83	m2	113	G
83	m2	114	G
83	m2	115	U
83	m2	118	C
83	m2	126	G
83	m2	127	C
83	m2	129	C
83	m2	130	G
83	m2	142	C
83	m2	143	U
83	m2	144	U
83	m2	145	G
83	m2	146	G
83	m2	155	G
83	m2	157	U
83	m2	159	A
83	m2	160	U
83	m2	161	U
83	m2	162	C
83	m2	168	C
83	m2	170	A
83	m2	171	A
83	m2	172	OMU
83	m2	180	G
83	m2	181	A
83	m2	182	C
83	m2	189	U
83	m2	190	G
83	m2	193	C
83	m2	195	C
83	m2	198	U
83	m2	199	U
83	m2	200	C
83	m2	201	C
83	m2	202	C
83	m2	204	G
83	m2	207	G
83	m2	208	G
83	m2	209	G

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Mol	Chain	Res	Type
83	m2	210	G
83	m2	211	A
83	m2	212	U
83	m2	215	G
83	m2	217	G
83	m2	296	U
83	m2	297	C
83	m2	304	A
83	m2	306	C
83	m2	307	U
83	m2	308	C
83	m2	310	G
83	m2	311	G
83	m2	313	C
83	m2	314	G
83	m2	315	A
83	m2	316	U
83	m2	319	C
83	m2	321	C
83	m2	324	C
83	m2	332	G
83	m2	333	C
83	m2	334	G
83	m2	336	C
83	m2	337	G
83	m2	342	C
83	m2	345	A
83	m2	349	G
83	m2	362	A
83	m2	364	C
83	m2	365	A
83	m2	366	A
83	m2	370	U
83	m2	372	G
83	m2	387	G
83	m2	388	C
83	m2	394	A
83	m2	395	U
83	m2	401	C
83	m2	402	C
83	m2	411	C
83	m2	415	G

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Mol	Chain	Res	Type
83	m2	423	G
83	m2	439	G
83	m2	440	G
83	m2	442	G
83	m2	450	A
83	m2	451	A
83	m2	452	C
83	m2	454	G
83	m2	466	A
83	m2	467	A
83	m2	473	G
83	m2	474	C
83	m2	476	G
83	m2	484	G
83	m2	487	A
83	m2	489	U
83	m2	494	C
83	m2	495	A
83	m2	502	A
83	m2	504	C
83	m2	518	A
83	m2	524	A
83	m2	525	A
83	m2	527	A
83	m2	530	A
83	m2	532	U
83	m2	534	C
83	m2	557	A
83	m2	558	U
83	m2	559	U
83	m2	561	G
83	m2	565	G
83	m2	568	U
83	m2	570	C
83	m2	575	U
83	m2	576	A
83	m2	578	A2M
83	m2	585	A
83	m2	589	A
83	m2	591	G
83	m2	592	A
83	m2	593	U

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Mol	Chain	Res	Type
83	m2	599	G
83	m2	600	G
83	m2	602	G
83	m2	606	A
83	m2	608	G
83	m2	609	U
83	m2	616	C
83	m2	617	C
83	m2	619	G
83	m2	622	G
83	m2	625	G
83	m2	630	A
83	m2	633	U
83	m2	645	A
83	m2	657	A
83	m2	662	C
83	m2	670	A2M
83	m2	671	A
83	m2	673	A
83	m2	674	A
83	m2	675	G
83	m2	686	G
83	m2	803	U
83	m2	813	A
83	m2	823	G
83	m2	824	PSU
83	m2	832	A
83	m2	837	C
83	m2	839	A
83	m2	840	G
83	m2	841	C
83	m2	842	C
83	m2	843	G
83	m2	845	C
83	m2	849	A
83	m2	870	G
83	m2	871	A
83	m2	872	A
83	m2	875	G
83	m2	882	G
83	m2	888	A
83	m2	889	U

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Mol	Chain	Res	Type
83	m2	890	U
83	m2	891	U
83	m2	892	U
83	m2	893	G
83	m2	894	U
83	m2	898	U
83	m2	900	U
83	m2	901	U
83	m2	902	C
83	m2	906	A
83	m2	911	G
83	m2	912	G
83	m2	915	A
83	m2	916	U
83	m2	918	A
83	m2	919	U
83	m2	921	A
83	m2	922	A
83	m2	924	A
83	m2	935	G
83	m2	936	G
83	m2	965	A
83	m2	972	G
83	m2	973	G
83	m2	980	G
83	m2	992	A
83	m2	994	A
83	m2	999	A
83	m2	1001	G
83	m2	1004	U
83	m2	1019	U
83	m2	1024	U
83	m2	1025	A
83	m2	1029	A
83	m2	1030	A
83	m2	1047	U
83	m2	1063	U
83	m2	1064	A
83	m2	1085	A
83	m2	1087	C
83	m2	1091	G
83	m2	1095	A

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Mol	Chain	Res	Type
83	m2	1101	G
83	m2	1102	A
83	m2	1111	C
83	m2	1117	U
83	m2	1118	C
83	m2	1123	G
83	m2	1132	G
83	m2	1135	A
83	m2	1138	U
83	m2	1140	C
83	m2	1141	C
83	m2	1150	A
83	m2	1152	A
83	m2	1155	C
83	m2	1156	U
83	m2	1157	U
83	m2	1168	G
83	m2	1184	A
83	m2	1197	A
83	m2	1209	G
83	m2	1210	A
83	m2	1212	G
83	m2	1217	C
83	m2	1219	A
83	m2	1226	G
83	m2	1233	C
83	m2	1244	U
83	m2	1245	PSU
83	m2	1250	B8N
83	m2	1252	A
83	m2	1253	A
83	m2	1255	A
83	m2	1258	G
83	m2	1259	G
83	m2	1261	A
83	m2	1266	C
83	m2	1268	C
83	m2	1273	C
83	m2	1274	C
83	m2	1275	C
83	m2	1276	G
83	m2	1282	G

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Mol	Chain	Res	Type
83	m2	1284	A
83	m2	1285	C
83	m2	1287	G
83	m2	1288	G
83	m2	1302	U
83	m2	1304	G
83	m2	1305	C
83	m2	1307	C
83	m2	1310	U
83	m2	1316	U
83	m2	1322	G
83	m2	1324	G
83	m2	1335	U
83	m2	1343	C
83	m2	1346	A
83	m2	1351	G
83	m2	1352	U
83	m2	1353	G
83	m2	1360	U
83	m2	1366	U
83	m2	1373	U
83	m2	1374	U
83	m2	1377	G
83	m2	1380	A
83	m2	1383	G
83	m2	1384	A
83	m2	1397	C
83	m2	1399	U
83	m2	1400	G
83	m2	1403	A
83	m2	1406	U
83	m2	1408	G
83	m2	1412	C
83	m2	1415	G
83	m2	1416	A
83	m2	1417	C
83	m2	1418	C
83	m2	1419	C
83	m2	1420	C
83	m2	1421	C
83	m2	1423	A
83	m2	1424	G

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Mol	Chain	Res	Type
83	m2	1425	C
83	m2	1431	G
83	m2	1435	C
83	m2	1437	C
83	m2	1438	C
83	m2	1439	C
83	m2	1440	A
83	m2	1451	G
83	m2	1454	A
83	m2	1456	A
83	m2	1465	U
83	m2	1466	C
83	m2	1468	G
83	m2	1491	A
83	m2	1492	G
83	m2	1497	G
83	m2	1499	G
83	m2	1500	A
83	m2	1509	G
83	m2	1512	G
83	m2	1522	G
83	m2	1523	C
83	m2	1525	C
83	m2	1530	G
83	m2	1532	U
83	m2	1533	A
83	m2	1534	C
83	m2	1535	A
83	m2	1542	G
83	m2	1546	C
83	m2	1554	G
83	m2	1557	U
83	m2	1559	C
83	m2	1560	C
83	m2	1562	U
83	m2	1565	G
83	m2	1570	C
83	m2	1572	G
83	m2	1581	A
83	m2	1582	A
83	m2	1587	U
83	m2	1590	A

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Mol	Chain	Res	Type
83	m2	1600	G
83	m2	1603	A
83	m2	1605	G
83	m2	1606	G
83	m2	1608	G
83	m2	1623	U
83	m2	1625	A
83	m2	1627	U
83	m2	1634	G
83	m2	1636	A
83	m2	1641	G
83	m2	1648	C
83	m2	1649	A
83	m2	1650	G
83	m2	1656	G
83	m2	1664	U
83	m2	1665	A
83	m2	1667	G
83	m2	1685	C
83	m2	1698	C
83	m2	1700	C
83	m2	1702	C
83	m2	1724	G
83	m2	1746	G
83	m2	1750	G
83	m2	1787	C
83	m2	1788	U
83	m2	1789	G
83	m2	1807	G
83	m2	1821	A
83	m2	1825	A
83	m2	1826	A
83	m2	1827	A
83	m2	1828	G
83	m2	1831	G
83	m2	1833	A
83	m2	1840	U
83	m2	1851	G
83	m2	1853	A
83	m2	1854	C
83	m2	1863	G
83	m2	1864	G

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Mol	Chain	Res	Type
83	m2	1865	A
83	m2	1866	U
83	m2	1867	C

All (12) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
2	A2	236	G
2	A2	406	C
2	A2	1446	G
2	A2	2259	C
2	A2	2382	C
2	A2	2430	G
2	A2	2463	U
2	A2	3253	G
2	A2	4277	C
2	A2	4351	U
2	A2	4582	U
11	C2	59	A

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

108 non-standard protein/DNA/RNA residues are 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	$\# Z > 2$
83	OMC	m2	519	83	19,22,23	0.28	0	26,31,34	0.46	0
2	A2M	A2	1337	2	22,25,26	0.12	0	31,36,39	0.28	0
2	OMG	A2	4151	2	23,26,27	0.37	0	33,38,41	0.53	0
2	A2M	A2	398	2	22,25,26	0.10	0	31,36,39	0.29	0
2	OMC	A2	3497	2	19,22,23	0.33	0	26,31,34	0.42	0
2	PSU	A2	3385	2	18,21,22	0.55	0	22,30,33	0.57	0
2	OMG	A2	3880	2	23,26,27	0.33	0	33,38,41	0.51	0
2	A2M	A2	1673	2	22,25,26	0.11	0	31,36,39	0.50	0
83	A2M	m2	486	83	22,25,26	0.08	0	31,36,39	0.21	0

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
83	A2M	m2	670	84,83	22,25,26	0.09	0	31,36,39	0.34	0
2	A2M	A2	4175	2,84	22,25,26	0.11	0	31,36,39	0.38	0
83	OMG	m2	685	83	23,26,27	0.27	0	33,38,41	0.49	0
2	PSU	A2	3420	2	18,21,22	0.54	0	22,30,33	0.37	0
2	A2M	A2	3374	2	22,25,26	0.11	0	31,36,39	0.29	0
83	OMG	m2	646	83	23,26,27	0.27	0	33,38,41	0.35	0
2	OMC	A2	2559	2	19,22,23	0.32	0	26,31,34	0.43	0
2	A2M	A2	2118	2,84	22,25,26	0.10	0	31,36,39	0.28	0
2	OMU	A2	3958	2	19,22,23	0.35	0	26,31,34	0.46	0
2	OMG	A2	4044	2,84	23,26,27	0.39	0	33,38,41	0.39	0
83	OMG	m2	1330	83	23,26,27	0.31	0	33,38,41	0.34	0
2	PSU	A2	3945	2	18,21,22	0.52	0	22,30,33	0.59	0
2	PSU	A2	4183	2	18,21,22	0.52	0	22,30,33	0.43	0
2	OMC	A2	3543	2	19,22,23	0.33	0	26,31,34	0.45	0
2	OMG	A2	4146	2	23,26,27	0.34	0	33,38,41	0.43	0
2	A2M	A2	3481	2	22,25,26	0.09	0	31,36,39	0.48	0
83	A2M	m2	514	83	22,25,26	0.09	0	31,36,39	0.38	0
2	OMU	A2	4150	2	19,22,23	0.31	0	26,31,34	0.47	0
83	PSU	m2	614	83	18,21,22	0.50	0	22,30,33	0.62	1 (4%)
83	PSU	m2	1083	83	18,21,22	0.59	1 (5%)	22,30,33	0.67	0
2	OMG	A2	2179	2	23,26,27	0.31	0	33,38,41	0.32	0
2	OMC	A2	1683	2,84	19,22,23	0.32	0	26,31,34	0.54	0
2	A2M	A2	2542	2	22,25,26	0.09	0	31,36,39	0.35	0
83	PSU	m2	824	83	18,21,22	0.60	1 (5%)	22,30,33	0.67	1 (4%)
2	OMU	A2	2592	2	19,22,23	0.33	0	26,31,34	0.63	0
83	A2M	m2	1033	83	22,25,26	0.07	0	31,36,39	0.19	0
2	OMG	A2	1130	2	23,26,27	0.42	0	33,38,41	0.55	0
83	OMG	m2	869	83	23,26,27	0.27	0	33,38,41	0.34	0
2	1MA	A2	4067	2	21,25,26	0.33	0	31,37,40	0.47	0
2	OMG	A2	4289	2	23,26,27	0.36	0	33,38,41	0.38	0
2	PSU	A2	4280	2	18,21,22	0.54	0	22,30,33	0.56	0
2	OMG	A2	1438	2	23,26,27	0.27	0	33,38,41	0.40	0
2	OMC	A2	2106	2	19,22,23	0.32	0	26,31,34	0.38	0
2	OMC	A2	2177	2,84	19,22,23	0.32	0	26,31,34	0.44	0
83	OMU	m2	121	83	19,22,23	0.30	0	26,31,34	0.46	0
2	5MC	A2	3438	2,84	18,22,23	0.30	0	26,32,35	0.40	0
83	OMG	m2	511	84,83	23,26,27	0.29	0	33,38,41	0.40	0
2	5MC	A2	4099	2	18,22,23	0.36	0	26,32,35	0.56	0
2	OMG	A2	4275	2	23,26,27	0.39	0	33,38,41	0.56	0
2	A2M	A2	3441	2	22,25,26	0.10	0	31,36,39	0.36	0
83	A2M	m2	578	83	22,25,26	0.08	0	31,36,39	0.27	0
2	A2M	A2	1140	2,84	22,25,26	0.09	0	31,36,39	0.19	0

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
83	OMG	m2	438	83	23,26,27	0.28	0	33,38,41	0.44	0
2	OMC	A2	3464	2	19,22,23	0.32	0	26,31,34	0.38	0
83	PSU	m2	825	83	18,21,22	0.49	0	22,30,33	0.54	0
2	PSU	A2	1490	2	18,21,22	0.91	1 (5%)	22,30,33	0.60	0
2	A2M	A2	3380	2	22,25,26	0.12	0	31,36,39	0.44	0
2	OMC	A2	3525	2	19,22,23	0.33	0	26,31,34	0.39	0
2	PSU	A2	1395	2	18,21,22	0.58	0	22,30,33	0.57	0
2	OMG	A2	3555	2,84	23,26,27	0.39	0	33,38,41	0.54	0
83	B8T	m2	1339	83	19,22,23	0.38	0	26,31,34	0.36	0
2	OMU	A2	4272	2	19,22,23	0.39	0	26,31,34	0.67	0
2	A2M	A2	1347	2,84	22,25,26	0.11	0	31,36,39	0.56	1 (3%)
83	OMU	m2	430	83	19,22,23	0.25	0	26,31,34	0.44	0
2	OMU	A2	3474	2	19,22,23	0.36	0	26,31,34	0.39	0
83	A2M	m2	1680	83	22,25,26	0.08	0	31,36,39	0.16	0
83	PSU	m2	1245	83	18,21,22	0.54	0	22,30,33	0.56	0
2	OMG	A2	2119	2,84	23,26,27	0.33	0	33,38,41	0.43	0
11	OMG	C2	75	11	23,26,27	0.31	0	33,38,41	0.37	0
2	OMU	A2	3581	2	19,22,23	0.33	0	26,31,34	0.46	0
83	OMC	m2	355	83	19,22,23	0.28	0	26,31,34	0.41	0
83	OMU	m2	172	83	19,22,23	0.26	0	26,31,34	0.64	1 (3%)
2	OMG	A2	1335	2	23,26,27	0.35	0	33,38,41	0.45	0
2	PSU	A2	4152	2	18,21,22	0.53	0	22,30,33	0.65	0
2	PSU	A2	4094	2	18,21,22	0.59	1 (5%)	22,30,33	0.64	1 (4%)
2	OMC	A2	4108	2	19,22,23	0.33	0	26,31,34	0.34	0
83	4AC	m2	1844	83	21,24,25	0.31	0	29,34,37	0.30	0
2	A2M	A2	4223	2	22,25,26	0.11	0	31,36,39	0.26	0
83	OMC	m2	174	83	19,22,23	0.34	0	26,31,34	0.56	0
2	PSU	A2	4288	2	18,21,22	0.51	0	22,30,33	0.62	0
2	A2M	A2	3486	2	22,25,26	0.09	0	31,36,39	0.42	0
2	OMC	A2	4188	2	19,22,23	0.34	0	26,31,34	0.50	0
2	OMG	A2	3448	2	23,26,27	0.34	0	33,38,41	0.34	0
2	A2M	A2	2156	2	22,25,26	0.10	0	31,36,39	0.22	0
2	OMC	A2	1154	2	19,22,23	0.33	0	26,31,34	0.46	0
83	A2M	m2	99	84,83	22,25,26	0.07	0	31,36,39	0.21	0
2	PSU	A2	2263	2	18,21,22	0.53	0	22,30,33	0.59	0
2	OMC	A2	2616	2	19,22,23	0.31	0	26,31,34	0.37	0
83	A2M	m2	27	83	22,25,26	0.12	0	31,36,39	0.32	0
2	OMC	A2	2579	2	19,22,23	0.29	0	26,31,34	0.39	0
2	OMC	A2	2120	2	19,22,23	0.32	0	26,31,34	0.48	0
2	PSU	A2	1496	2	18,21,22	0.53	0	22,30,33	0.57	0
2	A2M	A2	4270	2	22,25,26	0.11	0	31,36,39	0.31	0
2	A2M	A2	1137	2	22,25,26	0.09	0	31,36,39	0.23	0

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
83	UR3	m2	1832	83	19,22,23	0.28	0	26,32,35	0.43	0
2	PSU	A2	4055	2	18,21,22	0.54	0	22,30,33	0.63	1 (4%)
2	A2M	A2	2570	2	22,25,26	0.10	0	31,36,39	0.20	0
2	PSU	A2	3371	2	18,21,22	0.56	0	22,30,33	0.63	1 (4%)
2	OMG	A2	4022	2	23,26,27	0.31	0	33,38,41	0.39	0
2	OMG	A2	3848	7,2	23,26,27	0.35	0	33,38,41	0.34	0
2	PSU	A2	4102	2,84	18,21,22	0.53	0	22,30,33	0.41	0
83	OMU	m2	116	83	19,22,23	0.26	0	26,31,34	0.39	0
83	OMG	m2	603	83	23,26,27	0.27	0	33,38,41	0.33	0
2	OMG	A2	3283	2	23,26,27	0.34	0	33,38,41	0.48	0
83	OMC	m2	1705	83	19,22,23	0.27	0	26,31,34	0.36	0
2	2MG	A2	1330	2	23,26,27	0.31	0	32,38,41	0.35	0
83	B8N	m2	1250	83	24,29,30	0.58	0	29,42,45	0.63	0
2	OMG	A2	3400	2	23,26,27	0.31	0	33,38,41	0.44	0
2	OMC	A2	3357	2	19,22,23	0.25	0	26,31,34	0.42	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
83	OMC	m2	519	83	-	2/9/27/28	0/2/2/2
2	A2M	A2	1337	2	-	1/9/27/28	0/3/3/3
2	OMG	A2	4151	2	-	0/9/27/28	0/3/3/3
2	A2M	A2	398	2	-	0/9/27/28	0/3/3/3
2	OMC	A2	3497	2	-	0/9/27/28	0/2/2/2
2	PSU	A2	3385	2	-	0/7/25/26	0/2/2/2
2	OMG	A2	3880	2	-	0/9/27/28	0/3/3/3
2	A2M	A2	1673	2	-	0/9/27/28	0/3/3/3
83	A2M	m2	486	83	-	0/9/27/28	0/3/3/3
83	A2M	m2	670	84,83	-	2/9/27/28	0/3/3/3
2	A2M	A2	4175	2,84	-	1/9/27/28	0/3/3/3
83	OMG	m2	685	83	-	0/9/27/28	0/3/3/3
2	PSU	A2	3420	2	-	2/7/25/26	0/2/2/2
2	A2M	A2	3374	2	-	0/9/27/28	0/3/3/3
83	OMG	m2	646	83	-	1/9/27/28	0/3/3/3
2	OMC	A2	2559	2	-	0/9/27/28	0/2/2/2
2	A2M	A2	2118	2,84	-	1/9/27/28	0/3/3/3
2	OMU	A2	3958	2	-	0/9/27/28	0/2/2/2
2	OMG	A2	4044	2,84	-	0/9/27/28	0/3/3/3

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
83	OMG	m2	1330	83	-	2/9/27/28	0/3/3/3
2	PSU	A2	3945	2	-	0/7/25/26	0/2/2/2
2	PSU	A2	4183	2	-	2/7/25/26	0/2/2/2
2	OMC	A2	3543	2	-	0/9/27/28	0/2/2/2
2	OMG	A2	4146	2	-	1/9/27/28	0/3/3/3
2	A2M	A2	3481	2	-	0/9/27/28	0/3/3/3
83	A2M	m2	514	83	-	0/9/27/28	0/3/3/3
2	OMU	A2	4150	2	-	0/9/27/28	0/2/2/2
83	PSU	m2	614	83	-	0/7/25/26	0/2/2/2
83	PSU	m2	1083	83	-	1/7/25/26	0/2/2/2
2	OMG	A2	2179	2	-	1/9/27/28	0/3/3/3
2	OMC	A2	1683	2,84	-	0/9/27/28	0/2/2/2
2	A2M	A2	2542	2	-	3/9/27/28	0/3/3/3
83	PSU	m2	824	83	-	0/7/25/26	0/2/2/2
2	OMU	A2	2592	2	-	0/9/27/28	0/2/2/2
83	A2M	m2	1033	83	-	1/9/27/28	0/3/3/3
2	OMG	A2	1130	2	-	0/9/27/28	0/3/3/3
83	OMG	m2	869	83	-	1/9/27/28	0/3/3/3
2	1MA	A2	4067	2	-	2/7/25/26	0/3/3/3
2	OMG	A2	4289	2	-	3/9/27/28	0/3/3/3
2	PSU	A2	4280	2	-	0/7/25/26	0/2/2/2
2	OMG	A2	1438	2	-	2/9/27/28	0/3/3/3
2	OMC	A2	2106	2	-	2/9/27/28	0/2/2/2
2	OMC	A2	2177	2,84	-	1/9/27/28	0/2/2/2
83	OMU	m2	121	83	-	0/9/27/28	0/2/2/2
2	5MC	A2	3438	2,84	-	0/7/25/26	0/2/2/2
83	OMG	m2	511	84,83	-	0/9/27/28	0/3/3/3
2	5MC	A2	4099	2	-	4/7/25/26	0/2/2/2
2	OMG	A2	4275	2	-	0/9/27/28	0/3/3/3
2	A2M	A2	3441	2	-	4/9/27/28	0/3/3/3
83	A2M	m2	578	83	-	1/9/27/28	0/3/3/3
2	A2M	A2	1140	2,84	-	3/9/27/28	0/3/3/3
83	OMG	m2	438	83	-	0/9/27/28	0/3/3/3
2	OMC	A2	3464	2	-	0/9/27/28	0/2/2/2
83	PSU	m2	825	83	-	0/7/25/26	0/2/2/2
2	PSU	A2	1490	2	-	0/7/25/26	0/2/2/2
2	A2M	A2	3380	2	-	1/9/27/28	0/3/3/3
2	OMC	A2	3525	2	-	1/9/27/28	0/2/2/2
2	PSU	A2	1395	2	-	0/7/25/26	0/2/2/2
2	OMG	A2	3555	2,84	-	0/9/27/28	0/3/3/3

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
83	B8T	m2	1339	83	-	0/7/27/28	0/2/2/2
2	OMU	A2	4272	2	-	3/9/27/28	0/2/2/2
2	A2M	A2	1347	2,84	-	3/9/27/28	0/3/3/3
83	OMU	m2	430	83	-	4/9/27/28	0/2/2/2
2	OMU	A2	3474	2	-	2/9/27/28	0/2/2/2
83	A2M	m2	1680	83	-	1/9/27/28	0/3/3/3
83	PSU	m2	1245	83	-	1/7/25/26	0/2/2/2
2	OMG	A2	2119	2,84	-	1/9/27/28	0/3/3/3
11	OMG	C2	75	11	-	3/9/27/28	0/3/3/3
2	OMU	A2	3581	2	-	0/9/27/28	0/2/2/2
83	OMC	m2	355	83	-	1/9/27/28	0/2/2/2
83	OMU	m2	172	83	-	3/9/27/28	0/2/2/2
2	OMG	A2	1335	2	-	0/9/27/28	0/3/3/3
2	PSU	A2	4152	2	-	1/7/25/26	0/2/2/2
2	PSU	A2	4094	2	-	0/7/25/26	0/2/2/2
2	OMC	A2	4108	2	-	0/9/27/28	0/2/2/2
83	4AC	m2	1844	83	-	0/11/29/30	0/2/2/2
2	A2M	A2	4223	2	-	0/9/27/28	0/3/3/3
83	OMC	m2	174	83	-	2/9/27/28	0/2/2/2
2	PSU	A2	4288	2	-	3/7/25/26	0/2/2/2
2	A2M	A2	3486	2	-	0/9/27/28	0/3/3/3
2	OMC	A2	4188	2	-	0/9/27/28	0/2/2/2
2	OMG	A2	3448	2	-	0/9/27/28	0/3/3/3
2	A2M	A2	2156	2	-	1/9/27/28	0/3/3/3
2	OMC	A2	1154	2	-	1/9/27/28	0/2/2/2
83	A2M	m2	99	84,83	-	1/9/27/28	0/3/3/3
2	PSU	A2	2263	2	-	0/7/25/26	0/2/2/2
2	OMC	A2	2616	2	-	0/9/27/28	0/2/2/2
83	A2M	m2	27	83	-	2/9/27/28	0/3/3/3
2	OMC	A2	2579	2	-	1/9/27/28	0/2/2/2
2	OMC	A2	2120	2	-	0/9/27/28	0/2/2/2
2	PSU	A2	1496	2	-	0/7/25/26	0/2/2/2
2	A2M	A2	4270	2	-	0/9/27/28	0/3/3/3
2	A2M	A2	1137	2	-	0/9/27/28	0/3/3/3
83	UR3	m2	1832	83	-	2/7/25/26	0/2/2/2
2	PSU	A2	4055	2	-	0/7/25/26	0/2/2/2
2	A2M	A2	2570	2	-	0/9/27/28	0/3/3/3
2	PSU	A2	3371	2	-	0/7/25/26	0/2/2/2
2	OMG	A2	4022	2	-	1/9/27/28	0/3/3/3
2	OMG	A2	3848	7,2	-	1/9/27/28	0/3/3/3

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
2	PSU	A2	4102	2,84	-	3/7/25/26	0/2/2/2
83	OMU	m2	116	83	-	1/9/27/28	0/2/2/2
83	OMG	m2	603	83	-	1/9/27/28	0/3/3/3
2	OMG	A2	3283	2	-	0/9/27/28	0/3/3/3
83	OMC	m2	1705	83	-	1/9/27/28	0/2/2/2
2	2MG	A2	1330	2	-	0/9/27/28	0/3/3/3
83	B8N	m2	1250	83	-	3/16/34/35	0/2/2/2
2	OMG	A2	3400	2	-	0/9/27/28	0/3/3/3
2	OMC	A2	3357	2	-	4/9/27/28	0/2/2/2

All (4) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
2	A2	1490	PSU	O4'-C1'	-3.40	1.39	1.43
83	m2	824	PSU	O4'-C1'	-2.17	1.40	1.43
83	m2	1083	PSU	O4'-C1'	-2.15	1.40	1.43
2	A2	4094	PSU	O4'-C1'	-2.03	1.41	1.43

All (7) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
83	m2	172	OMU	O2'-C2'-C1'	2.48	113.92	109.08
83	m2	614	PSU	O4'-C1'-C2'	2.21	108.26	105.14
2	A2	1347	A2M	C3'-C2'-C1'	-2.20	98.75	102.89
2	A2	3371	PSU	O4'-C1'-C2'	2.10	108.10	105.14
2	A2	4094	PSU	O4'-C1'-C2'	2.10	108.10	105.14
2	A2	4055	PSU	O4'-C1'-C2'	2.07	108.07	105.14
83	m2	824	PSU	O4'-C1'-C2'	2.05	108.03	105.14

There are no chirality outliers.

All (97) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
11	C2	75	OMG	C1'-C2'-O2'-CM2
2	A2	1140	A2M	O4'-C4'-C5'-O5'
2	A2	1140	A2M	C3'-C4'-C5'-O5'
2	A2	1154	OMC	C1'-C2'-O2'-CM2
2	A2	1347	A2M	C4'-C5'-O5'-P
2	A2	1347	A2M	O4'-C4'-C5'-O5'
2	A2	1347	A2M	C3'-C4'-C5'-O5'

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Mol	Chain	Res	Type	Atoms
2	A2	1438	OMG	O4'-C4'-C5'-O5'
2	A2	1438	OMG	C3'-C4'-C5'-O5'
2	A2	2106	OMC	C1'-C2'-O2'-CM2
2	A2	2119	OMG	C1'-C2'-O2'-CM2
2	A2	2179	OMG	C1'-C2'-O2'-CM2
2	A2	2579	OMC	C1'-C2'-O2'-CM2
2	A2	3357	OMC	C2'-C1'-N1-C2
2	A2	3357	OMC	C2'-C1'-N1-C6
2	A2	3420	PSU	C2'-C1'-C5-C4
2	A2	3848	OMG	C1'-C2'-O2'-CM2
2	A2	4102	PSU	C2'-C1'-C5-C4
2	A2	4175	A2M	C1'-C2'-O2'-CM'
2	A2	4183	PSU	O4'-C1'-C5-C4
2	A2	4183	PSU	O4'-C1'-C5-C6
2	A2	4272	OMU	C1'-C2'-O2'-CM2
2	A2	4272	OMU	O4'-C4'-C5'-O5'
2	A2	4288	PSU	C3'-C4'-C5'-O5'
2	A2	4289	OMG	C1'-C2'-O2'-CM2
83	m2	116	OMU	C1'-C2'-O2'-CM2
83	m2	172	OMU	C1'-C2'-O2'-CM2
83	m2	174	OMC	O4'-C1'-N1-C2
83	m2	174	OMC	O4'-C1'-N1-C6
83	m2	430	OMU	C2'-C1'-N1-C2
83	m2	430	OMU	C2'-C1'-N1-C6
83	m2	603	OMG	C1'-C2'-O2'-CM2
83	m2	869	OMG	C1'-C2'-O2'-CM2
83	m2	1033	A2M	C1'-C2'-O2'-CM'
83	m2	1250	B8N	C3'-C4'-C5'-O5'
83	m2	1250	B8N	N3-C31-C32-C33
83	m2	1832	UR3	O4'-C1'-N1-C2
2	A2	3441	A2M	O4'-C4'-C5'-O5'
2	A2	4272	OMU	C3'-C4'-C5'-O5'
2	A2	4289	OMG	O4'-C4'-C5'-O5'
83	m2	27	A2M	C3'-C4'-C5'-O5'
83	m2	1250	B8N	O4'-C4'-C5'-O5'
83	m2	1832	UR3	O4'-C1'-N1-C6
2	A2	4288	PSU	O4'-C4'-C5'-O5'
2	A2	4289	OMG	C3'-C4'-C5'-O5'
83	m2	27	A2M	O4'-C4'-C5'-O5'
83	m2	1330	OMG	C3'-C4'-C5'-O5'
2	A2	4099	5MC	C2'-C1'-N1-C6
2	A2	4152	PSU	C4'-C5'-O5'-P

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Mol	Chain	Res	Type	Atoms
11	C2	75	OMG	O4'-C4'-C5'-O5'
83	m2	519	OMC	O4'-C4'-C5'-O5'
83	m2	1330	OMG	O4'-C4'-C5'-O5'
83	m2	519	OMC	C3'-C4'-C5'-O5'
2	A2	3441	A2M	C2'-C1'-N9-C8
2	A2	4067	1MA	O4'-C4'-C5'-O5'
83	m2	670	A2M	O4'-C4'-C5'-O5'
2	A2	4099	5MC	O4'-C1'-N1-C6
83	m2	578	A2M	C4'-C5'-O5'-P
2	A2	4067	1MA	C3'-C4'-C5'-O5'
83	m2	99	A2M	C1'-C2'-O2'-CM'
83	m2	1680	A2M	C1'-C2'-O2'-CM'
2	A2	3441	A2M	C3'-C4'-C5'-O5'
2	A2	2542	A2M	C2'-C1'-N9-C8
2	A2	2118	A2M	C3'-C2'-O2'-CM'
2	A2	4022	OMG	C3'-C2'-O2'-CM2
2	A2	4146	OMG	C3'-C2'-O2'-CM2
2	A2	4099	5MC	C2'-C1'-N1-C2
2	A2	3474	OMU	C4'-C5'-O5'-P
2	A2	4099	5MC	O4'-C1'-N1-C2
83	m2	646	OMG	C4'-C5'-O5'-P
83	m2	172	OMU	C4'-C5'-O5'-P
2	A2	3357	OMC	O4'-C1'-N1-C6
83	m2	430	OMU	O4'-C1'-N1-C6
2	A2	3474	OMU	C3'-C2'-O2'-CM2
83	m2	1083	PSU	C4'-C5'-O5'-P
83	m2	1245	PSU	C4'-C5'-O5'-P
2	A2	2542	A2M	C2'-C1'-N9-C4
2	A2	3357	OMC	O4'-C1'-N1-C2
11	C2	75	OMG	C3'-C4'-C5'-O5'
83	m2	670	A2M	C3'-C4'-C5'-O5'
2	A2	3420	PSU	O4'-C1'-C5-C4
2	A2	4102	PSU	O4'-C1'-C5-C4
2	A2	1140	A2M	C4'-C5'-O5'-P
83	m2	430	OMU	O4'-C1'-N1-C2
2	A2	1337	A2M	O4'-C1'-N9-C8
2	A2	2542	A2M	O4'-C1'-N9-C8
2	A2	3441	A2M	O4'-C1'-N9-C8
83	m2	1705	OMC	O4'-C4'-C5'-O5'
83	m2	355	OMC	C1'-C2'-O2'-CM2
2	A2	2106	OMC	O4'-C4'-C5'-O5'
2	A2	4102	PSU	O4'-C1'-C5-C6

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Mol	Chain	Res	Type	Atoms
2	A2	4288	PSU	O4'-C1'-C5-C6
2	A2	2177	OMC	O4'-C4'-C5'-O5'
2	A2	2156	A2M	C3'-C2'-O2'-CM'
2	A2	3380	A2M	C3'-C2'-O2'-CM'
2	A2	3525	OMC	C3'-C2'-O2'-CM2
83	m2	172	OMU	O4'-C4'-C5'-O5'

There are no ring outliers.

58 monomers are involved in 86 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
83	m2	519	OMC	1	0
2	A2	1337	A2M	1	0
2	A2	398	A2M	1	0
2	A2	3385	PSU	1	0
2	A2	3880	OMG	1	0
2	A2	1673	A2M	1	0
83	m2	486	A2M	2	0
83	m2	685	OMG	1	0
2	A2	3374	A2M	1	0
2	A2	2559	OMC	1	0
2	A2	2118	A2M	2	0
2	A2	3958	OMU	3	0
2	A2	4044	OMG	2	0
83	m2	1330	OMG	1	0
2	A2	4146	OMG	1	0
83	m2	514	A2M	1	0
2	A2	2179	OMG	1	0
83	m2	1033	A2M	2	0
83	m2	869	OMG	1	0
2	A2	4067	1MA	1	0
2	A2	4289	OMG	2	0
2	A2	1438	OMG	1	0
2	A2	2106	OMC	2	0
83	m2	511	OMG	4	0
2	A2	4099	5MC	2	0
2	A2	3441	A2M	1	0
83	m2	578	A2M	1	0
2	A2	1140	A2M	1	0
2	A2	1490	PSU	1	0
2	A2	3380	A2M	1	0
2	A2	3525	OMC	1	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
2	A2	3555	OMG	1	0
83	m2	1339	B8T	1	0
2	A2	4272	OMU	3	0
2	A2	1347	A2M	2	0
2	A2	3474	OMU	1	0
83	m2	1680	A2M	2	0
2	A2	2119	OMG	2	0
11	C2	75	OMG	1	0
2	A2	3581	OMU	1	0
83	m2	355	OMC	2	0
83	m2	172	OMU	1	0
2	A2	4094	PSU	1	0
83	m2	1844	4AC	4	0
83	m2	174	OMC	1	0
2	A2	4188	OMC	2	0
2	A2	1154	OMC	3	0
2	A2	2616	OMC	1	0
83	m2	27	A2M	2	0
2	A2	1496	PSU	1	0
2	A2	4270	A2M	1	0
2	A2	4022	OMG	1	0
2	A2	3848	OMG	1	0
83	m2	116	OMU	3	0
83	m2	603	OMG	3	0
83	m2	1705	OMC	2	0
2	A2	1330	2MG	1	0
83	m2	1250	B8N	2	0

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

Of 124 ligands modelled in this entry, 124 are monoatomic - leaving 0 for Mogul analysis.

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

No monomer is involved in short contacts.

5.7 Other polymers ⓘ

There are no such residues in this entry.

5.8 Polymer linkage issues ⓘ

The following chains have linkage breaks:

Mol	Chain	Number of breaks
2	A2	21
83	m2	9

All chain breaks are listed below:

Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	m2	130:G	O3'	141:A	P	24.99
1	A2	1512:U	O3'	1521:A	P	24.63
1	A2	2658:G	O3'	3240:C	P	19.23
1	m2	690:U	O3'	801:U	P	17.52
1	A2	891:C	O3'	917:G	P	17.46
1	A2	4437:C	O3'	4493:G	P	17.16
1	A2	3790:G	O3'	3796:G	P	16.55
1	A2	770:G	O3'	799:C	P	16.50
1	A2	3752:C	O3'	3758:G	P	16.18
1	A2	520:C	O3'	650:G	P	16.02
1	m2	536:G	O3'	554:U	P	15.88
1	A2	859:G	O3'	866:A	P	15.08
1	m2	324:C	O3'	331:G	P	14.90
1	A2	1564:C	O3'	1572:A	P	14.68
1	A2	1772:A	O3'	1820:C	P	13.84
1	A2	1914:C	O3'	2004:G	P	13.40
1	m2	1753:C	O3'	1786:G	P	12.16
1	A2	481:G	O3'	485:U	P	11.94
1	A2	1055:G	O3'	1059:C	P	11.91
1	A2	4668:U	O3'	4674:G	P	11.84
1	A2	3633:C	O3'	3685:G	P	9.84
1	A2	1072:U	O3'	1078:G	P	8.29
1	A2	866:A	O3'	868:C	P	8.22
1	A2	956:C	O3'	999:C	P	7.38
1	m2	227:G	O3'	289:U	P	7.38

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Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	A2	501:G	O3'	506:G	P	4.71
1	A2	4422:G	O3'	4424:U	P	4.57
1	m2	1692:U	O3'	1693:U	P	3.87
1	m2	1339:B8T	O3'	1340:G	P	3.35
1	m2	1350:G	O3'	1351:G	P	0.48

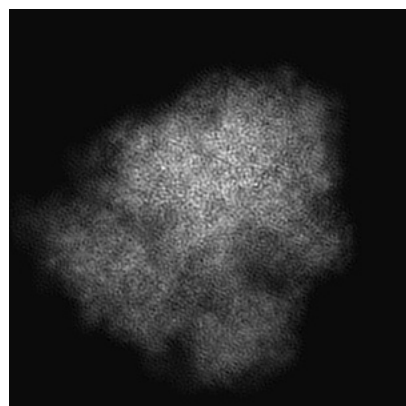
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-53473. These allow visual inspection of the internal detail of the map and identification of artifacts.

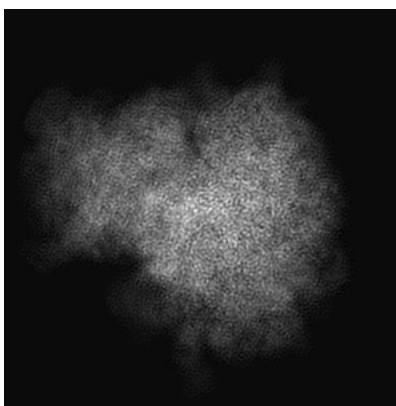
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections [i](#)

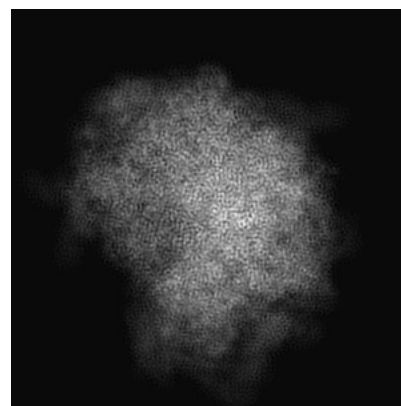
6.1.1 Primary map



X

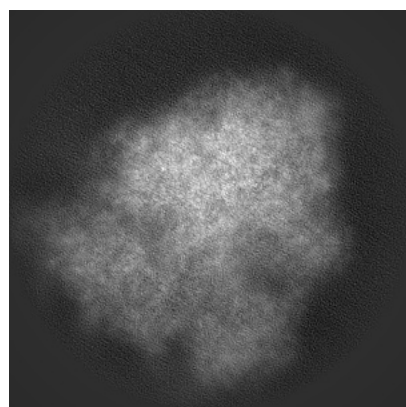


Y

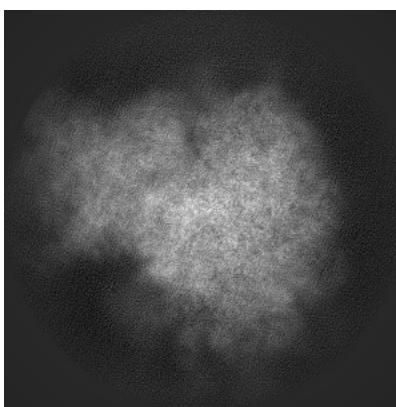


Z

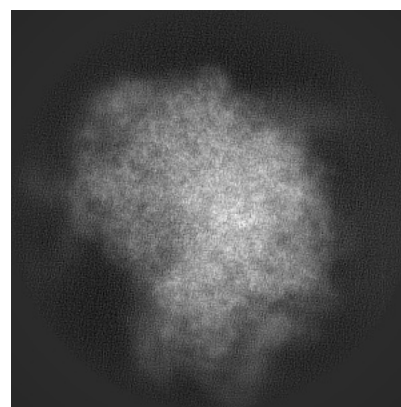
6.1.2 Raw map



X



Y

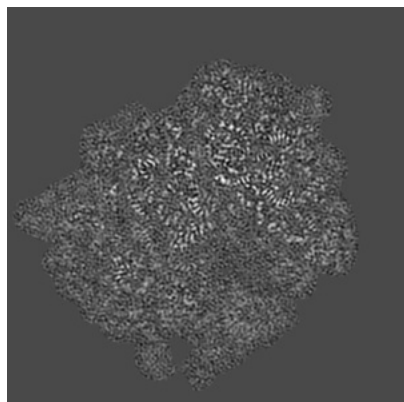


Z

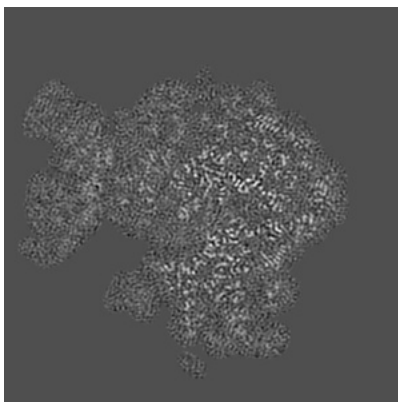
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

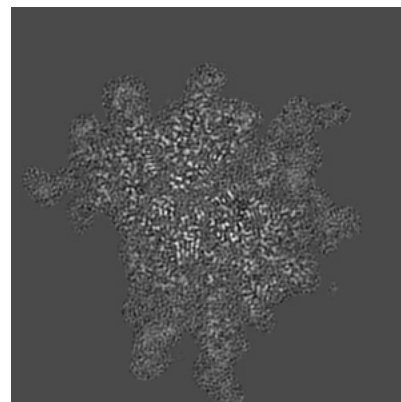
6.2.1 Primary map



X Index: 165

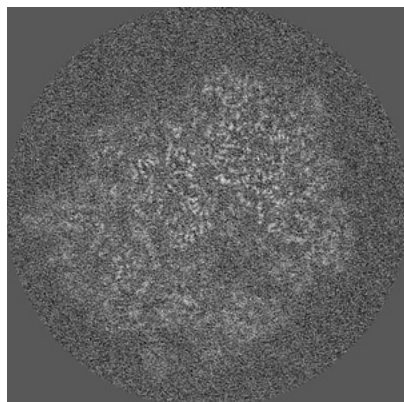


Y Index: 165

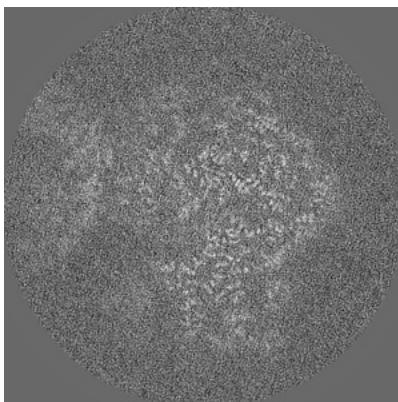


Z Index: 165

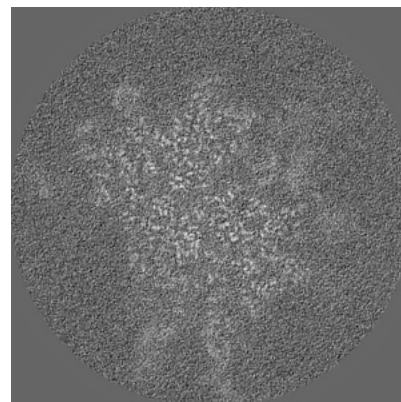
6.2.2 Raw map



X Index: 165



Y Index: 165

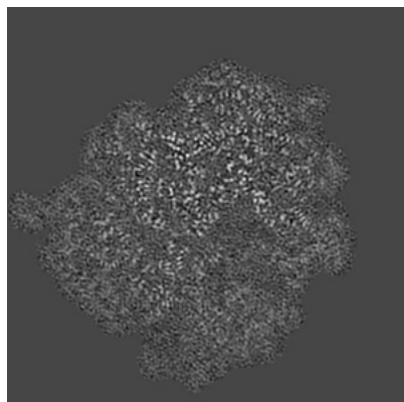


Z Index: 165

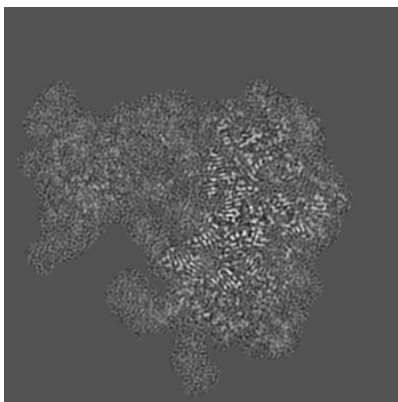
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

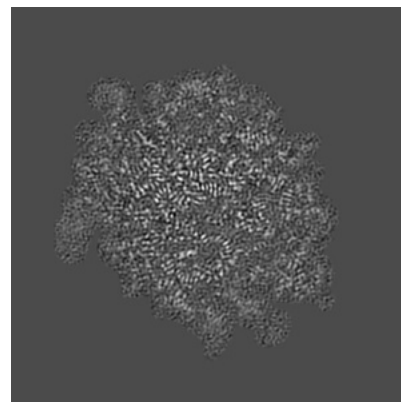
6.3.1 Primary map



X Index: 173

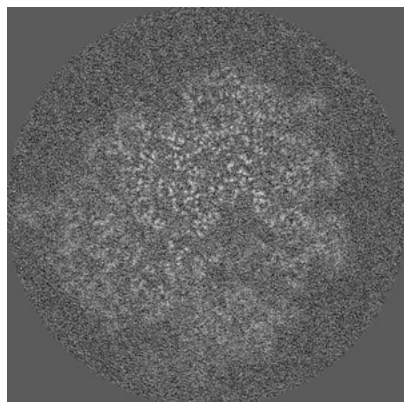


Y Index: 181

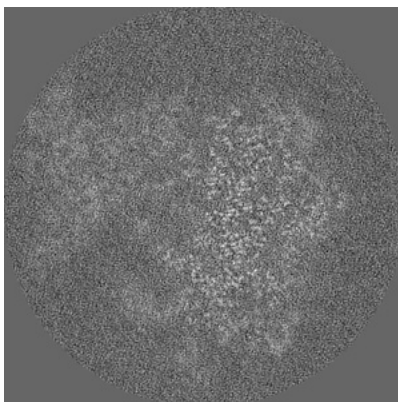


Z Index: 185

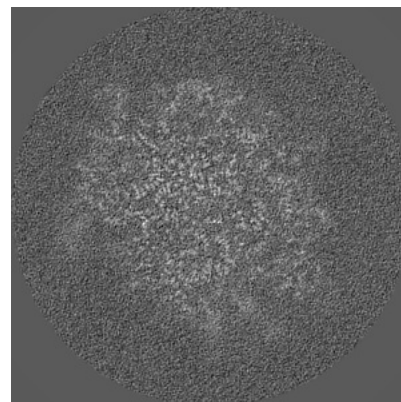
6.3.2 Raw map



X Index: 173



Y Index: 180

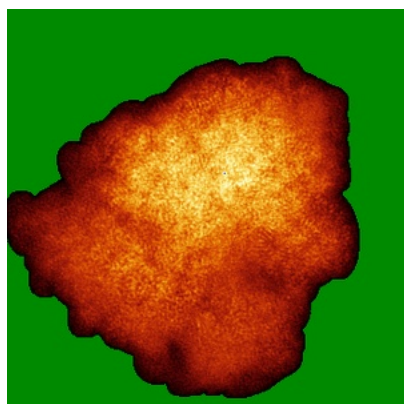


Z Index: 184

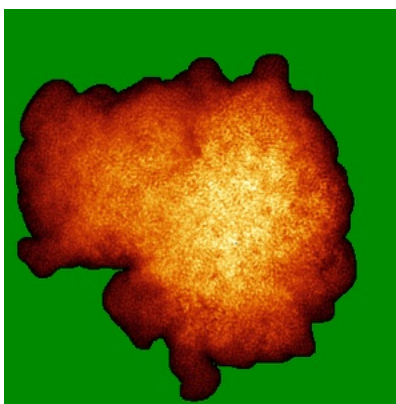
The images above show the largest variance slices of the map in three orthogonal directions.

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

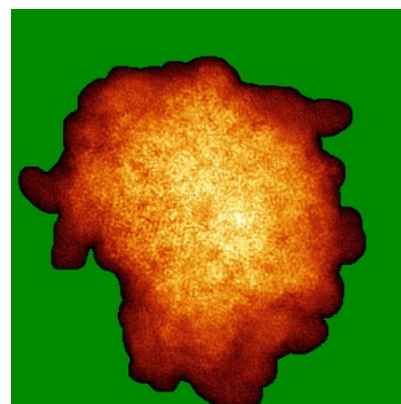
6.4.1 Primary map



X

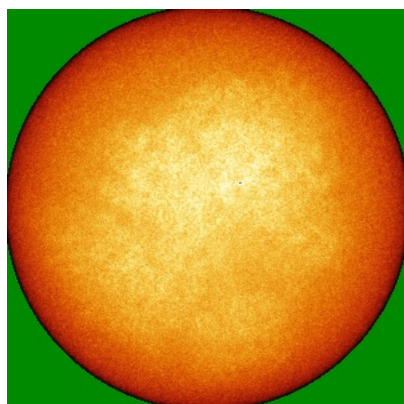


Y

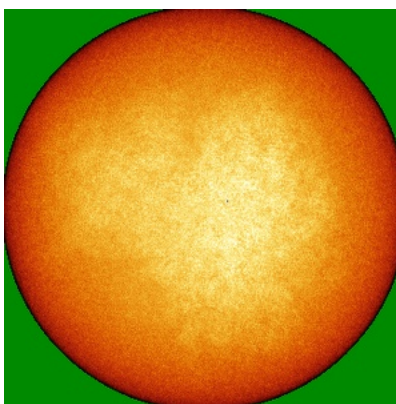


Z

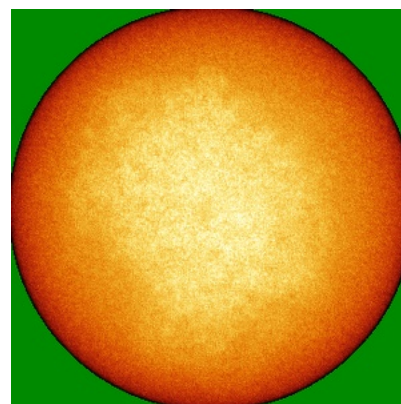
6.4.2 Raw map



X



Y

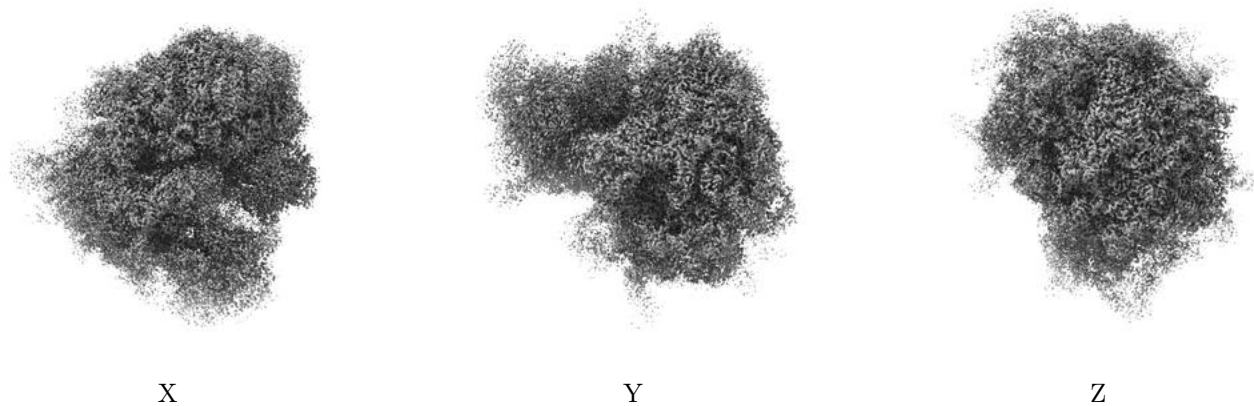


Z

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

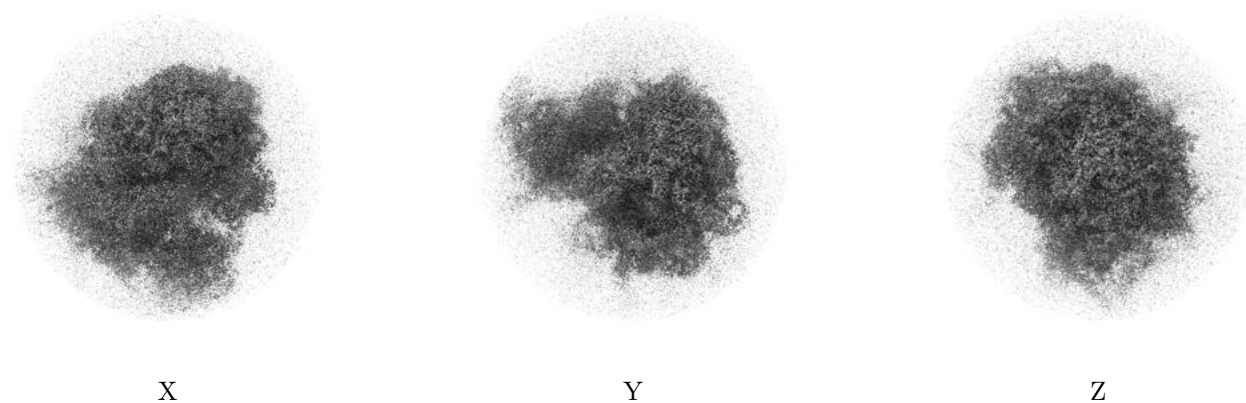
6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.037. 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.

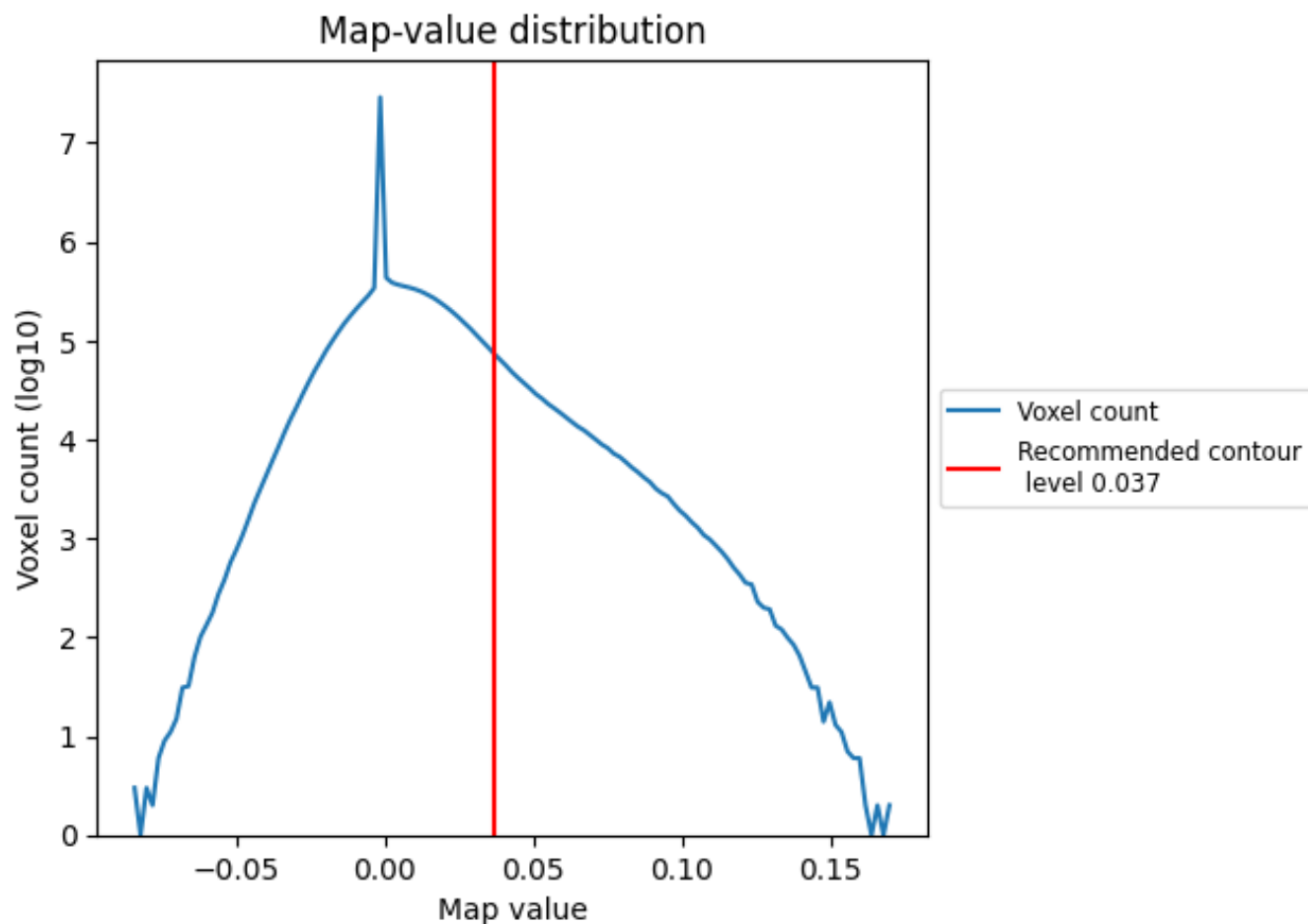
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

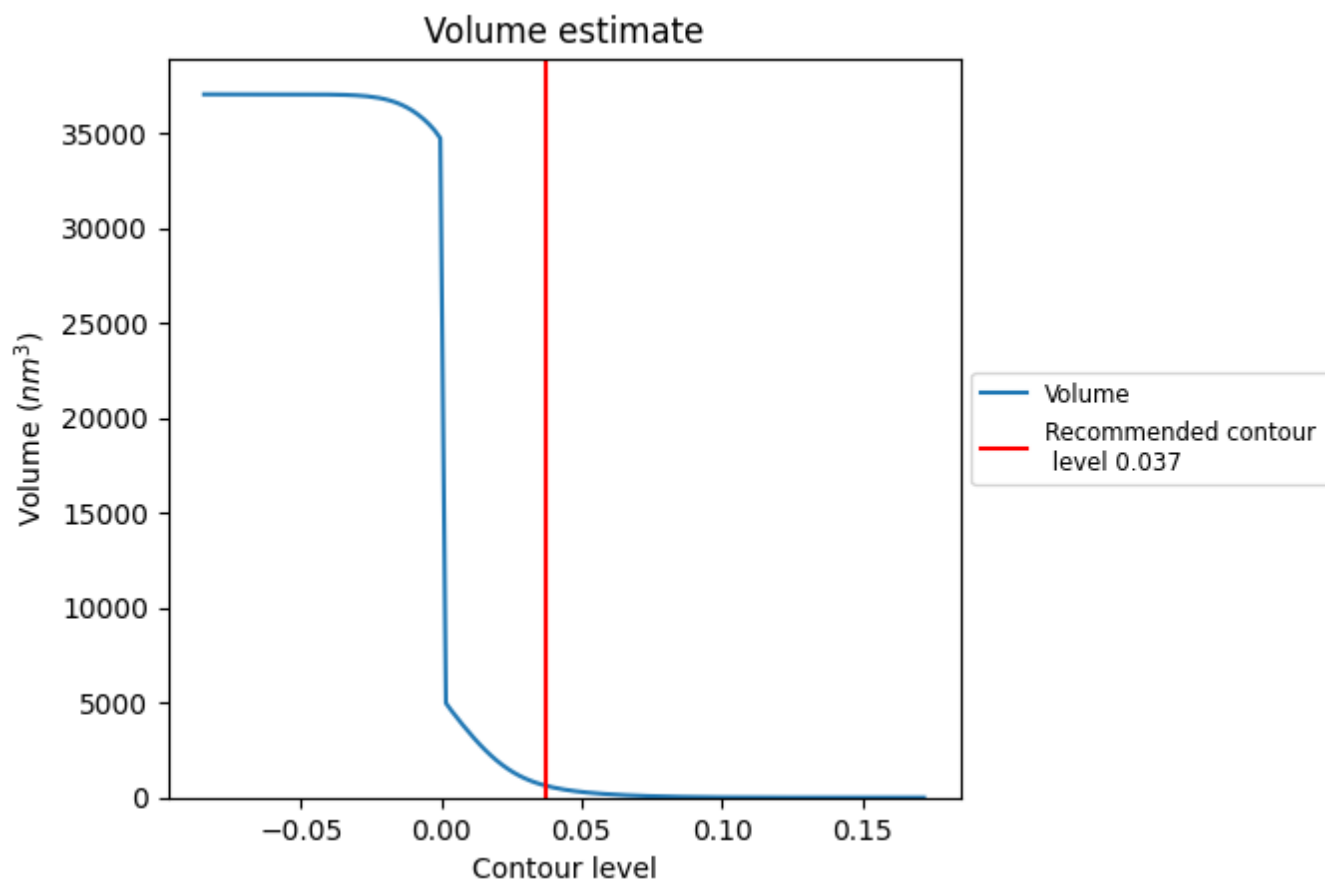
This section contains the results of statistical analysis of the map.

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.

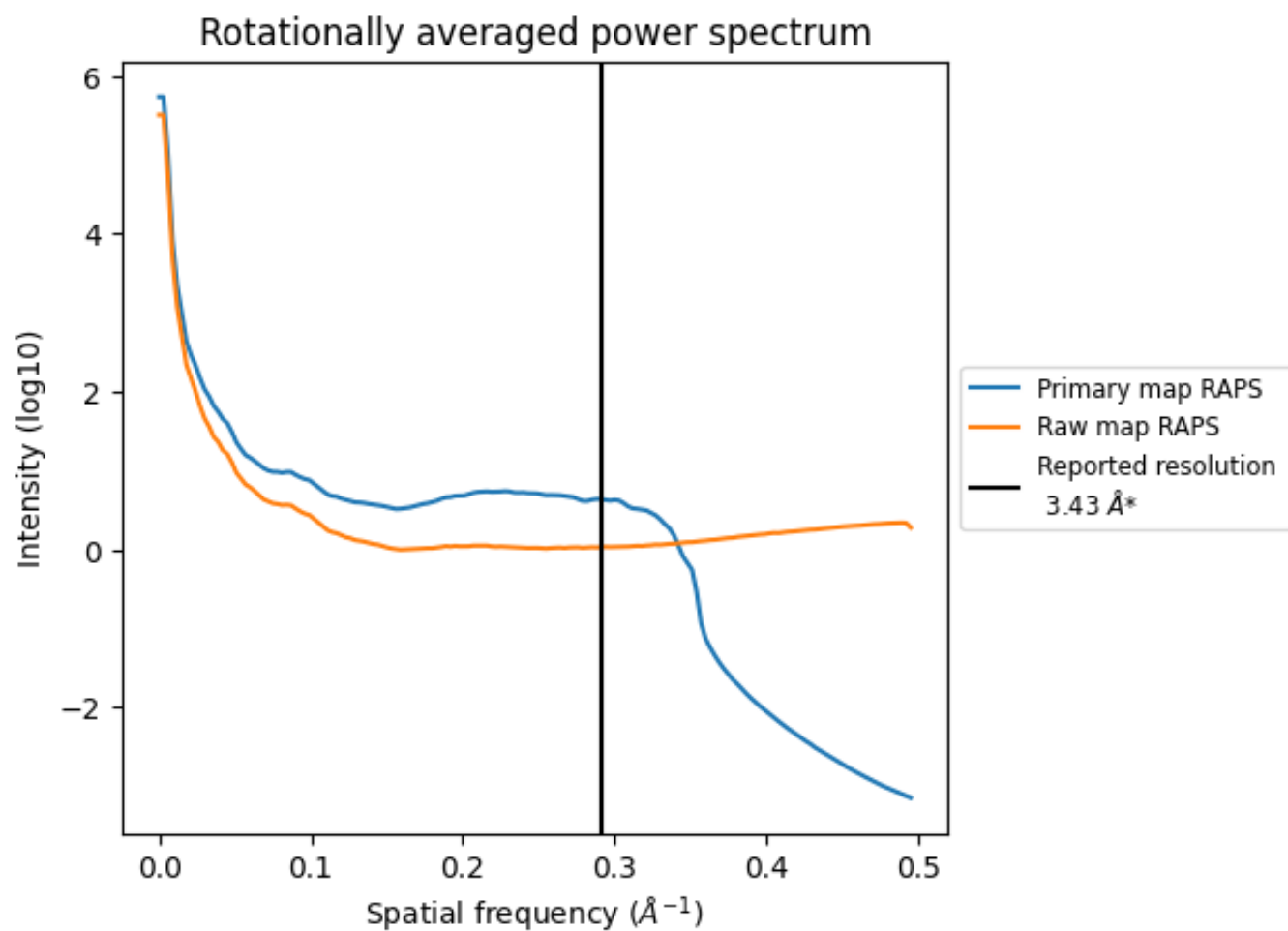
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 639 nm^3 ; this corresponds to an approximate mass of 577 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 ⓘ

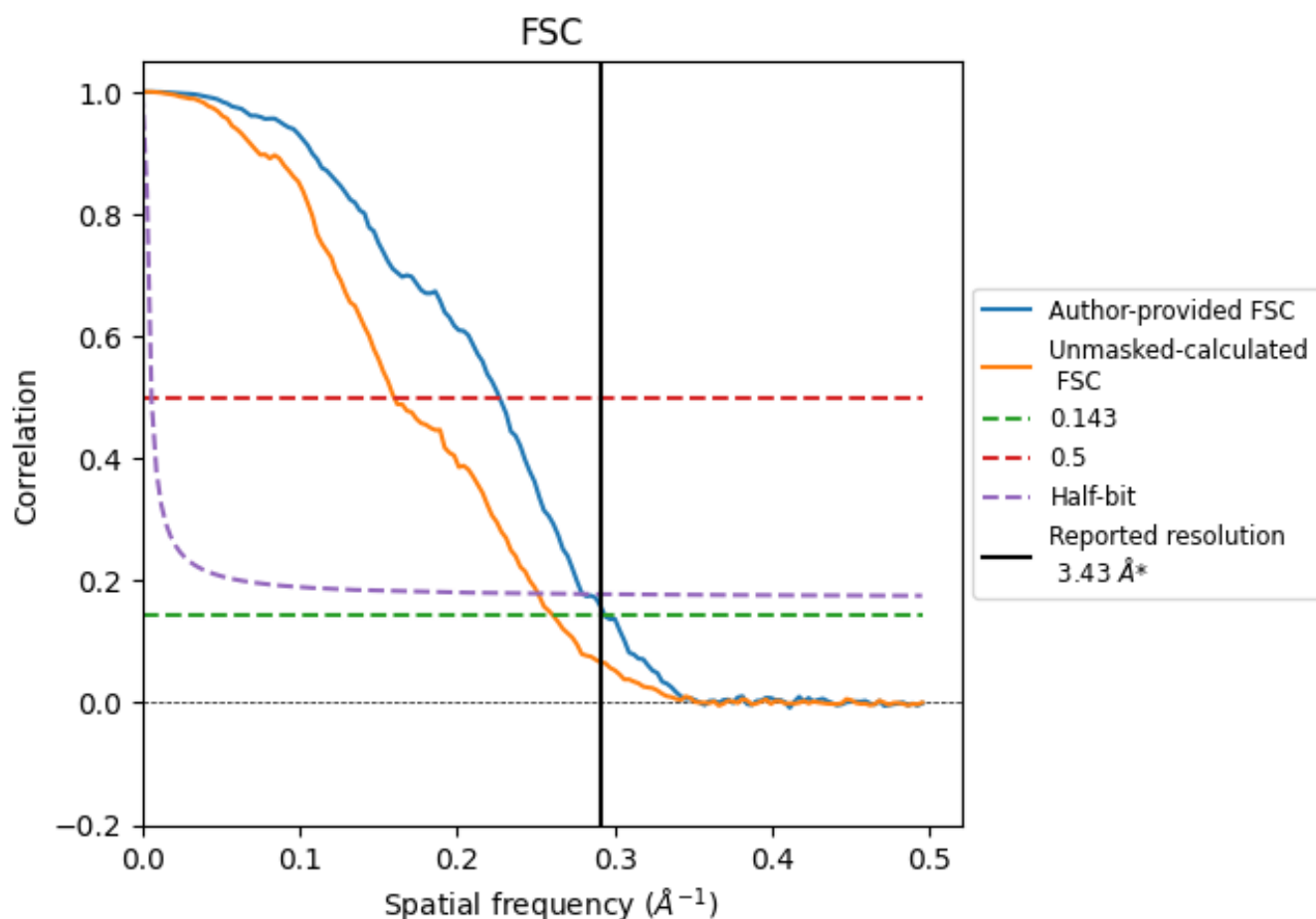


*Reported resolution corresponds to spatial frequency of 0.292 \AA^{-1}

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 0.292 \AA^{-1}

8.2 Resolution estimates [i](#)

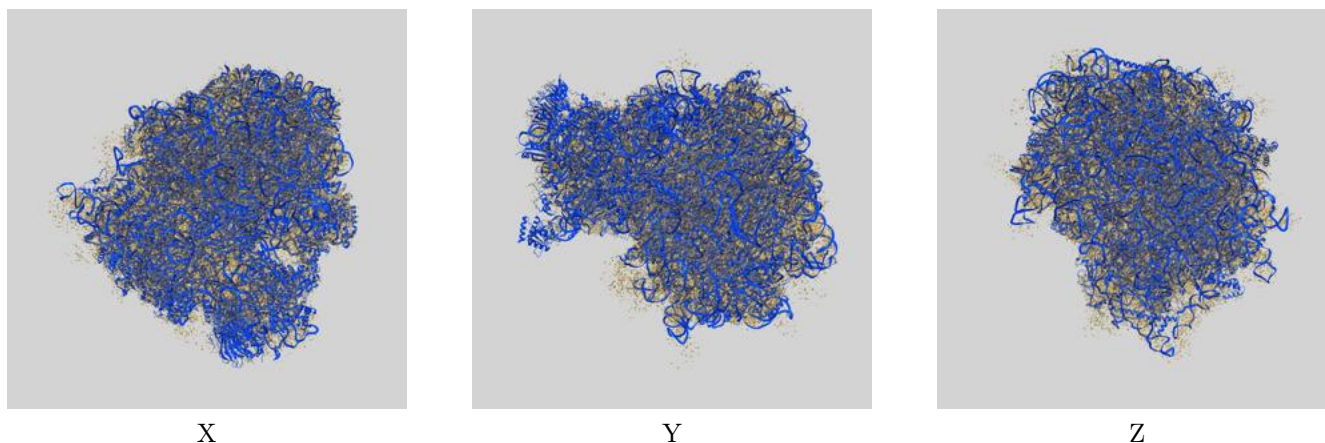
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	3.43	-	-
Author-provided FSC curve	3.41	4.41	3.57
Unmasked-calculated*	3.84	6.26	3.98

*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 3.84 differs from the reported value 3.43 by more than 10 %

9 Map-model fit [i](#)

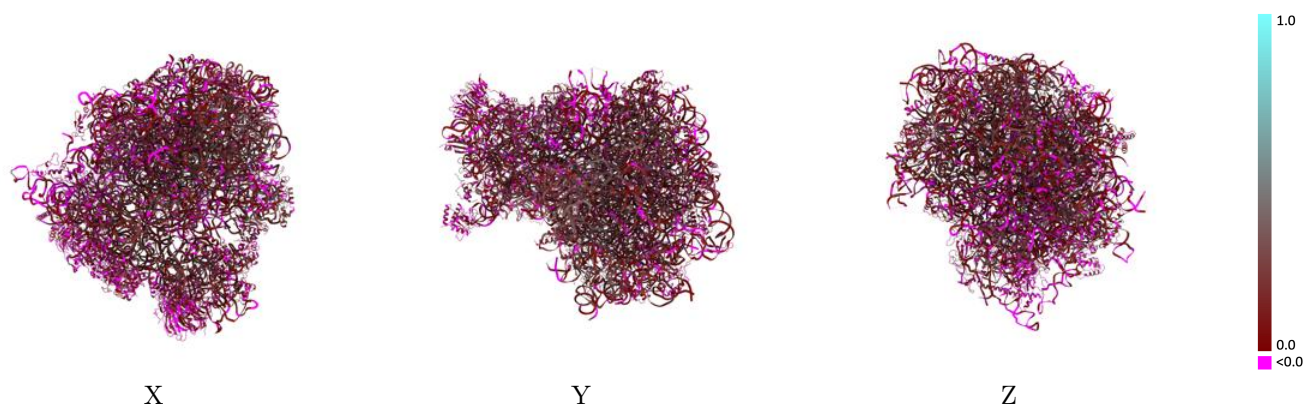
This section contains information regarding the fit between EMDB map EMD-53473 and PDB model 9QZP. Per-residue inclusion information can be found in section 3 on page 21.

9.1 Map-model overlay [i](#)



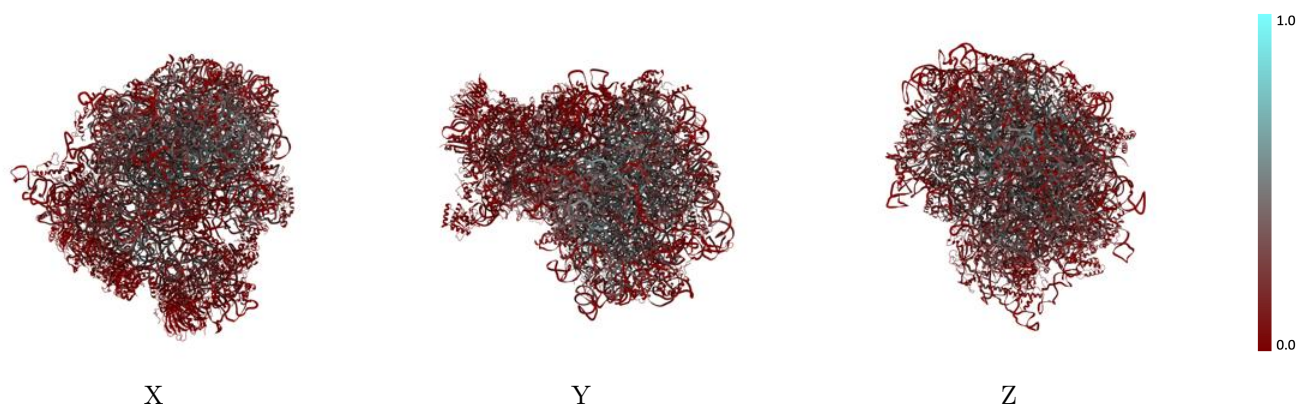
The images above show the 3D surface view of the map at the recommended contour level 0.037 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



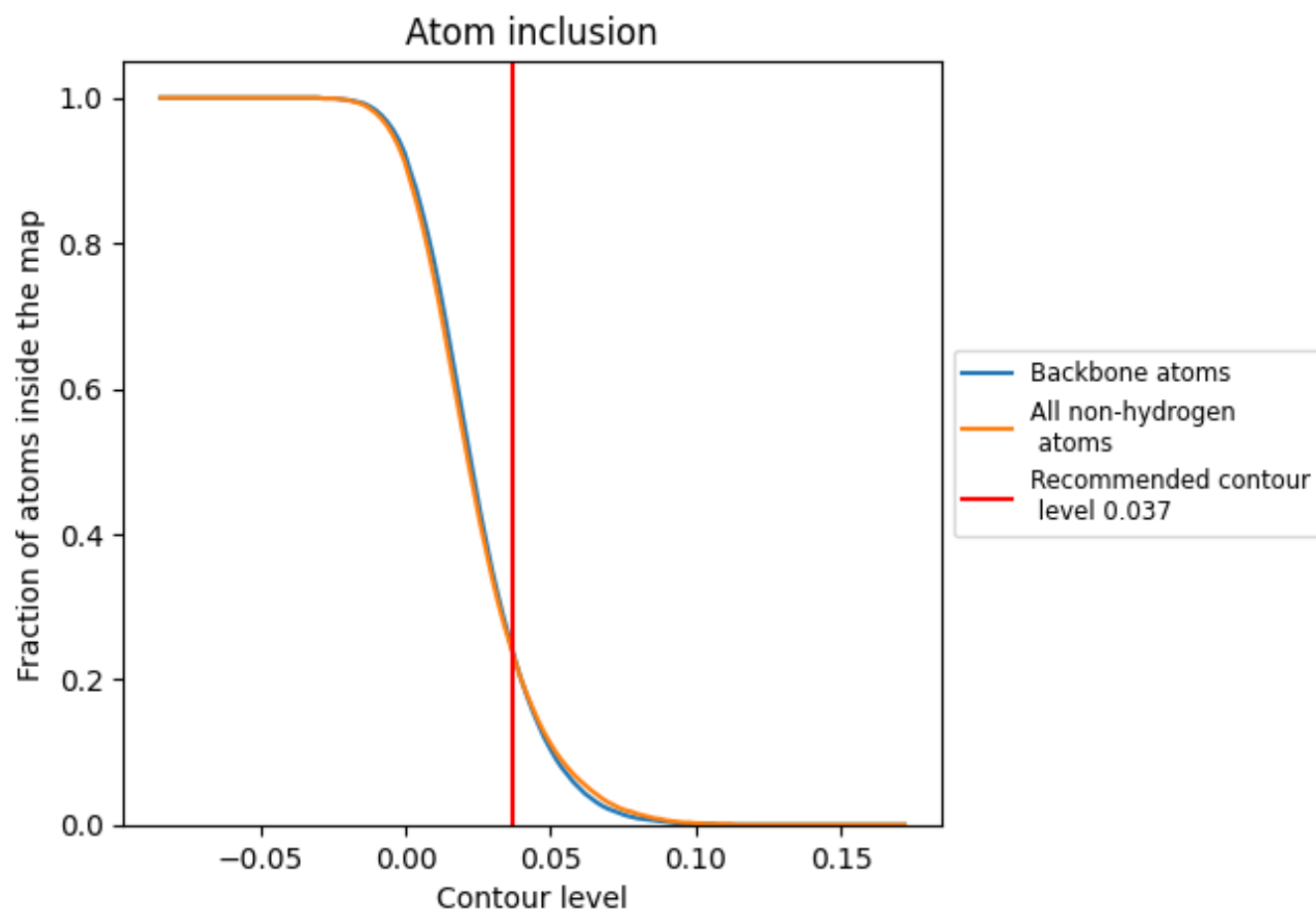
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions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

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



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (0.037).







































































9.4 Atom inclusion [i](#)



At the recommended contour level, 24% of all backbone atoms, 24% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary

The table lists the average atom inclusion at the recommended contour level (0.037) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.2350	 0.1470
A1	 0.3230	 0.1960
A2	 0.3130	 0.1510
A3	 0.0660	 0.1040
B1	 0.1860	 0.1490
B2	 0.4370	 0.3050
B3	 0.0890	 0.1280
Bv	 0.0710	 0.0640
Bx	 0.1400	 0.0730
By	 0.0450	 0.0530
Bz	 0.0420	 0.0610
C1	 0.2260	 0.2210
C2	 0.3270	 0.1570
C3	 0.0470	 0.0990
D1	 0.2650	 0.2210
D2	 0.2960	 0.1690
D3	 0.0420	 0.0790
E1	 0.1430	 0.1800
E2	 0.2410	 0.1500
E3	 0.2030	 0.1720
F1	 0.2360	 0.1680
F2	 0.2640	 0.1720
F3	 0.1650	 0.1570
G1	 0.2250	 0.1890
G2	 0.2120	 0.2410
G3	 0.0620	 0.1030
H1	 0.3710	 0.1970
H2	 0.1430	 0.1250
H3	 0.0850	 0.1010
I2	 0.2800	 0.1790
I3	 0.0140	 0.0420
J2	 0.2600	 0.1320
J3	 0.0930	 0.0780
K2	 0.3100	 0.2250
K3	 0.0720	 0.0920





















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Chain	Atom inclusion	Q-score
L1	0.0050	0.0160
L2	0.2070	0.0900
L3	0.0620	0.0500
M2	0.2930	0.2410
M3	0.0040	0.0240
N2	0.3030	0.2370
N3	0.1190	0.1440
O2	0.1000	0.0940
O3	0.0830	0.1500
P2	0.2640	0.1460
P3	0.1100	0.1030
Q2	0.2020	0.0720
Q3	0.0540	0.0820
R2	0.2030	0.1180
R3	0.0490	0.1140
S2	0.2220	0.1350
S3	0.0550	0.0920
T2	0.2140	0.1970
T3	0.0690	0.1070
U2	0.3270	0.2070
U3	0.0070	0.0230
V2	0.1920	0.1670
W2	0.2490	0.2470
X2	0.2610	0.1740
Y2	0.3140	0.2150
Z2	0.2910	0.1880
a2	0.2640	0.1600
b2	0.2070	0.1220
c2	0.2140	0.1870
d2	0.3040	0.1260
e2	0.1150	0.1340
f2	0.3030	0.1180
g2	0.2690	0.1930
h2	0.2440	0.1220
i2	0.2880	0.2260
j2	0.2430	0.1410
k2	0.2190	0.1470
m2	0.2370	0.1580
n2	0.0220	0.0420
o2	0.0430	0.0870
p2	0.0920	0.1540
q2	0.0520	0.1040

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Chain	Atom inclusion	Q-score
r2	 0.0660	 0.0390
s2	 0.0680	 0.1190
t2	 0.0250	 0.0500
u2	 0.0970	 0.0450
v2	 0.0180	 0.1010
w2	 0.1120	 0.0770
x2	 0.0620	 0.1350
y2	 0.0720	 0.1290
z2	 0.0480	 0.1180