



Full wwPDB EM Validation Report ⓘ

Jun 15, 2026 – 06:48 PM EDT

PDB ID : 8VL1 / pdb_00008vl1
EMDB ID : EMD-43335
Title : Escherichia coli transcription-translation loosely coupled complex (TTC-LC) containing mRNA with a 36 nt long spacer, ops signal, RfaH, NusA, and fMet-tRNAs in E-site and P-site
Authors : Molodtsov, V.; Wang, C.; Ebright, R.H.
Deposited on : 2024-01-11
Resolution : 5.30 Å(reported)

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

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

A user guide is available at

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

The types of validation reports are described at

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

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

EMDB validation analysis : 0.0.1.dev132
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

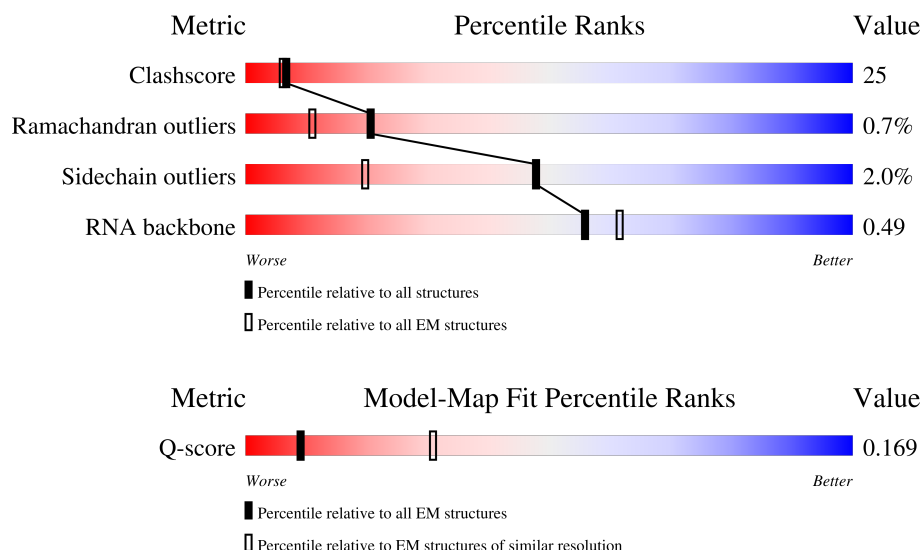
1 Overall quality at a glance i

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 5.30 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.



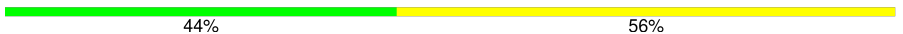
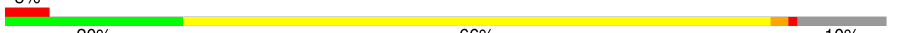
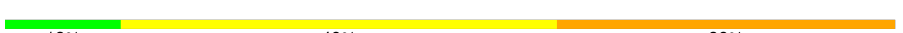
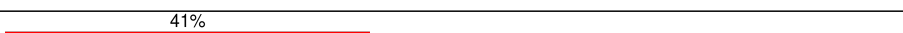
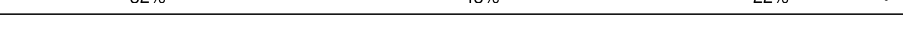
Metric	Whole archive (#Entries)	EM structures (#Entries)	Similar EM resolution (#Entries, resolution range(Å))
Clashscore	229148	23984	-
Ramachandran outliers	224038	23583	-
Sidechain outliers	223484	23102	-
RNA backbone	8273	3508	-
Q-score	-	25397	625 (4.80 - 5.80)

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	0	103	60% 40%
2	1	110	60% 40%
3	2	100	52% 41% 6%

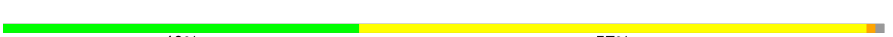
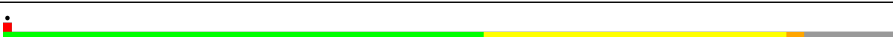
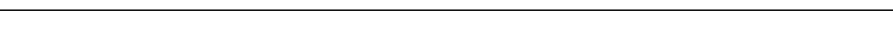
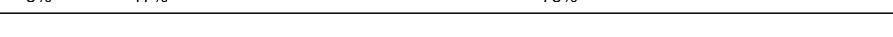
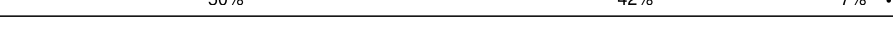
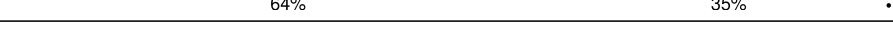
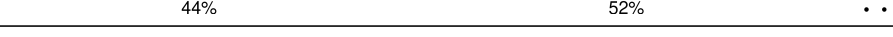
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Mol	Chain	Length	Quality of chain
4	3	104	
5	4	94	
6	5	38	
7	6	38	
8	7	53	
9	9	165	
10	A	76	
10	B	76	
11	AA	1342	
12	AB	162	
13	AC	329	
13	AD	329	
14	AE	1407	
15	AF	91	
16	AG	495	
17	C	75	
18	D	1542	
19	E	87	
20	F	71	
21	G	241	
22	H	557	
23	I	233	
24	J	206	
25	K	167	
26	L	135	

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Mol	Chain	Length	Quality of chain
27	M	179	
28	N	130	
29	O	130	
30	P	103	
31	Q	129	
32	R	124	
33	S	101	
34	T	89	
35	U	82	
36	V	84	
37	W	92	
38	X	118	
39	Y	142	
40	Z	121	
41	a	2904	
42	b	85	
43	c	78	
44	d	120	
45	e	63	
46	f	59	
47	g	70	
48	h	273	
49	i	57	
50	j	209	
51	k	55	

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Mol	Chain	Length	Quality of chain
52	l	201	
53	m	46	
54	n	179	
55	o	65	
56	p	177	
57	q	38	
58	r	149	
59	s	142	
60	t	123	
61	u	144	
62	v	136	
63	w	127	
64	x	117	
65	y	115	
66	z	118	

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

Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
68	ZN	AE	1502	-	-	X	-

2 Entry composition

There are 68 unique types of molecules in this entry. The entry contains 280046 atoms, of which 98638 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 Ribosomal protein L21.

Mol	Chain	Residues	Atoms						AltConf	Trace
1	0	103	Total	C	H	N	O	S	0	0
			1655	516	839	153	145	2		

- Molecule 2 is a protein called 50S ribosomal protein L22.

Mol	Chain	Residues	Atoms						AltConf	Trace
2	1	110	Total	C	H	N	O	S	0	0
			1779	532	922	166	156	3		

- Molecule 3 is a protein called 50S ribosomal protein L23.

Mol	Chain	Residues	Atoms						AltConf	Trace
3	2	94	Total	C	H	N	O	S	0	0
			1557	470	811	140	134	2		

- Molecule 4 is a protein called 50S ribosomal protein L24.

Mol	Chain	Residues	Atoms						AltConf	Trace
4	3	103	Total	C	H	N	O		0	0
			1632	498	844	148	142			

- Molecule 5 is a protein called 50S ribosomal protein L25.

Mol	Chain	Residues	Atoms						AltConf	Trace
5	4	94	Total	C	H	N	O	S	0	0
			1533	479	780	137	134	3		

- Molecule 6 is a DNA chain called NT DNA ops.

Mol	Chain	Residues	Atoms						AltConf	Trace
6	5	35	Total	C	N	O	P		0	0
			726	342	141	208	35			

- Molecule 7 is a DNA chain called T DNA ops.

Mol	Chain	Residues	Atoms					AltConf	Trace
7	6	35	Total	C	N	O	P	0	0
			703	336	117	215	35		

- Molecule 8 is a RNA chain called mRNA with 36 nt long spacer.

Mol	Chain	Residues	Atoms					AltConf	Trace
8	7	25	Total	C	N	O	P	0	0
			527	235	80	187	25		

- Molecule 9 is a protein called 50S ribosomal protein L10.

Mol	Chain	Residues	Atoms					AltConf	Trace
9	9	148	Total	C	N	O	S	0	0
			1117	705	196	209	7		

- Molecule 10 is a RNA chain called E-site and P-site tRNA (fMet).

Mol	Chain	Residues	Atoms						AltConf	Trace
10	A	76	Total 2446	C 723	H 826	N 295	O 527	P 75	0	0
10	B	76	Total 2433	C 723	H 813	N 295	O 527	P 75	0	0

- Molecule 11 is a protein called DNA-directed RNA polymerase subunit beta.

Mol	Chain	Residues	Atoms					AltConf	Trace
11	AA	1316	Total	C	N	O	S	0	0
			10381	6514	1810	2014	43		

- Molecule 12 is a protein called Transcription antitermination protein RfaH.

Mol	Chain	Residues	Atoms					AltConf	Trace
12	AB	161	Total	C	N	O	S	0	0
			1286	828	222	232	4		

- Molecule 13 is a protein called DNA-directed RNA polymerase subunit alpha.

Mol	Chain	Residues	Atoms					AltConf	Trace
13	AC	221	Total	C	N	O	S	0	0
			1698	1060	299	333	6		

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Mol	Chain	Residues	Atoms					AltConf	Trace
13	AD	299	Total	C	N	O	S	0	0
			2078	1287	378	407	6		

- Molecule 14 is a protein called DNA-directed RNA polymerase subunit beta'.

Mol	Chain	Residues	Atoms					AltConf	Trace
14	AE	1337	Total	C	N	O	S	0	0
			10404	6535	1856	1963	50		

- Molecule 15 is a protein called DNA-directed RNA polymerase subunit omega.

Mol	Chain	Residues	Atoms					AltConf	Trace
15	AF	82	Total	C	N	O	S	0	0
			650	396	122	131	1		

- Molecule 16 is a protein called Transcription termination/antitermination protein NusA.

Mol	Chain	Residues	Atoms					AltConf	Trace
16	AG	495	Total	C	N	O	S	0	0
			3852	2396	669	774	13		

- Molecule 17 is a protein called 30S ribosomal protein S18.

Mol	Chain	Residues	Atoms					AltConf	Trace
17	C	66	Total	C	H	N	O	S	0
			1103	344	559	102	97	1	0

- Molecule 18 is a RNA chain called 16S rRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
18	D	1524	Total	C	H	N	O	P	0
			49126	14585	16423	6003	10591	1524	0

- Molecule 19 is a protein called 30S ribosomal protein S20.

Mol	Chain	Residues	Atoms					AltConf	Trace
19	E	86	Total	C	H	N	O	S	0
			1388	414	719	138	114	3	0

- Molecule 20 is a protein called 30S ribosomal protein S21.

Mol	Chain	Residues	Atoms						AltConf	Trace
20	F	70	Total	C	H	N	O	S	0	0
			1218	366	629	125	97	1		

- Molecule 21 is a protein called 30S ribosomal protein S2.

Mol	Chain	Residues	Atoms						AltConf	Trace
21	G	225	Total	C	H	N	O	S	0	0
			3545	1113	1785	316	323	8		

- Molecule 22 is a protein called 30S ribosomal protein S1.

Mol	Chain	Residues	Atoms						AltConf	Trace
22	H	259	Total	C	H	N	O	S	0	0
			3184	1073	1454	305	349	3		

- Molecule 23 is a protein called 30S ribosomal protein S3.

Mol	Chain	Residues	Atoms						AltConf	Trace
23	I	208	Total	C	H	N	O	S	0	0
			3346	1036	1710	307	290	3		

- Molecule 24 is a protein called 30S ribosomal protein S4.

Mol	Chain	Residues	Atoms						AltConf	Trace
24	J	205	Total	C	H	N	O	S	0	0
			3350	1026	1707	315	298	4		

- Molecule 25 is a protein called 30S ribosomal protein S5.

Mol	Chain	Residues	Atoms						AltConf	Trace
25	K	156	Total	C	H	N	O	S	0	0
			2348	717	1196	217	212	6		

- Molecule 26 is a protein called 30S ribosomal protein S6.

Mol	Chain	Residues	Atoms						AltConf	Trace
26	L	104	Total	C	H	N	O	S	0	0
			1694	536	846	153	152	7		

- Molecule 27 is a protein called 30S ribosomal protein S7.

Mol	Chain	Residues	Atoms						AltConf	Trace
27	M	151	Total	C	H	N	O	S	0	0
			2416	735	1235	227	215	4		

- Molecule 28 is a protein called 30S ribosomal protein S8.

Mol	Chain	Residues	Atoms						AltConf	Trace
28	N	129	Total	C	H	N	O	S	0	0
			2010	616	1031	173	184	6		

- Molecule 29 is a protein called 30S ribosomal protein S9.

Mol	Chain	Residues	Atoms						AltConf	Trace
29	O	127	Total	C	H	N	O	S	0	0
			2092	634	1070	206	179	3		

- Molecule 30 is a protein called 30S ribosomal protein S10.

Mol	Chain	Residues	Atoms						AltConf	Trace
30	P	99	Total	C	H	N	O	S	0	0
			1621	495	831	151	143	1		

- Molecule 31 is a protein called 30S ribosomal protein S11.

Mol	Chain	Residues	Atoms						AltConf	Trace
31	Q	117	Total	C	H	N	O	S	0	0
			1764	540	887	174	160	3		

- Molecule 32 is a protein called 30S ribosomal protein S12.

Mol	Chain	Residues	Atoms						AltConf	Trace
32	R	121	Total	C	H	N	O	S	0	0
			1940	580	1001	194	161	4		

- Molecule 33 is a protein called 30S ribosomal protein S14.

Mol	Chain	Residues	Atoms						AltConf	Trace
33	S	100	Total	C	H	N	O	S	0	0
			1649	499	844	164	139	3		

- Molecule 34 is a protein called Small ribosomal subunit protein uS15.

Mol	Chain	Residues	Atoms						AltConf	Trace
34	T	88	Total	C	H	N	O	S	0	0
			1448	439	734	144	130	1		

- Molecule 35 is a protein called 30S ribosomal protein S16.

Mol	Chain	Residues	Atoms						AltConf	Trace
35	U	82	Total	C	H	N	O	S	0	0
			1315	406	666	128	114	1		

- Molecule 36 is a protein called 30S ribosomal protein S17.

Mol	Chain	Residues	Atoms						AltConf	Trace
36	V	80	Total	C	H	N	O	S	0	0
			1339	411	691	121	113	3		

- Molecule 37 is a protein called 30S ribosomal protein S19.

Mol	Chain	Residues	Atoms						AltConf	Trace
37	W	83	Total	C	H	N	O	S	0	0
			1351	424	688	126	111	2		

- Molecule 38 is a protein called 30S ribosomal protein S13.

Mol	Chain	Residues	Atoms						AltConf	Trace
38	X	116	Total	C	H	N	O	S	0	0
			1864	558	964	181	158	3		

- Molecule 39 is a protein called 50S ribosomal protein L11.

Mol	Chain	Residues	Atoms					AltConf	Trace
39	Y	141	Total	C	N	O	S	0	0
			1032	651	179	196	6		

- Molecule 40 is a protein called 50S ribosomal protein L7/L12.

Mol	Chain	Residues	Atoms					AltConf	Trace
40	Z	30	Total	C	N	O	S	0	0
			227	144	33	47	3		

- Molecule 41 is a RNA chain called 23S rRNA.

Mol	Chain	Residues	Atoms						AltConf	Trace
41	a	2880	Total	C	H	N	O	P	0	0
			92918	27587	31077	11398	19976	2880		

There is a discrepancy between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
a	887	A	U	conflict	GB 937521852

- Molecule 42 is a protein called 50S ribosomal protein L27.

Mol	Chain	Residues	Atoms						AltConf	Trace
42	b	76	Total	C	H	N	O	S	0	0
			1181	360	599	117	104	1		

- Molecule 43 is a protein called 50S ribosomal protein L28.

Mol	Chain	Residues	Atoms						AltConf	Trace
43	c	77	Total	C	H	N	O	S	0	0
			1277	388	652	129	106	2		

- Molecule 44 is a RNA chain called 5S rRNA.

Mol	Chain	Residues	Atoms						AltConf	Trace
44	d	120	Total	C	H	N	O	P	0	0
			3870	1144	1301	468	837	120		

- Molecule 45 is a protein called 50S ribosomal protein L29.

Mol	Chain	Residues	Atoms						AltConf	Trace
45	e	62	Total	C	H	N	O	S	0	0
			1032	308	531	98	94	1		

- Molecule 46 is a protein called 50S ribosomal protein L30.

Mol	Chain	Residues	Atoms						AltConf	Trace
46	f	58	Total	C	H	N	O	S	0	0
			936	281	488	87	78	2		

- Molecule 47 is a protein called 50S ribosomal protein L31.

Mol	Chain	Residues	Atoms						AltConf	Trace
47	g	66	Total	C	H	N	O	S	0	0
			1042	323	520	99	94	6		

- Molecule 48 is a protein called 50S ribosomal protein L2.

Mol	Chain	Residues	Atoms						AltConf	Trace
48	h	271	Total	C	H	N	O	S	0	0
			4236	1288	2154	423	364	7		

- Molecule 49 is a protein called 50S ribosomal protein L32.

Mol	Chain	Residues	Atoms						AltConf	Trace
49	i	56	Total	C	H	N	O	S	0	0
			903	269	459	94	80	1		

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

Mol	Chain	Residues	Atoms						AltConf	Trace
50	j	209	Total	C	H	N	O	S	0	0
			3182	979	1617	288	294	4		

- Molecule 51 is a protein called 50S ribosomal protein L33.

Mol	Chain	Residues	Atoms						AltConf	Trace
51	k	52	Total	C	H	N	O		0	0
			890	275	464	78	73			

- Molecule 52 is a protein called 50S ribosomal protein L4.

Mol	Chain	Residues	Atoms						AltConf	Trace
52	l	201	Total	C	H	N	O	S	0	0
			3171	974	1619	283	290	5		

- Molecule 53 is a protein called 50S ribosomal protein L34.

Mol	Chain	Residues	Atoms						AltConf	Trace
53	m	46	Total	C	H	N	O	S	0	0
			795	228	418	90	57	2		

- Molecule 54 is a protein called 50S ribosomal protein L5.

Mol	Chain	Residues	Atoms						AltConf	Trace
54	n	177	Total	C	H	N	O	S	0	0
			2853	899	1443	249	256	6		

- Molecule 55 is a protein called 50S ribosomal protein L35.

Mol	Chain	Residues	Atoms						AltConf	Trace
55	o	64	Total	C	H	N	O	S	0	0
			1076	323	572	105	74	2		

- Molecule 56 is a protein called 50S ribosomal protein L6.

Mol	Chain	Residues	Atoms						AltConf	Trace
56	p	175	Total	C	H	N	O	S	0	0
			2671	826	1358	241	244	2		

- Molecule 57 is a protein called 50S ribosomal protein L36.

Mol	Chain	Residues	Atoms						AltConf	Trace
57	q	38	Total	C	H	N	O	S	0	0
			645	185	343	65	48	4		

- Molecule 58 is a protein called 50S ribosomal protein L9.

Mol	Chain	Residues	Atoms						AltConf	Trace
58	r	149	Total	C	H	N	O	S	0	0
			2259	699	1148	197	214	1		

- Molecule 59 is a protein called 50S ribosomal protein L13.

Mol	Chain	Residues	Atoms						AltConf	Trace
59	s	142	Total	C	H	N	O	S	0	0
			2291	714	1162	212	199	4		

- Molecule 60 is a protein called 50S ribosomal protein L14.

Mol	Chain	Residues	Atoms						AltConf	Trace
60	t	123	Total	C	H	N	O	S	0	0
			1969	593	1023	181	166	6		

- Molecule 61 is a protein called 50S ribosomal protein L15.

Mol	Chain	Residues	Atoms						AltConf	Trace
61	u	144	Total	C	H	N	O	S	0	0
			2182	654	1129	207	190	2		

- Molecule 62 is a protein called 50S ribosomal protein L16.

Mol	Chain	Residues	Atoms						AltConf	Trace
62	v	136	Total	C	H	N	O	S	0	0
			2231	686	1157	205	177	6		

- Molecule 63 is a protein called 50S ribosomal protein L17.

Mol	Chain	Residues	Atoms						AltConf	Trace
63	w	119	Total	C	H	N	O	S	0	0
			1945	588	994	195	163	5		

- Molecule 64 is a protein called 50S ribosomal protein L18.

Mol	Chain	Residues	Atoms					AltConf	Trace
64	x	116	Total	C	H	N	O	0	0
			1815	552	923	178	162		

- Molecule 65 is a protein called 50S ribosomal protein L19.

Mol	Chain	Residues	Atoms						AltConf	Trace
65	y	114	Total	C	H	N	O	S	0	0
			1879	574	962	179	163	1		

- Molecule 66 is a protein called 50S ribosomal protein L20.

Mol	Chain	Residues	Atoms					AltConf	Trace
66	z	117	Total	C	H	N	O	0	0
			1967	604	1020	192	151		

- Molecule 67 is MAGNESIUM ION (CCD ID: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		AltConf
67	AE	1	Total	Mg	0
			1	1	

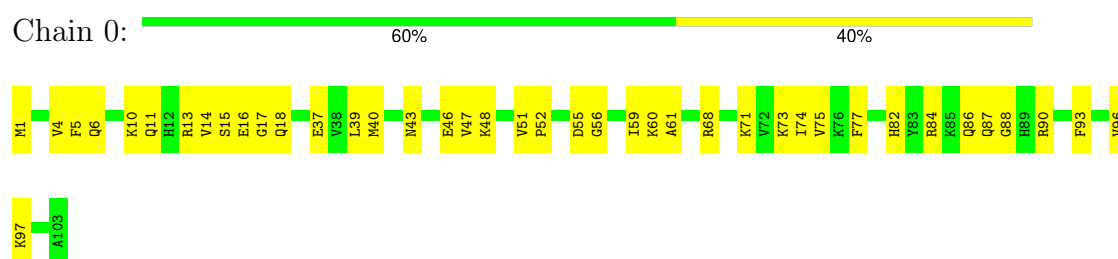
- Molecule 68 is ZINC ION (CCD ID: ZN) (formula: Zn).

Mol	Chain	Residues	Atoms		AltConf
68	AE	2	Total	Zn	0
			2	2	

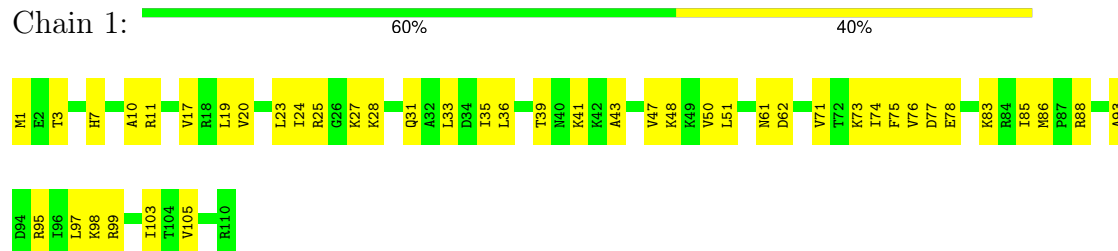
3 Residue-property plots [i](#)

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.

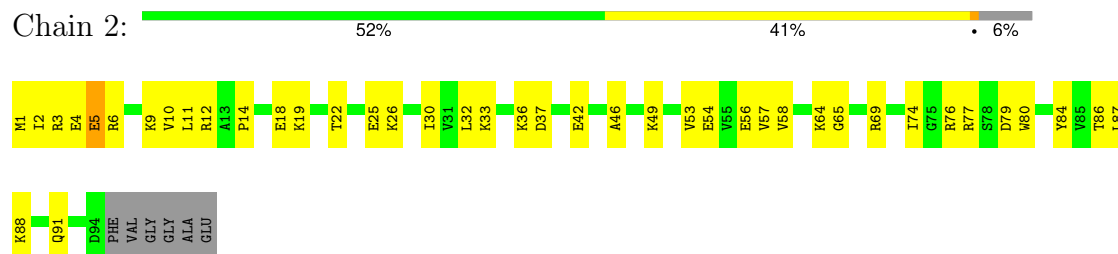
• Molecule 1: Ribosomal protein L21



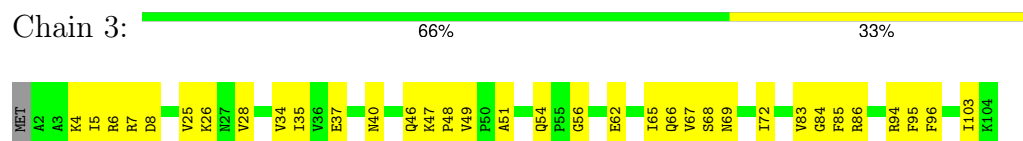
• Molecule 2: 50S ribosomal protein L22



• Molecule 3: 50S ribosomal protein L23



• Molecule 4: 50S ribosomal protein L24



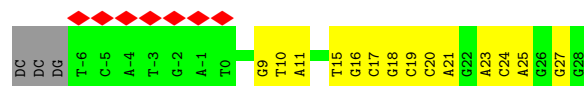
- Molecule 5: 50S ribosomal protein L25

Chain 4: 



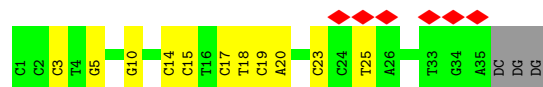
- Molecule 6: NT DNA ops

Chain 5: 



- Molecule 7: T DNA ops

Chain 6: 

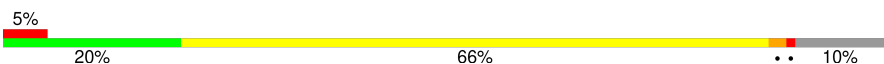


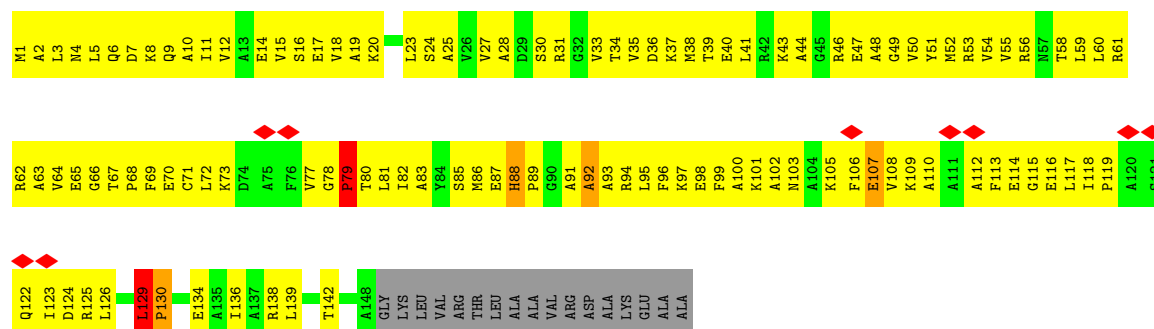
- Molecule 8: mRNA with 36 nt long spacer

Chain 7: 



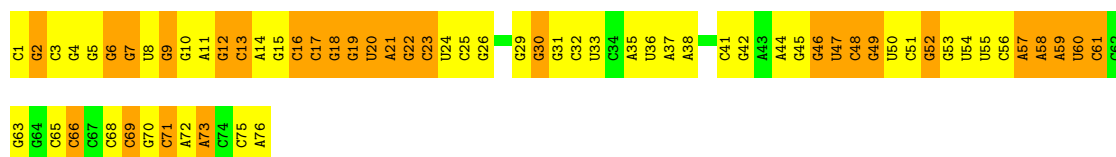
- Molecule 9: 50S ribosomal protein L10

Chain 9: 

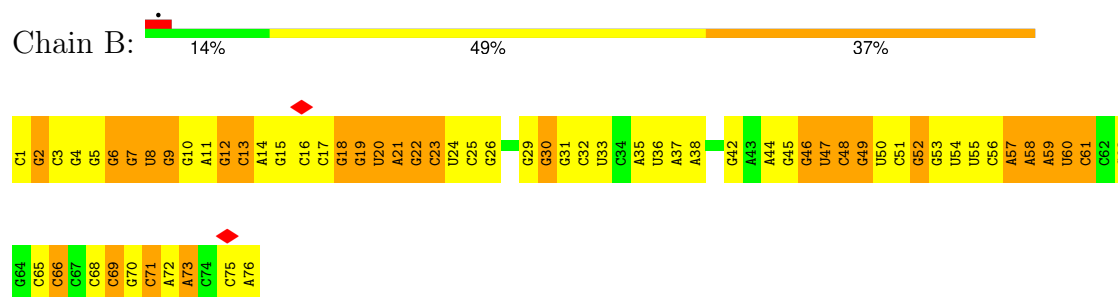


- Molecule 10: E-site and P-site tRNA (fMet)

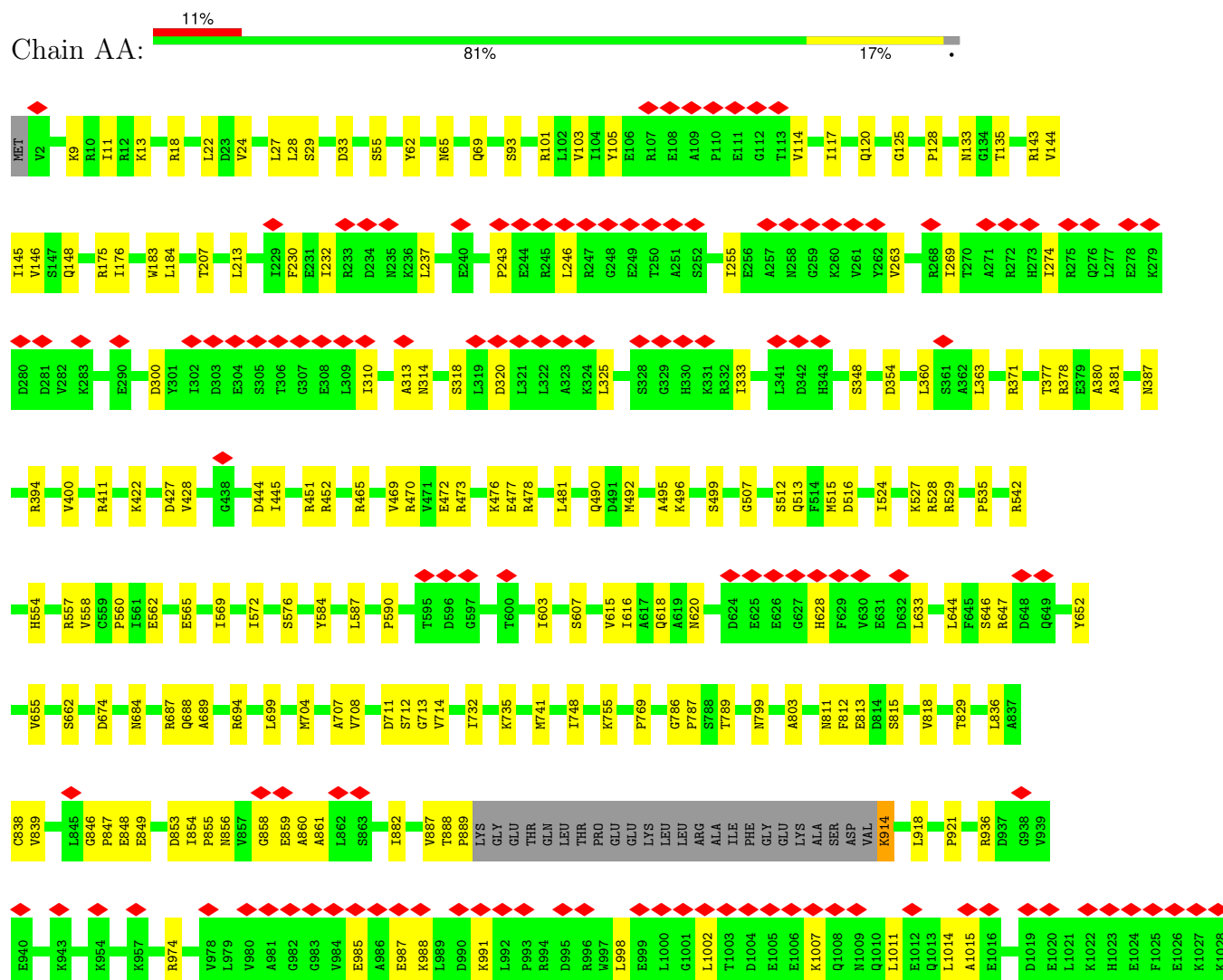
Chain A: 

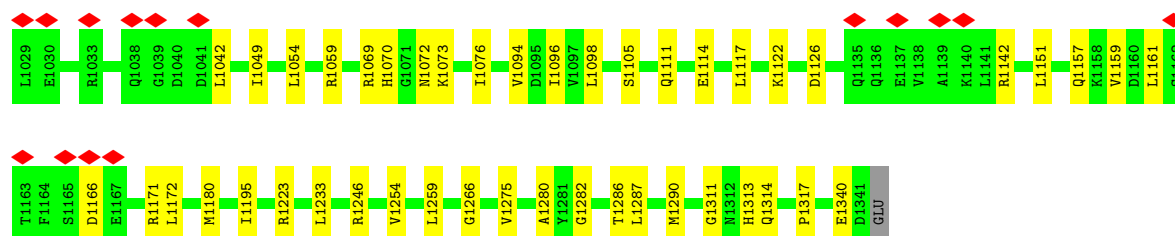


• Molecule 10: E-site and P-site tRNA (fMet)

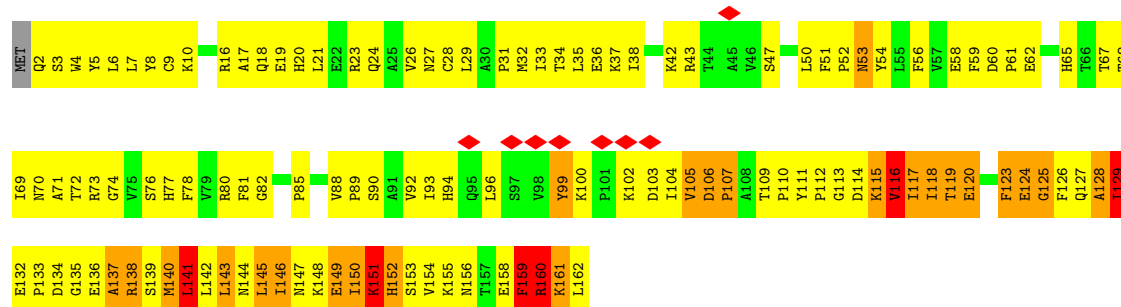


• Molecule 11: DNA-directed RNA polymerase subunit beta

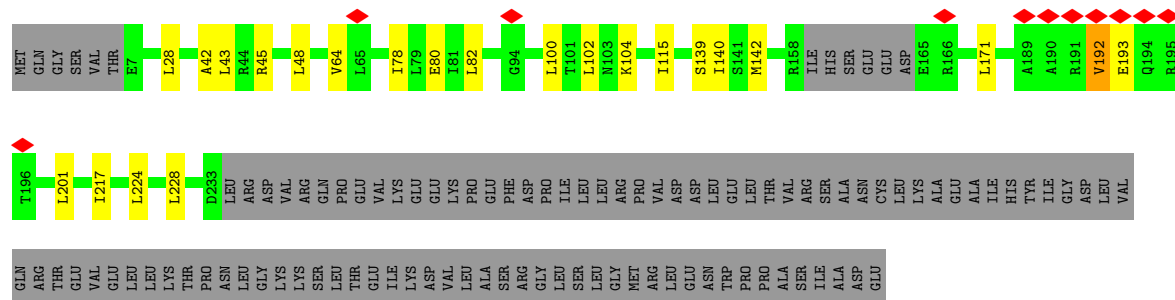




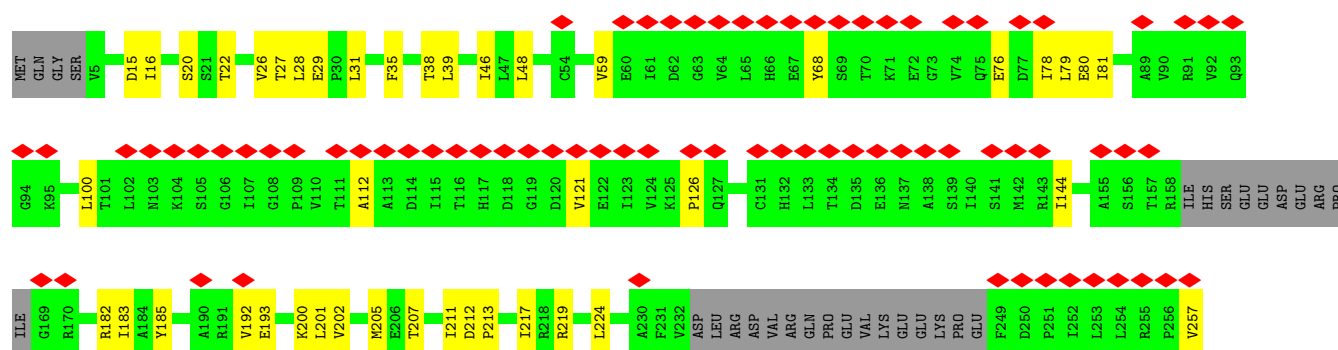
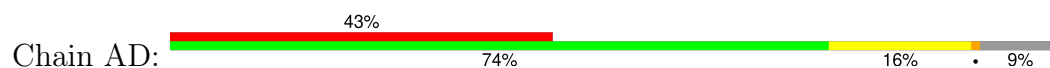
• Molecule 12: Transcription antitermination protein RfaH

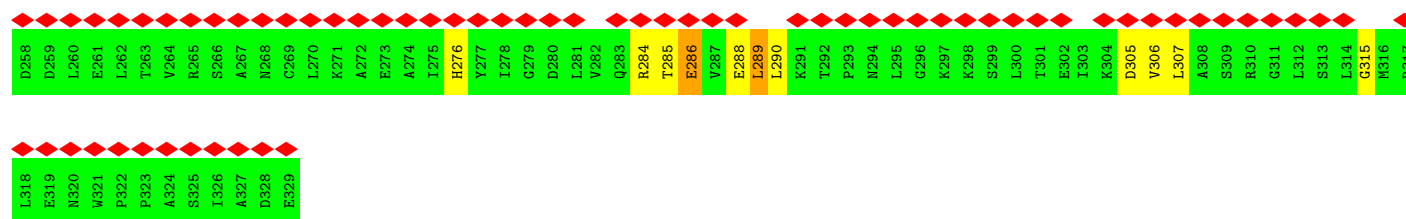


• Molecule 13: DNA-directed RNA polymerase subunit alpha

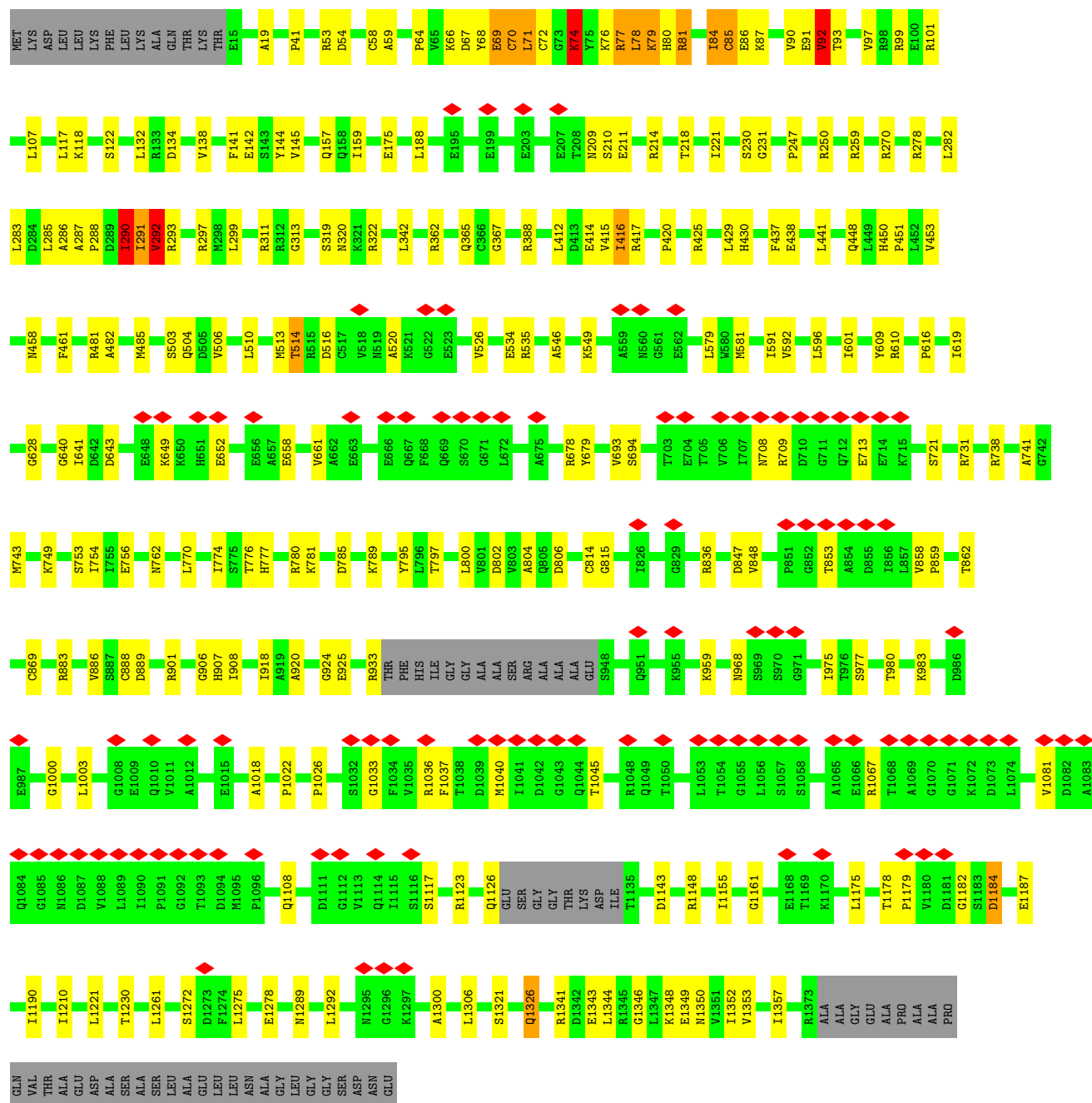
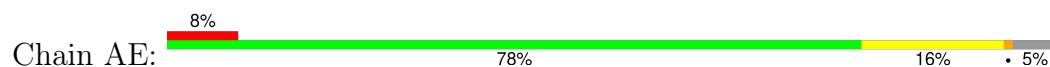


• Molecule 13: DNA-directed RNA polymerase subunit alpha

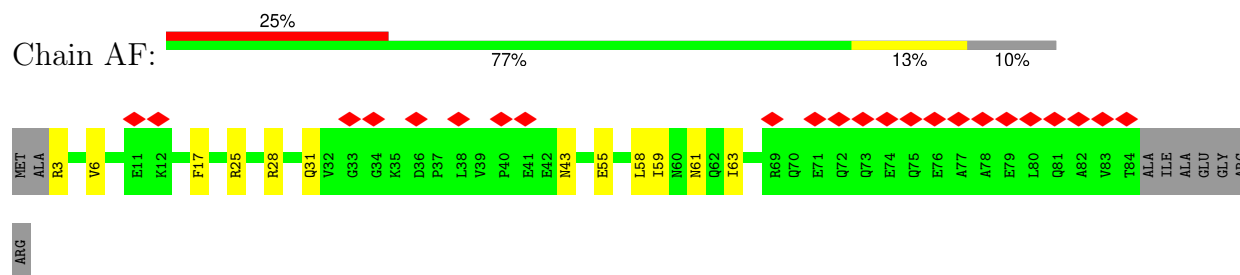




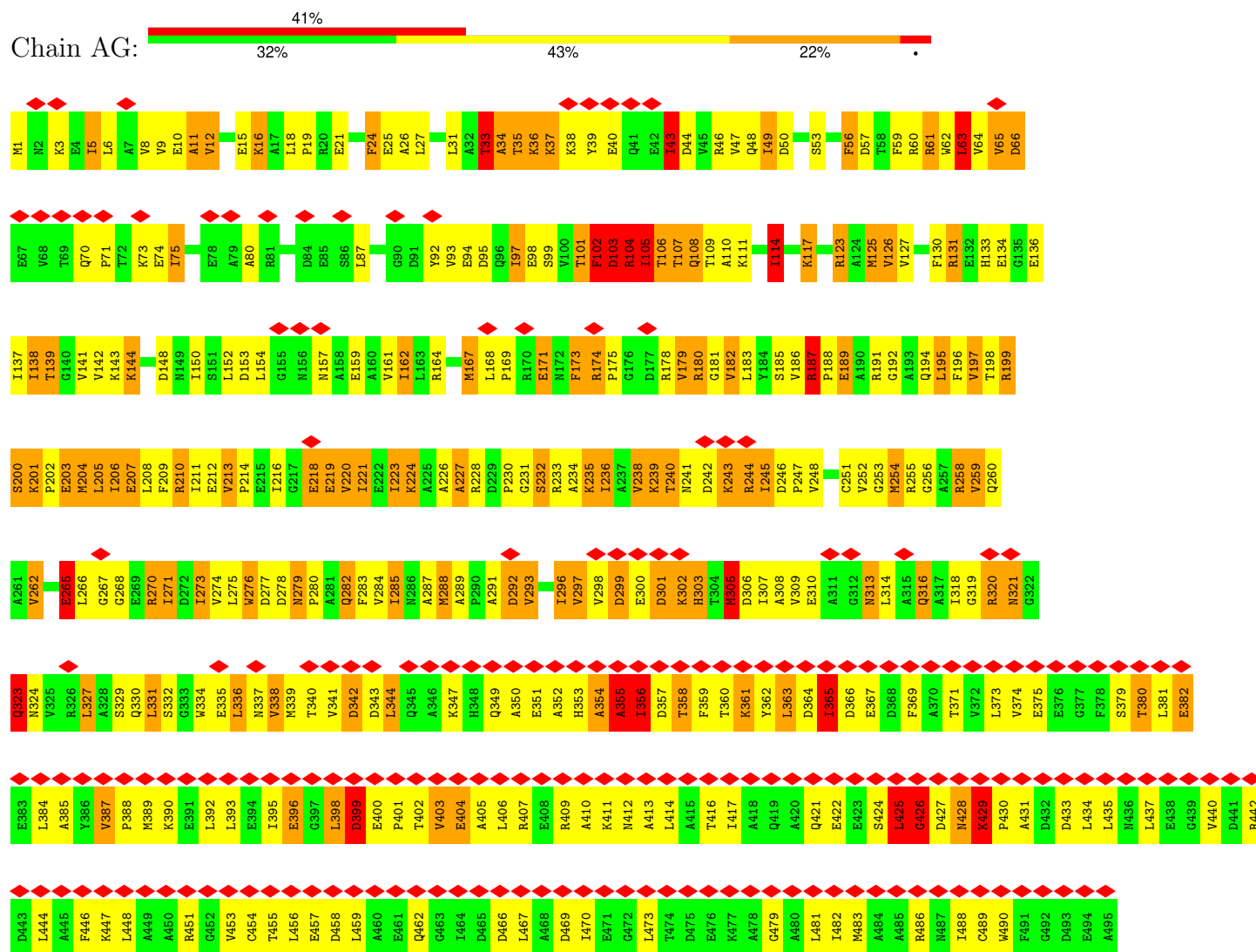
• Molecule 14: DNA-directed RNA polymerase subunit beta'



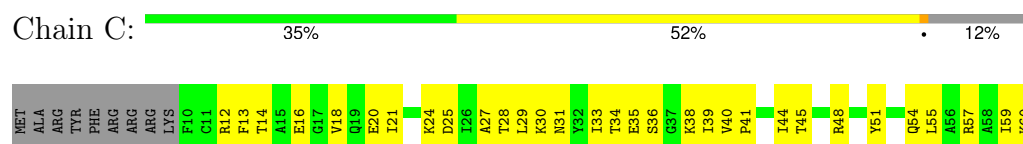
- Molecule 15: DNA-directed RNA polymerase subunit omega

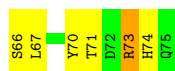


- Molecule 16: Transcription termination/antitermination protein NusA



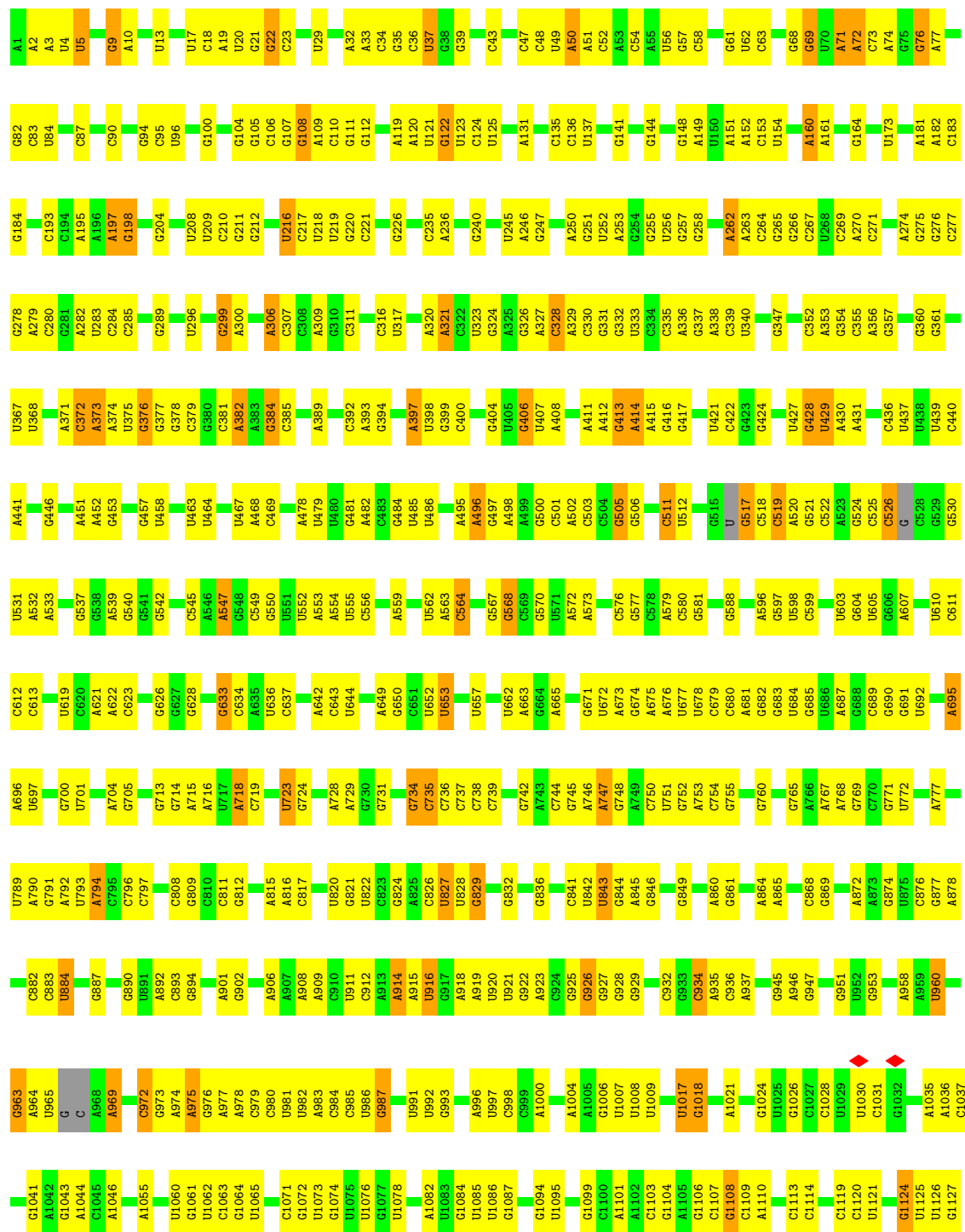
- Molecule 17: 30S ribosomal protein S18

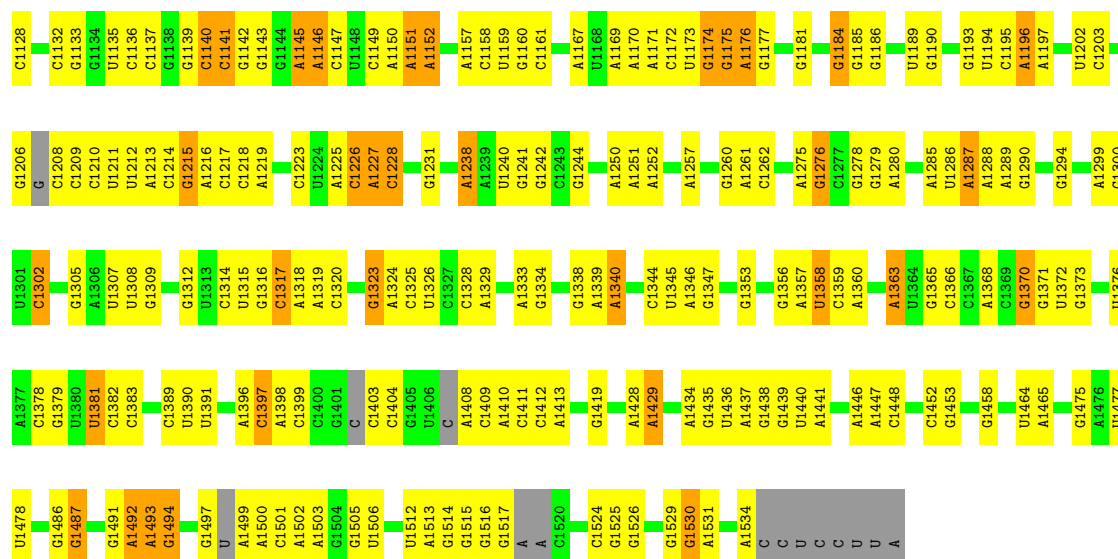




• Molecule 18: 16S rRNA

Chain D: 49% 43% 6%





• Molecule 19: 30S ribosomal protein S20

Chain E: 47% 52%



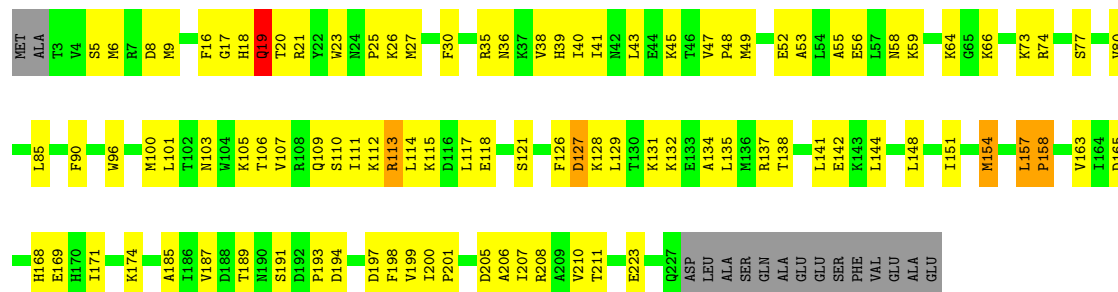
• Molecule 20: 30S ribosomal protein S21

Chain F: 41% 58%

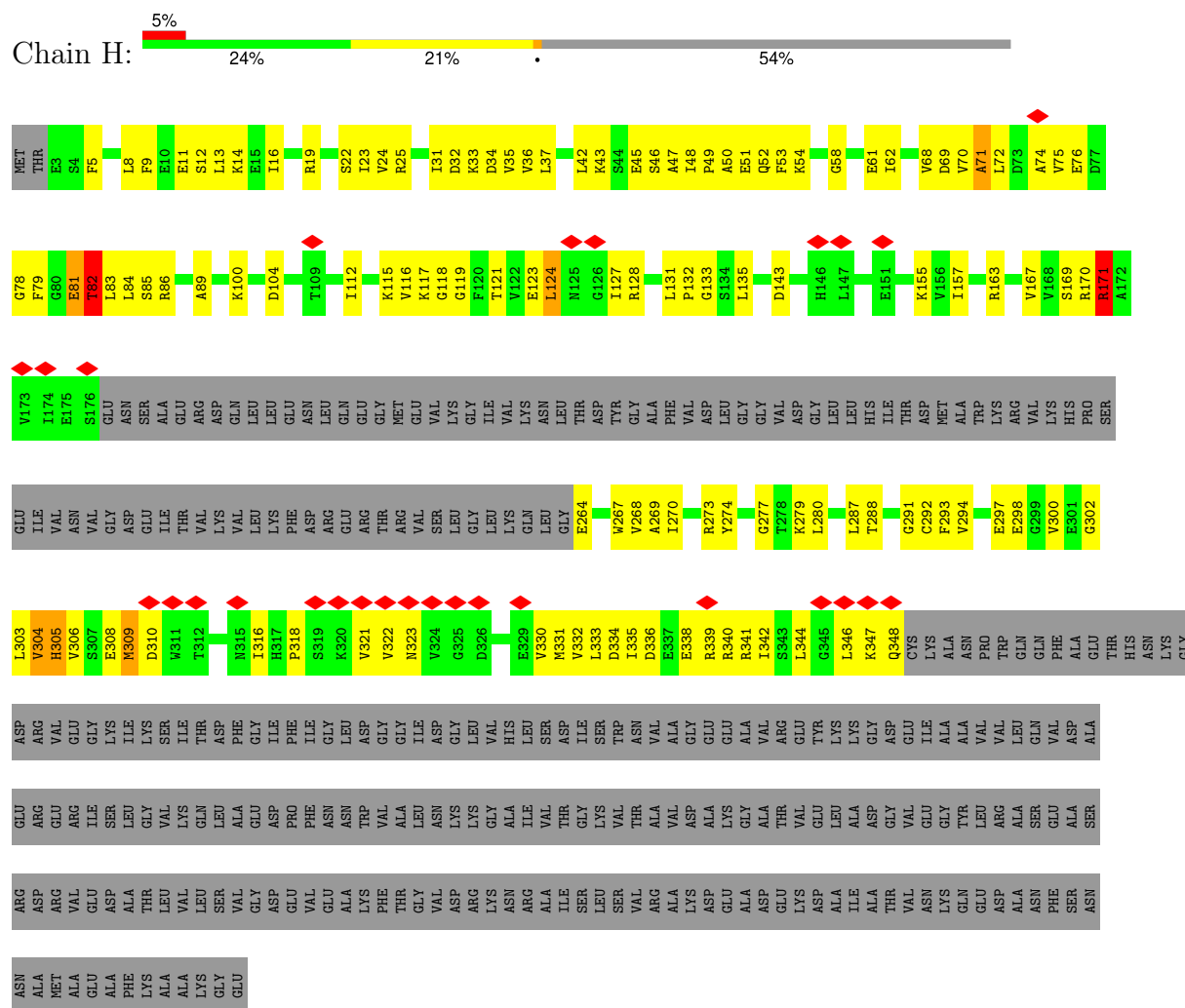


• Molecule 21: 30S ribosomal protein S2

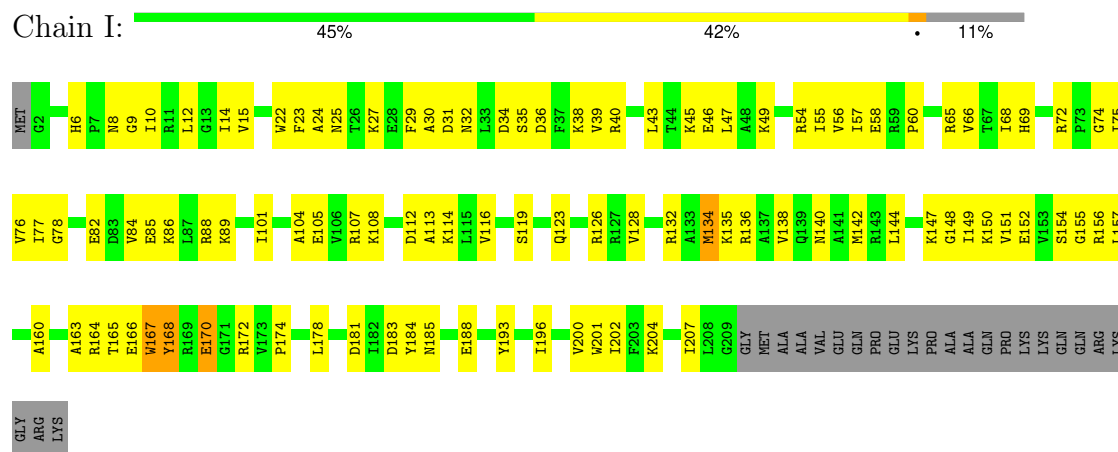
Chain G: 52% 39% 7%



• Molecule 22: 30S ribosomal protein S1

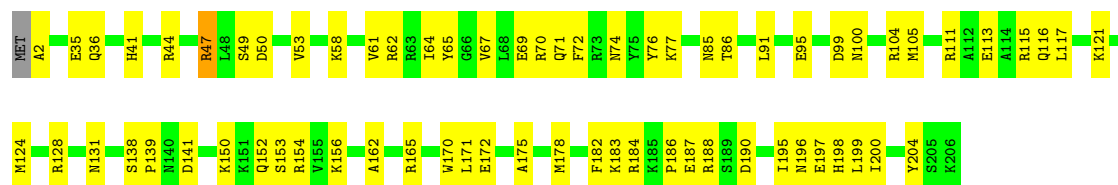


• Molecule 23: 30S ribosomal protein S3



• Molecule 24: 30S ribosomal protein S4





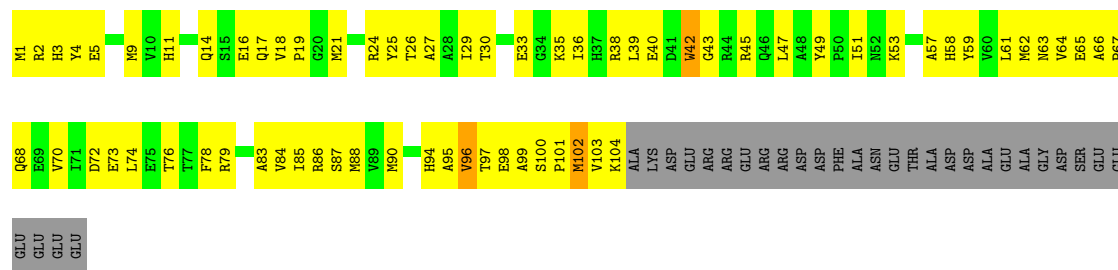
• Molecule 25: 30S ribosomal protein S5

Chain K: 44% 47% 7%



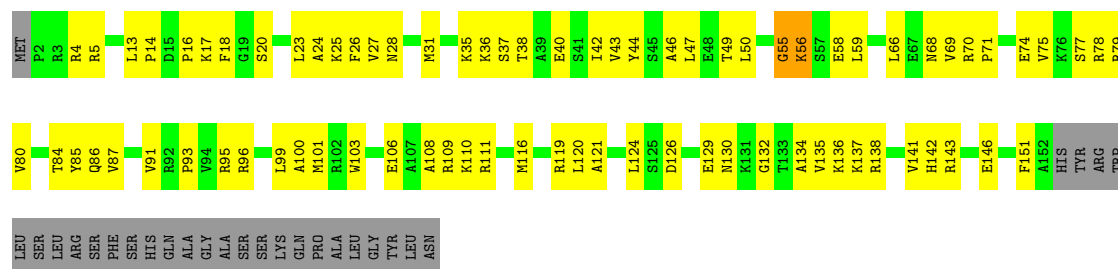
• Molecule 26: 30S ribosomal protein S6

Chain L: 27% 48% 23%



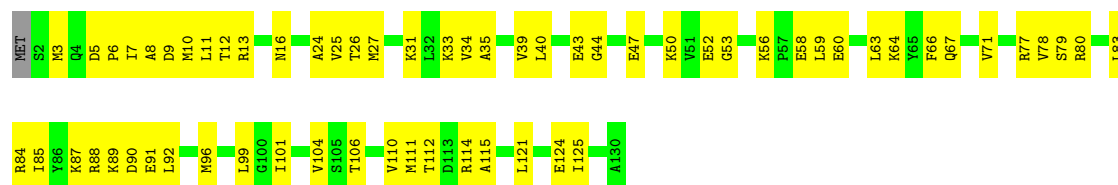
• Molecule 27: 30S ribosomal protein S7

Chain M: 41% 42% 16%

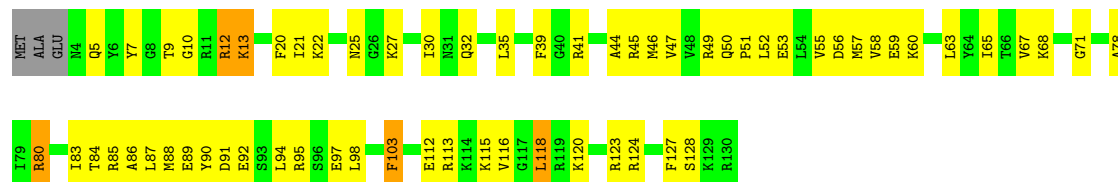


• Molecule 28: 30S ribosomal protein S8

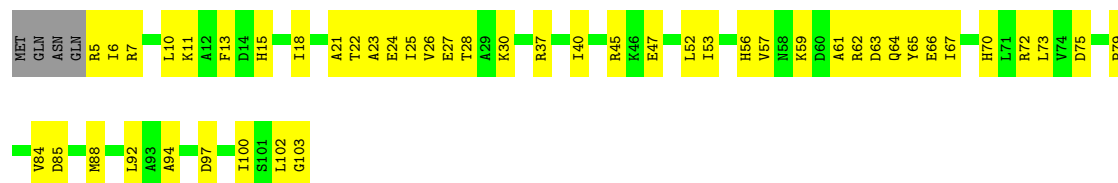
Chain N: 52% 48%



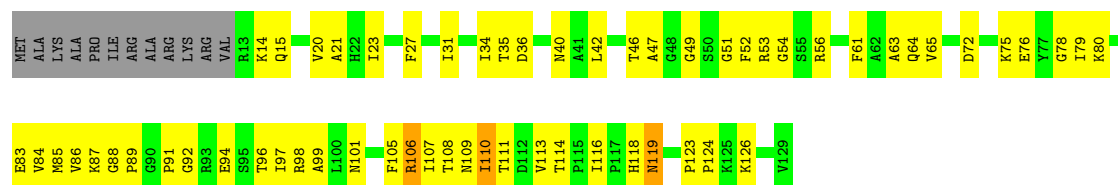
• Molecule 29: 30S ribosomal protein S9



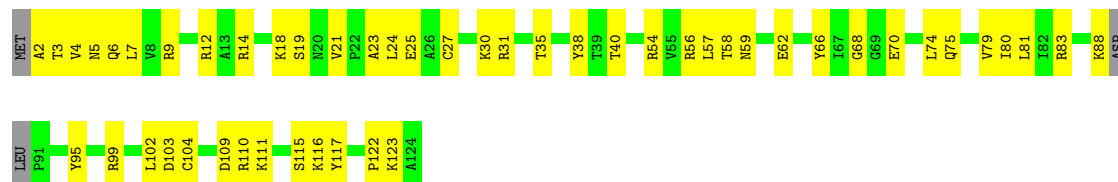
• Molecule 30: 30S ribosomal protein S10



• Molecule 31: 30S ribosomal protein S11



• Molecule 32: 30S ribosomal protein S12



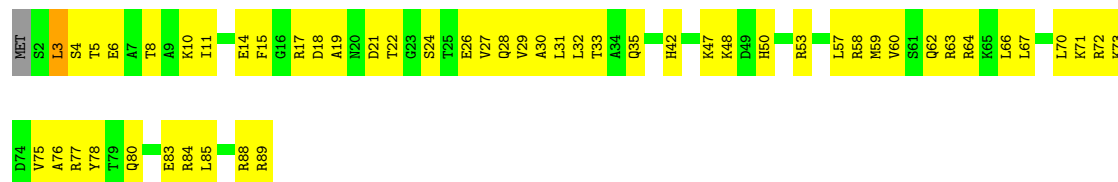
• Molecule 33: 30S ribosomal protein S14





- Molecule 34: Small ribosomal subunit protein uS15

Chain T: 40% 57% ..



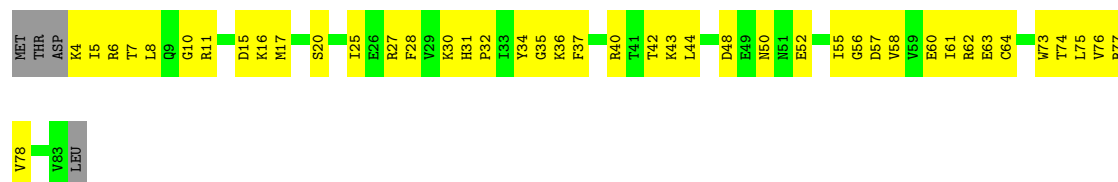
- Molecule 35: 30S ribosomal protein S16

Chain U: 54% 46%



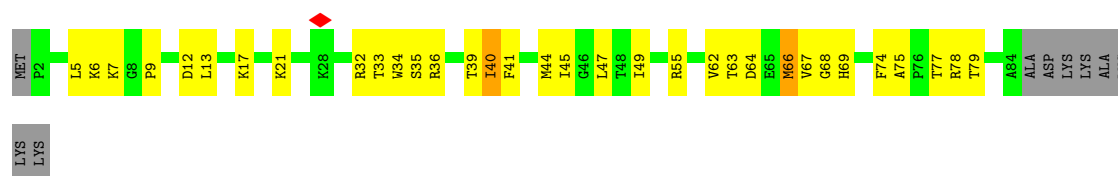
- Molecule 36: 30S ribosomal protein S17

Chain V: 44% 51% 5%



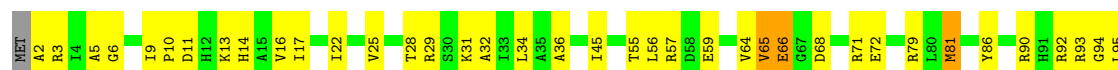
- Molecule 37: 30S ribosomal protein S19

Chain W: 54% 34% 10%



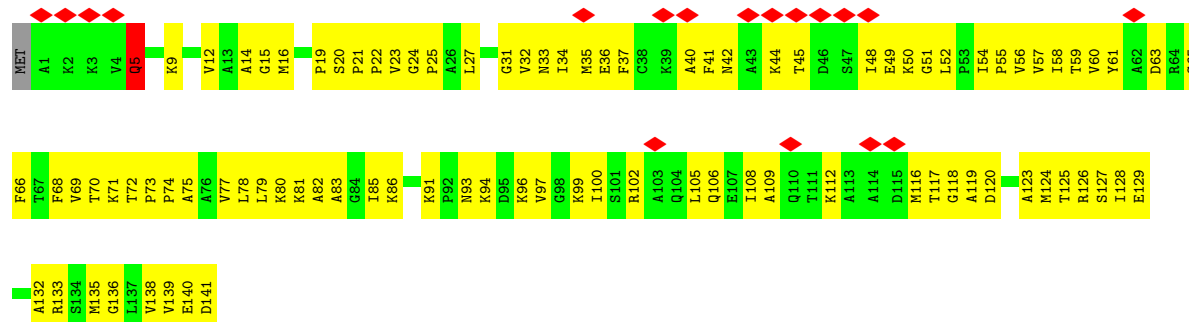
- Molecule 38: 30S ribosomal protein S13

Chain X: 55% 41%

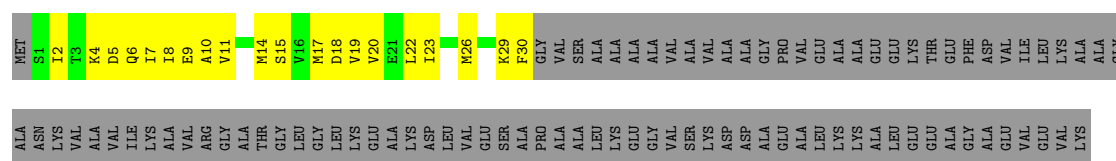




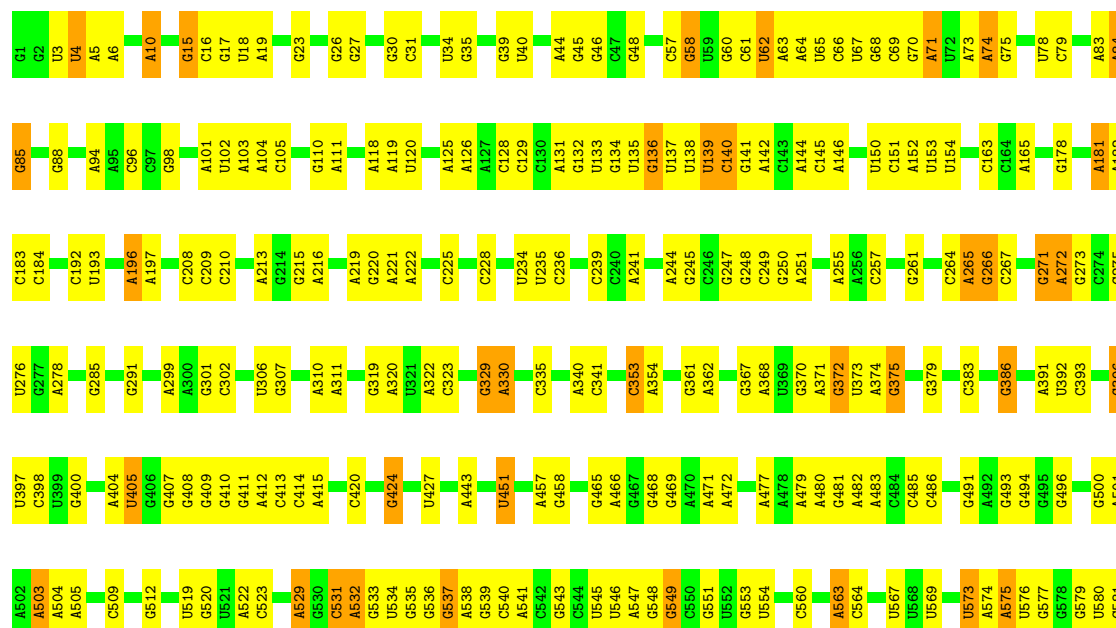
• Molecule 39: 50S ribosomal protein L11



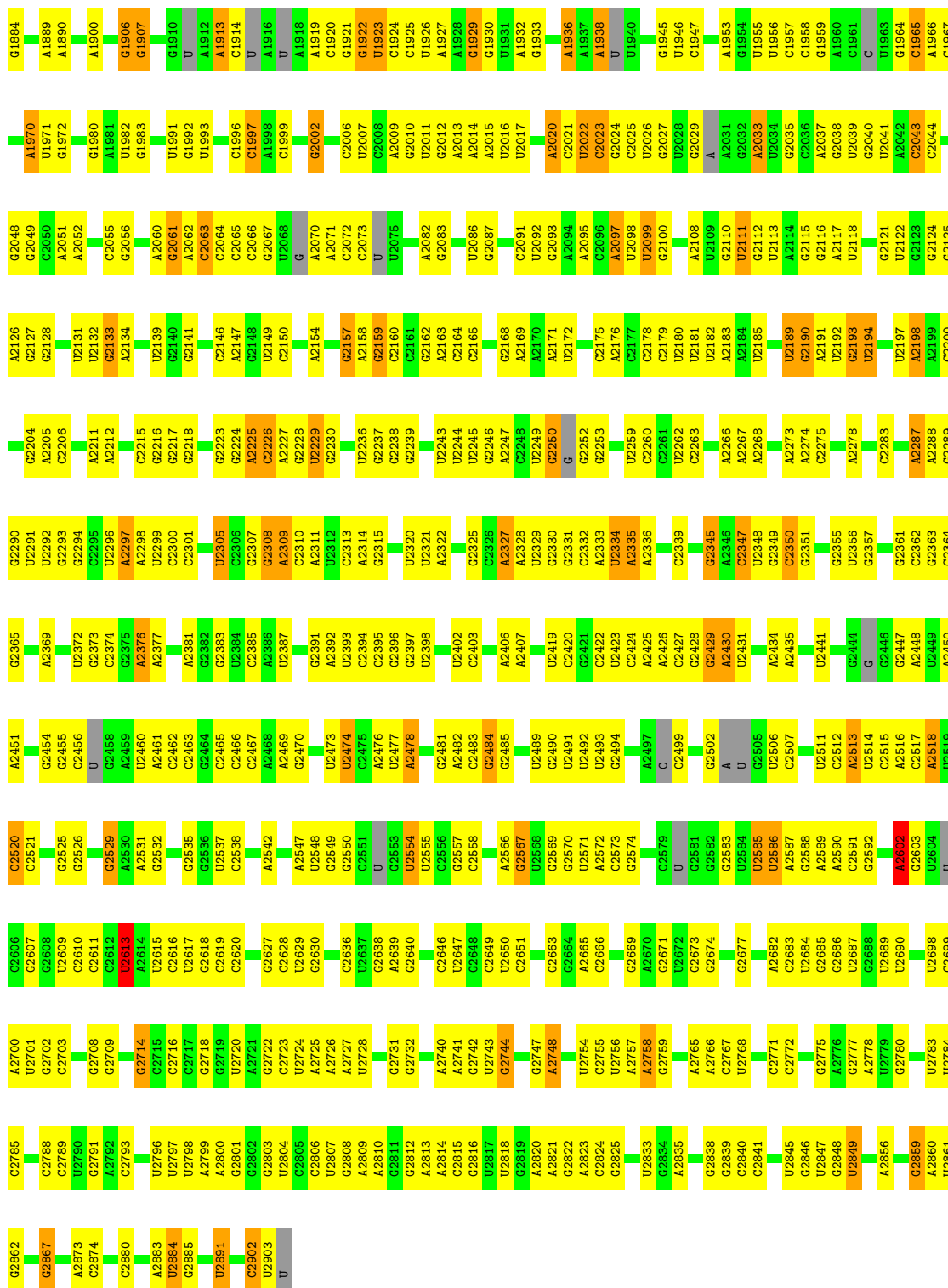
• Molecule 40: 50S ribosomal protein L7/L12



• Molecule 41: 23S rRNA



G1811	G1710	C1607	G1514	A1415	A1342	C1146	G1074	C1005	U846	A752	U646	A582
G1816	U1714	A1608	A1515	G1416	G1343	A1147	C1075	C1006	C851	A753	G647	G583
U1818	G1715	A1610	U1520	G1417	U1344	C1153	C1076	C1007	U852	U754	A654	C584
A1819	U1720	G1613	G1521	G1418	C1345	G1154	A1077	A1008	G856	U755	A654	G585
U1820	G1721	A1614	U1529	A1420	A1419	U1249	U1078	A1009	G857	G756	U657	A586
A1821	U1729	C1617	U1534	G1421	A1347	G1250	A1080	U1012	G858	G760	U658	U588
G1822	C1730	A	C1428	G1422	C1348	G1252	U1081	U1013	G859	A761	G659	U589
G1823	A1535	G1619	U1536	A1427	C1349	A1253	U1082	C1013	G860	A762	A666	U591
U1825	C1732	G1622	G1537	G1429	U1352	C1257	A1083	A1014	U861	A764	U667	A592
G1826	G1738	U1636	U1542	G1430	A1353	A1262	A1084	U1019	G862	C765	A668	U593
G1828	A1746	A1637	G1543	A1431	A1354	A1262	A1086	A1020	A670	U773	A670	U594
C1830	U1747	U1647	G1546	A1432	G1355	G1266	G1087	A1021	G869	G774	C671	C595
G1831	G1750	U1648	U1554	A1433	G1356	A1269	A1088	U1022	U870	G775	C672	U596
C1832	G1753	G1649	U1558	A1434	A1360	C1270	A1090	G1023	U871	G776	C672	U598
U1834	A1754	A1650	C1558	G1441	G1361	A1271	A1095	U1026	U872	A777	A677	A599
G1836	A1755	G1651	U1559	U1442	C1362	A1272	A1096	A1027	G874	G780	A685	G600
C1837	A1756	A1652	G1560	U1443	A1365	U1273	U1101	A1028	A878	A781	U686	C601
G1838	G1756	G1653	A1561	G1444	A1366	C1276	C1102	A1029	G879	A782	U686	A602
A1839	A1757	A1654	U1562	G1445	A1367	A1277	A1103	U1033	G880	G783	A689	A603
G1840	U1758	A1655	U1563	G1452	G1368	G1277	U1181	G1036	G881	G785	G690	U607
U1841	A1759	C1656	C1564	U1453	G1369	C1278	G1106	U1037	U884	C786	C691	A608
C1842	C1764	U1657	U1565	U1454	A1370	C1278	G1107	G1037	C885	C787	A609	A609
G1843	G1764	C1658	A1566	G1454	A1371	G1279	U1108	U1038	C886	A789	C610	C610
G1845	U1765	G1659	U1567	G1455	A1378	G1281	C1109	A1039	G888	G792	G701	A613
G1846	G1770	U1659	C1568	U1460	U1379	U1282	G1110	A1040	C889	U703	U702	A614
A1847	A1773	A1664	A1569	U1467	G1380	G1283	A1111	G1041	C890	C964	G704	U615
U1848	C1774	G1666	U1570	U1468	A1383	C1289	G1112	G1042	C891	A705	A705	A616
G1853	U1778	A1668	A1572	A1470	A1385	C1290	G1113	C1043	A892	C806	A706	G617
A1854	U1779	A1674	U1578	A1477	A1386	C1296	G1115	C1044	U893	U807	G707	G620
U1855	G1784	C1675	A1579	G1482	A1387	G1296	U1119	C1045	U895	U811	U710	A621
G1856	A1785	A1676	G1580	U1482	A1392	G1300	G1120	C1046	C896	C812	G622	A622
G1857	A1786	A1677	C1581	A1490	U1394	A1301	C1121	A1047	U897	U813	U714	C623
U1859	U1789	G1681	A1583	A1495	A1395	G1310	G1122	G1048	A900	C814	C717	C624
G1862	C1790	U1682	U1584	A1496	U1396	U1312	G1125	C1053	G818	G815	U717	G625
G1863	A1791	U1683	C1585	A1496	U1397	U1312	U1128	G1055	A819	A819	G726	G628
U1864	G1792	G1684	G1587	A1497	C1398	U1313	G1131	A1056	U826	U827	G728	G629
G1869	C1793	U1688	U1588	C1498	U1399	C1314	U1132	U1058	U827	U827	G729	G630
C1870	A1794	U1688	U1589	C1499	U1400	C1315	A1133	G1059	C908	U828	G731	A631
U1871	C1795	U1688	A1590	G1500	A1403	U1318	A1134	U1060	A909	U832	A633	A632
A1872	U1796	U1693	A1591	A1503	C1404	C1319	C1135	U1061	A910	U833	G738	C634
G1873	G1797	G1695	C1592	U1506	U1405	C1320	G1136	A983	C912	A833	G738	C635
C1874	U1798	A1700	A1597	U1507	U1406	A1321	U1137	A984	C913	A834	A742	G636
G1875	C1800	A1700	A1598	C1507	G1407	A1322	G1138	C985	U913	A743	A743	A637
A1801	A1801	G1703	U1599	U1508	G1408	C1323	U1139	C986	G914	U744	G744	G638
A1803	A1802	U1600	A1600	A1509	U1409	G1333	G1140	A996	C915	G	U	U639
U1880	U1802	G1601	U1601	G1510	U1411	G1334	U1141	C997	G916	U	U	C640
C1881	A1808	U1602	U1602	G1511	U1412	G1334	A1142	C998	A920	G748	G748	U642
U1882	A1808	A1603	A1413	G1512	A1412	U1340	A1143	U999	C921	A749	A749	A643
U1883	A1808	U1709	C1414	U1513	C1414	G1341	C1145	A1001	G923	U842	A751	A644
										A845		C645



• Molecule 42: 50S ribosomal protein L27

Chain b:



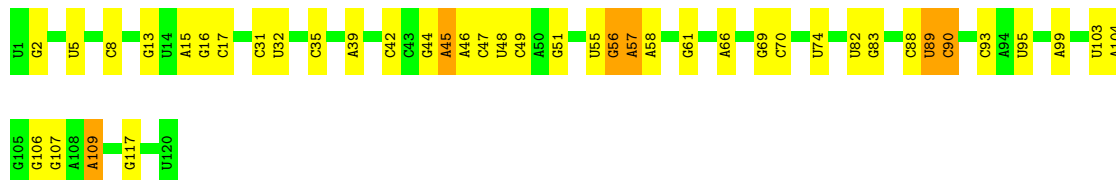
- Molecule 43: 50S ribosomal protein L28

Chain c: 



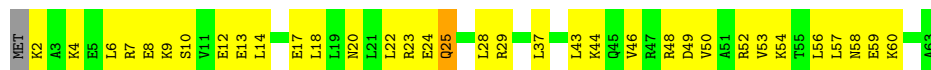
- Molecule 44: 5S rRNA

Chain d: 



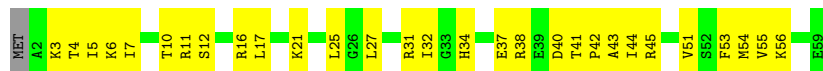
- Molecule 45: 50S ribosomal protein L29

Chain e: 



- Molecule 46: 50S ribosomal protein L30

Chain f: 



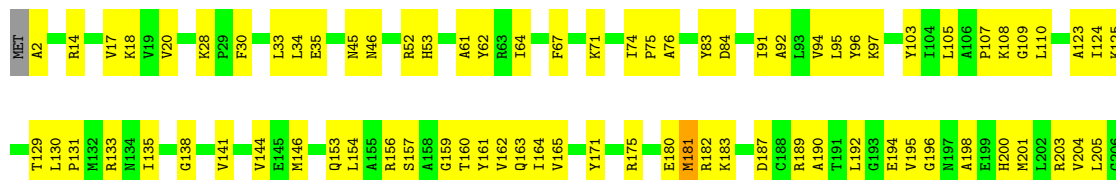
- Molecule 47: 50S ribosomal protein L31

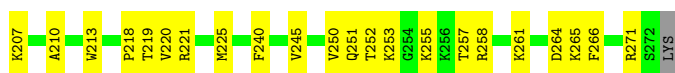
Chain g: 



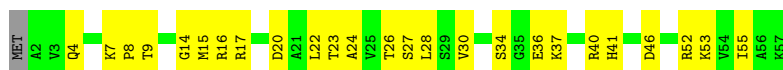
- Molecule 48: 50S ribosomal protein L2

Chain h: 

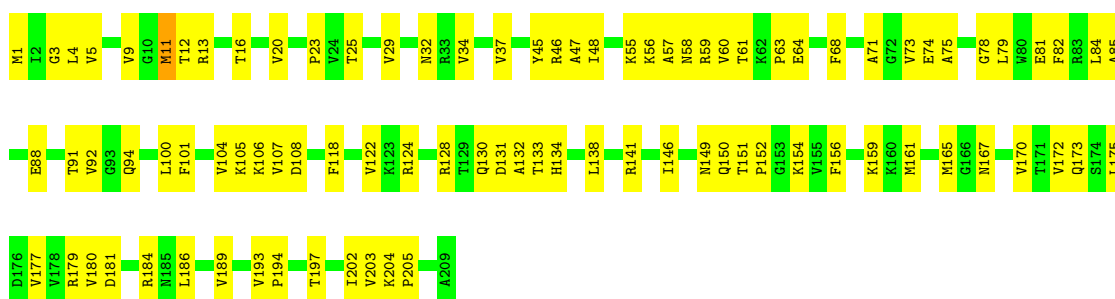




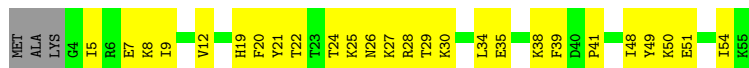
- Molecule 49: 50S ribosomal protein L32



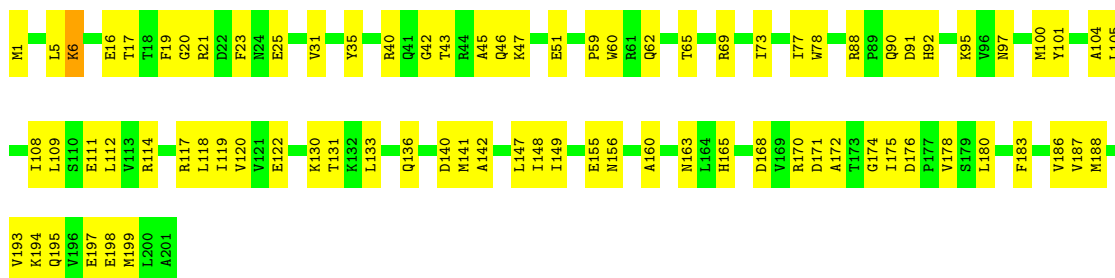
- Molecule 50: 50S ribosomal protein L3



- Molecule 51: 50S ribosomal protein L33



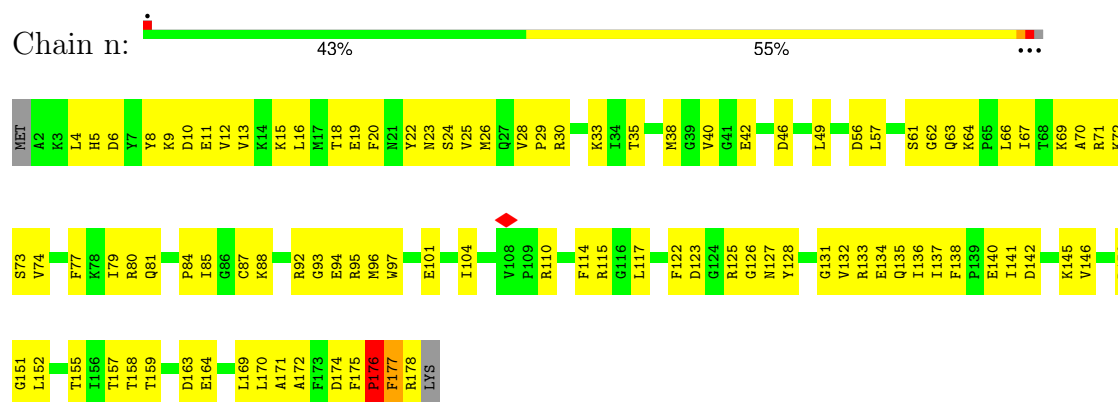
- Molecule 52: 50S ribosomal protein L4



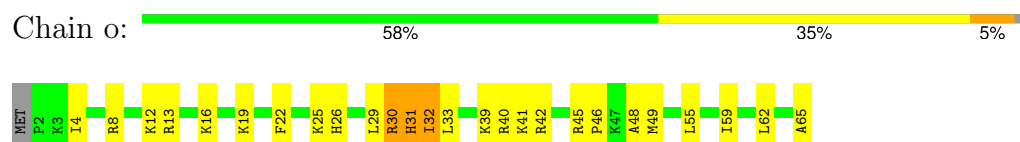
- Molecule 53: 50S ribosomal protein L34



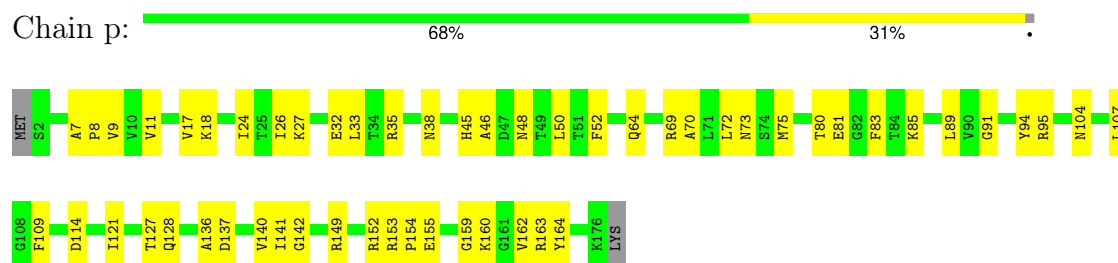
- Molecule 54: 50S ribosomal protein L5



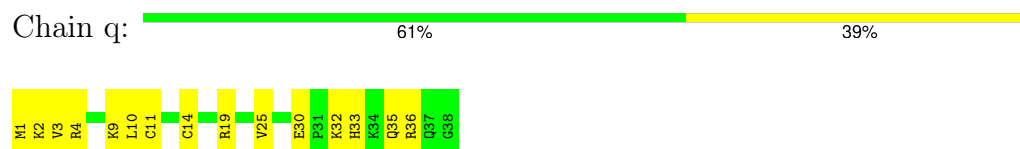
- Molecule 55: 50S ribosomal protein L35



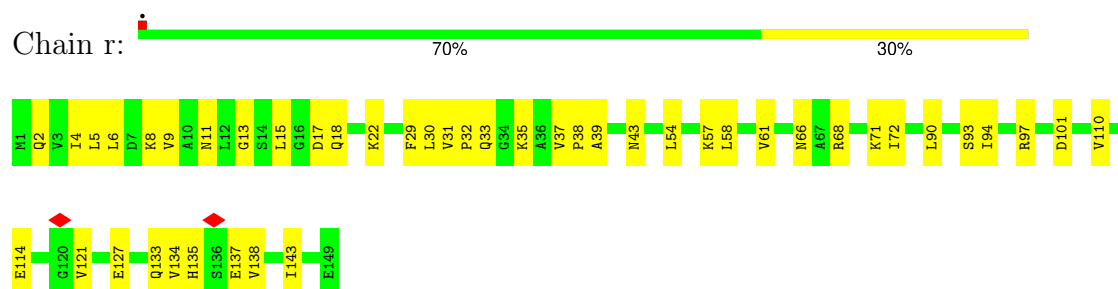
- Molecule 56: 50S ribosomal protein L6



- Molecule 57: 50S ribosomal protein L36

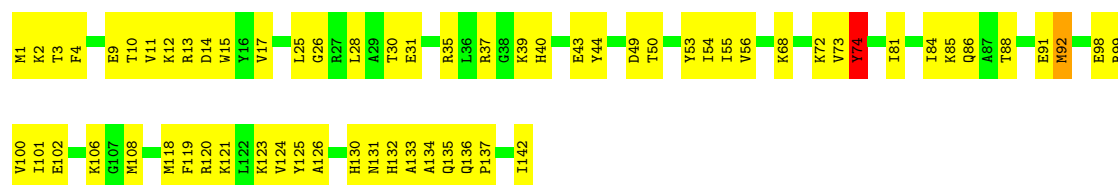


- Molecule 58: 50S ribosomal protein L9



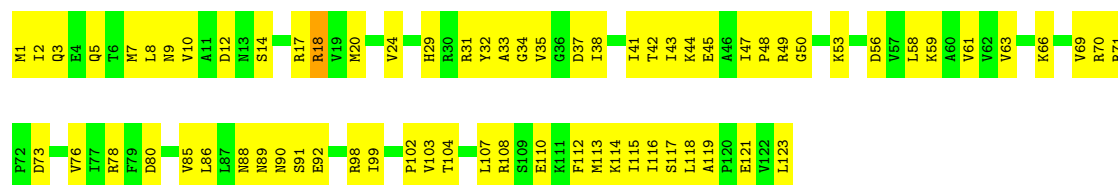
- Molecule 59: 50S ribosomal protein L13

Chain s:  55% 44% ..



• Molecule 60: 50S ribosomal protein L14

Chain t:  43% 56% .



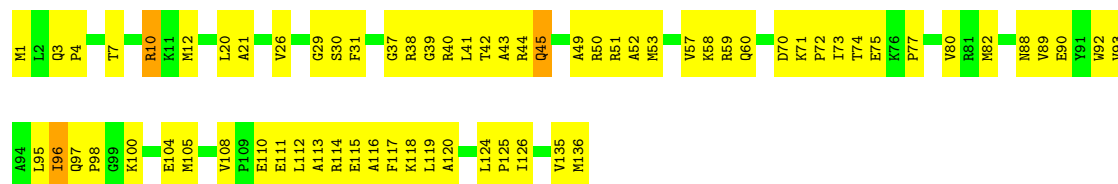
• Molecule 61: 50S ribosomal protein L15

Chain u:  62% 37% .

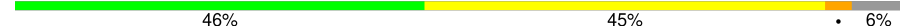


• Molecule 62: 50S ribosomal protein L16

Chain v:  50% 48% .

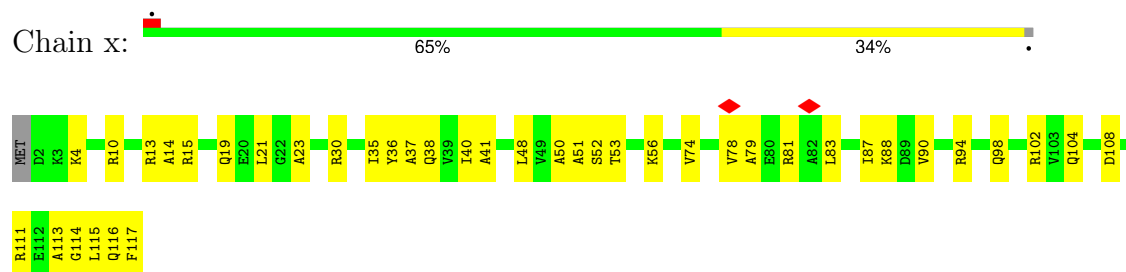


• Molecule 63: 50S ribosomal protein L17

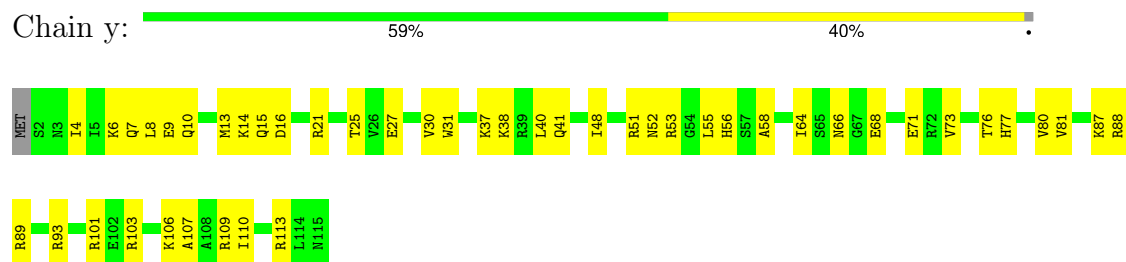
Chain w:  46% 45% 6% .



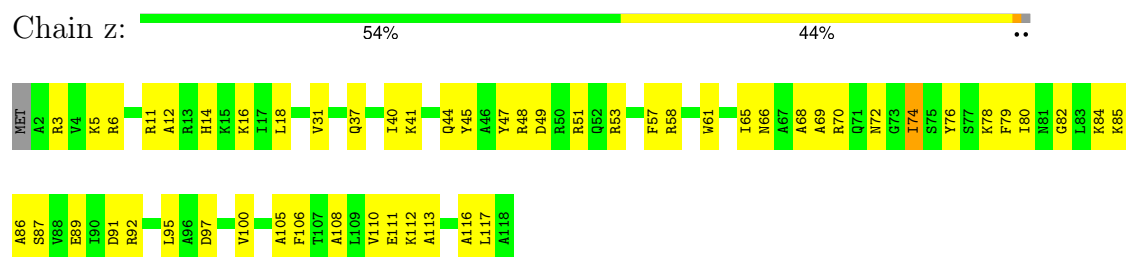
• Molecule 64: 50S ribosomal protein L18



- Molecule 65: 50S ribosomal protein L19



- Molecule 66: 50S ribosomal protein L20



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, Not provided	
Number of particles used	20048	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 ($e^-/\text{\AA}^2$)	50	Depositor
Minimum defocus (nm)	1250	Depositor
Maximum defocus (nm)	2500	Depositor
Magnification	Not provided	
Image detector	GATAN K2 SUMMIT (4k x 4k)	Depositor
Maximum map value	0.035	Depositor
Minimum map value	-0.007	Depositor
Average map value	-0.000	Depositor
Map value standard deviation	0.002	Depositor
Recommended contour level	0.0032	Depositor
Map size ($\text{\AA}$)	532.48, 532.48, 532.48	wwPDB
Map dimensions	512, 512, 512	wwPDB
Map angles ($^\circ$)	90.0, 90.0, 90.0	wwPDB
Pixel spacing ($\text{\AA}$)	1.04, 1.04, 1.04	Depositor

5 Model quality ⓘ

5.1 Standard geometry ⓘ

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

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

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# $ Z > 5$	RMSZ	# $ Z > 5$
1	0	0.43	0/829	0.51	0/1107
2	1	0.60	0/864	0.63	0/1156
3	2	0.55	1/752 (0.1%)	0.67	0/1005
4	3	0.35	0/796	0.48	0/1062
5	4	0.57	0/766	0.63	0/1025
6	5	0.57	0/816	0.78	0/1259
7	6	0.57	0/783	0.78	0/1203
8	7	0.61	2/585 (0.3%)	0.91	3/906 (0.3%)
9	9	0.37	0/1131	0.73	2/1524 (0.1%)
10	A	0.27	0/1810	0.58	0/2821
10	B	0.27	0/1810	0.58	0/2821
11	AA	0.38	0/10547	0.62	0/14232
12	AB	0.60	0/1317	1.21	17/1786 (1.0%)
13	AC	0.38	0/1718	0.62	0/2328
13	AD	0.37	0/2096	0.69	3/2854 (0.1%)
14	AE	0.39	0/10561	0.64	5/14258 (0.0%)
15	AF	0.30	0/652	0.61	0/879
16	AG	0.97	6/3897 (0.2%)	1.40	49/5273 (0.9%)
17	C	0.68	0/553	0.81	1/743 (0.1%)
18	D	0.29	0/36610	0.41	9/57091 (0.0%)
19	E	0.61	0/675	0.72	0/895
20	F	0.53	0/597	0.61	0/792
21	G	0.68	5/1791 (0.3%)	0.79	8/2413 (0.3%)
22	H	0.38	0/1746	0.81	0/2382
23	I	0.57	1/1663 (0.1%)	0.68	4/2241 (0.2%)
24	J	0.46	0/1665	0.58	0/2227
25	K	0.72	2/1165 (0.2%)	0.86	6/1568 (0.4%)
26	L	0.68	1/867 (0.1%)	0.79	1/1171 (0.1%)
27	M	0.56	0/1195	0.69	2/1602 (0.1%)
28	N	0.50	0/989	0.58	0/1326
29	O	0.66	3/1034 (0.3%)	0.79	3/1375 (0.2%)
30	P	0.51	0/800	0.73	0/1082

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
31	Q	0.72	2/893 (0.2%)	0.76	2/1205 (0.2%)
32	R	0.49	0/952	0.59	0/1274
33	S	0.52	0/817	0.61	0/1088
34	T	0.57	1/722 (0.1%)	0.68	0/964
35	U	0.41	0/659	0.53	0/884
36	V	0.49	0/657	0.62	0/881
37	W	0.55	1/680 (0.1%)	0.62	1/915 (0.1%)
38	X	0.45	0/909	0.72	1/1215 (0.1%)
39	Y	0.38	2/1046 (0.2%)	0.66	2/1410 (0.1%)
40	Z	0.13	0/227	0.41	0/304
41	a	0.30	0/69247	0.43	10/107985 (0.0%)
42	b	0.41	0/589	0.55	0/779
43	c	0.51	0/635	0.61	0/848
44	d	0.25	0/2872	0.37	0/4478
45	e	0.69	2/502 (0.4%)	0.65	1/667 (0.1%)
46	f	0.52	0/452	0.57	0/605
47	g	0.45	1/531 (0.2%)	0.64	0/709
48	h	0.49	1/2121 (0.0%)	0.59	1/2852 (0.0%)
49	i	0.41	0/450	0.55	0/599
50	j	0.51	0/1586	0.57	1/2134 (0.0%)
51	k	0.45	0/433	0.63	0/576
52	l	0.51	1/1571 (0.1%)	0.62	0/2113
53	m	0.45	0/380	0.56	0/498
54	n	0.46	0/1434	0.66	1/1926 (0.1%)
55	o	0.51	0/513	0.71	1/676 (0.1%)
56	p	0.41	0/1333	0.62	2/1805 (0.1%)
57	q	0.45	0/303	0.64	0/397
58	r	0.28	0/1122	0.48	0/1515
59	s	0.70	2/1152 (0.2%)	0.74	2/1551 (0.1%)
60	t	0.52	0/955	0.75	3/1279 (0.2%)
61	u	0.43	0/1062	0.59	0/1413
62	v	0.60	1/1093 (0.1%)	0.77	3/1460 (0.2%)
63	w	0.83	2/964 (0.2%)	0.80	3/1289 (0.2%)
64	x	0.34	0/902	0.50	0/1209
65	y	0.41	0/929	0.54	0/1242
66	z	0.62	2/960 (0.2%)	0.62	0/1278
All	All	0.41	39/194733 (0.0%)	0.56	147/286430 (0.1%)

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

Mol	Chain	#Chirality outliers	#Planarity outliers
9	9	0	3
12	AB	0	2
13	AC	0	1
13	AD	0	2
14	AE	0	4
16	AG	0	6
21	G	0	1
22	H	0	5
23	I	0	1
25	K	0	2
29	O	0	1
38	X	0	1
39	Y	0	1
54	n	0	1
55	o	0	1
59	s	0	1
61	u	0	2
All	All	0	35

All (39) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
16	AG	429	LYS	C-N	13.90	1.51	1.33
63	w	35	LYS	CE-NZ	-12.73	1.11	1.49
29	O	80	ARG	CD-NE	-10.50	1.31	1.46
63	w	42	LYS	CD-CE	-9.41	1.24	1.52
25	K	45	ARG	CG-CD	7.73	1.75	1.52
66	z	74	ILE	CG1-CD1	7.60	1.81	1.51
8	7	52	G	C1'-N9	-7.53	1.36	1.48
21	G	19	GLN	CB-CG	-7.52	1.29	1.52
29	O	118	LEU	CG-CD2	-7.32	1.28	1.52
37	W	66	MET	CG-SD	-7.20	1.62	1.80
26	L	42	TRP	CB-CG	-7.15	1.28	1.50
59	s	74	TYR	CZ-OH	-7.15	1.23	1.38
45	e	46	VAL	CB-CG1	-7.12	1.29	1.52
8	7	2	U	C1'-N1	7.02	1.58	1.48
3	2	5	GLU	CG-CD	-7.00	1.34	1.52
16	AG	246	ASP	C-N	7.00	1.50	1.33
31	Q	106	ARG	CD-NE	-6.41	1.37	1.46
21	G	158	PRO	CA-C	-6.27	1.44	1.52
16	AG	355	ALA	N-CA	6.26	1.54	1.46
29	O	80	ARG	CZ-NH2	6.25	1.41	1.33
66	z	74	ILE	CB-CG2	-6.04	1.32	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
48	h	181	MET	CG-SD	-6.03	1.65	1.80
21	G	158	PRO	CG-CD	6.02	1.71	1.50
39	Y	5	GLN	CB-CG	-6.02	1.34	1.52
59	s	92	MET	CG-SD	-5.95	1.65	1.80
16	AG	355	ALA	CA-C	5.88	1.60	1.52
25	K	45	ARG	CD-NE	5.83	1.54	1.46
52	l	6	LYS	CE-NZ	-5.79	1.31	1.49
21	G	157	LEU	C-O	-5.77	1.15	1.24
34	T	3	LEU	CG-CD2	-5.61	1.34	1.52
39	Y	5	GLN	CA-C	-5.58	1.45	1.52
47	g	62	LYS	CE-NZ	-5.53	1.32	1.49
45	e	25	GLN	CB-CG	-5.37	1.36	1.52
23	I	134	MET	CB-CG	-5.30	1.36	1.52
21	G	19	GLN	CD-NE2	-5.14	1.22	1.33
31	Q	119	ASN	CG-ND2	5.12	1.44	1.33
62	v	45	GLN	CB-CG	-5.12	1.37	1.52
16	AG	99	SER	N-CA	5.10	1.49	1.46
16	AG	293	VAL	N-CA	5.08	1.52	1.46

All (147) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
16	AG	104	ARG	CA-C-N	17.96	154.29	121.97
16	AG	104	ARG	C-N-CA	17.96	154.29	121.97
16	AG	246	ASP	CA-C-N	14.50	137.97	119.84
16	AG	246	ASP	C-N-CA	14.50	137.97	119.84
16	AG	354	ALA	O-C-N	-13.70	107.29	122.09
16	AG	429	LYS	CA-C-N	11.46	132.19	119.92
16	AG	429	LYS	C-N-CA	11.46	132.19	119.92
25	K	71	MET	CG-SD-CE	-11.36	75.91	100.90
21	G	158	PRO	N-CD-CG	-10.97	86.75	103.20
16	AG	24	PHE	CA-CB-CG	-10.56	103.24	113.80
16	AG	105	ILE	CA-C-N	10.17	136.87	120.60
16	AG	105	ILE	C-N-CA	10.17	136.87	120.60
12	AB	141	LEU	CA-C-N	-9.96	106.41	122.34
12	AB	141	LEU	C-N-CA	-9.96	106.41	122.34
26	L	102	MET	CA-CB-CG	-9.61	94.87	114.10
59	s	92	MET	CG-SD-CE	-9.37	80.28	100.90
21	G	113	ARG	NE-CZ-NH2	9.23	127.51	119.20
16	AG	103	ASP	CA-CB-CG	9.00	121.60	112.60
8	7	1	A	O3'-P-O5'	-8.64	91.03	104.00
25	K	78	ASN	N-CA-CB	-8.52	96.09	110.49

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
12	AB	129	ILE	N-CA-C	8.41	120.22	107.77
8	7	4	U	C2'-C3'-O3'	8.29	121.93	109.50
16	AG	35	THR	CA-CB-OG1	-8.25	97.23	109.60
16	AG	292	ASP	O-C-N	-8.21	107.49	122.44
16	AG	109	THR	CB-CA-C	8.18	124.75	110.85
16	AG	108	GLN	N-CA-CB	8.08	123.56	110.40
60	t	18	ARG	NE-CZ-NH2	8.07	126.46	119.20
12	AB	128	ALA	CA-C-N	-7.84	111.84	122.98
12	AB	128	ALA	C-N-CA	-7.84	111.84	122.98
55	o	30	ARG	NE-CZ-NH1	-7.65	113.85	121.50
17	C	73	ARG	NE-CZ-NH2	7.53	125.98	119.20
16	AG	33	THR	CA-C-N	7.48	135.83	121.54
16	AG	33	THR	C-N-CA	7.48	135.83	121.54
16	AG	102	PHE	CA-C-N	7.48	135.83	121.54
16	AG	102	PHE	C-N-CA	7.48	135.83	121.54
12	AB	159	PHE	CA-C-N	7.42	135.71	121.54
12	AB	159	PHE	C-N-CA	7.42	135.71	121.54
63	w	112	TYR	CZ-CE2-CD2	-7.39	106.29	119.60
29	O	80	ARG	CG-CD-NE	7.38	128.23	112.00
16	AG	108	GLN	N-CA-C	-7.32	103.81	112.89
23	I	134	MET	CB-CG-SD	-7.20	91.10	112.70
16	AG	426	GLY	O-C-N	-7.18	113.37	122.70
21	G	158	PRO	CA-N-CD	-7.09	102.07	112.00
62	v	96	ILE	CA-C-N	7.06	132.33	121.03
62	v	96	ILE	C-N-CA	7.06	132.33	121.03
60	t	18	ARG	NE-CZ-NH1	-7.03	114.47	121.50
21	G	19	GLN	CA-CB-CG	7.00	128.10	114.10
29	O	80	ARG	CB-CG-CD	-6.94	95.33	111.30
41	a	2334	U	OP2-P-O3'	-6.94	87.19	108.00
12	AB	159	PHE	CB-CA-C	6.86	124.07	110.42
16	AG	102	PHE	N-CA-C	6.77	125.22	110.80
18	D	37	U	N3-C4-O4	-6.63	99.49	119.40
16	AG	104	ARG	O-C-N	6.63	131.41	122.59
63	w	112	TYR	OH-CZ-CE2	-6.57	100.19	119.90
18	D	718	A	P-O5'-C5'	-6.54	111.08	120.90
31	Q	110	ILE	N-CA-C	-6.51	98.99	108.11
16	AG	106	THR	OG1-CB-CG2	-6.49	96.33	109.30
13	AD	289	LEU	CA-C-N	-6.46	108.99	121.58
13	AD	289	LEU	C-N-CA	-6.46	108.99	121.58
16	AG	355	ALA	CB-CA-C	6.43	123.21	110.42
18	D	827	U	N3-C4-O4	-6.41	100.16	119.40
41	a	1019	U	N3-C4-O4	-6.38	100.26	119.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	D	1358	U	N3-C4-O4	-6.36	100.31	119.40
16	AG	103	ASP	CA-C-N	6.34	133.65	121.54
16	AG	103	ASP	C-N-CA	6.34	133.65	121.54
37	W	40	ILE	CA-CB-CG1	-6.32	99.66	110.40
16	AG	109	THR	N-CA-C	-6.32	104.47	111.36
12	AB	99	TYR	CA-C-N	6.31	137.19	121.80
12	AB	99	TYR	C-N-CA	6.31	137.19	121.80
59	s	92	MET	CB-CG-SD	-6.30	93.81	112.70
16	AG	403	VAL	N-CA-C	-6.23	104.20	111.00
41	a	1019	U	C5-C4-O4	6.14	144.31	125.90
12	AB	106	ASP	CA-C-N	6.09	127.45	119.84
12	AB	106	ASP	C-N-CA	6.09	127.45	119.84
16	AG	108	GLN	CB-CA-C	-6.08	97.28	109.99
16	AG	404	GLU	N-CA-C	-6.08	104.97	112.38
16	AG	71	PRO	N-CA-C	6.07	122.61	113.81
29	O	103	PHE	CD1-CE1-CZ	-6.04	109.13	120.00
16	AG	33	THR	CA-CB-CG2	6.00	120.71	110.50
41	a	1082	U	N3-C4-O4	-5.93	101.60	119.40
41	a	2613	U	N3-C4-O4	-5.93	101.62	119.40
41	a	1141	U	C5-C4-O4	5.89	143.59	125.90
18	D	37	U	C5-C4-O4	5.89	143.57	125.90
21	G	157	LEU	C-N-CD	-5.87	100.92	125.00
18	D	1358	U	C5-C4-O4	5.87	143.50	125.90
54	n	176	PRO	N-CA-C	5.85	121.22	114.20
23	I	168	TYR	N-CA-CB	5.84	120.23	110.59
38	X	81	MET	CG-SD-CE	-5.82	88.09	100.90
16	AG	66	ASP	N-CA-C	-5.80	103.93	111.71
27	M	55	GLY	CA-C-N	5.79	132.59	121.54
27	M	55	GLY	C-N-CA	5.79	132.59	121.54
18	D	884	U	N3-C4-O4	-5.76	102.12	119.40
14	AE	853	THR	CA-C-N	5.75	131.02	122.74
14	AE	853	THR	C-N-CA	5.75	131.02	122.74
18	D	827	U	C5-C4-O4	5.72	143.05	125.90
21	G	154	MET	CA-CB-CG	-5.71	102.67	114.10
16	AG	220	VAL	N-CA-C	-5.68	108.31	113.71
23	I	134	MET	CG-SD-CE	-5.68	88.41	100.90
12	AB	107	PRO	O-C-N	5.65	130.26	122.64
25	K	45	ARG	CB-CG-CD	-5.64	98.32	111.30
21	G	154	MET	CG-SD-CE	-5.64	88.49	100.90
9	9	130	PRO	CA-N-CD	-5.63	104.12	112.00
25	K	45	ARG	CA-C-N	5.59	130.28	122.23
25	K	45	ARG	C-N-CA	5.59	130.28	122.23

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
12	AB	151	LYS	CA-C-N	5.58	132.21	121.54
12	AB	151	LYS	C-N-CA	5.58	132.21	121.54
16	AG	101	THR	CA-CB-OG1	5.58	117.97	109.60
60	t	18	ARG	CG-CD-NE	-5.56	99.76	112.00
41	a	1141	U	N3-C4-O4	-5.52	102.84	119.40
50	j	11	MET	CB-CG-SD	-5.50	96.19	112.70
14	AE	1184	ASP	N-CA-C	5.45	121.85	109.81
16	AG	74	GLU	CB-CA-C	-5.45	101.75	110.14
62	v	10	ARG	NE-CZ-NH1	-5.44	116.06	121.50
16	AG	354	ALA	CA-C-N	-5.43	111.17	121.54
16	AG	354	ALA	C-N-CA	-5.43	111.17	121.54
9	9	129	LEU	C-N-CD	-5.42	102.80	125.00
16	AG	355	ALA	N-CA-CB	5.41	119.62	110.49
63	w	38	LEU	N-CA-C	-5.35	105.18	112.35
31	Q	106	ARG	NE-CZ-NH1	-5.32	116.18	121.50
8	7	10	U	C2'-C3'-O3'	5.27	117.41	109.50
16	AG	105	ILE	N-CA-CB	-5.24	102.58	111.23
39	Y	5	GLN	CB-CA-C	-5.24	99.99	110.42
25	K	45	ARG	O-C-N	-5.23	116.83	123.01
16	AG	114	ILE	N-CA-C	-5.23	105.40	110.42
56	p	46	ALA	CA-C-N	5.22	131.52	121.54
56	p	46	ALA	C-N-CA	5.22	131.52	121.54
41	a	2613	U	C5-C4-O4	5.21	141.52	125.90
16	AG	107	THR	CA-C-N	-5.18	112.08	121.14
16	AG	107	THR	C-N-CA	-5.18	112.08	121.14
39	Y	5	GLN	CA-CB-CG	5.17	124.45	114.10
45	e	25	GLN	CG-CD-NE2	-5.15	108.68	116.40
23	I	170	GLU	CB-CG-CD	5.12	121.30	112.60
16	AG	105	ILE	O-C-N	5.11	128.96	122.57
21	G	158	PRO	N-CA-CB	-5.11	98.02	103.38
41	a	2334	U	C2'-C3'-O3'	-5.10	101.85	109.50
41	a	2602	A	C4'-C3'-O3'	5.10	117.05	109.40
14	AE	709	ARG	CA-C-N	5.09	135.45	126.45
14	AE	709	ARG	C-N-CA	5.09	135.45	126.45
18	D	884	U	C5-C4-O4	5.09	141.16	125.90
12	AB	138	ARG	CA-C-N	5.08	131.25	121.54
12	AB	138	ARG	C-N-CA	5.08	131.25	121.54
13	AD	289	LEU	CA-C-O	-5.07	113.27	120.51
16	AG	107	THR	CA-C-O	-5.06	113.16	119.18
16	AG	65	VAL	CA-C-N	5.05	131.60	121.81
16	AG	65	VAL	C-N-CA	5.05	131.60	121.81
16	AG	356	ILE	N-CA-C	-5.03	107.83	111.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
48	h	181	MET	CG-SD-CE	-5.01	89.89	100.90

There are no chirality outliers.

All (35) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
9	9	107	GLU	Peptide
9	9	79	PRO	Peptide
9	9	92	ALA	Peptide
12	AB	140	MET	Peptide
12	AB	151	LYS	Peptide
13	AC	192	VAL	Peptide
13	AD	20	SER	Peptide
13	AD	284	ARG	Mainchain
14	AE	1326	GLN	Peptide
14	AE	1344	LEU	Peptide
14	AE	313	GLY	Peptide
14	AE	416	ILE	Peptide
16	AG	102	PHE	Peptide
16	AG	104	ARG	Peptide,Mainchain
16	AG	11	ALA	Peptide
16	AG	292	ASP	Mainchain
16	AG	426	GLY	Mainchain
21	G	19	GLN	Sidechain
22	H	124	LEU	Peptide
22	H	171	ARG	Peptide
22	H	274	TYR	Peptide
22	H	81	GLU	Peptide
22	H	82	THR	Peptide
23	I	167	TRP	Mainchain
25	K	45	ARG	Mainchain
25	K	77	ASN	Peptide
29	O	12	ARG	Peptide
38	X	65	VAL	Peptide
39	Y	5	GLN	Peptide
54	n	176	PRO	Peptide
55	o	31	HIS	Peptide
59	s	74	TYR	Sidechain
61	u	35	HIS	Peptide
61	u	62	PRO	Peptide

5.2 Too-close contacts ⓘ

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	0	816	839	839	54	0
2	1	857	922	922	68	0
3	2	746	811	811	59	0
4	3	788	844	844	43	0
5	4	753	780	780	88	0
6	5	726	0	392	112	0
7	6	703	0	396	34	0
8	7	527	0	262	83	0
9	9	1117	0	1155	258	0
10	A	1620	826	827	108	0
10	B	1620	813	827	112	0
11	AA	10381	0	10388	386	0
12	AB	1286	0	1300	437	0
13	AC	1698	0	1718	13	0
13	AD	2078	0	1891	37	0
14	AE	10404	0	10625	278	0
15	AF	650	0	658	10	0
16	AG	3852	0	3828	900	0
17	C	544	559	560	106	0
18	D	32703	16423	16453	734	0
19	E	669	719	719	78	0
20	F	589	629	629	58	0
21	G	1760	1785	1787	155	0
22	H	1730	1454	1455	168	0
23	I	1636	1710	1710	130	0
24	J	1643	1707	1707	85	0
25	K	1152	1196	1196	122	0
26	L	848	846	846	105	0
27	M	1181	1235	1238	124	0
28	N	979	1031	1031	94	0
29	O	1022	1070	1070	88	0
30	P	790	831	831	103	0
31	Q	877	887	887	96	0
32	R	939	1001	1001	92	0
33	S	805	844	844	73	0
34	T	714	734	734	91	0
35	U	649	666	666	58	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
36	V	648	691	691	68	0
37	W	663	688	688	48	0
38	X	900	964	965	113	0
39	Y	1032	0	1088	211	0
40	Z	227	0	237	31	0
41	a	61841	31077	31120	1317	0
42	b	582	599	599	46	0
43	c	625	652	652	35	0
44	d	2569	1301	1301	40	0
45	e	501	531	531	46	0
46	f	448	488	488	37	0
47	g	522	520	520	64	0
48	h	2082	2154	2154	138	0
49	i	444	459	458	49	0
50	j	1565	1617	1616	121	0
51	k	426	464	464	46	0
52	l	1552	1619	1619	125	0
53	m	377	418	418	30	0
54	n	1410	1443	1444	157	0
55	o	504	572	572	53	0
56	p	1313	1358	1358	74	0
57	q	302	343	343	25	0
58	r	1111	1148	1148	49	0
59	s	1129	1162	1162	127	0
60	t	946	1023	1023	104	0
61	u	1053	1129	1129	77	0
62	v	1074	1157	1157	122	0
63	w	951	994	994	103	0
64	x	892	923	923	44	0
65	y	917	962	962	56	0
66	z	947	1020	1019	84	0
67	AE	1	0	0	0	0
68	AE	2	0	0	3	0
All	All	181408	98638	132670	7742	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 25.

All (7742) 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)
25:K:45:ARG:CD	25:K:45:ARG:CG	1.75	1.65
6:5:11:DA:C5'	12:AB:67:THR:HG22	1.20	1.60
6:5:10:DT:C7	11:AA:62:TYR:CD2	1.84	1.59
16:AG:425:LEU:CD2	16:AG:429:LYS:HE2	1.21	1.59
16:AG:287:ALA:CA	16:AG:331:LEU:HD12	1.19	1.59
66:z:74:ILE:CG1	66:z:74:ILE:CD1	1.81	1.57
11:AA:846:GLY:C	16:AG:104:ARG:HH12	1.02	1.57
16:AG:393:LEU:CB	16:AG:398:LEU:CD2	1.83	1.56
11:AA:855:PRO:HB2	16:AG:35:THR:CG2	1.37	1.54
6:5:9:DG:C5'	11:AA:473:ARG:HB3	1.34	1.54
12:AB:141:LEU:CB	12:AB:143:LEU:HD12	1.32	1.54
6:5:9:DG:H5''	11:AA:473:ARG:CB	1.29	1.53
16:AG:168:LEU:CD2	16:AG:231:GLY:HA3	1.34	1.53
12:AB:118:ILE:CG2	12:AB:141:LEU:HD11	1.09	1.52
6:5:10:DT:C5	11:AA:62:TYR:CD2	1.95	1.52
7:6:23:DC:C5'	14:AE:259:ARG:HH12	1.21	1.52
14:AE:79:LYS:CA	16:AG:144:LYS:HZ2	1.23	1.51
16:AG:287:ALA:CB	16:AG:331:LEU:HD12	1.41	1.49
6:5:10:DT:C6	11:AA:62:TYR:CE2	1.99	1.48
16:AG:425:LEU:CA	16:AG:429:LYS:HE3	1.40	1.48
16:AG:287:ALA:HA	16:AG:331:LEU:CD1	1.38	1.48
14:AE:72:CYS:SG	68:AE:1502:ZN:ZN	1.03	1.48
11:AA:855:PRO:HB2	16:AG:35:THR:CB	1.38	1.47
12:AB:148:LYS:CE	30:P:100:ILE:HG23	1.40	1.47
11:AA:887:VAL:CB	16:AG:105:ILE:HD12	1.42	1.47
14:AE:79:LYS:HA	16:AG:144:LYS:CE	1.42	1.47
16:AG:363:LEU:CD2	16:AG:410:ALA:H	1.24	1.47
12:AB:141:LEU:HB3	12:AB:143:LEU:CD1	1.42	1.46
11:AA:861:ALA:H	16:AG:38:LYS:NZ	1.00	1.45
16:AG:425:LEU:HD23	16:AG:429:LYS:CE	1.45	1.44
12:AB:115:LYS:NZ	12:AB:129:ILE:CG2	1.82	1.43
16:AG:168:LEU:HD22	16:AG:231:GLY:CA	1.49	1.42
6:5:11:DA:P	12:AB:68:THR:HA	1.58	1.41
12:AB:81:PHE:CD2	14:AE:293:ARG:NH1	1.77	1.41
16:AG:187:ARG:HB3	16:AG:188:PRO:CD	1.45	1.40
16:AG:393:LEU:CD2	16:AG:398:LEU:CD2	1.99	1.39
11:AA:887:VAL:HB	16:AG:105:ILE:CD1	1.52	1.39
16:AG:363:LEU:CD2	16:AG:410:ALA:N	1.82	1.38
14:AE:79:LYS:CA	16:AG:144:LYS:NZ	1.81	1.37
12:AB:115:LYS:NZ	12:AB:129:ILE:HG21	1.05	1.37
6:5:10:DT:C5	11:AA:62:TYR:CE2	2.04	1.37
16:AG:437:LEU:HD21	16:AG:456:LEU:CD1	1.54	1.37

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AA:380:ALA:N	12:AB:65:HIS:CE1	1.84	1.36
16:AG:363:LEU:HD23	16:AG:410:ALA:N	1.38	1.36
6:5:9:DG:OP2	11:AA:473:ARG:CB	1.73	1.36
16:AG:168:LEU:CD2	16:AG:231:GLY:CA	1.98	1.36
11:AA:861:ALA:N	16:AG:38:LYS:HZ1	1.22	1.35
12:AB:148:LYS:NZ	30:P:100:ILE:HG23	1.36	1.35
12:AB:118:ILE:CG2	12:AB:141:LEU:CD1	2.02	1.35
11:AA:846:GLY:C	16:AG:104:ARG:NH1	1.82	1.35
12:AB:148:LYS:NZ	30:P:100:ILE:CG2	1.90	1.35
16:AG:385:ALA:CB	16:AG:411:LYS:HE3	1.56	1.34
16:AG:387:VAL:HG12	16:AG:388:PRO:CD	1.57	1.34
14:AE:79:LYS:HA	16:AG:144:LYS:NZ	1.03	1.33
11:AA:856:ASN:N	16:AG:35:THR:HB	1.42	1.33
11:AA:858:GLY:HA3	16:AG:38:LYS:CB	1.40	1.33
16:AG:393:LEU:CA	16:AG:398:LEU:CD2	2.05	1.33
12:AB:138:ARG:HG3	12:AB:154:VAL:C	1.51	1.32
16:AG:434:LEU:HD22	16:AG:459:LEU:CD1	1.58	1.32
16:AG:393:LEU:CA	16:AG:398:LEU:HD22	1.59	1.32
12:AB:152:HIS:CE1	12:AB:159:PHE:HD2	1.47	1.31
16:AG:434:LEU:CD2	16:AG:456:LEU:HA	1.58	1.31
16:AG:393:LEU:HB3	16:AG:398:LEU:CD2	1.51	1.31
12:AB:138:ARG:HG3	12:AB:154:VAL:O	1.20	1.30
11:AA:858:GLY:CA	16:AG:38:LYS:HB2	1.62	1.30
12:AB:81:PHE:CE1	14:AE:293:ARG:HB3	1.32	1.30
12:AB:51:PHE:HE2	14:AE:288:PRO:CG	1.44	1.30
16:AG:448:LEU:HD21	16:AG:473:LEU:CD1	1.61	1.29
14:AE:287:ALA:CB	14:AE:292:VAL:HG13	1.61	1.29
11:AA:377:THR:OG1	12:AB:62:GLU:CB	1.80	1.29
12:AB:81:PHE:CE1	14:AE:293:ARG:CB	1.92	1.29
16:AG:123:ARG:HH12	16:AG:191:ARG:CA	1.36	1.29
12:AB:115:LYS:HZ2	12:AB:129:ILE:CG2	1.41	1.29
6:5:11:DA:OP2	12:AB:68:THR:CA	1.81	1.28
12:AB:82:GLY:CA	14:AE:142:GLU:HB2	1.62	1.28
12:AB:115:LYS:CE	12:AB:129:ILE:HG21	1.61	1.28
6:5:11:DA:C3'	12:AB:67:THR:HG21	1.63	1.28
12:AB:148:LYS:CE	30:P:100:ILE:CG2	2.09	1.28
16:AG:393:LEU:CG	16:AG:398:LEU:HD21	1.63	1.28
11:AA:856:ASN:ND2	16:AG:35:THR:HG22	1.47	1.27
14:AE:79:LYS:CB	16:AG:144:LYS:HZ2	1.44	1.27
6:5:10:DT:C1'	12:AB:71:ALA:H	1.46	1.27
12:AB:148:LYS:HE2	30:P:100:ILE:CG2	1.64	1.27

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:425:LEU:HA	16:AG:429:LYS:CE	1.64	1.27
6:5:11:DA:C5'	12:AB:67:THR:CG2	2.12	1.27
11:AA:387:ASN:CG	11:AA:394:ARG:NH1	1.92	1.26
12:AB:152:HIS:CD2	12:AB:159:PHE:CE2	2.24	1.26
16:AG:353:HIS:O	16:AG:356:ILE:HG23	1.13	1.26
11:AA:846:GLY:HA3	16:AG:104:ARG:NH2	1.48	1.26
11:AA:887:VAL:CB	16:AG:105:ILE:CD1	2.11	1.26
12:AB:81:PHE:HA	14:AE:141:PHE:O	1.27	1.26
11:AA:481:LEU:CD2	12:AB:16:ARG:CD	2.01	1.25
11:AA:377:THR:OG1	12:AB:62:GLU:HB3	1.20	1.25
11:AA:856:ASN:OD1	16:AG:35:THR:HA	1.08	1.25
16:AG:387:VAL:CG1	16:AG:388:PRO:HD2	1.67	1.25
12:AB:145:LEU:HA	30:P:88:MET:CG	1.66	1.25
16:AG:393:LEU:CD2	16:AG:398:LEU:HD21	1.57	1.25
12:AB:115:LYS:CD	12:AB:129:ILE:HG13	1.67	1.24
16:AG:353:HIS:C	16:AG:356:ILE:HG23	1.62	1.24
16:AG:434:LEU:HA	16:AG:456:LEU:CD2	1.64	1.24
14:AE:288:PRO:HD2	14:AE:291:ILE:CG2	1.66	1.24
12:AB:118:ILE:HG23	12:AB:141:LEU:CD1	1.59	1.24
12:AB:145:LEU:CA	30:P:88:MET:HG2	1.65	1.24
6:5:11:DA:OP2	12:AB:68:THR:HA	1.36	1.23
12:AB:51:PHE:CE2	14:AE:288:PRO:CG	2.22	1.23
8:7:11:U:H4'	25:K:20:ARG:NH2	1.50	1.23
7:6:25:DT:OP1	11:AA:496:LYS:CA	1.80	1.23
16:AG:355:ALA:CB	16:AG:382:GLU:CD	2.10	1.23
16:AG:363:LEU:HD23	16:AG:409:ARG:C	1.63	1.23
11:AA:855:PRO:CB	16:AG:35:THR:CB	2.16	1.23
16:AG:287:ALA:CB	16:AG:331:LEU:CD1	2.16	1.22
16:AG:425:LEU:CG	16:AG:429:LYS:HE2	1.69	1.22
6:5:10:DT:H71	11:AA:62:TYR:CG	1.75	1.22
16:AG:123:ARG:NH1	16:AG:191:ARG:C	1.98	1.22
16:AG:362:TYR:O	16:AG:413:ALA:HB2	1.35	1.21
6:5:10:DT:H1'	12:AB:71:ALA:N	1.21	1.21
11:AA:853:ASP:O	16:AG:103:ASP:OD1	1.58	1.21
12:AB:138:ARG:CG	12:AB:154:VAL:O	1.87	1.20
6:5:11:DA:OP2	12:AB:68:THR:N	1.73	1.20
16:AG:287:ALA:CA	16:AG:331:LEU:CD1	2.02	1.20
16:AG:393:LEU:HD22	16:AG:398:LEU:CD2	1.71	1.20
16:AG:433:ASP:O	16:AG:456:LEU:HD11	1.35	1.20
12:AB:144:ASN:O	30:P:88:MET:CE	1.88	1.20
16:AG:381:LEU:HA	16:AG:384:LEU:HD12	1.21	1.19

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AA:855:PRO:C	16:AG:35:THR:HB	1.68	1.19
12:AB:115:LYS:HD3	12:AB:129:ILE:CD1	1.72	1.19
16:AG:355:ALA:HA	16:AG:382:GLU:OE2	1.37	1.19
16:AG:442:ARG:O	16:AG:446:PHE:CD2	1.95	1.18
16:AG:437:LEU:CD2	16:AG:456:LEU:HD13	1.74	1.18
11:AA:855:PRO:CB	16:AG:35:THR:OG1	1.90	1.18
11:AA:887:VAL:CA	16:AG:105:ILE:HD12	1.73	1.18
56:p:159:GLY:O	56:p:163:ARG:NH1	1.77	1.18
12:AB:152:HIS:NE2	12:AB:159:PHE:CE2	2.12	1.18
16:AG:393:LEU:HA	16:AG:398:LEU:CD2	1.69	1.18
58:r:135:HIS:ND1	58:r:137:GLU:OE1	1.77	1.18
11:AA:887:VAL:C	16:AG:105:ILE:CD1	2.16	1.17
7:6:23:DC:C5'	14:AE:259:ARG:NH1	2.05	1.17
16:AG:393:LEU:HD22	16:AG:398:LEU:CD1	1.73	1.17
6:5:9:DG:C5'	11:AA:473:ARG:CG	2.21	1.17
11:AA:856:ASN:N	16:AG:35:THR:CB	2.02	1.17
12:AB:138:ARG:HA	12:AB:154:VAL:O	1.43	1.16
12:AB:152:HIS:CE1	12:AB:159:PHE:CD2	2.32	1.16
16:AG:434:LEU:HA	16:AG:456:LEU:HD21	1.25	1.16
16:AG:355:ALA:HB2	16:AG:382:GLU:OE1	1.40	1.16
12:AB:144:ASN:O	30:P:88:MET:HE2	0.99	1.16
16:AG:353:HIS:O	16:AG:357:ASP:N	1.77	1.16
26:L:9:MET:HE1	26:L:85:ILE:HB	1.17	1.15
12:AB:81:PHE:CE2	14:AE:293:ARG:NH1	1.88	1.15
38:X:81:MET:HE1	38:X:92:ARG:HB3	1.20	1.15
12:AB:152:HIS:CD2	12:AB:159:PHE:CD2	2.34	1.15
16:AG:355:ALA:HA	16:AG:382:GLU:CD	1.71	1.15
7:6:25:DT:C3'	11:AA:496:LYS:HG3	1.74	1.14
6:5:9:DG:H5'	11:AA:473:ARG:CG	1.76	1.14
16:AG:123:ARG:NH1	16:AG:191:ARG:O	1.80	1.14
16:AG:355:ALA:HB1	16:AG:382:GLU:CG	1.77	1.14
16:AG:424:SER:O	16:AG:429:LYS:HG2	1.47	1.14
41:a:2335:A:OP2	64:x:13:ARG:NE	1.81	1.14
37:W:40:ILE:HD13	37:W:66:MET:SD	1.89	1.13
12:AB:102:LYS:CE	14:AE:285:LEU:HG	1.78	1.13
12:AB:148:LYS:HE2	30:P:100:ILE:HG23	1.16	1.13
6:5:9:DG:OP2	11:AA:473:ARG:HB2	0.97	1.13
16:AG:123:ARG:HH12	16:AG:191:ARG:C	1.57	1.13
6:5:10:DT:H71	11:AA:62:TYR:CD2	1.77	1.12
11:AA:861:ALA:N	16:AG:38:LYS:NZ	1.84	1.12
14:AE:287:ALA:HB2	14:AE:292:VAL:CG1	1.78	1.12

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:5:9:DG:C5'	11:AA:473:ARG:CB	2.02	1.12
6:5:10:DT:O3'	12:AB:67:THR:O	1.67	1.12
6:5:11:DA:H5'	12:AB:67:THR:HG22	1.14	1.12
16:AG:285:ILE:CG1	16:AG:293:VAL:HG11	1.79	1.12
12:AB:82:GLY:HA2	14:AE:142:GLU:HB2	1.13	1.12
12:AB:115:LYS:HA	12:AB:129:ILE:HD12	1.28	1.12
16:AG:279:ASN:OD1	16:AG:280:PRO:HD3	1.50	1.12
16:AG:393:LEU:CD2	16:AG:398:LEU:CD1	2.27	1.12
16:AG:425:LEU:CD2	16:AG:429:LYS:CE	2.13	1.12
11:AA:887:VAL:HB	16:AG:105:ILE:CG1	1.80	1.11
16:AG:393:LEU:HD22	16:AG:398:LEU:HD11	1.28	1.11
6:5:10:DT:C7	11:AA:62:TYR:CG	2.31	1.11
10:A:76:A:O2'	41:a:2394:C:N3	1.81	1.11
12:AB:54:TYR:CE2	14:AE:291:ILE:HG23	1.84	1.11
16:AG:353:HIS:O	16:AG:356:ILE:CG2	1.98	1.11
12:AB:115:LYS:HD3	12:AB:129:ILE:HD12	1.31	1.11
16:AG:422:GLU:HA	16:AG:426:GLY:CA	1.79	1.11
6:5:10:DT:H73	11:AA:62:TYR:CD2	1.77	1.11
12:AB:35:LEU:HA	12:AB:106:ASP:OD1	1.47	1.11
16:AG:187:ARG:HB3	16:AG:188:PRO:HD3	1.32	1.10
16:AG:187:ARG:HB3	16:AG:188:PRO:HD2	1.21	1.10
16:AG:425:LEU:HA	16:AG:429:LYS:CG	1.80	1.10
11:AA:846:GLY:CA	16:AG:104:ARG:HH22	1.63	1.10
12:AB:152:HIS:NE2	12:AB:159:PHE:CD2	2.19	1.10
11:AA:380:ALA:N	12:AB:65:HIS:HE1	1.29	1.10
11:AA:887:VAL:C	16:AG:105:ILE:HD12	1.72	1.10
16:AG:422:GLU:HA	16:AG:426:GLY:HA3	1.33	1.10
11:AA:855:PRO:HB2	16:AG:35:THR:HG21	1.15	1.09
10:B:75:C:OP2	41:a:2602:A:O2'	1.67	1.09
46:f:4:THR:OG1	46:f:37:GLU:OE2	1.68	1.09
9:9:46:ARG:HG3	9:9:94:ARG:HH21	1.12	1.09
11:AA:855:PRO:CB	16:AG:35:THR:CG2	2.30	1.09
12:AB:118:ILE:HG21	12:AB:141:LEU:HD11	1.18	1.09
16:AG:285:ILE:HG23	16:AG:293:VAL:HG12	1.25	1.09
36:V:7:THR:OG1	36:V:60:GLU:OE2	1.66	1.09
39:Y:5:GLN:HG2	39:Y:61:TYR:CE1	1.87	1.09
16:AG:181:GLY:HA2	16:AG:204:MET:SD	1.93	1.09
58:r:135:HIS:HD1	58:r:137:GLU:CD	1.59	1.09
16:AG:434:LEU:CD2	16:AG:459:LEU:HD12	1.81	1.09
11:AA:855:PRO:CB	16:AG:35:THR:HG21	1.82	1.09
6:5:9:DG:H5'	11:AA:473:ARG:HG3	1.08	1.08

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:6:23:DC:H5''	14:AE:259:ARG:HH12	1.14	1.08
12:AB:51:PHE:CZ	14:AE:288:PRO:HD3	1.88	1.08
8:7:11:U:C4'	25:K:20:ARG:NH2	2.16	1.08
11:AA:387:ASN:CG	11:AA:394:ARG:HH11	1.55	1.08
16:AG:393:LEU:HB3	16:AG:398:LEU:HD23	1.26	1.08
16:AG:430:PRO:HG2	16:AG:435:LEU:HD21	1.29	1.08
12:AB:28:CYS:HA	12:AB:56:PHE:O	1.52	1.08
16:AG:442:ARG:O	16:AG:446:PHE:HD2	1.34	1.08
9:9:50:VAL:HG22	39:Y:119:ALA:HB3	1.33	1.07
16:AG:451:ARG:NH1	16:AG:467:LEU:HA	1.68	1.07
39:Y:96:LYS:HE3	39:Y:136:GLY:HA3	1.29	1.07
7:6:23:DC:O5'	14:AE:259:ARG:NH1	1.88	1.07
12:AB:102:LYS:HE2	14:AE:285:LEU:HG	1.07	1.07
16:AG:434:LEU:CD2	16:AG:456:LEU:CA	2.33	1.07
6:5:9:DG:C5'	11:AA:473:ARG:HG3	1.82	1.07
11:AA:469:VAL:O	11:AA:473:ARG:HG2	1.53	1.07
12:AB:115:LYS:HD2	12:AB:129:ILE:HG13	1.13	1.07
14:AE:287:ALA:HB2	14:AE:292:VAL:HG13	1.08	1.07
16:AG:365:ILE:HD13	16:AG:406:LEU:CD2	1.85	1.07
16:AG:385:ALA:CB	16:AG:411:LYS:CE	2.33	1.07
16:AG:434:LEU:HD21	16:AG:456:LEU:HA	1.26	1.07
6:5:11:DA:H5'	12:AB:67:THR:CG2	1.82	1.07
11:AA:914:LYS:N	16:AG:105:ILE:HA	1.68	1.06
14:AE:288:PRO:HD2	14:AE:291:ILE:HG21	1.10	1.06
53:m:22:MET:O	53:m:28:ARG:NH1	1.87	1.06
11:AA:847:PRO:HD2	16:AG:104:ARG:NH2	1.70	1.06
12:AB:82:GLY:HA2	14:AE:142:GLU:CB	1.85	1.06
16:AG:393:LEU:CB	16:AG:398:LEU:HD21	1.58	1.06
11:AA:887:VAL:CG2	16:AG:105:ILE:HG13	1.83	1.06
16:AG:434:LEU:HD23	16:AG:456:LEU:HD23	1.10	1.06
16:AG:451:ARG:HH12	16:AG:467:LEU:C	1.62	1.06
27:M:4:ARG:HG3	27:M:5:ARG:HG3	1.33	1.06
12:AB:115:LYS:CE	12:AB:129:ILE:CG2	2.25	1.06
11:AA:856:ASN:CG	16:AG:35:THR:HA	1.80	1.05
11:AA:889:PRO:HG3	16:AG:104:ARG:HD3	1.39	1.05
16:AG:385:ALA:HB1	16:AG:411:LYS:HE3	1.35	1.05
12:AB:115:LYS:CD	12:AB:129:ILE:CG1	2.33	1.05
18:D:1309:G:OP2	38:X:98:ARG:NE	1.88	1.05
16:AG:387:VAL:HG12	16:AG:388:PRO:HD2	1.06	1.05
16:AG:393:LEU:CB	16:AG:398:LEU:HD23	1.80	1.05
16:AG:361:LYS:HE2	16:AG:417:ILE:HG12	1.39	1.05

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:F:37:PHE:HB2	31:Q:126:LYS:HE2	1.35	1.05
8:7:11:U:O2'	25:K:33:PHE:CE1	2.10	1.05
16:AG:285:ILE:HG13	16:AG:293:VAL:CG1	1.86	1.05
26:L:9:MET:HE1	26:L:85:ILE:CB	1.86	1.05
16:AG:363:LEU:HD21	16:AG:410:ALA:H	0.94	1.04
9:9:103:ASN:ND2	9:9:107:GLU:OE2	1.91	1.04
61:u:122:VAL:HG12	61:u:125:LEU:HD21	1.35	1.04
56:p:149:ARG:NH2	56:p:152:ARG:O	1.91	1.04
16:AG:434:LEU:HD23	16:AG:456:LEU:HA	1.34	1.04
20:F:25:LYS:NZ	22:H:336:ASP:OD1	1.90	1.04
47:g:58:ASP:OD1	47:g:62:LYS:NZ	1.90	1.04
6:5:11:DA:H3'	12:AB:67:THR:HG21	1.05	1.03
11:AA:856:ASN:OD1	16:AG:35:THR:CA	2.04	1.03
11:AA:887:VAL:HG21	16:AG:105:ILE:HG13	1.39	1.03
16:AG:168:LEU:HD21	16:AG:231:GLY:CA	1.82	1.03
50:j:179:ARG:NH1	50:j:180:VAL:O	1.92	1.03
9:9:31:ARG:NH1	41:a:1054:A:O4'	1.91	1.03
16:AG:285:ILE:HG23	16:AG:293:VAL:CG1	1.87	1.03
16:AG:425:LEU:CG	16:AG:429:LYS:CE	2.34	1.03
18:D:564:C:OP2	32:R:12:ARG:NH2	1.91	1.03
9:9:48:ALA:HB3	9:9:51:TYR:HE1	1.20	1.03
12:AB:115:LYS:HD3	12:AB:129:ILE:CG1	1.87	1.03
16:AG:302:LYS:H	16:AG:302:LYS:HE3	1.16	1.03
16:AG:448:LEU:CD2	16:AG:473:LEU:CD1	2.36	1.03
65:y:51:ARG:NH1	65:y:56:HIS:O	1.91	1.03
8:7:4:U:O2'	18:D:1403:C:O2	1.75	1.03
16:AG:425:LEU:HA	16:AG:429:LYS:HG3	1.33	1.03
50:j:184:ARG:NH2	65:y:7:GLN:OE1	1.90	1.03
6:5:11:DA:P	12:AB:68:THR:CA	2.45	1.03
12:AB:124:GLU:O	12:AB:126:PHE:N	1.92	1.03
12:AB:146:ILE:N	30:P:88:MET:HA	1.72	1.03
23:I:152:GLU:OE2	23:I:154:SER:OG	1.77	1.03
23:I:123:GLN:OE1	23:I:126:ARG:NH2	1.92	1.02
12:AB:51:PHE:HE2	14:AE:288:PRO:HG2	1.20	1.02
26:L:42:TRP:CH2	26:L:102:MET:HG3	1.94	1.02
39:Y:93:ASN:HA	39:Y:135:MET:CE	1.89	1.02
7:6:18:DT:H3	8:7:49:A:N6	1.55	1.02
16:AG:425:LEU:C	16:AG:429:LYS:HE3	1.84	1.02
12:AB:141:LEU:HD23	12:AB:143:LEU:HG	1.42	1.02
16:AG:363:LEU:HD23	16:AG:409:ARG:CA	1.89	1.02
59:s:72:LYS:HD3	59:s:74:TYR:OH	1.59	1.02

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:9:136:ILE:HD12	9:9:139:LEU:HD12	1.38	1.02
12:AB:81:PHE:CA	14:AE:141:PHE:O	2.06	1.02
12:AB:102:LYS:HE2	14:AE:285:LEU:CG	1.89	1.02
16:AG:365:ILE:HD13	16:AG:406:LEU:HD22	1.37	1.02
39:Y:56:VAL:HA	39:Y:71:LYS:HZ1	1.21	1.02
16:AG:393:LEU:HD22	16:AG:398:LEU:HD21	1.25	1.01
16:AG:422:GLU:HA	16:AG:426:GLY:N	1.74	1.01
11:AA:481:LEU:HD23	12:AB:16:ARG:CD	1.87	1.01
12:AB:146:ILE:H	30:P:88:MET:HA	0.86	1.01
16:AG:131:ARG:O	16:AG:134:GLU:HG3	1.61	1.01
11:AA:481:LEU:HD21	12:AB:16:ARG:CD	1.60	1.01
16:AG:287:ALA:HA	16:AG:331:LEU:HD11	1.41	1.01
39:Y:14:ALA:HB1	39:Y:50:LYS:HD2	1.37	1.01
8:7:3:G:H5'	18:D:1500:A:N6	1.75	1.01
16:AG:365:ILE:CG2	16:AG:409:ARG:NH1	2.23	1.01
16:AG:393:LEU:CG	16:AG:398:LEU:CD2	2.30	1.01
6:5:10:DT:C7	11:AA:62:TYR:HD2	1.69	1.01
9:9:43:LYS:HG3	9:9:46:ARG:HH12	1.22	1.01
16:AG:363:LEU:CD2	16:AG:409:ARG:HB2	1.91	1.01
39:Y:60:VAL:HG22	39:Y:66:PHE:HB3	1.43	1.01
16:AG:363:LEU:HG	16:AG:410:ALA:HB2	1.37	1.00
52:l:1:MET:HE2	52:l:16:GLU:HA	1.40	1.00
6:5:11:DA:H3'	12:AB:67:THR:CG2	1.91	1.00
9:9:30:SER:O	9:9:31:ARG:NE	1.94	1.00
16:AG:361:LYS:CE	16:AG:417:ILE:HG12	1.90	1.00
14:AE:70:CYS:SG	14:AE:85:CYS:SG	2.60	1.00
14:AE:288:PRO:CD	14:AE:291:ILE:HG21	1.92	1.00
16:AG:168:LEU:HD21	16:AG:231:GLY:N	1.76	1.00
21:G:49:MET:HE1	21:G:201:PRO:HD3	1.42	1.00
18:D:377:G:OP1	35:U:5:ARG:NH1	1.94	1.00
16:AG:187:ARG:CB	16:AG:188:PRO:CD	2.37	1.00
16:AG:434:LEU:HD21	16:AG:456:LEU:CA	1.89	1.00
12:AB:142:LEU:HA	12:AB:150:ILE:O	1.62	1.00
16:AG:336:LEU:HD12	16:AG:336:LEU:H	1.26	1.00
16:AG:425:LEU:CA	16:AG:429:LYS:CE	2.30	0.99
12:AB:82:GLY:CA	14:AE:142:GLU:CB	2.38	0.99
26:L:42:TRP:HH2	26:L:102:MET:HG3	1.25	0.99
7:6:25:DT:H3'	11:AA:496:LYS:HG3	1.41	0.99
11:AA:887:VAL:HG11	16:AG:105:ILE:HG21	1.41	0.99
12:AB:81:PHE:CE1	14:AE:293:ARG:HB2	1.91	0.99
12:AB:140:MET:N	12:AB:152:HIS:O	1.96	0.99

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:393:LEU:HD22	16:AG:398:LEU:CG	1.93	0.99
21:G:154:MET:HE2	21:G:158:PRO:HD3	1.44	0.99
16:AG:434:LEU:HD23	16:AG:456:LEU:CD2	1.91	0.99
12:AB:146:ILE:H	30:P:88:MET:CA	1.76	0.98
16:AG:434:LEU:CD2	16:AG:456:LEU:HD23	1.92	0.98
14:AE:79:LYS:HE2	16:AG:153:ASP:OD1	1.63	0.98
39:Y:93:ASN:CA	39:Y:135:MET:HE1	1.94	0.98
52:l:1:MET:HE1	52:l:20:GLY:HA3	1.43	0.98
16:AG:362:TYR:O	16:AG:413:ALA:CB	2.10	0.98
14:AE:70:CYS:SG	14:AE:72:CYS:SG	2.61	0.98
16:AG:287:ALA:HB2	16:AG:331:LEU:CD1	1.92	0.98
16:AG:393:LEU:CD2	16:AG:398:LEU:HD11	1.90	0.98
16:AG:393:LEU:HD23	16:AG:398:LEU:HD22	1.43	0.98
16:AG:168:LEU:HD22	16:AG:231:GLY:HA2	1.40	0.97
33:S:86:GLU:OE1	33:S:90:ARG:NH1	1.96	0.97
9:9:3:LEU:HD12	9:9:4:ASN:H	1.25	0.97
16:AG:168:LEU:HD21	16:AG:230:PRO:C	1.89	0.97
14:AE:79:LYS:CA	16:AG:144:LYS:CE	2.32	0.97
16:AG:355:ALA:HB1	16:AG:382:GLU:HG3	1.41	0.97
6:5:10:DT:O5'	11:AA:62:TYR:OH	1.80	0.97
6:5:11:DA:H5''	12:AB:67:THR:CG2	1.86	0.97
12:AB:6:LEU:HD13	14:AE:291:ILE:HD11	1.46	0.97
14:AE:79:LYS:HA	16:AG:144:LYS:HE3	1.40	0.97
16:AG:279:ASN:CG	16:AG:280:PRO:CD	2.38	0.97
6:5:11:DA:C3'	12:AB:67:THR:CG2	2.42	0.97
8:7:2:U:P	18:D:1505:G:H8	1.88	0.97
11:AA:387:ASN:ND2	11:AA:394:ARG:HH12	1.61	0.97
39:Y:93:ASN:HA	39:Y:135:MET:HE1	1.00	0.97
41:a:2093:G:OP1	58:r:22:LYS:NZ	1.98	0.97
16:AG:228:ARG:HA	16:AG:327:LEU:HD11	1.46	0.97
18:D:1344:C:OP1	29:O:124:ARG:NH1	1.98	0.97
9:9:88:HIS:HB3	9:9:89:PRO:HD3	1.44	0.97
58:r:38:PRO:O	58:r:43:ASN:ND2	1.96	0.96
59:s:12:LYS:NZ	59:s:14:ASP:OD1	1.96	0.96
9:9:95:LEU:HD13	9:9:98:GLU:OE2	1.65	0.96
16:AG:380:THR:O	16:AG:384:LEU:CD1	2.13	0.96
16:AG:433:ASP:O	16:AG:456:LEU:CD1	2.12	0.96
11:AA:846:GLY:CA	16:AG:104:ARG:HH12	1.78	0.96
12:AB:141:LEU:CB	12:AB:143:LEU:CD1	2.15	0.96
16:AG:448:LEU:HD21	16:AG:473:LEU:HD11	1.48	0.96
12:AB:138:ARG:CA	12:AB:154:VAL:O	2.12	0.96

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AB:51:PHE:CE2	14:AE:288:PRO:CD	2.48	0.96
14:AE:287:ALA:CB	14:AE:292:VAL:CG1	2.38	0.96
16:AG:355:ALA:HB2	16:AG:382:GLU:CD	1.83	0.96
16:AG:393:LEU:CA	16:AG:398:LEU:HD23	1.88	0.96
33:S:92:GLU:OE1	33:S:92:GLU:N	1.97	0.96
18:D:562:U:O2'	32:R:14:ARG:NH1	1.99	0.96
6:5:11:DA:H5''	12:AB:67:THR:HG22	0.97	0.95
16:AG:437:LEU:CD2	16:AG:456:LEU:CD1	2.34	0.95
64:x:111:ARG:NH2	64:x:117:PHE:OXT	1.99	0.95
12:AB:51:PHE:CE2	14:AE:288:PRO:HG3	1.98	0.95
41:a:1042:G:HO2'	41:a:1043:C:H6	1.09	0.95
9:9:31:ARG:HD2	41:a:1053:C:O2'	1.66	0.95
12:AB:118:ILE:HG22	12:AB:141:LEU:HD11	1.49	0.95
16:AG:387:VAL:HG12	16:AG:388:PRO:HD3	1.47	0.95
16:AG:424:SER:O	16:AG:429:LYS:CG	2.13	0.95
39:Y:33:ASN:HB3	39:Y:36:GLU:HG2	1.48	0.95
39:Y:75:ALA:HB2	39:Y:112:LYS:HE3	1.48	0.95
9:9:61:ARG:NH1	41:a:1047:G:N7	2.15	0.95
23:I:164:ARG:NH2	23:I:166:GLU:OE2	1.99	0.95
16:AG:440:VAL:CG2	16:AG:481:LEU:HD22	1.97	0.95
12:AB:103:ASP:HB3	12:AB:105:VAL:HG23	1.48	0.95
16:AG:214:PRO:O	16:AG:218:GLU:OE2	1.83	0.95
11:AA:887:VAL:CG2	16:AG:105:ILE:CD1	2.45	0.95
16:AG:448:LEU:HD21	16:AG:473:LEU:HD13	1.48	0.95
36:V:7:THR:OG1	36:V:61:ILE:O	1.85	0.95
9:9:3:LEU:HD12	9:9:4:ASN:N	1.82	0.94
12:AB:152:HIS:CG	12:AB:159:PHE:CD2	2.55	0.94
40:Z:14:MET:HB3	40:Z:17:MET:HG2	1.46	0.94
12:AB:152:HIS:NE2	12:AB:159:PHE:HE2	1.63	0.94
16:AG:365:ILE:CD1	16:AG:406:LEU:HD22	1.97	0.94
41:a:30:G:OP2	66:z:5:LYS:NZ	2.00	0.94
16:AG:393:LEU:HA	16:AG:398:LEU:HD22	0.96	0.94
39:Y:5:GLN:HG2	39:Y:61:TYR:HE1	1.31	0.94
9:9:50:VAL:HG22	39:Y:119:ALA:CB	1.98	0.94
11:AA:387:ASN:ND2	11:AA:394:ARG:NH1	2.15	0.94
12:AB:82:GLY:N	14:AE:142:GLU:CB	2.30	0.94
16:AG:258:ARG:HH11	16:AG:258:ARG:HG2	1.31	0.94
6:5:9:DG:OP2	11:AA:473:ARG:CG	2.16	0.94
11:AA:858:GLY:CA	16:AG:38:LYS:CB	2.27	0.94
12:AB:152:HIS:ND1	12:AB:159:PHE:HD2	1.64	0.94
56:p:38:ASN:ND2	56:p:64:GLN:OE1	2.00	0.94

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:9:23:LEU:HD13	9:9:92:ALA:CB	1.98	0.94
11:AA:481:LEU:HD21	12:AB:16:ARG:HD3	1.47	0.94
18:D:526:C:OP2	32:R:88:LYS:NZ	2.01	0.94
39:Y:14:ALA:O	39:Y:50:LYS:HE3	1.67	0.94
6:5:11:DA:OP2	12:AB:67:THR:C	2.11	0.94
16:AG:451:ARG:NH1	16:AG:467:LEU:CA	2.30	0.93
11:AA:858:GLY:HA3	16:AG:38:LYS:HB2	0.94	0.93
16:AG:363:LEU:HG	16:AG:410:ALA:CB	1.97	0.93
16:AG:171:GLU:OE2	16:AG:267:GLY:HA3	1.68	0.93
9:9:88:HIS:HB3	9:9:89:PRO:CD	1.97	0.93
8:7:11:U:H4'	25:K:20:ARG:HH22	1.18	0.93
11:AA:846:GLY:HA3	16:AG:104:ARG:HH22	1.12	0.93
12:AB:81:PHE:HE1	14:AE:293:ARG:CB	1.55	0.93
12:AB:154:VAL:HG11	12:AB:159:PHE:HB3	1.47	0.93
12:AB:154:VAL:CG1	12:AB:159:PHE:HB3	1.98	0.93
12:AB:115:LYS:HD2	12:AB:129:ILE:CG1	1.97	0.93
16:AG:363:LEU:CG	16:AG:410:ALA:HB2	1.99	0.93
26:L:9:MET:CE	26:L:85:ILE:HB	1.97	0.93
11:AA:481:LEU:CD2	12:AB:16:ARG:HD3	1.98	0.93
12:AB:110:PRO:HA	12:AB:114:ASP:OD2	1.69	0.93
16:AG:240:THR:OG1	16:AG:247:PRO:HG3	1.68	0.93
7:6:18:DT:H3	8:7:49:A:H61	1.05	0.92
14:AE:59:ALA:CB	14:AE:71:LEU:HD21	1.99	0.92
9:9:58:THR:HG21	9:9:81:LEU:HD23	1.48	0.92
16:AG:123:ARG:CZ	16:AG:191:ARG:O	2.17	0.92
16:AG:285:ILE:HG13	16:AG:293:VAL:HG11	1.45	0.92
7:6:25:DT:OP1	11:AA:496:LYS:HA	1.12	0.92
11:AA:846:GLY:HA3	16:AG:104:ARG:CZ	1.97	0.92
12:AB:141:LEU:HD23	12:AB:143:LEU:CG	1.99	0.92
12:AB:154:VAL:HG11	12:AB:159:PHE:CB	1.98	0.92
7:6:25:DT:H2'	11:AA:496:LYS:HE3	1.51	0.92
16:AG:434:LEU:HD21	16:AG:455:THR:O	1.69	0.92
16:AG:451:ARG:NH1	16:AG:466:ASP:O	2.02	0.92
10:B:76:A:N3	41:a:2493:U:H4'	1.84	0.92
27:M:50:LEU:HD21	27:M:121:ALA:HA	1.52	0.92
31:Q:51:GLY:O	31:Q:56:ARG:NH2	2.03	0.92
16:AG:279:ASN:CG	16:AG:280:PRO:HD3	1.93	0.92
12:AB:115:LYS:HA	12:AB:129:ILE:CD1	1.99	0.92
11:AA:887:VAL:HB	16:AG:105:ILE:HD12	0.97	0.92
1:0:18:GLN:N	1:0:18:GLN:OE1	2.01	0.91
16:AG:123:ARG:HH12	16:AG:191:ARG:HA	1.32	0.91

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:453:VAL:HG13	16:AG:458:ASP:CB	2.00	0.91
14:AE:85:CYS:HG	68:AE:1502:ZN:ZN	0.70	0.91
16:AG:434:LEU:HD22	16:AG:459:LEU:HD12	0.93	0.91
6:5:10:DT:C6	11:AA:62:TYR:HE2	1.85	0.91
41:a:2293:G:OP1	64:x:94:ARG:NH1	2.04	0.91
36:V:27:ARG:NH1	36:V:42:THR:OG1	2.03	0.91
8:7:44:U:H5'	11:AA:1259:LEU:HD21	1.52	0.91
12:AB:118:ILE:HG23	12:AB:141:LEU:HD11	0.92	0.91
16:AG:183:LEU:HG	16:AG:197:VAL:HG13	1.51	0.91
18:D:1526:G:N7	20:F:40:LYS:NZ	2.18	0.91
60:t:90:ASN:OD1	60:t:91:SER:N	2.02	0.91
16:AG:174:ARG:HB3	16:AG:175:PRO:HD2	1.51	0.91
16:AG:363:LEU:HD23	16:AG:409:ARG:CB	2.00	0.91
41:a:1818:U:OP2	48:h:156:ARG:NH1	2.04	0.91
12:AB:115:LYS:CD	12:AB:129:ILE:HG21	2.01	0.91
16:AG:434:LEU:HD23	16:AG:456:LEU:CA	2.00	0.91
17:C:38:LYS:NZ	18:D:718:A:N1	2.19	0.91
6:5:9:DG:P	11:AA:473:ARG:HB2	2.10	0.91
16:AG:385:ALA:HB3	16:AG:411:LYS:CE	2.00	0.91
16:AG:434:LEU:HA	16:AG:456:LEU:HD23	1.53	0.91
6:5:10:DT:C1'	12:AB:71:ALA:N	2.10	0.90
12:AB:3:SER:O	12:AB:58:GLU:HA	1.71	0.90
16:AG:363:LEU:HD22	16:AG:409:ARG:HB2	1.53	0.90
41:a:2682:A:O4'	50:j:11:MET:SD	2.29	0.90
12:AB:140:MET:C	12:AB:152:HIS:O	2.13	0.90
10:B:30:G:HO2'	18:D:1339:A:H2	1.16	0.90
62:v:111:GLU:O	62:v:114:ARG:HG2	1.71	0.90
8:7:2:U:P	18:D:1505:G:C8	2.64	0.90
16:AG:425:LEU:HA	16:AG:429:LYS:CD	2.00	0.90
9:9:92:ALA:O	9:9:129:LEU:HD13	1.71	0.90
14:AE:79:LYS:CA	16:AG:144:LYS:HE3	1.99	0.90
16:AG:453:VAL:HG23	16:AG:462:GLN:NE2	1.87	0.90
39:Y:79:LEU:HD21	39:Y:105:LEU:HD21	1.52	0.90
47:g:27:THR:OG1	54:n:140:GLU:OE1	1.88	0.90
6:5:11:DA:P	12:AB:67:THR:O	2.30	0.90
11:AA:889:PRO:HG3	16:AG:104:ARG:HH11	1.36	0.90
16:AG:393:LEU:HB3	16:AG:398:LEU:HD21	1.25	0.90
39:Y:129:GLU:HB3	39:Y:133:ARG:NH1	1.87	0.90
16:AG:200:SER:O	16:AG:230:PRO:HG2	1.71	0.90
25:K:45:ARG:CD	25:K:45:ARG:CB	2.49	0.90
41:a:2393:U:OP1	55:o:30:ARG:NE	2.05	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:9:23:LEU:HD13	9:9:92:ALA:HB1	1.49	0.90
14:AE:79:LYS:HA	16:AG:144:LYS:HZ1	1.35	0.90
61:u:93:ASN:OD1	61:u:94:THR:N	2.05	0.90
65:y:89:ARG:NE	65:y:113:ARG:O	2.05	0.90
6:5:10:DT:C2	12:AB:71:ALA:HB3	1.84	0.89
11:AA:855:PRO:CG	16:AG:105:ILE:HG22	2.03	0.89
12:AB:51:PHE:HE2	14:AE:288:PRO:CD	1.85	0.89
52:l:170:ARG:NH1	52:l:174:GLY:O	2.05	0.89
16:AG:422:GLU:CA	16:AG:426:GLY:HA3	2.03	0.89
17:C:60:LYS:HZ2	18:D:734:G:C2'	1.86	0.89
41:a:78:U:OP1	45:e:4:LYS:NZ	2.06	0.89
16:AG:240:THR:OG1	16:AG:247:PRO:CG	2.20	0.89
60:t:20:MET:HB3	60:t:44:LYS:HZ3	1.33	0.89
12:AB:82:GLY:N	14:AE:142:GLU:HB3	1.86	0.89
14:AE:79:LYS:CG	16:AG:144:LYS:HZ2	1.86	0.89
10:B:38:A:H4'	18:D:790:A:O4'	1.71	0.89
64:x:36:TYR:OH	64:x:38:GLN:NE2	2.05	0.89
9:9:8:LYS:O	9:9:12:VAL:HG23	1.73	0.89
12:AB:51:PHE:CE2	14:AE:288:PRO:HD3	2.08	0.89
16:AG:365:ILE:HG21	16:AG:409:ARG:NH1	1.87	0.89
31:Q:15:GLN:NE2	31:Q:76:GLU:O	2.06	0.89
45:e:23:ARG:NH1	45:e:24:GLU:OE2	2.06	0.89
11:AA:856:ASN:HD21	16:AG:35:THR:HG22	1.37	0.89
16:AG:422:GLU:HA	16:AG:426:GLY:H	1.33	0.89
17:C:60:LYS:NZ	18:D:735:C:O4'	2.06	0.89
27:M:66:LEU:HD13	27:M:101:MET:CE	2.02	0.89
10:A:76:A:N6	41:a:2422:C:O4'	2.06	0.89
11:AA:887:VAL:CG2	16:AG:105:ILE:CG1	2.51	0.89
14:AE:59:ALA:CB	14:AE:71:LEU:CD2	2.51	0.89
39:Y:129:GLU:HB3	39:Y:133:ARG:HH12	1.38	0.88
12:AB:117:ILE:HG23	12:AB:160:ARG:O	1.73	0.88
27:M:13:LEU:HD12	27:M:14:PRO:HD2	1.54	0.88
16:AG:425:LEU:CA	16:AG:429:LYS:HG3	2.02	0.88
6:5:10:DT:H2''	12:AB:70:ASN:N	1.87	0.88
13:AD:286:GLU:H	13:AD:315:GLY:HA2	1.36	0.88
16:AG:171:GLU:CD	16:AG:267:GLY:HA3	1.99	0.88
18:D:545:C:H5'	24:J:69:GLU:HG3	1.55	0.88
39:Y:45:THR:CG2	39:Y:50:LYS:HD3	2.02	0.88
41:a:585:G:N7	66:z:6:ARG:NH1	2.21	0.88
16:AG:434:LEU:HD21	16:AG:455:THR:C	1.97	0.88
41:a:404:A:O2'	41:a:405:U:OP2	1.90	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:2:54:GLU:OE1	3:2:91:GLN:NE2	2.06	0.88
24:J:67:VAL:HB	24:J:72:PHE:CZ	2.09	0.88
38:X:81:MET:HE2	41:a:888:C:C5	2.09	0.88
11:AA:387:ASN:CG	11:AA:394:ARG:HH12	1.81	0.88
16:AG:363:LEU:HG	16:AG:410:ALA:CA	2.04	0.88
11:AA:380:ALA:H	12:AB:65:HIS:CE1	1.61	0.88
16:AG:123:ARG:NH1	16:AG:191:ARG:CA	2.10	0.88
16:AG:168:LEU:HD22	16:AG:231:GLY:HA3	1.00	0.88
11:AA:847:PRO:O	16:AG:104:ARG:NH1	2.07	0.87
16:AG:362:TYR:C	16:AG:413:ALA:HB2	1.99	0.87
41:a:2756:U:OP2	57:q:19:ARG:NE	2.06	0.87
18:D:280:C:O2	36:V:40:ARG:CZ	2.21	0.87
11:AA:887:VAL:CB	16:AG:105:ILE:CG1	2.49	0.87
18:D:1439:G:OP1	19:E:33:LYS:NZ	2.05	0.87
52:l:170:ARG:NH1	52:l:176:ASP:OD1	2.08	0.87
59:s:13:ARG:NH1	59:s:49:ASP:O	2.07	0.87
61:u:90:VAL:HG13	61:u:95:LEU:HD21	1.56	0.87
11:AA:887:VAL:CG1	16:AG:105:ILE:CG2	2.53	0.87
26:L:102:MET:SD	26:L:102:MET:N	2.46	0.87
28:N:5:ASP:OD2	28:N:77:ARG:NH1	2.07	0.87
17:C:27:ALA:O	17:C:30:LYS:HG2	1.74	0.87
39:Y:100:ILE:HD12	39:Y:105:LEU:HD11	1.54	0.87
47:g:16:CYS:HB2	47:g:37:CYS:HB3	1.54	0.87
63:w:24:MET:HA	63:w:24:MET:HE3	1.54	0.87
16:AG:302:LYS:O	16:AG:302:LYS:NZ	2.08	0.87
27:M:4:ARG:HG2	27:M:5:ARG:CZ	2.03	0.87
59:s:37:ARG:HA	59:s:118:MET:HE3	1.54	0.87
59:s:98:GLU:OE2	59:s:126:ALA:N	2.07	0.87
16:AG:302:LYS:HE3	16:AG:302:LYS:N	1.87	0.87
23:I:58:GLU:OE1	23:I:65:ARG:NH1	2.08	0.87
12:AB:6:LEU:CD1	14:AE:291:ILE:HD11	2.05	0.87
12:AB:148:LYS:NZ	30:P:100:ILE:HG22	1.89	0.87
14:AE:79:LYS:CE	16:AG:153:ASP:OD1	2.23	0.87
16:AG:254:MET:HA	16:AG:254:MET:HE3	1.55	0.87
20:F:31:GLU:OE2	20:F:35:ARG:NE	2.06	0.87
16:AG:486:ARG:O	16:AG:490:TRP:HD1	1.57	0.87
16:AG:430:PRO:HG2	16:AG:435:LEU:CD2	2.05	0.86
6:5:10:DT:H73	11:AA:62:TYR:HD2	1.29	0.86
6:5:10:DT:O4'	11:AA:62:TYR:OH	1.93	0.86
16:AG:453:VAL:HG13	16:AG:458:ASP:HB2	1.55	0.86
44:d:55:U:H1'	54:n:26:MET:HE2	1.57	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:87:GLN:NE2	1:0:88:GLY:O	2.09	0.86
7:6:25:DT:C2'	11:AA:496:LYS:HG3	2.05	0.86
11:AA:846:GLY:CA	16:AG:104:ARG:NH1	2.36	0.86
36:V:77:ARG:NH2	36:V:78:VAL:O	2.08	0.86
16:AG:362:TYR:C	16:AG:413:ALA:CB	2.48	0.86
62:v:77:PRO:HG2	62:v:80:VAL:HG21	1.57	0.86
41:a:2780:G:OP2	59:s:120:ARG:NH1	2.07	0.86
62:v:95:LEU:O	62:v:100:LYS:NZ	2.08	0.86
9:9:48:ALA:HB3	9:9:51:TYR:CE1	2.09	0.86
16:AG:248:VAL:HG13	16:AG:273:ILE:HB	1.56	0.86
33:S:90:ARG:HG3	33:S:92:GLU:OE1	1.73	0.86
12:AB:141:LEU:HA	12:AB:153:SER:N	1.91	0.85
12:AB:147:ASN:ND2	30:P:100:ILE:HG22	1.91	0.85
16:AG:61:ARG:HD3	16:AG:63:LEU:HD21	1.57	0.85
16:AG:387:VAL:CB	16:AG:388:PRO:HD2	2.06	0.85
39:Y:15:GLY:HA2	39:Y:50:LYS:HG3	1.57	0.85
6:5:10:DT:C5	11:AA:62:TYR:HD2	1.81	0.85
11:AA:846:GLY:CA	16:AG:104:ARG:NH2	2.25	0.85
16:AG:285:ILE:CG2	16:AG:293:VAL:HG12	2.06	0.85
12:AB:51:PHE:HZ	14:AE:288:PRO:HD3	1.40	0.85
16:AG:434:LEU:HD12	16:AG:454:CYS:O	1.75	0.85
12:AB:115:LYS:HZ3	12:AB:129:ILE:CG2	1.87	0.85
54:n:23:ASN:OD1	54:n:24:SER:N	2.09	0.85
16:AG:451:ARG:HH11	16:AG:467:LEU:HA	1.42	0.85
12:AB:145:LEU:HD11	30:P:84:VAL:HG12	1.57	0.85
19:E:9:LYS:O	19:E:13:GLN:NE2	2.10	0.85
48:h:175:ARG:HG3	48:h:181:MET:HE1	1.58	0.85
11:AA:853:ASP:O	16:AG:103:ASP:CG	2.19	0.85
39:Y:100:ILE:HD12	39:Y:105:LEU:CD1	2.05	0.85
11:AA:887:VAL:HG23	16:AG:105:ILE:HD11	1.57	0.85
9:9:58:THR:CG2	9:9:81:LEU:HD23	2.06	0.85
11:AA:854:ILE:O	16:AG:102:PHE:CE2	2.30	0.85
11:AA:855:PRO:CA	16:AG:35:THR:OG1	2.25	0.85
16:AG:243:LYS:HB3	16:AG:245:ILE:HD11	1.59	0.85
16:AG:393:LEU:HD23	16:AG:398:LEU:CD2	1.93	0.85
39:Y:75:ALA:CB	39:Y:112:LYS:HE3	2.07	0.85
6:5:10:DT:C6	11:AA:62:TYR:CZ	2.65	0.84
16:AG:141:VAL:HG22	16:AG:178:ARG:HG2	1.57	0.84
16:AG:210:ARG:HG3	16:AG:216:ILE:HB	1.59	0.84
39:Y:48:ILE:HG13	39:Y:49:GLU:H	1.41	0.84
36:V:25:ILE:O	36:V:27:ARG:NH2	2.10	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:Y:99:LYS:HD2	39:Y:140:GLU:OE1	1.75	0.84
8:7:2:U:OP2	18:D:1505:G:C8	2.30	0.84
47:g:26:SER:OG	54:n:140:GLU:OE2	1.94	0.84
11:AA:860:ALA:H	16:AG:38:LYS:CE	1.90	0.84
59:s:102:GLU:OE1	59:s:102:GLU:N	2.10	0.84
11:AA:855:PRO:C	16:AG:35:THR:CB	2.47	0.84
29:O:84:THR:HG21	29:O:103:PHE:HB3	1.59	0.84
63:w:38:LEU:HD13	63:w:111:ALA:HB2	1.59	0.84
6:5:9:DG:P	11:AA:473:ARG:CB	2.66	0.84
16:AG:187:ARG:CB	16:AG:188:PRO:HD2	2.03	0.84
10:A:33:U:OP1	27:M:77:SER:OG	1.94	0.84
38:X:81:MET:CE	38:X:92:ARG:HB3	2.06	0.84
39:Y:45:THR:HG22	39:Y:50:LYS:HD3	1.59	0.84
16:AG:425:LEU:HG	16:AG:429:LYS:CE	2.08	0.84
11:AA:861:ALA:H	16:AG:38:LYS:HZ3	1.22	0.84
16:AG:213:VAL:HG11	16:AG:251:CYS:HA	1.59	0.84
39:Y:116:MET:HG2	39:Y:124:MET:HG2	1.59	0.84
6:5:10:DT:C2	12:AB:72:THR:OG1	2.30	0.84
12:AB:115:LYS:CD	12:AB:129:ILE:CG2	2.55	0.84
61:u:122:VAL:CG1	61:u:125:LEU:HD21	2.07	0.84
16:AG:365:ILE:HG21	16:AG:409:ARG:HH12	1.42	0.83
54:n:73:SER:OG	54:n:79:ILE:O	1.96	0.83
11:AA:856:ASN:ND2	16:AG:35:THR:CG2	2.38	0.83
11:AA:377:THR:CB	12:AB:62:GLU:HB3	2.07	0.83
12:AB:145:LEU:HD21	30:P:84:VAL:O	1.76	0.83
16:AG:287:ALA:HB2	16:AG:331:LEU:HD13	1.58	0.83
16:AG:448:LEU:CD2	16:AG:473:LEU:HD11	2.05	0.83
12:AB:141:LEU:HD23	12:AB:143:LEU:CD1	2.07	0.83
16:AG:355:ALA:CA	16:AG:382:GLU:CD	2.49	0.83
6:5:9:DG:H5''	11:AA:473:ARG:CG	1.95	0.83
6:5:11:DA:C2'	12:AB:67:THR:HG21	2.08	0.83
16:AG:287:ALA:HA	16:AG:331:LEU:HD12	0.87	0.83
9:9:1:MET:HE1	9:9:7:ASP:HB3	1.61	0.83
12:AB:81:PHE:HD2	14:AE:293:ARG:HH11	0.90	0.83
16:AG:362:TYR:HA	16:AG:413:ALA:CB	2.09	0.83
66:z:78:LYS:HD3	66:z:117:LEU:CD2	2.09	0.83
16:AG:127:VAL:CG2	16:AG:192:GLY:HA2	2.09	0.83
16:AG:354:ALA:HA	16:AG:357:ASP:HB3	1.60	0.83
18:D:742:G:OP1	34:T:58:ARG:NH2	2.11	0.83
16:AG:252:VAL:HA	16:AG:259:VAL:HG23	1.61	0.83
42:b:36:ILE:HG21	42:b:39:ARG:HD3	1.59	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
50:j:152:PRO:HG3	50:j:156:PHE:CZ	2.14	0.83
62:v:7:THR:OG1	62:v:10:ARG:NH1	2.11	0.83
5:4:58:SER:OG	5:4:59:GLU:OE1	1.95	0.82
12:AB:110:PRO:CA	12:AB:114:ASP:OD2	2.26	0.82
16:AG:285:ILE:CB	16:AG:293:VAL:HG11	2.09	0.82
41:a:1257:C:H4'	52:l:78:TRP:CE2	2.14	0.82
41:a:1613:G:O2'	53:m:3:ARG:NH2	2.12	0.82
9:9:43:LYS:O	9:9:46:ARG:HG2	1.78	0.82
12:AB:124:GLU:C	12:AB:126:PHE:H	1.86	0.82
16:AG:365:ILE:CG2	16:AG:409:ARG:HH12	1.91	0.82
1:0:68:ARG:NH1	41:a:1223:G:OP1	2.10	0.82
16:AG:226:ALA:HA	16:AG:236:ILE:HG13	1.61	0.82
16:AG:451:ARG:NE	16:AG:470:ILE:HG13	1.95	0.82
23:I:22:TRP:CZ3	23:I:24:ALA:HB2	2.14	0.82
9:9:43:LYS:HG3	9:9:46:ARG:NH1	1.94	0.82
12:AB:117:ILE:HD13	12:AB:161:LYS:HZ1	1.44	0.82
11:AA:847:PRO:CD	16:AG:104:ARG:NH2	2.42	0.82
41:a:2298:A:OP1	54:n:72:LYS:NZ	2.11	0.82
52:l:112:LEU:HD12	52:l:117:ARG:HB2	1.62	0.82
56:p:104:ASN:ND2	56:p:114:ASP:OD1	2.12	0.82
16:AG:425:LEU:HG	16:AG:429:LYS:CD	2.10	0.82
10:A:6:G:O2'	10:A:7:G:O5'	1.96	0.82
16:AG:168:LEU:HD23	16:AG:231:GLY:HA3	1.56	0.82
16:AG:285:ILE:CG2	16:AG:293:VAL:CG1	2.57	0.82
10:B:6:G:O2'	10:B:7:G:O5'	1.96	0.82
21:G:105:LYS:O	21:G:109:GLN:NE2	2.12	0.82
60:t:12:ASP:OD2	60:t:85:VAL:HG13	1.79	0.82
61:u:122:VAL:HG13	61:u:125:LEU:HD11	1.61	0.82
16:AG:203:GLU:HA	16:AG:206:ILE:HG12	1.60	0.82
16:AG:433:ASP:C	16:AG:456:LEU:HG	2.03	0.82
41:a:2682:A:O2'	50:j:13:ARG:NE	2.12	0.82
56:p:7:ALA:O	56:p:69:ARG:NE	2.12	0.82
16:AG:362:TYR:HA	16:AG:413:ALA:HB1	1.62	0.82
18:D:311:C:OP1	35:U:31:ARG:NH2	2.13	0.82
39:Y:74:PRO:HG2	39:Y:77:VAL:HB	1.60	0.82
41:a:2393:U:H5''	55:o:30:ARG:CZ	2.09	0.82
58:r:137:GLU:HG2	58:r:138:VAL:HG23	1.61	0.82
16:AG:361:LYS:CD	16:AG:417:ILE:HG12	2.10	0.82
66:z:74:ILE:HD11	66:z:79:PHE:HA	1.60	0.82
14:AE:79:LYS:C	16:AG:144:LYS:HE3	2.04	0.81
16:AG:353:HIS:CA	16:AG:356:ILE:HG23	2.10	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:b:28:GLY:O	42:b:66:LYS:NZ	2.13	0.81
60:t:7:MET:HG3	60:t:18:ARG:CZ	2.10	0.81
9:9:46:ARG:HG3	9:9:94:ARG:NH2	1.94	0.81
14:AE:79:LYS:HG3	16:AG:144:LYS:NZ	1.94	0.81
20:F:37:PHE:HB2	31:Q:126:LYS:CE	2.10	0.81
56:p:121:ILE:HD12	56:p:141:ILE:HG22	1.60	0.81
6:5:10:DT:C4	11:AA:62:TYR:CD2	2.68	0.81
9:9:52:MET:HG2	9:9:95:LEU:HD11	1.62	0.81
11:AA:387:ASN:CB	11:AA:394:ARG:HH11	1.93	0.81
11:AA:887:VAL:HG23	16:AG:105:ILE:CD1	2.08	0.81
14:AE:79:LYS:CG	16:AG:144:LYS:NZ	2.44	0.81
16:AG:279:ASN:CG	16:AG:280:PRO:HD2	2.04	0.81
18:D:107:G:O6	19:E:10:ARG:NE	2.13	0.81
41:a:1655:A:H1'	50:j:118:PHE:HE2	1.46	0.81
8:7:2:U:OP1	18:D:1505:G:C8	2.33	0.81
11:AA:889:PRO:CG	16:AG:104:ARG:HH11	1.92	0.81
16:AG:133:HIS:ND1	16:AG:136:GLU:OE1	2.13	0.81
16:AG:363:LEU:HD23	16:AG:409:ARG:HB2	1.62	0.81
48:h:53:HIS:HE2	48:h:219:THR:HA	1.45	0.81
16:AG:12:VAL:HB	16:AG:16:LYS:HG3	1.61	0.81
39:Y:96:LYS:CE	39:Y:136:GLY:HA3	2.08	0.81
41:a:2356:U:O3'	42:b:20:ARG:NH1	2.14	0.81
9:9:25:ALA:HB3	9:9:99:PHE:CD2	2.16	0.81
11:AA:847:PRO:HD2	16:AG:104:ARG:HH22	1.43	0.81
6:5:11:DA:C4'	12:AB:67:THR:HG22	2.11	0.81
8:7:3:G:C5'	18:D:1500:A:H62	1.92	0.81
11:AA:481:LEU:HD23	12:AB:16:ARG:HD2	1.60	0.81
12:AB:38:ILE:HG12	12:AB:43:ARG:HG2	1.61	0.81
12:AB:118:ILE:HG21	12:AB:141:LEU:CD1	1.84	0.81
12:AB:141:LEU:HA	12:AB:152:HIS:C	2.06	0.81
16:AG:381:LEU:CA	16:AG:384:LEU:HD12	2.08	0.81
16:AG:430:PRO:CG	16:AG:435:LEU:HD21	2.11	0.81
16:AG:434:LEU:CG	16:AG:455:THR:C	2.54	0.81
39:Y:9:LYS:HE2	39:Y:55:PRO:HB3	1.63	0.81
39:Y:14:ALA:HB3	39:Y:51:GLY:H	1.45	0.81
11:AA:855:PRO:HB2	16:AG:35:THR:HB	1.61	0.81
14:AE:287:ALA:HB1	14:AE:292:VAL:HG13	1.60	0.81
66:z:69:ALA:HB1	66:z:74:ILE:HG23	1.63	0.81
11:AA:380:ALA:H	12:AB:65:HIS:HE1	1.00	0.81
11:AA:855:PRO:HG3	16:AG:105:ILE:HG22	1.62	0.81
11:AA:855:PRO:HB3	16:AG:35:THR:OG1	1.81	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:AE:81:ARG:HH21	14:AE:81:ARG:HG3	1.44	0.81
26:L:70:VAL:O	26:L:74:LEU:HD23	1.81	0.81
38:X:81:MET:HE1	38:X:92:ARG:CB	2.08	0.81
39:Y:40:ALA:HB1	39:Y:68:PHE:CE1	2.16	0.81
1:O:86:GLN:OE1	41:a:1225:G:H4'	1.81	0.80
4:3:46:GLN:OE1	4:3:56:GLY:HA2	1.82	0.80
39:Y:58:ILE:HG22	39:Y:60:VAL:HG23	1.63	0.80
41:a:1818:U:O5'	48:h:156:ARG:NH2	2.14	0.80
11:AA:377:THR:OG1	12:AB:62:GLU:CA	2.30	0.80
12:AB:81:PHE:HD2	14:AE:293:ARG:NH1	1.35	0.80
16:AG:379:SER:OG	16:AG:384:LEU:HD21	1.81	0.80
12:AB:120:GLU:HB2	12:AB:162:LEU:HG	1.63	0.80
12:AB:140:MET:CA	12:AB:152:HIS:O	2.29	0.80
16:AG:380:THR:O	16:AG:384:LEU:HD11	1.80	0.80
18:D:652:U:O3'	28:N:56:LYS:NZ	2.13	0.80
60:t:98:ARG:NH2	60:t:118:LEU:O	2.15	0.80
9:9:3:LEU:HD11	9:9:5:LEU:HD23	1.61	0.80
12:AB:145:LEU:HG	30:P:88:MET:HG3	1.62	0.80
12:AB:152:HIS:HB3	12:AB:154:VAL:HG12	1.62	0.80
16:AG:451:ARG:HH12	16:AG:467:LEU:CA	1.92	0.80
16:AG:425:LEU:HD23	16:AG:429:LYS:HE2	0.81	0.80
48:h:130:LEU:HA	48:h:189:ARG:HH11	1.46	0.80
41:a:2393:U:P	55:o:30:ARG:HE	2.04	0.80
11:AA:856:ASN:CG	16:AG:35:THR:CA	2.54	0.80
41:a:2356:U:H5''	42:b:20:ARG:HD2	1.61	0.80
9:9:77:VAL:HG13	9:9:114:GLU:OE1	1.81	0.80
54:n:117:LEU:HD13	54:n:176:PRO:HG2	1.64	0.80
28:N:9:ASP:OD2	28:N:13:ARG:NH1	2.15	0.80
16:AG:451:ARG:NH2	16:AG:470:ILE:HG13	1.97	0.80
52:l:111:GLU:OE2	52:l:114:ARG:NH1	2.14	0.80
9:9:118:ILE:HB	9:9:119:PRO:HD3	1.64	0.79
11:AA:855:PRO:CG	16:AG:35:THR:HG21	2.12	0.79
16:AG:440:VAL:CG2	16:AG:481:LEU:CD2	2.60	0.79
12:AB:144:ASN:ND2	12:AB:149:GLU:OE1	2.15	0.79
41:a:1276:A:O2'	63:w:20:MET:CE	2.30	0.79
9:9:112:ALA:O	9:9:123:ILE:HD11	1.83	0.79
11:AA:887:VAL:HG11	16:AG:105:ILE:CG2	2.12	0.79
16:AG:425:LEU:CA	16:AG:429:LYS:CG	2.60	0.79
18:D:275:G:H4'	36:V:16:LYS:HD2	1.64	0.79
41:a:2393:U:H5''	55:o:30:ARG:NH1	1.97	0.79
49:i:46:ASP:OD2	49:i:53:LYS:NZ	2.15	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:l:1:MET:HE1	52:l:20:GLY:CA	2.11	0.79
66:z:74:ILE:HD13	66:z:79:PHE:HD1	1.46	0.79
6:5:10:DT:C2'	12:AB:71:ALA:H	1.95	0.79
16:AG:451:ARG:NH1	16:AG:467:LEU:C	2.40	0.79
19:E:28:MET:HE1	19:E:58:VAL:HA	1.63	0.79
9:9:34:THR:O	9:9:38:MET:HG2	1.83	0.79
9:9:129:LEU:H	9:9:130:PRO:HD2	1.45	0.79
11:AA:887:VAL:CG1	16:AG:105:ILE:HG23	2.13	0.79
16:AG:235:LYS:HD3	16:AG:327:LEU:HG	1.65	0.79
27:M:18:PHE:HE2	27:M:47:LEU:HD21	1.48	0.79
8:7:2:U:OP1	18:D:1505:G:H8	1.65	0.79
12:AB:9:CYS:H	12:AB:53:ASN:HB3	1.48	0.79
16:AG:365:ILE:CD1	16:AG:406:LEU:CD2	2.57	0.79
29:O:56:ASP:O	29:O:60:LYS:NZ	2.16	0.79
16:AG:133:HIS:HD1	16:AG:136:GLU:CD	1.90	0.79
16:AG:447:LYS:O	16:AG:470:ILE:CD1	2.31	0.79
17:C:60:LYS:HZ3	18:D:735:C:P	2.05	0.79
41:a:2682:A:C8	50:j:11:MET:CE	2.65	0.79
47:g:18:CYS:HB3	47:g:40:CYS:CB	2.13	0.79
7:6:25:DT:H3'	11:AA:496:LYS:CG	2.13	0.79
11:AA:855:PRO:CA	16:AG:35:THR:CB	2.61	0.79
12:AB:138:ARG:CB	12:AB:154:VAL:O	2.30	0.79
11:AA:860:ALA:N	16:AG:38:LYS:CE	2.45	0.79
16:AG:363:LEU:CG	16:AG:410:ALA:N	2.45	0.79
27:M:20:SER:HB3	27:M:23:LEU:HD13	1.63	0.79
2:1:98:LYS:HD2	41:a:2012:G:OP1	1.83	0.79
9:9:3:LEU:CD1	9:9:4:ASN:H	1.96	0.79
16:AG:381:LEU:HA	16:AG:384:LEU:CD1	2.09	0.79
41:a:2636:C:H4'	50:j:81:GLU:OE1	1.82	0.79
12:AB:136:GLU:CD	12:AB:160:ARG:NH1	2.42	0.78
16:AG:412:ASN:O	16:AG:416:THR:HG23	1.84	0.78
28:N:27:MET:N	28:N:58:GLU:OE2	2.16	0.78
11:AA:887:VAL:C	16:AG:105:ILE:HD11	2.08	0.78
54:n:142:ASP:HB3	54:n:145:LYS:HE2	1.65	0.78
12:AB:110:PRO:HB3	12:AB:114:ASP:OD2	1.83	0.78
37:W:40:ILE:HG23	37:W:44:MET:SD	2.24	0.78
40:Z:14:MET:HG3	40:Z:15:SER:H	1.48	0.78
9:9:5:LEU:O	9:9:9:GLN:NE2	2.15	0.78
12:AB:54:TYR:HE2	14:AE:291:ILE:HG23	1.44	0.78
59:s:37:ARG:HA	59:s:118:MET:CE	2.14	0.78
64:x:104:GLN:NE2	64:x:108:ASP:OD2	2.16	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:353:HIS:C	16:AG:356:ILE:CG2	2.50	0.78
17:C:38:LYS:HE2	17:C:38:LYS:HA	1.64	0.78
41:a:1453:A:N1	63:w:74:GLU:OE1	2.16	0.78
12:AB:120:GLU:HB2	12:AB:162:LEU:CG	2.14	0.78
14:AE:79:LYS:HB3	16:AG:159:GLU:OE2	1.83	0.78
16:AG:284:VAL:HG12	16:AG:296:ILE:HD13	1.66	0.78
41:a:1406:U:O2'	41:a:1407:G:O5'	2.02	0.78
56:p:107:LEU:HB3	56:p:152:ARG:HE	1.47	0.78
16:AG:425:LEU:HG	16:AG:429:LYS:HE2	1.65	0.78
27:M:40:GLU:OE2	29:O:41:ARG:NH2	2.17	0.78
35:U:3:THR:HG21	35:U:5:ARG:CZ	2.13	0.78
10:A:18:G:N2	10:A:55:U:O2	2.17	0.78
16:AG:189:GLU:HB3	16:AG:194:GLN:HG3	1.64	0.78
22:H:306:VAL:HA	22:H:309:MET:SD	2.23	0.78
39:Y:9:LYS:HE2	39:Y:55:PRO:CG	2.14	0.78
39:Y:102:ARG:HD2	39:Y:141:ASP:CA	2.13	0.78
8:7:3:G:C5'	18:D:1500:A:N6	2.46	0.78
33:S:12:LYS:O	33:S:16:LEU:HD23	1.84	0.78
43:c:30:LEU:HD12	43:c:31:PRO:HD2	1.63	0.78
18:D:673:A:H2'	18:D:674:G:C8	2.18	0.78
7:6:23:DC:H5'	14:AE:259:ARG:HH12	1.43	0.77
9:9:60:LEU:HD23	9:9:64:VAL:HG21	1.65	0.77
9:9:81:LEU:HD21	41:a:1107:G:H4'	1.64	0.77
16:AG:385:ALA:HB1	16:AG:411:LYS:CE	2.04	0.77
24:J:62:ARG:NH1	24:J:69:GLU:OE2	2.16	0.77
60:t:8:LEU:C	60:t:18:ARG:HH12	1.92	0.77
16:AG:434:LEU:CD2	16:AG:456:LEU:N	2.46	0.77
39:Y:93:ASN:C	39:Y:94:LYS:HD2	2.10	0.77
57:q:2:LYS:NZ	57:q:32:LYS:O	2.17	0.77
12:AB:51:PHE:CZ	14:AE:288:PRO:CD	2.66	0.77
12:AB:139:SER:O	12:AB:153:SER:HA	1.85	0.77
16:AG:434:LEU:HD11	16:AG:455:THR:O	1.85	0.77
9:9:27:VAL:HG13	9:9:83:ALA:HB3	1.66	0.77
16:AG:354:ALA:HA	16:AG:357:ASP:CB	2.14	0.77
48:h:258:ARG:NH1	48:h:264:ASP:OD1	2.14	0.77
60:t:20:MET:HB3	60:t:44:LYS:NZ	1.98	0.77
16:AG:307:ILE:HB	16:AG:338:VAL:HA	1.66	0.77
6:5:10:DT:C2	12:AB:71:ALA:CB	2.61	0.77
9:9:134:GLU:OE2	9:9:136:ILE:HG22	1.85	0.77
12:AB:54:TYR:CE2	14:AE:291:ILE:CG2	2.65	0.77
26:L:42:TRP:HZ3	26:L:45:ARG:HH22	1.32	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:U:3:THR:HG21	35:U:5:ARG:NH1	1.99	0.77
5:4:50:MET:O	5:4:56:PHE:CE2	2.38	0.77
9:9:19:ALA:HA	9:9:70:GLU:HG2	1.66	0.77
9:9:94:ARG:HB2	9:9:97:LYS:HE2	1.67	0.77
16:AG:387:VAL:CG1	16:AG:388:PRO:CD	2.41	0.77
17:C:38:LYS:HD3	18:D:719:C:H1'	1.67	0.77
31:Q:87:LYS:CG	31:Q:113:VAL:HG23	2.15	0.77
12:AB:152:HIS:HB3	12:AB:154:VAL:CG1	2.15	0.77
41:a:196:A:OP2	61:u:47:ARG:NH1	2.18	0.77
41:a:2297:A:N1	41:a:2321:U:C4	2.53	0.77
16:AG:437:LEU:HD21	16:AG:456:LEU:HD13	0.81	0.77
11:AA:847:PRO:N	16:AG:104:ARG:NH1	2.33	0.76
11:AA:887:VAL:HB	16:AG:105:ILE:CG2	2.14	0.76
62:v:1:MET:HA	62:v:1:MET:HE3	1.67	0.76
2:1:11:ARG:CZ	2:1:98:LYS:HE3	2.15	0.76
12:AB:125:GLY:O	14:AE:87:LYS:C	2.27	0.76
16:AG:451:ARG:CZ	16:AG:470:ILE:HG13	2.16	0.76
18:D:550:G:O2'	32:R:116:LYS:NZ	2.18	0.76
41:a:2511:U:C4'	50:j:130:GLN:OE1	2.33	0.76
9:9:37:LYS:O	9:9:41:LEU:HG	1.86	0.76
18:D:375:U:P	35:U:70:ARG:HE	2.07	0.76
21:G:20:THR:HA	21:G:39:HIS:CE1	2.21	0.76
41:a:2636:C:HO2'	50:j:45:TYR:HH	1.26	0.76
59:s:15:TRP:HZ3	59:s:135:GLN:HA	1.50	0.76
16:AG:233:ARG:HB2	16:AG:327:LEU:HD23	1.66	0.76
16:AG:385:ALA:HB3	16:AG:411:LYS:HE3	1.57	0.76
6:5:11:DA:C4'	12:AB:67:THR:CG2	2.64	0.76
16:AG:283:PHE:CZ	16:AG:331:LEU:O	2.39	0.76
16:AG:402:THR:O	16:AG:405:ALA:HB3	1.84	0.76
42:b:68:LYS:NZ	42:b:70:GLU:OE1	2.19	0.76
12:AB:141:LEU:O	12:AB:151:LYS:CA	2.34	0.76
16:AG:131:ARG:HA	16:AG:186:VAL:HG11	1.68	0.76
16:AG:425:LEU:CB	16:AG:429:LYS:HE3	2.16	0.76
21:G:154:MET:CE	21:G:158:PRO:HD3	2.16	0.76
22:H:52:GLN:OE1	22:H:86:ARG:CB	2.34	0.76
38:X:57:ARG:NH2	47:g:35:ASP:OD1	2.19	0.76
39:Y:126:ARG:NH2	41:a:1083:U:OP1	2.19	0.76
41:a:1405:U:H2'	41:a:1406:U:C6	2.20	0.76
6:5:10:DT:C4	11:AA:62:TYR:CE2	2.74	0.76
16:AG:354:ALA:O	16:AG:356:ILE:N	2.19	0.76
16:AG:434:LEU:CD2	16:AG:455:THR:C	2.58	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:C:60:LYS:HZ2	18:D:734:G:H2'	1.49	0.76
18:D:1227:A:OP1	38:X:93:ARG:NH2	2.18	0.76
9:9:77:VAL:HG12	9:9:82:ILE:HG21	1.68	0.76
11:AA:860:ALA:H	16:AG:38:LYS:CD	1.98	0.76
18:D:376:G:H5''	35:U:5:ARG:HD2	1.67	0.76
12:AB:133:PRO:HA	12:AB:159:PHE:HZ	1.51	0.76
14:AE:59:ALA:HB1	14:AE:71:LEU:CD2	2.16	0.76
41:a:2335:A:OP2	64:x:13:ARG:CZ	2.33	0.76
65:y:71:GLU:OE1	65:y:101:ARG:NE	2.19	0.76
12:AB:141:LEU:CD2	12:AB:143:LEU:CD1	2.64	0.75
27:M:18:PHE:CE2	27:M:47:LEU:HD21	2.21	0.75
54:n:4:LEU:HD11	54:n:101:GLU:CA	2.16	0.75
16:AG:425:LEU:HD23	16:AG:429:LYS:NZ	1.99	0.75
16:AG:425:LEU:HA	16:AG:429:LYS:HE3	0.95	0.75
16:AG:425:LEU:O	16:AG:429:LYS:HE3	1.87	0.75
10:B:18:G:N2	10:B:55:U:O2	2.17	0.75
21:G:30:PHE:CD1	21:G:201:PRO:HG3	2.21	0.75
41:a:1801:A:P	48:h:146:MET:HE1	2.25	0.75
59:s:15:TRP:CZ3	59:s:135:GLN:HA	2.21	0.75
12:AB:125:GLY:O	14:AE:87:LYS:O	2.05	0.75
37:W:44:MET:SD	37:W:62:VAL:HG11	2.25	0.75
42:b:21:LEU:HD21	42:b:41:ARG:HE	1.51	0.75
11:AA:846:GLY:C	16:AG:104:ARG:CZ	2.58	0.75
31:Q:84:VAL:HG23	31:Q:107:ILE:HD11	1.67	0.75
41:a:686:U:O4	53:m:12:ARG:HD3	1.86	0.75
63:w:12:ARG:O	63:w:17:ARG:NH2	2.19	0.75
5:4:57:TYR:HB3	62:v:136:MET:CE	2.16	0.75
11:AA:856:ASN:CG	16:AG:35:THR:HG22	2.11	0.75
16:AG:336:LEU:HD12	16:AG:336:LEU:N	2.01	0.75
16:AG:385:ALA:HB3	16:AG:411:LYS:HE2	1.69	0.75
16:AG:434:LEU:CA	16:AG:456:LEU:HD21	2.12	0.75
18:D:739:C:O2'	34:T:42:HIS:ND1	2.18	0.75
41:a:1276:A:O2'	63:w:20:MET:HE2	1.87	0.75
8:7:3:G:H5''	18:D:1500:A:H62	1.49	0.75
16:AG:313:ASN:N	16:AG:313:ASN:OD1	2.19	0.75
22:H:36:VAL:O	22:H:37:LEU:HD12	1.87	0.75
31:Q:87:LYS:HG2	31:Q:113:VAL:HG23	1.67	0.75
39:Y:100:ILE:HD11	39:Y:139:VAL:HB	1.69	0.75
16:AG:240:THR:CG2	16:AG:247:PRO:HG3	2.16	0.75
16:AG:365:ILE:HG23	16:AG:409:ARG:HH11	1.51	0.75
18:D:375:U:OP2	35:U:70:ARG:NH2	2.17	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:1217:C:OP2	33:S:9:ARG:NH2	2.20	0.75
41:a:1828:G:O6	48:h:221:ARG:NH1	2.19	0.75
9:9:5:LEU:C	9:9:9:GLN:HE22	1.94	0.75
9:9:36:ASP:O	9:9:39:THR:HG22	1.87	0.75
16:AG:354:ALA:C	16:AG:356:ILE:H	1.93	0.75
16:AG:355:ALA:CB	16:AG:382:GLU:OE1	2.18	0.75
17:C:73:ARG:NH1	31:Q:113:VAL:HG12	2.02	0.75
41:a:742:A:H2'	41:a:743:A:C8	2.22	0.75
6:5:10:DT:H2''	12:AB:69:ILE:C	2.11	0.74
16:AG:354:ALA:C	16:AG:356:ILE:N	2.43	0.74
16:AG:451:ARG:HE	16:AG:470:ILE:HG13	1.50	0.74
18:D:404:G:OP1	24:J:115:ARG:NH1	2.19	0.74
21:G:49:MET:CE	21:G:201:PRO:HD3	2.15	0.74
41:a:2314:A:OP1	54:n:88:LYS:NZ	2.18	0.74
50:j:46:ARG:NH2	50:j:88:GLU:O	2.19	0.74
16:AG:371:THR:O	16:AG:375:GLU:HG3	1.86	0.74
22:H:303:LEU:O	22:H:305:HIS:N	2.20	0.74
24:J:105:MET:CE	24:J:171:LEU:HD13	2.18	0.74
39:Y:14:ALA:CB	39:Y:50:LYS:HD2	2.16	0.74
16:AG:62:TRP:HB3	16:AG:75:ILE:HD11	1.69	0.74
16:AG:365:ILE:HG23	16:AG:409:ARG:NH1	2.02	0.74
17:C:60:LYS:NZ	18:D:734:G:O3'	2.20	0.74
20:F:63:GLU:OE1	20:F:67:ARG:NH1	2.20	0.74
34:T:64:ARG:CZ	34:T:89:ARG:HH22	2.00	0.74
40:Z:2:ILE:HG22	40:Z:4:LYS:H	1.50	0.74
41:a:1278:C:H4'	63:w:24:MET:HE1	1.70	0.74
11:AA:387:ASN:OD1	11:AA:394:ARG:NH1	2.21	0.74
12:AB:145:LEU:HG	30:P:88:MET:CG	2.17	0.74
2:1:24:ILE:HG21	2:1:36:LEU:HD11	1.68	0.74
38:X:28:THR:HG23	38:X:31:LYS:HE3	1.69	0.74
39:Y:81:LYS:HZ2	39:Y:86:LYS:HE2	1.51	0.74
11:AA:855:PRO:HD2	16:AG:105:ILE:HG21	1.67	0.74
11:AA:887:VAL:CG1	16:AG:105:ILE:HG21	2.11	0.74
52:l:90:GLN:NE2	52:l:91:ASP:O	2.20	0.74
58:r:6:LEU:HD11	58:r:37:VAL:HG23	1.69	0.74
24:J:100:ASN:OD1	24:J:111:ARG:NH1	2.21	0.74
16:AG:393:LEU:C	16:AG:398:LEU:HD23	2.13	0.74
39:Y:9:LYS:HE2	39:Y:55:PRO:CB	2.17	0.74
47:g:18:CYS:CB	47:g:40:CYS:HB3	2.18	0.74
2:1:25:ARG:HH22	41:a:520:G:P	2.10	0.74
11:AA:377:THR:CG2	12:AB:62:GLU:HB3	2.18	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:248:VAL:HG22	16:AG:273:ILE:HG22	1.68	0.74
2:1:25:ARG:NH2	41:a:520:G:H5'	2.03	0.74
12:AB:89:PRO:CD	14:AE:290:ILE:HG21	2.18	0.74
12:AB:89:PRO:HD2	14:AE:290:ILE:HG21	1.68	0.74
12:AB:137:ALA:O	12:AB:155:LYS:C	2.30	0.74
16:AG:201:LYS:HD2	16:AG:201:LYS:N	2.01	0.74
16:AG:434:LEU:CA	16:AG:456:LEU:CD2	2.58	0.74
25:K:142:ASP:O	25:K:145:GLU:HG3	1.86	0.74
63:w:33:ILE:HD11	63:w:112:TYR:CD1	2.23	0.74
16:AG:228:ARG:HD2	16:AG:236:ILE:HB	1.67	0.73
17:C:18:VAL:O	17:C:51:TYR:OH	2.06	0.73
5:4:75:GLN:HG3	5:4:92:VAL:HG23	1.70	0.73
6:5:10:DT:N3	12:AB:72:THR:OG1	2.20	0.73
63:w:60:VAL:HG22	63:w:63:ARG:NH1	2.03	0.73
9:9:23:LEU:HD11	9:9:96:PHE:CD2	2.23	0.73
9:9:59:LEU:HD22	9:9:62:ARG:NE	2.02	0.73
16:AG:363:LEU:CD1	16:AG:410:ALA:HB2	2.17	0.73
38:X:81:MET:HE2	41:a:888:C:C6	2.24	0.73
54:n:56:ASP:OD2	54:n:150:ARG:NH1	2.20	0.73
60:t:17:ARG:HB2	60:t:45:GLU:OE1	1.88	0.73
12:AB:115:LYS:CE	12:AB:129:ILE:HG23	2.14	0.73
23:I:22:TRP:HZ3	23:I:24:ALA:HB2	1.54	0.73
39:Y:100:ILE:HG13	39:Y:139:VAL:CG2	2.18	0.73
56:p:69:ARG:NH1	56:p:73:ASN:OD1	2.21	0.73
9:9:43:LYS:NZ	9:9:98:GLU:HB2	2.04	0.73
9:9:78:GLY:N	9:9:79:PRO:HD2	2.04	0.73
11:AA:847:PRO:CD	16:AG:104:ARG:HH22	2.00	0.73
34:T:76:ALA:O	34:T:80:GLN:NE2	2.22	0.73
40:Z:6:GLN:OE1	40:Z:10:ALA:HB2	1.88	0.73
56:p:17:VAL:O	56:p:18:LYS:HE2	1.89	0.73
11:AA:887:VAL:HB	16:AG:105:ILE:HG23	1.69	0.73
12:AB:36:GLU:H	12:AB:106:ASP:CG	1.96	0.73
16:AG:198:THR:HB	16:AG:201:LYS:HD3	1.70	0.73
10:B:76:A:C2	41:a:2493:U:H4'	2.24	0.73
18:D:375:U:OP1	35:U:70:ARG:HG2	1.88	0.73
20:F:13:ASP:OD1	20:F:14:VAL:N	2.22	0.73
10:A:53:G:C2	10:A:54:U:C5	2.77	0.73
12:AB:21:LEU:HB3	12:AB:26:VAL:HG11	1.70	0.73
2:1:86:MET:HE1	2:1:88:ARG:HD3	1.70	0.73
16:AG:219:GLU:HA	16:AG:219:GLU:OE2	1.87	0.73
16:AG:258:ARG:HG2	16:AG:258:ARG:NH1	1.99	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:B:53:G:C2	10:B:54:U:C5	2.77	0.73
62:v:97:GLN:N	62:v:100:LYS:HE3	2.03	0.73
9:9:94:ARG:HB3	9:9:97:LYS:HZ3	1.54	0.73
12:AB:138:ARG:HD2	12:AB:155:LYS:HB2	1.70	0.73
16:AG:254:MET:HA	16:AG:254:MET:CE	2.17	0.73
16:AG:285:ILE:HG13	16:AG:293:VAL:HG13	1.71	0.73
16:AG:413:ALA:O	16:AG:416:THR:OG1	2.05	0.73
16:AG:434:LEU:HG	16:AG:455:THR:C	2.14	0.73
18:D:1158:C:O2'	21:G:132:LYS:NZ	2.21	0.73
19:E:21:ASN:C	19:E:25:ARG:HE	1.97	0.73
12:AB:18:GLN:HG3	12:AB:21:LEU:HD12	1.69	0.73
14:AE:79:LYS:CB	16:AG:144:LYS:NZ	2.33	0.73
16:AG:429:LYS:H	16:AG:429:LYS:HD2	1.53	0.73
14:AE:59:ALA:HB1	14:AE:71:LEU:HD22	1.71	0.72
16:AG:223:ILE:O	16:AG:223:ILE:HD13	1.89	0.72
41:a:537:G:OP2	59:s:2:LYS:NZ	2.20	0.72
66:z:92:ARG:HG3	66:z:95:LEU:HD12	1.71	0.72
5:4:45:ASP:OD1	5:4:46:LYS:N	2.21	0.72
16:AG:442:ARG:HB3	16:AG:446:PHE:HE2	1.53	0.72
41:a:465:G:OP1	53:m:12:ARG:NH2	2.22	0.72
41:a:2884:U:OP2	49:i:40:ARG:NH2	2.22	0.72
48:h:123:ALA:HB1	48:h:125:LYS:NZ	2.04	0.72
48:h:130:LEU:HA	48:h:189:ARG:NH1	2.03	0.72
49:i:55:ILE:HD13	63:w:112:TYR:CZ	2.23	0.72
9:9:129:LEU:HD12	9:9:130:PRO:HD3	1.70	0.72
16:AG:393:LEU:CD2	16:AG:398:LEU:CG	2.58	0.72
18:D:636:U:H4'	36:V:6:ARG:CZ	2.19	0.72
26:L:36:ILE:CD1	26:L:64:VAL:HG22	2.19	0.72
47:g:18:CYS:CB	47:g:40:CYS:CB	2.67	0.72
11:AA:860:ALA:H	16:AG:38:LYS:HD3	1.54	0.72
12:AB:120:GLU:HA	12:AB:124:GLU:HB3	1.72	0.72
16:AG:403:VAL:HG22	16:AG:406:LEU:HD12	1.71	0.72
16:AG:434:LEU:HD21	16:AG:456:LEU:N	2.03	0.72
20:F:32:VAL:HG12	20:F:36:GLU:OE1	1.90	0.72
34:T:21:ASP:O	34:T:22:THR:OG1	2.07	0.72
39:Y:102:ARG:CZ	39:Y:139:VAL:HG11	2.19	0.72
62:v:50:ARG:HA	62:v:53:MET:SD	2.29	0.72
18:D:13:U:C4	18:D:915:A:N6	2.57	0.72
54:n:134:GLU:HG2	54:n:137:ILE:HG23	1.72	0.72
9:9:59:LEU:HD13	9:9:62:ARG:HD2	1.70	0.72
14:AE:64:PRO:HG3	14:AE:90:VAL:CG1	2.20	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:Y:102:ARG:CD	39:Y:141:ASP:HA	2.18	0.72
9:9:47:GLU:HB3	9:9:51:TYR:CZ	2.24	0.72
12:AB:154:VAL:CG2	12:AB:159:PHE:HB3	2.20	0.72
24:J:58:LYS:NZ	24:J:69:GLU:OE2	2.15	0.72
24:J:184:ARG:NH1	24:J:187:GLU:OE1	2.22	0.72
39:Y:81:LYS:HD2	39:Y:81:LYS:N	2.05	0.72
2:1:23:LEU:HD11	49:i:24:ALA:HB2	1.70	0.72
12:AB:110:PRO:CB	12:AB:114:ASP:OD2	2.38	0.72
16:AG:365:ILE:HD13	16:AG:406:LEU:HD21	1.70	0.72
41:a:987:C:O3'	46:f:11:ARG:NH1	2.23	0.72
41:a:1825:U:OP2	48:h:52:ARG:NH1	2.23	0.72
43:c:71:LEU:HB3	43:c:76:GLU:OE2	1.88	0.72
62:v:73:ILE:HD11	62:v:93:VAL:HG22	1.70	0.72
16:AG:167:MET:HE1	16:AG:199:ARG:HG3	1.72	0.72
18:D:653:U:C6	28:N:56:LYS:HE3	2.25	0.72
24:J:76:TYR:CE2	24:J:204:TYR:HB2	2.24	0.72
38:X:66:GLU:OE1	38:X:66:GLU:N	2.23	0.72
47:g:5:ILE:HD12	54:n:64:LYS:HE3	1.72	0.72
56:p:85:LYS:HE3	56:p:164:TYR:CD1	2.24	0.72
59:s:15:TRP:CZ3	59:s:135:GLN:HG3	2.25	0.72
62:v:39:GLY:N	62:v:96:ILE:O	2.22	0.72
8:7:11:U:O2'	25:K:33:PHE:HE1	1.71	0.72
12:AB:137:ALA:O	12:AB:155:LYS:HA	1.90	0.72
27:M:55:GLY:O	27:M:56:LYS:O	2.07	0.72
5:4:77:VAL:CG2	5:4:79:ARG:HE	2.03	0.71
11:AA:846:GLY:CA	16:AG:104:ARG:CZ	2.63	0.71
12:AB:24:GLN:HB2	12:AB:26:VAL:HG23	1.70	0.71
12:AB:140:MET:SD	30:P:102:LEU:C	2.73	0.71
16:AG:297:VAL:HG23	16:AG:306:ASP:HB2	1.70	0.71
16:AG:365:ILE:CG2	16:AG:409:ARG:HH11	2.01	0.71
18:D:1328:C:O2'	38:X:29:ARG:NH2	2.23	0.71
23:I:166:GLU:OE1	23:I:166:GLU:N	2.23	0.71
41:a:1816:C:N4	48:h:35:GLU:OE2	2.20	0.71
41:a:2682:A:C8	50:j:11:MET:HE2	2.25	0.71
9:9:31:ARG:HD2	41:a:1053:C:HO2'	1.51	0.71
16:AG:127:VAL:HG22	16:AG:192:GLY:C	2.15	0.71
19:E:75:HIS:CE1	19:E:79:LEU:HD11	2.25	0.71
24:J:74:ASN:HA	24:J:77:LYS:HZ3	1.55	0.71
32:R:56:ARG:HG2	32:R:62:GLU:OE2	1.89	0.71
9:9:142:THR:OG1	40:Z:7:ILE:HG21	1.90	0.71
16:AG:262:VAL:HA	16:AG:265:GLU:HB2	1.72	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:C:38:LYS:NZ	18:D:718:A:C2	2.57	0.71
23:I:39:VAL:O	23:I:43:LEU:HD23	1.90	0.71
25:K:69:ARG:CZ	25:K:69:ARG:HA	2.20	0.71
12:AB:141:LEU:N	12:AB:153:SER:HB3	2.05	0.71
29:O:47:VAL:HA	29:O:50:GLN:HG2	1.70	0.71
41:a:2636:C:O2'	50:j:45:TYR:OH	1.99	0.71
10:B:13:C:OP2	41:a:1907:G:N2	2.23	0.71
23:I:142:MET:HE1	23:I:148:GLY:HA2	1.72	0.71
23:I:142:MET:HE3	23:I:170:GLU:OE1	1.89	0.71
41:a:1824:G:OP1	48:h:52:ARG:NH2	2.23	0.71
52:l:6:LYS:NZ	52:l:120:VAL:O	2.24	0.71
9:9:27:VAL:CG1	9:9:83:ALA:HB3	2.20	0.71
9:9:56:ARG:O	9:9:56:ARG:HG2	1.88	0.71
30:P:18:ILE:O	30:P:22:THR:HG23	1.90	0.71
60:t:8:LEU:C	60:t:18:ARG:NH1	2.48	0.71
11:AA:377:THR:OG1	12:AB:62:GLU:CG	2.38	0.71
14:AE:85:CYS:SG	68:AE:1502:ZN:ZN	1.79	0.71
16:AG:302:LYS:H	16:AG:302:LYS:CE	2.00	0.71
27:M:16:PRO:HB3	29:O:46:MET:HG3	1.73	0.71
41:a:2328:A:H2'	41:a:2329:U:C6	2.25	0.71
41:a:2469:A:N6	41:a:2481:G:O2'	2.23	0.71
41:a:2780:G:P	59:s:120:ARG:HH11	2.13	0.71
9:9:6:GLN:HA	9:9:9:GLN:NE2	2.06	0.71
9:9:56:ARG:NH2	41:a:1084:A:O4'	2.23	0.71
11:AA:889:PRO:HG3	16:AG:104:ARG:NH1	2.04	0.71
14:AE:77:ARG:NE	14:AE:77:ARG:HA	2.04	0.71
16:AG:352:ALA:O	16:AG:356:ILE:HG22	1.90	0.71
16:AG:486:ARG:O	16:AG:490:TRP:CD1	2.41	0.71
17:C:57:ARG:HE	17:C:61:ARG:HH12	1.37	0.71
18:D:311:C:P	35:U:31:ARG:HH22	2.13	0.71
25:K:111:MET:O	25:K:115:LEU:HD23	1.91	0.71
41:a:871:U:H2'	41:a:872:U:C6	2.26	0.71
2:1:83:LYS:HB2	2:1:95:ARG:HD3	1.72	0.71
8:7:13:U:H3	24:J:47:ARG:HH11	1.37	0.71
9:9:122:GLN:HG2	9:9:123:ILE:N	2.06	0.71
11:AA:887:VAL:O	16:AG:105:ILE:HD11	1.90	0.71
16:AG:220:VAL:O	16:AG:240:THR:HG22	1.90	0.71
33:S:90:ARG:HG3	33:S:92:GLU:CD	2.15	0.71
34:T:63:ARG:O	34:T:67:LEU:HD23	1.90	0.71
4:3:46:GLN:NE2	4:3:47:LYS:O	2.24	0.71
11:AA:854:ILE:C	16:AG:102:PHE:CD2	2.69	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AB:38:ILE:CD1	12:AB:160:ARG:NH1	2.54	0.71
16:AG:287:ALA:HB1	16:AG:331:LEU:HD12	1.62	0.71
18:D:636:U:H4'	36:V:6:ARG:NH2	2.06	0.71
47:g:16:CYS:CB	47:g:37:CYS:HB3	2.21	0.71
54:n:127:ASN:OD1	54:n:158:THR:N	2.18	0.71
55:o:25:LYS:HB3	61:u:62:PRO:HG2	1.71	0.71
12:AB:133:PRO:HA	12:AB:159:PHE:CZ	2.26	0.70
16:AG:282:GLN:NE2	16:AG:282:GLN:HA	2.05	0.70
16:AG:393:LEU:CD2	16:AG:398:LEU:HD13	2.19	0.70
39:Y:57:VAL:O	39:Y:69:VAL:HG22	1.91	0.70
11:AA:856:ASN:CG	16:AG:35:THR:CB	2.64	0.70
16:AG:354:ALA:O	16:AG:355:ALA:C	2.26	0.70
18:D:375:U:OP1	35:U:70:ARG:NE	2.24	0.70
18:D:973:G:O4'	30:P:57:VAL:HG22	1.90	0.70
31:Q:52:PHE:HD2	31:Q:56:ARG:HG2	1.56	0.70
1:O:10:LYS:HE2	41:a:994:C:O2'	1.90	0.70
12:AB:152:HIS:ND1	12:AB:159:PHE:CD2	2.49	0.70
14:AE:72:CYS:HG	14:AE:85:CYS:HG	1.37	0.70
16:AG:422:GLU:CB	16:AG:426:GLY:HA3	2.21	0.70
18:D:13:U:O4	18:D:21:G:C2	2.44	0.70
21:G:19:GLN:NE2	22:H:75:VAL:O	2.23	0.70
39:Y:19:PRO:HG2	39:Y:23:VAL:HG23	1.73	0.70
60:t:42:THR:O	60:t:44:LYS:NZ	2.19	0.70
12:AB:51:PHE:CE2	14:AE:288:PRO:HG2	2.06	0.70
12:AB:136:GLU:HA	12:AB:159:PHE:HE1	1.57	0.70
12:AB:141:LEU:H	12:AB:153:SER:HB3	1.56	0.70
39:Y:116:MET:CG	39:Y:124:MET:HG2	2.21	0.70
41:a:1012:U:O4	59:s:30:THR:HG21	1.92	0.70
44:d:42:C:O5'	54:n:64:LYS:NZ	2.20	0.70
10:A:6:G:HO2'	10:A:7:G:C5'	2.05	0.70
11:AA:65:ASN:HB3	11:AA:105:TYR:HB2	1.73	0.70
12:AB:141:LEU:CD2	12:AB:143:LEU:HD11	2.21	0.70
18:D:274:A:H4'	36:V:16:LYS:CE	2.22	0.70
18:D:736:C:C5'	26:L:90:MET:HE2	2.21	0.70
18:D:1376:U:OP2	27:M:25:LYS:NZ	2.24	0.70
26:L:42:TRP:CH2	26:L:102:MET:CG	2.73	0.70
38:X:71:ARG:CZ	38:X:72:GLU:HG3	2.22	0.70
14:AE:59:ALA:HB2	14:AE:71:LEU:HD21	1.74	0.70
16:AG:389:MET:O	16:AG:404:GLU:OE2	2.09	0.70
10:B:6:G:HO2'	10:B:7:G:C5'	2.04	0.70
18:D:1208:C:H2'	18:D:1209:C:H6	1.55	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:O:112:GLU:OE2	29:O:115:LYS:NZ	2.23	0.70
34:T:64:ARG:NH1	34:T:64:ARG:HA	2.05	0.70
41:a:2314:A:H1'	54:n:155:THR:HG21	1.74	0.70
41:a:2356:U:H5''	42:b:20:ARG:CD	2.21	0.70
52:l:6:LYS:NZ	52:l:120:VAL:N	2.39	0.70
14:AE:68:TYR:HA	14:AE:92:VAL:HG23	1.72	0.70
16:AG:362:TYR:CA	16:AG:413:ALA:CB	2.69	0.70
32:R:5:ASN:OD1	36:V:36:LYS:NZ	2.24	0.70
39:Y:100:ILE:HG13	39:Y:139:VAL:HG23	1.74	0.70
41:a:2350:C:OP1	55:o:45:ARG:NH2	2.25	0.70
58:r:33:GLN:OE1	58:r:35:LYS:NZ	2.24	0.70
63:w:42:LYS:N	63:w:42:LYS:HD3	2.05	0.70
12:AB:104:ILE:O	12:AB:111:TYR:OH	2.10	0.70
19:E:9:LYS:C	19:E:13:GLN:HE22	2.00	0.70
22:H:332:VAL:HG22	22:H:342:ILE:HD11	1.73	0.70
39:Y:19:PRO:O	39:Y:24:GLY:HA2	1.92	0.70
48:h:180:GLU:C	48:h:181:MET:HE2	2.17	0.70
59:s:73:VAL:C	59:s:74:TYR:CD2	2.70	0.70
3:2:56:GLU:OE2	3:2:88:LYS:HA	1.91	0.70
11:AA:854:ILE:O	16:AG:102:PHE:CD2	2.45	0.70
11:AA:855:PRO:CD	16:AG:105:ILE:HG21	2.22	0.70
11:AA:860:ALA:H	16:AG:38:LYS:HE2	1.56	0.70
24:J:76:TYR:HE2	24:J:204:TYR:HB2	1.57	0.70
6:5:9:DG:OP2	11:AA:470:ARG:HA	1.90	0.70
9:9:52:MET:HE1	9:9:85:SER:C	2.17	0.70
12:AB:18:GLN:HA	12:AB:21:LEU:HB2	1.74	0.70
12:AB:140:MET:SD	30:P:102:LEU:O	2.50	0.70
16:AG:279:ASN:CB	16:AG:280:PRO:CD	2.66	0.70
16:AG:353:HIS:HA	16:AG:356:ILE:CG2	2.21	0.70
16:AG:434:LEU:HD11	16:AG:455:THR:C	2.17	0.70
19:E:25:ARG:CD	19:E:66:LEU:HD11	2.22	0.70
23:I:24:ALA:HB3	23:I:29:PHE:CE2	2.27	0.70
29:O:7:TYR:N	29:O:89:GLU:OE2	2.24	0.70
38:X:28:THR:HG23	38:X:31:LYS:CE	2.22	0.70
54:n:56:ASP:HB3	54:n:150:ARG:NH2	2.07	0.70
12:AB:141:LEU:HB2	12:AB:143:LEU:CD1	2.18	0.69
16:AG:60:ARG:HG3	16:AG:98:GLU:HG3	1.74	0.69
25:K:93:ARG:HB3	25:K:93:ARG:CZ	2.21	0.69
56:p:107:LEU:O	56:p:152:ARG:NH1	2.25	0.69
2:1:7:HIS:CD2	2:1:10:ALA:HB2	2.27	0.69
11:AA:887:VAL:CB	16:AG:105:ILE:HG13	2.21	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:I:147:LYS:NZ	23:I:204:LYS:O	2.22	0.69
28:N:87:LYS:HD2	28:N:91:GLU:HB3	1.74	0.69
16:AG:433:ASP:C	16:AG:456:LEU:CG	2.64	0.69
19:E:45:ALA:O	19:E:48:GLN:HG3	1.92	0.69
21:G:26:LYS:HD2	21:G:193:PRO:HG2	1.74	0.69
36:V:16:LYS:HD3	36:V:16:LYS:C	2.17	0.69
62:v:105:MET:SD	62:v:117:PHE:CZ	2.85	0.69
12:AB:148:LYS:CE	30:P:100:ILE:HG22	2.17	0.69
16:AG:107:THR:HA	16:AG:110:ALA:HB3	1.75	0.69
16:AG:171:GLU:OE1	16:AG:267:GLY:HA3	1.92	0.69
16:AG:363:LEU:CD2	16:AG:409:ARG:CB	2.62	0.69
19:E:78:ASN:O	19:E:82:GLN:OE1	2.10	0.69
23:I:134:MET:HE3	23:I:151:VAL:HG23	1.75	0.69
41:a:2742:G:OP1	57:q:36:ARG:NH1	2.25	0.69
51:k:21:TYR:OH	51:k:39:PHE:O	2.10	0.69
16:AG:285:ILE:CD1	16:AG:293:VAL:HG11	2.22	0.69
16:AG:429:LYS:CD	16:AG:429:LYS:H	2.05	0.69
18:D:564:C:P	32:R:12:ARG:NH2	2.64	0.69
25:K:111:MET:CE	25:K:125:ALA:HB1	2.23	0.69
10:A:58:A:O2'	10:A:59:A:H3'	1.93	0.69
12:AB:148:LYS:HE2	30:P:100:ILE:HG21	1.70	0.69
32:R:27:CYS:HB2	32:R:30:LYS:HE2	1.75	0.69
41:a:1653:G:H3'	63:w:2:ARG:HD2	1.74	0.69
52:l:17:THR:HA	52:l:21:ARG:HH21	1.58	0.69
7:6:25:DT:H2'	11:AA:496:LYS:HG3	1.73	0.69
12:AB:128:ALA:HB1	12:AB:142:LEU:O	1.92	0.69
16:AG:239:LYS:HG3	16:AG:276:TRP:HB3	1.75	0.69
10:B:38:A:O2'	18:D:790:A:OP1	2.09	0.69
20:F:8:GLU:OE2	20:F:9:ASN:ND2	2.26	0.69
21:G:27:MET:SD	21:G:30:PHE:HB2	2.33	0.69
32:R:7:LEU:HD21	32:R:12:ARG:NH2	2.07	0.69
40:Z:2:ILE:HG22	40:Z:5:ASP:H	1.58	0.69
40:Z:14:MET:HB3	40:Z:17:MET:CG	2.21	0.69
52:l:101:TYR:CE2	52:l:105:LEU:HD11	2.28	0.69
12:AB:119:THR:HB	12:AB:123:PHE:O	1.93	0.69
12:AB:137:ALA:O	12:AB:155:LYS:CA	2.41	0.69
16:AG:171:GLU:OE2	16:AG:267:GLY:CA	2.41	0.69
16:AG:355:ALA:C	16:AG:380:THR:HG21	2.18	0.69
10:B:37:A:O2'	10:B:38:A:H5'	1.93	0.69
26:L:9:MET:HE1	26:L:85:ILE:CG2	2.22	0.69
27:M:66:LEU:HG	27:M:70:ARG:HE	1.58	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
62:v:1:MET:HE1	62:v:43:ALA:C	2.18	0.69
16:AG:279:ASN:CB	16:AG:280:PRO:HD2	2.22	0.69
22:H:22:SER:N	22:H:69:ASP:OD2	2.26	0.69
39:Y:118:GLY:O	39:Y:124:MET:HE3	1.93	0.69
46:f:16:ARG:NE	46:f:54:MET:SD	2.66	0.69
57:q:25:VAL:HB	57:q:35:GLN:OE1	1.92	0.69
61:u:123:ARG:NH2	61:u:143:GLU:OE2	2.25	0.69
2:l:25:ARG:HH22	41:a:520:G:C5'	2.04	0.69
9:9:92:ALA:HB1	9:9:129:LEU:HD22	1.74	0.69
12:AB:115:LYS:NZ	12:AB:129:ILE:HG22	2.04	0.69
12:AB:130:PHE:HA	12:AB:152:HIS:NE2	2.08	0.69
12:AB:154:VAL:HG21	12:AB:159:PHE:CB	2.23	0.69
16:AG:425:LEU:CB	16:AG:429:LYS:CE	2.71	0.69
16:AG:433:ASP:O	16:AG:437:LEU:HG	1.93	0.69
21:G:100:MET:HE1	21:G:101:LEU:HD21	1.75	0.69
41:a:2511:U:H4'	50:j:130:GLN:OE1	1.91	0.69
48:h:133:ARG:NE	48:h:187:ASP:OD2	2.25	0.69
8:7:11:U:H4'	25:K:20:ARG:HH21	1.53	0.68
9:9:43:LYS:HA	9:9:46:ARG:CZ	2.24	0.68
9:9:139:LEU:HD21	40:Z:11:VAL:HG22	1.75	0.68
16:AG:102:PHE:HB2	16:AG:106:THR:HG21	1.75	0.68
16:AG:353:HIS:C	16:AG:357:ASP:H	2.00	0.68
16:AG:462:GLN:OE1	16:AG:467:LEU:HD21	1.93	0.68
21:G:126:PHE:O	21:G:127:ASP:O	2.10	0.68
54:n:69:LYS:HD3	54:n:84:PRO:HA	1.74	0.68
10:A:37:A:O2'	10:A:38:A:H5'	1.93	0.68
12:AB:102:LYS:HE2	14:AE:285:LEU:HA	1.74	0.68
12:AB:113:GLY:H	12:AB:133:PRO:HD2	1.58	0.68
18:D:1308:U:P	38:X:100:GLN:HE22	2.16	0.68
34:T:76:ALA:C	34:T:80:GLN:HE22	2.00	0.68
38:X:90:ARG:CD	38:X:97:VAL:HA	2.23	0.68
39:Y:23:VAL:HG23	39:Y:24:GLY:H	1.58	0.68
39:Y:100:ILE:CG1	39:Y:139:VAL:HB	2.22	0.68
59:s:15:TRP:NE1	59:s:53:TYR:HD2	1.90	0.68
5:4:63:ILE:HD11	5:4:91:PHE:CD2	2.28	0.68
6:5:10:DT:H73	11:AA:62:TYR:CG	2.18	0.68
9:9:44:ALA:HB1	9:9:52:MET:HB3	1.74	0.68
41:a:1056:G:O2'	41:a:1103:A:N6	2.27	0.68
47:g:5:ILE:HB	54:n:64:LYS:HD3	1.75	0.68
63:w:45:ARG:O	63:w:49:GLU:OE1	2.10	0.68
5:4:53:LYS:HD2	5:4:56:PHE:CE2	2.28	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:7:11:U:C4'	25:K:20:ARG:HH22	1.93	0.68
11:AA:887:VAL:CB	16:AG:105:ILE:CG2	2.70	0.68
16:AG:440:VAL:HG23	16:AG:481:LEU:HD22	1.76	0.68
41:a:1190:G:OP1	61:u:30:THR:OG1	2.10	0.68
5:4:77:VAL:HG23	5:4:89:ILE:HG12	1.76	0.68
9:9:24:SER:CB	9:9:116:GLU:HG3	2.24	0.68
13:AD:48:LEU:HB2	13:AD:183:ILE:HD11	1.76	0.68
16:AG:298:VAL:HA	16:AG:303:HIS:HD1	1.58	0.68
22:H:304:VAL:O	22:H:304:VAL:HG12	1.94	0.68
31:Q:72:ASP:O	31:Q:75:LYS:NZ	2.21	0.68
34:T:27:VAL:O	34:T:31:LEU:HD23	1.93	0.68
39:Y:91:LYS:NZ	39:Y:97:VAL:HG21	2.07	0.68
41:a:1655:A:H1'	50:j:118:PHE:CE2	2.28	0.68
60:t:5:GLN:HA	60:t:20:MET:HE2	1.74	0.68
8:7:8:U:OP2	18:D:1397:C:N3	2.25	0.68
12:AB:147:ASN:HD22	30:P:100:ILE:HG22	1.57	0.68
18:D:1308:U:OP1	38:X:100:GLN:NE2	2.26	0.68
19:E:79:LEU:HA	19:E:82:GLN:OE1	1.94	0.68
27:M:4:ARG:HG2	27:M:5:ARG:NH1	2.08	0.68
31:Q:80:LYS:HB3	31:Q:106:ARG:NH1	2.09	0.68
33:S:19:LYS:HG3	33:S:20:TYR:CD1	2.29	0.68
62:v:111:GLU:O	62:v:115:GLU:OE1	2.10	0.68
2:1:19:LEU:HD21	49:i:22:LEU:HB2	1.75	0.68
12:AB:6:LEU:HD13	14:AE:291:ILE:CD1	2.21	0.68
18:D:564:C:OP1	32:R:12:ARG:NH1	2.27	0.68
34:T:28:GLN:O	34:T:32:LEU:HG	1.93	0.68
48:h:76:ALA:HB1	48:h:94:VAL:HG12	1.76	0.68
62:v:53:MET:HE3	62:v:120:ALA:HB2	1.74	0.68
66:z:68:ALA:HB1	66:z:106:PHE:CE2	2.29	0.68
1:0:52:PRO:HB3	66:z:89:GLU:OE2	1.94	0.68
2:1:25:ARG:HH22	41:a:520:G:H5'	1.58	0.68
4:3:72:ILE:HD12	4:3:83:VAL:HG21	1.76	0.68
12:AB:115:LYS:HE2	12:AB:129:ILE:HG23	1.76	0.68
18:D:1344:C:P	29:O:124:ARG:HH11	2.16	0.68
20:F:12:PHE:CE1	31:Q:107:ILE:HG22	2.29	0.68
22:H:308:GLU:O	22:H:310:ASP:N	2.26	0.68
25:K:156:LYS:HE3	28:N:71:VAL:HG13	1.76	0.68
26:L:18:VAL:HA	26:L:21:MET:HE2	1.75	0.68
63:w:24:MET:HE2	63:w:34:ILE:CD1	2.24	0.68
5:4:50:MET:HA	5:4:56:PHE:CZ	2.29	0.68
19:E:12:ILE:O	19:E:15:GLU:HG3	1.92	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:168:LEU:HB2	16:AG:171:GLU:HB2	1.76	0.68
16:AG:393:LEU:HD23	16:AG:398:LEU:CD1	2.20	0.68
10:B:24:U:O2'	41:a:1922:G:O3'	2.12	0.68
46:f:41:THR:O	46:f:44:ILE:HG22	1.94	0.68
56:p:85:LYS:CE	56:p:164:TYR:HD1	2.07	0.68
14:AE:1033:GLY:HA3	14:AE:1081:VAL:O	1.93	0.67
23:I:74:GLY:O	23:I:78:GLY:N	2.24	0.67
27:M:20:SER:CB	27:M:23:LEU:HD13	2.24	0.67
27:M:23:LEU:HD21	27:M:59:LEU:HD21	1.76	0.67
33:S:20:TYR:HE2	33:S:52:PRO:HG2	1.58	0.67
58:r:37:VAL:HG12	58:r:43:ASN:ND2	2.09	0.67
61:u:55:MET:HE1	61:u:60:ARG:HG2	1.75	0.67
62:v:1:MET:HE1	62:v:44:ARG:HA	1.76	0.67
5:4:75:GLN:HG3	5:4:92:VAL:CG2	2.24	0.67
5:4:89:ILE:HG21	5:4:91:PHE:CE1	2.29	0.67
11:AA:887:VAL:CB	16:AG:105:ILE:HG23	2.23	0.67
16:AG:266:LEU:H	16:AG:266:LEU:CD2	2.06	0.67
10:B:58:A:O2'	10:B:59:A:H3'	1.93	0.67
23:I:134:MET:HE1	23:I:167:TRP:HA	1.76	0.67
33:S:42:TRP:CZ3	47:g:66:ILE:HG23	2.29	0.67
48:h:205:LEU:HD12	48:h:210:ALA:HB1	1.75	0.67
60:t:112:PHE:HD2	60:t:115:ILE:HD12	1.59	0.67
5:4:57:TYR:OH	5:4:79:ARG:NH2	2.24	0.67
9:9:43:LYS:HZ2	9:9:101:LYS:HE2	1.58	0.67
9:9:47:GLU:HG3	9:9:94:ARG:CZ	2.24	0.67
12:AB:154:VAL:HG21	12:AB:159:PHE:CA	2.24	0.67
17:C:13:PHE:HB3	17:C:51:TYR:CE1	2.30	0.67
18:D:673:A:H5''	26:L:86:ARG:NH1	2.09	0.67
24:J:121:LYS:HG2	24:J:131:ASN:HD21	1.59	0.67
27:M:109:ARG:HD2	27:M:116:MET:HE1	1.76	0.67
39:Y:56:VAL:CA	39:Y:71:LYS:HZ1	2.01	0.67
46:f:40:ASP:C	46:f:45:ARG:HH22	2.03	0.67
54:n:94:GLU:HA	54:n:97:TRP:HE3	1.58	0.67
62:v:1:MET:HE1	62:v:44:ARG:N	2.09	0.67
63:w:38:LEU:HD13	63:w:111:ALA:CB	2.24	0.67
9:9:44:ALA:HA	9:9:47:GLU:OE1	1.95	0.67
12:AB:35:LEU:CA	12:AB:106:ASP:OD1	2.36	0.67
16:AG:453:VAL:CG2	16:AG:462:GLN:NE2	2.56	0.67
20:F:31:GLU:OE1	20:F:34:ARG:NH2	2.28	0.67
28:N:67:GLN:OE1	28:N:67:GLN:N	2.28	0.67
39:Y:99:LYS:NZ	39:Y:140:GLU:OE2	2.26	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:Z:20:VAL:O	40:Z:20:VAL:HG12	1.94	0.67
41:a:927:A:H2'	41:a:928:A:C8	2.30	0.67
41:a:1800:C:OP2	48:h:182:ARG:NH1	2.27	0.67
14:AE:288:PRO:HD2	14:AE:291:ILE:HG22	1.74	0.67
18:D:588:G:H4'	28:N:3:MET:HE1	1.77	0.67
18:D:945:G:C2	18:D:946:A:C8	2.83	0.67
27:M:46:ALA:CB	27:M:120:LEU:HD22	2.25	0.67
37:W:36:ARG:NH2	37:W:75:ALA:O	2.26	0.67
39:Y:37:PHE:CZ	39:Y:56:VAL:HG11	2.30	0.67
52:l:118:LEU:HD11	52:l:188:MET:SD	2.35	0.67
9:9:23:LEU:HD13	9:9:92:ALA:HB2	1.77	0.67
12:AB:89:PRO:CG	14:AE:290:ILE:HG21	2.24	0.67
25:K:115:LEU:HD12	25:K:120:VAL:HG11	1.75	0.67
27:M:111:ARG:O	27:M:119:ARG:NH2	2.27	0.67
39:Y:85:ILE:H	39:Y:85:ILE:HD12	1.59	0.67
39:Y:102:ARG:HD2	39:Y:141:ASP:HA	1.75	0.67
12:AB:34:THR:HA	12:AB:47:SER:HA	1.76	0.67
16:AG:207:GLU:HA	16:AG:210:ARG:HB2	1.76	0.67
16:AG:453:VAL:HG13	16:AG:458:ASP:HB3	1.77	0.67
22:H:23:ILE:N	22:H:69:ASP:OD2	2.27	0.67
26:L:42:TRP:HZ2	26:L:102:MET:SD	2.17	0.67
38:X:81:MET:CE	41:a:888:C:C5	2.78	0.67
66:z:79:PHE:HE1	66:z:110:VAL:HA	1.60	0.67
8:7:8:U:H5'	18:D:1397:C:N3	2.10	0.67
9:9:43:LYS:NZ	9:9:101:LYS:HE2	2.09	0.67
12:AB:154:VAL:HG21	12:AB:159:PHE:HB3	1.77	0.67
16:AG:284:VAL:CG1	16:AG:296:ILE:HD13	2.25	0.67
17:C:70:TYR:CE2	18:D:674:G:H4'	2.30	0.67
30:P:47:GLU:N	30:P:47:GLU:OE1	2.23	0.67
35:U:45:GLU:HG3	35:U:46:LYS:HD3	1.76	0.67
39:Y:116:MET:HG2	39:Y:124:MET:CG	2.24	0.67
41:a:111:A:O3'	45:e:58:ASN:ND2	2.28	0.67
41:a:2420:C:H5''	51:k:8:LYS:CE	2.25	0.67
43:c:71:LEU:HB3	43:c:76:GLU:CD	2.20	0.67
52:l:6:LYS:HE3	52:l:119:ILE:HG23	1.75	0.67
7:6:20:DA:N6	8:7:48:G:N1	2.42	0.67
9:9:50:VAL:CG2	39:Y:119:ALA:HB3	2.20	0.67
9:9:73:LYS:HG2	9:9:117:LEU:HD12	1.76	0.67
10:A:32:C:H4'	27:M:86:GLN:NE2	2.10	0.67
16:AG:235:LYS:HZ2	16:AG:331:LEU:HG	1.59	0.67
19:E:25:ARG:HD2	19:E:66:LEU:HD11	1.75	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:L:35:LYS:NZ	26:L:36:ILE:O	2.26	0.67
39:Y:32:VAL:HG22	39:Y:60:VAL:HG21	1.77	0.67
41:a:666:A:H1'	55:o:4:ILE:HD12	1.76	0.67
60:t:9:ASN:OD1	60:t:10:VAL:N	2.27	0.67
60:t:103:VAL:HG22	60:t:104:THR:H	1.58	0.67
5:4:84:PRO:O	5:4:85:LYS:HE2	1.95	0.67
10:B:75:C:OP2	41:a:2602:A:C2'	2.43	0.67
39:Y:45:THR:HG21	39:Y:50:LYS:HD3	1.76	0.67
57:q:1:MET:HE1	57:q:35:GLN:CA	2.25	0.67
1:0:55:ASP:OD1	1:0:56:GLY:N	2.27	0.66
10:A:71:C:H3'	10:A:72:A:C8	2.31	0.66
18:D:274:A:H4'	36:V:16:LYS:HE3	1.77	0.66
41:a:752:A:OP1	53:m:1:MET:HE1	1.95	0.66
5:4:50:MET:HB3	5:4:56:PHE:CE1	2.31	0.66
11:AA:860:ALA:N	16:AG:38:LYS:HD3	2.02	0.66
16:AG:235:LYS:NZ	16:AG:331:LEU:HG	2.11	0.66
16:AG:363:LEU:CD2	16:AG:409:ARG:CA	2.71	0.66
16:AG:447:LYS:O	16:AG:470:ILE:HD13	1.95	0.66
37:W:62:VAL:HG13	37:W:66:MET:HE3	1.77	0.66
18:D:1151:A:HO2'	18:D:1152:A:P	2.18	0.66
26:L:38:ARG:NH1	26:L:99:ALA:HB2	2.09	0.66
29:O:86:ALA:O	29:O:89:GLU:HG2	1.95	0.66
37:W:40:ILE:CD1	37:W:66:MET:SD	2.77	0.66
39:Y:48:ILE:HG13	39:Y:49:GLU:N	2.10	0.66
41:a:2682:A:C8	50:j:11:MET:HE1	2.30	0.66
59:s:15:TRP:NE1	59:s:53:TYR:CD2	2.63	0.66
4:3:26:LYS:HB2	4:3:35:ILE:HG23	1.78	0.66
12:AB:119:THR:O	12:AB:124:GLU:HB3	1.95	0.66
41:a:1665:A:O5'	60:t:66:LYS:NZ	2.28	0.66
41:a:2420:C:H5''	51:k:8:LYS:NZ	2.09	0.66
41:a:2682:A:O4'	50:j:11:MET:HE1	1.95	0.66
52:l:149:ILE:HD11	52:l:188:MET:HG2	1.78	0.66
6:5:10:DT:H1'	12:AB:71:ALA:H	0.83	0.66
11:AA:847:PRO:O	16:AG:104:ARG:CZ	2.44	0.66
11:AA:889:PRO:HG3	16:AG:104:ARG:CD	2.23	0.66
16:AG:321:ASN:OD1	16:AG:321:ASN:N	2.25	0.66
41:a:1802:A:H2'	41:a:1803:A:C8	2.30	0.66
60:t:1:MET:HE1	60:t:32:TYR:CD1	2.31	0.66
11:AA:889:PRO:CG	16:AG:104:ARG:HD3	2.23	0.66
16:AG:323:GLN:HA	16:AG:323:GLN:HE21	1.59	0.66
10:B:71:C:H3'	10:B:72:A:C8	2.30	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:1228:C:OP2	38:X:110:LYS:NZ	2.28	0.66
19:E:21:ASN:C	19:E:25:ARG:NE	2.54	0.66
22:H:303:LEU:O	22:H:303:LEU:HG	1.96	0.66
24:J:196:ASN:OD1	24:J:198:HIS:CE1	2.48	0.66
48:h:129:THR:HG22	48:h:189:ARG:HD2	1.77	0.66
54:n:93:GLY:O	54:n:96:MET:N	2.29	0.66
59:s:43:GLU:OE1	59:s:43:GLU:N	2.29	0.66
6:5:10:DT:H73	11:AA:62:TYR:CB	2.25	0.66
12:AB:27:ASN:CG	12:AB:58:GLU:HB2	2.20	0.66
16:AG:448:LEU:CD2	16:AG:473:LEU:HD12	2.24	0.66
17:C:73:ARG:HH12	31:Q:113:VAL:CB	2.09	0.66
18:D:876:C:HI'	28:N:12:THR:HG21	1.76	0.66
21:G:101:LEU:HD22	21:G:157:LEU:CD1	2.26	0.66
60:t:38:ILE:HD11	60:t:112:PHE:HZ	1.60	0.66
62:v:45:GLN:NE2	62:v:125:PRO:HD3	2.10	0.66
9:9:139:LEU:CD2	40:Z:11:VAL:HG22	2.26	0.66
21:G:90:PHE:CD2	21:G:154:MET:HE3	2.31	0.66
23:I:151:VAL:HG12	23:I:200:VAL:HG23	1.77	0.66
58:r:137:GLU:HG2	58:r:138:VAL:N	2.11	0.66
59:s:13:ARG:HH22	59:s:50:THR:HA	1.60	0.66
18:D:375:U:P	35:U:70:ARG:NE	2.69	0.66
22:H:308:GLU:HG2	22:H:309:MET:H	1.61	0.66
34:T:64:ARG:NH1	34:T:89:ARG:HH22	1.94	0.66
41:a:2175:C:H2'	41:a:2176:A:C8	2.31	0.66
50:j:3:GLY:O	50:j:4:LEU:HD12	1.95	0.66
60:t:1:MET:HE1	60:t:32:TYR:CG	2.31	0.66
9:9:25:ALA:HB2	9:9:96:PHE:CE1	2.31	0.66
11:AA:860:ALA:N	16:AG:38:LYS:HZ3	1.94	0.66
27:M:26:PHE:HD1	27:M:101:MET:SD	2.19	0.66
29:O:5:GLN:NE2	29:O:20:PHE:HB3	2.11	0.66
38:X:90:ARG:HD2	38:X:97:VAL:HA	1.77	0.66
41:a:617:G:H4'	52:l:199:MET:HE1	1.78	0.66
5:4:56:PHE:O	5:4:61:LEU:HD11	1.95	0.65
9:9:28:ALA:H	9:9:110:ALA:HA	1.61	0.65
9:9:47:GLU:HG2	9:9:95:LEU:HD21	1.78	0.65
14:AE:58:CYS:SG	14:AE:59:ALA:N	2.69	0.65
18:D:246:A:C2	18:D:282:A:C5	2.84	0.65
19:E:54:MET:HE1	19:E:75:HIS:CE1	2.31	0.65
20:F:58:LYS:O	20:F:62:ARG:HG2	1.96	0.65
21:G:134:ALA:O	21:G:137:ARG:HG2	1.96	0.65
22:H:163:ARG:N	22:H:298:GLU:OE2	2.28	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:O:83:ILE:O	29:O:87:LEU:HD23	1.96	0.65
41:a:614:A:O2'	41:a:615:U:OP2	2.12	0.65
6:5:10:DT:C7	11:AA:62:TYR:CB	2.74	0.65
39:Y:79:LEU:HD21	39:Y:105:LEU:CD2	2.24	0.65
66:z:85:LYS:HD2	66:z:116:ALA:HB1	1.78	0.65
9:9:43:LYS:HD3	9:9:98:GLU:HG3	1.78	0.65
12:AB:120:GLU:HB2	12:AB:162:LEU:HB2	1.77	0.65
16:AG:213:VAL:HG22	16:AG:216:ILE:HG13	1.78	0.65
16:AG:232:SER:HB2	16:AG:323:GLN:OE1	1.96	0.65
10:B:22:G:H2'	10:B:23:C:C6	2.32	0.65
17:C:51:TYR:CD1	17:C:54:GLN:NE2	2.63	0.65
21:G:23:TRP:CD1	21:G:39:HIS:HE1	2.14	0.65
21:G:27:MET:HG2	21:G:189:THR:HA	1.78	0.65
32:R:30:LYS:HB3	32:R:57:LEU:HD11	1.78	0.65
33:S:10:GLU:O	33:S:14:VAL:HG23	1.97	0.65
41:a:1454:C:O4'	63:w:63:ARG:NE	2.28	0.65
41:a:1596:A:H2'	41:a:1597:A:C8	2.32	0.65
52:l:194:LYS:O	52:l:198:GLU:OE1	2.15	0.65
60:t:99:ILE:HD13	60:t:118:LEU:HB2	1.77	0.65
6:5:11:DA:H5'	12:AB:67:THR:CB	2.27	0.65
25:K:137:VAL:O	25:K:140:THR:OG1	2.15	0.65
32:R:80:ILE:HD12	32:R:104:CYS:HB2	1.79	0.65
41:a:2334:U:H4'	41:a:2335:A:OP2	1.95	0.65
61:u:122:VAL:CG1	61:u:125:LEU:HD11	2.27	0.65
9:9:14:GLU:O	9:9:18:VAL:HG23	1.96	0.65
10:A:22:G:H2'	10:A:23:C:C6	2.32	0.65
16:AG:308:ALA:HB2	16:AG:344:LEU:HD11	1.77	0.65
16:AG:385:ALA:HB2	16:AG:411:LYS:HE3	1.70	0.65
39:Y:20:SER:HB2	39:Y:21:PRO:HD3	1.79	0.65
39:Y:81:LYS:NZ	39:Y:86:LYS:HE2	2.12	0.65
41:a:1824:G:OP1	48:h:52:ARG:CZ	2.44	0.65
41:a:2356:U:O3'	42:b:20:ARG:CZ	2.45	0.65
2:1:25:ARG:NH2	41:a:520:G:C5'	2.60	0.65
12:AB:81:PHE:HE1	14:AE:293:ARG:HB3	0.85	0.65
16:AG:228:ARG:HA	16:AG:327:LEU:CD1	2.22	0.65
39:Y:102:ARG:HD2	39:Y:141:ASP:CB	2.27	0.65
41:a:2419:U:H4'	51:k:22:THR:HG21	1.79	0.65
56:p:107:LEU:O	56:p:152:ARG:CZ	2.45	0.65
59:s:53:TYR:CE1	59:s:121:LYS:HD2	2.32	0.65
3:2:9:LYS:O	3:2:12:ARG:NH1	2.30	0.65
4:3:34:VAL:HG13	4:3:67:VAL:HG22	1.78	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:5:9:DG:O5'	11:AA:473:ARG:HB3	1.95	0.65
9:9:24:SER:OG	9:9:86:MET:HE2	1.96	0.65
16:AG:123:ARG:HG2	16:AG:191:ARG:O	1.96	0.65
16:AG:174:ARG:HB3	16:AG:175:PRO:CD	2.25	0.65
16:AG:425:LEU:HG	16:AG:429:LYS:HD3	1.78	0.65
41:a:2308:G:C5	54:n:77:PHE:CZ	2.85	0.65
41:a:2723:C:H5''	63:w:1:MET:HE3	1.78	0.65
48:h:53:HIS:NE2	48:h:219:THR:HA	2.11	0.65
52:l:40:ARG:HD2	52:l:92:HIS:ND1	2.11	0.65
6:5:16:DG:H1	11:AA:451:ARG:NH2	1.94	0.65
18:D:738:C:H5''	26:L:68:GLN:NE2	2.12	0.65
20:F:58:LYS:O	20:F:62:ARG:CG	2.45	0.65
26:L:2:ARG:CZ	26:L:68:GLN:NE2	2.60	0.65
34:T:80:GLN:O	34:T:83:GLU:HG3	1.95	0.65
39:Y:14:ALA:C	39:Y:50:LYS:HE3	2.22	0.65
39:Y:102:ARG:HG3	39:Y:140:GLU:O	1.96	0.65
40:Z:22:LEU:O	40:Z:26:MET:HE1	1.97	0.65
41:a:2572:A:N7	50:j:150:GLN:NE2	2.45	0.65
43:c:66:THR:HG22	43:c:70:GLU:OE2	1.97	0.65
55:o:30:ARG:HH12	61:u:62:PRO:CB	2.09	0.65
62:v:111:GLU:HG2	62:v:112:LEU:HD12	1.78	0.65
12:AB:9:CYS:HB3	12:AB:74:GLY:HA2	1.78	0.65
16:AG:361:LYS:CD	16:AG:417:ILE:CG1	2.75	0.65
17:C:34:THR:N	17:C:38:LYS:O	2.30	0.65
18:D:280:C:H1'	36:V:40:ARG:NH2	2.11	0.65
18:D:714:G:H2'	18:D:715:A:C8	2.32	0.65
20:F:10:GLU:OE2	20:F:18:ARG:NE	2.30	0.65
21:G:90:PHE:CE2	21:G:154:MET:HE3	2.31	0.65
28:N:10:MET:HG3	28:N:27:MET:HE1	1.79	0.65
33:S:42:TRP:HH2	47:g:66:ILE:HA	1.61	0.65
41:a:2355:G:C2'	41:a:2356:U:H5'	2.26	0.65
52:l:147:LEU:HB2	52:l:183:PHE:HD2	1.62	0.65
54:n:117:LEU:HD13	54:n:176:PRO:CG	2.26	0.65
62:v:45:GLN:NE2	62:v:125:PRO:CD	2.59	0.65
9:9:129:LEU:N	9:9:130:PRO:HD2	2.12	0.65
12:AB:136:GLU:O	12:AB:156:ASN:HA	1.97	0.65
19:E:81:ALA:C	19:E:85:LYS:HZ3	2.04	0.65
32:R:25:GLU:CD	32:R:59:ASN:HD22	2.05	0.65
35:U:5:ARG:HG2	35:U:68:SER:HB2	1.79	0.65
46:f:10:THR:HB	46:f:56:LYS:HZ2	1.61	0.65
54:n:135:GLN:HG3	54:n:150:ARG:NH2	2.12	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:9:24:SER:OG	9:9:116:GLU:HG3	1.97	0.64
12:AB:139:SER:C	12:AB:152:HIS:O	2.40	0.64
12:AB:152:HIS:HE2	12:AB:159:PHE:HE2	1.44	0.64
16:AG:64:VAL:HG22	16:AG:75:ILE:HD12	1.78	0.64
16:AG:355:ALA:CB	16:AG:382:GLU:CG	2.54	0.64
16:AG:393:LEU:CD2	16:AG:398:LEU:HD22	1.90	0.64
16:AG:393:LEU:HD23	16:AG:398:LEU:HD13	1.78	0.64
24:J:138:SER:O	24:J:141:ASP:CG	2.40	0.64
26:L:25:TYR:O	26:L:29:ILE:HD12	1.97	0.64
27:M:23:LEU:HD21	27:M:59:LEU:CD2	2.27	0.64
33:S:53:ARG:HB3	33:S:59:ARG:NH1	2.11	0.64
33:S:87:ALA:CA	33:S:90:ARG:NH1	2.60	0.64
36:V:60:GLU:OE1	36:V:76:VAL:CG2	2.45	0.64
38:X:16:VAL:HG23	38:X:17:ILE:HD12	1.80	0.64
39:Y:102:ARG:NE	39:Y:102:ARG:HA	2.11	0.64
41:a:196:A:C2	61:u:50:PHE:HZ	2.15	0.64
50:j:55:LYS:HZ1	50:j:59:ARG:C	2.03	0.64
56:p:149:ARG:CZ	56:p:162:VAL:O	2.45	0.64
18:D:675:A:H1'	31:Q:118:HIS:CD2	2.33	0.64
41:a:2682:A:H2'	41:a:2683:C:O4'	1.96	0.64
10:A:19:G:C2	41:a:2112:G:O4'	2.50	0.64
12:AB:123:PHE:H	12:AB:123:PHE:HD2	1.46	0.64
12:AB:154:VAL:HG21	12:AB:159:PHE:N	2.13	0.64
12:AB:155:LYS:HG3	12:AB:156:ASN:H	1.62	0.64
16:AG:162:ILE:HD11	16:AG:199:ARG:HD2	1.80	0.64
16:AG:393:LEU:O	16:AG:398:LEU:HD23	1.95	0.64
17:C:73:ARG:HH12	31:Q:113:VAL:HA	1.62	0.64
21:G:53:ALA:HA	21:G:56:GLU:OE1	1.96	0.64
52:l:45:ALA:HB1	52:l:88:ARG:NH2	2.12	0.64
62:v:105:MET:CE	62:v:108:VAL:HG21	2.28	0.64
66:z:41:LYS:HG3	66:z:45:TYR:CE2	2.32	0.64
9:9:100:ALA:HB2	9:9:125:ARG:NE	2.13	0.64
12:AB:38:ILE:HD13	12:AB:160:ARG:NH1	2.13	0.64
13:AD:286:GLU:H	13:AD:315:GLY:CA	2.11	0.64
16:AG:227:ALA:HB3	16:AG:331:LEU:HA	1.78	0.64
25:K:148:ASN:OD1	28:N:96:MET:CE	2.45	0.64
34:T:53:ARG:O	34:T:57:LEU:HD23	1.97	0.64
52:l:112:LEU:HG	52:l:118:LEU:HB2	1.79	0.64
56:p:52:PHE:CD1	56:p:69:ARG:HD3	2.32	0.64
1:0:51:VAL:HG23	66:z:86:ALA:O	1.98	0.64
3:2:36:LYS:HE3	3:2:79:ASP:OD2	1.98	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AB:38:ILE:HD12	12:AB:160:ARG:HH11	1.63	0.64
18:D:1308:U:P	38:X:100:GLN:NE2	2.70	0.64
18:D:1323:G:H2'	18:D:1324:A:C8	2.33	0.64
19:E:16:LYS:O	19:E:19:LYS:HG2	1.97	0.64
21:G:26:LYS:NZ	21:G:193:PRO:HG2	2.12	0.64
25:K:61:GLN:HA	25:K:64:MET:CE	2.27	0.64
38:X:28:THR:HA	38:X:31:LYS:HE3	1.78	0.64
48:h:52:ARG:HE	48:h:53:HIS:HE1	1.44	0.64
52:l:6:LYS:HE3	52:l:119:ILE:CG2	2.28	0.64
52:l:62:GLN:O	52:l:69:ARG:NE	2.30	0.64
60:t:42:THR:CG2	60:t:44:LYS:HZ1	2.10	0.64
3:2:3:ARG:NH1	3:2:5:GLU:OE2	2.30	0.64
10:A:76:A:N6	41:a:2422:C:O5'	2.29	0.64
14:AE:79:LYS:HG3	16:AG:144:LYS:HZ3	1.61	0.64
10:B:54:U:H3	10:B:58:A:N6	1.96	0.64
18:D:1158:C:O2'	21:G:132:LYS:CE	2.45	0.64
19:E:28:MET:SD	19:E:61:GLN:HG3	2.36	0.64
27:M:23:LEU:HD12	27:M:23:LEU:N	2.13	0.64
39:Y:99:LYS:HD2	39:Y:140:GLU:CD	2.22	0.64
45:e:13:GLU:O	45:e:17:GLU:OE1	2.16	0.64
60:t:8:LEU:N	60:t:18:ARG:NH1	2.46	0.64
4:3:68:SER:OG	41:a:335:C:O2	2.15	0.64
8:7:44:U:C5'	11:AA:1259:LEU:HD21	2.27	0.64
9:9:56:ARG:H	9:9:56:ARG:HD3	1.63	0.64
12:AB:31:PRO:HB2	12:AB:50:LEU:HB3	1.80	0.64
16:AG:183:LEU:HA	16:AG:197:VAL:HA	1.78	0.64
18:D:1151:A:O2'	18:D:1152:A:O5'	2.13	0.64
22:H:277:GLY:HA2	22:H:331:MET:CE	2.27	0.64
24:J:62:ARG:HG2	24:J:72:PHE:CZ	2.33	0.64
25:K:111:MET:HE1	25:K:125:ALA:HB1	1.80	0.64
33:S:77:PHE:HE1	33:S:93:ILE:HG21	1.63	0.64
33:S:87:ALA:HA	33:S:90:ARG:CZ	2.28	0.64
39:Y:24:GLY:O	39:Y:27:LEU:HG	1.96	0.64
41:a:998:C:OP2	66:z:92:ARG:NH2	2.29	0.64
41:a:1278:C:C4'	63:w:24:MET:HE1	2.27	0.64
16:AG:233:ARG:CB	16:AG:327:LEU:HD23	2.28	0.64
16:AG:434:LEU:HD12	16:AG:454:CYS:C	2.22	0.64
21:G:49:MET:HE1	21:G:201:PRO:CD	2.23	0.64
34:T:3:LEU:HD22	34:T:8:THR:CG2	2.27	0.64
39:Y:21:PRO:HB2	39:Y:22:PRO:HD3	1.80	0.64
39:Y:102:ARG:NE	39:Y:139:VAL:HG21	2.13	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1839:G:H1'	41:a:1927:A:C8	2.32	0.64
47:g:58:ASP:C	47:g:62:LYS:NZ	2.55	0.64
52:l:69:ARG:NH1	52:l:69:ARG:HA	2.13	0.64
8:7:9:U:N3	18:D:1196:A:N7	2.45	0.64
9:9:73:LYS:NZ	9:9:115:GLY:O	2.31	0.64
16:AG:131:ARG:HG2	16:AG:186:VAL:HG12	1.80	0.64
18:D:1226:C:H2'	38:X:102:THR:OG1	1.98	0.64
23:I:132:ARG:O	23:I:135:LYS:HG2	1.97	0.64
39:Y:9:LYS:HG2	39:Y:55:PRO:HB3	1.80	0.64
39:Y:86:LYS:HD3	39:Y:86:LYS:C	2.23	0.64
60:t:42:THR:HG23	60:t:44:LYS:NZ	2.12	0.64
62:v:31:PHE:N	62:v:104:GLU:OE2	2.30	0.64
9:9:47:GLU:HB3	9:9:51:TYR:OH	1.97	0.64
16:AG:63:LEU:HD22	16:AG:92:TYR:HD1	1.63	0.64
22:H:119:GLY:HA2	22:H:133:GLY:HA2	1.80	0.64
30:P:24:GLU:HA	30:P:27:GLU:OE1	1.98	0.64
39:Y:102:ARG:HG3	39:Y:141:ASP:HA	1.80	0.64
39:Y:126:ARG:NH2	41:a:1083:U:P	2.70	0.64
64:x:114:GLY:O	64:x:116:GLN:NE2	2.31	0.64
65:y:64:ILE:HG13	65:y:64:ILE:O	1.97	0.64
2:1:71:VAL:HB	2:1:74:ILE:HD11	1.79	0.63
7:6:25:DT:H2'	11:AA:496:LYS:CE	2.28	0.63
10:A:32:C:H4'	27:M:86:GLN:HE21	1.62	0.63
11:AA:314:ASN:HD21	11:AA:348:SER:HA	1.62	0.63
11:AA:855:PRO:CA	16:AG:35:THR:HB	2.28	0.63
16:AG:425:LEU:O	16:AG:429:LYS:CE	2.45	0.63
10:B:30:G:O2'	18:D:1339:A:H2	1.80	0.63
18:D:718:A:H5'	31:Q:119:ASN:ND2	2.13	0.63
36:V:31:HIS:CD2	36:V:34:TYR:H	2.15	0.63
39:Y:5:GLN:HG2	39:Y:61:TYR:CD1	2.32	0.63
39:Y:55:PRO:HB2	39:Y:71:LYS:HE2	1.80	0.63
9:9:50:VAL:HG23	39:Y:120:ASP:OD2	1.98	0.63
9:9:117:LEU:O	9:9:117:LEU:HD23	1.98	0.63
12:AB:38:ILE:HD12	12:AB:160:ARG:NH1	2.13	0.63
12:AB:136:GLU:OE1	12:AB:160:ARG:NH1	2.31	0.63
16:AG:354:ALA:CA	16:AG:357:ASP:H	2.10	0.63
16:AG:433:ASP:O	16:AG:456:LEU:CG	2.45	0.63
22:H:19:ARG:O	22:H:72:LEU:HB2	1.98	0.63
22:H:70:VAL:HG12	22:H:83:LEU:O	1.98	0.63
38:X:22:ILE:HB	38:X:25:VAL:HG12	1.80	0.63
39:Y:31:GLY:C	39:Y:60:VAL:HG11	2.23	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:995:C:O2	59:s:3:THR:OG1	2.17	0.63
41:a:2682:A:O4'	50:j:11:MET:CE	2.47	0.63
11:AA:855:PRO:CG	16:AG:105:ILE:CG2	2.76	0.63
14:AE:84:ILE:HG22	14:AE:91:GLU:HG3	1.79	0.63
16:AG:133:HIS:ND1	16:AG:136:GLU:CD	2.55	0.63
23:I:156:ARG:HH11	23:I:193:TYR:HB3	1.63	0.63
16:AG:226:ALA:O	16:AG:228:ARG:HD3	1.97	0.63
16:AG:349:GLN:C	16:AG:351:GLU:H	2.06	0.63
10:B:31:G:O3'	18:D:1340:A:O2'	2.17	0.63
38:X:16:VAL:HG12	38:X:34:LEU:HD12	1.80	0.63
39:Y:60:VAL:HG22	39:Y:66:PHE:CB	2.23	0.63
41:a:1277:G:N3	63:w:23:ASN:ND2	2.47	0.63
41:a:2311:A:N3	54:n:85:ILE:HD11	2.13	0.63
41:a:2393:U:P	55:o:30:ARG:NE	2.70	0.63
9:9:77:VAL:HG12	9:9:77:VAL:O	1.99	0.63
11:AA:120:GLN:NE2	11:AA:490:GLN:OE1	2.31	0.63
11:AA:387:ASN:HB3	11:AA:394:ARG:HH11	1.63	0.63
11:AA:859:GLU:C	16:AG:38:LYS:HZ3	2.06	0.63
12:AB:102:LYS:HE2	14:AE:285:LEU:CB	2.29	0.63
16:AG:205:LEU:HA	16:AG:208:LEU:HB2	1.80	0.63
16:AG:223:ILE:HA	16:AG:238:VAL:HG12	1.81	0.63
18:D:43:C:OP1	35:U:12:LYS:NZ	2.29	0.63
18:D:1228:C:OP1	38:X:107:ARG:NH1	2.32	0.63
24:J:99:ASP:OD1	24:J:100:ASN:N	2.31	0.63
39:Y:37:PHE:CE1	39:Y:56:VAL:HG11	2.33	0.63
48:h:123:ALA:HB1	48:h:125:LYS:HZ1	1.63	0.63
52:l:23:PHE:HB2	52:l:111:GLU:OE2	1.99	0.63
8:7:9:U:N3	18:D:1196:A:C8	2.66	0.63
11:AA:528:ARG:NH2	11:AA:576:SER:O	2.29	0.63
14:AE:79:LYS:NZ	16:AG:153:ASP:OD1	2.30	0.63
16:AG:323:GLN:HE21	16:AG:323:GLN:CA	2.12	0.63
16:AG:353:HIS:CA	16:AG:356:ILE:CG2	2.76	0.63
16:AG:424:SER:C	16:AG:429:LYS:CG	2.72	0.63
17:C:33:ILE:O	22:H:338:GLU:CD	2.42	0.63
24:J:150:LYS:HE2	24:J:178:MET:HE2	1.79	0.63
41:a:537:G:OP1	59:s:2:LYS:HE2	1.98	0.63
41:a:1067:A:O2'	41:a:1068:G:O4'	2.16	0.63
41:a:2013:A:N1	41:a:2613:U:O4	2.31	0.63
41:a:2682:A:O5'	50:j:11:MET:HE1	1.98	0.63
61:u:84:LYS:HZ2	61:u:98:ALA:C	2.04	0.63
63:w:33:ILE:CG1	63:w:112:TYR:HD1	2.11	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:84:ARG:NH1	41:a:813:U:OP1	2.31	0.63
16:AG:287:ALA:HB1	16:AG:331:LEU:HB3	1.81	0.63
18:D:406:G:H21	24:J:116:GLN:HE22	1.46	0.63
18:D:1152:A:OP1	30:P:70:HIS:ND1	2.29	0.63
33:S:13:ARG:HB3	33:S:60:GLN:HG3	1.81	0.63
34:T:88:ARG:NH2	41:a:714:U:OP2	2.31	0.63
39:Y:5:GLN:HE22	39:Y:60:VAL:N	1.96	0.63
5:4:50:MET:O	5:4:52:ALA:N	2.31	0.63
5:4:80:HIS:CE1	5:4:82:TYR:H	2.17	0.63
9:9:25:ALA:HB3	9:9:99:PHE:HD2	1.61	0.63
10:A:33:U:O3'	27:M:84:THR:OG1	2.16	0.63
16:AG:301:ASP:OD1	16:AG:301:ASP:N	2.32	0.63
18:D:932:C:H5'	27:M:4:ARG:HH11	1.63	0.63
21:G:114:LEU:HD13	21:G:144:LEU:CB	2.28	0.63
27:M:4:ARG:CG	27:M:5:ARG:CZ	2.75	0.63
29:O:123:ARG:HG2	29:O:123:ARG:O	1.98	0.63
32:R:75:GLN:OE1	32:R:75:GLN:N	2.32	0.63
35:U:18:GLN:OE1	35:U:37:GLY:O	2.17	0.63
65:y:31:TRP:CZ3	65:y:38:LYS:HD3	2.33	0.63
6:5:11:DA:P	12:AB:67:THR:C	2.79	0.63
8:7:9:U:O4	18:D:1196:A:N9	2.32	0.63
16:AG:354:ALA:HA	16:AG:357:ASP:CA	2.28	0.63
16:AG:422:GLU:HG2	16:AG:426:GLY:HA3	1.80	0.63
21:G:19:GLN:HG3	22:H:76:GLU:HB3	1.78	0.63
22:H:16:ILE:HG22	22:H:25:ARG:NH2	2.14	0.63
41:a:2335:A:P	64:x:13:ARG:CZ	2.87	0.63
49:i:17:ARG:HA	49:i:20:ASP:OD1	1.99	0.63
9:9:7:ASP:O	9:9:11:ILE:HD12	1.98	0.62
10:B:76:A:H1'	41:a:2494:G:OP1	1.98	0.62
17:C:13:PHE:CE2	17:C:21:ILE:HD11	2.34	0.62
39:Y:14:ALA:HB3	39:Y:51:GLY:N	2.12	0.62
5:4:77:VAL:HG11	5:4:79:ARG:HH21	1.64	0.62
8:7:11:U:O4'	25:K:20:ARG:NH2	2.32	0.62
16:AG:361:LYS:CE	16:AG:417:ILE:CG1	2.74	0.62
16:AG:380:THR:C	16:AG:384:LEU:HD11	2.24	0.62
21:G:49:MET:HE3	21:G:199:VAL:O	2.00	0.62
21:G:126:PHE:O	21:G:126:PHE:CD1	2.53	0.62
39:Y:9:LYS:HE2	39:Y:55:PRO:HG3	1.82	0.62
39:Y:100:ILE:CD1	39:Y:139:VAL:HB	2.29	0.62
41:a:754:U:H2'	41:a:755:U:C6	2.34	0.62
51:k:21:TYR:CE2	51:k:38:LYS:HG2	2.34	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:n:33:LYS:HA	54:n:96:MET:SD	2.39	0.62
63:w:35:LYS:HZ2	63:w:112:TYR:HE2	1.45	0.62
9:9:47:GLU:HA	9:9:94:ARG:HH22	1.65	0.62
10:A:54:U:H3	10:A:58:A:N6	1.96	0.62
16:AG:380:THR:O	16:AG:384:LEU:HD12	1.95	0.62
25:K:61:GLN:HA	25:K:64:MET:HE3	1.81	0.62
39:Y:102:ARG:HA	39:Y:102:ARG:HE	1.64	0.62
41:a:2725:A:C4	41:a:2727:A:C8	2.87	0.62
12:AB:141:LEU:O	12:AB:151:LYS:C	2.42	0.62
16:AG:389:MET:HE3	16:AG:407:ARG:HD2	1.81	0.62
16:AG:434:LEU:CD1	16:AG:454:CYS:C	2.72	0.62
18:D:13:U:O2	18:D:914:A:C8	2.52	0.62
21:G:73:LYS:O	21:G:74:ARG:HG2	2.00	0.62
23:I:14:ILE:HG22	23:I:15:VAL:HG13	1.82	0.62
27:M:17:LYS:HG3	27:M:18:PHE:CD1	2.35	0.62
41:a:1138:G:N3	59:s:108:MET:HE2	2.14	0.62
48:h:251:GLN:HE22	48:h:253:LYS:C	2.07	0.62
58:r:5:LEU:HD12	58:r:17:ASP:O	2.00	0.62
63:w:49:GLU:OE1	63:w:49:GLU:N	2.26	0.62
6:5:10:DT:C5	11:AA:62:TYR:HE2	1.97	0.62
9:9:52:MET:C	9:9:52:MET:HE3	2.25	0.62
12:AB:119:THR:HG23	12:AB:141:LEU:HD13	1.81	0.62
16:AG:288:MET:SD	16:AG:293:VAL:N	2.73	0.62
21:G:118:GLU:O	21:G:121:SER:OG	2.15	0.62
25:K:45:ARG:HB3	25:K:71:MET:HE2	1.81	0.62
37:W:12:ASP:OD2	37:W:35:SER:OG	2.14	0.62
63:w:37:THR:OG1	63:w:40:LYS:HD2	1.99	0.62
3:2:56:GLU:OE2	3:2:87:LEU:C	2.42	0.62
6:5:9:DG:P	11:AA:473:ARG:CG	2.88	0.62
11:AA:861:ALA:N	16:AG:38:LYS:HZ3	1.82	0.62
16:AG:428:ASN:O	16:AG:428:ASN:ND2	2.32	0.62
16:AG:433:ASP:C	16:AG:456:LEU:CD1	2.72	0.62
16:AG:453:VAL:HG21	16:AG:462:GLN:CD	2.25	0.62
21:G:30:PHE:CD1	21:G:201:PRO:CG	2.83	0.62
62:v:88:ASN:OD1	62:v:89:VAL:N	2.32	0.62
5:4:77:VAL:HG22	5:4:79:ARG:HE	1.64	0.62
9:9:23:LEU:HD11	9:9:96:PHE:CE2	2.34	0.62
22:H:61:GLU:HG2	22:H:62:ILE:HG23	1.81	0.62
39:Y:80:LYS:HG3	39:Y:86:LYS:O	2.00	0.62
60:t:8:LEU:N	60:t:18:ARG:HH12	1.98	0.62
1:0:82:HIS:CG	1:0:82:HIS:O	2.52	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:5:11:DA:H2'	12:AB:67:THR:HG21	1.82	0.62
16:AG:422:GLU:CA	16:AG:426:GLY:CA	2.65	0.62
22:H:135:LEU:CB	22:H:167:VAL:CB	2.78	0.62
25:K:113:ALA:HA	25:K:116:GLU:OE1	2.00	0.62
27:M:50:LEU:HD21	27:M:121:ALA:CA	2.28	0.62
63:w:42:LYS:HE2	63:w:97:ILE:HG21	1.82	0.62
3:2:56:GLU:OE2	3:2:87:LEU:O	2.18	0.62
16:AG:266:LEU:H	16:AG:266:LEU:HD23	1.64	0.62
16:AG:425:LEU:O	16:AG:429:LYS:NZ	2.33	0.62
33:S:87:ALA:HA	33:S:90:ARG:NH1	2.15	0.62
37:W:66:MET:SD	37:W:66:MET:O	2.57	0.62
41:a:2307:G:O4'	41:a:2308:G:C5	2.52	0.62
62:v:38:ARG:HA	62:v:96:ILE:O	2.00	0.62
11:AA:1282:GLY:HA3	15:AF:17:PHE:HE1	1.63	0.62
16:AG:27:LEU:HD22	16:AG:114:ILE:HG23	1.82	0.62
17:C:45:THR:HA	22:H:340:ARG:NH2	2.15	0.62
18:D:1309:G:P	38:X:98:ARG:HE	2.20	0.62
25:K:69:ARG:HA	25:K:69:ARG:NE	2.15	0.62
27:M:66:LEU:HD13	27:M:101:MET:HE1	1.81	0.62
33:S:19:LYS:HG3	33:S:20:TYR:CE1	2.35	0.62
38:X:28:THR:CG2	38:X:31:LYS:HE3	2.30	0.62
39:Y:79:LEU:CD1	39:Y:132:ALA:HA	2.29	0.62
59:s:37:ARG:NH2	59:s:44:TYR:OH	2.32	0.62
62:v:1:MET:HE1	62:v:44:ARG:CA	2.30	0.62
11:AA:860:ALA:N	16:AG:38:LYS:NZ	2.48	0.61
12:AB:154:VAL:CG2	12:AB:159:PHE:N	2.62	0.61
16:AG:437:LEU:CD2	16:AG:456:LEU:HD11	2.27	0.61
21:G:38:VAL:CG2	22:H:75:VAL:HG21	2.30	0.61
25:K:140:THR:O	25:K:144:LEU:HD23	1.99	0.61
39:Y:99:LYS:NZ	39:Y:138:VAL:HG13	2.14	0.61
41:a:985:C:C2	41:a:986:C:C5	2.88	0.61
41:a:2315:G:H4'	54:n:127:ASN:ND2	2.14	0.61
55:o:32:ILE:HG13	55:o:32:ILE:O	2.00	0.61
58:r:114:GLU:OE2	58:r:134:VAL:N	2.33	0.61
2:1:98:LYS:NZ	41:a:2011:U:O3'	2.34	0.61
6:5:10:DT:C4	11:AA:62:TYR:HD2	2.12	0.61
16:AG:127:VAL:CG2	16:AG:192:GLY:CA	2.78	0.61
10:B:36:U:H2'	10:B:37:A:C8	2.35	0.61
44:d:5:U:OP1	44:d:61:G:O2'	2.16	0.61
54:n:38:MET:SD	54:n:150:ARG:HD2	2.39	0.61
10:A:65:C:H2'	10:A:66:C:H5'	1.83	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AB:80:ARG:HB3	12:AB:82:GLY:O	1.99	0.61
16:AG:285:ILE:CG2	16:AG:293:VAL:HG11	2.28	0.61
18:D:1124:G:O2'	18:D:1145:A:N6	2.33	0.61
18:D:1302:C:OP1	38:X:13:LYS:NZ	2.30	0.61
25:K:86:LYS:HZ2	25:K:95:PHE:HE1	1.49	0.61
27:M:18:PHE:HD2	27:M:59:LEU:HD21	1.65	0.61
28:N:24:ALA:HB1	28:N:60:GLU:OE2	2.00	0.61
41:a:1824:G:C3'	48:h:52:ARG:HH12	2.13	0.61
44:d:55:U:H1'	54:n:26:MET:CE	2.26	0.61
55:o:13:ARG:HD3	61:u:58:TYR:O	2.00	0.61
9:9:113:PHE:O	9:9:123:ILE:HG13	1.99	0.61
12:AB:118:ILE:HD12	12:AB:141:LEU:HD12	1.80	0.61
16:AG:451:ARG:CZ	16:AG:469:ASP:HB2	2.31	0.61
18:D:61:G:N7	19:E:6:SER:OG	2.33	0.61
25:K:81:LEU:HG	25:K:96:MET:HE1	1.82	0.61
31:Q:84:VAL:HG23	31:Q:107:ILE:CD1	2.30	0.61
38:X:55:THR:O	38:X:59:GLU:OE1	2.18	0.61
62:v:75:GLU:HG3	62:v:90:GLU:OE1	2.01	0.61
6:5:10:DT:C5'	11:AA:62:TYR:OH	2.48	0.61
12:AB:82:GLY:HA2	14:AE:142:GLU:OE1	2.01	0.61
12:AB:102:LYS:CD	14:AE:285:LEU:HA	2.31	0.61
10:B:30:G:O2'	18:D:1339:A:C2	2.52	0.61
27:M:111:ARG:O	27:M:119:ARG:NH1	2.34	0.61
40:Z:8:ILE:HG13	40:Z:9:GLU:N	2.15	0.61
63:w:47:VAL:O	63:w:51:LEU:HD23	2.01	0.61
9:9:23:LEU:HA	9:9:118:ILE:CG1	2.31	0.61
12:AB:140:MET:HA	12:AB:153:SER:HA	1.83	0.61
16:AG:353:HIS:HA	16:AG:356:ILE:HG23	1.78	0.61
16:AG:362:TYR:CA	16:AG:413:ALA:HB1	2.30	0.61
16:AG:393:LEU:CB	16:AG:398:LEU:HD22	1.90	0.61
16:AG:402:THR:O	16:AG:406:LEU:N	2.33	0.61
16:AG:440:VAL:HG23	16:AG:481:LEU:CD2	2.29	0.61
18:D:337:G:H2'	18:D:338:A:C8	2.34	0.61
41:a:413:C:H2'	41:a:414:C:H6	1.66	0.61
41:a:1406:U:C2	41:a:1407:G:C8	2.88	0.61
48:h:141:VAL:HG23	48:h:162:VAL:CG1	2.30	0.61
60:t:20:MET:CB	60:t:44:LYS:HZ3	2.10	0.61
8:7:7:U:H6	8:7:7:U:H5''	1.64	0.61
11:AA:889:PRO:CG	16:AG:104:ARG:NH1	2.62	0.61
16:AG:433:ASP:HB3	16:AG:456:LEU:HD12	1.83	0.61
18:D:687:A:C2	18:D:704:A:C5	2.88	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:N:43:GLU:CD	28:N:101:ILE:HG21	2.25	0.61
51:k:34:LEU:HB3	51:k:51:GLU:HG3	1.82	0.61
11:AA:888:THR:N	16:AG:105:ILE:CD1	2.62	0.61
12:AB:35:LEU:HB3	12:AB:37:LYS:HD3	1.82	0.61
12:AB:140:MET:HE1	30:P:103:GLY:N	2.16	0.61
12:AB:142:LEU:HA	12:AB:151:LYS:HA	1.83	0.61
14:AE:888:CYS:SG	14:AE:889:ASP:N	2.74	0.61
14:AE:1161:GLY:HA3	14:AE:1179:PRO:HA	1.83	0.61
16:AG:285:ILE:HD13	16:AG:285:ILE:N	2.15	0.61
10:B:65:C:H2'	10:B:66:C:H5'	1.83	0.61
18:D:1145:A:O2'	18:D:1146:A:O5'	2.18	0.61
18:D:1160:G:C2	18:D:1161:C:C6	2.89	0.61
28:N:88:ARG:HB2	28:N:91:GLU:OE1	2.00	0.61
39:Y:79:LEU:CD2	39:Y:105:LEU:HD21	2.27	0.61
41:a:548:G:H3'	41:a:549:G:O4'	2.01	0.61
51:k:34:LEU:HB3	51:k:51:GLU:CG	2.31	0.61
55:o:30:ARG:HH12	61:u:62:PRO:CA	2.14	0.61
8:7:11:U:O2'	25:K:33:PHE:CZ	2.52	0.61
16:AG:424:SER:C	16:AG:429:LYS:HG3	2.26	0.61
3:2:1:MET:N	41:a:142:A:O2'	2.34	0.61
6:5:9:DG:O5'	11:AA:473:ARG:CB	2.49	0.61
11:AA:472:GLU:HG3	11:AA:473:ARG:HD3	1.82	0.61
14:AE:1275:LEU:HB3	14:AE:1278:GLU:HB2	1.83	0.61
18:D:563:A:C2	18:D:567:G:C5	2.89	0.61
18:D:1157:A:C2	18:D:1181:G:C4	2.88	0.61
22:H:135:LEU:O	22:H:169:SER:HA	2.00	0.61
23:I:45:LYS:NZ	23:I:46:GLU:OE1	2.29	0.61
30:P:23:ALA:O	30:P:27:GLU:OE1	2.18	0.61
48:h:164:ILE:HG23	48:h:164:ILE:O	2.01	0.61
59:s:98:GLU:O	59:s:102:GLU:OE1	2.18	0.61
61:u:76:GLU:HG3	61:u:111:ILE:CD1	2.31	0.61
62:v:111:GLU:OE1	62:v:111:GLU:N	2.32	0.61
3:2:56:GLU:OE1	3:2:86:THR:O	2.19	0.60
9:9:23:LEU:HD21	9:9:96:PHE:CG	2.36	0.60
9:9:25:ALA:HB2	9:9:96:PHE:CD1	2.36	0.60
9:9:98:GLU:HA	9:9:101:LYS:HG2	1.83	0.60
17:C:29:LEU:HD23	17:C:59:ILE:HD13	1.83	0.60
23:I:58:GLU:OE2	30:P:94:ALA:CB	2.48	0.60
25:K:16:ILE:HD11	25:K:38:VAL:HG12	1.83	0.60
41:a:2191:A:H2'	41:a:2192:U:C6	2.36	0.60
50:j:1:MET:HE2	50:j:100:LEU:HD11	1.83	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:s:11:VAL:HG11	59:s:13:ARG:NH1	2.15	0.60
62:v:111:GLU:HA	62:v:114:ARG:HE	1.66	0.60
11:AA:855:PRO:CD	16:AG:105:ILE:CG2	2.79	0.60
14:AE:785:ASP:O	14:AE:789:LYS:HB2	2.01	0.60
16:AG:253:GLY:HA3	16:AG:258:ARG:HD2	1.82	0.60
18:D:718:A:O2'	20:F:34:ARG:NH2	2.34	0.60
22:H:155:LYS:O	22:H:169:SER:N	2.34	0.60
28:N:104:VAL:HG23	28:N:125:ILE:HD13	1.83	0.60
29:O:94:LEU:HB3	29:O:97:GLU:OE2	2.01	0.60
41:a:1839:G:C8	41:a:1927:A:H1'	2.35	0.60
47:g:1:MET:HE2	47:g:6:HIS:CD2	2.35	0.60
64:x:90:VAL:HG21	64:x:115:LEU:HD21	1.81	0.60
12:AB:152:HIS:CD2	12:AB:159:PHE:HE2	1.91	0.60
16:AG:63:LEU:HG	16:AG:73:LYS:HB3	1.83	0.60
17:C:73:ARG:CZ	31:Q:113:VAL:HG12	2.31	0.60
18:D:562:U:C2'	32:R:14:ARG:HH11	2.13	0.60
29:O:57:MET:HA	29:O:60:LYS:HE2	1.83	0.60
30:P:26:VAL:HG12	30:P:30:LYS:NZ	2.16	0.60
34:T:67:LEU:HD12	34:T:78:TYR:CE1	2.35	0.60
41:a:1070:A:N7	41:a:1096:A:O2'	2.35	0.60
54:n:122:PHE:CZ	54:n:170:LEU:HD12	2.36	0.60
66:z:74:ILE:HD11	66:z:79:PHE:CA	2.30	0.60
1:O:51:VAL:O	1:O:51:VAL:HG13	2.02	0.60
9:9:31:ARG:CD	41:a:1053:C:O2'	2.47	0.60
9:9:43:LYS:HZ3	9:9:98:GLU:HB2	1.65	0.60
9:9:123:ILE:O	9:9:123:ILE:HG23	2.02	0.60
10:A:5:G:H2'	10:A:6:G:H5'	1.82	0.60
10:A:36:U:H2'	10:A:37:A:C8	2.36	0.60
10:A:69:C:H2'	10:A:70:G:O4'	2.02	0.60
14:AE:814:CYS:SG	14:AE:883:ARG:NH2	2.74	0.60
27:M:137:LYS:O	27:M:141:VAL:HG23	2.01	0.60
36:V:61:ILE:HG13	36:V:73:TRP:CZ3	2.36	0.60
41:a:639:U:H2'	41:a:640:C:C6	2.36	0.60
41:a:1322:A:C5	41:a:1323:C:C5	2.90	0.60
54:n:122:PHE:O	54:n:123:ASP:OD1	2.19	0.60
59:s:74:TYR:CD2	59:s:74:TYR:N	2.69	0.60
61:u:76:GLU:HG3	61:u:111:ILE:HD13	1.84	0.60
9:9:61:ARG:NE	41:a:1046:A:O5'	2.34	0.60
11:AA:914:LYS:N	16:AG:105:ILE:CA	2.58	0.60
12:AB:138:ARG:CG	12:AB:154:VAL:C	2.47	0.60
14:AE:79:LYS:HG2	16:AG:144:LYS:HD2	1.84	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:279:ASN:HB3	16:AG:280:PRO:HD2	1.83	0.60
10:B:69:C:H2'	10:B:70:G:O4'	2.02	0.60
21:G:19:GLN:OE1	21:G:21:ARG:HG2	2.00	0.60
25:K:112:ARG:O	25:K:116:GLU:OE1	2.19	0.60
37:W:45:ILE:HA	37:W:62:VAL:CG1	2.31	0.60
41:a:1277:G:H5'	63:w:20:MET:HE2	1.84	0.60
48:h:84:ASP:HB2	48:h:91:ILE:HD12	1.84	0.60
1:O:73:LYS:NZ	41:a:1225:G:OP1	2.34	0.60
7:6:18:DT:N3	8:7:49:A:N6	2.36	0.60
8:7:53:G:OP1	11:AA:1073:LYS:NZ	2.24	0.60
9:9:14:GLU:OE1	9:9:62:ARG:O	2.19	0.60
9:9:33:VAL:HG22	9:9:36:ASP:CG	2.27	0.60
12:AB:92:VAL:HA	12:AB:96:LEU:HB2	1.83	0.60
13:AD:211:ILE:HG12	13:AD:219:ARG:HH12	1.65	0.60
14:AE:86:GLU:OE1	14:AE:86:GLU:N	2.34	0.60
17:C:14:THR:HG21	17:C:48:ARG:NE	2.17	0.60
17:C:60:LYS:NZ	18:D:734:G:C2'	2.64	0.60
18:D:652:U:H4'	28:N:56:LYS:NZ	2.16	0.60
18:D:1125:U:C2	18:D:1127:G:C8	2.89	0.60
35:U:5:ARG:HH22	35:U:24:SER:HA	1.66	0.60
37:W:67:VAL:HG21	47:g:57:VAL:HG22	1.82	0.60
41:a:1257:C:H4'	52:l:78:TRP:CD2	2.37	0.60
41:a:1817:G:H3'	48:h:156:ARG:HH22	1.67	0.60
47:g:65:ASN:OD1	47:g:66:ILE:N	2.30	0.60
63:w:60:VAL:HA	63:w:63:ARG:NH1	2.17	0.60
2:1:1:MET:SD	2:1:3:THR:OG1	2.59	0.60
12:AB:115:LYS:HD2	12:AB:129:ILE:HG21	1.82	0.60
12:AB:129:ILE:O	12:AB:131:THR:N	2.35	0.60
14:AE:506:VAL:HG23	14:AE:628:GLY:HA3	1.83	0.60
16:AG:168:LEU:HD21	16:AG:230:PRO:O	2.00	0.60
16:AG:434:LEU:CD2	16:AG:459:LEU:CD1	2.54	0.60
31:Q:46:THR:OG1	31:Q:49:GLY:N	2.34	0.60
39:Y:36:GLU:HG3	39:Y:66:PHE:CE1	2.36	0.60
39:Y:72:THR:OG1	39:Y:73:PRO:HD2	2.01	0.60
41:a:1406:U:O2'	41:a:1407:G:O4'	2.19	0.60
60:t:42:THR:HG23	60:t:44:LYS:HZ1	1.65	0.60
11:AA:55:SER:OG	11:AA:465:ARG:NH1	2.33	0.60
16:AG:212:GLU:HB3	16:AG:258:ARG:HB3	1.84	0.60
16:AG:422:GLU:CA	16:AG:426:GLY:H	2.08	0.60
16:AG:434:LEU:CD1	16:AG:455:THR:C	2.74	0.60
18:D:1006:G:H2'	18:D:1007:U:C6	2.36	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:F:49:LYS:O	20:F:53:VAL:HG23	2.02	0.60
28:N:43:GLU:HG2	28:N:101:ILE:HD13	1.84	0.60
39:Y:5:GLN:HE22	39:Y:60:VAL:C	2.10	0.60
8:7:9:U:O4	18:D:1196:A:C8	2.55	0.60
16:AG:320:ARG:CG	16:AG:320:ARG:HH11	2.14	0.60
16:AG:421:GLN:O	16:AG:425:LEU:N	2.34	0.60
18:D:878:A:OP2	28:N:80:ARG:NH1	2.35	0.60
21:G:110:SER:HA	21:G:113:ARG:HE	1.65	0.60
27:M:103:TRP:O	27:M:106:GLU:HG3	2.01	0.60
39:Y:5:GLN:CG	39:Y:61:TYR:CE1	2.76	0.60
39:Y:19:PRO:O	39:Y:24:GLY:CA	2.48	0.60
41:a:271:G:C4	41:a:272:A:C8	2.89	0.60
47:g:5:ILE:HD12	54:n:64:LYS:CE	2.31	0.60
60:t:8:LEU:CA	60:t:18:ARG:HH12	2.15	0.60
61:u:51:GLU:OE1	61:u:51:GLU:N	2.34	0.60
9:9:47:GLU:HG3	9:9:94:ARG:NH1	2.16	0.60
9:9:97:LYS:NZ	9:9:130:PRO:HA	2.17	0.60
12:AB:89:PRO:CG	14:AE:290:ILE:CG2	2.79	0.60
14:AE:92:VAL:HG22	14:AE:92:VAL:O	2.01	0.60
41:a:567:U:OP2	61:u:29:LYS:NZ	2.35	0.60
41:a:1065:U:O2'	41:a:1066:U:O5'	2.16	0.60
59:s:13:ARG:HH22	59:s:50:THR:CA	2.14	0.60
7:6:23:DC:H5''	14:AE:259:ARG:NH1	1.91	0.59
12:AB:115:LYS:CA	12:AB:129:ILE:CD1	2.79	0.59
16:AG:139:THR:HG22	16:AG:180:ARG:HB3	1.83	0.59
16:AG:303:HIS:HB3	16:AG:334:TRP:HZ3	1.66	0.59
17:C:12:ARG:CG	22:H:264:GLU:CB	2.80	0.59
18:D:13:U:N3	18:D:915:A:N6	2.50	0.59
18:D:890:G:O2'	18:D:906:A:N6	2.35	0.59
18:D:1169:A:H2'	18:D:1170:A:C8	2.37	0.59
21:G:35:ARG:HD2	22:H:14:LYS:HE3	1.84	0.59
22:H:81:GLU:O	22:H:82:THR:HG23	2.02	0.59
23:I:142:MET:HE2	23:I:149:ILE:HG22	1.83	0.59
41:a:2189:U:P	41:a:2189:U:O4'	2.60	0.59
41:a:2363:G:OP2	55:o:40:ARG:HD2	2.01	0.59
46:f:10:THR:HG22	46:f:54:MET:C	2.27	0.59
63:w:33:ILE:CG1	63:w:112:TYR:CD1	2.85	0.59
3:2:56:GLU:OE1	3:2:56:GLU:N	2.33	0.59
5:4:63:ILE:HD11	5:4:91:PHE:CE2	2.37	0.59
9:9:51:TYR:HB2	9:9:88:HIS:CB	2.32	0.59
11:AA:371:ARG:NH1	12:AB:67:THR:OG1	2.34	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:B:5:G:H2'	10:B:6:G:H5'	1.82	0.59
22:H:131:LEU:CB	22:H:167:VAL:HA	2.32	0.59
34:T:64:ARG:CZ	34:T:64:ARG:HB3	2.32	0.59
39:Y:99:LYS:HD3	39:Y:138:VAL:O	2.02	0.59
40:Z:8:ILE:HG13	40:Z:9:GLU:H	1.68	0.59
41:a:856:G:H2'	41:a:857:G:C8	2.37	0.59
44:d:42:C:P	54:n:64:LYS:HZ1	2.25	0.59
50:j:25:THR:HG21	50:j:193:VAL:HG22	1.84	0.59
54:n:25:VAL:O	54:n:28:VAL:HG12	2.02	0.59
6:5:10:DT:N1	11:AA:62:TYR:CE2	2.65	0.59
11:AA:1072:ASN:ND2	11:AA:1111:GLN:OE1	2.34	0.59
16:AG:353:HIS:O	16:AG:356:ILE:C	2.43	0.59
16:AG:434:LEU:HG	16:AG:456:LEU:N	2.16	0.59
17:C:73:ARG:HH12	31:Q:113:VAL:CA	2.16	0.59
22:H:333:LEU:O	22:H:334:ASP:OD1	2.19	0.59
23:I:24:ALA:C	23:I:29:PHE:HE2	2.11	0.59
23:I:54:ARG:NH2	23:I:56:VAL:HG21	2.18	0.59
25:K:10:GLU:OE1	25:K:10:GLU:N	2.35	0.59
29:O:56:ASP:O	29:O:57:MET:HE2	2.02	0.59
41:a:1028:A:H2'	41:a:1029:A:C8	2.37	0.59
41:a:2460:U:C2	41:a:2461:A:C8	2.91	0.59
41:a:2682:A:H8	50:j:11:MET:HE1	1.65	0.59
48:h:67:PHE:CE1	48:h:156:ARG:HD2	2.37	0.59
59:s:102:GLU:HB3	59:s:119:PHE:HZ	1.66	0.59
4:3:7:ARG:HH22	41:a:98:G:N2	2.00	0.59
9:9:103:ASN:HA	9:9:107:GLU:OE2	2.02	0.59
9:9:136:ILE:CD1	9:9:139:LEU:HD12	2.25	0.59
16:AG:10:GLU:HA	16:AG:12:VAL:HG22	1.84	0.59
16:AG:354:ALA:HA	16:AG:357:ASP:N	2.18	0.59
16:AG:363:LEU:HD21	16:AG:410:ALA:N	1.79	0.59
18:D:692:U:N3	18:D:695:A:OP2	2.34	0.59
18:D:1290:G:H4'	27:M:35:LYS:NZ	2.17	0.59
21:G:106:THR:HA	21:G:109:GLN:NE2	2.18	0.59
28:N:47:GLU:OE2	28:N:64:LYS:HA	2.02	0.59
29:O:55:VAL:O	29:O:57:MET:N	2.34	0.59
36:V:10:GLY:HA3	36:V:25:ILE:HD13	1.83	0.59
38:X:81:MET:HE1	38:X:92:ARG:HD3	1.83	0.59
41:a:323:C:H2'	52:l:163:ASN:OD1	2.02	0.59
41:a:575:A:C2	41:a:576:U:C5	2.90	0.59
41:a:753:A:C2	41:a:754:U:C6	2.89	0.59
41:a:2225:A:H4'	41:a:2226:C:O5'	2.03	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:g:58:ASP:C	47:g:62:LYS:HZ3	2.11	0.59
59:s:13:ARG:NH2	59:s:49:ASP:O	2.34	0.59
59:s:55:ILE:HG13	59:s:123:LYS:NZ	2.17	0.59
6:5:11:DA:H2'	12:AB:67:THR:CG2	2.32	0.59
8:7:10:U:H5''	8:7:10:U:O2	2.03	0.59
11:AA:1142:ARG:NH1	11:AA:1161:LEU:O	2.36	0.59
16:AG:434:LEU:HD22	16:AG:459:LEU:CG	2.31	0.59
16:AG:434:LEU:N	16:AG:456:LEU:HG	2.17	0.59
17:C:51:TYR:HD1	17:C:54:GLN:NE2	2.00	0.59
18:D:739:C:HO2'	34:T:42:HIS:CE1	2.15	0.59
26:L:42:TRP:CZ2	26:L:102:MET:HG3	2.36	0.59
41:a:2298:A:C4	41:a:2321:U:C5	2.90	0.59
41:a:2780:G:P	59:s:120:ARG:NH1	2.74	0.59
49:i:22:LEU:HG	49:i:23:THR:N	2.17	0.59
1:0:68:ARG:HH12	41:a:1223:G:P	2.25	0.59
9:9:73:LYS:CG	9:9:117:LEU:HD12	2.32	0.59
9:9:78:GLY:N	9:9:79:PRO:CD	2.66	0.59
11:AA:846:GLY:O	16:AG:104:ARG:NH1	2.13	0.59
12:AB:141:LEU:O	12:AB:150:ILE:C	2.45	0.59
14:AE:64:PRO:HG3	14:AE:90:VAL:HG12	1.83	0.59
16:AG:437:LEU:HD22	16:AG:488:ILE:CD1	2.33	0.59
28:N:106:THR:HB	28:N:121:LEU:HD13	1.82	0.59
34:T:60:VAL:O	34:T:64:ARG:HG2	2.03	0.59
39:Y:23:VAL:HG23	39:Y:24:GLY:N	2.17	0.59
41:a:993:G:OP2	66:z:51:ARG:NH2	2.36	0.59
51:k:19:HIS:ND1	51:k:41:PRO:CD	2.65	0.59
56:p:85:LYS:NZ	56:p:164:TYR:HD1	2.01	0.59
59:s:73:VAL:HG13	59:s:86:GLN:HG2	1.84	0.59
60:t:5:GLN:HA	60:t:20:MET:CE	2.32	0.59
61:u:61:LEU:HB3	61:u:62:PRO:HD2	1.85	0.59
5:4:76:ASP:OD1	5:4:77:VAL:N	2.32	0.59
7:6:25:DT:OP1	11:AA:496:LYS:C	2.43	0.59
11:AA:69:GLN:HE21	11:AA:101:ARG:HD2	1.67	0.59
14:AE:1221:LEU:HD22	14:AE:1306:LEU:HB2	1.84	0.59
18:D:1107:C:P	23:I:172:ARG:HG3	2.43	0.59
19:E:28:MET:SD	19:E:57:ILE:HG22	2.42	0.59
21:G:96:TRP:CH2	21:G:100:MET:HE3	2.38	0.59
25:K:12:GLN:HB3	25:K:14:LYS:NZ	2.18	0.59
41:a:468:G:N7	53:m:39:ARG:NH2	2.42	0.59
41:a:1172:C:H2'	41:a:1173:U:O4'	2.02	0.59
41:a:2547:A:H2'	41:a:2548:U:C6	2.38	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:f:40:ASP:C	46:f:45:ARG:NH2	2.60	0.59
47:g:18:CYS:HB3	47:g:40:CYS:HB2	1.85	0.59
3:2:56:GLU:OE1	3:2:87:LEU:C	2.46	0.59
9:9:94:ARG:CB	9:9:97:LYS:HE2	2.32	0.59
13:AD:112:ALA:HB3	13:AD:126:PRO:HA	1.83	0.59
16:AG:127:VAL:HG21	16:AG:192:GLY:HA2	1.84	0.59
10:B:75:C:P	41:a:2602:A:HO2'	2.15	0.59
25:K:159:LYS:NZ	28:N:44:GLY:HA3	2.16	0.59
27:M:66:LEU:HD11	27:M:100:ALA:HB3	1.85	0.59
28:N:8:ALA:O	28:N:12:THR:HG23	2.03	0.59
33:S:20:TYR:HD2	33:S:55:SER:OG	1.85	0.59
36:V:60:GLU:OE1	36:V:76:VAL:HG23	2.01	0.59
38:X:25:VAL:HG23	38:X:29:ARG:CB	2.32	0.59
38:X:93:ARG:HG2	38:X:95:LEU:HD23	1.84	0.59
39:Y:70:THR:O	39:Y:71:LYS:HG3	2.03	0.59
41:a:947:A:H2'	41:a:948:C:C6	2.37	0.59
48:h:62:TYR:CE2	48:h:64:ILE:HD13	2.38	0.59
58:r:135:HIS:ND1	58:r:137:GLU:CD	2.44	0.59
1:0:61:ALA:HB1	1:0:96:VAL:HG22	1.84	0.59
7:6:20:DA:C6	8:7:48:G:C2	2.91	0.59
16:AG:197:VAL:HG11	16:AG:199:ARG:HH21	1.66	0.59
18:D:978:A:C5	18:D:1319:A:C2	2.90	0.59
21:G:114:LEU:HD13	21:G:144:LEU:HB2	1.84	0.59
51:k:7:GLU:CD	51:k:7:GLU:H	2.11	0.59
54:n:4:LEU:HD11	54:n:101:GLU:HA	1.84	0.59
54:n:114:PHE:O	54:n:115:ARG:NE	2.34	0.59
1:0:16:GLU:C	1:0:18:GLN:OE1	2.46	0.59
12:AB:19:GLU:O	12:AB:23:ARG:HB2	2.03	0.59
12:AB:90:SER:HB3	12:AB:93:ILE:HB	1.84	0.59
12:AB:154:VAL:HG11	12:AB:159:PHE:HB2	1.82	0.59
18:D:588:G:H4'	28:N:3:MET:CE	2.33	0.59
18:D:1055:A:C2'	23:I:156:ARG:HH21	2.15	0.59
19:E:32:ILE:HG12	19:E:75:HIS:CE1	2.37	0.59
27:M:17:LYS:HZ2	27:M:18:PHE:HE1	1.50	0.59
30:P:22:THR:O	30:P:26:VAL:HG23	2.03	0.59
39:Y:102:ARG:CG	39:Y:141:ASP:HA	2.33	0.59
41:a:15:G:O2'	49:i:15:MET:CE	2.51	0.59
41:a:2002:G:OP1	63:w:17:ARG:NH2	2.26	0.59
48:h:30:PHE:O	48:h:34:LEU:HD23	2.02	0.59
54:n:38:MET:HG3	54:n:57:LEU:HD21	1.84	0.59
61:u:91:ASP:O	61:u:95:LEU:HD23	2.03	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
66:z:74:ILE:HD13	66:z:79:PHE:CD1	2.34	0.59
2:1:86:MET:CE	2:1:88:ARG:HD3	2.32	0.58
6:5:10:DT:O3'	12:AB:68:THR:CA	2.51	0.58
12:AB:120:GLU:HB2	12:AB:162:LEU:CB	2.32	0.58
16:AG:287:ALA:CB	16:AG:331:LEU:HB3	2.33	0.58
18:D:439:U:O2	18:D:440:C:C5	2.56	0.58
18:D:736:C:O2'	26:L:88:MET:HE1	2.02	0.58
18:D:829:G:O3'	21:G:23:TRP:CZ2	2.56	0.58
21:G:23:TRP:HZ3	21:G:25:PRO:HA	1.67	0.58
22:H:273:ARG:CZ	22:H:297:GLU:HG3	2.33	0.58
24:J:91:LEU:HD21	24:J:195:ILE:HD12	1.84	0.58
29:O:49:ARG:HB3	29:O:53:GLU:OE2	2.02	0.58
35:U:19:VAL:HG12	35:U:37:GLY:C	2.28	0.58
38:X:22:ILE:HG23	38:X:66:GLU:OE2	2.03	0.58
41:a:70:G:H4'	41:a:71:A:OP1	2.02	0.58
53:m:1:MET:HE3	53:m:3:ARG:HH11	1.67	0.58
62:v:97:GLN:H	62:v:100:LYS:HZ1	1.51	0.58
66:z:82:GLY:CA	66:z:117:LEU:CD1	2.81	0.58
5:4:60:VAL:HG12	5:4:71:LYS:HE2	1.86	0.58
16:AG:202:PRO:O	16:AG:205:LEU:HD12	2.02	0.58
18:D:736:C:H5''	26:L:90:MET:HE2	1.85	0.58
21:G:26:LYS:CD	21:G:193:PRO:HG2	2.32	0.58
26:L:9:MET:HE2	26:L:51:ILE:HD11	1.83	0.58
28:N:7:ILE:HB	28:N:77:ARG:NH1	2.18	0.58
33:S:42:TRP:CH2	47:g:66:ILE:HA	2.38	0.58
36:V:11:ARG:HE	36:V:58:VAL:CG2	2.16	0.58
41:a:1432:G:H2'	41:a:1433:A:C8	2.37	0.58
41:a:1798:U:OP2	48:h:271:ARG:NH2	2.36	0.58
41:a:2259:U:N3	41:a:2260:C:C5	2.71	0.58
59:s:102:GLU:CD	59:s:124:VAL:HG11	2.28	0.58
6:5:11:DA:C2'	12:AB:67:THR:CG2	2.77	0.58
9:9:17:GLU:HG2	9:9:88:HIS:NE2	2.18	0.58
9:9:60:LEU:HA	9:9:64:VAL:HB	1.84	0.58
12:AB:77:HIS:NE2	14:AE:297:ARG:NH2	2.51	0.58
12:AB:99:TYR:O	12:AB:100:LYS:HG2	2.03	0.58
16:AG:25:GLU:HG3	16:AG:49:ILE:HD11	1.86	0.58
16:AG:221:ILE:HD13	16:AG:247:PRO:HA	1.85	0.58
16:AG:240:THR:CB	16:AG:247:PRO:HG3	2.32	0.58
16:AG:453:VAL:CG2	16:AG:462:GLN:CD	2.76	0.58
10:B:15:G:N1	10:B:20:U:O2	2.37	0.58
18:D:110:C:O2'	35:U:25:ARG:O	2.21	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:718:A:C4	31:Q:118:HIS:ND1	2.71	0.58
22:H:273:ARG:HD3	22:H:280:LEU:HD11	1.85	0.58
24:J:62:ARG:HG2	24:J:72:PHE:CE1	2.37	0.58
27:M:50:LEU:HD11	27:M:124:LEU:HB2	1.84	0.58
30:P:25:ILE:O	30:P:28:THR:OG1	2.19	0.58
32:R:7:LEU:CD2	32:R:12:ARG:NH2	2.65	0.58
39:Y:126:ARG:HH22	41:a:1083:U:P	2.25	0.58
40:Z:14:MET:HG3	40:Z:15:SER:N	2.17	0.58
41:a:563:A:OP1	66:z:41:LYS:NZ	2.35	0.58
41:a:2243:U:H2'	41:a:2244:U:C6	2.38	0.58
41:a:2395:C:H2'	41:a:2396:G:O4'	2.03	0.58
45:e:2:LYS:CE	45:e:52:ARG:HE	2.16	0.58
52:l:119:ILE:O	52:l:187:VAL:HA	2.03	0.58
52:l:170:ARG:CZ	52:l:175:ILE:HA	2.33	0.58
53:m:24:THR:HG23	53:m:27:GLY:H	1.68	0.58
60:t:121:GLU:OE2	65:y:41:GLN:NE2	2.36	0.58
65:y:7:GLN:CD	65:y:7:GLN:C	2.71	0.58
5:4:46:LYS:O	5:4:50:MET:HG2	2.04	0.58
16:AG:240:THR:OG1	16:AG:247:PRO:HG2	2.01	0.58
10:B:70:G:H2'	10:B:71:C:H5''	1.85	0.58
18:D:522:C:OP2	32:R:66:TYR:OH	2.17	0.58
18:D:657:U:C4'	34:T:28:GLN:HE22	2.15	0.58
18:D:972:C:O2'	30:P:57:VAL:HG13	2.03	0.58
21:G:128:LYS:HD2	21:G:128:LYS:O	2.03	0.58
22:H:291:GLY:HA2	22:H:305:HIS:HA	1.85	0.58
27:M:26:PHE:CD1	27:M:101:MET:SD	2.97	0.58
41:a:654:A:H61	55:o:19:LYS:HD2	1.68	0.58
41:a:1869:G:N2	41:a:1871:A:O2'	2.37	0.58
56:p:85:LYS:CE	56:p:164:TYR:CD1	2.86	0.58
58:r:68:ARG:O	58:r:72:ILE:HD12	2.03	0.58
5:4:2:PHE:HE2	5:4:55:GLU:HG2	1.69	0.58
9:9:7:ASP:OD1	9:9:8:LYS:N	2.35	0.58
12:AB:125:GLY:O	14:AE:87:LYS:CA	2.51	0.58
14:AE:79:LYS:CA	16:AG:144:LYS:HZ1	2.00	0.58
14:AE:741:ALA:O	14:AE:762:ASN:ND2	2.32	0.58
16:AG:63:LEU:HD23	16:AG:63:LEU:H	1.69	0.58
16:AG:233:ARG:HG2	16:AG:270:ARG:HB2	1.84	0.58
16:AG:318:ILE:HG12	16:AG:338:VAL:HG11	1.85	0.58
16:AG:352:ALA:O	16:AG:356:ILE:CG2	2.51	0.58
17:C:16:GLU:OE1	17:C:18:VAL:N	2.36	0.58
18:D:1308:U:P	38:X:98:ARG:HG3	2.43	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:Q:51:GLY:O	31:Q:56:ARG:CZ	2.51	0.58
39:Y:109:ALA:HB2	39:Y:128:ILE:HG13	1.86	0.58
44:d:106:G:H2'	44:d:107:G:O4'	2.04	0.58
48:h:181:MET:HE2	48:h:181:MET:N	2.18	0.58
59:s:31:GLU:OE2	59:s:35:ARG:NH1	2.37	0.58
12:AB:89:PRO:HG2	14:AE:290:ILE:CG2	2.33	0.58
16:AG:33:THR:HG23	16:AG:43:ILE:HD11	1.85	0.58
10:B:75:C:P	41:a:2602:A:O2'	2.61	0.58
23:I:25:ASN:N	23:I:29:PHE:HE2	2.01	0.58
25:K:156:LYS:HA	28:N:66:PHE:CD2	2.39	0.58
27:M:18:PHE:HB3	27:M:59:LEU:HD11	1.84	0.58
34:T:11:ILE:HG21	34:T:31:LEU:HD13	1.85	0.58
41:a:832:U:H2'	41:a:833:A:H8	1.69	0.58
41:a:1197:G:H2'	41:a:1198:U:H6	1.68	0.58
58:r:6:LEU:HD11	58:r:37:VAL:CG2	2.34	0.58
58:r:37:VAL:HG12	58:r:43:ASN:HD21	1.68	0.58
60:t:88:ASN:ND2	60:t:90:ASN:OD1	2.36	0.58
2:1:73:LYS:C	2:1:74:ILE:HD12	2.29	0.58
9:9:23:LEU:N	9:9:118:ILE:HD11	2.19	0.58
11:AA:855:PRO:HB2	16:AG:35:THR:OG1	1.69	0.58
11:AA:856:ASN:CG	16:AG:35:THR:CG2	2.75	0.58
11:AA:861:ALA:CA	16:AG:38:LYS:HZ1	2.12	0.58
12:AB:6:LEU:H	12:AB:88:VAL:HG21	1.69	0.58
14:AE:1143:ASP:OD1	14:AE:1148:ARG:NH1	2.37	0.58
16:AG:413:ALA:HA	16:AG:416:THR:OG1	2.03	0.58
16:AG:453:VAL:HG11	16:AG:459:LEU:HG	1.84	0.58
18:D:921:U:H4'	25:K:23:LYS:NZ	2.18	0.58
24:J:74:ASN:HA	24:J:77:LYS:NZ	2.17	0.58
25:K:146:ASN:OD1	25:K:147:MET:N	2.37	0.58
27:M:24:ALA:O	27:M:27:VAL:HG22	2.04	0.58
41:a:580:U:H2'	41:a:581:C:H6	1.69	0.58
41:a:607:U:C5	41:a:620:G:C4	2.91	0.58
41:a:782:A:C2	48:h:225:MET:HE2	2.39	0.58
41:a:2349:G:C6	41:a:2369:A:C6	2.92	0.58
41:a:2815:C:C2	41:a:2816:G:C8	2.91	0.58
46:f:40:ASP:OD1	46:f:45:ARG:NH2	2.37	0.58
56:p:149:ARG:NH2	56:p:162:VAL:N	2.52	0.58
59:s:39:LYS:HA	59:s:44:TYR:CD1	2.39	0.58
59:s:132:HIS:O	59:s:135:GLN:HB2	2.03	0.58
60:t:112:PHE:CD2	60:t:115:ILE:HD12	2.39	0.58
62:v:49:ALA:O	62:v:52:ALA:N	2.37	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:49:ILE:HG22	16:AG:49:ILE:O	2.03	0.58
17:C:73:ARG:NH1	31:Q:113:VAL:HA	2.18	0.58
27:M:28:ASN:HA	27:M:31:MET:HE3	1.85	0.58
35:U:12:LYS:CD	35:U:13:LYS:HG2	2.33	0.58
41:a:2822:G:O2'	41:a:2824:C:OP2	2.20	0.58
58:r:97:ARG:NH1	58:r:101:ASP:OD2	2.37	0.58
9:9:54:VAL:HG13	9:9:54:VAL:O	2.03	0.58
10:A:15:G:N1	10:A:20:U:O2	2.37	0.58
11:AA:707:ALA:O	11:AA:711:ASP:HB2	2.03	0.58
12:AB:135:GLY:O	12:AB:137:ALA:N	2.37	0.58
16:AG:233:ARG:O	16:AG:327:LEU:HD21	2.04	0.58
18:D:974:A:OP1	33:S:69:ARG:NH1	2.37	0.58
21:G:66:LYS:HB3	21:G:154:MET:CE	2.34	0.58
27:M:103:TRP:CG	27:M:137:LYS:HZ2	2.22	0.58
41:a:5:A:H2'	41:a:6:A:C8	2.39	0.58
41:a:1568:G:N7	48:h:28:LYS:NZ	2.52	0.58
46:f:41:THR:O	46:f:44:ILE:N	2.30	0.58
51:k:34:LEU:H	51:k:51:GLU:CD	2.12	0.58
61:u:55:MET:CE	61:u:60:ARG:HG2	2.33	0.58
2:1:11:ARG:NH1	2:1:98:LYS:HE3	2.19	0.58
6:5:25:DA:C2	7:6:5:DG:C2	2.92	0.58
16:AG:266:LEU:HD23	16:AG:266:LEU:N	2.19	0.58
25:K:148:ASN:OD1	28:N:96:MET:HE1	2.03	0.58
34:T:32:LEU:HD11	34:T:62:GLN:OE1	2.04	0.58
39:Y:32:VAL:HG22	39:Y:60:VAL:CG2	2.34	0.58
41:a:593:U:H2'	41:a:594:U:C6	2.39	0.58
41:a:857:G:H5''	42:b:78:LYS:HZ2	1.69	0.58
41:a:2335:A:OP2	64:x:13:ARG:CD	2.51	0.58
52:l:148:ILE:HD13	52:l:187:VAL:HB	1.85	0.58
58:r:30:LEU:HD23	58:r:35:LYS:HE2	1.85	0.58
59:s:53:TYR:CD1	59:s:121:LYS:HD2	2.39	0.58
12:AB:141:LEU:HD23	12:AB:143:LEU:HD11	1.81	0.57
16:AG:233:ARG:HH12	16:AG:324:ASN:N	2.02	0.57
16:AG:440:VAL:HG22	16:AG:481:LEU:CD2	2.34	0.57
17:C:13:PHE:HB3	17:C:51:TYR:CD1	2.39	0.57
18:D:915:A:N6	18:D:916:U:C4	2.72	0.57
30:P:21:ALA:O	30:P:24:GLU:HG3	2.04	0.57
31:Q:20:VAL:CG1	31:Q:85:MET:HE1	2.33	0.57
36:V:4:LYS:HD2	36:V:5:ILE:HG23	1.85	0.57
41:a:628:G:C6	41:a:636:G:C2	2.92	0.57
41:a:2328:A:H2'	41:a:2329:U:H6	1.68	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:k:35:GLU:OE2	51:k:48:ILE:HG23	2.03	0.57
16:AG:130:PHE:C	16:AG:186:VAL:HG11	2.28	0.57
18:D:496:A:N3	18:D:496:A:H2'	2.17	0.57
21:G:35:ARG:HG2	21:G:40:ILE:HG13	1.86	0.57
33:S:90:ARG:C	33:S:92:GLU:OE1	2.47	0.57
36:V:11:ARG:HE	36:V:58:VAL:HG22	1.69	0.57
36:V:42:THR:HG22	36:V:44:LEU:CD1	2.34	0.57
38:X:25:VAL:HG23	38:X:29:ARG:HB2	1.85	0.57
39:Y:116:MET:HG2	39:Y:124:MET:CB	2.34	0.57
5:4:48:MET:HE3	5:4:84:PRO:O	2.04	0.57
6:5:9:DG:P	11:AA:470:ARG:HA	2.44	0.57
11:AA:13:LYS:HB2	11:AA:1180:MET:HE2	1.86	0.57
12:AB:31:PRO:O	12:AB:50:LEU:N	2.31	0.57
12:AB:158:GLU:HB3	12:AB:162:LEU:HD22	1.85	0.57
16:AG:354:ALA:CA	16:AG:357:ASP:HB3	2.32	0.57
16:AG:363:LEU:HG	16:AG:410:ALA:N	2.16	0.57
16:AG:363:LEU:CG	16:AG:410:ALA:CB	2.71	0.57
16:AG:387:VAL:CB	16:AG:388:PRO:CD	2.76	0.57
19:E:61:GLN:NE2	19:E:66:LEU:HD22	2.18	0.57
21:G:35:ARG:HB3	22:H:14:LYS:HZ3	1.69	0.57
25:K:93:ARG:HB3	25:K:93:ARG:NH2	2.18	0.57
32:R:110:ARG:NE	32:R:117:TYR:HD2	2.02	0.57
35:U:39:PHE:HD1	35:U:50:THR:HG22	1.70	0.57
41:a:1198:U:H2'	41:a:1199:U:C6	2.40	0.57
41:a:2189:U:O4'	41:a:2189:U:OP1	2.21	0.57
66:z:53:ARG:CG	66:z:57:PHE:HE2	2.18	0.57
66:z:74:ILE:HG12	66:z:79:PHE:HB2	1.86	0.57
1:0:15:SER:HB2	1:0:18:GLN:NE2	2.19	0.57
4:3:6:ARG:NH1	41:a:84:A:O2'	2.36	0.57
12:AB:102:LYS:CE	14:AE:285:LEU:HA	2.34	0.57
12:AB:102:LYS:HE2	14:AE:285:LEU:CA	2.34	0.57
16:AG:434:LEU:HD11	16:AG:455:THR:N	2.19	0.57
18:D:662:U:O2'	18:D:836:G:OP1	2.21	0.57
18:D:911:U:H2'	18:D:912:C:C6	2.40	0.57
22:H:51:GLU:OE1	22:H:51:GLU:N	2.38	0.57
22:H:75:VAL:HG23	22:H:81:GLU:OE2	2.04	0.57
23:I:82:GLU:O	23:I:86:LYS:HD3	2.04	0.57
28:N:66:PHE:CG	28:N:67:GLN:OE1	2.57	0.57
34:T:10:LYS:O	34:T:14:GLU:OE1	2.23	0.57
34:T:28:GLN:HB2	34:T:66:LEU:HD21	1.86	0.57
41:a:1682:G:H2'	41:a:1683:U:C6	2.39	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
45:e:18:LEU:HD11	45:e:54:LYS:HE3	1.86	0.57
52:l:62:GLN:O	52:l:69:ARG:HD3	2.03	0.57
52:l:136:GLN:NE2	52:l:140:ASP:OD2	2.37	0.57
62:v:26:VAL:HG13	62:v:104:GLU:CD	2.30	0.57
63:w:9:GLN:O	63:w:17:ARG:HD3	2.05	0.57
64:x:21:LEU:HD22	64:x:23:ALA:HB2	1.87	0.57
9:9:124:ASP:OD2	9:9:126:LEU:HD21	2.05	0.57
10:A:70:G:H2'	10:A:71:C:H5''	1.85	0.57
11:AA:481:LEU:CD2	12:AB:16:ARG:HD2	2.12	0.57
14:AE:144:TYR:OH	14:AE:293:ARG:NH2	2.37	0.57
16:AG:302:LYS:NZ	16:AG:302:LYS:HB2	2.20	0.57
21:G:19:GLN:CG	22:H:76:GLU:HB3	2.34	0.57
23:I:6:HIS:HB3	33:S:89:MET:SD	2.45	0.57
37:W:40:ILE:HG21	37:W:66:MET:SD	2.45	0.57
39:Y:91:LYS:HZ3	39:Y:97:VAL:HG21	1.70	0.57
41:a:1047:G:HO2'	41:a:1110:G:H1	1.51	0.57
41:a:1266:G:OP1	49:i:16:ARG:NE	2.35	0.57
54:n:4:LEU:HD11	54:n:101:GLU:N	2.18	0.57
62:v:41:LEU:CD2	62:v:45:GLN:HE22	2.18	0.57
65:y:9:GLU:OE2	65:y:55:LEU:N	2.37	0.57
9:9:38:MET:SD	39:Y:117:THR:HG22	2.45	0.57
16:AG:162:ILE:HD13	16:AG:167:MET:HE2	1.86	0.57
16:AG:210:ARG:HE	16:AG:216:ILE:HG22	1.69	0.57
16:AG:379:SER:HG	16:AG:384:LEU:HD21	1.69	0.57
10:B:60:U:H4'	10:B:61:C:OP2	2.04	0.57
17:C:55:LEU:O	17:C:59:ILE:HG12	2.04	0.57
18:D:1356:G:H2'	18:D:1357:A:C8	2.40	0.57
21:G:19:GLN:HG3	22:H:76:GLU:CB	2.35	0.57
27:M:68:ASN:ND2	27:M:130:ASN:OD1	2.38	0.57
29:O:5:GLN:NE2	29:O:20:PHE:CB	2.67	0.57
30:P:63:ASP:OD2	30:P:64:GLN:N	2.38	0.57
31:Q:52:PHE:CD2	31:Q:56:ARG:HG2	2.39	0.57
33:S:63:ARG:NH2	33:S:68:GLY:O	2.32	0.57
34:T:32:LEU:CD1	34:T:62:GLN:OE1	2.53	0.57
41:a:3:U:H2'	41:a:4:U:H6	1.69	0.57
41:a:500:G:N1	41:a:503:A:OP2	2.37	0.57
5:4:57:TYR:CD2	62:v:136:MET:HE2	2.40	0.57
10:A:60:U:H4'	10:A:61:C:OP2	2.04	0.57
10:A:75:C:O2'	10:A:76:A:C8	2.58	0.57
12:AB:38:ILE:HD13	12:AB:160:ARG:HH12	1.70	0.57
16:AG:61:ARG:HD2	16:AG:73:LYS:HD3	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:137:ILE:O	16:AG:137:ILE:HG22	2.03	0.57
16:AG:422:GLU:CG	16:AG:426:GLY:HA3	2.35	0.57
18:D:695:A:H8	18:D:695:A:H5''	1.68	0.57
20:F:16:LEU:C	20:F:20:LYS:HZ3	2.13	0.57
36:V:64:CYS:HG	36:V:74:THR:HG1	1.51	0.57
39:Y:58:ILE:HG23	39:Y:66:PHE:CE2	2.40	0.57
39:Y:116:MET:HB3	39:Y:124:MET:HE2	1.86	0.57
41:a:685:A:O2'	41:a:773:U:O4	2.17	0.57
41:a:1021:A:N1	41:a:1141:U:O4	2.37	0.57
41:a:2291:U:H2'	41:a:2292:U:C6	2.40	0.57
42:b:55:ARG:NE	42:b:55:ARG:HA	2.20	0.57
50:j:3:GLY:N	50:j:204:LYS:HE2	2.20	0.57
62:v:49:ALA:C	62:v:53:MET:HE1	2.30	0.57
65:y:51:ARG:HD3	65:y:58:ALA:HB3	1.86	0.57
3:2:19:LYS:NZ	41:a:1340:U:OP1	2.34	0.57
8:7:9:U:O4	18:D:1196:A:O4'	2.21	0.57
9:9:4:ASN:HB3	9:9:8:LYS:NZ	2.19	0.57
11:AA:888:THR:O	16:AG:105:ILE:HD13	2.05	0.57
18:D:1439:G:P	19:E:33:LYS:HZ1	2.25	0.57
21:G:52:GLU:O	21:G:56:GLU:OE1	2.23	0.57
26:L:14:GLN:OE1	26:L:83:ALA:HA	2.05	0.57
41:a:2071:A:H2'	41:a:2072:C:C6	2.40	0.57
41:a:2189:U:O2'	41:a:2190:G:O4'	2.22	0.57
41:a:2483:C:O2	62:v:51:ARG:NH2	2.38	0.57
43:c:66:THR:O	43:c:70:GLU:OE1	2.23	0.57
46:f:7:ILE:CG2	46:f:55:VAL:HG21	2.35	0.57
60:t:59:LYS:HD2	60:t:89:ASN:HA	1.87	0.57
8:7:7:U:H5''	8:7:7:U:C6	2.39	0.57
11:AA:847:PRO:N	16:AG:104:ARG:CZ	2.67	0.57
13:AC:45:ARG:HD3	13:AD:38:THR:HB	1.86	0.57
21:G:35:ARG:HD2	22:H:14:LYS:CE	2.34	0.57
29:O:94:LEU:O	29:O:98:LEU:HD13	2.04	0.57
31:Q:15:GLN:OE1	31:Q:78:GLY:HA3	2.05	0.57
33:S:87:ALA:HB1	33:S:93:ILE:HG13	1.87	0.57
41:a:659:G:H1'	52:l:100:MET:HE1	1.87	0.57
41:a:839:U:C2	41:a:840:C:C5	2.93	0.57
47:g:47:LYS:HG2	54:n:115:ARG:NH2	2.20	0.57
52:l:42:GLY:CA	52:l:92:HIS:HE1	2.18	0.57
54:n:46:ASP:OD1	54:n:49:LEU:HD13	2.03	0.57
59:s:120:ARG:CZ	59:s:120:ARG:HB2	2.34	0.57
66:z:74:ILE:CD1	66:z:74:ILE:CB	2.76	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:A:5:G:H2'	10:A:6:G:C5'	2.35	0.57
11:AA:856:ASN:OD1	16:AG:34:ALA:O	2.23	0.57
12:AB:130:PHE:CD2	12:AB:152:HIS:HE1	2.23	0.57
16:AG:31:LEU:HD23	16:AG:110:ALA:HB1	1.85	0.57
16:AG:367:GLU:O	16:AG:371:THR:HG23	2.04	0.57
16:AG:457:GLU:HG2	16:AG:489:CYS:SG	2.45	0.57
22:H:297:GLU:HG2	22:H:300:VAL:HG21	1.86	0.57
28:N:7:ILE:O	28:N:10:MET:HB3	2.05	0.57
31:Q:87:LYS:HG3	31:Q:113:VAL:HG23	1.86	0.57
34:T:21:ASP:OD1	34:T:22:THR:N	2.34	0.57
41:a:5:A:H2'	41:a:6:A:H8	1.70	0.57
41:a:16:C:C4'	49:i:15:MET:HE3	2.35	0.57
41:a:811:U:H2'	61:u:21:ARG:HA	1.86	0.57
41:a:2514:U:H2'	41:a:2515:C:C6	2.40	0.57
51:k:7:GLU:CG	51:k:27:LYS:HZ3	2.18	0.57
62:v:105:MET:HE2	62:v:108:VAL:HG21	1.87	0.57
4:3:94:ARG:HB3	4:3:103:ILE:HD12	1.86	0.56
8:7:1:A:H2'	8:7:2:U:H6	1.70	0.56
9:9:24:SER:HB2	9:9:86:MET:HA	1.86	0.56
11:AA:856:ASN:CA	16:AG:35:THR:HB	2.30	0.56
11:AA:1223:ARG:NH2	14:AE:721:SER:OG	2.38	0.56
16:AG:392:LEU:CD2	16:AG:395:ILE:HD11	2.34	0.56
18:D:193:C:O2'	19:E:59:ASP:OD2	2.23	0.56
18:D:280:C:O2	36:V:40:ARG:NH2	2.37	0.56
22:H:280:LEU:N	22:H:330:VAL:O	2.31	0.56
26:L:95:ALA:O	26:L:96:VAL:C	2.47	0.56
28:N:52:GLU:OE1	28:N:53:GLY:N	2.37	0.56
32:R:4:VAL:HG13	32:R:5:ASN:N	2.20	0.56
38:X:28:THR:HG22	38:X:29:ARG:NH1	2.21	0.56
41:a:839:U:H2'	41:a:840:C:C6	2.40	0.56
41:a:858:G:OP2	42:b:78:LYS:NZ	2.35	0.56
41:a:2191:A:H2'	41:a:2192:U:H6	1.70	0.56
41:a:2720:U:OP1	65:y:53:ARG:NH2	2.38	0.56
41:a:2849:U:P	65:y:93:ARG:NH2	2.78	0.56
44:d:42:C:C5'	54:n:64:LYS:HZ1	2.18	0.56
59:s:73:VAL:HG12	59:s:74:TYR:N	2.20	0.56
2:1:7:HIS:HD2	2:1:10:ALA:HB2	1.68	0.56
5:4:57:TYR:CB	62:v:136:MET:CE	2.84	0.56
11:AA:1254:VAL:O	14:AE:99:ARG:NH2	2.37	0.56
12:AB:117:ILE:HD13	12:AB:161:LYS:NZ	2.18	0.56
16:AG:173:PHE:N	16:AG:173:PHE:CD2	2.73	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:283:PHE:CE2	16:AG:331:LEU:O	2.58	0.56
16:AG:434:LEU:CD2	16:AG:455:THR:O	2.51	0.56
16:AG:447:LYS:O	16:AG:470:ILE:HD11	2.05	0.56
18:D:512:U:OP1	24:J:44:ARG:NH1	2.38	0.56
18:D:1290:G:H4'	27:M:35:LYS:HZ3	1.68	0.56
25:K:148:ASN:ND2	25:K:153:VAL:HG12	2.20	0.56
39:Y:77:VAL:HG13	39:Y:81:LYS:HZ3	1.69	0.56
41:a:64:A:H2'	41:a:65:U:C6	2.40	0.56
41:a:1570:A:H2'	41:a:1571:A:C8	2.41	0.56
53:m:12:ARG:NE	53:m:44:VAL:HG21	2.19	0.56
55:o:30:ARG:HH12	61:u:62:PRO:HB3	1.70	0.56
62:v:70:ASP:C	62:v:92:TRP:HZ3	2.13	0.56
65:y:106:LYS:HA	65:y:109:ARG:NH2	2.20	0.56
3:2:19:LYS:O	3:2:22:THR:OG1	2.22	0.56
3:2:56:GLU:CD	3:2:87:LEU:C	2.73	0.56
11:AA:838:CYS:HB2	11:AA:918:LEU:HD22	1.85	0.56
11:AA:1122:LYS:NZ	11:AA:1126:ASP:OD1	2.38	0.56
12:AB:130:PHE:CZ	12:AB:160:ARG:HG3	2.41	0.56
16:AG:9:VAL:HA	16:AG:24:PHE:CZ	2.40	0.56
16:AG:434:LEU:HD23	16:AG:456:LEU:CG	2.35	0.56
18:D:552:U:O2'	32:R:83:ARG:O	2.19	0.56
18:D:562:U:C3'	32:R:14:ARG:HH11	2.17	0.56
21:G:131:LYS:HG3	21:G:132:LYS:HD2	1.87	0.56
22:H:116:VAL:CA	22:H:279:LYS:HD2	2.35	0.56
22:H:117:LYS:CB	22:H:279:LYS:O	2.54	0.56
24:J:121:LYS:HG2	24:J:131:ASN:ND2	2.20	0.56
24:J:196:ASN:ND2	24:J:199:LEU:HG	2.20	0.56
27:M:66:LEU:HD23	27:M:70:ARG:HH21	1.69	0.56
27:M:95:ARG:O	27:M:99:LEU:HG	2.05	0.56
27:M:134:ALA:O	27:M:137:LYS:HG2	2.05	0.56
29:O:118:LEU:CD2	29:O:124:ARG:HA	2.36	0.56
31:Q:87:LYS:HG3	31:Q:113:VAL:CG2	2.36	0.56
41:a:78:U:H5''	45:e:4:LYS:NZ	2.20	0.56
41:a:228:C:N4	41:a:2407:A:N3	2.52	0.56
41:a:1693:U:O2'	48:h:14:ARG:NH2	2.38	0.56
41:a:2192:U:H2'	41:a:2193:G:C8	2.40	0.56
41:a:2683:C:H4'	50:j:13:ARG:CZ	2.34	0.56
48:h:52:ARG:HE	48:h:53:HIS:CE1	2.23	0.56
63:w:35:LYS:NZ	63:w:112:TYR:HE2	2.04	0.56
5:4:53:LYS:HB2	5:4:56:PHE:CE2	2.40	0.56
9:9:27:VAL:HG13	9:9:27:VAL:O	2.05	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:9:47:GLU:O	9:9:51:TYR:CE1	2.59	0.56
16:AG:130:PHE:HB3	16:AG:186:VAL:HG22	1.86	0.56
18:D:439:U:O2	18:D:440:C:C6	2.58	0.56
19:E:61:GLN:HE21	19:E:66:LEU:HD22	1.69	0.56
23:I:140:ASN:O	23:I:144:LEU:HD23	2.06	0.56
26:L:42:TRP:CZ2	26:L:102:MET:CG	2.88	0.56
37:W:17:LYS:HB3	37:W:21:LYS:HZ3	1.70	0.56
41:a:1360:G:N7	41:a:1361:G:C8	2.73	0.56
41:a:1818:U:OP2	48:h:156:ARG:CZ	2.53	0.56
41:a:2816:G:O4'	49:i:41:HIS:CE1	2.57	0.56
44:d:48:U:H2'	44:d:49:C:C6	2.40	0.56
45:e:2:LYS:HD2	45:e:6:LEU:HD11	1.85	0.56
52:l:17:THR:HA	52:l:21:ARG:NH2	2.19	0.56
54:n:163:ASP:OD1	54:n:164:GLU:N	2.39	0.56
55:o:55:LEU:HD21	55:o:59:ILE:HD11	1.87	0.56
61:u:90:VAL:CG1	61:u:95:LEU:HD21	2.33	0.56
4:3:6:ARG:CZ	41:a:84:A:H2'	2.35	0.56
8:7:6:U:C2	18:D:1493:A:H2	2.24	0.56
13:AC:82:LEU:HD11	13:AC:171:LEU:HD23	1.86	0.56
14:AE:79:LYS:HZ3	16:AG:153:ASP:CG	2.13	0.56
16:AG:63:LEU:HD23	16:AG:63:LEU:N	2.21	0.56
19:E:61:GLN:HE21	19:E:66:LEU:CD2	2.18	0.56
23:I:66:VAL:HB	23:I:101:ILE:HD12	1.88	0.56
30:P:15:HIS:HA	30:P:18:ILE:HG22	1.87	0.56
38:X:71:ARG:NE	47:g:50:ASP:O	2.39	0.56
39:Y:45:THR:HG21	39:Y:50:LYS:CD	2.36	0.56
41:a:17:G:H2'	41:a:18:U:C6	2.41	0.56
62:v:37:GLY:C	62:v:126:ILE:HD11	2.30	0.56
12:AB:26:VAL:HA	12:AB:58:GLU:OE1	2.05	0.56
12:AB:141:LEU:O	12:AB:151:LYS:HA	2.03	0.56
16:AG:320:ARG:HB2	16:AG:323:GLN:HB2	1.88	0.56
17:C:60:LYS:HZ1	18:D:735:C:C4'	2.16	0.56
18:D:440:C:C4	18:D:441:A:N7	2.73	0.56
21:G:107:VAL:O	21:G:111:ILE:HG12	2.04	0.56
25:K:99:ALA:HB2	25:K:124:LEU:HG	1.88	0.56
25:K:136:VAL:O	25:K:140:THR:HG23	2.04	0.56
41:a:64:A:H2'	41:a:65:U:H6	1.71	0.56
41:a:1141:U:H5	59:s:68:LYS:NZ	2.02	0.56
41:a:1824:G:H5''	48:h:52:ARG:HH22	1.69	0.56
6:5:10:DT:O3'	12:AB:68:THR:HA	2.03	0.56
16:AG:287:ALA:HB1	16:AG:331:LEU:CB	2.36	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:362:TYR:CZ	16:AG:382:GLU:HG2	2.40	0.56
16:AG:392:LEU:HD22	16:AG:395:ILE:HD11	1.87	0.56
26:L:102:MET:HG2	26:L:103:VAL:N	2.06	0.56
38:X:16:VAL:CG2	38:X:17:ILE:HD12	2.35	0.56
41:a:589:U:H2'	41:a:590:A:H8	1.71	0.56
41:a:1442:U:H2'	41:a:1443:U:C6	2.40	0.56
41:a:1565:C:OP1	48:h:18:LYS:NZ	2.26	0.56
41:a:1857:G:O2'	41:a:1884:G:N2	2.38	0.56
41:a:2299:U:H2'	41:a:2300:C:H6	1.71	0.56
42:b:45:PHE:CD2	42:b:80:ILE:HD11	2.40	0.56
52:l:1:MET:HE1	52:l:20:GLY:N	2.21	0.56
54:n:62:GLY:C	54:n:95:ARG:NH2	2.63	0.56
56:p:85:LYS:HE3	56:p:164:TYR:CE1	2.41	0.56
8:7:11:U:O2	8:7:11:U:H2'	2.06	0.56
10:A:16:C:O2	41:a:2181:U:H5'	2.06	0.56
12:AB:123:PHE:HD2	12:AB:123:PHE:N	2.02	0.56
16:AG:187:ARG:CB	16:AG:188:PRO:HD3	2.20	0.56
16:AG:285:ILE:O	16:AG:288:MET:CE	2.54	0.56
18:D:521:G:O5'	32:R:70:GLU:OE1	2.23	0.56
19:E:9:LYS:C	19:E:13:GLN:NE2	2.61	0.56
21:G:5:SER:OG	21:G:8:ASP:OD2	2.24	0.56
22:H:116:VAL:HA	22:H:279:LYS:HD2	1.88	0.56
41:a:1009:A:OP2	59:s:39:LYS:NZ	2.37	0.56
41:a:1668:A:O2'	41:a:1674:G:N7	2.32	0.56
41:a:2335:A:OP2	64:x:13:ARG:HB3	2.06	0.56
51:k:19:HIS:ND1	51:k:41:PRO:HD2	2.21	0.56
62:v:77:PRO:HG2	62:v:80:VAL:CG2	2.31	0.56
64:x:36:TYR:CD2	64:x:52:SER:HB2	2.41	0.56
8:7:12:U:H3'	8:7:12:U:H6	1.70	0.56
10:A:22:G:H2'	10:A:23:C:H6	1.70	0.56
12:AB:34:THR:O	12:AB:106:ASP:OD1	2.24	0.56
16:AG:437:LEU:HD22	16:AG:488:ILE:HD12	1.88	0.56
18:D:671:G:O3'	26:L:79:ARG:NH2	2.39	0.56
18:D:1381:U:C2	18:D:1382:C:C6	2.94	0.56
25:K:74:VAL:HG11	25:K:144:LEU:HD12	1.88	0.56
27:M:23:LEU:HD12	27:M:23:LEU:H	1.70	0.56
28:N:7:ILE:O	28:N:11:LEU:HD13	2.06	0.56
34:T:35:GLN:CB	34:T:59:MET:HE1	2.36	0.56
35:U:14:ARG:HD3	35:U:42:ILE:HG21	1.86	0.56
38:X:93:ARG:CG	38:X:95:LEU:HD23	2.35	0.56
41:a:2517:C:C2	41:a:2542:A:N6	2.74	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:b:51:VAL:C	42:b:62:LYS:HZ3	2.14	0.56
45:e:2:LYS:HE2	45:e:52:ARG:HE	1.71	0.56
47:g:2:LYS:HB2	47:g:5:ILE:HD11	1.88	0.56
49:i:40:ARG:HG3	49:i:41:HIS:ND1	2.21	0.56
52:l:62:GLN:HA	52:l:69:ARG:NH2	2.20	0.56
56:p:109:PHE:CE1	56:p:152:ARG:NH1	2.74	0.56
58:r:54:LEU:HA	58:r:57:LYS:NZ	2.20	0.56
60:t:20:MET:HB3	60:t:44:LYS:CE	2.35	0.56
61:u:125:LEU:HD23	61:u:125:LEU:N	2.21	0.56
3:2:2:ILE:HG21	3:2:42:GLU:OE1	2.06	0.56
11:AA:516:ASP:OD1	11:AA:516:ASP:O	2.22	0.56
11:AA:818:VAL:HG22	11:AA:1096:ILE:HG12	1.87	0.56
11:AA:985:GLU:HB3	11:AA:988:LYS:HB2	1.88	0.56
14:AE:679:TYR:OH	14:AE:754:ILE:O	2.23	0.56
18:D:652:U:C3'	28:N:56:LYS:HZ1	2.18	0.56
34:T:3:LEU:HB2	34:T:35:GLN:CD	2.31	0.56
41:a:1038:G:H2'	41:a:1039:A:C8	2.41	0.56
41:a:2376:A:N3	64:x:111:ARG:NH1	2.54	0.56
42:b:51:VAL:C	42:b:62:LYS:NZ	2.63	0.56
45:e:56:LEU:O	45:e:59:GLU:HG3	2.06	0.56
47:g:5:ILE:CD1	54:n:64:LYS:HE3	2.36	0.56
48:h:183:LYS:HE2	48:h:266:PHE:HA	1.88	0.56
50:j:91:THR:HB	50:j:94:GLN:HG2	1.87	0.56
51:k:5:ILE:HG23	51:k:28:ARG:NH1	2.21	0.56
58:r:4:ILE:HD12	58:r:17:ASP:O	2.05	0.56
63:w:67:PHE:O	63:w:71:ARG:N	2.39	0.56
9:9:24:SER:CB	9:9:86:MET:HE2	2.36	0.55
9:9:73:LYS:HD2	9:9:73:LYS:O	2.06	0.55
11:AA:860:ALA:N	16:AG:38:LYS:HE2	2.17	0.55
16:AG:387:VAL:HB	16:AG:388:PRO:HD2	1.85	0.55
18:D:588:G:C5'	28:N:3:MET:HE1	2.36	0.55
18:D:1071:C:H2'	18:D:1072:G:H8	1.71	0.55
22:H:163:ARG:CB	22:H:298:GLU:HG3	2.35	0.55
23:I:150:LYS:HE3	23:I:201:TRP:CZ3	2.41	0.55
31:Q:61:PHE:O	31:Q:64:GLN:HG2	2.06	0.55
40:Z:2:ILE:HB	40:Z:5:ASP:HB3	1.88	0.55
41:a:1321:A:C4	41:a:1322:A:C8	2.95	0.55
41:a:1365:A:O2'	43:c:11:ARG:NH1	2.39	0.55
41:a:2803:G:H2'	41:a:2804:U:C6	2.41	0.55
50:j:55:LYS:NZ	50:j:60:VAL:N	2.55	0.55
52:l:1:MET:CE	52:l:16:GLU:HA	2.26	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:p:94:TYR:CE1	56:p:107:LEU:O	2.59	0.55
59:s:37:ARG:HD3	59:s:39:LYS:HD2	1.88	0.55
2:1:33:LEU:HD23	2:1:51:LEU:HD23	1.89	0.55
5:4:80:HIS:CD2	5:4:81:PRO:HD2	2.41	0.55
8:7:8:U:H5'	18:D:1397:C:C2	2.41	0.55
16:AG:354:ALA:HA	16:AG:357:ASP:H	1.70	0.55
10:B:18:G:H1'	10:B:58:A:C2	2.41	0.55
10:B:22:G:H2'	10:B:23:C:H6	1.70	0.55
18:D:276:G:H5'	36:V:17:MET:CE	2.35	0.55
18:D:1082:A:OP1	25:K:23:LYS:NZ	2.34	0.55
18:D:1366:C:O2'	30:P:62:ARG:NH2	2.37	0.55
18:D:1515:G:H2'	18:D:1516:G:H8	1.70	0.55
25:K:45:ARG:C	25:K:71:MET:HE2	2.32	0.55
25:K:112:ARG:HG3	25:K:116:GLU:OE2	2.06	0.55
38:X:28:THR:O	38:X:31:LYS:HG2	2.07	0.55
41:a:1289:C:C2	41:a:1290:C:C5	2.94	0.55
41:a:1469:A:H2'	41:a:1470:A:C8	2.42	0.55
50:j:46:ARG:NH1	50:j:85:ALA:O	2.32	0.55
63:w:24:MET:HA	63:w:24:MET:CE	2.33	0.55
3:2:56:GLU:CD	3:2:88:LYS:HA	2.32	0.55
9:9:56:ARG:NH2	41:a:1084:A:C1'	2.69	0.55
9:9:122:GLN:HG2	9:9:123:ILE:H	1.70	0.55
12:AB:120:GLU:CB	12:AB:162:LEU:HD12	2.37	0.55
12:AB:134:ASP:OD1	12:AB:135:GLY:N	2.37	0.55
13:AC:28:LEU:HD22	13:AC:201:LEU:HD23	1.88	0.55
17:C:40:VAL:HG13	22:H:339:ARG:HD3	1.88	0.55
19:E:32:ILE:HG23	19:E:75:HIS:NE2	2.21	0.55
20:F:26:ALA:HB3	20:F:28:VAL:HG23	1.88	0.55
23:I:156:ARG:NH1	23:I:193:TYR:O	2.39	0.55
27:M:4:ARG:HG3	27:M:5:ARG:CG	2.22	0.55
34:T:80:GLN:O	34:T:84:ARG:HG2	2.05	0.55
41:a:1348:C:C5	41:a:1349:C:C5	2.95	0.55
41:a:1799:G:C2	48:h:154:LEU:HD23	2.41	0.55
59:s:133:ALA:C	59:s:135:GLN:N	2.60	0.55
4:3:47:LYS:HE3	41:a:483:A:OP1	2.06	0.55
5:4:55:GLU:O	5:4:59:GLU:OE1	2.24	0.55
5:4:80:HIS:CE1	5:4:83:LYS:HG3	2.41	0.55
6:5:9:DG:OP2	11:AA:473:ARG:HG2	2.05	0.55
6:5:16:DG:OP1	11:AA:175:ARG:NH2	2.39	0.55
9:9:46:ARG:CG	9:9:94:ARG:HH21	2.02	0.55
9:9:112:ALA:O	9:9:123:ILE:CD1	2.54	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AA:103:VAL:HB	11:AA:114:VAL:HG11	1.89	0.55
11:AA:858:GLY:O	16:AG:38:LYS:NZ	2.39	0.55
16:AG:131:ARG:CA	16:AG:186:VAL:HG11	2.35	0.55
10:B:22:G:O2'	10:B:23:C:P	2.65	0.55
18:D:502:A:O3'	32:R:116:LYS:HE2	2.06	0.55
18:D:827:U:O4	18:D:872:A:N1	2.39	0.55
22:H:118:GLY:O	22:H:133:GLY:HA2	2.06	0.55
23:I:132:ARG:HA	23:I:135:LYS:HE3	1.89	0.55
34:T:3:LEU:HD22	34:T:8:THR:HG22	1.88	0.55
41:a:728:G:C4	41:a:730:A:C8	2.95	0.55
41:a:923:G:O4'	42:b:29:GLU:OE2	2.24	0.55
41:a:2189:U:O2'	41:a:2190:G:C5'	2.54	0.55
45:e:20:ASN:O	45:e:23:ARG:HG2	2.05	0.55
54:n:134:GLU:N	54:n:134:GLU:OE1	2.40	0.55
59:s:13:ARG:CZ	59:s:49:ASP:O	2.54	0.55
62:v:96:ILE:HA	62:v:100:LYS:NZ	2.21	0.55
2:1:25:ARG:NH2	41:a:519:U:O3'	2.40	0.55
6:5:10:DT:O3'	12:AB:68:THR:C	2.47	0.55
9:9:43:LYS:HD3	9:9:98:GLU:CG	2.35	0.55
12:AB:4:TRP:CD2	12:AB:27:ASN:ND2	2.75	0.55
16:AG:46:ARG:HH11	16:AG:61:ARG:HB3	1.71	0.55
16:AG:235:LYS:HG2	16:AG:235:LYS:O	2.07	0.55
16:AG:285:ILE:HD12	16:AG:293:VAL:HG11	1.88	0.55
16:AG:307:ILE:HD11	16:AG:336:LEU:HB2	1.87	0.55
16:AG:425:LEU:N	16:AG:429:LYS:HG3	2.20	0.55
18:D:50:A:O2'	18:D:360:G:N2	2.40	0.55
18:D:1370:G:C2	18:D:1371:G:C8	2.95	0.55
19:E:6:SER:O	19:E:10:ARG:HG2	2.07	0.55
22:H:100:LYS:O	22:H:104:ASP:N	2.38	0.55
24:J:62:ARG:HA	24:J:72:PHE:CE1	2.41	0.55
41:a:888:C:N4	41:a:889:C:N4	2.55	0.55
41:a:1720:U:H2'	41:a:1721:G:O4'	2.06	0.55
48:h:107:PRO:HD2	48:h:110:LEU:HD22	1.89	0.55
50:j:78:GLY:C	50:j:79:LEU:HD22	2.32	0.55
60:t:18:ARG:HB3	60:t:45:GLU:HB3	1.89	0.55
1:0:10:LYS:HE2	41:a:996:A:OP1	2.07	0.55
5:4:77:VAL:HG11	5:4:79:ARG:NH2	2.21	0.55
9:9:98:GLU:HA	9:9:101:LYS:NZ	2.22	0.55
12:AB:8:TYR:OH	14:AE:278:ARG:NH2	2.40	0.55
16:AG:133:HIS:CE1	16:AG:136:GLU:OE1	2.59	0.55
18:D:1329:A:OP1	38:X:28:THR:HB	2.06	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:I:151:VAL:HG12	23:I:200:VAL:CG2	2.35	0.55
30:P:26:VAL:O	30:P:30:LYS:HG3	2.07	0.55
38:X:95:LEU:O	38:X:109:ARG:NH2	2.39	0.55
50:j:11:MET:SD	50:j:11:MET:C	2.90	0.55
54:n:5:HIS:HD2	54:n:97:TRP:CE2	2.24	0.55
60:t:12:ASP:CG	60:t:85:VAL:HG13	2.31	0.55
8:7:1:A:C4	8:7:2:U:C5	2.95	0.55
10:A:37:A:H2'	10:A:38:A:O4'	2.07	0.55
12:AB:123:PHE:N	12:AB:123:PHE:CD2	2.73	0.55
18:D:280:C:H1'	36:V:40:ARG:NH1	2.21	0.55
18:D:672:U:O2'	18:D:673:A:H5'	2.06	0.55
18:D:1208:C:H2'	18:D:1209:C:C6	2.40	0.55
24:J:162:ALA:HA	24:J:165:ARG:CZ	2.36	0.55
38:X:95:LEU:HB3	38:X:96:PRO:CD	2.37	0.55
41:a:960:A:H61	62:v:82:MET:HE1	1.71	0.55
41:a:1059:G:H2'	41:a:1060:U:C5	2.41	0.55
41:a:2298:A:P	54:n:72:LYS:HZ3	2.29	0.55
44:d:45:A:H5'	54:n:92:ARG:NH1	2.20	0.55
45:e:23:ARG:HG3	45:e:24:GLU:N	2.21	0.55
55:o:30:ARG:NH1	61:u:62:PRO:HB3	2.21	0.55
59:s:74:TYR:CE2	59:s:92:MET:HE1	2.42	0.55
61:u:57:LEU:HA	61:u:60:ARG:HD2	1.89	0.55
1:0:13:ARG:NH2	66:z:97:ASP:OD1	2.40	0.55
3:2:54:GLU:OE1	3:2:54:GLU:N	2.38	0.55
9:9:61:ARG:NH2	41:a:1045:C:O3'	2.40	0.55
10:A:60:U:P	10:A:61:C:H41	2.30	0.55
11:AA:858:GLY:HA2	16:AG:38:LYS:HB2	1.78	0.55
12:AB:141:LEU:CA	12:AB:152:HIS:C	2.80	0.55
16:AG:162:ILE:HG22	16:AG:162:ILE:O	2.06	0.55
18:D:750:C:N3	18:D:751:U:C5	2.75	0.55
22:H:5:PHE:O	22:H:9:PHE:HD1	1.90	0.55
22:H:268:VAL:HG23	22:H:269:ALA:H	1.72	0.55
31:Q:20:VAL:HB	31:Q:35:THR:HG23	1.89	0.55
38:X:96:PRO:HG3	38:X:102:THR:HG22	1.89	0.55
41:a:150:U:H2'	41:a:151:C:H6	1.72	0.55
41:a:930:G:H1'	46:f:25:LEU:HD21	1.87	0.55
41:a:1708:C:C2	41:a:1709:U:C5	2.95	0.55
41:a:2315:G:H4'	54:n:127:ASN:HD22	1.71	0.55
41:a:2351:G:O6	55:o:39:LYS:CE	2.55	0.55
41:a:2638:G:O2'	41:a:2775:G:N2	2.37	0.55
45:e:28:LEU:HD23	45:e:43:LEU:HD22	1.89	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
45:e:53:VAL:O	45:e:57:LEU:HD23	2.06	0.55
59:s:15:TRP:HE3	59:s:135:GLN:O	1.90	0.55
60:t:59:LYS:CD	60:t:89:ASN:HA	2.36	0.55
61:u:76:GLU:CG	61:u:111:ILE:HD13	2.37	0.55
10:A:56:C:H1'	41:a:2112:G:C6	2.41	0.55
11:AA:143:ARG:NH2	11:AA:512:SER:O	2.40	0.55
11:AA:360:LEU:HD13	11:AA:378:ARG:HH11	1.72	0.55
16:AG:168:LEU:CD2	16:AG:230:PRO:O	2.55	0.55
16:AG:305:MET:HG3	16:AG:336:LEU:HB3	1.89	0.55
16:AG:320:ARG:NH1	16:AG:320:ARG:HG2	2.22	0.55
18:D:1078:U:O3'	25:K:138:ARG:NH2	2.40	0.55
21:G:27:MET:CE	21:G:187:VAL:HG12	2.37	0.55
24:J:138:SER:O	24:J:141:ASP:OD1	2.24	0.55
25:K:157:ARG:NE	28:N:43:GLU:O	2.40	0.55
39:Y:44:LYS:O	39:Y:44:LYS:HD2	2.06	0.55
41:a:1082:U:O4	41:a:1086:A:N1	2.39	0.55
62:v:45:GLN:HE21	62:v:124:LEU:HD23	1.72	0.55
63:w:94:TYR:C	63:w:116:VAL:HG23	2.32	0.55
66:z:78:LYS:CD	66:z:117:LEU:CD2	2.84	0.55
2:1:62:ASP:OD1	2:1:62:ASP:O	2.24	0.55
4:3:26:LYS:HE2	4:3:26:LYS:HA	1.89	0.55
5:4:80:HIS:ND1	5:4:83:LYS:N	2.55	0.55
9:9:58:THR:HG21	9:9:81:LEU:CD2	2.31	0.55
10:A:22:G:O2'	10:A:23:C:P	2.65	0.55
11:AA:839:VAL:HG12	11:AA:1049:ILE:HG12	1.87	0.55
10:B:37:A:H2'	10:B:38:A:O4'	2.07	0.55
17:C:54:GLN:HG3	17:C:55:LEU:N	2.22	0.55
22:H:268:VAL:HG23	22:H:269:ALA:N	2.22	0.55
33:S:42:TRP:CH2	47:g:66:ILE:HG23	2.42	0.55
37:W:64:ASP:OD2	47:g:57:VAL:HG23	2.06	0.55
41:a:2741:A:O3'	57:q:36:ARG:NH1	2.40	0.55
48:h:157:SER:O	48:h:195:VAL:HG11	2.07	0.55
50:j:180:VAL:O	50:j:180:VAL:HG13	2.05	0.55
56:p:33:LEU:HD11	56:p:136:ALA:HB1	1.89	0.55
62:v:1:MET:HE3	62:v:1:MET:CA	2.34	0.55
2:1:43:ALA:O	2:1:47:VAL:HG12	2.07	0.54
11:AA:856:ASN:CA	16:AG:35:THR:CA	2.81	0.54
12:AB:141:LEU:CB	12:AB:143:LEU:HD11	2.29	0.54
16:AG:130:PHE:O	16:AG:186:VAL:HG11	2.07	0.54
16:AG:288:MET:HE2	16:AG:293:VAL:HB	1.88	0.54
10:B:60:U:P	10:B:61:C:H41	2.30	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:1076:U:OP1	21:G:174:LYS:NZ	2.40	0.54
19:E:19:LYS:HG3	19:E:20:HIS:N	2.22	0.54
24:J:175:ALA:O	24:J:178:MET:SD	2.65	0.54
28:N:7:ILE:O	28:N:11:LEU:CD1	2.55	0.54
39:Y:33:ASN:HB3	39:Y:36:GLU:CG	2.30	0.54
39:Y:123:ALA:O	39:Y:126:ARG:HG3	2.07	0.54
41:a:134:G:H2'	41:a:135:U:C6	2.43	0.54
41:a:608:A:H2'	41:a:609:A:C8	2.42	0.54
41:a:1665:A:O3'	60:t:66:LYS:HD3	2.07	0.54
41:a:2683:C:H4'	50:j:13:ARG:NH2	2.22	0.54
41:a:2742:G:P	57:q:36:ARG:NH1	2.80	0.54
57:q:1:MET:HE1	57:q:35:GLN:C	2.32	0.54
2:1:31:GLN:O	2:1:35:ILE:HG12	2.07	0.54
2:1:41:LYS:HZ2	41:a:2009:A:P	2.30	0.54
9:9:23:LEU:HD21	9:9:96:PHE:CD1	2.42	0.54
10:A:18:G:H1'	10:A:58:A:C2	2.41	0.54
11:AA:377:THR:HB	12:AB:65:HIS:HE2	1.72	0.54
11:AA:1142:ARG:NH2	11:AA:1166:ASP:OD1	2.38	0.54
16:AG:223:ILE:HD13	16:AG:223:ILE:C	2.32	0.54
10:B:5:G:H2'	10:B:6:G:C5'	2.35	0.54
10:B:32:C:H5'	18:D:1340:A:O3'	2.06	0.54
18:D:946:A:H2'	18:D:947:G:C8	2.41	0.54
19:E:78:ASN:C	19:E:82:GLN:OE1	2.50	0.54
21:G:30:PHE:CD1	21:G:45:LYS:NZ	2.74	0.54
22:H:71:ALA:C	22:H:72:LEU:HD12	2.32	0.54
27:M:27:VAL:HG12	27:M:43:VAL:HG21	1.90	0.54
30:P:5:ARG:HG3	30:P:7:ARG:NH1	2.23	0.54
36:V:31:HIS:ND1	36:V:32:PRO:HD2	2.23	0.54
41:a:16:C:O4'	49:i:15:MET:CE	2.55	0.54
41:a:151:C:C2	41:a:152:A:C8	2.95	0.54
41:a:2591:C:H2'	41:a:2592:G:C8	2.42	0.54
45:e:2:LYS:HE2	45:e:49:ASP:HA	1.88	0.54
47:g:61:ASN:OD1	47:g:65:ASN:ND2	2.40	0.54
60:t:41:ILE:HD11	60:t:86:LEU:HD22	1.89	0.54
63:w:24:MET:HE2	63:w:34:ILE:HD13	1.87	0.54
6:5:15:DT:C2	11:AA:183:TRP:CD1	2.94	0.54
12:AB:78:PHE:CD1	12:AB:85:PRO:HB3	2.42	0.54
14:AE:886:VAL:HG11	14:AE:1230:THR:HG21	1.89	0.54
16:AG:448:LEU:HD22	16:AG:467:LEU:HD22	1.89	0.54
17:C:34:THR:HG22	17:C:36:SER:H	1.72	0.54
20:F:19:PHE:CE1	20:F:23:CYS:SG	3.00	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:F:19:PHE:CE2	31:Q:110:ILE:HG22	2.42	0.54
26:L:99:ALA:HB3	26:L:104:LYS:NZ	2.22	0.54
30:P:6:ILE:HG21	30:P:100:ILE:HD11	1.89	0.54
30:P:47:GLU:OE1	30:P:67:ILE:O	2.24	0.54
41:a:3:U:H2'	41:a:4:U:C6	2.42	0.54
41:a:151:C:H2'	41:a:152:A:H8	1.72	0.54
41:a:784:G:H5'	41:a:785:G:OP1	2.06	0.54
41:a:1353:A:H2'	41:a:1354:A:C8	2.43	0.54
41:a:2063:C:O2	41:a:2063:C:H2'	2.06	0.54
41:a:2766:A:C2	41:a:2767:C:C6	2.96	0.54
41:a:2838:G:C4	41:a:2839:G:C8	2.95	0.54
48:h:17:VAL:HB	48:h:204:VAL:HG12	1.89	0.54
50:j:156:PHE:CD2	59:s:81:ILE:HD13	2.42	0.54
65:y:15:GLN:NE2	65:y:16:ASP:OD1	2.40	0.54
7:6:20:DA:C6	8:7:48:G:N2	2.76	0.54
8:7:3:G:C3'	18:D:1500:A:N6	2.71	0.54
11:AA:29:SER:O	11:AA:33:ASP:HB2	2.07	0.54
11:AA:1117:LEU:HD12	11:AA:1195:ILE:HG12	1.90	0.54
16:AG:448:LEU:HD11	16:AG:481:LEU:HD13	1.89	0.54
10:B:31:G:O2'	18:D:1340:A:H4'	2.08	0.54
18:D:17:U:H2'	18:D:18:C:C6	2.42	0.54
18:D:280:C:H1'	36:V:40:ARG:CZ	2.37	0.54
18:D:375:U:OP1	35:U:70:ARG:CG	2.54	0.54
19:E:22:ALA:CA	19:E:25:ARG:HH21	2.21	0.54
20:F:31:GLU:O	20:F:34:ARG:HG2	2.08	0.54
23:I:155:GLY:O	23:I:196:ILE:HG12	2.07	0.54
25:K:14:LYS:HD3	25:K:117:VAL:CG1	2.38	0.54
26:L:39:LEU:HB3	26:L:62:MET:HE1	1.89	0.54
39:Y:15:GLY:HA2	39:Y:50:LYS:CG	2.34	0.54
41:a:15:G:H2'	49:i:15:MET:HE1	1.90	0.54
41:a:537:G:OP2	59:s:2:LYS:CE	2.55	0.54
41:a:1349:C:C2	41:a:1350:C:C5	2.95	0.54
41:a:1563:U:H2'	41:a:1564:C:C6	2.43	0.54
41:a:1774:C:O2	41:a:1774:C:H2'	2.07	0.54
41:a:1790:C:H2'	41:a:1791:A:C5	2.43	0.54
41:a:2020:A:C2	41:a:2022:U:O4'	2.61	0.54
52:l:69:ARG:HA	52:l:69:ARG:CZ	2.37	0.54
55:o:16:LYS:HD2	55:o:65:ALA:HB1	1.88	0.54
56:p:149:ARG:NE	56:p:162:VAL:O	2.40	0.54
57:q:9:LYS:HZ3	57:q:11:CYS:C	2.16	0.54
65:y:31:TRP:CZ3	65:y:40:LEU:HD11	2.42	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
66:z:78:LYS:HD3	66:z:117:LEU:HD23	1.88	0.54
5:4:57:TYR:HB3	62:v:136:MET:HE3	1.87	0.54
6:5:10:DT:C7	11:AA:62:TYR:HB3	2.37	0.54
10:A:76:A:N1	41:a:2422:C:C6	2.75	0.54
11:AA:243:PRO:HB2	11:AA:274:ILE:HG23	1.90	0.54
11:AA:887:VAL:HB	16:AG:105:ILE:CB	2.36	0.54
16:AG:220:VAL:HG12	16:AG:245:ILE:HG21	1.88	0.54
16:AG:374:VAL:O	16:AG:374:VAL:HG12	2.08	0.54
17:C:13:PHE:CZ	17:C:21:ILE:HG12	2.42	0.54
18:D:106:C:N4	19:E:10:ARG:NH2	2.56	0.54
18:D:1399:C:O2	18:D:1502:A:N6	2.40	0.54
22:H:9:PHE:O	22:H:13:LEU:HD23	2.06	0.54
27:M:31:MET:SD	27:M:36:LYS:HA	2.48	0.54
27:M:79:ARG:O	27:M:79:ARG:HD2	2.06	0.54
27:M:79:ARG:HB2	27:M:84:THR:HG23	1.90	0.54
28:N:31:LYS:HE3	28:N:31:LYS:HA	1.89	0.54
29:O:88:MET:HE1	29:O:95:ARG:HA	1.89	0.54
32:R:74:LEU:HD11	32:R:80:ILE:HD11	1.90	0.54
41:a:2038:G:H2'	41:a:2039:U:O4'	2.08	0.54
52:l:188:MET:HE2	52:l:193:VAL:HA	1.89	0.54
54:n:46:ASP:CG	54:n:49:LEU:HD13	2.33	0.54
58:r:71:LYS:HZ2	58:r:72:ILE:HA	1.72	0.54
4:3:48:PRO:HG3	4:3:56:GLY:HA3	1.88	0.54
9:9:94:ARG:CB	9:9:97:LYS:HZ3	2.20	0.54
11:AA:855:PRO:HD3	16:AG:103:ASP:OD2	2.08	0.54
13:AC:100:LEU:HD23	13:AC:115:ILE:HG21	1.89	0.54
14:AE:901:ARG:HH21	14:AE:906:GLY:HA2	1.73	0.54
18:D:564:C:P	32:R:12:ARG:HH22	2.30	0.54
18:D:695:A:OP1	31:Q:53:ARG:HD3	2.07	0.54
18:D:713:G:H2'	18:D:714:G:C8	2.42	0.54
21:G:18:HIS:HA	22:H:43:LYS:HD2	1.90	0.54
21:G:19:GLN:CD	22:H:76:GLU:HB3	2.33	0.54
23:I:164:ARG:CZ	23:I:166:GLU:OE2	2.54	0.54
25:K:131:THR:O	25:K:131:THR:HG22	2.08	0.54
27:M:71:PRO:HG3	27:M:103:TRP:CH2	2.42	0.54
28:N:78:VAL:HG12	28:N:85:ILE:HG21	1.89	0.54
41:a:2454:G:C4	41:a:2455:G:C8	2.96	0.54
41:a:2649:C:H2'	41:a:2650:U:H6	1.73	0.54
48:h:62:TYR:CD2	48:h:64:ILE:HD13	2.41	0.54
60:t:1:MET:CE	60:t:32:TYR:HB3	2.38	0.54
5:4:35:GLU:OE1	5:4:93:ARG:CZ	2.56	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:AE:1346:GLY:O	14:AE:1350:ASN:ND2	2.41	0.54
16:AG:168:LEU:CD2	16:AG:231:GLY:HA2	2.09	0.54
18:D:399:G:H2'	18:D:400:C:C6	2.43	0.54
19:E:25:ARG:HD3	19:E:66:LEU:HD21	1.90	0.54
21:G:187:VAL:HG21	21:G:199:VAL:HG23	1.89	0.54
25:K:86:LYS:N	25:K:86:LYS:HD2	2.23	0.54
38:X:28:THR:CA	38:X:31:LYS:HE3	2.37	0.54
41:a:851:C:H2'	41:a:852:U:H6	1.72	0.54
41:a:1000:A:H2'	41:a:1001:A:C8	2.43	0.54
41:a:1779:U:H5	41:a:1784:A:N7	2.06	0.54
50:j:4:LEU:HG	50:j:32:ASN:OD1	2.08	0.54
50:j:55:LYS:HZ2	50:j:60:VAL:HB	1.71	0.54
61:u:122:VAL:HG13	61:u:125:LEU:CD1	2.34	0.54
3:2:56:GLU:OE1	3:2:87:LEU:CA	2.56	0.54
5:4:6:ALA:HB3	5:4:65:VAL:CG2	2.37	0.54
9:9:3:LEU:HD12	9:9:4:ASN:CA	2.37	0.54
9:9:3:LEU:CG	9:9:4:ASN:H	2.21	0.54
10:A:41:C:O4'	27:M:143:ARG:NH1	2.41	0.54
11:AA:811:ASN:HA	11:AA:815:SER:HB2	1.90	0.54
11:AA:860:ALA:HB3	16:AG:38:LYS:HE2	1.89	0.54
12:AB:112:PRO:HB3	12:AB:132:GLU:OE1	2.07	0.54
14:AE:514:THR:HG21	14:AE:596:LEU:HD12	1.90	0.54
17:C:39:ILE:HD12	18:D:719:C:O2	2.07	0.54
17:C:60:LYS:HD2	18:D:734:G:O2'	2.07	0.54
18:D:37:U:O4	18:D:397:A:N1	2.40	0.54
18:D:73:C:O2	18:D:73:C:H2'	2.07	0.54
18:D:76:G:C4	18:D:77:A:C8	2.95	0.54
18:D:404:G:O2'	18:D:498:A:N1	2.38	0.54
18:D:736:C:H2'	18:D:737:C:C6	2.43	0.54
25:K:111:MET:HE2	25:K:125:ALA:HB1	1.90	0.54
26:L:3:HIS:CG	26:L:65:GLU:OE1	2.61	0.54
31:Q:20:VAL:HG13	31:Q:85:MET:HE1	1.90	0.54
33:S:27:LEU:HD21	33:S:48:LEU:N	2.21	0.54
38:X:81:MET:CE	38:X:92:ARG:HD3	2.38	0.54
41:a:251:A:OP1	61:u:58:TYR:OH	2.23	0.54
41:a:265:A:N1	41:a:427:U:O2'	2.40	0.54
41:a:1656:C:OP1	50:j:141:ARG:CD	2.55	0.54
41:a:1830:C:C2	41:a:1831:G:C8	2.95	0.54
48:h:67:PHE:CE1	48:h:105:LEU:HD11	2.42	0.54
50:j:124:ARG:NH1	50:j:161:MET:O	2.41	0.54
51:k:8:LYS:NZ	51:k:54:ILE:HG12	2.23	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:l:60:TRP:CD1	52:l:65:THR:HG21	2.43	0.54
52:l:175:ILE:CG2	52:l:180:LEU:HD11	2.38	0.54
62:v:1:MET:CE	62:v:44:ARG:HA	2.38	0.54
18:D:679:C:C2	18:D:680:C:C5	2.95	0.54
23:I:31:ASP:HA	33:S:65:ARG:HH22	1.72	0.54
30:P:27:GLU:HA	30:P:30:LYS:NZ	2.23	0.54
41:a:84:A:N1	41:a:98:G:O2'	2.37	0.54
41:a:1817:G:C6	41:a:1818:U:C5	2.95	0.54
41:a:2387:U:O4'	42:b:41:ARG:NH1	2.41	0.54
41:a:2537:U:H2'	41:a:2538:C:C6	2.43	0.54
59:s:133:ALA:C	59:s:135:GLN:H	2.16	0.54
62:v:96:ILE:HA	62:v:100:LYS:CE	2.38	0.54
64:x:48:LEU:CD2	64:x:87:ILE:HD11	2.38	0.54
2:l:41:LYS:HE3	41:a:2010:G:P	2.48	0.54
4:3:7:ARG:HB3	41:a:85:G:OP2	2.08	0.54
5:4:31:TYR:OH	44:d:103:U:O2'	2.20	0.54
5:4:80:HIS:ND1	5:4:83:LYS:HB2	2.23	0.54
7:6:14:DC:H2'	7:6:15:DC:C6	2.43	0.54
7:6:20:DA:N6	8:7:48:G:C2	2.76	0.54
8:7:6:U:C2	18:D:1493:A:C2	2.95	0.54
9:9:36:ASP:N	9:9:36:ASP:OD1	2.40	0.54
9:9:97:LYS:HD3	9:9:130:PRO:HG3	1.89	0.54
9:9:129:LEU:HD12	9:9:129:LEU:C	2.32	0.54
18:D:547:A:OP1	24:J:70:ARG:NH2	2.41	0.54
18:D:1358:U:O4	18:D:1363:A:N1	2.41	0.54
19:E:24:ARG:O	19:E:28:MET:HG2	2.08	0.54
19:E:25:ARG:HD3	19:E:66:LEU:HD11	1.90	0.54
19:E:32:ILE:CG2	19:E:75:HIS:NE2	2.71	0.54
21:G:58:ASN:OD1	21:G:223:GLU:OE1	2.26	0.54
25:K:77:ASN:HB2	25:K:82:GLN:HG3	1.89	0.54
28:N:83:LEU:HD21	32:R:4:VAL:HG11	1.89	0.54
35:U:23:ASP:HB3	35:U:26:ASN:OD1	2.07	0.54
41:a:1386:C:H2'	41:a:1387:A:C8	2.42	0.54
41:a:2259:U:C2	41:a:2260:C:C6	2.96	0.54
41:a:2297:A:C2	41:a:2320:U:O4'	2.61	0.54
41:a:2557:G:H2'	41:a:2558:C:C6	2.42	0.54
46:f:10:THR:HB	46:f:56:LYS:NZ	2.22	0.54
58:r:143:ILE:O	58:r:143:ILE:HG13	2.08	0.54
60:t:35:VAL:HG23	60:t:63:VAL:HA	1.89	0.54
60:t:73:ASP:OD2	65:y:80:VAL:HG11	2.08	0.54
62:v:45:GLN:NE2	62:v:125:PRO:HD2	2.22	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
63:w:33:ILE:CD1	63:w:112:TYR:CD1	2.91	0.54
5:4:19:ARG:NH1	44:d:95:U:OP2	2.41	0.53
11:AA:684:ASN:OD1	11:AA:687:ARG:NH2	2.41	0.53
12:AB:129:ILE:O	12:AB:142:LEU:CB	2.57	0.53
16:AG:299:ASP:HB2	16:AG:303:HIS:HA	1.89	0.53
16:AG:334:TRP:N	16:AG:334:TRP:CD1	2.73	0.53
22:H:279:LYS:HA	22:H:331:MET:HB3	1.90	0.53
27:M:4:ARG:CG	27:M:5:ARG:NH1	2.72	0.53
31:Q:84:VAL:CG2	31:Q:107:ILE:HD11	2.37	0.53
31:Q:84:VAL:HG21	31:Q:97:ILE:CD1	2.39	0.53
39:Y:78:LEU:HD13	39:Y:108:ILE:HG12	1.90	0.53
41:a:1019:U:O4	41:a:1142:A:N1	2.40	0.53
41:a:2064:C:H2'	41:a:2065:C:C6	2.42	0.53
41:a:2287:A:C8	41:a:2289:G:C8	2.96	0.53
45:e:4:LYS:HZ2	45:e:7:ARG:HH12	1.54	0.53
50:j:55:LYS:NZ	50:j:59:ARG:HB3	2.23	0.53
54:n:16:LEU:HD11	54:n:169:LEU:CD1	2.38	0.53
59:s:73:VAL:C	59:s:74:TYR:HD2	2.14	0.53
65:y:27:GLU:HB3	65:y:87:LYS:NZ	2.22	0.53
3:2:30:ILE:HG22	3:2:32:LEU:CD2	2.38	0.53
9:9:51:TYR:HB2	9:9:88:HIS:HB2	1.90	0.53
12:AB:27:ASN:ND2	12:AB:58:GLU:HB2	2.24	0.53
14:AE:1000:GLY:HA3	14:AE:1026:PRO:HG2	1.90	0.53
14:AE:1045:THR:O	14:AE:1067:ARG:NH2	2.41	0.53
16:AG:363:LEU:CG	16:AG:410:ALA:CA	2.81	0.53
10:B:75:C:O2'	10:B:76:A:C8	2.58	0.53
18:D:197:A:HI'	18:D:198:G:OP2	2.08	0.53
18:D:750:C:C2	18:D:751:U:C5	2.97	0.53
22:H:45:GLU:OE1	22:H:45:GLU:N	2.42	0.53
27:M:38:THR:O	27:M:42:ILE:HG12	2.08	0.53
32:R:109:ASP:C	32:R:111:LYS:HZ3	2.16	0.53
39:Y:77:VAL:O	39:Y:77:VAL:HG12	2.08	0.53
41:a:150:U:H2'	41:a:151:C:C6	2.44	0.53
41:a:1433:A:H2'	41:a:1434:A:O4'	2.07	0.53
41:a:1657:U:O2'	50:j:138:LEU:HD22	2.08	0.53
41:a:2747:G:O6	41:a:2755:C:H5''	2.08	0.53
48:h:74:ILE:CG2	48:h:96:TYR:HD1	2.21	0.53
52:l:119:ILE:O	52:l:188:MET:N	2.35	0.53
60:t:14:SER:OG	60:t:86:LEU:HD12	2.08	0.53
62:v:1:MET:HA	62:v:1:MET:CE	2.38	0.53
2:1:61:ASN:OD1	41:a:496:G:HI'	2.09	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:3:6:ARG:HG3	41:a:84:A:H3'	1.91	0.53
6:5:9:DG:H2'	12:AB:71:ALA:HA	1.90	0.53
11:AA:628:HIS:HB3	11:AA:647:ARG:HH21	1.73	0.53
11:AA:633:LEU:HD13	11:AA:644:LEU:HD23	1.89	0.53
12:AB:10:LYS:HB2	12:AB:73:ARG:HB3	1.90	0.53
14:AE:287:ALA:CB	14:AE:292:VAL:HG22	2.37	0.53
19:E:56:PRO:HG2	19:E:57:ILE:HD12	1.90	0.53
21:G:23:TRP:HZ3	21:G:25:PRO:CA	2.21	0.53
27:M:23:LEU:O	27:M:27:VAL:HG13	2.07	0.53
41:a:588:U:H2'	41:a:589:U:C6	2.43	0.53
41:a:2833:U:O2	50:j:58:ASN:ND2	2.41	0.53
50:j:4:LEU:HD23	50:j:29:VAL:HG11	1.89	0.53
50:j:48:ILE:HG12	50:j:84:LEU:HD21	1.90	0.53
52:l:147:LEU:HD11	52:l:170:ARG:HD2	1.89	0.53
60:t:2:ILE:CG2	60:t:8:LEU:HD11	2.38	0.53
66:z:92:ARG:HA	66:z:95:LEU:HD12	1.89	0.53
1:0:5:PHE:HB3	1:0:59:ILE:HD12	1.91	0.53
2:1:24:ILE:O	2:1:71:VAL:HG21	2.08	0.53
8:7:9:U:C4	18:D:1196:A:C8	2.97	0.53
13:AD:28:LEU:HD12	13:AD:201:LEU:HD23	1.91	0.53
16:AG:434:LEU:CG	16:AG:456:LEU:N	2.71	0.53
18:D:274:A:H4'	36:V:16:LYS:NZ	2.24	0.53
21:G:23:TRP:CD1	21:G:39:HIS:CE1	2.97	0.53
34:T:11:ILE:HA	34:T:14:GLU:OE1	2.09	0.53
41:a:78:U:H2'	41:a:79:C:C6	2.44	0.53
41:a:839:U:H2'	41:a:840:C:H6	1.73	0.53
41:a:1084:A:N7	41:a:1085:A:C6	2.76	0.53
41:a:1278:C:H5'	63:w:24:MET:SD	2.49	0.53
41:a:1818:U:O2'	48:h:153:GLN:O	2.23	0.53
62:v:97:GLN:H	62:v:100:LYS:CE	2.21	0.53
10:A:16:C:O2	41:a:2181:U:C5'	2.57	0.53
11:AA:377:THR:HB	12:AB:65:HIS:NE2	2.23	0.53
12:AB:54:TYR:HE2	14:AE:291:ILE:CG2	2.13	0.53
14:AE:968:ASN:HA	14:AE:1117:SER:HB2	1.90	0.53
16:AG:327:LEU:HD12	16:AG:327:LEU:O	2.07	0.53
16:AG:380:THR:C	16:AG:384:LEU:CD1	2.79	0.53
21:G:148:LEU:HD22	21:G:151:ILE:HD11	1.89	0.53
22:H:321:VAL:HG13	22:H:322:VAL:HG23	1.91	0.53
23:I:58:GLU:OE2	23:I:60:PRO:HD3	2.09	0.53
23:I:85:GLU:HG3	23:I:89:LYS:NZ	2.23	0.53
26:L:62:MET:HA	26:L:62:MET:HE3	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:M:103:TRP:HB3	27:M:137:LYS:HD3	1.89	0.53
28:N:87:LYS:HG2	28:N:92:LEU:HA	1.90	0.53
31:Q:79:ILE:O	31:Q:105:PHE:HE1	1.91	0.53
32:R:56:ARG:HA	32:R:62:GLU:OE2	2.08	0.53
39:Y:102:ARG:HG3	39:Y:140:GLU:C	2.33	0.53
41:a:372:G:O2'	41:a:400:G:O6	2.16	0.53
41:a:1412:U:C4	41:a:1413:A:N7	2.77	0.53
41:a:2867:G:C6	65:y:21:ARG:NH1	2.76	0.53
47:g:47:LYS:HD3	54:n:178:ARG:HH22	1.74	0.53
52:l:6:LYS:HZ2	52:l:119:ILE:HA	1.73	0.53
52:l:60:TRP:NE1	52:l:69:ARG:NH1	2.56	0.53
54:n:71:ARG:O	54:n:81:GLN:NE2	2.40	0.53
58:r:33:GLN:HB3	58:r:35:LYS:HZ3	1.74	0.53
3:2:6:ARG:NH2	3:2:10:VAL:HG22	2.24	0.53
5:4:35:GLU:OE1	5:4:36:ALA:O	2.27	0.53
12:AB:144:ASN:ND2	12:AB:149:GLU:CD	2.67	0.53
14:AE:526:VAL:HG12	14:AE:549:LYS:HB2	1.91	0.53
16:AG:362:TYR:CE1	16:AG:414:LEU:HD21	2.43	0.53
20:F:63:GLU:HG2	20:F:67:ARG:NH2	2.23	0.53
32:R:102:LEU:O	32:R:103:ASP:OD1	2.26	0.53
41:a:589:U:H2'	41:a:590:A:C8	2.42	0.53
41:a:1379:U:H4'	41:a:1380:G:OP1	2.09	0.53
41:a:2473:U:C4	41:a:2474:U:C5	2.97	0.53
54:n:42:GLU:HG3	54:n:49:LEU:HD23	1.91	0.53
60:t:99:ILE:HD13	60:t:118:LEU:CB	2.39	0.53
61:u:111:ILE:HD12	61:u:111:ILE:N	2.23	0.53
63:w:14:SER:HA	63:w:17:ARG:CZ	2.39	0.53
1:0:10:LYS:CE	41:a:996:A:OP1	2.56	0.53
11:AA:855:PRO:HD2	11:AA:887:VAL:HG11	1.91	0.53
12:AB:5:TYR:HA	12:AB:88:VAL:HG22	1.90	0.53
12:AB:141:LEU:HA	12:AB:152:HIS:CA	2.39	0.53
12:AB:147:ASN:ND2	30:P:100:ILE:CG2	2.70	0.53
16:AG:131:ARG:HA	16:AG:186:VAL:CG1	2.38	0.53
16:AG:402:THR:HA	16:AG:405:ALA:HB3	1.91	0.53
17:C:12:ARG:NH2	17:C:13:PHE:CE1	2.77	0.53
18:D:339:C:H2'	18:D:340:U:H6	1.73	0.53
18:D:1308:U:H3'	38:X:98:ARG:CZ	2.39	0.53
22:H:332:VAL:HG12	22:H:334:ASP:H	1.73	0.53
29:O:118:LEU:HD23	29:O:124:ARG:HA	1.91	0.53
32:R:25:GLU:OE1	32:R:59:ASN:ND2	2.42	0.53
32:R:35:THR:OG1	32:R:56:ARG:HG3	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:S:54:ASP:HA	33:S:59:ARG:HD2	1.90	0.53
39:Y:80:LYS:CG	39:Y:86:LYS:O	2.56	0.53
41:a:813:U:H2'	41:a:814:C:H6	1.73	0.53
41:a:1340:U:C5	41:a:1603:A:C8	2.97	0.53
56:p:32:GLU:C	56:p:33:LEU:HD12	2.34	0.53
61:u:84:LYS:NZ	61:u:98:ALA:O	2.20	0.53
63:w:44:LEU:HD22	63:w:113:ILE:HD13	1.90	0.53
4:3:69:ASN:ND2	41:a:329:G:OP2	2.37	0.53
14:AE:649:LYS:NZ	14:AE:652:GLU:OE1	2.37	0.53
18:D:842:U:H3'	18:D:843:U:C5'	2.39	0.53
20:F:17:ARG:HA	20:F:20:LYS:NZ	2.24	0.53
21:G:23:TRP:HB3	21:G:39:HIS:CE1	2.44	0.53
23:I:23:PHE:HD2	30:P:97:ASP:HB2	1.74	0.53
24:J:105:MET:HE1	24:J:171:LEU:HB3	1.89	0.53
26:L:39:LEU:O	26:L:39:LEU:HD12	2.07	0.53
28:N:50:LYS:HE2	28:N:50:LYS:HA	1.89	0.53
29:O:67:VAL:C	29:O:68:LYS:HD2	2.33	0.53
38:X:68:ASP:O	38:X:71:ARG:HG3	2.09	0.53
41:a:414:C:H2'	41:a:415:A:C8	2.44	0.53
41:a:1665:A:C3'	60:t:66:LYS:HZ2	2.21	0.53
41:a:2308:G:C6	54:n:77:PHE:CZ	2.97	0.53
41:a:2649:C:C2	41:a:2650:U:C5	2.97	0.53
52:l:62:GLN:O	52:l:69:ARG:CD	2.56	0.53
53:m:26:ASN:O	53:m:30:VAL:HG23	2.08	0.53
60:t:38:ILE:HD11	60:t:112:PHE:CZ	2.43	0.53
66:z:108:ALA:O	66:z:111:GLU:HG3	2.09	0.53
9:9:87:GLU:OE2	9:9:95:LEU:HG	2.09	0.53
11:AA:246:LEU:HB3	11:AA:269:ILE:HD13	1.91	0.53
11:AA:847:PRO:N	16:AG:104:ARG:NH2	2.57	0.53
12:AB:38:ILE:CD1	12:AB:160:ARG:HH11	2.21	0.53
13:AD:16:ILE:HG23	13:AD:26:VAL:HG22	1.91	0.53
14:AE:77:ARG:HD3	14:AE:78:LEU:H	1.73	0.53
18:D:932:C:C5'	27:M:4:ARG:HH11	2.22	0.53
29:O:21:ILE:HD11	29:O:87:LEU:HD22	1.90	0.53
29:O:47:VAL:O	29:O:80:ARG:HD2	2.09	0.53
32:R:24:LEU:C	32:R:25:GLU:OE1	2.52	0.53
32:R:110:ARG:NE	32:R:117:TYR:CD2	2.76	0.53
38:X:22:ILE:HB	38:X:25:VAL:CG1	2.39	0.53
41:a:590:A:H2'	41:a:591:U:C6	2.44	0.53
41:a:888:C:C4	41:a:889:C:C4	2.97	0.53
42:b:51:VAL:N	42:b:62:LYS:HZ3	2.07	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:k:29:THR:HG23	51:k:30:LYS:HG2	1.90	0.53
54:n:71:ARG:C	54:n:81:GLN:HE22	2.17	0.53
54:n:132:VAL:CG2	54:n:152:LEU:HD23	2.39	0.53
56:p:89:LEU:HD22	56:p:162:VAL:HG22	1.91	0.53
57:q:25:VAL:CB	57:q:35:GLN:OE1	2.56	0.53
1:0:10:LYS:CE	41:a:994:C:O2'	2.56	0.53
4:3:26:LYS:HE3	4:3:37:GLU:HB2	1.90	0.53
8:7:3:G:H3'	18:D:1500:A:N6	2.24	0.53
9:9:52:MET:HE1	9:9:85:SER:CA	2.38	0.53
9:9:136:ILE:HG13	9:9:136:ILE:O	2.09	0.53
12:AB:21:LEU:HB3	12:AB:26:VAL:CG1	2.38	0.53
14:AE:66:LYS:HB2	14:AE:69:GLU:CB	2.38	0.53
14:AE:1175:LEU:O	14:AE:1187:GLU:HA	2.08	0.53
10:B:38:A:O2'	18:D:790:A:P	2.66	0.53
18:D:62:U:H2'	18:D:63:C:C6	2.44	0.53
18:D:324:G:N1	18:D:327:A:OP2	2.39	0.53
24:J:61:VAL:HG21	24:J:200:ILE:HD11	1.90	0.53
24:J:74:ASN:O	24:J:77:LYS:HG2	2.09	0.53
25:K:137:VAL:O	25:K:141:ILE:HG12	2.09	0.53
25:K:158:GLY:C	25:K:159:LYS:HD2	2.34	0.53
28:N:87:LYS:HD2	28:N:91:GLU:CB	2.37	0.53
29:O:10:GLY:HA3	29:O:78:ALA:O	2.08	0.53
30:P:47:GLU:OE1	30:P:67:ILE:HG23	2.09	0.53
34:T:15:PHE:CD2	34:T:30:ALA:CB	2.92	0.53
37:W:66:MET:HG2	37:W:74:PHE:CZ	2.44	0.53
39:Y:32:VAL:HA	39:Y:60:VAL:HG21	1.90	0.53
41:a:607:U:C5	41:a:620:G:C5	2.97	0.53
41:a:1243:C:H1'	61:u:4:ASN:O	2.09	0.53
41:a:1801:A:OP2	48:h:146:MET:HE1	2.09	0.53
41:a:2849:U:O5'	65:y:93:ARG:NH2	2.42	0.53
59:s:9:GLU:HG2	59:s:10:THR:HG23	1.90	0.53
60:t:71:ARG:NH2	60:t:123:LEU:O	2.42	0.53
1:0:4:VAL:HG22	1:0:40:MET:HB3	1.91	0.52
3:2:33:LYS:HG3	3:2:80:TRP:CE3	2.45	0.52
9:9:59:LEU:HB2	9:9:62:ARG:HG3	1.92	0.52
9:9:94:ARG:HB2	9:9:97:LYS:CE	2.36	0.52
11:AA:786:GLY:N	11:AA:789:THR:OG1	2.41	0.52
21:G:26:LYS:HD2	21:G:193:PRO:CG	2.39	0.52
21:G:127:ASP:O	21:G:128:LYS:HB3	2.09	0.52
23:I:24:ALA:C	23:I:29:PHE:CE2	2.87	0.52
23:I:49:LYS:NZ	23:I:75:ILE:HD11	2.24	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:I:85:GLU:OE2	23:I:88:ARG:NH1	2.26	0.52
23:I:105:GLU:OE1	23:I:107:ARG:NE	2.42	0.52
25:K:134:ILE:HG23	25:K:135:ASN:N	2.24	0.52
27:M:85:TYR:CD2	27:M:151:PHE:CZ	2.97	0.52
31:Q:86:VAL:O	31:Q:113:VAL:HG22	2.08	0.52
38:X:13:LYS:HE3	38:X:17:ILE:HG21	1.91	0.52
40:Z:18:ASP:HB3	40:Z:22:LEU:CD1	2.39	0.52
41:a:1820:U:C4	48:h:159:GLY:HA3	2.44	0.52
41:a:2315:G:O2'	54:n:125:ARG:HD3	2.09	0.52
41:a:2356:U:H4'	42:b:20:ARG:HH11	1.73	0.52
41:a:2473:U:O2	41:a:2473:U:H2'	2.09	0.52
41:a:2847:U:H2'	41:a:2848:G:O4'	2.09	0.52
46:f:10:THR:HG23	46:f:11:ARG:HG2	1.90	0.52
52:l:62:GLN:HA	52:l:69:ARG:HH21	1.74	0.52
54:n:72:LYS:HA	54:n:81:GLN:HE22	1.74	0.52
61:u:58:TYR:CE2	61:u:59:ARG:HG2	2.44	0.52
11:AA:846:GLY:C	16:AG:104:ARG:NH2	2.65	0.52
12:AB:89:PRO:HG2	14:AE:290:ILE:HG22	1.91	0.52
14:AE:79:LYS:CG	16:AG:144:LYS:HD2	2.39	0.52
14:AE:342:LEU:HD23	14:AE:1352:ILE:HG23	1.90	0.52
16:AG:285:ILE:HG23	16:AG:293:VAL:HG11	1.82	0.52
18:D:58:C:O2	18:D:58:C:H2'	2.09	0.52
18:D:1175:G:N3	18:D:1176:A:C8	2.77	0.52
18:D:1329:A:P	38:X:28:THR:HB	2.48	0.52
21:G:53:ALA:CA	21:G:56:GLU:OE1	2.57	0.52
24:J:50:ASP:O	24:J:53:VAL:HG22	2.09	0.52
24:J:67:VAL:HB	24:J:72:PHE:CE1	2.43	0.52
41:a:67:U:N3	41:a:74:A:C6	2.77	0.52
41:a:372:G:O2'	43:c:54:LYS:HE3	2.09	0.52
41:a:581:C:H2'	41:a:582:A:H8	1.73	0.52
41:a:1167:C:C2	41:a:1168:G:C8	2.97	0.52
41:a:1266:G:O2'	41:a:2012:G:O6	2.22	0.52
41:a:2093:G:OP2	58:r:22:LYS:HE3	2.09	0.52
41:a:2356:U:C5'	42:b:20:ARG:HD2	2.36	0.52
48:h:34:LEU:HD12	48:h:61:ALA:HB1	1.91	0.52
59:s:25:LEU:CD1	59:s:101:ILE:HD13	2.39	0.52
62:v:20:LEU:H	62:v:20:LEU:HD23	1.73	0.52
11:AA:855:PRO:CB	16:AG:35:THR:HG1	2.19	0.52
11:AA:887:VAL:HG12	16:AG:105:ILE:HG23	1.88	0.52
13:AD:205:MET:HE3	13:AD:213:PRO:HB3	1.91	0.52
14:AE:122:SER:HB2	14:AE:132:LEU:HD12	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:233:ARG:C	16:AG:327:LEU:HD21	2.34	0.52
18:D:512:U:P	24:J:44:ARG:NH1	2.83	0.52
23:I:85:GLU:O	23:I:89:LYS:HE2	2.09	0.52
26:L:36:ILE:O	26:L:36:ILE:HG23	2.09	0.52
26:L:88:MET:HE2	26:L:90:MET:HE3	1.91	0.52
34:T:24:SER:O	34:T:28:GLN:HG2	2.09	0.52
39:Y:33:ASN:HB2	39:Y:36:GLU:OE2	2.08	0.52
39:Y:126:ARG:NH1	41:a:1083:U:P	2.83	0.52
41:a:271:G:C4	41:a:272:A:N7	2.77	0.52
41:a:1145:C:N3	41:a:1146:C:C5	2.77	0.52
41:a:1204:A:O4'	41:a:1206:G:C8	2.63	0.52
41:a:1656:C:OP1	50:j:141:ARG:HD3	2.10	0.52
41:a:1980:G:O2'	41:a:1982:U:OP2	2.22	0.52
41:a:2049:G:N2	50:j:161:MET:HE1	2.24	0.52
47:g:11:GLU:HA	47:g:25:ARG:HA	1.90	0.52
64:x:38:GLN:HG2	64:x:50:ALA:CB	2.38	0.52
8:7:8:U:OP2	18:D:1397:C:C2	2.62	0.52
12:AB:120:GLU:N	12:AB:162:LEU:HG	2.24	0.52
13:AD:100:LEU:HD21	13:AD:121:VAL:HG11	1.90	0.52
16:AG:21:GLU:HB3	16:AG:49:ILE:HD12	1.92	0.52
21:G:9:MET:HE1	21:G:47:VAL:CG2	2.40	0.52
22:H:32:ASP:OD1	22:H:35:VAL:CG2	2.57	0.52
29:O:112:GLU:HG3	29:O:115:LYS:HZ3	1.75	0.52
41:a:1145:C:C2	41:a:1146:C:C5	2.97	0.52
41:a:1314:C:C2	41:a:1315:C:C5	2.98	0.52
41:a:2355:G:H2'	41:a:2356:U:H5'	1.91	0.52
41:a:2812:G:H2'	41:a:2813:A:C8	2.45	0.52
50:j:131:ASP:OD1	50:j:134:HIS:HB2	2.10	0.52
53:m:25:LYS:HG3	53:m:29:GLN:HE22	1.73	0.52
54:n:93:GLY:HA2	54:n:97:TRP:CZ3	2.44	0.52
56:p:9:VAL:HG22	56:p:69:ARG:NH1	2.24	0.52
60:t:1:MET:CE	60:t:32:TYR:CB	2.87	0.52
60:t:34:GLY:N	60:t:37:ASP:OD2	2.39	0.52
61:u:14:LYS:HE2	61:u:14:LYS:HA	1.91	0.52
9:9:88:HIS:O	9:9:89:PRO:C	2.51	0.52
12:AB:144:ASN:HD22	12:AB:149:GLU:CD	2.17	0.52
14:AE:81:ARG:HH21	14:AE:81:ARG:CG	2.18	0.52
16:AG:11:ALA:HB3	16:AG:12:VAL:HA	1.92	0.52
16:AG:179:VAL:HB	16:AG:199:ARG:HH11	1.73	0.52
16:AG:285:ILE:CD1	16:AG:285:ILE:N	2.73	0.52
16:AG:329:SER:HA	16:AG:336:LEU:HG	1.92	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:679:C:H2'	18:D:680:C:H6	1.74	0.52
18:D:1086:U:C2	18:D:1087:G:C8	2.97	0.52
18:D:1240:U:C5	27:M:116:MET:HE2	2.44	0.52
21:G:26:LYS:HE3	21:G:194:ASP:OD1	2.10	0.52
23:I:142:MET:SD	23:I:170:GLU:OE1	2.67	0.52
23:I:183:ASP:HB3	23:I:204:LYS:NZ	2.25	0.52
26:L:42:TRP:CZ2	26:L:102:MET:SD	3.01	0.52
28:N:66:PHE:CD2	28:N:67:GLN:OE1	2.62	0.52
29:O:5:GLN:NE2	29:O:22:LYS:NZ	2.56	0.52
30:P:7:ARG:HD3	30:P:75:ASP:OD1	2.10	0.52
30:P:24:GLU:O	30:P:28:THR:HG23	2.10	0.52
34:T:63:ARG:HD2	34:T:67:LEU:CD2	2.40	0.52
41:a:537:G:P	59:s:2:LYS:HE2	2.49	0.52
41:a:560:C:O2	66:z:48:ARG:NH1	2.37	0.52
41:a:963:U:C2	41:a:964:C:C5	2.97	0.52
41:a:1138:G:C2	59:s:108:MET:HE2	2.45	0.52
41:a:1295:C:C2	41:a:1296:G:C8	2.97	0.52
41:a:1607:C:N4	41:a:1622:G:OP2	2.35	0.52
41:a:1666:G:OP1	60:t:66:LYS:HD3	2.10	0.52
41:a:1818:U:C6	48:h:156:ARG:NH2	2.77	0.52
50:j:152:PRO:CG	50:j:156:PHE:CZ	2.91	0.52
52:l:170:ARG:CZ	52:l:176:ASP:OD1	2.58	0.52
55:o:19:LYS:HE2	55:o:19:LYS:HA	1.91	0.52
63:w:37:THR:OG1	63:w:40:LYS:CD	2.57	0.52
12:AB:147:ASN:HD22	30:P:100:ILE:CG2	2.22	0.52
13:AC:43:LEU:HD13	13:AC:217:ILE:HD11	1.90	0.52
16:AG:365:ILE:HD11	16:AG:406:LEU:HD22	1.87	0.52
16:AG:444:LEU:HD22	16:AG:473:LEU:HD21	1.92	0.52
26:L:33:GLU:HG3	26:L:33:GLU:O	2.10	0.52
33:S:79:LEU:HB2	33:S:84:VAL:HG22	1.91	0.52
41:a:2758:A:H1'	56:p:38:ASN:HD21	1.74	0.52
52:l:60:TRP:NE1	52:l:69:ARG:HH12	2.07	0.52
6:5:17:DC:H5''	6:5:17:DC:H6	1.74	0.52
11:AA:855:PRO:HA	16:AG:35:THR:OG1	2.06	0.52
16:AG:213:VAL:HG22	16:AG:213:VAL:O	2.08	0.52
16:AG:282:GLN:NE2	16:AG:282:GLN:CA	2.73	0.52
16:AG:285:ILE:HD12	16:AG:293:VAL:HG21	1.91	0.52
16:AG:398:LEU:HG	16:AG:398:LEU:O	2.09	0.52
17:C:70:TYR:CE2	26:L:49:TYR:CZ	2.98	0.52
23:I:178:LEU:HD23	23:I:178:LEU:C	2.35	0.52
29:O:88:MET:HE1	29:O:95:ARG:HG3	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:V:7:THR:CB	36:V:60:GLU:OE2	2.56	0.52
38:X:71:ARG:HH12	47:g:52:ALA:HB3	1.75	0.52
39:Y:139:VAL:HG22	39:Y:140:GLU:N	2.25	0.52
41:a:1154:G:OP2	66:z:58:ARG:CZ	2.58	0.52
41:a:1198:U:C2	41:a:1199:U:C5	2.98	0.52
41:a:1348:C:C4	41:a:1349:C:C6	2.97	0.52
48:h:175:ARG:CG	48:h:181:MET:HE1	2.34	0.52
54:n:29:PRO:HB2	54:n:169:LEU:HD22	1.91	0.52
62:v:21:ALA:HB2	62:v:100:LYS:HG3	1.90	0.52
5:4:2:PHE:CE2	5:4:55:GLU:HG2	2.45	0.52
9:9:47:GLU:O	9:9:49:GLY:N	2.43	0.52
9:9:93:ALA:O	9:9:129:LEU:O	2.27	0.52
16:AG:355:ALA:HB1	16:AG:382:GLU:CB	2.36	0.52
16:AG:362:TYR:CE2	16:AG:382:GLU:HG2	2.45	0.52
18:D:537:G:OP1	32:R:110:ARG:NH2	2.38	0.52
18:D:769:G:H4'	18:D:1513:A:H4'	1.90	0.52
18:D:1227:A:H5'	38:X:110:LYS:HZ1	1.75	0.52
18:D:1439:G:P	19:E:33:LYS:NZ	2.82	0.52
30:P:22:THR:O	30:P:25:ILE:HG22	2.10	0.52
34:T:3:LEU:HD22	34:T:8:THR:HG23	1.92	0.52
34:T:64:ARG:NH2	34:T:89:ARG:HH22	2.07	0.52
37:W:47:LEU:O	37:W:62:VAL:HG23	2.08	0.52
38:X:96:PRO:CG	38:X:102:THR:HG22	2.40	0.52
41:a:833:A:H2'	41:a:834:G:C8	2.44	0.52
41:a:1681:G:H2'	41:a:1757:A:N1	2.25	0.52
48:h:76:ALA:HB1	48:h:94:VAL:CG1	2.39	0.52
56:p:149:ARG:CZ	56:p:154:PRO:HD3	2.40	0.52
8:7:3:G:C4'	18:D:1500:A:H62	2.22	0.52
13:AC:140:ILE:HD12	13:AC:142:MET:HE3	1.91	0.52
14:AE:78:LEU:C	16:AG:144:LYS:HZ1	2.18	0.52
14:AE:959:LYS:HB3	14:AE:983:LYS:HB2	1.91	0.52
10:B:38:A:O2'	18:D:790:A:O5'	2.27	0.52
18:D:501:C:H2'	18:D:502:A:H8	1.75	0.52
18:D:925:G:C2	18:D:927:G:C8	2.98	0.52
29:O:9:THR:HB	29:O:85:ARG:NH1	2.25	0.52
31:Q:61:PHE:CD1	31:Q:64:GLN:NE2	2.78	0.52
39:Y:58:ILE:HG23	39:Y:66:PHE:CD2	2.45	0.52
39:Y:77:VAL:HA	39:Y:80:LYS:HD3	1.91	0.52
39:Y:78:LEU:HD12	39:Y:112:LYS:NZ	2.25	0.52
41:a:1394:U:H4'	41:a:1603:A:H4'	1.92	0.52
41:a:2428:G:H21	61:u:60:ARG:CZ	2.23	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:d:15:A:O4'	44:d:109:A:C8	2.63	0.52
49:i:26:THR:HG23	49:i:27:SER:N	2.25	0.52
51:k:19:HIS:CE1	51:k:41:PRO:HD3	2.45	0.52
62:v:115:GLU:HA	62:v:118:LYS:NZ	2.25	0.52
8:7:13:U:C6	8:7:13:U:H5''	2.45	0.52
9:9:78:GLY:H	9:9:79:PRO:HD2	1.75	0.52
10:A:31:G:H2'	10:A:32:C:O4'	2.10	0.52
11:AA:714:VAL:HB	11:AA:787:PRO:HD2	1.92	0.52
11:AA:974:ARG:HD2	11:AA:1014:LEU:HD21	1.92	0.52
16:AG:233:ARG:NH1	16:AG:324:ASN:HB2	2.25	0.52
16:AG:399:ASP:OD2	16:AG:399:ASP:N	2.40	0.52
10:B:48:C:C5	10:B:59:A:C8	2.98	0.52
18:D:512:U:P	24:J:44:ARG:HH12	2.34	0.52
20:F:37:PHE:HB3	31:Q:124:PRO:O	2.10	0.52
21:G:49:MET:CE	21:G:201:PRO:CD	2.87	0.52
22:H:48:ILE:HG22	22:H:49:PRO:O	2.10	0.52
27:M:111:ARG:N	27:M:119:ARG:NH2	2.58	0.52
29:O:123:ARG:O	29:O:123:ARG:CG	2.57	0.52
31:Q:52:PHE:CZ	31:Q:65:VAL:HG21	2.45	0.52
34:T:3:LEU:HD12	34:T:35:GLN:OE1	2.10	0.52
39:Y:9:LYS:CG	39:Y:55:PRO:HB3	2.39	0.52
39:Y:116:MET:HG2	39:Y:124:MET:HB3	1.91	0.52
39:Y:126:ARG:NH1	41:a:1082:U:C5'	2.73	0.52
41:a:414:C:H2'	41:a:415:A:H8	1.75	0.52
41:a:617:G:C4'	52:l:199:MET:HE1	2.40	0.52
41:a:632:A:H2'	41:a:633:A:C8	2.44	0.52
41:a:743:A:O2'	41:a:1659:G:OP1	2.25	0.52
41:a:1707:G:C8	41:a:1756:G:C5	2.98	0.52
41:a:1996:C:H4'	50:j:128:ARG:HH12	1.75	0.52
58:r:2:GLN:HG3	58:r:39:ALA:CB	2.40	0.52
59:s:15:TRP:CE3	59:s:135:GLN:O	2.63	0.52
62:v:45:GLN:HE21	62:v:124:LEU:HA	1.75	0.52
4:3:85:PHE:CE1	4:3:94:ARG:HG2	2.46	0.51
10:A:56:C:C5	41:a:2168:G:C6	2.98	0.51
12:AB:155:LYS:HG3	12:AB:156:ASN:N	2.25	0.51
18:D:564:C:H5'	36:V:34:TYR:HE1	1.75	0.51
21:G:90:PHE:CD2	21:G:154:MET:CE	2.94	0.51
22:H:36:VAL:O	22:H:48:ILE:HB	2.10	0.51
28:N:79:SER:HB2	28:N:85:ILE:HG22	1.92	0.51
29:O:30:ILE:HD13	29:O:39:PHE:HE2	1.75	0.51
35:U:76:LYS:C	35:U:80:LYS:HZ3	2.17	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:Y:16:MET:HG2	39:Y:16:MET:O	2.09	0.51
41:a:69:C:O2	41:a:73:A:O2'	2.26	0.51
41:a:645:C:H2'	41:a:647:G:C8	2.45	0.51
41:a:872:U:H2'	41:a:873:C:H6	1.75	0.51
41:a:2093:G:O2'	41:a:2198:A:N1	2.41	0.51
41:a:2311:A:C5	54:n:77:PHE:CD2	2.98	0.51
42:b:21:LEU:HD21	42:b:41:ARG:NE	2.20	0.51
60:t:17:ARG:NH2	60:t:47:ILE:HG23	2.25	0.51
66:z:74:ILE:HG13	66:z:78:LYS:CE	2.40	0.51
2:1:27:LYS:O	2:1:71:VAL:HG22	2.10	0.51
5:4:48:MET:CE	5:4:86:LEU:HG	2.40	0.51
10:A:6:G:C4	10:A:7:G:C8	2.98	0.51
11:AA:524:ILE:HG21	11:AA:708:VAL:HG13	1.92	0.51
11:AA:1070:HIS:NE2	11:AA:1114:GLU:OE1	2.36	0.51
11:AA:1287:LEU:HD13	14:AE:1357:ILE:HD11	1.91	0.51
14:AE:71:LEU:HD22	14:AE:90:VAL:HG21	1.93	0.51
14:AE:1179:PRO:HB2	14:AE:1182:GLY:H	1.75	0.51
16:AG:363:LEU:HD11	16:AG:410:ALA:HB2	1.92	0.51
10:B:6:G:C4	10:B:7:G:C8	2.98	0.51
18:D:934:C:C4	18:D:1345:U:C5	2.99	0.51
25:K:36:LEU:HD13	25:K:134:ILE:CD1	2.41	0.51
25:K:62:LYS:HA	25:K:62:LYS:HE3	1.93	0.51
33:S:87:ALA:N	33:S:90:ARG:NH1	2.58	0.51
35:U:39:PHE:CD1	35:U:50:THR:HG22	2.45	0.51
37:W:41:PHE:O	37:W:44:MET:HG3	2.11	0.51
41:a:65:U:H2'	41:a:66:C:H6	1.75	0.51
41:a:1024:G:OP2	41:a:1025:G:O2'	2.25	0.51
41:a:2331:G:O2'	41:a:2336:A:N1	2.40	0.51
41:a:2420:C:H5''	51:k:8:LYS:HE2	1.92	0.51
48:h:18:LYS:HE3	48:h:20:VAL:CG2	2.41	0.51
3:2:53:VAL:HG21	3:2:87:LEU:HD23	1.91	0.51
8:7:2:U:H2'	8:7:3:G:C8	2.45	0.51
9:9:61:ARG:NE	41:a:1046:A:P	2.83	0.51
11:AA:255:ILE:HB	11:AA:263:VAL:HB	1.92	0.51
14:AE:795:TYR:OH	14:AE:1326:GLN:NE2	2.42	0.51
16:AG:9:VAL:HG22	16:AG:24:PHE:CE2	2.44	0.51
17:C:60:LYS:CE	18:D:735:C:C4'	2.88	0.51
18:D:936:C:C2	18:D:937:A:C8	2.98	0.51
18:D:1371:G:O3'	29:O:71:GLY:HA3	2.10	0.51
26:L:11:HIS:HB3	26:L:14:GLN:OE1	2.11	0.51
26:L:67:PRO:O	26:L:70:VAL:HG22	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:T:5:THR:HG23	34:T:6:GLU:N	2.25	0.51
41:a:139:U:OP2	41:a:140:C:N4	2.43	0.51
41:a:628:G:C5	41:a:636:G:N2	2.79	0.51
41:a:995:C:N4	59:s:2:LYS:HG3	2.24	0.51
41:a:1853:A:H2'	41:a:1854:A:C8	2.46	0.51
41:a:2329:U:H2'	41:a:2330:G:C8	2.45	0.51
41:a:2813:A:C4	41:a:2814:A:C8	2.99	0.51
47:g:47:LYS:HE2	54:n:115:ARG:NE	2.25	0.51
54:n:136:ILE:HD12	54:n:141:ILE:HG23	1.92	0.51
60:t:2:ILE:HG23	60:t:8:LEU:HD11	1.91	0.51
62:v:53:MET:HE3	62:v:120:ALA:CB	2.40	0.51
1:0:17:GLY:HA2	1:0:97:LYS:HE2	1.92	0.51
3:2:10:VAL:HG12	3:2:11:LEU:HD12	1.93	0.51
6:5:9:DG:OP2	11:AA:470:ARG:O	2.28	0.51
7:6:17:DC:H42	8:7:50:G:H1	1.58	0.51
8:7:13:U:H5''	8:7:13:U:H6	1.73	0.51
11:AA:860:ALA:N	16:AG:38:LYS:CD	2.62	0.51
16:AG:353:HIS:O	16:AG:356:ILE:CB	2.58	0.51
17:C:12:ARG:HG3	22:H:264:GLU:CB	2.41	0.51
18:D:868:C:H2'	18:D:869:G:O4'	2.11	0.51
19:E:75:HIS:CE1	19:E:79:LEU:CD1	2.93	0.51
22:H:279:LYS:HA	22:H:331:MET:CB	2.41	0.51
22:H:304:VAL:HG11	22:H:346:LEU:HB2	1.93	0.51
26:L:70:VAL:O	26:L:74:LEU:CD2	2.56	0.51
27:M:111:ARG:NH2	27:M:126:ASP:OD2	2.43	0.51
31:Q:23:ILE:N	31:Q:87:LYS:HZ1	2.08	0.51
41:a:413:C:C2	41:a:414:C:C5	2.99	0.51
41:a:1346:G:C6	41:a:1601:G:C6	2.98	0.51
41:a:1791:A:C2	41:a:1829:A:C4'	2.94	0.51
41:a:1996:C:H5''	50:j:128:ARG:NH1	2.25	0.51
41:a:2086:U:H2'	41:a:2087:G:C8	2.46	0.51
41:a:2646:C:OP2	41:a:2732:G:O2'	2.25	0.51
48:h:124:ILE:O	48:h:125:LYS:HD3	2.10	0.51
51:k:21:TYR:CE2	51:k:49:TYR:CE2	2.99	0.51
51:k:21:TYR:CE2	51:k:49:TYR:CD2	2.98	0.51
58:r:29:PHE:CE2	58:r:35:LYS:HE3	2.45	0.51
62:v:97:GLN:N	62:v:100:LYS:CE	2.72	0.51
64:x:53:THR:HG23	64:x:74:VAL:HG21	1.92	0.51
66:z:53:ARG:HG3	66:z:57:PHE:HE2	1.75	0.51
8:7:43:U:O2	8:7:43:U:H2'	2.11	0.51
11:AA:125:GLY:HA2	11:AA:499:SER:HB2	1.92	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AB:135:GLY:C	12:AB:137:ALA:H	2.17	0.51
16:AG:244:ARG:CD	16:AG:244:ARG:N	2.73	0.51
18:D:564:C:H5'	36:V:34:TYR:CE1	2.46	0.51
18:D:728:A:H2'	18:D:729:A:C8	2.46	0.51
21:G:168:HIS:HB3	21:G:169:GLU:OE1	2.10	0.51
29:O:57:MET:HA	29:O:60:LYS:CE	2.40	0.51
41:a:1666:G:P	60:t:66:LYS:HD3	2.50	0.51
41:a:1824:G:O2'	48:h:252:THR:HG21	2.11	0.51
41:a:2093:G:OP2	58:r:22:LYS:HG2	2.11	0.51
41:a:2133:G:O2'	41:a:2157:G:N2	2.44	0.51
41:a:2387:U:H4'	42:b:41:ARG:HH12	1.76	0.51
50:j:131:ASP:OD2	50:j:132:ALA:N	2.42	0.51
52:l:168:ASP:HB3	52:l:183:PHE:HE2	1.75	0.51
54:n:5:HIS:HD2	54:n:97:TRP:CZ2	2.29	0.51
61:u:56:PRO:O	61:u:60:ARG:HG3	2.10	0.51
62:v:112:LEU:HA	62:v:115:GLU:OE1	2.10	0.51
62:v:115:GLU:O	62:v:119:LEU:HD23	2.10	0.51
63:w:35:LYS:NZ	63:w:112:TYR:CE2	2.76	0.51
66:z:65:ILE:HB	66:z:76:TYR:CE1	2.45	0.51
4:3:66:GLN:OE1	4:3:69:ASN:ND2	2.44	0.51
5:4:2:PHE:CE1	5:4:56:PHE:CE1	2.99	0.51
5:4:63:ILE:CD1	5:4:91:PHE:CD2	2.93	0.51
9:9:55:VAL:CG1	9:9:56:ARG:N	2.74	0.51
10:A:48:C:C5	10:A:59:A:C8	2.98	0.51
11:AA:853:ASP:O	16:AG:103:ASP:OD2	2.29	0.51
12:AB:78:PHE:CG	12:AB:85:PRO:HB3	2.46	0.51
12:AB:81:PHE:CB	14:AE:141:PHE:O	2.58	0.51
14:AE:287:ALA:HB2	14:AE:292:VAL:HG11	1.85	0.51
16:AG:63:LEU:HA	16:AG:92:TYR:HA	1.91	0.51
18:D:335:C:C2	18:D:336:A:C8	2.99	0.51
18:D:977:A:O2'	18:D:979:C:OP2	2.25	0.51
20:F:25:LYS:NZ	22:H:336:ASP:CG	2.68	0.51
22:H:342:ILE:HG12	22:H:344:LEU:HD12	1.93	0.51
24:J:74:ASN:CA	24:J:77:LYS:HZ3	2.23	0.51
30:P:15:HIS:HB3	30:P:70:HIS:CE1	2.46	0.51
33:S:10:GLU:OE2	33:S:63:ARG:HD2	2.11	0.51
34:T:3:LEU:HD23	34:T:3:LEU:C	2.36	0.51
38:X:31:LYS:HG3	38:X:32:ALA:N	2.25	0.51
40:Z:15:SER:O	40:Z:19:VAL:HG23	2.11	0.51
41:a:756:A:H2'	41:a:757:G:O4'	2.11	0.51
41:a:2391:G:P	55:o:32:ILE:HD12	2.50	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:h:52:ARG:NE	48:h:53:HIS:HE1	2.07	0.51
48:h:108:LYS:HD3	48:h:194:GLU:HB2	1.92	0.51
50:j:68:PHE:CD2	50:j:75:ALA:HA	2.46	0.51
50:j:154:LYS:HE3	50:j:156:PHE:CE1	2.46	0.51
2:1:25:ARG:NH1	41:a:519:U:O3'	2.43	0.51
5:4:50:MET:CA	5:4:56:PHE:CZ	2.94	0.51
7:6:19:DC:H2'	7:6:20:DA:H8	1.74	0.51
9:9:106:PHE:CG	9:9:106:PHE:O	2.63	0.51
18:D:21:G:C2	18:D:22:G:C5	2.99	0.51
18:D:975:A:H8	18:D:1357:A:HO2'	1.57	0.51
18:D:978:A:C4	18:D:1319:A:C2	2.98	0.51
21:G:38:VAL:HG22	22:H:75:VAL:HG21	1.92	0.51
21:G:187:VAL:HG13	21:G:191:SER:HB2	1.93	0.51
30:P:73:LEU:HD21	30:P:75:ASP:OD2	2.10	0.51
31:Q:27:PHE:CE2	31:Q:89:PRO:HG2	2.46	0.51
32:R:3:THR:O	32:R:6:GLN:N	2.44	0.51
36:V:62:ARG:CA	36:V:73:TRP:CZ3	2.93	0.51
37:W:45:ILE:HA	37:W:62:VAL:HG11	1.93	0.51
41:a:851:C:H2'	41:a:852:U:C6	2.46	0.51
41:a:2006:C:C2	41:a:2007:U:C5	2.99	0.51
41:a:2305:U:H4'	54:n:133:ARG:NH2	2.26	0.51
41:a:2308:G:C5	54:n:77:PHE:HZ	2.26	0.51
41:a:2466:C:N3	41:a:2467:C:C5	2.79	0.51
41:a:2727:A:O2'	60:t:70:ARG:NH2	2.43	0.51
54:n:30:ARG:HB2	54:n:30:ARG:NH2	2.25	0.51
56:p:162:VAL:O	56:p:162:VAL:HG12	2.11	0.51
59:s:31:GLU:HG2	59:s:142:ILE:HG12	1.92	0.51
62:v:105:MET:HE2	62:v:108:VAL:CG2	2.40	0.51
63:w:58:ASP:CG	63:w:80:PHE:CZ	2.89	0.51
1:0:37:GLU:O	1:0:37:GLU:HG3	2.11	0.51
2:1:31:GLN:OE1	2:1:31:GLN:N	2.29	0.51
9:9:98:GLU:HA	9:9:101:LYS:HZ3	1.73	0.51
14:AE:510:LEU:HD22	14:AE:601:ILE:HD12	1.91	0.51
10:B:23:C:H2'	10:B:24:U:C6	2.45	0.51
10:B:76:A:H1'	41:a:2494:G:P	2.50	0.51
18:D:452:A:H2'	18:D:453:G:O4'	2.10	0.51
18:D:767:A:H2'	18:D:768:A:O4'	2.11	0.51
18:D:1106:G:O3'	23:I:172:ARG:HG3	2.11	0.51
23:I:108:LYS:N	23:I:108:LYS:HD2	2.26	0.51
39:Y:41:PHE:O	39:Y:45:THR:OG1	2.22	0.51
39:Y:45:THR:HG21	39:Y:50:LYS:HZ2	1.74	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:Z:26:MET:HA	40:Z:30:PHE:CD1	2.46	0.51
41:a:642:U:O2'	41:a:644:A:N7	2.43	0.51
41:a:1064:C:O2'	41:a:1065:U:C6	2.62	0.51
41:a:2588:G:O6	41:a:2607:G:C6	2.64	0.51
41:a:2884:U:H3	49:i:40:ARG:NH1	2.09	0.51
45:e:20:ASN:HA	45:e:23:ARG:HE	1.76	0.51
54:n:15:LYS:O	54:n:18:THR:N	2.43	0.51
55:o:31:HIS:O	55:o:32:ILE:C	2.54	0.51
59:s:30:THR:HG23	59:s:31:GLU:N	2.26	0.51
59:s:72:LYS:HD3	59:s:74:TYR:CZ	2.43	0.51
61:u:54:GLN:NE2	61:u:60:ARG:HH12	2.07	0.51
61:u:58:TYR:CD2	61:u:59:ARG:HG2	2.46	0.51
9:9:10:ALA:O	9:9:14:GLU:HG3	2.11	0.51
10:A:19:G:N2	41:a:2112:G:O4'	2.44	0.51
11:AA:232:ILE:HG12	11:AA:237:LEU:HG	1.93	0.51
12:AB:141:LEU:HA	12:AB:152:HIS:HA	1.92	0.51
14:AE:67:ASP:C	14:AE:69:GLU:H	2.18	0.51
14:AE:209:ASN:HB3	14:AE:214:ARG:HH21	1.76	0.51
18:D:500:G:H2'	18:D:501:C:C6	2.46	0.51
18:D:1226:C:C2'	38:X:102:THR:OG1	2.58	0.51
20:F:25:LYS:HE3	22:H:336:ASP:OD2	2.11	0.51
23:I:132:ARG:HG3	23:I:135:LYS:HE3	1.92	0.51
27:M:71:PRO:HG3	27:M:103:TRP:HH2	1.76	0.51
32:R:62:GLU:OE2	32:R:62:GLU:HA	2.10	0.51
41:a:1095:A:H2'	41:a:1096:A:C8	2.45	0.51
47:g:7:PRO:HD2	54:n:62:GLY:O	2.11	0.51
65:y:9:GLU:CG	65:y:55:LEU:HB2	2.41	0.51
65:y:106:LYS:HD3	65:y:109:ARG:NH2	2.26	0.51
14:AE:430:HIS:HB3	14:AE:925:GLU:HG3	1.93	0.51
10:B:31:G:H2'	10:B:32:C:O4'	2.10	0.51
18:D:356:A:N3	18:D:368:U:O2'	2.37	0.51
18:D:1158:C:O2'	21:G:132:LYS:HE2	2.09	0.51
18:D:1492:A:O2'	18:D:1493:A:H5''	2.11	0.51
22:H:335:ILE:HG12	22:H:342:ILE:HD12	1.92	0.51
23:I:142:MET:HE3	23:I:170:GLU:C	2.36	0.51
25:K:156:LYS:NZ	28:N:71:VAL:O	2.42	0.51
27:M:35:LYS:HE3	27:M:38:THR:OG1	2.11	0.51
29:O:7:TYR:H	29:O:89:GLU:CD	2.19	0.51
31:Q:63:ALA:HB1	31:Q:96:THR:CG2	2.41	0.51
32:R:54:ARG:HH11	32:R:62:GLU:HG3	1.76	0.51
34:T:11:ILE:HD11	34:T:30:ALA:C	2.36	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:T:76:ALA:C	34:T:80:GLN:NE2	2.68	0.51
39:Y:126:ARG:HD2	39:Y:126:ARG:C	2.36	0.51
40:Z:18:ASP:HB3	40:Z:22:LEU:HD12	1.93	0.51
41:a:372:G:C8	43:c:61:LYS:HD2	2.46	0.51
41:a:580:U:H2'	41:a:581:C:C6	2.46	0.51
41:a:626:A:N3	61:u:78:ARG:CZ	2.74	0.51
41:a:703:U:H2'	41:a:704:G:O4'	2.10	0.51
41:a:910:A:C8	62:v:12:MET:CE	2.93	0.51
41:a:1453:A:C8	63:w:73:ASN:OD1	2.64	0.51
41:a:1956:U:C4	41:a:1957:C:C5	2.99	0.51
41:a:2259:U:C6	41:a:2427:C:C4	2.99	0.51
41:a:2682:A:H8	50:j:11:MET:CE	2.15	0.51
44:d:42:C:P	54:n:64:LYS:NZ	2.83	0.51
56:p:24:ILE:HG22	56:p:35:ARG:O	2.11	0.51
59:s:15:TRP:CH2	59:s:135:GLN:HG3	2.46	0.51
59:s:84:ILE:HG23	59:s:84:ILE:O	2.10	0.51
59:s:98:GLU:O	59:s:99:ARG:C	2.53	0.51
63:w:103:ARG:HA	63:w:110:MET:HE2	1.93	0.51
66:z:65:ILE:HB	66:z:76:TYR:HE1	1.75	0.51
5:4:62:THR:OG1	5:4:71:LYS:HE3	2.11	0.50
6:5:10:DT:N1	11:AA:62:TYR:HE2	2.07	0.50
10:A:56:C:H2'	10:A:57:A:H5''	1.93	0.50
11:AA:93:SER:HA	11:AA:128:PRO:HA	1.93	0.50
16:AG:216:ILE:HG12	16:AG:221:ILE:HB	1.93	0.50
16:AG:453:VAL:HG11	16:AG:459:LEU:CD2	2.41	0.50
17:C:71:THR:H	17:C:74:HIS:HE1	1.58	0.50
17:C:73:ARG:HH12	31:Q:113:VAL:HB	1.76	0.50
18:D:107:G:C5	19:E:10:ARG:NH2	2.79	0.50
18:D:107:G:C6	19:E:10:ARG:NH2	2.79	0.50
18:D:122:G:C5	18:D:123:U:C5	2.98	0.50
18:D:276:G:C5'	36:V:17:MET:CE	2.89	0.50
36:V:15:ASP:OD1	36:V:55:ILE:HD11	2.11	0.50
39:Y:86:LYS:HD3	39:Y:86:LYS:O	2.11	0.50
41:a:125:A:C5	53:m:10:LEU:HD13	2.46	0.50
41:a:1318:U:C2	41:a:1319:C:C5	2.99	0.50
41:a:1945:G:C4	41:a:1946:U:C5	3.00	0.50
43:c:71:LEU:HB3	43:c:76:GLU:CG	2.41	0.50
50:j:122:VAL:HG21	50:j:141:ARG:HG3	1.92	0.50
59:s:35:ARG:HA	59:s:40:HIS:CE1	2.46	0.50
60:t:114:LYS:O	60:t:118:LEU:HD23	2.11	0.50
63:w:12:ARG:NE	63:w:20:MET:HE1	2.26	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:9:43:LYS:O	9:9:47:GLU:OE1	2.28	0.50
11:AA:856:ASN:O	16:AG:37:LYS:HB2	2.11	0.50
12:AB:27:ASN:OD1	12:AB:58:GLU:HB2	2.11	0.50
15:AF:3:ARG:NH1	15:AF:55:GLU:OE2	2.40	0.50
18:D:678:U:H2'	18:D:679:C:H6	1.76	0.50
18:D:1176:A:H2'	18:D:1177:G:C8	2.45	0.50
19:E:54:MET:CE	19:E:75:HIS:HE1	2.24	0.50
21:G:109:GLN:C	21:G:113:ARG:HE	2.19	0.50
26:L:9:MET:O	26:L:84:VAL:HA	2.12	0.50
30:P:24:GLU:CA	30:P:27:GLU:OE1	2.59	0.50
30:P:66:GLU:CG	33:S:99:ALA:HB2	2.41	0.50
41:a:4:U:C2	41:a:5:A:C8	3.00	0.50
55:o:25:LYS:HB3	61:u:62:PRO:CG	2.40	0.50
59:s:13:ARG:HH22	59:s:49:ASP:C	2.20	0.50
59:s:15:TRP:CD1	59:s:53:TYR:HB2	2.46	0.50
63:w:82:GLU:OE1	63:w:82:GLU:N	2.44	0.50
66:z:82:GLY:HA3	66:z:117:LEU:CD1	2.41	0.50
2:1:36:LEU:HD23	2:1:48:LYS:HB2	1.94	0.50
6:5:16:DG:O6	11:AA:451:ARG:NH1	2.45	0.50
14:AE:157:GLN:HE22	14:AE:188:LEU:HD22	1.77	0.50
14:AE:693:VAL:HG21	14:AE:743:MET:HE3	1.93	0.50
16:AG:12:VAL:HG23	16:AG:16:LYS:HE2	1.92	0.50
17:C:24:LYS:HB3	26:L:102:MET:HE1	1.93	0.50
18:D:657:U:C1'	34:T:28:GLN:HE22	2.24	0.50
18:D:718:A:H5'	31:Q:119:ASN:HD22	1.76	0.50
18:D:958:A:O4'	37:W:55:ARG:NH1	2.45	0.50
21:G:17:GLY:HA3	21:G:41:ILE:HD11	1.92	0.50
25:K:111:MET:O	25:K:114:VAL:HG12	2.11	0.50
39:Y:57:VAL:HG23	39:Y:71:LYS:HD3	1.93	0.50
41:a:16:C:O4'	49:i:15:MET:HE3	2.12	0.50
41:a:44:A:H2'	41:a:45:G:O4'	2.12	0.50
41:a:532:A:H4'	41:a:533:G:C8	2.46	0.50
41:a:993:G:C6	41:a:1162:G:C6	2.99	0.50
41:a:2063:C:C2	41:a:2064:C:C6	2.99	0.50
49:i:40:ARG:NH1	49:i:41:HIS:CD2	2.80	0.50
51:k:8:LYS:HZ1	51:k:54:ILE:HG12	1.76	0.50
59:s:123:LYS:CE	59:s:132:HIS:HD2	2.25	0.50
62:v:105:MET:SD	62:v:113:ALA:HB1	2.51	0.50
63:w:58:ASP:OD1	63:w:63:ARG:NH2	2.44	0.50
3:2:11:LEU:HD12	3:2:11:LEU:N	2.26	0.50
12:AB:50:LEU:HD21	14:AE:285:LEU:HB3	1.94	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:AE:287:ALA:HB3	14:AE:292:VAL:HG22	1.94	0.50
16:AG:422:GLU:O	16:AG:427:ASP:N	2.45	0.50
18:D:555:U:H2'	18:D:556:C:C6	2.47	0.50
21:G:16:PHE:HB3	22:H:43:LYS:HA	1.93	0.50
22:H:273:ARG:NH1	22:H:297:GLU:HB2	2.25	0.50
23:I:142:MET:CE	23:I:170:GLU:OE1	2.58	0.50
26:L:4:TYR:OH	26:L:68:GLN:HG2	2.11	0.50
26:L:45:ARG:HA	26:L:45:ARG:NE	2.26	0.50
29:O:88:MET:SD	29:O:95:ARG:CZ	2.99	0.50
31:Q:87:LYS:CG	31:Q:113:VAL:CG2	2.89	0.50
41:a:813:U:H2'	41:a:814:C:C6	2.46	0.50
41:a:1278:C:O2'	63:w:27:SER:OG	2.16	0.50
41:a:2814:A:H2'	41:a:2815:C:H6	1.75	0.50
42:b:55:ARG:HA	42:b:55:ARG:CZ	2.41	0.50
43:c:65:ASP:CG	43:c:66:THR:N	2.70	0.50
49:i:30:VAL:HG22	49:i:37:LYS:CD	2.41	0.50
54:n:8:TYR:HE2	54:n:30:ARG:HE	1.59	0.50
59:s:74:TYR:N	59:s:74:TYR:HD2	2.10	0.50
60:t:7:MET:SD	60:t:18:ARG:NE	2.85	0.50
62:v:31:PHE:O	62:v:104:GLU:CD	2.54	0.50
62:v:39:GLY:O	62:v:96:ILE:N	2.33	0.50
62:v:50:ARG:CA	62:v:53:MET:SD	3.00	0.50
63:w:24:MET:HE3	63:w:24:MET:CA	2.34	0.50
1:0:71:LYS:HG3	1:0:73:LYS:HE3	1.92	0.50
8:7:12:U:H5''	25:K:56:VAL:HG21	1.93	0.50
9:9:87:GLU:OE2	9:9:95:LEU:HB2	2.12	0.50
10:A:15:G:H2'	10:A:15:G:N3	2.27	0.50
11:AA:400:VAL:HG11	11:AA:452:ARG:HD2	1.94	0.50
12:AB:130:PHE:CD2	12:AB:152:HIS:CE1	2.99	0.50
16:AG:285:ILE:HD13	16:AG:285:ILE:H	1.76	0.50
16:AG:296:ILE:HD11	16:AG:307:ILE:HD13	1.93	0.50
16:AG:360:THR:HG22	16:AG:360:THR:O	2.10	0.50
10:B:23:C:H2'	10:B:24:U:H6	1.77	0.50
17:C:25:ASP:O	17:C:29:LEU:HD13	2.11	0.50
18:D:280:C:H1'	36:V:40:ARG:HH22	1.75	0.50
18:D:580:C:H2'	18:D:581:G:O4'	2.11	0.50
18:D:612:C:C2	18:D:613:C:C5	2.99	0.50
18:D:739:C:O2	34:T:42:HIS:HE1	1.94	0.50
18:D:754:C:OP1	34:T:72:ARG:NH1	2.44	0.50
18:D:864:A:H2'	18:D:865:A:C8	2.46	0.50
20:F:63:GLU:O	20:F:67:ARG:HD2	2.12	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:G:35:ARG:HB3	22:H:14:LYS:NZ	2.26	0.50
21:G:109:GLN:O	21:G:113:ARG:NE	2.45	0.50
22:H:163:ARG:CA	22:H:298:GLU:HG3	2.41	0.50
23:I:188:GLU:OE1	23:I:188:GLU:N	2.45	0.50
26:L:65:GLU:OE1	26:L:65:GLU:HA	2.12	0.50
32:R:110:ARG:HE	32:R:117:TYR:HD2	1.57	0.50
38:X:92:ARG:HD2	41:a:888:C:OP1	2.12	0.50
41:a:1598:A:C5	41:a:1599:U:C5	2.99	0.50
41:a:1799:G:N2	48:h:154:LEU:HD23	2.26	0.50
41:a:2082:A:H2'	41:a:2083:G:O4'	2.12	0.50
41:a:2266:A:H4'	41:a:2267:A:N3	2.26	0.50
41:a:2848:G:O2'	41:a:2867:G:N2	2.36	0.50
48:h:261:LYS:HA	48:h:264:ASP:OD2	2.12	0.50
52:l:155:GLU:HG2	52:l:156:ASN:N	2.26	0.50
59:s:54:ILE:O	59:s:123:LYS:HG2	2.12	0.50
9:9:31:ARG:HH22	41:a:1106:G:N2	2.10	0.50
9:9:55:VAL:HG12	9:9:56:ARG:N	2.25	0.50
10:A:23:C:H2'	10:A:24:U:C6	2.46	0.50
10:A:48:C:O2'	10:A:49:G:OP2	2.27	0.50
13:AD:182:ARG:NH1	14:AE:534:GLU:OE1	2.44	0.50
14:AE:417:ARG:NH1	15:AF:43:ASN:O	2.45	0.50
16:AG:123:ARG:NH1	16:AG:123:ARG:HG2	2.26	0.50
16:AG:171:GLU:OE2	16:AG:267:GLY:C	2.54	0.50
16:AG:448:LEU:CG	16:AG:473:LEU:HD11	2.42	0.50
17:C:41:PRO:HD2	22:H:339:ARG:NE	2.26	0.50
17:C:51:TYR:HD1	17:C:54:GLN:HE22	1.58	0.50
18:D:36:C:C4	18:D:37:U:C5	3.00	0.50
18:D:49:U:C2	18:D:361:G:N2	2.79	0.50
22:H:303:LEU:O	22:H:303:LEU:CG	2.60	0.50
28:N:110:VAL:C	28:N:111:MET:SD	2.95	0.50
31:Q:31:ILE:HD12	31:Q:46:THR:HG22	1.94	0.50
34:T:33:THR:OG1	34:T:63:ARG:NH1	2.45	0.50
41:a:672:C:O2'	52:l:77:ILE:HD11	2.11	0.50
41:a:1199:U:H2'	41:a:1200:C:C6	2.47	0.50
41:a:1999:C:O3'	41:a:2722:G:N2	2.44	0.50
41:a:2698:U:H2'	41:a:2699:C:C6	2.46	0.50
52:l:6:LYS:CE	52:l:119:ILE:HG23	2.40	0.50
65:y:25:THR:OG1	65:y:88:ARG:HB3	2.12	0.50
3:2:3:ARG:NH2	3:2:4:GLU:HB2	2.26	0.50
9:9:30:SER:O	9:9:31:ARG:CZ	2.59	0.50
9:9:97:LYS:HZ1	9:9:130:PRO:HA	1.76	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:440:VAL:HG21	16:AG:481:LEU:HD22	1.89	0.50
17:C:60:LYS:HE3	18:D:735:C:C4'	2.42	0.50
17:C:66:SER:HA	26:L:49:TYR:CE1	2.47	0.50
18:D:10:A:OP2	25:K:131:THR:HG21	2.12	0.50
18:D:109:A:H2'	18:D:326:G:N2	2.27	0.50
18:D:160:A:H2'	18:D:161:A:O4'	2.12	0.50
18:D:501:C:H2'	18:D:502:A:C8	2.46	0.50
18:D:1017:U:O2'	18:D:1018:G:O4'	2.30	0.50
21:G:126:PHE:C	21:G:127:ASP:O	2.55	0.50
23:I:68:ILE:HG13	23:I:101:ILE:HD11	1.93	0.50
25:K:113:ALA:CA	25:K:116:GLU:OE1	2.60	0.50
38:X:71:ARG:HD2	38:X:72:GLU:N	2.26	0.50
39:Y:126:ARG:CZ	41:a:1083:U:P	3.00	0.50
41:a:65:U:C2	41:a:66:C:C6	3.00	0.50
41:a:960:A:H61	62:v:82:MET:CE	2.24	0.50
41:a:1199:U:H2'	41:a:1200:C:H6	1.77	0.50
41:a:1348:C:O2	41:a:1348:C:H2'	2.11	0.50
41:a:2063:C:O2	41:a:2450:A:N1	2.45	0.50
41:a:2230:G:C5'	43:c:30:LEU:HD23	2.42	0.50
41:a:2489:U:H2'	41:a:2490:G:O4'	2.12	0.50
52:l:112:LEU:CD1	52:l:117:ARG:HB2	2.39	0.50
54:n:122:PHE:CE2	54:n:128:TYR:CD1	3.00	0.50
62:v:73:ILE:HD12	62:v:73:ILE:N	2.26	0.50
5:4:6:ALA:HB3	5:4:65:VAL:HG22	1.93	0.50
9:9:52:MET:HG2	9:9:95:LEU:CD1	2.39	0.50
10:A:23:C:H2'	10:A:24:U:H6	1.77	0.50
16:AG:305:MET:HG2	16:AG:336:LEU:HA	1.94	0.50
10:B:68:C:C3'	10:B:69:C:H5''	2.42	0.50
17:C:60:LYS:NZ	18:D:735:C:C5'	2.75	0.50
18:D:339:C:C2	18:D:340:U:C5	2.99	0.50
18:D:821:G:H2'	18:D:822:U:C6	2.46	0.50
21:G:169:GLU:OE1	21:G:169:GLU:N	2.44	0.50
22:H:112:ILE:HA	22:H:121:THR:O	2.12	0.50
30:P:7:ARG:HD3	30:P:75:ASP:CG	2.36	0.50
30:P:37:ARG:HB3	30:P:75:ASP:HB2	1.93	0.50
34:T:33:THR:HG21	34:T:85:LEU:HD12	1.94	0.50
35:U:73:ALA:HA	35:U:76:LYS:CE	2.41	0.50
41:a:234:U:C2	41:a:235:U:C6	3.00	0.50
41:a:413:C:H2'	41:a:414:C:C6	2.46	0.50
41:a:636:G:N2	61:u:76:GLU:OE2	2.45	0.50
41:a:807:U:OP2	61:u:41:ARG:NH1	2.42	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1791:A:C2	41:a:1829:A:O4'	2.65	0.50
41:a:1823:G:O2'	41:a:1824:G:H5'	2.12	0.50
41:a:2065:C:H2'	41:a:2066:C:H6	1.77	0.50
41:a:2230:G:H5'	43:c:30:LEU:HD23	1.94	0.50
41:a:2252:G:H2'	41:a:2253:G:H8	1.76	0.50
41:a:2387:U:C4'	42:b:41:ARG:HH12	2.24	0.50
41:a:2392:A:C8	41:a:2429:G:C2	3.00	0.50
41:a:2682:A:C2	50:j:23:PRO:HB3	2.46	0.50
41:a:2723:C:H4'	63:w:1:MET:HE3	1.92	0.50
42:b:47:ALA:HB2	42:b:59:LEU:HD11	1.94	0.50
48:h:124:ILE:C	48:h:125:LYS:HD3	2.36	0.50
48:h:130:LEU:HD21	48:h:192:LEU:HD21	1.93	0.50
52:l:170:ARG:HH12	52:l:176:ASP:N	2.09	0.50
56:p:137:ASP:OD2	56:p:140:VAL:HG23	2.12	0.50
4:3:6:ARG:NH1	41:a:84:A:H2'	2.27	0.50
6:5:9:DG:P	11:AA:473:ARG:HG3	2.51	0.50
8:7:3:G:C4'	18:D:1500:A:N6	2.74	0.50
8:7:11:U:C4'	25:K:20:ARG:HH21	2.16	0.50
9:9:67:THR:N	9:9:68:PRO:CD	2.75	0.50
11:AA:846:GLY:C	16:AG:104:ARG:HH22	2.18	0.50
12:AB:29:LEU:HB3	12:AB:56:PHE:HD2	1.77	0.50
16:AG:139:THR:HA	16:AG:180:ARG:HA	1.92	0.50
16:AG:369:PHE:O	16:AG:373:LEU:HG	2.12	0.50
10:B:15:G:H2'	10:B:15:G:N3	2.27	0.50
17:C:60:LYS:NZ	18:D:735:C:C4'	2.74	0.50
23:I:82:GLU:O	23:I:86:LYS:CD	2.60	0.50
25:K:134:ILE:O	25:K:137:VAL:HG12	2.12	0.50
32:R:80:ILE:HG22	32:R:81:LEU:N	2.27	0.50
39:Y:55:PRO:C	39:Y:71:LYS:HE2	2.36	0.50
41:a:1454:C:H4'	63:w:63:ARG:NH1	2.26	0.50
41:a:1790:C:C3'	41:a:1791:A:C8	2.94	0.50
41:a:2063:C:C4	41:a:2064:C:C5	3.00	0.50
41:a:2682:A:C5'	50:j:11:MET:HE1	2.42	0.50
51:k:34:LEU:CB	51:k:51:GLU:OE2	2.60	0.50
59:s:98:GLU:HB3	59:s:102:GLU:OE2	2.12	0.50
9:9:43:LYS:HA	9:9:46:ARG:NH1	2.27	0.49
10:A:68:C:C3'	10:A:69:C:H5''	2.42	0.49
11:AA:400:VAL:HG21	11:AA:452:ARG:HE	1.77	0.49
14:AE:814:CYS:SG	14:AE:815:GLY:N	2.85	0.49
21:G:23:TRP:CZ3	21:G:25:PRO:HA	2.47	0.49
21:G:165:ASP:O	21:G:169:GLU:OE1	2.28	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:W:33:THR:HG21	37:W:49:ILE:HD11	1.94	0.49
37:W:49:ILE:HG23	37:W:49:ILE:O	2.11	0.49
38:X:65:VAL:HG23	38:X:66:GLU:OE1	2.12	0.49
39:Y:99:LYS:HZ3	39:Y:138:VAL:HG13	1.75	0.49
41:a:15:G:C2'	49:i:15:MET:HE1	2.42	0.49
41:a:271:G:H4'	41:a:272:A:OP1	2.10	0.49
41:a:2298:A:O3'	54:n:72:LYS:HE2	2.11	0.49
46:f:41:THR:N	46:f:45:ARG:HH22	2.09	0.49
58:r:90:LEU:HD21	58:r:94:ILE:CD1	2.42	0.49
60:t:1:MET:HE3	60:t:32:TYR:HB3	1.94	0.49
60:t:103:VAL:HG21	60:t:107:LEU:CD2	2.41	0.49
61:u:55:MET:HE1	61:u:60:ARG:HA	1.94	0.49
2:1:35:ILE:O	2:1:39:THR:HG23	2.12	0.49
3:2:49:LYS:HE2	3:2:49:LYS:HA	1.93	0.49
5:4:48:MET:HE2	5:4:86:LEU:HG	1.93	0.49
12:AB:129:ILE:O	12:AB:142:LEU:HB2	2.12	0.49
13:AD:288:GLU:C	13:AD:290:LEU:N	2.69	0.49
16:AG:183:LEU:HD23	16:AG:197:VAL:HG22	1.94	0.49
16:AG:248:VAL:HG21	16:AG:275:LEU:HG	1.94	0.49
10:B:32:C:H2'	10:B:33:U:H5'	1.94	0.49
18:D:908:A:H2'	18:D:909:A:C8	2.48	0.49
25:K:46:VAL:N	25:K:71:MET:HE2	2.27	0.49
39:Y:20:SER:CB	39:Y:21:PRO:HD3	2.42	0.49
39:Y:21:PRO:CB	39:Y:22:PRO:HD3	2.41	0.49
39:Y:126:ARG:HD2	39:Y:127:SER:N	2.26	0.49
41:a:132:G:H2'	41:a:133:U:C6	2.47	0.49
41:a:780:G:C2	41:a:782:A:C2	3.00	0.49
41:a:1801:A:OP1	48:h:146:MET:HE1	2.11	0.49
41:a:2193:G:O2'	41:a:2194:U:OP1	2.28	0.49
53:m:25:LYS:C	53:m:29:GLN:HE22	2.20	0.49
57:q:2:LYS:CE	57:q:35:GLN:HE21	2.25	0.49
3:2:2:ILE:HG12	3:2:42:GLU:CD	2.37	0.49
3:2:12:ARG:HH21	45:e:29:ARG:CZ	2.25	0.49
4:3:94:ARG:CB	4:3:103:ILE:HD12	2.43	0.49
8:7:3:G:H5''	18:D:1501:C:N4	2.27	0.49
9:9:52:MET:CE	9:9:85:SER:HB2	2.42	0.49
9:9:108:VAL:O	9:9:109:LYS:HB3	2.12	0.49
10:A:33:U:H5''	27:M:84:THR:HG21	1.93	0.49
12:AB:19:GLU:HG2	12:AB:23:ARG:CZ	2.42	0.49
14:AE:77:ARG:HB3	14:AE:80:HIS:HB2	1.93	0.49
16:AG:240:THR:HG21	16:AG:247:PRO:HG3	1.92	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:320:ARG:CG	16:AG:320:ARG:NH1	2.73	0.49
10:B:47:U:O2'	10:B:50:U:P	2.70	0.49
18:D:373:A:C2	18:D:482:A:C6	3.00	0.49
18:D:744:C:H2'	18:D:745:G:C8	2.47	0.49
18:D:877:G:H5''	28:N:80:ARG:HD2	1.93	0.49
18:D:922:G:H2'	18:D:923:A:C8	2.47	0.49
20:F:6:VAL:HG12	31:Q:108:THR:O	2.11	0.49
20:F:36:GLU:OE2	20:F:36:GLU:HA	2.11	0.49
23:I:49:LYS:HZ3	23:I:75:ILE:HD11	1.75	0.49
24:J:104:ARG:NE	24:J:104:ARG:HA	2.27	0.49
28:N:106:THR:HB	28:N:121:LEU:CD1	2.41	0.49
34:T:5:THR:O	34:T:8:THR:OG1	2.27	0.49
39:Y:129:GLU:CB	39:Y:133:ARG:HH12	2.16	0.49
41:a:3:U:C2	41:a:4:U:C5	3.01	0.49
41:a:125:A:OP2	53:m:19:ARG:NH2	2.39	0.49
41:a:247:G:O6	55:o:12:LYS:NZ	2.39	0.49
41:a:372:G:O4'	43:c:61:LYS:NZ	2.44	0.49
41:a:998:C:OP1	66:z:84:LYS:NZ	2.39	0.49
41:a:1021:A:H3'	41:a:1021:A:N3	2.27	0.49
41:a:2290:G:H2'	41:a:2291:U:C6	2.47	0.49
50:j:55:LYS:NZ	50:j:59:ARG:C	2.70	0.49
54:n:8:TYR:HA	54:n:12:VAL:HB	1.95	0.49
54:n:134:GLU:CG	54:n:137:ILE:HG23	2.40	0.49
58:r:137:GLU:CG	58:r:138:VAL:HG23	2.39	0.49
59:s:1:MET:HE2	66:z:97:ASP:OD2	2.13	0.49
59:s:131:ASN:OD1	59:s:131:ASN:O	2.30	0.49
64:x:4:LYS:N	64:x:4:LYS:HD3	2.26	0.49
64:x:90:VAL:HG22	64:x:115:LEU:HD11	1.94	0.49
9:9:67:THR:O	9:9:72:LEU:HG	2.12	0.49
14:AE:81:ARG:HG3	14:AE:81:ARG:NH2	2.17	0.49
14:AE:429:LEU:HB3	14:AE:925:GLU:HG2	1.94	0.49
16:AG:323:GLN:CA	16:AG:323:GLN:NE2	2.75	0.49
16:AG:442:ARG:C	16:AG:446:PHE:HD2	2.15	0.49
18:D:321:A:N7	18:D:328:C:O2'	2.38	0.49
18:D:1073:U:O2'	21:G:105:LYS:HE3	2.11	0.49
18:D:1215:G:C2	18:D:1216:A:C8	3.01	0.49
24:J:170:TRP:CD2	24:J:186:PRO:HB3	2.47	0.49
28:N:35:ALA:O	28:N:39:VAL:HG23	2.12	0.49
28:N:96:MET:HG2	28:N:96:MET:O	2.13	0.49
30:P:26:VAL:HG12	30:P:30:LYS:HZ2	1.78	0.49
35:U:18:GLN:NE2	35:U:35:ARG:HE	2.10	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:57:C:H2'	41:a:58:G:O4'	2.12	0.49
41:a:851:C:C2	41:a:852:U:C5	3.00	0.49
41:a:1590:A:H2'	41:a:1591:A:C8	2.48	0.49
41:a:2189:U:O2'	41:a:2190:G:O5'	2.31	0.49
41:a:2362:C:O5'	55:o:40:ARG:NH2	2.45	0.49
41:a:2849:U:OP2	65:y:93:ARG:CZ	2.60	0.49
47:g:63:ARG:HG3	47:g:64:PHE:CD1	2.48	0.49
50:j:184:ARG:HH12	65:y:7:GLN:NE2	2.09	0.49
58:r:6:LEU:CD1	58:r:37:VAL:HG23	2.39	0.49
59:s:13:ARG:NH2	59:s:50:THR:HA	2.27	0.49
59:s:55:ILE:CG1	59:s:123:LYS:HE2	2.42	0.49
2:1:17:VAL:HG12	2:1:76:VAL:HG21	1.94	0.49
3:2:14:PRO:HA	3:2:32:LEU:HD13	1.94	0.49
4:3:6:ARG:HG2	4:3:7:ARG:H	1.77	0.49
7:6:23:DC:H5'	14:AE:259:ARG:NH1	2.11	0.49
10:A:32:C:H2'	10:A:33:U:H5'	1.94	0.49
10:A:72:A:H2'	10:A:73:A:H5''	1.95	0.49
11:AA:143:ARG:NH1	11:AA:507:GLY:O	2.32	0.49
11:AA:411:ARG:NH2	11:AA:427:ASP:OD2	2.44	0.49
12:AB:6:LEU:O	12:AB:7:LEU:HD23	2.13	0.49
18:D:73:C:O2	18:D:73:C:C2'	2.57	0.49
18:D:925:G:C6	18:D:927:G:N7	2.81	0.49
18:D:1458:G:OP1	19:E:30:THR:OG1	2.30	0.49
23:I:164:ARG:HD2	23:I:166:GLU:OE2	2.12	0.49
25:K:34:THR:HG22	25:K:52:LYS:HD3	1.95	0.49
36:V:62:ARG:HA	36:V:73:TRP:CZ3	2.47	0.49
38:X:3:ARG:HH22	47:g:36:VAL:HG12	1.77	0.49
39:Y:37:PHE:HE1	39:Y:58:ILE:HD11	1.76	0.49
41:a:111:A:O2'	45:e:58:ASN:ND2	2.46	0.49
41:a:920:A:C5	41:a:921:C:C5	3.01	0.49
41:a:1843:C:C2	41:a:1844:C:C5	3.00	0.49
41:a:2093:G:N7	41:a:2225:A:H2'	2.28	0.49
49:i:28:LEU:O	49:i:37:LYS:NZ	2.46	0.49
52:l:97:ASN:ND2	52:l:100:MET:HE2	2.27	0.49
52:l:194:LYS:O	52:l:197:GLU:N	2.45	0.49
56:p:26:ILE:HG22	56:p:75:MET:HE2	1.93	0.49
56:p:127:THR:HG22	56:p:128:GLN:N	2.26	0.49
59:s:125:TYR:HE2	59:s:130:HIS:HB2	1.78	0.49
9:9:25:ALA:CB	9:9:99:PHE:HD2	2.25	0.49
9:9:52:MET:HE2	9:9:85:SER:HB2	1.95	0.49
12:AB:51:PHE:CZ	14:AE:288:PRO:HG3	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:AD:285:THR:HA	13:AD:315:GLY:HA2	1.93	0.49
14:AE:425:ARG:NH1	14:AE:458:ASN:O	2.45	0.49
16:AG:5:ILE:HA	16:AG:8:VAL:HB	1.93	0.49
16:AG:65:VAL:HG12	16:AG:66:ASP:H	1.76	0.49
16:AG:283:PHE:CE2	16:AG:332:SER:HA	2.47	0.49
16:AG:448:LEU:HD23	16:AG:473:LEU:HD12	1.95	0.49
10:B:56:C:H2'	10:B:57:A:H5''	1.93	0.49
18:D:311:C:O5'	18:D:311:C:H6	1.94	0.49
18:D:373:A:C2	18:D:374:A:C8	3.01	0.49
18:D:653:U:O5'	28:N:56:LYS:HE2	2.11	0.49
18:D:816:A:OP1	18:D:1526:G:O2'	2.27	0.49
18:D:1314:C:H2'	18:D:1315:U:C6	2.48	0.49
21:G:110:SER:CA	21:G:113:ARG:HE	2.25	0.49
25:K:13:GLU:C	25:K:14:LYS:HD2	2.37	0.49
29:O:52:LEU:HB3	29:O:58:VAL:HG12	1.94	0.49
31:Q:98:ARG:HG2	31:Q:98:ARG:HH11	1.77	0.49
32:R:7:LEU:HD23	36:V:34:TYR:OH	2.12	0.49
32:R:38:TYR:HE2	32:R:40:THR:CG2	2.25	0.49
35:U:12:LYS:HD3	35:U:13:LYS:HG2	1.93	0.49
36:V:17:MET:HG3	36:V:20:SER:HB2	1.95	0.49
41:a:1142:A:C4	41:a:1144:A:N7	2.80	0.49
41:a:1830:C:N3	41:a:1831:G:N7	2.61	0.49
41:a:2014:A:H2'	41:a:2015:A:C8	2.47	0.49
41:a:2159:G:H2'	41:a:2160:C:C6	2.47	0.49
41:a:2179:C:C2	41:a:2180:U:C5	3.00	0.49
41:a:2515:C:H2'	41:a:2516:A:H8	1.78	0.49
45:e:14:LEU:HD13	45:e:57:LEU:HD22	1.95	0.49
47:g:36:VAL:HG21	47:g:41:HIS:ND1	2.27	0.49
54:n:16:LEU:HD12	54:n:28:VAL:CG2	2.43	0.49
59:s:73:VAL:CG1	59:s:74:TYR:N	2.75	0.49
63:w:114:GLU:N	63:w:114:GLU:OE1	2.46	0.49
66:z:79:PHE:CE1	66:z:110:VAL:HG22	2.48	0.49
5:4:1:MET:O	5:4:1:MET:HE3	2.13	0.49
5:4:89:ILE:CG2	5:4:91:PHE:CE1	2.95	0.49
12:AB:32:MET:HE2	12:AB:32:MET:HA	1.95	0.49
12:AB:141:LEU:O	12:AB:151:LYS:N	2.46	0.49
12:AB:158:GLU:HB3	12:AB:162:LEU:CD2	2.41	0.49
16:AG:18:LEU:HB3	16:AG:19:PRO:HD2	1.95	0.49
16:AG:130:PHE:C	16:AG:186:VAL:CG1	2.86	0.49
18:D:500:G:H2'	18:D:501:C:H6	1.77	0.49
18:D:736:C:H2'	18:D:737:C:H6	1.77	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:1071:C:H2'	18:D:1072:G:C8	2.47	0.49
25:K:88:VAL:HG12	25:K:93:ARG:HA	1.94	0.49
29:O:51:PRO:HA	29:O:103:PHE:HE2	1.76	0.49
36:V:50:ASN:HB2	36:V:52:GLU:OE2	2.13	0.49
39:Y:60:VAL:CG2	39:Y:66:PHE:HB3	2.28	0.49
39:Y:99:LYS:CD	39:Y:138:VAL:O	2.61	0.49
41:a:145:C:H2'	41:a:146:A:H8	1.78	0.49
41:a:1567:G:OP2	48:h:83:TYR:OH	2.19	0.49
41:a:1921:G:C2'	41:a:1922:G:H5'	2.43	0.49
41:a:2314:A:P	54:n:88:LYS:HZ2	2.35	0.49
41:a:2485:G:O3'	62:v:125:PRO:HB3	2.12	0.49
41:a:2684:U:C4	41:a:2685:G:N7	2.81	0.49
41:a:2740:A:H2'	41:a:2741:A:C8	2.47	0.49
43:c:6:GLN:CD	43:c:49:LEU:HD22	2.38	0.49
49:i:40:ARG:NH1	49:i:41:HIS:CG	2.81	0.49
57:q:10:LEU:HD23	57:q:33:HIS:CD2	2.48	0.49
63:w:82:GLU:O	63:w:86:ARG:HG2	2.13	0.49
63:w:87:PHE:HZ	63:w:115:LEU:HG	1.77	0.49
66:z:58:ARG:HA	66:z:61:TRP:CE3	2.48	0.49
3:2:11:LEU:HD11	3:2:46:ALA:HB3	1.95	0.49
5:4:35:GLU:OE1	5:4:93:ARG:NH2	2.46	0.49
6:5:9:DG:H5''	11:AA:473:ARG:HB3	0.54	0.49
9:9:24:SER:HB3	9:9:116:GLU:HG3	1.94	0.49
11:AA:9:LYS:HG2	11:AA:1171:ARG:HH12	1.77	0.49
11:AA:732:ILE:HD11	11:AA:769:PRO:HB3	1.95	0.49
14:AE:749:LYS:HD3	14:AE:753:SER:HB2	1.94	0.49
10:B:21:A:H4'	10:B:21:A:OP1	2.13	0.49
17:C:51:TYR:O	17:C:54:GLN:HG3	2.11	0.49
18:D:56:U:H2'	18:D:57:G:C8	2.48	0.49
18:D:953:G:O6	38:X:103:LYS:CE	2.60	0.49
21:G:114:LEU:HD13	21:G:144:LEU:HB3	1.94	0.49
27:M:138:ARG:HG2	27:M:142:HIS:CE1	2.48	0.49
29:O:12:ARG:HD3	29:O:13:LYS:H	1.78	0.49
32:R:74:LEU:HD11	32:R:80:ILE:CD1	2.43	0.49
41:a:537:G:P	59:s:2:LYS:CE	3.00	0.49
41:a:1036:G:C6	41:a:1120:G:C5	3.01	0.49
41:a:2547:A:H2'	41:a:2548:U:H6	1.77	0.49
45:e:10:SER:OG	45:e:13:GLU:HG2	2.13	0.49
52:l:195:GLN:HA	52:l:198:GLU:OE1	2.12	0.49
62:v:57:VAL:HG22	62:v:60:GLN:O	2.13	0.49
11:AA:887:VAL:O	16:AG:105:ILE:CD1	2.48	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AB:161:LYS:HA	12:AB:161:LYS:HD3	1.55	0.49
16:AG:59:PHE:CD1	16:AG:97:ILE:HG23	2.48	0.49
10:B:12:G:C2'	10:B:13:C:OP1	2.61	0.49
17:C:13:PHE:CB	17:C:51:TYR:CD1	2.95	0.49
17:C:60:LYS:NZ	18:D:734:G:C3'	2.76	0.49
18:D:136:C:C2	18:D:137:U:C6	3.00	0.49
25:K:157:ARG:HH12	28:N:43:GLU:CD	2.20	0.49
27:M:74:GLU:OE1	27:M:75:VAL:O	2.30	0.49
31:Q:94:GLU:O	31:Q:97:ILE:HG22	2.13	0.49
34:T:64:ARG:CZ	34:T:64:ARG:CB	2.91	0.49
34:T:77:ARG:HH21	34:T:80:GLN:NE2	2.11	0.49
38:X:81:MET:HG2	41:a:888:C:C4	2.48	0.49
39:Y:77:VAL:HG13	39:Y:81:LYS:NZ	2.27	0.49
39:Y:126:ARG:HH12	41:a:1082:U:C3'	2.26	0.49
41:a:235:U:C2	41:a:236:C:C5	3.00	0.49
41:a:594:U:H2'	41:a:595:C:C6	2.47	0.49
48:h:52:ARG:HG3	48:h:53:HIS:ND1	2.27	0.49
62:v:29:GLY:HA3	62:v:104:GLU:HG3	1.94	0.49
65:y:7:GLN:NE2	65:y:10:GLN:HE21	2.11	0.49
2:1:95:ARG:NH1	2:1:97:LEU:HG	2.28	0.49
5:4:2:PHE:HB2	5:4:61:LEU:HG	1.94	0.49
10:A:47:U:O2'	10:A:50:U:P	2.70	0.49
11:AA:855:PRO:HB3	16:AG:35:THR:HG1	1.78	0.49
16:AG:384:LEU:O	16:AG:387:VAL:HG23	2.12	0.49
10:B:72:A:H2'	10:B:73:A:H5''	1.95	0.49
18:D:653:U:O4'	28:N:56:LYS:CE	2.60	0.49
18:D:718:A:C5	31:Q:118:HIS:ND1	2.81	0.49
18:D:1314:C:H2'	18:D:1315:U:H6	1.78	0.49
23:I:77:ILE:HA	23:I:84:VAL:HG13	1.94	0.49
33:S:57:PRO:HA	33:S:60:GLN:OE1	2.13	0.49
39:Y:91:LYS:HZ2	39:Y:97:VAL:HG21	1.78	0.49
41:a:583:G:OP1	66:z:11:ARG:NH2	2.46	0.49
41:a:1385:A:O2'	41:a:1396:U:O2	2.31	0.49
41:a:2677:G:C6	41:a:2731:G:C6	3.01	0.49
41:a:2754:U:O3'	57:q:19:ARG:NH2	2.46	0.49
41:a:2902:C:N4	41:a:2903:U:O4	2.45	0.49
48:h:131:PRO:HD3	48:h:189:ARG:NH1	2.27	0.49
54:n:67:ILE:HD13	54:n:87:CYS:HB3	1.95	0.49
59:s:15:TRP:CZ3	59:s:135:GLN:CA	2.93	0.49
2:1:20:VAL:O	2:1:23:LEU:HB3	2.13	0.48
6:5:17:DC:C5	11:AA:542:ARG:HD3	2.48	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AA:1280:ALA:HB1	14:AE:918:ILE:HG22	1.96	0.48
12:AB:136:GLU:OE1	12:AB:160:ARG:CZ	2.61	0.48
12:AB:140:MET:HE1	30:P:102:LEU:C	2.38	0.48
12:AB:155:LYS:CG	12:AB:156:ASN:H	2.26	0.48
13:AD:100:LEU:HB2	13:AD:144:ILE:HG23	1.94	0.48
14:AE:412:LEU:HD22	14:AE:441:LEU:HD21	1.94	0.48
16:AG:453:VAL:HG11	16:AG:459:LEU:HD23	1.95	0.48
18:D:674:G:P	26:L:86:ARG:HH22	2.35	0.48
18:D:1218:C:H2'	18:D:1219:A:C8	2.48	0.48
18:D:1428:A:H2'	18:D:1429:A:O4'	2.13	0.48
19:E:22:ALA:HA	19:E:25:ARG:HE	1.78	0.48
26:L:66:ALA:HB1	26:L:67:PRO:HD2	1.94	0.48
29:O:32:GLN:HG3	29:O:32:GLN:O	2.13	0.48
41:a:1571:A:H2'	41:a:1572:A:C8	2.47	0.48
46:f:41:THR:N	46:f:45:ARG:NH2	2.61	0.48
47:g:47:LYS:NZ	54:n:115:ARG:HA	2.27	0.48
51:k:21:TYR:CE2	51:k:38:LYS:CG	2.96	0.48
52:l:40:ARG:HD2	52:l:92:HIS:CE1	2.48	0.48
54:n:70:ALA:O	54:n:81:GLN:OE1	2.31	0.48
56:p:94:TYR:CE1	56:p:152:ARG:HD2	2.48	0.48
60:t:121:GLU:HG3	60:t:123:LEU:HD12	1.94	0.48
62:v:70:ASP:C	62:v:92:TRP:CZ3	2.90	0.48
65:y:9:GLU:HG3	65:y:55:LEU:CB	2.42	0.48
5:4:55:GLU:OE1	5:4:55:GLU:N	2.42	0.48
8:7:11:U:HO2'	25:K:33:PHE:HE1	1.49	0.48
9:9:95:LEU:HD12	9:9:99:PHE:CZ	2.48	0.48
11:AA:146:VAL:HG21	11:AA:513:GLN:HE21	1.77	0.48
11:AA:380:ALA:HA	12:AB:65:HIS:ND1	2.28	0.48
11:AA:554:HIS:HD2	11:AA:558:VAL:HB	1.77	0.48
14:AE:291:ILE:O	14:AE:293:ARG:N	2.46	0.48
16:AG:64:VAL:HA	16:AG:75:ILE:HB	1.94	0.48
16:AG:440:VAL:HG22	16:AG:481:LEU:HD21	1.95	0.48
17:C:21:ILE:HG13	17:C:54:GLN:NE2	2.29	0.48
18:D:110:C:H2'	18:D:111:G:O4'	2.14	0.48
18:D:123:U:H5''	18:D:311:C:O2'	2.13	0.48
18:D:429:U:N3	18:D:431:A:N6	2.61	0.48
18:D:718:A:C8	31:Q:118:HIS:HB3	2.47	0.48
22:H:267:TRP:CE3	22:H:340:ARG:HG3	2.47	0.48
30:P:63:ASP:OD1	30:P:65:TYR:CZ	2.66	0.48
35:U:26:ASN:ND2	35:U:31:ARG:HE	2.11	0.48
41:a:26:G:H2'	41:a:27:G:O4'	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:962:G:C5	41:a:963:U:C5	3.01	0.48
41:a:1790:C:H2'	41:a:1791:A:C8	2.48	0.48
41:a:2297:A:N1	41:a:2321:U:O4	2.47	0.48
41:a:2351:G:O6	55:o:39:LYS:HE3	2.13	0.48
52:l:149:ILE:CG2	52:l:175:ILE:HD11	2.43	0.48
57:q:1:MET:HE1	57:q:35:GLN:HA	1.95	0.48
65:y:40:LEU:HD12	65:y:40:LEU:N	2.28	0.48
66:z:51:ARG:N	66:z:51:ARG:HD2	2.28	0.48
1:0:61:ALA:HB1	1:0:96:VAL:CG2	2.43	0.48
1:0:68:ARG:NH2	1:0:90:ARG:HD3	2.29	0.48
5:4:1:MET:SD	5:4:3:THR:OG1	2.71	0.48
9:9:24:SER:O	9:9:116:GLU:HB2	2.14	0.48
10:A:21:A:H4'	10:A:21:A:OP1	2.13	0.48
10:A:22:G:O2'	10:A:23:C:O5'	2.31	0.48
11:AA:1340:GLU:HB2	14:AE:19:ALA:HB3	1.94	0.48
13:AD:182:ARG:O	13:AD:205:MET:HA	2.14	0.48
16:AG:200:SER:C	16:AG:230:PRO:HG2	2.38	0.48
16:AG:453:VAL:HG11	16:AG:459:LEU:CG	2.42	0.48
18:D:429:U:O2	18:D:430:A:C8	2.66	0.48
20:F:4:ILE:CD1	20:F:19:PHE:HA	2.43	0.48
24:J:124:MET:HE2	24:J:128:ARG:CA	2.43	0.48
27:M:109:ARG:O	27:M:119:ARG:NH2	2.46	0.48
29:O:49:ARG:O	29:O:53:GLU:OE1	2.32	0.48
34:T:64:ARG:HA	34:T:64:ARG:HH11	1.75	0.48
38:X:9:ILE:HG13	38:X:10:PRO:HD2	1.95	0.48
41:a:208:C:C2	41:a:209:C:C5	3.01	0.48
41:a:1825:U:P	48:h:52:ARG:NH1	2.86	0.48
41:a:2455:G:H2'	41:a:2456:C:C6	2.48	0.48
46:f:4:THR:CB	46:f:37:GLU:OE2	2.58	0.48
46:f:41:THR:CA	46:f:45:ARG:HH22	2.26	0.48
51:k:7:GLU:HG3	51:k:27:LYS:HZ3	1.77	0.48
56:p:94:TYR:C	56:p:95:ARG:HD2	2.38	0.48
63:w:38:LEU:N	63:w:39:PRO:CD	2.76	0.48
2:1:86:MET:HE1	2:1:88:ARG:CD	2.42	0.48
11:AA:529:ARG:HH11	11:AA:572:ILE:HG22	1.77	0.48
14:AE:59:ALA:HB3	14:AE:71:LEU:CD2	2.42	0.48
14:AE:81:ARG:CG	14:AE:81:ARG:NH2	2.77	0.48
14:AE:107:LEU:HD22	14:AE:299:LEU:HD21	1.95	0.48
16:AG:131:ARG:O	16:AG:134:GLU:CG	2.47	0.48
16:AG:349:GLN:C	16:AG:351:GLU:N	2.71	0.48
16:AG:353:HIS:HA	16:AG:356:ILE:HG21	1.94	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:71:A:N1	18:D:100:G:C8	2.81	0.48
18:D:893:C:C2	18:D:894:G:C8	3.01	0.48
18:D:1439:G:OP1	19:E:33:LYS:CE	2.61	0.48
23:I:142:MET:HG3	23:I:170:GLU:OE1	2.13	0.48
27:M:18:PHE:CD2	27:M:59:LEU:HD21	2.46	0.48
27:M:132:GLY:O	27:M:136:LYS:NZ	2.45	0.48
27:M:142:HIS:O	27:M:146:GLU:HG2	2.13	0.48
39:Y:52:LEU:O	39:Y:54:ILE:HG12	2.14	0.48
41:a:208:C:H2'	41:a:209:C:H6	1.78	0.48
41:a:760:G:H2'	41:a:761:A:O4'	2.14	0.48
41:a:879:G:H2'	41:a:880:G:O4'	2.14	0.48
41:a:1042:G:C4	41:a:1043:C:C6	3.02	0.48
41:a:1205:A:C2	52:l:165:HIS:CE1	3.00	0.48
41:a:2420:C:C5'	51:k:8:LYS:NZ	2.76	0.48
41:a:2809:A:H2'	41:a:2810:A:C8	2.48	0.48
47:g:47:LYS:HE2	54:n:115:ARG:HA	1.95	0.48
47:g:58:ASP:CG	47:g:62:LYS:NZ	2.68	0.48
48:h:160:THR:O	48:h:195:VAL:HG12	2.13	0.48
51:k:7:GLU:HG2	51:k:27:LYS:NZ	2.29	0.48
59:s:15:TRP:HD1	59:s:53:TYR:HB2	1.78	0.48
59:s:98:GLU:OE1	59:s:124:VAL:HG13	2.13	0.48
62:v:42:THR:HA	62:v:93:VAL:HG12	1.95	0.48
63:w:73:ASN:C	63:w:73:ASN:HD22	2.17	0.48
64:x:115:LEU:HD23	64:x:117:PHE:CE1	2.49	0.48
1:0:6:GLN:OE1	1:0:11:GLN:HG2	2.14	0.48
6:5:10:DT:C4'	11:AA:62:TYR:OH	2.61	0.48
9:9:31:ARG:NE	9:9:31:ARG:HA	2.28	0.48
11:AA:18:ARG:HE	11:AA:620:ASN:HA	1.77	0.48
11:AA:565:GLU:HA	11:AA:569:ILE:HG12	1.95	0.48
14:AE:678:ARG:NH1	14:AE:756:GLU:OE1	2.46	0.48
14:AE:977:SER:OG	14:AE:980:THR:OG1	2.31	0.48
16:AG:36:LYS:HE3	16:AG:43:ILE:HG12	1.95	0.48
16:AG:277:ASP:HB2	16:AG:282:GLN:HG3	1.95	0.48
16:AG:393:LEU:CG	16:AG:398:LEU:HD22	2.22	0.48
16:AG:422:GLU:CA	16:AG:426:GLY:N	2.62	0.48
17:C:45:THR:HA	22:H:340:ARG:HH21	1.77	0.48
18:D:43:C:P	35:U:12:LYS:NZ	2.86	0.48
18:D:1319:A:C8	18:D:1323:G:C6	3.01	0.48
39:Y:126:ARG:NH1	41:a:1082:U:O5'	2.46	0.48
40:Z:29:LYS:HG3	40:Z:30:PHE:CE2	2.49	0.48
41:a:1321:A:C5	41:a:1322:A:C8	3.02	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2091:C:O2	43:c:34:HIS:CE1	2.67	0.48
41:a:2093:G:P	58:r:22:LYS:HZ2	2.24	0.48
42:b:26:PHE:H	42:b:29:GLU:HG3	1.79	0.48
50:j:101:PHE:CZ	50:j:203:VAL:HG12	2.48	0.48
59:s:39:LYS:HA	59:s:44:TYR:HD1	1.79	0.48
59:s:88:THR:O	59:s:92:MET:HE2	2.13	0.48
62:v:41:LEU:HD22	62:v:45:GLN:NE2	2.27	0.48
66:z:78:LYS:CD	66:z:117:LEU:HD21	2.43	0.48
2:1:1:MET:CG	2:1:62:ASP:OD2	2.61	0.48
2:1:23:LEU:HD21	2:1:35:ILE:HG22	1.95	0.48
5:4:69:GLU:H	5:4:69:GLU:CD	2.22	0.48
10:A:70:G:C3'	10:A:71:C:H5''	2.43	0.48
11:AA:363:LEU:HB3	11:AA:381:ALA:HB1	1.96	0.48
12:AB:142:LEU:HD12	12:AB:150:ILE:O	2.13	0.48
16:AG:62:TRP:CD1	16:AG:95:ASP:HB2	2.49	0.48
17:C:38:LYS:HE2	17:C:38:LYS:CA	2.37	0.48
18:D:35:G:H2'	18:D:36:C:C6	2.48	0.48
18:D:106:C:N4	19:E:10:ARG:HH22	2.12	0.48
18:D:123:U:C2	18:D:124:C:C5	3.01	0.48
23:I:119:SER:O	23:I:123:GLN:HG2	2.13	0.48
24:J:76:TYR:HE2	24:J:204:TYR:CB	2.25	0.48
25:K:157:ARG:CZ	28:N:43:GLU:HG3	2.43	0.48
37:W:44:MET:SD	37:W:62:VAL:CG1	2.99	0.48
38:X:86:TYR:CE2	38:X:90:ARG:NE	2.82	0.48
39:Y:74:PRO:O	39:Y:78:LEU:HG	2.13	0.48
40:Z:26:MET:O	40:Z:30:PHE:HB2	2.14	0.48
41:a:272:A:N3	41:a:273:G:C8	2.80	0.48
41:a:379:G:N1	41:a:396:G:C6	2.82	0.48
41:a:840:C:C2	41:a:841:G:C8	3.01	0.48
41:a:1342:A:O2'	41:a:1344:U:OP2	2.31	0.48
41:a:1664:A:C2'	41:a:1665:A:H5'	2.43	0.48
41:a:1792:G:H5''	48:h:204:VAL:HG23	1.95	0.48
41:a:2349:G:O6	41:a:2369:A:N6	2.46	0.48
54:n:94:GLU:HA	54:n:97:TRP:CE3	2.43	0.48
61:u:77:ILE:HD11	61:u:108:ALA:HB1	1.95	0.48
62:v:41:LEU:HD22	62:v:45:GLN:HE22	1.78	0.48
5:4:59:GLU:OE1	5:4:59:GLU:N	2.47	0.48
11:AA:560:PRO:O	14:AE:780:ARG:NH2	2.47	0.48
11:AA:618:GLN:HG3	14:AE:770:LEU:HD13	1.95	0.48
11:AA:887:VAL:CA	16:AG:105:ILE:CD1	2.57	0.48
12:AB:102:LYS:NZ	14:AE:285:LEU:HG	2.27	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AB:118:ILE:HG12	12:AB:130:PHE:CD2	2.49	0.48
14:AE:520:ALA:HB3	14:AE:546:ALA:HB2	1.95	0.48
16:AG:433:ASP:O	16:AG:456:LEU:HD21	2.12	0.48
16:AG:467:LEU:HD12	16:AG:482:ILE:HD11	1.95	0.48
17:C:12:ARG:CZ	17:C:13:PHE:CE1	2.97	0.48
17:C:60:LYS:CD	18:D:734:G:O2'	2.62	0.48
18:D:376:G:OP2	35:U:70:ARG:HD3	2.12	0.48
18:D:406:G:C5	18:D:495:A:C5	3.01	0.48
18:D:563:A:N1	18:D:884:U:O4	2.46	0.48
18:D:1127:G:C2	18:D:1128:C:C6	3.01	0.48
18:D:1227:A:H5'	38:X:110:LYS:NZ	2.28	0.48
18:D:1530:G:H2'	18:D:1531:A:C8	2.49	0.48
21:G:112:LYS:HE2	21:G:112:LYS:HA	1.95	0.48
24:J:170:TRP:CZ3	24:J:190:ASP:HB3	2.48	0.48
26:L:35:LYS:HG3	26:L:36:ILE:N	2.27	0.48
29:O:52:LEU:O	29:O:55:VAL:O	2.31	0.48
32:R:27:CYS:SG	32:R:30:LYS:NZ	2.68	0.48
33:S:12:LYS:HD2	33:S:16:LEU:CD2	2.44	0.48
33:S:87:ALA:HA	33:S:90:ARG:HG2	1.94	0.48
54:n:22:TYR:HE2	54:n:29:PRO:HD3	1.79	0.48
61:u:117:THR:HG23	61:u:118:THR:N	2.28	0.48
65:y:13:MET:HG3	65:y:77:HIS:CE1	2.49	0.48
66:z:74:ILE:HG13	66:z:78:LYS:HE3	1.95	0.48
2:1:71:VAL:O	2:1:71:VAL:HG23	2.13	0.48
7:6:19:DC:H2'	7:6:20:DA:C8	2.48	0.48
10:A:19:G:C6	41:a:2112:G:O5'	2.67	0.48
16:AG:27:LEU:HD22	16:AG:114:ILE:HG12	1.96	0.48
16:AG:148:ASP:HA	16:AG:164:ARG:CD	2.44	0.48
16:AG:209:PHE:CE1	16:AG:262:VAL:HG21	2.49	0.48
16:AG:220:VAL:CG1	16:AG:245:ILE:HG21	2.44	0.48
17:C:73:ARG:NH1	31:Q:113:VAL:CG1	2.76	0.48
18:D:744:C:H2'	18:D:745:G:H8	1.78	0.48
21:G:111:ILE:HD13	21:G:148:LEU:HD13	1.95	0.48
22:H:37:LEU:HD11	22:H:47:ALA:HA	1.96	0.48
22:H:70:VAL:HA	22:H:85:SER:HA	1.95	0.48
23:I:134:MET:HE2	23:I:168:TYR:CG	2.49	0.48
27:M:16:PRO:CB	29:O:46:MET:HG3	2.42	0.48
28:N:40:LEU:O	28:N:44:GLY:N	2.47	0.48
29:O:86:ALA:HA	29:O:89:GLU:HG2	1.95	0.48
29:O:118:LEU:HD22	29:O:123:ARG:C	2.39	0.48
36:V:42:THR:HG22	36:V:44:LEU:HD12	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:X:16:VAL:HG23	38:X:17:ILE:N	2.29	0.48
41:a:581:C:C2	41:a:582:A:C8	3.02	0.48
41:a:1048:A:C4	41:a:1049:C:C5	3.02	0.48
41:a:1666:G:H1'	60:t:3:GLN:NE2	2.29	0.48
41:a:1808:A:O2'	43:c:3:ARG:NH1	2.46	0.48
41:a:2314:A:C1'	54:n:155:THR:HG21	2.42	0.48
41:a:2683:C:OP1	65:y:51:ARG:NH2	2.46	0.48
48:h:141:VAL:HG23	48:h:162:VAL:HG12	1.94	0.48
51:k:19:HIS:CD2	51:k:20:PHE:N	2.82	0.48
52:l:6:LYS:HZ1	52:l:120:VAL:N	2.11	0.48
57:q:1:MET:CA	57:q:1:MET:HE3	2.43	0.48
62:v:95:LEU:HD23	62:v:95:LEU:H	1.78	0.48
63:w:33:ILE:HD11	63:w:112:TYR:CE1	2.49	0.48
5:4:21:ARG:HA	5:4:25:LYS:O	2.14	0.48
9:9:99:PHE:O	9:9:103:ASN:OD1	2.32	0.48
11:AA:836:LEU:HD13	11:AA:1054:LEU:HD13	1.95	0.48
12:AB:112:PRO:HA	12:AB:133:PRO:HG2	1.95	0.48
16:AG:87:LEU:HD22	16:AG:93:VAL:HG21	1.96	0.48
16:AG:288:MET:O	16:AG:288:MET:HG2	2.13	0.48
16:AG:299:ASP:CB	16:AG:303:HIS:HA	2.44	0.48
16:AG:342:ASP:N	16:AG:342:ASP:OD1	2.45	0.48
16:AG:389:MET:HA	16:AG:407:ARG:NH1	2.29	0.48
10:B:65:C:C2'	10:B:66:C:H5'	2.44	0.48
10:B:70:G:C3'	10:B:71:C:H5''	2.44	0.48
18:D:562:U:O3'	32:R:14:ARG:NH1	2.46	0.48
18:D:1172:C:H2'	18:D:1173:U:C6	2.49	0.48
18:D:1325:C:C2	18:D:1326:U:C5	3.02	0.48
19:E:19:LYS:CG	19:E:20:HIS:N	2.76	0.48
23:I:134:MET:HG3	23:I:151:VAL:CG2	2.44	0.48
25:K:112:ARG:C	25:K:116:GLU:OE1	2.57	0.48
25:K:157:ARG:NH1	28:N:43:GLU:HA	2.29	0.48
28:N:83:LEU:HD11	32:R:4:VAL:HB	1.95	0.48
30:P:5:ARG:HG3	30:P:7:ARG:HH12	1.78	0.48
30:P:64:GLN:OE1	30:P:65:TYR:O	2.31	0.48
33:S:74:LEU:HD23	33:S:77:PHE:CD2	2.49	0.48
41:a:624:C:O2'	41:a:657:U:OP1	2.29	0.48
41:a:2012:G:H8	41:a:2012:G:O5'	1.96	0.48
42:b:51:VAL:HG12	42:b:59:LEU:HD12	1.96	0.48
52:l:5:LEU:HD23	52:l:122:GLU:HG3	1.95	0.48
52:l:147:LEU:HD11	52:l:170:ARG:CD	2.43	0.48
60:t:88:ASN:OD1	60:t:89:ASN:N	2.46	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
62:v:58:LYS:O	62:v:59:ARG:HG3	2.13	0.48
64:x:115:LEU:HD23	64:x:117:PHE:HE1	1.79	0.48
4:3:28:VAL:HG23	4:3:34:VAL:HG12	1.96	0.48
6:5:10:DT:C4	11:AA:62:TYR:HE2	2.27	0.48
9:9:40:GLU:CD	9:9:54:VAL:HG11	2.39	0.48
10:A:12:G:C2'	10:A:13:C:OP1	2.61	0.48
12:AB:2:GLN:HB2	12:AB:60:ASP:HB2	1.95	0.48
12:AB:115:LYS:HZ2	12:AB:129:ILE:HG21	0.65	0.48
16:AG:243:LYS:O	16:AG:243:LYS:HD3	2.14	0.48
16:AG:442:ARG:C	16:AG:446:PHE:CD2	2.85	0.48
10:B:35:A:H2'	10:B:36:U:C6	2.49	0.48
18:D:330:C:O2	18:D:330:C:H2'	2.14	0.48
18:D:436:C:C2	18:D:437:U:C5	3.02	0.48
18:D:864:A:H5'	25:K:90:THR:O	2.14	0.48
20:F:62:ARG:HD2	20:F:63:GLU:N	2.29	0.48
24:J:113:GLU:O	24:J:117:LEU:HG	2.13	0.48
26:L:38:ARG:HD2	26:L:40:GLU:OE2	2.13	0.48
29:O:49:ARG:C	29:O:53:GLU:OE1	2.57	0.48
34:T:71:LYS:HE2	34:T:78:TYR:CE2	2.48	0.48
36:V:62:ARG:C	36:V:73:TRP:CE3	2.92	0.48
38:X:94:GLY:C	38:X:95:LEU:HD22	2.38	0.48
39:Y:19:PRO:HG2	39:Y:24:GLY:H	1.79	0.48
39:Y:78:LEU:HD12	39:Y:112:LYS:HZ1	1.77	0.48
41:a:272:A:C2	41:a:273:G:C5	3.02	0.48
41:a:880:G:C2	41:a:881:G:C8	3.02	0.48
41:a:1932:A:H2'	41:a:1933:G:O4'	2.13	0.48
41:a:2259:U:C2	41:a:2260:C:C5	3.01	0.48
41:a:2299:U:H2'	41:a:2300:C:C6	2.49	0.48
41:a:2591:C:H2'	41:a:2592:G:H8	1.78	0.48
41:a:2771:C:C2	41:a:2772:C:C5	3.01	0.48
43:c:67:VAL:HA	43:c:70:GLU:OE1	2.14	0.48
44:d:42:C:C5	54:n:66:LEU:HD12	2.49	0.48
46:f:3:LYS:HE2	46:f:3:LYS:HA	1.95	0.48
50:j:124:ARG:HA	50:j:165:MET:CE	2.44	0.48
52:l:141:MET:O	52:l:142:ALA:HB3	2.13	0.48
58:r:18:GLN:HE22	58:r:39:ALA:HB2	1.79	0.48
62:v:41:LEU:HG	62:v:96:ILE:HG13	1.95	0.48
65:y:9:GLU:HG3	65:y:55:LEU:HB2	1.95	0.48
4:3:40:ASN:OD1	4:3:65:ILE:HB	2.14	0.47
5:4:77:VAL:HG13	5:4:77:VAL:O	2.14	0.47
7:6:10:DG:OP1	14:AE:311:ARG:NH2	2.47	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:9:38:MET:SD	39:Y:117:THR:CG2	3.02	0.47
9:9:103:ASN:HA	9:9:107:GLU:HG3	1.96	0.47
12:AB:50:LEU:HD21	14:AE:285:LEU:CB	2.44	0.47
12:AB:144:ASN:O	30:P:88:MET:SD	2.70	0.47
13:AD:15:ASP:OD1	13:AD:27:THR:OG1	2.30	0.47
14:AE:437:PHE:HZ	14:AE:453:VAL:HG11	1.79	0.47
14:AE:804:ALA:O	14:AE:806:ASP:N	2.47	0.47
16:AG:429:LYS:HD2	16:AG:429:LYS:N	2.24	0.47
18:D:611:C:C2	18:D:612:C:C5	3.02	0.47
18:D:626:G:O3'	35:U:51:ARG:NH1	2.45	0.47
18:D:687:A:C2	18:D:704:A:C4	3.02	0.47
21:G:26:LYS:O	21:G:26:LYS:HD3	2.14	0.47
21:G:110:SER:HA	21:G:113:ARG:HH21	1.79	0.47
22:H:5:PHE:CE1	22:H:9:PHE:HE1	2.31	0.47
23:I:132:ARG:CA	23:I:135:LYS:HE3	2.44	0.47
23:I:156:ARG:HG2	23:I:193:TYR:HB2	1.95	0.47
25:K:148:ASN:ND2	25:K:153:VAL:CG1	2.77	0.47
25:K:157:ARG:NH1	28:N:43:GLU:CD	2.72	0.47
28:N:3:MET:O	28:N:3:MET:HE3	2.14	0.47
28:N:34:VAL:HG22	28:N:59:LEU:HD21	1.96	0.47
39:Y:19:PRO:HG2	39:Y:23:VAL:CG2	2.42	0.47
39:Y:56:VAL:HA	39:Y:71:LYS:NZ	2.08	0.47
41:a:576:U:H2'	41:a:577:G:C8	2.48	0.47
41:a:1841:U:C2	41:a:1842:G:C8	3.02	0.47
41:a:2070:A:H2'	41:a:2071:A:O4'	2.14	0.47
41:a:2314:A:P	54:n:88:LYS:NZ	2.87	0.47
41:a:2685:G:OP1	60:t:78:ARG:NH2	2.47	0.47
56:p:69:ARG:NH1	56:p:73:ASN:HD21	2.11	0.47
4:3:84:GLY:HA3	4:3:95:PHE:CZ	2.48	0.47
9:9:3:LEU:O	9:9:4:ASN:OD1	2.32	0.47
11:AA:318:SER:OG	11:AA:320:ASP:OD1	2.31	0.47
13:AC:102:LEU:HD12	13:AC:115:ILE:HG12	1.96	0.47
14:AE:1108:GLN:HE21	14:AE:1123:ARG:HH12	1.62	0.47
16:AG:61:ARG:HD2	16:AG:73:LYS:CD	2.44	0.47
16:AG:305:MET:HE3	16:AG:305:MET:HB3	1.77	0.47
16:AG:361:LYS:HD2	16:AG:417:ILE:HG12	1.92	0.47
16:AG:437:LEU:CG	16:AG:456:LEU:CD1	2.92	0.47
10:B:26:G:H1	10:B:44:A:N6	2.12	0.47
18:D:375:U:H1'	35:U:28:ARG:HH22	1.79	0.47
18:D:1397:C:OP1	18:D:1397:C:H2'	2.13	0.47
18:D:1412:C:H2'	18:D:1413:A:C8	2.49	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:G:117:LEU:HD21	21:G:141:LEU:HB2	1.97	0.47
25:K:156:LYS:HE3	28:N:71:VAL:CG1	2.41	0.47
32:R:74:LEU:HD11	32:R:80:ILE:HG13	1.96	0.47
41:a:6:A:O3'	59:s:132:HIS:CE1	2.67	0.47
41:a:39:G:H1'	52:l:43:THR:HG21	1.95	0.47
41:a:689:A:H2'	41:a:690:G:C8	2.50	0.47
41:a:1141:U:H4'	41:a:1142:A:O5'	2.13	0.47
41:a:1333:G:C2	41:a:1334:G:C8	3.03	0.47
41:a:1656:C:P	50:j:141:ARG:HD2	2.55	0.47
41:a:1790:C:H2'	41:a:1791:A:N7	2.29	0.47
41:a:2345:G:C5	41:a:2347:C:C5	3.02	0.47
45:e:2:LYS:HZ1	45:e:49:ASP:C	2.21	0.47
47:g:47:LYS:HE2	54:n:115:ARG:CZ	2.43	0.47
56:p:94:TYR:HD1	56:p:107:LEU:HD23	1.79	0.47
59:s:123:LYS:HD2	59:s:132:HIS:HD2	1.78	0.47
61:u:59:ARG:HH11	61:u:59:ARG:HG3	1.78	0.47
62:v:57:VAL:HG22	62:v:57:VAL:O	2.14	0.47
64:x:41:ALA:HB2	64:x:48:LEU:HD11	1.95	0.47
64:x:90:VAL:HG13	64:x:115:LEU:HD11	1.96	0.47
66:z:72:ASN:ND2	66:z:106:PHE:CE2	2.81	0.47
10:B:22:G:O2'	10:B:23:C:O5'	2.31	0.47
10:B:48:C:C2'	10:B:49:G:OP2	2.62	0.47
18:D:61:G:O6	19:E:10:ARG:NH2	2.47	0.47
18:D:72:A:C5	18:D:73:C:C5	3.02	0.47
18:D:377:G:OP1	35:U:5:ARG:CZ	2.60	0.47
18:D:610:U:O2	18:D:610:U:H2'	2.12	0.47
18:D:808:C:OP1	34:T:48:LYS:HE3	2.14	0.47
18:D:1176:A:H2'	18:D:1177:G:O4'	2.14	0.47
18:D:1307:U:H2'	18:D:1308:U:C6	2.49	0.47
19:E:9:LYS:O	19:E:12:ILE:HG12	2.14	0.47
27:M:71:PRO:O	27:M:96:ARG:NH1	2.47	0.47
27:M:95:ARG:CZ	27:M:99:LEU:HD21	2.45	0.47
32:R:99:ARG:NE	32:R:104:CYS:SG	2.87	0.47
41:a:963:U:H2'	41:a:964:C:H6	1.79	0.47
41:a:1442:U:H2'	41:a:1443:U:H6	1.79	0.47
41:a:1844:C:C2	41:a:1845:G:C8	3.02	0.47
41:a:2061:G:C2	41:a:2063:C:C5	3.01	0.47
41:a:2299:U:P	54:n:72:LYS:HE2	2.54	0.47
41:a:2313:C:H5''	54:n:88:LYS:HD2	1.95	0.47
41:a:2537:U:H2'	41:a:2538:C:H6	1.79	0.47
44:d:117:G:OP1	64:x:56:LYS:HD3	2.13	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:n:171:ALA:O	54:n:174:ASP:N	2.47	0.47
57:q:14:CYS:SG	57:q:33:HIS:HB3	2.53	0.47
62:v:53:MET:HB2	62:v:116:ALA:HB1	1.96	0.47
2:1:105:VAL:HG13	2:1:105:VAL:O	2.14	0.47
5:4:11:GLU:OE2	5:4:19:ARG:NH2	2.47	0.47
8:7:7:U:C6	8:7:7:U:C3'	2.97	0.47
9:9:69:PHE:C	9:9:72:LEU:HD11	2.40	0.47
12:AB:92:VAL:HG13	12:AB:96:LEU:O	2.13	0.47
12:AB:120:GLU:H	12:AB:162:LEU:HG	1.79	0.47
16:AG:444:LEU:HD22	16:AG:473:LEU:CD2	2.43	0.47
17:C:16:GLU:OE1	17:C:18:VAL:HB	2.14	0.47
18:D:35:G:N3	32:R:115:SER:OG	2.47	0.47
18:D:520:A:O2'	32:R:70:GLU:CD	2.58	0.47
18:D:983:A:H5'	18:D:984:C:OP2	2.15	0.47
18:D:1227:A:C2	18:D:1228:C:C6	3.02	0.47
18:D:1324:A:H2'	18:D:1325:C:H6	1.78	0.47
26:L:29:ILE:HG21	26:L:64:VAL:HG13	1.97	0.47
34:T:35:GLN:HB3	34:T:59:MET:HE1	1.95	0.47
36:V:11:ARG:HD2	36:V:56:GLY:C	2.39	0.47
38:X:86:TYR:HE2	38:X:90:ARG:NH2	2.11	0.47
41:a:644:A:H2'	41:a:645:C:O4'	2.15	0.47
41:a:1197:G:C4	41:a:1198:U:C5	3.02	0.47
41:a:1791:A:C2	41:a:1829:A:H4'	2.49	0.47
41:a:2298:A:H5''	54:n:72:LYS:CE	2.45	0.47
41:a:2300:C:H2'	41:a:2301:C:H6	1.79	0.47
41:a:2419:U:O2'	41:a:2420:C:H5'	2.14	0.47
47:g:45:THR:O	47:g:49:ARG:CZ	2.62	0.47
50:j:9:VAL:O	50:j:197:THR:HG23	2.15	0.47
53:m:25:LYS:HE2	53:m:28:ARG:HH21	1.78	0.47
56:p:8:PRO:C	56:p:69:ARG:HH21	2.23	0.47
62:v:59:ARG:CZ	62:v:59:ARG:O	2.63	0.47
5:4:51:GLN:OE1	5:4:57:TYR:OH	2.27	0.47
9:9:16:SER:HB2	9:9:20:LYS:NZ	2.29	0.47
10:A:48:C:C2'	10:A:49:G:OP2	2.62	0.47
10:A:65:C:C2'	10:A:66:C:H5'	2.44	0.47
10:A:75:C:O2'	10:A:76:A:N7	2.48	0.47
11:AA:1311:GLY:O	15:AF:31:GLN:NE2	2.47	0.47
11:AA:1314:GLN:HB2	15:AF:28:ARG:HH12	1.79	0.47
12:AB:141:LEU:CG	12:AB:143:LEU:CD1	2.88	0.47
13:AD:59:VAL:HG22	13:AD:144:ILE:HA	1.97	0.47
14:AE:797:THR:HG22	14:AE:924:GLY:HA3	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:61:ARG:HA	16:AG:94:GLU:HA	1.97	0.47
16:AG:314:LEU:HB3	16:AG:318:ILE:HD12	1.97	0.47
18:D:378:G:H2'	18:D:379:C:C6	2.49	0.47
21:G:19:GLN:OE1	21:G:21:ARG:CG	2.63	0.47
24:J:139:PRO:N	24:J:182:PHE:HE2	2.12	0.47
24:J:195:ILE:HG13	24:J:195:ILE:O	2.14	0.47
26:L:62:MET:HE3	26:L:62:MET:CA	2.43	0.47
36:V:27:ARG:HH12	36:V:42:THR:CB	2.26	0.47
39:Y:102:ARG:HH11	39:Y:141:ASP:CG	2.23	0.47
39:Y:126:ARG:NH1	41:a:1083:U:OP2	2.47	0.47
41:a:104:A:C5	41:a:105:C:C5	3.02	0.47
41:a:1936:A:N7	41:a:1945:G:C8	2.83	0.47
41:a:2043:C:C2	41:a:2044:C:C5	3.02	0.47
41:a:2327:A:H2'	41:a:2328:A:C8	2.49	0.47
41:a:2513:A:H2'	41:a:2514:U:C6	2.50	0.47
41:a:2748:A:C2	41:a:2757:A:C5	3.03	0.47
41:a:2803:G:H2'	41:a:2804:U:C5	2.50	0.47
52:l:42:GLY:HA3	52:l:92:HIS:CE1	2.49	0.47
56:p:9:VAL:HG23	56:p:69:ARG:HD2	1.95	0.47
56:p:18:LYS:HE2	56:p:18:LYS:HA	1.96	0.47
2:1:23:LEU:HD21	2:1:35:ILE:CG2	2.44	0.47
2:1:28:LYS:HB2	2:1:31:GLN:OE1	2.14	0.47
5:4:2:PHE:CE1	5:4:56:PHE:CZ	3.03	0.47
8:7:13:U:H3	24:J:47:ARG:NH1	2.10	0.47
10:A:47:U:H2'	10:A:48:C:H4'	1.97	0.47
11:AA:103:VAL:HG12	11:AA:117:ILE:HG22	1.97	0.47
14:AE:1350:ASN:HA	14:AE:1353:VAL:HG12	1.97	0.47
16:AG:1:MET:SD	16:AG:53:SER:HA	2.55	0.47
18:D:636:U:H5'	36:V:6:ARG:NE	2.29	0.47
18:D:642:A:H2'	18:D:643:C:H6	1.80	0.47
18:D:1225:A:H2'	18:D:1226:C:C5	2.49	0.47
18:D:1308:U:P	38:X:98:ARG:CG	3.03	0.47
18:D:1323:G:H2'	18:D:1324:A:H8	1.78	0.47
18:D:1359:C:OP2	33:S:75:ARG:NE	2.42	0.47
21:G:110:SER:CA	21:G:113:ARG:HH21	2.27	0.47
23:I:25:ASN:O	23:I:27:LYS:N	2.47	0.47
25:K:96:MET:HG2	25:K:125:ALA:HB2	1.97	0.47
29:O:7:TYR:O	29:O:89:GLU:CD	2.58	0.47
31:Q:85:MET:HB3	31:Q:113:VAL:HG11	1.95	0.47
33:S:63:ARG:HD3	33:S:68:GLY:O	2.15	0.47
34:T:15:PHE:CD2	34:T:30:ALA:HB2	2.49	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:T:77:ARG:NH2	34:T:80:GLN:NE2	2.62	0.47
36:V:42:THR:HG22	36:V:44:LEU:HD11	1.95	0.47
41:a:626:A:C2	61:u:78:ARG:CZ	2.98	0.47
41:a:1069:A:O2'	41:a:1073:A:N6	2.38	0.47
41:a:2190:G:H2'	41:a:2191:A:C8	2.49	0.47
41:a:2362:C:OP2	55:o:40:ARG:CZ	2.63	0.47
42:b:45:PHE:HD2	42:b:80:ILE:HD11	1.77	0.47
47:g:5:ILE:CD1	54:n:64:LYS:CE	2.92	0.47
52:l:148:ILE:CD1	52:l:187:VAL:HB	2.44	0.47
62:v:30:SER:N	62:v:104:GLU:OE2	2.48	0.47
62:v:111:GLU:O	62:v:112:LEU:C	2.57	0.47
1:0:73:LYS:HE2	1:0:73:LYS:HA	1.95	0.47
2:1:25:ARG:HH12	41:a:520:G:P	2.37	0.47
3:2:1:MET:HE1	41:a:139:U:C4	2.50	0.47
4:3:96:PHE:CE1	4:3:103:ILE:HG12	2.50	0.47
10:A:52:G:C6	10:A:63:G:O6	2.68	0.47
11:AA:689:ALA:HB2	11:AA:1233:LEU:HD23	1.97	0.47
11:AA:861:ALA:H	16:AG:38:LYS:HZ1	0.48	0.47
11:AA:889:PRO:HA	16:AG:104:ARG:O	2.15	0.47
12:AB:115:LYS:CD	12:AB:129:ILE:CB	2.92	0.47
14:AE:66:LYS:HB2	14:AE:69:GLU:HB3	1.96	0.47
14:AE:77:ARG:HA	14:AE:77:ARG:CZ	2.44	0.47
14:AE:86:GLU:N	14:AE:86:GLU:CD	2.73	0.47
14:AE:616:PRO:HA	14:AE:619:ILE:HG22	1.96	0.47
16:AG:353:HIS:O	16:AG:356:ILE:CA	2.63	0.47
16:AG:479:GLY:O	16:AG:483:MET:HG2	2.15	0.47
10:B:68:C:H3'	10:B:69:C:H5''	1.97	0.47
17:C:65:LEU:HD23	17:C:67:LEU:HD11	1.96	0.47
18:D:306:A:C5	18:D:307:C:C5	3.02	0.47
18:D:331:G:H2'	19:E:4:ILE:HD11	1.95	0.47
18:D:737:C:C2	18:D:738:C:C5	3.03	0.47
18:D:745:G:H2'	18:D:746:A:C8	2.50	0.47
18:D:1458:G:H5''	19:E:26:SER:OG	2.14	0.47
20:F:33:ARG:O	20:F:36:GLU:HG2	2.15	0.47
21:G:131:LYS:HE3	21:G:132:LYS:NZ	2.29	0.47
23:I:107:ARG:HB3	23:I:108:LYS:HD2	1.97	0.47
24:J:61:VAL:HA	24:J:64:ILE:HD12	1.96	0.47
24:J:62:ARG:CZ	24:J:69:GLU:HG2	2.44	0.47
24:J:76:TYR:CE2	24:J:204:TYR:CB	2.95	0.47
25:K:16:ILE:H	25:K:16:ILE:HD12	1.79	0.47
25:K:159:LYS:HZ1	28:N:44:GLY:HA3	1.80	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:L:9:MET:HE2	26:L:51:ILE:CD1	2.44	0.47
27:M:69:VAL:HA	27:M:138:ARG:HD3	1.97	0.47
27:M:129:GLU:O	27:M:130:ASN:HB2	2.13	0.47
28:N:112:THR:HG23	28:N:115:ALA:H	1.80	0.47
30:P:10:LEU:C	30:P:11:LYS:HD3	2.40	0.47
30:P:15:HIS:O	30:P:18:ILE:HG22	2.14	0.47
33:S:63:ARG:NH2	33:S:70:PRO:HD3	2.30	0.47
39:Y:106:GLN:OE1	39:Y:125:THR:HG21	2.14	0.47
41:a:247:G:H4'	41:a:386:G:C5	2.49	0.47
41:a:600:G:H2'	41:a:601:C:C6	2.50	0.47
41:a:783:A:N3	41:a:783:A:H2'	2.29	0.47
41:a:1065:U:HO2'	41:a:1066:U:C5'	2.26	0.47
41:a:1209:U:O2'	41:a:1237:A:N1	2.46	0.47
41:a:1770:G:C6	41:a:1983:G:C5	3.03	0.47
41:a:1858:A:N6	41:a:1884:G:H1'	2.29	0.47
41:a:2193:G:O2'	41:a:2194:U:P	2.72	0.47
41:a:2243:U:H2'	41:a:2244:U:H6	1.80	0.47
41:a:2298:A:C4	41:a:2321:U:H5	2.31	0.47
41:a:2784:U:H2'	41:a:2785:C:H6	1.80	0.47
42:b:26:PHE:O	42:b:29:GLU:HG2	2.15	0.47
51:k:25:LYS:HD3	51:k:51:GLU:OE2	2.15	0.47
54:n:38:MET:HE3	54:n:57:LEU:HG	1.96	0.47
58:r:58:LEU:O	58:r:61:VAL:HG22	2.14	0.47
59:s:72:LYS:HD3	59:s:74:TYR:HH	1.76	0.47
60:t:20:MET:HB3	60:t:44:LYS:HE2	1.95	0.47
61:u:19:LEU:CD2	61:u:27:LEU:HD12	2.45	0.47
63:w:44:LEU:C	63:w:44:LEU:HD23	2.40	0.47
63:w:56:LYS:NZ	63:w:90:ARG:O	2.48	0.47
2:1:50:VAL:HG12	2:1:105:VAL:CG1	2.45	0.47
3:2:18:GLU:O	3:2:22:THR:HG23	2.15	0.47
9:9:51:TYR:CD2	9:9:88:HIS:HB2	2.49	0.47
10:A:33:U:H4'	27:M:84:THR:HB	1.97	0.47
12:AB:111:TYR:HB3	12:AB:112:PRO:HD2	1.96	0.47
12:AB:141:LEU:O	12:AB:150:ILE:O	2.33	0.47
14:AE:290:ILE:O	14:AE:290:ILE:HG12	2.14	0.47
14:AE:640:GLY:N	14:AE:643:ASP:OD2	2.42	0.47
16:AG:26:ALA:HB1	16:AG:117:LYS:HG2	1.96	0.47
16:AG:233:ARG:HG2	16:AG:270:ARG:CB	2.45	0.47
10:B:48:C:O2'	10:B:49:G:OP2	2.27	0.47
18:D:771:G:C6	18:D:809:G:C6	3.03	0.47
18:D:821:G:H2'	18:D:822:U:H6	1.79	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:1149:C:H2'	18:D:1150:A:H8	1.79	0.47
21:G:110:SER:HA	21:G:113:ARG:NE	2.29	0.47
22:H:115:LYS:HA	22:H:119:GLY:C	2.40	0.47
27:M:66:LEU:HD13	27:M:101:MET:HE3	1.93	0.47
31:Q:20:VAL:HB	31:Q:35:THR:CG2	2.45	0.47
31:Q:116:ILE:HG13	31:Q:116:ILE:O	2.14	0.47
41:a:658:U:O2'	52:l:97:ASN:OD1	2.28	0.47
41:a:1349:C:N3	41:a:1350:C:C5	2.83	0.47
41:a:1385:A:C4	41:a:1386:C:C5	3.02	0.47
41:a:1429:G:H2'	41:a:1430:G:H8	1.79	0.47
41:a:2246:G:H2'	41:a:2247:A:C8	2.49	0.47
41:a:2373:G:H2'	41:a:2374:C:C6	2.49	0.47
41:a:2517:C:O2	41:a:2542:A:N6	2.48	0.47
48:h:138:GLY:H	48:h:164:ILE:HG23	1.79	0.47
51:k:7:GLU:HG2	51:k:27:LYS:HZ3	1.80	0.47
52:l:23:PHE:CE2	52:l:25:GLU:HA	2.50	0.47
52:l:112:LEU:HD21	52:l:186:VAL:HB	1.97	0.47
52:l:168:ASP:HB3	52:l:183:PHE:CE2	2.49	0.47
59:s:25:LEU:HD11	59:s:101:ILE:HD13	1.97	0.47
63:w:24:MET:HG3	63:w:44:LEU:HD12	1.97	0.47
3:2:56:GLU:OE2	3:2:88:LYS:CA	2.61	0.47
10:A:35:A:H2'	10:A:36:U:C6	2.49	0.47
11:AA:207:THR:OG1	11:AA:354:ASP:OD2	2.32	0.47
12:AB:7:LEU:HD22	12:AB:78:PHE:HA	1.96	0.47
14:AE:516:ASP:OD1	14:AE:516:ASP:N	2.47	0.47
16:AG:138:ILE:HD12	16:AG:183:LEU:CD1	2.45	0.47
16:AG:336:LEU:N	16:AG:336:LEU:CD1	2.73	0.47
10:B:75:C:O2'	10:B:76:A:N7	2.48	0.47
18:D:109:A:C6	18:D:326:G:C6	3.02	0.47
18:D:610:U:N3	18:D:611:C:C5	2.83	0.47
22:H:302:GLY:HA3	22:H:344:LEU:HD13	1.96	0.47
26:L:36:ILE:HD12	26:L:63:ASN:O	2.14	0.47
29:O:21:ILE:HD12	29:O:86:ALA:HB1	1.96	0.47
29:O:63:LEU:HD13	29:O:65:ILE:HD11	1.97	0.47
29:O:127:PHE:CD1	29:O:128:SER:N	2.83	0.47
32:R:2:ALA:N	32:R:7:LEU:HD11	2.29	0.47
32:R:5:ASN:HB3	32:R:9:ARG:HH22	1.80	0.47
36:V:31:HIS:HD2	36:V:34:TYR:H	1.62	0.47
37:W:9:PRO:HD3	47:g:64:PHE:CD2	2.49	0.47
41:a:909:A:H2'	41:a:912:C:H5	1.80	0.47
41:a:1006:C:C2	41:a:1138:G:N2	2.83	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1278:C:H2'	41:a:1279:G:H8	1.80	0.47
41:a:1467:U:C4	41:a:1546:G:C2	3.03	0.47
41:a:1794:A:H2'	41:a:1795:C:H6	1.79	0.47
41:a:2297:A:C6	41:a:2321:U:O4	2.68	0.47
41:a:2884:U:N3	49:i:40:ARG:NH1	2.62	0.47
49:i:4:GLN:OE1	49:i:8:PRO:CD	2.63	0.47
51:k:9:ILE:CG2	51:k:27:LYS:HZ1	2.27	0.47
54:n:93:GLY:C	54:n:97:TRP:CZ3	2.92	0.47
56:p:52:PHE:CZ	56:p:72:LEU:HD22	2.50	0.47
58:r:18:GLN:OE1	58:r:39:ALA:HB1	2.15	0.47
59:s:56:VAL:O	59:s:124:VAL:HA	2.14	0.47
59:s:136:GLN:N	59:s:137:PRO:CD	2.78	0.47
62:v:31:PHE:O	62:v:104:GLU:OE1	2.33	0.47
63:w:115:LEU:HD23	63:w:117:ASP:N	2.30	0.47
2:1:11:ARG:NH1	2:1:11:ARG:HB3	2.30	0.47
2:1:95:ARG:NH1	2:1:97:LEU:CD2	2.77	0.47
4:3:47:LYS:NZ	41:a:482:A:O5'	2.48	0.47
5:4:53:LYS:HD2	5:4:56:PHE:CZ	2.50	0.47
5:4:80:HIS:CD2	5:4:83:LYS:HZ3	2.32	0.47
9:9:38:MET:HA	9:9:41:LEU:HD12	1.96	0.47
12:AB:116:VAL:N	12:AB:129:ILE:CD1	2.78	0.47
14:AE:67:ASP:C	14:AE:69:GLU:N	2.71	0.47
10:B:70:G:C2'	10:B:71:C:H5''	2.45	0.47
18:D:216:U:H2'	18:D:217:C:C6	2.50	0.47
24:J:41:HIS:ND1	24:J:44:ARG:NH1	2.63	0.47
24:J:138:SER:O	24:J:139:PRO:C	2.57	0.47
27:M:69:VAL:HG23	27:M:100:ALA:HB1	1.97	0.47
29:O:89:GLU:O	29:O:92:GLU:OE1	2.32	0.47
36:V:7:THR:CG2	36:V:60:GLU:OE2	2.63	0.47
41:a:78:U:OP1	45:e:4:LYS:CE	2.63	0.47
41:a:400:G:OP2	43:c:57:ARG:NH1	2.46	0.47
41:a:584:C:N4	41:a:585:G:C6	2.83	0.47
41:a:1441:G:H2'	41:a:1442:U:C6	2.50	0.47
41:a:1477:A:N6	41:a:1514:G:O2'	2.45	0.47
41:a:2520:C:C6	41:a:2567:G:H1'	2.50	0.47
41:a:2788:C:H2'	41:a:2789:C:C6	2.49	0.47
45:e:12:GLU:CG	45:e:13:GLU:N	2.78	0.47
48:h:130:LEU:HG	48:h:135:ILE:HG13	1.96	0.47
49:i:55:ILE:HD13	63:w:112:TYR:CE1	2.50	0.47
55:o:62:LEU:HG	55:o:65:ALA:HA	1.96	0.47
60:t:59:LYS:HZ3	60:t:92:GLU:HG2	1.80	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
62:v:97:GLN:H	62:v:100:LYS:NZ	2.13	0.47
1:0:71:LYS:HE2	1:0:90:ARG:HD2	1.97	0.46
3:2:5:GLU:HA	45:e:22:LEU:CD2	2.45	0.46
3:2:57:VAL:HG12	3:2:86:THR:OG1	2.15	0.46
5:4:23:ALA:O	5:4:24:ASN:OD1	2.33	0.46
5:4:48:MET:HE3	5:4:85:LYS:HA	1.97	0.46
10:A:26:G:H1	10:A:44:A:N6	2.12	0.46
11:AA:400:VAL:HG22	11:AA:584:TYR:HB3	1.96	0.46
12:AB:131:THR:HG23	12:AB:142:LEU:HD22	1.97	0.46
16:AG:207:GLU:HG3	16:AG:210:ARG:HB3	1.96	0.46
18:D:35:G:H2'	18:D:36:C:H6	1.80	0.46
18:D:307:C:O2	18:D:307:C:H2'	2.15	0.46
18:D:829:G:O3'	21:G:23:TRP:HZ2	1.95	0.46
18:D:977:A:H1'	18:D:982:U:O4	2.15	0.46
18:D:1382:C:N3	18:D:1383:C:C5	2.83	0.46
20:F:19:PHE:HE2	31:Q:110:ILE:C	2.23	0.46
23:I:181:ASP:HB2	23:I:207:ILE:HD11	1.97	0.46
23:I:183:ASP:HB3	23:I:204:LYS:HZ2	1.80	0.46
27:M:46:ALA:O	27:M:49:THR:OG1	2.25	0.46
27:M:66:LEU:CD1	27:M:101:MET:CE	2.84	0.46
28:N:6:PRO:O	28:N:33:LYS:NZ	2.48	0.46
30:P:13:PHE:CE1	30:P:67:ILE:HD11	2.50	0.46
38:X:5:ALA:HB3	38:X:66:GLU:OE2	2.14	0.46
39:Y:96:LYS:HD2	39:Y:136:GLY:O	2.15	0.46
41:a:396:G:C6	41:a:397:U:C4	3.04	0.46
41:a:654:A:N6	55:o:19:LYS:HD2	2.29	0.46
41:a:1499:C:C2	41:a:1500:G:C8	3.03	0.46
41:a:2197:U:C5	41:a:2224:G:C6	3.03	0.46
41:a:2236:U:H2'	41:a:2237:G:O4'	2.15	0.46
41:a:2294:G:OP1	64:x:98:GLN:OE1	2.33	0.46
41:a:2454:G:C6	41:a:2499:C:N4	2.83	0.46
41:a:2725:A:C4	41:a:2727:A:N7	2.83	0.46
45:e:20:ASN:HA	45:e:23:ARG:NE	2.30	0.46
47:g:8:LYS:NZ	47:g:10:GLU:HB2	2.30	0.46
49:i:52:ARG:HA	49:i:52:ARG:NE	2.31	0.46
50:j:1:MET:O	50:j:204:LYS:HD2	2.15	0.46
54:n:12:VAL:O	54:n:16:LEU:HG	2.14	0.46
58:r:31:VAL:HB	58:r:32:PRO:HD3	1.97	0.46
60:t:24:VAL:HG23	60:t:33:ALA:HB2	1.97	0.46
62:v:29:GLY:H	62:v:104:GLU:HG2	1.79	0.46
66:z:113:ALA:O	66:z:117:LEU:HD13	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:4:53:LYS:HD2	5:4:56:PHE:HE2	1.77	0.46
5:4:53:LYS:HB3	5:4:55:GLU:OE1	2.16	0.46
5:4:53:LYS:O	5:4:56:PHE:HD2	1.97	0.46
11:AA:848:GLU:HG2	11:AA:888:THR:HG22	1.96	0.46
14:AE:513:MET:HE2	14:AE:513:MET:HB3	1.79	0.46
16:AG:233:ARG:O	16:AG:327:LEU:CD2	2.62	0.46
16:AG:285:ILE:O	16:AG:288:MET:HE3	2.14	0.46
10:B:24:U:H2'	10:B:25:C:C6	2.51	0.46
18:D:653:U:N1	28:N:56:LYS:HE3	2.29	0.46
18:D:1324:A:H2'	18:D:1325:C:C6	2.51	0.46
20:F:12:PHE:CE1	31:Q:107:ILE:CG2	2.98	0.46
20:F:16:LEU:HD12	31:Q:98:ARG:HH21	1.81	0.46
22:H:303:LEU:C	22:H:305:HIS:N	2.73	0.46
24:J:62:ARG:HA	24:J:72:PHE:CZ	2.50	0.46
24:J:62:ARG:NE	24:J:72:PHE:CE2	2.81	0.46
25:K:57:PRO:HA	25:K:60:ILE:HG12	1.96	0.46
25:K:165:LEU:HD12	28:N:114:ARG:NH1	2.30	0.46
32:R:2:ALA:CA	32:R:7:LEU:HD11	2.45	0.46
41:a:15:G:O2'	49:i:15:MET:HE2	2.15	0.46
41:a:590:A:H2'	41:a:591:U:H6	1.80	0.46
41:a:1223:G:N2	41:a:1226:A:OP2	2.43	0.46
41:a:1421:G:C2	41:a:1422:G:C8	3.03	0.46
41:a:1790:C:O3'	41:a:1791:A:C8	2.68	0.46
41:a:2294:G:OP1	64:x:10:ARG:HD3	2.15	0.46
41:a:2700:A:H2'	41:a:2701:U:C6	2.50	0.46
41:a:2814:A:C4	41:a:2815:C:C5	3.02	0.46
42:b:53:CYS:SG	42:b:57:HIS:HA	2.55	0.46
45:e:10:SER:OG	45:e:12:GLU:HG2	2.15	0.46
46:f:12:SER:C	46:f:54:MET:HE1	2.40	0.46
47:g:56:ARG:HH21	47:g:59:ARG:HH21	1.62	0.46
55:o:31:HIS:O	55:o:32:ILE:O	2.33	0.46
56:p:121:ILE:HD12	56:p:141:ILE:CG2	2.39	0.46
56:p:155:GLU:O	56:p:159:GLY:HA2	2.15	0.46
61:u:110:VAL:HG23	61:u:125:LEU:HD12	1.97	0.46
62:v:41:LEU:HD22	62:v:124:LEU:HD22	1.97	0.46
66:z:69:ALA:O	66:z:74:ILE:HG22	2.15	0.46
66:z:108:ALA:O	66:z:112:LYS:HG2	2.15	0.46
2:1:11:ARG:CZ	2:1:11:ARG:HB3	2.45	0.46
4:3:46:GLN:NE2	4:3:54:GLN:HG2	2.31	0.46
9:9:77:VAL:CG1	9:9:82:ILE:HG21	2.41	0.46
11:AA:28:LEU:HD21	11:AA:524:ILE:HG13	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AB:130:PHE:HA	12:AB:152:HIS:HE2	1.77	0.46
14:AE:1037:PHE:HB3	14:AE:1040:MET:HB2	1.97	0.46
18:D:1145:A:HO2'	18:D:1146:A:P	2.38	0.46
21:G:207:ILE:HG23	21:G:208:ARG:N	2.30	0.46
27:M:109:ARG:C	27:M:119:ARG:HH21	2.23	0.46
34:T:48:LYS:O	34:T:50:HIS:N	2.48	0.46
37:W:32:ARG:NH2	37:W:34:TRP:CZ3	2.84	0.46
41:a:67:U:N3	41:a:74:A:N6	2.62	0.46
41:a:257:C:O2	61:u:104:GLN:NE2	2.48	0.46
41:a:392:U:C2	41:a:393:C:C5	3.04	0.46
41:a:609:A:H3'	41:a:610:C:H6	1.80	0.46
41:a:1786:A:H1'	41:a:1938:A:N6	2.30	0.46
41:a:1796:U:H2'	41:a:1797:G:C8	2.50	0.46
41:a:2351:G:O6	55:o:39:LYS:NZ	2.48	0.46
41:a:2646:C:O5'	41:a:2646:C:H6	1.98	0.46
46:f:41:THR:O	46:f:42:PRO:C	2.58	0.46
47:g:16:CYS:SG	47:g:37:CYS:N	2.89	0.46
50:j:5:VAL:HG22	50:j:202:ILE:CD1	2.45	0.46
57:q:1:MET:HE3	57:q:1:MET:HA	1.97	0.46
60:t:76:VAL:HG12	65:y:73:VAL:HB	1.97	0.46
1:0:4:VAL:CG2	1:0:40:MET:HB3	2.45	0.46
3:2:64:LYS:HD2	41:a:1312:U:C4	2.51	0.46
3:2:69:ARG:HG3	3:2:74:ILE:HD13	1.96	0.46
9:9:6:GLN:CA	9:9:9:GLN:NE2	2.78	0.46
9:9:18:VAL:CG1	9:9:71:CYS:HB2	2.45	0.46
9:9:59:LEU:HB2	9:9:62:ARG:CG	2.45	0.46
10:A:30:G:O2'	10:A:31:G:H5'	2.15	0.46
11:AA:1286:THR:O	11:AA:1290:MET:HB2	2.14	0.46
12:AB:6:LEU:CD1	14:AE:291:ILE:CD1	2.84	0.46
12:AB:141:LEU:HB2	12:AB:153:SER:OG	2.16	0.46
14:AE:287:ALA:HB3	14:AE:292:VAL:CG2	2.46	0.46
16:AG:183:LEU:CD2	16:AG:197:VAL:HG22	2.45	0.46
16:AG:434:LEU:HD11	16:AG:454:CYS:C	2.39	0.46
10:B:46:G:H1'	10:B:47:U:C5	2.51	0.46
10:B:75:C:P	41:a:2602:A:H2'	2.55	0.46
18:D:197:A:C6	18:D:221:C:H4'	2.50	0.46
18:D:928:G:H2'	18:D:929:G:H8	1.81	0.46
18:D:1055:A:H1'	23:I:156:ARG:NH2	2.31	0.46
18:D:1189:U:OP1	33:S:98:LYS:NZ	2.48	0.46
18:D:1308:U:H3'	38:X:98:ARG:NH2	2.29	0.46
22:H:277:GLY:CA	22:H:331:MET:CE	2.93	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:H:287:LEU:HG	22:H:292:CYS:HA	1.97	0.46
26:L:17:GLN:C	26:L:21:MET:HE1	2.40	0.46
26:L:36:ILE:HD12	26:L:36:ILE:HA	1.81	0.46
27:M:111:ARG:O	27:M:119:ARG:CZ	2.63	0.46
30:P:27:GLU:HA	30:P:30:LYS:HZ1	1.79	0.46
33:S:79:LEU:HB2	33:S:84:VAL:CG2	2.45	0.46
34:T:11:ILE:HG21	34:T:31:LEU:CD1	2.45	0.46
34:T:29:VAL:HG22	34:T:66:LEU:HD23	1.96	0.46
34:T:47:LYS:HD3	34:T:47:LYS:N	2.30	0.46
39:Y:5:GLN:NE2	39:Y:60:VAL:C	2.74	0.46
41:a:592:A:C2	55:o:4:ILE:HD11	2.49	0.46
41:a:872:U:H2'	41:a:873:C:C6	2.50	0.46
41:a:923:G:H4'	42:b:29:GLU:OE1	2.16	0.46
41:a:1656:C:H2'	41:a:1657:U:H6	1.81	0.46
41:a:1708:C:H2'	41:a:1709:U:H6	1.79	0.46
41:a:2296:U:H4'	41:a:2297:A:OP1	2.14	0.46
41:a:2529:G:N7	57:q:32:LYS:NZ	2.63	0.46
41:a:2616:C:N3	41:a:2617:U:C5	2.83	0.46
41:a:2683:C:H4'	50:j:13:ARG:NH1	2.31	0.46
49:i:40:ARG:HH12	49:i:41:HIS:CD2	2.34	0.46
52:l:59:PRO:HB2	52:l:60:TRP:CE3	2.50	0.46
52:l:194:LYS:HG3	52:l:198:GLU:OE2	2.16	0.46
62:v:41:LEU:HG	62:v:96:ILE:CG1	2.45	0.46
66:z:69:ALA:HA	66:z:106:PHE:HZ	1.80	0.46
1:O:1:MET:O	1:O:15:SER:OG	2.19	0.46
8:7:44:U:H3'	8:7:44:U:H6	1.80	0.46
9:9:6:GLN:HA	9:9:9:GLN:HE22	1.74	0.46
11:AA:735:LYS:HA	11:AA:748:ILE:HG22	1.97	0.46
12:AB:9:CYS:N	12:AB:53:ASN:HB3	2.24	0.46
14:AE:81:ARG:CZ	16:AG:144:LYS:HE2	2.45	0.46
16:AG:63:LEU:HD22	16:AG:92:TYR:CD1	2.48	0.46
16:AG:302:LYS:HZ1	16:AG:302:LYS:C	2.10	0.46
17:C:60:LYS:HE3	18:D:735:C:H4'	1.97	0.46
18:D:915:A:C5	18:D:916:U:C6	3.04	0.46
18:D:1328:C:O3'	38:X:28:THR:HB	2.16	0.46
18:D:1363:A:C5	18:D:1365:G:C6	3.03	0.46
20:F:12:PHE:CD1	31:Q:101:ASN:ND2	2.84	0.46
21:G:19:GLN:NE2	22:H:75:VAL:HG13	2.30	0.46
25:K:143:GLY:O	25:K:146:ASN:OD1	2.34	0.46
41:a:132:G:H2'	41:a:133:U:H6	1.80	0.46
41:a:493:G:H2'	41:a:494:G:O4'	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:984:A:N3	41:a:984:A:H2'	2.29	0.46
41:a:1405:U:H2'	41:a:1406:U:C5	2.49	0.46
41:a:2013:A:C2	41:a:2613:U:O4	2.68	0.46
41:a:2200:C:OP2	43:c:37:ARG:NH2	2.48	0.46
41:a:2840:C:C2	41:a:2841:C:C5	3.03	0.46
44:d:83:G:H1'	46:f:53:PHE:CE1	2.51	0.46
48:h:181:MET:HE2	48:h:181:MET:CA	2.45	0.46
50:j:55:LYS:NZ	50:j:56:LYS:O	2.49	0.46
5:4:50:MET:C	5:4:56:PHE:CZ	2.94	0.46
9:9:11:ILE:HG12	9:9:65:GLU:HG2	1.98	0.46
10:A:6:G:O2'	10:A:7:G:H8	1.98	0.46
10:A:24:U:H2'	10:A:25:C:C6	2.51	0.46
10:A:46:G:H1'	10:A:47:U:C5	2.51	0.46
11:AA:856:ASN:CA	16:AG:35:THR:CB	2.89	0.46
12:AB:6:LEU:N	12:AB:88:VAL:HG21	2.31	0.46
15:AF:25:ARG:NH1	15:AF:61:ASN:OD1	2.49	0.46
10:B:30:G:O2'	10:B:31:G:H5'	2.15	0.46
10:B:47:U:H2'	10:B:48:C:H4'	1.97	0.46
17:C:31:ASN:OD1	22:H:268:VAL:HG12	2.15	0.46
18:D:22:G:C5	18:D:23:C:C5	3.03	0.46
18:D:440:C:C2	18:D:441:A:C8	3.04	0.46
18:D:696:A:H2'	18:D:697:U:H6	1.81	0.46
18:D:878:A:P	28:N:80:ARG:HH11	2.39	0.46
18:D:1149:C:C2	18:D:1150:A:C8	3.03	0.46
18:D:1209:C:C2	18:D:1210:C:C5	3.04	0.46
20:F:63:GLU:HG2	20:F:67:ARG:CZ	2.46	0.46
22:H:303:LEU:O	22:H:304:VAL:C	2.59	0.46
23:I:142:MET:HE3	23:I:170:GLU:HB3	1.98	0.46
28:N:8:ALA:N	28:N:77:ARG:HH11	2.14	0.46
30:P:63:ASP:OD2	30:P:63:ASP:C	2.59	0.46
36:V:30:LYS:HD3	36:V:35:GLY:HA2	1.97	0.46
36:V:61:ILE:C	36:V:73:TRP:HZ3	2.23	0.46
37:W:63:THR:HG22	37:W:64:ASP:N	2.30	0.46
37:W:77:THR:OG1	37:W:78:ARG:HD2	2.15	0.46
41:a:57:C:C2	41:a:58:G:C8	3.04	0.46
41:a:921:C:C2	41:a:922:C:C5	3.03	0.46
41:a:1019:U:O4	41:a:1142:A:C2	2.69	0.46
41:a:1068:G:O6	41:a:1069:A:N6	2.48	0.46
41:a:1200:C:C2	41:a:1201:U:C5	3.03	0.46
45:e:12:GLU:HG2	45:e:13:GLU:H	1.80	0.46
49:i:4:GLN:OE1	49:i:7:LYS:HA	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:l:35:TYR:CD2	52:l:178:VAL:HG11	2.51	0.46
1:0:82:HIS:CD2	61:u:23:ILE:HG12	2.50	0.46
8:7:8:U:OP2	18:D:1397:C:O2	2.33	0.46
9:9:94:ARG:CB	9:9:97:LYS:CE	2.94	0.46
11:AA:812:PHE:O	14:AE:504:GLN:NE2	2.40	0.46
12:AB:140:MET:CE	30:P:102:LEU:C	2.89	0.46
13:AD:35:PHE:HA	13:AD:38:THR:HG22	1.97	0.46
16:AG:36:LYS:HB2	16:AG:101:THR:HG21	1.97	0.46
16:AG:197:VAL:HG11	16:AG:199:ARG:NH2	2.30	0.46
10:B:52:G:C6	10:B:63:G:O6	2.68	0.46
17:C:12:ARG:NH2	17:C:28:THR:CG2	2.79	0.46
18:D:13:U:C4	18:D:21:G:C2	3.03	0.46
18:D:110:C:H1'	35:U:25:ARG:NH1	2.31	0.46
18:D:110:C:O4'	35:U:25:ARG:NH2	2.49	0.46
18:D:413:G:N2	18:D:428:G:H1'	2.31	0.46
18:D:520:A:O3'	32:R:70:GLU:OE2	2.32	0.46
19:E:82:GLN:HA	19:E:85:LYS:NZ	2.31	0.46
20:F:37:PHE:HA	31:Q:123:PRO:HB2	1.97	0.46
20:F:64:ASN:HA	20:F:67:ARG:HD3	1.97	0.46
22:H:304:VAL:O	22:H:304:VAL:CG1	2.63	0.46
24:J:95:GLU:HG3	24:J:186:PRO:HG3	1.98	0.46
26:L:38:ARG:NE	26:L:97:THR:O	2.49	0.46
27:M:74:GLU:CG	27:M:91:VAL:CG1	2.94	0.46
27:M:134:ALA:HA	27:M:137:LYS:CD	2.46	0.46
33:S:20:TYR:CD2	33:S:55:SER:OG	2.67	0.46
41:a:136:G:C6	41:a:144:A:C6	3.04	0.46
41:a:374:A:C2	41:a:375:G:H1'	2.51	0.46
41:a:669:G:H2'	41:a:670:A:N7	2.31	0.46
41:a:1370:C:H2'	41:a:1371:G:O4'	2.16	0.46
41:a:2702:G:C5	41:a:2703:C:C5	3.03	0.46
44:d:57:A:C5	54:n:26:MET:CE	2.99	0.46
47:g:18:CYS:HB2	47:g:40:CYS:HB3	1.96	0.46
52:l:42:GLY:HA3	52:l:92:HIS:HE1	1.80	0.46
54:n:142:ASP:CB	54:n:145:LYS:HE2	2.40	0.46
56:p:85:LYS:HA	56:p:85:LYS:HD2	1.77	0.46
60:t:8:LEU:N	60:t:8:LEU:HD12	2.31	0.46
66:z:53:ARG:HG2	66:z:57:PHE:CE2	2.49	0.46
14:AE:218:THR:HA	14:AE:221:ILE:HG22	1.96	0.46
17:C:33:ILE:O	22:H:338:GLU:CG	2.64	0.46
18:D:826:C:O2	28:N:16:ASN:ND2	2.49	0.46
18:D:829:G:H4'	21:G:23:TRP:CH2	2.51	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:1086:U:O2	18:D:1086:U:H2'	2.16	0.46
18:D:1108:G:C5	18:D:1109:C:C5	3.04	0.46
18:D:1238:A:H2	18:D:1241:G:N3	2.13	0.46
19:E:54:MET:O	19:E:58:VAL:HG23	2.16	0.46
22:H:270:ILE:HG23	22:H:270:ILE:O	2.15	0.46
26:L:94:HIS:CG	26:L:95:ALA:H	2.34	0.46
27:M:108:ALA:O	27:M:119:ARG:NH2	2.45	0.46
27:M:116:MET:HA	27:M:119:ARG:HD3	1.97	0.46
30:P:10:LEU:HD22	30:P:22:THR:HG22	1.98	0.46
32:R:18:LYS:HD3	32:R:19:SER:O	2.16	0.46
34:T:15:PHE:CE2	34:T:30:ALA:HB1	2.51	0.46
34:T:75:VAL:HG13	34:T:76:ALA:N	2.31	0.46
41:a:6:A:H5'	59:s:131:ASN:OD1	2.15	0.46
41:a:396:G:C5	41:a:397:U:C5	3.03	0.46
41:a:753:A:C6	41:a:754:U:C5	3.04	0.46
41:a:1131:G:OP1	59:s:81:ILE:O	2.34	0.46
41:a:1406:U:O2'	41:a:1407:G:P	2.74	0.46
41:a:1433:A:H2'	41:a:1434:A:C1'	2.46	0.46
41:a:1598:A:C4	41:a:1599:U:C6	3.03	0.46
41:a:1613:G:O2'	53:m:3:ARG:CZ	2.64	0.46
41:a:1797:G:O2'	48:h:257:THR:HG22	2.16	0.46
41:a:2364:C:H2'	41:a:2365:G:O4'	2.15	0.46
48:h:108:LYS:HZ3	48:h:194:GLU:N	2.12	0.46
52:l:60:TRP:CD1	52:l:69:ARG:HH12	2.33	0.46
52:l:175:ILE:HG21	52:l:180:LEU:HD11	1.97	0.46
57:q:1:MET:HE3	57:q:2:LYS:N	2.31	0.46
60:t:1:MET:CE	60:t:32:TYR:CG	2.98	0.46
9:9:25:ALA:HB3	9:9:99:PHE:CE2	2.50	0.46
9:9:43:LYS:HE3	9:9:102:ALA:HB2	1.97	0.46
9:9:87:GLU:OE2	9:9:95:LEU:CG	2.64	0.46
11:AA:562:GLU:OE1	11:AA:662:SER:OG	2.28	0.46
13:AC:64:VAL:HG11	13:AC:78:ILE:HG21	1.97	0.46
13:AD:185:TYR:HA	13:AD:202:VAL:O	2.16	0.46
14:AE:420:PRO:HA	14:AE:437:PHE:O	2.16	0.46
14:AE:482:ALA:HA	15:AF:6:VAL:HG21	1.98	0.46
16:AG:227:ALA:CB	16:AG:331:LEU:HA	2.46	0.46
16:AG:425:LEU:C	16:AG:429:LYS:CE	2.74	0.46
16:AG:434:LEU:HD11	16:AG:455:THR:CA	2.46	0.46
10:B:30:G:HO2'	10:B:31:G:H5'	1.81	0.46
17:C:14:THR:HG21	17:C:48:ARG:CD	2.46	0.46
18:D:277:C:OP1	36:V:43:LYS:HD2	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:915:A:C6	18:D:916:U:C5	3.04	0.46
21:G:117:LEU:HD23	21:G:141:LEU:HA	1.98	0.46
21:G:197:ASP:OD2	21:G:198:PHE:CE2	2.68	0.46
22:H:277:GLY:HA2	22:H:331:MET:HE3	1.97	0.46
22:H:288:THR:O	22:H:288:THR:HG22	2.16	0.46
23:I:22:TRP:CZ2	23:I:32:ASN:HB2	2.50	0.46
23:I:134:MET:HE2	23:I:168:TYR:CB	2.46	0.46
25:K:12:GLN:HB3	25:K:14:LYS:HZ2	1.81	0.46
25:K:16:ILE:HD11	25:K:38:VAL:CG1	2.46	0.46
27:M:138:ARG:O	27:M:142:HIS:ND1	2.46	0.46
29:O:58:VAL:HG23	29:O:59:GLU:N	2.31	0.46
30:P:5:ARG:HE	30:P:7:ARG:CZ	2.29	0.46
30:P:13:PHE:CZ	30:P:67:ILE:HD11	2.50	0.46
32:R:56:ARG:HG2	32:R:62:GLU:CD	2.41	0.46
33:S:86:GLU:OE2	33:S:90:ARG:HD3	2.16	0.46
41:a:152:A:H2'	41:a:153:U:C6	2.51	0.46
41:a:404:A:H4'	41:a:405:U:O5'	2.15	0.46
41:a:1406:U:O2	41:a:1407:G:C8	2.69	0.46
41:a:1818:U:H6	48:h:156:ARG:NH2	2.13	0.46
41:a:2373:G:H2'	41:a:2374:C:H6	1.81	0.46
41:a:2766:A:C5	41:a:2767:C:C5	3.03	0.46
43:c:41:GLU:O	43:c:44:LYS:HD2	2.16	0.46
47:g:47:LYS:CD	54:n:178:ARG:HH22	2.29	0.46
47:g:58:ASP:C	47:g:62:LYS:HZ1	2.21	0.46
49:i:28:LEU:HB2	49:i:37:LYS:HE3	1.98	0.46
50:j:91:THR:HB	50:j:94:GLN:CG	2.44	0.46
65:y:103:ARG:HD3	65:y:107:ALA:C	2.41	0.46
3:2:64:LYS:CE	41:a:1312:U:O4	2.64	0.46
9:9:34:THR:HA	9:9:37:LYS:HZ3	1.81	0.46
10:A:70:G:C2'	10:A:71:C:H5''	2.45	0.46
11:AA:230:PHE:HB2	11:AA:333:ILE:HB	1.97	0.46
11:AA:1002:LEU:HD21	11:AA:1007:LYS:HB2	1.98	0.46
14:AE:287:ALA:CB	14:AE:292:VAL:CG2	2.93	0.46
14:AE:848:VAL:HB	14:AE:858:VAL:HG22	1.98	0.46
10:B:6:G:O2'	10:B:7:G:H8	1.98	0.46
18:D:371:A:H2'	18:D:372:C:O4'	2.15	0.46
18:D:736:C:H5'	26:L:90:MET:HE2	1.97	0.46
18:D:915:A:C5	18:D:916:U:C5	3.04	0.46
18:D:980:C:C5	18:D:981:U:C2	3.04	0.46
18:D:1175:G:HO2'	18:D:1176:A:P	2.38	0.46
18:D:1287:A:H2'	18:D:1288:A:C8	2.51	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:I:84:VAL:HG23	23:I:85:GLU:N	2.31	0.46
25:K:150:PRO:O	25:K:153:VAL:HG22	2.16	0.46
26:L:100:SER:HB2	26:L:101:PRO:HD2	1.98	0.46
27:M:44:TYR:C	27:M:44:TYR:CD2	2.94	0.46
30:P:59:LYS:HE2	30:P:62:ARG:NH2	2.31	0.46
33:S:6:MET:HE3	33:S:63:ARG:NH1	2.31	0.46
37:W:9:PRO:HB3	47:g:64:PHE:CZ	2.51	0.46
37:W:32:ARG:NH2	37:W:34:TRP:HZ3	2.14	0.46
38:X:104:THR:O	38:X:104:THR:HG22	2.16	0.46
39:Y:59:THR:O	39:Y:66:PHE:HB2	2.16	0.46
39:Y:79:LEU:O	39:Y:83:ALA:HB3	2.16	0.46
39:Y:102:ARG:HD3	39:Y:141:ASP:OD1	2.15	0.46
41:a:15:G:O2'	49:i:15:MET:HE1	2.16	0.46
41:a:372:G:O2'	43:c:54:LYS:CE	2.64	0.46
41:a:582:A:H5''	66:z:11:ARG:HH12	1.81	0.46
41:a:591:U:C2	41:a:592:A:C8	3.03	0.46
41:a:685:A:C8	41:a:773:U:C4	3.04	0.46
41:a:1585:C:H2'	41:a:1586:A:O4'	2.15	0.46
41:a:1707:G:C4	41:a:1708:C:C5	3.04	0.46
41:a:2357:G:P	42:b:20:ARG:CZ	3.04	0.46
41:a:2650:U:H2'	41:a:2651:C:H6	1.81	0.46
48:h:108:LYS:HZ3	48:h:194:GLU:H	1.64	0.46
52:l:6:LYS:CE	52:l:119:ILE:CG2	2.94	0.46
54:n:127:ASN:OD1	54:n:157:THR:HA	2.16	0.46
57:q:2:LYS:HE2	57:q:35:GLN:HG3	1.97	0.46
59:s:123:LYS:HD2	59:s:132:HIS:CD2	2.50	0.46
64:x:88:LYS:O	64:x:115:LEU:HD12	2.16	0.46
9:9:60:LEU:O	9:9:64:VAL:HB	2.15	0.45
9:9:122:GLN:OE1	9:9:122:GLN:N	2.49	0.45
12:AB:17:ALA:O	12:AB:20:HIS:HB2	2.17	0.45
14:AE:388:ARG:NH2	14:AE:414:GLU:OE1	2.49	0.45
16:AG:80:ALA:HB1	16:AG:87:LEU:HB2	1.98	0.45
16:AG:365:ILE:CD1	16:AG:406:LEU:HD21	2.37	0.45
17:C:12:ARG:NH1	17:C:13:PHE:CE1	2.84	0.45
17:C:34:THR:HB	17:C:38:LYS:H	1.81	0.45
18:D:878:A:P	28:N:80:ARG:NH1	2.89	0.45
18:D:1512:U:H2'	18:D:1513:A:C8	2.51	0.45
25:K:157:ARG:NH2	28:N:43:GLU:OE2	2.46	0.45
26:L:43:GLY:HA2	26:L:58:HIS:CE1	2.51	0.45
27:M:138:ARG:HG2	27:M:142:HIS:HE1	1.80	0.45
28:N:114:ARG:HD2	28:N:114:ARG:C	2.41	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:R:122:PRO:C	32:R:123:LYS:HE2	2.42	0.45
35:U:6:LEU:HD23	35:U:17:TYR:CB	2.47	0.45
37:W:44:MET:HE1	37:W:62:VAL:CG2	2.46	0.45
38:X:13:LYS:HZ2	38:X:17:ILE:HG21	1.81	0.45
39:Y:59:THR:C	39:Y:66:PHE:HB2	2.41	0.45
41:a:67:U:N3	41:a:68:G:C5	2.84	0.45
41:a:607:U:C6	41:a:620:G:N7	2.85	0.45
41:a:1083:U:O2'	41:a:1085:A:OP2	2.31	0.45
41:a:1398:C:C2	41:a:1399:C:C5	3.04	0.45
41:a:1803:A:O2'	48:h:257:THR:HG21	2.16	0.45
41:a:2478:A:C8	41:a:2529:G:C5	3.04	0.45
41:a:2743:U:O2'	56:p:153:ARG:NH1	2.49	0.45
41:a:2756:U:N3	41:a:2758:A:N6	2.64	0.45
52:l:130:LYS:HB2	52:l:133:LEU:HD12	1.97	0.45
55:o:55:LEU:CD2	55:o:59:ILE:HD11	2.45	0.45
58:r:72:ILE:HG22	58:r:72:ILE:O	2.16	0.45
4:3:49:VAL:HG23	4:3:51:ALA:O	2.17	0.45
6:5:10:DT:OP1	11:AA:476:LYS:HD2	2.16	0.45
9:9:3:LEU:HD12	9:9:5:LEU:H	1.81	0.45
9:9:4:ASN:OD1	41:a:1046:A:N7	2.50	0.45
10:A:19:G:O6	41:a:2112:G:O5'	2.34	0.45
11:AA:812:PHE:HZ	14:AE:503:SER:HB2	1.81	0.45
11:AA:1246:ARG:HH11	11:AA:1266:GLY:HA2	1.82	0.45
14:AE:53:ARG:HA	14:AE:54:ASP:HA	1.59	0.45
14:AE:781:LYS:HD2	14:AE:933:ARG:HH12	1.81	0.45
16:AG:33:THR:CG2	16:AG:43:ILE:HD11	2.44	0.45
16:AG:102:PHE:HB3	16:AG:103:ASP:CG	2.41	0.45
16:AG:402:THR:C	16:AG:405:ALA:HB3	2.41	0.45
16:AG:434:LEU:CD1	16:AG:455:THR:O	2.61	0.45
10:B:50:U:H2'	10:B:51:C:C6	2.51	0.45
17:C:21:ILE:CD1	17:C:54:GLN:NE2	2.79	0.45
18:D:13:U:O2	18:D:915:A:N7	2.49	0.45
18:D:151:A:C5	18:D:152:A:C8	3.04	0.45
18:D:827:U:O4	18:D:872:A:C2	2.69	0.45
18:D:1109:C:C2	18:D:1110:A:C8	3.04	0.45
18:D:1435:G:H2'	18:D:1436:U:C6	2.51	0.45
21:G:18:HIS:CE1	21:G:205:ASP:OD1	2.68	0.45
27:M:87:VAL:HG12	27:M:151:PHE:HD2	1.81	0.45
27:M:135:VAL:HG23	27:M:136:LYS:HD2	1.96	0.45
34:T:8:THR:O	34:T:11:ILE:HG22	2.16	0.45
38:X:56:LEU:HA	38:X:59:GLU:OE1	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:Y:126:ARG:HH12	41:a:1082:U:C5'	2.29	0.45
41:a:1794:A:H2'	41:a:1795:C:C6	2.50	0.45
41:a:2064:C:H2'	41:a:2065:C:H6	1.79	0.45
41:a:2289:G:C2	41:a:2290:G:C8	3.04	0.45
41:a:2723:C:H2'	41:a:2724:U:O4'	2.16	0.45
46:f:51:VAL:O	46:f:55:VAL:HG12	2.17	0.45
48:h:2:ALA:N	48:h:20:VAL:O	2.48	0.45
48:h:33:LEU:O	48:h:64:ILE:HG12	2.16	0.45
50:j:11:MET:HE3	50:j:11:MET:HB3	1.63	0.45
52:l:1:MET:CE	52:l:20:GLY:N	2.79	0.45
52:l:195:GLN:CA	52:l:198:GLU:OE1	2.64	0.45
54:n:16:LEU:HD11	54:n:169:LEU:HD13	1.99	0.45
59:s:55:ILE:HG13	59:s:123:LYS:CE	2.46	0.45
60:t:50:GLY:O	60:t:53:LYS:NZ	2.47	0.45
65:y:106:LYS:CD	65:y:109:ARG:HH22	2.29	0.45
1:0:4:VAL:HG23	1:0:39:LEU:HB2	1.97	0.45
2:1:20:VAL:CG1	2:1:47:VAL:HG11	2.46	0.45
3:2:3:ARG:CZ	3:2:5:GLU:OE1	2.64	0.45
6:5:10:DT:H2''	12:AB:71:ALA:H	1.79	0.45
9:9:61:ARG:CD	41:a:1046:A:O5'	2.64	0.45
10:A:32:C:C2'	10:A:33:U:H5'	2.46	0.45
11:AA:557:ARG:NH2	11:AA:607:SER:O	2.50	0.45
12:AB:16:ARG:C	12:AB:19:GLU:H	2.25	0.45
17:C:12:ARG:HB2	22:H:264:GLU:CB	2.46	0.45
17:C:63:ARG:HB3	17:C:70:TYR:CE1	2.51	0.45
18:D:269:C:H2'	18:D:270:A:C8	2.51	0.45
18:D:377:G:H2'	18:D:378:G:H8	1.81	0.45
18:D:983:A:H2	18:D:984:C:C6	2.34	0.45
18:D:1226:C:H4'	18:D:1227:A:OP1	2.16	0.45
21:G:100:MET:SD	21:G:101:LEU:HD23	2.57	0.45
22:H:24:VAL:HG23	22:H:24:VAL:O	2.16	0.45
32:R:68:GLY:O	32:R:99:ARG:NH1	2.49	0.45
34:T:3:LEU:CD2	34:T:8:THR:HG23	2.47	0.45
36:V:30:LYS:HB2	36:V:37:PHE:CE1	2.52	0.45
39:Y:102:ARG:HH21	39:Y:105:LEU:HD12	1.81	0.45
41:a:6:A:N3	59:s:135:GLN:NE2	2.59	0.45
41:a:182:A:H2'	41:a:183:C:H6	1.81	0.45
41:a:581:C:H2'	41:a:582:A:C8	2.50	0.45
41:a:666:A:H4'	61:u:48:ARG:HD2	1.97	0.45
41:a:893:C:H2'	41:a:894:U:C5	2.51	0.45
41:a:952:G:C6	41:a:966:G:C6	3.04	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:985:C:N3	41:a:986:C:C5	2.84	0.45
41:a:2193:G:HO2'	41:a:2194:U:P	2.40	0.45
41:a:2314:A:O2'	54:n:155:THR:HG21	2.15	0.45
41:a:2393:U:C5'	55:o:30:ARG:NH1	2.76	0.45
41:a:2755:C:H3'	57:q:19:ARG:NH2	2.31	0.45
44:d:44:G:H3'	54:n:92:ARG:HH22	1.81	0.45
44:d:82:U:O2'	46:f:53:PHE:HE2	1.98	0.45
47:g:1:MET:SD	47:g:3:LYS:HE2	2.56	0.45
47:g:30:HIS:CG	47:g:31:ASP:N	2.85	0.45
48:h:52:ARG:NE	48:h:53:HIS:CE1	2.83	0.45
54:n:4:LEU:HD21	54:n:101:GLU:HG3	1.98	0.45
56:p:69:ARG:NH1	56:p:73:ASN:CG	2.74	0.45
59:s:31:GLU:HG2	59:s:142:ILE:HG23	1.98	0.45
60:t:42:THR:HG22	60:t:44:LYS:HZ1	1.81	0.45
62:v:114:ARG:C	62:v:118:LYS:HZ3	2.23	0.45
63:w:48:VAL:CG1	63:w:49:GLU:OE1	2.64	0.45
64:x:15:ARG:C	64:x:19:GLN:HE22	2.24	0.45
7:6:25:DT:OP1	11:AA:496:LYS:O	2.34	0.45
8:7:47:U:H4'	14:AE:322:ARG:CZ	2.46	0.45
10:B:32:C:C2'	10:B:33:U:H5'	2.46	0.45
18:D:198:G:OP2	18:D:198:G:C8	2.70	0.45
18:D:1436:U:H2'	18:D:1437:A:C8	2.52	0.45
22:H:32:ASP:OD1	22:H:35:VAL:HG23	2.17	0.45
23:I:39:VAL:HG23	23:I:40:ARG:N	2.32	0.45
23:I:55:ILE:HD12	23:I:68:ILE:HG12	1.99	0.45
23:I:165:THR:C	23:I:166:GLU:OE1	2.60	0.45
26:L:85:ILE:HG22	26:L:86:ARG:HG2	1.99	0.45
27:M:78:ARG:CG	27:M:80:VAL:HG13	2.46	0.45
29:O:94:LEU:O	29:O:97:GLU:CD	2.60	0.45
30:P:26:VAL:C	30:P:30:LYS:HZ3	2.25	0.45
31:Q:84:VAL:HG21	31:Q:97:ILE:HD11	1.99	0.45
34:T:3:LEU:HD23	34:T:4:SER:O	2.15	0.45
37:W:5:LEU:HD12	37:W:6:LYS:N	2.32	0.45
41:a:152:A:H2'	41:a:153:U:H6	1.82	0.45
41:a:704:G:O2'	41:a:726:G:N2	2.45	0.45
41:a:1177:G:O2'	41:a:1178:C:O5'	2.28	0.45
43:c:43:GLU:OE1	43:c:45:ARG:CB	2.65	0.45
48:h:130:LEU:CA	48:h:189:ARG:HH11	2.24	0.45
50:j:74:GLU:HG2	50:j:75:ALA:N	2.31	0.45
63:w:37:THR:HG21	63:w:40:LYS:HZ2	1.81	0.45
64:x:35:ILE:HD11	64:x:102:ARG:HH21	1.81	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
64:x:83:LEU:HD23	64:x:88:LYS:HE3	1.97	0.45
65:y:7:GLN:HE21	65:y:10:GLN:HE21	1.65	0.45
4:3:4:LYS:O	4:3:94:ARG:NH2	2.40	0.45
10:A:18:G:N7	10:A:57:A:N6	2.65	0.45
12:AB:104:ILE:C	12:AB:106:ASP:N	2.72	0.45
12:AB:115:LYS:CD	12:AB:129:ILE:CD1	2.63	0.45
12:AB:140:MET:HE1	30:P:103:GLY:CA	2.47	0.45
13:AD:182:ARG:HD2	14:AE:581:MET:HE1	1.97	0.45
14:AE:1341:ARG:NH1	14:AE:1343:GLU:OE2	2.49	0.45
16:AG:288:MET:HE3	16:AG:289:ALA:N	2.32	0.45
16:AG:339:MET:HE2	16:AG:343:ASP:HB3	1.99	0.45
16:AG:403:VAL:C	16:AG:405:ALA:N	2.73	0.45
18:D:339:C:H2'	18:D:340:U:C6	2.50	0.45
18:D:611:C:C2	18:D:612:C:C6	3.04	0.45
18:D:739:C:O2'	34:T:42:HIS:CE1	2.68	0.45
18:D:1308:U:H3'	38:X:98:ARG:NE	2.32	0.45
18:D:1368:A:OP1	29:O:113:ARG:NH2	2.48	0.45
19:E:16:LYS:HG3	19:E:19:LYS:HE3	1.98	0.45
20:F:58:LYS:HD2	20:F:58:LYS:C	2.42	0.45
21:G:9:MET:HE1	21:G:47:VAL:HG22	1.99	0.45
21:G:77:SER:O	21:G:80:VAL:HG12	2.17	0.45
21:G:128:LYS:C	21:G:129:LEU:HD12	2.42	0.45
23:I:135:LYS:O	23:I:138:VAL:HG12	2.17	0.45
26:L:35:LYS:HD2	26:L:36:ILE:H	1.82	0.45
26:L:47:LEU:HD13	26:L:51:ILE:HG12	1.98	0.45
26:L:98:GLU:HA	26:L:98:GLU:OE1	2.16	0.45
26:L:99:ALA:HB3	26:L:104:LYS:HZ3	1.81	0.45
28:N:26:THR:OG1	28:N:58:GLU:CG	2.65	0.45
30:P:47:GLU:HB2	30:P:67:ILE:HG22	1.99	0.45
34:T:11:ILE:HD11	34:T:30:ALA:HB1	1.99	0.45
37:W:62:VAL:HG13	37:W:66:MET:CE	2.44	0.45
39:Y:126:ARG:NH2	41:a:1082:U:O3'	2.47	0.45
41:a:219:A:H2'	41:a:220:G:C8	2.52	0.45
41:a:391:A:C5	41:a:392:U:C5	3.04	0.45
41:a:730:A:C2	41:a:731:C:C6	3.05	0.45
41:a:783:A:O2'	41:a:785:G:OP1	2.35	0.45
41:a:832:U:H2'	41:a:833:A:C8	2.50	0.45
41:a:1042:G:C5	41:a:1043:C:C5	3.05	0.45
41:a:2313:C:O3'	54:n:88:LYS:NZ	2.48	0.45
45:e:24:GLU:O	45:e:25:GLN:C	2.57	0.45
52:l:21:ARG:HA	52:l:21:ARG:NE	2.31	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:n:10:ASP:OD1	54:n:11:GLU:HG3	2.17	0.45
55:o:26:HIS:CE1	55:o:48:ALA:HB2	2.51	0.45
55:o:30:ARG:HH12	61:u:62:PRO:N	2.15	0.45
60:t:108:ARG:CG	60:t:116:ILE:HD13	2.46	0.45
63:w:12:ARG:HE	63:w:20:MET:HE1	1.82	0.45
1:0:71:LYS:CG	1:0:73:LYS:HE3	2.47	0.45
2:1:20:VAL:HG11	2:1:47:VAL:HG11	1.99	0.45
4:3:47:LYS:HZ3	41:a:482:A:P	2.40	0.45
10:A:50:U:H2'	10:A:51:C:C6	2.51	0.45
11:AA:1314:GLN:HA	15:AF:28:ARG:HH22	1.80	0.45
14:AE:79:LYS:CD	16:AG:159:GLU:HB3	2.44	0.45
14:AE:1321:SER:OG	14:AE:1349:GLU:OE2	2.32	0.45
16:AG:228:ARG:HB3	16:AG:234:ALA:HA	1.98	0.45
18:D:984:C:C2	18:D:985:C:C5	3.05	0.45
18:D:1173:U:H2'	18:D:1174:G:O4'	2.16	0.45
19:E:49:LYS:O	19:E:53:GLU:HG2	2.16	0.45
22:H:54:LYS:HD3	22:H:58:GLY:C	2.42	0.45
22:H:297:GLU:OE1	22:H:297:GLU:N	2.49	0.45
25:K:145:GLU:CD	25:K:146:ASN:N	2.74	0.45
26:L:9:MET:C	26:L:9:MET:SD	3.00	0.45
26:L:17:GLN:CB	26:L:21:MET:HE1	2.47	0.45
27:M:4:ARG:HG2	27:M:5:ARG:NH2	2.32	0.45
29:O:113:ARG:HD2	33:S:101:TRP:CD1	2.52	0.45
30:P:64:GLN:HG2	33:S:101:TRP:CH2	2.52	0.45
32:R:31:ARG:O	32:R:57:LEU:HD12	2.17	0.45
39:Y:102:ARG:HE	39:Y:139:VAL:HG21	1.79	0.45
41:a:396:G:OP2	43:c:10:LYS:NZ	2.45	0.45
41:a:1212:G:O2'	41:a:1236:G:N2	2.47	0.45
41:a:1432:G:H2'	41:a:1433:A:H8	1.80	0.45
41:a:2061:G:N2	41:a:2063:C:C6	2.84	0.45
41:a:2065:C:C2	41:a:2066:C:C5	3.04	0.45
41:a:2572:A:OP2	50:j:149:ASN:O	2.35	0.45
41:a:2700:A:H2'	41:a:2701:U:H6	1.81	0.45
41:a:2766:A:C6	41:a:2767:C:C5	3.05	0.45
43:c:76:GLU:OE1	43:c:77:LYS:O	2.34	0.45
48:h:52:ARG:CD	48:h:53:HIS:CE1	3.00	0.45
51:k:7:GLU:OE2	51:k:27:LYS:HG3	2.16	0.45
51:k:34:LEU:HB2	51:k:51:GLU:OE2	2.16	0.45
52:l:117:ARG:NH1	61:u:1:MET:HE2	2.31	0.45
63:w:60:VAL:HG22	63:w:63:ARG:HH12	1.78	0.45
2:1:24:ILE:HG21	2:1:36:LEU:CD1	2.42	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:2:3:ARG:HH22	3:2:4:GLU:HB2	1.81	0.45
9:9:14:GLU:OE1	9:9:63:ALA:HA	2.17	0.45
9:9:24:SER:HG	9:9:85:SER:C	2.25	0.45
10:A:53:G:N3	10:A:54:U:C5	2.85	0.45
10:A:68:C:H3'	10:A:69:C:H5''	1.97	0.45
12:AB:129:ILE:H	12:AB:142:LEU:HB3	1.81	0.45
12:AB:140:MET:O	12:AB:151:LYS:O	2.34	0.45
14:AE:283:LEU:HD23	14:AE:292:VAL:HG11	1.99	0.45
16:AG:362:TYR:HA	16:AG:413:ALA:HB3	1.94	0.45
18:D:309:A:O2'	18:D:607:A:N1	2.45	0.45
18:D:604:G:H2'	18:D:605:U:O4'	2.17	0.45
18:D:735:C:C2	18:D:736:C:C5	3.05	0.45
18:D:1006:G:H2'	18:D:1007:U:H6	1.81	0.45
18:D:1132:C:O2	18:D:1132:C:H2'	2.16	0.45
18:D:1152:A:OP1	30:P:72:ARG:NH1	2.43	0.45
22:H:37:LEU:HG	22:H:46:SER:O	2.17	0.45
23:I:58:GLU:OE2	30:P:94:ALA:HB1	2.16	0.45
23:I:85:GLU:O	23:I:89:LYS:CE	2.64	0.45
25:K:50:TYR:O	25:K:66:LYS:HE3	2.17	0.45
25:K:76:LEU:HD12	25:K:76:LEU:O	2.17	0.45
27:M:134:ALA:O	27:M:137:LYS:CG	2.65	0.45
31:Q:88:GLY:H	31:Q:114:THR:HG22	1.81	0.45
32:R:110:ARG:N	32:R:111:LYS:HZ3	2.15	0.45
33:S:6:MET:HE3	33:S:63:ARG:HH11	1.82	0.45
33:S:13:ARG:HB3	33:S:60:GLN:HE21	1.81	0.45
33:S:77:PHE:CE1	33:S:93:ILE:HD13	2.51	0.45
34:T:11:ILE:HD13	34:T:31:LEU:HA	1.99	0.45
34:T:15:PHE:CE2	34:T:30:ALA:CB	3.00	0.45
39:Y:60:VAL:HG22	39:Y:66:PHE:CD1	2.52	0.45
39:Y:99:LYS:NZ	39:Y:138:VAL:O	2.44	0.45
41:a:575:A:C2	41:a:576:U:C6	3.05	0.45
41:a:861:A:H2'	41:a:862:G:O4'	2.17	0.45
41:a:1588:G:C6	41:a:1589:U:O4	2.69	0.45
41:a:1665:A:C5'	60:t:66:LYS:NZ	2.79	0.45
41:a:2297:A:C2	41:a:2320:U:C1'	3.00	0.45
43:c:71:LEU:O	43:c:76:GLU:HG3	2.16	0.45
46:f:41:THR:C	46:f:45:ARG:NH2	2.75	0.45
51:k:19:HIS:CE1	51:k:41:PRO:CD	2.99	0.45
54:n:63:GLN:HG3	54:n:95:ARG:HD3	1.99	0.45
56:p:11:VAL:HG12	56:p:48:ASN:O	2.17	0.45
56:p:18:LYS:HG3	56:p:27:LYS:NZ	2.32	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
61:u:95:LEU:HD12	61:u:100:ILE:HG21	1.99	0.45
65:y:14:LYS:NZ	65:y:76:THR:O	2.49	0.45
1:0:74:ILE:HD12	1:0:74:ILE:N	2.32	0.45
2:1:86:MET:SD	2:1:88:ARG:HD2	2.56	0.45
12:AB:31:PRO:C	12:AB:50:LEU:H	2.22	0.45
14:AE:774:ILE:HA	14:AE:777:HIS:HD2	1.82	0.45
16:AG:123:ARG:HG2	16:AG:123:ARG:HH11	1.82	0.45
16:AG:142:VAL:HB	16:AG:175:PRO:HA	1.99	0.45
10:B:8:U:O2'	10:B:47:U:C4	2.67	0.45
18:D:280:C:O2	36:V:40:ARG:NE	2.48	0.45
18:D:723:U:O2	20:F:49:LYS:HE3	2.16	0.45
18:D:789:U:O2'	18:D:791:G:N7	2.45	0.45
18:D:958:A:C4	37:W:55:ARG:HD3	2.52	0.45
18:D:982:U:P	33:S:6:MET:HE2	2.56	0.45
18:D:1202:U:C5	18:D:1203:C:C5	3.04	0.45
18:D:1436:U:H2'	18:D:1437:A:H8	1.82	0.45
18:D:1437:A:C2	18:D:1438:G:C8	3.04	0.45
18:D:1513:A:H2'	18:D:1514:G:C8	2.51	0.45
19:E:54:MET:C	19:E:54:MET:SD	3.00	0.45
22:H:53:PHE:CE1	22:H:68:VAL:HG21	2.52	0.45
22:H:332:VAL:O	22:H:344:LEU:HA	2.17	0.45
25:K:93:ARG:NH2	25:K:93:ARG:CB	2.80	0.45
26:L:18:VAL:N	26:L:19:PRO:CD	2.80	0.45
27:M:132:GLY:O	27:M:136:LYS:HD3	2.17	0.45
28:N:25:VAL:HG23	28:N:63:LEU:CD1	2.46	0.45
29:O:80:ARG:NH2	29:O:103:PHE:CE1	2.85	0.45
38:X:3:ARG:HB3	47:g:35:ASP:OD2	2.15	0.45
39:Y:24:GLY:HA3	39:Y:25:PRO:HD3	1.85	0.45
39:Y:126:ARG:HH12	41:a:1083:U:P	2.40	0.45
40:Z:14:MET:CG	40:Z:15:SER:H	2.25	0.45
41:a:235:U:N3	41:a:236:C:C5	2.84	0.45
41:a:306:U:H2'	41:a:307:G:O4'	2.16	0.45
41:a:598:U:H2'	41:a:599:A:C8	2.52	0.45
41:a:607:U:C6	41:a:620:G:C5	3.05	0.45
41:a:782:A:N1	48:h:225:MET:HE2	2.32	0.45
41:a:1753:G:N2	41:a:1755:A:H3'	2.32	0.45
41:a:1796:U:H2'	41:a:1797:G:H8	1.82	0.45
41:a:1889:A:H2'	41:a:1890:A:C8	2.51	0.45
41:a:2347:C:C2	41:a:2348:U:C5	3.05	0.45
50:j:146:ILE:HA	50:j:159:LYS:HZ1	1.82	0.45
52:l:23:PHE:CE2	52:l:25:GLU:HB2	2.52	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:l:23:PHE:CD1	52:l:111:GLU:CD	2.94	0.45
52:l:147:LEU:HB2	52:l:183:PHE:CD2	2.48	0.45
59:s:17:VAL:HG23	59:s:137:PRO:HB2	1.98	0.45
60:t:17:ARG:NH2	60:t:47:ILE:CG2	2.80	0.45
63:w:13:ASN:O	63:w:17:ARG:HG3	2.17	0.45
3:2:65:GLY:H	3:2:76:ARG:HH12	1.64	0.45
5:4:80:HIS:CE1	5:4:83:LYS:HZ3	2.33	0.45
10:A:18:G:C5	10:A:57:A:N6	2.85	0.45
11:AA:176:ILE:HD11	11:AA:428:VAL:HG21	1.98	0.45
11:AA:712:SER:OG	11:AA:713:GLY:N	2.49	0.45
12:AB:6:LEU:CD2	14:AE:291:ILE:HD11	2.47	0.45
16:AG:440:VAL:CG2	16:AG:481:LEU:HD21	2.42	0.45
10:B:18:G:N7	10:B:57:A:N6	2.65	0.45
10:B:72:A:C3'	10:B:73:A:H5''	2.47	0.45
17:C:70:TYR:C	17:C:71:THR:HG23	2.42	0.45
18:D:511:C:C2	18:D:512:U:C5	3.05	0.45
20:F:12:PHE:CD1	31:Q:101:ASN:CG	2.95	0.45
22:H:49:PRO:HD3	22:H:84:LEU:HD21	1.99	0.45
23:I:72:ARG:NE	23:I:75:ILE:HB	2.32	0.45
25:K:13:GLU:OE1	25:K:13:GLU:N	2.50	0.45
32:R:30:LYS:C	32:R:81:LEU:HD12	2.42	0.45
34:T:71:LYS:HE2	34:T:78:TYR:CD2	2.52	0.45
41:a:1013:C:C2	41:a:1014:A:C8	3.04	0.45
41:a:1107:G:C5	41:a:1108:U:C5	3.04	0.45
41:a:1144:A:C4	41:a:1145:C:C5	3.05	0.45
41:a:1386:C:H2'	41:a:1387:A:H8	1.82	0.45
41:a:2649:C:H2'	41:a:2650:U:C6	2.51	0.45
48:h:123:ALA:HB1	48:h:125:LYS:CE	2.47	0.45
56:p:8:PRO:C	56:p:69:ARG:NH2	2.75	0.45
56:p:149:ARG:CZ	56:p:162:VAL:C	2.89	0.45
65:y:15:GLN:NE2	65:y:16:ASP:CG	2.75	0.45
66:z:66:ASN:CG	66:z:76:TYR:HB2	2.42	0.45
66:z:76:TYR:O	66:z:80:ILE:HG12	2.16	0.45
4:3:66:GLN:OE1	41:a:329:G:OP2	2.34	0.45
6:5:10:DT:H71	11:AA:62:TYR:CB	2.40	0.45
9:9:23:LEU:HA	9:9:118:ILE:HD11	1.99	0.45
9:9:58:THR:HG23	41:a:1108:U:OP1	2.17	0.45
9:9:61:ARG:HE	41:a:1046:A:P	2.39	0.45
9:9:118:ILE:HB	9:9:119:PRO:CD	2.42	0.45
12:AB:6:LEU:HD13	14:AE:291:ILE:CG1	2.46	0.45
16:AG:19:PRO:HG2	16:AG:19:PRO:O	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:362:TYR:HE1	16:AG:414:LEU:HD21	1.80	0.45
16:AG:429:LYS:CB	16:AG:430:PRO:HD2	2.47	0.45
10:B:3:C:H2'	10:B:4:G:H8	1.82	0.45
17:C:34:THR:OG1	17:C:38:LYS:HB2	2.17	0.45
17:C:38:LYS:NZ	18:D:718:A:C6	2.79	0.45
18:D:1060:U:OP1	33:S:85:ARG:NH1	2.48	0.45
18:D:1333:A:H2'	18:D:1334:G:O4'	2.17	0.45
21:G:52:GLU:C	21:G:56:GLU:OE1	2.60	0.45
22:H:32:ASP:OD1	22:H:35:VAL:HG22	2.17	0.45
22:H:74:ALA:HB3	22:H:83:LEU:HD23	1.99	0.45
25:K:86:LYS:HE3	25:K:95:PHE:CD1	2.50	0.45
28:N:110:VAL:O	28:N:111:MET:SD	2.75	0.45
29:O:5:GLN:HE21	29:O:20:PHE:HB3	1.81	0.45
31:Q:21:ALA:HB3	31:Q:84:VAL:HG22	1.99	0.45
32:R:30:LYS:O	32:R:81:LEU:HD12	2.16	0.45
38:X:2:ALA:HB1	38:X:9:ILE:CG2	2.47	0.45
39:Y:44:LYS:HG2	39:Y:68:PHE:HE2	1.83	0.45
41:a:78:U:P	45:e:4:LYS:NZ	2.89	0.45
41:a:980:A:N7	41:a:1136:G:H5'	2.32	0.45
41:a:1007:C:H5''	59:s:37:ARG:NE	2.31	0.45
41:a:1444:G:C4	41:a:1445:G:C8	3.04	0.45
41:a:2098:U:H2'	41:a:2099:U:O4'	2.17	0.45
41:a:2747:G:C2	41:a:2756:U:C5	3.05	0.45
43:c:33:LEU:O	43:c:34:HIS:CG	2.70	0.45
48:h:146:MET:HE2	48:h:153:GLN:OE1	2.17	0.45
48:h:265:LYS:HG3	48:h:266:PHE:CD2	2.52	0.45
49:i:30:VAL:HG22	49:i:37:LYS:HD2	1.99	0.45
50:j:48:ILE:CD1	50:j:84:LEU:HD11	2.46	0.45
50:j:71:ALA:HB3	50:j:73:VAL:HG22	1.98	0.45
52:l:90:GLN:OE1	52:l:92:HIS:CD2	2.70	0.45
56:p:141:ILE:HG13	56:p:142:GLY:N	2.32	0.45
63:w:60:VAL:HG22	63:w:63:ARG:HH11	1.80	0.45
65:y:31:TRP:CH2	65:y:40:LEU:HD11	2.52	0.45
5:4:16:ALA:O	5:4:20:LEU:HG	2.17	0.44
10:A:3:C:H2'	10:A:4:G:C8	2.53	0.44
11:AA:674:ASP:OD2	11:AA:1070:HIS:ND1	2.50	0.44
12:AB:6:LEU:HD22	14:AE:291:ILE:HD11	1.99	0.44
12:AB:37:LYS:HE3	12:AB:109:THR:HG21	1.99	0.44
12:AB:123:PHE:O	12:AB:123:PHE:CD2	2.71	0.44
12:AB:126:PHE:CD2	12:AB:126:PHE:O	2.71	0.44
16:AG:314:LEU:HD22	16:AG:318:ILE:HD11	1.98	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:B:53:G:N3	10:B:54:U:C5	2.85	0.44
18:D:675:A:C1'	31:Q:118:HIS:CD2	2.99	0.44
18:D:765:G:C6	18:D:812:G:C4	3.04	0.44
18:D:918:A:H2'	18:D:919:A:C8	2.51	0.44
18:D:1408:A:H2'	18:D:1409:C:C6	2.53	0.44
20:F:32:VAL:HG11	31:Q:89:PRO:HG3	1.99	0.44
21:G:64:LYS:HE2	21:G:64:LYS:HA	1.99	0.44
21:G:66:LYS:HB3	21:G:154:MET:HE3	1.99	0.44
23:I:134:MET:HE1	23:I:167:TRP:CA	2.43	0.44
23:I:156:ARG:CG	23:I:193:TYR:HB2	2.47	0.44
27:M:31:MET:HE1	27:M:36:LYS:CD	2.46	0.44
31:Q:14:LYS:HD3	31:Q:15:GLN:N	2.32	0.44
31:Q:52:PHE:CE1	31:Q:65:VAL:HG21	2.52	0.44
32:R:23:ALA:O	32:R:25:GLU:OE1	2.36	0.44
35:U:8:ARG:H	35:U:28:ARG:HD3	1.81	0.44
38:X:14:HIS:HB2	38:X:17:ILE:HD13	1.99	0.44
39:Y:19:PRO:CG	39:Y:23:VAL:CG2	2.95	0.44
39:Y:45:THR:HG22	39:Y:45:THR:O	2.16	0.44
41:a:644:A:C2	41:a:2369:A:H1'	2.51	0.44
41:a:1408:G:N3	41:a:1408:G:H2'	2.31	0.44
41:a:1656:C:C2	41:a:1657:U:C5	3.05	0.44
41:a:1997:C:OP1	50:j:128:ARG:NH1	2.51	0.44
41:a:2049:G:N2	50:j:161:MET:CE	2.80	0.44
41:a:2095:A:O5'	58:r:11:ASN:OD1	2.35	0.44
41:a:2356:U:H4'	42:b:20:ARG:HD2	1.99	0.44
56:p:91:GLY:HA3	56:p:94:TYR:CD2	2.52	0.44
57:q:30:GLU:HB3	57:q:33:HIS:ND1	2.31	0.44
60:t:48:PRO:HB2	60:t:49:ARG:HE	1.81	0.44
65:y:31:TRP:CZ3	65:y:40:LEU:CD1	3.00	0.44
4:3:5:ILE:HG13	4:3:72:ILE:HD11	1.99	0.44
9:9:23:LEU:O	9:9:87:GLU:CD	2.60	0.44
12:AB:76:SER:O	12:AB:77:HIS:HB3	2.16	0.44
14:AE:1289:ASN:ND2	14:AE:1300:ALA:O	2.46	0.44
10:B:3:C:H2'	10:B:4:G:C8	2.52	0.44
10:B:18:G:C5	10:B:57:A:N6	2.85	0.44
17:C:65:LEU:HD23	17:C:67:LEU:CD1	2.47	0.44
17:C:65:LEU:HG	17:C:65:LEU:O	2.16	0.44
18:D:653:U:O4'	28:N:56:LYS:HE3	2.16	0.44
18:D:715:A:H2'	18:D:716:A:C8	2.52	0.44
18:D:824:G:C6	18:D:877:G:C6	3.05	0.44
21:G:109:GLN:C	21:G:113:ARG:NE	2.75	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:H:163:ARG:N	22:H:298:GLU:CG	2.81	0.44
23:I:24:ALA:HB3	23:I:29:PHE:CD2	2.51	0.44
25:K:13:GLU:HG2	25:K:13:GLU:O	2.16	0.44
26:L:96:VAL:O	26:L:96:VAL:HG23	2.16	0.44
27:M:17:LYS:NZ	27:M:18:PHE:CE1	2.78	0.44
28:N:84:ARG:HD2	28:N:124:GLU:OE1	2.17	0.44
28:N:89:LYS:HG3	28:N:90:ASP:N	2.32	0.44
33:S:12:LYS:NZ	33:S:16:LEU:HD21	2.31	0.44
36:V:27:ARG:HH22	36:V:42:THR:N	2.16	0.44
37:W:13:LEU:O	37:W:17:LYS:HG2	2.17	0.44
37:W:44:MET:HE1	37:W:66:MET:HE1	1.98	0.44
38:X:3:ARG:NH2	54:n:110:ARG:HD2	2.33	0.44
38:X:55:THR:HG22	38:X:59:GLU:OE2	2.17	0.44
41:a:837:C:N3	41:a:941:A:N6	2.65	0.44
41:a:1182:G:H2'	41:a:1183:U:O4'	2.17	0.44
41:a:1565:C:O2'	41:a:1567:G:N7	2.47	0.44
41:a:1709:U:H2'	41:a:1710:G:H8	1.82	0.44
41:a:2149:U:H2'	41:a:2150:C:C6	2.52	0.44
41:a:2743:U:H2'	41:a:2744:G:O4'	2.17	0.44
41:a:2814:A:C5	41:a:2815:C:C5	3.05	0.44
48:h:164:ILE:O	48:h:164:ILE:CG2	2.65	0.44
50:j:3:GLY:O	50:j:82:PHE:CE2	2.69	0.44
53:m:12:ARG:CG	53:m:44:VAL:HG21	2.47	0.44
62:v:104:GLU:OE1	62:v:104:GLU:HA	2.17	0.44
66:z:82:GLY:CA	66:z:117:LEU:HD11	2.48	0.44
1:0:46:GLU:HG2	1:0:48:LYS:NZ	2.32	0.44
6:5:11:DA:C5'	12:AB:67:THR:O	2.66	0.44
8:7:2:U:OP2	18:D:926:G:N2	2.50	0.44
9:9:6:GLN:CA	9:9:9:GLN:HE22	2.30	0.44
11:AA:148:GLN:NE2	11:AA:535:PRO:O	2.40	0.44
12:AB:124:GLU:C	12:AB:126:PHE:N	2.53	0.44
14:AE:708:ASN:OD1	14:AE:708:ASN:N	2.50	0.44
16:AG:143:LYS:HG3	16:AG:153:ASP:HB2	1.99	0.44
16:AG:226:ALA:HB1	16:AG:228:ARG:HH11	1.81	0.44
16:AG:429:LYS:HB2	16:AG:430:PRO:HD2	1.99	0.44
18:D:524:G:H2'	18:D:525:C:C6	2.52	0.44
18:D:682:G:C2	18:D:683:G:C8	3.06	0.44
18:D:753:A:OP1	34:T:73:LYS:NZ	2.50	0.44
18:D:1240:U:OP1	27:M:116:MET:HB2	2.17	0.44
20:F:38:TYR:O	31:Q:126:LYS:HE3	2.17	0.44
22:H:308:GLU:C	22:H:310:ASP:H	2.25	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:I:72:ARG:HD3	23:I:75:ILE:HG22	1.99	0.44
23:I:114:LYS:HB2	23:I:185:ASN:ND2	2.32	0.44
24:J:49:SER:O	24:J:53:VAL:HG13	2.18	0.44
24:J:172:GLU:CD	24:J:183:LYS:HD2	2.42	0.44
26:L:5:GLU:CD	26:L:61:LEU:HD11	2.42	0.44
27:M:106:GLU:O	27:M:110:LYS:HG2	2.18	0.44
31:Q:51:GLY:O	31:Q:56:ARG:NH1	2.50	0.44
38:X:55:THR:C	38:X:59:GLU:OE1	2.60	0.44
38:X:79:ARG:NH1	47:g:56:ARG:NH1	2.66	0.44
40:Z:2:ILE:CG2	40:Z:5:ASP:H	2.28	0.44
41:a:6:A:O2'	59:s:132:HIS:ND1	2.50	0.44
41:a:27:G:C4	41:a:512:G:N2	2.86	0.44
41:a:30:G:H2'	41:a:31:C:H6	1.82	0.44
41:a:68:G:N2	41:a:74:A:OP2	2.51	0.44
41:a:1146:C:C2	41:a:1147:A:C8	3.05	0.44
41:a:1965:C:OP1	41:a:1966:A:O2'	2.35	0.44
41:a:2021:C:OP1	49:i:9:THR:HG21	2.18	0.44
41:a:2492:U:C2'	41:a:2493:U:H5'	2.48	0.44
41:a:2617:U:C2	41:a:2618:G:C8	3.05	0.44
41:a:2718:G:O2'	41:a:2847:U:H5''	2.18	0.44
41:a:2727:A:O3'	60:t:70:ARG:NH2	2.50	0.44
41:a:2747:G:N2	41:a:2756:U:C6	2.85	0.44
44:d:45:A:C5'	54:n:92:ARG:NH1	2.79	0.44
48:h:200:HIS:O	48:h:203:ARG:HG2	2.17	0.44
48:h:240:PHE:CD2	48:h:240:PHE:O	2.71	0.44
52:l:1:MET:HE3	52:l:19:PHE:C	2.43	0.44
54:n:22:TYR:CE2	54:n:28:VAL:HA	2.52	0.44
54:n:40:VAL:HG13	54:n:42:GLU:CD	2.43	0.44
59:s:28:LEU:HD12	59:s:142:ILE:CG2	2.47	0.44
59:s:132:HIS:HA	59:s:135:GLN:OE1	2.17	0.44
63:w:73:ASN:C	63:w:73:ASN:ND2	2.76	0.44
63:w:79:LEU:HD23	63:w:83:LEU:HD12	1.99	0.44
65:y:27:GLU:HB3	65:y:87:LYS:HZ1	1.83	0.44
3:2:6:ARG:NE	3:2:42:GLU:CD	2.76	0.44
5:4:31:TYR:O	5:4:92:VAL:HA	2.17	0.44
8:7:12:U:C3'	8:7:12:U:C6	3.00	0.44
8:7:51:A:O3'	11:AA:688:GLN:NE2	2.51	0.44
10:A:71:C:H2'	10:A:72:A:N9	2.32	0.44
16:AG:15:GLU:HA	16:AG:18:LEU:HB2	2.00	0.44
17:C:41:PRO:HD2	22:H:339:ARG:CZ	2.47	0.44
18:D:415:A:C6	18:D:416:G:C5	3.05	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:G:27:MET:HG3	21:G:27:MET:O	2.16	0.44
26:L:27:ALA:O	26:L:30:THR:OG1	2.29	0.44
31:Q:85:MET:HG3	31:Q:111:THR:HB	2.00	0.44
33:S:82:ILE:HG23	33:S:83:LYS:N	2.31	0.44
37:W:68:GLY:HA3	47:g:59:ARG:HH22	1.82	0.44
38:X:71:ARG:NH1	38:X:72:GLU:HG3	2.33	0.44
38:X:90:ARG:HD3	38:X:96:PRO:O	2.17	0.44
39:Y:33:ASN:OD1	39:Y:35:MET:HE2	2.17	0.44
39:Y:102:ARG:NE	39:Y:139:VAL:CG2	2.78	0.44
41:a:66:C:N4	41:a:74:A:N6	2.66	0.44
41:a:535:G:H2'	41:a:536:G:O4'	2.17	0.44
41:a:677:A:O2'	41:a:2071:A:H5'	2.18	0.44
41:a:841:G:C6	41:a:938:G:C6	3.06	0.44
41:a:1020:A:C2	41:a:1141:U:C2	3.06	0.44
41:a:1434:A:H2'	41:a:1435:G:C8	2.53	0.44
41:a:1441:G:H2'	41:a:1442:U:H6	1.83	0.44
41:a:1790:C:C2'	41:a:1791:A:C8	3.00	0.44
41:a:1824:G:P	48:h:52:ARG:CZ	3.06	0.44
41:a:2548:U:C2	41:a:2549:G:C8	3.05	0.44
45:e:10:SER:HG	45:e:12:GLU:HG2	1.82	0.44
45:e:24:GLU:O	45:e:28:LEU:HD13	2.17	0.44
46:f:6:LYS:HD3	46:f:6:LYS:N	2.33	0.44
46:f:12:SER:CA	46:f:32:ILE:HD11	2.46	0.44
50:j:5:VAL:HB	50:j:32:ASN:ND2	2.31	0.44
50:j:61:THR:OG1	50:j:63:PRO:HD2	2.18	0.44
52:l:95:LYS:HE2	52:l:97:ASN:OD1	2.17	0.44
53:m:41:ARG:NE	53:m:41:ARG:HA	2.33	0.44
54:n:117:LEU:N	54:n:177:PHE:O	2.51	0.44
55:o:31:HIS:C	55:o:33:LEU:HG	2.43	0.44
59:s:98:GLU:O	59:s:100:VAL:N	2.50	0.44
64:x:90:VAL:CG2	64:x:115:LEU:HD21	2.46	0.44
65:y:106:LYS:CD	65:y:109:ARG:NH2	2.79	0.44
66:z:78:LYS:HG2	66:z:117:LEU:HD21	2.00	0.44
11:AA:24:VAL:HG11	11:AA:704:MET:HE1	1.99	0.44
11:AA:444:ASP:OD1	11:AA:444:ASP:N	2.51	0.44
11:AA:694:ARG:HH22	13:AC:80:GLU:HG3	1.82	0.44
12:AB:52:PRO:O	12:AB:54:TYR:N	2.50	0.44
14:AE:1022:PRO:O	14:AE:1126:GLN:NE2	2.45	0.44
16:AG:126:VAL:HG13	16:AG:130:PHE:CE2	2.53	0.44
10:B:71:C:H2'	10:B:72:A:C1'	2.47	0.44
18:D:323:U:H4'	19:E:17:ALA:HB3	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:429:U:O2	18:D:430:A:N7	2.50	0.44
18:D:676:A:H2'	18:D:677:U:H6	1.82	0.44
18:D:746:A:H2'	18:D:747:A:C8	2.53	0.44
18:D:1125:U:N3	18:D:1127:G:C8	2.86	0.44
18:D:1315:U:O2	18:D:1360:A:H2	2.00	0.44
20:F:26:ALA:CB	20:F:28:VAL:HG23	2.47	0.44
21:G:163:VAL:CG1	21:G:185:ALA:HA	2.47	0.44
21:G:171:ILE:H	21:G:171:ILE:HD12	1.82	0.44
23:I:152:GLU:OE2	23:I:165:THR:HG23	2.18	0.44
26:L:53:LYS:HD2	26:L:53:LYS:O	2.18	0.44
26:L:72:ASP:O	26:L:76:THR:HG23	2.17	0.44
29:O:46:MET:HB3	29:O:50:GLN:NE2	2.31	0.44
29:O:51:PRO:CA	29:O:103:PHE:HE2	2.30	0.44
29:O:86:ALA:C	29:O:89:GLU:HG2	2.43	0.44
32:R:74:LEU:HD11	32:R:80:ILE:CG1	2.48	0.44
34:T:10:LYS:CD	34:T:14:GLU:OE2	2.65	0.44
34:T:26:GLU:HG3	34:T:27:VAL:N	2.33	0.44
35:U:45:GLU:HG3	35:U:46:LYS:CD	2.45	0.44
36:V:15:ASP:OD1	36:V:15:ASP:N	2.49	0.44
39:Y:93:ASN:O	39:Y:94:LYS:HD2	2.16	0.44
41:a:145:C:H2'	41:a:146:A:C8	2.53	0.44
41:a:208:C:H2'	41:a:209:C:C6	2.53	0.44
41:a:479:A:H4'	41:a:480:A:OP1	2.18	0.44
41:a:536:G:H4'	66:z:57:PHE:HZ	1.82	0.44
41:a:900:A:C5	41:a:901:C:C5	3.06	0.44
41:a:957:C:H5'	62:v:75:GLU:OE1	2.18	0.44
41:a:1169:A:O4'	41:a:1169:A:N3	2.51	0.44
41:a:2262:U:N3	41:a:2263:C:C5	2.85	0.44
41:a:2309:A:H2'	41:a:2310:C:O4'	2.16	0.44
41:a:2806:C:H2'	41:a:2807:U:O4'	2.18	0.44
54:n:79:ILE:O	54:n:79:ILE:HG13	2.17	0.44
55:o:39:LYS:HG2	55:o:42:ARG:HH12	1.82	0.44
9:9:34:THR:HA	9:9:37:LYS:NZ	2.32	0.44
10:A:3:C:H2'	10:A:4:G:H8	1.82	0.44
10:A:71:C:H2'	10:A:72:A:C1'	2.47	0.44
11:AA:133:ASN:O	11:AA:527:LYS:NZ	2.42	0.44
11:AA:855:PRO:HG2	16:AG:105:ILE:HG22	1.94	0.44
14:AE:288:PRO:HD2	14:AE:291:ILE:CB	2.42	0.44
14:AE:907:HIS:ND1	14:AE:908:ILE:O	2.48	0.44
16:AG:167:MET:HE3	16:AG:173:PHE:CZ	2.53	0.44
16:AG:227:ALA:CB	16:AG:330:GLN:C	2.90	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:C:18:VAL:HG13	17:C:51:TYR:HE1	1.83	0.44
17:C:54:GLN:HA	17:C:57:ARG:NH1	2.33	0.44
17:C:60:LYS:CE	18:D:735:C:O4'	2.64	0.44
17:C:70:TYR:CE2	26:L:49:TYR:CE1	3.05	0.44
18:D:219:U:C2	18:D:220:G:C8	3.05	0.44
18:D:278:G:OP2	36:V:43:LYS:NZ	2.49	0.44
18:D:511:C:C2	18:D:512:U:C6	3.05	0.44
18:D:622:A:C8	18:D:623:C:C6	3.06	0.44
18:D:704:A:C4	18:D:705:G:C8	3.06	0.44
19:E:16:LYS:HA	19:E:19:LYS:HE3	2.00	0.44
21:G:30:PHE:CE1	21:G:201:PRO:HG3	2.53	0.44
22:H:124:LEU:O	22:H:127:ILE:O	2.36	0.44
22:H:332:VAL:CG1	22:H:335:ILE:HG13	2.48	0.44
28:N:35:ALA:HB1	28:N:110:VAL:HG11	2.00	0.44
29:O:63:LEU:CD1	29:O:65:ILE:HD11	2.48	0.44
30:P:40:ILE:HD12	30:P:73:LEU:CD2	2.48	0.44
30:P:73:LEU:C	30:P:73:LEU:HD23	2.42	0.44
33:S:87:ALA:CA	33:S:90:ARG:CZ	2.96	0.44
34:T:3:LEU:HD23	34:T:4:SER:N	2.33	0.44
36:V:31:HIS:NE2	36:V:34:TYR:HD2	2.15	0.44
41:a:128:C:C2	41:a:129:C:C5	3.06	0.44
41:a:234:U:C4	41:a:235:U:C5	3.05	0.44
41:a:564:C:H1'	66:z:37:GLN:NE2	2.32	0.44
41:a:963:U:H2'	41:a:964:C:C6	2.53	0.44
41:a:1310:G:N2	41:a:1313:U:C4	2.85	0.44
41:a:1406:U:O2'	41:a:1407:G:C5'	2.66	0.44
41:a:1443:U:H2'	41:a:1444:G:H8	1.82	0.44
41:a:1820:U:C2	48:h:201:MET:HE2	2.53	0.44
41:a:1839:G:C4	41:a:1840:G:C8	3.06	0.44
41:a:1856:U:H2'	41:a:1857:G:O4'	2.17	0.44
41:a:2702:G:C4	41:a:2703:C:C5	3.06	0.44
41:a:2783:U:H2'	41:a:2784:U:C6	2.52	0.44
46:f:41:THR:O	46:f:43:ALA:N	2.51	0.44
48:h:91:ILE:HG22	48:h:92:ALA:N	2.33	0.44
52:l:108:ILE:HD11	52:l:180:LEU:HD13	1.97	0.44
52:l:149:ILE:CD1	52:l:188:MET:HG2	2.46	0.44
55:o:55:LEU:C	55:o:55:LEU:HD23	2.43	0.44
56:p:11:VAL:HG13	56:p:11:VAL:O	2.17	0.44
61:u:101:ILE:C	61:u:105:ILE:HD12	2.42	0.44
65:y:106:LYS:CA	65:y:109:ARG:NH2	2.81	0.44
2:1:78:GLU:OE2	41:a:1262:A:OP1	2.35	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:3:7:ARG:NH2	41:a:84:A:O2'	2.51	0.44
5:4:31:TYR:HE2	44:d:104:A:O4'	2.01	0.44
8:7:1:A:C6	10:B:37:A:N1	2.86	0.44
9:9:31:ARG:NH1	41:a:1054:A:C1'	2.80	0.44
10:A:18:G:N2	10:A:58:A:N7	2.65	0.44
11:AA:481:LEU:HD23	12:AB:16:ARG:HD3	1.79	0.44
11:AA:914:LYS:N	16:AG:105:ILE:C	2.76	0.44
12:AB:118:ILE:HG21	12:AB:141:LEU:CG	2.44	0.44
12:AB:130:PHE:HZ	12:AB:160:ARG:HG3	1.81	0.44
12:AB:138:ARG:HG3	12:AB:155:LYS:N	2.20	0.44
13:AD:78:ILE:HA	13:AD:81:ILE:HG22	2.00	0.44
13:AD:192:VAL:HG12	13:AD:193:GLU:H	1.83	0.44
16:AG:3:LYS:HB3	16:AG:6:LEU:HD12	1.99	0.44
18:D:13:U:O4	18:D:20:U:O4	2.34	0.44
18:D:598:U:C2	18:D:599:C:C5	3.06	0.44
18:D:636:U:H2'	18:D:637:C:C6	2.52	0.44
18:D:1120:C:H2'	18:D:1121:U:H6	1.83	0.44
22:H:12:SER:O	22:H:16:ILE:HG13	2.17	0.44
22:H:53:PHE:HE1	22:H:68:VAL:HG21	1.82	0.44
22:H:76:GLU:OE1	22:H:79:PHE:HA	2.18	0.44
22:H:303:LEU:CD2	22:H:305:HIS:HB3	2.48	0.44
25:K:134:ILE:CG2	25:K:135:ASN:N	2.81	0.44
29:O:39:PHE:CD1	29:O:44:ALA:HB1	2.53	0.44
30:P:24:GLU:OE2	30:P:92:LEU:CD2	2.66	0.44
30:P:52:LEU:HB2	33:S:81:ARG:NH1	2.33	0.44
32:R:24:LEU:HD23	32:R:95:TYR:HE1	1.83	0.44
34:T:63:ARG:HD2	34:T:67:LEU:HD21	2.00	0.44
41:a:221:A:C4	41:a:266:G:N7	2.85	0.44
41:a:580:U:O3'	66:z:31:VAL:HG13	2.17	0.44
41:a:607:U:N3	41:a:608:A:N7	2.66	0.44
41:a:730:A:C2	41:a:731:C:C5	3.05	0.44
41:a:782:A:C6	48:h:225:MET:HE2	2.53	0.44
41:a:1936:A:C8	41:a:1945:G:N7	2.86	0.44
41:a:2345:G:N3	41:a:2381:A:H2'	2.33	0.44
41:a:2839:G:C5	41:a:2840:C:C5	3.05	0.44
42:b:21:LEU:HD23	42:b:40:GLN:HA	2.00	0.44
44:d:48:U:P	64:x:30:ARG:HH22	2.41	0.44
47:g:59:ARG:HH11	47:g:63:ARG:CD	2.31	0.44
48:h:45:ASN:OD1	48:h:46:ASN:N	2.51	0.44
55:o:45:ARG:HB3	55:o:46:PRO:CD	2.48	0.44
56:p:27:LYS:HG2	56:p:32:GLU:HG3	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:s:73:VAL:CG1	59:s:86:GLN:HG2	2.46	0.44
61:u:91:ASP:HB2	61:u:93:ASN:OD1	2.17	0.44
63:w:38:LEU:HA	63:w:38:LEU:HD12	1.71	0.44
64:x:50:ALA:O	64:x:81:ARG:NH2	2.51	0.44
66:z:89:GLU:N	66:z:89:GLU:OE1	2.50	0.44
2:1:36:LEU:N	2:1:36:LEU:HD12	2.33	0.44
5:4:50:MET:CB	5:4:56:PHE:CE1	2.98	0.44
8:7:9:U:OP2	8:7:9:U:C6	2.71	0.44
12:AB:31:PRO:HG2	12:AB:51:PHE:CE2	2.53	0.44
12:AB:117:ILE:HG21	12:AB:161:LYS:HE2	1.99	0.44
16:AG:402:THR:C	16:AG:405:ALA:H	2.26	0.44
16:AG:433:ASP:HB3	16:AG:456:LEU:CD1	2.47	0.44
16:AG:440:VAL:HG13	16:AG:444:LEU:HB2	2.00	0.44
10:B:75:C:P	41:a:2602:A:C2'	3.05	0.44
17:C:34:THR:HG22	17:C:35:GLU:N	2.33	0.44
18:D:153:C:C2	18:D:154:U:C5	3.06	0.44
18:D:633:G:H2'	18:D:634:C:H6	1.82	0.44
18:D:986:U:H2'	18:D:987:G:C8	2.53	0.44
18:D:1103:C:O2	21:G:106:THR:HG21	2.18	0.44
18:D:1120:C:C2	18:D:1121:U:C5	3.06	0.44
21:G:9:MET:HE2	21:G:43:LEU:HD22	2.00	0.44
22:H:294:VAL:HG11	22:H:330:VAL:HG21	2.00	0.44
23:I:14:ILE:HG13	23:I:178:LEU:HD21	1.99	0.44
25:K:50:TYR:C	25:K:66:LYS:HE3	2.43	0.44
29:O:51:PRO:HB3	29:O:103:PHE:HD2	1.83	0.44
29:O:89:GLU:HG3	29:O:90:TYR:N	2.33	0.44
31:Q:63:ALA:CB	31:Q:92:GLY:HA3	2.48	0.44
31:Q:83:GLU:OE1	31:Q:83:GLU:N	2.51	0.44
31:Q:84:VAL:HG11	31:Q:97:ILE:HD13	2.00	0.44
36:V:28:PHE:CZ	36:V:37:PHE:HB3	2.53	0.44
38:X:16:VAL:HG23	38:X:17:ILE:CD1	2.47	0.44
41:a:598:U:H2'	41:a:599:A:H8	1.82	0.44
41:a:1141:U:H4'	41:a:1142:A:O4'	2.18	0.44
41:a:1282:U:H2'	41:a:1283:G:O4'	2.18	0.44
41:a:1414:C:H2'	41:a:1415:U:C6	2.52	0.44
41:a:1996:C:OP1	60:t:31:ARG:NH2	2.51	0.44
41:a:2175:C:C2'	41:a:2176:A:C8	3.00	0.44
41:a:2514:U:H2'	41:a:2515:C:H6	1.81	0.44
41:a:2616:C:C2	41:a:2617:U:C6	3.06	0.44
46:f:4:THR:CG2	46:f:37:GLU:OE2	2.65	0.44
48:h:74:ILE:HB	48:h:96:TYR:CD1	2.53	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:i:23:THR:O	49:i:23:THR:HG23	2.17	0.44
54:n:175:PHE:CG	54:n:176:PRO:HD2	2.52	0.44
55:o:49:MET:H	55:o:49:MET:HE2	1.81	0.44
58:r:90:LEU:HD21	58:r:94:ILE:HD11	2.00	0.44
59:s:92:MET:HE3	59:s:92:MET:HB2	1.13	0.44
8:7:4:U:C5	18:D:1403:C:H1'	2.52	0.44
9:9:138:ARG:NH1	40:Z:10:ALA:O	2.51	0.44
10:A:19:G:O6	41:a:2111:U:H3'	2.18	0.44
11:AA:11:ILE:HG22	11:AA:1172:LEU:HD11	1.99	0.44
12:AB:141:LEU:HB2	12:AB:143:LEU:HD11	1.98	0.44
13:AD:257:VAL:N	13:AD:276:HIS:O	2.34	0.44
14:AE:41:PRO:HB2	14:AE:270:ARG:HG3	1.99	0.44
14:AE:71:LEU:HA	14:AE:71:LEU:HD12	1.78	0.44
14:AE:641:ILE:HD12	14:AE:641:ILE:HA	1.89	0.44
16:AG:233:ARG:HB2	16:AG:327:LEU:CD2	2.42	0.44
16:AG:363:LEU:CD2	16:AG:409:ARG:N	2.81	0.44
16:AG:363:LEU:HG	16:AG:410:ALA:HA	1.94	0.44
16:AG:365:ILE:HB	16:AG:366:ASP:H	1.73	0.44
10:B:18:G:N2	10:B:58:A:N7	2.65	0.44
18:D:76:G:C5	18:D:77:A:C8	3.06	0.44
18:D:360:G:H2'	18:D:361:G:C8	2.53	0.44
18:D:1160:G:C6	18:D:1161:C:C5	3.06	0.44
18:D:1437:A:N3	18:D:1438:G:C8	2.86	0.44
18:D:1486:G:H2'	18:D:1487:G:O4'	2.18	0.44
25:K:132:ASN:OD1	25:K:134:ILE:HG22	2.18	0.44
29:O:91:ASP:HB3	29:O:94:LEU:HG	2.00	0.44
32:R:38:TYR:CE2	32:R:40:THR:CG2	3.01	0.44
34:T:32:LEU:HD12	34:T:63:ARG:HA	1.98	0.44
35:U:14:ARG:CD	35:U:42:ILE:HG21	2.47	0.44
35:U:73:ALA:HA	35:U:76:LYS:HE3	1.98	0.44
36:V:63:GLU:HB2	36:V:73:TRP:CZ2	2.53	0.44
37:W:79:THR:HG23	37:W:79:THR:O	2.17	0.44
39:Y:14:ALA:O	39:Y:16:MET:N	2.51	0.44
41:a:536:G:H4'	66:z:57:PHE:CZ	2.53	0.44
41:a:594:U:C2	41:a:595:C:C5	3.05	0.44
41:a:1349:C:O2	41:a:1349:C:H2'	2.17	0.44
41:a:2092:U:C2	41:a:2225:A:O2'	2.71	0.44
41:a:2387:U:C4'	42:b:41:ARG:NH1	2.81	0.44
41:a:2391:G:P	55:o:32:ILE:CD1	3.06	0.44
41:a:2466:C:C2	41:a:2467:C:C5	3.06	0.44
41:a:2646:C:H2'	41:a:2647:U:O4'	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2848:G:C2	41:a:2867:G:C4	3.05	0.44
45:e:13:GLU:C	45:e:17:GLU:OE1	2.61	0.44
52:l:23:PHE:CE2	52:l:25:GLU:CB	3.01	0.44
52:l:23:PHE:CD1	52:l:111:GLU:CG	3.00	0.44
53:m:26:ASN:HA	53:m:29:GLN:HE21	1.83	0.44
54:n:126:GLY:HA2	54:n:163:ASP:HA	2.00	0.44
55:o:13:ARG:HG2	61:u:62:PRO:O	2.18	0.44
56:p:94:TYR:HE1	56:p:152:ARG:HD2	1.83	0.44
58:r:127:GLU:OE2	58:r:143:ILE:HB	2.17	0.44
63:w:12:ARG:NH1	63:w:20:MET:HE1	2.33	0.44
1:0:93:PHE:CD1	1:0:93:PHE:C	2.96	0.43
2:1:1:MET:SD	2:1:62:ASP:OD2	2.76	0.43
9:9:68:PRO:O	9:9:72:LEU:HD21	2.16	0.43
9:9:92:ALA:HB3	9:9:129:LEU:HB2	1.99	0.43
10:A:32:C:O3'	27:M:86:GLN:NE2	2.51	0.43
11:AA:560:PRO:HB2	14:AE:776:THR:HG21	1.99	0.43
11:AA:811:ASN:ND2	11:AA:1098:LEU:O	2.51	0.43
11:AA:856:ASN:CB	16:AG:35:THR:HB	2.48	0.43
12:AB:38:ILE:HA	12:AB:42:LYS:O	2.17	0.43
12:AB:120:GLU:CB	12:AB:162:LEU:HB2	2.46	0.43
13:AD:288:GLU:C	13:AD:290:LEU:H	2.26	0.43
14:AE:117:LEU:HG	14:AE:118:LYS:HG3	2.00	0.43
10:B:11:A:H2'	10:B:12:G:O4'	2.17	0.43
10:B:35:A:H2'	10:B:36:U:O4'	2.18	0.43
18:D:262:A:H2'	18:D:263:A:C8	2.53	0.43
18:D:284:C:C2	18:D:285:C:C5	3.06	0.43
18:D:414:A:C4	18:D:415:A:C8	3.05	0.43
18:D:545:C:C5'	24:J:69:GLU:HG3	2.38	0.43
18:D:953:G:O6	38:X:103:LYS:HE3	2.18	0.43
18:D:1146:A:C5	18:D:1147:C:C5	3.06	0.43
19:E:25:ARG:O	19:E:28:MET:HB2	2.18	0.43
19:E:45:ALA:O	19:E:48:GLN:NE2	2.49	0.43
21:G:117:LEU:HD21	21:G:141:LEU:CA	2.48	0.43
21:G:206:ALA:O	21:G:210:VAL:HG23	2.18	0.43
23:I:22:TRP:CH2	23:I:32:ASN:HB2	2.53	0.43
23:I:34:ASP:HB2	33:S:65:ARG:CZ	2.48	0.43
25:K:15:LEU:HD23	25:K:15:LEU:H	1.83	0.43
28:N:10:MET:HB2	28:N:33:LYS:HZ2	1.83	0.43
31:Q:91:PRO:HD2	31:Q:92:GLY:H	1.83	0.43
34:T:15:PHE:HE2	34:T:30:ALA:HB1	1.82	0.43
34:T:15:PHE:HD2	34:T:30:ALA:CB	2.31	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:X:11:ASP:HA	38:X:45:ILE:HB	1.99	0.43
39:Y:12:VAL:HG22	39:Y:54:ILE:O	2.17	0.43
39:Y:14:ALA:O	39:Y:15:GLY:C	2.61	0.43
39:Y:82:ALA:HB3	39:Y:108:ILE:CD1	2.48	0.43
41:a:65:U:N3	41:a:66:C:C5	2.86	0.43
41:a:66:C:H42	41:a:74:A:H62	1.67	0.43
41:a:150:U:C2	41:a:151:C:C5	3.06	0.43
41:a:579:G:H2'	41:a:580:U:C6	2.53	0.43
41:a:597:G:H2'	41:a:598:U:C6	2.52	0.43
41:a:685:A:N1	41:a:787:C:H1'	2.33	0.43
41:a:841:G:H2'	41:a:842:U:C6	2.52	0.43
41:a:1319:C:O2'	41:a:1320:C:H5'	2.18	0.43
41:a:2025:C:H2'	41:a:2026:U:C6	2.53	0.43
41:a:2063:C:N3	41:a:2064:C:C5	2.86	0.43
41:a:2477:U:O2	57:q:4:ARG:NH2	2.51	0.43
41:a:2884:U:O2	49:i:40:ARG:NH2	2.51	0.43
44:d:45:A:C4	44:d:46:A:C8	3.05	0.43
48:h:251:GLN:OE1	48:h:255:LYS:CB	2.65	0.43
52:l:73:ILE:HG13	52:l:73:ILE:O	2.17	0.43
54:n:15:LYS:C	54:n:19:GLU:OE1	2.61	0.43
54:n:93:GLY:O	54:n:94:GLU:C	2.61	0.43
56:p:149:ARG:HD2	56:p:162:VAL:O	2.18	0.43
60:t:18:ARG:NE	60:t:18:ARG:HA	2.33	0.43
61:u:55:MET:SD	61:u:60:ARG:N	2.90	0.43
62:v:53:MET:O	62:v:57:VAL:HG12	2.18	0.43
66:z:74:ILE:CD1	66:z:79:PHE:HA	2.40	0.43
1:0:13:ARG:CZ	66:z:97:ASP:CG	2.91	0.43
3:2:56:GLU:OE1	3:2:86:THR:C	2.61	0.43
5:4:29:ILE:HD13	44:d:74:U:O2	2.17	0.43
13:AD:29:GLU:HB3	13:AD:200:LYS:HG3	1.99	0.43
16:AG:104:ARG:N	16:AG:105:ILE:HG12	2.32	0.43
16:AG:168:LEU:CB	16:AG:171:GLU:HB2	2.46	0.43
16:AG:430:PRO:CG	16:AG:435:LEU:CD2	2.85	0.43
18:D:689:C:H2'	18:D:690:G:O4'	2.19	0.43
18:D:882:C:O2'	18:D:883:C:H5'	2.18	0.43
18:D:892:A:H2'	18:D:893:C:H6	1.83	0.43
18:D:1250:A:H2'	18:D:1251:A:C8	2.52	0.43
21:G:48:PRO:O	21:G:52:GLU:HG2	2.18	0.43
22:H:5:PHE:CZ	22:H:9:PHE:CE1	3.06	0.43
22:H:49:PRO:CD	22:H:84:LEU:HD21	2.47	0.43
26:L:78:PHE:HD1	26:L:87:SER:HB3	1.83	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:O:88:MET:CE	29:O:95:ARG:HG3	2.46	0.43
39:Y:81:LYS:N	39:Y:81:LYS:CD	2.79	0.43
41:a:858:G:OP1	42:b:78:LYS:HE3	2.18	0.43
41:a:1021:A:C2	41:a:1141:U:O4	2.71	0.43
41:a:2093:G:P	58:r:22:LYS:HG2	2.58	0.43
41:a:2450:A:C2	41:a:2451:A:C4	3.06	0.43
44:d:89:U:O2'	44:d:90:C:P	2.76	0.43
46:f:31:ARG:HG2	46:f:34:HIS:HB2	2.00	0.43
49:i:17:ARG:CA	49:i:20:ASP:OD1	2.66	0.43
50:j:131:ASP:OD1	50:j:133:THR:O	2.36	0.43
59:s:35:ARG:HA	59:s:40:HIS:ND1	2.32	0.43
60:t:42:THR:CG2	60:t:44:LYS:NZ	2.78	0.43
62:v:96:ILE:HA	62:v:100:LYS:HE3	2.00	0.43
63:w:60:VAL:HA	63:w:63:ARG:CZ	2.47	0.43
63:w:60:VAL:CG2	63:w:63:ARG:NH1	2.78	0.43
1:0:73:LYS:HZ1	1:0:88:GLY:HA3	1.83	0.43
4:3:6:ARG:CG	4:3:7:ARG:H	2.31	0.43
6:5:20:DC:H1'	6:5:21:DA:H5'	2.00	0.43
9:9:3:LEU:CD1	9:9:4:ASN:N	2.63	0.43
10:A:49:G:H8	10:A:49:G:O5'	2.01	0.43
11:AA:813:GLU:HB2	14:AE:461:PHE:HD2	1.83	0.43
16:AG:162:ILE:HD13	16:AG:167:MET:CE	2.49	0.43
16:AG:316:GLN:HE21	16:AG:316:GLN:HB2	1.49	0.43
16:AG:318:ILE:CG1	16:AG:338:VAL:HG11	2.48	0.43
16:AG:320:ARG:HB2	16:AG:323:GLN:CB	2.48	0.43
16:AG:331:LEU:HA	16:AG:331:LEU:HD23	1.79	0.43
10:B:31:G:H5'	18:D:1339:A:C2	2.54	0.43
10:B:71:C:H2'	10:B:72:A:N9	2.32	0.43
18:D:108:G:H5''	18:D:108:G:N3	2.33	0.43
18:D:320:A:H2'	18:D:321:A:O4'	2.19	0.43
18:D:564:C:OP1	32:R:12:ARG:CZ	2.66	0.43
18:D:1060:U:H5''	30:P:53:ILE:HD12	1.99	0.43
18:D:1062:U:H2'	18:D:1063:C:C6	2.53	0.43
18:D:1208:C:C2	18:D:1209:C:C6	3.07	0.43
18:D:1389:C:H2'	18:D:1390:U:O4'	2.17	0.43
21:G:101:LEU:HD22	21:G:157:LEU:HD11	1.99	0.43
23:I:112:ASP:O	23:I:116:VAL:HG23	2.18	0.43
23:I:157:LEU:CD1	23:I:166:GLU:OE1	2.66	0.43
24:J:35:GLU:HG2	24:J:36:GLN:N	2.32	0.43
26:L:9:MET:HE3	26:L:86:ARG:N	2.33	0.43
28:N:35:ALA:HB1	28:N:110:VAL:CG1	2.48	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:S:86:GLU:O	33:S:90:ARG:HG2	2.17	0.43
35:U:9:HIS:O	35:U:16:PHE:HB3	2.18	0.43
39:Y:32:VAL:HG13	39:Y:58:ILE:HG21	2.00	0.43
39:Y:81:LYS:HD2	39:Y:81:LYS:H	1.83	0.43
41:a:133:U:H2'	41:a:134:G:C8	2.54	0.43
41:a:153:U:H2'	41:a:154:U:C6	2.53	0.43
41:a:299:A:N1	41:a:322:A:O2'	2.42	0.43
41:a:587:C:O2	61:u:33:ARG:NH2	2.52	0.43
41:a:1084:A:N6	41:a:1085:A:N1	2.66	0.43
41:a:1170:C:O2	41:a:1170:C:H2'	2.18	0.43
41:a:1665:A:O3'	60:t:66:LYS:NZ	2.44	0.43
41:a:1695:G:N7	48:h:14:ARG:NH2	2.66	0.43
41:a:1839:G:O4'	41:a:1927:A:O4'	2.37	0.43
41:a:2845:U:H5''	65:y:52:ASN:O	2.19	0.43
41:a:2846:G:H2'	41:a:2847:U:O4'	2.19	0.43
43:c:4:VAL:HG22	43:c:11:ARG:HG2	2.00	0.43
44:d:31:C:C2	44:d:32:U:C5	3.06	0.43
48:h:130:LEU:CD2	48:h:192:LEU:HD21	2.48	0.43
50:j:204:LYS:HD3	50:j:205:PRO:HD2	2.00	0.43
51:k:21:TYR:CE2	51:k:38:LYS:HB3	2.54	0.43
54:n:15:LYS:O	54:n:19:GLU:OE1	2.35	0.43
55:o:45:ARG:HD2	55:o:45:ARG:C	2.43	0.43
56:p:69:ARG:NH1	56:p:73:ASN:ND2	2.66	0.43
56:p:83:PHE:HB2	56:p:141:ILE:HD13	2.00	0.43
58:r:6:LEU:C	58:r:15:LEU:HD13	2.43	0.43
60:t:44:LYS:N	60:t:44:LYS:HD2	2.33	0.43
60:t:59:LYS:NZ	60:t:92:GLU:CG	2.81	0.43
62:v:115:GLU:O	62:v:116:ALA:C	2.61	0.43
63:w:96:ARG:O	63:w:113:ILE:HA	2.19	0.43
1:0:73:LYS:HD3	1:0:86:GLN:HE21	1.82	0.43
3:2:56:GLU:CD	3:2:88:LYS:N	2.76	0.43
4:3:7:ARG:O	4:3:25:VAL:O	2.36	0.43
10:A:11:A:H2'	10:A:12:G:O4'	2.17	0.43
11:AA:829:THR:HG23	11:AA:1059:ARG:HG2	2.00	0.43
14:AE:362:ARG:H	14:AE:365:GLN:HE21	1.66	0.43
14:AE:802:ASP:OD1	14:AE:1348:LYS:NZ	2.42	0.43
16:AG:252:VAL:HG13	16:AG:256:GLY:HA2	1.99	0.43
16:AG:316:GLN:H	16:AG:316:GLN:HG3	1.54	0.43
16:AG:358:THR:O	16:AG:358:THR:OG1	2.24	0.43
17:C:27:ALA:CA	17:C:30:LYS:HE3	2.49	0.43
18:D:316:C:C2	18:D:317:U:C5	3.06	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:1104:G:H4'	21:G:113:ARG:HH22	1.83	0.43
18:D:1329:A:OP1	38:X:28:THR:CB	2.66	0.43
18:D:1410:A:H2'	18:D:1411:C:C6	2.54	0.43
21:G:23:TRP:CZ3	21:G:25:PRO:CA	3.01	0.43
21:G:30:PHE:CG	21:G:45:LYS:NZ	2.87	0.43
21:G:131:LYS:HE3	21:G:132:LYS:HZ3	1.84	0.43
22:H:305:HIS:CG	22:H:306:VAL:H	2.36	0.43
23:I:22:TRP:CZ2	23:I:32:ASN:CB	3.01	0.43
23:I:149:ILE:HG13	23:I:202:ILE:HG12	2.01	0.43
23:I:151:VAL:CG1	23:I:200:VAL:HG23	2.47	0.43
27:M:23:LEU:N	27:M:23:LEU:CD1	2.81	0.43
27:M:23:LEU:H	27:M:23:LEU:CD1	2.30	0.43
27:M:108:ALA:O	27:M:119:ARG:NE	2.50	0.43
32:R:18:LYS:HD3	32:R:19:SER:N	2.33	0.43
33:S:13:ARG:HB3	33:S:60:GLN:CG	2.46	0.43
38:X:101:ARG:C	38:X:103:LYS:H	2.26	0.43
39:Y:14:ALA:C	39:Y:50:LYS:HD2	2.44	0.43
40:Z:29:LYS:O	40:Z:30:PHE:HD2	2.02	0.43
41:a:966:G:O4'	41:a:2267:A:N6	2.51	0.43
41:a:1141:U:H5	59:s:68:LYS:HZ1	1.67	0.43
41:a:1792:G:C5'	48:h:204:VAL:HG23	2.48	0.43
41:a:2016:U:H2'	41:a:2017:U:C6	2.52	0.43
41:a:2465:C:O2	41:a:2465:C:H2'	2.18	0.43
41:a:2768:U:OP1	59:s:85:LYS:HE3	2.19	0.43
44:d:39:A:C2	44:d:44:G:C2	3.07	0.43
50:j:181:ASP:HB3	50:j:186:LEU:HB2	1.99	0.43
59:s:88:THR:OG1	59:s:91:GLU:HG3	2.18	0.43
62:v:74:THR:HA	62:v:88:ASN:O	2.19	0.43
62:v:112:LEU:HD12	62:v:112:LEU:N	2.34	0.43
63:w:20:MET:HB2	63:w:20:MET:HE3	1.83	0.43
63:w:58:ASP:HB2	63:w:80:PHE:CE1	2.54	0.43
63:w:87:PHE:HE2	63:w:115:LEU:HD11	1.83	0.43
65:y:55:LEU:HA	65:y:77:HIS:CD2	2.54	0.43
66:z:14:HIS:O	66:z:18:LEU:HD23	2.18	0.43
11:AA:300:ASP:OD1	11:AA:313:ALA:N	2.52	0.43
14:AE:975:ILE:HG22	14:AE:977:SER:H	1.83	0.43
16:AG:227:ALA:HB2	16:AG:330:GLN:O	2.18	0.43
16:AG:431:ALA:O	16:AG:435:LEU:HG	2.18	0.43
17:C:29:LEU:CD2	17:C:59:ILE:HD13	2.48	0.43
18:D:376:G:C2	18:D:389:A:C2	3.06	0.43
18:D:414:A:C2	18:D:415:A:H1'	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:562:U:C3'	32:R:14:ARG:NH1	2.81	0.43
18:D:622:A:C8	18:D:623:C:C5	3.06	0.43
18:D:771:G:H2'	18:D:772:U:C6	2.54	0.43
18:D:877:G:H5''	28:N:80:ARG:CD	2.49	0.43
18:D:1184:G:C2	18:D:1185:G:C8	3.07	0.43
18:D:1240:U:H5	27:M:116:MET:HE2	1.83	0.43
18:D:1244:G:C6	18:D:1294:G:C6	3.06	0.43
18:D:1464:U:C2	18:D:1465:A:C8	3.07	0.43
19:E:22:ALA:CA	19:E:25:ARG:HE	2.31	0.43
21:G:187:VAL:N	21:G:200:ILE:O	2.40	0.43
22:H:19:ARG:HD3	22:H:19:ARG:N	2.32	0.43
23:I:69:HIS:CD2	23:I:104:ALA:HB3	2.54	0.43
26:L:25:TYR:O	26:L:26:THR:C	2.61	0.43
40:Z:2:ILE:HG22	40:Z:4:LYS:N	2.28	0.43
41:a:39:G:H2'	41:a:40:U:C6	2.53	0.43
41:a:1144:A:C5	41:a:1145:C:C5	3.07	0.43
41:a:1360:G:C8	41:a:1361:G:C8	3.06	0.43
41:a:1409:U:C2	41:a:1410:G:C8	3.07	0.43
41:a:1506:U:H2'	41:a:1507:C:C6	2.54	0.43
41:a:2590:A:H2'	41:a:2591:C:H6	1.83	0.43
44:d:55:U:O2'	54:n:26:MET:CE	2.67	0.43
48:h:220:VAL:HG23	48:h:225:MET:HE3	2.00	0.43
54:n:92:ARG:CZ	54:n:92:ARG:HB3	2.47	0.43
54:n:132:VAL:CG2	54:n:152:LEU:CD2	2.97	0.43
56:p:94:TYR:CE1	56:p:152:ARG:CD	3.01	0.43
1:0:77:PHE:CD2	66:z:40:ILE:HG21	2.53	0.43
6:5:18:DG:C4	6:5:19:DC:C5	3.06	0.43
10:A:35:A:H2'	10:A:36:U:O4'	2.18	0.43
11:AA:557:ARG:HB3	11:AA:587:LEU:HD13	2.00	0.43
11:AA:646:SER:OG	11:AA:647:ARG:N	2.52	0.43
12:AB:116:VAL:H	12:AB:129:ILE:CD1	2.32	0.43
14:AE:79:LYS:CB	16:AG:159:GLU:OE2	2.60	0.43
14:AE:438:GLU:OE2	14:AE:481:ARG:NH1	2.44	0.43
15:AF:58:LEU:HD12	15:AF:59:ILE:HG12	2.00	0.43
16:AG:226:ALA:HB1	16:AG:228:ARG:NH1	2.34	0.43
18:D:57:G:H2'	18:D:58:C:C6	2.54	0.43
18:D:337:G:H2'	18:D:338:A:H8	1.80	0.43
18:D:404:G:N7	24:J:2:ALA:HB3	2.33	0.43
18:D:611:C:N3	18:D:612:C:C5	2.87	0.43
18:D:690:G:H2'	18:D:691:G:C8	2.54	0.43
18:D:1125:U:O2'	18:D:1126:U:O5'	2.32	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:E:58:VAL:HG12	19:E:72:ALA:HB1	1.99	0.43
22:H:71:ALA:HB3	22:H:83:LEU:O	2.18	0.43
22:H:76:GLU:OE2	22:H:78:GLY:C	2.61	0.43
25:K:97:GLN:NE2	25:K:98:PRO:O	2.52	0.43
27:M:78:ARG:HG3	27:M:80:VAL:HG13	1.99	0.43
28:N:66:PHE:C	28:N:67:GLN:OE1	2.62	0.43
30:P:85:ASP:HA	30:P:88:MET:HE3	1.99	0.43
32:R:57:LEU:HD23	32:R:59:ASN:CG	2.43	0.43
38:X:106:ALA:HB3	38:X:110:LYS:CD	2.48	0.43
41:a:307:G:N1	41:a:310:A:OP2	2.47	0.43
41:a:404:A:HO2'	41:a:405:U:P	2.38	0.43
41:a:451:U:C5'	52:l:47:LYS:HZ3	2.32	0.43
41:a:969:G:H2'	41:a:970:U:C6	2.54	0.43
41:a:1114:C:H2'	41:a:1115:G:C8	2.54	0.43
41:a:1289:C:H2'	41:a:1290:C:H6	1.84	0.43
41:a:1913:A:OP2	41:a:1913:A:H3'	2.18	0.43
41:a:1946:U:H2'	41:a:1947:C:H6	1.84	0.43
41:a:2216:G:H2'	41:a:2217:G:H8	1.82	0.43
41:a:2392:A:N3	61:u:55:MET:HE2	2.34	0.43
41:a:2516:A:O2'	41:a:2517:C:H5'	2.18	0.43
45:e:8:GLU:O	45:e:9:LYS:HD3	2.19	0.43
47:g:37:CYS:N	47:g:40:CYS:SG	2.71	0.43
49:i:55:ILE:CD1	63:w:112:TYR:CD2	3.02	0.43
50:j:172:VAL:HG11	50:j:175:LEU:HD11	1.99	0.43
52:l:112:LEU:HD23	52:l:118:LEU:HD13	1.99	0.43
57:q:3:VAL:HG12	57:q:36:ARG:HB3	2.00	0.43
58:r:93:SER:OG	58:r:121:VAL:HG13	2.18	0.43
59:s:28:LEU:HD12	59:s:142:ILE:HG21	2.01	0.43
60:t:118:LEU:O	60:t:119:ALA:C	2.62	0.43
61:u:95:LEU:HD13	61:u:100:ILE:HD12	2.00	0.43
63:w:98:LEU:N	63:w:112:TYR:O	2.51	0.43
8:7:7:U:H6	8:7:7:U:H3'	1.84	0.43
9:9:18:VAL:HG13	9:9:71:CYS:HB2	2.00	0.43
9:9:24:SER:O	9:9:96:PHE:CE1	2.72	0.43
9:9:35:VAL:O	9:9:35:VAL:HG22	2.19	0.43
10:A:72:A:C3'	10:A:73:A:H5''	2.47	0.43
11:AA:858:GLY:H	16:AG:38:LYS:HE3	1.84	0.43
12:AB:2:GLN:HA	12:AB:59:PHE:C	2.43	0.43
14:AE:412:LEU:HA	14:AE:415:VAL:HG22	2.01	0.43
10:B:75:C:OP1	41:a:2602:A:C8	2.71	0.43
17:C:60:LYS:NZ	18:D:735:C:O5'	2.52	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:296:U:O2'	18:D:556:C:O2	2.34	0.43
18:D:416:G:C4	18:D:417:G:C8	3.07	0.43
18:D:611:C:H2'	18:D:611:C:O2	2.18	0.43
18:D:642:A:C5	18:D:643:C:C5	3.06	0.43
18:D:1172:C:H2'	18:D:1173:U:H6	1.82	0.43
18:D:1317:C:OP2	33:S:28:LYS:CE	2.66	0.43
18:D:1328:C:H5''	38:X:28:THR:HG21	2.01	0.43
20:F:31:GLU:CD	20:F:34:ARG:NH2	2.77	0.43
20:F:39:GLU:OE2	20:F:44:GLU:HA	2.18	0.43
21:G:96:TRP:CZ2	21:G:100:MET:HE3	2.52	0.43
22:H:19:ARG:O	22:H:72:LEU:HD13	2.18	0.43
24:J:69:GLU:HA	24:J:72:PHE:HD2	1.83	0.43
24:J:117:LEU:HD23	24:J:154:ARG:HH21	1.84	0.43
26:L:1:MET:HE2	26:L:67:PRO:N	2.34	0.43
26:L:16:GLU:H	26:L:16:GLU:CD	2.26	0.43
30:P:66:GLU:HG3	33:S:99:ALA:HB2	2.00	0.43
31:Q:47:ALA:O	31:Q:52:PHE:HB2	2.19	0.43
32:R:80:ILE:CG2	32:R:81:LEU:N	2.82	0.43
34:T:71:LYS:NZ	34:T:78:TYR:CD2	2.87	0.43
36:V:48:ASP:HB3	36:V:75:LEU:CD2	2.49	0.43
38:X:22:ILE:HG23	38:X:66:GLU:CD	2.43	0.43
39:Y:139:VAL:HG22	39:Y:140:GLU:H	1.82	0.43
41:a:1385:A:C4	41:a:1403:A:C2	3.07	0.43
41:a:2297:A:C2	41:a:2321:U:C5	3.07	0.43
41:a:2484:G:OP1	62:v:44:ARG:HD2	2.18	0.43
41:a:2821:A:C2	41:a:2822:G:C4	3.07	0.43
45:e:37:LEU:HD21	45:e:43:LEU:CD2	2.49	0.43
46:f:10:THR:CB	46:f:56:LYS:NZ	2.81	0.43
52:l:131:THR:HG22	52:l:160:ALA:O	2.18	0.43
52:l:193:VAL:O	52:l:194:LYS:C	2.62	0.43
53:m:4:THR:HG23	53:m:5:PHE:N	2.34	0.43
54:n:6:ASP:O	54:n:10:ASP:OD2	2.36	0.43
54:n:66:LEU:HD23	54:n:67:ILE:N	2.34	0.43
54:n:132:VAL:HG11	54:n:137:ILE:HD11	2.01	0.43
58:r:137:GLU:OE1	58:r:137:GLU:N	2.41	0.43
59:s:1:MET:SD	59:s:2:LYS:C	3.02	0.43
60:t:108:ARG:HG2	60:t:116:ILE:HD13	2.01	0.43
61:u:101:ILE:HG13	61:u:102:GLY:H	1.82	0.43
61:u:103:ILE:HG23	61:u:104:GLN:HG2	1.99	0.43
11:AA:861:ALA:HB1	11:AA:882:ILE:HD13	2.01	0.43
12:AB:33:ILE:HD11	12:AB:104:ILE:HA	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:AD:212:ASP:OD1	13:AD:212:ASP:N	2.48	0.43
16:AG:224:LYS:HB2	16:AG:276:TRP:CH2	2.54	0.43
16:AG:353:HIS:C	16:AG:357:ASP:HB2	2.43	0.43
16:AG:442:ARG:O	16:AG:446:PHE:CE2	2.63	0.43
18:D:33:A:H2'	18:D:34:C:C6	2.53	0.43
18:D:43:C:P	35:U:12:LYS:HE3	2.59	0.43
18:D:381:C:H2'	18:D:382:A:O4'	2.19	0.43
18:D:563:A:C2	18:D:567:G:C6	3.07	0.43
18:D:678:U:H2'	18:D:679:C:C6	2.53	0.43
18:D:1325:C:N3	18:D:1326:U:C5	2.86	0.43
18:D:1381:U:C4	18:D:1382:C:C5	3.07	0.43
18:D:1513:A:H2'	18:D:1514:G:H8	1.84	0.43
18:D:1526:G:P	20:F:42:THR:HG23	2.59	0.43
19:E:78:ASN:O	19:E:82:GLN:CD	2.61	0.43
20:F:14:VAL:O	20:F:18:ARG:HG3	2.19	0.43
20:F:51:SER:OG	20:F:55:ARG:NH2	2.52	0.43
22:H:330:VAL:HB	22:H:344:LEU:HD23	2.01	0.43
23:I:147:LYS:HE2	23:I:204:LYS:C	2.42	0.43
25:K:68:ARG:O	25:K:69:ARG:NH1	2.52	0.43
29:O:21:ILE:CD1	29:O:86:ALA:CB	2.97	0.43
29:O:80:ARG:O	29:O:84:THR:HG23	2.18	0.43
30:P:64:GLN:HG2	33:S:101:TRP:HH2	1.84	0.43
31:Q:85:MET:HB3	31:Q:113:VAL:HG21	2.00	0.43
37:W:17:LYS:O	37:W:21:LYS:HD3	2.19	0.43
37:W:39:THR:HG22	37:W:40:ILE:N	2.33	0.43
38:X:64:VAL:O	38:X:64:VAL:HG13	2.19	0.43
38:X:90:ARG:NH1	38:X:96:PRO:O	2.51	0.43
41:a:39:G:H2'	41:a:40:U:H6	1.83	0.43
41:a:407:G:C2	41:a:408:G:C8	3.07	0.43
41:a:910:A:H8	62:v:12:MET:CE	2.31	0.43
41:a:1198:U:H2'	41:a:1199:U:H6	1.83	0.43
41:a:1269:A:H2'	41:a:1270:C:C6	2.54	0.43
41:a:1657:U:C2	41:a:1658:C:C5	3.06	0.43
41:a:2175:C:H2'	41:a:2176:A:N7	2.33	0.43
44:d:69:G:C5	44:d:70:C:C6	3.06	0.43
52:l:105:LEU:O	52:l:109:LEU:HG	2.18	0.43
54:n:38:MET:HE3	54:n:57:LEU:HD23	2.01	0.43
60:t:1:MET:HE1	60:t:32:TYR:CB	2.48	0.43
62:v:3:GLN:OE1	62:v:4:PRO:O	2.36	0.43
62:v:111:GLU:HG2	62:v:112:LEU:N	2.33	0.43
8:7:1:A:H2'	8:7:2:U:C6	2.50	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:9:25:ALA:HB2	9:9:96:PHE:HE1	1.82	0.43
11:AA:998:LEU:HG	11:AA:1011:LEU:HB3	1.99	0.43
11:AA:1157:GLN:HG3	11:AA:1159:VAL:HG13	2.00	0.43
12:AB:4:TRP:CE2	12:AB:27:ASN:ND2	2.87	0.43
14:AE:708:ASN:HA	14:AE:713:GLU:HA	2.00	0.43
16:AG:320:ARG:HH11	16:AG:320:ARG:HG2	1.80	0.43
10:B:71:C:C3'	10:B:72:A:C8	2.99	0.43
18:D:13:U:N3	18:D:915:A:C6	2.87	0.43
18:D:246:A:C2	18:D:282:A:C4	3.06	0.43
18:D:588:G:C4'	28:N:3:MET:HE1	2.48	0.43
18:D:621:A:H2'	18:D:622:A:C8	2.54	0.43
18:D:736:C:H4'	26:L:88:MET:HE2	1.99	0.43
18:D:1073:U:C2	18:D:1074:G:C8	3.07	0.43
18:D:1217:C:P	33:S:9:ARG:HE	2.41	0.43
18:D:1261:A:C5	18:D:1262:C:C6	3.07	0.43
18:D:1434:A:H2'	18:D:1435:G:O4'	2.18	0.43
22:H:8:LEU:HD12	22:H:11:GLU:OE2	2.18	0.43
26:L:29:ILE:HD12	26:L:29:ILE:H	1.83	0.43
35:U:4:ILE:N	35:U:4:ILE:HD12	2.34	0.43
37:W:69:HIS:ND1	37:W:74:PHE:HZ	2.17	0.43
39:Y:99:LYS:CE	39:Y:140:GLU:OE2	2.66	0.43
41:a:61:C:C2	41:a:94:A:C2	3.06	0.43
41:a:78:U:C5'	45:e:4:LYS:NZ	2.81	0.43
41:a:126:A:OP1	53:m:45:SER:OG	2.37	0.43
41:a:367:G:C4	41:a:368:A:C8	3.07	0.43
41:a:538:A:H2'	41:a:539:G:O4'	2.18	0.43
41:a:818:G:N1	41:a:1188:U:OP2	2.36	0.43
41:a:870:U:N3	41:a:871:U:C5	2.86	0.43
41:a:997:G:H5''	66:z:92:ARG:NH1	2.34	0.43
41:a:1028:A:N6	41:a:1125:G:H2'	2.34	0.43
41:a:1074:G:C6	41:a:1075:C:N4	2.87	0.43
41:a:1565:C:OP1	48:h:18:LYS:HD2	2.19	0.43
41:a:2049:G:H21	50:j:161:MET:HE2	1.84	0.43
41:a:2287:A:N7	41:a:2289:G:C8	2.87	0.43
41:a:2547:A:H5'	60:t:29:HIS:NE2	2.34	0.43
41:a:2665:A:C2	41:a:2666:C:C6	3.07	0.43
43:c:76:GLU:CD	43:c:76:GLU:C	2.87	0.43
45:e:59:GLU:CD	45:e:60:LYS:N	2.76	0.43
48:h:94:VAL:HG12	48:h:95:LEU:N	2.34	0.43
50:j:81:GLU:O	50:j:82:PHE:HD1	2.01	0.43
51:k:7:GLU:OE2	51:k:27:LYS:N	2.50	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:k:34:LEU:HB3	51:k:51:GLU:OE2	2.19	0.43
54:n:20:PHE:HB3	54:n:22:TYR:CE1	2.54	0.43
54:n:136:ILE:HA	54:n:141:ILE:HG21	2.01	0.43
66:z:12:ALA:O	66:z:16:LYS:HG2	2.19	0.43
1:0:1:MET:HE3	1:0:43:ASN:OD1	2.18	0.43
1:0:52:PRO:HB3	66:z:89:GLU:CD	2.44	0.43
3:2:42:GLU:OE1	3:2:42:GLU:HA	2.19	0.43
5:4:80:HIS:HE1	5:4:82:TYR:CG	2.37	0.43
9:9:23:LEU:HA	9:9:118:ILE:HG13	2.01	0.43
12:AB:104:ILE:C	12:AB:106:ASP:H	2.27	0.43
16:AG:38:LYS:HB3	16:AG:39:TYR:H	1.55	0.43
16:AG:183:LEU:HD22	16:AG:195:LEU:HD13	2.00	0.43
16:AG:201:LYS:HB3	16:AG:201:LYS:HZ3	1.83	0.43
18:D:5:U:O2	18:D:5:U:O4'	2.37	0.43
18:D:119:A:C2	18:D:240:G:C8	3.07	0.43
18:D:723:U:C2	20:F:49:LYS:HE3	2.54	0.43
18:D:745:G:H2'	18:D:746:A:H8	1.84	0.43
18:D:754:C:O5'	34:T:72:ARG:NH1	2.51	0.43
18:D:964:A:C2	18:D:969:A:N3	2.87	0.43
18:D:1408:A:H2'	18:D:1409:C:H6	1.84	0.43
20:F:5:LYS:HD3	31:Q:109:ASN:OD1	2.19	0.43
21:G:26:LYS:CE	21:G:193:PRO:HG2	2.49	0.43
22:H:31:ILE:O	22:H:31:ILE:HG13	2.18	0.43
24:J:76:TYR:HE1	24:J:86:THR:HG21	1.84	0.43
28:N:31:LYS:N	28:N:31:LYS:HD2	2.34	0.43
28:N:47:GLU:OE1	28:N:63:LEU:C	2.62	0.43
30:P:56:HIS:C	30:P:57:VAL:HG23	2.44	0.43
34:T:10:LYS:HE2	34:T:10:LYS:HA	2.00	0.43
37:W:40:ILE:HG22	37:W:67:VAL:HA	2.01	0.43
38:X:25:VAL:O	38:X:25:VAL:HG13	2.19	0.43
38:X:71:ARG:CZ	47:g:50:ASP:O	2.67	0.43
38:X:92:ARG:NH1	41:a:888:C:OP1	2.38	0.43
41:a:320:A:N3	52:l:163:ASN:ND2	2.67	0.43
41:a:838:C:C2	41:a:839:U:C6	3.07	0.43
41:a:1141:U:O2	41:a:1142:A:N6	2.52	0.43
41:a:1657:U:H2'	41:a:1658:C:H6	1.83	0.43
41:a:1770:G:C5	41:a:1983:G:C6	3.07	0.43
41:a:1792:G:H5'	48:h:204:VAL:CG2	2.49	0.43
41:a:1879:C:H2'	41:a:1880:U:O4'	2.18	0.43
41:a:2061:G:N3	41:a:2063:C:C5	2.87	0.43
41:a:2179:C:H2'	41:a:2180:U:H6	1.83	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2273:A:H2'	41:a:2274:A:C8	2.54	0.43
41:a:2547:A:H4'	60:t:29:HIS:NE2	2.34	0.43
41:a:2615:U:C2	49:i:4:GLN:HA	2.53	0.43
48:h:183:LYS:CE	48:h:265:LYS:O	2.67	0.43
49:i:55:ILE:HD13	63:w:112:TYR:CE2	2.54	0.43
50:j:37:VAL:HG13	50:j:48:ILE:CD1	2.49	0.43
52:l:6:LYS:HZ1	52:l:119:ILE:C	2.26	0.43
59:s:124:VAL:O	59:s:125:TYR:HD1	2.01	0.43
62:v:72:PRO:C	62:v:73:ILE:HD12	2.43	0.43
62:v:105:MET:HG2	62:v:117:PHE:CZ	2.54	0.43
65:y:106:LYS:CB	65:y:109:ARG:HH22	2.32	0.43
2:1:19:LEU:HD23	2:1:19:LEU:C	2.43	0.42
2:1:36:LEU:HD23	2:1:48:LYS:CA	2.48	0.42
3:2:6:ARG:HE	3:2:42:GLU:CD	2.27	0.42
5:4:57:TYR:HD2	62:v:136:MET:HE2	1.81	0.42
6:5:16:DG:N2	11:AA:445:ILE:HG21	2.34	0.42
9:9:94:ARG:N	9:9:94:ARG:HD3	2.34	0.42
10:A:11:A:H2'	10:A:12:G:C8	2.54	0.42
11:AA:125:GLY:H	11:AA:495:ALA:HB1	1.84	0.42
11:AA:853:ASP:HA	16:AG:102:PHE:CE1	2.54	0.42
11:AA:855:PRO:HG3	16:AG:105:ILE:CG2	2.40	0.42
12:AB:5:TYR:HA	12:AB:88:VAL:CG2	2.49	0.42
12:AB:119:THR:HG21	12:AB:123:PHE:CE1	2.54	0.42
13:AD:68:TYR:HE1	13:AD:79:LEU:HD13	1.82	0.42
14:AE:658:GLU:HA	14:AE:661:VAL:HG22	2.00	0.42
16:AG:21:GLU:HB3	16:AG:49:ILE:CD1	2.49	0.42
16:AG:161:VAL:HB	16:AG:196:PHE:CE1	2.54	0.42
18:D:68:G:H3'	18:D:69:G:H5''	2.01	0.42
18:D:375:U:OP2	35:U:70:ARG:NE	2.51	0.42
18:D:384:G:H2'	18:D:385:C:C6	2.54	0.42
18:D:415:A:C4	18:D:416:G:C8	3.07	0.42
18:D:427:U:OP2	18:D:428:G:O2'	2.31	0.42
18:D:1157:A:C2	18:D:1181:G:C5	3.06	0.42
18:D:1315:U:H2'	18:D:1316:G:O4'	2.19	0.42
19:E:21:ASN:HB3	19:E:25:ARG:CZ	2.48	0.42
20:F:3:VAL:HG22	31:Q:111:THR:HG23	2.01	0.42
22:H:336:ASP:HB2	22:H:341:ARG:H	1.84	0.42
23:I:72:ARG:O	23:I:75:ILE:HG22	2.19	0.42
23:I:149:ILE:HG13	23:I:201:TRP:O	2.18	0.42
26:L:2:ARG:HB2	26:L:4:TYR:CE1	2.53	0.42
32:R:25:GLU:OE2	32:R:30:LYS:HE3	2.18	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:W:12:ASP:OD2	37:W:35:SER:CB	2.66	0.42
37:W:45:ILE:HA	37:W:62:VAL:HG12	2.00	0.42
38:X:71:ARG:NH1	47:g:51:VAL:O	2.52	0.42
39:Y:82:ALA:HB3	39:Y:108:ILE:HD12	2.01	0.42
39:Y:99:LYS:CE	39:Y:138:VAL:O	2.67	0.42
41:a:74:A:N7	41:a:88:G:C5	2.87	0.42
41:a:125:A:C6	53:m:10:LEU:HD13	2.54	0.42
41:a:1139:G:O3'	59:s:26:GLY:HA3	2.19	0.42
41:a:1145:C:H2'	41:a:1146:C:H6	1.84	0.42
41:a:1511:G:H2'	41:a:1512:C:H6	1.84	0.42
41:a:1956:U:C2	41:a:1957:C:C6	3.06	0.42
41:a:1964:G:H4'	41:a:1965:C:OP2	2.19	0.42
41:a:2037:A:H2'	41:a:2038:G:C8	2.54	0.42
41:a:2557:G:H2'	41:a:2558:C:H6	1.84	0.42
41:a:2708:G:O4'	63:w:71:ARG:NH1	2.52	0.42
41:a:2756:U:N3	41:a:2758:A:C6	2.87	0.42
43:c:30:LEU:CD1	43:c:31:PRO:HD2	2.42	0.42
44:d:57:A:C4	44:d:58:A:C8	3.07	0.42
50:j:61:THR:HG23	50:j:63:PRO:HD2	2.00	0.42
50:j:156:PHE:CE2	59:s:81:ILE:HD13	2.54	0.42
50:j:184:ARG:NH2	65:y:7:GLN:CD	2.72	0.42
53:m:25:LYS:HG3	53:m:29:GLN:NE2	2.34	0.42
55:o:8:ARG:HA	55:o:8:ARG:HE	1.82	0.42
62:v:38:ARG:HB2	62:v:98:PRO:HD3	2.00	0.42
62:v:71:LYS:C	62:v:92:TRP:CE3	2.97	0.42
66:z:47:TYR:CZ	66:z:51:ARG:NH2	2.87	0.42
1:0:60:LYS:N	1:0:60:LYS:HD3	2.33	0.42
3:2:76:ARG:CZ	3:2:77:ARG:O	2.67	0.42
5:4:77:VAL:CG2	5:4:79:ARG:NE	2.79	0.42
9:9:3:LEU:CD1	9:9:5:LEU:H	2.31	0.42
9:9:19:ALA:CB	9:9:69:PHE:HE2	2.31	0.42
9:9:69:PHE:O	9:9:72:LEU:HD11	2.19	0.42
9:9:70:GLU:HG3	9:9:71:CYS:N	2.34	0.42
10:A:19:G:N3	10:A:57:A:C4	2.87	0.42
11:AA:22:LEU:HB3	11:AA:655:VAL:HG11	2.01	0.42
11:AA:699:LEU:HG	11:AA:799:ASN:HD22	1.84	0.42
11:AA:755:LYS:HD2	11:AA:755:LYS:HA	1.86	0.42
11:AA:861:ALA:CB	16:AG:38:LYS:HZ1	2.32	0.42
12:AB:26:VAL:HG13	12:AB:58:GLU:H	1.84	0.42
12:AB:29:LEU:H	12:AB:56:PHE:HB2	1.84	0.42
13:AC:228:LEU:HD21	13:AD:224:LEU:HD23	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:296:ILE:CD1	16:AG:307:ILE:HA	2.49	0.42
16:AG:434:LEU:HD23	16:AG:456:LEU:CB	2.49	0.42
17:C:33:ILE:O	22:H:338:GLU:HB3	2.19	0.42
17:C:40:VAL:HG22	22:H:339:ARG:HE	1.83	0.42
17:C:44:ILE:CG2	22:H:340:ARG:HB2	2.50	0.42
18:D:250:A:O4'	18:D:252:U:C6	2.72	0.42
18:D:283:U:C2	18:D:284:C:C6	3.06	0.42
18:D:671:G:H4'	26:L:79:ARG:HH12	1.83	0.42
18:D:687:A:C2	18:D:704:A:C6	3.07	0.42
18:D:1276:G:O5'	18:D:1276:G:H8	2.02	0.42
22:H:123:GLU:HA	22:H:128:ARG:HA	2.01	0.42
23:I:54:ARG:NH1	23:I:69:HIS:NE2	2.66	0.42
23:I:72:ARG:HD3	23:I:75:ILE:H	1.84	0.42
34:T:10:LYS:HA	34:T:10:LYS:CE	2.49	0.42
35:U:7:ALA:HA	35:U:28:ARG:HD3	2.01	0.42
39:Y:42:ASN:OD1	39:Y:50:LYS:NZ	2.46	0.42
39:Y:45:THR:HG21	39:Y:50:LYS:NZ	2.34	0.42
41:a:299:A:C5	41:a:322:A:C2	3.08	0.42
41:a:1080:A:C6	41:a:1081:U:C4	3.08	0.42
41:a:2305:U:H5''	54:n:131:GLY:HA3	2.00	0.42
41:a:2469:A:C6	41:a:2482:A:C8	3.07	0.42
41:a:2615:U:H2'	41:a:2616:C:H6	1.83	0.42
45:e:25:GLN:OE1	45:e:50:VAL:HG21	2.19	0.42
50:j:57:ALA:O	50:j:60:VAL:HG12	2.19	0.42
52:l:6:LYS:NZ	52:l:119:ILE:C	2.77	0.42
52:l:47:LYS:HA	52:l:51:GLU:CD	2.44	0.42
52:l:170:ARG:NH1	52:l:175:ILE:HA	2.34	0.42
53:m:25:LYS:O	53:m:28:ARG:N	2.52	0.42
59:s:56:VAL:O	59:s:125:TYR:N	2.50	0.42
9:9:80:THR:O	41:a:1108:U:OP1	2.37	0.42
9:9:129:LEU:H	9:9:129:LEU:HG	1.72	0.42
11:AA:590:PRO:HB2	11:AA:655:VAL:HG21	2.01	0.42
14:AE:210:SER:O	14:AE:214:ARG:N	2.43	0.42
16:AG:87:LEU:CD2	16:AG:93:VAL:HG21	2.49	0.42
10:B:11:A:H2'	10:B:12:G:C8	2.54	0.42
10:B:12:G:H2'	10:B:13:C:OP1	2.20	0.42
10:B:22:G:C8	10:B:22:G:OP2	2.72	0.42
10:B:57:A:O5'	10:B:58:A:P	2.77	0.42
18:D:21:G:H2'	18:D:22:G:C8	2.54	0.42
18:D:105:G:C5	18:D:106:C:C4	3.08	0.42
18:D:235:C:H2'	18:D:236:A:H8	1.83	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:311:C:P	35:U:31:ARG:NH2	2.86	0.42
18:D:505:G:C2	18:D:506:G:C5	3.07	0.42
18:D:610:U:C2	18:D:611:C:C6	3.06	0.42
18:D:679:C:N3	18:D:680:C:C5	2.87	0.42
18:D:921:U:H4'	25:K:23:LYS:HZ3	1.82	0.42
18:D:1071:C:C5'	25:K:54:ARG:HH21	2.33	0.42
18:D:1107:C:OP1	23:I:172:ARG:HG3	2.19	0.42
18:D:1194:U:H2'	18:D:1195:C:C6	2.54	0.42
21:G:49:MET:HB3	21:G:49:MET:HE2	1.79	0.42
23:I:54:ARG:NH2	23:I:56:VAL:CG2	2.82	0.42
26:L:18:VAL:HA	26:L:21:MET:CE	2.47	0.42
32:R:30:LYS:CB	32:R:57:LEU:HD11	2.47	0.42
35:U:19:VAL:HG11	35:U:52:LEU:CD2	2.50	0.42
36:V:43:LYS:C	36:V:44:LEU:HD12	2.45	0.42
39:Y:96:LYS:CD	39:Y:136:GLY:HA3	2.50	0.42
41:a:301:G:C4	41:a:302:C:C5	3.07	0.42
41:a:533:G:H2'	41:a:534:U:C6	2.54	0.42
41:a:580:U:C2	41:a:581:C:C5	3.07	0.42
41:a:814:C:C2	41:a:815:C:C5	3.07	0.42
41:a:996:A:N6	41:a:1160:G:C6	2.87	0.42
41:a:1683:U:H2'	41:a:1684:G:H8	1.83	0.42
41:a:1688:U:O2'	41:a:1700:A:N7	2.44	0.42
41:a:1906:G:C8	41:a:1929:G:C5	3.08	0.42
41:a:2066:C:O2'	41:a:2067:G:H5'	2.20	0.42
41:a:2215:C:H2'	41:a:2216:G:C8	2.54	0.42
41:a:2332:C:OP1	42:b:77:ARG:NH2	2.52	0.42
41:a:2859:G:H2'	41:a:2860:A:C8	2.54	0.42
46:f:4:THR:O	46:f:5:ILE:HD13	2.19	0.42
54:n:104:ILE:HD11	54:n:174:ASP:O	2.19	0.42
58:r:8:LYS:NZ	58:r:9:VAL:O	2.52	0.42
58:r:114:GLU:OE2	58:r:133:GLN:N	2.53	0.42
60:t:7:MET:SD	60:t:18:ARG:HG3	2.59	0.42
60:t:59:LYS:HZ3	60:t:92:GLU:CG	2.33	0.42
64:x:79:ALA:CB	64:x:113:ALA:HB3	2.49	0.42
66:z:111:GLU:OE2	66:z:112:LYS:HD2	2.19	0.42
1:0:40:MET:HE1	66:z:105:ALA:HB1	2.01	0.42
2:1:75:PHE:HE2	2:1:77:ASP:OD2	2.02	0.42
4:3:7:ARG:HG3	4:3:8:ASP:N	2.33	0.42
8:7:12:U:H4'	25:K:56:VAL:HG11	2.02	0.42
8:7:46:G:C2'	8:7:47:U:H5'	2.49	0.42
10:A:22:G:OP2	10:A:22:G:C8	2.72	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AB:35:LEU:HA	12:AB:106:ASP:CG	2.33	0.42
12:AB:112:PRO:HA	12:AB:133:PRO:CG	2.49	0.42
14:AE:694:SER:OG	14:AE:738:ARG:NE	2.52	0.42
16:AG:314:LEU:HD23	16:AG:314:LEU:HA	1.85	0.42
16:AG:354:ALA:O	16:AG:358:THR:HG22	2.19	0.42
16:AG:451:ARG:NH2	16:AG:470:ILE:CG1	2.67	0.42
17:C:30:LYS:HA	17:C:33:ILE:HG12	2.01	0.42
18:D:539:A:H2'	18:D:540:G:C8	2.54	0.42
18:D:674:G:H2'	18:D:675:A:H8	1.84	0.42
18:D:946:A:C2	18:D:947:G:C5	3.08	0.42
18:D:1149:C:N3	18:D:1150:A:N7	2.67	0.42
18:D:1172:C:C2	18:D:1173:U:C5	3.08	0.42
18:D:1202:U:N3	33:S:82:ILE:HD13	2.35	0.42
18:D:1436:U:C2	18:D:1437:A:C8	3.07	0.42
18:D:1477:U:H2'	18:D:1478:U:C6	2.53	0.42
18:D:1493:A:O2'	18:D:1494:G:P	2.78	0.42
18:D:1499:A:O2'	18:D:1500:A:H5'	2.20	0.42
18:D:1499:A:H2'	18:D:1500:A:H8	1.85	0.42
21:G:131:LYS:O	21:G:135:LEU:HD13	2.19	0.42
21:G:134:ALA:HA	21:G:137:ARG:HE	1.83	0.42
22:H:74:ALA:HB3	22:H:83:LEU:CD2	2.50	0.42
23:I:58:GLU:OE1	23:I:65:ARG:CZ	2.65	0.42
23:I:132:ARG:CB	23:I:135:LYS:HE3	2.49	0.42
23:I:156:ARG:O	23:I:156:ARG:HG3	2.20	0.42
25:K:123:VAL:C	25:K:124:LEU:HD23	2.45	0.42
26:L:102:MET:CG	26:L:103:VAL:N	2.78	0.42
28:N:27:MET:CA	28:N:58:GLU:OE2	2.67	0.42
29:O:86:ALA:O	29:O:89:GLU:CG	2.64	0.42
30:P:45:ARG:HD3	30:P:47:GLU:OE2	2.19	0.42
30:P:59:LYS:HE2	30:P:62:ARG:CZ	2.50	0.42
32:R:27:CYS:CB	32:R:30:LYS:HE2	2.47	0.42
32:R:59:ASN:OD1	32:R:59:ASN:C	2.62	0.42
32:R:116:LYS:O	32:R:117:TYR:HD1	2.02	0.42
39:Y:99:LYS:HD2	39:Y:140:GLU:OE2	2.19	0.42
41:a:471:A:H2'	41:a:472:A:O4'	2.19	0.42
41:a:523:C:O2	41:a:554:U:O2'	2.37	0.42
41:a:553:G:C5	41:a:554:U:C5	3.07	0.42
41:a:666:A:H1'	55:o:4:ILE:CD1	2.47	0.42
41:a:751:A:C6	41:a:789:A:C5	3.08	0.42
41:a:973:A:O4'	41:a:1188:U:C6	2.72	0.42
41:a:1542:U:H2'	41:a:1543:G:O4'	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1709:U:H2'	41:a:1710:G:C8	2.55	0.42
41:a:1789:A:OP2	48:h:221:ARG:NH1	2.52	0.42
41:a:1863:G:H2'	41:a:1864:U:H6	1.83	0.42
41:a:2462:C:C2	41:a:2463:C:C5	3.07	0.42
41:a:2856:A:C6	41:a:2862:G:C6	3.08	0.42
41:a:2884:U:C2	49:i:40:ARG:CZ	3.02	0.42
44:d:46:A:C5	44:d:47:C:C5	3.06	0.42
44:d:83:G:O4'	46:f:53:PHE:CE2	2.72	0.42
49:i:4:GLN:OE1	49:i:8:PRO:HD3	2.19	0.42
50:j:16:THR:HG23	50:j:20:VAL:O	2.19	0.42
50:j:100:LEU:O	50:j:100:LEU:HD12	2.19	0.42
50:j:107:VAL:HG13	50:j:177:VAL:HG11	2.00	0.42
54:n:4:LEU:HD11	54:n:101:GLU:CB	2.50	0.42
54:n:33:LYS:CA	54:n:96:MET:SD	3.07	0.42
56:p:45:HIS:HA	56:p:50:LEU:HD13	2.01	0.42
56:p:89:LEU:HD22	56:p:162:VAL:HA	2.02	0.42
59:s:133:ALA:O	59:s:134:ALA:C	2.62	0.42
62:v:105:MET:HE1	62:v:113:ALA:HA	2.01	0.42
66:z:76:TYR:OH	66:z:92:ARG:HG2	2.19	0.42
2:1:99:ARG:CZ	2:1:99:ARG:HB2	2.49	0.42
3:2:11:LEU:HD11	3:2:46:ALA:CB	2.50	0.42
9:9:4:ASN:HB3	9:9:8:LYS:HZ2	1.83	0.42
10:A:12:G:H2'	10:A:13:C:OP1	2.20	0.42
12:AB:90:SER:HB2	12:AB:94:HIS:CE1	2.54	0.42
13:AD:211:ILE:HD12	13:AD:211:ILE:HA	1.92	0.42
14:AE:859:PRO:HD2	14:AE:862:THR:HG21	2.00	0.42
16:AG:320:ARG:HA	16:AG:320:ARG:HD3	1.83	0.42
17:C:20:GLU:OE1	17:C:21:ILE:O	2.36	0.42
18:D:197:A:N6	18:D:221:C:H4'	2.35	0.42
18:D:792:A:H1'	18:D:794:A:N7	2.34	0.42
19:E:60:ARG:HG3	19:E:64:LYS:NZ	2.34	0.42
21:G:55:ALA:O	21:G:59:LYS:HG2	2.19	0.42
23:I:184:TYR:O	23:I:185:ASN:CG	2.63	0.42
26:L:14:GLN:NE2	26:L:83:ALA:HB2	2.35	0.42
30:P:7:ARG:HA	30:P:75:ASP:OD1	2.20	0.42
33:S:63:ARG:HH22	33:S:70:PRO:HD3	1.85	0.42
35:U:12:LYS:HD2	35:U:13:LYS:HG2	2.00	0.42
39:Y:102:ARG:NH1	39:Y:139:VAL:HG11	2.33	0.42
41:a:531:C:C5	41:a:2035:G:C2	3.07	0.42
41:a:639:U:H2'	41:a:640:C:H6	1.82	0.42
41:a:729:G:C8	48:h:207:LYS:HD2	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1560:G:C5	41:a:1561:C:C5	3.08	0.42
41:a:1900:A:N1	41:a:1970:A:C6	2.87	0.42
41:a:2002:G:OP1	63:w:13:ASN:HA	2.19	0.42
41:a:2072:C:C2	41:a:2073:C:C5	3.08	0.42
41:a:2093:G:P	58:r:22:LYS:CG	3.07	0.42
41:a:2228:G:H2'	41:a:2229:U:C6	2.54	0.42
44:d:39:A:C2	44:d:44:G:C4	3.07	0.42
44:d:57:A:N7	54:n:26:MET:CE	2.82	0.42
49:i:14:GLY:HA2	49:i:17:ARG:HG2	2.02	0.42
50:j:91:THR:HG22	50:j:92:VAL:N	2.35	0.42
50:j:104:VAL:HG12	50:j:105:LYS:N	2.33	0.42
52:l:6:LYS:NZ	52:l:119:ILE:CG2	2.83	0.42
54:n:73:SER:OG	54:n:80:ARG:HA	2.19	0.42
64:x:14:ALA:O	64:x:15:ARG:C	2.61	0.42
64:x:88:LYS:O	64:x:116:GLN:OE1	2.38	0.42
66:z:76:TYR:CZ	66:z:80:ILE:HD11	2.53	0.42
1:0:47:VAL:C	1:0:48:LYS:HD3	2.45	0.42
5:4:2:PHE:CD1	5:4:56:PHE:CE1	3.08	0.42
9:9:31:ARG:HH11	41:a:1054:A:C4'	2.31	0.42
10:A:59:A:C5	10:A:60:U:C5	3.08	0.42
11:AA:478:ARG:HD2	11:AA:492:MET:HA	2.02	0.42
11:AA:741:MET:SD	11:AA:974:ARG:NH2	2.93	0.42
11:AA:889:PRO:HG2	16:AG:104:ARG:NH1	2.34	0.42
13:AD:46:ILE:HD11	13:AD:224:LEU:HD13	2.02	0.42
16:AG:226:ALA:O	16:AG:228:ARG:N	2.52	0.42
16:AG:239:LYS:CG	16:AG:276:TRP:HB3	2.46	0.42
16:AG:343:ASP:HB3	16:AG:347:LYS:HE3	2.01	0.42
16:AG:398:LEU:C	16:AG:398:LEU:HD12	2.44	0.42
17:C:44:ILE:HG22	22:H:340:ARG:HB2	2.02	0.42
17:C:60:LYS:NZ	18:D:734:G:H2'	2.25	0.42
18:D:2:A:O2'	18:D:3:A:O4'	2.36	0.42
18:D:218:U:H2'	18:D:219:U:O4'	2.20	0.42
18:D:256:U:H2'	18:D:257:G:C8	2.55	0.42
18:D:374:A:C6	18:D:375:U:C4	3.08	0.42
18:D:393:A:C2	18:D:394:G:C8	3.07	0.42
18:D:501:C:H1'	18:D:549:C:H1'	2.01	0.42
18:D:1035:A:C6	18:D:1036:A:N7	2.88	0.42
20:F:39:GLU:OE1	20:F:44:GLU:OE1	2.38	0.42
22:H:310:ASP:HB2	22:H:316:ILE:HD11	2.00	0.42
23:I:113:ALA:HB1	23:I:200:VAL:HG13	2.02	0.42
23:I:134:MET:HG3	23:I:151:VAL:HG21	2.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:K:100:SER:O	25:K:103:THR:HG22	2.20	0.42
27:M:99:LEU:O	27:M:103:TRP:CG	2.73	0.42
27:M:103:TRP:CG	27:M:137:LYS:NZ	2.87	0.42
29:O:50:GLN:OE1	29:O:50:GLN:HA	2.20	0.42
31:Q:21:ALA:HA	31:Q:34:ILE:HD13	2.01	0.42
32:R:25:GLU:O	32:R:25:GLU:HG2	2.20	0.42
32:R:66:TYR:CE2	32:R:68:GLY:HA2	2.55	0.42
34:T:70:LEU:HG	34:T:78:TYR:HB2	2.02	0.42
41:a:196:A:C5	41:a:805:G:C6	3.07	0.42
41:a:250:G:H2'	41:a:251:A:C8	2.54	0.42
41:a:534:U:O2'	66:z:49:ASP:OD2	2.37	0.42
41:a:579:G:H2'	41:a:580:U:H6	1.85	0.42
41:a:833:A:OP2	61:u:39:LYS:NZ	2.49	0.42
41:a:1361:G:H2'	41:a:1362:C:C6	2.54	0.42
41:a:1365:A:H4'	43:c:11:ARG:HH12	1.85	0.42
41:a:1707:G:C5	41:a:1708:C:C5	3.07	0.42
41:a:1773:A:C8	41:a:1829:A:C8	3.08	0.42
41:a:2029:G:N1	41:a:2033:A:OP2	2.35	0.42
41:a:2249:U:H3'	41:a:2250:G:C5'	2.49	0.42
41:a:2722:G:C2'	41:a:2723:C:H5'	2.50	0.42
41:a:2813:A:H2'	41:a:2814:A:H8	1.84	0.42
43:c:66:THR:O	43:c:69:ALA:N	2.52	0.42
44:d:82:U:O2	46:f:53:PHE:HZ	2.02	0.42
47:g:57:VAL:O	47:g:58:ASP:C	2.62	0.42
50:j:68:PHE:CE1	50:j:79:LEU:HD21	2.54	0.42
54:n:94:GLU:CA	54:n:97:TRP:HE3	2.31	0.42
59:s:15:TRP:CE3	59:s:135:GLN:HB3	2.55	0.42
60:t:7:MET:C	60:t:18:ARG:NH1	2.77	0.42
62:v:1:MET:SD	62:v:44:ARG:HA	2.59	0.42
62:v:31:PHE:C	62:v:104:GLU:OE1	2.62	0.42
9:9:23:LEU:CA	9:9:118:ILE:HD11	2.50	0.42
12:AB:6:LEU:HB2	12:AB:56:PHE:CE1	2.55	0.42
12:AB:120:GLU:H	12:AB:162:LEU:CG	2.31	0.42
13:AD:22:THR:OG1	13:AD:207:THR:OG1	2.36	0.42
14:AE:1036:ARG:HE	14:AE:1081:VAL:HG21	1.84	0.42
16:AG:244:ARG:N	16:AG:244:ARG:HE	2.18	0.42
16:AG:277:ASP:HB2	16:AG:282:GLN:CB	2.49	0.42
16:AG:277:ASP:HB2	16:AG:282:GLN:CG	2.50	0.42
16:AG:307:ILE:HG13	16:AG:338:VAL:HG23	2.02	0.42
10:B:31:G:H2'	10:B:32:C:C6	2.55	0.42
17:C:13:PHE:CG	17:C:51:TYR:CD1	3.08	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:13:U:C2	18:D:914:A:C8	3.08	0.42
18:D:76:G:C5	18:D:77:A:N7	2.88	0.42
18:D:119:A:C4	18:D:240:G:N7	2.88	0.42
18:D:195:A:H5''	19:E:64:LYS:HZ1	1.84	0.42
18:D:796:C:N3	18:D:797:C:C5	2.87	0.42
18:D:1241:G:H2'	18:D:1242:G:H8	1.85	0.42
19:E:50:ALA:O	19:E:54:MET:HG3	2.20	0.42
22:H:308:GLU:OE1	22:H:308:GLU:N	2.53	0.42
22:H:331:MET:SD	22:H:348:GLN:HG2	2.60	0.42
22:H:332:VAL:O	22:H:333:LEU:HB2	2.17	0.42
26:L:42:TRP:HB2	26:L:59:TYR:O	2.20	0.42
26:L:94:HIS:CG	26:L:95:ALA:N	2.88	0.42
31:Q:79:ILE:O	31:Q:105:PHE:CE1	2.72	0.42
34:T:64:ARG:CZ	34:T:89:ARG:HH12	2.33	0.42
39:Y:82:ALA:CB	39:Y:108:ILE:HD12	2.50	0.42
41:a:136:G:H2'	41:a:137:U:C6	2.54	0.42
41:a:192:C:C5	41:a:193:U:C6	3.07	0.42
41:a:271:G:C2	41:a:272:A:C4	3.08	0.42
41:a:271:G:C2	41:a:272:A:C5	3.07	0.42
41:a:1076:C:H2'	41:a:1077:A:C8	2.54	0.42
41:a:1172:C:C2'	41:a:1173:U:O4'	2.68	0.42
41:a:1874:C:H2'	41:a:1875:G:O4'	2.20	0.42
41:a:2040:G:H2'	41:a:2041:U:O4'	2.20	0.42
41:a:2554:U:H2'	41:a:2555:U:C6	2.55	0.42
48:h:205:LEU:HD12	48:h:210:ALA:CB	2.44	0.42
49:i:17:ARG:CB	49:i:20:ASP:OD1	2.67	0.42
52:l:6:LYS:HE2	52:l:6:LYS:HB2	1.75	0.42
54:n:12:VAL:HG13	54:n:172:ALA:CB	2.49	0.42
54:n:38:MET:HE3	54:n:57:LEU:CG	2.49	0.42
56:p:149:ARG:HH12	56:p:154:PRO:HB3	1.85	0.42
61:u:123:ARG:C	61:u:125:LEU:HD22	2.45	0.42
62:v:135:VAL:HG12	62:v:136:MET:CE	2.50	0.42
63:w:70:THR:O	63:w:71:ARG:HB2	2.20	0.42
65:y:110:ILE:HG13	65:y:110:ILE:O	2.20	0.42
6:5:10:DT:H2''	12:AB:69:ILE:CA	2.50	0.42
6:5:16:DG:N1	11:AA:451:ARG:NH2	2.66	0.42
9:9:33:VAL:H	9:9:36:ASP:CG	2.28	0.42
10:A:9:G:O2'	10:A:45:G:H2'	2.20	0.42
10:A:71:C:C3'	10:A:72:A:C8	2.99	0.42
11:AA:144:VAL:HG23	11:AA:515:MET:HB2	2.02	0.42
11:AA:1313:HIS:CD2	15:AF:31:GLN:HE22	2.37	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AB:120:GLU:H	12:AB:162:LEU:CD1	2.33	0.42
13:AD:31:LEU:HD21	13:AD:39:LEU:HD12	2.00	0.42
14:AE:609:TYR:HD2	14:AE:610:ARG:HD2	1.84	0.42
16:AG:10:GLU:C	16:AG:12:VAL:HG22	2.44	0.42
16:AG:234:ALA:HB3	16:AG:271:ILE:HD13	2.01	0.42
10:B:59:A:C5	10:B:60:U:C5	3.08	0.42
18:D:323:U:H2'	18:D:324:G:O4'	2.19	0.42
18:D:619:U:H1'	24:J:128:ARG:NH1	2.34	0.42
18:D:678:U:C2	18:D:679:C:C5	3.08	0.42
18:D:1073:U:O2	21:G:103:ASN:ND2	2.52	0.42
18:D:1127:G:C6	18:D:1128:C:C5	3.07	0.42
21:G:208:ARG:NH1	21:G:211:THR:CG2	2.82	0.42
22:H:52:GLN:OE1	22:H:52:GLN:HA	2.18	0.42
22:H:135:LEU:CB	22:H:157:ILE:CB	2.97	0.42
23:I:66:VAL:HB	23:I:101:ILE:CD1	2.50	0.42
25:K:111:MET:O	25:K:112:ARG:C	2.63	0.42
29:O:50:GLN:HA	29:O:53:GLU:OE1	2.20	0.42
32:R:3:THR:O	32:R:6:GLN:HB2	2.20	0.42
32:R:18:LYS:NZ	32:R:19:SER:O	2.47	0.42
34:T:70:LEU:CD2	34:T:78:TYR:HA	2.49	0.42
34:T:77:ARG:HA	34:T:77:ARG:NE	2.35	0.42
35:U:8:ARG:HD3	35:U:17:TYR:CE1	2.54	0.42
41:a:340:A:H2'	41:a:341:C:O4'	2.19	0.42
41:a:573:U:O2'	41:a:574:A:H3'	2.19	0.42
41:a:592:A:N3	55:o:4:ILE:HD11	2.35	0.42
41:a:874:G:C6	41:a:904:G:C6	3.08	0.42
41:a:1588:G:N1	41:a:1589:U:O4	2.53	0.42
41:a:1665:A:O3'	60:t:66:LYS:CD	2.67	0.42
41:a:1778:U:C4	41:a:1784:A:C4	3.07	0.42
41:a:1827:U:O2'	41:a:1828:G:H5'	2.19	0.42
41:a:1996:C:OP1	60:t:31:ARG:NE	2.52	0.42
41:a:2097:A:H2'	41:a:2098:U:C6	2.55	0.42
41:a:2420:C:C5'	51:k:8:LYS:HZ1	2.32	0.42
45:e:56:LEU:C	45:e:59:GLU:HG3	2.44	0.42
47:g:47:LYS:CE	54:n:115:ARG:HA	2.49	0.42
48:h:144:VAL:HG21	48:h:162:VAL:HG11	2.01	0.42
48:h:196:GLY:O	48:h:198:ALA:N	2.53	0.42
50:j:37:VAL:HG22	50:j:48:ILE:HD12	2.02	0.42
50:j:175:LEU:HB3	50:j:189:VAL:CG1	2.50	0.42
52:l:46:GLN:O	52:l:88:ARG:NH2	2.53	0.42
54:n:38:MET:HE3	54:n:57:LEU:CD2	2.50	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:s:125:TYR:CD2	59:s:130:HIS:HB3	2.54	0.42
62:v:95:LEU:C	62:v:96:ILE:HD13	2.45	0.42
65:y:37:LYS:CA	65:y:37:LYS:HE3	2.49	0.42
2:1:36:LEU:CD2	2:1:48:LYS:HA	2.48	0.42
2:1:95:ARG:C	2:1:95:ARG:HD2	2.44	0.42
3:2:37:ASP:OD2	3:2:37:ASP:C	2.62	0.42
3:2:57:VAL:HG22	3:2:58:VAL:N	2.35	0.42
9:9:17:GLU:OE2	9:9:53:ARG:NH2	2.53	0.42
9:9:105:LYS:O	9:9:106:PHE:HB3	2.18	0.42
11:AA:855:PRO:HG3	16:AG:106:THR:N	2.34	0.42
12:AB:81:PHE:CD1	14:AE:141:PHE:O	2.72	0.42
12:AB:130:PHE:HB3	12:AB:133:PRO:HD3	2.00	0.42
12:AB:132:GLU:HB3	12:AB:151:LYS:HE3	2.02	0.42
12:AB:136:GLU:HG3	12:AB:160:ARG:HH12	1.84	0.42
12:AB:141:LEU:CA	12:AB:153:SER:N	2.75	0.42
16:AG:233:ARG:CB	16:AG:327:LEU:CD2	2.98	0.42
16:AG:271:ILE:H	16:AG:271:ILE:HG12	1.49	0.42
17:C:34:THR:HB	17:C:38:LYS:N	2.35	0.42
17:C:40:VAL:HG13	22:H:339:ARG:CD	2.49	0.42
18:D:34:C:H2'	18:D:35:G:H8	1.83	0.42
18:D:511:C:O3'	24:J:44:ARG:NH1	2.53	0.42
18:D:552:U:H2'	18:D:553:A:H8	1.85	0.42
18:D:553:A:H2'	18:D:554:A:H8	1.84	0.42
18:D:768:A:C5	18:D:769:G:C8	3.08	0.42
18:D:1152:A:P	30:P:72:ARG:HH22	2.43	0.42
18:D:1174:G:H2'	18:D:1175:G:H5'	2.02	0.42
18:D:1309:G:C8	38:X:98:ARG:NH2	2.88	0.42
18:D:1318:A:OP1	37:W:7:LYS:NZ	2.52	0.42
18:D:1437:A:C4	18:D:1438:G:C8	3.08	0.42
19:E:48:GLN:CD	19:E:49:LYS:N	2.78	0.42
24:J:65:TYR:CE1	24:J:100:ASN:CG	2.98	0.42
24:J:67:VAL:C	24:J:72:PHE:CE2	2.98	0.42
24:J:76:TYR:HE1	24:J:86:THR:CG2	2.33	0.42
24:J:170:TRP:CD1	24:J:171:LEU:HG	2.54	0.42
25:K:77:ASN:O	25:K:80:THR:O	2.37	0.42
26:L:88:MET:SD	26:L:90:MET:HE3	2.60	0.42
29:O:112:GLU:HG3	29:O:115:LYS:NZ	2.35	0.42
31:Q:64:GLN:HB2	31:Q:99:ALA:HB2	2.01	0.42
34:T:63:ARG:HD2	34:T:67:LEU:HD23	2.01	0.42
37:W:44:MET:HE1	37:W:62:VAL:HG22	2.01	0.42
39:Y:9:LYS:CE	39:Y:55:PRO:HB3	2.43	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:Z:20:VAL:O	40:Z:20:VAL:CG1	2.65	0.42
41:a:134:G:H2'	41:a:135:U:H6	1.85	0.42
41:a:151:C:H2'	41:a:152:A:C8	2.53	0.42
41:a:197:A:N6	41:a:2430:A:H2'	2.34	0.42
41:a:353:C:C2	41:a:354:A:C8	3.07	0.42
41:a:667:U:H2'	41:a:668:A:O4'	2.20	0.42
41:a:780:G:O2'	41:a:783:A:N6	2.53	0.42
41:a:1008:A:OP2	59:s:37:ARG:NH1	2.53	0.42
41:a:1065:U:O2'	41:a:1066:U:P	2.77	0.42
41:a:1343:G:C8	41:a:1597:A:C6	3.08	0.42
41:a:1580:A:C8	41:a:1581:G:C8	3.08	0.42
41:a:2585:U:O2	41:a:2585:U:O4'	2.37	0.42
41:a:2586:U:H2'	41:a:2587:A:O4'	2.20	0.42
46:f:27:LEU:O	46:f:38:ARG:NE	2.44	0.42
48:h:18:LYS:HE3	48:h:20:VAL:HG21	2.02	0.42
48:h:108:LYS:CG	48:h:109:GLY:N	2.83	0.42
51:k:9:ILE:HD12	51:k:51:GLU:HG2	2.01	0.42
52:l:149:ILE:HG21	52:l:175:ILE:HD11	2.01	0.42
52:l:170:ARG:HH22	52:l:176:ASP:H	1.68	0.42
56:p:137:ASP:OD2	56:p:140:VAL:N	2.45	0.42
58:r:8:LYS:HD2	58:r:13:GLY:O	2.20	0.42
60:t:113:MET:SD	60:t:113:MET:C	3.03	0.42
64:x:48:LEU:CD2	64:x:87:ILE:CD1	2.98	0.42
66:z:79:PHE:CE1	66:z:110:VAL:HA	2.46	0.42
1:0:74:ILE:HD13	1:0:87:GLN:OE1	2.20	0.42
3:2:26:LYS:CE	3:2:26:LYS:HA	2.50	0.42
4:3:6:ARG:HG2	4:3:7:ARG:N	2.35	0.42
9:9:87:GLU:OE1	9:9:91:ALA:O	2.38	0.42
10:A:5:G:C2'	10:A:6:G:H5'	2.50	0.42
10:A:19:G:N2	41:a:2112:G:C1'	2.82	0.42
10:A:31:G:H2'	10:A:32:C:C6	2.55	0.42
10:A:57:A:O5'	10:A:58:A:P	2.77	0.42
11:AA:741:MET:HE3	11:AA:974:ARG:HH12	1.85	0.42
12:AB:3:SER:HB3	12:AB:5:TYR:CE2	2.55	0.42
14:AE:134:ASP:HB3	14:AE:159:ILE:HG21	2.02	0.42
16:AG:152:LEU:HD11	16:AG:162:ILE:HD12	2.01	0.42
10:B:68:C:C4	10:B:69:C:C5	3.08	0.42
17:C:27:ALA:HB1	17:C:30:LYS:HE3	2.00	0.42
19:E:60:ARG:O	19:E:64:LYS:HD3	2.19	0.42
21:G:101:LEU:HD22	21:G:157:LEU:HD12	1.99	0.42
21:G:106:THR:HA	21:G:109:GLN:HE21	1.85	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:H:277:GLY:C	22:H:331:MET:HE3	2.45	0.42
22:H:322:VAL:HG12	22:H:323:ASN:HB2	2.02	0.42
23:I:172:ARG:NH1	23:I:174:PRO:HG3	2.35	0.42
24:J:41:HIS:ND1	24:J:44:ARG:CZ	2.82	0.42
25:K:123:VAL:O	25:K:124:LEU:HD23	2.20	0.42
28:N:10:MET:CG	28:N:27:MET:HE1	2.48	0.42
29:O:116:VAL:HG11	30:P:62:ARG:HD3	2.02	0.42
32:R:4:VAL:CG1	32:R:5:ASN:N	2.83	0.42
39:Y:60:VAL:CG2	39:Y:66:PHE:CD1	3.03	0.42
39:Y:102:ARG:CZ	39:Y:139:VAL:HG21	2.50	0.42
41:a:104:A:C4	41:a:105:C:C6	3.08	0.42
41:a:136:G:H2'	41:a:137:U:H6	1.85	0.42
41:a:1281:G:H2'	41:a:1282:U:C6	2.55	0.42
41:a:1443:U:H2'	41:a:1444:G:C8	2.54	0.42
41:a:1587:G:H2'	41:a:1588:G:C8	2.54	0.42
41:a:1906:G:C8	41:a:1929:G:C4	3.08	0.42
41:a:2783:U:H2'	41:a:2784:U:H6	1.85	0.42
50:j:34:VAL:HG13	50:j:48:ILE:CG2	2.50	0.42
50:j:152:PRO:HG3	50:j:156:PHE:CE1	2.53	0.42
52:l:77:ILE:O	52:l:77:ILE:HG22	2.19	0.42
52:l:170:ARG:CZ	52:l:174:GLY:O	2.66	0.42
56:p:70:ALA:HA	56:p:73:ASN:HD22	1.84	0.42
56:p:91:GLY:HA2	56:p:160:LYS:CD	2.50	0.42
59:s:125:TYR:HE2	59:s:130:HIS:CB	2.32	0.42
62:v:112:LEU:CA	62:v:115:GLU:OE1	2.68	0.42
63:w:33:ILE:HG13	63:w:112:TYR:HD1	1.83	0.42
64:x:36:TYR:CD2	64:x:37:ALA:N	2.88	0.42
3:2:3:ARG:HG3	3:2:5:GLU:OE1	2.20	0.41
4:3:35:ILE:HD11	4:3:62:GLU:C	2.44	0.41
5:4:77:VAL:HG21	5:4:79:ARG:HE	1.84	0.41
9:9:43:LYS:HZ2	9:9:98:GLU:HB2	1.82	0.41
9:9:95:LEU:O	9:9:98:GLU:HG2	2.20	0.41
9:9:98:GLU:CB	9:9:101:LYS:HZ1	2.33	0.41
14:AE:975:ILE:HD11	14:AE:1003:LEU:HG	2.02	0.41
16:AG:43:ILE:HB	16:AG:44:ASP:H	1.54	0.41
16:AG:105:ILE:HB	16:AG:106:THR:H	1.51	0.41
16:AG:204:MET:SD	16:AG:204:MET:O	2.78	0.41
16:AG:206:ILE:HD11	16:AG:226:ALA:HB2	2.02	0.41
16:AG:434:LEU:HD22	16:AG:459:LEU:HD11	1.80	0.41
10:B:19:G:N3	10:B:57:A:C4	2.87	0.41
17:C:44:ILE:HG21	22:H:339:ARG:C	2.45	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:36:C:N4	18:D:37:U:C4	2.88	0.41
18:D:619:U:O4'	24:J:128:ARG:CZ	2.67	0.41
18:D:684:U:H2'	18:D:685:G:O4'	2.19	0.41
18:D:820:U:H4'	18:D:821:G:OP2	2.20	0.41
18:D:980:C:H2'	18:D:981:U:H5'	2.01	0.41
18:D:1290:G:C4'	27:M:35:LYS:NZ	2.83	0.41
18:D:1381:U:N3	18:D:1382:C:C5	2.88	0.41
19:E:53:GLU:O	19:E:57:ILE:CD1	2.68	0.41
20:F:66:ARG:NH2	20:F:67:ARG:NH2	2.66	0.41
23:I:6:HIS:CD2	33:S:89:MET:HB3	2.55	0.41
23:I:36:ASP:OD1	23:I:57:ILE:HG21	2.19	0.41
25:K:115:LEU:O	25:K:120:VAL:HG12	2.20	0.41
30:P:40:ILE:HD12	30:P:73:LEU:HD22	2.02	0.41
30:P:45:ARG:CD	30:P:47:GLU:OE2	2.67	0.41
32:R:21:VAL:HG23	32:R:21:VAL:O	2.19	0.41
32:R:79:VAL:O	32:R:103:ASP:HB2	2.20	0.41
39:Y:79:LEU:HD11	39:Y:132:ALA:HA	2.02	0.41
40:Z:6:GLN:CD	40:Z:10:ALA:HB2	2.45	0.41
41:a:209:C:N3	41:a:210:C:C5	2.88	0.41
41:a:920:A:H2'	41:a:921:C:H6	1.85	0.41
41:a:1322:A:C4	41:a:1323:C:C6	3.08	0.41
41:a:1399:C:C2	41:a:1400:U:C5	3.08	0.41
41:a:2218:G:C5'	43:c:45:ARG:HH12	2.33	0.41
41:a:2244:U:O2'	41:a:2245:U:H5'	2.20	0.41
41:a:2507:C:C2	41:a:2583:G:C2	3.07	0.41
42:b:40:GLN:NE2	42:b:59:LEU:HD23	2.35	0.41
45:e:14:LEU:HA	45:e:17:GLU:OE1	2.20	0.41
48:h:71:LYS:HB2	48:h:96:TYR:CZ	2.54	0.41
52:l:31:VAL:HG21	52:l:104:ALA:CB	2.50	0.41
52:l:97:ASN:CG	52:l:100:MET:HE2	2.45	0.41
52:l:108:ILE:HD11	52:l:180:LEU:CD1	2.50	0.41
52:l:198:GLU:OE1	52:l:198:GLU:N	2.43	0.41
54:n:8:TYR:O	54:n:13:VAL:HG23	2.20	0.41
55:o:29:LEU:O	55:o:29:LEU:HG	2.19	0.41
58:r:66:ASN:CG	58:r:135:HIS:HB2	2.45	0.41
60:t:110:GLU:OE1	60:t:110:GLU:HA	2.19	0.41
61:u:98:ALA:O	61:u:100:ILE:HG13	2.20	0.41
63:w:33:ILE:CD1	63:w:112:TYR:HD1	2.32	0.41
63:w:98:LEU:CD1	63:w:114:GLU:OE2	2.68	0.41
64:x:15:ARG:C	64:x:19:GLN:NE2	2.78	0.41
65:y:106:LYS:HB3	65:y:109:ARG:HH22	1.84	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:77:PHE:CD1	1:0:77:PHE:C	2.98	0.41
5:4:53:LYS:HB2	5:4:56:PHE:HE2	1.84	0.41
6:5:9:DG:H5'	11:AA:473:ARG:CD	2.45	0.41
6:5:10:DT:C2	11:AA:62:TYR:HE2	2.38	0.41
9:9:129:LEU:HD12	9:9:130:PRO:CD	2.43	0.41
11:AA:469:VAL:HA	11:AA:472:GLU:HG2	2.01	0.41
11:AA:616:ILE:HG13	11:AA:652:TYR:HB2	2.02	0.41
11:AA:836:LEU:HD21	11:AA:921:PRO:HD3	2.02	0.41
12:AB:118:ILE:CG2	12:AB:128:ALA:HB3	2.50	0.41
14:AE:211:GLU:HA	14:AE:214:ARG:HD2	2.01	0.41
14:AE:230:SER:OG	14:AE:231:GLY:N	2.49	0.41
14:AE:513:MET:HE1	14:AE:579:LEU:HD13	2.01	0.41
14:AE:1272:SER:HB2	14:AE:1292:LEU:HD11	2.02	0.41
16:AG:288:MET:HE3	16:AG:288:MET:C	2.45	0.41
17:C:12:ARG:NH1	17:C:13:PHE:CD1	2.89	0.41
18:D:299:G:H2'	18:D:300:A:C8	2.55	0.41
18:D:356:A:C5	18:D:357:G:C8	3.08	0.41
18:D:520:A:C2'	32:R:70:GLU:OE1	2.67	0.41
18:D:652:U:C4'	28:N:56:LYS:HZ1	2.33	0.41
18:D:662:U:H2'	18:D:663:A:C8	2.55	0.41
18:D:951:G:C6	18:D:1231:G:C6	3.08	0.41
18:D:1113:C:C2	18:D:1114:C:C5	3.08	0.41
18:D:1251:A:H2'	18:D:1252:A:O4'	2.20	0.41
22:H:86:ARG:O	22:H:89:ALA:HB3	2.20	0.41
22:H:309:MET:HE1	22:H:318:PRO:HG3	2.02	0.41
23:I:14:ILE:HB	23:I:178:LEU:HD21	2.01	0.41
23:I:23:PHE:CD2	30:P:97:ASP:HB2	2.53	0.41
23:I:147:LYS:CE	23:I:204:LYS:O	2.69	0.41
24:J:67:VAL:CG1	24:J:71:GLN:HB3	2.50	0.41
24:J:139:PRO:N	24:J:182:PHE:CE2	2.88	0.41
25:K:82:GLN:OE1	25:K:147:MET:SD	2.78	0.41
25:K:133:PRO:HA	25:K:136:VAL:HG12	2.02	0.41
27:M:109:ARG:CD	27:M:116:MET:HE1	2.49	0.41
29:O:21:ILE:HD12	29:O:86:ALA:CB	2.50	0.41
29:O:35:LEU:CD2	29:O:45:ARG:HG2	2.50	0.41
29:O:50:GLN:CA	29:O:53:GLU:OE1	2.69	0.41
32:R:57:LEU:HG	32:R:58:THR:N	2.35	0.41
33:S:10:GLU:OE1	33:S:63:ARG:HG3	2.20	0.41
33:S:47:LYS:O	33:S:50:THR:OG1	2.25	0.41
39:Y:23:VAL:HG23	39:Y:27:LEU:HD23	2.01	0.41
39:Y:60:VAL:CG2	39:Y:66:PHE:CB	2.96	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:Y:63:ASP:C	39:Y:65:SER:H	2.28	0.41
41:a:6:A:O3'	59:s:132:HIS:HE1	2.03	0.41
41:a:310:A:C2	41:a:330:A:C4	3.07	0.41
41:a:409:G:H2'	41:a:410:G:C8	2.55	0.41
41:a:1064:C:H2'	41:a:1065:U:C6	2.55	0.41
41:a:1365:A:O5'	43:c:28:ARG:NH2	2.51	0.41
41:a:1509:A:N3	41:a:1510:G:C8	2.87	0.41
41:a:1829:A:C8	41:a:1830:C:C5	3.08	0.41
41:a:2072:C:C2	41:a:2073:C:C6	3.08	0.41
41:a:2226:C:H2'	41:a:2227:A:O4'	2.20	0.41
41:a:2259:U:C5	41:a:2427:C:N4	2.88	0.41
41:a:2759:G:O2'	56:p:35:ARG:NH1	2.53	0.41
48:h:71:LYS:CB	48:h:96:TYR:CZ	3.03	0.41
48:h:91:ILE:HG21	48:h:103:TYR:HB3	2.03	0.41
50:j:47:ALA:O	50:j:48:ILE:HD13	2.20	0.41
50:j:151:THR:O	50:j:151:THR:HG23	2.19	0.41
55:o:33:LEU:O	55:o:41:LYS:NZ	2.45	0.41
59:s:118:MET:HB3	59:s:118:MET:HE2	1.81	0.41
3:2:5:GLU:OE1	3:2:5:GLU:N	2.52	0.41
5:4:2:PHE:CZ	5:4:56:PHE:CZ	3.08	0.41
5:4:48:MET:CE	5:4:85:LYS:HA	2.51	0.41
5:4:78:GLN:OE1	5:4:88:HIS:HB3	2.21	0.41
9:9:3:LEU:CD1	9:9:5:LEU:HD23	2.43	0.41
10:A:1:C:C5	10:A:2:G:N7	2.89	0.41
11:AA:849:GLU:HB2	11:AA:887:VAL:HG23	2.02	0.41
12:AB:152:HIS:HB3	12:AB:154:VAL:HG11	2.00	0.41
14:AE:416:ILE:HD13	14:AE:416:ILE:HA	1.92	0.41
16:AG:185:SER:O	16:AG:185:SER:OG	2.37	0.41
16:AG:266:LEU:CD2	16:AG:266:LEU:N	2.73	0.41
18:D:124:C:H2'	18:D:125:U:C6	2.56	0.41
18:D:603:U:C2	18:D:604:G:C8	3.08	0.41
18:D:958:A:H5'	37:W:55:ARG:HH12	1.86	0.41
18:D:986:U:H2'	18:D:987:G:O4'	2.19	0.41
18:D:1411:C:H2'	18:D:1412:C:H6	1.85	0.41
18:D:1524:C:H2'	18:D:1525:G:C8	2.55	0.41
19:E:9:LYS:CG	19:E:13:GLN:NE2	2.82	0.41
21:G:187:VAL:HG11	21:G:193:PRO:HB3	2.02	0.41
22:H:116:VAL:O	22:H:279:LYS:HG2	2.21	0.41
22:H:267:TRP:CD2	22:H:340:ARG:HG3	2.56	0.41
27:M:31:MET:HE3	27:M:31:MET:HB2	1.76	0.41
29:O:97:GLU:HG2	29:O:98:LEU:N	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:Q:20:VAL:HG12	31:Q:21:ALA:N	2.35	0.41
31:Q:35:THR:HA	31:Q:42:LEU:HG	2.01	0.41
33:S:10:GLU:CD	33:S:63:ARG:HD2	2.45	0.41
34:T:26:GLU:CG	34:T:27:VAL:N	2.83	0.41
35:U:5:ARG:HA	35:U:71:VAL:HG21	2.01	0.41
36:V:7:THR:C	36:V:8:LEU:HD12	2.46	0.41
38:X:3:ARG:CZ	54:n:110:ARG:HD2	2.49	0.41
39:Y:126:ARG:NH1	41:a:1082:U:O3'	2.51	0.41
41:a:30:G:H2'	41:a:31:C:C6	2.56	0.41
41:a:235:U:C2	41:a:236:C:C6	3.08	0.41
41:a:485:C:N3	41:a:486:C:C5	2.89	0.41
41:a:635:C:OP1	61:u:126:ARG:CZ	2.68	0.41
41:a:700:G:H2'	41:a:701:G:O4'	2.20	0.41
41:a:1026:G:C8	41:a:1134:A:C4	3.08	0.41
41:a:1128:G:C6	41:a:2518:A:C6	3.08	0.41
41:a:1248:G:O2'	66:z:3:ARG:HA	2.21	0.41
41:a:1843:C:H2'	41:a:1844:C:H6	1.85	0.41
41:a:2332:C:P	42:b:77:ARG:HH21	2.42	0.41
41:a:2589:A:H2'	41:a:2590:A:H8	1.86	0.41
41:a:2708:G:C2	41:a:2709:G:C8	3.08	0.41
41:a:2784:U:H2'	41:a:2785:C:C6	2.55	0.41
41:a:2838:G:C6	41:a:2839:G:C5	3.08	0.41
43:c:3:ARG:CD	43:c:30:LEU:HD11	2.50	0.41
48:h:165:VAL:HG12	48:h:175:ARG:NH2	2.35	0.41
49:i:22:LEU:CG	49:i:23:THR:N	2.84	0.41
54:n:122:PHE:HZ	54:n:170:LEU:HD12	1.81	0.41
56:p:52:PHE:HZ	56:p:72:LEU:HD22	1.85	0.41
60:t:116:ILE:HG13	60:t:117:SER:N	2.34	0.41
61:u:96:LYS:HG2	61:u:102:GLY:O	2.21	0.41
62:v:45:GLN:HE22	62:v:125:PRO:HD2	1.84	0.41
62:v:105:MET:HG2	62:v:117:PHE:HZ	1.86	0.41
64:x:88:LYS:HB3	64:x:116:GLN:OE1	2.20	0.41
1:0:51:VAL:CG2	66:z:87:SER:HG	2.33	0.41
4:3:86:ARG:CZ	4:3:95:PHE:CD2	3.04	0.41
10:A:18:G:H2'	10:A:57:A:C2	2.55	0.41
10:A:33:U:H5''	27:M:84:THR:CG2	2.50	0.41
11:AA:310:ILE:HG21	11:AA:325:LEU:HB3	2.03	0.41
12:AB:120:GLU:HB3	12:AB:162:LEU:HD12	2.02	0.41
13:AD:76:GLU:HB3	13:AD:80:GLU:HB3	2.01	0.41
14:AE:74:LYS:HE2	14:AE:74:LYS:HB2	1.59	0.41
14:AE:97:VAL:HG12	14:AE:101:ARG:HG3	2.01	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:AE:319:SER:HA	14:AE:320:ASN:HA	1.73	0.41
18:D:21:G:N2	18:D:22:G:C6	2.88	0.41
18:D:122:G:C4	18:D:123:U:C6	3.08	0.41
18:D:124:C:H2'	18:D:125:U:H6	1.84	0.41
18:D:1064:G:O2'	18:D:1190:G:N2	2.53	0.41
18:D:1119:C:C2	18:D:1120:C:C5	3.09	0.41
18:D:1227:A:OP2	38:X:110:LYS:HE3	2.20	0.41
18:D:1439:G:H2'	18:D:1440:U:O4'	2.20	0.41
21:G:115:LYS:C	21:G:115:LYS:HD3	2.45	0.41
22:H:293:PHE:HA	22:H:302:GLY:O	2.20	0.41
23:I:47:LEU:HD22	23:I:76:VAL:HG22	2.03	0.41
24:J:76:TYR:O	24:J:76:TYR:CD1	2.73	0.41
26:L:1:MET:CE	26:L:67:PRO:HD3	2.50	0.41
26:L:2:ARG:C	26:L:65:GLU:OE1	2.62	0.41
27:M:18:PHE:CD2	27:M:59:LEU:HG	2.56	0.41
30:P:73:LEU:HD23	30:P:73:LEU:O	2.21	0.41
30:P:79:PRO:HG2	30:P:102:LEU:HD11	2.02	0.41
31:Q:20:VAL:CG1	31:Q:21:ALA:N	2.82	0.41
32:R:3:THR:O	32:R:4:VAL:C	2.62	0.41
35:U:73:ALA:HA	35:U:76:LYS:HE2	2.02	0.41
41:a:66:C:H42	41:a:74:A:N6	2.19	0.41
41:a:152:A:C6	41:a:153:U:C4	3.09	0.41
41:a:241:A:N1	41:a:255:A:H5''	2.36	0.41
41:a:975:A:H1'	41:a:990:A:C2	2.55	0.41
41:a:1095:A:O2'	41:a:1096:A:O4'	2.36	0.41
41:a:1246:A:H2'	41:a:1247:A:O4'	2.20	0.41
41:a:1398:C:H2'	41:a:1399:C:H6	1.86	0.41
41:a:2049:G:H21	50:j:161:MET:CE	2.33	0.41
41:a:2511:U:C5'	50:j:130:GLN:OE1	2.68	0.41
41:a:2520:C:O2'	41:a:2521:C:H5'	2.20	0.41
41:a:2636:C:O2'	50:j:45:TYR:CZ	2.70	0.41
41:a:2884:U:P	49:i:40:ARG:NH2	2.93	0.41
43:c:67:VAL:CA	43:c:70:GLU:OE1	2.69	0.41
44:d:55:U:H2'	44:d:56:G:C8	2.56	0.41
50:j:11:MET:SD	50:j:12:THR:N	2.94	0.41
50:j:55:LYS:HZ2	50:j:60:VAL:CA	2.33	0.41
52:l:194:LYS:C	52:l:198:GLU:OE1	2.63	0.41
55:o:30:ARG:CZ	61:u:62:PRO:HB3	2.50	0.41
56:p:109:PHE:CD1	56:p:152:ARG:NH1	2.88	0.41
59:s:123:LYS:CD	59:s:132:HIS:HD2	2.33	0.41
62:v:49:ALA:O	62:v:50:ARG:C	2.62	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
62:v:53:MET:SD	62:v:53:MET:N	2.86	0.41
62:v:57:VAL:CG2	62:v:60:GLN:O	2.69	0.41
1:0:4:VAL:HG23	1:0:4:VAL:O	2.20	0.41
2:1:74:ILE:HD12	2:1:74:ILE:N	2.34	0.41
3:2:5:GLU:HA	45:e:22:LEU:HD21	2.02	0.41
3:2:9:LYS:C	3:2:12:ARG:HH22	2.28	0.41
3:2:19:LYS:HE2	3:2:84:TYR:OH	2.21	0.41
6:5:27:DG:N2	7:6:3:DC:O2	2.53	0.41
7:6:25:DT:H2'	11:AA:496:LYS:CG	2.44	0.41
9:9:33:VAL:O	9:9:36:ASP:OD1	2.39	0.41
9:9:41:LEU:CD1	39:Y:117:THR:HG22	2.50	0.41
9:9:47:GLU:C	9:9:49:GLY:H	2.28	0.41
9:9:77:VAL:O	9:9:77:VAL:CG1	2.67	0.41
10:A:5:G:C2'	10:A:6:G:C5'	2.98	0.41
10:A:47:U:O2'	10:A:50:U:OP1	2.37	0.41
11:AA:135:THR:HG22	11:AA:144:VAL:HG22	2.02	0.41
12:AB:104:ILE:O	12:AB:106:ASP:N	2.53	0.41
16:AG:26:ALA:CA	16:AG:117:LYS:HG2	2.50	0.41
16:AG:56:PHE:CD1	16:AG:56:PHE:N	2.88	0.41
16:AG:131:ARG:HG2	16:AG:186:VAL:CG1	2.48	0.41
16:AG:285:ILE:O	16:AG:288:MET:HE2	2.19	0.41
16:AG:363:LEU:HD21	16:AG:407:ARG:C	2.46	0.41
10:B:49:G:O5'	10:B:49:G:H8	2.01	0.41
17:C:70:TYR:CD2	18:D:674:G:H4'	2.55	0.41
17:C:70:TYR:HE2	26:L:49:TYR:CZ	2.38	0.41
18:D:112:G:N1	18:D:330:C:C5	2.88	0.41
18:D:306:A:C4	18:D:307:C:C6	3.08	0.41
18:D:338:A:C5	18:D:339:C:C5	3.08	0.41
18:D:673:A:H2'	18:D:674:G:H8	1.79	0.41
18:D:1319:A:C8	18:D:1323:G:C5	3.08	0.41
20:F:5:LYS:CD	31:Q:109:ASN:OD1	2.68	0.41
21:G:100:MET:CE	21:G:101:LEU:HD21	2.49	0.41
21:G:185:ALA:O	21:G:200:ILE:N	2.54	0.41
22:H:33:LYS:HA	22:H:34:ASP:HA	1.80	0.41
22:H:303:LEU:O	22:H:305:HIS:HA	2.21	0.41
27:M:74:GLU:HG3	27:M:91:VAL:CG1	2.50	0.41
30:P:6:ILE:CG2	30:P:100:ILE:HG13	2.51	0.41
35:U:26:ASN:ND2	35:U:31:ARG:CD	2.83	0.41
37:W:34:TRP:O	37:W:36:ARG:N	2.54	0.41
39:Y:34:ILE:O	39:Y:34:ILE:HG22	2.20	0.41
40:Z:29:LYS:C	40:Z:30:PHE:HD2	2.29	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:16:C:O4'	49:i:15:MET:HE1	2.19	0.41
41:a:126:A:O5'	53:m:19:ARG:HG3	2.21	0.41
41:a:458:G:N2	41:a:469:G:H2'	2.36	0.41
41:a:749:A:H2'	41:a:749:A:N3	2.35	0.41
41:a:826:U:OP1	41:a:2428:G:H3'	2.20	0.41
41:a:1009:A:N3	41:a:1153:C:O2'	2.48	0.41
41:a:1139:G:O2'	41:a:1143:A:N1	2.49	0.41
41:a:1746:A:H2'	41:a:1747:U:C6	2.56	0.41
41:a:1953:A:H2	41:a:2549:G:N3	2.18	0.41
41:a:2310:C:O2'	54:n:74:VAL:HG12	2.20	0.41
41:a:2590:A:C2	41:a:2591:C:C5	3.08	0.41
44:d:47:C:C4	44:d:48:U:C5	3.09	0.41
45:e:13:GLU:O	45:e:14:LEU:C	2.63	0.41
45:e:20:ASN:O	45:e:24:GLU:HG2	2.20	0.41
48:h:35:GLU:CD	48:h:64:ILE:HD11	2.46	0.41
49:i:34:SER:OG	49:i:36:GLU:HG2	2.20	0.41
54:n:138:PHE:O	54:n:141:ILE:HG22	2.20	0.41
56:p:149:ARG:CZ	56:p:154:PRO:CD	2.98	0.41
60:t:43:ILE:HD12	60:t:56:ASP:HB2	2.01	0.41
63:w:18:GLN:HE22	63:w:22:ARG:NE	2.18	0.41
63:w:38:LEU:HB3	63:w:39:PRO:HD3	2.02	0.41
2:1:23:LEU:CD2	2:1:39:THR:HG21	2.50	0.41
5:4:18:ARG:NH1	44:d:93:C:OP2	2.53	0.41
9:9:15:VAL:HG22	9:9:66:GLY:HA3	2.00	0.41
9:9:117:LEU:O	9:9:117:LEU:CD2	2.67	0.41
11:AA:1069:ARG:NH2	11:AA:1114:GLU:OE2	2.44	0.41
12:AB:31:PRO:HG2	12:AB:51:PHE:CD2	2.55	0.41
12:AB:72:THR:HG22	12:AB:73:ARG:N	2.36	0.41
12:AB:149:GLU:H	12:AB:149:GLU:HG2	1.63	0.41
13:AC:104:LYS:O	13:AC:139:SER:OG	2.36	0.41
13:AD:205:MET:HE1	13:AD:217:ILE:HG13	2.03	0.41
14:AE:138:VAL:HG21	14:AE:145:VAL:HB	2.03	0.41
14:AE:247:PRO:HA	14:AE:250:ARG:HE	1.86	0.41
14:AE:367:GLY:HA3	14:AE:448:GLN:HB2	2.03	0.41
16:AG:207:GLU:HG3	16:AG:210:ARG:CB	2.50	0.41
18:D:429:U:H3	18:D:431:A:N6	2.18	0.41
18:D:517:G:O2'	18:D:530:G:H4'	2.21	0.41
18:D:1140:C:O2'	18:D:1141:C:O5'	2.39	0.41
18:D:1189:U:H4'	23:I:10:ILE:CD1	2.50	0.41
18:D:1411:C:H2'	18:D:1412:C:C6	2.56	0.41
19:E:43:ASP:CG	19:E:46:ALA:HB3	2.46	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:G:20:THR:O	21:G:23:TRP:CD1	2.74	0.41
21:G:134:ALA:HA	21:G:137:ARG:NE	2.35	0.41
22:H:54:LYS:HD3	22:H:58:GLY:HA2	2.03	0.41
23:I:72:ARG:CD	23:I:75:ILE:HB	2.51	0.41
24:J:153:SER:OG	24:J:156:LYS:NZ	2.53	0.41
28:N:83:LEU:HD11	32:R:4:VAL:CB	2.51	0.41
28:N:96:MET:HE2	28:N:99:LEU:HD12	2.03	0.41
31:Q:20:VAL:HG13	31:Q:85:MET:CE	2.51	0.41
33:S:86:GLU:CD	33:S:90:ARG:HD3	2.45	0.41
34:T:32:LEU:HD13	34:T:62:GLN:OE1	2.21	0.41
41:a:319:G:H2'	41:a:320:A:O4'	2.21	0.41
41:a:443:A:N1	52:l:40:ARG:NH2	2.69	0.41
41:a:1427:A:H4'	41:a:1428:C:O4'	2.21	0.41
41:a:1589:U:O2'	41:a:1590:A:O5'	2.29	0.41
41:a:1958:C:O2'	41:a:1959:G:H5'	2.20	0.41
41:a:2024:G:C5	41:a:2040:G:C2	3.09	0.41
41:a:2526:G:O2'	57:q:1:MET:HG2	2.21	0.41
41:a:2569:G:C2	41:a:2570:G:C8	3.09	0.41
41:a:2756:U:C4	41:a:2759:G:O6	2.73	0.41
48:h:163:GLN:HB3	48:h:175:ARG:HB3	2.02	0.41
50:j:68:PHE:CZ	50:j:79:LEU:HD21	2.56	0.41
50:j:175:LEU:HB3	50:j:189:VAL:HG12	2.02	0.41
51:k:19:HIS:ND1	51:k:41:PRO:HD3	2.36	0.41
54:n:66:LEU:HD23	54:n:67:ILE:C	2.45	0.41
54:n:141:ILE:HD11	54:n:146:VAL:CG1	2.50	0.41
56:p:69:ARG:HH12	56:p:73:ASN:CG	2.25	0.41
60:t:24:VAL:O	60:t:24:VAL:HG13	2.20	0.41
62:v:112:LEU:HD12	62:v:112:LEU:H	1.86	0.41
63:w:24:MET:CG	63:w:44:LEU:HD12	2.51	0.41
8:7:1:A:H61	10:B:37:A:N6	2.18	0.41
10:A:68:C:C4	10:A:69:C:C5	3.08	0.41
11:AA:524:ILE:HD12	11:AA:712:SER:HB2	2.02	0.41
11:AA:1105:SER:HB2	14:AE:731:ARG:HB3	2.03	0.41
12:AB:90:SER:HB2	12:AB:94:HIS:NE2	2.35	0.41
12:AB:155:LYS:HB3	12:AB:158:GLU:CD	2.45	0.41
13:AC:42:ALA:HB1	13:AC:224:LEU:HD11	2.03	0.41
14:AE:66:LYS:HB2	14:AE:69:GLU:HB2	2.01	0.41
16:AG:284:VAL:HG22	16:AG:305:MET:CE	2.51	0.41
16:AG:425:LEU:HD23	16:AG:429:LYS:HZ3	1.81	0.41
16:AG:433:ASP:O	16:AG:456:LEU:CD2	2.69	0.41
16:AG:434:LEU:CD1	16:AG:455:THR:CA	2.98	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:B:9:G:O2'	10:B:45:G:H2'	2.20	0.41
18:D:124:C:C2	18:D:125:U:C5	3.08	0.41
18:D:437:U:H5''	24:J:152:GLN:CD	2.46	0.41
18:D:1202:U:C2	33:S:82:ILE:HD13	2.55	0.41
18:D:1368:A:OP1	29:O:116:VAL:HG23	2.20	0.41
20:F:4:ILE:HD12	20:F:19:PHE:HA	2.03	0.41
21:G:137:ARG:HG3	21:G:138:THR:N	2.36	0.41
22:H:287:LEU:HD23	22:H:318:PRO:HB2	2.03	0.41
22:H:308:GLU:HB2	22:H:347:LYS:HB3	2.02	0.41
23:I:155:GLY:HA2	23:I:163:ALA:HB1	2.02	0.41
23:I:156:ARG:HD3	23:I:160:ALA:O	2.21	0.41
25:K:157:ARG:HG2	25:K:159:LYS:HD3	2.01	0.41
26:L:36:ILE:HD11	26:L:62:MET:HE2	2.03	0.41
27:M:16:PRO:HB3	29:O:46:MET:CG	2.47	0.41
29:O:56:ASP:OD1	29:O:56:ASP:N	2.50	0.41
31:Q:36:ASP:OD1	31:Q:40:ASN:N	2.53	0.41
35:U:4:ILE:HG22	35:U:71:VAL:HG11	2.03	0.41
38:X:90:ARG:HD3	38:X:97:VAL:HA	2.01	0.41
39:Y:37:PHE:HZ	39:Y:56:VAL:HG11	1.79	0.41
39:Y:100:ILE:HG13	39:Y:139:VAL:CB	2.51	0.41
41:a:184:C:C2	41:a:213:A:N1	2.89	0.41
41:a:397:U:H2'	41:a:398:C:C6	2.56	0.41
41:a:1430:G:H2'	41:a:1431:A:C8	2.56	0.41
41:a:1494:A:H2'	41:a:1495:A:C8	2.56	0.41
41:a:2357:G:OP1	42:b:20:ARG:CZ	2.68	0.41
41:a:2571:U:O2'	50:j:151:THR:HG23	2.20	0.41
48:h:84:ASP:OD2	48:h:91:ILE:HD11	2.21	0.41
48:h:220:VAL:CG2	48:h:225:MET:HE3	2.51	0.41
50:j:170:VAL:HG13	50:j:194:PRO:CG	2.50	0.41
54:n:170:LEU:HB2	54:n:177:PHE:HZ	1.86	0.41
58:r:110:VAL:HG21	58:r:114:GLU:OE1	2.21	0.41
60:t:20:MET:CB	60:t:44:LYS:HE2	2.51	0.41
60:t:61:VAL:HG11	60:t:112:PHE:CE2	2.56	0.41
63:w:18:GLN:CD	63:w:18:GLN:C	2.88	0.41
63:w:28:LEU:HD23	63:w:48:VAL:HG21	2.03	0.41
63:w:49:GLU:OE2	63:w:95:THR:CG2	2.69	0.41
63:w:87:PHE:O	63:w:88:ALA:C	2.64	0.41
66:z:66:ASN:OD1	66:z:70:ARG:NE	2.54	0.41
2:1:25:ARG:NH2	41:a:520:G:P	2.87	0.41
2:1:85:ILE:O	2:1:85:ILE:HG13	2.19	0.41
9:9:96:PHE:HB3	9:9:125:ARG:HH11	1.86	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:9:129:LEU:N	9:9:130:PRO:CD	2.81	0.41
10:A:19:G:O6	41:a:2112:G:P	2.79	0.41
10:A:25:C:C4	10:A:26:G:C8	3.09	0.41
11:AA:998:LEU:HD21	11:AA:1015:ALA:HB2	2.02	0.41
11:AA:1275:VAL:HG13	11:AA:1287:LEU:HD11	2.01	0.41
13:AD:286:GLU:N	13:AD:315:GLY:HA2	2.18	0.41
14:AE:836:ARG:HG3	14:AE:869:CYS:HB3	2.02	0.41
14:AE:1003:LEU:HD23	14:AE:1018:ALA:HB2	2.02	0.41
16:AG:125:MET:SD	16:AG:125:MET:O	2.79	0.41
16:AG:323:GLN:HE21	16:AG:323:GLN:C	2.28	0.41
16:AG:448:LEU:CD2	16:AG:467:LEU:HD22	2.51	0.41
10:B:1:C:C5	10:B:2:G:N7	2.89	0.41
10:B:18:G:H2'	10:B:57:A:C2	2.55	0.41
17:C:12:ARG:CB	22:H:264:GLU:CB	2.98	0.41
18:D:104:G:C2	18:D:105:G:C5	3.09	0.41
18:D:376:G:O3'	35:U:5:ARG:NE	2.54	0.41
18:D:960:U:O2'	18:D:1223:C:H4'	2.20	0.41
18:D:963:G:C2	18:D:964:A:C8	3.09	0.41
18:D:985:C:C2	18:D:986:U:C5	3.08	0.41
18:D:997:U:O2'	18:D:998:C:H5'	2.21	0.41
18:D:1158:C:O2	18:D:1158:C:C2'	2.69	0.41
18:D:1512:U:H2'	18:D:1513:A:H8	1.85	0.41
22:H:309:MET:HE1	22:H:318:PRO:CA	2.51	0.41
23:I:6:HIS:CD2	23:I:9:GLY:H	2.39	0.41
23:I:35:SER:HA	23:I:38:LYS:CE	2.51	0.41
26:L:3:HIS:CD2	26:L:95:ALA:HA	2.56	0.41
29:O:83:ILE:O	29:O:87:LEU:CD2	2.68	0.41
32:R:25:GLU:CD	32:R:30:LYS:HE3	2.46	0.41
35:U:67:ILE:O	35:U:67:ILE:HG13	2.20	0.41
37:W:66:MET:SD	37:W:66:MET:C	3.03	0.41
38:X:16:VAL:HG12	38:X:34:LEU:CD1	2.50	0.41
40:Z:6:GLN:HG3	40:Z:6:GLN:O	2.21	0.41
41:a:62:U:O2	41:a:62:U:H2'	2.20	0.41
41:a:138:U:C5	41:a:142:A:C6	3.08	0.41
41:a:532:A:N7	41:a:2021:C:O2'	2.43	0.41
41:a:638:G:H2'	41:a:639:U:C6	2.56	0.41
41:a:938:G:C2	41:a:939:G:C8	3.08	0.41
41:a:1048:A:C5	41:a:1049:C:C5	3.09	0.41
41:a:1824:G:O5'	48:h:52:ARG:NH1	2.53	0.41
41:a:2040:G:OP1	59:s:106:LYS:NZ	2.38	0.41
41:a:2377:A:O2'	64:x:117:PHE:O	2.36	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2461:A:H2'	41:a:2462:C:C6	2.55	0.41
41:a:2686:G:H2'	41:a:2687:U:C6	2.56	0.41
42:b:30:SER:HA	42:b:66:LYS:HD2	2.03	0.41
45:e:44:LYS:HZ3	45:e:48:ARG:HE	1.69	0.41
51:k:26:ASN:OD1	51:k:28:ARG:CZ	2.69	0.41
60:t:80:ASP:OD1	65:y:68:GLU:OE2	2.39	0.41
63:w:45:ARG:O	63:w:49:GLU:CD	2.62	0.41
63:w:60:VAL:CA	63:w:63:ARG:NH1	2.84	0.41
63:w:73:ASN:ND2	63:w:73:ASN:O	2.50	0.41
64:x:51:ALA:HB3	64:x:78:VAL:HB	2.02	0.41
64:x:83:LEU:HD23	64:x:88:LYS:HD3	2.02	0.41
66:z:91:ASP:O	66:z:95:LEU:HG	2.20	0.41
3:2:6:ARG:NH2	3:2:10:VAL:CG2	2.84	0.41
3:2:22:THR:O	3:2:25:GLU:HG2	2.20	0.41
4:3:5:ILE:HA	4:3:72:ILE:HD11	2.03	0.41
8:7:10:U:O2	8:7:10:U:C3'	2.69	0.41
8:7:44:U:C3'	8:7:44:U:C6	3.04	0.41
8:7:47:U:H4'	14:AE:322:ARG:NH1	2.35	0.41
9:9:46:ARG:O	9:9:48:ALA:N	2.53	0.41
9:9:59:LEU:HD22	9:9:62:ARG:CD	2.51	0.41
9:9:94:ARG:O	9:9:98:GLU:HG2	2.21	0.41
12:AB:36:GLU:O	12:AB:37:LYS:HD2	2.21	0.41
12:AB:154:VAL:CG2	12:AB:158:GLU:HB2	2.51	0.41
13:AD:305:ASP:C	13:AD:307:LEU:H	2.28	0.41
14:AE:450:HIS:HA	14:AE:451:PRO:HD3	1.92	0.41
16:AG:37:LYS:HA	16:AG:37:LYS:HD3	1.67	0.41
16:AG:148:ASP:HA	16:AG:164:ARG:HD2	2.02	0.41
16:AG:437:LEU:CD2	16:AG:488:ILE:CD1	2.98	0.41
10:B:25:C:C4	10:B:26:G:C8	3.09	0.41
10:B:47:U:O2'	10:B:50:U:OP1	2.37	0.41
18:D:71:A:C6	18:D:100:G:N7	2.89	0.41
18:D:107:G:C6	18:D:108:G:C4	3.09	0.41
18:D:135:C:H2'	35:U:1:MET:CE	2.51	0.41
18:D:264:C:H2'	18:D:265:G:O4'	2.21	0.41
18:D:407:U:C2	18:D:408:A:C8	3.08	0.41
18:D:407:U:O4'	24:J:116:GLN:OE1	2.39	0.41
18:D:526:C:P	32:R:88:LYS:HZ3	2.43	0.41
18:D:687:A:C8	18:D:701:U:C4	3.08	0.41
18:D:695:A:OP1	31:Q:54:GLY:HA3	2.21	0.41
18:D:695:A:O5'	31:Q:53:ARG:CZ	2.69	0.41
18:D:718:A:C2	31:Q:118:HIS:CE1	3.09	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:920:U:H2'	18:D:921:U:C6	2.56	0.41
18:D:1186:G:H4'	29:O:112:GLU:OE2	2.21	0.41
18:D:1227:A:C8	38:X:116:ILE:HD11	2.56	0.41
18:D:1288:A:H2'	18:D:1289:A:O4'	2.21	0.41
20:F:66:ARG:NH2	20:F:67:ARG:HH22	2.19	0.41
21:G:47:VAL:N	21:G:48:PRO:HD2	2.35	0.41
21:G:56:GLU:HG2	21:G:198:PHE:CZ	2.55	0.41
21:G:117:LEU:CD2	21:G:141:LEU:HA	2.51	0.41
21:G:142:GLU:OE2	21:G:142:GLU:HA	2.21	0.41
22:H:70:VAL:O	22:H:71:ALA:HB2	2.20	0.41
22:H:297:GLU:HG2	22:H:300:VAL:CG2	2.51	0.41
23:I:8:ASN:O	23:I:12:LEU:HD23	2.21	0.41
23:I:8:ASN:O	23:I:12:LEU:CD2	2.69	0.41
23:I:135:LYS:HG3	23:I:136:ARG:N	2.35	0.41
23:I:147:LYS:CE	23:I:204:LYS:C	2.94	0.41
24:J:188:ARG:NH2	24:J:195:ILE:HG13	2.36	0.41
25:K:45:ARG:CG	25:K:45:ARG:NE	2.72	0.41
25:K:86:LYS:HE3	25:K:95:PHE:HD1	1.86	0.41
25:K:88:VAL:HG12	25:K:93:ARG:HG2	2.02	0.41
26:L:42:TRP:HB3	26:L:59:TYR:HB2	2.03	0.41
27:M:37:SER:HA	29:O:41:ARG:HH22	1.85	0.41
31:Q:20:VAL:CG1	31:Q:85:MET:CE	2.99	0.41
31:Q:49:GLY:C	31:Q:51:GLY:H	2.29	0.41
31:Q:94:GLU:O	31:Q:94:GLU:OE2	2.39	0.41
37:W:40:ILE:CG2	37:W:44:MET:SD	3.04	0.41
38:X:13:LYS:CE	38:X:17:ILE:HG21	2.51	0.41
38:X:55:THR:CG2	38:X:59:GLU:OE2	2.69	0.41
38:X:71:ARG:NH1	38:X:72:GLU:CG	2.84	0.41
41:a:272:A:C2	41:a:273:G:C8	3.09	0.41
41:a:630:G:N2	41:a:633:A:OP2	2.45	0.41
41:a:690:G:H2'	41:a:691:C:H6	1.85	0.41
41:a:958:U:C4	62:v:40:ARG:NH1	2.88	0.41
41:a:985:C:O2	41:a:986:C:C6	2.73	0.41
41:a:1252:G:H1	66:z:37:GLN:HE21	1.68	0.41
41:a:1355:G:C2	41:a:1356:G:C8	3.09	0.41
41:a:1366:A:H2'	41:a:1367:A:O4'	2.21	0.41
41:a:1589:U:O2'	41:a:1590:A:P	2.79	0.41
41:a:1792:G:C5'	48:h:204:VAL:CG2	2.99	0.41
41:a:1820:U:C2	48:h:201:MET:CE	3.03	0.41
41:a:1824:G:H5''	48:h:52:ARG:NH2	2.36	0.41
41:a:2205:A:H2'	41:a:2206:C:C6	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2397:G:H2'	41:a:2398:U:H6	1.86	0.41
41:a:2454:G:C5	41:a:2455:G:C8	3.09	0.41
41:a:2511:U:O4'	50:j:130:GLN:OE1	2.39	0.41
41:a:2649:C:C2	41:a:2650:U:C6	3.08	0.41
41:a:2673:G:C2	41:a:2674:G:C8	3.08	0.41
41:a:2808:G:N1	41:a:2891:U:C5	2.89	0.41
42:b:17:GLU:OE1	42:b:19:LYS:NZ	2.54	0.41
42:b:55:ARG:CZ	42:b:55:ARG:CA	2.99	0.41
42:b:68:LYS:HZ2	42:b:70:GLU:CD	2.23	0.41
44:d:8:C:O2'	64:x:40:ILE:HD13	2.21	0.41
44:d:57:A:N7	54:n:26:MET:HE3	2.36	0.41
48:h:189:ARG:HG3	48:h:190:ALA:N	2.36	0.41
48:h:210:ALA:HA	48:h:213:TRP:CE3	2.56	0.41
50:j:108:ASP:OD1	50:j:173:GLN:HA	2.21	0.41
51:k:8:LYS:HA	51:k:24:THR:HA	2.02	0.41
52:l:23:PHE:CD1	52:l:111:GLU:HG3	2.55	0.41
52:l:171:ASP:OD1	52:l:172:ALA:N	2.53	0.41
54:n:67:ILE:CD1	54:n:87:CYS:HB3	2.51	0.41
55:o:22:PHE:CE1	55:o:62:LEU:HD23	2.55	0.41
55:o:45:ARG:HB3	55:o:46:PRO:HD3	2.02	0.41
56:p:80:THR:HG23	56:p:81:GLU:N	2.36	0.41
62:v:31:PHE:CZ	62:v:110:GLU:HG3	2.56	0.41
65:y:6:LYS:HD2	65:y:6:LYS:C	2.46	0.41
65:y:48:ILE:HD11	65:y:101:ARG:NH1	2.36	0.41
1:0:13:ARG:C	1:0:14:VAL:HG13	2.46	0.41
11:AA:856:ASN:CG	16:AG:35:THR:HB	2.43	0.41
12:AB:4:TRP:CZ3	12:AB:93:ILE:HG13	2.56	0.41
12:AB:32:MET:O	12:AB:50:LEU:HD12	2.21	0.41
12:AB:60:ASP:HA	12:AB:61:PRO:HD3	1.88	0.41
12:AB:72:THR:HG22	12:AB:73:ARG:H	1.86	0.41
12:AB:141:LEU:HD22	12:AB:143:LEU:HD11	2.01	0.41
14:AE:591:ILE:HG23	14:AE:592:VAL:HG13	2.03	0.41
14:AE:800:LEU:HB3	14:AE:920:ALA:HB1	2.02	0.41
16:AG:105:ILE:HD13	16:AG:105:ILE:H	1.85	0.41
16:AG:296:ILE:HD11	16:AG:307:ILE:CD1	2.50	0.41
16:AG:374:VAL:O	16:AG:374:VAL:CG1	2.68	0.41
18:D:18:C:C2	18:D:19:A:C8	3.09	0.41
18:D:567:G:H2'	18:D:568:G:O4'	2.21	0.41
18:D:570:G:H1'	18:D:820:U:C4	2.56	0.41
18:D:1000:A:N1	18:D:1041:G:C6	2.89	0.41
21:G:6:MET:HA	21:G:9:MET:SD	2.61	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:G:20:THR:HA	21:G:39:HIS:HE1	1.82	0.41
21:G:36:ASN:ND2	22:H:19:ARG:HD2	2.36	0.41
21:G:85:LEU:HD23	21:G:85:LEU:C	2.46	0.41
21:G:138:THR:O	21:G:142:GLU:HG2	2.22	0.41
28:N:8:ALA:N	28:N:77:ARG:NH1	2.69	0.41
29:O:25:ASN:HB2	29:O:27:LYS:NZ	2.36	0.41
29:O:56:ASP:O	29:O:60:LYS:CE	2.69	0.41
30:P:26:VAL:HG12	30:P:30:LYS:HZ3	1.86	0.41
32:R:24:LEU:CD2	32:R:95:TYR:CE1	3.04	0.41
38:X:3:ARG:NH2	47:g:35:ASP:HB2	2.36	0.41
39:Y:19:PRO:CG	39:Y:23:VAL:HG23	2.49	0.41
40:Z:19:VAL:O	40:Z:23:ILE:HG12	2.21	0.41
41:a:18:U:H2'	41:a:19:A:H8	1.86	0.41
41:a:1443:U:C2	41:a:1444:G:C8	3.09	0.41
41:a:1597:A:H5''	41:a:1598:A:H5'	2.01	0.41
41:a:2483:C:H1'	62:v:51:ARG:CZ	2.51	0.41
41:a:2531:A:C4	41:a:2532:G:C8	3.08	0.41
42:b:70:GLU:HG3	42:b:72:LYS:HG2	2.03	0.41
48:h:53:HIS:NE2	48:h:219:THR:HG23	2.35	0.41
48:h:74:ILE:HG22	48:h:96:TYR:HD1	1.86	0.41
48:h:75:PRO:O	48:h:97:LYS:HD3	2.21	0.41
51:k:26:ASN:OD1	51:k:28:ARG:NH1	2.54	0.41
59:s:1:MET:SD	59:s:2:LYS:O	2.79	0.41
59:s:49:ASP:OD1	59:s:121:LYS:NZ	2.54	0.41
60:t:58:LEU:N	60:t:58:LEU:HD23	2.36	0.41
62:v:104:GLU:CD	62:v:105:MET:H	2.28	0.41
62:v:115:GLU:HA	62:v:118:LYS:HE2	2.02	0.41
3:2:26:LYS:HA	3:2:26:LYS:HE3	2.02	0.40
6:5:23:DA:H2''	6:5:24:DC:C6	2.56	0.40
6:5:25:DA:N3	7:6:5:DG:N2	2.68	0.40
8:7:10:U:O2	8:7:10:U:H3'	2.21	0.40
8:7:13:U:C6	8:7:13:U:C3'	3.03	0.40
9:9:2:ALA:HB3	9:9:6:GLN:HB2	2.01	0.40
9:9:43:LYS:HA	9:9:46:ARG:NH2	2.36	0.40
9:9:123:ILE:O	9:9:123:ILE:CG2	2.68	0.40
11:AA:27:LEU:O	11:AA:528:ARG:NH1	2.43	0.40
11:AA:936:ARG:HB2	11:AA:1042:LEU:HD12	2.03	0.40
12:AB:89:PRO:HD2	14:AE:290:ILE:CG2	2.46	0.40
12:AB:142:LEU:CA	12:AB:151:LYS:HA	2.50	0.40
16:AG:27:LEU:CD2	16:AG:114:ILE:HG23	2.51	0.40
16:AG:108:GLN:HA	16:AG:111:LYS:HD2	2.03	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:244:ARG:N	16:AG:244:ARG:NE	2.69	0.40
16:AG:291:ALA:HA	16:AG:316:GLN:HE22	1.87	0.40
16:AG:434:LEU:HG	16:AG:455:THR:CA	2.50	0.40
18:D:19:A:C6	18:D:20:U:C4	3.10	0.40
18:D:597:G:N7	18:D:598:U:C5	2.89	0.40
18:D:643:C:C2	18:D:644:U:C5	3.08	0.40
18:D:751:U:H2'	18:D:752:G:O4'	2.22	0.40
21:G:30:PHE:HA	21:G:45:LYS:NZ	2.35	0.40
22:H:308:GLU:HG3	22:H:346:LEU:H	1.86	0.40
23:I:123:GLN:HB3	23:I:128:VAL:HG21	2.02	0.40
26:L:9:MET:CE	26:L:86:ARG:HG2	2.52	0.40
27:M:47:LEU:HD21	27:M:58:GLU:HB3	2.03	0.40
34:T:18:ASP:O	34:T:19:ALA:HB3	2.20	0.40
34:T:70:LEU:HD23	34:T:78:TYR:HD1	1.85	0.40
36:V:11:ARG:HD3	36:V:57:ASP:C	2.46	0.40
36:V:31:HIS:HD2	36:V:34:TYR:HB2	1.85	0.40
37:W:17:LYS:HB3	37:W:21:LYS:NZ	2.36	0.40
41:a:10:A:C6	41:a:2800:A:C2	3.10	0.40
41:a:67:U:C4	41:a:74:A:N6	2.89	0.40
41:a:209:C:C2	41:a:210:C:C6	3.09	0.40
41:a:465:G:H2'	41:a:466:A:C8	2.56	0.40
41:a:529:A:C4	41:a:2023:C:C5	3.09	0.40
41:a:654:A:H61	55:o:19:LYS:CD	2.31	0.40
41:a:789:A:N1	53:m:3:ARG:NH2	2.68	0.40
41:a:841:G:H2'	41:a:842:U:H6	1.86	0.40
41:a:888:C:C4	41:a:889:C:N4	2.88	0.40
41:a:967:U:H2'	41:a:968:C:C6	2.56	0.40
41:a:1069:A:C2	41:a:1073:A:C8	3.08	0.40
41:a:1177:G:O2'	41:a:1178:C:O4'	2.39	0.40
41:a:1239:G:H2'	41:a:1240:U:O4'	2.21	0.40
41:a:1351:C:H2'	41:a:1352:U:O4'	2.21	0.40
41:a:1558:C:O4'	41:a:1560:G:C8	2.74	0.40
41:a:1571:A:H2'	41:a:1572:A:H8	1.86	0.40
41:a:1676:A:H2'	41:a:1677:A:C8	2.56	0.40
41:a:1707:G:H2'	41:a:1708:C:H6	1.87	0.40
41:a:1759:A:O2'	41:a:2714:G:O2'	2.38	0.40
41:a:1778:U:H2'	41:a:1784:A:N6	2.36	0.40
41:a:1822:C:O2'	41:a:1823:G:H5'	2.20	0.40
41:a:1827:U:H2'	41:a:1828:G:O4'	2.20	0.40
41:a:2262:U:OP1	41:a:2387:U:O2'	2.33	0.40
41:a:2393:U:H2'	41:a:2394:C:C6	2.56	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2420:C:H5''	51:k:8:LYS:HZ1	1.85	0.40
41:a:2821:A:H4'	50:j:167:ASN:ND2	2.36	0.40
42:b:17:GLU:OE1	42:b:19:LYS:HD3	2.21	0.40
48:h:245:VAL:HA	48:h:250:VAL:O	2.20	0.40
50:j:55:LYS:HD3	50:j:56:LYS:N	2.36	0.40
50:j:104:VAL:CG1	50:j:106:LYS:O	2.69	0.40
53:m:12:ARG:HG3	53:m:44:VAL:HG21	2.03	0.40
53:m:26:ASN:HA	53:m:29:GLN:NE2	2.36	0.40
54:n:61:SER:O	54:n:63:GLN:N	2.54	0.40
56:p:17:VAL:O	56:p:18:LYS:CE	2.65	0.40
63:w:98:LEU:HD12	63:w:114:GLU:OE2	2.21	0.40
65:y:31:TRP:CZ3	65:y:38:LYS:CD	3.03	0.40
1:0:13:ARG:NH2	66:z:97:ASP:CG	2.79	0.40
2:1:10:ALA:CB	2:1:103:ILE:CD1	2.99	0.40
2:1:93:ALA:HB2	41:a:1614:A:C2	2.56	0.40
4:3:84:GLY:C	4:3:95:PHE:CE2	2.99	0.40
11:AA:22:LEU:HD22	11:AA:603:ILE:HG21	2.02	0.40
11:AA:176:ILE:HD12	11:AA:184:LEU:HD23	2.04	0.40
11:AA:803:ALA:HB2	11:AA:1094:VAL:HG21	2.03	0.40
12:AB:82:GLY:HA2	14:AE:142:GLU:CG	2.49	0.40
13:AC:48:LEU:HA	13:AC:48:LEU:HD23	1.86	0.40
13:AD:48:LEU:HG	14:AE:535:ARG:HG3	2.02	0.40
14:AE:1175:LEU:HD22	14:AE:1190:ILE:HD11	2.03	0.40
16:AG:433:ASP:C	16:AG:456:LEU:HD11	2.23	0.40
10:B:5:G:C2'	10:B:6:G:C5'	2.98	0.40
10:B:9:G:H5''	10:B:11:A:OP2	2.21	0.40
10:B:38:A:H4'	18:D:790:A:C1'	2.49	0.40
18:D:255:G:H2'	18:D:256:U:C6	2.55	0.40
18:D:262:A:C4	18:D:263:A:C8	3.10	0.40
18:D:276:G:H5''	36:V:17:MET:HE3	2.03	0.40
18:D:333:U:OP1	19:E:2:ALA:N	2.55	0.40
18:D:436:C:N3	18:D:437:U:C5	2.89	0.40
18:D:505:G:H2'	18:D:506:G:C8	2.57	0.40
18:D:519:C:H2'	18:D:520:A:O4'	2.20	0.40
18:D:545:C:OP2	24:J:62:ARG:NH1	2.53	0.40
18:D:562:U:O3'	32:R:12:ARG:HD2	2.20	0.40
18:D:671:G:O2'	26:L:79:ARG:NH1	2.54	0.40
18:D:811:C:O2'	18:D:901:A:N1	2.50	0.40
18:D:1372:U:H2'	18:D:1373:G:O4'	2.22	0.40
18:D:1382:C:O2	18:D:1382:C:H2'	2.19	0.40
18:D:1499:A:OP2	18:D:1499:A:H3'	2.22	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:E:16:LYS:CG	19:E:19:LYS:HE3	2.52	0.40
21:G:17:GLY:C	22:H:43:LYS:HD2	2.46	0.40
22:H:42:LEU:HD23	22:H:81:GLU:HB3	2.03	0.40
24:J:85:ASN:ND2	25:K:101:GLU:OE1	2.52	0.40
24:J:197:GLU:HA	24:J:200:ILE:HD12	2.03	0.40
26:L:24:ARG:HA	26:L:24:ARG:NE	2.36	0.40
26:L:38:ARG:CZ	26:L:99:ALA:HB2	2.52	0.40
29:O:91:ASP:C	29:O:92:GLU:OE2	2.64	0.40
30:P:22:THR:HG21	30:P:72:ARG:HG3	2.03	0.40
36:V:61:ILE:HA	36:V:75:LEU:HA	2.02	0.40
38:X:65:VAL:CG2	38:X:66:GLU:OE1	2.70	0.40
41:a:16:C:C4'	49:i:15:MET:CE	2.99	0.40
41:a:826:U:O2'	61:u:53:GLY:HA3	2.20	0.40
41:a:1289:C:H2'	41:a:1290:C:C6	2.56	0.40
41:a:2065:C:N3	41:a:2066:C:C5	2.89	0.40
41:a:2072:C:N3	41:a:2073:C:C5	2.90	0.40
41:a:2223:G:OP1	48:h:171:TYR:OH	2.38	0.40
41:a:2250:G:OP1	41:a:2275:C:O2'	2.25	0.40
41:a:2639:A:H2'	41:a:2640:G:O4'	2.21	0.40
41:a:2650:U:C2	41:a:2651:C:C5	3.08	0.40
41:a:2848:G:N2	41:a:2867:G:C4	2.89	0.40
42:b:55:ARG:NE	42:b:55:ARG:CA	2.84	0.40
46:f:25:LEU:HD23	46:f:25:LEU:C	2.46	0.40
47:g:58:ASP:O	47:g:59:ARG:C	2.64	0.40
48:h:53:HIS:CD2	48:h:218:PRO:O	2.74	0.40
48:h:124:ILE:O	48:h:124:ILE:HG22	2.21	0.40
48:h:180:GLU:O	48:h:181:MET:HE2	2.21	0.40
50:j:55:LYS:HD3	50:j:55:LYS:C	2.45	0.40
51:k:21:TYR:CZ	51:k:38:LYS:CG	3.04	0.40
56:p:149:ARG:NH1	56:p:154:PRO:CG	2.84	0.40
58:r:5:LEU:CD1	58:r:17:ASP:HB3	2.50	0.40
59:s:53:TYR:HD1	59:s:121:LYS:HB3	1.86	0.40
60:t:59:LYS:NZ	60:t:92:GLU:HG2	2.37	0.40
60:t:102:PRO:HD3	65:y:66:ASN:HB2	2.02	0.40
61:u:123:ARG:HG3	61:u:143:GLU:OE2	2.21	0.40
63:w:45:ARG:CZ	63:w:45:ARG:HB3	2.50	0.40
65:y:4:ILE:O	65:y:8:LEU:HD13	2.21	0.40
66:z:53:ARG:HG2	66:z:57:PHE:HE2	1.83	0.40
9:9:11:ILE:O	9:9:15:VAL:HG23	2.20	0.40
9:9:59:LEU:HD22	9:9:62:ARG:HE	1.81	0.40
9:9:96:PHE:O	9:9:125:ARG:NH1	2.54	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AA:213:LEU:HD13	11:AA:422:LYS:HG2	2.03	0.40
11:AA:987:GLU:HG2	11:AA:991:LYS:HE3	2.03	0.40
12:AB:116:VAL:N	12:AB:129:ILE:HD11	2.36	0.40
12:AB:120:GLU:C	12:AB:162:LEU:HD12	2.46	0.40
14:AE:1155:ILE:HG13	14:AE:1210:ILE:HB	2.02	0.40
14:AE:1178:THR:HG23	14:AE:1184:ASP:HB3	2.04	0.40
16:AG:127:VAL:HG22	16:AG:192:GLY:CA	2.46	0.40
18:D:9:G:OP2	25:K:126:LYS:HD2	2.21	0.40
18:D:982:U:H4'	18:D:983:A:O4'	2.21	0.40
18:D:983:A:C2	18:D:984:C:C6	3.10	0.40
19:E:15:GLU:CD	19:E:16:LYS:N	2.80	0.40
21:G:17:GLY:CA	21:G:41:ILE:HD11	2.52	0.40
22:H:34:ASP:OD1	22:H:50:ALA:HB3	2.20	0.40
24:J:76:TYR:CD1	24:J:76:TYR:C	2.98	0.40
24:J:172:GLU:HG3	24:J:183:LYS:HD2	2.04	0.40
25:K:46:VAL:C	25:K:71:MET:HG3	2.47	0.40
26:L:51:ILE:CG2	26:L:86:ARG:HH21	2.34	0.40
26:L:73:GLU:O	26:L:76:THR:OG1	2.33	0.40
27:M:93:PRO:O	27:M:96:ARG:HG2	2.20	0.40
28:N:83:LEU:HD21	32:R:4:VAL:HG21	2.02	0.40
35:U:69:ASP:OD1	35:U:69:ASP:C	2.63	0.40
38:X:81:MET:HE1	38:X:92:ARG:CD	2.48	0.40
41:a:151:C:C2	41:a:152:A:N7	2.89	0.40
41:a:235:U:H2'	41:a:236:C:H6	1.86	0.40
41:a:239:C:O2'	41:a:622:G:O2'	2.32	0.40
41:a:244:A:H2'	41:a:245:G:O4'	2.21	0.40
41:a:320:A:H2'	52:l:131:THR:HG21	2.03	0.40
41:a:370:G:C6	41:a:424:G:N7	2.89	0.40
41:a:392:U:N3	41:a:393:C:C5	2.89	0.40
41:a:540:C:C2	41:a:541:A:C8	3.09	0.40
41:a:706:A:C2	41:a:707:G:H1'	2.57	0.40
41:a:859:G:O2'	41:a:916:G:O6	2.37	0.40
41:a:871:U:H2'	41:a:872:U:H6	1.78	0.40
41:a:873:C:C2	41:a:874:G:C8	3.09	0.40
41:a:1141:U:C5	59:s:68:LYS:NZ	2.87	0.40
41:a:1250:G:OP2	61:u:21:ARG:NE	2.47	0.40
41:a:1923:U:OP2	41:a:1923:U:H2'	2.21	0.40
41:a:1945:G:H2'	41:a:1946:U:C6	2.56	0.40
41:a:2048:G:C6	41:a:2049:G:C5	3.10	0.40
41:a:2063:C:O2	41:a:2063:C:C2'	2.69	0.40
41:a:2454:G:C5	41:a:2455:G:N7	2.89	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2627:G:H2'	41:a:2628:C:C6	2.56	0.40
41:a:2728:U:H5'	60:t:70:ARG:HH22	1.85	0.40
45:e:2:LYS:HZ1	45:e:49:ASP:CA	2.34	0.40
47:g:7:PRO:CG	54:n:62:GLY:O	2.69	0.40
50:j:61:THR:CG2	50:j:64:GLU:HG2	2.52	0.40
51:k:19:HIS:CD2	51:k:20:PHE:H	2.39	0.40
54:n:30:ARG:O	54:n:159:THR:HG23	2.22	0.40
56:p:52:PHE:CE1	56:p:69:ARG:HD3	2.55	0.40
59:s:4:PHE:CG	66:z:100:VAL:HG11	2.57	0.40
62:v:50:ARG:O	62:v:53:MET:SD	2.80	0.40
63:w:47:VAL:C	63:w:50:PRO:HD2	2.47	0.40
65:y:48:ILE:HD11	65:y:101:ARG:HH11	1.86	0.40
66:z:111:GLU:OE2	66:z:112:LYS:HG2	2.20	0.40
1:0:75:VAL:HG12	66:z:44:GLN:OE1	2.22	0.40
4:3:85:PHE:HE1	4:3:94:ARG:HG2	1.86	0.40
9:9:19:ALA:HB2	9:9:69:PHE:CE2	2.57	0.40
9:9:38:MET:SD	9:9:41:LEU:HD12	2.61	0.40
10:A:29:G:C6	10:A:42:G:C6	3.10	0.40
11:AA:477:GLU:OE1	12:AB:71:ALA:O	2.40	0.40
12:AB:2:GLN:HA	12:AB:59:PHE:O	2.21	0.40
12:AB:16:ARG:O	12:AB:19:GLU:HB3	2.22	0.40
14:AE:68:TYR:CA	14:AE:92:VAL:HG23	2.45	0.40
14:AE:438:GLU:HG3	14:AE:485:MET:HE1	2.03	0.40
14:AE:847:ASP:OD1	14:AE:847:ASP:N	2.54	0.40
16:AG:182:VAL:CG2	16:AG:204:MET:HA	2.52	0.40
16:AG:381:LEU:CA	16:AG:384:LEU:CD1	2.86	0.40
18:D:338:A:H2'	18:D:339:C:H6	1.85	0.40
18:D:502:A:H2'	18:D:503:C:H6	1.86	0.40
18:D:553:A:H2'	18:D:554:A:C8	2.56	0.40
18:D:680:C:H2'	18:D:681:A:H8	1.85	0.40
18:D:860:A:H2'	18:D:861:G:O4'	2.20	0.40
19:E:9:LYS:HG2	19:E:13:GLN:NE2	2.36	0.40
21:G:100:MET:HE1	21:G:101:LEU:CD2	2.49	0.40
22:H:32:ASP:OD1	22:H:32:ASP:C	2.64	0.40
22:H:170:ARG:O	22:H:171:ARG:CB	2.69	0.40
25:K:34:THR:HB	25:K:50:TYR:CE2	2.57	0.40
28:N:83:LEU:HD23	36:V:36:LYS:CD	2.52	0.40
29:O:118:LEU:HB3	29:O:120:LYS:O	2.21	0.40
31:Q:75:LYS:HG3	31:Q:76:GLU:N	2.36	0.40
32:R:57:LEU:HD23	32:R:59:ASN:OD1	2.20	0.40
33:S:37:SER:O	33:S:41:ARG:HG3	2.22	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:X:36:ALA:HB3	38:X:59:GLU:HG2	2.03	0.40
41:a:48:G:C2	41:a:178:G:C6	3.09	0.40
41:a:181:A:H2'	41:a:182:A:C8	2.56	0.40
41:a:814:C:N3	41:a:815:C:C5	2.89	0.40
41:a:966:G:C1'	41:a:2267:A:H62	2.35	0.40
41:a:1005:C:H1'	41:a:1012:U:N3	2.37	0.40
41:a:1406:U:C2	41:a:1407:G:N7	2.89	0.40
41:a:1709:U:C2	41:a:1710:G:C8	3.10	0.40
41:a:1996:C:H4'	50:j:128:ARG:NH1	2.35	0.40
41:a:2065:C:C2	41:a:2066:C:C6	3.09	0.40
41:a:2216:G:H2'	41:a:2217:G:C8	2.57	0.40
41:a:2313:C:O2'	54:n:35:THR:HG21	2.21	0.40
41:a:2549:G:C2	41:a:2550:G:C5	3.10	0.40
41:a:2723:C:C5'	63:w:1:MET:HE3	2.48	0.40
45:e:14:LEU:CA	45:e:17:GLU:OE1	2.70	0.40
53:m:22:MET:SD	53:m:28:ARG:HD3	2.62	0.40
53:m:25:LYS:O	53:m:26:ASN:C	2.64	0.40
54:n:16:LEU:HD11	54:n:169:LEU:HD12	2.03	0.40
54:n:136:ILE:HA	54:n:141:ILE:CG2	2.51	0.40
60:t:35:VAL:HG11	60:t:69:VAL:HG11	2.04	0.40
60:t:103:VAL:HG21	60:t:107:LEU:HD22	2.03	0.40
62:v:110:GLU:H	62:v:110:GLU:CD	2.29	0.40
63:w:78:LYS:HA	63:w:82:GLU:OE1	2.21	0.40
5:4:50:MET:O	5:4:56:PHE:CZ	2.75	0.40
5:4:75:GLN:CG	5:4:92:VAL:HG23	2.47	0.40
10:A:16:C:H3'	10:A:17:C:H5''	2.04	0.40
11:AA:888:THR:C	16:AG:105:ILE:HD13	2.46	0.40
14:AE:282:LEU:HD23	14:AE:282:LEU:HA	1.92	0.40
16:AG:282:GLN:HA	16:AG:282:GLN:HE21	1.82	0.40
16:AG:302:LYS:HE3	16:AG:302:LYS:CA	2.51	0.40
10:B:29:G:C6	10:B:42:G:C6	3.10	0.40
17:C:27:ALA:HA	17:C:30:LYS:HE3	2.04	0.40
18:D:104:G:C2	18:D:105:G:C8	3.10	0.40
18:D:619:U:C1'	24:J:128:ARG:NH1	2.85	0.40
18:D:915:A:H8	18:D:915:A:O5'	2.04	0.40
18:D:1061:G:H5'	30:P:61:ALA:HB2	2.02	0.40
18:D:1084:G:OP1	18:D:1086:U:C6	2.75	0.40
19:E:59:ASP:OD1	19:E:76:LYS:HE2	2.21	0.40
21:G:90:PHE:CD2	21:G:154:MET:SD	3.14	0.40
22:H:16:ILE:O	22:H:25:ARG:NH1	2.55	0.40
22:H:119:GLY:HA3	22:H:132:PRO:C	2.47	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:I:30:ALA:O	33:S:65:ARG:NH2	2.55	0.40
24:J:71:GLN:O	24:J:74:ASN:HB3	2.22	0.40
25:K:99:ALA:HB1	25:K:103:THR:HG21	2.03	0.40
25:K:156:LYS:HA	28:N:66:PHE:CG	2.57	0.40
26:L:9:MET:SD	26:L:57:ALA:HB1	2.60	0.40
27:M:20:SER:OG	27:M:23:LEU:HD13	2.22	0.40
27:M:35:LYS:CE	27:M:38:THR:OG1	2.69	0.40
34:T:17:ARG:HG3	34:T:18:ASP:N	2.36	0.40
41:a:909:A:C4	41:a:912:C:C5	3.09	0.40
41:a:1057:A:C2	41:a:1082:U:N3	2.89	0.40
41:a:1065:U:HO2'	41:a:1066:U:P	2.42	0.40
41:a:1520:U:O4	41:a:1521:G:C6	2.75	0.40
41:a:1636:U:H2'	41:a:1637:A:C8	2.57	0.40
41:a:1651:G:P	63:w:40:LYS:NZ	2.94	0.40
41:a:1881:C:C4	41:a:1882:U:C5	3.09	0.40
41:a:2291:U:H2'	41:a:2292:U:H6	1.85	0.40
41:a:2314:A:O2'	54:n:155:THR:CG2	2.69	0.40
41:a:2619:C:O2'	41:a:2620:C:H5'	2.21	0.40
41:a:2839:G:H2'	41:a:2840:C:H6	1.86	0.40
46:f:17:LEU:O	46:f:21:LYS:HG2	2.22	0.40
48:h:161:TYR:HB3	48:h:194:GLU:HG2	2.03	0.40
49:i:28:LEU:CB	49:i:37:LYS:HE3	2.51	0.40
50:j:5:VAL:HB	50:j:32:ASN:HD21	1.86	0.40
50:j:45:TYR:CE2	50:j:81:GLU:OE2	2.75	0.40
51:k:12:VAL:HG12	51:k:50:LYS:O	2.22	0.40
52:l:40:ARG:HD2	52:l:92:HIS:CG	2.57	0.40
54:n:9:LYS:HE3	54:n:9:LYS:HB3	1.91	0.40
54:n:150:ARG:HG2	54:n:151:GLY:N	2.36	0.40
59:s:9:GLU:HG2	59:s:10:THR:N	2.37	0.40
61:u:95:LEU:HD13	61:u:100:ILE:CD1	2.51	0.40
62:v:73:ILE:HD13	62:v:93:VAL:HG13	2.04	0.40
65:y:30:VAL:HG12	65:y:81:VAL:HG12	2.03	0.40
66:z:86:ALA:O	66:z:87:SER:OG	2.35	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	0	101/103 (98%)	97 (96%)	4 (4%)	0	100	100
2	1	108/110 (98%)	107 (99%)	1 (1%)	0	100	100
3	2	92/100 (92%)	87 (95%)	5 (5%)	0	100	100
4	3	101/104 (97%)	98 (97%)	3 (3%)	0	100	100
5	4	92/94 (98%)	89 (97%)	3 (3%)	0	100	100
9	9	146/165 (88%)	100 (68%)	43 (30%)	3 (2%)	5	28
11	AA	1312/1342 (98%)	1197 (91%)	114 (9%)	1 (0%)	48	82
12	AB	159/162 (98%)	107 (67%)	41 (26%)	11 (7%)	1	10
13	AC	217/329 (66%)	203 (94%)	12 (6%)	2 (1%)	14	49
13	AD	293/329 (89%)	263 (90%)	27 (9%)	3 (1%)	12	47
14	AE	1331/1407 (95%)	1211 (91%)	114 (9%)	6 (0%)	24	63
15	AF	80/91 (88%)	77 (96%)	3 (4%)	0	100	100
16	AG	493/495 (100%)	375 (76%)	86 (17%)	32 (6%)	1	12
17	C	64/75 (85%)	63 (98%)	1 (2%)	0	100	100
19	E	84/87 (97%)	83 (99%)	1 (1%)	0	100	100
20	F	68/71 (96%)	68 (100%)	0	0	100	100
21	G	223/241 (92%)	212 (95%)	10 (4%)	1 (0%)	30	67
22	H	255/557 (46%)	182 (71%)	66 (26%)	7 (3%)	4	24
23	I	206/233 (88%)	193 (94%)	13 (6%)	0	100	100
24	J	203/206 (98%)	201 (99%)	2 (1%)	0	100	100
25	K	154/167 (92%)	145 (94%)	8 (5%)	1 (1%)	21	58
26	L	102/135 (76%)	97 (95%)	4 (4%)	1 (1%)	12	47
27	M	149/179 (83%)	140 (94%)	8 (5%)	1 (1%)	18	55
28	N	127/130 (98%)	123 (97%)	4 (3%)	0	100	100
29	O	125/130 (96%)	116 (93%)	8 (6%)	1 (1%)	16	52

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
30	P	97/103 (94%)	89 (92%)	8 (8%)	0	100	100
31	Q	115/129 (89%)	107 (93%)	8 (7%)	0	100	100
32	R	117/124 (94%)	112 (96%)	5 (4%)	0	100	100
33	S	98/101 (97%)	97 (99%)	1 (1%)	0	100	100
34	T	86/89 (97%)	83 (96%)	3 (4%)	0	100	100
35	U	80/82 (98%)	76 (95%)	4 (5%)	0	100	100
36	V	78/84 (93%)	74 (95%)	4 (5%)	0	100	100
37	W	81/92 (88%)	80 (99%)	1 (1%)	0	100	100
38	X	114/118 (97%)	103 (90%)	9 (8%)	2 (2%)	6	32
39	Y	139/142 (98%)	101 (73%)	38 (27%)	0	100	100
40	Z	28/121 (23%)	22 (79%)	6 (21%)	0	100	100
42	b	74/85 (87%)	73 (99%)	1 (1%)	0	100	100
43	c	75/78 (96%)	72 (96%)	3 (4%)	0	100	100
45	e	60/63 (95%)	57 (95%)	3 (5%)	0	100	100
46	f	56/59 (95%)	52 (93%)	4 (7%)	0	100	100
47	g	64/70 (91%)	62 (97%)	2 (3%)	0	100	100
48	h	269/273 (98%)	255 (95%)	14 (5%)	0	100	100
49	i	54/57 (95%)	49 (91%)	5 (9%)	0	100	100
50	j	207/209 (99%)	198 (96%)	9 (4%)	0	100	100
51	k	50/55 (91%)	50 (100%)	0	0	100	100
52	l	199/201 (99%)	188 (94%)	11 (6%)	0	100	100
53	m	44/46 (96%)	43 (98%)	1 (2%)	0	100	100
54	n	175/179 (98%)	160 (91%)	14 (8%)	1 (1%)	21	58
55	o	62/65 (95%)	57 (92%)	4 (6%)	1 (2%)	7	36
56	p	173/177 (98%)	162 (94%)	11 (6%)	0	100	100
57	q	36/38 (95%)	35 (97%)	1 (3%)	0	100	100
58	r	147/149 (99%)	139 (95%)	8 (5%)	0	100	100
59	s	140/142 (99%)	133 (95%)	7 (5%)	0	100	100
60	t	121/123 (98%)	114 (94%)	7 (6%)	0	100	100
61	u	142/144 (99%)	134 (94%)	8 (6%)	0	100	100
62	v	134/136 (98%)	129 (96%)	5 (4%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
63	w	117/127 (92%)	112 (96%)	5 (4%)	0	100	100
64	x	114/117 (97%)	107 (94%)	7 (6%)	0	100	100
65	y	112/115 (97%)	105 (94%)	7 (6%)	0	100	100
66	z	115/118 (98%)	111 (96%)	4 (4%)	0	100	100
All	All	10058/11053 (91%)	9175 (91%)	809 (8%)	74 (1%)	20	55

All (74) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
12	AB	107	PRO
12	AB	125	GLY
12	AB	130	PHE
12	AB	137	ALA
12	AB	141	LEU
12	AB	152	HIS
12	AB	160	ARG
13	AD	286	GLU
16	AG	34	ALA
16	AG	105	ILE
16	AG	187	ARG
16	AG	227	ALA
21	G	127	ASP
22	H	304	VAL
27	M	56	LYS
55	o	32	ILE
12	AB	105	VAL
14	AE	92	VAL
14	AE	175	GLU
16	AG	63	LEU
16	AG	104	ARG
16	AG	242	ASP
16	AG	268	GLY
16	AG	279	ASN
16	AG	365	ILE
22	H	171	ARG
22	H	305	HIS
22	H	309	MET
12	AB	127	GLN
16	AG	33	THR
16	AG	102	PHE
16	AG	200	SER

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Mol	Chain	Res	Type
16	AG	305	MET
16	AG	319	GLY
16	AG	350	ALA
16	AG	355	ALA
16	AG	396	GLU
16	AG	425	LEU
25	K	78	ASN
26	L	96	VAL
29	O	13	LYS
54	n	177	PHE
13	AC	193	GLU
16	AG	48	GLN
16	AG	265	GLU
16	AG	363	LEU
16	AG	401	PRO
22	H	82	THR
12	AB	53	ASN
13	AD	289	LEU
14	AE	74	LYS
14	AE	286	ALA
16	AG	224	LYS
16	AG	399	ASP
22	H	143	ASP
38	X	6	GLY
38	X	66	GLU
9	9	79	PRO
9	9	88	HIS
13	AC	192	VAL
13	AD	306	VAL
14	AE	290	ILE
16	AG	43	ILE
16	AG	70	GLN
16	AG	169	PRO
16	AG	171	GLU
16	AG	323	GLN
16	AG	364	ASP
16	AG	382	GLU
22	H	71	ALA
14	AE	292	VAL
9	9	129	LEU
11	AA	1317	PRO
12	AB	116	VAL

5.3.2 Protein sidechains ⓘ

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	0	84/84 (100%)	84 (100%)	0	100	100
2	1	93/93 (100%)	93 (100%)	0	100	100
3	2	81/84 (96%)	81 (100%)	0	100	100
4	3	84/85 (99%)	84 (100%)	0	100	100
5	4	78/78 (100%)	78 (100%)	0	100	100
9	9	112/123 (91%)	112 (100%)	0	100	100
11	AA	1135/1157 (98%)	1130 (100%)	5 (0%)	84	83
12	AB	141/142 (99%)	124 (88%)	17 (12%)	5	17
13	AC	186/286 (65%)	186 (100%)	0	100	100
13	AD	185/286 (65%)	185 (100%)	0	100	100
14	AE	1122/1168 (96%)	1104 (98%)	18 (2%)	55	69
15	AF	70/75 (93%)	69 (99%)	1 (1%)	59	71
16	AG	409/409 (100%)	288 (70%)	121 (30%)	0	2
17	C	57/65 (88%)	57 (100%)	0	100	100
19	E	65/66 (98%)	65 (100%)	0	100	100
20	F	60/61 (98%)	60 (100%)	0	100	100
21	G	187/199 (94%)	187 (100%)	0	100	100
22	H	137/461 (30%)	137 (100%)	0	100	100
23	I	171/190 (90%)	171 (100%)	0	100	100
24	J	172/173 (99%)	171 (99%)	1 (1%)	78	81
25	K	119/126 (94%)	119 (100%)	0	100	100
26	L	91/116 (78%)	91 (100%)	0	100	100
27	M	124/147 (84%)	124 (100%)	0	100	100
28	N	104/105 (99%)	104 (100%)	0	100	100
29	O	105/107 (98%)	105 (100%)	0	100	100
30	P	86/90 (96%)	86 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
31	Q	90/99 (91%)	90 (100%)	0	100	100
32	R	101/104 (97%)	101 (100%)	0	100	100
33	S	83/84 (99%)	83 (100%)	0	100	100
34	T	76/77 (99%)	76 (100%)	0	100	100
35	U	65/65 (100%)	65 (100%)	0	100	100
36	V	74/78 (95%)	74 (100%)	0	100	100
37	W	72/79 (91%)	72 (100%)	0	100	100
38	X	94/96 (98%)	94 (100%)	0	100	100
39	Y	109/110 (99%)	109 (100%)	0	100	100
40	Z	26/85 (31%)	26 (100%)	0	100	100
42	b	58/63 (92%)	58 (100%)	0	100	100
43	c	67/68 (98%)	67 (100%)	0	100	100
45	e	54/55 (98%)	54 (100%)	0	100	100
46	f	48/49 (98%)	48 (100%)	0	100	100
47	g	59/62 (95%)	59 (100%)	0	100	100
48	h	216/218 (99%)	216 (100%)	0	100	100
49	i	47/48 (98%)	47 (100%)	0	100	100
50	j	164/164 (100%)	164 (100%)	0	100	100
51	k	47/49 (96%)	47 (100%)	0	100	100
52	l	165/165 (100%)	165 (100%)	0	100	100
53	m	38/38 (100%)	38 (100%)	0	100	100
54	n	148/150 (99%)	148 (100%)	0	100	100
55	o	51/52 (98%)	51 (100%)	0	100	100
56	p	136/138 (99%)	136 (100%)	0	100	100
57	q	34/34 (100%)	34 (100%)	0	100	100
58	r	114/114 (100%)	114 (100%)	0	100	100
59	s	116/116 (100%)	116 (100%)	0	100	100
60	t	104/104 (100%)	104 (100%)	0	100	100
61	u	103/103 (100%)	103 (100%)	0	100	100
62	v	109/109 (100%)	109 (100%)	0	100	100
63	w	99/103 (96%)	99 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
64	x	86/87 (99%)	86 (100%)	0	100	100
65	y	99/100 (99%)	99 (100%)	0	100	100
66	z	89/90 (99%)	89 (100%)	0	100	100
All	All	8299/9132 (91%)	8136 (98%)	163 (2%)	48	65

All (163) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
11	AA	145	ILE
11	AA	615	VAL
11	AA	914	LYS
11	AA	1076	ILE
11	AA	1151	LEU
12	AB	115	LYS
12	AB	116	VAL
12	AB	117	ILE
12	AB	118	ILE
12	AB	119	THR
12	AB	120	GLU
12	AB	123	PHE
12	AB	124	GLU
12	AB	129	ILE
12	AB	143	LEU
12	AB	145	LEU
12	AB	146	ILE
12	AB	149	GLU
12	AB	150	ILE
12	AB	159	PHE
12	AB	160	ARG
12	AB	161	LYS
14	AE	69	GLU
14	AE	70	CYS
14	AE	71	LEU
14	AE	74	LYS
14	AE	76	LYS
14	AE	77	ARG
14	AE	78	LEU
14	AE	79	LYS
14	AE	81	ARG
14	AE	84	ILE
14	AE	85	CYS

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Mol	Chain	Res	Type
14	AE	92	VAL
14	AE	93	THR
14	AE	290	ILE
14	AE	291	ILE
14	AE	292	VAL
14	AE	514	THR
14	AE	1261	LEU
15	AF	63	ILE
16	AG	5	ILE
16	AG	12	VAL
16	AG	16	LYS
16	AG	36	LYS
16	AG	37	LYS
16	AG	40	GLU
16	AG	43	ILE
16	AG	47	VAL
16	AG	49	ILE
16	AG	50	ASP
16	AG	56	PHE
16	AG	57	ASP
16	AG	61	ARG
16	AG	63	LEU
16	AG	75	ILE
16	AG	97	ILE
16	AG	103	ASP
16	AG	104	ARG
16	AG	114	ILE
16	AG	117	LYS
16	AG	123	ARG
16	AG	125	MET
16	AG	126	VAL
16	AG	131	ARG
16	AG	138	ILE
16	AG	139	THR
16	AG	144	LYS
16	AG	150	ILE
16	AG	154	LEU
16	AG	157	ASN
16	AG	162	ILE
16	AG	167	MET
16	AG	173	PHE
16	AG	174	ARG

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Mol	Chain	Res	Type
16	AG	179	VAL
16	AG	180	ARG
16	AG	182	VAL
16	AG	187	ARG
16	AG	189	GLU
16	AG	195	LEU
16	AG	197	VAL
16	AG	199	ARG
16	AG	201	LYS
16	AG	203	GLU
16	AG	204	MET
16	AG	205	LEU
16	AG	206	ILE
16	AG	207	GLU
16	AG	210	ARG
16	AG	211	ILE
16	AG	213	VAL
16	AG	218	GLU
16	AG	219	GLU
16	AG	221	ILE
16	AG	223	ILE
16	AG	232	SER
16	AG	235	LYS
16	AG	236	ILE
16	AG	238	VAL
16	AG	239	LYS
16	AG	240	THR
16	AG	241	ASN
16	AG	243	LYS
16	AG	244	ARG
16	AG	245	ILE
16	AG	254	MET
16	AG	255	ARG
16	AG	258	ARG
16	AG	259	VAL
16	AG	260	GLN
16	AG	262	VAL
16	AG	265	GLU
16	AG	270	ARG
16	AG	271	ILE
16	AG	273	ILE
16	AG	274	VAL

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Mol	Chain	Res	Type
16	AG	276	TRP
16	AG	278	ASP
16	AG	282	GLN
16	AG	285	ILE
16	AG	288	MET
16	AG	296	ILE
16	AG	297	VAL
16	AG	299	ASP
16	AG	300	GLU
16	AG	301	ASP
16	AG	302	LYS
16	AG	303	HIS
16	AG	305	MET
16	AG	309	VAL
16	AG	310	GLU
16	AG	313	ASN
16	AG	316	GLN
16	AG	320	ARG
16	AG	321	ASN
16	AG	323	GLN
16	AG	327	LEU
16	AG	331	LEU
16	AG	335	GLU
16	AG	336	LEU
16	AG	337	ASN
16	AG	338	VAL
16	AG	340	THR
16	AG	341	VAL
16	AG	342	ASP
16	AG	344	LEU
16	AG	356	ILE
16	AG	358	THR
16	AG	359	PHE
16	AG	361	LYS
16	AG	365	ILE
16	AG	380	THR
16	AG	387	VAL
16	AG	390	LYS
16	AG	396	GLU
16	AG	398	LEU
16	AG	399	ASP
16	AG	400	GLU

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Mol	Chain	Res	Type
16	AG	425	LEU
16	AG	428	ASN
16	AG	429	LYS
24	J	47	ARG

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (124) such sidechains are listed below:

Mol	Chain	Res	Type
1	0	6	GLN
1	0	82	HIS
2	1	7	HIS
2	1	9	HIS
3	2	15	HIS
9	9	9	GLN
11	AA	69	GLN
11	AA	314	ASN
11	AA	330	HIS
11	AA	343	HIS
11	AA	387	ASN
11	AA	513	GLN
11	AA	554	HIS
11	AA	688	GLN
11	AA	762	ASN
11	AA	808	ASN
11	AA	856	ASN
11	AA	1008	GLN
11	AA	1013	GLN
11	AA	1237	HIS
11	AA	1288	GLN
11	AA	1313	HIS
12	AB	27	ASN
12	AB	65	HIS
12	AB	144	ASN
12	AB	147	ASN
13	AC	23	HIS
13	AC	147	GLN
13	AD	37	HIS
13	AD	66	HIS
13	AD	84	ASN
13	AD	117	HIS
13	AD	128	HIS
13	AD	227	GLN

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Mol	Chain	Res	Type
14	AE	80	HIS
14	AE	157	GLN
14	AE	164	GLN
14	AE	232	ASN
14	AE	365	GLN
14	AE	424	ASN
14	AE	450	HIS
14	AE	469	HIS
14	AE	777	HIS
14	AE	910	ASN
14	AE	1108	GLN
14	AE	1197	ASN
14	AE	1235	ASN
14	AE	1238	GLN
14	AE	1326	GLN
15	AF	62	GLN
16	AG	108	GLN
16	AG	194	GLN
16	AG	260	GLN
16	AG	282	GLN
16	AG	316	GLN
16	AG	323	GLN
16	AG	324	ASN
16	AG	337	ASN
16	AG	428	ASN
17	C	54	GLN
19	E	13	GLN
19	E	61	GLN
20	F	9	ASN
21	G	18	HIS
21	G	39	HIS
21	G	94	HIS
22	H	286	ASN
23	I	3	GLN
23	I	6	HIS
23	I	8	ASN
23	I	190	HIS
24	J	36	GLN
24	J	131	ASN
24	J	198	HIS
25	K	89	HIS
26	L	3	HIS

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Mol	Chain	Res	Type
26	L	68	GLN
27	M	68	ASN
27	M	86	GLN
29	O	5	GLN
29	O	81	HIS
29	O	126	GLN
30	P	58	ASN
32	R	77	HIS
33	S	4	GLN
34	T	80	GLN
35	U	26	ASN
35	U	59	HIS
37	W	83	HIS
39	Y	5	GLN
39	Y	104	GLN
45	e	25	GLN
45	e	31	GLN
46	f	20	HIS
46	f	49	ASN
47	g	33	ASN
48	h	53	HIS
48	h	90	ASN
49	i	6	ASN
50	j	173	GLN
51	k	19	HIS
52	l	92	HIS
52	l	156	ASN
52	l	163	ASN
53	m	29	GLN
54	n	5	HIS
54	n	81	GLN
55	o	24	HIS
55	o	28	ASN
57	q	13	ASN
58	r	133	GLN
59	s	132	HIS
62	v	13	HIS
62	v	45	GLN
62	v	97	GLN
63	w	18	GLN
64	x	19	GLN
64	x	29	HIS

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Mol	Chain	Res	Type
64	x	38	GLN
65	y	10	GLN
65	y	77	HIS
66	z	37	GLN
66	z	72	ASN
66	z	81	ASN

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
10	A	75/76 (98%)	29 (38%)	8 (10%)
10	B	75/76 (98%)	29 (38%)	8 (10%)
18	D	1514/1542 (98%)	289 (19%)	20 (1%)
41	a	2859/2904 (98%)	510 (17%)	0
44	d	119/120 (99%)	15 (12%)	0
8	7	23/53 (43%)	14 (60%)	3 (13%)
All	All	4665/4771 (97%)	886 (18%)	39 (0%)

All (886) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
8	7	3	G
8	7	4	U
8	7	5	U
8	7	7	U
8	7	8	U
8	7	9	U
8	7	10	U
8	7	11	U
8	7	12	U
8	7	13	U
8	7	14	U
8	7	44	U
8	7	47	U
8	7	49	A
10	A	2	G
10	A	6	G
10	A	7	G
10	A	8	U
10	A	10	G
10	A	13	C

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Mol	Chain	Res	Type
10	A	14	A
10	A	16	C
10	A	17	C
10	A	18	G
10	A	19	G
10	A	20	U
10	A	21	A
10	A	22	G
10	A	23	C
10	A	30	G
10	A	46	G
10	A	47	U
10	A	48	C
10	A	49	G
10	A	52	G
10	A	57	A
10	A	58	A
10	A	59	A
10	A	61	C
10	A	66	C
10	A	69	C
10	A	71	C
10	A	73	A
10	B	2	G
10	B	6	G
10	B	7	G
10	B	8	U
10	B	10	G
10	B	13	C
10	B	14	A
10	B	16	C
10	B	17	C
10	B	18	G
10	B	19	G
10	B	20	U
10	B	21	A
10	B	22	G
10	B	23	C
10	B	30	G
10	B	46	G
10	B	47	U
10	B	48	C

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Mol	Chain	Res	Type
10	B	49	G
10	B	52	G
10	B	57	A
10	B	58	A
10	B	59	A
10	B	61	C
10	B	66	C
10	B	69	C
10	B	71	C
10	B	73	A
18	D	4	U
18	D	5	U
18	D	9	G
18	D	22	G
18	D	29	U
18	D	32	A
18	D	39	G
18	D	47	C
18	D	48	C
18	D	50	A
18	D	51	A
18	D	52	C
18	D	54	C
18	D	69	G
18	D	71	A
18	D	72	A
18	D	74	A
18	D	76	G
18	D	82	G
18	D	83	C
18	D	84	U
18	D	87	C
18	D	90	C
18	D	94	G
18	D	95	C
18	D	96	U
18	D	108	G
18	D	120	A
18	D	121	U
18	D	122	G
18	D	131	A
18	D	141	G

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Mol	Chain	Res	Type
18	D	144	G
18	D	148	G
18	D	149	A
18	D	160	A
18	D	164	G
18	D	173	U
18	D	181	A
18	D	182	A
18	D	183	C
18	D	184	G
18	D	197	A
18	D	198	G
18	D	204	G
18	D	208	U
18	D	209	U
18	D	210	C
18	D	211	G
18	D	212	G
18	D	216	U
18	D	226	G
18	D	245	U
18	D	247	G
18	D	251	G
18	D	253	A
18	D	258	G
18	D	262	A
18	D	266	G
18	D	267	C
18	D	271	C
18	D	279	A
18	D	289	G
18	D	299	G
18	D	306	A
18	D	321	A
18	D	328	C
18	D	329	A
18	D	332	G
18	D	347	G
18	D	352	C
18	D	353	A
18	D	354	G
18	D	355	C

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Mol	Chain	Res	Type
18	D	367	U
18	D	372	C
18	D	373	A
18	D	376	G
18	D	382	A
18	D	384	G
18	D	392	C
18	D	397	A
18	D	398	U
18	D	406	G
18	D	411	A
18	D	412	A
18	D	413	G
18	D	414	A
18	D	421	U
18	D	422	C
18	D	424	G
18	D	429	U
18	D	446	G
18	D	451	A
18	D	457	G
18	D	458	U
18	D	463	U
18	D	464	U
18	D	467	U
18	D	468	A
18	D	469	C
18	D	478	A
18	D	479	U
18	D	481	G
18	D	484	G
18	D	485	U
18	D	486	U
18	D	496	A
18	D	497	G
18	D	505	G
18	D	511	C
18	D	518	C
18	D	519	C
18	D	526	C
18	D	531	U
18	D	532	A

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Mol	Chain	Res	Type
18	D	533	A
18	D	542	G
18	D	547	A
18	D	559	A
18	D	564	C
18	D	568	G
18	D	572	A
18	D	573	A
18	D	576	C
18	D	577	G
18	D	579	A
18	D	596	A
18	D	628	G
18	D	633	G
18	D	649	A
18	D	650	G
18	D	653	U
18	D	665	A
18	D	695	A
18	D	700	G
18	D	723	U
18	D	724	G
18	D	731	G
18	D	734	G
18	D	735	C
18	D	747	A
18	D	748	G
18	D	755	G
18	D	760	G
18	D	777	A
18	D	793	U
18	D	794	A
18	D	815	A
18	D	817	C
18	D	828	U
18	D	829	G
18	D	832	G
18	D	841	C
18	D	843	U
18	D	844	G
18	D	845	A
18	D	846	G

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Mol	Chain	Res	Type
18	D	849	G
18	D	874	G
18	D	887	G
18	D	902	G
18	D	914	A
18	D	916	U
18	D	926	G
18	D	934	C
18	D	935	A
18	D	960	U
18	D	963	G
18	D	965	U
18	D	969	A
18	D	972	C
18	D	975	A
18	D	976	G
18	D	987	G
18	D	991	U
18	D	992	U
18	D	993	G
18	D	996	A
18	D	1004	A
18	D	1008	U
18	D	1009	U
18	D	1017	U
18	D	1018	G
18	D	1021	A
18	D	1024	G
18	D	1026	G
18	D	1028	C
18	D	1030	U
18	D	1031	C
18	D	1037	C
18	D	1043	G
18	D	1044	A
18	D	1046	A
18	D	1065	U
18	D	1085	U
18	D	1094	G
18	D	1095	U
18	D	1099	G
18	D	1101	A

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Mol	Chain	Res	Type
18	D	1108	G
18	D	1124	G
18	D	1133	G
18	D	1135	U
18	D	1136	C
18	D	1137	C
18	D	1139	G
18	D	1140	C
18	D	1141	C
18	D	1142	G
18	D	1143	G
18	D	1145	A
18	D	1146	A
18	D	1151	A
18	D	1152	A
18	D	1159	U
18	D	1167	A
18	D	1171	A
18	D	1174	G
18	D	1175	G
18	D	1176	A
18	D	1184	G
18	D	1193	G
18	D	1196	A
18	D	1197	A
18	D	1206	G
18	D	1211	U
18	D	1212	U
18	D	1213	A
18	D	1214	C
18	D	1215	G
18	D	1226	C
18	D	1227	A
18	D	1228	C
18	D	1238	A
18	D	1257	A
18	D	1260	G
18	D	1275	A
18	D	1276	G
18	D	1278	G
18	D	1279	G
18	D	1280	A

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Mol	Chain	Res	Type
18	D	1285	A
18	D	1286	U
18	D	1287	A
18	D	1299	A
18	D	1300	G
18	D	1302	C
18	D	1305	G
18	D	1312	G
18	D	1317	C
18	D	1320	C
18	D	1323	G
18	D	1338	G
18	D	1340	A
18	D	1346	A
18	D	1347	G
18	D	1353	G
18	D	1363	A
18	D	1370	G
18	D	1378	C
18	D	1379	G
18	D	1381	U
18	D	1391	U
18	D	1396	A
18	D	1397	C
18	D	1398	A
18	D	1404	C
18	D	1419	G
18	D	1429	A
18	D	1441	A
18	D	1446	A
18	D	1447	A
18	D	1448	C
18	D	1452	C
18	D	1453	G
18	D	1475	G
18	D	1487	G
18	D	1491	G
18	D	1492	A
18	D	1493	A
18	D	1494	G
18	D	1497	G
18	D	1503	A

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Mol	Chain	Res	Type
18	D	1506	U
18	D	1517	G
18	D	1529	G
18	D	1530	G
18	D	1534	A
41	a	4	U
41	a	10	A
41	a	15	G
41	a	23	G
41	a	34	U
41	a	35	G
41	a	46	G
41	a	58	G
41	a	60	G
41	a	62	U
41	a	63	A
41	a	71	A
41	a	74	A
41	a	75	G
41	a	83	A
41	a	84	A
41	a	85	G
41	a	96	C
41	a	101	A
41	a	102	U
41	a	103	A
41	a	110	G
41	a	118	A
41	a	119	A
41	a	120	U
41	a	131	A
41	a	136	G
41	a	139	U
41	a	140	C
41	a	141	G
41	a	163	C
41	a	165	A
41	a	181	A
41	a	196	A
41	a	215	G
41	a	216	A
41	a	222	A

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Mol	Chain	Res	Type
41	a	225	C
41	a	248	G
41	a	249	C
41	a	261	G
41	a	264	C
41	a	265	A
41	a	266	G
41	a	267	C
41	a	271	G
41	a	272	A
41	a	275	C
41	a	276	U
41	a	278	A
41	a	285	G
41	a	291	G
41	a	311	A
41	a	329	G
41	a	330	A
41	a	353	C
41	a	361	G
41	a	362	A
41	a	371	A
41	a	372	G
41	a	373	U
41	a	375	G
41	a	383	C
41	a	386	G
41	a	396	G
41	a	405	U
41	a	411	G
41	a	412	A
41	a	420	C
41	a	424	G
41	a	451	U
41	a	457	A
41	a	477	A
41	a	481	G
41	a	491	G
41	a	501	A
41	a	503	A
41	a	504	A
41	a	505	A

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Mol	Chain	Res	Type
41	a	509	C
41	a	522	A
41	a	529	A
41	a	531	C
41	a	532	A
41	a	537	G
41	a	543	G
41	a	545	U
41	a	546	U
41	a	547	A
41	a	549	G
41	a	551	G
41	a	563	A
41	a	569	U
41	a	573	U
41	a	575	A
41	a	588	U
41	a	603	A
41	a	609	A
41	a	613	A
41	a	614	A
41	a	615	U
41	a	616	A
41	a	627	A
41	a	637	A
41	a	645	C
41	a	647	G
41	a	654	A
41	a	668	A
41	a	686	U
41	a	710	U
41	a	717	C
41	a	730	A
41	a	738	G
41	a	757	G
41	a	764	A
41	a	765	C
41	a	775	G
41	a	776	G
41	a	782	A
41	a	784	G
41	a	785	G

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Mol	Chain	Res	Type
41	a	800	A
41	a	805	G
41	a	812	C
41	a	819	A
41	a	827	U
41	a	828	U
41	a	845	A
41	a	846	U
41	a	858	G
41	a	859	G
41	a	869	G
41	a	878	A
41	a	881	G
41	a	884	U
41	a	885	C
41	a	888	C
41	a	891	G
41	a	895	U
41	a	896	A
41	a	897	C
41	a	899	A
41	a	907	G
41	a	910	A
41	a	914	G
41	a	915	C
41	a	931	U
41	a	941	A
41	a	945	A
41	a	946	C
41	a	953	G
41	a	961	C
41	a	974	G
41	a	983	A
41	a	995	C
41	a	996	A
41	a	999	U
41	a	1005	C
41	a	1012	U
41	a	1013	C
41	a	1022	G
41	a	1023	U
41	a	1026	G

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Mol	Chain	Res	Type
41	a	1033	U
41	a	1041	G
41	a	1045	C
41	a	1046	A
41	a	1047	G
41	a	1060	U
41	a	1061	U
41	a	1064	C
41	a	1065	U
41	a	1066	U
41	a	1067	A
41	a	1068	G
41	a	1070	A
41	a	1071	G
41	a	1073	A
41	a	1074	G
41	a	1079	C
41	a	1080	A
41	a	1081	U
41	a	1082	U
41	a	1083	U
41	a	1084	A
41	a	1087	G
41	a	1088	A
41	a	1090	A
41	a	1095	A
41	a	1101	U
41	a	1107	G
41	a	1110	G
41	a	1111	A
41	a	1112	G
41	a	1119	U
41	a	1122	G
41	a	1128	G
41	a	1132	U
41	a	1134	A
41	a	1135	C
41	a	1142	A
41	a	1169	A
41	a	1170	C
41	a	1173	U
41	a	1174	U

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Mol	Chain	Res	Type
41	a	1175	A
41	a	1176	U
41	a	1177	G
41	a	1178	C
41	a	1179	G
41	a	1180	U
41	a	1186	G
41	a	1238	G
41	a	1248	G
41	a	1253	A
41	a	1256	G
41	a	1266	G
41	a	1271	G
41	a	1272	A
41	a	1273	U
41	a	1300	G
41	a	1301	A
41	a	1321	A
41	a	1345	C
41	a	1352	U
41	a	1365	A
41	a	1368	G
41	a	1378	A
41	a	1379	U
41	a	1380	G
41	a	1383	A
41	a	1392	A
41	a	1395	A
41	a	1406	U
41	a	1408	G
41	a	1409	U
41	a	1414	C
41	a	1416	G
41	a	1417	C
41	a	1419	A
41	a	1420	A
41	a	1428	C
41	a	1452	G
41	a	1453	A
41	a	1460	U
41	a	1482	G
41	a	1490	A

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Mol	Chain	Res	Type
41	a	1497	U
41	a	1503	A
41	a	1508	A
41	a	1509	A
41	a	1510	G
41	a	1515	A
41	a	1529	G
41	a	1534	U
41	a	1535	A
41	a	1536	C
41	a	1537	G
41	a	1554	U
41	a	1559	U
41	a	1566	A
41	a	1569	A
41	a	1578	U
41	a	1580	A
41	a	1581	G
41	a	1582	C
41	a	1583	A
41	a	1584	U
41	a	1585	C
41	a	1589	U
41	a	1590	A
41	a	1608	A
41	a	1610	A
41	a	1613	G
41	a	1647	U
41	a	1648	U
41	a	1649	G
41	a	1651	G
41	a	1665	A
41	a	1674	G
41	a	1677	A
41	a	1703	G
41	a	1714	U
41	a	1715	G
41	a	1729	U
41	a	1730	C
41	a	1732	C
41	a	1738	G
41	a	1750	G

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Mol	Chain	Res	Type
41	a	1758	U
41	a	1764	C
41	a	1773	A
41	a	1791	A
41	a	1800	C
41	a	1808	A
41	a	1811	G
41	a	1816	C
41	a	1829	A
41	a	1833	C
41	a	1847	A
41	a	1848	A
41	a	1858	A
41	a	1859	U
41	a	1862	G
41	a	1864	U
41	a	1869	G
41	a	1870	C
41	a	1872	A
41	a	1873	G
41	a	1906	G
41	a	1907	G
41	a	1913	A
41	a	1914	C
41	a	1919	A
41	a	1920	C
41	a	1922	G
41	a	1923	U
41	a	1924	C
41	a	1925	C
41	a	1926	U
41	a	1929	G
41	a	1930	G
41	a	1936	A
41	a	1938	A
41	a	1955	U
41	a	1965	C
41	a	1967	C
41	a	1970	A
41	a	1971	U
41	a	1972	G
41	a	1991	U

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Mol	Chain	Res	Type
41	a	1992	G
41	a	1993	U
41	a	1997	C
41	a	2002	G
41	a	2020	A
41	a	2022	U
41	a	2023	C
41	a	2027	G
41	a	2033	A
41	a	2043	C
41	a	2051	A
41	a	2052	A
41	a	2055	C
41	a	2056	G
41	a	2060	A
41	a	2061	G
41	a	2062	A
41	a	2063	C
41	a	2097	A
41	a	2099	U
41	a	2100	G
41	a	2108	A
41	a	2110	G
41	a	2111	U
41	a	2113	U
41	a	2115	G
41	a	2116	G
41	a	2117	A
41	a	2118	U
41	a	2121	G
41	a	2122	U
41	a	2124	G
41	a	2125	G
41	a	2126	A
41	a	2127	G
41	a	2128	G
41	a	2131	U
41	a	2132	U
41	a	2133	G
41	a	2134	A
41	a	2139	U
41	a	2141	G

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Mol	Chain	Res	Type
41	a	2146	C
41	a	2147	A
41	a	2154	A
41	a	2157	G
41	a	2158	A
41	a	2159	G
41	a	2162	G
41	a	2163	A
41	a	2164	C
41	a	2165	C
41	a	2169	A
41	a	2171	A
41	a	2172	U
41	a	2178	C
41	a	2182	U
41	a	2183	A
41	a	2185	U
41	a	2189	U
41	a	2190	G
41	a	2193	G
41	a	2194	U
41	a	2198	A
41	a	2204	G
41	a	2211	A
41	a	2212	A
41	a	2225	A
41	a	2226	C
41	a	2229	U
41	a	2238	G
41	a	2239	G
41	a	2250	G
41	a	2268	A
41	a	2278	A
41	a	2283	C
41	a	2287	A
41	a	2288	A
41	a	2297	A
41	a	2305	U
41	a	2308	G
41	a	2309	A
41	a	2322	A
41	a	2325	G

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Mol	Chain	Res	Type
41	a	2327	A
41	a	2333	A
41	a	2335	A
41	a	2339	C
41	a	2345	G
41	a	2347	C
41	a	2350	C
41	a	2361	G
41	a	2372	U
41	a	2376	A
41	a	2383	G
41	a	2385	C
41	a	2402	U
41	a	2403	C
41	a	2406	A
41	a	2423	U
41	a	2424	C
41	a	2425	A
41	a	2426	A
41	a	2429	G
41	a	2430	A
41	a	2431	U
41	a	2434	A
41	a	2435	A
41	a	2441	U
41	a	2447	G
41	a	2448	A
41	a	2470	G
41	a	2474	U
41	a	2476	A
41	a	2478	A
41	a	2484	G
41	a	2491	U
41	a	2502	G
41	a	2506	U
41	a	2512	C
41	a	2513	A
41	a	2518	A
41	a	2520	C
41	a	2525	G
41	a	2529	G
41	a	2535	G

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Mol	Chain	Res	Type
41	a	2554	U
41	a	2566	A
41	a	2567	G
41	a	2573	C
41	a	2574	G
41	a	2585	U
41	a	2586	U
41	a	2602	A
41	a	2603	G
41	a	2609	U
41	a	2610	C
41	a	2611	C
41	a	2613	U
41	a	2629	U
41	a	2630	G
41	a	2663	G
41	a	2669	G
41	a	2671	G
41	a	2689	U
41	a	2690	U
41	a	2714	G
41	a	2716	C
41	a	2726	A
41	a	2744	G
41	a	2748	A
41	a	2758	A
41	a	2765	A
41	a	2777	G
41	a	2778	A
41	a	2791	G
41	a	2793	C
41	a	2796	U
41	a	2797	U
41	a	2798	U
41	a	2799	A
41	a	2801	G
41	a	2818	U
41	a	2820	A
41	a	2823	A
41	a	2825	G
41	a	2835	A
41	a	2849	U

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Mol	Chain	Res	Type
41	a	2859	G
41	a	2861	U
41	a	2867	G
41	a	2873	A
41	a	2874	C
41	a	2880	C
41	a	2883	A
41	a	2884	U
41	a	2885	G
41	a	2891	U
41	a	2902	C
44	d	2	G
44	d	13	G
44	d	16	G
44	d	17	C
44	d	35	C
44	d	45	A
44	d	51	G
44	d	56	G
44	d	57	A
44	d	66	A
44	d	88	C
44	d	89	U
44	d	90	C
44	d	99	A
44	d	109	A

All (39) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
8	7	4	U
8	7	7	U
8	7	10	U
10	A	6	G
10	A	7	G
10	A	9	G
10	A	12	G
10	A	21	A
10	A	22	G
10	A	57	A
10	A	60	U
10	B	6	G

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Mol	Chain	Res	Type
10	B	7	G
10	B	9	G
10	B	12	G
10	B	21	A
10	B	22	G
10	B	57	A
10	B	60	U
18	D	121	U
18	D	181	A
18	D	183	C
18	D	197	A
18	D	209	U
18	D	428	G
18	D	496	A
18	D	517	G
18	D	991	U
18	D	992	U
18	D	1145	A
18	D	1196	A
18	D	1211	U
18	D	1212	U
18	D	1213	A
18	D	1214	C
18	D	1447	A
18	D	1491	G
18	D	1492	A
18	D	1493	A

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

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

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

Of 3 ligands modelled in this entry, 3 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

There are no chain breaks in this entry.

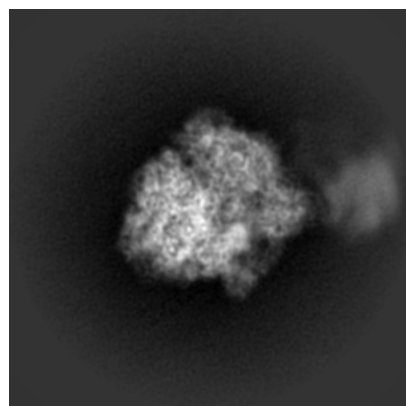
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-43335. These allow visual inspection of the internal detail of the map and identification of artifacts.

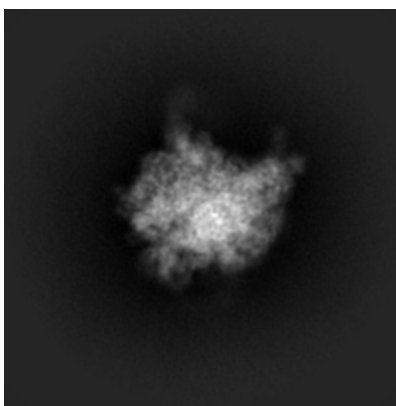
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections [i](#)

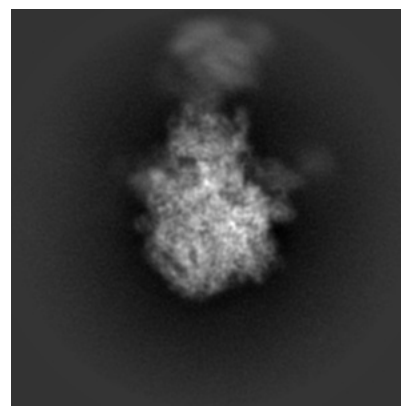
6.1.1 Primary map



X

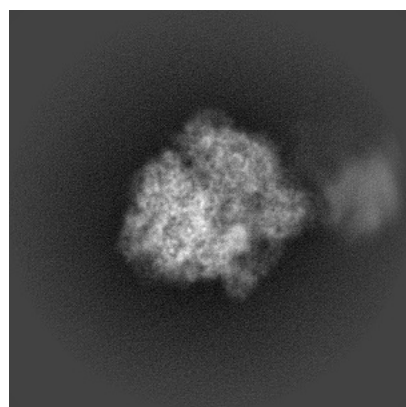


Y

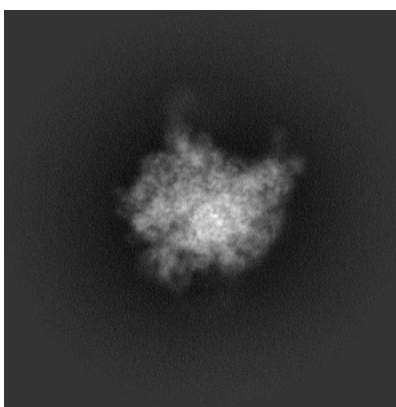


Z

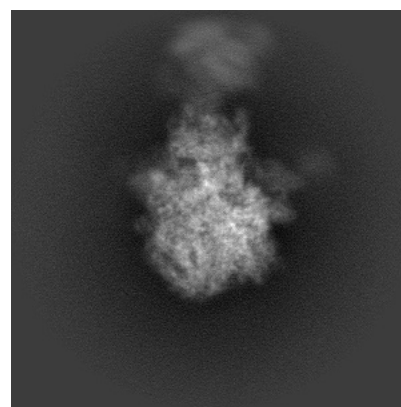
6.1.2 Raw map



X



Y

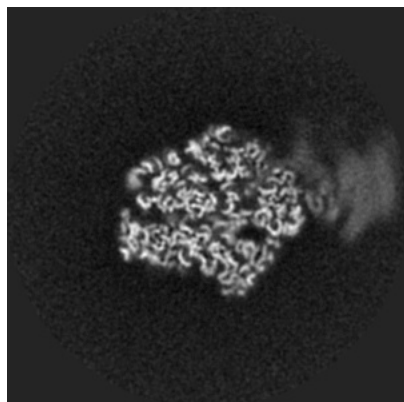


Z

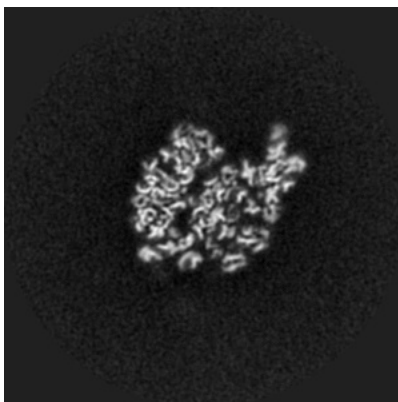
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

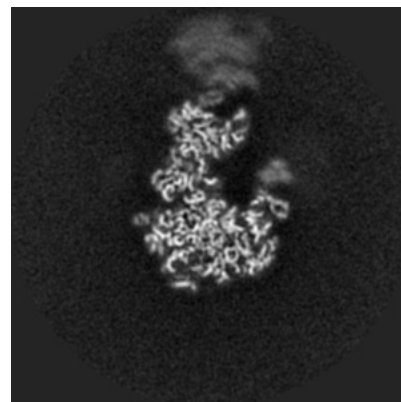
6.2.1 Primary map



X Index: 256

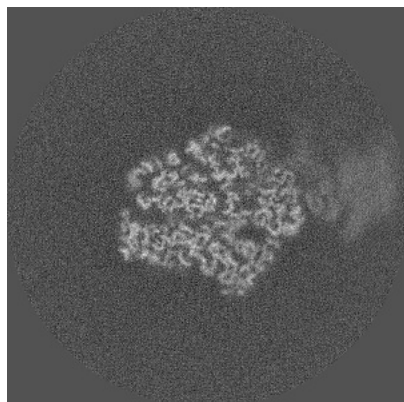


Y Index: 256

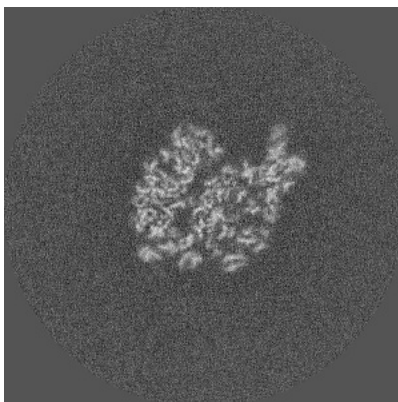


Z Index: 256

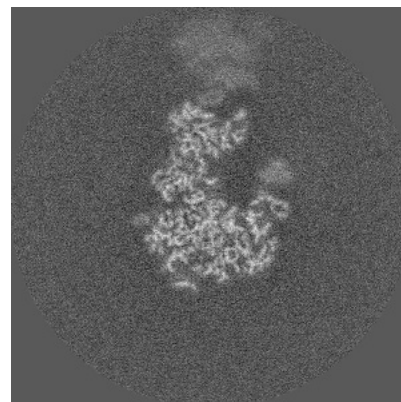
6.2.2 Raw map



X Index: 256



Y Index: 256

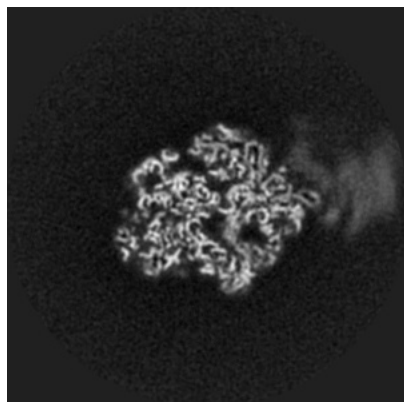


Z Index: 256

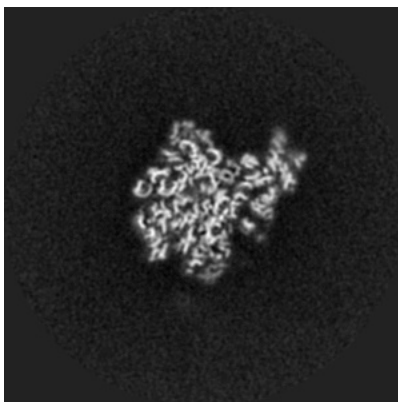
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

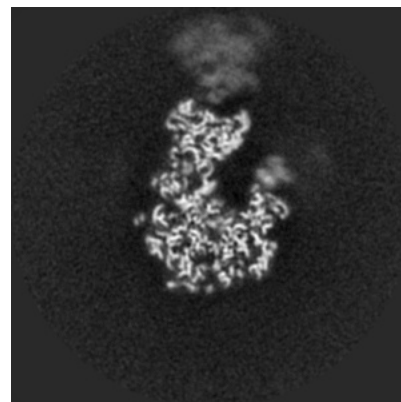
6.3.1 Primary map



X Index: 249

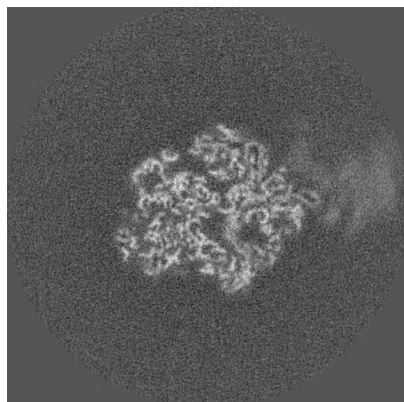


Y Index: 246

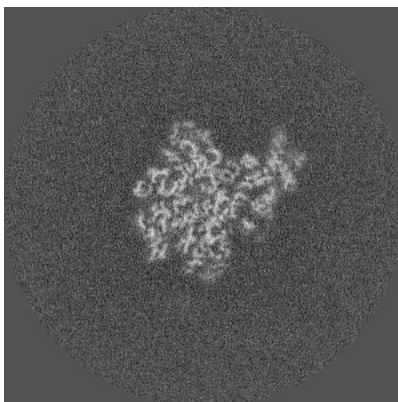


Z Index: 250

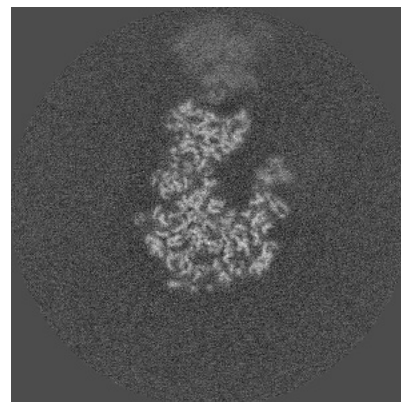
6.3.2 Raw map



X Index: 249



Y Index: 246



Z Index: 252

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

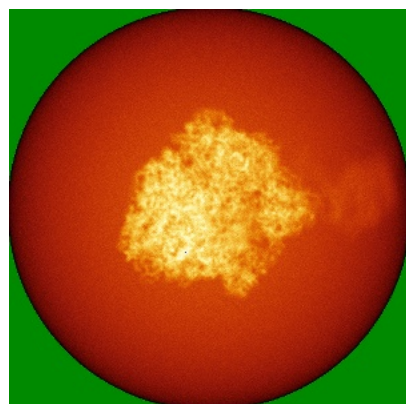


Y

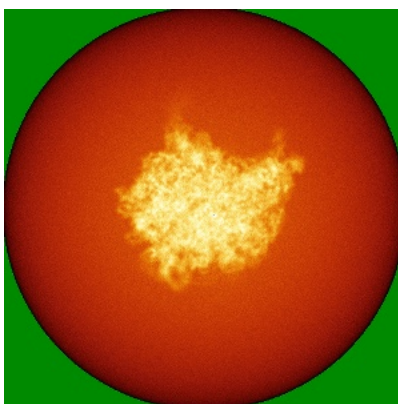


Z

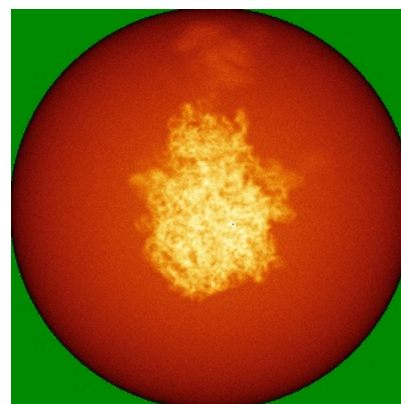
6.4.2 Raw map



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



X



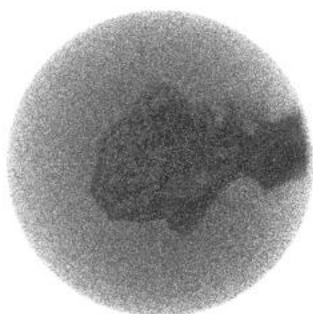
Y



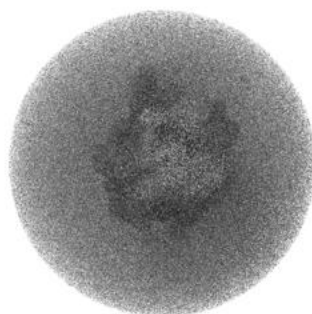
Z

The images above show the 3D surface view of the map at the recommended contour level 0.0032. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

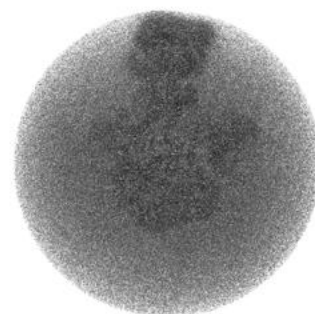
6.5.2 Raw map



X



Y



Z

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.

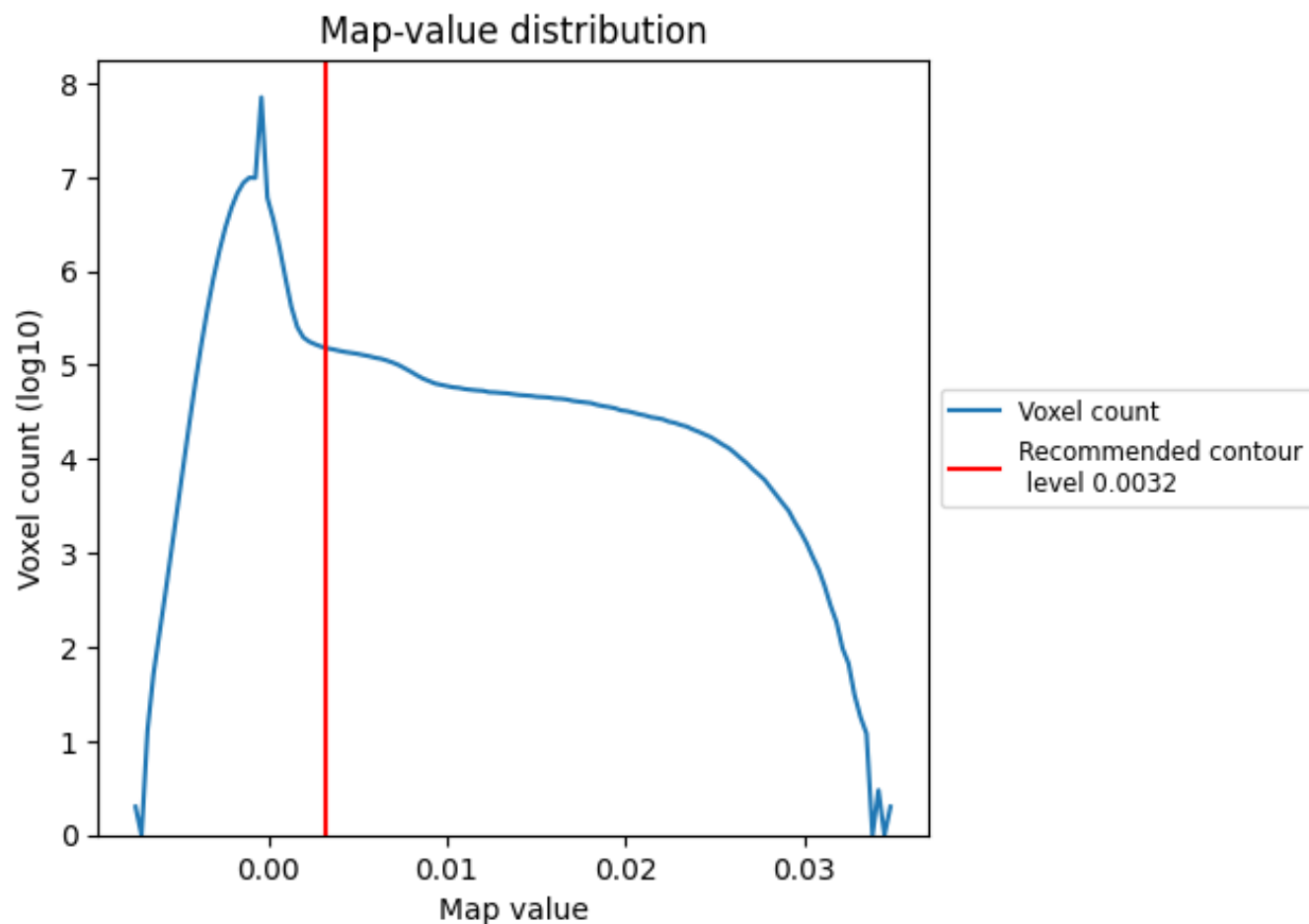
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

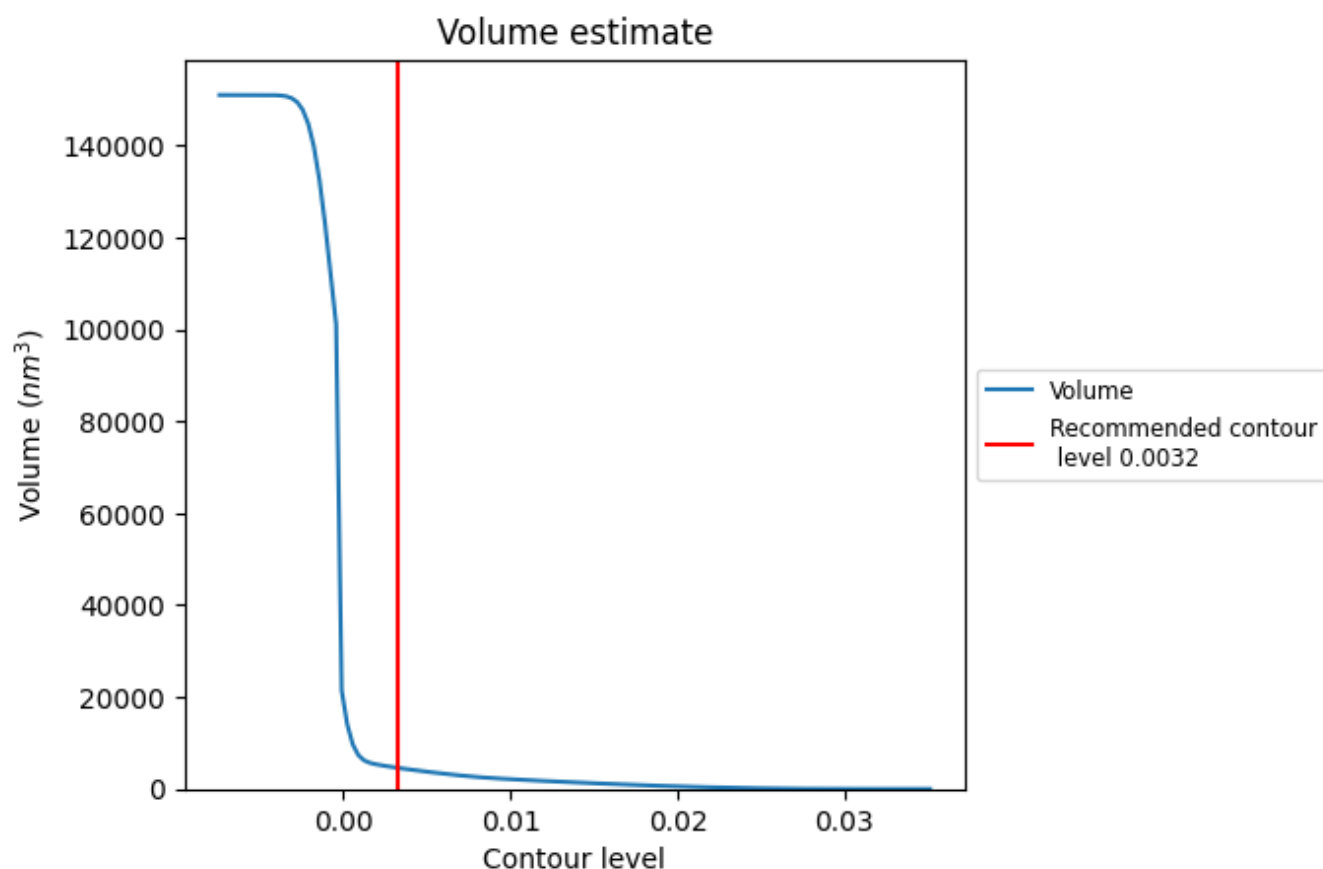
This section contains the results of statistical analysis of the map.

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.

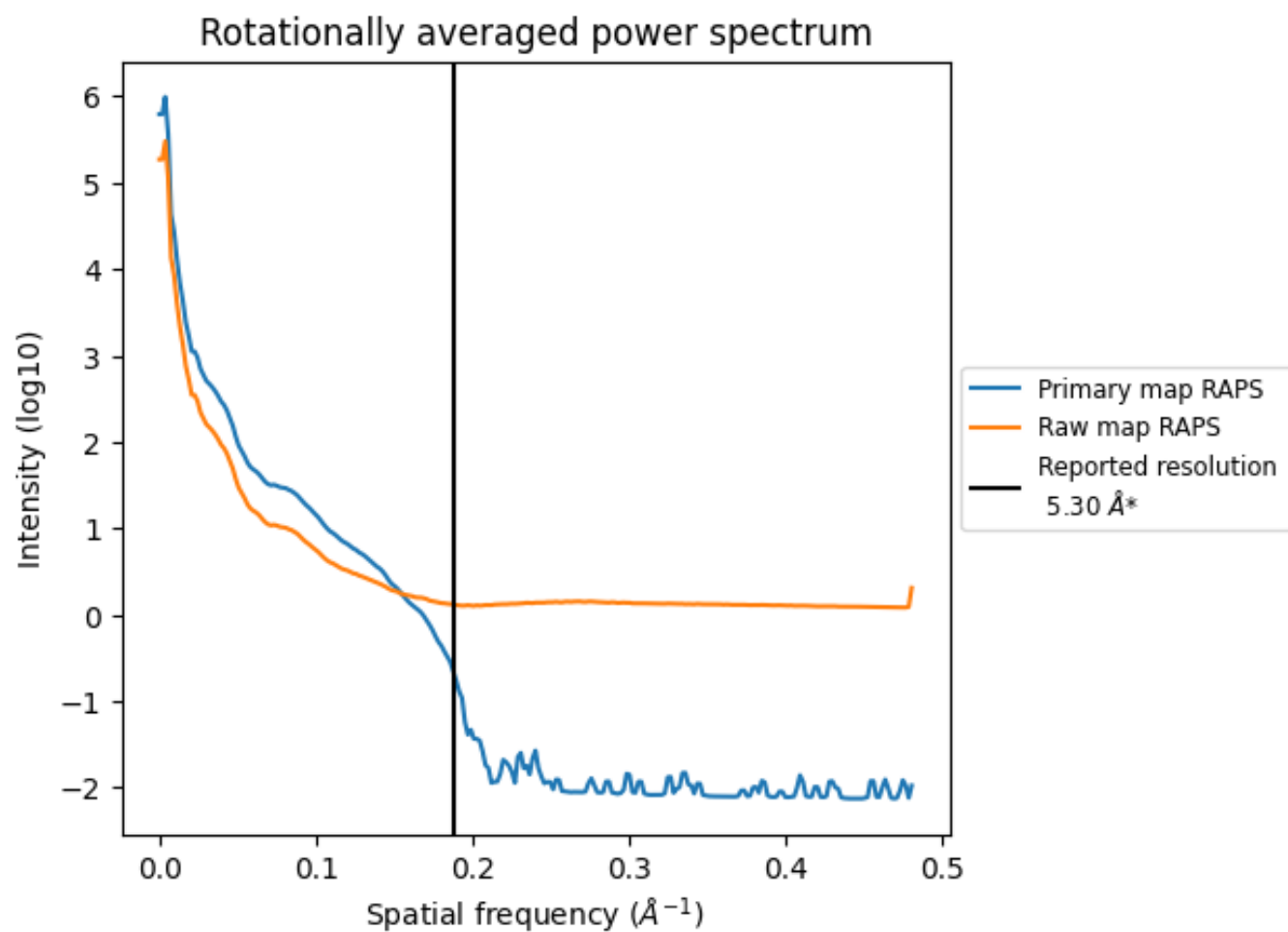
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 4617 nm^3 ; this corresponds to an approximate mass of 4171 kDa.

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

7.3 Rotationally averaged power spectrum [i](#)

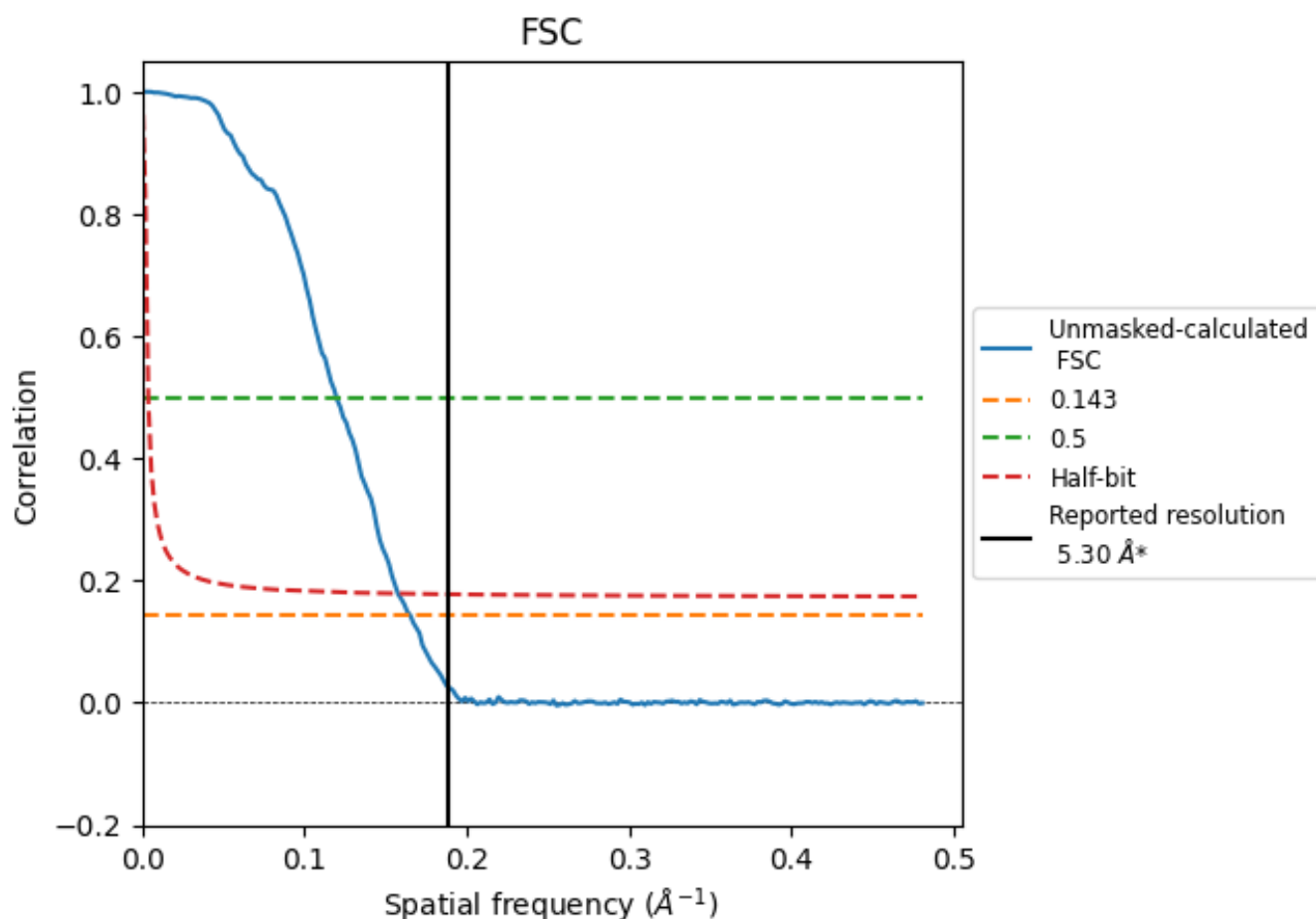


*Reported resolution corresponds to spatial frequency of 0.189 $\text{\AA}^{-1}$

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 0.189 $\text{\AA}^{-1}$

8.2 Resolution estimates [i](#)

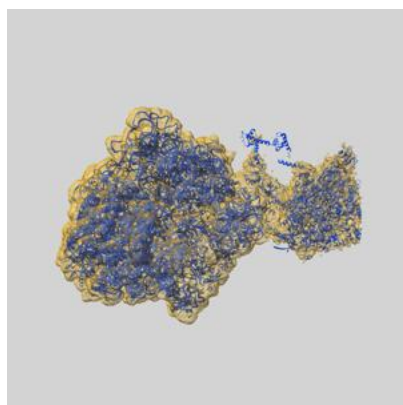
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	5.30	-	-
Author-provided FSC curve	-	-	-
Unmasked-calculated*	6.06	8.35	6.35

*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 6.06 differs from the reported value 5.3 by more than 10 %

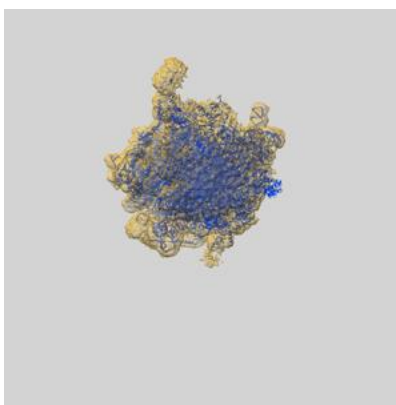
9 Map-model fit [i](#)

This section contains information regarding the fit between EMDB map EMD-43335 and PDB model 8VL1. Per-residue inclusion information can be found in section [3](#) on page [17](#).

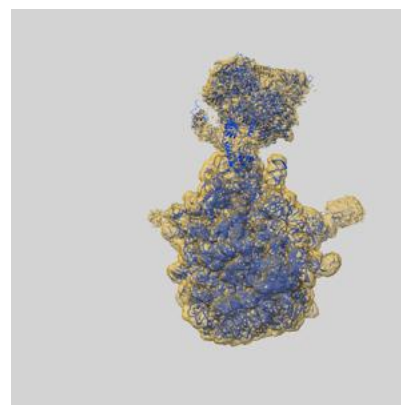
9.1 Map-model overlay [i](#)



X



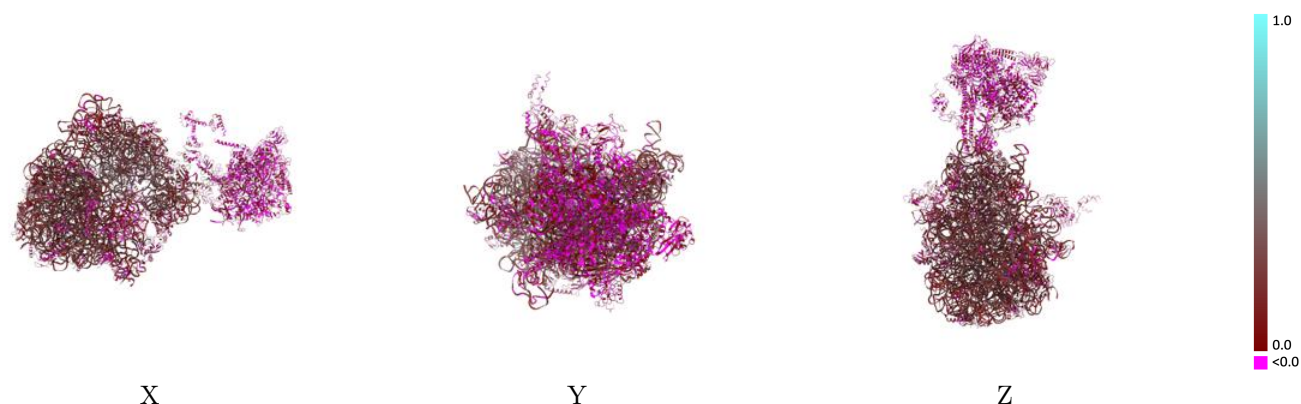
Y



Z

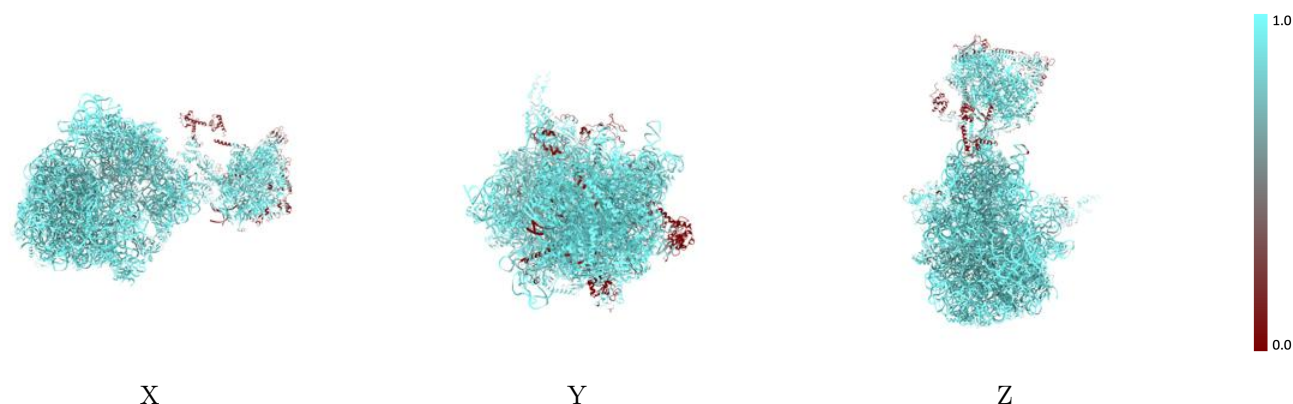
The images above show the 3D surface view of the map at the recommended contour level 0.0032 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



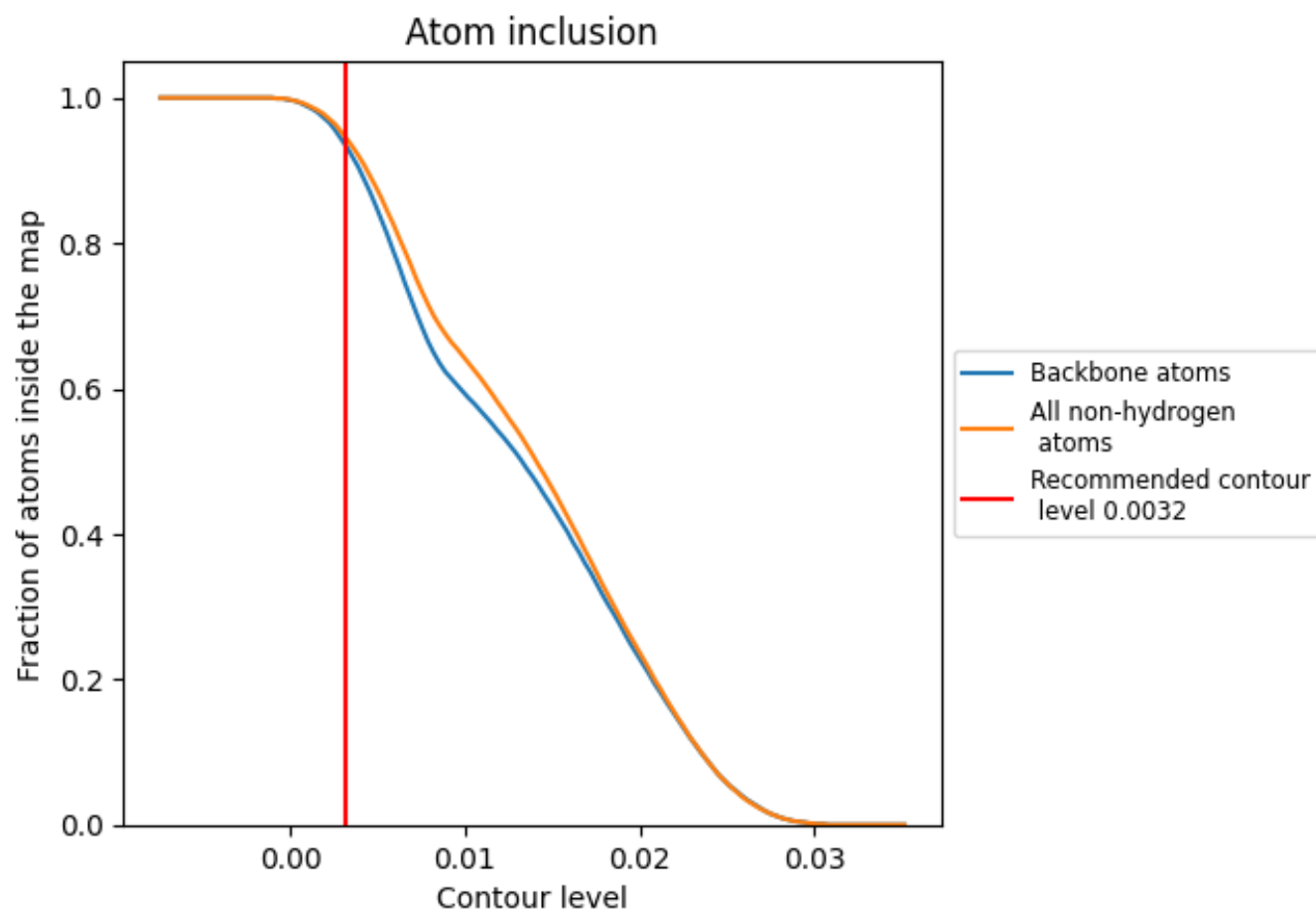
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions 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 (0.0032).

9.4 Atom inclusion [i](#)



At the recommended contour level, 93% of all backbone atoms, 94% 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 (0.0032) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.9450	 0.1690
0	 0.9860	 0.1640
1	 0.9800	 0.1890
2	 0.9740	 0.1670
3	 0.9850	 0.1450
4	 0.9820	 0.1480
5	 0.7960	 0.0610
6	 0.8010	 0.0410
7	 0.9830	 0.0840
9	 0.9400	 0.0720
A	 0.9990	 0.1890
AA	 0.8670	 0.0310
AB	 0.8960	 0.0830
AC	 0.9080	 0.0330
AD	 0.5300	 0.0270
AE	 0.8970	 0.0400
AF	 0.6910	 0.0140
AG	 0.5570	 0.0550
B	 0.9490	 0.1000
C	 0.9940	 0.1750
D	 0.9970	 0.2270
E	 0.9880	 0.1450
F	 0.9820	 0.1930
G	 0.9750	 0.1590
H	 0.8660	 0.0740
I	 0.9710	 0.1820
J	 0.9860	 0.1610
K	 0.9890	 0.2160
L	 0.9880	 0.1690
M	 0.9840	 0.1640
N	 0.9760	 0.1830
O	 0.9810	 0.1480
P	 0.9740	 0.1640
Q	 0.9870	 0.1760
R	 0.9910	 0.2180



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Chain	Atom inclusion	Q-score
S	 0.9830	 0.1590
T	 0.9860	 0.1650
U	 0.9860	 0.1450
V	 0.9780	 0.1870
W	 0.9290	 0.1100
X	 0.9680	 0.1290
Y	 0.8430	 0.0730
Z	 0.9340	 0.0500
a	 0.9980	 0.2190
b	 0.9750	 0.1500
c	 0.9820	 0.1810
d	 0.9980	 0.1840
e	 0.9900	 0.1330
f	 0.9860	 0.1770
g	 0.9770	 0.0970
h	 0.9860	 0.1680
i	 0.9740	 0.2050
j	 0.9810	 0.1510
k	 0.9900	 0.1420
l	 0.9740	 0.1510
m	 0.9890	 0.1910
n	 0.9690	 0.1210
o	 0.9820	 0.1500
p	 0.9840	 0.1350
q	 0.9830	 0.1490
r	 0.9170	 0.1190
s	 0.9780	 0.1690
t	 0.9590	 0.1590
u	 0.9870	 0.1610
v	 0.9870	 0.1810
w	 0.9770	 0.1650
x	 0.9700	 0.0960
y	 0.9760	 0.1460
z	 0.9870	 0.1550