



Full wwPDB EM Validation Report ⓘ

Nov 3, 2022 – 01:43 AM EDT

PDB ID : 5UYL
EMDB ID : EMD-8616
Title : 70S ribosome bound with cognate ternary complex base-paired to A site codon (Structure II)
Authors : Loveland, A.B.; Demo, G.; Grigorieff, N.; Korostelev, A.A.
Deposited on : 2017-02-24
Resolution : 3.60 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.dev43
Mogul : 1.8.5 (274361), CSD as541be (2020)
MolProbity : 4.02b-467
buster-report : 1.1.7 (2018)
Percentile statistics : 20191225.v01 (using entries in the PDB archive December 25th 2019)
MapQ : **FAILED**
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.31.2

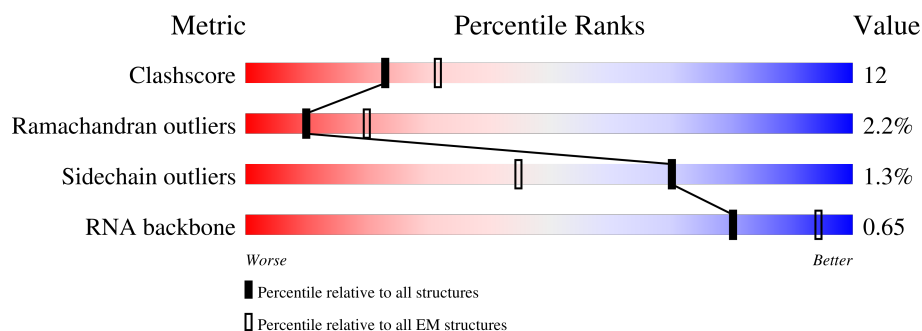
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 3.60 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)
Clashscore	158937	4297
Ramachandran outliers	154571	4023
Sidechain outliers	154315	3826
RNA backbone	4643	859

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 $\leq 5\%$.

Mol	Chain	Length	Quality of chain
1	04	271	68% 30% .
2	05	209	67% 33%
3	06	201	71% 27% .
4	07	177	61% 38% .
5	08	176	70% 28% .
6	09	149	61% 38% .
7	10	131	46% 49% 5%


























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Mol	Chain	Length	Quality of chain
8	11	141	 52% 45%
9	12	142	 73% 26%
10	13	122	 71% 27%
11	14	143	 69% 29%
12	15	136	 69% 29%
13	16	120	 68% 32%
14	17	116	 72% 28%
15	18	114	 68% 32%
16	19	117	 76% 22%
17	20	103	 62% 37%
18	21	110	 68% 31%
19	22	93	 69% 31%
20	23	102	 68% 29%
21	24	94	 65% 34%
22	25	75	 75% 25%
23	26	77	 58% 42%
24	27	63	 76% 22%
25	28	58	 74% 26%
26	29	66	 77% 23%
27	30	56	 61% 39%
28	31	50	 78% 22%
29	32	46	 76% 24%
30	33	64	 77% 22%
31	34	38	 74% 24%
32	B	218	 52% 45%

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Mol	Chain	Length	Quality of chain
33	C	206	
34	D	205	
35	E	157	
36	F	100	
37	G	151	
38	H	129	
39	I	127	
40	J	98	
41	K	116	
42	L	123	
43	M	114	
44	N	100	
45	O	88	
46	P	82	
47	Q	80	
48	R	65	
49	S	79	
50	T	85	
51	U	65	
52	03	234	
53	A	1539	
54	01	2903	
55	02	120	
56	W	77	
56	X	77	

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Mol	Chain	Length	Quality of chain
57	V	18	<div><div></div><div>72%</div><div>22%</div><div>6%</div></div>
58	Y	76	<div><div></div><div>43%</div><div>46%</div><div>11%</div></div>
59	Z	392	<div><div></div><div>33%</div><div>63%</div><div>..</div></div>

2 Entry composition

There are 62 unique types of molecules in this entry. The entry contains 153753 atoms, of which 0 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a protein called 50S ribosomal protein L2.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	04	271	Total	C	N	O	S	0	0
			2083	1288	423	365	7		

- Molecule 2 is a protein called 50S ribosomal protein L3.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	05	209	Total	C	N	O	S	0	0
			1565	979	288	294	4		

- Molecule 3 is a protein called 50S ribosomal protein L4.

Mol	Chain	Residues	Atoms					AltConf	Trace
3	06	201	Total	C	N	O	S	0	0
			1552	974	283	290	5		

- Molecule 4 is a protein called 50S ribosomal protein L5.

Mol	Chain	Residues	Atoms					AltConf	Trace
4	07	177	Total	C	N	O	S	0	0
			1411	899	249	257	6		

- Molecule 5 is a protein called 50S ribosomal protein L6.

Mol	Chain	Residues	Atoms					AltConf	Trace
5	08	176	Total	C	N	O	S	0	0
			1323	832	243	246	2		

- Molecule 6 is a protein called 50S ribosomal protein L9.

Mol	Chain	Residues	Atoms					AltConf	Trace
6	09	149	Total	C	N	O	S	0	0
			1111	699	197	214	1		

- Molecule 7 is a protein called 50S ribosomal protein L10.

Mol	Chain	Residues	Atoms					AltConf	Trace
7	10	131	Total	C	N	O	S	0	0
			989	625	175	184	5		

- Molecule 8 is a protein called 50S ribosomal protein L11.

Mol	Chain	Residues	Atoms					AltConf	Trace
8	11	141	Total	C	N	O	S	0	0
			1032	651	179	196	6		

- Molecule 9 is a protein called 50S ribosomal protein L13.

Mol	Chain	Residues	Atoms					AltConf	Trace
9	12	142	Total	C	N	O	S	0	0
			1129	714	212	199	4		

- Molecule 10 is a protein called 50S ribosomal protein L14.

Mol	Chain	Residues	Atoms					AltConf	Trace
10	13	122	Total	C	N	O	S	0	0
			939	587	180	166	6		

- Molecule 11 is a protein called 50S ribosomal protein L15.

Mol	Chain	Residues	Atoms					AltConf	Trace
11	14	143	Total	C	N	O	S	0	0
			1045	649	206	189	1		

- Molecule 12 is a protein called 50S ribosomal protein L16.

Mol	Chain	Residues	Atoms					AltConf	Trace
12	15	136	Total	C	N	O	S	0	0
			1074	686	205	177	6		

- Molecule 13 is a protein called 50S ribosomal protein L17.

Mol	Chain	Residues	Atoms					AltConf	Trace
13	16	120	Total	C	N	O	S	0	0
			961	593	196	167	5		

- Molecule 14 is a protein called 50S ribosomal protein L18.

Mol	Chain	Residues	Atoms				AltConf	Trace
14	17	116	Total	C	N	O	0	0
			892	552	178	162		

- Molecule 15 is a protein called 50S ribosomal protein L19.

Mol	Chain	Residues	Atoms					AltConf	Trace
15	18	114	Total	C	N	O	S	0	0
			917	574	179	163	1		

- Molecule 16 is a protein called 50S ribosomal protein L20.

Mol	Chain	Residues	Atoms				AltConf	Trace
16	19	117	Total	C	N	O	0	0
			947	604	192	151		

- Molecule 17 is a protein called 50S ribosomal protein L21.

Mol	Chain	Residues	Atoms					AltConf	Trace
17	20	103	Total	C	N	O	S	0	0
			816	516	153	145	2		

- Molecule 18 is a protein called 50S ribosomal protein L22.

Mol	Chain	Residues	Atoms					AltConf	Trace
18	21	110	Total	C	N	O	S	0	0
			857	532	166	156	3		

- Molecule 19 is a protein called 50S ribosomal protein L23.

Mol	Chain	Residues	Atoms					AltConf	Trace
19	22	93	Total	C	N	O	S	0	0
			739	466	139	132	2		

- Molecule 20 is a protein called 50S ribosomal protein L24.

Mol	Chain	Residues	Atoms				AltConf	Trace
20	23	102	Total	C	N	O	0	0
			780	492	146	142		

- Molecule 21 is a protein called 50S ribosomal protein L25.

Mol	Chain	Residues	Atoms					AltConf	Trace
21	24	94	Total	C	N	O	S	0	0
			753	479	137	134	3		

- Molecule 22 is a protein called 50S ribosomal protein L27.

Mol	Chain	Residues	Atoms					AltConf	Trace
22	25	75	Total	C	N	O	S	0	0
			575	356	116	102	1		

- Molecule 23 is a protein called 50S ribosomal protein L28.

Mol	Chain	Residues	Atoms					AltConf	Trace
23	26	77	Total	C	N	O	S	0	0
			625	388	129	106	2		

- Molecule 24 is a protein called 50S ribosomal protein L29.

Mol	Chain	Residues	Atoms					AltConf	Trace
24	27	63	Total	C	N	O	S	0	0
			509	313	99	95	2		

- Molecule 25 is a protein called 50S ribosomal protein L30.

Mol	Chain	Residues	Atoms					AltConf	Trace
25	28	58	Total	C	N	O	S	0	0
			449	281	87	79	2		

- Molecule 26 is a protein called 50S ribosomal protein L31.

Mol	Chain	Residues	Atoms					AltConf	Trace
26	29	66	Total	C	N	O	S	0	0
			523	323	99	95	6		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
27	30	56	Total	C	N	O	S	0	0
			444	269	94	80	1		

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

Mol	Chain	Residues	Atoms				AltConf	Trace
28	31	50	Total	C	N	O	0	0
			410	263	75	72		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
29	32	46	Total	C	N	O	S	0	0
			377	228	90	57	2		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
30	33	64	Total	C	N	O	S	0	0
			504	323	105	74	2		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
31	34	38	Total	C	N	O	S	0	0
			302	185	65	48	4		

- Molecule 32 is a protein called 30S ribosomal protein S2.

Mol	Chain	Residues	Atoms					AltConf	Trace
32	B	218	Total	C	N	O	S	0	0
			1705	1081	305	312	7		

- Molecule 33 is a protein called 30S ribosomal protein S3.

Mol	Chain	Residues	Atoms					AltConf	Trace
33	C	206	Total	C	N	O	S	0	0
			1625	1028	305	289	3		

- Molecule 34 is a protein called 30S ribosomal protein S4.

Mol	Chain	Residues	Atoms					AltConf	Trace
34	D	205	Total	C	N	O	S	0	0
			1643	1026	315	298	4		

- Molecule 35 is a protein called 30S ribosomal protein S5.

Mol	Chain	Residues	Atoms					AltConf	Trace
35	E	157	Total	C	N	O	S	0	0
			1157	719	218	214	6		

- Molecule 36 is a protein called 30S ribosomal protein S6.

Mol	Chain	Residues	Atoms					AltConf	Trace
36	F	100	Total	C	N	O	S	0	0
			818	515	148	149	6		

- Molecule 37 is a protein called 30S ribosomal protein S7.

Mol	Chain	Residues	Atoms					AltConf	Trace
37	G	151	Total	C	N	O	S	0	0
			1182	735	227	216	4		

- Molecule 38 is a protein called 30S ribosomal protein S8.

Mol	Chain	Residues	Atoms					AltConf	Trace
38	H	129	Total	C	N	O	S	0	0
			979	616	173	184	6		

- Molecule 39 is a protein called 30S ribosomal protein S9.

Mol	Chain	Residues	Atoms					AltConf	Trace
39	I	127	Total	C	N	O	S	0	0
			1022	634	206	179	3		

- Molecule 40 is a protein called 30S ribosomal protein S10.

Mol	Chain	Residues	Atoms					AltConf	Trace
40	J	98	Total	C	N	O	S	0	0
			787	493	150	143	1		

- Molecule 41 is a protein called 30S ribosomal protein S11.

Mol	Chain	Residues	Atoms					AltConf	Trace
41	K	116	Total	C	N	O	S	0	0
			870	535	173	159	3		

- Molecule 42 is a protein called 30S ribosomal protein S12.

Mol	Chain	Residues	Atoms					AltConf	Trace
42	L	123	Total	C	N	O	S	0	0
			955	590	196	165	4		

- Molecule 43 is a protein called 30S ribosomal protein S13.

Mol	Chain	Residues	Atoms					AltConf	Trace
43	M	114	Total	C	N	O	S	0	0
			884	546	178	157	3		

- Molecule 44 is a protein called 30S ribosomal protein S14.

Mol	Chain	Residues	Atoms					AltConf	Trace
44	N	100	Total	C	N	O	S	0	0
			805	499	164	139	3		

- Molecule 45 is a protein called 30S ribosomal protein S15.

Mol	Chain	Residues	Atoms					AltConf	Trace
45	O	88	Total	C	N	O	S	0	0
			714	439	144	130	1		

- Molecule 46 is a protein called 30S ribosomal protein S16.

Mol	Chain	Residues	Atoms					AltConf	Trace
46	P	82	Total	C	N	O	S	0	0
			649	406	128	114	1		

- Molecule 47 is a protein called 30S ribosomal protein S17.

Mol	Chain	Residues	Atoms					AltConf	Trace
47	Q	80	Total	C	N	O	S	0	0
			649	411	121	114	3		

- Molecule 48 is a protein called 30S ribosomal protein S18.

Mol	Chain	Residues	Atoms					AltConf	Trace
48	R	65	Total	C	N	O	S	0	0
			536	339	100	96	1		

- Molecule 49 is a protein called 30S ribosomal protein S19.

Mol	Chain	Residues	Atoms					AltConf	Trace
49	S	79	Total	C	N	O	S	0	0
			638	408	120	108	2		

- Molecule 50 is a protein called 30S ribosomal protein S20.

Mol	Chain	Residues	Atoms					AltConf	Trace
50	T	85	Total	C	N	O	S	0	0
			665	411	137	114	3		

- Molecule 51 is a protein called 30S ribosomal protein S21.

Mol	Chain	Residues	Atoms					AltConf	Trace
51	U	65	Total	C	N	O	S	0	0
			545	335	117	92	1		

- Molecule 52 is a protein called 50S ribosomal protein L1.

Mol	Chain	Residues	Atoms					AltConf	Trace
52	03	134	Total	C	N	O	S	0	0
			1026	645	186	193	2		

- Molecule 53 is a RNA chain called 16S ribosomal RNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
53	A	1539	Total	C	N	O	P	0	0
			33012	14725	6052	10697	1538		

- Molecule 54 is a RNA chain called 23S ribosomal RNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
54	01	2903	Total	C	N	O	P	0	0
			62317	27801	11468	20146	2902		

There is a discrepancy between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
01	747	C	U	conflict	GB 802133627

- Molecule 55 is a RNA chain called 5S ribosomal RNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
55	02	120	Total	C	N	O	P	0	0
			2568	1145	471	833	119		

- Molecule 56 is a RNA chain called tRNAfMet.

Mol	Chain	Residues	Atoms					AltConf	Trace
56	X	77	Total	C	N	O	P	0	0
			1640	732	297	535	76		
56	W	77	Total	C	N	O	P	0	0
			1640	732	297	535	76		

- Molecule 57 is a RNA chain called mRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
57	V	18	Total	C	N	O	P	0	0
			388	175	76	120	17		

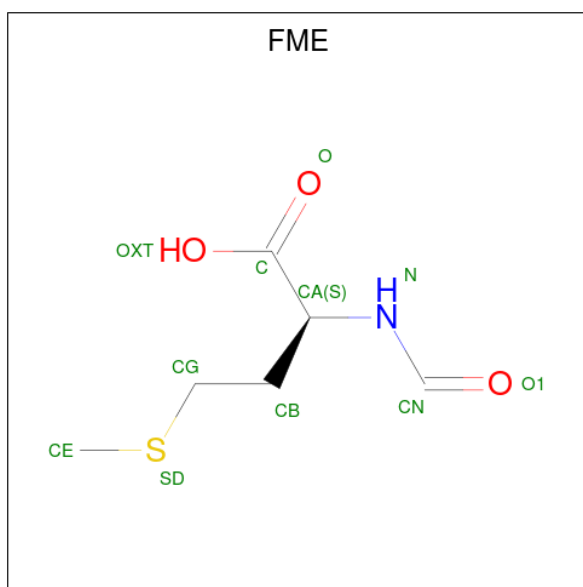
- Molecule 58 is a RNA chain called tRNAPhe.

Mol	Chain	Residues	Atoms					AltConf	Trace
58	Y	76	Total	C	N	O	P	0	0
			1619	723	290	531	75		

- Molecule 59 is a protein called Elongation factor Tu 2.

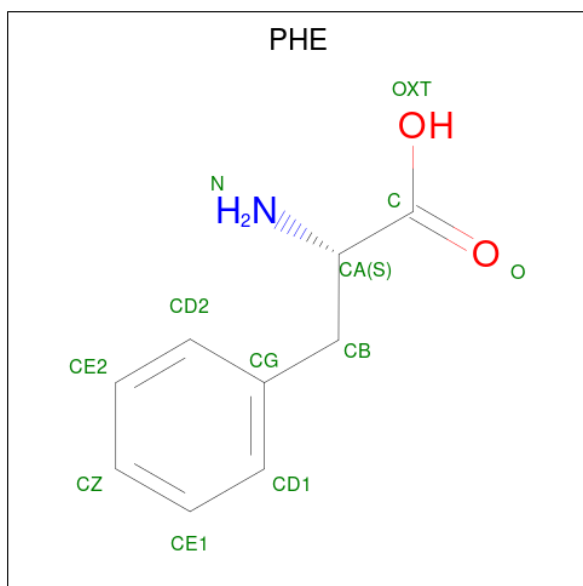
Mol	Chain	Residues	Atoms					AltConf	Trace
59	Z	392	Total	C	N	O	S	0	0
			3029	1915	521	580	13		

- Molecule 60 is N-FORMYLMETHIONINE (three-letter code: FME) (formula: C₆H₁₁NO₃S).



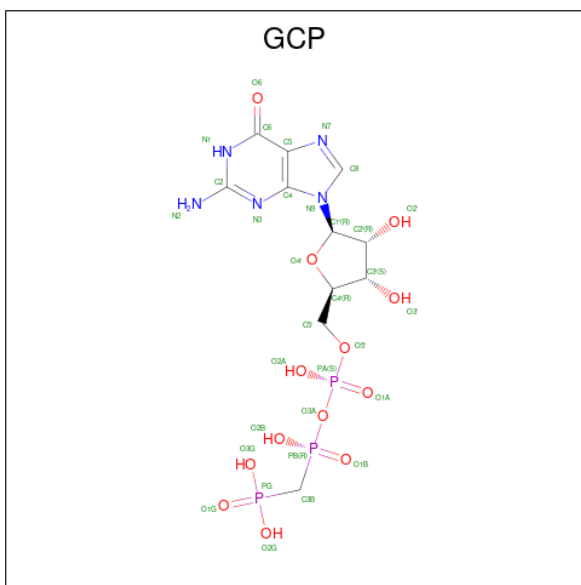
Mol	Chain	Residues	Atoms					AltConf
60	W	1	Total	C	N	O	S	0
			10	6	1	2	1	

- Molecule 61 is PHENYLALANINE (three-letter code: PHE) (formula: $C_9H_{11}NO_2$).



Mol	Chain	Residues	Atoms					AltConf
61	Y	1	Total	C	N	O		0
			11	9	1	1		

- Molecule 62 is PHOSPHOMETHYLPHOSPHONIC ACID GUANYLATE ESTER (three-letter code: GCP) (formula: $C_{11}H_{18}N_5O_{13}P_3$).

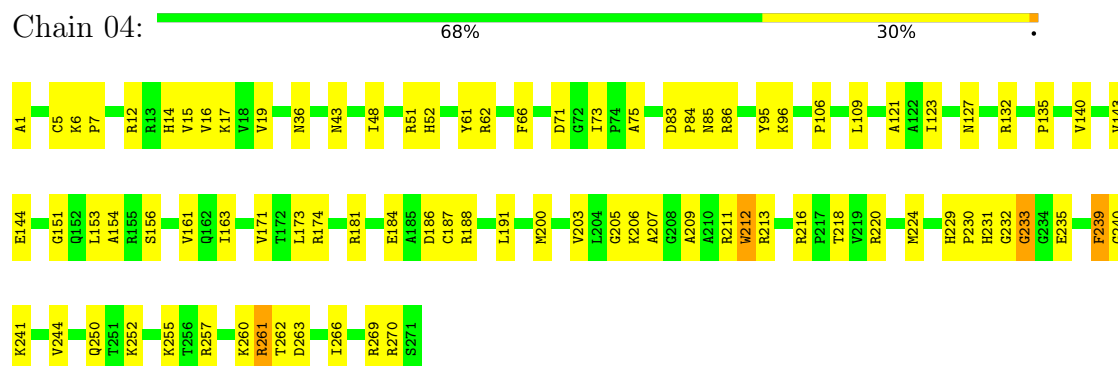


Mol	Chain	Residues	Atoms					AltConf
62	Z	1	Total	C	N	O	P	0
			32	11	5	13	3	

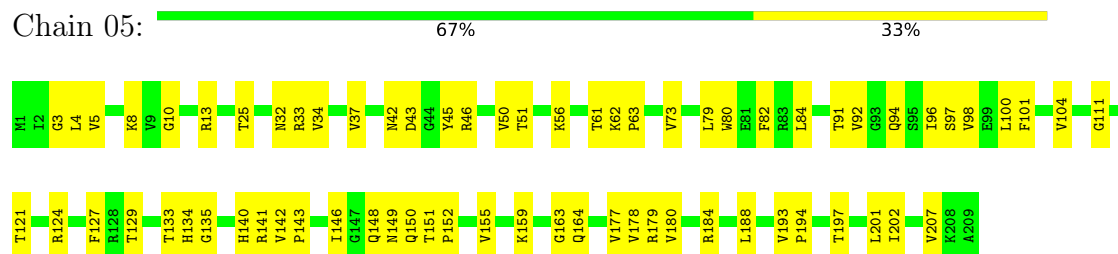
3 Residue-property plots [i](#)

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. 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. 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.

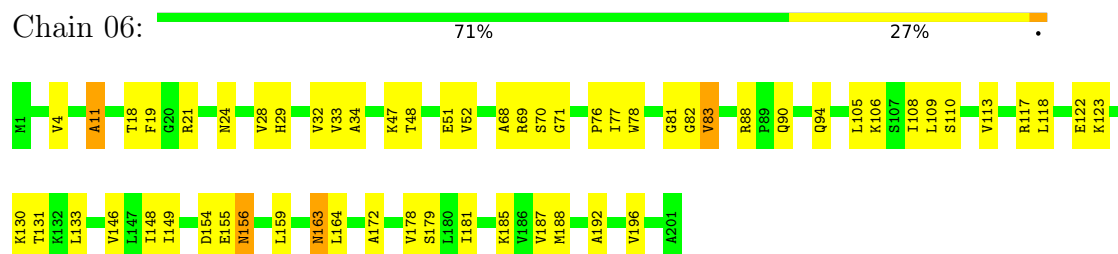
- Molecule 1: 50S ribosomal protein L2



- Molecule 2: 50S ribosomal protein L3

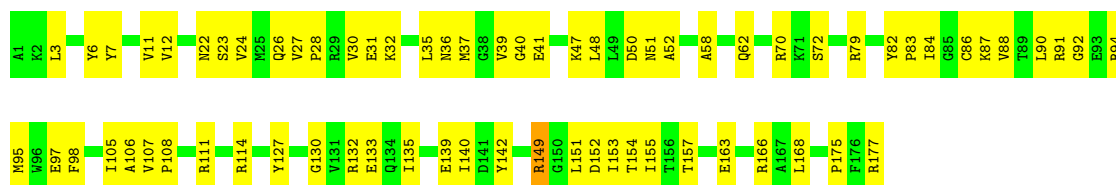


- Molecule 3: 50S ribosomal protein L4



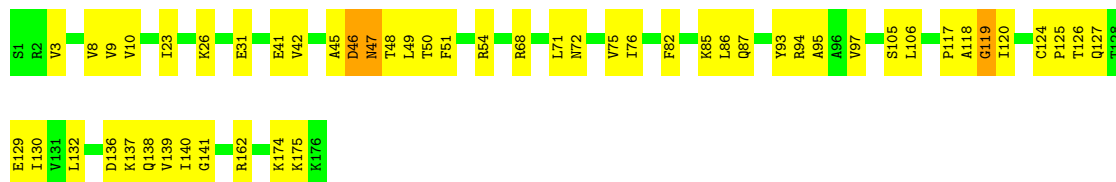
- Molecule 4: 50S ribosomal protein L5





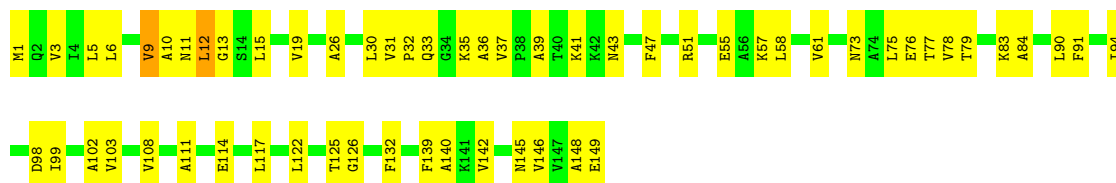
• Molecule 5: 50S ribosomal protein L6

Chain 08: 70% 28% .



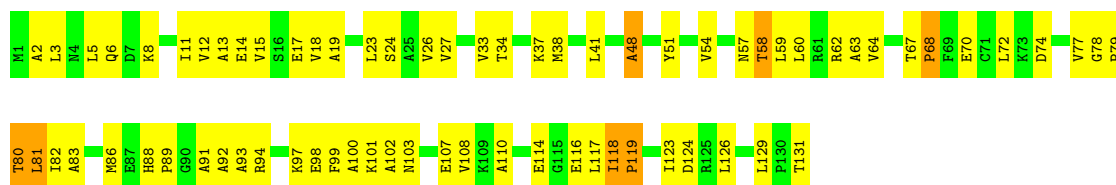
• Molecule 6: 50S ribosomal protein L9

Chain 09: 61% 38% .



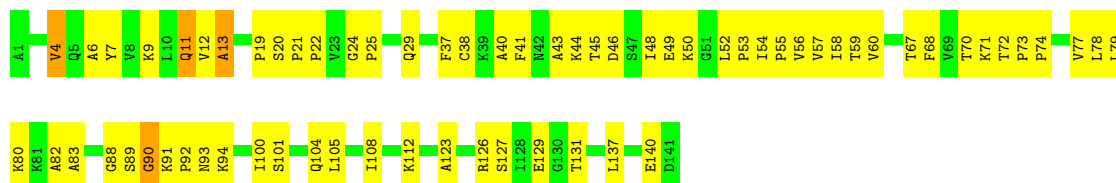
• Molecule 7: 50S ribosomal protein L10

Chain 10: 46% 49% 5% .



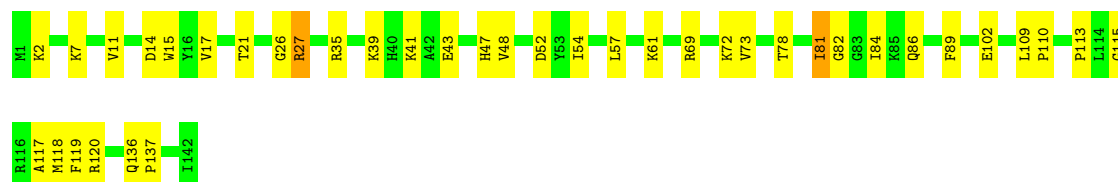
• Molecule 8: 50S ribosomal protein L11

Chain 11: 52% 45% .



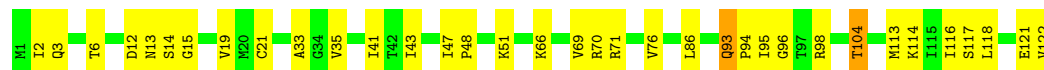
• Molecule 9: 50S ribosomal protein L13

Chain 12: 73% 26% .



- Molecule 10: 50S ribosomal protein L14

Chain 13: 71% 27% .



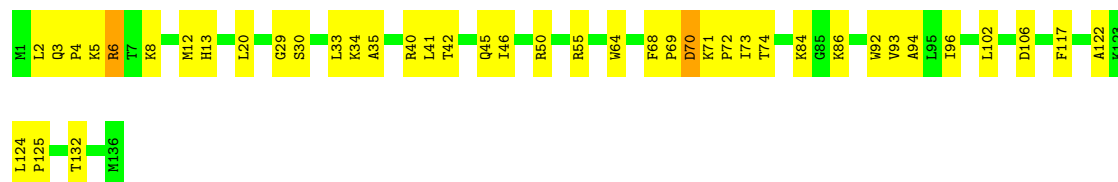
- Molecule 11: 50S ribosomal protein L15

Chain 14: 69% 29% .



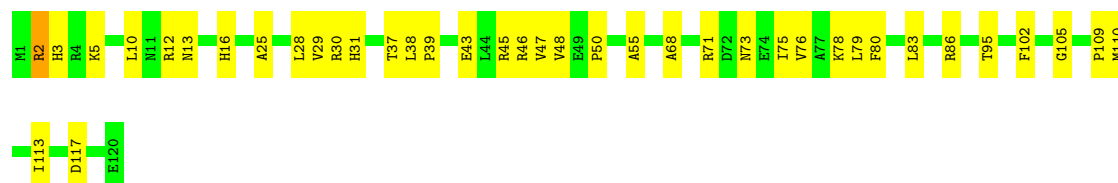
- Molecule 12: 50S ribosomal protein L16

Chain 15: 69% 29% .



- Molecule 13: 50S ribosomal protein L17

Chain 16: 68% 32% .



- Molecule 14: 50S ribosomal protein L18

Chain 17: 72% 28%




- Molecule 15: 50S ribosomal protein L19

Chain 18:  68% 32%



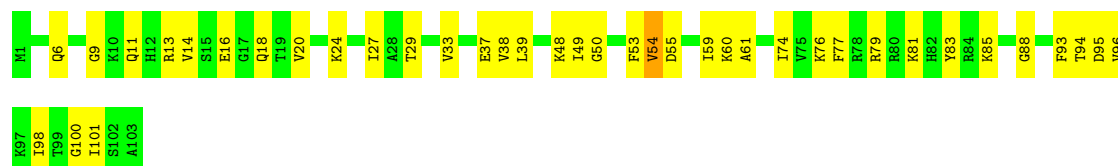
- Molecule 16: 50S ribosomal protein L20

Chain 19:  76% 22% .



- Molecule 17: 50S ribosomal protein L21

Chain 20:  62% 37% .



- Molecule 18: 50S ribosomal protein L22

Chain 21:  68% 31% .



- Molecule 19: 50S ribosomal protein L23

Chain 22:  69% 31%



- Molecule 20: 50S ribosomal protein L24

Chain 23:  68% 29% .



- Molecule 21: 50S ribosomal protein L25

Chain 24:  65% 34% .



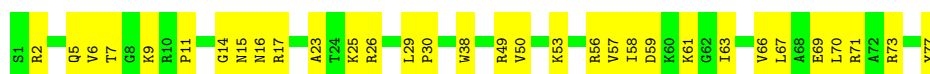
- Molecule 22: 50S ribosomal protein L27

Chain 25: 75% 25%



- Molecule 23: 50S ribosomal protein L28

Chain 26: 58% 42%



- Molecule 24: 50S ribosomal protein L29

Chain 27: 76% 22%



- Molecule 25: 50S ribosomal protein L30

Chain 28: 74% 26%



- Molecule 26: 50S ribosomal protein L31

Chain 29: 77% 23%



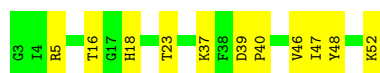
- Molecule 27: 50S ribosomal protein L32

Chain 30: 61% 39%



- Molecule 28: 50S ribosomal protein L33

Chain 31: 78% 22%



- Molecule 29: 50S ribosomal protein L34

Chain 32: 76% 24%



- Molecule 30: 50S ribosomal protein L35

Chain 33: 77% 22%



- Molecule 31: 50S ribosomal protein L36

Chain 34: 74% 24%



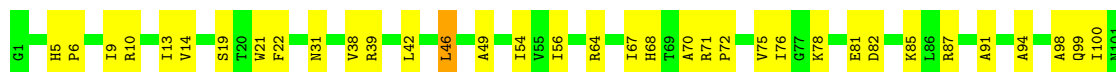
- Molecule 32: 30S ribosomal protein S2

Chain B: 52% 45%

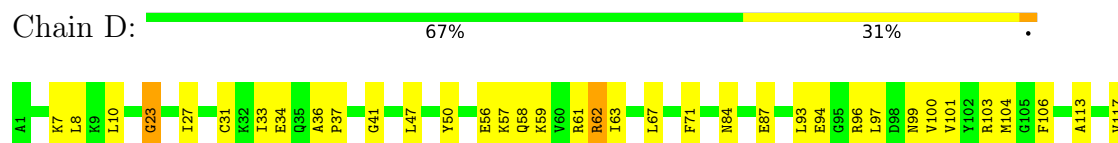


- Molecule 33: 30S ribosomal protein S3

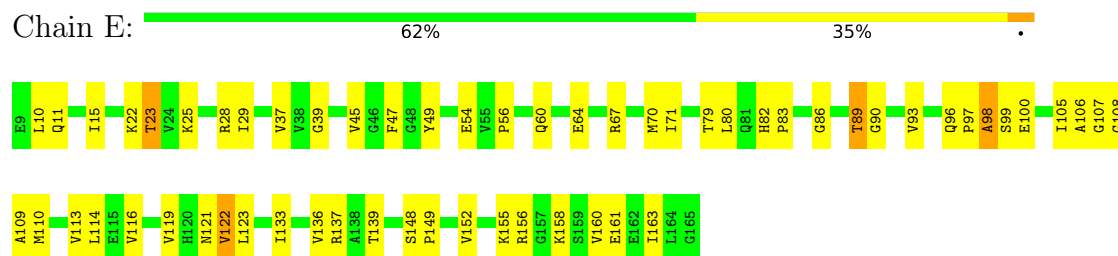
Chain C: 67% 31%



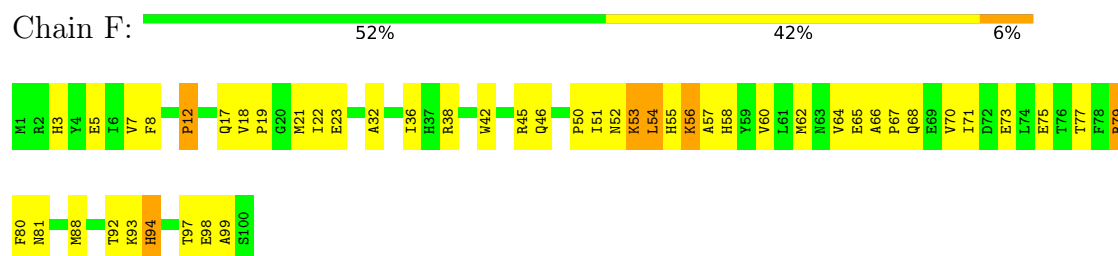
- Molecule 34: 30S ribosomal protein S4



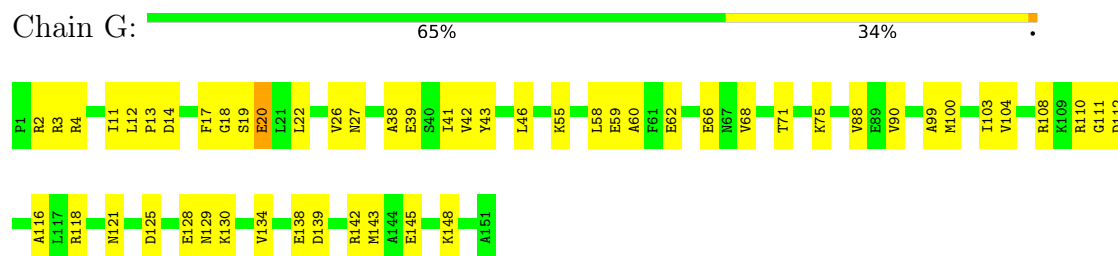
- Molecule 35: 30S ribosomal protein S5



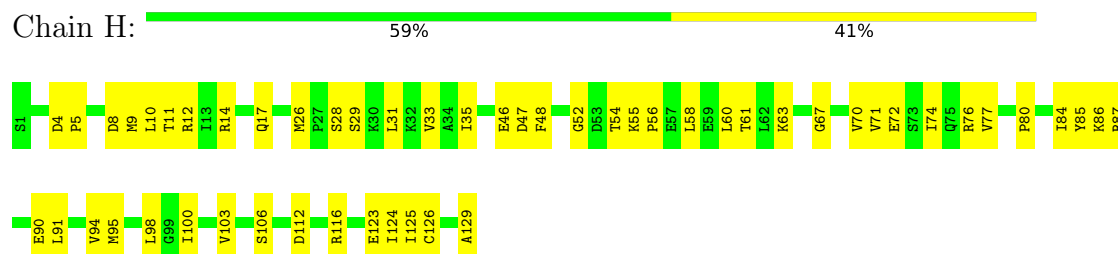
- Molecule 36: 30S ribosomal protein S6



- Molecule 37: 30S ribosomal protein S7

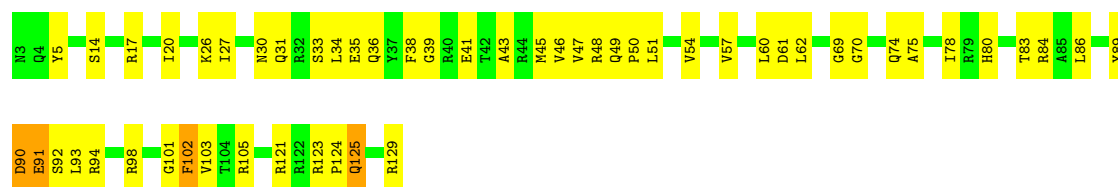


- Molecule 38: 30S ribosomal protein S8



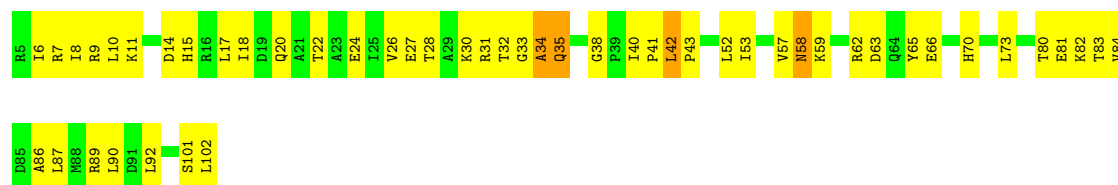
- Molecule 39: 30S ribosomal protein S9

Chain I:  58% 39% .



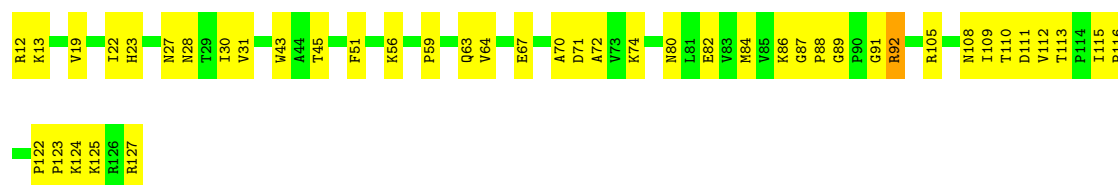
- Molecule 40: 30S ribosomal protein S10

Chain J:  49% 47% .



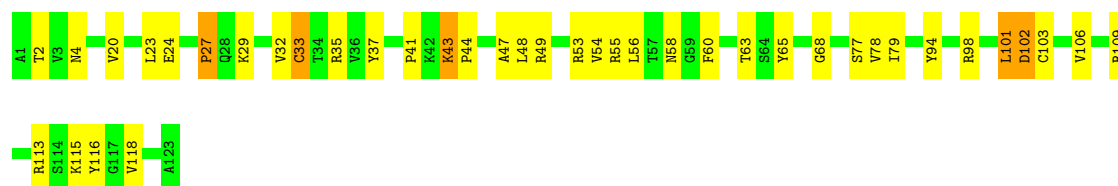
- Molecule 41: 30S ribosomal protein S11

Chain K:  62% 37% .



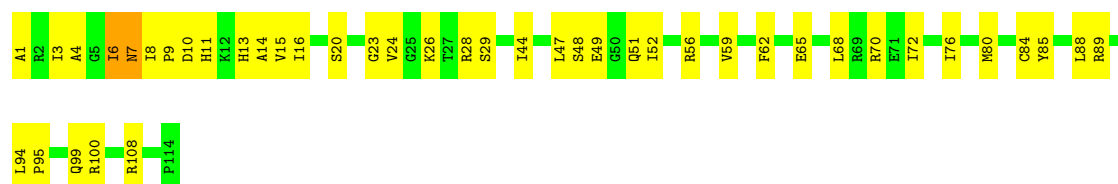
- Molecule 42: 30S ribosomal protein S12

Chain L:  67% 28% .



- Molecule 43: 30S ribosomal protein S13

Chain M:  62% 36% .



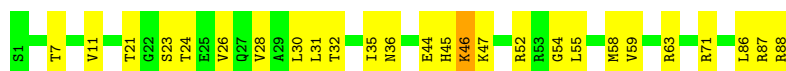
- Molecule 44: 30S ribosomal protein S14

Chain N:  61% 37%



- Molecule 45: 30S ribosomal protein S15

Chain O:  70% 28%



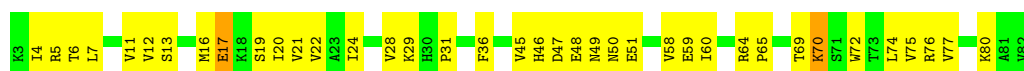
- Molecule 46: 30S ribosomal protein S16

Chain P:  62% 35%



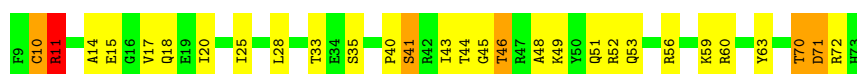
- Molecule 47: 30S ribosomal protein S17

Chain Q:  52% 45%



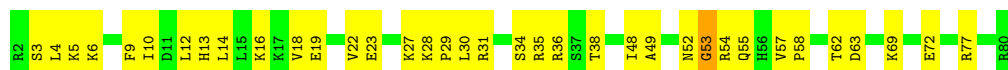
- Molecule 48: 30S ribosomal protein S18

Chain R:  55% 35% 8%



- Molecule 49: 30S ribosomal protein S19

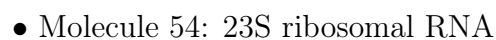
Chain S:  54% 44%



- Molecule 50: 30S ribosomal protein S20

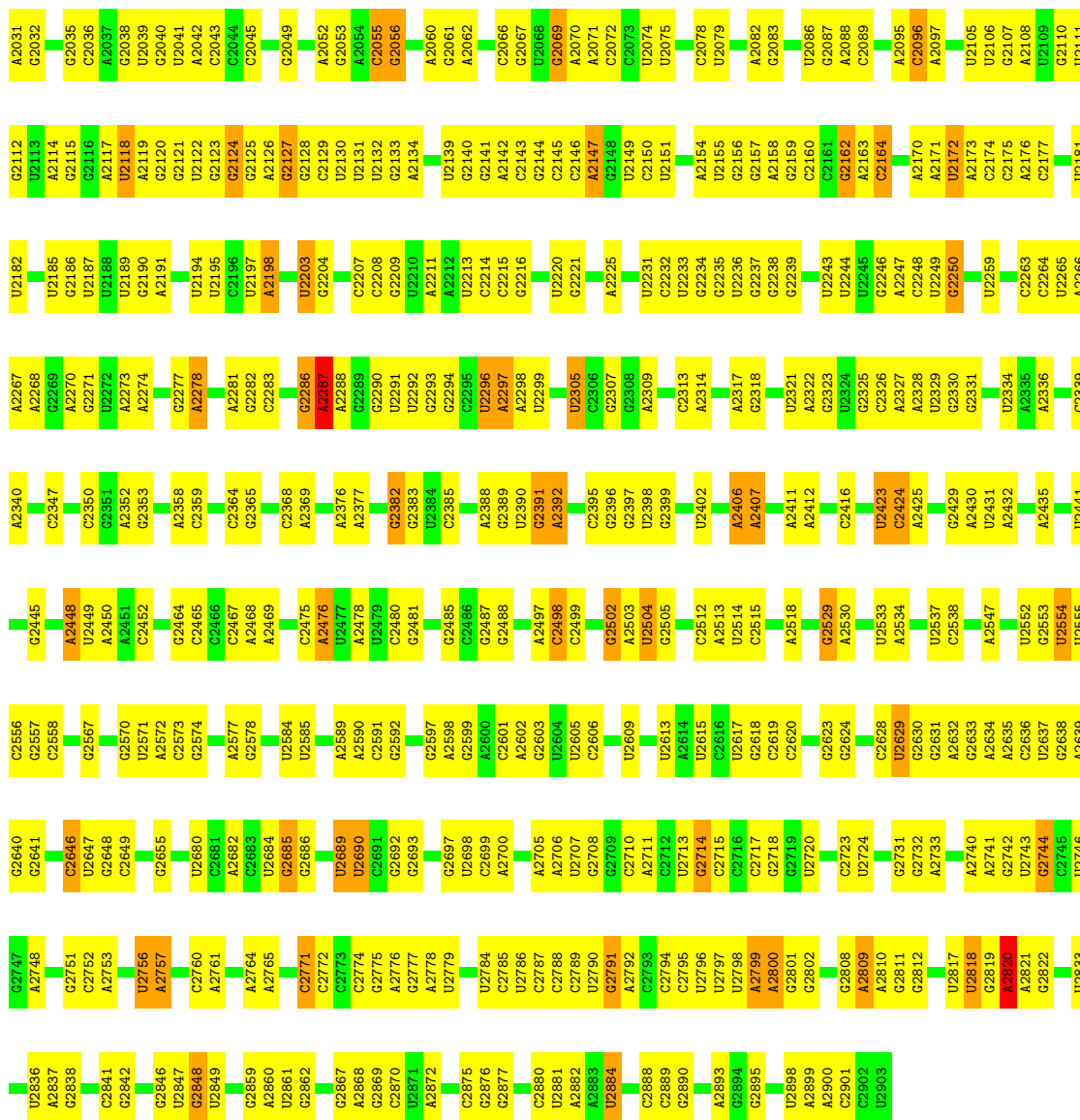
Chain T:  71% 28%



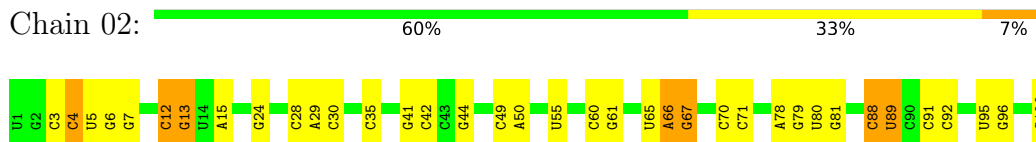


G446	G329	C225	U138	G1
U451	A330	A226	U139	C8
C455	G333	A227	G141	G9
G456	A340	C228	A142	A10
A457		C229	C143	G11
	C353		A144	U12
A460	C354	G232	C145	C16
C461	U354	A233	A146	G17
	U355	U234	C147	G10
U464	G356		U148	C20
G465	C357	C239	A149	A21
A466	U358	C240	U150	G27
G467		A241	C151	
A477	G361	C242	A152	U34
G481	A362	U243		G35
C490	C363	A244	A155	
A491	C364	G245	A156	C45
A492	U365		C157	G46
C493	C366	G248	U158	
A495		C249		
	A371		A161	A49
A499	G372	A255	U162	U50
G500	U373	A256	C163	G51
	G381	C257	C164	C57
A504	G388			G58
A505	G389	C264	G168	A63
G506	U390	A265	G169	G68
A507		G276		C69
A508	C394	C281	G176	G70
	U395	A282	G177	A71
	G400	G283	A181	A74
G512			A182	G75
C517	A404	G287	C183	G85
U518	U405	U288	G186	C96
U519	G406	A294	G187	C97
G520		G295	G188	U100
	G411	U296	G189	U101
A528		G297		U102
A529	C414		A195	C116
G530	A415	G301	A196	G117
C531		C302	A197	A118
A532	C421	G303	C198	A119
G533	A422			U120
U534	G423	A310	G205	G121
C535	U534	A311	C211	C128
G536	G536	G312	G212	C129
G537	U437		A213	C130
A538	C438	A322	G214	A131
G539	U439	C323	C215	
C540	C440	A216	A216	
A541	U441	G326	G327	
C542	G442	U329	A221	
G543			A222	
G544	C445			

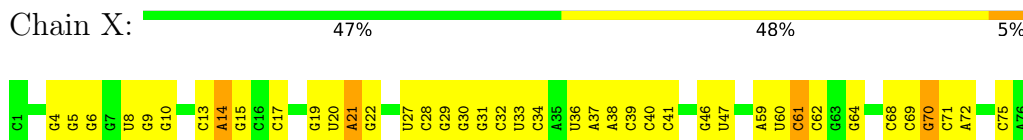
C1924	A1829	C1732	A1597	G1482	A1378	A1287	U1181	G1093	A1020	U826	U741	A633	U545
C1830	C1830	U1736	A1598	G1483	U1379	G1288	G1182	A1096	A1021	U827	U742	C634	U546
G1831	G1831	G1737	A1603	U1486	A1383	C1295	U1183	U1097	G1022	U828	A743	G635	A547
C1837	C1837	A1738	A1603	U1487	A1383	G1296	G1186	A1098	C915	A829	G744	G636	G548
G1842	G1842	G1740	A1608	U1491	C1386	C1297	G1187	G1099	C922	G831	U745	A637	G549
C1843	C1843	A1744	G1611	G1491	A1387	G1300	G1190	A1103	G1026	U832	U746	G638	G553
A1847	A1847	A1745	A1616	U1498	G1388	A1301	G1191	C1104	A1027	A833	C747	C640	U554
A1936	A1936	A1746	A1616	C1498	G1389	A1307	C1196	U1105	U1033	G834	C748	C641	U555
G1849	G1849	U1747	A1634	C1499	U1394	A1308	G1197	G1106	U1035	U839	A749	C645	A556
G1850	G1850	C1748	G1500	G1500	U1396	G1309	G1203	G1107	G1036	C840	A752	U646	C557
U1943	U1943	G1753	A1637	U1506	U1397	G1310	U1203	G1110	G1037	A845	U755	C560	U558
U1944	U1944	U1754	A1637	C1507	G1311	A1311	U1206	A1111	G1038	U846	A756	C561	G559
U1955	U1955	A1755	G1645	A1508	U1409	C1314	G1206	G1112	A1039	U847	U757	U652	C560
C1967	C1967	A1759	U1647	A1509	G1410	G1315	G1212	C1113	A1040	C848	U766	A563	A563
C1967	C1967	C1760	U1648	G1510	U1411	G1316	A1213	C1114	A1041	A849	U767	A564	U566
A1970	A1970	C1764	G1651	A1515	U1412	U1317	U1222	G1115	G1042	A851	G768	A565	U567
U1971	U1971	U1765	G1652	U1522	G1416	U1318	G1223	G1116	C1043	U852	G770	A566	U568
G1972	G1972	G1766	G1653	U1523	A1419	C1319	U1224	U1119	C1044	C853	G771	U658	U569
U1981	U1981	A1773	G1659	G1524	A1420	C1320	G1225	G1120	A1046	C854	G772	G570	G570
U1982	U1982	C1774	G1660	C1526	G1421	A1321	G1225	G1121	G1047	G855	U773	U571	U571
G1989	G1989	U1775	A1664	A1532	C1428	U1326	G1236	C1123	C1053	C857	G776	U573	U573
C1990	C1990	G1776	A1665	U1533	G1429	A1327	A1237	G1124	A1054	G858	U780	A574	A574
G1991	G1991	U1777	G1666	U1534	G1430	A1329	G1238	U1130	G1055	G859	G781	U576	U576
U1993	U1993	U1778	G1666	U1535	A1431	C1330	G1245	G1131	A1057	U860	A782	G672	G672
C1996	C1996	A1780	G1674	C1536	A1434	G1331	G1248	U1132	U1060	G862	G783	G674	G674
C1997	C1997	U1789	C1675	G1537	G1435	A1332	U1249	C1135	U1061	A863	G784	A675	G579
C1998	C1998	G1790	A1677	G1538	U1436	A1336	G1250	G1138	U1062	G864	G785	A676	U580
C1999	C1999	A1791	A1678	G1540	U1438	G1337	C1252	U1141	G1063	C865	A788	A677	C581
C2000	C2000	U1794	A1679	G1547	A1439	G1338	G1253	C1140	U1064	G869	U789	A685	A582
C2001	C2001	C1795	U1683	C1548	U1440	A1342	A1254	C1142	U1065	G873	C796	U686	G583
A2009	A2009	U1796	G1684	A1549	G1441	C1345	U1255	A1143	U1066	G874	G797	C687	U588
U2011	U2011	G1797	A1684	A1549	U1442	C1345	G1256	G1144	A1067	G875	G799	G695	U593
U2012	U2012	U1798	G1695	U1554	U1443	A1354	C1257	A1145	U1068	C876	A800	U594	U594
A2013	A2013	G1800	G1696	G1555	G1445	A1355	U1258	C1145	A1069	A877	G801	U703	C595
A2014	A2014	A1801	A1701	G1560	G1446	G1356	G1259	U1148	U1071	A878	A802	G704	U598
A2015	A2015	C1806	G1702	U1563	C1447	C1357	G1266	G1149	A1077	G879	G805	A705	A599
G2018	G2018	A1807	C1704	C1564	G1448	G1358	U1267	U1156	U1078	C885	C806	G708	A599
A2020	A2020	A1808	A1705	C1565	C1454	A1359	A1268	G1157	C1079	A886	U807	A603	A603
C2021	C2021	A1809	A1709	A1566	C1461	G1360	A1269	G1170	A1080	U887	G808	U720	A609
U2022	U2022	G1810	G1709	U1569	U1468	C1362	G1270	C1171	U1081	C889	U810	A721	C610
C2023	C2023	C1811	G1710	A1569	U1469	G1363	G1271	G1172	U1082	C890	U811	U724	C611
G2024	G2024	A1812	A1713	A1579	A1469	C1364	A1272	A1173	A1084	A896	C812	G725	U615
C2025	C2025	U1817	G1714	A1580	A1470	A1365	C1278	U1174	A1085	C897	U813	G726	U616
U2026	U2026	U1818	G1715	G1581	G1471	A1366	G1279	A1175	A1086	C898	C814	A727	A616
G2027	G2027	U1826	U1729	U1584	C1472	G1368	U1282	A1176	G1087	A899	C817	G728	G617
U2028	U2028	U1827	C1730	C1595	G1475	G1369	G1283	G1177	A1088	A900	C818	G729	A627
A2030	A2030	G1828	G1731	A1596	U1476	G1370	A1284	C1178	A1089	C901	A819	A730	G628
							A1286	U1180	A1090	G907	A825	C736	G628
									C1092			C737	



• Molecule 55: 5S ribosomal RNA

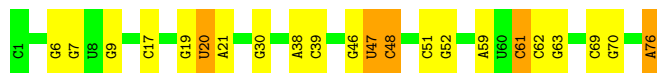


• Molecule 56: tRNA^{fMet}



- Molecule 56: tRNA^{fMet}

Chain W:  71% 22% 6%



- Molecule 57: mRNA

Chain V:  72% 22% 6%



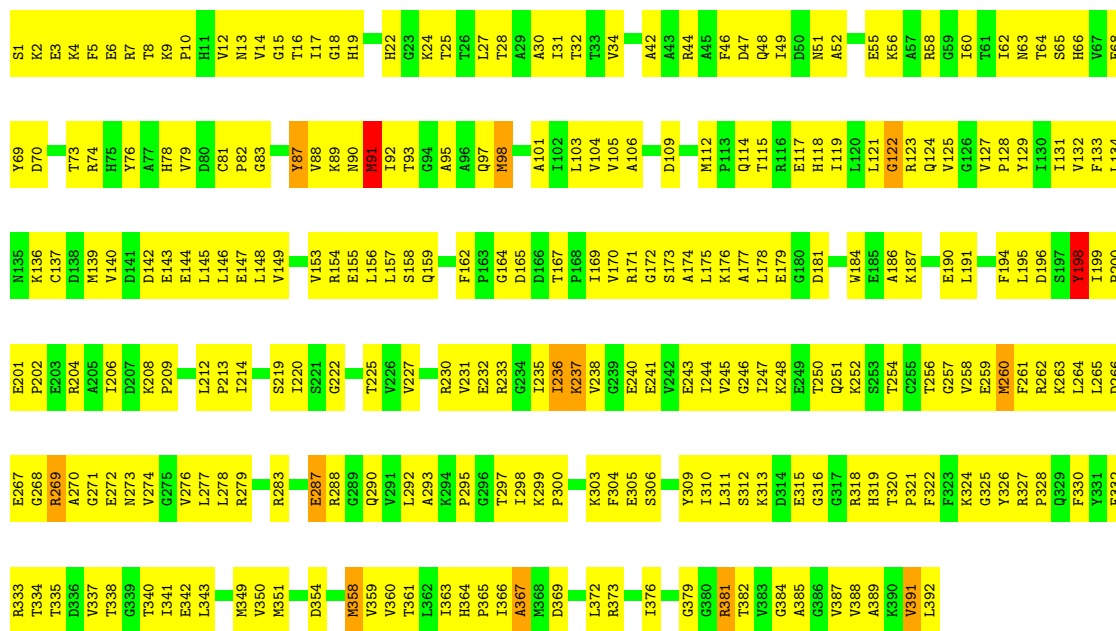
- Molecule 58: tRNA^{Phe}

Chain Y:  43% 46% 11%



- Molecule 59: Elongation factor Tu 2

Chain Z:  33% 63% 4%



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C1	Depositor
Number of particles used	10431	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION; CTFFIND3 was used to determine CTF values. FREALIGN applied CTF correction.	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	1.0	Depositor
Minimum defocus (nm)	500	Depositor
Maximum defocus (nm)	5000	Depositor
Magnification	60976	Depositor
Image detector	GATAN K2 SUMMIT (4k x 4k)	Depositor

5 Model quality ⓘ

5.1 Standard geometry ⓘ

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

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	04	0.34	0/2122	0.61	0/2852
2	05	0.37	0/1586	0.57	0/2134
3	06	0.35	0/1571	0.59	1/2113 (0.0%)
4	07	0.38	0/1435	0.56	0/1926
5	08	0.34	0/1343	0.59	1/1816 (0.1%)
6	09	0.40	0/1122	0.64	0/1515
7	10	0.46	0/1002	0.74	2/1350 (0.1%)
8	11	0.42	0/1046	0.64	0/1410
9	12	0.35	0/1152	0.58	0/1551
10	13	0.35	0/948	0.60	0/1268
11	14	0.34	0/1054	0.62	0/1403
12	15	0.39	0/1093	0.57	0/1460
13	16	0.37	0/974	0.56	0/1301
14	17	0.33	0/902	0.53	0/1209
15	18	0.35	0/929	0.58	0/1242
16	19	0.39	0/960	0.52	0/1278
17	20	0.38	0/829	0.66	1/1107 (0.1%)
18	21	0.32	0/864	0.59	0/1156
19	22	0.34	0/745	0.55	0/994
20	23	0.38	0/788	0.62	0/1051
21	24	0.38	0/766	0.55	0/1025
22	25	0.39	0/582	0.54	0/769
23	26	0.36	0/635	0.56	0/848
24	27	0.35	0/510	0.58	0/677
25	28	0.33	0/453	0.56	0/605
26	29	0.42	0/532	0.57	0/709
27	30	0.31	0/450	0.52	0/599
28	31	0.38	0/417	0.52	0/554
29	32	0.41	0/380	0.60	0/498
30	33	0.35	0/513	0.61	0/676
31	34	0.30	0/303	0.54	0/397
32	B	0.41	1/1736 (0.1%)	0.67	2/2338 (0.1%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
33	C	0.36	0/1652	0.57	0/2225
34	D	0.35	0/1665	0.60	0/2227
35	E	0.35	0/1170	0.62	0/1573
36	F	0.37	0/836	0.70	0/1128
37	G	0.34	0/1196	0.58	0/1602
38	H	0.34	0/989	0.62	1/1326 (0.1%)
39	I	0.37	0/1034	0.66	0/1375
40	J	0.37	0/797	0.65	0/1077
41	K	0.37	0/886	0.60	0/1195
42	L	0.36	0/969	0.72	1/1300 (0.1%)
43	M	0.31	0/893	0.57	0/1193
44	N	0.36	0/817	0.53	0/1088
45	O	0.35	0/722	0.55	0/964
46	P	0.37	0/659	0.62	0/884
47	Q	0.36	0/658	0.65	0/881
48	R	0.41	0/545	0.67	0/731
49	S	0.39	0/653	0.59	0/877
50	T	0.34	0/671	0.54	0/888
51	U	0.44	0/551	0.70	1/728 (0.1%)
52	03	1.71	1/1033 (0.1%)	0.83	0/1387
53	A	0.44	0/36963	0.69	7/57662 (0.0%)
54	01	0.45	0/69796	0.68	9/108888 (0.0%)
55	02	0.37	0/2872	0.69	0/4479
56	W	0.42	0/1832	0.68	0/2855
56	X	0.62	0/1832	0.71	0/2855
57	V	0.50	0/436	0.67	0/679
58	Y	0.57	0/1809	0.70	0/2819
59	Z	1.75	6/3085 (0.2%)	0.77	0/4173
All	All	0.50	8/166763 (0.0%)	0.67	26/248890 (0.0%)

All (8) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
32	B	160	LEU	C-N	5.97	1.47	1.34
59	Z	91	MET	SD-CE	5.73	2.10	1.77
59	Z	287	GLU	CB-CG	5.66	1.62	1.52
59	Z	287	GLU	CG-CD	5.54	1.60	1.51
59	Z	260	MET	SD-CE	5.31	2.07	1.77
59	Z	198	TYR	CE2-CZ	5.17	1.45	1.38
59	Z	87	TYR	CD2-CE2	5.06	1.47	1.39
52	03	38	PHE	CE1-CZ	5.06	1.47	1.37

All (26) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
32	B	160	LEU	O-C-N	-8.61	108.92	122.70
17	20	50	GLY	N-CA-C	-6.51	96.82	113.10
53	A	1301	U	N1-C1'-C2'	6.23	122.09	114.00
54	01	301	G	N9-C1'-C2'	6.21	122.07	114.00
32	B	160	LEU	CA-C-N	6.06	130.52	117.20
38	H	67	GLY	N-CA-C	-6.00	98.10	113.10
54	01	974	G	N9-C1'-C2'	5.69	121.40	114.00
5	08	119	GLY	N-CA-C	5.53	126.91	113.10
53	A	1346	A	N9-C1'-C2'	5.52	121.18	114.00
53	A	1256	A	N9-C1'-C2'	5.52	121.17	114.00
7	10	117	LEU	N-CA-C	5.50	125.86	111.00
54	01	1818	U	N1-C1'-C2'	5.50	121.14	114.00
3	06	81	GLY	N-CA-C	-5.46	99.46	113.10
54	01	2287	A	N9-C1'-C2'	5.45	121.08	114.00
53	A	429	U	N1-C1'-C2'	5.42	121.05	114.00
54	01	242	G	C1'-O4'-C4'	-5.41	105.58	109.90
51	U	10	PRO	N-CA-C	5.40	126.14	112.10
54	01	1320	C	N1-C1'-C2'	5.40	121.02	114.00
54	01	2820	A	N9-C1'-C2'	5.23	120.80	114.00
54	01	100	U	N1-C1'-C2'	5.20	120.76	114.00
53	A	1300	G	N9-C1'-C2'	5.18	120.74	114.00
53	A	1540	U	N1-C1'-C2'	5.10	120.63	114.00
7	10	80	THR	N-CA-C	5.09	124.74	111.00
54	01	669	G	N9-C1'-C2'	5.09	120.61	114.00
42	L	43	LYS	N-CA-C	5.01	124.53	111.00
53	A	818	G	N9-C1'-C2'	5.01	120.51	114.00

There are no chirality outliers.

There are no planarity outliers.

5.2 Too-close contacts

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	04	2083	0	2157	73	0
2	05	1565	0	1616	47	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
3	06	1552	0	1619	49	0
4	07	1411	0	1447	64	0
5	08	1323	0	1374	34	0
6	09	1111	0	1148	37	0
7	10	989	0	1025	62	0
8	11	1032	0	1088	63	0
9	12	1129	0	1162	30	0
10	13	939	0	1012	27	0
11	14	1045	0	1117	34	0
12	15	1074	0	1157	38	0
13	16	961	0	1000	30	0
14	17	892	0	923	20	0
15	18	917	0	965	33	0
16	19	947	0	1022	31	0
17	20	816	0	839	29	0
18	21	857	0	922	23	0
19	22	739	0	807	20	0
20	23	780	0	834	22	0
21	24	753	0	780	28	0
22	25	575	0	592	18	0
23	26	625	0	655	19	0
24	27	509	0	543	9	0
25	28	449	0	491	16	0
26	29	523	0	524	17	0
27	30	444	0	461	21	0
28	31	410	0	440	7	0
29	32	377	0	418	10	0
30	33	504	0	574	13	0
31	34	302	0	343	9	0
32	B	1705	0	1732	92	0
33	C	1625	0	1699	49	0
34	D	1643	0	1710	50	0
35	E	1157	0	1199	47	0
36	F	818	0	808	39	0
37	G	1182	0	1240	40	0
38	H	979	0	1034	46	0
39	I	1022	0	1070	38	0
40	J	787	0	828	44	0
41	K	870	0	878	36	0
42	L	955	0	1019	26	0
43	M	884	0	944	35	0
44	N	805	0	847	34	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
45	O	714	0	737	17	0
46	P	649	0	666	26	0
47	Q	649	0	691	29	0
48	R	536	0	552	24	0
49	S	638	0	665	31	0
50	T	665	0	714	20	0
51	U	545	0	579	44	0
52	03	1026	0	1092	121	0
53	A	33012	0	16618	454	0
54	01	62317	0	31346	855	0
55	02	2568	0	1303	34	0
56	W	1640	0	836	15	0
56	X	1640	0	837	28	0
57	V	388	0	196	4	0
58	Y	1619	0	821	38	0
59	Z	3029	0	3043	291	0
60	W	10	0	10	3	0
61	Y	11	0	8	3	0
62	Z	32	0	14	0	0
All	All	153753	0	104791	3217	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 12.

All (3217) 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)
59:Z:112:MET:CE	59:Z:112:MET:SD	2.02	1.47
59:Z:358:MET:SD	59:Z:358:MET:CE	2.05	1.45
52:03:218:MET:CE	52:03:218:MET:SD	2.05	1.44
56:W:76:A:O3'	60:W:101:FME:C	1.63	1.43
59:Z:260:MET:CE	59:Z:260:MET:SD	2.07	1.43
59:Z:91:MET:CE	59:Z:91:MET:SD	2.10	1.39
54:01:45:G:H5''	54:01:46:G:H5'	1.32	1.12
51:U:9:GLU:HG2	51:U:10:PRO:HD3	1.10	1.08
52:03:51:ASP:HB3	52:03:57:GLN:HG3	1.39	1.04
35:E:80:LEU:HD13	35:E:122:VAL:HG11	1.39	1.03
7:10:57:ASN:HB2	7:10:62:ARG:HD2	1.39	1.02
54:01:2277:G:H2'	54:01:2278:A:H5''	1.43	0.99
7:10:118:ILE:H	7:10:119:PRO:CD	1.71	0.98
36:F:12:PRO:HD2	36:F:54:LEU:HD21	1.43	0.98

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:01:507:A:H5''	54:01:508:A:H5''	1.46	0.97
40:J:42:LEU:HD11	40:J:73:LEU:HG	1.42	0.97
59:Z:303:LYS:HG2	59:Z:361:THR:HG22	1.46	0.97
59:Z:369:ASP:HB2	59:Z:372:LEU:HB2	1.48	0.96
7:10:118:ILE:H	7:10:119:PRO:HD2	1.30	0.96
8:11:55:PRO:HG2	8:11:71:LYS:HB2	1.46	0.96
51:U:9:GLU:HG2	51:U:10:PRO:CD	1.96	0.95
48:R:70:THR:HG23	48:R:71:ASP:H	1.31	0.95
52:03:200:LYS:HD2	52:03:201:PRO:HD2	1.45	0.95
59:Z:7:ARG:HH22	59:Z:269:ARG:HE	1.12	0.94
55:02:3:C:H2'	55:02:4:C:H5''	1.46	0.94
36:F:38:ARG:HD3	36:F:97:THR:HA	1.49	0.93
52:03:46:VAL:HG22	52:03:212:VAL:HA	1.48	0.93
59:Z:256:THR:HG21	59:Z:279:ARG:HD3	1.52	0.92
1:04:48:ILE:HD11	1:04:51:ARG:HA	1.52	0.91
59:Z:214:ILE:HD12	59:Z:290:GLN:HB3	1.51	0.90
9:12:109:LEU:HD13	9:12:118:MET:HG3	1.49	0.90
1:04:106:PRO:HD2	1:04:109:LEU:HD22	1.52	0.89
3:06:34:ALA:HA	3:06:94:GLN:HE21	1.35	0.89
54:01:1597:A:H5''	54:01:1598:A:H5'	1.53	0.89
11:14:95:LEU:HD22	11:14:100:ILE:HD11	1.52	0.89
8:11:91:LYS:HG3	8:11:94:LYS:HE2	1.53	0.89
32:B:112:ARG:HH12	32:B:116:LEU:HD13	1.36	0.88
59:Z:206:ILE:HG22	59:Z:270:ALA:H	1.39	0.88
12:15:33:LEU:HD13	12:15:117:PHE:HB3	1.56	0.88
7:10:88:HIS:HB2	7:10:89:PRO:HD3	1.56	0.88
58:Y:6:G:H3'	58:Y:7:A:H5''	1.56	0.87
8:11:48:ILE:HG13	8:11:49:GLU:H	1.39	0.87
59:Z:10:PRO:HB2	59:Z:74:ARG:HD3	1.57	0.87
8:11:123:ALA:HB1	54:01:1081:U:H4'	1.56	0.86
51:U:16:ARG:HB2	51:U:19:LYS:HD3	1.57	0.86
59:Z:6:GLU:HG2	59:Z:8:THR:HG23	1.54	0.86
38:H:46:GLU:HB3	38:H:61:THR:HB	1.56	0.86
56:W:76:A:C3'	60:W:101:FME:C	2.53	0.86
21:24:72:VAL:HG12	21:24:93:ARG:HA	1.58	0.85
59:Z:58:ARG:HG3	59:Z:60:ILE:HG12	1.57	0.85
53:A:1259:C:H3'	53:A:1260:G:H5''	1.57	0.85
59:Z:246:GLY:HA3	59:Z:290:GLN:HG3	1.57	0.85
8:11:20:SER:HB3	8:11:21:PRO:HD3	1.57	0.85
46:P:31:ARG:HH21	53:A:230:G:H5''	1.40	0.85
52:03:31:LYS:HA	52:03:34:ALA:HB3	1.59	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:10:118:ILE:HG22	7:10:119:PRO:HD3	1.59	0.84
59:Z:14:VAL:HG11	59:Z:195:LEU:HD21	1.58	0.84
43:M:15:VAL:HG23	43:M:16:ILE:HD12	1.57	0.84
59:Z:119:ILE:HD12	59:Z:156:LEU:HG	1.57	0.84
35:E:107:GLY:HA3	53:A:9:G:H5'	1.60	0.84
53:A:1029:U:H2'	53:A:1031:C:H1'	1.58	0.84
49:S:5:LYS:HG3	49:S:6:LYS:HG2	1.59	0.84
49:S:29:PRO:HG2	49:S:31:ARG:HH12	1.42	0.83
52:03:4:LEU:HD12	52:03:9:ARG:HG2	1.58	0.83
12:15:45:GLN:HE21	54:01:2485:G:H5''	1.43	0.83
42:L:109:ARG:HH12	53:A:537:G:H5''	1.42	0.82
54:01:2800:A:H3'	54:01:2801:G:H5'	1.61	0.82
53:A:112:G:H21	53:A:354:G:H5'	1.42	0.82
50:T:70:LYS:HA	50:T:73:ARG:HE	1.42	0.82
59:Z:154:ARG:HD3	59:Z:165:ASP:HA	1.61	0.82
59:Z:326:TYR:O	59:Z:341:ILE:HG12	1.80	0.82
10:13:21:CYS:HA	10:13:41:ILE:HG22	1.62	0.82
7:10:19:ALA:HA	7:10:70:GLU:HG3	1.60	0.81
36:F:52:ASN:O	36:F:53:LYS:HG3	1.79	0.81
54:01:121:G:H4'	54:01:149:A:H5'	1.62	0.81
2:05:155:VAL:HG21	54:01:2618:G:H21	1.45	0.81
54:01:2452:C:H42	54:01:2504:U:H3	1.29	0.81
52:03:4:LEU:HB3	52:03:9:ARG:HE	1.44	0.81
1:04:257:ARG:HH21	1:04:266:ILE:HD12	1.45	0.81
40:J:17:LEU:HG	40:J:20:GLN:HE21	1.46	0.80
37:G:128:GLU:HG3	37:G:130:LYS:HE2	1.64	0.80
8:11:29:GLN:HE22	54:01:1096:A:H61	1.30	0.80
59:Z:89:LYS:HD3	59:Z:288:ARG:HH22	1.46	0.80
59:Z:245:VAL:HG13	59:Z:250:THR:HG21	1.63	0.80
39:I:83:THR:HG21	39:I:102:PHE:HB3	1.64	0.79
56:X:13:C:H2'	56:X:14:A:H5''	1.64	0.79
20:23:42:LYS:HE2	54:01:499:U:H5''	1.64	0.79
35:E:155:LYS:HB3	38:H:70:VAL:HG13	1.64	0.79
35:E:152:VAL:HG11	38:H:98:LEU:HD13	1.65	0.79
59:Z:105:VAL:HB	59:Z:134:LEU:HD23	1.64	0.79
59:Z:309:TYR:HE2	59:Z:311:LEU:HD13	1.48	0.79
59:Z:184:TRP:HA	59:Z:187:LYS:HD3	1.66	0.78
53:A:405:U:H3'	53:A:406:G:H5'	1.65	0.78
10:13:121:GLU:HG2	10:13:122:VAL:HG23	1.64	0.78
30:33:18:LYS:HG3	54:01:651:G:H5'	1.64	0.78
51:U:66:ARG:HG3	53:A:1099:G:H4'	1.64	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:B:18:GLN:HG3	32:B:187:ASP:OD2	1.84	0.78
6:09:1:MET:HB3	6:09:3:VAL:HG23	1.65	0.78
54:01:189:G:H2'	54:01:205:G:H22	1.48	0.78
39:I:20:ILE:HD11	39:I:60:LEU:HD22	1.65	0.77
46:P:20:VAL:HG23	46:P:35:ARG:HA	1.66	0.77
10:13:13:ASN:HD21	10:13:98:ARG:HB2	1.50	0.76
52:03:41:SER:HA	52:03:177:LYS:HA	1.67	0.76
2:05:33:ARG:HD3	2:05:73:VAL:HB	1.67	0.76
59:Z:93:THR:HG23	59:Z:334:THR:HA	1.67	0.76
59:Z:63:ASN:HA	59:Z:90:ASN:ND2	1.99	0.76
9:12:117:ALA:HA	9:12:120:ARG:HH21	1.50	0.76
47:Q:45:VAL:HG21	47:Q:60:ILE:HD13	1.67	0.76
42:L:32:VAL:HB	42:L:55:ARG:HB3	1.67	0.76
8:11:38:CYS:HA	8:11:41:PHE:HB3	1.68	0.76
46:P:4:ILE:HG12	46:P:21:VAL:HG22	1.67	0.76
8:11:101:SER:HB3	8:11:104:GLN:HG3	1.66	0.76
33:C:116:ALA:HB3	33:C:184:ASN:HD22	1.47	0.76
8:11:126:ARG:HA	8:11:129:GLU:HG3	1.68	0.75
44:N:25:GLU:HB2	44:N:29:ILE:HD12	1.68	0.75
48:R:11:ARG:HG3	48:R:14:ALA:HB3	1.68	0.75
8:11:11:GLN:HB2	8:11:56:VAL:HG12	1.69	0.75
54:01:2114:A:H61	54:01:2117:A:H62	1.32	0.75
54:01:2799:A:H2'	54:01:2800:A:H5'	1.68	0.75
41:K:124:LYS:HE2	53:A:1523:G:H5''	1.67	0.75
54:01:1053:C:H2'	54:01:1054:A:H5''	1.66	0.75
54:01:1645:G:H5''	54:01:1646:C:H5'	1.68	0.75
32:B:73:ARG:HH22	32:B:94:ARG:HH22	1.31	0.75
38:H:77:VAL:HG12	38:H:84:ILE:HD12	1.67	0.75
54:01:2162:G:H2'	54:01:2163:A:H8	1.51	0.75
35:E:54:GLU:HG2	35:E:56:PRO:HD2	1.68	0.75
54:01:265:A:H4'	54:01:266:G:OP1	1.85	0.75
52:03:57:GLN:HE22	52:03:203:GLN:HB3	1.52	0.75
32:B:71:THR:HG22	32:B:72:LYS:H	1.52	0.75
13:16:37:THR:HG22	13:16:39:PRO:HD2	1.67	0.74
36:F:66:ALA:HB1	36:F:67:PRO:HD2	1.69	0.74
51:U:28:LEU:HA	51:U:31:VAL:HG12	1.69	0.74
22:25:36:GLN:NE2	22:25:39:THR:HA	2.01	0.74
42:L:79:ILE:HG22	42:L:103:CYS:HB2	1.67	0.74
15:18:52:ARG:NH2	54:01:2720:U:H5''	2.01	0.74
59:Z:260:MET:HB2	59:Z:274:VAL:HG12	1.68	0.74
3:06:105:LEU:HD23	3:06:108:ILE:HD11	1.69	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:A:1206:G:H2'	53:A:1207:G:H5''	1.68	0.74
12:15:12:MET:HA	54:01:910:A:H62	1.52	0.74
21:24:77:VAL:HG23	21:24:89:ILE:HG12	1.68	0.74
59:Z:305:GLU:HA	59:Z:359:VAL:HG22	1.70	0.73
21:24:86:LEU:HD13	21:24:89:ILE:HD11	1.68	0.73
39:I:123:ARG:HD3	39:I:124:PRO:HD2	1.69	0.73
54:01:528:A:N1	54:01:2042:A:H2'	2.03	0.73
7:10:103:ASN:ND2	7:10:110:ALA:HB3	2.02	0.73
43:M:28:ARG:HH21	43:M:62:PHE:HB2	1.52	0.73
44:N:68:ARG:HG3	53:A:1202:U:H4'	1.70	0.73
37:G:12:LEU:HD12	37:G:13:PRO:HD2	1.69	0.73
51:U:16:ARG:HH21	51:U:19:LYS:HG2	1.54	0.73
53:A:1137:C:H5'	53:A:1138:G:H5'	1.71	0.73
59:Z:343:LEU:HA	59:Z:358:MET:HB2	1.70	0.73
13:16:78:LYS:HE2	13:16:83:LEU:HD21	1.71	0.73
53:A:1414:U:H2'	53:A:1415:G:H8	1.54	0.73
52:03:193:LEU:HD23	52:03:196:LEU:HD22	1.70	0.72
41:K:30:ILE:HB	41:K:45:THR:HG22	1.72	0.72
10:13:71:ARG:HH12	15:18:71:ARG:HH21	1.37	0.72
32:B:183:PHE:HB3	32:B:197:PHE:HB2	1.71	0.72
59:Z:237:LYS:H	59:Z:237:LYS:HD2	1.52	0.72
8:11:72:THR:HG21	8:11:112:LYS:HG3	1.72	0.72
11:14:96:LYS:HE3	11:14:103:ILE:HA	1.72	0.72
55:02:3:C:C2'	55:02:4:C:H5''	2.18	0.72
59:Z:88:VAL:O	59:Z:92:ILE:HG13	1.90	0.72
17:20:14:VAL:HG23	17:20:18:GLN:HE21	1.53	0.72
32:B:118:THR:HA	32:B:121:GLN:HE21	1.55	0.72
59:Z:313:LYS:HD3	59:Z:319:HIS:O	1.89	0.72
21:24:42:LEU:HD13	21:24:47:VAL:HG21	1.72	0.72
53:A:1033:G:H3'	53:A:1034:G:H5''	1.72	0.72
32:B:16:GLY:O	32:B:17:HIS:HB2	1.89	0.72
51:U:66:ARG:HG3	53:A:1099:G:C4'	2.20	0.72
4:07:22:ASN:HB2	4:07:26:GLN:HE22	1.55	0.72
2:05:13:ARG:HH11	15:18:55:HIS:HA	1.53	0.71
5:08:138:GLN:HE22	54:01:2746:U:H1'	1.54	0.71
59:Z:49:ILE:HG23	59:Z:65:SER:HB3	1.72	0.71
10:13:76:VAL:H	15:18:72:VAL:HG22	1.55	0.71
32:B:165:ALA:HB3	32:B:190:SER:HB3	1.72	0.71
19:22:2:ILE:HD11	54:01:144:A:H4'	1.71	0.71
39:I:91:GLU:HA	39:I:94:ARG:HB2	1.72	0.71
8:11:45:THR:HG22	8:11:50:LYS:HG2	1.72	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:01:2591:C:H2'	54:01:2592:G:C8	2.26	0.71
34:D:144:ILE:HD13	34:D:177:MET:HB3	1.71	0.71
37:G:111:GLY:HA2	37:G:118:ARG:HH11	1.56	0.71
41:K:124:LYS:HE3	53:A:780:A:H5''	1.72	0.71
49:S:14:LEU:O	49:S:18:VAL:HG23	1.91	0.71
54:01:885:C:H2'	54:01:886:A:H5'	1.72	0.71
7:10:27:VAL:HG13	7:10:83:ALA:HB3	1.73	0.70
54:01:2277:G:C2'	54:01:2278:A:H5''	2.18	0.70
10:13:13:ASN:ND2	10:13:98:ARG:HB2	2.06	0.70
59:Z:149:VAL:O	59:Z:153:VAL:HG23	1.92	0.70
34:D:97:LEU:HB2	34:D:134:TYR:HB3	1.74	0.70
54:01:2141:G:H22	54:01:2151:U:H1'	1.56	0.70
53:A:484:G:H4'	53:A:485:U:H5''	1.73	0.70
32:B:18:GLN:O	32:B:19:THR:HB	1.89	0.70
56:X:69:C:H2'	56:X:70:G:H5'	1.73	0.70
59:Z:260:MET:O	59:Z:263:LYS:HB2	1.92	0.70
11:14:101:ILE:HG13	11:14:102:GLY:H	1.56	0.70
53:A:960:U:H4'	53:A:961:U:O5'	1.91	0.70
29:32:34:ARG:HH21	29:32:39:ARG:HD2	1.56	0.70
54:01:1055:G:H1'	54:01:1084:A:H61	1.57	0.70
20:23:36:GLU:HA	20:23:61:GLU:HG2	1.74	0.69
22:25:39:THR:H	54:01:2331:G:H4'	1.56	0.69
33:C:116:ALA:HB3	33:C:184:ASN:ND2	2.07	0.69
39:I:35:GLU:HA	39:I:39:GLY:HA3	1.73	0.69
54:01:275:C:H2'	54:01:276:U:H4'	1.72	0.69
39:I:48:ARG:HA	39:I:51:LEU:HD12	1.73	0.69
43:M:23:GLY:HA2	43:M:68:LEU:HD22	1.75	0.69
46:P:5:ARG:HB2	53:A:376:G:H5''	1.74	0.69
38:H:9:MET:HG3	38:H:26:MET:SD	2.32	0.69
5:08:87:GLN:HE21	5:08:162:ARG:HD2	1.57	0.69
32:B:16:GLY:HA2	32:B:40:ILE:HG13	1.73	0.69
54:01:118:A:H5'	54:01:119:A:C8	2.27	0.69
59:Z:206:ILE:HG22	59:Z:270:ALA:N	2.06	0.69
32:B:182:VAL:HG23	32:B:195:VAL:HA	1.72	0.69
46:P:43:ALA:HA	46:P:46:LYS:HE3	1.73	0.69
16:19:49:ARG:HD2	54:01:993:G:OP1	1.93	0.69
23:26:5:GLN:HG2	23:26:49:ARG:HB2	1.73	0.69
40:J:57:VAL:O	40:J:58:ASN:HB2	1.93	0.69
43:M:13:HIS:HB2	43:M:16:ILE:HD13	1.75	0.69
8:11:79:LEU:HD13	8:11:137:LEU:HD13	1.74	0.69
59:Z:237:LYS:HD3	59:Z:240:GLU:HB2	1.75	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:Z:364:HIS:HB3	59:Z:365:PRO:HD2	1.75	0.69
19:22:35:ALA:HB3	19:22:38:ALA:HB2	1.74	0.68
20:23:10:VAL:HG12	20:23:71:ILE:HA	1.75	0.68
33:C:82:ASP:HA	33:C:85:LYS:HE3	1.75	0.68
52:03:193:LEU:HA	52:03:196:LEU:HB2	1.75	0.68
54:01:2106:U:H2'	54:01:2107:G:H8	1.58	0.68
21:24:25:LYS:HG2	21:24:43:ASP:HA	1.75	0.68
36:F:64:VAL:HG22	36:F:65:GLU:H	1.56	0.68
53:A:1218:C:H2'	53:A:1219:A:C8	2.28	0.68
59:Z:212:LEU:HD12	59:Z:231:VAL:HG22	1.75	0.68
52:03:27:ILE:HB	52:03:182:ALA:HB1	1.76	0.68
54:01:2296:U:H5''	54:01:2297:A:OP1	1.93	0.68
2:05:4:LEU:HD23	2:05:32:ASN:HD22	1.57	0.68
34:D:131:ILE:H	34:D:131:ILE:HD12	1.57	0.68
39:I:27:ILE:HG21	39:I:34:LEU:HD22	1.75	0.68
41:K:19:VAL:HG13	41:K:82:GLU:HB2	1.75	0.68
7:10:23:LEU:HD13	7:10:118:ILE:HB	1.74	0.68
16:19:105:PHE:O	16:19:109:VAL:HG23	1.94	0.68
34:D:172:VAL:HG22	34:D:174:ALA:H	1.58	0.68
4:07:127:TYR:HB3	4:07:155:ILE:HB	1.75	0.68
54:01:876:C:H42	54:01:901:C:H42	1.42	0.68
59:Z:63:ASN:HD22	59:Z:90:ASN:HD21	1.42	0.68
59:Z:83:GLY:HA2	59:Z:118:HIS:NE2	2.09	0.68
37:G:148:LYS:O	37:G:148:LYS:HD3	1.93	0.68
56:X:39:C:H2'	56:X:40:C:C6	2.28	0.68
32:B:33:ALA:HB2	32:B:39:ILE:HG13	1.76	0.68
39:I:89:TYR:HB3	39:I:93:LEU:HD12	1.75	0.68
59:Z:129:TYR:HB3	59:Z:199:ILE:HD13	1.76	0.67
59:Z:231:VAL:HG21	59:Z:236:ILE:HD11	1.75	0.67
54:01:2699:C:H2'	54:01:2700:A:H8	1.59	0.67
59:Z:19:HIS:HB3	59:Z:22:HIS:CE1	2.29	0.67
29:32:24:THR:HG23	29:32:27:GLY:H	1.58	0.67
48:R:48:ALA:O	48:R:52:ARG:HG3	1.94	0.67
50:T:83:ASN:HA	50:T:86:ALA:HB3	1.77	0.67
54:01:534:U:H2'	54:01:535:G:H8	1.58	0.67
4:07:58:ALA:O	4:07:139:GLU:HG2	1.94	0.67
54:01:322:A:H5'	54:01:340:A:H1'	1.76	0.67
54:01:2267:A:H5''	54:01:2268:A:H5'	1.76	0.67
1:04:261:ARG:HD3	1:04:262:THR:HG23	1.77	0.67
4:07:87:LYS:HD2	54:01:2313:C:H5''	1.74	0.67
52:03:34:ALA:HB1	52:03:178:VAL:HG11	1.76	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:01:677:A:O2'	54:01:2071:A:H5'	1.94	0.67
8:11:48:ILE:HG13	8:11:49:GLU:N	2.10	0.67
46:P:13:LYS:HE3	53:A:392:C:H4'	1.75	0.67
54:01:742:A:H2'	54:01:743:A:C8	2.30	0.67
32:B:66:ILE:CD1	32:B:159:ALA:HB3	2.25	0.67
59:Z:237:LYS:HB3	59:Z:267:GLU:OE2	1.95	0.67
11:14:111:ILE:H	11:14:111:ILE:HD12	1.58	0.67
34:D:84:ASN:HB3	34:D:87:GLU:HB2	1.77	0.67
6:09:84:ALA:HA	6:09:91:PHE:H	1.59	0.67
9:12:7:LYS:HG2	54:01:538:A:H4'	1.77	0.67
53:A:769:G:H4'	53:A:1513:A:H4'	1.77	0.67
59:Z:97:GLN:NE2	59:Z:230:ARG:H	1.93	0.67
36:F:68:GLN:NE2	53:A:738:C:H5''	2.09	0.66
43:M:47:LEU:HD21	43:M:51:GLN:HB2	1.77	0.66
54:01:118:A:H5'	54:01:119:A:H8	1.61	0.66
54:01:1790:C:H2'	54:01:1791:A:C5	2.30	0.66
50:T:73:ARG:HH22	53:A:263:A:P	2.18	0.66
18:21:83:LYS:HG2	18:21:95:ARG:HH12	1.61	0.66
45:O:87:ARG:HE	45:O:88:ARG:H	1.41	0.66
59:Z:112:MET:H	59:Z:115:THR:HB	1.60	0.66
59:Z:332:PHE:H	59:Z:335:THR:HB	1.59	0.66
6:09:9:VAL:HB	6:09:13:GLY:HA3	1.76	0.66
46:P:7:ALA:HB3	46:P:18:GLN:HB2	1.78	0.66
53:A:225:C:H2'	53:A:226:G:H5''	1.77	0.66
6:09:12:LEU:HD13	6:09:19:VAL:HG21	1.76	0.66
9:12:117:ALA:HA	9:12:120:ARG:NH2	2.11	0.66
23:26:70:LEU:HD23	23:26:73:ARG:HH21	1.59	0.66
47:Q:64:ARG:HD2	53:A:264:C:H4'	1.77	0.66
54:01:1394:U:H4'	54:01:1603:A:H4'	1.77	0.66
54:01:2118:U:H5	54:01:2149:U:H1'	1.59	0.66
59:Z:19:HIS:HA	59:Z:83:GLY:HA3	1.76	0.66
59:Z:212:LEU:HG	59:Z:231:VAL:HA	1.77	0.66
3:06:146:VAL:HG12	3:06:185:LYS:HB2	1.78	0.66
12:15:41:LEU:HG	12:15:96:ILE:HG13	1.77	0.66
42:L:78:VAL:O	42:L:102:ASP:HB2	1.94	0.66
54:01:2553:G:H3'	54:01:2554:U:H5''	1.77	0.66
59:Z:186:ALA:O	59:Z:190:GLU:HG3	1.96	0.66
7:10:118:ILE:N	7:10:119:PRO:CD	2.51	0.65
17:20:49:ILE:HG22	17:20:54:VAL:HA	1.78	0.65
54:01:2771:C:H2'	54:01:2772:C:C6	2.32	0.65
20:23:4:ILE:HD12	20:23:4:ILE:H	1.61	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:D:100:VAL:HG21	34:D:136:VAL:HG21	1.77	0.65
9:12:7:LYS:O	9:12:11:VAL:HG23	1.96	0.65
52:03:37:LYS:HD2	54:01:2127:G:H5'	1.78	0.65
4:07:28:PRO:HB2	4:07:168:LEU:HD22	1.78	0.65
5:08:8:VAL:HB	5:08:49:LEU:HB2	1.77	0.65
33:C:46:LEU:HB3	33:C:49:ALA:HB3	1.79	0.65
53:A:664:G:H22	53:A:741:G:H1	1.45	0.65
54:01:858:G:H5'	54:01:859:G:OP2	1.96	0.65
59:Z:322:PHE:CE1	59:Z:350:VAL:HB	2.32	0.65
32:B:137:THR:O	32:B:141:GLU:HG3	1.97	0.65
52:03:57:GLN:NE2	52:03:203:GLN:HB3	2.12	0.65
53:A:202:G:H21	53:A:466:A:H61	1.45	0.65
54:01:639:U:H2'	54:01:640:C:C6	2.31	0.65
2:05:151:THR:HB	2:05:152:PRO:HD3	1.78	0.65
11:14:30:THR:O	11:14:33:ARG:HG2	1.97	0.65
12:15:45:GLN:NE2	54:01:2485:G:H5''	2.11	0.65
32:B:129:THR:HB	32:B:132:GLU:HG2	1.79	0.65
34:D:94:GLU:HA	34:D:99:ASN:ND2	2.11	0.65
4:07:47:LYS:HA	4:07:50:ASP:OD2	1.96	0.65
52:03:165:ASN:HB3	52:03:171:ILE:HB	1.77	0.65
54:01:780:G:H2'	54:01:782:A:N7	2.12	0.65
54:01:2467:C:H2'	54:01:2468:A:O4'	1.96	0.65
44:N:92:ILE:H	44:N:92:ILE:HD12	1.61	0.64
10:13:69:VAL:HG21	10:13:104:THR:HG21	1.80	0.64
42:L:101:LEU:O	42:L:103:CYS:N	2.30	0.64
53:A:212:G:H2'	53:A:213:G:H8	1.61	0.64
5:08:94:ARG:HB2	5:08:105:SER:HB2	1.78	0.64
14:17:80:GLU:O	14:17:84:GLU:HG3	1.97	0.64
8:11:53:PRO:HG2	8:11:77:VAL:HG11	1.79	0.64
15:18:38:ARG:HH22	15:18:40:GLN:HB3	1.61	0.64
53:A:352:C:H4'	53:A:354:G:OP1	1.96	0.64
11:14:63:LYS:HA	30:33:12:ARG:HG2	1.79	0.64
21:24:80:HIS:ND1	21:24:81:PRO:HD2	2.12	0.64
34:D:96:ARG:O	34:D:100:VAL:HG23	1.97	0.64
50:T:79:THR:HG22	50:T:83:ASN:ND2	2.13	0.64
53:A:112:G:N2	53:A:354:G:H5'	2.12	0.64
1:04:144:GLU:HG3	1:04:188:ARG:O	1.98	0.64
32:B:174:GLU:HA	32:B:177:ASN:ND2	2.13	0.64
33:C:87:ARG:HG3	33:C:98:ALA:HB3	1.79	0.64
42:L:41:PRO:HD2	42:L:47:ALA:H	1.62	0.64
13:16:38:LEU:HB3	13:16:39:PRO:HD3	1.79	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:Z:5:PHE:HB2	59:Z:263:LYS:HB3	1.79	0.64
59:Z:140:VAL:HG23	59:Z:142:ASP:H	1.63	0.64
1:04:144:GLU:HB2	1:04:187:CYS:HB3	1.80	0.64
40:J:52:LEU:HD21	40:J:59:LYS:HD2	1.80	0.64
32:B:129:THR:HG22	32:B:131:LYS:H	1.63	0.64
38:H:29:SER:O	38:H:33:VAL:HG23	1.98	0.64
43:M:24:VAL:HG23	43:M:28:ARG:HB3	1.80	0.64
46:P:57:ILE:O	46:P:61:VAL:HG23	1.98	0.64
58:Y:25:C:H3'	58:Y:26:A:H5''	1.80	0.64
17:20:38:VAL:HG13	17:20:54:VAL:HG23	1.80	0.63
35:E:10:LEU:HD22	35:E:67:ARG:HH22	1.61	0.63
49:S:5:LYS:HD2	49:S:6:LYS:HE3	1.79	0.63
53:A:1412:C:H2'	53:A:1413:A:C8	2.33	0.63
54:01:189:G:H2'	54:01:205:G:N2	2.11	0.63
55:02:88:C:H5''	55:02:89:U:OP1	1.97	0.63
58:Y:7:A:H3'	58:Y:8:U:H5''	1.80	0.63
48:R:49:LYS:O	48:R:53:GLN:HG3	1.98	0.63
7:10:8:LYS:O	7:10:12:VAL:HG23	1.98	0.63
45:O:45:HIS:O	45:O:47:LYS:N	2.31	0.63
18:21:29:VAL:HG21	18:21:69:LEU:HD23	1.81	0.63
54:01:2799:A:C2'	54:01:2800:A:H5'	2.29	0.63
59:Z:300:PRO:HG2	59:Z:365:PRO:HB2	1.81	0.63
49:S:62:THR:HG22	49:S:63:ASP:H	1.64	0.63
3:06:69:ARG:HH22	54:01:2445:G:P	2.22	0.63
54:01:554:U:H2'	54:01:555:G:O4'	1.98	0.63
54:01:839:U:H2'	54:01:840:C:C6	2.34	0.63
59:Z:88:VAL:HG13	59:Z:92:ILE:HD11	1.81	0.63
53:A:129:A:H1'	53:A:130:A:N7	2.14	0.63
59:Z:63:ASN:ND2	59:Z:90:ASN:HD21	1.95	0.63
36:F:12:PRO:HG3	36:F:57:ALA:HA	1.80	0.63
54:01:1386:C:H2'	54:01:1387:A:C8	2.33	0.63
54:01:2141:G:H2'	54:01:2142:A:H8	1.64	0.63
6:09:94:ILE:HG23	6:09:98:ASP:HB2	1.81	0.63
8:11:77:VAL:HA	8:11:80:LYS:HG3	1.81	0.63
19:22:58:VAL:HG13	19:22:85:VAL:HG22	1.80	0.63
20:23:12:VAL:HA	20:23:69:VAL:HG12	1.79	0.63
29:32:34:ARG:HD3	54:01:467:G:OP2	1.99	0.63
59:Z:27:LEU:O	59:Z:31:ILE:HG13	1.99	0.63
44:N:7:ALA:HA	44:N:10:VAL:HG12	1.80	0.62
52:03:207:VAL:HB	52:03:210:LYS:HG3	1.80	0.62
59:Z:191:LEU:HA	59:Z:194:PHE:CD2	2.34	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:03:27:ILE:HA	52:03:30:LEU:CD2	2.29	0.62
52:03:40:GLU:HB2	52:03:217:THR:OG1	1.98	0.62
42:L:109:ARG:NH1	53:A:537:G:H5''	2.13	0.62
59:Z:297:THR:HG23	59:Z:298:ILE:HD12	1.81	0.62
8:11:74:PRO:HG2	8:11:77:VAL:HG22	1.82	0.62
36:F:3:HIS:O	36:F:92:THR:HG22	1.99	0.62
54:01:1173:U:H2'	54:01:1174:U:H4'	1.81	0.62
1:04:16:VAL:HB	1:04:203:VAL:HG22	1.81	0.62
34:D:127:ARG:HH21	53:A:619:U:H4'	1.64	0.62
43:M:52:ILE:HG22	43:M:56:ARG:NH1	2.15	0.62
59:Z:129:TYR:HB3	59:Z:199:ILE:CD1	2.29	0.62
59:Z:231:VAL:HB	59:Z:270:ALA:HA	1.81	0.62
13:16:2:ARG:HG2	54:01:1653:G:H3'	1.82	0.62
14:17:56:LYS:O	14:17:60:GLU:HG3	2.00	0.62
25:28:37:ARG:HH21	54:01:929:U:H4'	1.65	0.62
54:01:1509:A:H2'	54:01:1510:G:C8	2.34	0.62
52:03:65:LEU:HD22	52:03:188:ASN:HA	1.81	0.62
58:Y:25:C:C3'	58:Y:26:A:H5''	2.29	0.62
59:Z:89:LYS:HD3	59:Z:288:ARG:NH2	2.14	0.62
3:06:122:GLU:HG3	3:06:123:LYS:H	1.65	0.62
32:B:216:VAL:O	32:B:220:VAL:HG23	1.98	0.62
37:G:112:ASP:H	37:G:118:ARG:HD3	1.63	0.62
44:N:85:GLU:HG3	44:N:89:ARG:HH22	1.63	0.62
52:03:31:LYS:HE3	52:03:181:ASP:HA	1.82	0.62
53:A:81:A:H2	53:A:88:U:H3	1.46	0.62
56:W:47:U:H3'	56:W:48:C:H5'	1.81	0.62
17:20:24:LYS:HA	17:20:94:THR:OG1	1.99	0.62
32:B:22:TRP:CZ3	32:B:24:PRO:HA	2.35	0.62
43:M:52:ILE:HG22	43:M:56:ARG:HH12	1.64	0.62
53:A:552:U:H2'	53:A:553:A:C8	2.34	0.62
54:01:1869:G:H3'	54:01:1870:C:C5'	2.30	0.62
59:Z:213:PRO:HD3	59:Z:232:GLU:HB2	1.81	0.62
54:01:310:A:C2'	54:01:311:A:H5''	2.30	0.62
59:Z:14:VAL:CG1	59:Z:195:LEU:HD21	2.29	0.62
2:05:135:GLY:HA2	54:01:743:A:OP1	1.99	0.61
53:A:1129:C:H2'	53:A:1139:G:N7	2.15	0.61
37:G:22:LEU:O	37:G:26:VAL:HG23	1.99	0.61
54:01:296:U:H2'	54:01:297:G:C8	2.36	0.61
54:01:2800:A:C2	54:01:2895:G:H1'	2.35	0.61
59:Z:7:ARG:NH2	59:Z:269:ARG:HE	1.93	0.61
59:Z:9:LYS:HB3	59:Z:74:ARG:HA	1.83	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:Z:155:GLU:HA	59:Z:158:SER:HB2	1.82	0.61
4:07:114:ARG:HG3	4:07:177:ARG:HE	1.65	0.61
49:S:3:SER:HA	53:A:1314:C:H41	1.65	0.61
8:11:89:SER:HB2	8:11:92:PRO:HG3	1.82	0.61
16:19:24:TYR:HE1	54:01:17:G:H4'	1.66	0.61
20:23:80:ASP:OD2	20:23:95:PHE:HB3	2.00	0.61
33:C:39:ARG:HG3	33:C:54:ILE:HD11	1.83	0.61
40:J:40:ILE:HB	40:J:73:LEU:HB2	1.80	0.61
59:Z:7:ARG:HH12	59:Z:272:GLU:HG3	1.64	0.61
3:06:154:ASP:OD2	3:06:156:ASN:HB3	2.01	0.61
6:09:47:PHE:HA	6:09:51:ARG:HD2	1.81	0.61
26:29:11:GLU:HA	26:29:25:ARG:HA	1.83	0.61
45:O:23:SER:HB3	45:O:26:VAL:HG23	1.82	0.61
54:01:296:U:H2'	54:01:297:G:H8	1.64	0.61
54:01:1068:G:H21	54:01:1096:A:H5'	1.65	0.61
59:Z:106:ALA:HB3	59:Z:109:ASP:OD2	2.01	0.61
58:Y:51:U:H4'	59:Z:325:GLY:O	2.00	0.61
36:F:64:VAL:HG22	36:F:65:GLU:N	2.16	0.61
47:Q:12:VAL:HB	47:Q:21:VAL:HG13	1.83	0.61
53:A:513:C:H2'	53:A:514:C:C6	2.36	0.61
9:12:17:VAL:HG23	9:12:137:PRO:HB2	1.83	0.61
15:18:4:ILE:O	15:18:8:GLU:HG3	2.01	0.61
18:21:88:ARG:HG3	18:21:94:ASP:OD2	2.01	0.61
59:Z:24:LYS:HG2	59:Z:104:VAL:HG21	1.81	0.61
44:N:53:ASP:HA	44:N:58:ARG:HD3	1.82	0.61
52:03:193:LEU:CD2	52:03:196:LEU:HD22	2.30	0.61
54:01:594:U:H2'	54:01:595:C:C6	2.35	0.61
4:07:40:GLY:HA3	54:01:2307:G:O6	2.00	0.60
8:11:52:LEU:O	8:11:54:ILE:HD12	2.01	0.60
54:01:2130:U:H5'	54:01:2159:G:N2	2.16	0.60
51:U:25:ALA:HB1	57:V:9:G:H4'	1.82	0.60
54:01:1683:U:H2'	54:01:1684:G:C8	2.36	0.60
59:Z:299:LYS:HD3	59:Z:300:PRO:O	2.01	0.60
5:08:95:ALA:HB1	5:08:130:ILE:HD11	1.82	0.60
22:25:33:ILE:HD11	22:25:78:ILE:HD11	1.83	0.60
37:G:41:ILE:HG23	37:G:116:ALA:HA	1.83	0.60
44:N:39:ASP:HA	44:N:42:ASN:HD22	1.67	0.60
56:X:14:A:H2'	56:X:15:G:O4'	2.00	0.60
61:Y:101:PHE:O	59:Z:261:PHE:HA	2.01	0.60
3:06:71:GLY:N	54:01:674:G:H5''	2.16	0.60
9:12:41:LYS:HB3	9:12:43:GLU:OE1	2.01	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:15:3:GLN:HE21	12:15:92:TRP:HE1	1.50	0.60
18:21:66:ILE:HD12	18:21:66:ILE:N	2.15	0.60
33:C:9:ILE:HG23	33:C:10:ARG:HG3	1.81	0.60
54:01:1077:A:H2'	54:01:1078:U:H5'	1.83	0.60
59:Z:27:LEU:HD21	59:Z:104:VAL:HG21	1.84	0.60
59:Z:82:PRO:HD2	59:Z:91:MET:HB2	1.83	0.60
12:15:40:ARG:NH1	12:15:73:ILE:HG13	2.17	0.60
53:A:1259:C:H3'	53:A:1260:G:C5'	2.31	0.60
53:A:1513:A:H2'	53:A:1514:G:H8	1.66	0.60
54:01:878:A:H3'	54:01:879:G:H8	1.66	0.60
54:01:1026:G:H2'	54:01:1027:A:H8	1.65	0.60
54:01:1055:G:H2'	54:01:1056:G:O4'	2.00	0.60
54:01:1807:G:H2'	54:01:1808:A:H5'	1.84	0.60
59:Z:19:HIS:HD2	59:Z:114:GLN:HB2	1.67	0.60
17:20:60:LYS:HB2	17:20:100:GLY:HA3	1.84	0.60
21:24:28:ALA:HB3	21:24:40:ILE:HB	1.83	0.60
42:L:56:LEU:HD12	42:L:60:PHE:HB2	1.82	0.60
59:Z:343:LEU:HA	59:Z:358:MET:CB	2.30	0.60
10:13:3:GLN:NE2	54:01:1666:G:H1'	2.16	0.60
15:18:7:LEU:HA	15:18:10:GLU:OE2	2.02	0.60
34:D:7:LYS:HG2	53:A:430:A:OP2	2.01	0.60
54:01:1056:G:H4'	54:01:1086:A:H8	1.67	0.60
54:01:2172:U:H4'	54:01:2174:C:H5	1.65	0.60
53:A:552:U:H2'	53:A:553:A:H8	1.66	0.60
59:Z:1:SER:HB2	59:Z:2:LYS:HD2	1.83	0.60
7:10:78:GLY:N	7:10:79:PRO:HD2	2.16	0.60
35:E:82:HIS:HB2	35:E:83:PRO:HD2	1.82	0.60
41:K:63:GLN:O	41:K:67:GLU:HG3	2.02	0.60
42:L:48:LEU:HB3	53:A:520:A:OP1	2.01	0.60
53:A:695:A:H2'	53:A:696:A:C8	2.36	0.60
55:02:66:A:H5''	55:02:67:G:OP1	2.01	0.60
56:X:21:A:H61	56:X:46:G:H2'	1.66	0.60
2:05:124:ARG:NH1	2:05:163:GLY:HA3	2.17	0.60
16:19:16:ILE:HG13	16:19:31:TYR:HE1	1.65	0.60
21:24:47:VAL:O	21:24:51:GLN:HG2	2.01	0.60
25:28:35:VAL:HG22	25:28:36:GLU:H	1.66	0.60
39:I:33:SER:H	39:I:36:GLN:HE21	1.49	0.60
45:O:71:ARG:NH2	53:A:754:C:H5'	2.16	0.60
47:Q:58:VAL:HG23	47:Q:77:VAL:HA	1.83	0.60
54:01:2817:U:H3'	54:01:2818:U:H5''	1.83	0.60
53:A:219:U:H2'	53:A:220:G:H8	1.67	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:A:399:G:H2'	53:A:400:C:C6	2.37	0.59
54:01:310:A:O2'	54:01:311:A:H5''	2.01	0.59
54:01:1068:G:N2	54:01:1096:A:H5'	2.16	0.59
54:01:1251:C:O2'	54:01:1252:G:H3'	2.02	0.59
54:01:1386:C:H2'	54:01:1387:A:H8	1.66	0.59
18:21:66:ILE:HD12	18:21:66:ILE:H	1.68	0.59
21:24:4:ILE:HD13	21:24:47:VAL:HG22	1.83	0.59
31:34:32:LYS:HE3	54:01:2478:A:H5'	1.84	0.59
45:O:32:THR:HG22	45:O:36:ASN:HD21	1.67	0.59
51:U:66:ARG:CG	53:A:1099:G:H4'	2.31	0.59
54:01:215:G:H4'	54:01:216:A:H4'	1.83	0.59
59:Z:6:GLU:CG	59:Z:8:THR:HG23	2.31	0.59
13:16:37:THR:HA	13:16:110:MET:HE2	1.83	0.59
32:B:22:TRP:HA	32:B:189:ASN:HB3	1.84	0.59
37:G:142:ARG:HB3	56:X:41:C:H4'	1.84	0.59
43:M:94:LEU:HB3	43:M:95:PRO:HD2	1.83	0.59
53:A:1425:U:H2'	53:A:1426:G:H8	1.68	0.59
58:Y:28:G:H2'	58:Y:29:G:H8	1.67	0.59
11:14:101:ILE:HG13	11:14:102:GLY:N	2.16	0.59
5:08:41:GLU:HG3	5:08:54:ARG:HH21	1.66	0.59
14:17:43:ASN:ND2	14:17:45:SER:HB3	2.17	0.59
33:C:67:ILE:HD11	33:C:100:ILE:HD11	1.84	0.59
35:E:133:ILE:HD12	35:E:133:ILE:H	1.67	0.59
51:U:32:ARG:HG3	51:U:33:ARG:HG3	1.84	0.59
59:Z:277:LEU:HD12	59:Z:278:LEU:N	2.18	0.59
1:04:6:LYS:HD2	54:01:705:A:H4'	1.85	0.59
4:07:175:PRO:HB3	26:29:47:LYS:NZ	2.17	0.59
5:08:137:LYS:HE2	54:01:2746:U:H5''	1.84	0.59
6:09:39:ALA:HA	6:09:43:ASN:HB2	1.84	0.59
52:03:65:LEU:HD11	52:03:175:ILE:HG22	1.84	0.59
48:R:17:VAL:HG22	48:R:18:GLN:H	1.68	0.59
54:01:2512:C:H2'	54:01:2513:A:O4'	2.03	0.59
1:04:144:GLU:HA	1:04:151:GLY:HA2	1.85	0.59
1:04:153:LEU:HD11	1:04:181:ARG:NH2	2.18	0.59
4:07:111:ARG:HD3	43:M:6:ILE:HG23	1.84	0.59
8:11:89:SER:O	8:11:91:LYS:N	2.35	0.59
32:B:221:ARG:HH11	32:B:224:ARG:HH11	1.49	0.59
33:C:113:LYS:HA	33:C:184:ASN:ND2	2.17	0.59
40:J:53:ILE:HD12	40:J:63:ASP:OD2	2.03	0.59
48:R:70:THR:HG23	48:R:71:ASP:N	2.12	0.59
53:A:235:C:H2'	53:A:236:A:C8	2.37	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:A:1206:G:C2'	53:A:1207:G:H5''	2.31	0.59
59:Z:60:ILE:HG13	59:Z:62:ILE:HG23	1.84	0.59
59:Z:128:PRO:HA	59:Z:162:PHE:HE1	1.68	0.59
59:Z:243:GLU:HG2	59:Z:245:VAL:HG22	1.85	0.59
41:K:88:PRO:HD3	51:U:28:LEU:HD22	1.85	0.59
53:A:695:A:H2'	53:A:696:A:H8	1.68	0.59
53:A:760:G:H2'	53:A:761:G:H5'	1.83	0.59
54:01:2760:C:O2'	54:01:2761:A:H5'	2.03	0.59
1:04:52:HIS:NE2	1:04:218:THR:HG23	2.18	0.59
54:01:1020:A:H1'	54:01:1021:A:OP2	2.03	0.59
59:Z:227:VAL:HG13	59:Z:276:VAL:HB	1.85	0.59
1:04:239:PHE:HZ	54:01:1826:G:H4'	1.68	0.58
10:13:113:MET:SD	10:13:116:ILE:HD11	2.43	0.58
54:01:1713:A:H61	54:01:1745:A:H61	1.51	0.58
59:Z:208:LYS:HB3	59:Z:233:ARG:NH2	2.18	0.58
7:10:14:GLU:O	7:10:18:VAL:HG23	2.03	0.58
39:I:98:ARG:HG3	39:I:103:VAL:HG21	1.85	0.58
59:Z:133:PHE:HD1	59:Z:170:VAL:HG23	1.67	0.58
9:12:81:ILE:HG23	9:12:82:GLY:N	2.18	0.58
48:R:59:LYS:HD3	53:A:735:C:H5'	1.83	0.58
53:A:1170:A:H2'	53:A:1171:A:O4'	2.03	0.58
54:01:45:G:H5''	54:01:46:G:C5'	2.21	0.58
54:01:1857:G:H2'	54:01:1884:G:N2	2.18	0.58
6:09:30:LEU:HB3	6:09:36:ALA:HB3	1.85	0.58
15:18:5:LYS:O	15:18:9:GLN:HG2	2.03	0.58
40:J:10:LEU:HD12	40:J:10:LEU:O	2.03	0.58
44:N:9:GLU:O	44:N:13:VAL:HG23	2.03	0.58
53:A:1005:A:H2'	53:A:1006:G:O4'	2.03	0.58
54:01:2452:C:N4	54:01:2504:U:H3	1.98	0.58
18:21:55:ILE:O	18:21:59:GLU:HG2	2.03	0.58
22:25:36:GLN:HE22	22:25:39:THR:HA	1.65	0.58
38:H:54:THR:HG23	38:H:55:LYS:HG3	1.84	0.58
54:01:1222:U:H2'	54:01:1223:G:C8	2.38	0.58
59:Z:244:ILE:HG23	59:Z:244:ILE:O	2.03	0.58
52:03:5:THR:HB	54:01:2129:C:H5'	1.86	0.58
54:01:720:U:H2'	54:01:721:A:H8	1.69	0.58
59:Z:258:VAL:HG22	59:Z:276:VAL:HG22	1.86	0.58
1:04:163:ILE:HA	1:04:173:LEU:HD23	1.85	0.58
32:B:16:GLY:O	32:B:17:HIS:CB	2.51	0.58
32:B:35:ASN:HB3	32:B:37:VAL:HG12	1.84	0.58
52:03:9:ARG:O	52:03:13:GLU:HG3	2.04	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:01:1801:A:H5''	54:01:2203:U:H2'	1.85	0.58
54:01:2699:C:H2'	54:01:2700:A:C8	2.38	0.58
8:11:105:LEU:HD23	8:11:108:ILE:HD12	1.86	0.58
15:18:59:THR:HG22	15:18:72:VAL:HG12	1.85	0.58
21:24:30:ILE:HG13	21:24:40:ILE:HG13	1.86	0.58
33:C:13:ILE:HG22	33:C:14:VAL:HG23	1.86	0.58
52:03:19:LYS:HD3	52:03:21:TYR:HE1	1.69	0.58
54:01:2208:C:H2'	54:01:2209:G:C8	2.39	0.58
58:Y:62:C:H2'	58:Y:63:G:O4'	2.03	0.58
8:11:59:THR:HB	8:11:67:THR:HG23	1.85	0.58
16:19:43:GLN:HE21	17:20:77:PHE:HB3	1.69	0.58
38:H:85:TYR:CE1	38:H:123:GLU:HB2	2.39	0.58
39:I:17:ARG:HH22	53:A:1129:C:H5''	1.69	0.58
52:03:69:THR:HG22	52:03:161:VAL:HG12	1.86	0.58
53:A:662:U:H2'	53:A:663:A:C8	2.38	0.58
54:01:1437:C:H2'	54:01:1438:U:C6	2.39	0.58
54:01:1440:U:H2'	54:01:1441:G:C8	2.39	0.58
54:01:2537:U:H2'	54:01:2538:C:C6	2.39	0.58
2:05:179:ARG:HB3	2:05:188:LEU:HD12	1.86	0.58
41:K:23:HIS:HB3	41:K:30:ILE:HG23	1.85	0.58
41:K:111:ASP:HB3	51:U:3:ILE:HG23	1.85	0.58
23:26:17:ARG:HE	23:26:23:ALA:HB2	1.69	0.57
51:U:65:ARG:HH12	53:A:1088:G:H5'	1.68	0.57
54:01:2514:U:H2'	54:01:2515:C:C6	2.39	0.57
16:19:43:GLN:NE2	17:20:77:PHE:HB3	2.19	0.57
47:Q:69:THR:HG22	47:Q:70:LYS:H	1.68	0.57
56:X:32:C:H2'	56:X:33:U:C2	2.39	0.57
58:Y:25:C:H2'	58:Y:26:A:H5''	1.85	0.57
36:F:73:GLU:O	36:F:77:THR:HG23	2.04	0.57
51:U:4:LYS:HD3	51:U:6:ARG:NH1	2.19	0.57
53:A:70:U:H5''	53:A:71:A:OP1	2.05	0.57
53:A:580:C:H2'	53:A:581:G:O4'	2.04	0.57
54:01:1440:U:H2'	54:01:1441:G:H8	1.68	0.57
35:E:93:VAL:HG11	35:E:139:THR:HG22	1.86	0.57
38:H:12:ARG:NH2	53:A:826:C:H5'	2.20	0.57
52:03:33:LEU:HD13	52:03:220:ALA:H	1.68	0.57
54:01:1053:C:C2'	54:01:1054:A:H5''	2.33	0.57
54:01:2287:A:O2'	54:01:2288:A:H2'	2.04	0.57
54:01:2638:G:HO2'	54:01:2639:A:H8	1.51	0.57
32:B:20:ARG:HG3	32:B:21:TYR:N	2.17	0.57
54:01:640:C:H2'	54:01:641:U:H6	1.69	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:Z:259:GLU:HG3	59:Z:263:LYS:C	2.25	0.57
6:09:76:GLU:HB3	6:09:142:VAL:HG22	1.85	0.57
43:M:24:VAL:HA	53:A:1329:A:H5''	1.87	0.57
54:01:1675:C:H2'	54:01:1676:A:O4'	2.04	0.57
59:Z:191:LEU:HA	59:Z:194:PHE:HD2	1.68	0.57
32:B:56:LEU:HD21	32:B:183:PHE:HZ	1.69	0.57
46:P:36:VAL:HG23	46:P:53:ASP:HB3	1.87	0.57
54:01:2074:U:H2'	54:01:2075:U:C6	2.39	0.57
54:01:2629:U:O2'	54:01:2630:G:H5''	2.05	0.57
55:02:104:A:H2'	55:02:105:G:O4'	2.04	0.57
2:05:91:THR:HG23	2:05:94:GLN:HB2	1.87	0.57
10:13:3:GLN:HE22	54:01:1666:G:H1'	1.70	0.57
12:15:6:ARG:O	12:15:6:ARG:HD3	2.05	0.57
35:E:113:VAL:HG13	35:E:114:LEU:HD12	1.86	0.57
56:X:31:G:H2'	56:X:32:C:H5'	1.86	0.57
40:J:20:GLN:O	40:J:24:GLU:HG3	2.04	0.57
46:P:46:LYS:HD2	53:A:617:G:H4'	1.86	0.57
54:01:703:U:H2'	54:01:704:G:O4'	2.05	0.57
54:01:1539:U:H2'	54:01:1540:G:H8	1.70	0.57
54:01:2818:U:H2'	54:01:2819:G:H8	1.70	0.57
59:Z:14:VAL:HG21	59:Z:69:TYR:OH	2.05	0.57
32:B:174:GLU:HA	32:B:177:ASN:HD22	1.68	0.57
36:F:5:GLU:O	36:F:7:VAL:HG23	2.05	0.57
37:G:58:LEU:HD12	37:G:59:GLU:N	2.20	0.57
37:G:129:ASN:HB2	37:G:134:VAL:HG11	1.87	0.57
53:A:80:A:H62	53:A:86:G:H21	1.53	0.57
53:A:737:C:H2'	53:A:738:C:C6	2.40	0.57
53:A:1417:G:H2'	53:A:1482:G:N2	2.20	0.57
54:01:161:A:H3'	54:01:162:U:H5''	1.86	0.57
54:01:609:A:H2'	54:01:610:C:O4'	2.05	0.57
33:C:134:LYS:O	33:C:138:GLN:HG3	2.05	0.56
41:K:110:THR:HB	48:R:72:ARG:HH12	1.69	0.56
46:P:6:LEU:HD22	46:P:17:TYR:HB3	1.86	0.56
3:06:155:GLU:HG2	3:06:159:LEU:HG	1.86	0.56
26:29:58:ASP:O	26:29:62:LYS:HG3	2.05	0.56
49:S:28:LYS:HE3	49:S:29:PRO:HD2	1.87	0.56
53:A:1052:U:H2'	53:A:1200:C:H41	1.70	0.56
53:A:1070:U:H2'	53:A:1071:C:C6	2.40	0.56
54:01:1060:U:H5'	54:01:1062:G:H5'	1.86	0.56
54:01:1367:A:H2'	54:01:1368:G:H5'	1.86	0.56
54:01:2070:A:H2'	54:01:2071:A:C8	2.40	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:10:26:VAL:HB	7:10:82:ILE:HD12	1.87	0.56
8:11:11:GLN:HB3	8:11:55:PRO:HA	1.87	0.56
10:13:76:VAL:H	15:18:72:VAL:CG2	2.19	0.56
16:19:108:LEU:HA	17:20:48:LYS:HE2	1.88	0.56
35:E:156:ARG:NH1	35:E:163:ILE:HB	2.20	0.56
38:H:28:SER:HB3	38:H:56:PRO:HB2	1.87	0.56
39:I:80:HIS:HE1	39:I:84:ARG:HH11	1.53	0.56
52:03:7:ARG:HD2	52:03:7:ARG:O	2.04	0.56
52:03:19:LYS:HD3	52:03:21:TYR:CE1	2.41	0.56
53:A:1342:C:H2'	53:A:1343:G:C8	2.41	0.56
54:01:1532:A:H1'	54:01:1540:G:N2	2.21	0.56
54:01:1880:U:H2'	54:01:1881:C:C6	2.41	0.56
59:Z:106:ALA:HB2	59:Z:136:LYS:HD3	1.86	0.56
2:05:46:ARG:HG2	2:05:84:LEU:HD12	1.88	0.56
35:E:96:GLN:HG2	35:E:98:ALA:H	1.70	0.56
41:K:51:PHE:CE2	41:K:64:VAL:HG11	2.41	0.56
46:P:36:VAL:HG21	46:P:57:ILE:HG13	1.86	0.56
52:03:4:LEU:HB3	52:03:9:ARG:NE	2.16	0.56
58:Y:26:A:H61	58:Y:45:U:H3	1.51	0.56
59:Z:147:GLU:HA	59:Z:171:ARG:HH22	1.68	0.56
1:04:15:VAL:HG22	1:04:205:GLY:HA3	1.87	0.56
4:07:107:VAL:HB	4:07:108:PRO:HD3	1.88	0.56
19:22:59:ASN:HD22	54:01:1342:A:H5''	1.69	0.56
22:25:66:GLU:HB3	22:25:68:LYS:HG3	1.87	0.56
24:27:49:ASP:O	24:27:53:VAL:HG23	2.05	0.56
31:34:4:ARG:O	31:34:37:GLN:HB3	2.06	0.56
43:M:3:ILE:HG13	43:M:7:ASN:HB3	1.87	0.56
45:O:7:THR:O	45:O:11:VAL:HG23	2.06	0.56
51:U:17:ARG:HA	51:U:20:ARG:NH1	2.21	0.56
53:A:663:A:H5'	53:A:836:G:OP1	2.05	0.56
54:01:1326:U:H2'	54:01:1327:A:H8	1.71	0.56
54:01:2170:A:H2'	54:01:2171:A:O4'	2.05	0.56
54:01:2698:U:H2'	54:01:2699:C:C6	2.40	0.56
54:01:581:C:H2'	54:01:582:A:C8	2.41	0.56
54:01:2785:C:H2'	54:01:2786:U:C6	2.41	0.56
59:Z:28:THR:HG23	59:Z:78:HIS:HD2	1.69	0.56
2:05:10:GLY:H	2:05:197:THR:HG23	1.70	0.56
11:14:64:PHE:HB2	54:01:2416:C:OP1	2.05	0.56
40:J:22:THR:O	40:J:26:VAL:HG23	2.06	0.56
53:A:714:G:H2'	53:A:715:A:C8	2.41	0.56
59:Z:219:SER:HB2	59:Z:283:ARG:HD2	1.87	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:10:34:THR:O	7:10:38:MET:HG3	2.06	0.56
9:12:78:THR:HG22	54:01:2641:G:H5''	1.88	0.56
18:21:42:LYS:HB2	54:01:2010:G:H5''	1.87	0.56
54:01:704:G:H2'	54:01:726:G:N2	2.21	0.56
54:01:2121:G:H2'	54:01:2122:U:O4'	2.06	0.56
59:Z:132:VAL:HB	59:Z:169:ILE:HD12	1.87	0.56
59:Z:167:THR:HG22	59:Z:169:ILE:HD13	1.88	0.56
14:17:43:ASN:HD21	14:17:46:GLU:HG2	1.70	0.56
15:18:24:THR:HB	15:18:87:ARG:H	1.71	0.56
32:B:89:PHE:HE1	32:B:152:ASP:HB2	1.70	0.56
32:B:212:TYR:O	32:B:216:VAL:HG23	2.06	0.56
54:01:1936:A:H2	54:01:1943:U:H3	1.54	0.56
59:Z:4:LYS:HG2	59:Z:264:LEU:HB2	1.87	0.56
3:06:149:ILE:HD11	3:06:172:ALA:HA	1.88	0.56
10:13:12:ASP:HB2	10:13:96:GLY:HA3	1.88	0.56
12:15:20:LEU:HD23	21:24:81:PRO:HG2	1.87	0.56
12:15:29:GLY:HA2	12:15:106:ASP:HB2	1.88	0.56
15:18:38:ARG:NH2	15:18:40:GLN:HB3	2.21	0.56
32:B:89:PHE:HB3	32:B:149:GLY:O	2.05	0.56
53:A:304:U:H2'	53:A:305:G:C8	2.41	0.56
53:A:1352:C:H2'	53:A:1353:G:C8	2.41	0.56
3:06:4:VAL:HG12	3:06:11:ALA:HB2	1.88	0.55
3:06:24:ASN:O	3:06:28:VAL:HG23	2.06	0.55
6:09:79:THR:HG23	6:09:145:ASN:O	2.06	0.55
10:13:14:SER:OG	10:13:86:LEU:HD12	2.06	0.55
14:17:52:SER:OG	14:17:54:VAL:HG12	2.06	0.55
19:22:21:SER:O	19:22:25:GLU:HG2	2.05	0.55
27:30:30:ASP:HB3	27:30:34:GLY:H	1.70	0.55
31:34:15:LYS:HB2	31:34:15:LYS:NZ	2.21	0.55
34:D:61:ARG:HH21	34:D:67:LEU:HA	1.71	0.55
53:A:797:C:H2'	53:A:798:U:C6	2.41	0.55
54:01:640:C:H2'	54:01:641:U:C6	2.41	0.55
54:01:2875:C:H2'	54:01:2876:G:H8	1.71	0.55
59:Z:322:PHE:CZ	59:Z:350:VAL:HB	2.41	0.55
2:05:148:GLN:HB2	2:05:152:PRO:HG2	1.88	0.55
3:06:181:ILE:HG23	11:14:2:ARG:HG3	1.88	0.55
4:07:97:GLU:HG2	26:29:25:ARG:HB3	1.88	0.55
16:19:36:GLN:HE21	54:01:1252:G:H1	1.54	0.55
24:27:9:LYS:HB3	24:27:12:GLU:HB2	1.88	0.55
25:28:23:LEU:HD22	25:28:28:LEU:HD12	1.86	0.55
32:B:27:LYS:HB3	32:B:28:PRO:HD3	1.89	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:K:71:ASP:HA	41:K:74:LYS:HE2	1.88	0.55
42:L:33:CYS:HA	42:L:54:VAL:HA	1.88	0.55
52:03:193:LEU:O	52:03:197:LYS:HG3	2.06	0.55
53:A:935:A:H2'	53:A:936:C:C6	2.42	0.55
53:A:1286:U:H2'	53:A:1287:A:H5'	1.88	0.55
54:01:275:C:H3'	54:01:276:U:H5''	1.88	0.55
54:01:833:A:H2'	54:01:834:G:H8	1.71	0.55
59:Z:214:ILE:CD1	59:Z:290:GLN:HB3	2.31	0.55
59:Z:300:PRO:HB2	59:Z:366:ILE:N	2.21	0.55
1:04:71:ASP:O	1:04:73:ILE:HG13	2.06	0.55
5:08:136:ASP:OD2	5:08:139:VAL:HG23	2.06	0.55
52:03:214:ILE:CD1	52:03:222:VAL:HB	2.36	0.55
53:A:666:G:H2'	53:A:667:G:H8	1.72	0.55
54:01:1796:U:H2'	54:01:1797:G:H8	1.71	0.55
59:Z:306:SER:HB3	59:Z:358:MET:CE	2.37	0.55
8:11:91:LYS:HG3	8:11:94:LYS:CE	2.32	0.55
12:15:3:GLN:NE2	12:15:92:TRP:HE1	2.04	0.55
23:26:61:LYS:HE3	54:01:372:G:OP1	2.06	0.55
38:H:17:GLN:HE21	38:H:71:VAL:H	1.54	0.55
49:S:77:ARG:HD2	53:A:1225:A:H1'	1.88	0.55
53:A:1412:C:H2'	53:A:1413:A:H8	1.70	0.55
54:01:481:G:H2'	54:01:507:A:N1	2.20	0.55
54:01:1837:C:H2'	54:01:1899:A:H61	1.71	0.55
59:Z:243:GLU:HG3	59:Z:251:GLN:O	2.07	0.55
32:B:45:THR:HG22	32:B:49:PHE:HD1	1.70	0.55
51:U:49:ALA:O	51:U:53:LYS:HG3	2.06	0.55
54:01:100:U:H4'	54:01:101:A:O4'	2.07	0.55
1:04:257:ARG:HH11	1:04:257:ARG:HG2	1.71	0.55
6:09:132:PHE:O	6:09:139:PHE:HB3	2.07	0.55
38:H:76:ARG:NH1	38:H:125:ILE:HG23	2.21	0.55
41:K:88:PRO:HG2	41:K:89:GLY:H	1.71	0.55
52:03:222:VAL:HG12	52:03:224:VAL:HG13	1.89	0.55
53:A:123:U:H5''	53:A:311:C:O2'	2.06	0.55
53:A:909:A:H2'	53:A:910:C:O4'	2.07	0.55
53:A:1342:C:H2'	53:A:1343:G:H8	1.72	0.55
54:01:1170:C:H2'	54:01:1171:G:C8	2.42	0.55
59:Z:238:VAL:HG13	59:Z:257:GLY:N	2.21	0.55
6:09:6:LEU:HD11	6:09:37:VAL:HG23	1.88	0.55
14:17:76:LYS:O	14:17:80:GLU:HG3	2.07	0.55
17:20:83:TYR:CZ	54:01:1187:G:H5''	2.42	0.55
54:01:1278:C:H2'	54:01:1279:G:H8	1.71	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:Z:173:SER:N	59:Z:184:TRP:CE3	2.73	0.55
1:04:140:VAL:HG12	1:04:191:LEU:HD23	1.88	0.55
12:15:5:LYS:O	12:15:6:ARG:HG3	2.07	0.55
25:28:15:ARG:HE	25:28:52:PHE:HE2	1.55	0.55
32:B:183:PHE:CB	32:B:197:PHE:HB2	2.35	0.55
38:H:86:LYS:HD2	38:H:90:GLU:HG2	1.88	0.55
53:A:20:U:H2'	53:A:21:G:O4'	2.07	0.55
30:33:6:VAL:HG21	30:33:60:CYS:HB2	1.87	0.55
33:C:13:ILE:H	33:C:13:ILE:HD12	1.72	0.55
42:L:49:ARG:HB3	42:L:65:TYR:HE1	1.72	0.55
49:S:16:LYS:NZ	49:S:16:LYS:HB3	2.22	0.55
52:03:5:THR:O	52:03:9:ARG:HG3	2.06	0.55
53:A:181:A:N6	53:A:194:C:H2'	2.22	0.55
53:A:1513:A:H2'	53:A:1514:G:C8	2.42	0.55
54:01:685:A:H5''	54:01:788:A:H62	1.72	0.55
59:Z:7:ARG:NH1	59:Z:272:GLU:HG3	2.22	0.55
59:Z:155:GLU:O	59:Z:159:GLN:HG2	2.07	0.55
1:04:240:GLY:HA3	54:01:2597:G:C5'	2.36	0.55
2:05:82:PHE:HE1	2:05:202:ILE:HG23	1.71	0.55
15:18:105:LYS:O	15:18:108:ARG:HG2	2.06	0.55
28:31:18:HIS:HB3	28:31:39:ASP:OD1	2.06	0.55
54:01:2185:U:H2'	54:01:2186:G:H8	1.72	0.55
54:01:2233:U:H2'	54:01:2234:G:H8	1.72	0.55
59:Z:213:PRO:HB2	59:Z:334:THR:HG21	1.88	0.55
4:07:70:ARG:HG2	4:07:70:ARG:HH21	1.71	0.54
6:09:84:ALA:HB2	6:09:90:LEU:HD12	1.89	0.54
12:15:42:THR:OG1	12:15:45:GLN:HG3	2.07	0.54
19:22:56:GLU:OE2	19:22:88:LYS:HG2	2.07	0.54
22:25:39:THR:HG21	54:01:2336:A:H61	1.71	0.54
46:P:2:VAL:HG23	46:P:65:ALA:HA	1.88	0.54
53:A:312:C:H2'	53:A:313:A:H8	1.72	0.54
53:A:1206:G:C3'	53:A:1207:G:H5''	2.36	0.54
54:01:414:C:H2'	54:01:415:A:C8	2.41	0.54
54:01:2286:G:H5''	54:01:2287:A:OP1	2.07	0.54
59:Z:332:PHE:HB3	59:Z:372:LEU:HD21	1.89	0.54
7:10:124:ASP:HB3	7:10:126:LEU:HG	1.89	0.54
8:11:46:ASP:OD1	8:11:50:LYS:HD3	2.07	0.54
52:03:14:LYS:HD2	52:03:33:LEU:HD23	1.89	0.54
52:03:177:LYS:H	52:03:180:PHE:HD2	1.53	0.54
59:Z:170:VAL:HG21	59:Z:191:LEU:HB2	1.89	0.54
59:Z:176:LYS:HD3	59:Z:184:TRP:NE1	2.22	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:J:15:HIS:HB3	40:J:70:HIS:NE2	2.22	0.54
47:Q:48:GLU:HB2	47:Q:51:GLU:OE2	2.07	0.54
51:U:18:PHE:HA	51:U:21:SER:HB3	1.89	0.54
53:A:1062:U:H2'	53:A:1063:C:C6	2.42	0.54
54:01:2286:G:H5'	54:01:2287:A:H1'	1.89	0.54
57:V:19:U:H2'	57:V:20:U:H6	1.71	0.54
8:11:56:VAL:HG23	8:11:70:THR:HA	1.88	0.54
39:I:125:GLN:HE22	53:A:1342:C:H1'	1.71	0.54
41:K:80:ASN:HB3	41:K:105:ARG:HH11	1.72	0.54
44:N:20:PHE:O	44:N:21:ALA:HB3	2.07	0.54
47:Q:13:SER:H	47:Q:21:VAL:HG13	1.73	0.54
52:03:26:ALA:O	52:03:29:LEU:HG	2.08	0.54
53:A:335:C:H2'	53:A:336:A:C8	2.42	0.54
53:A:918:A:H2'	53:A:919:A:C8	2.42	0.54
54:01:460:A:H2'	54:01:461:C:O4'	2.07	0.54
54:01:1872:A:H2'	54:01:1873:G:O4'	2.08	0.54
54:01:2105:U:H2'	54:01:2106:U:O4'	2.07	0.54
54:01:2432:A:H1'	56:X:75:C:H5'	1.88	0.54
54:01:2636:C:H2'	54:01:2637:U:C6	2.42	0.54
55:02:5:U:H2'	55:02:6:G:C8	2.43	0.54
12:15:122:ALA:HB1	54:01:2467:C:H1'	1.90	0.54
13:16:39:PRO:HG2	54:01:1651:G:H4'	1.90	0.54
21:24:75:GLN:HB2	21:24:92:VAL:HG23	1.89	0.54
37:G:125:ASP:HA	37:G:128:GLU:OE2	2.08	0.54
50:T:30:PHE:HB3	50:T:53:MET:HB3	1.89	0.54
53:A:1448:C:H2'	53:A:1449:C:H6	1.72	0.54
54:01:873:C:H2'	54:01:874:G:H8	1.73	0.54
54:01:2159:G:H2'	54:01:2160:C:O4'	2.07	0.54
54:01:2573:C:H5''	54:01:2574:G:H5''	1.89	0.54
59:Z:12:VAL:HB	59:Z:76:TYR:CE1	2.43	0.54
1:04:12:ARG:HH21	54:01:728:G:H4'	1.71	0.54
7:10:59:LEU:HB2	7:10:62:ARG:HB2	1.90	0.54
18:21:20:VAL:HG21	18:21:43:ALA:HB3	1.88	0.54
36:F:18:VAL:HB	36:F:19:PRO:HD3	1.89	0.54
53:A:1448:C:H2'	53:A:1449:C:C6	2.42	0.54
54:01:144:A:H2'	54:01:145:C:C6	2.43	0.54
54:01:1213:A:N6	54:01:1236:G:H1'	2.23	0.54
12:15:4:PRO:HG2	12:15:70:ASP:HA	1.90	0.54
15:18:99:LEU:O	15:18:99:LEU:HD23	2.08	0.54
22:25:22:PHE:CD2	54:01:922:C:H1'	2.43	0.54
41:K:30:ILE:HD11	41:K:43:TRP:HB2	1.90	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:U:39:LYS:O	51:U:43:GLU:HG2	2.08	0.54
53:A:392:C:H2'	53:A:393:A:H8	1.72	0.54
54:01:2123:G:H2'	54:01:2124:G:O4'	2.07	0.54
54:01:2233:U:H2'	54:01:2234:G:C8	2.43	0.54
4:07:175:PRO:HB3	26:29:47:LYS:HZ1	1.73	0.54
13:16:2:ARG:O	13:16:2:ARG:HD3	2.08	0.54
27:30:24:VAL:HG13	27:30:25:THR:H	1.73	0.54
51:U:33:ARG:O	51:U:33:ARG:HD2	2.08	0.54
54:01:155:A:H2'	54:01:156:A:C8	2.43	0.54
54:01:2788:C:H2'	54:01:2789:C:C6	2.42	0.54
1:04:207:ALA:HB2	54:01:1790:C:O2'	2.07	0.54
4:07:37:MET:HB3	4:07:86:CYS:SG	2.48	0.54
7:10:99:PHE:HA	7:10:102:ALA:HB3	1.90	0.54
10:13:48:PRO:HB3	53:A:1422:G:H5'	1.90	0.54
23:26:63:ILE:O	23:26:67:LEU:HD13	2.08	0.54
34:D:59:LYS:O	34:D:63:ILE:HG13	2.08	0.54
54:01:27:G:N2	54:01:512:G:H1'	2.23	0.54
59:Z:63:ASN:HD22	59:Z:90:ASN:ND2	2.06	0.54
1:04:229:HIS:ND1	1:04:230:PRO:HD2	2.23	0.54
4:07:90:LEU:HD13	4:07:95:MET:HA	1.89	0.54
7:10:94:ARG:HD3	7:10:131:THR:HG22	1.90	0.54
8:11:126:ARG:HA	8:11:129:GLU:CG	2.38	0.54
13:16:29:VAL:HG11	13:16:75:ILE:HG23	1.89	0.54
32:B:66:ILE:HD12	32:B:159:ALA:HB3	1.89	0.54
36:F:32:ALA:HB2	36:F:70:VAL:HG21	1.89	0.54
48:R:56:ARG:HB3	48:R:60:ARG:NH1	2.23	0.54
52:03:21:TYR:HD2	52:03:222:VAL:HG13	1.71	0.54
53:A:935:A:H2'	53:A:936:C:H6	1.71	0.54
53:A:1207:G:H2'	53:A:1208:C:O4'	2.07	0.54
54:01:828:U:H2'	54:01:829:A:C8	2.43	0.54
59:Z:304:PHE:CD2	59:Z:388:VAL:HG13	2.43	0.54
59:Z:341:ILE:HA	59:Z:360:VAL:HG22	1.89	0.54
10:13:15:GLY:O	10:13:47:ILE:HG12	2.07	0.53
24:27:2:LYS:HE2	54:01:102:U:H1'	1.89	0.53
24:27:21:LEU:HA	24:27:25:GLN:HB3	1.90	0.53
36:F:19:PRO:O	36:F:23:GLU:HG3	2.08	0.53
47:Q:11:VAL:CG1	47:Q:20:ILE:HD11	2.38	0.53
53:A:82:G:H2'	53:A:83:C:O4'	2.08	0.53
53:A:1118:U:H2'	53:A:1119:C:C6	2.42	0.53
54:01:2875:C:H2'	54:01:2876:G:C8	2.43	0.53
5:08:71:LEU:O	5:08:75:VAL:HG23	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:10:37:LYS:HG2	7:10:41:LEU:HD12	1.91	0.53
7:10:64:VAL:O	7:10:68:PRO:HD3	2.08	0.53
30:33:57:VAL:HA	30:33:60:CYS:SG	2.48	0.53
53:A:412:A:H5''	53:A:413:G:OP2	2.08	0.53
54:01:1141:U:H4'	54:01:1142:A:O4'	2.08	0.53
54:01:2358:A:H2'	54:01:2359:C:O4'	2.08	0.53
59:Z:287:GLU:HG3	59:Z:290:GLN:HB2	1.88	0.53
33:C:156:LEU:HD11	33:C:163:ARG:HE	1.73	0.53
34:D:149:LYS:HZ2	34:D:177:MET:HB2	1.72	0.53
36:F:46:GLN:HA	36:F:56:LYS:HA	1.91	0.53
53:A:784:A:H2'	53:A:785:G:C8	2.43	0.53
53:A:825:A:H2'	53:A:826:C:C6	2.43	0.53
53:A:1395:C:H6	53:A:1395:C:H5'	1.72	0.53
54:01:239:C:H2'	54:01:240:C:O4'	2.08	0.53
54:01:362:A:H3'	54:01:363:G:H8	1.73	0.53
54:01:2597:G:H2'	54:01:2598:A:C8	2.44	0.53
54:01:2808:G:H2'	54:01:2890:G:C6	2.43	0.53
5:08:126:THR:HG22	5:08:127:GLN:H	1.74	0.53
19:22:39:THR:OG1	19:22:42:GLU:HG3	2.09	0.53
20:23:12:VAL:HB	20:23:18:LYS:HA	1.90	0.53
27:30:14:MET:SD	54:01:2045:C:H5''	2.49	0.53
33:C:19:SER:HB3	33:C:21:TRP:NE1	2.24	0.53
33:C:120:THR:HG23	33:C:188:ALA:HB2	1.88	0.53
41:K:27:ASN:HB3	41:K:56:LYS:HZ1	1.73	0.53
52:03:38:PHE:CE1	54:01:2127:G:H4'	2.43	0.53
53:A:10:A:H2'	53:A:11:G:H8	1.73	0.53
53:A:1070:U:H2'	53:A:1071:C:H6	1.74	0.53
54:01:2281:A:O2'	54:01:2282:G:H5'	2.08	0.53
54:01:2328:A:H2'	54:01:2329:U:C6	2.43	0.53
54:01:2339:C:H2'	54:01:2340:A:C8	2.44	0.53
56:X:36:U:H2'	56:X:37:A:O4'	2.08	0.53
58:Y:4:C:H2'	58:Y:5:G:C8	2.43	0.53
59:Z:332:PHE:HB2	59:Z:335:THR:OG1	2.09	0.53
3:06:48:THR:O	3:06:52:VAL:HG23	2.08	0.53
8:11:44:LYS:O	8:11:48:ILE:HG12	2.07	0.53
18:21:82:MET:HB2	18:21:98:LYS:HB2	1.90	0.53
54:01:1664:A:H61	54:01:1996:C:H42	1.57	0.53
54:01:1794:A:H2'	54:01:1795:C:C6	2.44	0.53
55:02:95:U:H2'	55:02:96:G:C8	2.43	0.53
56:X:13:C:C2'	56:X:14:A:H5''	2.37	0.53
59:Z:98:MET:HG2	59:Z:101:ALA:HB2	1.89	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:Z:209:PRO:HD2	59:Z:233:ARG:NH2	2.23	0.53
59:Z:305:GLU:HB3	59:Z:389:ALA:HB3	1.90	0.53
1:04:62:ARG:NH1	1:04:84:PRO:HD2	2.24	0.53
4:07:24:VAL:O	4:07:27:VAL:HG12	2.07	0.53
16:19:47:ARG:HG2	16:19:47:ARG:HH21	1.74	0.53
26:29:9:TYR:O	26:29:25:ARG:HD2	2.09	0.53
32:B:20:ARG:NH1	53:A:831:A:H5'	2.24	0.53
42:L:113:ARG:HD2	42:L:118:VAL:O	2.09	0.53
56:X:28:C:H2'	56:X:29:G:H8	1.73	0.53
5:08:97:VAL:HG23	5:08:124:CYS:SG	2.49	0.53
12:15:55:ARG:HD3	54:01:2469:A:H4'	1.91	0.53
22:25:22:PHE:HD2	54:01:922:C:H1'	1.72	0.53
27:30:54:ILE:HG23	27:30:56:LYS:H	1.74	0.53
39:I:47:VAL:HG12	39:I:78:ILE:HB	1.89	0.53
45:O:24:THR:O	45:O:28:VAL:HG23	2.09	0.53
54:01:1932:A:H2'	54:01:1933:G:O4'	2.09	0.53
54:01:2246:G:H2'	54:01:2247:A:C8	2.44	0.53
59:Z:16:THR:HG22	59:Z:24:LYS:HD2	1.91	0.53
59:Z:122:GLY:O	59:Z:125:VAL:HG12	2.08	0.53
59:Z:230:ARG:HB3	59:Z:273:ASN:HA	1.91	0.53
3:06:109:LEU:O	3:06:113:VAL:HG23	2.09	0.53
4:07:133:GLU:HB3	4:07:135:ILE:HG13	1.90	0.53
4:07:139:GLU:H	4:07:139:GLU:CD	2.12	0.53
8:11:79:LEU:O	8:11:83:ALA:HB3	2.07	0.53
13:16:43:GLU:OE2	13:16:46:ARG:HD3	2.09	0.53
22:25:17:LEU:HD11	22:25:37:ARG:HH21	1.73	0.53
23:26:58:ILE:HG12	23:26:66:VAL:HG21	1.90	0.53
27:30:10:SER:O	27:30:14:MET:HG3	2.08	0.53
53:A:1273:C:H2'	53:A:1274:A:O4'	2.08	0.53
54:01:2533:U:H2'	54:01:2534:A:O4'	2.08	0.53
14:17:15:ARG:O	14:17:19:GLN:HG2	2.08	0.53
17:20:9:GLY:O	54:01:996:A:H1'	2.08	0.53
18:21:4:ILE:HG22	18:21:106:VAL:HG22	1.91	0.53
41:K:109:ILE:HG21	51:U:16:ARG:NH1	2.24	0.53
45:O:63:ARG:HH12	45:O:87:ARG:HH22	1.56	0.53
49:S:30:LEU:H	49:S:48:ILE:HG22	1.74	0.53
54:01:517:C:H2'	54:01:518:G:O4'	2.09	0.53
54:01:1038:G:H2'	54:01:1039:A:C8	2.43	0.53
54:01:1047:G:H2'	54:01:1110:G:C2	2.44	0.53
54:01:2480:C:H2'	54:01:2481:G:O4'	2.09	0.53
16:19:49:ARG:NH1	16:19:49:ARG:HB3	2.23	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:G:27:ASN:HB3	53:A:1375:A:H5'	1.91	0.53
48:R:56:ARG:HB3	48:R:60:ARG:HH12	1.74	0.53
52:03:4:LEU:HB3	52:03:9:ARG:HH21	1.74	0.53
54:01:996:A:H2'	54:01:997:G:H8	1.74	0.53
54:01:1035:U:H2'	54:01:1036:G:H8	1.74	0.53
54:01:2623:G:H2'	54:01:2624:G:H8	1.73	0.53
55:02:65:U:H3'	55:02:108:A:N6	2.24	0.53
59:Z:124:GLN:HG3	59:Z:385:ALA:HB1	1.91	0.53
59:Z:143:GLU:O	59:Z:147:GLU:HG3	2.09	0.53
59:Z:269:ARG:HD2	59:Z:272:GLU:HG2	1.89	0.53
20:23:4:ILE:HD12	20:23:4:ILE:N	2.23	0.52
31:34:19:ARG:HB2	31:34:24:ARG:HD2	1.91	0.52
35:E:149:PRO:HA	35:E:152:VAL:HG22	1.91	0.52
53:A:235:C:H2'	53:A:236:A:H8	1.74	0.52
53:A:797:C:H2'	53:A:798:U:H6	1.74	0.52
53:A:1379:G:O2'	53:A:1380:U:H5'	2.10	0.52
54:01:1683:U:H2'	54:01:1684:G:H8	1.74	0.52
54:01:1709:U:H2'	54:01:1710:G:C8	2.44	0.52
54:01:1830:C:H2'	54:01:1831:G:H8	1.74	0.52
58:Y:25:C:C2'	58:Y:26:A:H5''	2.39	0.52
59:Z:79:VAL:HG12	59:Z:81:CYS:SG	2.49	0.52
9:12:113:PRO:HD2	54:01:558:U:P	2.48	0.52
24:27:32:ALA:HB2	24:27:37:LEU:HD23	1.92	0.52
33:C:81:GLU:OE2	33:C:85:LYS:HB2	2.09	0.52
37:G:19:SER:O	37:G:22:LEU:HG	2.09	0.52
40:J:81:GLU:HA	40:J:84:VAL:HG12	1.91	0.52
41:K:108:ASN:HB3	51:U:6:ARG:HG2	1.90	0.52
53:A:225:C:C3'	53:A:226:G:H5''	2.39	0.52
53:A:381:C:H2'	53:A:382:A:O4'	2.09	0.52
54:01:742:A:H2'	54:01:743:A:H8	1.74	0.52
54:01:2339:C:H2'	54:01:2340:A:H8	1.73	0.52
54:01:2632:A:H2'	54:01:2633:G:H8	1.75	0.52
56:X:28:C:H2'	56:X:29:G:C8	2.44	0.52
59:Z:309:TYR:CE2	59:Z:311:LEU:HA	2.44	0.52
1:04:260:LYS:HA	1:04:263:ASP:OD2	2.09	0.52
2:05:4:LEU:HD21	2:05:96:ILE:HG22	1.91	0.52
12:15:69:PRO:HA	12:15:94:ALA:HB2	1.90	0.52
29:32:3:ARG:HD3	29:32:4:THR:H	1.74	0.52
32:B:18:GLN:O	32:B:19:THR:CB	2.55	0.52
32:B:60:ALA:CB	32:B:223:GLY:HA3	2.39	0.52
40:J:17:LEU:HG	40:J:20:GLN:NE2	2.21	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:S:19:GLU:O	49:S:23:GLU:HG2	2.10	0.52
52:03:180:PHE:HB2	52:03:185:LEU:HD21	1.92	0.52
53:A:560:A:H4'	53:A:561:U:H5'	1.90	0.52
2:05:8:LYS:HB2	2:05:201:LEU:CD1	2.39	0.52
10:13:114:LYS:HE3	10:13:118:LEU:HD11	1.91	0.52
16:19:5:ARG:NH2	16:19:5:ARG:HG3	2.24	0.52
22:25:36:GLN:HE21	22:25:53:HIS:HB3	1.73	0.52
29:32:4:THR:HG22	54:01:687:C:H1'	1.92	0.52
30:33:3:ILE:HG21	30:33:62:PRO:HG2	1.92	0.52
53:A:225:C:C2'	53:A:226:G:H5''	2.38	0.52
53:A:312:C:H2'	53:A:313:A:C8	2.44	0.52
54:01:634:C:H2'	54:01:635:C:C6	2.44	0.52
54:01:2106:U:H2'	54:01:2107:G:C8	2.42	0.52
54:01:2554:U:H2'	54:01:2555:U:C6	2.44	0.52
56:W:20:U:H3'	56:W:21:A:H5'	1.90	0.52
59:Z:245:VAL:HA	59:Z:250:THR:HG22	1.91	0.52
9:12:14:ASP:CG	9:12:15:TRP:H	2.13	0.52
16:19:50:ARG:HG2	54:01:1156:A:C8	2.45	0.52
19:22:13:ALA:HB1	19:22:14:PRO:HD2	1.92	0.52
40:J:57:VAL:O	40:J:58:ASN:CB	2.57	0.52
44:N:63:CYS:HB3	44:N:67:GLY:H	1.75	0.52
51:U:33:ARG:O	51:U:34:ARG:HB2	2.08	0.52
58:Y:45:U:H3'	58:Y:46:G:H5''	1.91	0.52
4:07:90:LEU:HD12	4:07:90:LEU:O	2.10	0.52
12:15:8:LYS:HD2	54:01:869:G:H1'	1.90	0.52
27:30:39:ARG:CZ	54:01:2884:U:H3	2.23	0.52
32:B:83:ALA:HB3	32:B:90:PHE:HB3	1.91	0.52
33:C:19:SER:HB3	33:C:21:TRP:HE1	1.73	0.52
34:D:27:ILE:HD12	34:D:27:ILE:N	2.24	0.52
35:E:133:ILE:O	35:E:136:VAL:HG12	2.08	0.52
53:A:1527:U:H2'	53:A:1528:U:C6	2.44	0.52
54:01:1139:G:O2'	54:01:1140:C:H5'	2.09	0.52
54:01:2774:C:H2'	54:01:2775:G:O4'	2.09	0.52
54:01:2811:G:H2'	54:01:2812:G:C8	2.43	0.52
55:02:41:G:H2'	55:02:41:G:N3	2.25	0.52
59:Z:142:ASP:OD2	59:Z:145:LEU:HB2	2.09	0.52
3:06:192:ALA:O	3:06:196:VAL:HG23	2.09	0.52
7:10:27:VAL:CG1	7:10:83:ALA:HB3	2.38	0.52
12:15:74:THR:HG21	12:15:86:LYS:HE2	1.90	0.52
21:24:80:HIS:CE1	21:24:81:PRO:HD2	2.44	0.52
30:33:61:LEU:HD12	30:33:61:LEU:O	2.10	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:F:17:GLN:O	36:F:21:MET:HG3	2.10	0.52
40:J:52:LEU:HB2	44:N:80:ARG:HD2	1.91	0.52
44:N:45:LEU:HG	49:S:12:LEU:HD21	1.91	0.52
53:A:212:G:H2'	53:A:213:G:C8	2.42	0.52
54:01:528:A:C2	54:01:2042:A:H2'	2.45	0.52
54:01:1857:G:H2'	54:01:1884:G:H22	1.74	0.52
54:01:2388:A:H5'	54:01:2389:G:OP2	2.10	0.52
59:Z:330:PHE:CE2	59:Z:360:VAL:HG11	2.44	0.52
5:08:23:ILE:HD11	5:08:42:VAL:HG11	1.91	0.52
10:13:71:ARG:HH12	15:18:71:ARG:NH2	2.05	0.52
38:H:106:SER:HA	53:A:642:A:C8	2.45	0.52
50:T:34:VAL:HG21	50:T:78:LEU:HD21	1.91	0.52
53:A:1409:C:H2'	53:A:1410:A:C8	2.45	0.52
1:04:17:LYS:NZ	1:04:17:LYS:HB2	2.25	0.52
5:08:51:PHE:CE2	5:08:68:ARG:HA	2.45	0.52
9:12:81:ILE:HG23	9:12:82:GLY:H	1.74	0.52
33:C:175:HIS:ND1	53:A:1108:G:H5'	2.25	0.52
35:E:148:SER:HB2	35:E:149:PRO:HD2	1.92	0.52
38:H:77:VAL:HG23	38:H:126:CYS:HA	1.92	0.52
50:T:8:LYS:HE3	50:T:12:GLN:OE1	2.09	0.52
51:U:9:GLU:CG	51:U:10:PRO:HD3	2.06	0.52
52:03:53:ARG:HB3	56:X:62:C:H4'	1.90	0.52
53:A:52:C:H2'	53:A:53:A:C8	2.45	0.52
53:A:78:A:H2'	53:A:79:G:O4'	2.10	0.52
53:A:802:A:H2'	53:A:803:G:O4'	2.10	0.52
54:01:176:A:O2'	54:01:177:G:H5'	2.10	0.52
54:01:441:U:H2'	54:01:442:G:C8	2.45	0.52
54:01:662:G:H2'	54:01:663:G:H8	1.75	0.52
54:01:1056:G:H4'	54:01:1086:A:C8	2.43	0.52
54:01:1357:C:H2'	54:01:1358:G:O4'	2.10	0.52
54:01:1869:G:H3'	54:01:1870:C:H5''	1.92	0.52
54:01:2297:A:N1	54:01:2321:U:H5	2.07	0.52
58:Y:54:U:OP1	59:Z:320:THR:HG21	2.10	0.52
59:Z:300:PRO:CG	59:Z:365:PRO:HB2	2.40	0.52
2:05:34:VAL:HG22	2:05:50:VAL:HG12	1.92	0.52
3:06:76:PRO:HA	3:06:82:GLY:HA2	1.92	0.52
53:A:1399:C:H4'	53:A:1400:C:O5'	2.09	0.52
54:01:1297:C:OP1	54:01:2710:C:H4'	2.09	0.52
54:01:2130:U:H4'	54:01:2134:A:H4'	1.92	0.52
54:01:2710:C:H2'	54:01:2711:A:C8	2.45	0.52
11:14:93:ASN:O	11:14:94:THR:HB	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:33:63:TYR:CE2	54:01:242:G:H5''	2.45	0.51
36:F:3:HIS:HB2	36:F:92:THR:HA	1.92	0.51
45:O:44:GLU:C	45:O:46:LYS:H	2.12	0.51
52:03:184:LYS:HA	52:03:187:GLU:OE1	2.10	0.51
53:A:304:U:H2'	53:A:305:G:H8	1.73	0.51
53:A:868:C:H2'	53:A:869:G:O4'	2.10	0.51
53:A:1414:U:H2'	53:A:1415:G:C8	2.41	0.51
54:01:749:A:H4'	54:01:1271:G:N3	2.25	0.51
54:01:817:C:H2'	54:01:818:G:O4'	2.09	0.51
54:01:833:A:H2'	54:01:834:G:C8	2.45	0.51
54:01:848:C:H2'	54:01:849:A:C8	2.45	0.51
54:01:1268:A:H2'	54:01:1269:A:O4'	2.10	0.51
54:01:1796:U:H2'	54:01:1797:G:C8	2.45	0.51
54:01:2514:U:H2'	54:01:2515:C:H6	1.75	0.51
56:X:69:C:C2'	56:X:70:G:H5'	2.38	0.51
56:W:76:A:H3'	60:W:101:FME:C	2.39	0.51
2:05:3:GLY:C	2:05:4:LEU:HD12	2.30	0.51
4:07:22:ASN:HB2	4:07:26:GLN:NE2	2.24	0.51
15:18:52:ARG:HH22	54:01:2720:U:H5''	1.74	0.51
32:B:76:SER:HA	32:B:79:VAL:HG23	1.91	0.51
33:C:72:PRO:O	33:C:76:ILE:HG12	2.11	0.51
52:03:51:ASP:OD1	52:03:53:ARG:HB2	2.10	0.51
52:03:56:ASP:OD1	52:03:57:GLN:HG2	2.11	0.51
53:A:110:C:H2'	53:A:111:G:O4'	2.10	0.51
53:A:1256:A:H1'	53:A:1258:G:C4	2.45	0.51
53:A:1354:U:H2'	53:A:1355:G:H8	1.74	0.51
54:01:1098:A:H2'	54:01:1099:G:H5'	1.91	0.51
58:Y:10:G:C2	58:Y:26:A:H1'	2.45	0.51
52:03:30:LEU:HD12	52:03:31:LYS:N	2.26	0.51
53:A:1198:G:H5'	53:A:1198:G:H8	1.76	0.51
54:01:71:A:H5'	54:01:71:A:N3	2.25	0.51
17:20:33:VAL:HG23	17:20:61:ALA:HB3	1.91	0.51
32:B:183:PHE:HA	32:B:195:VAL:HG13	1.91	0.51
35:E:49:TYR:HE1	35:E:133:ILE:HG12	1.75	0.51
53:A:434:U:H2'	53:A:435:A:C8	2.44	0.51
54:01:2801:G:H2'	54:01:2802:G:H8	1.75	0.51
59:Z:140:VAL:HG23	59:Z:142:ASP:N	2.26	0.51
59:Z:324:LYS:HE3	59:Z:342:GLU:HA	1.92	0.51
1:04:224:MET:SD	1:04:229:HIS:HB2	2.51	0.51
32:B:153:MET:HE1	32:B:157:PRO:HG3	1.91	0.51
52:03:27:ILE:HA	52:03:30:LEU:HD21	1.92	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:03:214:ILE:HD12	52:03:222:VAL:HB	1.92	0.51
53:A:309:A:H2'	53:A:310:G:H8	1.75	0.51
53:A:1506:U:O2'	53:A:1507:A:H5'	2.10	0.51
54:01:2646:C:H2'	54:01:2647:U:O4'	2.11	0.51
55:02:28:C:H2'	55:02:29:A:C8	2.45	0.51
56:W:21:A:H61	56:W:46:G:H2'	1.75	0.51
59:Z:309:TYR:CE2	59:Z:311:LEU:HD13	2.36	0.51
59:Z:321:PRO:HG2	59:Z:349:MET:HE2	1.93	0.51
1:04:153:LEU:HD11	1:04:181:ARG:HH21	1.75	0.51
3:06:34:ALA:CA	3:06:94:GLN:HE21	2.17	0.51
16:19:24:TYR:CE1	54:01:17:G:H4'	2.46	0.51
25:28:40:THR:HG23	25:28:43:ILE:H	1.76	0.51
32:B:46:VAL:HB	32:B:47:PRO:HD3	1.93	0.51
38:H:28:SER:HB2	38:H:58:LEU:HB2	1.91	0.51
53:A:514:C:H2'	53:A:515:G:H8	1.76	0.51
54:01:182:A:O2'	54:01:183:C:H5'	2.11	0.51
54:01:969:G:H2'	54:01:970:U:C6	2.45	0.51
54:01:1777:U:O2'	54:01:1778:U:H5'	2.11	0.51
59:Z:16:THR:OG1	59:Z:78:HIS:NE2	2.44	0.51
59:Z:206:ILE:HB	59:Z:235:ILE:HG23	1.92	0.51
14:17:29:HIS:CD2	55:02:7:G:H5''	2.46	0.51
32:B:45:THR:HG22	32:B:49:PHE:CD1	2.46	0.51
40:J:27:GLU:OE2	40:J:31:ARG:HD2	2.11	0.51
53:A:880:C:H2'	53:A:881:G:H8	1.76	0.51
54:01:2570:G:H2'	54:01:2571:U:O4'	2.11	0.51
54:01:2884:U:O2	54:01:2884:U:H3'	2.11	0.51
58:Y:1:G:H1'	59:Z:63:ASN:HD21	1.76	0.51
59:Z:237:LYS:HE3	59:Z:267:GLU:OE2	2.10	0.51
6:09:76:GLU:O	6:09:142:VAL:HG13	2.10	0.51
11:14:3:LEU:HD23	54:01:1203:U:H5'	1.93	0.51
35:E:108:GLY:O	35:E:109:ALA:HB3	2.11	0.51
39:I:49:GLN:N	39:I:50:PRO:HD2	2.25	0.51
53:A:17:U:H2'	53:A:18:C:C6	2.46	0.51
54:01:2684:U:H2'	54:01:2685:G:O4'	2.11	0.51
55:02:95:U:H2'	55:02:96:G:H8	1.76	0.51
56:W:62:C:H2'	56:W:63:G:C8	2.46	0.51
59:Z:49:ILE:HG23	59:Z:65:SER:CB	2.39	0.51
59:Z:147:GLU:HA	59:Z:171:ARG:NH2	2.26	0.51
4:07:114:ARG:HG3	4:07:177:ARG:NE	2.25	0.51
6:09:125:THR:HG21	6:09:148:ALA:HB2	1.93	0.51
38:H:9:MET:HE1	38:H:35:ILE:HD12	1.93	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:J:83:THR:O	40:J:87:LEU:HG	2.10	0.51
49:S:38:THR:HG22	49:S:69:LYS:HG2	1.91	0.51
52:03:44:VAL:HG11	52:03:192:LEU:HD23	1.92	0.51
53:A:29:U:O2'	53:A:30:U:H5'	2.11	0.51
53:A:219:U:H2'	53:A:220:G:C8	2.46	0.51
53:A:224:U:H2'	53:A:225:C:C6	2.46	0.51
53:A:1258:G:H2'	53:A:1259:C:C6	2.45	0.51
54:01:2487:G:H2'	54:01:2488:G:C8	2.46	0.51
59:Z:157:LEU:HB3	59:Z:164:GLY:HA3	1.93	0.51
59:Z:324:LYS:HE3	59:Z:343:LEU:N	2.25	0.51
16:19:54:ARG:O	16:19:58:GLN:HG2	2.11	0.51
21:24:20:LEU:HD11	21:24:41:GLU:HG3	1.92	0.51
52:03:191:ALA:O	52:03:194:VAL:HB	2.10	0.51
53:A:824:G:H2'	53:A:825:A:H8	1.76	0.51
53:A:1227:A:H2'	53:A:1228:C:H5'	1.92	0.51
54:01:807:U:H2'	54:01:808:G:C8	2.46	0.51
54:01:1370:C:H2'	54:01:1371:G:O4'	2.11	0.51
54:01:2795:C:H2'	54:01:2796:U:O4'	2.11	0.51
54:01:2818:U:H2'	54:01:2819:G:C8	2.46	0.51
54:01:2848:G:O2'	54:01:2849:U:H5'	2.11	0.51
59:Z:27:LEU:HD12	59:Z:28:THR:N	2.25	0.51
59:Z:287:GLU:CG	59:Z:290:GLN:HB2	2.40	0.51
3:06:131:THR:HB	3:06:164:LEU:HD11	1.94	0.50
7:10:74:ASP:HA	7:10:77:VAL:HG23	1.92	0.50
15:18:3:ILE:N	15:18:3:ILE:HD12	2.26	0.50
34:D:56:GLU:HG2	34:D:198:LEU:HB2	1.93	0.50
37:G:111:GLY:HA2	37:G:118:ARG:HD3	1.93	0.50
52:03:19:LYS:HZ2	52:03:21:TYR:HD1	1.59	0.50
52:03:47:ASN:HB3	52:03:211:LYS:H	1.76	0.50
53:A:726:C:H2'	53:A:727:G:C8	2.46	0.50
54:01:74:A:H4'	54:01:75:G:O5'	2.11	0.50
54:01:579:G:H4'	54:01:2018:G:H5''	1.92	0.50
54:01:1837:C:H2'	54:01:1899:A:N6	2.26	0.50
54:01:1856:U:H2'	54:01:1857:G:O4'	2.11	0.50
56:X:5:G:H2'	56:X:6:G:C8	2.46	0.50
4:07:31:GLU:H	4:07:157:THR:HA	1.76	0.50
6:09:77:THR:HA	6:09:142:VAL:HG13	1.93	0.50
7:10:24:SER:HB3	7:10:86:MET:SD	2.52	0.50
21:24:48:MET:CE	21:24:51:GLN:HE21	2.24	0.50
38:H:17:GLN:HG3	38:H:71:VAL:HB	1.93	0.50
43:M:10:ASP:O	43:M:11:HIS:HB2	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
43:M:28:ARG:HG2	43:M:62:PHE:CE2	2.46	0.50
53:A:67:C:H2'	53:A:68:G:C8	2.45	0.50
53:A:80:A:N6	53:A:86:G:H21	2.08	0.50
55:02:13:G:O2'	55:02:15:A:H2'	2.11	0.50
59:Z:137:CYS:HB3	59:Z:184:TRP:CZ3	2.47	0.50
5:08:42:VAL:HG23	5:08:50:THR:O	2.12	0.50
7:10:23:LEU:HD22	7:10:118:ILE:HG21	1.92	0.50
23:26:53:LYS:O	23:26:57:VAL:HG23	2.11	0.50
25:28:35:VAL:HG22	25:28:36:GLU:N	2.26	0.50
25:28:40:THR:OG1	25:28:41:PRO:HD2	2.11	0.50
47:Q:58:VAL:HG21	47:Q:74:LEU:HD11	1.93	0.50
53:A:744:C:H2'	53:A:745:G:C8	2.47	0.50
53:A:1228:C:H2'	53:A:1229:A:H8	1.76	0.50
53:A:1402:C:H2'	53:A:1403:C:O4'	2.10	0.50
54:01:655:A:H4'	54:01:656:G:H5'	1.93	0.50
2:05:184:ARG:HE	15:18:6:GLN:NE2	2.09	0.50
4:07:35:LEU:HD11	4:07:98:PHE:CE2	2.46	0.50
4:07:114:ARG:HH11	43:M:70:ARG:CZ	2.25	0.50
7:10:62:ARG:HG3	54:01:1046:A:O2'	2.12	0.50
11:14:20:GLY:HA2	11:14:28:GLY:HA2	1.92	0.50
15:18:92:ARG:HD3	54:01:1753:G:H5''	1.93	0.50
29:32:12:ARG:HH11	29:32:44:VAL:HG21	1.76	0.50
34:D:205:LYS:HB3	53:A:8:A:C5	2.46	0.50
37:G:110:ARG:HH22	37:G:121:ASN:HB3	1.76	0.50
45:O:55:LEU:O	45:O:59:VAL:HG23	2.11	0.50
54:01:310:A:H2'	54:01:311:A:H5''	1.91	0.50
54:01:676:A:H62	54:01:802:A:H61	1.60	0.50
54:01:720:U:H2'	54:01:721:A:C8	2.46	0.50
54:01:2176:A:H2'	54:01:2177:C:C6	2.47	0.50
55:02:29:A:H2'	55:02:30:C:O4'	2.12	0.50
56:X:21:A:N6	56:X:46:G:H2'	2.26	0.50
59:Z:206:ILE:HB	59:Z:235:ILE:CG2	2.42	0.50
59:Z:312:SER:HB2	59:Z:315:GLU:HG3	1.93	0.50
5:08:138:GLN:NE2	54:01:2746:U:H1'	2.22	0.50
16:19:5:ARG:HG3	16:19:5:ARG:HH21	1.76	0.50
16:19:93:ILE:HD12	17:20:13:ARG:HB2	1.94	0.50
35:E:158:LYS:HZ1	38:H:63:LYS:HE3	1.75	0.50
50:T:42:ASP:HB3	50:T:45:ALA:HB3	1.93	0.50
52:03:11:ILE:HG22	52:03:220:ALA:HB2	1.94	0.50
52:03:27:ILE:O	52:03:31:LYS:HG3	2.11	0.50
52:03:221:GLY:HA3	54:01:2176:A:H5''	1.93	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:A:216:U:H2'	53:A:217:C:C6	2.46	0.50
53:A:884:U:H4'	53:A:885:G:H5''	1.93	0.50
54:01:2088:A:H2'	54:01:2089:C:C6	2.47	0.50
55:02:70:C:H2'	55:02:71:C:H6	1.77	0.50
59:Z:269:ARG:CZ	59:Z:272:GLU:HG2	2.41	0.50
4:07:72:SER:OG	4:07:79:ARG:HA	2.10	0.50
7:10:18:VAL:HG12	7:10:70:GLU:HB2	1.93	0.50
16:19:93:ILE:O	16:19:97:ILE:HG13	2.12	0.50
25:28:50:VAL:HB	25:28:53:MET:HG2	1.94	0.50
35:E:39:GLY:HA3	35:E:45:VAL:HG12	1.93	0.50
46:P:33:ILE:N	46:P:33:ILE:HD12	2.26	0.50
53:A:950:U:H2'	53:A:951:G:H8	1.76	0.50
54:01:546:U:H2'	54:01:547:A:O4'	2.10	0.50
54:01:1548:A:H2'	54:01:1549:A:C8	2.46	0.50
54:01:2487:G:H2'	54:01:2488:G:H8	1.77	0.50
54:01:2705:A:H2'	54:01:2706:A:O4'	2.11	0.50
59:Z:129:TYR:CD2	59:Z:199:ILE:HG23	2.46	0.50
3:06:149:ILE:CG2	3:06:188:MET:HG2	2.41	0.50
22:25:38:GLY:HA2	54:01:2330:G:H21	1.77	0.50
37:G:62:GLU:O	37:G:66:GLU:HG2	2.12	0.50
40:J:7:ARG:H	40:J:101:SER:HB2	1.77	0.50
53:A:1014:A:H2'	53:A:1015:G:O4'	2.12	0.50
54:01:172:A:H2'	54:01:173:A:C8	2.46	0.50
54:01:582:A:H2'	54:01:583:G:C8	2.47	0.50
59:Z:30:ALA:O	59:Z:34:VAL:HG23	2.11	0.50
59:Z:170:VAL:HG13	59:Z:194:PHE:HE2	1.76	0.50
7:10:81:LEU:HD12	54:01:1107:G:H1'	1.92	0.50
34:D:36:ALA:N	34:D:37:PRO:HD3	2.27	0.50
34:D:100:VAL:O	34:D:104:MET:HG2	2.12	0.50
36:F:18:VAL:O	36:F:22:ILE:HG13	2.12	0.50
39:I:43:ALA:O	39:I:46:VAL:HG22	2.12	0.50
53:A:382:A:H2'	53:A:383:A:C8	2.47	0.50
53:A:407:U:H2'	53:A:408:A:C8	2.46	0.50
53:A:513:C:H2'	53:A:514:C:H6	1.74	0.50
53:A:613:C:H2'	53:A:614:C:C6	2.47	0.50
54:01:644:A:H2'	54:01:645:C:C4'	2.42	0.50
54:01:2450:A:OP1	54:01:2497:A:H2'	2.12	0.50
54:01:2810:A:H2'	54:01:2811:G:O4'	2.12	0.50
57:V:19:U:H2'	57:V:20:U:C6	2.47	0.50
59:Z:243:GLU:HG2	59:Z:245:VAL:CG2	2.42	0.50
7:10:59:LEU:HB2	7:10:62:ARG:CB	2.42	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:30:39:ARG:NH1	54:01:2884:U:H3	2.10	0.50
32:B:73:ARG:HH22	32:B:94:ARG:NH2	2.05	0.50
33:C:156:LEU:HD12	33:C:156:LEU:O	2.11	0.50
34:D:187:ARG:HD2	34:D:190:LEU:HD11	1.94	0.50
35:E:23:THR:HA	35:E:28:ARG:HA	1.94	0.50
52:03:175:ILE:HB	52:03:188:ASN:HB3	1.93	0.50
54:01:2185:U:H2'	54:01:2186:G:C8	2.47	0.50
54:01:2248:C:H2'	54:01:2249:U:H5'	1.93	0.50
58:Y:37:A:H2'	58:Y:38:A:O4'	2.12	0.50
1:04:123:ILE:HG21	36:F:80:PHE:CZ	2.47	0.49
10:13:19:VAL:HG12	10:13:43:ILE:HA	1.94	0.49
17:20:76:LYS:HB2	17:20:85:LYS:HB3	1.93	0.49
25:28:40:THR:CG2	25:28:43:ILE:HG12	2.42	0.49
33:C:111:ASP:O	33:C:115:VAL:HG23	2.11	0.49
40:J:84:VAL:HA	40:J:87:LEU:HD12	1.94	0.49
52:03:54:LYS:HB2	56:X:62:C:O2'	2.12	0.49
53:A:1148:U:H2'	53:A:1149:C:O4'	2.11	0.49
53:A:1458:G:H2'	53:A:1459:G:C8	2.47	0.49
54:01:886:A:H2'	54:01:886:A:N3	2.26	0.49
54:01:971:G:H2'	54:01:972:A:O4'	2.12	0.49
54:01:2556:C:H2'	54:01:2557:G:O4'	2.12	0.49
56:X:59:A:H2'	56:X:60:U:O4'	2.12	0.49
56:W:17:C:H5'	56:W:61:C:OP1	2.11	0.49
1:04:83:ASP:HB3	1:04:86:ARG:HB2	1.94	0.49
2:05:56:LYS:HB2	2:05:56:LYS:NZ	2.27	0.49
11:14:13:LYS:HE2	54:01:1245:G:OP1	2.12	0.49
34:D:10:LEU:HD13	34:D:62:ARG:HG3	1.94	0.49
35:E:105:ILE:HD11	35:E:123:LEU:HD22	1.93	0.49
53:A:916:U:H2'	53:A:917:G:H8	1.77	0.49
54:01:979:A:H2'	54:01:982:C:H42	1.78	0.49
54:01:1447:C:H2'	54:01:1448:G:H8	1.77	0.49
54:01:1506:U:H2'	54:01:1507:C:C6	2.47	0.49
54:01:2156:G:H2'	54:01:2157:G:H5'	1.93	0.49
54:01:2158:A:H4'	54:01:2159:G:O4'	2.12	0.49
54:01:2286:G:H5'	54:01:2287:A:C1'	2.42	0.49
54:01:2286:G:H4'	54:01:2287:A:O4'	2.12	0.49
59:Z:139:MET:HA	59:Z:139:MET:CE	2.42	0.49
1:04:5:CYS:SG	1:04:17:LYS:HE3	2.52	0.49
3:06:149:ILE:HG23	3:06:188:MET:HG2	1.94	0.49
17:20:16:GLU:HB2	17:20:101:ILE:HG13	1.94	0.49
33:C:38:VAL:O	33:C:42:LEU:HD13	2.12	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:D:158:LEU:HD23	34:D:158:LEU:O	2.12	0.49
46:P:36:VAL:HG13	46:P:36:VAL:O	2.12	0.49
53:A:601:G:H2'	53:A:602:A:C8	2.47	0.49
53:A:1256:A:O2'	53:A:1257:A:H5''	2.12	0.49
54:01:555:G:HO2'	54:01:556:A:H8	1.58	0.49
54:01:2584:U:H2'	54:01:2585:U:H2'	1.94	0.49
59:Z:176:LYS:NZ	59:Z:184:TRP:HE1	2.10	0.49
3:06:117:ARG:HH12	11:14:2:ARG:HG2	1.77	0.49
5:08:82:PHE:CE2	5:08:137:LYS:HB2	2.47	0.49
9:12:84:ILE:HG23	9:12:84:ILE:O	2.13	0.49
13:16:45:ARG:HG2	13:16:95:THR:HG21	1.94	0.49
18:21:8:ARG:HA	18:21:102:HIS:ND1	2.27	0.49
52:03:29:LEU:HD12	52:03:30:LEU:N	2.27	0.49
53:A:1082:A:H2'	53:A:1083:U:O4'	2.13	0.49
54:01:329:G:O4'	54:01:477:A:H1'	2.13	0.49
54:01:372:G:H2'	54:01:400:G:O6	2.13	0.49
54:01:873:C:H2'	54:01:874:G:C8	2.47	0.49
54:01:1810:A:H2'	54:01:1811:G:O4'	2.12	0.49
54:01:2186:G:H2'	54:01:2187:U:O4'	2.11	0.49
54:01:2801:G:H2'	54:01:2802:G:C8	2.48	0.49
54:01:2881:U:H2'	54:01:2882:A:C8	2.48	0.49
59:Z:47:ASP:HA	59:Z:56:LYS:HE3	1.94	0.49
59:Z:112:MET:HB3	59:Z:112:MET:HE2	1.95	0.49
3:06:148:ILE:HD13	3:06:187:VAL:HG11	1.93	0.49
7:10:77:VAL:C	7:10:79:PRO:HD2	2.33	0.49
12:15:30:SER:O	12:15:132:THR:HG23	2.12	0.49
27:30:24:VAL:HG13	27:30:25:THR:N	2.27	0.49
29:32:11:LYS:HE3	54:01:686:U:H5''	1.93	0.49
33:C:39:ARG:HH21	33:C:56:ILE:HG13	1.77	0.49
38:H:72:GLU:HB3	38:H:129:ALA:HB3	1.93	0.49
46:P:5:ARG:HD2	53:A:376:G:H4'	1.94	0.49
47:Q:59:GLU:HB2	47:Q:76:ARG:H	1.77	0.49
52:03:67:HIS:HB2	52:03:188:ASN:OD1	2.12	0.49
52:03:200:LYS:HD3	52:03:208:TYR:CD2	2.48	0.49
53:A:501:C:H2'	53:A:502:A:C8	2.47	0.49
53:A:533:A:O2'	53:A:534:U:H5''	2.12	0.49
53:A:1228:C:H2'	53:A:1229:A:C8	2.48	0.49
54:01:2270:A:H2'	54:01:2271:G:O4'	2.13	0.49
54:01:2837:A:H2'	54:01:2838:G:H8	1.77	0.49
56:W:38:A:H2'	56:W:39:C:O4'	2.12	0.49
59:Z:144:GLU:HA	59:Z:147:GLU:CG	2.42	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:Z:316:GLY:O	59:Z:381:ARG:HG3	2.12	0.49
7:10:13:ALA:O	7:10:17:GLU:HG3	2.13	0.49
10:13:51:LYS:HD2	10:13:95:ILE:HG22	1.95	0.49
32:B:153:MET:CE	32:B:157:PRO:HG3	2.43	0.49
37:G:2:ARG:HB3	53:A:933:G:OP2	2.12	0.49
41:K:86:LYS:HB2	41:K:112:VAL:HG23	1.95	0.49
44:N:3:GLN:HG3	53:A:1047:G:H5'	1.94	0.49
51:U:33:ARG:O	51:U:34:ARG:CB	2.61	0.49
53:A:56:U:H2'	53:A:57:G:H8	1.78	0.49
53:A:129:A:H1'	53:A:130:A:C8	2.48	0.49
54:01:233:A:H2'	54:01:234:U:O4'	2.13	0.49
54:01:437:U:H2'	54:01:438:G:C8	2.47	0.49
54:01:540:C:H2'	54:01:541:A:H8	1.78	0.49
54:01:2141:G:N2	54:01:2151:U:H1'	2.26	0.49
2:05:43:ASP:HB3	2:05:45:TYR:CE1	2.48	0.49
16:19:2:ARG:HB2	54:01:1248:G:C5	2.48	0.49
20:23:31:GLY:O	20:23:66:VAL:HG23	2.12	0.49
20:23:83:GLY:HA3	20:23:96:LYS:HE3	1.94	0.49
37:G:38:ALA:O	37:G:42:VAL:HG23	2.13	0.49
51:U:65:ARG:NE	51:U:65:ARG:HA	2.28	0.49
53:A:460:A:H2'	53:A:461:A:H8	1.77	0.49
53:A:1372:U:H2'	53:A:1373:G:O4'	2.12	0.49
54:01:598:U:H2'	54:01:599:A:H8	1.78	0.49
54:01:1611:C:H5'	54:01:1611:C:H6	1.75	0.49
59:Z:98:MET:O	59:Z:127:VAL:HG22	2.12	0.49
59:Z:175:LEU:O	59:Z:179:GLU:HG3	2.12	0.49
4:07:7:TYR:HA	4:07:11:VAL:CG2	2.43	0.49
8:11:55:PRO:HG3	54:01:1060:U:OP2	2.12	0.49
23:26:11:PRO:HG3	23:26:30:PRO:HD2	1.94	0.49
33:C:71:ARG:O	33:C:75:VAL:HG23	2.12	0.49
33:C:78:LYS:HB2	33:C:78:LYS:NZ	2.28	0.49
36:F:68:GLN:HA	36:F:71:ILE:HG22	1.94	0.49
39:I:5:TYR:HB2	39:I:20:ILE:HG23	1.95	0.49
53:A:67:C:H2'	53:A:68:G:H8	1.77	0.49
54:01:2498:C:O2'	54:01:2499:C:H5'	2.12	0.49
58:Y:49:C:H2'	58:Y:50:U:C6	2.47	0.49
7:10:48:ALA:HB3	7:10:51:TYR:CE1	2.47	0.49
19:22:80:TRP:CZ3	19:22:82:LYS:HB3	2.48	0.49
47:Q:28:VAL:HG22	47:Q:29:LYS:N	2.28	0.49
51:U:29:ALA:HA	51:U:32:ARG:HD3	1.94	0.49
51:U:64:ALA:HA	51:U:66:ARG:HH22	1.78	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:A:286:C:H2'	53:A:287:U:C6	2.48	0.49
53:A:604:G:H2'	53:A:605:U:O4'	2.13	0.49
53:A:1007:U:H2'	53:A:1008:U:C6	2.48	0.49
54:01:1443:U:H2'	54:01:1444:G:C8	2.48	0.49
54:01:1447:C:H2'	54:01:1448:G:C8	2.47	0.49
54:01:1866:A:H2'	54:01:1867:G:O4'	2.12	0.49
54:01:2163:A:H2'	54:01:2163:A:N3	2.27	0.49
54:01:2231:U:H2'	54:01:2232:C:C6	2.48	0.49
54:01:2811:G:H2'	54:01:2812:G:H8	1.77	0.49
59:Z:176:LYS:HE2	59:Z:181:ASP:OD2	2.11	0.49
59:Z:304:PHE:HD2	59:Z:388:VAL:HG13	1.75	0.49
16:19:49:ARG:HB3	16:19:49:ARG:HH11	1.78	0.49
32:B:98:GLY:O	32:B:102:ASN:HB3	2.13	0.49
41:K:19:VAL:HG22	41:K:82:GLU:OE1	2.13	0.49
53:A:1456:A:H2'	53:A:1457:G:O4'	2.11	0.49
54:01:1524:G:H2'	54:01:1525:A:H8	1.78	0.49
59:Z:30:ALA:HB2	59:Z:178:LEU:HB2	1.94	0.49
59:Z:74:ARG:HH12	59:Z:201:GLU:HA	1.78	0.49
5:08:9:VAL:HA	5:08:48:THR:HA	1.94	0.48
5:08:118:ALA:O	5:08:120:ILE:N	2.46	0.48
5:08:174:LYS:HE3	54:01:2529:G:H4'	1.94	0.48
13:16:102:PHE:HE1	13:16:109:PRO:HG3	1.77	0.48
25:28:40:THR:HG22	25:28:43:ILE:HG12	1.94	0.48
35:E:47:PHE:CZ	35:E:137:ARG:HG2	2.47	0.48
52:03:15:VAL:HG23	52:03:21:TYR:OH	2.13	0.48
52:03:64:VAL:HG22	52:03:160:GLN:HG2	1.94	0.48
53:A:70:U:H4'	53:A:71:A:H8	1.77	0.48
53:A:505:G:P	53:A:535:A:H5'	2.53	0.48
53:A:694:A:H2'	53:A:695:A:O4'	2.13	0.48
53:A:946:A:H2'	53:A:947:G:C8	2.48	0.48
53:A:1004:A:H2'	53:A:1005:A:O4'	2.12	0.48
55:02:3:C:C3'	55:02:4:C:H5''	2.43	0.48
18:21:31:GLN:O	18:21:35:ILE:HG13	2.11	0.48
41:K:92:ARG:HE	51:U:24:LYS:NZ	2.10	0.48
44:N:92:ILE:HD12	44:N:92:ILE:N	2.27	0.48
46:P:31:ARG:NH2	53:A:230:G:H5''	2.18	0.48
49:S:27:LYS:HG2	49:S:28:LYS:H	1.78	0.48
52:03:8:MET:O	52:03:12:ARG:HB2	2.12	0.48
53:A:1340:A:H2'	53:A:1341:U:O4'	2.13	0.48
1:04:235:GLU:HG2	54:01:2599:G:C8	2.47	0.48
6:09:55:GLU:HA	6:09:58:LEU:HD12	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:09:126:GLY:H	6:09:146:VAL:HB	1.79	0.48
12:15:41:LEU:HA	12:15:45:GLN:OE1	2.13	0.48
13:16:102:PHE:CE1	13:16:109:PRO:HG3	2.48	0.48
15:18:3:ILE:HD12	15:18:3:ILE:H	1.77	0.48
22:25:45:ALA:O	22:25:47:VAL:HG23	2.13	0.48
39:I:121:ARG:HD2	53:A:1348:U:H4'	1.95	0.48
40:J:14:ASP:HB3	40:J:17:LEU:HB3	1.94	0.48
40:J:59:LYS:HE2	40:J:62:ARG:HH21	1.78	0.48
52:03:62:ALA:HB1	52:03:160:GLN:HE22	1.77	0.48
52:03:207:VAL:HB	52:03:210:LYS:CG	2.42	0.48
53:A:253:A:H2'	53:A:254:G:C8	2.48	0.48
53:A:299:G:H2'	53:A:300:A:C8	2.48	0.48
54:01:215:G:C4'	54:01:216:A:H4'	2.44	0.48
54:01:1547:C:H2'	54:01:1548:A:H8	1.78	0.48
12:15:41:LEU:HD22	12:15:125:PRO:HD2	1.95	0.48
17:20:37:GLU:HA	17:20:53:PHE:CD1	2.48	0.48
32:B:80:LYS:HB2	32:B:92:ASN:HD22	1.78	0.48
35:E:98:ALA:HB2	35:E:123:LEU:HG	1.96	0.48
36:F:66:ALA:HB1	36:F:67:PRO:CD	2.43	0.48
39:I:121:ARG:NH1	53:A:1345:U:H5''	2.29	0.48
42:L:98:ARG:HB2	42:L:116:TYR:HA	1.94	0.48
44:N:61:ASN:HB3	44:N:72:PHE:CE2	2.48	0.48
52:03:43:ASP:OD2	52:03:217:THR:HG22	2.13	0.48
52:03:46:VAL:O	52:03:171:ILE:HG22	2.14	0.48
53:A:52:C:H2'	53:A:53:A:H8	1.78	0.48
53:A:265:G:H2'	53:A:267:C:H5	1.78	0.48
53:A:297:G:H4'	53:A:557:G:H4'	1.94	0.48
53:A:530:G:H3'	53:A:531:U:C5'	2.44	0.48
54:01:1316:U:H2'	54:01:1317:G:C8	2.48	0.48
59:Z:304:PHE:CE2	59:Z:306:SER:HB2	2.48	0.48
4:07:140:ILE:HG22	4:07:142:TYR:H	1.79	0.48
7:10:41:LEU:HD13	54:01:1083:U:O5'	2.12	0.48
11:14:77:ILE:N	11:14:77:ILE:HD12	2.29	0.48
23:26:56:ARG:HA	23:26:59:ASP:HB2	1.96	0.48
27:30:8:THR:HG21	54:01:2020:A:H5'	1.95	0.48
38:H:116:ARG:HG3	38:H:116:ARG:HH11	1.79	0.48
39:I:70:GLY:O	39:I:74:GLN:HG3	2.13	0.48
39:I:90:ASP:O	39:I:92:SER:N	2.46	0.48
51:U:36:PHE:C	51:U:38:GLU:H	2.17	0.48
53:A:1202:U:H2'	53:A:1203:C:O4'	2.14	0.48
53:A:1406:U:H2'	53:A:1407:C:O4'	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:A:1458:G:H2'	53:A:1459:G:H8	1.78	0.48
54:01:1507:C:H2'	54:01:1508:A:O4'	2.13	0.48
54:01:2637:U:H2'	54:01:2638:G:O4'	2.12	0.48
54:01:2785:C:H2'	54:01:2786:U:H6	1.77	0.48
59:Z:259:GLU:CD	59:Z:262:ARG:HA	2.33	0.48
1:04:240:GLY:HA3	54:01:2597:G:H5''	1.95	0.48
1:04:252:LYS:HB2	1:04:252:LYS:NZ	2.29	0.48
5:08:46:ASP:O	5:08:48:THR:N	2.46	0.48
6:09:55:GLU:HA	6:09:58:LEU:HB2	1.94	0.48
21:24:79:ARG:HG3	21:24:86:LEU:HD23	1.96	0.48
28:31:47:ILE:HD12	28:31:47:ILE:N	2.28	0.48
32:B:114:LYS:HB2	32:B:114:LYS:NZ	2.29	0.48
36:F:18:VAL:HG21	36:F:58:HIS:CD2	2.48	0.48
40:J:6:ILE:HG22	40:J:8:ILE:HG13	1.94	0.48
52:03:170:ILE:HG21	54:01:2177:C:O2	2.13	0.48
54:01:364:C:H2'	54:01:365:U:C6	2.49	0.48
54:01:2898:U:H2'	54:01:2899:A:C8	2.49	0.48
59:Z:125:VAL:HG13	59:Z:127:VAL:HG23	1.95	0.48
6:09:99:ILE:O	6:09:103:VAL:HG23	2.13	0.48
12:15:2:LEU:HB3	12:15:68:PHE:CE1	2.49	0.48
17:20:6:GLN:HG2	17:20:11:GLN:HG3	1.95	0.48
32:B:26:MET:HE3	32:B:192:PRO:HD3	1.96	0.48
38:H:12:ARG:HD3	38:H:26:MET:HB3	1.96	0.48
40:J:30:LYS:HB3	40:J:30:LYS:NZ	2.29	0.48
43:M:47:LEU:CD2	43:M:51:GLN:HB2	2.41	0.48
52:03:50:ILE:CD1	52:03:52:ALA:HB2	2.44	0.48
53:A:77:A:H2'	53:A:78:A:H8	1.79	0.48
53:A:195:A:H2'	53:A:196:A:C8	2.48	0.48
53:A:810:C:H2'	53:A:811:C:O4'	2.13	0.48
54:01:119:A:H4'	54:01:120:U:H5'	1.95	0.48
54:01:248:G:H5'	54:01:249:C:H5'	1.95	0.48
54:01:560:C:H2'	54:01:561:G:O4'	2.14	0.48
54:01:889:C:H2'	54:01:890:C:H5'	1.95	0.48
55:02:78:A:H2'	55:02:79:G:O4'	2.14	0.48
59:Z:73:THR:HG21	59:Z:196:ASP:HB3	1.95	0.48
59:Z:241:GLU:HA	59:Z:254:THR:HA	1.96	0.48
59:Z:243:GLU:HG3	59:Z:251:GLN:C	2.34	0.48
59:Z:245:VAL:HA	59:Z:250:THR:CG2	2.44	0.48
3:06:117:ARG:HA	3:06:185:LYS:HD3	1.94	0.48
7:10:59:LEU:HD22	7:10:62:ARG:HB2	1.95	0.48
10:13:2:ILE:HD12	10:13:6:THR:HG21	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:13:116:ILE:HD12	10:13:117:SER:N	2.29	0.48
13:16:55:ALA:HA	13:16:80:PHE:CE1	2.48	0.48
17:20:79:ARG:O	17:20:81:LYS:HG2	2.14	0.48
27:30:11:LYS:HA	27:30:14:MET:CE	2.44	0.48
29:32:12:ARG:NH1	29:32:44:VAL:HG21	2.28	0.48
32:B:66:ILE:HD13	32:B:159:ALA:HB3	1.93	0.48
34:D:37:PRO:HD2	34:D:41:GLY:HA3	1.96	0.48
34:D:120:LYS:HG2	34:D:130:ASN:HD21	1.78	0.48
34:D:131:ILE:HG22	34:D:133:SER:H	1.79	0.48
36:F:77:THR:O	36:F:81:ASN:HB2	2.14	0.48
53:A:977:A:H2'	53:A:978:A:H5''	1.95	0.48
53:A:1123:U:O2'	53:A:1124:G:H5'	2.14	0.48
54:01:181:A:H2'	54:01:182:A:C8	2.49	0.48
54:01:832:U:H2'	54:01:833:A:C8	2.49	0.48
54:01:1035:U:H2'	54:01:1036:G:C8	2.49	0.48
54:01:1043:C:H2'	54:01:1044:C:O4'	2.14	0.48
54:01:1354:A:H2'	54:01:1355:G:O4'	2.13	0.48
54:01:1536:C:H4'	54:01:1537:G:C2	2.49	0.48
54:01:1736:U:H2'	54:01:1737:G:O4'	2.13	0.48
54:01:1775:U:H2'	54:01:1776:G:O4'	2.13	0.48
54:01:2082:A:H2'	54:01:2083:G:O4'	2.13	0.48
54:01:2207:C:H2'	54:01:2208:C:C6	2.48	0.48
59:Z:306:SER:HB3	59:Z:358:MET:HE3	1.94	0.48
1:04:239:PHE:HB2	54:01:1903:G:OP1	2.13	0.48
8:11:79:LEU:HA	8:11:82:ALA:HB3	1.96	0.48
9:12:21:THR:HG23	9:12:61:LYS:HB3	1.96	0.48
11:14:79:LEU:HD12	11:14:112:LEU:HD12	1.96	0.48
13:16:73:ASN:HA	13:16:76:VAL:HG12	1.95	0.48
32:B:37:VAL:HG22	32:B:38:HIS:N	2.28	0.48
32:B:158:ASP:O	32:B:181:PRO:HD2	2.14	0.48
33:C:64:ARG:HG3	33:C:99:GLN:O	2.13	0.48
34:D:8:LEU:HD21	34:D:31:CYS:HB3	1.96	0.48
41:K:71:ASP:O	41:K:72:ALA:HB3	2.14	0.48
47:Q:4:ILE:HD12	47:Q:4:ILE:O	2.13	0.48
51:U:17:ARG:HA	51:U:20:ARG:HH11	1.78	0.48
52:03:50:ILE:HD11	52:03:52:ALA:HB2	1.95	0.48
53:A:10:A:H2'	53:A:11:G:C8	2.47	0.48
54:01:1078:U:H4'	54:01:1079:C:H5''	1.96	0.48
54:01:1524:G:H2'	54:01:1525:A:C8	2.49	0.48
54:01:2317:A:H2'	54:01:2318:G:O4'	2.14	0.48
4:07:62:GLN:NE2	4:07:90:LEU:HB3	2.29	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:11:37:PHE:HE1	8:11:58:ILE:HG13	1.79	0.48
32:B:162:VAL:HB	32:B:184:ALA:CB	2.43	0.48
43:M:6:ILE:HG13	43:M:7:ASN:H	1.79	0.48
43:M:7:ASN:ND2	43:M:20:SER:HB2	2.29	0.48
50:T:77:ASN:O	50:T:81:GLN:HG2	2.14	0.48
52:03:31:LYS:HD2	52:03:182:ALA:H	1.77	0.48
53:A:1106:G:H2'	53:A:1107:C:C6	2.49	0.48
54:01:168:G:H2'	54:01:169:G:H8	1.79	0.48
54:01:226:A:H5''	54:01:257:C:O2'	2.14	0.48
54:01:464:U:H2'	54:01:465:G:O4'	2.14	0.48
54:01:575:A:O2'	54:01:576:U:H5'	2.13	0.48
54:01:669:G:H2'	54:01:669:G:N3	2.28	0.48
54:01:2743:U:C3'	54:01:2744:G:H5''	2.44	0.48
59:Z:89:LYS:HA	59:Z:92:ILE:HD12	1.95	0.48
59:Z:117:GLU:HB2	59:Z:382:THR:OG1	2.14	0.48
3:06:71:GLY:H	54:01:674:G:H5''	1.79	0.47
5:08:8:VAL:O	5:08:49:LEU:N	2.47	0.47
5:08:72:ASN:O	5:08:76:ILE:HG12	2.14	0.47
20:23:32:LYS:HG2	20:23:65:GLN:OE1	2.14	0.47
25:28:23:LEU:HD21	25:28:53:MET:SD	2.54	0.47
26:29:11:GLU:HB2	26:29:25:ARG:HE	1.79	0.47
46:P:59:HIS:O	46:P:63:GLN:HG2	2.13	0.47
53:A:24:U:H2'	53:A:25:C:C6	2.49	0.47
53:A:831:A:H2'	53:A:832:G:O4'	2.14	0.47
55:02:4:C:H6	55:02:4:C:H5'	1.79	0.47
59:Z:260:MET:HG2	59:Z:261:PHE:HD2	1.80	0.47
2:05:127:PHE:CD1	54:01:2512:C:H5''	2.49	0.47
9:12:102:GLU:HG2	9:12:119:PHE:HZ	1.79	0.47
15:18:77:SER:OG	15:18:79:VAL:HG12	2.13	0.47
18:21:20:VAL:HA	27:30:21:LEU:HD12	1.95	0.47
34:D:144:ILE:HB	34:D:149:LYS:NZ	2.28	0.47
53:A:151:A:H2'	53:A:152:A:O4'	2.13	0.47
53:A:921:U:H2'	53:A:922:G:O4'	2.14	0.47
54:01:264:C:O3'	54:01:265:A:H2'	2.14	0.47
54:01:282:A:H2'	54:01:283:G:C8	2.49	0.47
54:01:1990:C:H2'	54:01:1991:U:C6	2.49	0.47
54:01:2134:A:C6	54:01:2157:G:H4'	2.49	0.47
7:10:27:VAL:HG12	7:10:83:ALA:O	2.15	0.47
12:15:50:ARG:HH21	12:15:50:ARG:HG2	1.79	0.47
13:16:28:LEU:HD23	13:16:48:VAL:HG21	1.96	0.47
25:28:37:ARG:NH2	54:01:929:U:H4'	2.30	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:D:170:LEU:HA	34:D:182:LYS:HE2	1.96	0.47
39:I:101:GLY:O	39:I:103:VAL:HG22	2.14	0.47
46:P:20:VAL:HG22	46:P:21:VAL:N	2.29	0.47
52:03:190:GLU:O	52:03:194:VAL:N	2.47	0.47
53:A:59:A:H3'	53:A:331:G:H22	1.80	0.47
53:A:929:G:H5''	53:A:1534:A:H1'	1.96	0.47
53:A:946:A:H2'	53:A:947:G:H8	1.79	0.47
53:A:1175:G:H2'	53:A:1176:A:C8	2.50	0.47
53:A:1409:C:H2'	53:A:1410:A:H8	1.79	0.47
53:A:1492:A:H2'	54:01:1913:A:C2	2.49	0.47
54:01:537:G:H22	54:01:555:G:H2'	1.79	0.47
54:01:813:U:H2'	54:01:814:C:C6	2.50	0.47
54:01:1258:U:H2'	54:01:1259:G:C8	2.50	0.47
54:01:1430:G:H2'	54:01:1431:A:O4'	2.14	0.47
54:01:2577:A:H5''	54:01:2578:G:H5'	1.94	0.47
54:01:2648:G:H2'	54:01:2649:C:O4'	2.14	0.47
59:Z:144:GLU:HA	59:Z:147:GLU:HG3	1.96	0.47
59:Z:172:GLY:HA2	59:Z:184:TRP:CZ3	2.49	0.47
59:Z:305:GLU:HA	59:Z:359:VAL:CG2	2.43	0.47
1:04:121:ALA:HB1	1:04:127:ASN:HB3	1.96	0.47
1:04:220:ARG:HG3	54:01:1789:A:OP1	2.14	0.47
2:05:61:THR:HB	2:05:63:PRO:HD2	1.97	0.47
3:06:155:GLU:OE2	3:06:159:LEU:HD11	2.14	0.47
8:11:4:VAL:HA	8:11:7:TYR:CE1	2.49	0.47
9:12:39:LYS:NZ	9:12:39:LYS:HB3	2.29	0.47
17:20:14:VAL:HG21	17:20:98:ILE:HG13	1.96	0.47
18:21:57:ASN:HD22	18:21:61:ASN:HD22	1.61	0.47
33:C:19:SER:HB3	33:C:21:TRP:CD1	2.49	0.47
37:G:41:ILE:HD11	53:A:1240:U:O4'	2.15	0.47
39:I:45:MET:O	39:I:49:GLN:HG3	2.14	0.47
42:L:33:CYS:HB2	42:L:54:VAL:HG22	1.95	0.47
52:03:163:TYR:HB2	52:03:171:ILE:HD11	1.96	0.47
54:01:704:G:H2'	54:01:726:G:H22	1.78	0.47
54:01:1388:G:H2'	54:01:1389:G:C8	2.50	0.47
59:Z:24:LYS:HA	59:Z:27:LEU:HD21	1.96	0.47
8:11:72:THR:OG1	8:11:73:PRO:HD2	2.15	0.47
8:11:79:LEU:HD23	8:11:82:ALA:HB3	1.96	0.47
11:14:30:THR:HG22	54:01:810:U:O4	2.15	0.47
18:21:58:ALA:HA	18:21:62:ASP:OD1	2.15	0.47
30:33:63:TYR:CD2	54:01:242:G:H5''	2.49	0.47
44:N:13:VAL:HA	44:N:59:GLN:HE22	1.80	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:N:68:ARG:HG2	44:N:68:ARG:HH11	1.79	0.47
49:S:36:ARG:HG2	53:A:1320:C:N4	2.30	0.47
53:A:622:A:H2'	53:A:623:C:H5'	1.96	0.47
53:A:726:C:H2'	53:A:727:G:H8	1.79	0.47
53:A:990:C:H2'	53:A:991:U:O4'	2.15	0.47
53:A:1023:U:H2'	53:A:1024:G:O4'	2.14	0.47
53:A:1391:U:H2'	53:A:1392:G:C8	2.49	0.47
54:01:1196:C:H2'	54:01:1197:G:H8	1.78	0.47
54:01:2697:G:H2'	54:01:2698:U:O4'	2.15	0.47
54:01:2841:C:H2'	54:01:2842:G:H8	1.78	0.47
58:Y:27:G:H2'	58:Y:28:G:O4'	2.14	0.47
58:Y:64:A:H4'	59:Z:379:GLY:HA3	1.97	0.47
1:04:240:GLY:HA3	54:01:2597:G:H5'	1.96	0.47
11:14:14:LYS:HG2	54:01:662:G:O2'	2.14	0.47
19:22:39:THR:O	19:22:43:ILE:HG13	2.15	0.47
21:24:26:PHE:CZ	21:24:42:LEU:HD11	2.49	0.47
34:D:101:VAL:HG13	34:D:106:PHE:HB2	1.95	0.47
35:E:156:ARG:NH2	35:E:163:ILE:HG22	2.29	0.47
38:H:10:LEU:HD22	38:H:74:ILE:CD1	2.45	0.47
40:J:40:ILE:HG23	40:J:41:PRO:HD2	1.96	0.47
42:L:54:VAL:HG21	42:L:79:ILE:HD11	1.96	0.47
52:03:214:ILE:HD12	52:03:214:ILE:O	2.14	0.47
53:A:644:U:H2'	53:A:645:G:H8	1.79	0.47
53:A:1137:C:C5'	53:A:1138:G:H5'	2.41	0.47
53:A:1508:A:H2'	53:A:1509:C:C6	2.50	0.47
54:01:211:C:H5'	54:01:1366:A:O2'	2.15	0.47
54:01:1409:U:H2'	54:01:1410:G:C8	2.50	0.47
54:01:2368:C:H2'	54:01:2369:A:H8	1.80	0.47
54:01:2630:G:H2'	54:01:2631:G:H8	1.79	0.47
54:01:2731:G:H2'	54:01:2732:G:C8	2.49	0.47
58:Y:27:G:O2'	58:Y:28:G:H5'	2.15	0.47
59:Z:134:LEU:HB2	59:Z:171:ARG:HG2	1.97	0.47
7:10:59:LEU:H	7:10:59:LEU:HD12	1.78	0.47
8:11:52:LEU:HB2	8:11:54:ILE:HD11	1.95	0.47
9:12:52:ASP:O	9:12:54:ILE:HG13	2.14	0.47
16:19:91:ARG:HH11	16:19:91:ARG:HG3	1.79	0.47
17:20:74:ILE:N	17:20:74:ILE:HD12	2.30	0.47
25:28:9:THR:HG22	25:28:53:MET:C	2.35	0.47
32:B:162:VAL:HB	32:B:184:ALA:HB2	1.97	0.47
33:C:54:ILE:HG22	33:C:67:ILE:HG12	1.97	0.47
35:E:70:MET:O	35:E:71:ILE:HD13	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:H:8:ASP:O	38:H:11:THR:HG22	2.15	0.47
41:K:116:PRO:HB3	53:A:676:A:C1'	2.45	0.47
47:Q:11:VAL:HG13	47:Q:20:ILE:HD11	1.97	0.47
52:03:19:LYS:NZ	52:03:21:TYR:HD1	2.11	0.47
52:03:175:ILE:HB	52:03:188:ASN:CB	2.44	0.47
52:03:190:GLU:O	52:03:194:VAL:HG23	2.15	0.47
53:A:31:G:N2	53:A:47:C:H5''	2.29	0.47
53:A:634:C:H2'	53:A:635:A:C8	2.50	0.47
53:A:876:C:H2'	53:A:877:G:H8	1.79	0.47
53:A:1075:U:H2'	53:A:1076:U:C6	2.50	0.47
53:A:1305:G:H22	53:A:1331:G:H2'	1.79	0.47
54:01:885:C:C2'	54:01:886:A:H5'	2.44	0.47
54:01:1499:C:H2'	54:01:1500:G:H8	1.79	0.47
54:01:1858:A:C2	54:01:1885:A:H1'	2.50	0.47
54:01:2243:U:H2'	54:01:2244:U:C6	2.49	0.47
54:01:2464:G:H2'	54:01:2465:C:H6	1.80	0.47
54:01:2591:C:H2'	54:01:2592:G:H8	1.77	0.47
54:01:2809:A:H2'	54:01:2810:A:C8	2.50	0.47
56:X:68:C:C2	56:X:69:C:H1'	2.50	0.47
58:Y:28:G:H2'	58:Y:29:G:C8	2.49	0.47
59:Z:10:PRO:O	59:Z:74:ARG:HB2	2.15	0.47
59:Z:173:SER:HB3	59:Z:184:TRP:CD2	2.50	0.47
59:Z:277:LEU:HD12	59:Z:278:LEU:H	1.79	0.47
59:Z:318:ARG:HH21	59:Z:319:HIS:CD2	2.32	0.47
59:Z:366:ILE:HD12	59:Z:367:ALA:H	1.79	0.47
3:06:105:LEU:HA	3:06:108:ILE:HG12	1.97	0.47
3:06:148:ILE:HD13	3:06:187:VAL:CG1	2.44	0.47
8:11:88:GLY:HA3	54:01:1063:G:O2'	2.15	0.47
8:11:123:ALA:HA	8:11:126:ARG:NH1	2.30	0.47
21:24:29:ILE:HA	21:24:40:ILE:HG12	1.96	0.47
53:A:56:U:H2'	53:A:57:G:C8	2.49	0.47
53:A:514:C:H2'	53:A:515:G:C8	2.50	0.47
54:01:1579:A:H2'	54:01:1580:A:O4'	2.14	0.47
54:01:1765:U:H2'	54:01:1766:G:H8	1.79	0.47
54:01:1849:G:H2'	54:01:1850:G:H8	1.79	0.47
54:01:2129:C:H2'	54:01:2159:G:H22	1.80	0.47
59:Z:7:ARG:HH12	59:Z:272:GLU:CG	2.27	0.47
3:06:130:LYS:HB2	3:06:133:LEU:HD12	1.96	0.47
4:07:94:ARG:NH2	26:29:1:MET:HB3	2.30	0.47
6:09:26:ALA:O	6:09:31:VAL:HG23	2.14	0.47
9:12:72:LYS:HB3	9:12:89:PHE:HB2	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:23:40:LEU:HB3	20:23:59:GLU:OE2	2.14	0.47
32:B:95:TRP:CH2	32:B:171:ALA:HA	2.49	0.47
34:D:31:CYS:HA	53:A:429:U:OP2	2.15	0.47
49:S:5:LYS:HG3	49:S:6:LYS:N	2.29	0.47
50:T:30:PHE:HD2	50:T:56:ILE:HG21	1.80	0.47
53:A:1171:A:H2'	53:A:1172:C:C6	2.49	0.47
54:01:856:G:H2'	54:01:857:G:C8	2.50	0.47
54:01:942:G:H2'	54:01:943:A:O4'	2.14	0.47
54:01:2529:G:OP2	54:01:2530:A:H5''	2.15	0.47
59:Z:167:THR:HG22	59:Z:169:ILE:CD1	2.45	0.47
59:Z:174:ALA:HA	59:Z:177:ALA:HB3	1.96	0.47
59:Z:321:PRO:HG2	59:Z:349:MET:CE	2.45	0.47
1:04:135:PRO:HG2	36:F:80:PHE:HD1	1.79	0.47
3:06:47:LYS:HB2	3:06:51:GLU:HB2	1.97	0.47
4:07:47:LYS:NZ	4:07:51:ASN:HD21	2.13	0.47
4:07:127:TYR:HD2	4:07:155:ILE:HD12	1.80	0.47
21:24:16:ALA:HA	21:24:19:ARG:NH2	2.30	0.47
22:25:39:THR:HG21	54:01:2336:A:N6	2.30	0.47
27:30:52:LYS:HE2	27:30:56:LYS:H	1.79	0.47
37:G:104:VAL:O	37:G:108:ARG:HG2	2.15	0.47
37:G:138:GLU:O	37:G:142:ARG:HG2	2.15	0.47
38:H:14:ARG:HD2	53:A:875:U:O2'	2.15	0.47
38:H:48:PHE:HB3	38:H:60:LEU:HD23	1.98	0.47
38:H:63:LYS:NZ	38:H:63:LYS:HB3	2.30	0.47
38:H:95:MET:HE2	38:H:98:LEU:HD11	1.97	0.47
49:S:35:ARG:NH1	49:S:35:ARG:HB3	2.30	0.47
49:S:54:ARG:HG3	49:S:55:GLN:HG2	1.97	0.47
51:U:50:SER:HA	51:U:53:LYS:HD2	1.97	0.47
52:03:15:VAL:HG11	52:03:220:ALA:HB1	1.97	0.47
53:A:86:G:H4'	53:A:87:C:C5	2.50	0.47
53:A:477:C:H2'	53:A:478:A:C8	2.50	0.47
53:A:880:C:O2'	53:A:881:G:H5'	2.15	0.47
54:01:807:U:H2'	54:01:808:G:H8	1.80	0.47
54:01:1119:U:H2'	54:01:1120:G:H8	1.80	0.47
54:01:2841:C:H2'	54:01:2842:G:C8	2.49	0.47
54:01:2859:G:H2'	54:01:2860:A:C8	2.50	0.47
59:Z:350:VAL:HG13	59:Z:354:ASP:HB2	1.97	0.47
7:10:2:ALA:HB1	7:10:6:GLN:HB2	1.97	0.46
32:B:209:VAL:HA	32:B:212:TYR:HD2	1.79	0.46
33:C:102:ILE:HD12	33:C:102:ILE:O	2.14	0.46
35:E:110:MET:HG3	35:E:139:THR:HG21	1.96	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:A:422:C:H4'	53:A:423:G:H5''	1.97	0.46
53:A:1033:G:C3'	53:A:1034:G:H5''	2.42	0.46
53:A:1476:A:H2'	53:A:1477:U:O4'	2.15	0.46
54:01:863:A:H2'	54:01:864:G:C8	2.50	0.46
54:01:975:A:H1'	54:01:990:A:C6	2.50	0.46
54:01:1285:A:H2	54:01:1328:A:H5''	1.80	0.46
54:01:1434:A:H2'	54:01:1435:G:C8	2.49	0.46
54:01:1744:A:H3'	54:01:1745:A:H8	1.80	0.46
54:01:2292:U:H2'	54:01:2293:G:C8	2.50	0.46
58:Y:76:A:H5'	59:Z:220:ILE:HD11	1.96	0.46
1:04:216:ARG:HG3	1:04:216:ARG:NH1	2.31	0.46
4:07:39:VAL:HG12	4:07:84:ILE:O	2.16	0.46
6:09:83:LYS:HA	6:09:149:GLU:HB3	1.98	0.46
32:B:46:VAL:HA	32:B:49:PHE:CE1	2.50	0.46
33:C:49:ALA:HB1	33:C:75:VAL:HG22	1.98	0.46
35:E:28:ARG:HH12	53:A:15:G:H4'	1.80	0.46
41:K:84:MET:SD	41:K:110:THR:OG1	2.73	0.46
48:R:41:SER:HB3	48:R:51:GLN:HG2	1.97	0.46
53:A:1502:A:H5'	53:A:1504:G:N7	2.30	0.46
54:01:662:G:O2'	54:01:663:G:H5'	2.15	0.46
54:01:736:C:H2'	54:01:737:C:C6	2.50	0.46
54:01:769:U:H2'	54:01:770:G:C8	2.49	0.46
54:01:1170:C:H2'	54:01:1171:G:N7	2.30	0.46
54:01:1388:G:H2'	54:01:1389:G:H8	1.80	0.46
54:01:2055:C:H5'	54:01:2056:G:O5'	2.15	0.46
54:01:2127:G:H2'	54:01:2128:G:O4'	2.15	0.46
54:01:2537:U:H2'	54:01:2538:C:H6	1.80	0.46
59:Z:243:GLU:HA	59:Z:252:LYS:HA	1.97	0.46
59:Z:373:ARG:HA	59:Z:387:VAL:HG23	1.97	0.46
3:06:21:ARG:HG2	3:06:110:SER:OG	2.16	0.46
4:07:105:ILE:C	4:07:108:PRO:HD2	2.36	0.46
5:08:41:GLU:HG3	5:08:54:ARG:NH2	2.30	0.46
6:09:31:VAL:N	6:09:32:PRO:HD2	2.30	0.46
37:G:99:ALA:O	37:G:103:ILE:HG13	2.15	0.46
38:H:29:SER:HB2	53:A:589:U:H5''	1.97	0.46
39:I:49:GLN:N	39:I:50:PRO:CD	2.78	0.46
43:M:85:TYR:O	43:M:89:ARG:HG2	2.14	0.46
43:M:88:LEU:HD23	43:M:88:LEU:O	2.16	0.46
45:O:21:THR:HG21	53:A:658:C:H1'	1.97	0.46
53:A:1203:C:H2'	53:A:1204:A:H8	1.80	0.46
54:01:57:C:H2'	54:01:58:G:O4'	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:01:244:A:H2'	54:01:245:G:O4'	2.15	0.46
54:01:302:C:H2'	54:01:303:G:H8	1.80	0.46
54:01:593:U:H2'	54:01:594:U:C6	2.49	0.46
54:01:796:C:H2'	54:01:797:G:C8	2.50	0.46
54:01:1594:U:H2'	54:01:1595:C:C6	2.50	0.46
54:01:1999:C:H5''	54:01:2723:C:O2'	2.16	0.46
54:01:2123:G:N2	54:01:2175:C:H42	2.13	0.46
59:Z:12:VAL:HG22	59:Z:202:PRO:HD3	1.97	0.46
59:Z:324:LYS:CE	59:Z:342:GLU:HA	2.45	0.46
2:05:8:LYS:HB2	2:05:201:LEU:HD11	1.97	0.46
2:05:159:LYS:HE2	54:01:2512:C:O2'	2.15	0.46
4:07:140:ILE:HD12	4:07:140:ILE:N	2.30	0.46
8:11:59:THR:HB	8:11:67:THR:CG2	2.45	0.46
14:17:74:VAL:O	14:17:78:VAL:HG23	2.15	0.46
15:18:80:VAL:HG23	15:18:80:VAL:O	2.15	0.46
32:B:94:ARG:HG2	53:A:1100:C:OP2	2.15	0.46
32:B:118:THR:HA	32:B:121:GLN:NE2	2.25	0.46
34:D:103:ARG:HD2	34:D:167:PRO:HG2	1.96	0.46
34:D:186:GLU:HB3	34:D:189:ASP:OD2	2.15	0.46
35:E:45:VAL:HG11	35:E:116:VAL:HG23	1.96	0.46
35:E:79:THR:OG1	35:E:80:LEU:N	2.48	0.46
36:F:51:ILE:C	36:F:53:LYS:H	2.17	0.46
38:H:80:PRO:HG2	53:A:878:A:C5'	2.45	0.46
40:J:53:ILE:HG13	44:N:84:ARG:CD	2.45	0.46
53:A:458:U:H2'	53:A:459:A:H8	1.81	0.46
53:A:579:A:H2'	53:A:580:C:C6	2.50	0.46
53:A:742:G:H2'	53:A:743:A:H8	1.80	0.46
53:A:784:A:H4'	54:01:1837:C:OP1	2.16	0.46
54:01:1278:C:H2'	54:01:1279:G:C8	2.49	0.46
54:01:1411:U:H2'	54:01:1412:U:C6	2.50	0.46
54:01:1989:G:H2'	54:01:1990:C:O4'	2.16	0.46
54:01:2001:C:H4'	54:01:2689:U:O2'	2.15	0.46
54:01:2292:U:H2'	54:01:2293:G:H8	1.81	0.46
54:01:2628:C:H3'	54:01:2629:U:H5'	1.97	0.46
56:W:6:G:O2'	56:W:7:G:H5'	2.15	0.46
56:W:48:C:H2'	56:W:59:A:H4'	1.97	0.46
59:Z:131:ILE:HG13	59:Z:199:ILE:HD12	1.98	0.46
2:05:133:THR:HG23	2:05:134:HIS:N	2.31	0.46
4:07:23:SER:HB3	4:07:26:GLN:HG3	1.98	0.46
4:07:127:TYR:CD2	4:07:155:ILE:HD12	2.51	0.46
42:L:20:VAL:HB	42:L:94:TYR:CE1	2.50	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:N:20:PHE:O	44:N:21:ALA:CB	2.64	0.46
47:Q:6:THR:C	47:Q:7:LEU:HD12	2.35	0.46
48:R:33:THR:HG23	48:R:35:SER:H	1.79	0.46
49:S:36:ARG:HG2	53:A:1320:C:H41	1.80	0.46
54:01:862:G:H2'	54:01:863:A:O4'	2.15	0.46
54:01:1765:U:H2'	54:01:1766:G:C8	2.51	0.46
54:01:1893:C:H2'	54:01:1894:C:H5'	1.97	0.46
54:01:2475:C:H2'	54:01:2476:A:H5'	1.97	0.46
55:02:70:C:H2'	55:02:71:C:C6	2.50	0.46
58:Y:7:A:H3'	58:Y:8:U:C5'	2.46	0.46
59:Z:206:ILE:HG22	59:Z:270:ALA:HB3	1.98	0.46
59:Z:332:PHE:O	59:Z:333:ARG:HG2	2.16	0.46
1:04:61:TYR:HE1	54:01:1816:C:H3'	1.80	0.46
2:05:51:THR:HB	2:05:79:LEU:HD23	1.98	0.46
7:10:11:ILE:HD11	7:10:62:ARG:HG2	1.96	0.46
9:12:69:ARG:HA	9:12:89:PHE:HD2	1.80	0.46
11:14:23:ILE:HD12	11:14:23:ILE:N	2.30	0.46
13:16:79:LEU:HD23	13:16:83:LEU:HB2	1.97	0.46
40:J:30:LYS:HA	40:J:34:ALA:HA	1.98	0.46
44:N:86:ALA:HA	44:N:89:ARG:NH1	2.30	0.46
53:A:206:C:H2'	53:A:207:C:O4'	2.16	0.46
53:A:418:C:H2'	53:A:419:C:C6	2.51	0.46
53:A:886:G:H2'	53:A:887:G:O4'	2.15	0.46
54:01:1468:U:H2'	54:01:1522:A:N6	2.31	0.46
54:01:2156:G:C2'	54:01:2157:G:H5'	2.45	0.46
55:02:5:U:H2'	55:02:6:G:H8	1.81	0.46
58:Y:15:G:N2	58:Y:21:A:H1'	2.31	0.46
59:Z:305:GLU:CB	59:Z:389:ALA:HB3	2.46	0.46
6:09:132:PHE:HB2	6:09:140:ALA:HB3	1.98	0.46
10:13:2:ILE:HB	10:13:33:ALA:HB3	1.97	0.46
13:16:30:ARG:HH21	13:16:31:HIS:HE1	1.64	0.46
17:20:37:GLU:O	17:20:39:LEU:HD12	2.16	0.46
17:20:54:VAL:HG12	17:20:55:ASP:N	2.31	0.46
32:B:95:TRP:HZ2	32:B:100:LEU:HG	1.80	0.46
33:C:133:MET:O	33:C:137:VAL:HG23	2.15	0.46
37:G:68:VAL:HG21	37:G:103:ILE:HD11	1.98	0.46
53:A:35:G:H2'	53:A:36:C:C6	2.49	0.46
53:A:358:U:H2'	53:A:359:G:C8	2.51	0.46
53:A:429:U:H4'	53:A:430:A:O5'	2.14	0.46
53:A:536:C:H2'	53:A:537:G:C8	2.51	0.46
53:A:1163:A:H2'	53:A:1164:G:C8	2.50	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:01:100:U:H5''	54:01:101:A:O5'	2.16	0.46
54:01:575:A:H2'	54:01:576:U:H6	1.79	0.46
54:01:2055:C:H2'	54:01:2504:U:H5''	1.98	0.46
54:01:2273:A:H2'	54:01:2274:A:C8	2.50	0.46
1:04:200:MET:CE	53:A:773:G:H4'	2.45	0.46
2:05:121:THR:HB	2:05:127:PHE:CE2	2.50	0.46
3:06:131:THR:HG23	54:01:321:U:H5''	1.97	0.46
4:07:163:GLU:HA	4:07:166:ARG:NH1	2.31	0.46
8:11:101:SER:HA	8:11:140:GLU:O	2.16	0.46
9:12:110:PRO:O	9:12:115:GLY:HA3	2.16	0.46
40:J:9:ARG:HB3	40:J:73:LEU:HD23	1.97	0.46
40:J:65:TYR:HB3	44:N:95:LEU:HD11	1.98	0.46
43:M:7:ASN:ND2	43:M:9:PRO:HD3	2.31	0.46
51:U:11:PHE:C	51:U:13:VAL:H	2.19	0.46
52:03:10:VAL:O	52:03:14:LYS:HG3	2.16	0.46
52:03:181:ASP:O	52:03:185:LEU:HG	2.16	0.46
53:A:59:A:H5''	53:A:387:U:H5''	1.98	0.46
54:01:49:A:H5'	54:01:51:G:O4'	2.16	0.46
54:01:163:C:H2'	54:01:164:C:O4'	2.16	0.46
54:01:582:A:H2'	54:01:583:G:H8	1.80	0.46
54:01:996:A:H2'	54:01:997:G:C8	2.51	0.46
54:01:1443:U:H2'	54:01:1444:G:H8	1.78	0.46
58:Y:47:U:H2'	58:Y:50:U:OP1	2.16	0.46
58:Y:49:C:H2'	58:Y:50:U:H6	1.81	0.46
1:04:174:ARG:HG2	1:04:174:ARG:HH11	1.80	0.46
1:04:206:LYS:HD3	54:01:729:G:C8	2.51	0.46
11:14:17:LYS:HD3	54:01:663:G:H5''	1.97	0.46
13:16:30:ARG:HE	13:16:31:HIS:CE1	2.33	0.46
16:19:12:ARG:HG2	16:19:12:ARG:HH21	1.81	0.46
20:23:5:ARG:HB2	54:01:85:G:OP1	2.16	0.46
30:33:51:LYS:HG3	30:33:52:GLY:N	2.30	0.46
45:O:54:GLY:O	45:O:58:MET:HG2	2.16	0.46
48:R:41:SER:CB	48:R:51:GLN:HE21	2.29	0.46
50:T:54:GLN:N	50:T:55:PRO:HD2	2.31	0.46
53:A:641:U:H4'	53:A:642:A:C8	2.51	0.46
54:01:172:A:H2'	54:01:173:A:H8	1.80	0.46
54:01:1041:G:H2'	54:01:1042:G:H8	1.81	0.46
54:01:1877:A:H2'	54:01:1878:G:O4'	2.16	0.46
54:01:2717:C:H2'	54:01:2718:G:O4'	2.16	0.46
59:Z:246:GLY:CA	59:Z:290:GLN:HG3	2.38	0.46
5:08:125:PRO:HD2	5:08:129:GLU:O	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:33:21:PHE:O	30:33:49:VAL:HG23	2.16	0.46
41:K:28:ASN:HD21	41:K:56:LYS:HD2	1.81	0.46
44:N:2:LYS:O	44:N:5:MET:N	2.48	0.46
52:03:177:LYS:N	52:03:180:PHE:HD2	2.13	0.46
53:A:757:U:H2'	53:A:758:C:O4'	2.16	0.46
53:A:1328:C:H2'	53:A:1329:A:H8	1.81	0.46
54:01:151:C:H2'	54:01:152:A:H8	1.80	0.46
54:01:543:G:H5'	54:01:543:G:H8	1.80	0.46
54:01:1678:A:H2'	54:01:1679:A:O4'	2.16	0.46
54:01:1917:U:C2'	54:01:1918:A:H5'	2.46	0.46
54:01:2234:G:O2'	54:01:2235:G:H5'	2.16	0.46
54:01:2347:C:C5	54:01:2382:G:H1'	2.51	0.46
1:04:75:ALA:HB2	1:04:95:TYR:CD1	2.50	0.45
3:06:163:ASN:HB2	54:01:322:A:OP2	2.16	0.45
6:09:32:PRO:HA	23:26:38:TRP:CD1	2.51	0.45
7:10:24:SER:O	7:10:116:GLU:HB2	2.16	0.45
7:10:78:GLY:N	7:10:79:PRO:CD	2.78	0.45
8:11:29:GLN:NE2	54:01:1096:A:H61	2.07	0.45
11:14:29:LYS:HG2	11:14:30:THR:HG23	1.97	0.45
11:14:36:LYS:HE2	54:01:808:G:OP2	2.17	0.45
12:15:13:HIS:HE1	54:01:2265:U:H4'	1.81	0.45
14:17:108:ASP:O	14:17:112:GLU:HG3	2.16	0.45
18:21:74:ILE:HG23	18:21:74:ILE:O	2.16	0.45
32:B:163:ILE:HG22	32:B:168:GLU:OE1	2.16	0.45
34:D:198:LEU:HA	34:D:201:GLU:OE1	2.15	0.45
37:G:100:MET:O	37:G:104:VAL:HG23	2.17	0.45
46:P:78:VAL:O	46:P:78:VAL:HG22	2.16	0.45
47:Q:60:ILE:CG2	47:Q:72:TRP:HB3	2.46	0.45
52:03:165:ASN:HA	52:03:171:ILE:HA	1.98	0.45
53:A:16:A:O2'	53:A:17:U:H5'	2.16	0.45
53:A:1291:U:H2'	53:A:1292:G:C8	2.51	0.45
53:A:1387:G:H2'	53:A:1388:C:C6	2.50	0.45
53:A:1496:C:H2'	53:A:1497:G:O4'	2.15	0.45
54:01:1366:A:H2'	54:01:1367:A:O4'	2.15	0.45
54:01:1444:G:H2'	54:01:1445:G:H8	1.81	0.45
54:01:2009:A:H2'	54:01:2010:G:H8	1.81	0.45
54:01:2086:U:H2'	54:01:2087:G:C8	2.50	0.45
54:01:2475:C:C2'	54:01:2476:A:H5'	2.46	0.45
54:01:2771:C:H2'	54:01:2772:C:H6	1.79	0.45
59:Z:338:THR:H	59:Z:363:ILE:HD11	1.81	0.45
59:Z:376:ILE:HB	59:Z:384:GLY:HA3	1.97	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:04:163:ILE:HG23	1:04:171:VAL:CG1	2.47	0.45
2:05:194:PRO:HA	54:01:2680:U:H5'	1.99	0.45
8:11:6:ALA:HB3	8:11:60:VAL:O	2.17	0.45
33:C:6:PRO:HG2	33:C:200:TRP:HE1	1.81	0.45
36:F:50:PRO:HG3	36:F:55:HIS:CE1	2.51	0.45
37:G:75:LYS:HD3	37:G:88:VAL:HG11	1.98	0.45
39:I:26:LYS:C	39:I:27:ILE:HD12	2.36	0.45
49:S:69:LYS:O	49:S:72:GLU:HB2	2.17	0.45
50:T:70:LYS:HA	50:T:73:ARG:NE	2.21	0.45
52:03:23:ILE:HA	52:03:26:ALA:HB3	1.98	0.45
53:A:1219:A:H2'	53:A:1220:G:C8	2.51	0.45
54:01:1196:C:H2'	54:01:1197:G:C8	2.51	0.45
54:01:1532:A:H1'	54:01:1540:G:H22	1.81	0.45
59:Z:237:LYS:HB3	59:Z:267:GLU:CD	2.37	0.45
1:04:14:HIS:O	1:04:203:VAL:HG21	2.16	0.45
2:05:25:THR:HG21	2:05:193:VAL:HG22	1.99	0.45
2:05:111:GLY:HA3	2:05:201:LEU:HD23	1.97	0.45
7:10:98:GLU:HA	7:10:101:LYS:HD3	1.99	0.45
7:10:107:GLU:HG2	7:10:107:GLU:O	2.17	0.45
10:13:41:ILE:C	10:13:41:ILE:HD12	2.37	0.45
15:18:91:VAL:HG21	15:18:96:LEU:HD21	1.96	0.45
32:B:20:ARG:HE	32:B:21:TYR:HD1	1.65	0.45
39:I:80:HIS:CD2	39:I:105:ARG:HA	2.52	0.45
53:A:211:G:H2'	53:A:212:G:H5'	1.99	0.45
53:A:220:G:O2'	53:A:221:C:H5'	2.16	0.45
53:A:1323:G:H2'	53:A:1324:A:C8	2.51	0.45
54:01:2639:A:H2'	54:01:2640:G:O4'	2.17	0.45
59:Z:64:THR:HG23	59:Z:90:ASN:HB3	1.99	0.45
59:Z:204:ARG:HH11	59:Z:271:GLY:HA3	1.82	0.45
7:10:48:ALA:HB3	7:10:51:TYR:HE1	1.82	0.45
12:15:35:ALA:HB2	12:15:102:LEU:HD11	1.97	0.45
13:16:47:VAL:C	13:16:50:PRO:HD2	2.36	0.45
14:17:81:ARG:O	14:17:85:LYS:HG2	2.16	0.45
19:22:47:VAL:HA	19:22:51:PHE:HD2	1.82	0.45
32:B:162:VAL:O	32:B:184:ALA:HB1	2.15	0.45
35:E:15:ILE:HD12	35:E:15:ILE:N	2.31	0.45
36:F:68:GLN:HE22	53:A:738:C:H5''	1.82	0.45
39:I:70:GLY:HA3	53:A:1371:G:O3'	2.16	0.45
48:R:70:THR:CG2	48:R:71:ASP:H	2.11	0.45
52:03:27:ILE:O	52:03:30:LEU:HG	2.17	0.45
52:03:192:LEU:O	52:03:196:LEU:HD13	2.15	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:A:306:A:O2'	53:A:307:C:H5'	2.17	0.45
53:A:1346:A:O2'	53:A:1347:G:H4'	2.16	0.45
54:01:248:G:C2	54:01:2431:U:H4'	2.51	0.45
54:01:388:G:N7	54:01:390:U:H2'	2.32	0.45
54:01:2146:C:H4'	54:01:2147:A:C5	2.52	0.45
54:01:2406:A:H5'	54:01:2407:A:OP1	2.16	0.45
4:07:36:ASN:HB3	4:07:152:ASP:OD1	2.15	0.45
8:11:79:LEU:CD1	8:11:137:LEU:HD13	2.45	0.45
23:26:25:LYS:HE2	23:26:25:LYS:HA	1.99	0.45
27:30:28:SER:OG	27:30:39:ARG:HD2	2.17	0.45
34:D:50:TYR:CD2	53:A:508:U:H4'	2.51	0.45
40:J:17:LEU:O	40:J:20:GLN:HG2	2.17	0.45
49:S:18:VAL:O	49:S:22:VAL:HG23	2.17	0.45
52:03:62:ALA:C	52:03:160:GLN:HE22	2.20	0.45
52:03:69:THR:HG23	52:03:159:GLY:C	2.37	0.45
53:A:599:C:H2'	53:A:600:A:H8	1.82	0.45
54:01:825:A:H2'	54:01:826:U:O4'	2.16	0.45
54:01:854:C:H2'	54:01:855:G:H8	1.81	0.45
54:01:878:A:H3'	54:01:879:G:C8	2.47	0.45
54:01:1138:G:H2'	54:01:1139:G:O4'	2.16	0.45
54:01:1181:U:H2'	54:01:1182:G:C8	2.51	0.45
54:01:1258:U:H2'	54:01:1259:G:H8	1.82	0.45
54:01:2096:C:H2'	54:01:2097:A:H8	1.81	0.45
54:01:2786:U:O2'	54:01:2787:C:H5'	2.16	0.45
59:Z:304:PHE:HA	59:Z:392:LEU:HD23	1.98	0.45
59:Z:322:PHE:O	59:Z:349:MET:SD	2.75	0.45
59:Z:335:THR:HG21	59:Z:366:ILE:HG12	1.99	0.45
3:06:68:ALA:HA	54:01:1255:U:C5	2.51	0.45
11:14:18:ARG:HH22	54:01:1249:U:H2'	1.81	0.45
14:17:106:LEU:O	14:17:106:LEU:HD23	2.16	0.45
15:18:29:VAL:HG13	15:18:79:VAL:O	2.16	0.45
16:19:78:PHE:HE1	16:19:109:VAL:HA	1.80	0.45
21:24:65:VAL:O	21:24:65:VAL:HG13	2.17	0.45
42:L:43:LYS:HB3	42:L:44:PRO:HD3	1.99	0.45
52:03:26:ALA:HB1	52:03:222:VAL:HG11	1.98	0.45
53:A:335:C:H2'	53:A:336:A:H8	1.79	0.45
53:A:1175:G:H2'	53:A:1176:A:H8	1.81	0.45
53:A:1434:A:H2'	53:A:1435:G:O4'	2.16	0.45
54:01:708:G:N2	54:01:724:U:H1'	2.31	0.45
54:01:1111:A:O2'	54:01:1112:G:H4'	2.17	0.45
54:01:2029:G:O6	54:01:2032:G:H5''	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:01:2110:G:H2'	54:01:2120:G:OP1	2.16	0.45
54:01:2220:U:H2'	54:01:2221:G:H8	1.81	0.45
54:01:2398:U:H2'	54:01:2399:G:C8	2.52	0.45
2:05:142:VAL:HB	2:05:143:PRO:HD2	1.98	0.45
4:07:88:VAL:HA	55:02:42:C:O2	2.17	0.45
7:10:11:ILE:O	7:10:15:VAL:HG23	2.17	0.45
15:18:30:TRP:CD2	15:18:37:LYS:HE3	2.51	0.45
24:27:7:ARG:CD	24:27:7:ARG:H	2.29	0.45
25:28:23:LEU:CD2	25:28:28:LEU:HD12	2.46	0.45
41:K:122:PRO:HG2	51:U:34:ARG:O	2.16	0.45
43:M:1:ALA:O	43:M:8:ILE:HG12	2.17	0.45
54:01:394:C:H2'	54:01:395:U:O4'	2.16	0.45
54:01:577:G:OP1	54:01:2502:G:H2'	2.17	0.45
54:01:766:U:H2'	54:01:767:U:C6	2.52	0.45
54:01:1535:A:H3'	54:01:1536:C:H5'	1.99	0.45
54:01:2154:A:H2'	54:01:2155:U:C6	2.51	0.45
54:01:2423:U:O2'	54:01:2425:A:H2'	2.17	0.45
54:01:2601:C:H2'	54:01:2603:G:C8	2.51	0.45
54:01:2623:G:H2'	54:01:2624:G:C8	2.52	0.45
54:01:2723:C:H2'	54:01:2724:U:O4'	2.17	0.45
58:Y:2:C:H4'	59:Z:87:TYR:HE1	1.81	0.45
59:Z:24:LYS:HG2	59:Z:104:VAL:CG2	2.45	0.45
59:Z:304:PHE:CZ	59:Z:360:VAL:HB	2.52	0.45
12:15:41:LEU:HD13	12:15:124:LEU:HD22	1.98	0.45
14:17:43:ASN:ND2	14:17:46:GLU:HG2	2.30	0.45
15:18:87:ARG:HB3	15:18:87:ARG:NH1	2.32	0.45
19:22:50:LEU:HD23	24:27:26:PHE:CE2	2.52	0.45
33:C:13:ILE:HD12	33:C:13:ILE:N	2.31	0.45
34:D:94:GLU:HA	34:D:99:ASN:HD22	1.79	0.45
39:I:105:ARG:HH11	39:I:105:ARG:HG3	1.82	0.45
39:I:129:ARG:HG2	39:I:129:ARG:HH11	1.81	0.45
44:N:58:ARG:NH2	53:A:980:C:H4'	2.31	0.45
49:S:9:PHE:CE2	53:A:1318:A:H4'	2.52	0.45
52:03:27:ILE:HB	52:03:182:ALA:CB	2.45	0.45
53:A:142:G:H2'	53:A:143:A:O4'	2.17	0.45
53:A:1118:U:H2'	53:A:1119:C:H6	1.82	0.45
53:A:1158:C:H2'	53:A:1159:U:H4'	1.98	0.45
53:A:1277:C:H2'	53:A:1278:G:H5''	1.98	0.45
54:01:616:A:H2'	54:01:617:G:O4'	2.17	0.45
54:01:808:G:H2'	54:01:809:G:H8	1.82	0.45
54:01:1482:G:H1'	54:01:1509:A:C2	2.52	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:01:2391:G:H4'	54:01:2392:A:OP1	2.17	0.45
1:04:270:ARG:HB3	1:04:270:ARG:NH1	2.32	0.45
9:12:27:ARG:HH22	54:01:1142:A:H4'	1.82	0.45
16:19:57:ARG:HG2	16:19:57:ARG:HH11	1.81	0.45
22:25:55:LEU:CD1	22:25:76:ILE:HD12	2.47	0.45
31:34:30:GLU:HG3	31:34:32:LYS:H	1.82	0.45
32:B:42:LEU:HA	32:B:45:THR:OG1	2.15	0.45
34:D:113:ALA:O	34:D:117:VAL:HG23	2.17	0.45
37:G:3:ARG:HD2	37:G:4:ARG:NH1	2.31	0.45
38:H:63:LYS:O	38:H:70:VAL:HG23	2.17	0.45
54:01:576:U:H2'	54:01:577:G:C8	2.52	0.45
54:01:1363:C:H2'	54:01:1364:G:H8	1.82	0.45
54:01:2236:U:H2'	54:01:2237:G:O4'	2.17	0.45
56:W:51:C:H2'	56:W:52:G:C8	2.52	0.45
61:Y:101:PHE:N	59:Z:274:VAL:HA	2.32	0.45
1:04:96:LYS:HD2	54:01:1490:A:N6	2.31	0.45
23:26:5:GLN:HG3	23:26:49:ARG:H	1.82	0.45
27:30:54:ILE:HG23	27:30:56:LYS:N	2.31	0.45
31:34:15:LYS:HB2	31:34:15:LYS:HZ2	1.82	0.45
32:B:221:ARG:NH1	32:B:224:ARG:HH11	2.14	0.45
35:E:25:LYS:HG2	53:A:923:A:OP1	2.17	0.45
35:E:47:PHE:CE2	35:E:137:ARG:HG2	2.52	0.45
38:H:103:VAL:HA	38:H:124:ILE:HA	1.99	0.45
39:I:86:LEU:O	39:I:94:ARG:HD2	2.16	0.45
41:K:59:PRO:HB3	41:K:91:GLY:HA2	1.97	0.45
53:A:1349:A:H2'	53:A:1350:A:O4'	2.17	0.45
53:A:1464:U:H2'	53:A:1465:A:C8	2.51	0.45
54:01:354:A:H2'	54:01:355:U:O4'	2.17	0.45
54:01:1709:U:H2'	54:01:1710:G:H8	1.81	0.45
54:01:2605:U:H2'	54:01:2606:C:C6	2.51	0.45
59:Z:321:PRO:HB2	59:Z:349:MET:HG3	1.98	0.45
59:Z:372:LEU:HB3	59:Z:388:VAL:CG2	2.47	0.45
1:04:233:GLY:HA3	54:01:2598:A:H5''	1.98	0.44
2:05:146:ILE:HA	2:05:159:LYS:HE3	1.98	0.44
5:08:140:ILE:HD12	5:08:141:GLY:N	2.32	0.44
11:14:51:GLU:HB2	11:14:54:GLN:HB3	1.98	0.44
12:15:12:MET:SD	12:15:72:PRO:HD2	2.57	0.44
13:16:10:LEU:O	13:16:12:ARG:HG3	2.17	0.44
13:16:25:ALA:O	13:16:29:VAL:HG23	2.17	0.44
13:16:113:ILE:HG23	13:16:113:ILE:O	2.16	0.44
23:26:7:THR:OG1	23:26:9:LYS:HG3	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:31:46:VAL:HG12	28:31:47:ILE:N	2.32	0.44
38:H:55:LYS:HE3	53:A:653:U:O4'	2.17	0.44
38:H:112:ASP:O	38:H:116:ARG:HG2	2.17	0.44
52:03:54:LYS:HE3	52:03:56:ASP:OD2	2.18	0.44
52:03:69:THR:HG23	52:03:159:GLY:HA2	1.99	0.44
53:A:490:C:H2'	53:A:491:G:H8	1.82	0.44
54:01:116:C:H2'	54:01:117:G:O4'	2.17	0.44
54:01:1053:C:C3'	54:01:1054:A:H5''	2.47	0.44
54:01:1097:U:H2'	54:01:1098:A:O4'	2.17	0.44
54:01:1739:A:H2'	54:01:1740:G:O4'	2.18	0.44
54:01:1827:U:O2'	54:01:1828:G:H5'	2.17	0.44
59:Z:15:GLY:HA2	59:Z:79:VAL:H	1.82	0.44
59:Z:19:HIS:CG	59:Z:112:MET:HG3	2.52	0.44
59:Z:184:TRP:HA	59:Z:187:LYS:CD	2.42	0.44
1:04:61:TYR:CE1	54:01:1816:C:H3'	2.52	0.44
1:04:216:ARG:HG3	1:04:216:ARG:HH11	1.82	0.44
4:07:130:GLY:HA2	4:07:152:ASP:HA	2.00	0.44
20:23:45:GLN:HB2	20:23:58:VAL:HG23	1.99	0.44
23:26:14:GLY:O	23:26:25:LYS:HE2	2.17	0.44
32:B:95:TRP:CZ3	32:B:171:ALA:HA	2.53	0.44
37:G:13:PRO:HA	37:G:20:GLU:HG3	1.99	0.44
44:N:80:ARG:O	44:N:83:VAL:HG12	2.17	0.44
49:S:49:ALA:HA	49:S:58:PRO:HA	1.99	0.44
53:A:79:G:H2'	53:A:80:A:O4'	2.17	0.44
53:A:169:C:H2'	53:A:170:U:C6	2.52	0.44
53:A:434:U:H2'	53:A:435:A:H8	1.81	0.44
53:A:715:A:H2'	53:A:716:A:C8	2.52	0.44
53:A:1463:U:H2'	53:A:1464:U:C6	2.52	0.44
54:01:615:U:H5''	54:01:616:A:OP2	2.16	0.44
54:01:1595:C:H2'	54:01:1596:A:H8	1.82	0.44
54:01:2552:U:C6	54:01:2554:U:H5'	2.52	0.44
56:W:69:C:H2'	56:W:70:G:C8	2.52	0.44
2:05:62:LYS:HB2	2:05:63:PRO:HD3	1.98	0.44
5:08:10:VAL:HG12	5:08:47:ASN:O	2.18	0.44
6:09:57:LYS:O	6:09:61:VAL:HG13	2.17	0.44
8:11:93:ASN:HD22	54:01:1077:A:H5'	1.82	0.44
41:K:110:THR:HB	48:R:72:ARG:NH1	2.32	0.44
41:K:116:PRO:HB3	53:A:676:A:H1'	1.99	0.44
48:R:40:PRO:HG2	48:R:43:ILE:HG12	1.98	0.44
53:A:1370:G:O2'	53:A:1371:G:H5'	2.17	0.44
54:01:322:A:H5'	54:01:340:A:C1'	2.44	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:01:610:C:H2'	54:01:611:C:H6	1.81	0.44
54:01:1123:C:H2'	54:01:1124:G:H8	1.81	0.44
58:Y:6:G:C3'	58:Y:7:A:H5''	2.37	0.44
59:Z:91:MET:O	59:Z:95:ALA:N	2.50	0.44
59:Z:204:ARG:HA	59:Z:204:ARG:HE	1.82	0.44
1:04:86:ARG:HG2	1:04:86:ARG:NH1	2.32	0.44
1:04:184:GLU:HG3	1:04:186:ASP:H	1.82	0.44
2:05:37:VAL:HG23	2:05:92:VAL:CG2	2.48	0.44
2:05:101:PHE:O	2:05:104:VAL:HG22	2.18	0.44
18:21:13:SER:O	18:21:17:VAL:HG23	2.17	0.44
21:24:51:GLN:OE1	21:24:86:LEU:HD11	2.18	0.44
26:29:33:ASN:HD21	43:M:49:GLU:HG3	1.83	0.44
28:31:52:LYS:NZ	28:31:52:LYS:HB3	2.33	0.44
37:G:68:VAL:HG23	37:G:99:ALA:HB1	2.00	0.44
53:A:55:A:C5	59:Z:222:GLY:HA3	2.52	0.44
53:A:147:G:H2'	53:A:148:G:C8	2.53	0.44
53:A:345:C:H4'	53:A:346:G:C4	2.52	0.44
54:01:745:G:O2'	54:01:748:G:H1'	2.17	0.44
54:01:1318:U:H2'	54:01:1319:C:C6	2.52	0.44
54:01:1476:U:H1'	54:01:1732:C:C2	2.53	0.44
54:01:2040:G:H2'	54:01:2041:U:O4'	2.17	0.44
59:Z:391:VAL:C	59:Z:392:LEU:HD22	2.38	0.44
1:04:207:ALA:HB1	54:01:1790:C:H4'	2.00	0.44
1:04:255:LYS:HD3	1:04:269:ARG:HH12	1.81	0.44
22:25:70:PRO:HB3	55:02:12:C:N4	2.33	0.44
27:30:11:LYS:HA	27:30:14:MET:HE3	1.99	0.44
34:D:71:PHE:HE1	34:D:93:LEU:HD11	1.82	0.44
35:E:60:GLN:O	35:E:64:GLU:HG3	2.18	0.44
36:F:50:PRO:HG3	36:F:55:HIS:NE2	2.32	0.44
45:O:86:LEU:HD12	45:O:86:LEU:O	2.17	0.44
53:A:79:G:O2'	53:A:80:A:H5'	2.17	0.44
53:A:1004:A:H1'	53:A:1026:G:C6	2.53	0.44
54:01:1659:G:H2'	54:01:1660:G:O4'	2.17	0.44
54:01:2096:C:H2'	54:01:2097:A:C8	2.52	0.44
54:01:2756:U:H1'	54:01:2757:A:H5''	1.98	0.44
54:01:2900:A:H2'	54:01:2901:C:O4'	2.17	0.44
59:Z:17:ILE:C	59:Z:17:ILE:HD12	2.37	0.44
59:Z:25:THR:HG22	59:Z:46:PHE:HD1	1.82	0.44
1:04:209:ALA:HA	1:04:212:TRP:CE2	2.52	0.44
2:05:97:SER:O	2:05:100:LEU:HD13	2.18	0.44
2:05:149:ASN:CG	2:05:150:GLN:H	2.21	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:06:77:ILE:HG13	3:06:78:TRP:CD1	2.53	0.44
10:13:48:PRO:HB3	53:A:1422:G:C5'	2.48	0.44
15:18:13:LYS:NZ	15:18:80:VAL:HG23	2.32	0.44
34:D:58:GLN:HG3	53:A:544:G:OP1	2.18	0.44
38:H:85:TYR:HE1	38:H:123:GLU:HB2	1.82	0.44
42:L:35:ARG:HH12	42:L:37:TYR:HB3	1.82	0.44
45:O:87:ARG:NE	45:O:87:ARG:HA	2.32	0.44
51:U:35:GLU:O	51:U:36:PHE:HB2	2.17	0.44
52:03:51:ASP:C	52:03:53:ARG:H	2.20	0.44
53:A:458:U:H2'	53:A:459:A:C8	2.53	0.44
53:A:865:A:H2'	53:A:866:C:C6	2.52	0.44
53:A:950:U:H2'	53:A:951:G:C8	2.53	0.44
54:01:96:C:H2'	54:01:97:C:C6	2.52	0.44
54:01:213:A:H2'	54:01:214:G:C8	2.52	0.44
54:01:2019:A:H2	54:01:2035:G:H22	1.65	0.44
54:01:2557:G:H2'	54:01:2558:C:C6	2.53	0.44
54:01:2776:A:H4'	54:01:2777:G:H5''	1.99	0.44
59:Z:293:ALA:HB1	59:Z:297:THR:CG2	2.48	0.44
3:06:70:SER:HB2	3:06:78:TRP:HZ2	1.83	0.44
4:07:32:LYS:HD3	4:07:91:ARG:HH22	1.82	0.44
18:21:16:LYS:HE3	54:01:1266:G:N7	2.33	0.44
21:24:3:THR:HG23	21:24:62:THR:O	2.18	0.44
32:B:55:GLU:O	32:B:59:ILE:HG12	2.18	0.44
32:B:60:ALA:HB3	32:B:223:GLY:HA3	1.99	0.44
35:E:22:LYS:HB3	35:E:29:ILE:CG2	2.47	0.44
42:L:115:LYS:O	42:L:116:TYR:HB2	2.17	0.44
52:03:65:LEU:HD13	52:03:188:ASN:HB3	1.99	0.44
53:A:211:G:C2'	53:A:212:G:H5'	2.47	0.44
53:A:358:U:H2'	53:A:359:G:H8	1.82	0.44
53:A:1301:U:O2	53:A:1301:U:H2'	2.17	0.44
53:A:1306:A:H62	53:A:1331:G:H1'	1.83	0.44
54:01:326:G:H2'	54:01:327:G:H8	1.82	0.44
54:01:2011:U:H2'	54:01:2012:G:O4'	2.17	0.44
59:Z:265:LEU:HD21	59:Z:268:GLY:HA2	2.00	0.44
1:04:156:SER:HB2	54:01:1818:U:H5'	2.00	0.44
1:04:257:ARG:HG3	54:01:1799:G:OP1	2.17	0.44
2:05:141:ARG:HG2	2:05:141:ARG:HH11	1.82	0.44
3:06:47:LYS:HE3	54:01:451:U:OP1	2.18	0.44
6:09:126:GLY:O	6:09:146:VAL:N	2.50	0.44
11:14:49:GLY:HA3	11:14:58:TYR:HE1	1.83	0.44
12:15:64:TRP:HZ3	12:15:106:ASP:HB3	1.83	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:19:91:ARG:HG3	16:19:91:ARG:NH1	2.33	0.44
32:B:33:ALA:HB3	32:B:37:VAL:O	2.17	0.44
32:B:80:LYS:HB2	32:B:92:ASN:ND2	2.33	0.44
34:D:61:ARG:NH2	34:D:67:LEU:HA	2.33	0.44
36:F:93:LYS:O	36:F:94:HIS:CB	2.65	0.44
40:J:42:LEU:HA	40:J:43:PRO:HD2	1.80	0.44
44:N:2:LYS:HE2	53:A:983:A:H5'	1.98	0.44
48:R:17:VAL:HG22	48:R:18:GLN:N	2.33	0.44
50:T:67:HIS:ND1	50:T:67:HIS:O	2.50	0.44
52:03:50:ILE:HB	52:03:57:GLN:HB3	2.00	0.44
52:03:65:LEU:HD22	52:03:188:ASN:CA	2.45	0.44
53:A:1472:U:H2'	53:A:1473:G:C8	2.53	0.44
54:01:1316:U:H2'	54:01:1317:G:H8	1.82	0.44
54:01:1469:A:H2'	54:01:1470:A:C8	2.53	0.44
54:01:2617:U:H2'	54:01:2618:G:O4'	2.18	0.44
54:01:2743:U:H3'	54:01:2744:G:H5''	1.99	0.44
58:Y:73:A:H2'	58:Y:73:A:N3	2.33	0.44
59:Z:66:HIS:CD2	59:Z:66:HIS:N	2.84	0.44
2:05:42:ASN:HB3	54:01:2784:U:H4'	1.98	0.44
4:07:30:VAL:HG22	4:07:95:MET:HE1	2.00	0.44
19:22:49:LYS:C	19:22:50:LEU:HD12	2.39	0.44
23:26:69:GLU:O	23:26:73:ARG:HG3	2.18	0.44
34:D:33:ILE:HG13	34:D:34:GLU:N	2.32	0.44
37:G:39:GLU:HG2	37:G:43:TYR:CE2	2.53	0.44
38:H:4:ASP:HB2	53:A:877:G:O2'	2.17	0.44
40:J:86:ALA:HA	40:J:90:LEU:HD12	2.00	0.44
44:N:27:LYS:O	44:N:31:SER:HB2	2.18	0.44
46:P:14:ARG:HH11	46:P:14:ARG:HG3	1.82	0.44
48:R:56:ARG:O	48:R:60:ARG:HG3	2.17	0.44
52:03:54:LYS:HE3	52:03:56:ASP:CG	2.39	0.44
53:A:211:G:H3'	53:A:211:G:N3	2.33	0.44
53:A:264:C:H2'	53:A:265:G:O4'	2.18	0.44
53:A:407:U:H2'	53:A:408:A:H8	1.82	0.44
53:A:707:U:H2'	53:A:708:C:C6	2.53	0.44
53:A:1354:U:H2'	53:A:1355:G:C8	2.52	0.44
54:01:744:U:H2'	54:01:745:G:O4'	2.17	0.44
54:01:2141:G:H2'	54:01:2142:A:C8	2.48	0.44
54:01:2231:U:H2'	54:01:2232:C:H6	1.83	0.44
54:01:2293:G:H2'	54:01:2294:G:C8	2.52	0.44
54:01:2743:U:H2'	54:01:2744:G:H5''	1.99	0.44
54:01:2836:U:H2'	54:01:2837:A:C8	2.53	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:07:105:ILE:HG13	4:07:106:ALA:N	2.33	0.43
11:14:55:MET:SD	11:14:59:ARG:HD3	2.58	0.43
16:19:48:ASP:O	16:19:52:ARG:N	2.47	0.43
38:H:5:PRO:O	38:H:8:ASP:HB3	2.17	0.43
40:J:34:ALA:O	40:J:35:GLN:HB3	2.18	0.43
47:Q:5:ARG:HD2	53:A:636:U:OP1	2.17	0.43
49:S:52:ASN:CG	49:S:53:GLY:H	2.21	0.43
54:01:68:G:H2'	54:01:69:C:O4'	2.18	0.43
54:01:1295:C:H2'	54:01:1296:G:C8	2.53	0.43
54:01:2144:G:H1'	54:01:2147:A:H61	1.83	0.43
54:01:2329:U:H2'	54:01:2330:G:C8	2.53	0.43
54:01:2869:G:H2'	54:01:2870:C:O4'	2.17	0.43
55:02:91:C:H2'	55:02:92:C:C6	2.52	0.43
1:04:66:PHE:HZ	1:04:86:ARG:HE	1.66	0.43
1:04:213:ARG:HH11	1:04:213:ARG:HG3	1.83	0.43
6:09:94:ILE:HB	6:09:122:LEU:HB2	1.99	0.43
8:11:56:VAL:HG21	8:11:68:PHE:HD2	1.84	0.43
11:14:79:LEU:HD23	11:14:82:LEU:HD22	2.00	0.43
32:B:187:ASP:HB3	32:B:190:SER:OG	2.18	0.43
33:C:143:LEU:HD23	33:C:143:LEU:O	2.18	0.43
37:G:139:ASP:O	37:G:143:MET:HG2	2.18	0.43
40:J:80:THR:HB	40:J:83:THR:OG1	2.19	0.43
45:O:31:LEU:O	45:O:35:ILE:HG13	2.18	0.43
52:03:170:ILE:HG21	54:01:2177:C:H1'	2.00	0.43
53:A:323:U:H2'	53:A:324:G:O4'	2.18	0.43
53:A:392:C:H2'	53:A:393:A:C8	2.51	0.43
53:A:1109:C:H2'	53:A:1110:A:O4'	2.18	0.43
53:A:1492:A:H2'	54:01:1913:A:H2	1.82	0.43
54:01:629:G:H5''	54:01:650:C:O2'	2.18	0.43
54:01:704:G:HO2'	54:01:705:A:H8	1.65	0.43
54:01:859:G:HO2'	54:01:860:U:H6	1.64	0.43
54:01:898:C:H2'	54:01:899:A:O4'	2.18	0.43
54:01:1001:A:H2'	54:01:1002:G:O4'	2.18	0.43
54:01:1105:U:C2'	54:01:1106:G:H5''	2.47	0.43
54:01:2194:U:H2'	54:01:2195:U:C6	2.53	0.43
59:Z:98:MET:O	59:Z:127:VAL:HG13	2.17	0.43
59:Z:238:VAL:HG23	59:Z:266:ASP:O	2.18	0.43
2:05:3:GLY:O	2:05:4:LEU:HD12	2.18	0.43
2:05:4:LEU:HD23	2:05:32:ASN:ND2	2.27	0.43
3:06:164:LEU:HD12	3:06:164:LEU:H	1.83	0.43
4:07:70:ARG:HD2	54:01:2298:A:OP1	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:15:34:LYS:HD3	21:24:82:TYR:HA	1.99	0.43
15:18:90:ALA:HB2	15:18:112:ARG:HA	2.00	0.43
17:20:13:ARG:HG2	17:20:13:ARG:HH11	1.84	0.43
17:20:59:ILE:HG23	17:20:101:ILE:HD13	1.99	0.43
18:21:57:ASN:ND2	18:21:61:ASN:HD22	2.16	0.43
26:29:42:PRO:O	26:29:46:GLY:HA3	2.17	0.43
31:34:1:MET:HE1	31:34:36:ARG:HB2	2.00	0.43
33:C:22:PHE:CE2	40:J:11:LYS:HG3	2.53	0.43
35:E:113:VAL:HG13	35:E:114:LEU:CD1	2.47	0.43
36:F:75:GLU:O	36:F:79:ARG:HD2	2.18	0.43
40:J:102:LEU:N	40:J:102:LEU:HD12	2.33	0.43
42:L:53:ARG:HA	42:L:63:THR:HA	2.00	0.43
48:R:44:THR:O	48:R:46:THR:N	2.44	0.43
52:03:31:LYS:HD2	52:03:182:ALA:N	2.34	0.43
53:A:77:A:H2'	53:A:78:A:C8	2.54	0.43
54:01:1113:U:H2'	54:01:1114:C:C6	2.53	0.43
54:01:2052:A:H2'	54:01:2053:G:H8	1.83	0.43
54:01:2208:C:H2'	54:01:2209:G:H8	1.80	0.43
54:01:2352:A:H2'	54:01:2353:G:O4'	2.18	0.43
58:Y:72:C:H2'	58:Y:73:A:O4'	2.18	0.43
1:04:86:ARG:HG2	1:04:86:ARG:HH11	1.82	0.43
4:07:154:THR:HG21	54:01:2314:A:O4'	2.18	0.43
10:13:66:LYS:NZ	10:13:66:LYS:HB3	2.33	0.43
30:33:31:ILE:O	30:33:31:ILE:HG22	2.19	0.43
33:C:70:ALA:HB2	33:C:114:LEU:HD13	2.00	0.43
39:I:14:SER:HB2	39:I:69:GLY:HA3	2.01	0.43
40:J:59:LYS:HD3	53:A:973:G:OP1	2.18	0.43
45:O:30:LEU:HD21	53:A:658:C:OP1	2.18	0.43
53:A:955:U:H3	53:A:1225:A:H61	1.66	0.43
53:A:1203:C:H2'	53:A:1204:A:C8	2.54	0.43
53:A:1306:A:H2'	53:A:1307:U:O4'	2.18	0.43
54:01:69:C:O2'	54:01:70:G:H5'	2.18	0.43
54:01:365:U:H2'	54:01:366:C:C6	2.53	0.43
54:01:1045:C:H5'	54:01:1046:A:H5'	1.98	0.43
54:01:1061:U:H4'	54:01:1070:A:N3	2.33	0.43
54:01:1186:G:H2'	54:01:1187:G:O4'	2.18	0.43
54:01:1565:C:O2'	54:01:1566:A:H2'	2.18	0.43
54:01:1764:C:H2'	54:01:1765:U:C6	2.54	0.43
54:01:2376:A:H2'	54:01:2377:A:O4'	2.18	0.43
54:01:2555:U:H2'	54:01:2556:C:H5'	2.01	0.43
54:01:2846:G:H2'	54:01:2847:U:O4'	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:Z:32:THR:HB	59:Z:42:ALA:HB3	2.00	0.43
59:Z:297:THR:CG2	59:Z:298:ILE:HD12	2.46	0.43
4:07:32:LYS:HD3	4:07:91:ARG:NH2	2.33	0.43
19:22:7:LEU:HD22	19:22:46:ALA:HA	2.01	0.43
21:24:26:PHE:HE2	21:24:89:ILE:HG13	1.83	0.43
23:26:71:ARG:HD2	23:26:77:TYR:OH	2.17	0.43
26:29:13:THR:HA	26:29:23:LYS:HA	1.99	0.43
32:B:221:ARG:HH11	32:B:221:ARG:HG2	1.83	0.43
44:N:4:SER:O	44:N:8:ARG:N	2.51	0.43
49:S:62:THR:HG22	49:S:63:ASP:N	2.33	0.43
51:U:13:VAL:HG13	51:U:15:LEU:HG	1.99	0.43
53:A:1094:G:H2'	53:A:1094:G:N3	2.34	0.43
54:01:1222:U:H2'	54:01:1223:G:H8	1.81	0.43
54:01:1326:U:H2'	54:01:1327:A:C8	2.53	0.43
54:01:1595:C:H2'	54:01:1596:A:C8	2.53	0.43
54:01:2478:A:C2	54:01:2529:G:H2'	2.53	0.43
59:Z:137:CYS:HB2	59:Z:172:GLY:O	2.19	0.43
1:04:43:ASN:HD21	54:01:1806:C:H1'	1.84	0.43
3:06:159:LEU:O	54:01:321:U:H4'	2.19	0.43
27:30:49:ARG:HG2	54:01:2884:U:H6	1.83	0.43
32:B:46:VAL:HA	32:B:49:PHE:CZ	2.54	0.43
32:B:96:LEU:HD22	53:A:1103:C:H5'	1.99	0.43
33:C:10:ARG:HA	33:C:13:ILE:HD13	2.01	0.43
34:D:59:LYS:HE3	34:D:194:ILE:HG22	2.00	0.43
35:E:89:THR:O	35:E:89:THR:HG22	2.18	0.43
53:A:32:A:H2'	53:A:33:A:C8	2.53	0.43
53:A:203:G:H1'	53:A:465:A:H61	1.84	0.43
53:A:666:G:H5'	53:A:726:C:H1'	2.01	0.43
53:A:691:G:H2'	53:A:692:U:C6	2.54	0.43
54:01:741:U:H2'	54:01:742:A:H8	1.82	0.43
54:01:1177:G:H2'	54:01:1178:C:O4'	2.18	0.43
54:01:1310:G:H3'	54:01:1311:G:C8	2.53	0.43
54:01:2139:U:H2'	54:01:2140:G:C8	2.54	0.43
54:01:2396:G:H2'	54:01:2397:G:H8	1.83	0.43
54:01:2634:A:H2'	54:01:2635:A:C8	2.54	0.43
59:Z:123:ARG:HB2	59:Z:162:PHE:CZ	2.53	0.43
59:Z:139:MET:HA	59:Z:139:MET:HE2	1.99	0.43
59:Z:241:GLU:OE2	59:Z:252:LYS:HD3	2.19	0.43
59:Z:269:ARG:CD	59:Z:272:GLU:HG2	2.48	0.43
3:06:18:THR:HG22	3:06:19:PHE:HD1	1.84	0.43
3:06:163:ASN:ND2	54:01:323:C:H5''	2.34	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:08:41:GLU:HB2	5:08:54:ARG:HE	1.84	0.43
6:09:73:ASN:HB3	6:09:108:VAL:HG23	2.01	0.43
11:14:96:LYS:HB2	11:14:96:LYS:NZ	2.33	0.43
12:15:40:ARG:HH11	12:15:93:VAL:HG11	1.84	0.43
13:16:3:HIS:CD2	54:01:2820:A:H4'	2.53	0.43
13:16:47:VAL:O	13:16:50:PRO:HD2	2.19	0.43
16:19:41:ALA:HB1	54:01:534:U:H5'	2.00	0.43
24:27:43:LEU:HD23	24:27:43:LEU:O	2.19	0.43
27:30:33:SER:HB3	27:30:35:GLU:HG3	2.00	0.43
41:K:22:ILE:HG23	41:K:31:VAL:HG22	2.00	0.43
52:03:27:ILE:CB	52:03:182:ALA:HB1	2.47	0.43
52:03:63:THR:O	52:03:160:GLN:NE2	2.52	0.43
53:A:269:C:H2'	53:A:270:A:C8	2.54	0.43
54:01:146:A:H2'	54:01:147:C:C6	2.53	0.43
54:01:494:G:H2'	54:01:495:G:H8	1.84	0.43
59:Z:70:ASP:N	59:Z:70:ASP:OD1	2.52	0.43
8:11:78:LEU:O	8:11:78:LEU:HD23	2.18	0.43
8:11:127:SER:HA	54:01:1080:A:H1'	2.01	0.43
12:15:12:MET:HG2	12:15:72:PRO:HG2	2.01	0.43
32:B:103:TRP:CH2	32:B:107:ARG:HD3	2.54	0.43
34:D:23:GLY:HA3	53:A:409:U:OP1	2.18	0.43
34:D:186:GLU:HG3	34:D:188:SER:H	1.82	0.43
35:E:160:VAL:HG13	35:E:161:GLU:N	2.33	0.43
40:J:41:PRO:HG3	53:A:1150:A:N3	2.34	0.43
47:Q:22:VAL:HG21	47:Q:60:ILE:HD11	2.01	0.43
52:03:41:SER:C	52:03:217:THR:HG23	2.39	0.43
53:A:505:G:OP2	53:A:535:A:H5'	2.18	0.43
53:A:772:U:H2'	53:A:773:G:C8	2.53	0.43
53:A:945:G:H2'	53:A:945:G:N3	2.33	0.43
54:01:1286:A:H1'	54:01:1288:G:OP2	2.18	0.43
54:01:1695:G:H2'	54:01:1696:G:O4'	2.19	0.43
54:01:1911:U:H2'	54:01:1918:A:N1	2.34	0.43
54:01:2014:A:H2'	54:01:2015:A:C8	2.54	0.43
59:Z:12:VAL:HB	59:Z:76:TYR:HE1	1.81	0.43
59:Z:269:ARG:NE	59:Z:272:GLU:HG2	2.33	0.43
59:Z:333:ARG:HH12	59:Z:372:LEU:HD13	1.83	0.43
59:Z:335:THR:HG22	59:Z:337:VAL:HG22	2.00	0.43
1:04:211:ARG:HH11	1:04:211:ARG:HG3	1.84	0.43
4:07:132:ARG:CZ	54:01:2305:U:H4'	2.49	0.43
8:11:11:GLN:HG2	8:11:54:ILE:O	2.18	0.43
12:15:4:PRO:HG3	12:15:68:PHE:HE2	1.83	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:15:71:LYS:HB3	12:15:93:VAL:O	2.18	0.43
13:16:83:LEU:HD22	13:16:86:ARG:HH22	1.83	0.43
14:17:110:ALA:HB1	14:17:115:LEU:HD23	2.01	0.43
32:B:26:MET:CE	32:B:192:PRO:HD3	2.49	0.43
33:C:39:ARG:HA	33:C:54:ILE:HD11	2.01	0.43
33:C:91:ALA:HA	33:C:94:ALA:HB3	2.01	0.43
35:E:133:ILE:HD12	35:E:133:ILE:N	2.33	0.43
40:J:28:THR:O	40:J:28:THR:HG22	2.19	0.43
43:M:16:ILE:HD12	43:M:16:ILE:H	1.84	0.43
44:N:12:ARG:HH21	44:N:58:ARG:HH21	1.67	0.43
47:Q:16:MET:O	47:Q:19:SER:OG	2.35	0.43
53:A:923:A:H2'	53:A:924:C:O4'	2.18	0.43
53:A:1286:U:C2'	53:A:1287:A:H5'	2.49	0.43
54:01:197:A:H4'	54:01:2069:G:OP2	2.18	0.43
54:01:507:A:C5'	54:01:508:A:H5''	2.33	0.43
54:01:657:U:H2'	54:01:658:U:C6	2.53	0.43
54:01:814:C:H1'	54:01:1225:G:N2	2.34	0.43
54:01:950:G:H2'	54:01:951:C:O4'	2.19	0.43
54:01:1070:A:HO2'	54:01:1071:G:P	2.41	0.43
54:01:1396:U:H5''	54:01:1397:U:OP2	2.19	0.43
54:01:2364:C:H2'	54:01:2365:G:O4'	2.17	0.43
54:01:2861:U:H2'	54:01:2862:G:C8	2.53	0.43
55:02:118:C:H2'	55:02:119:A:C8	2.54	0.43
59:Z:236:ILE:HD12	59:Z:268:GLY:O	2.19	0.43
2:05:5:VAL:HG21	2:05:80:TRP:CE3	2.53	0.43
3:06:118:LEU:HD11	3:06:188:MET:SD	2.59	0.43
5:08:3:VAL:CG2	54:01:2751:G:H4'	2.49	0.43
5:08:26:LYS:HB2	5:08:31:GLU:HG3	2.01	0.43
7:10:60:LEU:HD12	7:10:60:LEU:N	2.33	0.43
7:10:97:LYS:O	7:10:100:ALA:HB3	2.19	0.43
7:10:118:ILE:CG2	7:10:119:PRO:HD3	2.41	0.43
13:16:78:LYS:HG2	13:16:83:LEU:HG	2.01	0.43
32:B:153:MET:HG2	32:B:155:GLY:H	1.83	0.43
34:D:57:LYS:HD2	34:D:203:TYR:OH	2.19	0.43
35:E:133:ILE:O	35:E:137:ARG:HG3	2.19	0.43
38:H:86:LYS:HD2	38:H:90:GLU:CG	2.49	0.43
41:K:22:ILE:HG12	41:K:31:VAL:HG13	1.99	0.43
41:K:28:ASN:ND2	41:K:56:LYS:HD2	2.33	0.43
43:M:14:ALA:HA	43:M:44:ILE:HD11	2.00	0.43
50:T:82:ILE:HD12	50:T:83:ASN:N	2.33	0.43
51:U:16:ARG:NE	51:U:19:LYS:HE2	2.34	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:A:36:C:H2'	53:A:37:U:O4'	2.17	0.43
53:A:90:C:H2'	53:A:91:U:C6	2.54	0.43
54:01:581:C:H2'	54:01:582:A:H8	1.84	0.43
54:01:755:U:H2'	54:01:756:A:C8	2.54	0.43
54:01:1287:A:O2'	54:01:1288:G:H5'	2.18	0.43
54:01:1563:U:H2'	54:01:1564:C:C6	2.54	0.43
54:01:1827:U:H2'	54:01:1828:G:O4'	2.19	0.43
54:01:2298:A:H2'	54:01:2299:U:O4'	2.18	0.43
54:01:2423:U:H5'	54:01:2424:C:C5'	2.49	0.43
54:01:2589:A:H2'	54:01:2590:A:C8	2.53	0.43
54:01:2714:G:H2'	54:01:2715:C:C6	2.54	0.43
58:Y:16:U:H1'	58:Y:60:U:O2'	2.19	0.43
58:Y:54:U:H5'	59:Z:319:HIS:HE1	1.84	0.43
59:Z:219:SER:HB2	59:Z:283:ARG:CD	2.49	0.43
59:Z:306:SER:HB3	59:Z:358:MET:HE1	2.01	0.43
4:07:7:TYR:O	4:07:12:VAL:HG23	2.19	0.42
4:07:153:ILE:HD12	4:07:153:ILE:N	2.34	0.42
4:07:163:GLU:HA	4:07:166:ARG:HH12	1.84	0.42
7:10:27:VAL:HG11	7:10:54:VAL:HG23	2.01	0.42
9:12:35:ARG:HB2	9:12:54:ILE:HD11	2.01	0.42
16:19:71:ASN:OD1	16:19:109:VAL:HG21	2.19	0.42
19:22:74:ILE:HG13	19:22:74:ILE:O	2.19	0.42
32:B:56:LEU:HD23	32:B:59:ILE:HD11	2.01	0.42
34:D:131:ILE:HG12	53:A:620:C:C2	2.54	0.42
42:L:27:PRO:HG3	53:A:553:A:H1'	2.00	0.42
42:L:49:ARG:HH22	53:A:522:C:N4	2.17	0.42
43:M:85:TYR:CE2	53:A:1321:U:H5''	2.54	0.42
51:U:7:GLU:HG3	51:U:11:PHE:HB3	2.01	0.42
52:03:11:ILE:CG2	52:03:220:ALA:HB2	2.49	0.42
53:A:91:U:H2'	53:A:92:U:O4'	2.18	0.42
53:A:730:G:N2	53:A:765:G:H5''	2.33	0.42
53:A:900:A:H2'	53:A:901:A:C8	2.54	0.42
53:A:1399:C:H1'	53:A:1400:C:OP2	2.19	0.42
53:A:1469:C:H2'	53:A:1470:U:O4'	2.19	0.42
53:A:1486:G:H2'	53:A:1487:G:O4'	2.19	0.42
54:01:1213:A:H61	54:01:1236:G:H1'	1.83	0.42
54:01:1441:G:H2'	54:01:1442:U:C6	2.54	0.42
54:01:2027:G:H2'	54:01:2028:U:C6	2.54	0.42
54:01:2293:G:H2'	54:01:2294:G:H8	1.83	0.42
54:01:2690:U:O2'	54:01:2872:A:H1'	2.19	0.42
61:Y:101:PHE:N	59:Z:261:PHE:H	2.17	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:Z:25:THR:HG22	59:Z:46:PHE:CD1	2.54	0.42
59:Z:247:ILE:HG23	59:Z:364:HIS:CB	2.49	0.42
1:04:123:ILE:O	1:04:123:ILE:HG22	2.19	0.42
3:06:105:LEU:CD2	3:06:108:ILE:HD11	2.45	0.42
5:08:132:LEU:HD12	5:08:132:LEU:O	2.19	0.42
8:11:7:TYR:HA	8:11:58:ILE:O	2.19	0.42
16:19:56:PHE:CZ	54:01:536:G:H4'	2.54	0.42
16:19:90:ASP:OD2	16:19:92:LYS:HB3	2.20	0.42
47:Q:29:LYS:HB2	47:Q:36:PHE:CE1	2.53	0.42
53:A:303:A:H2'	53:A:304:U:O4'	2.19	0.42
53:A:991:U:C5	53:A:1212:U:H1'	2.54	0.42
54:01:499:U:H2'	54:01:500:G:O4'	2.19	0.42
54:01:1083:U:H2'	54:01:1085:A:OP2	2.19	0.42
54:01:1444:G:H2'	54:01:1445:G:C8	2.53	0.42
59:Z:52:ALA:HB3	59:Z:55:GLU:HB2	2.00	0.42
59:Z:304:PHE:O	59:Z:359:VAL:HG22	2.19	0.42
1:04:1:ALA:N	1:04:19:VAL:HB	2.34	0.42
5:08:85:LYS:C	5:08:86:LEU:HD12	2.39	0.42
7:10:3:LEU:HD12	7:10:5:LEU:H	1.85	0.42
7:10:72:LEU:HD12	7:10:72:LEU:H	1.85	0.42
21:24:21:ARG:HA	21:24:25:LYS:O	2.19	0.42
31:34:30:GLU:OE2	31:34:31:PRO:HD2	2.18	0.42
35:E:96:GLN:HG3	35:E:97:PRO:HD2	2.00	0.42
37:G:17:PHE:CZ	37:G:22:LEU:HD21	2.54	0.42
41:K:123:PRO:O	51:U:34:ARG:N	2.52	0.42
42:L:68:GLY:HA3	42:L:98:ARG:NH1	2.34	0.42
50:T:67:HIS:O	50:T:68:LYS:HG2	2.19	0.42
52:03:38:PHE:CD1	54:01:2127:G:H4'	2.53	0.42
53:A:607:A:H2'	53:A:608:A:H8	1.84	0.42
53:A:1325:C:O2'	53:A:1326:U:H5'	2.19	0.42
53:A:1481:U:H2'	53:A:1482:G:C8	2.54	0.42
54:01:151:C:H2'	54:01:152:A:C8	2.54	0.42
54:01:226:A:H2'	54:01:227:A:O4'	2.18	0.42
54:01:947:A:H2'	54:01:948:C:C6	2.54	0.42
54:01:1015:U:H2'	54:01:1016:G:C8	2.55	0.42
54:01:1547:C:H2'	54:01:1548:A:C8	2.55	0.42
54:01:1704:C:H2'	54:01:1705:A:C8	2.54	0.42
54:01:1747:U:H2'	54:01:1748:C:C6	2.54	0.42
54:01:1928:A:H2'	54:01:1929:G:O4'	2.19	0.42
54:01:2126:A:N1	54:01:2163:A:H1'	2.34	0.42
54:01:2322:A:H2'	54:01:2323:G:O4'	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:01:2789:C:H2'	54:01:2893:A:N7	2.35	0.42
59:Z:28:THR:HA	59:Z:31:ILE:HD12	2.00	0.42
59:Z:310:ILE:HG13	59:Z:350:VAL:HG11	2.01	0.42
59:Z:324:LYS:HE3	59:Z:342:GLU:CA	2.49	0.42
1:04:163:ILE:HG23	1:04:171:VAL:HG13	2.02	0.42
2:05:164:GLN:HE22	54:01:2822:G:H5''	1.84	0.42
3:06:33:VAL:HG12	3:06:94:GLN:HE22	1.84	0.42
6:09:5:LEU:CD2	6:09:13:GLY:HA2	2.50	0.42
6:09:78:VAL:HG11	6:09:102:ALA:HB1	2.02	0.42
7:10:58:THR:HA	7:10:63:ALA:HB3	2.02	0.42
8:11:90:GLY:C	8:11:91:LYS:HD3	2.39	0.42
9:12:26:GLY:HA3	54:01:1140:C:H5'	2.00	0.42
14:17:107:ALA:HB1	14:17:111:ARG:HH22	1.84	0.42
18:21:88:ARG:HG3	18:21:88:ARG:HH21	1.83	0.42
27:30:3:GLN:HA	54:01:2615:U:C2	2.54	0.42
32:B:19:THR:OG1	32:B:20:ARG:N	2.52	0.42
38:H:52:GLY:HA3	38:H:56:PRO:HA	2.00	0.42
38:H:87:ARG:O	38:H:91:LEU:HG	2.19	0.42
43:M:72:ILE:O	43:M:76:ILE:HG13	2.20	0.42
52:03:163:TYR:CD2	52:03:171:ILE:HD11	2.54	0.42
52:03:192:LEU:C	52:03:196:LEU:HD13	2.40	0.42
53:A:1098:C:H2'	53:A:1099:G:O4'	2.19	0.42
54:01:1060:U:H5'	54:01:1062:G:C5'	2.49	0.42
54:01:1891:G:H2'	54:01:1892:C:C6	2.54	0.42
54:01:2039:U:H2'	54:01:2040:G:C8	2.55	0.42
54:01:2395:C:H2'	54:01:2396:G:O4'	2.19	0.42
56:X:17:C:H5'	56:X:61:C:OP1	2.19	0.42
58:Y:26:A:H2'	58:Y:27:G:O4'	2.19	0.42
3:06:163:ASN:HD21	54:01:323:C:H5''	1.84	0.42
4:07:30:VAL:HG13	4:07:30:VAL:O	2.20	0.42
8:11:21:PRO:HB2	8:11:22:PRO:HD3	2.00	0.42
25:28:50:VAL:HB	25:28:53:MET:CG	2.49	0.42
30:33:60:CYS:SG	30:33:61:LEU:N	2.92	0.42
32:B:125:PHE:CE1	32:B:127:LYS:HE2	2.55	0.42
34:D:172:VAL:HG22	34:D:174:ALA:N	2.30	0.42
35:E:106:ALA:HB1	35:E:110:MET:HE3	2.01	0.42
37:G:58:LEU:O	37:G:62:GLU:N	2.49	0.42
38:H:116:ARG:HG3	38:H:116:ARG:NH1	2.35	0.42
49:S:13:HIS:HE1	49:S:34:SER:HB2	1.84	0.42
53:A:459:A:H2'	53:A:460:A:C8	2.54	0.42
53:A:664:G:N2	53:A:741:G:H1	2.14	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:A:882:C:O2'	53:A:883:C:H5'	2.19	0.42
54:01:519:U:H2'	54:01:520:G:H8	1.85	0.42
54:01:598:U:H2'	54:01:599:A:C8	2.54	0.42
54:01:851:C:O2'	54:01:852:U:H5'	2.19	0.42
54:01:1092:C:H2'	54:01:1093:G:O4'	2.18	0.42
54:01:2155:U:H2'	54:01:2156:G:O4'	2.20	0.42
54:01:2190:G:H2'	54:01:2191:A:C8	2.54	0.42
54:01:2411:A:H2'	54:01:2412:A:C8	2.53	0.42
54:01:2740:A:H2'	54:01:2741:A:C8	2.55	0.42
56:X:4:G:H2'	56:X:5:G:O4'	2.20	0.42
59:Z:350:VAL:CG1	59:Z:354:ASP:HB2	2.49	0.42
4:07:52:ALA:HA	4:07:149:ARG:NE	2.35	0.42
5:08:126:THR:HG22	5:08:127:GLN:N	2.34	0.42
6:09:33:GLN:HB2	6:09:35:LYS:HG2	2.02	0.42
6:09:84:ALA:CB	6:09:90:LEU:HD12	2.48	0.42
7:10:67:THR:N	7:10:68:PRO:CD	2.83	0.42
8:11:90:GLY:C	8:11:92:PRO:HD3	2.40	0.42
26:29:43:PHE:HB2	26:29:47:LYS:HE3	2.01	0.42
27:30:37:HIS:HB3	27:30:43:THR:HG22	2.02	0.42
30:33:61:LEU:HD13	30:33:64:ALA:HB3	2.02	0.42
32:B:96:LEU:HD13	53:A:1103:C:H4'	2.01	0.42
37:G:11:ILE:HG13	37:G:20:GLU:OE2	2.20	0.42
37:G:71:THR:O	37:G:90:VAL:HG12	2.20	0.42
39:I:89:TYR:HB3	39:I:93:LEU:CD1	2.46	0.42
40:J:66:GLU:HG2	44:N:98:ALA:HB2	2.00	0.42
42:L:98:ARG:NH1	42:L:106:VAL:HG22	2.34	0.42
43:M:48:SER:O	43:M:52:ILE:HG13	2.20	0.42
43:M:94:LEU:C	43:M:108:ARG:HG2	2.40	0.42
50:T:8:LYS:HA	50:T:11:ILE:HG12	2.01	0.42
53:A:634:C:H2'	53:A:635:A:H8	1.85	0.42
53:A:783:C:O2'	53:A:784:A:H5'	2.20	0.42
53:A:1366:C:H2'	53:A:1367:C:C6	2.55	0.42
54:01:1106:G:H2'	54:01:1107:G:O4'	2.19	0.42
54:01:2881:U:H2'	54:01:2882:A:H8	1.84	0.42
59:Z:7:ARG:NH2	59:Z:269:ARG:HG2	2.33	0.42
59:Z:321:PRO:HB3	59:Z:351:MET:HA	2.02	0.42
7:10:94:ARG:O	7:10:98:GLU:HG2	2.19	0.42
8:11:56:VAL:HG22	8:11:57:VAL:N	2.35	0.42
9:12:47:HIS:ND1	9:12:48:VAL:HG23	2.34	0.42
9:12:136:GLN:HE21	54:01:2899:A:C5'	2.33	0.42
17:20:20:VAL:O	17:20:96:VAL:HG22	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:21:24:ILE:HG13	18:21:24:ILE:O	2.20	0.42
34:D:96:ARG:HG2	34:D:96:ARG:HH11	1.84	0.42
36:F:8:PHE:CZ	36:F:60:VAL:HG11	2.54	0.42
36:F:97:THR:O	36:F:98:GLU:HB2	2.20	0.42
37:G:14:ASP:HB3	37:G:17:PHE:O	2.20	0.42
50:T:20:ASN:OD1	50:T:65:LEU:HD13	2.20	0.42
53:A:22:G:H5'	53:A:561:U:O2	2.19	0.42
53:A:188:C:H2'	53:A:189:A:O4'	2.20	0.42
53:A:518:C:H4'	53:A:519:C:H5''	2.01	0.42
53:A:621:A:H2'	53:A:622:A:C8	2.55	0.42
53:A:1516:G:H2'	53:A:1518:A:OP2	2.19	0.42
54:01:128:C:H2'	54:01:129:C:C6	2.55	0.42
54:01:130:C:H2'	54:01:131:A:O4'	2.19	0.42
54:01:275:C:H2'	54:01:276:U:C4'	2.47	0.42
54:01:519:U:H2'	54:01:520:G:C8	2.55	0.42
54:01:567:U:H2'	54:01:568:U:O4'	2.19	0.42
54:01:853:C:H2'	54:01:854:C:C6	2.54	0.42
54:01:1070:A:O2'	54:01:1071:G:P	2.78	0.42
54:01:1077:A:C2'	54:01:1078:U:H5'	2.47	0.42
54:01:1144:A:H2'	54:01:1145:C:C6	2.55	0.42
54:01:1525:A:H2'	54:01:1526:C:O4'	2.18	0.42
54:01:2632:A:H2'	54:01:2633:G:C8	2.53	0.42
55:02:106:G:H2'	55:02:107:G:O4'	2.20	0.42
59:Z:144:GLU:HG2	59:Z:147:GLU:OE1	2.20	0.42
59:Z:327:ARG:CD	59:Z:340:THR:HA	2.50	0.42
9:12:27:ARG:HD2	9:12:27:ARG:N	2.34	0.42
18:21:14:ALA:HB2	18:21:101:SER:HB3	2.01	0.42
19:22:61:LEU:HD11	19:22:82:LYS:HD3	2.02	0.42
42:L:58:ASN:OD1	42:L:60:PHE:HD2	2.03	0.42
43:M:76:ILE:O	43:M:80:MET:HG3	2.20	0.42
48:R:11:ARG:NH1	48:R:15:GLU:HB2	2.35	0.42
54:01:796:C:H2'	54:01:797:G:H8	1.83	0.42
54:01:864:G:O2'	54:01:865:C:H5'	2.19	0.42
54:01:1023:U:O2'	54:01:1122:G:H5'	2.19	0.42
54:01:1483:G:H4'	54:01:1510:G:H21	1.85	0.42
1:04:244:VAL:HG12	1:04:250:GLN:HA	2.02	0.42
14:17:26:LEU:HD23	14:17:92:PHE:HD1	1.85	0.42
16:19:49:ARG:HH11	16:19:49:ARG:CB	2.32	0.42
17:20:14:VAL:HG21	17:20:98:ILE:CG1	2.50	0.42
17:20:88:GLY:H	54:01:1225:G:H5'	1.85	0.42
28:31:5:ARG:HG2	28:31:23:THR:HB	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:B:104:LYS:NZ	53:A:1073:U:H4'	2.35	0.42
32:B:163:ILE:O	32:B:185:ILE:HG23	2.19	0.42
32:B:202:ASN:ND2	32:B:204:ASP:OD1	2.53	0.42
35:E:10:LEU:HD12	35:E:11:GLN:N	2.35	0.42
35:E:15:ILE:HD11	35:E:37:VAL:HG23	2.02	0.42
38:H:31:LEU:O	38:H:35:ILE:HG13	2.20	0.42
50:T:79:THR:HG22	50:T:83:ASN:HD22	1.83	0.42
53:A:184:G:H2'	53:A:185:U:C6	2.54	0.42
53:A:1208:C:H2'	53:A:1209:C:C6	2.55	0.42
53:A:1230:C:H5'	56:W:30:G:H5''	2.02	0.42
53:A:1417:G:N2	53:A:1482:G:H2'	2.35	0.42
53:A:1470:U:O2'	53:A:1471:U:H5'	2.19	0.42
53:A:1526:G:H2'	53:A:1527:U:C6	2.55	0.42
54:01:155:A:H2'	54:01:156:A:H8	1.84	0.42
54:01:302:C:H2'	54:01:303:G:C8	2.55	0.42
54:01:333:G:H2'	54:01:333:G:N3	2.35	0.42
54:01:386:G:H3'	54:01:387:U:C5'	2.49	0.42
54:01:445:C:O2'	54:01:446:G:H5'	2.20	0.42
54:01:863:A:H4'	55:02:100:G:N2	2.34	0.42
54:01:1471:G:H2'	54:01:1472:C:C6	2.55	0.42
54:01:2215:C:H2'	54:01:2216:G:H8	1.85	0.42
54:01:2464:G:H2'	54:01:2465:C:C6	2.54	0.42
59:Z:13:ASN:O	59:Z:98:MET:HA	2.20	0.42
59:Z:44:ARG:HD2	59:Z:48:GLN:OE1	2.20	0.42
59:Z:69:TYR:CE1	59:Z:78:HIS:HB2	2.54	0.42
59:Z:170:VAL:CG1	59:Z:194:PHE:HE2	2.32	0.42
59:Z:246:GLY:H	59:Z:250:THR:CG2	2.33	0.42
4:07:58:ALA:HB1	4:07:139:GLU:HB3	2.02	0.42
5:08:87:GLN:NE2	5:08:162:ARG:HD2	2.29	0.42
8:11:40:ALA:HA	8:11:43:ALA:HB3	2.02	0.42
9:12:73:VAL:CG1	9:12:86:GLN:HB2	2.50	0.42
11:14:110:VAL:HG21	11:14:127:VAL:HG22	2.02	0.42
14:17:105:ALA:HA	14:17:108:ASP:OD2	2.19	0.42
19:22:50:LEU:HD12	19:22:50:LEU:N	2.35	0.42
20:23:42:LYS:CE	54:01:499:U:H5''	2.44	0.42
28:31:37:LYS:HB2	28:31:48:TYR:CE2	2.55	0.42
40:J:15:HIS:HB3	40:J:70:HIS:CD2	2.55	0.42
44:N:2:LYS:HD2	53:A:1049:U:O2'	2.19	0.42
46:P:79:ASN:HB2	46:P:82:ALA:HB3	2.01	0.42
53:A:363:A:H2'	53:A:364:A:O4'	2.19	0.42
54:01:326:G:H2'	54:01:327:G:C8	2.55	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:01:492:A:H2'	54:01:493:G:O4'	2.20	0.42
54:01:828:U:H4'	54:01:831:G:N1	2.35	0.42
54:01:1123:C:H2'	54:01:1124:G:C8	2.55	0.42
54:01:1923:U:H2'	54:01:1924:C:H6	1.85	0.42
54:01:2277:G:C3'	54:01:2278:A:H5''	2.49	0.42
54:01:2790:U:H5''	54:01:2791:G:OP1	2.20	0.42
1:04:252:LYS:HD3	54:01:1901:A:H4'	2.02	0.41
4:07:35:LEU:HD21	4:07:98:PHE:CZ	2.55	0.41
4:07:151:LEU:HA	54:01:2305:U:C4	2.55	0.41
7:10:18:VAL:HA	7:10:86:MET:CE	2.50	0.41
8:11:25:PRO:HB2	58:Y:56:C:OP2	2.20	0.41
9:12:113:PRO:HB2	54:01:557:C:H5''	2.02	0.41
13:16:2:ARG:CB	13:16:5:LYS:HD2	2.49	0.41
23:26:15:ASN:HD22	54:01:381:G:H5''	1.85	0.41
24:27:42:LEU:O	24:27:46:VAL:HG23	2.20	0.41
31:34:36:ARG:HD3	54:01:2742:G:OP1	2.19	0.41
32:B:51:GLU:O	32:B:54:ALA:HB3	2.20	0.41
38:H:84:ILE:CG2	38:H:86:LYS:HE3	2.50	0.41
49:S:35:ARG:HB3	49:S:35:ARG:HH11	1.85	0.41
53:A:769:G:O2'	53:A:770:C:H5'	2.20	0.41
53:A:1173:U:H2'	53:A:1174:G:H8	1.85	0.41
53:A:1256:A:H1'	53:A:1258:G:N9	2.34	0.41
54:01:225:C:H2'	54:01:226:A:O4'	2.20	0.41
54:01:540:C:H2'	54:01:541:A:C8	2.55	0.41
54:01:948:C:H2'	54:01:949:G:H8	1.84	0.41
54:01:1314:C:H42	54:01:1338:G:H1	1.68	0.41
54:01:1701:A:H2'	54:01:1702:G:H5'	2.02	0.41
54:01:2038:G:H2'	54:01:2039:U:O4'	2.19	0.41
54:01:2215:C:H2'	54:01:2216:G:C8	2.54	0.41
54:01:2791:G:H2'	54:01:2792:A:O4'	2.20	0.41
59:Z:305:GLU:N	59:Z:392:LEU:HD23	2.35	0.41
1:04:257:ARG:HH21	1:04:266:ILE:CD1	2.23	0.41
4:07:41:GLU:O	4:07:48:LEU:HD23	2.20	0.41
8:11:83:ALA:HA	8:11:100:ILE:HD11	2.02	0.41
11:14:69:ARG:HH11	11:14:69:ARG:HG3	1.85	0.41
22:25:17:LEU:HD11	22:25:37:ARG:NH2	2.35	0.41
26:29:46:GLY:HA2	26:29:49:ARG:NH2	2.35	0.41
28:31:16:THR:HG21	28:31:39:ASP:OD2	2.21	0.41
36:F:12:PRO:HD2	36:F:54:LEU:CD2	2.32	0.41
39:I:46:VAL:HG21	39:I:75:ALA:HB1	2.01	0.41
39:I:54:VAL:HG23	39:I:54:VAL:O	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:Q:16:MET:HG3	47:Q:19:SER:OG	2.20	0.41
53:A:179:A:H2'	53:A:180:U:O4'	2.20	0.41
53:A:690:G:H2'	53:A:691:G:O4'	2.20	0.41
54:01:967:U:H2'	54:01:968:C:C6	2.55	0.41
54:01:1182:G:H2'	54:01:1183:U:O4'	2.20	0.41
54:01:1368:G:H2'	54:01:1369:G:H8	1.85	0.41
54:01:2150:C:H2'	54:01:2151:U:O4'	2.20	0.41
54:01:2197:U:H1'	54:01:2198:A:C8	2.55	0.41
55:02:60:C:H2'	55:02:61:G:H8	1.85	0.41
59:Z:198:TYR:O	59:Z:200:PRO:HD3	2.20	0.41
2:05:98:VAL:HG22	2:05:180:VAL:HG13	2.03	0.41
2:05:129:THR:HG23	2:05:140:HIS:O	2.20	0.41
6:09:57:LYS:O	6:09:61:VAL:HG22	2.20	0.41
8:11:55:PRO:CG	8:11:71:LYS:HE2	2.50	0.41
15:18:29:VAL:CG1	15:18:79:VAL:HG22	2.51	0.41
21:24:80:HIS:ND1	21:24:81:PRO:CD	2.81	0.41
35:E:155:LYS:HB2	35:E:155:LYS:NZ	2.35	0.41
40:J:53:ILE:HG13	44:N:84:ARG:NE	2.35	0.41
52:03:211:LYS:HG2	54:01:2177:C:H4'	2.02	0.41
53:A:46:G:OP1	53:A:307:C:H4'	2.20	0.41
53:A:756:C:H2'	53:A:757:U:O4'	2.21	0.41
54:01:214:G:H2'	54:01:215:G:C8	2.55	0.41
54:01:705:A:C2	54:01:727:A:H1'	2.56	0.41
54:01:736:C:H2'	54:01:737:C:H6	1.86	0.41
54:01:2266:A:H4'	54:01:2267:A:N3	2.35	0.41
54:01:2347:C:H5	54:01:2382:G:H1'	1.83	0.41
56:X:38:A:H2'	56:X:39:C:O4'	2.20	0.41
10:13:70:ARG:HG2	10:13:70:ARG:HH11	1.85	0.41
26:29:45:THR:O	26:29:49:ARG:HG2	2.20	0.41
32:B:67:LEU:HD12	32:B:153:MET:HE1	2.01	0.41
33:C:173:PRO:HB2	33:C:176:THR:OG1	2.20	0.41
34:D:7:LYS:HE2	53:A:408:A:OP2	2.21	0.41
36:F:36:ILE:HD11	36:F:62:MET:HB3	2.02	0.41
43:M:26:LYS:O	43:M:29:SER:HB3	2.20	0.41
47:Q:7:LEU:HD23	47:Q:24:ILE:HD13	2.01	0.41
47:Q:17:GLU:OE2	53:A:255:G:H1'	2.20	0.41
49:S:57:VAL:HA	49:S:58:PRO:HD3	1.91	0.41
52:03:53:ARG:CB	56:X:62:C:H4'	2.50	0.41
53:A:628:G:H2'	53:A:629:A:C8	2.54	0.41
53:A:985:C:H2'	53:A:986:U:C6	2.55	0.41
53:A:1052:U:H2'	53:A:1200:C:N4	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:A:1405:G:O2'	53:A:1406:U:H5'	2.20	0.41
54:01:168:G:H2'	54:01:169:G:C8	2.56	0.41
54:01:357:C:H2'	54:01:358:U:C6	2.55	0.41
54:01:634:C:H2'	54:01:635:C:H6	1.85	0.41
54:01:1106:G:H5'	54:01:1106:G:H8	1.85	0.41
54:01:1336:A:H2'	54:01:1337:G:H8	1.86	0.41
54:01:2590:A:H2'	54:01:2591:C:H6	1.84	0.41
56:W:47:U:H3'	56:W:48:C:C5'	2.50	0.41
59:Z:133:PHE:CD1	59:Z:170:VAL:HG23	2.53	0.41
5:08:93:TYR:CD1	5:08:106:LEU:HA	2.56	0.41
7:10:48:ALA:HB3	7:10:51:TYR:OH	2.20	0.41
8:11:7:TYR:HB3	8:11:59:THR:HG23	2.03	0.41
17:20:14:VAL:HG11	17:20:98:ILE:HG13	2.02	0.41
19:22:7:LEU:HD22	19:22:46:ALA:CA	2.50	0.41
20:23:48:VAL:C	20:23:50:ALA:H	2.24	0.41
25:28:46:MET:O	25:28:49:ALA:HB3	2.21	0.41
26:29:58:ASP:OD1	26:29:58:ASP:N	2.53	0.41
32:B:16:GLY:HA3	32:B:39:ILE:HA	2.02	0.41
33:C:146:LYS:N	33:C:146:LYS:HD2	2.36	0.41
33:C:147:GLY:HA3	33:C:171:ARG:O	2.20	0.41
37:G:42:VAL:O	37:G:46:LEU:HD13	2.21	0.41
39:I:30:ASN:O	39:I:31:GLN:HB2	2.20	0.41
51:U:11:PHE:O	51:U:13:VAL:N	2.54	0.41
52:03:31:LYS:HA	52:03:34:ALA:CB	2.41	0.41
53:A:170:U:O2'	53:A:171:A:H5'	2.21	0.41
54:01:570:G:O2'	54:01:571:U:H5'	2.20	0.41
54:01:1282:U:H2'	54:01:1283:G:O4'	2.21	0.41
54:01:1858:A:H1'	54:01:1885:A:C2	2.55	0.41
54:01:2448:A:H3'	54:01:2449:U:H2'	2.01	0.41
54:01:2713:U:H3'	54:01:2714:G:H5''	2.03	0.41
54:01:2888:C:H2'	54:01:2889:C:C6	2.55	0.41
55:02:100:G:H2'	55:02:101:A:O4'	2.20	0.41
56:X:27:U:H2'	56:X:28:C:C6	2.56	0.41
57:V:8:A:H2'	57:V:9:G:C8	2.56	0.41
58:Y:4:C:H2'	58:Y:5:G:H8	1.85	0.41
59:Z:22:HIS:HB3	59:Z:104:VAL:HG12	2.02	0.41
59:Z:115:THR:O	59:Z:119:ILE:HG13	2.19	0.41
59:Z:121:LEU:O	59:Z:125:VAL:N	2.54	0.41
59:Z:145:LEU:O	59:Z:149:VAL:HG23	2.21	0.41
59:Z:227:VAL:CG1	59:Z:276:VAL:HB	2.50	0.41
2:05:207:VAL:HG21	54:01:2771:C:H5''	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:07:70:ARG:HG2	4:07:70:ARG:NH2	2.36	0.41
6:09:111:ALA:HB3	6:09:114:GLU:OE1	2.21	0.41
7:10:118:ILE:HD12	7:10:118:ILE:HA	1.86	0.41
8:11:52:LEU:HA	8:11:53:PRO:HD3	1.90	0.41
9:12:2:LYS:HB3	54:01:995:C:H42	1.85	0.41
11:14:93:ASN:C	11:14:95:LEU:H	2.24	0.41
33:C:196:GLY:HA3	53:A:1057:G:O3'	2.20	0.41
34:D:165:GLU:O	34:D:166:LYS:HB2	2.21	0.41
40:J:32:THR:OG1	40:J:82:LYS:HB3	2.21	0.41
40:J:53:ILE:HG12	53:A:1060:U:H5'	2.02	0.41
53:A:778:G:H2'	53:A:779:C:O4'	2.20	0.41
53:A:1436:U:H2'	53:A:1437:A:C8	2.54	0.41
54:01:287:G:H2'	54:01:288:U:C6	2.56	0.41
54:01:922:C:H2'	54:01:923:G:H8	1.86	0.41
54:01:962:G:O2'	54:01:963:U:H5'	2.20	0.41
54:01:1039:A:H2	54:01:1116:G:H22	1.68	0.41
54:01:1283:G:H1'	54:01:1329:U:O2	2.21	0.41
54:01:1842:G:H2'	54:01:1843:C:C6	2.56	0.41
54:01:2163:A:N3	54:01:2164:C:H5'	2.36	0.41
2:05:177:VAL:HG22	2:05:178:VAL:N	2.36	0.41
4:07:3:LEU:HA	4:07:6:TYR:HB3	2.03	0.41
4:07:94:ARG:HG2	26:29:9:TYR:CE2	2.56	0.41
12:15:5:LYS:HD2	12:15:6:ARG:CG	2.50	0.41
17:20:27:ILE:HG13	17:20:33:VAL:HG11	2.02	0.41
20:23:9:GLU:OE2	20:23:21:ARG:HB3	2.20	0.41
33:C:112:ALA:HB2	33:C:182:ASP:HB3	2.03	0.41
37:G:145:GLU:HA	37:G:148:LYS:HB2	2.02	0.41
38:H:10:LEU:HD22	38:H:74:ILE:HD11	2.02	0.41
40:J:81:GLU:O	40:J:84:VAL:HG12	2.21	0.41
47:Q:46:HIS:HA	47:Q:70:LYS:HE3	2.03	0.41
52:03:164:ARG:HD2	52:03:165:ASN:O	2.20	0.41
53:A:39:G:O2'	53:A:40:C:H5'	2.20	0.41
54:01:566:U:H2'	54:01:567:U:O4'	2.21	0.41
54:01:633:A:H2'	54:01:634:C:H5'	2.03	0.41
54:01:799:G:H5''	54:01:800:A:H2'	2.01	0.41
54:01:2143:C:H2'	54:01:2144:G:C8	2.56	0.41
54:01:2286:G:H4'	54:01:2287:A:O5'	2.21	0.41
58:Y:65:G:H2'	58:Y:66:U:C6	2.55	0.41
59:Z:112:MET:HB2	59:Z:115:THR:OG1	2.21	0.41
59:Z:328:PRO:CD	59:Z:341:ILE:HD11	2.51	0.41
1:04:143:VAL:HB	1:04:153:LEU:HB2	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:04:154:ALA:HB2	1:04:161:VAL:HG23	2.01	0.41
7:10:18:VAL:HA	7:10:86:MET:HE3	2.03	0.41
20:23:6:ARG:HB2	54:01:85:G:P	2.60	0.41
20:23:82:VAL:HG12	20:23:83:GLY:N	2.35	0.41
33:C:150:VAL:HG12	33:C:199:VAL:HG23	2.03	0.41
35:E:86:GLY:N	35:E:93:VAL:O	2.53	0.41
36:F:64:VAL:CG2	36:F:65:GLU:H	2.30	0.41
39:I:61:ASP:C	39:I:62:LEU:HD12	2.41	0.41
41:K:70:ALA:O	41:K:74:LYS:HG3	2.20	0.41
41:K:115:ILE:HD12	51:U:23:GLU:OE1	2.20	0.41
42:L:29:LYS:CG	42:L:56:LEU:HD22	2.51	0.41
43:M:100:ARG:HH11	43:M:100:ARG:HG3	1.86	0.41
44:N:41:TRP:CZ2	49:S:10:ILE:HD11	2.56	0.41
47:Q:13:SER:H	47:Q:21:VAL:CG1	2.34	0.41
53:A:246:A:N3	53:A:247:G:H1'	2.35	0.41
53:A:272:C:H2'	53:A:273:U:C6	2.56	0.41
54:01:8:C:H2'	54:01:9:G:O4'	2.21	0.41
54:01:438:G:H2'	54:01:439:A:C8	2.56	0.41
54:01:783:A:H2'	54:01:784:G:H5'	2.03	0.41
54:01:2128:G:H2'	54:01:2129:C:O4'	2.20	0.41
54:01:2249:U:H3'	54:01:2250:G:C5'	2.50	0.41
54:01:2267:A:C5'	54:01:2268:A:H5'	2.48	0.41
54:01:2389:G:H5''	54:01:2390:U:O4'	2.21	0.41
59:Z:227:VAL:HG11	59:Z:292:LEU:HD11	2.02	0.41
3:06:88:ARG:O	3:06:90:GLN:HG2	2.20	0.41
3:06:178:VAL:HG13	3:06:179:SER:N	2.36	0.41
4:07:24:VAL:HG12	55:02:55:U:H4'	2.03	0.41
6:09:94:ILE:N	6:09:94:ILE:HD12	2.36	0.41
7:10:91:ALA:O	7:10:92:ALA:HB3	2.21	0.41
8:11:9:LYS:HG2	8:11:57:VAL:HG22	2.03	0.41
8:11:12:VAL:O	8:11:13:ALA:C	2.59	0.41
12:15:86:LYS:HG3	54:01:956:G:OP1	2.21	0.41
13:16:71:ARG:CZ	54:01:2708:G:H1'	2.51	0.41
14:17:30:ARG:HA	14:17:35:ILE:HG13	2.02	0.41
19:22:54:GLU:HB3	19:22:88:LYS:HE3	2.02	0.41
20:23:45:GLN:HE22	20:23:54:PRO:HD2	1.86	0.41
29:32:6:GLN:HA	29:32:7:PRO:HD2	1.93	0.41
29:32:34:ARG:NH2	29:32:39:ARG:HD2	2.31	0.41
32:B:96:LEU:H	32:B:99:MET:HE2	1.85	0.41
32:B:191:ASP:OD1	32:B:193:ASP:HB2	2.21	0.41
32:B:206:ILE:HA	32:B:209:VAL:HG22	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:C:5:HIS:CG	44:N:88:MET:HB3	2.56	0.41
38:H:94:VAL:HG21	38:H:100:ILE:O	2.21	0.41
40:J:15:HIS:O	40:J:18:ILE:HG22	2.21	0.41
41:K:127:ARG:HH22	53:A:1522:U:H5''	1.86	0.41
43:M:56:ARG:HA	43:M:59:VAL:HG22	2.02	0.41
45:O:52:ARG:HH11	45:O:52:ARG:HG2	1.85	0.41
47:Q:48:GLU:HB2	47:Q:51:GLU:CD	2.42	0.41
47:Q:65:PRO:O	53:A:264:C:O2'	2.38	0.41
48:R:41:SER:HB2	48:R:51:GLN:HE21	1.86	0.41
52:O3:33:LEU:HD22	52:O3:220:ALA:CB	2.50	0.41
52:O3:41:SER:O	52:O3:217:THR:HG23	2.21	0.41
53:A:163:C:H2'	53:A:164:G:O4'	2.21	0.41
53:A:599:C:H2'	53:A:600:A:C8	2.56	0.41
53:A:666:G:H2'	53:A:667:G:C8	2.52	0.41
53:A:704:A:H2'	53:A:705:G:O4'	2.21	0.41
53:A:1030:U:H2'	53:A:1031:C:H5'	2.03	0.41
53:A:1128:C:O2'	53:A:1129:C:H5'	2.21	0.41
53:A:1137:C:H5'	53:A:1138:G:C5'	2.45	0.41
53:A:1368:A:O2'	53:A:1369:C:H5'	2.20	0.41
53:A:1509:C:O2'	53:A:1510:C:H5'	2.20	0.41
54:O1:195:A:H2'	54:O1:198:C:N4	2.36	0.41
54:O1:974:G:H1'	54:O1:975:A:C8	2.56	0.41
54:O1:1363:C:H2'	54:O1:1364:G:C8	2.56	0.41
54:O1:1475:G:O2'	54:O1:1476:U:P	2.78	0.41
54:O1:2066:C:O2'	54:O1:2067:G:H5'	2.21	0.41
54:O1:2181:U:H2'	54:O1:2182:U:O4'	2.21	0.41
54:O1:2220:U:H2'	54:O1:2221:G:C8	2.56	0.41
54:O1:2692:G:O2'	54:O1:2693:G:H5'	2.21	0.41
54:O1:2876:G:H2'	54:O1:2877:G:O4'	2.20	0.41
56:X:71:C:H2'	56:X:72:A:C8	2.56	0.41
59:Z:129:TYR:HB3	59:Z:199:ILE:CG1	2.51	0.41
59:Z:134:LEU:HB3	59:Z:146:LEU:HD12	2.03	0.41
59:Z:137:CYS:HB3	59:Z:184:TRP:CH2	2.56	0.41
1:O4:96:LYS:HD2	54:O1:1490:A:H62	1.86	0.41
1:O4:231:HIS:HA	1:O4:241:LYS:HD2	2.03	0.41
3:O6:47:LYS:O	3:O6:83:VAL:HB	2.20	0.41
4:O7:82:TYR:CD1	4:O7:83:PRO:HD2	2.56	0.41
7:10:33:VAL:HA	54:O1:1055:G:H4'	2.02	0.41
10:13:93:GLN:HA	10:13:94:PRO:HD2	1.93	0.41
11:14:42:SER:HB2	54:O1:672:C:H5	1.86	0.41
14:17:33:ARG:HG2	14:17:34:HIS:CD2	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:19:5:ARG:HH21	16:19:5:ARG:CG	2.34	0.41
20:23:97:SER:O	20:23:98:ASN:CB	2.68	0.41
32:B:34:ARG:HG2	32:B:34:ARG:HH11	1.85	0.41
32:B:76:SER:HA	32:B:79:VAL:CG2	2.51	0.41
35:E:28:ARG:NH1	53:A:15:G:H4'	2.36	0.41
36:F:38:ARG:NH1	36:F:98:GLU:H	2.19	0.41
51:U:36:PHE:C	51:U:38:GLU:N	2.74	0.41
52:03:38:PHE:HB2	52:03:39:VAL:H	1.75	0.41
52:03:67:HIS:CD2	52:03:188:ASN:HD21	2.39	0.41
52:03:170:ILE:N	52:03:170:ILE:HD12	2.36	0.41
52:03:207:VAL:O	52:03:210:LYS:HG3	2.21	0.41
53:A:483:C:H2'	53:A:484:G:C8	2.56	0.41
53:A:608:A:H2'	53:A:609:A:O4'	2.21	0.41
53:A:1324:A:H2'	53:A:1325:C:O4'	2.21	0.41
53:A:1424:U:H2'	53:A:1425:U:O4'	2.21	0.41
54:01:1499:C:H2'	54:01:1500:G:C8	2.55	0.41
54:01:1853:A:H2'	54:01:1854:A:C8	2.56	0.41
54:01:1917:U:O2'	54:01:1918:A:H5'	2.21	0.41
55:02:49:C:H2'	55:02:50:A:C8	2.56	0.41
59:Z:103:LEU:HB3	59:Z:132:VAL:HG22	2.03	0.41
59:Z:225:THR:O	59:Z:225:THR:HG22	2.19	0.41
3:06:21:ARG:HH21	3:06:106:LYS:HE2	1.86	0.40
6:09:99:ILE:HD12	6:09:117:LEU:HD21	2.02	0.40
7:10:98:GLU:O	7:10:102:ALA:N	2.53	0.40
13:16:68:ALA:HA	54:01:2707:U:O2'	2.20	0.40
15:18:102:ARG:NH1	15:18:102:ARG:HG3	2.36	0.40
22:25:14:ALA:HB1	54:01:2271:G:OP1	2.21	0.40
33:C:123:LEU:HD13	33:C:195:ILE:HG21	2.02	0.40
36:F:42:TRP:HE3	36:F:45:ARG:NH2	2.19	0.40
36:F:88:MET:HB3	48:R:63:TYR:HE2	1.85	0.40
38:H:80:PRO:HG2	53:A:878:A:H5'	2.02	0.40
43:M:84:CYS:HB2	49:S:72:GLU:OE1	2.21	0.40
44:N:20:PHE:C	44:N:22:LYS:H	2.24	0.40
46:P:51:ARG:C	46:P:52:LEU:HD12	2.41	0.40
48:R:25:ILE:HA	48:R:28:LEU:HB3	2.03	0.40
51:U:34:ARG:HD3	51:U:34:ARG:HA	1.94	0.40
52:03:27:ILE:HA	52:03:30:LEU:HG	2.03	0.40
53:A:321:A:O2'	53:A:322:C:H5'	2.21	0.40
53:A:641:U:H4'	53:A:642:A:H8	1.85	0.40
53:A:744:C:H2'	53:A:745:G:H8	1.86	0.40
53:A:1026:G:H2'	53:A:1027:C:C6	2.55	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:A:1413:A:H2	53:A:1487:G:H22	1.69	0.40
54:01:186:G:H2'	54:01:187:G:H8	1.85	0.40
54:01:548:G:H2'	54:01:549:G:O4'	2.21	0.40
54:01:553:G:H2'	54:01:554:U:O4'	2.20	0.40
54:01:634:C:H6	54:01:634:C:O5'	2.04	0.40
54:01:772:C:O2'	54:01:773:U:H5'	2.20	0.40
54:01:808:G:H2'	54:01:809:G:C8	2.56	0.40
54:01:1554:U:O2'	54:01:1555:G:H5''	2.21	0.40
54:01:2214:C:H2'	54:01:2215:C:H5'	2.03	0.40
55:02:60:C:H2'	55:02:61:G:C8	2.56	0.40
55:02:65:U:H3'	55:02:108:A:H61	1.86	0.40
55:02:119:A:H2'	55:02:120:A:O4'	2.22	0.40
59:Z:18:GLY:O	59:Z:118:HIS:CD2	2.74	0.40
59:Z:214:ILE:HD12	59:Z:290:GLN:CB	2.37	0.40
59:Z:304:PHE:O	59:Z:359:VAL:HG13	2.21	0.40
7:10:93:ALA:O	7:10:129:LEU:HD13	2.20	0.40
9:12:136:GLN:HE21	54:01:2899:A:H5''	1.85	0.40
11:14:110:VAL:CG2	11:14:127:VAL:HG22	2.51	0.40
13:16:12:ARG:HD3	13:16:16:HIS:ND1	2.36	0.40
15:18:26:GLU:HA	15:18:43:GLU:HA	2.03	0.40
18:21:81:SER:O	18:21:83:LYS:HD2	2.21	0.40
23:26:2:ARG:HD2	23:26:29:LEU:HD22	2.02	0.40
23:26:6:VAL:HG23	23:26:50:VAL:HG12	2.03	0.40
27:30:13:GLY:HA3	54:01:16:C:H5''	2.04	0.40
34:D:131:ILE:HG23	53:A:403:C:H5'	2.03	0.40
34:D:183:ARG:HG2	34:D:183:ARG:HH11	1.85	0.40
37:G:12:LEU:CD1	37:G:13:PRO:HD2	2.44	0.40
43:M:7:ASN:HD22	43:M:9:PRO:HD3	1.85	0.40
49:S:77:ARG:HH11	53:A:1222:G:H5''	1.85	0.40
52:03:60:ARG:HG3	52:03:164:ARG:HB2	2.03	0.40
53:A:349:A:H2'	53:A:350:G:O4'	2.22	0.40
53:A:476:U:H2'	53:A:477:C:C6	2.57	0.40
53:A:986:U:H2'	53:A:987:G:O4'	2.20	0.40
54:01:1148:U:H2'	54:01:1149:G:O4'	2.22	0.40
54:01:1179:G:C4	54:01:1180:U:H1'	2.56	0.40
54:01:1300:G:H4'	54:01:1301:A:H5''	2.04	0.40
54:01:1336:A:H2'	54:01:1337:G:C8	2.56	0.40
54:01:1637:A:H5'	54:01:1760:C:O2'	2.21	0.40
59:Z:206:ILE:HG21	59:Z:269:ARG:HB2	2.03	0.40
59:Z:225:THR:HG22	59:Z:278:LEU:HB2	2.02	0.40
59:Z:259:GLU:HG3	59:Z:263:LYS:O	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:Z:303:LYS:CG	59:Z:361:THR:HG22	2.33	0.40
7:10:51:TYR:OH	7:10:89:PRO:HG2	2.22	0.40
8:11:24:GLY:HA3	8:11:25:PRO:HD3	1.93	0.40
11:14:109:LYS:HE2	54:01:636:G:N7	2.35	0.40
12:15:73:ILE:HD13	12:15:73:ILE:HA	1.94	0.40
15:18:105:LYS:HG3	53:A:1432:G:H5''	2.02	0.40
19:22:15:HIS:HE1	19:22:17:SER:HB3	1.86	0.40
35:E:114:LEU:O	35:E:119:VAL:HG22	2.21	0.40
37:G:3:ARG:HH11	37:G:4:ARG:NH1	2.18	0.40
37:G:55:LYS:HB2	37:G:60:ALA:HB2	2.01	0.40
39:I:121:ARG:HH11	53:A:1345:U:H5''	1.86	0.40
48:R:10:CYS:O	48:R:11:ARG:HB3	2.22	0.40
52:03:33:LEU:HD22	52:03:220:ALA:HB3	2.03	0.40
52:03:166:ASP:OD2	52:03:170:ILE:HB	2.21	0.40
53:A:5:U:H4'	53:A:6:G:C5	2.57	0.40
53:A:746:A:H2'	53:A:747:A:C8	2.57	0.40
53:A:1090:U:O2'	53:A:1091:U:H5'	2.22	0.40
54:01:542:C:H2'	54:01:543:G:H5'	2.02	0.40
54:01:839:U:H2'	54:01:840:C:H6	1.80	0.40
54:01:1060:U:C5'	54:01:1062:G:H5'	2.51	0.40
54:01:1190:G:H2'	54:01:1191:G:H8	1.86	0.40
54:01:1360:G:H2'	54:01:1361:G:H5'	2.04	0.40
54:01:1548:A:H2'	54:01:1549:A:H8	1.86	0.40
54:01:1981:A:H5''	54:01:1982:U:OP2	2.21	0.40
54:01:2290:G:H2'	54:01:2291:U:O4'	2.20	0.40
55:02:80:U:H2'	55:02:81:G:H8	1.85	0.40
56:X:21:A:N6	56:X:47:U:H5''	2.36	0.40
59:Z:148:LEU:HD12	59:Z:149:VAL:N	2.36	0.40
3:06:29:HIS:HA	3:06:32:VAL:HG12	2.03	0.40
4:07:94:ARG:HH22	26:29:1:MET:HB3	1.86	0.40
8:11:55:PRO:HG2	8:11:71:LYS:HE2	2.03	0.40
12:15:41:LEU:HB2	12:15:46:ILE:HD11	2.04	0.40
14:17:49:VAL:HG21	14:17:82:ALA:HA	2.03	0.40
15:18:1:SER:N	54:01:2875:C:O2'	2.52	0.40
17:20:93:PHE:HE2	17:20:95:ASP:OD2	2.05	0.40
20:23:45:GLN:NE2	20:23:55:GLY:H	2.18	0.40
21:24:56:PHE:CE1	21:24:61:LEU:HD21	2.56	0.40
27:30:9:ARG:HB2	54:01:17:G:OP1	2.22	0.40
32:B:89:PHE:CE1	32:B:152:ASP:HB2	2.52	0.40
33:C:68:HIS:HA	33:C:103:ALA:O	2.21	0.40
33:C:149:LYS:HG2	33:C:200:TRP:HB2	2.04	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:K:87:GLY:N	41:K:113:THR:HG22	2.35	0.40
46:P:6:LEU:HD13	46:P:17:TYR:CG	2.56	0.40
46:P:20:VAL:CG2	46:P:21:VAL:N	2.85	0.40
47:Q:65:PRO:HG3	53:A:234:C:O2'	2.21	0.40
50:T:8:LYS:O	50:T:12:GLN:N	2.50	0.40
52:O3:33:LEU:HD13	52:O3:220:ALA:HB3	2.03	0.40
53:A:460:A:H2'	53:A:461:A:C8	2.56	0.40
53:A:505:G:H2'	53:A:506:G:C8	2.57	0.40
53:A:751:U:H2'	53:A:752:G:O4'	2.21	0.40
54:O1:20:C:H2'	54:O1:21:A:C8	2.57	0.40
54:O1:878:A:H5'	54:O1:879:G:OP2	2.21	0.40
54:O1:948:C:OP1	54:O1:962:G:OP1	2.39	0.40
54:O1:1420:A:H5'	54:O1:1421:G:OP2	2.21	0.40
54:O1:1486:U:H2'	54:O1:1487:U:C6	2.56	0.40
54:O1:2024:G:H2'	54:O1:2025:C:O4'	2.21	0.40
54:O1:2263:C:H2'	54:O1:2264:C:O4'	2.21	0.40
58:Y:26:A:H3'	58:Y:27:G:C8	2.56	0.40
59:Z:261:PHE:O	59:Z:262:ARG:HG2	2.21	0.40
1:O4:135:PRO:HG2	36:F:80:PHE:CD1	2.55	0.40
4:O7:92:GLY:O	4:O7:95:MET:HG3	2.21	0.40
21:24:20:LEU:HD11	21:24:41:GLU:CG	2.50	0.40
22:25:30:GLY:N	22:25:57:ALA:O	2.49	0.40
39:I:38:PHE:O	39:I:41:GLU:HB2	2.22	0.40
46:P:40:ASN:ND2	46:P:43:ALA:HB2	2.36	0.40
47:Q:47:ASP:HB2	47:Q:74:LEU:HD22	2.04	0.40
47:Q:59:GLU:HB3	47:Q:75:VAL:HB	2.02	0.40
50:T:4:LYS:HZ1	53:A:61:G:P	2.44	0.40
51:U:19:LYS:C	51:U:21:SER:H	2.25	0.40
51:U:28:LEU:HD23	51:U:28:LEU:O	2.21	0.40
53:A:119:A:H4'	53:A:120:A:C8	2.56	0.40
53:A:357:G:O2'	53:A:358:U:H5'	2.22	0.40
53:A:687:A:N3	53:A:688:G:H1'	2.36	0.40
53:A:1251:A:H2'	53:A:1252:A:O4'	2.21	0.40
53:A:1254:A:H2'	53:A:1255:G:C8	2.56	0.40
54:O1:20:C:H2'	54:O1:21:A:H8	1.86	0.40
54:O1:222:A:N6	54:O1:232:G:H1'	2.36	0.40
54:O1:809:G:O2'	54:O1:810:U:H5'	2.21	0.40
54:O1:1045:C:H5'	54:O1:1046:A:C5'	2.51	0.40
54:O1:1307:A:H2'	54:O1:1308:A:H5'	2.04	0.40
54:O1:2078:C:O2'	54:O1:2079:U:H5'	2.21	0.40
54:O1:2146:C:H4'	54:O1:2147:A:C4	2.56	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:01:2619:C:O2'	54:01:2620:C:H5'	2.21	0.40
54:01:2685:G:H2'	54:01:2686:G:H8	1.86	0.40
54:01:2752:C:H2'	54:01:2753:A:O4'	2.22	0.40
56:X:59:A:C2'	56:X:60:U:H5'	2.51	0.40
58:Y:68:C:H2'	58:Y:69:G:C8	2.56	0.40

There are no symmetry-related clashes.

5.3 Torsion angles [i](#)

5.3.1 Protein backbone [i](#)

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	04	269/271 (99%)	233 (87%)	33 (12%)	3 (1%)	14	53
2	05	207/209 (99%)	184 (89%)	23 (11%)	0	100	100
3	06	199/201 (99%)	175 (88%)	22 (11%)	2 (1%)	15	55
4	07	175/177 (99%)	154 (88%)	20 (11%)	1 (1%)	25	64
5	08	174/176 (99%)	155 (89%)	13 (8%)	6 (3%)	3	31
6	09	147/149 (99%)	117 (80%)	26 (18%)	4 (3%)	5	35
7	10	129/131 (98%)	97 (75%)	22 (17%)	10 (8%)	1	11
8	11	139/141 (99%)	117 (84%)	17 (12%)	5 (4%)	3	29
9	12	140/142 (99%)	132 (94%)	7 (5%)	1 (1%)	22	61
10	13	120/122 (98%)	101 (84%)	17 (14%)	2 (2%)	9	45
11	14	141/143 (99%)	112 (79%)	23 (16%)	6 (4%)	2	24
12	15	134/136 (98%)	115 (86%)	18 (13%)	1 (1%)	22	61
13	16	118/120 (98%)	102 (86%)	14 (12%)	2 (2%)	9	45
14	17	114/116 (98%)	106 (93%)	8 (7%)	0	100	100
15	18	112/114 (98%)	104 (93%)	8 (7%)	0	100	100
16	19	115/117 (98%)	108 (94%)	7 (6%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
17	20	101/103 (98%)	83 (82%)	16 (16%)	2 (2%)	7	41
18	21	108/110 (98%)	95 (88%)	11 (10%)	2 (2%)	8	42
19	22	91/93 (98%)	78 (86%)	13 (14%)	0	100	100
20	23	100/102 (98%)	83 (83%)	14 (14%)	3 (3%)	4	33
21	24	92/94 (98%)	85 (92%)	7 (8%)	0	100	100
22	25	73/75 (97%)	65 (89%)	8 (11%)	0	100	100
23	26	75/77 (97%)	66 (88%)	9 (12%)	0	100	100
24	27	61/63 (97%)	56 (92%)	4 (7%)	1 (2%)	9	46
25	28	56/58 (97%)	51 (91%)	5 (9%)	0	100	100
26	29	64/66 (97%)	51 (80%)	13 (20%)	0	100	100
27	30	54/56 (96%)	52 (96%)	2 (4%)	0	100	100
28	31	48/50 (96%)	45 (94%)	2 (4%)	1 (2%)	7	40
29	32	44/46 (96%)	37 (84%)	7 (16%)	0	100	100
30	33	62/64 (97%)	54 (87%)	7 (11%)	1 (2%)	9	46
31	34	36/38 (95%)	29 (81%)	6 (17%)	1 (3%)	5	34
32	B	216/218 (99%)	180 (83%)	30 (14%)	6 (3%)	5	34
33	C	204/206 (99%)	187 (92%)	16 (8%)	1 (0%)	29	68
34	D	203/205 (99%)	167 (82%)	31 (15%)	5 (2%)	5	36
35	E	155/157 (99%)	133 (86%)	15 (10%)	7 (4%)	2	23
36	F	98/100 (98%)	75 (76%)	18 (18%)	5 (5%)	2	20
37	G	149/151 (99%)	129 (87%)	18 (12%)	2 (1%)	12	50
38	H	127/129 (98%)	120 (94%)	6 (5%)	1 (1%)	19	59
39	I	125/127 (98%)	98 (78%)	22 (18%)	5 (4%)	3	26
40	J	96/98 (98%)	78 (81%)	10 (10%)	8 (8%)	1	10
41	K	114/116 (98%)	92 (81%)	19 (17%)	3 (3%)	5	35
42	L	121/123 (98%)	95 (78%)	18 (15%)	8 (7%)	1	16
43	M	112/114 (98%)	95 (85%)	14 (12%)	3 (3%)	5	35
44	N	98/100 (98%)	86 (88%)	11 (11%)	1 (1%)	15	55
45	O	86/88 (98%)	74 (86%)	11 (13%)	1 (1%)	13	51
46	P	80/82 (98%)	69 (86%)	9 (11%)	2 (2%)	5	36
47	Q	78/80 (98%)	61 (78%)	11 (14%)	6 (8%)	1	12

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
48	R	63/65 (97%)	50 (79%)	7 (11%)	6 (10%)	0	8
49	S	77/79 (98%)	66 (86%)	9 (12%)	2 (3%)	5	35
50	T	83/85 (98%)	75 (90%)	7 (8%)	1 (1%)	13	51
51	U	63/65 (97%)	41 (65%)	15 (24%)	7 (11%)	0	6
52	03	130/234 (56%)	107 (82%)	22 (17%)	1 (1%)	19	59
59	Z	390/392 (100%)	329 (84%)	55 (14%)	6 (2%)	10	47
All	All	6366/6574 (97%)	5449 (86%)	776 (12%)	141 (2%)	10	39

All (141) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
3	06	83	VAL
5	08	46	ASP
5	08	47	ASN
5	08	119	GLY
6	09	9	VAL
7	10	58	THR
7	10	80	THR
7	10	108	VAL
7	10	118	ILE
8	11	90	GLY
10	13	35	VAL
17	20	54	VAL
20	23	98	ASN
24	27	24	GLU
32	B	17	HIS
32	B	19	THR
32	B	73	ARG
36	F	54	LEU
36	F	99	ALA
39	I	57	VAL
39	I	90	ASP
39	I	91	GLU
40	J	34	ALA
40	J	58	ASN
42	L	102	ASP
45	O	46	LYS
48	R	71	ASP
51	U	24	LYS
51	U	34	ARG

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Mol	Chain	Res	Type
59	Z	198	TYR
1	04	232	GLY
3	06	11	ALA
4	07	149	ARG
5	08	45	ALA
6	09	10	ALA
6	09	15	LEU
7	10	114	GLU
8	11	13	ALA
11	14	31	GLY
11	14	36	LYS
11	14	65	GLY
18	21	63	GLY
18	21	64	ALA
20	23	6	ARG
30	33	31	ILE
32	B	87	ASP
34	D	23	GLY
34	D	169	TRP
35	E	23	THR
35	E	89	THR
35	E	99	SER
35	E	122	VAL
36	F	53	LYS
36	F	94	HIS
37	G	18	GLY
37	G	20	GLU
38	H	47	ASP
39	I	125	GLN
40	J	35	GLN
40	J	38	GLY
40	J	92	LEU
41	K	125	LYS
42	L	77	SER
42	L	101	LEU
44	N	3	GLN
46	P	79	ASN
47	Q	17	GLU
47	Q	49	ASN
48	R	45	GLY
48	R	46	THR
50	T	68	LYS

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Mol	Chain	Res	Type
51	U	14	ALA
51	U	64	ALA
6	09	41	LYS
7	10	81	LEU
11	14	29	LYS
11	14	86	GLU
31	34	36	ARG
34	D	152	SER
35	E	90	GLY
40	J	89	ARG
42	L	2	THR
42	L	23	LEU
46	P	44	SER
47	Q	50	ASN
48	R	10	CYS
49	S	4	LEU
51	U	9	GLU
51	U	12	ASP
59	Z	3	GLU
59	Z	367	ALA
7	10	119	PRO
7	10	123	ILE
10	13	93	GLN
12	15	70	ASP
13	16	117	ASP
20	23	51	LEU
35	E	121	ASN
39	I	102	PHE
41	K	13	LYS
42	L	24	GLU
42	L	33	CYS
43	M	4	ALA
59	Z	51	ASN
5	08	117	PRO
5	08	175	LYS
7	10	48	ALA
7	10	68	PRO
17	20	29	THR
28	31	40	PRO
32	B	18	GLN
34	D	47	LEU
34	D	167	PRO

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Mol	Chain	Res	Type
41	K	92	ARG
42	L	27	PRO
47	Q	80	LYS
51	U	66	ARG
52	03	52	ALA
59	Z	295	PRO
1	04	233	GLY
8	11	11	GLN
11	14	88	GLY
32	B	88	GLN
33	C	46	LEU
35	E	98	ALA
36	F	56	LYS
40	J	42	LEU
43	M	65	GLU
47	Q	70	LYS
48	R	11	ARG
8	11	19	PRO
40	J	33	GLY
43	M	6	ILE
1	04	7	PRO
9	12	81	ILE
13	16	105	GLY
49	S	53	GLY
59	Z	122	GLY
8	11	4	VAL
47	Q	31	PRO
48	R	20	ILE

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	04	216/216 (100%)	210 (97%)	6 (3%)	43	72
2	05	164/164 (100%)	164 (100%)	0	100	100
3	06	165/165 (100%)	163 (99%)	2 (1%)	71	87

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
4	07	148/148 (100%)	148 (100%)	0	100	100
5	08	137/137 (100%)	137 (100%)	0	100	100
6	09	114/114 (100%)	111 (97%)	3 (3%)	46	74
7	10	100/100 (100%)	100 (100%)	0	100	100
8	11	109/109 (100%)	108 (99%)	1 (1%)	78	90
9	12	116/116 (100%)	114 (98%)	2 (2%)	60	82
10	13	103/103 (100%)	102 (99%)	1 (1%)	76	88
11	14	102/102 (100%)	101 (99%)	1 (1%)	76	88
12	15	109/109 (100%)	107 (98%)	2 (2%)	59	81
13	16	100/100 (100%)	98 (98%)	2 (2%)	55	79
14	17	86/86 (100%)	86 (100%)	0	100	100
15	18	99/99 (100%)	99 (100%)	0	100	100
16	19	89/89 (100%)	87 (98%)	2 (2%)	52	77
17	20	84/84 (100%)	84 (100%)	0	100	100
18	21	93/93 (100%)	91 (98%)	2 (2%)	52	77
19	22	80/80 (100%)	79 (99%)	1 (1%)	69	86
20	23	83/83 (100%)	82 (99%)	1 (1%)	71	87
21	24	78/78 (100%)	77 (99%)	1 (1%)	69	86
22	25	57/57 (100%)	57 (100%)	0	100	100
23	26	67/67 (100%)	65 (97%)	2 (3%)	41	71
24	27	55/55 (100%)	54 (98%)	1 (2%)	59	81
25	28	48/48 (100%)	48 (100%)	0	100	100
26	29	59/59 (100%)	59 (100%)	0	100	100
27	30	47/47 (100%)	47 (100%)	0	100	100
28	31	45/45 (100%)	45 (100%)	0	100	100
29	32	38/38 (100%)	38 (100%)	0	100	100
30	33	51/51 (100%)	51 (100%)	0	100	100
31	34	34/34 (100%)	34 (100%)	0	100	100
32	B	180/180 (100%)	178 (99%)	2 (1%)	73	88
33	C	170/170 (100%)	166 (98%)	4 (2%)	49	75
34	D	172/172 (100%)	168 (98%)	4 (2%)	50	76

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
35	E	119/119 (100%)	118 (99%)	1 (1%)	81	91
36	F	87/87 (100%)	85 (98%)	2 (2%)	50	76
37	G	124/124 (100%)	124 (100%)	0	100	100
38	H	104/104 (100%)	104 (100%)	0	100	100
39	I	105/105 (100%)	105 (100%)	0	100	100
40	J	86/86 (100%)	86 (100%)	0	100	100
41	K	89/89 (100%)	88 (99%)	1 (1%)	73	88
42	L	103/103 (100%)	102 (99%)	1 (1%)	76	88
43	M	92/92 (100%)	90 (98%)	2 (2%)	52	77
44	N	83/83 (100%)	82 (99%)	1 (1%)	71	87
45	O	76/76 (100%)	76 (100%)	0	100	100
46	P	65/65 (100%)	63 (97%)	2 (3%)	40	71
47	Q	74/74 (100%)	74 (100%)	0	100	100
48	R	56/56 (100%)	53 (95%)	3 (5%)	22	57
49	S	70/70 (100%)	70 (100%)	0	100	100
50	T	65/65 (100%)	65 (100%)	0	100	100
51	U	55/55 (100%)	51 (93%)	4 (7%)	14	46
52	03	110/181 (61%)	106 (96%)	4 (4%)	35	67
59	Z	324/325 (100%)	314 (97%)	10 (3%)	40	71
All	All	5285/5357 (99%)	5214 (99%)	71 (1%)	70	86

All (71) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	04	36	ASN
1	04	85	ASN
1	04	132	ARG
1	04	212	TRP
1	04	239	PHE
1	04	261	ARG
3	06	156	ASN
3	06	163	ASN
6	09	11	ASN
6	09	12	LEU
6	09	75	LEU

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Mol	Chain	Res	Type
8	11	131	THR
9	12	27	ARG
9	12	57	LEU
10	13	104	THR
11	14	27	LEU
12	15	6	ARG
12	15	84	LYS
13	16	2	ARG
13	16	13	ASN
16	19	49	ARG
16	19	57	ARG
18	21	3	THR
18	21	57	ASN
19	22	32	LEU
20	23	21	ARG
21	24	79	ARG
23	26	16	ASN
23	26	26	ARG
24	27	7	ARG
32	B	23	ASN
32	B	35	ASN
33	C	31	ASN
33	C	146	LYS
33	C	156	LEU
33	C	163	ARG
34	D	62	ARG
34	D	139	ASN
34	D	177	MET
34	D	183	ARG
35	E	100	GLU
36	F	12	PRO
36	F	79	ARG
41	K	12	ARG
42	L	4	ASN
43	M	7	ASN
43	M	99	GLN
44	N	68	ARG
46	P	13	LYS
46	P	19	VAL
48	R	11	ARG
48	R	41	SER
48	R	70	THR

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Mol	Chain	Res	Type
51	U	10	PRO
51	U	28	LEU
51	U	61	ARG
51	U	67	THR
52	03	25	GLU
52	03	33	LEU
52	03	41	SER
52	03	55	SER
59	Z	68	GLU
59	Z	91	MET
59	Z	98	MET
59	Z	236	ILE
59	Z	237	LYS
59	Z	248	LYS
59	Z	269	ARG
59	Z	358	MET
59	Z	381	ARG
59	Z	391	VAL

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (90) such sidechains are listed below:

Mol	Chain	Res	Type
1	04	24	HIS
1	04	36	ASN
1	04	89	ASN
1	04	127	ASN
1	04	133	ASN
1	04	152	GLN
2	05	32	ASN
2	05	49	GLN
3	06	94	GLN
3	06	156	ASN
3	06	163	ASN
4	07	26	GLN
4	07	51	ASN
5	08	37	ASN
5	08	63	GLN
5	08	87	GLN
5	08	138	GLN
6	09	11	ASN
6	09	66	ASN
6	09	133	GLN

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Mol	Chain	Res	Type
7	10	88	HIS
7	10	103	ASN
8	11	5	GLN
8	11	29	GLN
8	11	30	GLN
8	11	93	ASN
9	12	136	GLN
12	15	3	GLN
12	15	13	HIS
13	16	3	HIS
13	16	31	HIS
14	17	100	HIS
15	18	6	GLN
15	18	65	ASN
16	19	36	GLN
16	19	43	GLN
17	20	6	GLN
17	20	18	GLN
17	20	82	HIS
18	21	57	ASN
19	22	59	ASN
20	23	45	GLN
20	23	68	ASN
20	23	98	ASN
21	24	51	GLN
22	25	8	ASN
22	25	42	HIS
23	26	16	ASN
26	29	61	ASN
30	33	27	ASN
32	B	23	ASN
32	B	35	ASN
32	B	121	GLN
32	B	177	ASN
33	C	31	ASN
33	C	139	ASN
34	D	88	ASN
34	D	139	ASN
35	E	81	GLN
36	F	17	GLN
36	F	68	GLN
37	G	129	ASN

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Mol	Chain	Res	Type
39	I	30	ASN
39	I	36	GLN
39	I	74	GLN
39	I	80	HIS
39	I	125	GLN
40	J	20	GLN
40	J	58	ASN
41	K	28	ASN
41	K	80	ASN
42	L	4	ASN
43	M	7	ASN
44	N	42	ASN
44	N	48	GLN
45	O	36	ASN
45	O	45	HIS
46	P	26	ASN
46	P	63	GLN
48	R	51	GLN
49	S	55	GLN
50	T	51	ASN
52	03	67	HIS
52	03	160	GLN
59	Z	13	ASN
59	Z	63	ASN
59	Z	66	HIS
59	Z	97	GLN
59	Z	329	GLN
59	Z	355	ASN

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
53	A	1538/1539 (99%)	160 (10%)	9 (0%)
54	01	2902/2903 (99%)	358 (12%)	16 (0%)
55	02	119/120 (99%)	11 (9%)	2 (1%)
56	W	76/77 (98%)	7 (9%)	0
56	X	76/77 (98%)	13 (17%)	0
57	V	17/18 (94%)	2 (11%)	0
58	Y	75/76 (98%)	14 (18%)	0
All	All	4803/4810 (99%)	565 (11%)	27 (0%)

All (565) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
53	A	6	G
53	A	9	G
53	A	22	G
53	A	31	G
53	A	32	A
53	A	39	G
53	A	47	C
53	A	51	A
53	A	71	A
53	A	82	G
53	A	87	C
53	A	121	U
53	A	130	A
53	A	173	U
53	A	183	C
53	A	184	G
53	A	197	A
53	A	209	U
53	A	210	C
53	A	226	G
53	A	247	G
53	A	251	G
53	A	266	G
53	A	267	C
53	A	279	A
53	A	281	G
53	A	289	G
53	A	306	A
53	A	328	C
53	A	345	C
53	A	346	G
53	A	351	G
53	A	352	C
53	A	363	A
53	A	367	U
53	A	372	C
53	A	411	A
53	A	412	A
53	A	413	G
53	A	422	C
53	A	429	U
53	A	430	A
53	A	439	U

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Mol	Chain	Res	Type
53	A	467	U
53	A	484	G
53	A	485	U
53	A	486	U
53	A	496	A
53	A	497	G
53	A	509	A
53	A	510	A
53	A	527	G
53	A	531	U
53	A	532	A
53	A	533	A
53	A	547	A
53	A	559	A
53	A	561	U
53	A	572	A
53	A	573	A
53	A	575	G
53	A	576	C
53	A	577	G
53	A	607	A
53	A	633	G
53	A	642	A
53	A	665	A
53	A	703	G
53	A	713	G
53	A	724	G
53	A	731	G
53	A	755	G
53	A	777	A
53	A	815	A
53	A	817	C
53	A	818	G
53	A	819	A
53	A	821	G
53	A	832	G
53	A	842	U
53	A	843	U
53	A	844	G
53	A	846	G
53	A	873	A
53	A	890	G

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Mol	Chain	Res	Type
53	A	902	G
53	A	926	G
53	A	934	C
53	A	960	U
53	A	961	U
53	A	966	G
53	A	969	A
53	A	975	A
53	A	976	G
53	A	977	A
53	A	992	U
53	A	993	G
53	A	994	A
53	A	1004	A
53	A	1020	G
53	A	1028	C
53	A	1031	C
53	A	1033	G
53	A	1034	G
53	A	1053	G
53	A	1054	C
53	A	1085	U
53	A	1094	G
53	A	1101	A
53	A	1130	A
53	A	1137	C
53	A	1138	G
53	A	1139	G
53	A	1159	U
53	A	1160	G
53	A	1168	U
53	A	1182	G
53	A	1184	G
53	A	1191	A
53	A	1196	A
53	A	1198	G
53	A	1201	A
53	A	1202	U
53	A	1207	G
53	A	1213	A
53	A	1225	A
53	A	1238	A

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Mol	Chain	Res	Type
53	A	1241	G
53	A	1256	A
53	A	1257	A
53	A	1258	G
53	A	1260	G
53	A	1275	A
53	A	1278	G
53	A	1280	A
53	A	1282	C
53	A	1286	U
53	A	1287	A
53	A	1290	G
53	A	1300	G
53	A	1317	C
53	A	1323	G
53	A	1346	A
53	A	1347	G
53	A	1363	A
53	A	1381	U
53	A	1395	C
53	A	1400	C
53	A	1446	A
53	A	1448	C
53	A	1452	C
53	A	1475	G
53	A	1492	A
53	A	1503	A
53	A	1506	U
53	A	1517	G
53	A	1529	G
53	A	1530	G
53	A	1534	A
53	A	1540	U
54	01	10	A
54	01	12	U
54	01	34	U
54	01	35	G
54	01	46	G
54	01	50	U
54	01	51	G
54	01	63	A
54	01	71	A

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Mol	Chain	Res	Type
54	01	74	A
54	01	75	G
54	01	102	U
54	01	118	A
54	01	120	U
54	01	138	U
54	01	139	U
54	01	140	C
54	01	141	G
54	01	142	A
54	01	158	U
54	01	162	U
54	01	163	C
54	01	181	A
54	01	196	A
54	01	216	A
54	01	221	A
54	01	222	A
54	01	228	C
54	01	229	C
54	01	242	G
54	01	248	G
54	01	249	C
54	01	255	A
54	01	266	G
54	01	276	U
54	01	281	C
54	01	294	A
54	01	301	G
54	01	311	A
54	01	312	G
54	01	322	A
54	01	323	C
54	01	329	G
54	01	330	A
54	01	353	C
54	01	361	G
54	01	371	A
54	01	372	G
54	01	373	U
54	01	386	G
54	01	387	U

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Mol	Chain	Res	Type
54	01	404	A
54	01	406	G
54	01	411	G
54	01	422	A
54	01	424	G
54	01	451	U
54	01	455	C
54	01	457	A
54	01	481	G
54	01	491	G
54	01	504	A
54	01	505	A
54	01	506	G
54	01	508	A
54	01	529	A
54	01	531	C
54	01	532	A
54	01	542	C
54	01	543	G
54	01	545	U
54	01	547	A
54	01	563	A
54	01	572	A
54	01	573	U
54	01	575	A
54	01	588	U
54	01	603	A
54	01	616	A
54	01	627	A
54	01	637	A
54	01	646	U
54	01	654	A
54	01	669	G
54	01	677	A
54	01	686	U
54	01	687	C
54	01	695	G
54	01	730	A
54	01	747	C
54	01	748	G
54	01	752	A
54	01	776	G

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Mol	Chain	Res	Type
54	01	782	A
54	01	784	G
54	01	785	G
54	01	805	G
54	01	812	C
54	01	819	A
54	01	827	U
54	01	828	U
54	01	830	G
54	01	845	A
54	01	846	U
54	01	847	U
54	01	858	G
54	01	859	G
54	01	860	U
54	01	878	A
54	01	886	A
54	01	888	C
54	01	889	C
54	01	896	A
54	01	907	G
54	01	910	A
54	01	915	C
54	01	932	U
54	01	941	A
54	01	946	C
54	01	961	C
54	01	974	G
54	01	983	A
54	01	990	A
54	01	995	C
54	01	996	A
54	01	1012	U
54	01	1013	C
54	01	1021	A
54	01	1022	G
54	01	1026	G
54	01	1033	U
54	01	1045	C
54	01	1046	A
54	01	1054	A
54	01	1057	A

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Mol	Chain	Res	Type
54	01	1060	U
54	01	1062	G
54	01	1065	U
54	01	1066	U
54	01	1068	G
54	01	1071	G
54	01	1079	C
54	01	1084	A
54	01	1088	A
54	01	1090	A
54	01	1103	A
54	01	1104	C
54	01	1106	G
54	01	1111	A
54	01	1112	G
54	01	1131	G
54	01	1132	U
54	01	1135	C
54	01	1143	A
54	01	1157	G
54	01	1174	U
54	01	1176	U
54	01	1177	G
54	01	1179	G
54	01	1180	U
54	01	1206	G
54	01	1212	G
54	01	1238	G
54	01	1248	G
54	01	1250	G
54	01	1251	C
54	01	1253	A
54	01	1256	G
54	01	1271	G
54	01	1272	A
54	01	1301	A
54	01	1321	A
54	01	1329	U
54	01	1330	C
54	01	1332	G
54	01	1345	C
54	01	1365	A

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Mol	Chain	Res	Type
54	01	1378	A
54	01	1379	U
54	01	1383	A
54	01	1395	A
54	01	1416	G
54	01	1419	A
54	01	1420	A
54	01	1428	C
54	01	1454	C
54	01	1461	C
54	01	1476	U
54	01	1482	G
54	01	1490	A
54	01	1491	G
54	01	1498	C
54	01	1515	A
54	01	1524	G
54	01	1533	C
54	01	1535	A
54	01	1536	C
54	01	1555	G
54	01	1560	G
54	01	1569	A
54	01	1581	G
54	01	1608	A
54	01	1611	C
54	01	1616	A
54	01	1634	A
54	01	1646	C
54	01	1647	U
54	01	1648	U
54	01	1674	G
54	01	1715	G
54	01	1729	U
54	01	1730	C
54	01	1738	G
54	01	1758	U
54	01	1764	C
54	01	1773	A
54	01	1780	A
54	01	1800	C
54	01	1801	A

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Mol	Chain	Res	Type
54	01	1808	A
54	01	1816	C
54	01	1829	A
54	01	1847	A
54	01	1871	A
54	01	1901	A
54	01	1906	G
54	01	1913	A
54	01	1929	G
54	01	1930	G
54	01	1931	U
54	01	1937	A
54	01	1944	U
54	01	1955	U
54	01	1967	C
54	01	1970	A
54	01	1971	U
54	01	1972	G
54	01	1991	U
54	01	1992	G
54	01	1993	U
54	01	1997	C
54	01	2022	U
54	01	2023	C
54	01	2030	A
54	01	2031	A
54	01	2036	C
54	01	2043	C
54	01	2049	G
54	01	2055	C
54	01	2056	G
54	01	2060	A
54	01	2061	G
54	01	2062	A
54	01	2069	G
54	01	2072	C
54	01	2095	A
54	01	2096	C
54	01	2108	A
54	01	2111	U
54	01	2112	G
54	01	2115	G

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Mol	Chain	Res	Type
54	01	2118	U
54	01	2119	A
54	01	2124	G
54	01	2125	G
54	01	2127	G
54	01	2131	U
54	01	2132	U
54	01	2133	G
54	01	2145	C
54	01	2147	A
54	01	2162	G
54	01	2164	C
54	01	2172	U
54	01	2173	A
54	01	2189	U
54	01	2198	A
54	01	2203	U
54	01	2204	G
54	01	2211	A
54	01	2213	U
54	01	2225	A
54	01	2238	G
54	01	2239	G
54	01	2250	G
54	01	2259	U
54	01	2278	A
54	01	2283	C
54	01	2287	A
54	01	2297	A
54	01	2305	U
54	01	2309	A
54	01	2325	G
54	01	2327	A
54	01	2334	U
54	01	2350	C
54	01	2382	G
54	01	2383	G
54	01	2385	C
54	01	2392	A
54	01	2402	U
54	01	2406	A
54	01	2407	A

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Mol	Chain	Res	Type
54	01	2423	U
54	01	2424	C
54	01	2429	G
54	01	2430	A
54	01	2435	A
54	01	2441	U
54	01	2448	A
54	01	2476	A
54	01	2498	C
54	01	2502	G
54	01	2503	A
54	01	2504	U
54	01	2505	G
54	01	2518	A
54	01	2529	G
54	01	2547	A
54	01	2554	U
54	01	2567	G
54	01	2572	A
54	01	2602	A
54	01	2609	U
54	01	2613	U
54	01	2629	U
54	01	2646	C
54	01	2655	G
54	01	2682	A
54	01	2685	G
54	01	2689	U
54	01	2690	U
54	01	2714	G
54	01	2733	A
54	01	2744	G
54	01	2748	A
54	01	2757	A
54	01	2764	A
54	01	2765	A
54	01	2771	C
54	01	2778	A
54	01	2779	U
54	01	2791	G
54	01	2794	C
54	01	2797	U

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Mol	Chain	Res	Type
54	01	2798	U
54	01	2799	A
54	01	2800	A
54	01	2809	A
54	01	2818	U
54	01	2820	A
54	01	2821	A
54	01	2833	U
54	01	2848	G
54	01	2867	G
54	01	2868	A
54	01	2880	C
54	01	2884	U
55	02	4	C
55	02	12	C
55	02	13	G
55	02	24	G
55	02	35	C
55	02	44	G
55	02	67	G
55	02	89	U
55	02	108	A
55	02	109	A
55	02	116	G
56	X	8	U
56	X	9	G
56	X	10	G
56	X	14	A
56	X	19	G
56	X	20	U
56	X	21	A
56	X	22	G
56	X	30	G
56	X	34	C
56	X	61	C
56	X	64	G
56	X	70	G
57	V	12	A
57	V	19	U
56	W	9	G
56	W	19	G
56	W	20	U

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Mol	Chain	Res	Type
56	W	47	U
56	W	48	C
56	W	61	C
56	W	76	A
58	Y	7	A
58	Y	8	U
58	Y	9	A
58	Y	10	G
58	Y	18	G
58	Y	21	A
58	Y	26	A
58	Y	44	G
58	Y	45	U
58	Y	46	G
58	Y	48	C
58	Y	55	U
58	Y	61	C
58	Y	63	G

All (27) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
53	A	246	A
53	A	280	C
53	A	429	U
53	A	438	U
53	A	495	A
53	A	960	U
53	A	1129	C
53	A	1190	G
53	A	1399	C
54	01	265	A
54	01	372	G
54	01	421	C
54	01	490	C
54	01	859	G
54	01	1020	A
54	01	1070	A
54	01	1111	A
54	01	1130	U
54	01	1475	G
54	01	1930	G

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Mol	Chain	Res	Type
54	01	2286	G
54	01	2296	U
54	01	2326	C
54	01	2391	G
54	01	2756	U
55	02	66	A
55	02	88	C

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 monosaccharides in this entry.

5.6 Ligand geometry [i](#)

3 ligands are modelled in this entry.

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

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
62	GCP	Z	401	-	27,34,34	2.31	11 (40%)	34,54,54	3.88	17 (50%)
60	FME	W	101	-	8,9,10	0.82	0	7,9,11	1.33	1 (14%)
61	PHE	Y	101	58	10,11,12	1.01	0	10,13,15	0.34	0

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

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
62	GCP	Z	401	-	-	9/15/38/38	0/3/3/3
60	FME	W	101	-	-	3/7/9/11	-
61	PHE	Y	101	58	-	1/5/6/8	0/1/1/1

All (11) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
62	Z	401	GCP	O4'-C1'	5.13	1.48	1.41
62	Z	401	GCP	PB-O3A	-3.96	1.53	1.58
62	Z	401	GCP	C2'-C1'	3.96	1.59	1.53
62	Z	401	GCP	C5-C6	3.81	1.47	1.41
62	Z	401	GCP	C2-N2	3.72	1.41	1.33
62	Z	401	GCP	C6-N1	3.42	1.39	1.33
62	Z	401	GCP	C2'-C3'	2.94	1.61	1.53
62	Z	401	GCP	C2-N1	2.80	1.40	1.35
62	Z	401	GCP	C5'-C4'	2.45	1.59	1.51
62	Z	401	GCP	C3'-C4'	2.18	1.58	1.53
62	Z	401	GCP	O4'-C4'	2.02	1.49	1.45

All (18) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
62	Z	401	GCP	C1'-N9-C4	12.96	149.41	126.64
62	Z	401	GCP	C5-C6-N1	-9.02	111.10	123.43
62	Z	401	GCP	C2-N1-C6	7.33	127.57	115.93
62	Z	401	GCP	O1G-PG-C3B	-7.29	95.52	111.24
62	Z	401	GCP	O4'-C1'-C2'	-4.93	99.72	106.93
62	Z	401	GCP	C4-C5-C6	-4.37	116.63	120.80
62	Z	401	GCP	O5'-PA-O1A	-4.15	92.84	109.07
62	Z	401	GCP	O2B-PB-O1B	3.72	122.48	110.07
62	Z	401	GCP	N3-C2-N1	-3.51	122.54	127.22
62	Z	401	GCP	O3G-PG-O1G	3.01	120.34	112.39
62	Z	401	GCP	O3'-C3'-C4'	-2.92	102.59	111.05
62	Z	401	GCP	O2G-PG-C3B	2.89	113.42	106.40
62	Z	401	GCP	PB-O3A-PA	2.81	141.47	132.56
62	Z	401	GCP	C2-N3-C4	-2.70	112.27	115.36
60	W	101	FME	O-C-CA	-2.42	118.45	124.78
62	Z	401	GCP	O4'-C4'-C5'	2.34	117.09	109.37
62	Z	401	GCP	O2A-PA-O1A	2.20	123.09	112.24
62	Z	401	GCP	C4-C5-N7	2.18	111.67	109.40

There are no chirality outliers.

All (13) torsion outliers are listed below:

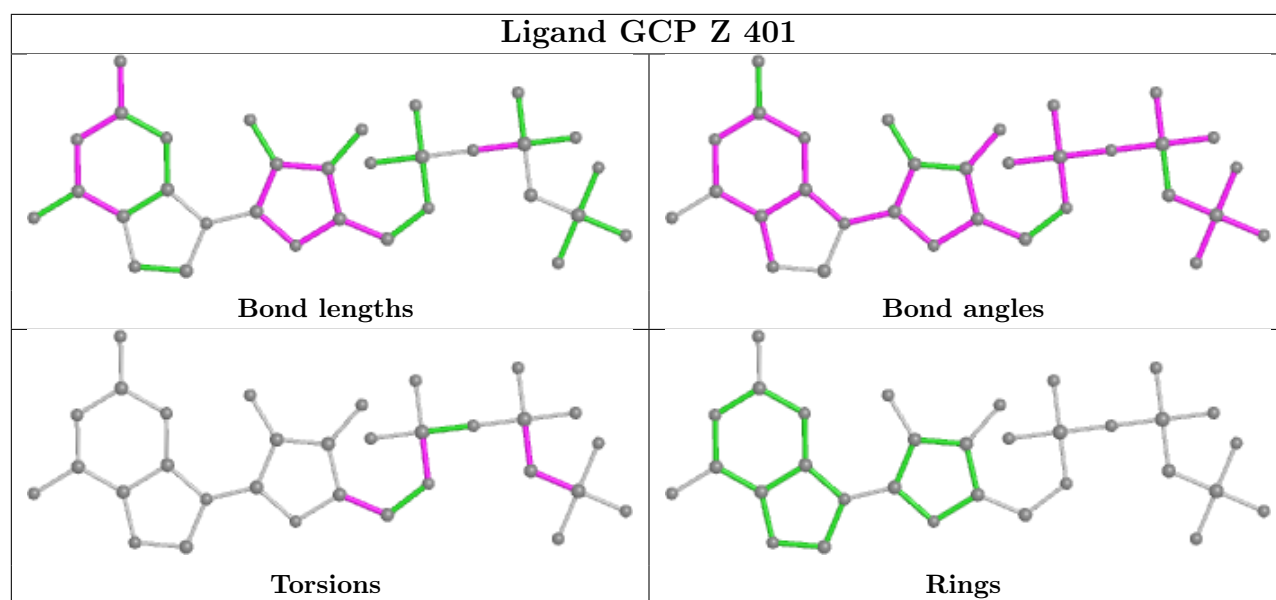
Mol	Chain	Res	Type	Atoms
60	W	101	FME	O1-CN-N-CA
60	W	101	FME	O-C-CA-CB
61	Y	101	PHE	O-C-CA-CB
62	Z	401	GCP	PB-C3B-PG-O1G
62	Z	401	GCP	PB-C3B-PG-O2G
62	Z	401	GCP	PG-C3B-PB-O1B
62	Z	401	GCP	C5'-O5'-PA-O3A
62	Z	401	GCP	O4'-C4'-C5'-O5'
60	W	101	FME	C-CA-CB-CG
62	Z	401	GCP	C5'-O5'-PA-O1A
62	Z	401	GCP	C5'-O5'-PA-O2A
62	Z	401	GCP	PB-C3B-PG-O3G
62	Z	401	GCP	C3'-C4'-C5'-O5'

There are no ring outliers.

2 monomers are involved in 6 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
60	W	101	FME	3	0
61	Y	101	PHE	3	0

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



5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

6 Map visualisation

This section contains visualisations of the EMDB entry EMD-8616. 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

This section was not generated.

6.2 Central slices

This section was not generated.

6.3 Largest variance slices

This section was not generated.

6.4 Orthogonal surface views

This section was not generated.

6.5 Mask visualisation

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

7 Map analysis ⓘ

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

7.1 Map-value distribution ⓘ

This section was not generated.

7.2 Volume estimate versus contour level ⓘ

This section was not generated.

7.3 Rotationally averaged power spectrum ⓘ

This section was not generated. The rotationally averaged power spectrum had issues being displayed.

8 Fourier-Shell correlation

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

9 Map-model fit

This section was not generated.