



Full wwPDB X-ray Structure Validation Report ⓘ

Feb 1, 2016 – 10:14 PM GMT

PDB ID : 4WF1
Title : Crystal structure of the E. coli ribosome bound to negamycin.
Authors : Olivier, N.B.; Altman, R.B.; Noeske, J.; Basarab, G.S.; Code, E.; Ferguson, A.D.; Gao, N.; Huang, J.; Juetten, M.F.; Livchak, S.; Miller, M.D.; Prince, D.B.; Cate, J.H.D.; Buurman, E.T.; Blanchard, S.C.
Deposited on : 2014-08-31
Resolution : 3.09 Å(reported)

This is a Full wwPDB X-ray Structure Validation Report for a publicly released PDB entry.
We welcome your comments at validation@mail.wwpdb.org
A user guide is available at
<http://wwpdb.org/validation/2016/XrayValidationReportHelp>
with specific help available everywhere you see the ⓘ symbol.

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

MolProbity : 4.02b-467
Mogul : 1.7 (RC4), CSD as536be (2015)
Xtriage (Phenix) : 1.9-1692
EDS : rb-20026688
Percentile statistics : 20151230.v01 (using entries in the PDB archive December 30th 2015)
Refmac : 5.8.0135
CCP4 : 6.5.0
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : trunk26865

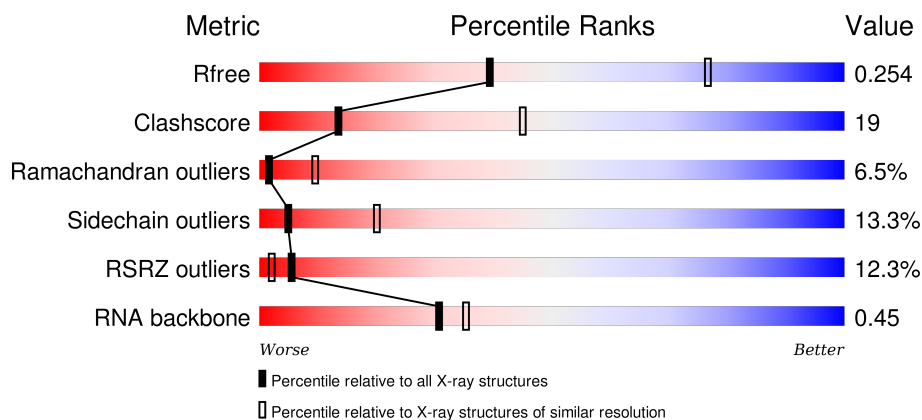
1 Overall quality at a glance i

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 3.09 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)	Similar resolution (#Entries, resolution range(Å))
R_{free}	91344	1114 (3.14-3.06)
Clashscore	102246	1222 (3.14-3.06)
Ramachandran outliers	100387	1174 (3.14-3.06)
Sidechain outliers	100360	1174 (3.14-3.06)
RSRZ outliers	91569	1119 (3.14-3.06)
RNA backbone	2183	1010 (3.52-2.68)

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

Mol	Chain	Length	Quality of chain
1	AA	1539	<div> <div>4%</div> <div>35%</div> <div>49%</div> <div>15%</div> </div>
1	CA	1539	<div> <div>9%</div> <div>35%</div> <div>51%</div> <div>13%</div> </div>
2	AB	218	<div> <div>6%</div> <div>31%</div> <div>51%</div> <div>16%</div> <div>•</div> </div>
2	CB	218	<div> <div>11%</div> <div>36%</div> <div>45%</div> <div>16%</div> <div>•</div> </div>

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Mol	Chain	Length	Quality of chain
3	AC	206	
3	CC	206	
4	AD	205	
4	CD	205	
5	AE	150	
5	CE	150	
6	AF	100	
6	CF	100	
7	AG	151	
7	CG	151	
8	AH	129	
8	CH	129	
9	AI	127	
9	CI	127	
10	AJ	98	
10	CJ	98	
11	AK	117	
11	CK	117	
12	AL	123	
12	CL	123	
13	AM	114	
13	CM	114	
14	AN	100	
14	CN	100	
15	AO	88	

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Mol	Chain	Length	Quality of chain
15	CO	88	
16	AP	82	
16	CP	82	
17	AQ	80	
17	CQ	80	
18	AR	55	
18	CR	55	
19	AS	79	
19	CS	79	
20	AT	85	
20	CT	85	
21	AU	51	
21	CU	51	
22	BA	2903	
22	DA	2903	
23	BB	119	
23	DB	119	
24	BC	271	
24	DC	271	
25	BD	209	
25	DD	209	
26	BE	201	
26	DE	201	
27	BF	177	
27	DF	177	

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Mol	Chain	Length	Quality of chain
28	BG	176	
28	DG	176	
29	BH	149	
29	DH	149	
30	BI	141	
30	DI	141	
31	BJ	142	
31	DJ	142	
32	BK	122	
32	DK	122	
33	BL	143	
33	DL	143	
34	BM	136	
34	DM	136	
35	BN	120	
35	DN	120	
36	BO	116	
36	DO	116	
37	BP	114	
37	DP	114	
38	BQ	117	
38	DQ	117	
39	BR	103	
39	DR	103	
40	BS	110	

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Mol	Chain	Length	Quality of chain
40	DS	110	
41	BT	93	
41	DT	93	
42	BU	102	
42	DU	102	
43	BV	94	
43	DV	94	
44	BW	76	
44	DW	76	
45	BX	77	
45	DX	77	
46	BY	63	
46	DY	63	
47	BZ	58	
47	DZ	58	
48	B0	56	
48	D0	56	
49	B1	50	
49	D1	50	
50	B2	46	
50	D2	46	
51	B3	64	
51	D3	64	
52	B4	38	
52	D4	38	

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Mol	Chain	Length	Quality of chain
53	B5	207	

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

Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
54	MG	AA	1617	-	-	-	X
54	MG	AA	1622	-	-	-	X
54	MG	AA	1630	-	-	-	X
54	MG	AA	1635	-	-	-	X
54	MG	AA	1662	-	-	-	X
54	MG	AA	1670	-	-	-	X
54	MG	AM	201	-	-	-	X
54	MG	BA	3040	-	-	-	X
54	MG	BA	3057	-	-	-	X
54	MG	BA	3083	-	-	-	X
54	MG	BA	3104	-	-	-	X
54	MG	BA	3124	-	-	-	X
54	MG	BA	3130	-	-	-	X
54	MG	BA	3131	-	-	-	X
54	MG	BA	3136	-	-	-	X
54	MG	BA	3146	-	-	-	X
54	MG	BA	3150	-	-	-	X
54	MG	BA	3152	-	-	-	X
54	MG	BA	3158	-	-	-	X
54	MG	BA	3160	-	-	-	X
54	MG	BA	3162	-	-	-	X
54	MG	BA	3167	-	-	-	X
54	MG	BA	3174	-	-	-	X
54	MG	BA	3177	-	-	-	X
54	MG	BA	3185	-	-	-	X
54	MG	CA	1614	-	-	-	X
54	MG	CA	1641	-	-	-	X
54	MG	DA	3027	-	-	-	X
54	MG	DA	3048	-	-	-	X
54	MG	DA	3057	-	-	-	X
54	MG	DA	3071	-	-	-	X
54	MG	DA	3104	-	-	-	X
54	MG	DA	3109	-	-	-	X
54	MG	DA	3114	-	-	-	X
54	MG	DA	3115	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
54	MG	DA	3127	-	-	-	X
54	MG	DA	3135	-	-	-	X
54	MG	DA	3147	-	-	-	X
54	MG	DA	3152	-	-	-	X
54	MG	DA	3155	-	-	-	X

2 Entry composition

There are 57 unique types of molecules in this entry. The entry contains 288204 atoms, of which 0 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a RNA chain called 16S rRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
1	AA	1538	Total	C	N	O	P	0	0	0
			32995	14716	6050	10691	1538			
1	CA	1539	Total	C	N	O	P	0	0	0
			33015	14725	6052	10699	1539			

- Molecule 2 is a protein called 30S ribosomal protein S2.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
2	AB	218	Total	C	N	O	S	0	0	0
			1705	1081	305	312	7			
2	CB	218	Total	C	N	O	S	0	0	0
			1705	1081	305	312	7			

- Molecule 3 is a protein called 30S ribosomal protein S3.

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

- Molecule 4 is a protein called 30S ribosomal protein S4.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
4	AD	205	Total	C	N	O	S	0	0	0
			1643	1026	315	298	4			
4	CD	205	Total	C	N	O	S	0	0	0
			1643	1026	315	298	4			

- Molecule 5 is a protein called 30S ribosomal protein S5.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
5	AE	150	Total	C	N	O	S	0	0	0
			1106	687	211	202	6			
5	CE	150	Total	C	N	O	S	0	0	0
			1106	687	211	202	6			

- Molecule 6 is a protein called 30S ribosomal protein S6.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
6	AF	100	Total	C	N	O	S	0	0	0
			818	515	148	149	6			
6	CF	100	Total	C	N	O	S	0	0	0
			818	515	148	149	6			

- Molecule 7 is a protein called 30S ribosomal protein S7.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
7	AG	151	Total	C	N	O	S	0	0	0
			1182	735	227	216	4			
7	CG	151	Total	C	N	O	S	0	0	0
			1182	735	227	216	4			

- Molecule 8 is a protein called 30S ribosomal protein S8.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
8	AH	129	Total	C	N	O	S	0	0	0
			979	616	173	184	6			
8	CH	129	Total	C	N	O	S	0	0	0
			979	616	173	184	6			

- Molecule 9 is a protein called 30S ribosomal protein S9.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
9	AI	127	Total	C	N	O	S	0	0	0
			1022	634	206	179	3			
9	CI	127	Total	C	N	O	S	0	0	0
			1022	634	206	179	3			

- Molecule 10 is a protein called 30S ribosomal protein S10.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	AJ	98	Total	C	N	O	S	0	0	0
			787	493	150	143	1			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	CJ	98	Total	C	N	O	S	0	0	0
			787	493	150	143	1			

- Molecule 11 is a protein called 30S ribosomal protein S11.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
11	AK	117	Total	C	N	O	S	0	0	0
			877	540	174	160	3			
11	CK	117	Total	C	N	O	S	0	0	0
			877	540	174	160	3			

- Molecule 12 is a protein called 30S ribosomal protein S12.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
12	AL	123	Total	C	N	O	S	0	0	0
			955	590	196	165	4			
12	CL	123	Total	C	N	O	S	0	0	0
			955	590	196	165	4			

- Molecule 13 is a protein called 30S ribosomal protein S13.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
13	AM	114	Total	C	N	O	S	0	0	0
			884	546	178	157	3			
13	CM	114	Total	C	N	O	S	0	0	0
			884	546	178	157	3			

- Molecule 14 is a protein called 30S ribosomal protein S14.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
14	AN	96	Total	C	N	O	S	0	0	0
			774	483	160	128	3			
14	CN	96	Total	C	N	O	S	0	0	0
			774	483	160	128	3			

- Molecule 15 is a protein called 30S ribosomal protein S15.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
15	AO	88	Total	C	N	O	S	0	0	0
			714	439	144	130	1			
15	CO	88	Total	C	N	O	S	0	0	0
			714	439	144	130	1			

- Molecule 16 is a protein called 30S ribosomal protein S16.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
16	AP	82	Total	C	N	O	S	0	0	0
			649	406	128	114	1			
16	CP	82	Total	C	N	O	S	0	0	0
			649	406	128	114	1			

- Molecule 17 is a protein called 30S ribosomal protein S17.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
17	AQ	80	Total	C	N	O	S	0	0	0
			649	411	121	114	3			
17	CQ	80	Total	C	N	O	S	0	0	0
			649	411	121	114	3			

- Molecule 18 is a protein called 30S ribosomal protein S18.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
18	AR	55	Total	C	N	O	0	0	0
			456	288	86	82			
18	CR	55	Total	C	N	O	0	0	0
			456	288	86	82			

- Molecule 19 is a protein called 30S ribosomal protein S19.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
19	AS	79	Total	C	N	O	S	0	0	0
			638	408	120	108	2			
19	CS	79	Total	C	N	O	S	0	0	0
			638	408	120	108	2			

- Molecule 20 is a protein called 30S ribosomal protein S20.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
20	AT	85	Total	C	N	O	S	0	0	0
			665	411	137	114	3			
20	CT	85	Total	C	N	O	S	0	0	0
			665	411	137	114	3			

- Molecule 21 is a protein called 30S ribosomal protein S21.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
21	AU	51	Total	C	N	O	S	0	0	0
			426	265	86	74	1			
21	CU	51	Total	C	N	O	S	0	0	0
			426	265	86	74	1			

- Molecule 22 is a RNA chain called 23S rRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
22	BA	2897	Total	C	N	O	P	0	0	0
			62195	27745	11446	20107	2897			
22	DA	2897	Total	C	N	O	P	0	0	0
			62195	27745	11446	20107	2897			

- Molecule 23 is a RNA chain called 5S rRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
23	BB	119	Total	C	N	O	P	0	0	0
			2549	1135	466	829	119			
23	DB	118	Total	C	N	O	P	0	0	0
			2529	1126	464	821	118			

- Molecule 24 is a protein called 50S ribosomal protein L2.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
24	BC	271	Total	C	N	O	S	0	0	0
			2083	1288	423	365	7			
24	DC	271	Total	C	N	O	S	0	0	0
			2083	1288	423	365	7			

- Molecule 25 is a protein called 50S ribosomal protein L3.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
25	BD	209	Total	C	N	O	S	0	0	0
			1565	979	288	294	4			
25	DD	209	Total	C	N	O	S	0	0	0
			1565	979	288	294	4			

- Molecule 26 is a protein called 50S ribosomal protein L4.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	BE	201	Total	C	N	O	S	0	0	0
			1552	974	283	290	5			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	DE	201	Total	C	N	O	S	0	0	0
			1552	974	283	290	5			

- Molecule 27 is a protein called 50S ribosomal protein L5.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
27	BF	177	Total	C	N	O	S	0	0	0
			1411	899	249	257	6			
27	DF	177	Total	C	N	O	S	0	0	0
			1411	899	249	257	6			

- Molecule 28 is a protein called 50S ribosomal protein L6.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
28	BG	176	Total	C	N	O	S	0	0	0
			1323	832	243	246	2			
28	DG	176	Total	C	N	O	S	0	0	0
			1323	832	243	246	2			

- Molecule 29 is a protein called 50S ribosomal protein L9.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
29	BH	149	Total	C	N	O	S	0	0	0
			1110	699	197	213	1			
29	DH	149	Total	C	N	O	S	0	0	0
			1110	699	197	213	1			

- Molecule 30 is a protein called 50S ribosomal protein L11.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
30	BI	141	Total	C	N	O	S	0	0	0
			1032	651	179	196	6			
30	DI	141	Total	C	N	O	S	0	0	0
			1032	651	179	196	6			

- Molecule 31 is a protein called 50S ribosomal protein L13.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
31	BJ	142	Total	C	N	O	S	0	0	0
			1129	714	212	199	4			
31	DJ	142	Total	C	N	O	S	0	0	0
			1129	714	212	199	4			

- Molecule 32 is a protein called 50S ribosomal protein L14.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
32	BK	122	Total	C	N	O	S	0	0	0
			939	587	180	166	6			
32	DK	122	Total	C	N	O	S	0	0	0
			939	587	180	166	6			

- Molecule 33 is a protein called 50S ribosomal protein L15.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
33	BL	143	Total	C	N	O	S	0	0	0
			1045	649	206	189	1			
33	DL	143	Total	C	N	O	S	0	0	0
			1045	649	206	189	1			

- Molecule 34 is a protein called 50S ribosomal protein L16.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
34	BM	136	Total	C	N	O	S	0	0	0
			1074	686	205	177	6			
34	DM	136	Total	C	N	O	S	0	0	0
			1074	686	205	177	6			

- Molecule 35 is a protein called 50S ribosomal protein L17.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
35	BN	120	Total	C	N	O	S	0	0	0
			961	593	196	167	5			
35	DN	120	Total	C	N	O	S	0	0	0
			961	593	196	167	5			

- Molecule 36 is a protein called 50S ribosomal protein L18.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
36	BO	116	Total	C	N	O	0	0	0
			892	552	178	162			
36	DO	116	Total	C	N	O	0	0	0
			892	552	178	162			

- Molecule 37 is a protein called 50S ribosomal protein L19.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
37	BP	114	Total	C	N	O	S	0	0	0
			917	574	179	163	1			
37	DP	114	Total	C	N	O	S	0	0	0
			917	574	179	163	1			

- Molecule 38 is a protein called 50S ribosomal protein L20.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
38	BQ	117	Total	C	N	O	S	0	0	0
			947	604	192	151				
38	DQ	117	Total	C	N	O	S	0	0	0
			947	604	192	151				

- Molecule 39 is a protein called 50S ribosomal protein L21.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
39	BR	103	Total	C	N	O	S	0	0	0
			816	516	153	145	2			
39	DR	103	Total	C	N	O	S	0	0	0
			816	516	153	145	2			

- Molecule 40 is a protein called 50S ribosomal protein L22.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
40	BS	110	Total	C	N	O	S	0	0	0
			857	532	166	156	3			
40	DS	110	Total	C	N	O	S	0	0	0
			857	532	166	156	3			

- Molecule 41 is a protein called 50S ribosomal protein L23.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
41	BT	93	Total	C	N	O	S	0	0	0
			739	466	139	132	2			
41	DT	93	Total	C	N	O	S	0	0	0
			739	466	139	132	2			

- Molecule 42 is a protein called 50S ribosomal protein L24.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
42	BU	102	Total	C	N	O	S	0	0	0
			780	492	146	142				

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Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
42	DU	102	Total	C	N	O			
			780	492	146	142	0	0	0

- Molecule 43 is a protein called 50S ribosomal protein L25.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
43	BV	94	Total	C	N	O	S			
			753	479	137	134	3	0	0	0
43	DV	94	Total	C	N	O	S			
			753	479	137	134	3	0	0	0

- Molecule 44 is a protein called 50S ribosomal protein L27.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
44	BW	76	Total	C	N	O	S			
			580	359	117	103	1	0	0	0
44	DW	75	Total	C	N	O	S			
			569	353	113	102	1	0	0	0

- Molecule 45 is a protein called 50S ribosomal protein L28.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
45	BX	77	Total	C	N	O	S			
			625	388	129	106	2	0	0	0
45	DX	77	Total	C	N	O	S			
			625	388	129	106	2	0	0	0

- Molecule 46 is a protein called 50S ribosomal protein L29.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
46	BY	63	Total	C	N	O	S			
			509	313	99	95	2	0	0	0
46	DY	63	Total	C	N	O	S			
			509	313	99	95	2	0	0	0

- Molecule 47 is a protein called 50S ribosomal protein L30.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
47	BZ	58	Total	C	N	O	S			
			449	281	87	79	2	0	0	0
47	DZ	58	Total	C	N	O	S			
			449	281	87	79	2	0	0	0

- Molecule 48 is a protein called 50S ribosomal protein L32.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
48	B0	56	Total	C	N	O	S	0	0	0
			444	269	94	80	1			
48	D0	56	Total	C	N	O	S	0	0	0
			444	269	94	80	1			

- Molecule 49 is a protein called 50S ribosomal protein L33.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
49	B1	50	Total	C	N	O	0	0	0
			410	263	75	72			
49	D1	50	Total	C	N	O	0	0	0
			410	263	75	72			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
50	B2	46	Total	C	N	O	S	0	0	0
			377	228	90	57	2			
50	D2	46	Total	C	N	O	S	0	0	0
			377	228	90	57	2			

- Molecule 51 is a protein called 50S ribosomal protein L35.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
51	B3	64	Total	C	N	O	S	0	0	0
			504	323	105	74	2			
51	D3	64	Total	C	N	O	S	0	0	0
			504	323	105	74	2			

- Molecule 52 is a protein called 50S ribosomal protein L36.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
52	B4	38	Total	C	N	O	S	0	0	0
			302	185	65	48	4			
52	D4	38	Total	C	N	O	S	0	0	0
			302	185	65	48	4			

- Molecule 53 is a protein called 50S ribosomal protein L1.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
53	B5	191	Total	C	N	O	0	0	1
			1142	691	221	230			

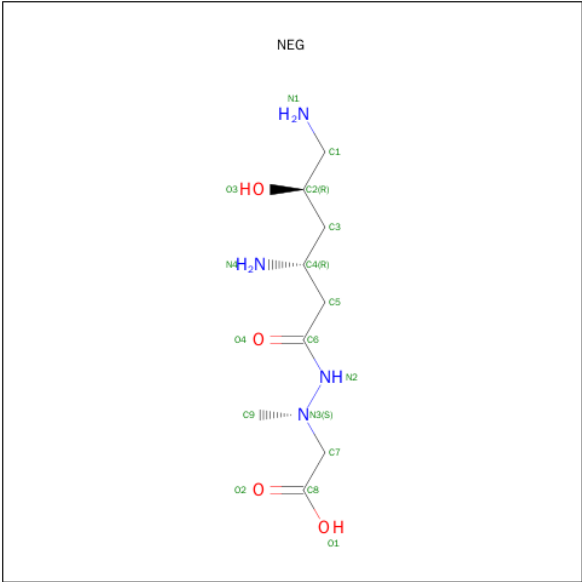
- Molecule 54 is MAGNESIUM ION (three-letter code: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
54	BB	4	Total	Mg	0	0
			4	4		
54	DQ	1	Total	Mg	0	0
			1	1		
54	BA	194	Total	Mg	0	0
			194	194		
54	CA	56	Total	Mg	0	0
			56	56		
54	CT	1	Total	Mg	0	0
			1	1		
54	DL	2	Total	Mg	0	0
			2	2		
54	D2	1	Total	Mg	0	0
			1	1		
54	AA	71	Total	Mg	0	0
			71	71		
54	BQ	1	Total	Mg	0	0
			1	1		
54	DA	164	Total	Mg	0	0
			164	164		
54	DB	3	Total	Mg	0	0
			3	3		
54	AM	1	Total	Mg	0	0
			1	1		

- Molecule 55 is ZINC ION (three-letter code: ZN) (formula: Zn).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
55	B4	1	Total	Zn	0	0
			1	1		
55	D4	1	Total	Zn	0	0
			1	1		

- Molecule 56 is NEGAMYCIN (three-letter code: NEG) (formula: C₉H₂₀N₄O₄).



Mol	Chain	Residues	Atoms				ZeroOcc	AltConf
56	CA	1	Total	C	N	O	0	0
			17	9	4	4		

• Molecule 57 is water.

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	AA	196	Total	O	0	0
			196	196		
57	AE	1	Total	O	0	0
			1	1		
57	AL	1	Total	O	0	0
			1	1		
57	AN	3	Total	O	0	0
			3	3		
57	AT	1	Total	O	0	0
			1	1		
57	AU	1	Total	O	0	0
			1	1		
57	BA	620	Total	O	0	0
			620	620		
57	BB	14	Total	O	0	0
			14	14		
57	BC	10	Total	O	0	0
			10	10		
57	BD	4	Total	O	0	0
			4	4		
57	BF	1	Total	O	0	0
			1	1		

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Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
57	BG	1	Total O 1 1	0	0
57	BL	6	Total O 6 6	0	0
57	BN	3	Total O 3 3	0	0
57	BS	1	Total O 1 1	0	0
57	B2	1	Total O 1 1	0	0
57	B3	3	Total O 3 3	0	0
57	B4	2	Total O 2 2	0	0
57	CA	186	Total O 186 186	0	0
57	CL	1	Total O 1 1	0	0
57	CN	3	Total O 3 3	0	0
57	CT	3	Total O 3 3	0	0
57	CU	1	Total O 1 1	0	0
57	DA	611	Total O 611 611	0	0
57	DB	13	Total O 13 13	0	0
57	DC	8	Total O 8 8	0	0
57	DD	3	Total O 3 3	0	0
57	DE	5	Total O 5 5	0	0
57	DJ	1	Total O 1 1	0	0
57	DL	4	Total O 4 4	0	0
57	DN	1	Total O 1 1	0	0
57	DS	1	Total O 1 1	0	0

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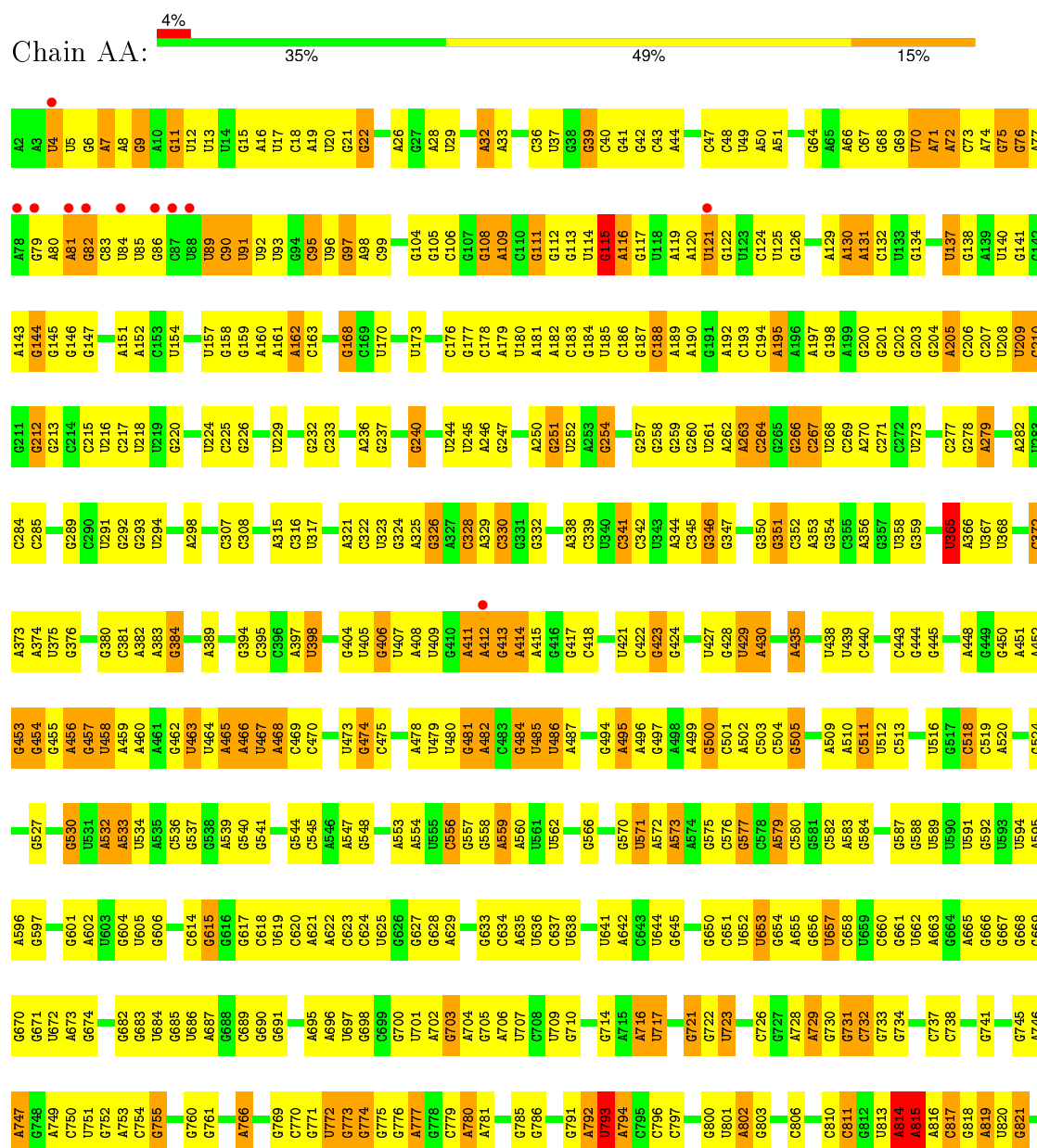
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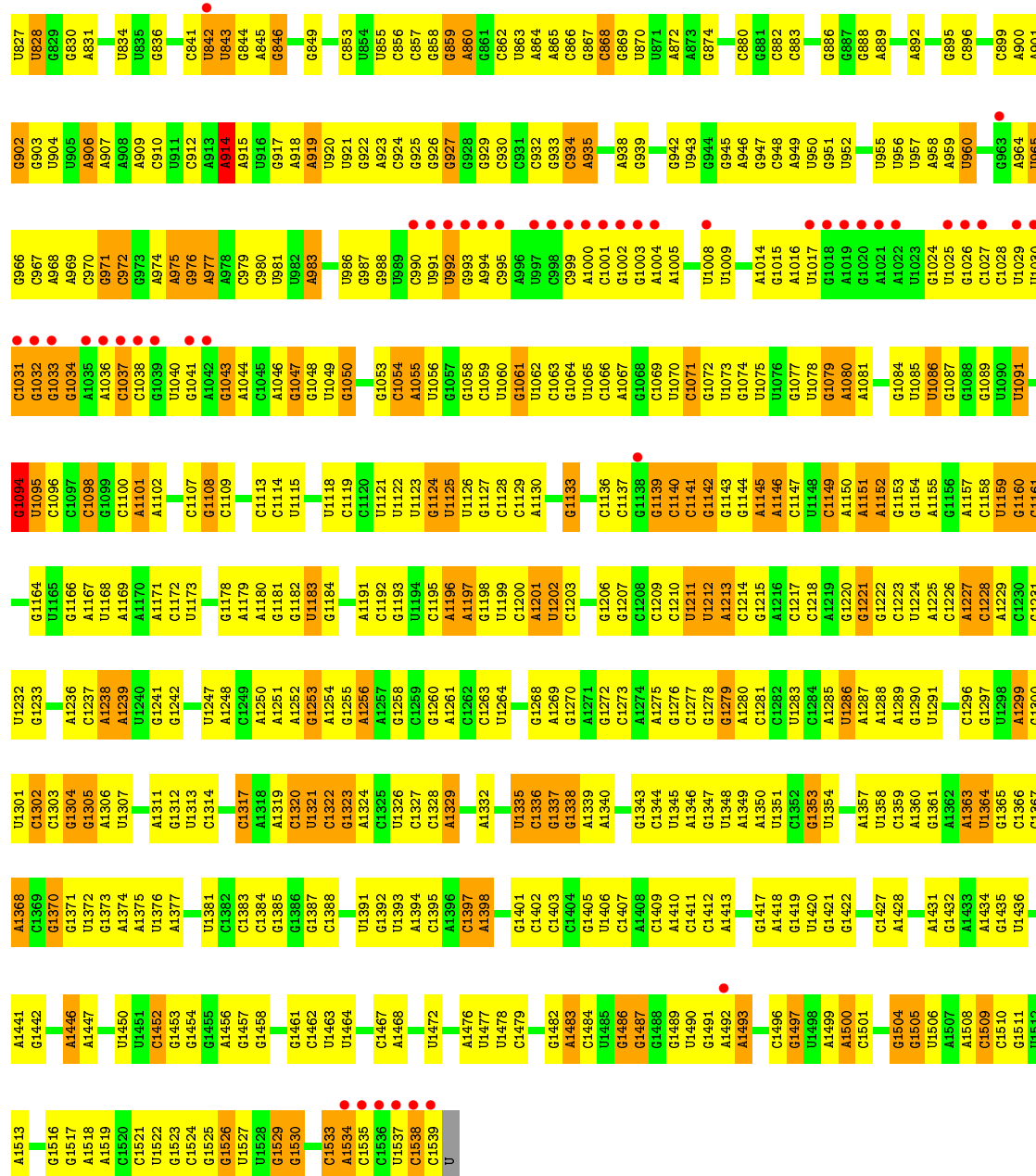
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
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57	DU	1	Total 1	O 1	0	0
57	DV	1	Total 1	O 1	0	0
57	D2	1	Total 1	O 1	0	0
57	D3	2	Total 2	O 2	0	0
57	D4	1	Total 1	O 1	0	0

3 Residue-property plots

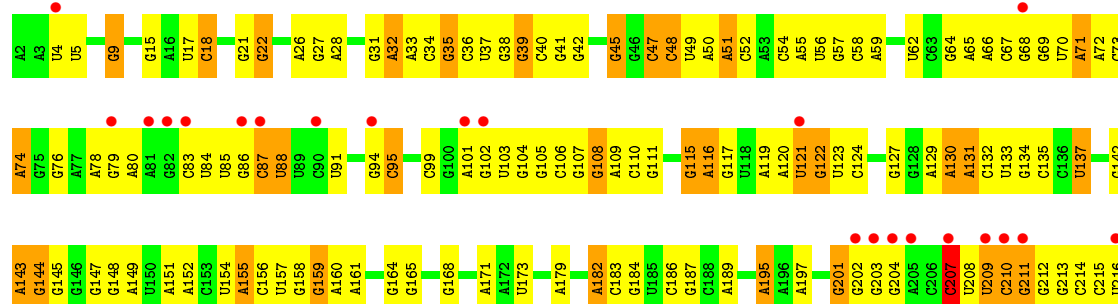
These plots are drawn for all protein, RNA and DNA chains in the entry. The first graphic for a chain summarises the proportions of errors displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and electron density. Residues are color-coded according to the number of geometric quality criteria for which they contain at least one outlier: green = 0, yellow = 1, orange = 2 and red = 3 or more. A red dot above a residue indicates a poor fit to the electron density ($RSRZ > 2$). 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: 16S rRNA

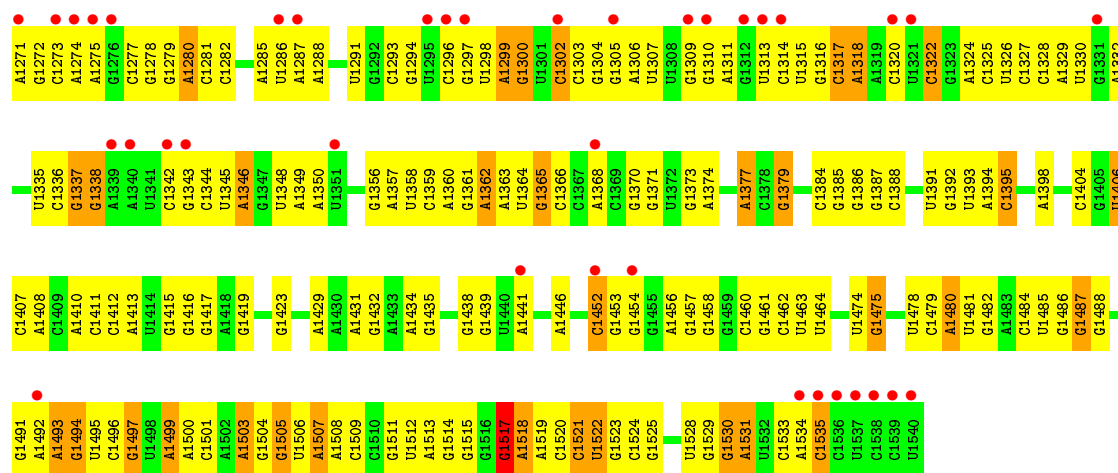




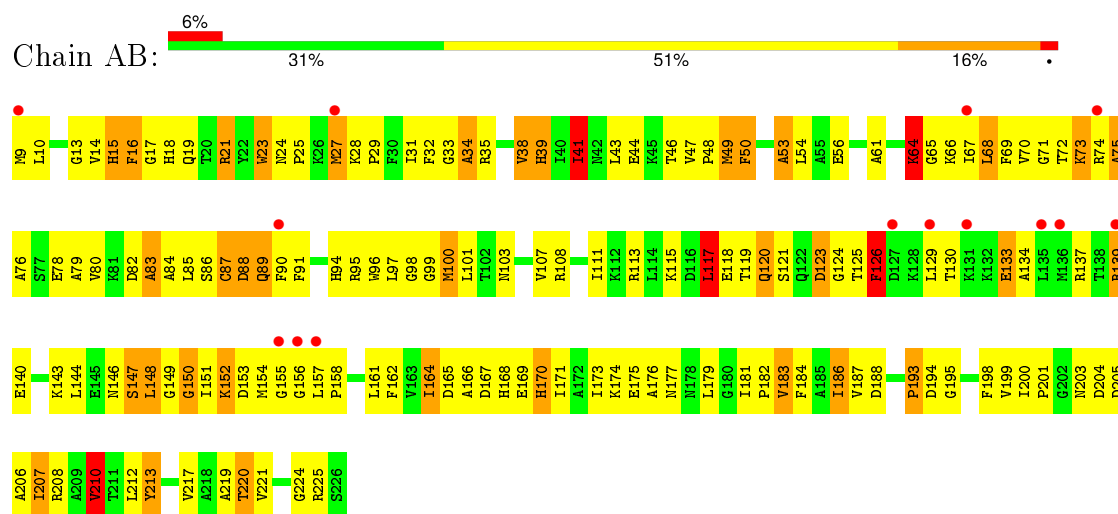
• Molecule 1: 16S rRNA



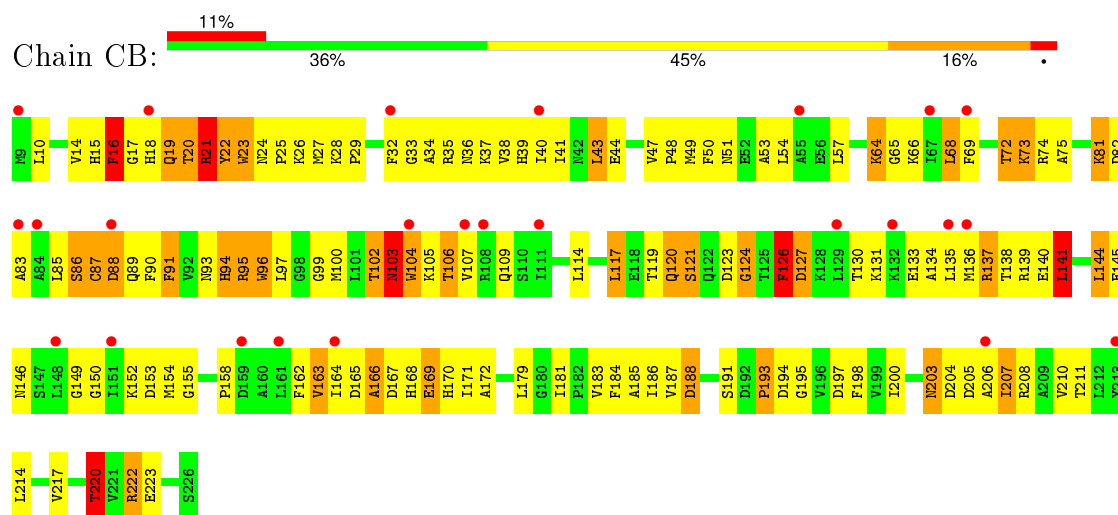




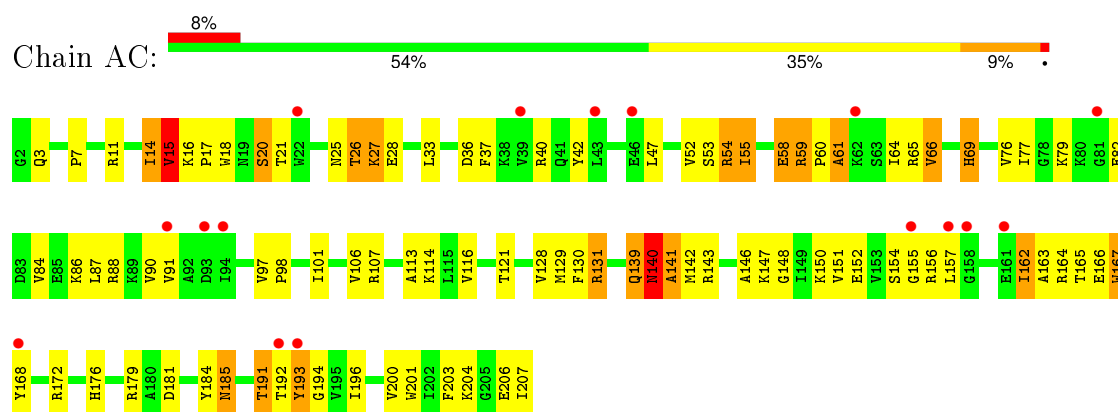
• Molecule 2: 30S ribosomal protein S2



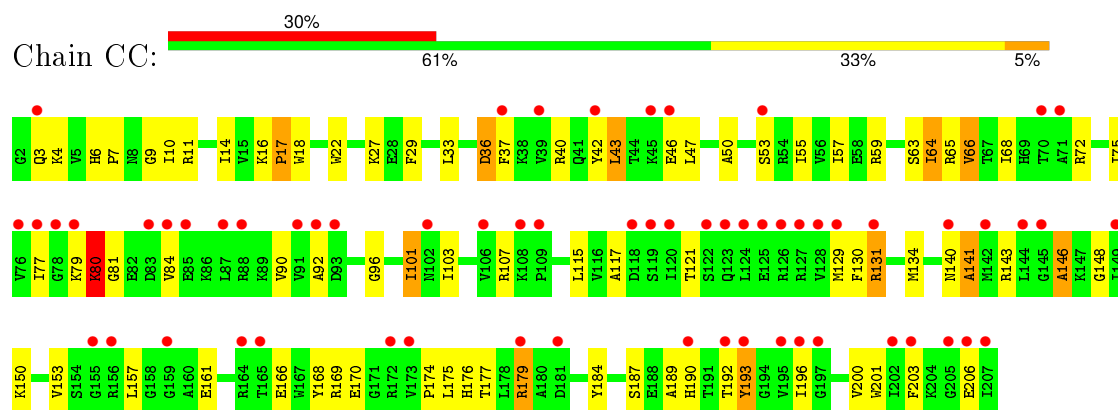
• Molecule 2: 30S ribosomal protein S2



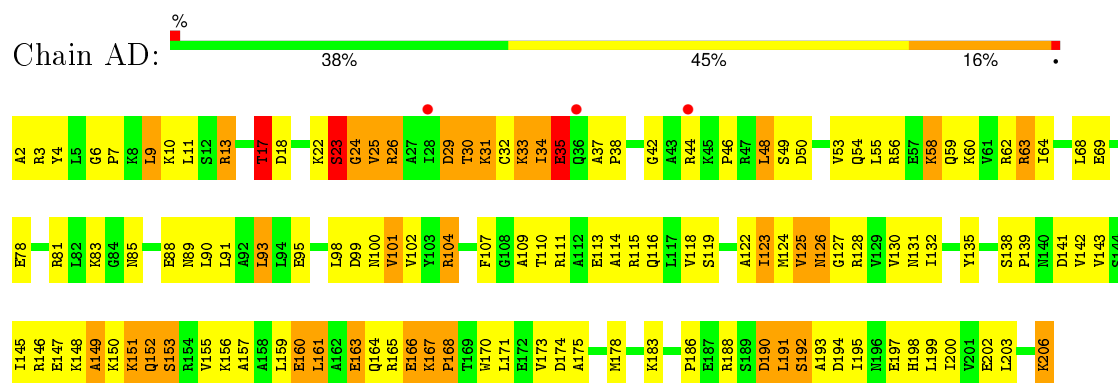
• Molecule 3: 30S ribosomal protein S3



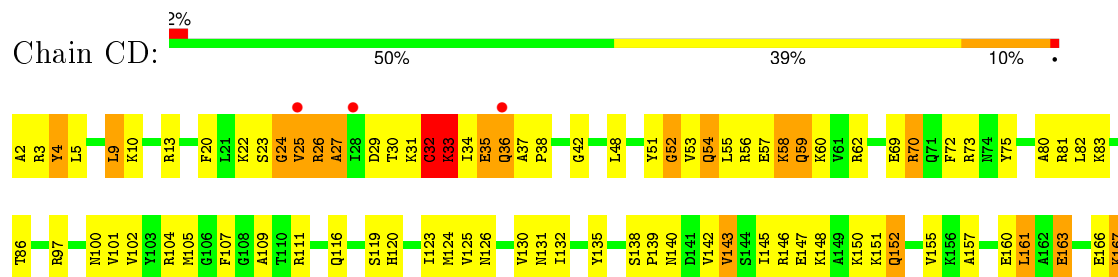
- Molecule 3: 30S ribosomal protein S3



- Molecule 4: 30S ribosomal protein S4

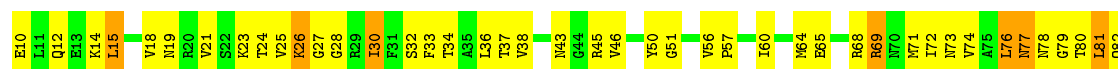
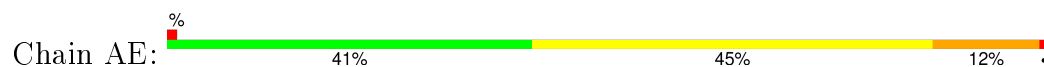


- Molecule 4: 30S ribosomal protein S4





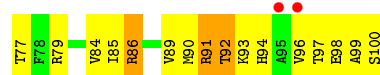
- Molecule 5: 30S ribosomal protein S5



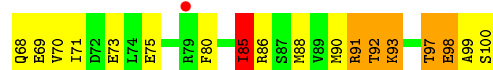
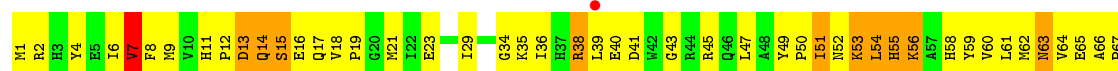
- Molecule 5: 30S ribosomal protein S5



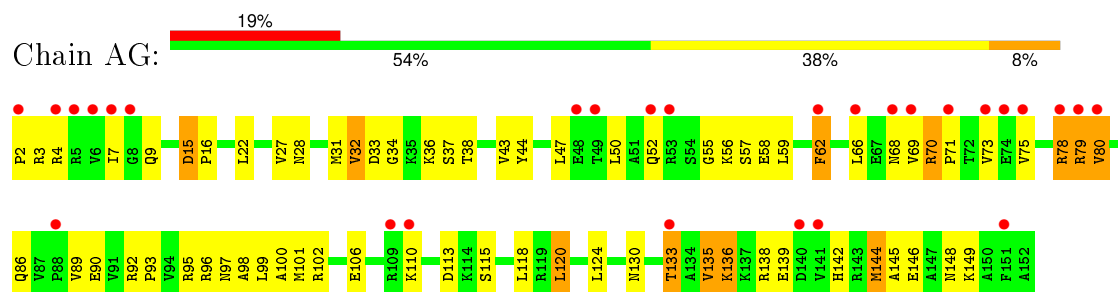
- Molecule 6: 30S ribosomal protein S6



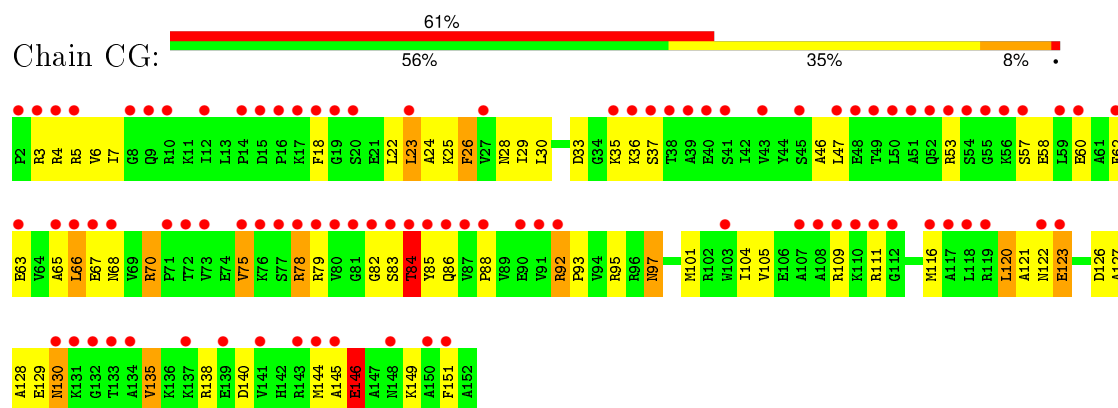
- Molecule 6: 30S ribosomal protein S6



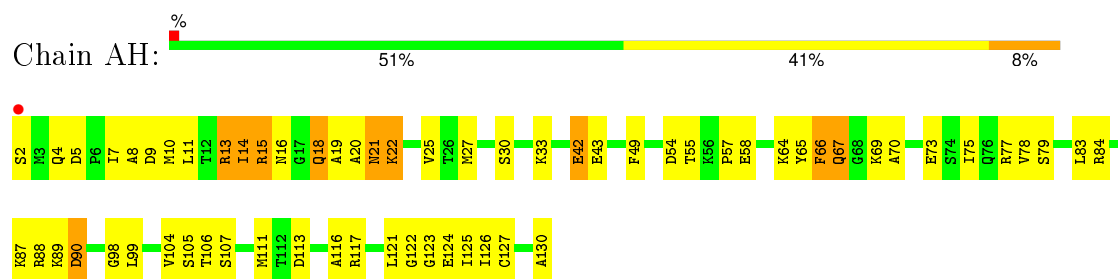
- Molecule 7: 30S ribosomal protein S7



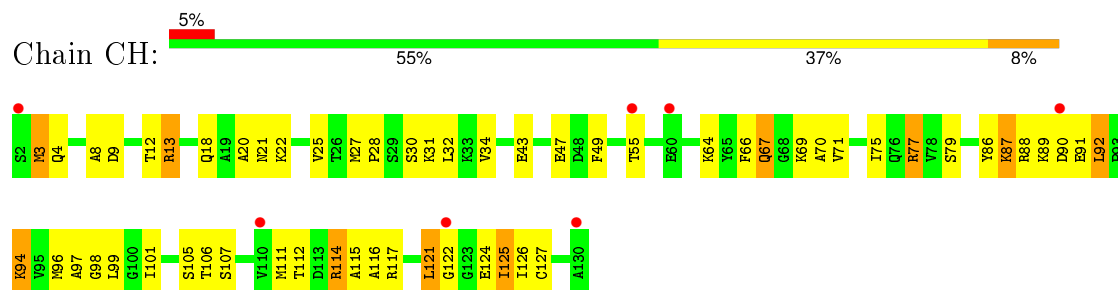
- Molecule 7: 30S ribosomal protein S7



- Molecule 8: 30S ribosomal protein S8

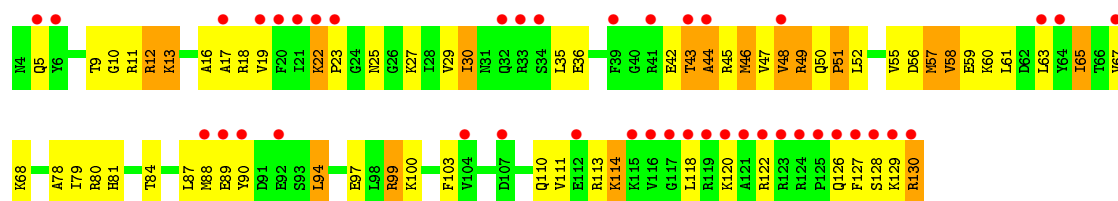


- Molecule 8: 30S ribosomal protein S8

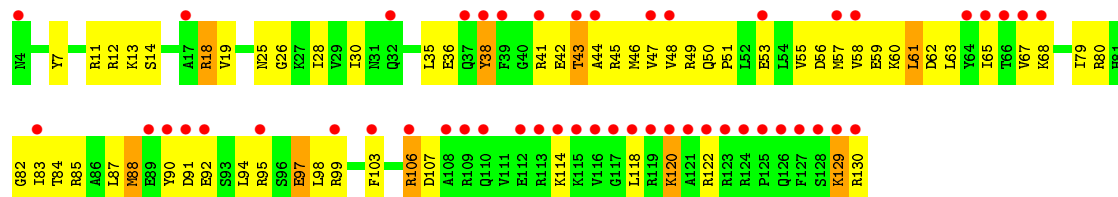
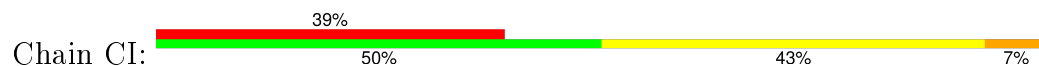


- Molecule 9: 30S ribosomal protein S9

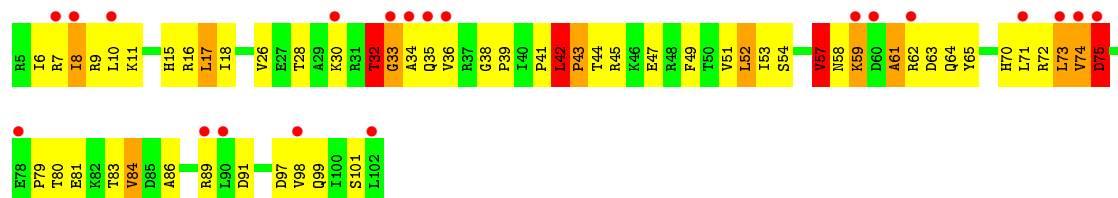
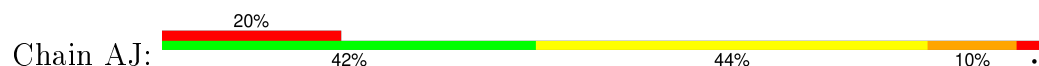




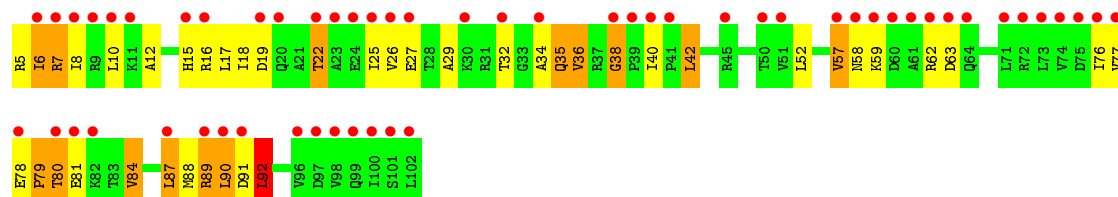
• Molecule 9: 30S ribosomal protein S9



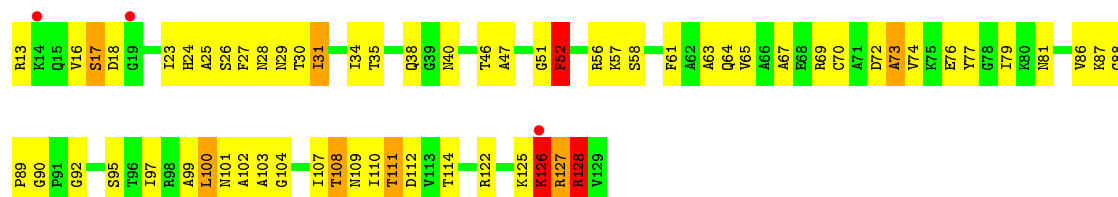
• Molecule 10: 30S ribosomal protein S10



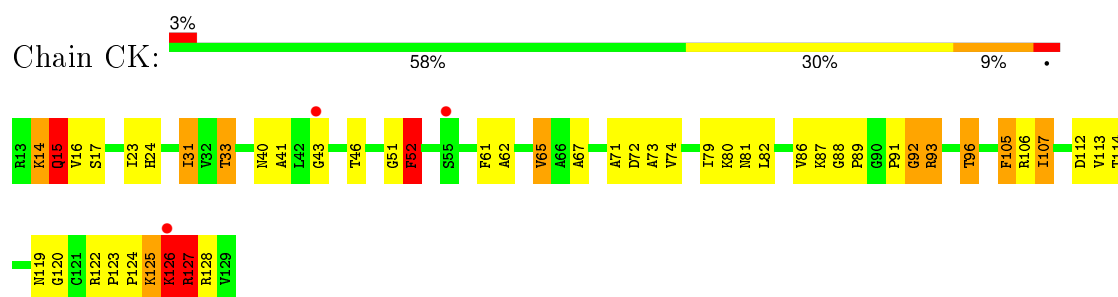
• Molecule 10: 30S ribosomal protein S10



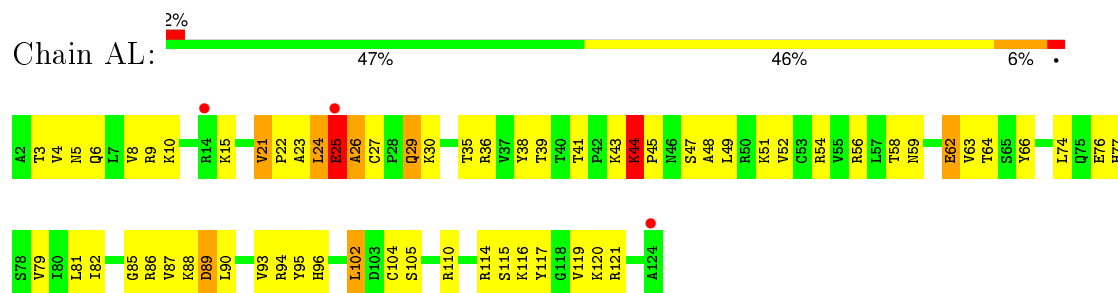
• Molecule 11: 30S ribosomal protein S11



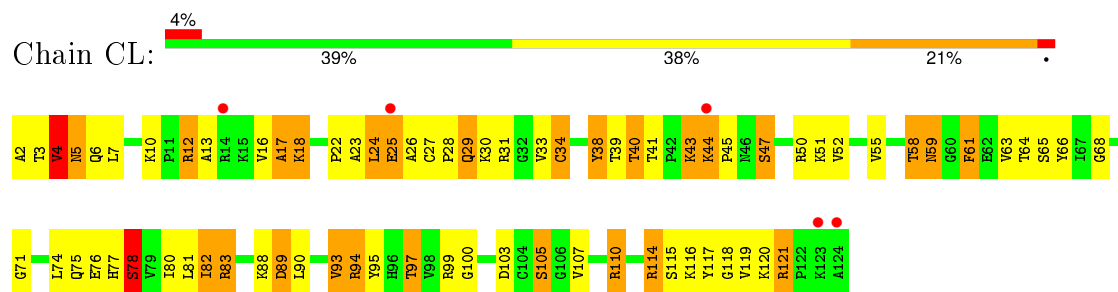
• Molecule 11: 30S ribosomal protein S11



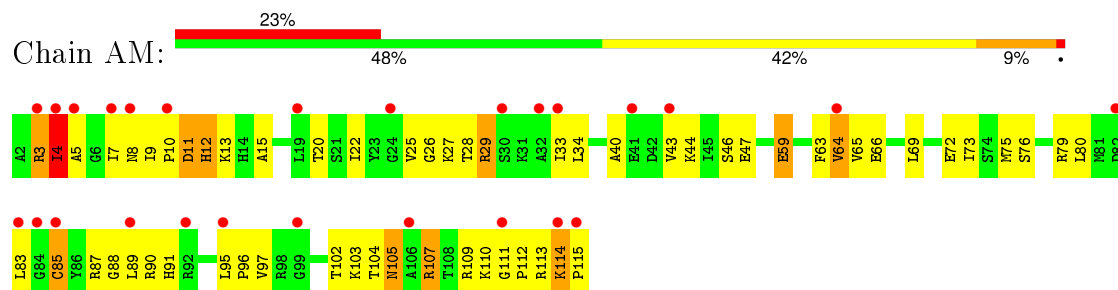
• Molecule 12: 30S ribosomal protein S12



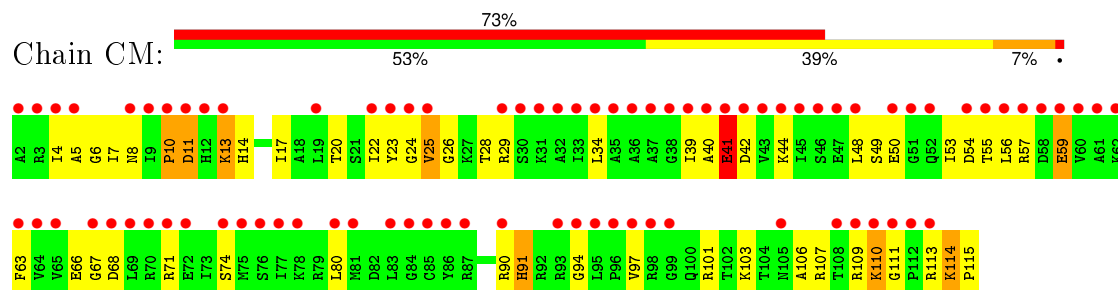
• Molecule 12: 30S ribosomal protein S12



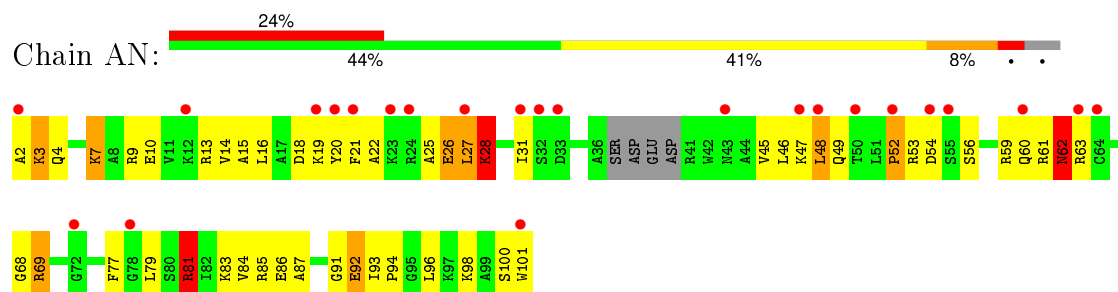
• Molecule 13: 30S ribosomal protein S13



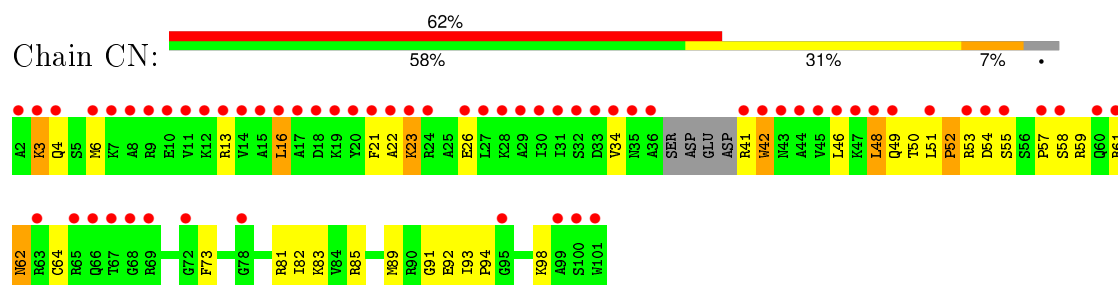
• Molecule 13: 30S ribosomal protein S13



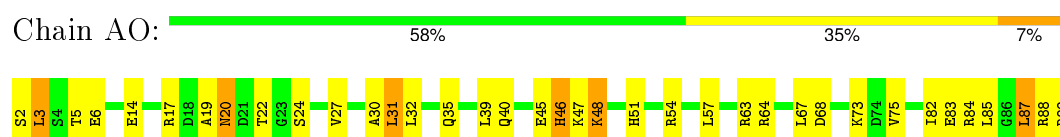
- Molecule 14: 30S ribosomal protein S14



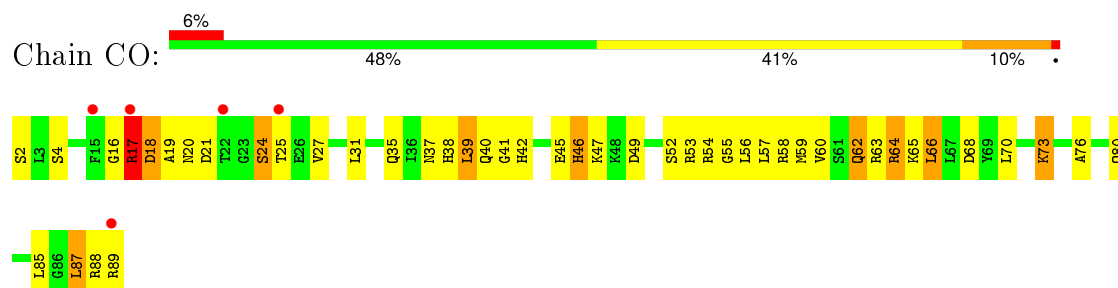
- Molecule 14: 30S ribosomal protein S14



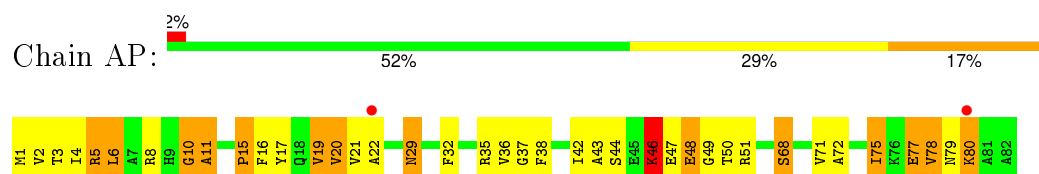
- Molecule 15: 30S ribosomal protein S15



- Molecule 15: 30S ribosomal protein S15

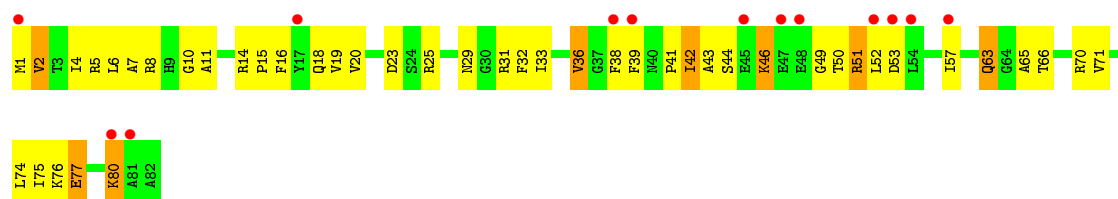


- Molecule 16: 30S ribosomal protein S16

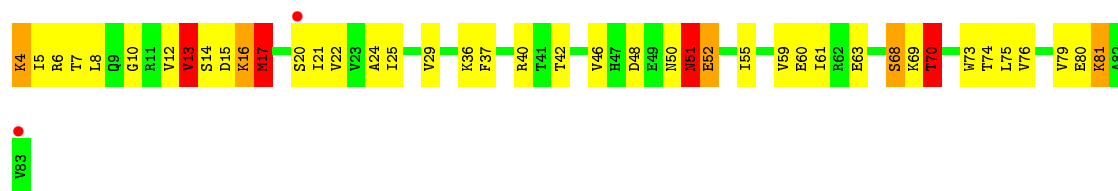


- Molecule 16: 30S ribosomal protein S16

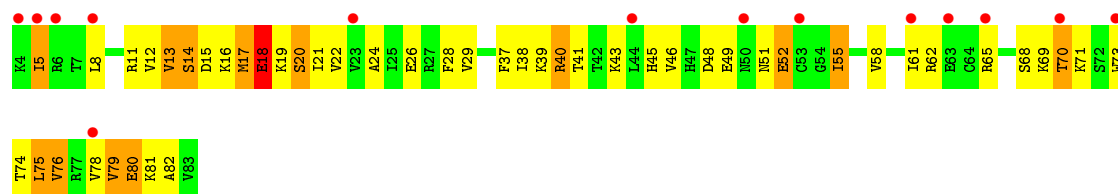




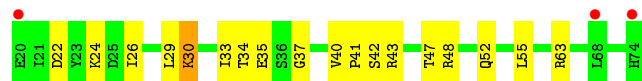
- Molecule 17: 30S ribosomal protein S17



- Molecule 17: 30S ribosomal protein S17



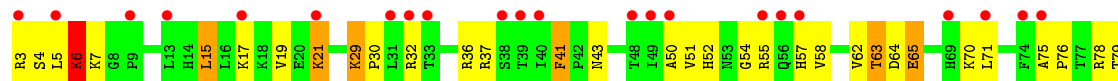
- Molecule 18: 30S ribosomal protein S18



- Molecule 18: 30S ribosomal protein S18

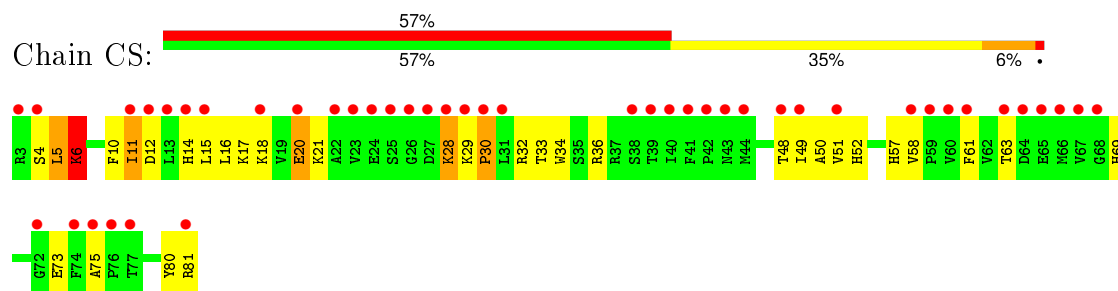


- Molecule 19: 30S ribosomal protein S19

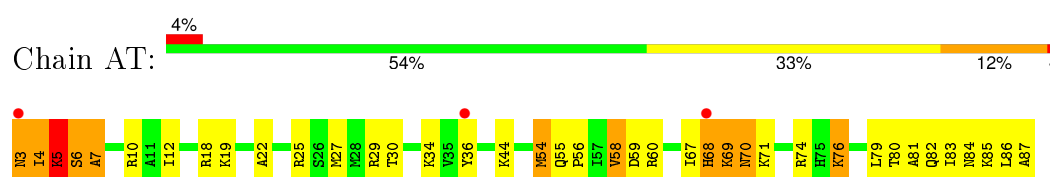


Y80
R81

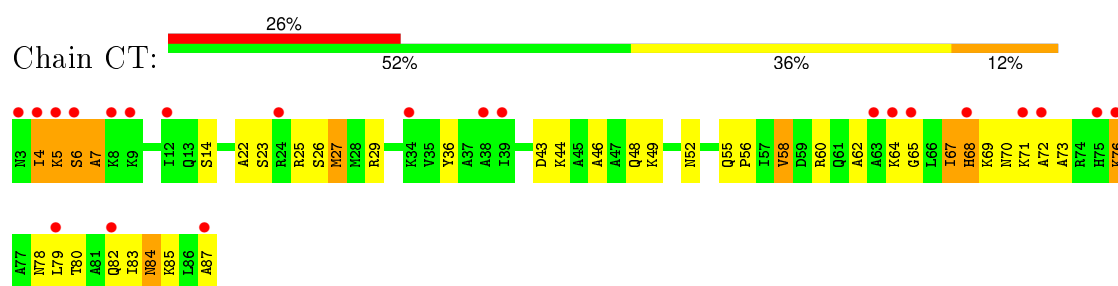
- Molecule 19: 30S ribosomal protein S19



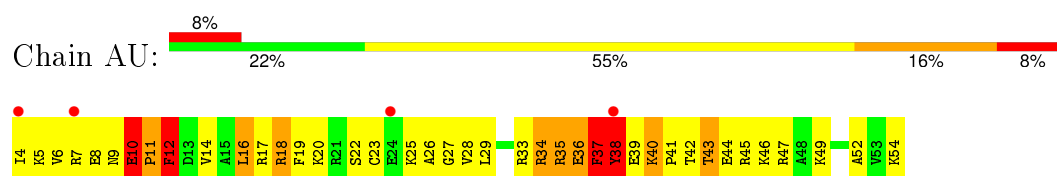
- Molecule 20: 30S ribosomal protein S20



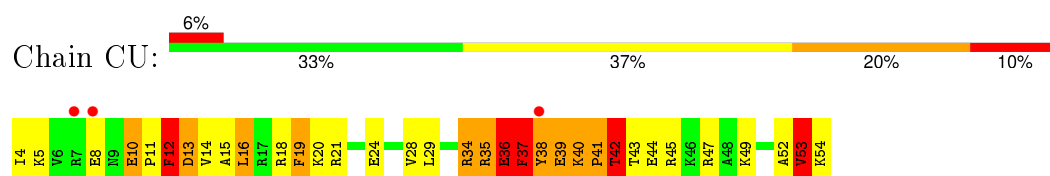
- Molecule 20: 30S ribosomal protein S20



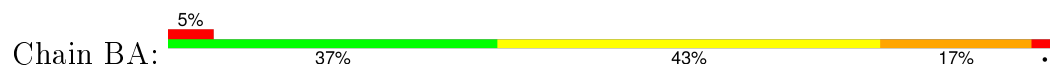
- Molecule 21: 30S ribosomal protein S21



- Molecule 21: 30S ribosomal protein S21

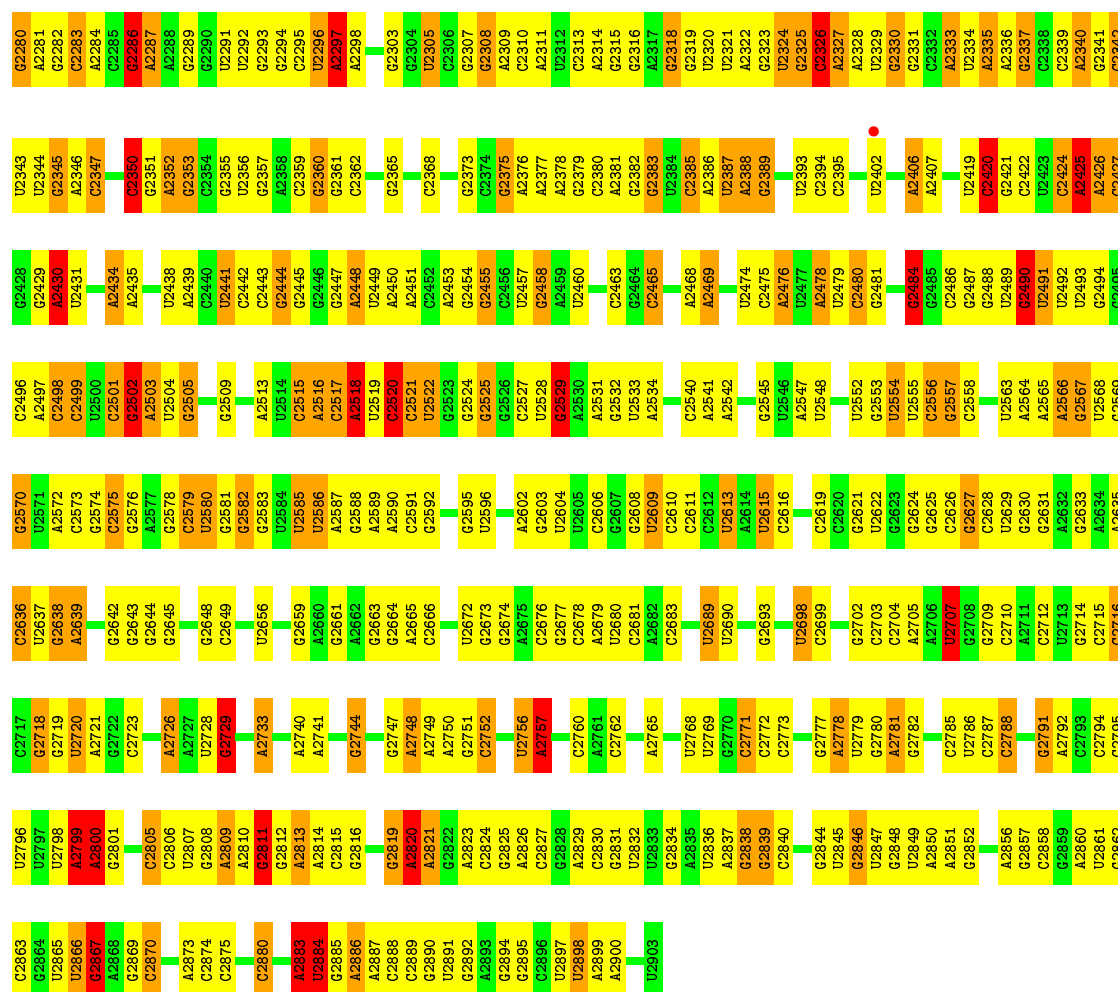


- Molecule 22: 23S rRNA

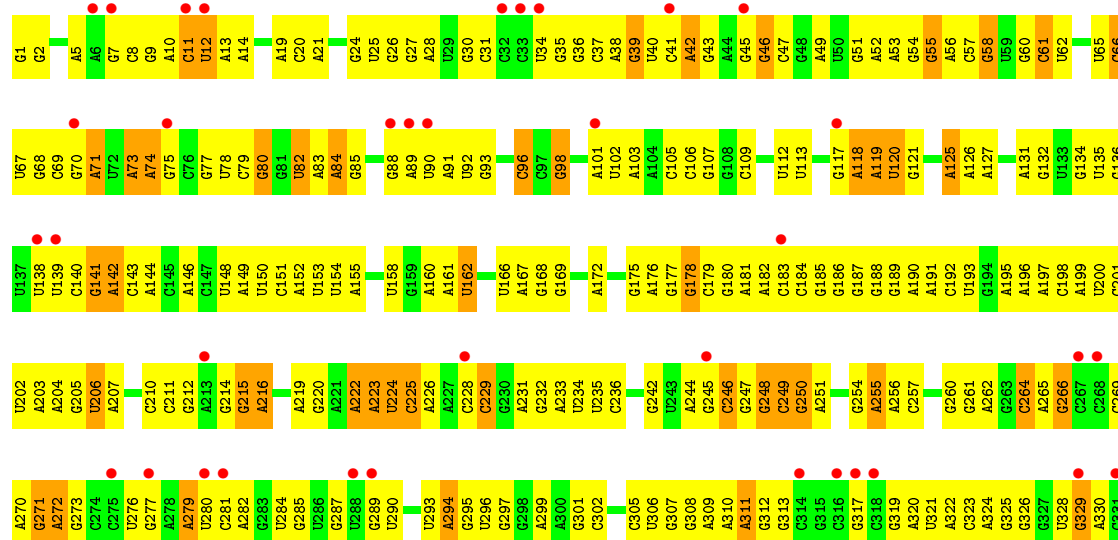


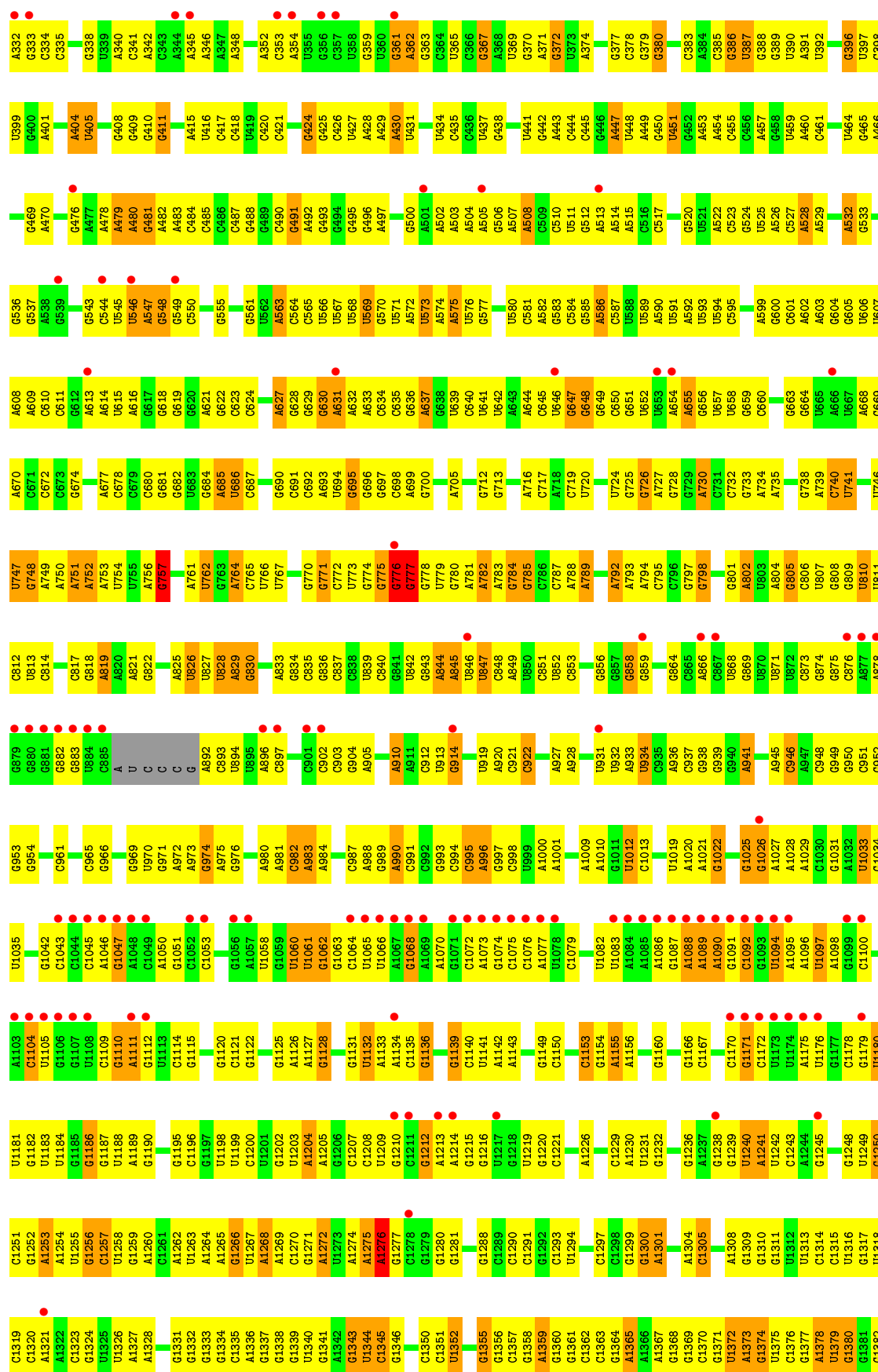
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C1109	G907	G971	G907	A844	A782	A716	B636	U569	A503	A256	A340	A255	U170	G80	G7
G1110	C1045	A973	C908	A845	A783	A717	A637	G570	A504	G424	C341	C257	U171	G81	G8
A1111	A1046	A974	A909	U846	G784	C717	U638	U571	G506	C440	A342	C258	A176	A82	G9
G1112	G1047	G974	A910	U847	G785	A722	U639	A572	G507	U441	C343	G258	G177	A84	A10
U1113	A1050	A975	U913	U850	C786	A723	C640	U573	A508	G442	A345	G259	G178	C11	U12
G1114	G976	G977	G914	C851	A787	U724	A644	A574	C509	C445	A346	G261	C179	G85	U11
G1116	G978	G978	C915	U852	A789	G725	C645	U576	C510	C446	G350	G265	G180	A91	A13
C1121	A1058	A980	A917	C854	A790	G726	U646	G577	U511	G447	C351	G266	A181	A91	A14
G1122	U1059	A981	A918	G855	A792	A727	G647	G578	G512	U448	A352	G270	A182	U92	G15
C1123	U1060	C982	U919	G856	A793	G728	A654	U580	A514	A449	C353	G271	C183	G93	G16
G1124	U1061	A983	A920	G857	A794	G729	A655	C581	A515	G450	C357	A272	C184	U96	G17
G1125	G1062	A984	C921	G858	C795	A730	U656	A575	C516	U451	U358	G273	G185	A19	U18
A1126	C985	C985	C922	G859	C796	G732	U657	G583	C517	G452	U359	G274	G186	C96	C20
A1127	C986	C986	C923	U860	G797	G733	U658	G584	C518	A453	G360	G275	G187	G98	G21
G1128	U1065	C987	G924	A861	G798	A734	U659	G585	U519	A454	U361	G276	G189	A20	A21
A1129	U1066	A988	G927	G862	A799	A735	G664	A586	G520	C455	G362	A278	A190	A101	C22
U1130	A1067	G989	A927	A863	A800	G738	U665	U592	A526	G463	A367	G289	A191	U102	G26
G1131	G1068	A990	A928	G864	G801	A739	U666	U593	C527	U464	A368	U290	A192	U103	G27
U1132	A1069	C991	U929	C865	A802	A740	A667	U594	A528	A466	U369	G291	A193	A104	A28
A1133	A1070	C992	G930	A866	U803	C741	U667	G595	A529	G467	G370	G292	U193	U34	U29
G1134	G1071	G993	U931	C867	A804	U742	A668	U596	G530	C462	A371	G293	A195	G108	G30
C1135	C1072	C994	U932	U868	G805	A743	G669	A597	C531	G463	G372	G294	A196	C109	G31
G1136	A1073	C995	A933	G869	C806	A744	U670	U598	A532	U464	U373	G301	A197	C32	G32
U1137	G1074	A996	U934	U870	A807	U745	A671	G600	G533	G473	A374	G302	A198	C33	G33
G1138	C1075	G997	C935	U871	U807	G746	U672	C601	U535	C474	C378	G307	A199	U34	U34
G1139	C1076	C998	A936	U872	U808	U747	C673	U596	G536	G476	G379	G308	A125	A44	A44
U1140	A1077	U999	C937	G875	A819	U748	U674	G597	C537	A477	G380	A309	A126	G45	G45
C1141	U1078	A1000	G938	C876	U811	G748	A675	U598	A538	A478	G381	A310	A127	G46	G46
A1142	C1079	A1001	U941	A877	C812	A749	U676	U599	G543	A479	G386	G312	G215	C47	C47
G1143	U1080	G1002	G942	A878	U813	A750	A677	G600	C544	G481	U387	G313	G216	G48	G48
A1144	U1081	G1003	G943	G879	C814	A751	C678	C601	U546	A482	A391	G314	A217	C57	C57
C1145	U1082	U1004	A943	G880	C815	A752	U678	A602	U547	A483	U392	G315	A218	G58	G58
G1146	U1083	C1005	C944	G881	C817	A753	C680	A603	A548	C484	U393	G316	A219	G59	G59
A1084	A1084	C1006	A945	G882	C818	U754	C681	G604	G549	C485	C394	G317	A221	G60	G60
A1085	A1085	C1007	C946	G883	A819	U755	U682	U595	U554	C486	U395	G318	A222	C61	C61
A1086	A1086	A1008	A947	U884	A820	U756	G683	G605	G555	C487	G396	G319	A223	U62	U62
G1087	A1087	A1009	C948	C885	A821	A757	U684	U596	A556	G488	G397	G320	A224	A63	A63
A1088	A1088	G949	G950	A	C823	G758	A685	A608	C557	G489	G400	G321	C225	A64	A64
A1090	A1090	C951	C951	U	U824	G759	U686	A609	C558	C491	A401	G322	A226	U65	U65
G1091	C1092	G952	G952	C	A825	G760	C687	C611	C559	A492	G404	G323	A227	C66	C66
G1093	G1093	G953	G953	G	U826	A761	C687	G612	U562	G493	U405	G324	C240	C69	C69
A1094	A1094	U955	U955	A830	U828	U762	C691	A613	G549	C494	U406	G325	G245	A71	A71
A1095	A1095	G1023	G956	C893	A829	A763	C692	U615	U554	G495	A404	G326	G246	U72	U72
A1096	A1096	G1024	C957	U894	G831	G765	U694	U616	G555	C496	U405	G327	G247	A73	A73
U1097	U1097	G1025	U958	U895	U832	U767	G696	A621	A556	C497	G401	G328	G248	A74	A74
G1102	G1102	G1026	G959	U896	A833	G768	G697	G622	C557	A491	A401	G329	G249	U158	U158
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		U1035	G904	G904	G841	G708			U566	G500	U419	C337	G250		

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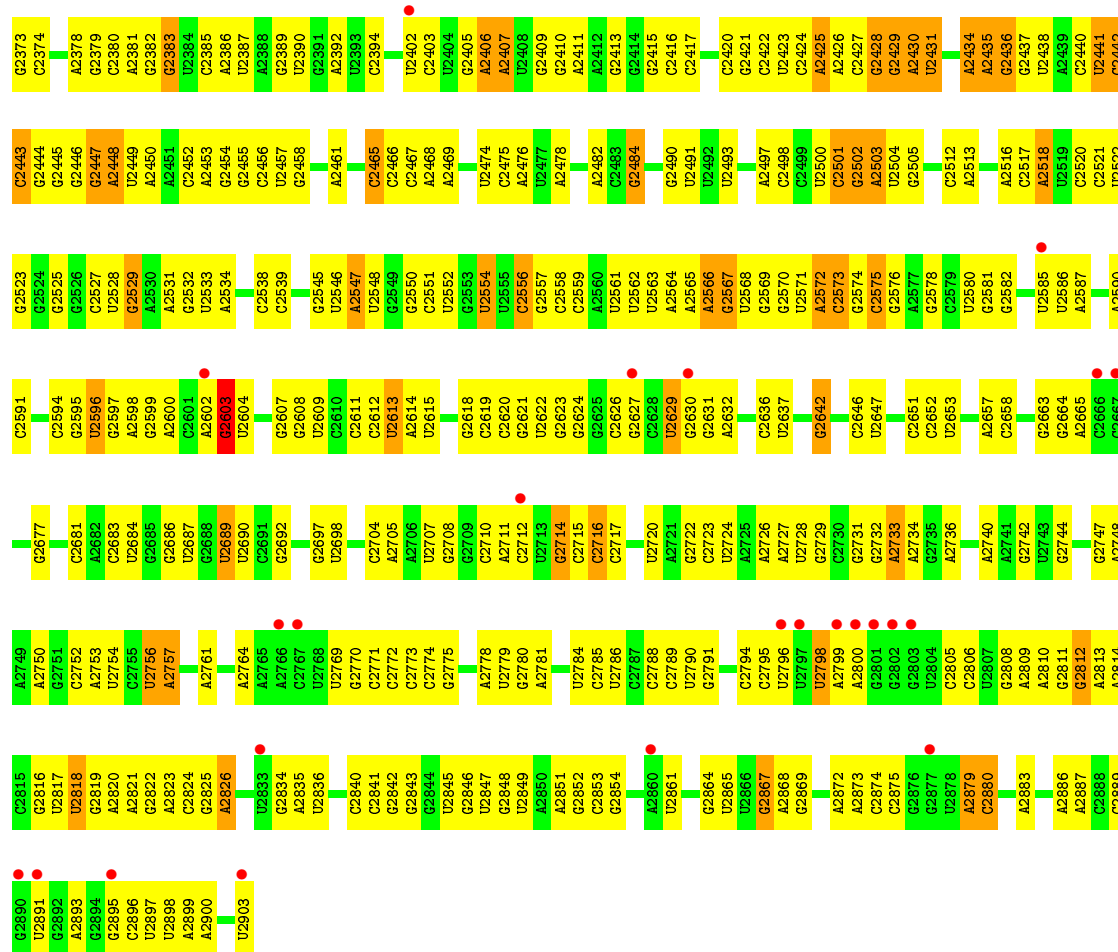


• Molecule 22: 23S rRNA



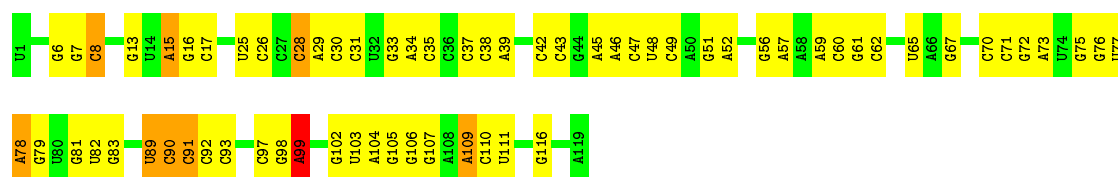


WORLDWIDE
PDB
PROTEIN DATA BANK



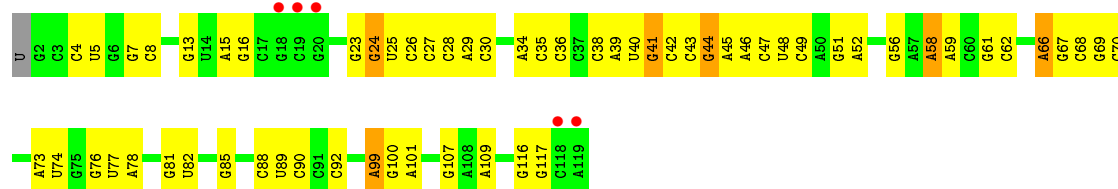
• Molecule 23: 5S rRNA

Chain BB: 45% 48% 7% •

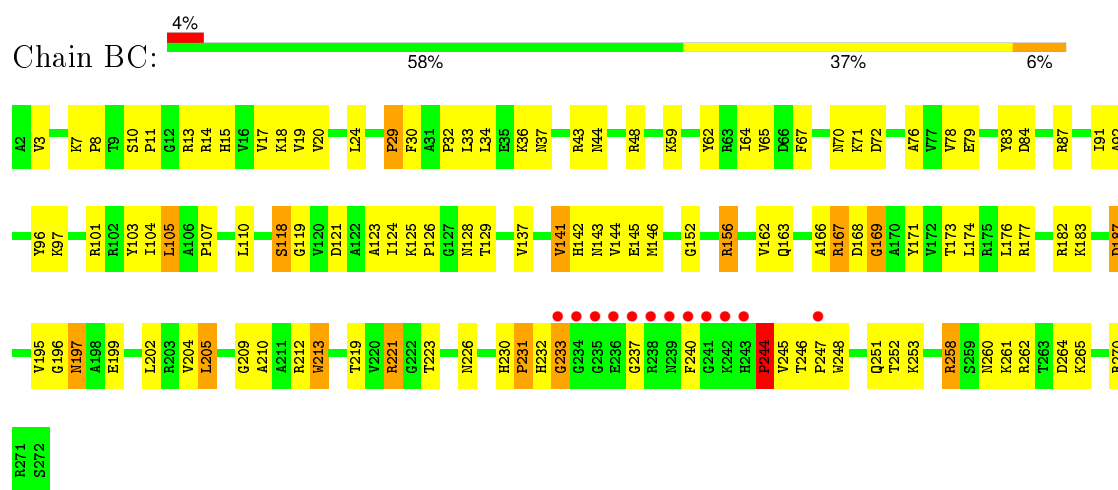


• Molecule 23: 5S rRNA

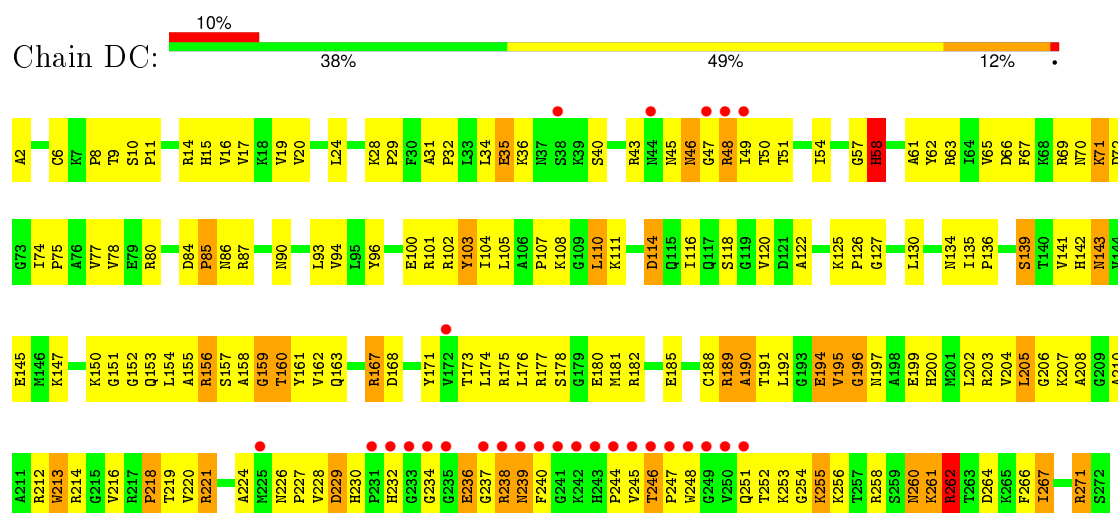
Chain DB: 48% 46% 5% •



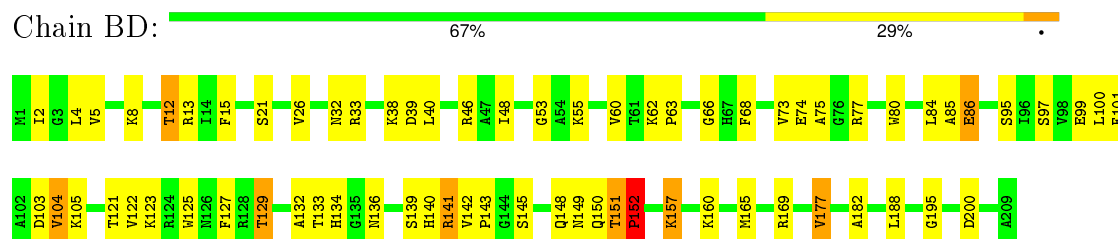
• Molecule 24: 50S ribosomal protein L2



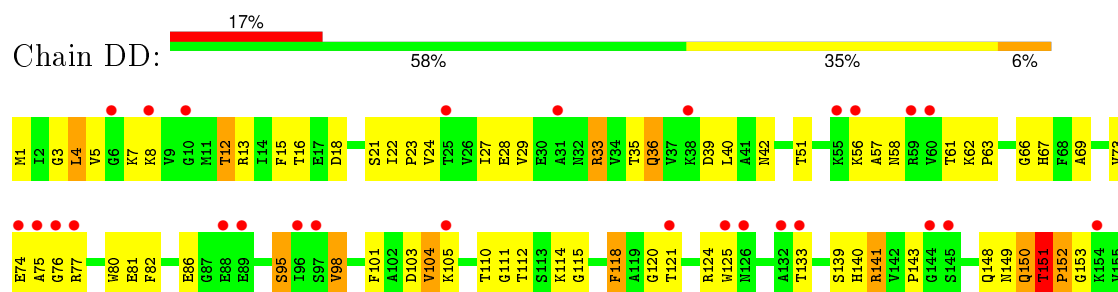
• Molecule 24: 50S ribosomal protein L2

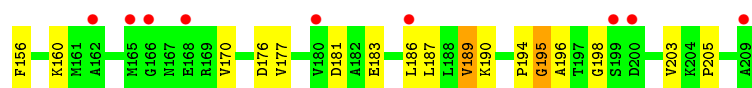


• Molecule 25: 50S ribosomal protein L3

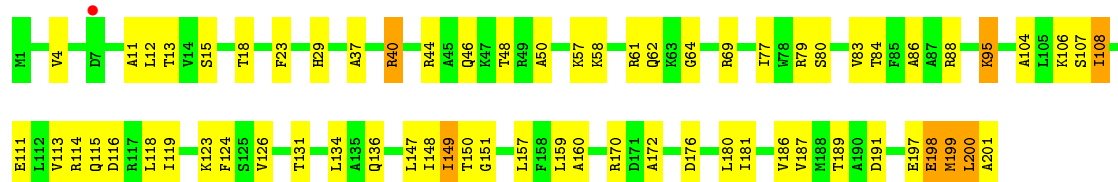


• Molecule 25: 50S ribosomal protein L3

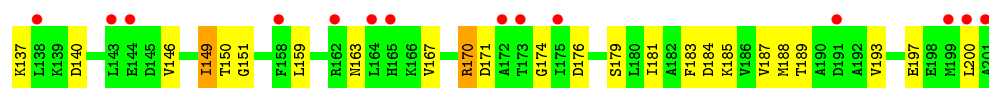
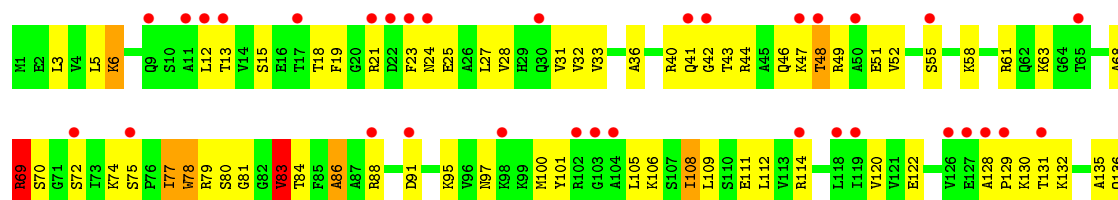




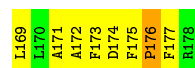
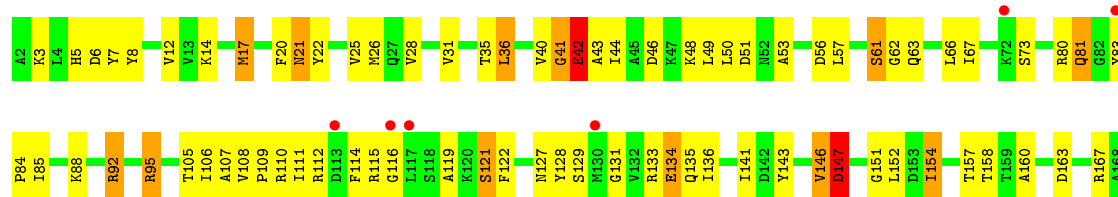
- Molecule 26: 50S ribosomal protein L4



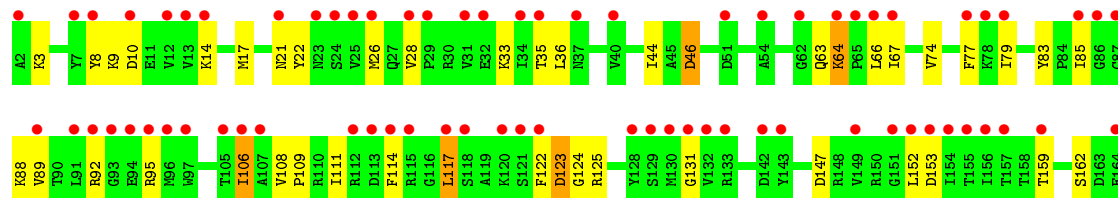
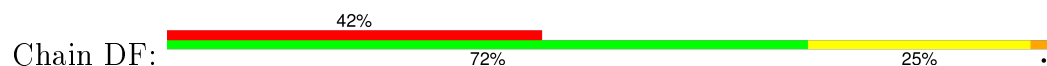
- Molecule 26: 50S ribosomal protein L4

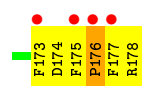


- Molecule 27: 50S ribosomal protein L5

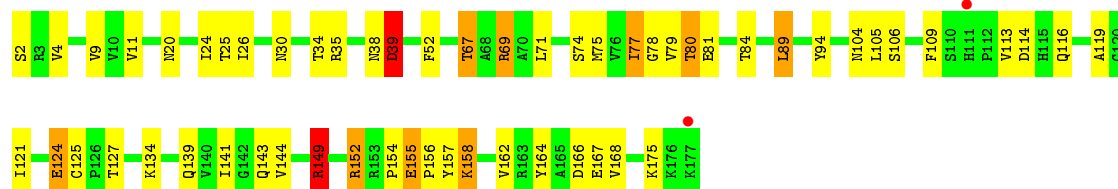


- Molecule 27: 50S ribosomal protein L5

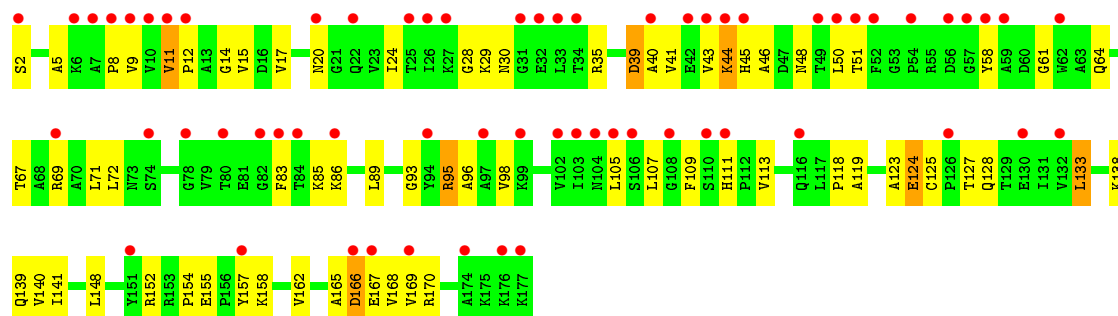




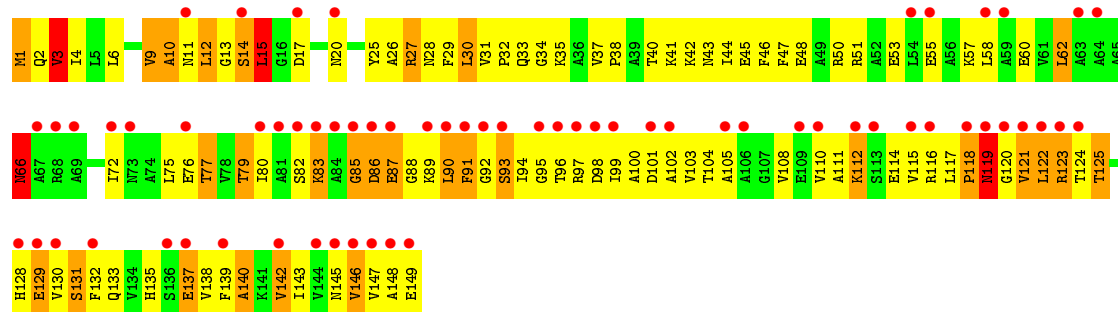
- Molecule 28: 50S ribosomal protein L6



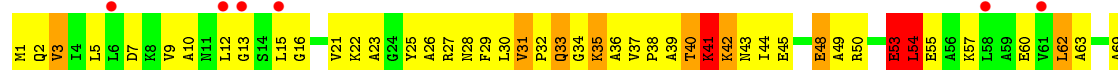
- Molecule 28: 50S ribosomal protein L6

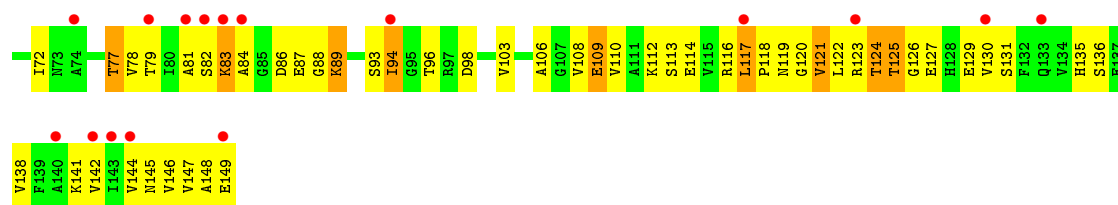


- Molecule 29: 50S ribosomal protein L9

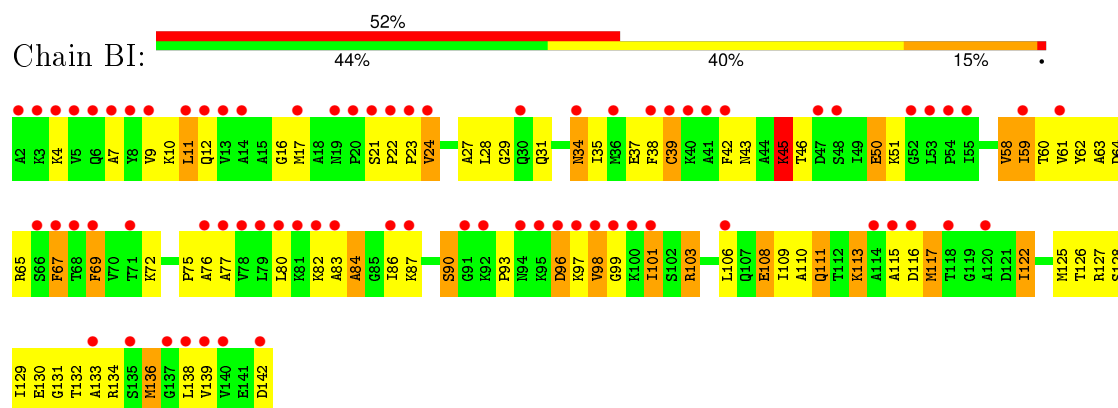


- Molecule 29: 50S ribosomal protein L9

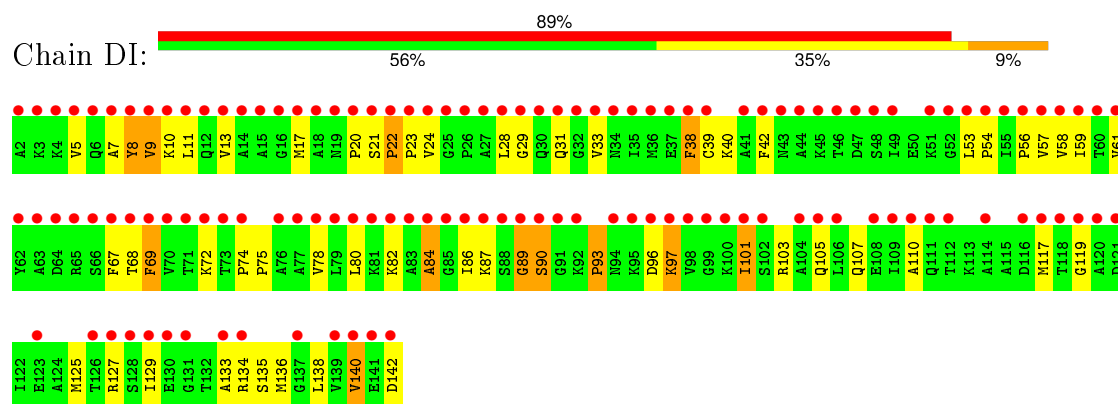




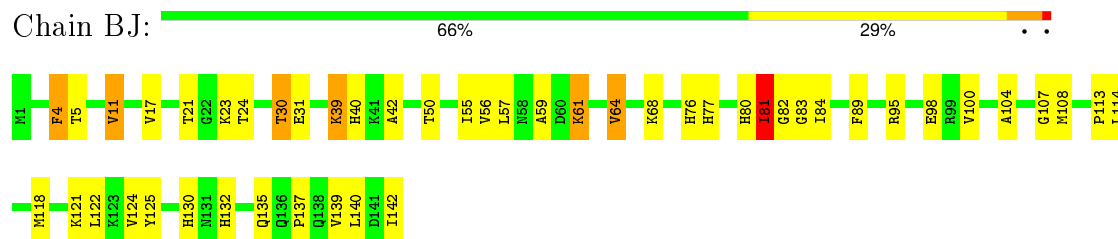
• Molecule 30: 50S ribosomal protein L11



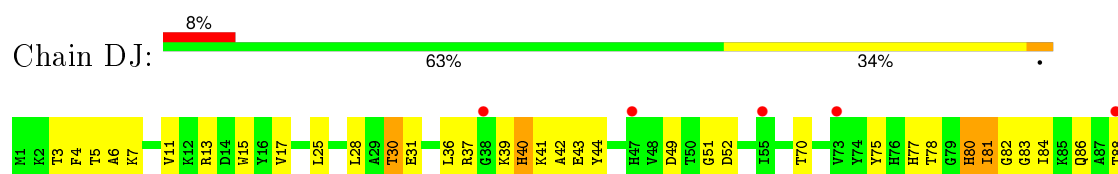
• Molecule 30: 50S ribosomal protein L11



• Molecule 31: 50S ribosomal protein L13



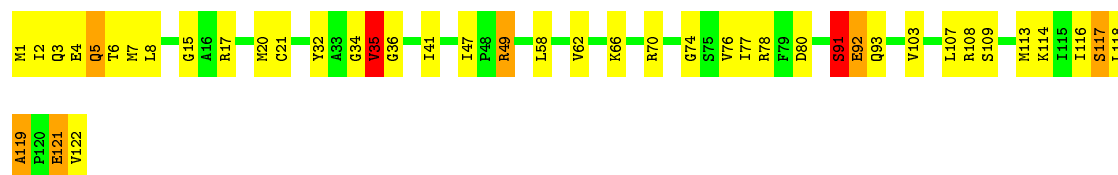
• Molecule 31: 50S ribosomal protein L13





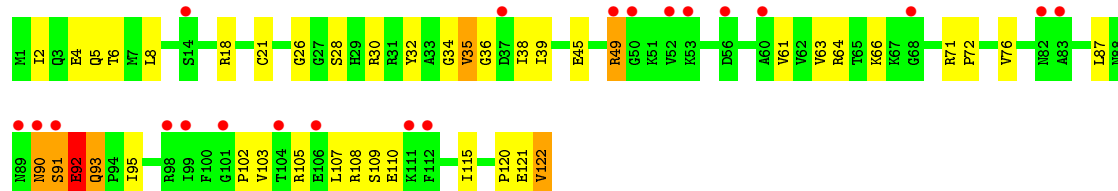
- Molecule 32: 50S ribosomal protein L14

Chain BK: 65% 29% 5% •



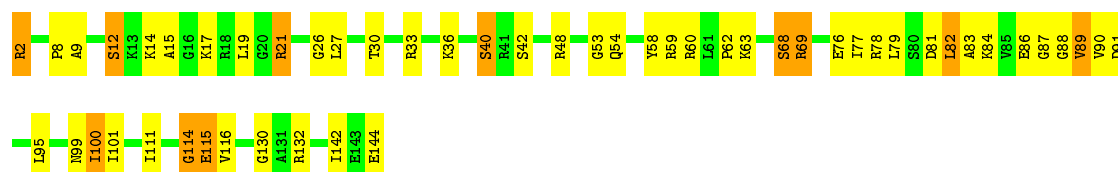
- Molecule 32: 50S ribosomal protein L14

Chain DK: 17% 66% 29% 5% •



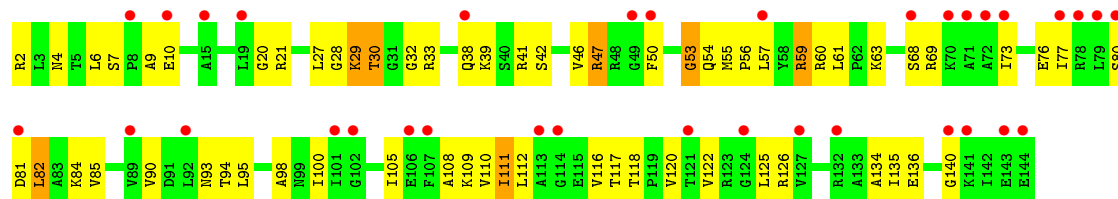
- Molecule 33: 50S ribosomal protein L15

Chain BL: 64% 29% 8%



- Molecule 33: 50S ribosomal protein L15

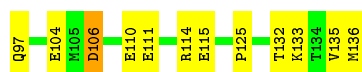
Chain DL: 24% 56% 39% 5%



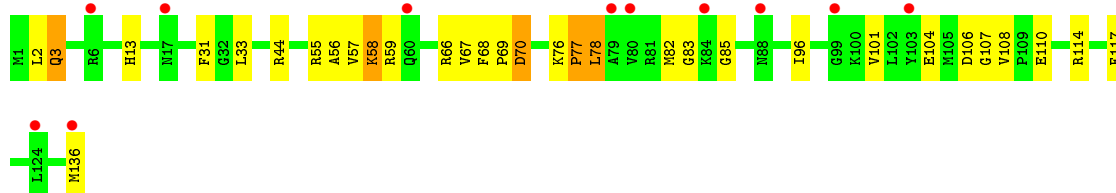
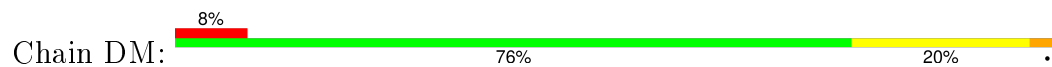
- Molecule 34: 50S ribosomal protein L16

Chain BM: 63% 32% 6%

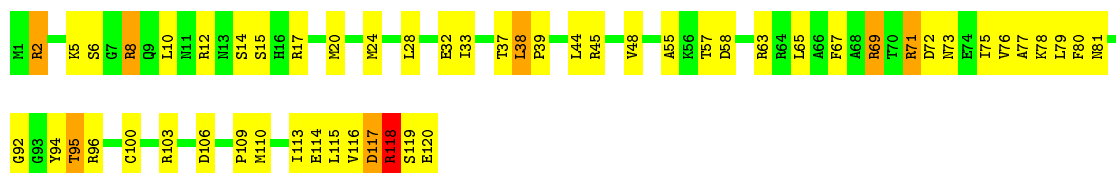




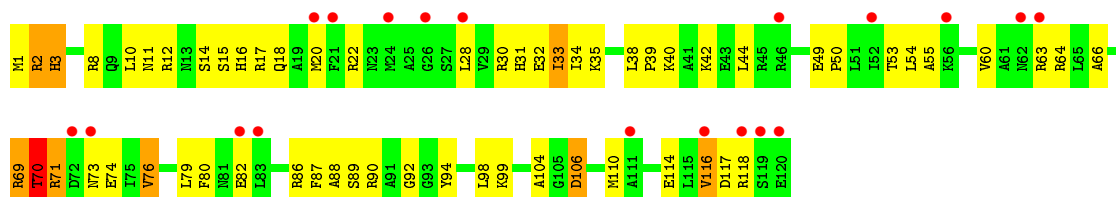
- Molecule 34: 50S ribosomal protein L16



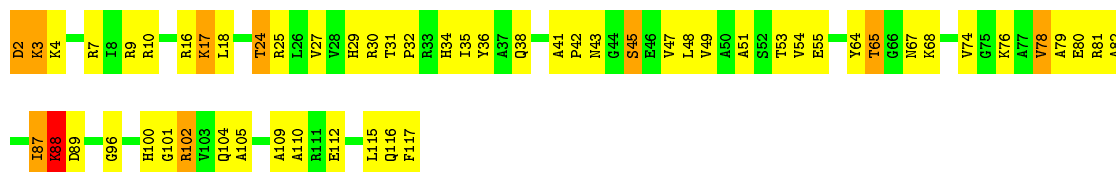
- Molecule 35: 50S ribosomal protein L17



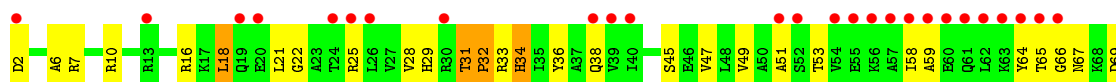
- Molecule 35: 50S ribosomal protein L17



- Molecule 36: 50S ribosomal protein L18



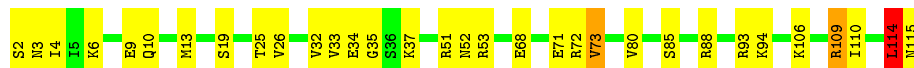
- Molecule 36: 50S ribosomal protein L18





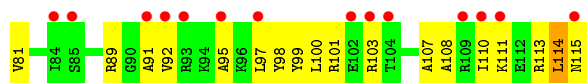
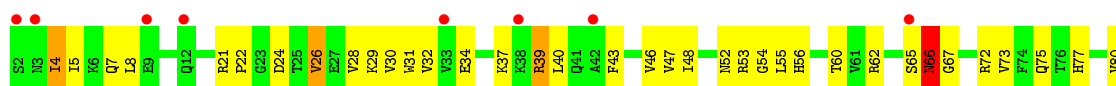
- Molecule 37: 50S ribosomal protein L19

Chain BP: 72% 25%



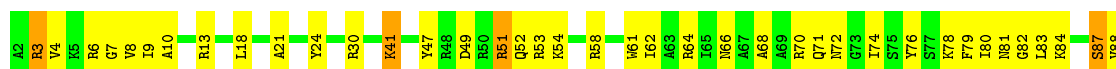
- Molecule 37: 50S ribosomal protein L19

Chain DP: 19% 53% 43%



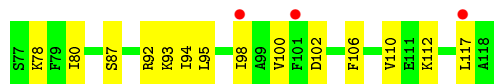
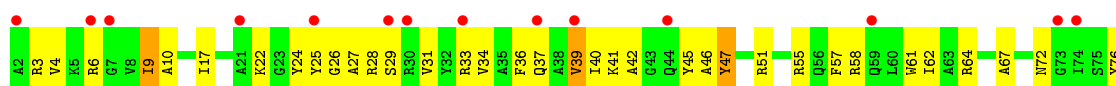
- Molecule 38: 50S ribosomal protein L20

Chain BQ: 56% 40%



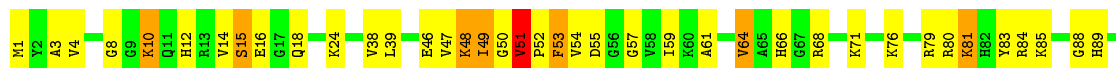
- Molecule 38: 50S ribosomal protein L20

Chain DQ: 15% 58% 39%

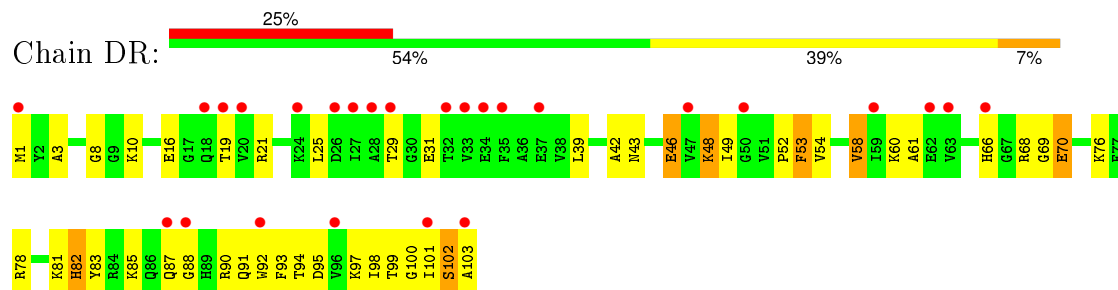


- Molecule 39: 50S ribosomal protein L21

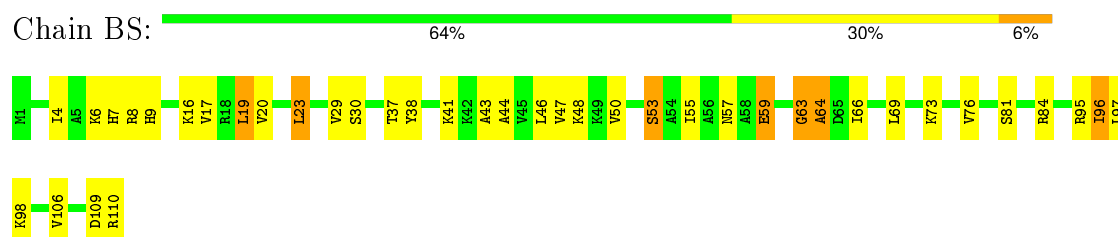
Chain BR: 57% 35% 7%



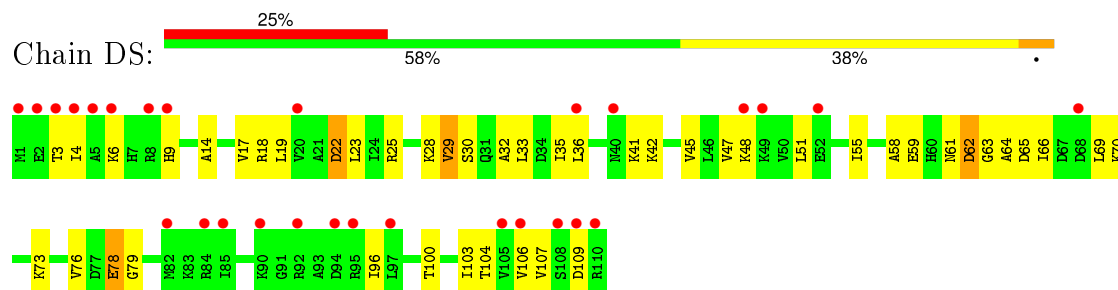
- Molecule 39: 50S ribosomal protein L21



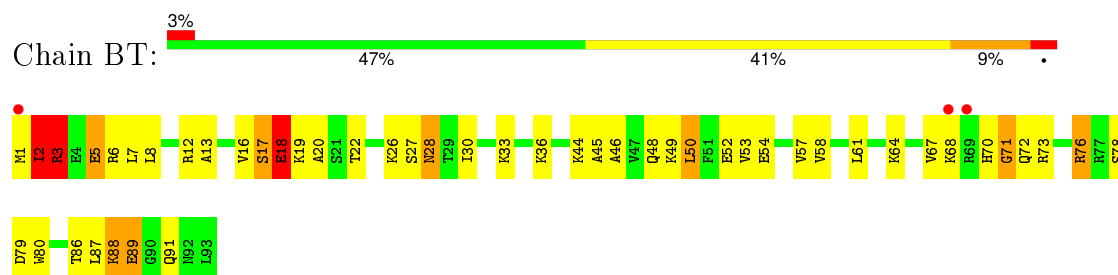
- Molecule 40: 50S ribosomal protein L22



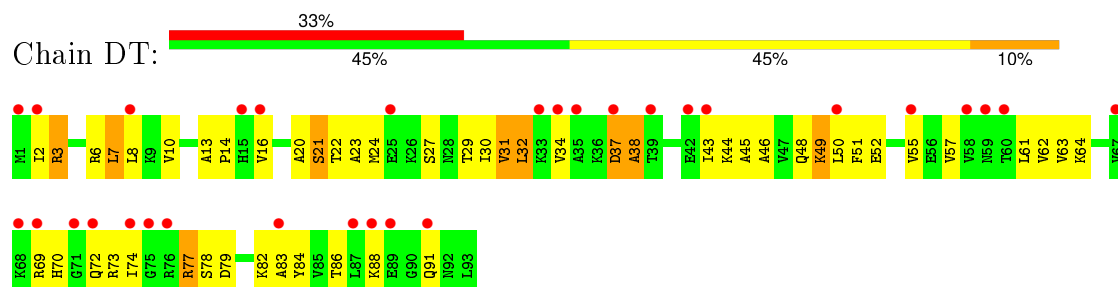
- Molecule 40: 50S ribosomal protein L22



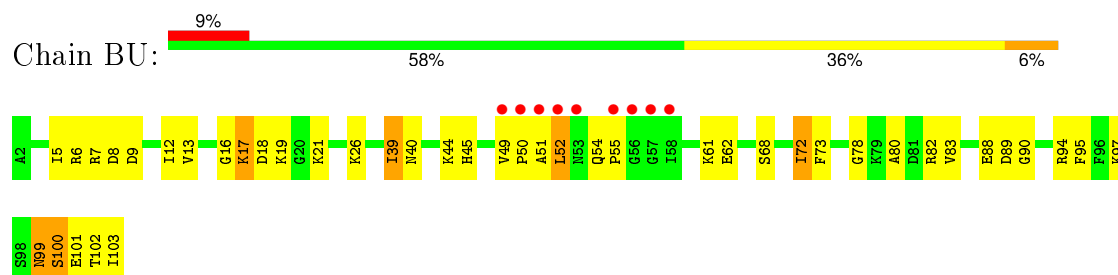
- Molecule 41: 50S ribosomal protein L23



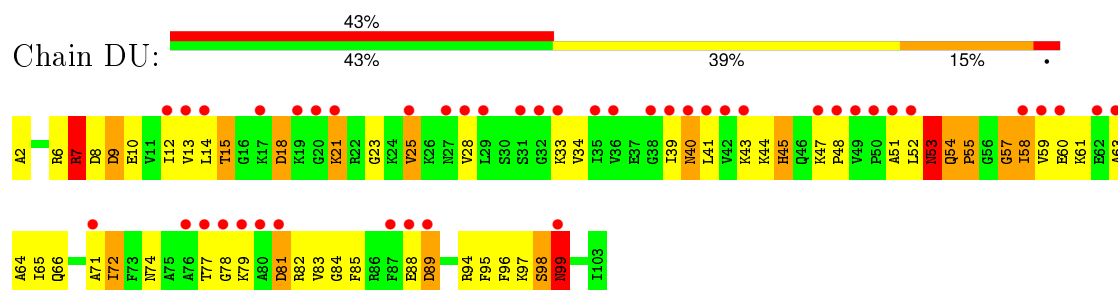
- Molecule 41: 50S ribosomal protein L23



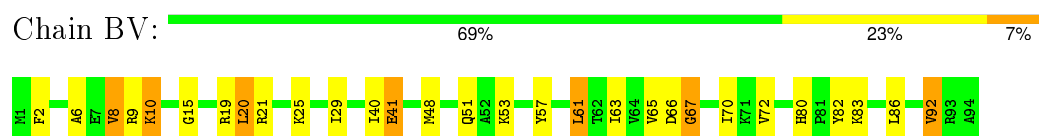
- Molecule 42: 50S ribosomal protein L24



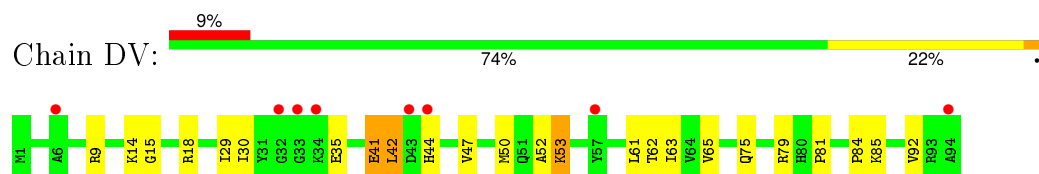
- Molecule 42: 50S ribosomal protein L24



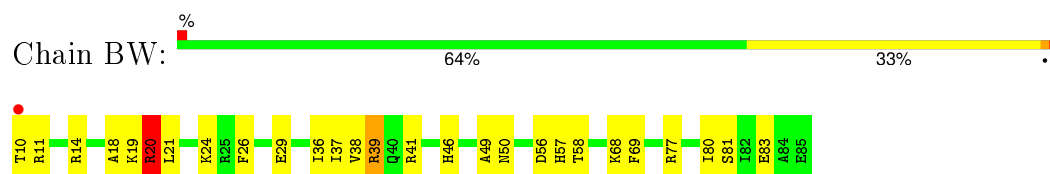
- Molecule 43: 50S ribosomal protein L25



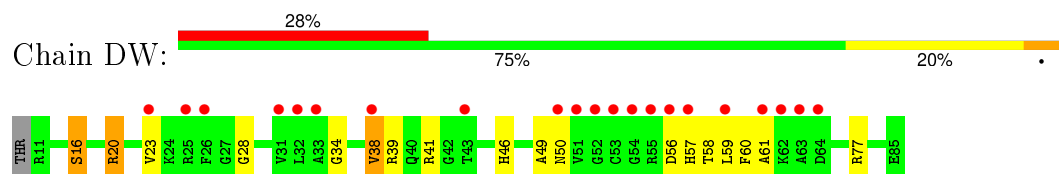
- Molecule 43: 50S ribosomal protein L25



- Molecule 44: 50S ribosomal protein L27

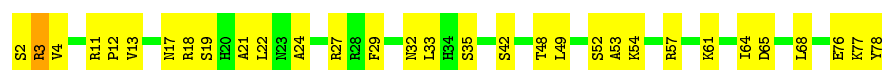


- Molecule 44: 50S ribosomal protein L27




- Molecule 45: 50S ribosomal protein L28

Chain BX:  60% 39%




- Molecule 45: 50S ribosomal protein L28

Chain DX:  12% 42% 49% 9%



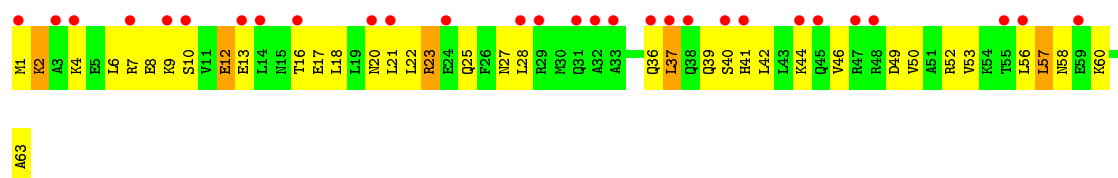
- Molecule 46: 50S ribosomal protein L29

Chain BY:  3% 43% 41% 14%



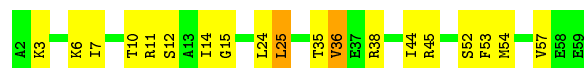
- Molecule 46: 50S ribosomal protein L29

Chain DY:  46% 41% 51% 8%



- Molecule 47: 50S ribosomal protein L30

Chain BZ:  67% 29%



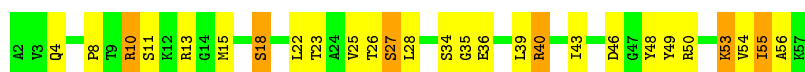
- Molecule 47: 50S ribosomal protein L30

Chain DZ:  10% 48% 43% 9%

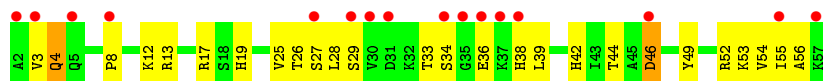


- Molecule 48: 50S ribosomal protein L32

Chain B0:  52% 38% 11%



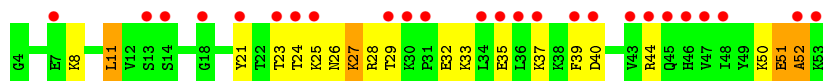
- Molecule 48: 50S ribosomal protein L32



- Molecule 49: 50S ribosomal protein L33



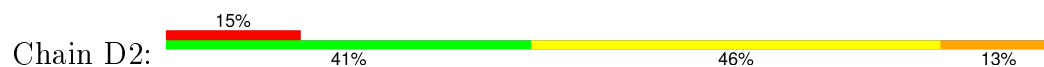
- Molecule 49: 50S ribosomal protein L33



- Molecule 50: 50S ribosomal protein L34



- Molecule 50: 50S ribosomal protein L34

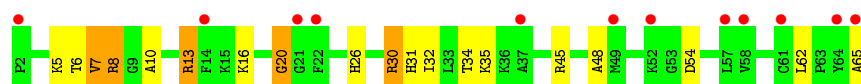


- Molecule 51: 50S ribosomal protein L35



- Molecule 51: 50S ribosomal protein L35

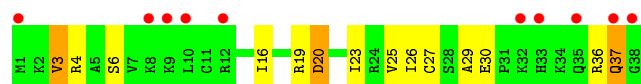




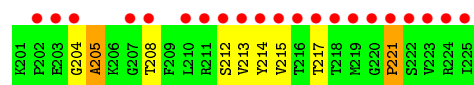
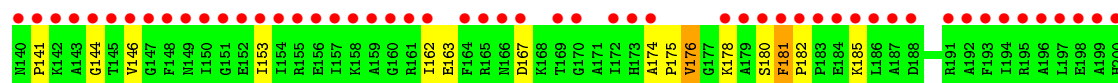
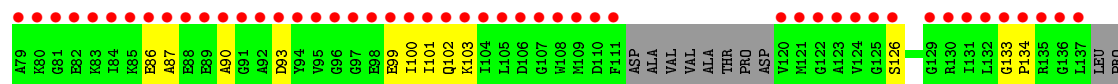
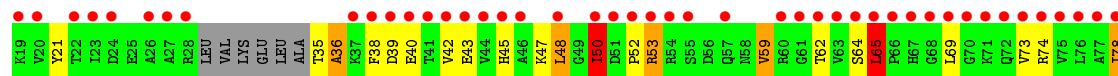
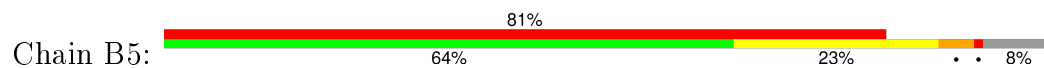
- Molecule 52: 50S ribosomal protein L36



- Molecule 52: 50S ribosomal protein L36



- Molecule 53: 50S ribosomal protein L1



4 Data and refinement statistics

Property	Value	Source
Space group	P 21 21 21	Depositor
Cell constants a, b, c, α , β , γ	211.91Å 434.31Å 624.40Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	41.14 – 3.09 41.14 – 3.09	Depositor EDS
% Data completeness (in resolution range)	99.9 (41.14-3.09) 99.9 (41.14-3.09)	Depositor EDS
R_{merge}	0.13	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	2.37 (at 3.06Å)	Xtriage
Refinement program	PHENIX (phenix.refine: 1.8.1_1168)	Depositor
R, R_{free}	0.202 , 0.244 0.211 , 0.254	Depositor DCC
R_{free} test set	4190 reflections (0.40%)	DCC
Wilson B-factor (Å ²)	72.7	Xtriage
Anisotropy	0.339	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.28 , 58.9	EDS
Estimated twinning fraction	No twinning to report.	Xtriage
L-test for twinning ²	$\langle L \rangle = 0.46$, $\langle L^2 \rangle = 0.29$	Xtriage
Outliers	1 of 1039051 reflections (0.000%)	Xtriage
F_o, F_c correlation	0.93	EDS
Total number of atoms	288204	wwPDB-VP
Average B, all atoms (Å ²)	67.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.50% of the height of the origin peak. No significant pseudotranslation is detected.*

¹Intensities estimated from amplitudes.

²Theoretical values of $\langle |L| \rangle$, $\langle L^2 \rangle$ for acentric reflections are 0.5, 0.375 respectively for untwinned datasets, and 0.333, 0.2 for perfectly twinned datasets.

5 Model quality ⓘ

5.1 Standard geometry ⓘ

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

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

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	$\# Z > 5$	RMSZ	$\# Z > 5$
1	AA	0.58	3/36944 (0.0%)	1.07	91/57632 (0.2%)
1	CA	0.51	3/36966 (0.0%)	1.00	57/57666 (0.1%)
2	AB	0.42	0/1736	0.60	0/2338
2	CB	0.35	0/1736	0.55	0/2338
3	AC	0.36	0/1652	0.52	0/2225
3	CC	0.36	0/1652	0.48	0/2225
4	AD	0.40	0/1665	0.60	0/2227
4	CD	0.50	0/1665	0.66	0/2227
5	AE	0.46	0/1119	0.71	0/1504
5	CE	0.44	0/1119	0.68	0/1504
6	AF	0.46	0/836	0.63	0/1128
6	CF	0.39	0/836	0.54	0/1128
7	AG	0.37	0/1196	0.53	0/1602
7	CG	0.37	0/1196	0.48	0/1602
8	AH	0.41	0/989	0.59	0/1326
8	CH	0.34	0/989	0.55	0/1326
9	AI	0.35	0/1034	0.54	0/1375
9	CI	0.35	0/1034	0.50	0/1375
10	AJ	0.35	0/797	0.54	0/1077
10	CJ	0.34	0/797	0.51	0/1077
11	AK	0.38	0/893	0.59	0/1205
11	CK	0.35	0/893	0.59	0/1205
12	AL	0.49	0/969	0.73	0/1300
12	CL	0.41	0/969	0.71	0/1300
13	AM	0.36	0/893	0.57	0/1193
13	CM	0.36	0/893	0.54	0/1193
14	AN	0.36	0/785	0.52	0/1043
14	CN	0.34	0/785	0.48	0/1043
15	AO	0.38	0/722	0.60	0/964
15	CO	0.34	0/722	0.54	0/964
16	AP	0.42	0/659	0.59	0/884
16	CP	0.40	0/659	0.62	0/884

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
17	AQ	0.47	0/658	0.62	0/881
17	CQ	0.37	0/658	0.56	0/881
18	AR	0.39	0/463	0.59	0/621
18	CR	0.37	0/463	0.57	0/621
19	AS	0.37	0/653	0.53	0/877
19	CS	0.35	0/653	0.48	0/877
20	AT	0.44	0/671	0.62	0/888
20	CT	0.37	0/671	0.54	0/888
21	AU	0.58	0/431	0.70	0/570
21	CU	0.57	0/431	0.67	0/570
22	BA	1.11	160/69659 (0.2%)	1.57	1390/108672 (1.3%)
22	DA	0.49	0/69659	0.96	49/108672 (0.0%)
23	BB	0.92	4/2850 (0.1%)	1.50	47/4444 (1.1%)
23	DB	0.39	0/2828	0.83	0/4410
24	BC	0.68	1/2122 (0.0%)	0.79	0/2852
24	DC	0.39	0/2122	0.60	0/2852
25	BD	0.80	1/1586 (0.1%)	0.98	4/2134 (0.2%)
25	DD	0.37	0/1586	0.56	0/2134
26	BE	0.60	0/1571	0.72	0/2113
26	DE	0.37	0/1571	0.53	0/2113
27	BF	0.43	0/1435	0.61	0/1926
27	DF	0.32	0/1435	0.46	0/1926
28	BG	0.50	0/1343	0.70	1/1816 (0.1%)
28	DG	0.32	0/1343	0.46	0/1816
29	BH	0.32	0/1121	0.63	0/1515
29	DH	0.34	0/1121	0.56	0/1515
30	BI	0.39	0/1046	0.51	0/1410
30	DI	0.38	0/1046	0.51	0/1410
31	BJ	0.77	0/1152	0.76	0/1551
31	DJ	0.36	0/1152	0.56	0/1551
32	BK	0.70	0/948	0.86	0/1268
32	DK	0.39	0/948	0.55	0/1268
33	BL	0.66	0/1054	0.80	0/1403
33	DL	0.36	0/1054	0.57	0/1403
34	BM	0.71	0/1093	0.86	1/1460 (0.1%)
34	DM	0.34	0/1093	0.53	0/1460
35	BN	0.71	0/974	0.89	0/1301
35	DN	0.36	0/974	0.53	0/1301
36	BO	0.50	0/902	0.67	0/1209
36	DO	0.32	0/902	0.46	0/1209
37	BP	0.64	0/929	0.74	0/1242
37	DP	0.39	0/929	0.55	0/1242
38	BQ	0.86	0/960	0.92	1/1278 (0.1%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
38	DQ	0.37	0/960	0.52	0/1278
39	BR	0.82	0/829	0.91	1/1107 (0.1%)
39	DR	0.35	0/829	0.55	0/1107
40	BS	0.85	0/864	0.91	1/1156 (0.1%)
40	DS	0.35	0/864	0.55	0/1156
41	BT	0.67	1/745 (0.1%)	0.77	1/994 (0.1%)
41	DT	0.38	0/745	0.57	0/994
42	BU	0.54	0/788	0.71	0/1051
42	DU	0.41	0/788	0.57	0/1051
43	BV	0.63	0/766	0.70	0/1025
43	DV	0.31	0/766	0.45	0/1025
44	BW	0.70	0/587	0.80	1/776 (0.1%)
44	DW	0.33	0/576	0.48	0/762
45	BX	0.52	0/635	0.71	0/848
45	DX	0.40	0/635	0.60	0/848
46	BY	0.55	0/510	0.74	0/677
46	DY	0.37	0/510	0.54	0/677
47	BZ	0.77	0/453	0.86	0/605
47	DZ	0.34	0/453	0.50	0/605
48	B0	0.77	0/450	0.81	0/599
48	D0	0.36	0/450	0.57	0/599
49	B1	0.50	0/417	0.60	0/554
49	D1	0.36	0/417	0.49	0/554
50	B2	0.77	0/380	0.95	0/498
50	D2	0.39	0/380	0.62	0/498
51	B3	0.63	0/513	0.86	1/676 (0.1%)
51	D3	0.33	0/513	0.53	0/676
52	B4	0.71	0/303	0.87	0/397
52	D4	0.32	0/303	0.51	0/397
53	B5	0.34	0/1145	0.47	0/1556
All	All	0.69	173/310634 (0.1%)	1.09	1646/464376 (0.4%)

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

Mol	Chain	#Chirality outliers	#Planarity outliers
5	CE	0	2
11	AK	0	1
11	CK	0	1
12	CL	0	1

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Mol	Chain	#Chirality outliers	#Planarity outliers
25	BD	0	1
25	DD	0	1
41	BT	0	1
47	BZ	0	1
All	All	0	9

All (173) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	BA	2777	G	N7-C5	15.93	1.48	1.39
22	BA	984	A	N9-C4	-14.44	1.29	1.37
22	BA	2777	G	N9-C8	13.80	1.47	1.37
22	BA	1142	A	N9-C4	-12.51	1.30	1.37
1	CA	500	G	N9-C8	10.22	1.45	1.37
22	BA	2777	G	C8-N7	10.10	1.37	1.30
22	BA	783	A	N9-C4	-10.06	1.31	1.37
22	BA	974	G	N9-C8	9.79	1.44	1.37
22	BA	984	A	C5-C6	-8.80	1.33	1.41
25	BD	151	THR	C-O	8.78	1.40	1.23
22	BA	2266	A	N9-C4	-8.26	1.32	1.37
22	BA	2572	A	N3-C4	-8.20	1.29	1.34
22	BA	984	A	N7-C5	-8.19	1.34	1.39
22	BA	752	A	N9-C4	-7.91	1.33	1.37
22	BA	528	A	N3-C4	-7.81	1.30	1.34
22	BA	528	A	N9-C4	-7.78	1.33	1.37
22	BA	948	C	N3-C4	-7.66	1.28	1.33
22	BA	1784	A	N9-C4	-7.63	1.33	1.37
1	AA	500	G	N9-C8	7.31	1.43	1.37
22	BA	2013	A	N9-C4	-7.30	1.33	1.37
22	BA	1299	G	N7-C5	-7.18	1.34	1.39
22	BA	1244	A	N9-C4	-7.15	1.33	1.37
22	BA	1785	A	N7-C5	-7.10	1.34	1.39
22	BA	1452	G	N9-C8	7.08	1.42	1.37
22	BA	2712	C	N1-C6	-7.07	1.32	1.37
22	BA	528	A	C5-C6	-7.06	1.34	1.41
22	BA	26	G	C6-N1	-6.95	1.34	1.39
22	BA	2250	G	N9-C8	6.91	1.42	1.37
22	BA	2065	C	N1-C6	-6.83	1.33	1.37
1	CA	500	G	C6-N1	6.83	1.44	1.39
1	CA	500	G	C8-N7	6.80	1.35	1.30
22	BA	572	A	N3-C4	-6.79	1.30	1.34
22	BA	783	A	C5-C6	-6.78	1.34	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	BA	1676	A	N3-C4	-6.77	1.30	1.34
22	BA	1127	A	N3-C4	-6.70	1.30	1.34
22	BA	19	A	N7-C5	-6.69	1.35	1.39
22	BA	2645	G	N7-C5	-6.65	1.35	1.39
22	BA	1783	A	N7-C5	-6.61	1.35	1.39
22	BA	974	G	N3-C4	-6.61	1.30	1.35
22	BA	2029	G	N9-C8	-6.61	1.33	1.37
22	BA	2626	C	N1-C6	-6.59	1.33	1.37
24	BC	213	TRP	CB-CG	-6.47	1.38	1.50
22	BA	789	A	N9-C4	-6.46	1.33	1.37
22	BA	981	A	N9-C4	-6.46	1.33	1.37
22	BA	1263	U	C4-O4	-6.43	1.18	1.23
22	BA	948	C	N1-C6	-6.42	1.33	1.37
22	BA	2030	A	C5-C6	6.39	1.46	1.41
22	BA	26	G	C6-O6	-6.28	1.18	1.24
22	BA	512	G	C8-N7	6.28	1.34	1.30
22	BA	1774	C	N1-C6	-6.26	1.33	1.37
22	BA	916	G	N7-C5	-6.25	1.35	1.39
22	BA	2447	G	N9-C8	-6.24	1.33	1.37
22	BA	2699	C	N1-C6	-6.22	1.33	1.37
22	BA	2014	A	N3-C4	-6.21	1.31	1.34
22	BA	677	A	N9-C4	-6.21	1.34	1.37
22	BA	1264	A	C6-N6	-6.20	1.28	1.33
22	BA	974	G	N9-C4	-6.15	1.33	1.38
22	BA	2025	C	N3-C4	-6.11	1.29	1.33
23	BB	78	A	N3-C4	-6.09	1.31	1.34
22	BA	561	G	N9-C4	-6.09	1.33	1.38
22	BA	2681	C	N1-C6	-6.08	1.33	1.37
22	BA	512	G	N7-C5	6.07	1.42	1.39
22	BA	1039	A	N9-C4	-6.07	1.34	1.37
22	BA	1976	U	C4-O4	-6.07	1.18	1.23
22	BA	802	A	C6-N1	-6.04	1.31	1.35
22	BA	1936	A	N9-C4	-6.01	1.34	1.37
23	BB	15	A	C5-C6	-6.01	1.35	1.41
22	BA	758	C	N1-C6	-6.01	1.33	1.37
22	BA	795	C	N3-C4	-6.00	1.29	1.33
22	BA	2458	G	C2-N3	-5.95	1.27	1.32
22	BA	993	G	C6-N1	-5.94	1.35	1.39
22	BA	1126	A	N7-C5	-5.91	1.35	1.39
22	BA	2826	A	C5-C4	-5.90	1.34	1.38
22	BA	2636	C	N1-C6	-5.89	1.33	1.37
22	BA	1676	A	C6-N1	-5.88	1.31	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	BA	2689	U	C2-N3	-5.88	1.33	1.37
22	BA	751	A	N3-C4	-5.87	1.31	1.34
22	BA	2570	G	N9-C8	-5.86	1.33	1.37
22	BA	1127	A	N9-C4	-5.84	1.34	1.37
22	BA	535	G	C6-N1	-5.83	1.35	1.39
22	BA	1452	G	N9-C4	-5.83	1.33	1.38
22	BA	2723	C	N3-C4	-5.82	1.29	1.33
22	BA	1617	C	N1-C6	-5.81	1.33	1.37
1	AA	500	G	C6-N1	5.80	1.43	1.39
22	BA	2250	G	C5-C6	-5.80	1.36	1.42
22	BA	1785	A	N9-C4	-5.79	1.34	1.37
23	BB	99	A	N9-C4	-5.77	1.34	1.37
22	BA	972	A	C5-C4	-5.76	1.34	1.38
22	BA	2570	G	C6-N1	-5.75	1.35	1.39
22	BA	198	C	N1-C6	-5.74	1.33	1.37
22	BA	450	G	N3-C4	-5.74	1.31	1.35
23	BB	78	A	N9-C4	-5.72	1.34	1.37
22	BA	2590	A	N9-C4	-5.71	1.34	1.37
22	BA	2564	A	N7-C5	-5.69	1.35	1.39
22	BA	2014	A	C5-C4	-5.68	1.34	1.38
1	AA	500	G	C8-N7	5.68	1.34	1.30
22	BA	1271	G	C6-N1	-5.63	1.35	1.39
22	BA	1296	G	N9-C8	-5.63	1.33	1.37
22	BA	1756	G	N7-C5	-5.60	1.35	1.39
22	BA	788	A	N7-C5	-5.57	1.35	1.39
22	BA	2030	A	N3-C4	-5.56	1.31	1.34
22	BA	1674	G	N7-C5	-5.55	1.35	1.39
22	BA	2250	G	C2-N2	5.54	1.40	1.34
22	BA	2740	A	N3-C4	-5.54	1.31	1.34
22	BA	255	A	N9-C4	-5.52	1.34	1.37
22	BA	2608	G	C6-N1	-5.50	1.35	1.39
22	BA	2771	C	N3-C4	-5.49	1.30	1.33
22	BA	752	A	N9-C8	5.49	1.42	1.37
22	BA	581	C	N3-C4	-5.48	1.30	1.33
22	BA	752	A	C5-C4	5.48	1.42	1.38
22	BA	1187	G	N7-C5	-5.46	1.35	1.39
22	BA	2628	C	N1-C6	-5.46	1.33	1.37
22	BA	2248	C	N3-C4	-5.44	1.30	1.33
22	BA	1002	G	N1-C2	-5.44	1.33	1.37
22	BA	1275	A	N9-C8	-5.43	1.33	1.37
22	BA	2053	G	N7-C5	-5.42	1.35	1.39
22	BA	1265	A	N3-C4	-5.42	1.31	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	BA	2777	G	C5-C4	5.42	1.42	1.38
22	BA	984	A	N3-C4	-5.41	1.31	1.34
22	BA	1309	G	C6-N1	-5.41	1.35	1.39
22	BA	578	G	N7-C5	-5.39	1.36	1.39
22	BA	2250	G	N9-C4	-5.35	1.33	1.38
22	BA	2389	G	N7-C5	-5.34	1.36	1.39
22	BA	526	A	N9-C8	-5.34	1.33	1.37
22	BA	2825	G	N7-C5	-5.32	1.36	1.39
22	BA	2570	G	N7-C5	-5.30	1.36	1.39
22	BA	1253	A	N9-C4	-5.29	1.34	1.37
22	BA	2463	C	N1-C6	-5.28	1.33	1.37
22	BA	401	A	N3-C4	-5.24	1.31	1.34
22	BA	979	A	N9-C4	-5.23	1.34	1.37
22	BA	2266	A	N3-C4	-5.22	1.31	1.34
22	BA	1039	A	N3-C4	-5.22	1.31	1.34
22	BA	1938	A	N9-C4	-5.22	1.34	1.37
41	BT	18	GLU	CG-CD	5.22	1.59	1.51
22	BA	568	U	C2-N3	-5.21	1.34	1.37
22	BA	1635	A	N3-C4	-5.21	1.31	1.34
22	BA	1779	U	N3-C4	-5.21	1.33	1.38
22	BA	2814	A	C5-C6	-5.20	1.36	1.41
22	BA	685	A	N9-C4	-5.20	1.34	1.37
22	BA	2038	G	N7-C5	-5.19	1.36	1.39
22	BA	125	A	N7-C5	-5.19	1.36	1.39
22	BA	2707	U	C2-N3	-5.18	1.34	1.37
22	BA	2279	G	C6-N1	-5.17	1.35	1.39
22	BA	572	A	C6-N1	-5.16	1.31	1.35
22	BA	1296	G	C5-C4	-5.16	1.34	1.38
22	BA	673	C	N1-C2	-5.15	1.34	1.40
22	BA	863	A	N3-C4	-5.14	1.31	1.34
22	BA	538	A	N7-C5	-5.13	1.36	1.39
22	BA	2045	C	N3-C4	-5.13	1.30	1.33
22	BA	2038	G	C5-C6	-5.13	1.37	1.42
22	BA	801	G	C8-N7	5.10	1.34	1.30
22	BA	1550	C	N1-C6	-5.09	1.34	1.37
22	BA	1817	G	C2-N3	5.08	1.36	1.32
22	BA	1251	C	N1-C6	-5.08	1.34	1.37
22	BA	1655	A	N7-C5	-5.07	1.36	1.39
22	BA	2029	G	N7-C5	-5.07	1.36	1.39
22	BA	2569	G	C6-N1	-5.07	1.36	1.39
22	BA	2883	A	C6-N1	-5.07	1.32	1.35
22	BA	2032	G	C6-N1	-5.06	1.36	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	BA	1671	U	C2-N3	-5.06	1.34	1.37
22	BA	2570	G	C5-C4	-5.06	1.34	1.38
22	BA	789	A	N3-C4	-5.05	1.31	1.34
22	BA	2276	G	C6-N1	-5.05	1.36	1.39
22	BA	2719	G	N9-C8	-5.05	1.34	1.37
22	BA	1635	A	N9-C4	-5.05	1.34	1.37
22	BA	2430	A	N3-C4	-5.04	1.31	1.34
22	BA	2227	A	N7-C5	-5.04	1.36	1.39
22	BA	2497	A	N3-C4	-5.03	1.31	1.34
22	BA	2070	A	N7-C5	-5.03	1.36	1.39
22	BA	1676	A	N9-C4	-5.01	1.34	1.37
22	BA	2622	U	N1-C2	-5.01	1.34	1.38
22	BA	990	A	N3-C4	-5.01	1.31	1.34
22	BA	1136	G	C5-C4	-5.00	1.34	1.38

All (1646) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
25	BD	151	THR	C-N-CD	-21.19	73.98	120.60
22	BA	2777	G	C5-N7-C8	-19.20	94.70	104.30
22	BA	984	A	C2-N3-C4	-18.54	101.33	110.60
22	BA	2777	G	C4-C5-C6	-17.34	108.40	118.80
22	BA	528	A	N1-C6-N6	16.96	128.78	118.60
22	BA	2250	G	C4-C5-N7	16.54	117.42	110.80
1	CA	500	G	C5-N7-C8	-15.57	96.51	104.30
22	BA	528	A	C2-N3-C4	-15.22	102.99	110.60
22	BA	984	A	N3-C4-C5	15.10	137.37	126.80
22	BA	528	A	C6-C5-N7	-15.02	121.78	132.30
22	BA	2250	G	C5-N7-C8	-14.98	96.81	104.30
22	BA	2777	G	N1-C6-O6	-14.32	111.31	119.90
22	BA	783	A	C5-N7-C8	-14.10	96.85	103.90
22	BA	1936	A	C2-N3-C4	-13.64	103.78	110.60
22	BA	984	A	N3-C4-N9	-13.60	116.52	127.40
22	BA	2777	G	N3-C4-N9	-13.45	117.93	126.00
23	BB	83	G	O5'-P-OP2	-13.29	93.74	105.70
22	BA	2777	G	N7-C8-N9	13.17	119.69	113.10
22	BA	1142	A	C2-N3-C4	-12.80	104.20	110.60
22	BA	532	A	O5'-P-OP1	-12.67	94.30	105.70
1	CA	500	G	C4-C5-N7	12.65	115.86	110.80
22	BA	2030	A	C5-C6-N6	12.44	133.65	123.70
22	BA	2777	G	C8-N9-C1'	12.43	143.16	127.00
22	BA	565	C	N1-C2-O2	-12.24	111.56	118.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	1142	A	N3-C4-C5	12.15	135.30	126.80
22	BA	561	G	N1-C6-O6	12.15	127.19	119.90
22	BA	1142	A	N3-C4-N9	-12.13	117.70	127.40
1	CA	500	G	N3-C4-C5	11.97	134.59	128.60
22	BA	783	A	C4-C5-N7	11.94	116.67	110.70
25	BD	151	THR	C-N-CA	11.91	172.01	122.00
22	BA	528	A	C5-N7-C8	-11.82	97.99	103.90
22	BA	2030	A	N9-C4-C5	11.68	110.47	105.80
22	BA	673	C	N1-C2-O2	-11.66	111.91	118.90
22	BA	819	A	O5'-P-OP1	-11.52	95.33	105.70
22	BA	783	A	C2-N3-C4	-11.50	104.85	110.60
22	BA	2824	C	N1-C2-O2	-11.49	112.01	118.90
22	BA	1779	U	C5-C6-N1	-11.45	116.97	122.70
1	AA	500	G	C5-N7-C8	-11.39	98.61	104.30
22	BA	752	A	C5-N7-C8	-11.37	98.22	103.90
22	BA	974	G	C5-N7-C8	-11.34	98.63	104.30
22	BA	2777	G	N3-C4-C5	11.30	134.25	128.60
22	BA	783	A	N1-C6-N6	11.27	125.36	118.60
22	BA	512	G	C5-N7-C8	-11.26	98.67	104.30
22	BA	528	A	C4-C5-N7	11.25	116.32	110.70
22	BA	1452	G	N3-C4-C5	11.23	134.22	128.60
22	BA	1300	G	O5'-P-OP2	-11.13	95.69	105.70
22	BA	950	G	O5'-P-OP2	-11.09	95.72	105.70
22	BA	528	A	N1-C2-N3	11.08	134.84	129.30
22	BA	1652	A	O5'-P-OP2	-11.02	95.78	105.70
22	BA	1663	G	C2-N3-C4	-10.88	106.46	111.90
1	CA	500	G	N3-C4-N9	-10.84	119.50	126.00
22	BA	1249	U	O5'-P-OP1	-10.76	96.02	105.70
22	BA	2814	A	N1-C6-N6	10.70	125.02	118.60
22	BA	944	C	O5'-P-OP2	-10.69	96.08	105.70
22	BA	752	A	C2-N3-C4	-10.49	105.35	110.60
22	BA	2889	C	N1-C2-O2	-10.49	112.61	118.90
22	BA	2777	G	C4-C5-N7	10.44	114.98	110.80
22	BA	2030	A	N1-C6-N6	-10.44	112.34	118.60
22	BA	2030	A	C4-C5-N7	-10.44	105.48	110.70
22	BA	974	G	N3-C4-C5	10.40	133.80	128.60
1	CA	500	G	N7-C8-N9	10.38	118.29	113.10
22	BA	2777	G	C6-C5-N7	10.37	136.62	130.40
22	BA	1251	C	C5-C4-N4	-10.30	112.99	120.20
22	BA	1452	G	N3-C4-N9	-10.21	119.87	126.00
22	BA	128	C	N1-C2-O2	-10.18	112.79	118.90
1	CA	500	G	C5-C6-O6	-10.03	122.58	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	AA	500	G	N3-C4-N9	-9.98	120.01	126.00
22	BA	2867	G	O5'-P-OP1	-9.97	96.72	105.70
22	BA	2689	U	C5-C4-O4	9.97	131.88	125.90
22	BA	1340	U	N3-C4-O4	-9.95	112.44	119.40
22	BA	974	G	C4-C5-N7	9.90	114.76	110.80
22	BA	2053	G	C5-C6-O6	-9.90	122.66	128.60
22	BA	1340	U	C5-C4-O4	9.87	131.82	125.90
1	AA	500	G	N3-C4-C5	9.85	133.52	128.60
22	BA	1694	C	N1-C2-O2	-9.84	113.00	118.90
22	BA	2777	G	C5-C6-N1	9.80	116.40	111.50
22	BA	574	A	O5'-P-OP2	9.79	122.45	110.70
22	BA	664	G	O5'-P-OP2	-9.79	96.89	105.70
22	BA	528	A	C4-C5-C6	9.78	121.89	117.00
22	BA	914	G	N1-C6-O6	9.77	125.76	119.90
22	BA	2840	C	O5'-P-OP2	-9.76	96.92	105.70
22	BA	2250	G	N3-C4-C5	9.75	133.48	128.60
22	BA	1692	U	N3-C2-O2	9.72	129.00	122.20
23	BB	15	A	N1-C6-N6	9.72	124.43	118.60
22	BA	2250	G	C6-C5-N7	-9.70	124.58	130.40
22	BA	752	A	N7-C8-N9	9.68	118.64	113.80
22	BA	1002	G	C5-C6-O6	9.66	134.40	128.60
22	BA	2359	C	N1-C2-O2	-9.63	113.12	118.90
22	BA	984	A	N1-C6-N6	9.63	124.38	118.60
22	BA	740	C	OP1-P-OP2	-9.62	105.17	119.60
22	BA	2777	G	C4-N9-C1'	-9.62	114.00	126.50
22	BA	2005	A	N1-C6-N6	9.56	124.34	118.60
22	BA	974	G	N3-C4-N9	-9.51	120.30	126.00
22	BA	1677	A	N1-C6-N6	9.49	124.29	118.60
22	BA	1619	G	O5'-P-OP2	-9.44	97.21	105.70
22	BA	2744	G	C5-C6-O6	-9.43	122.94	128.60
1	CA	500	G	N1-C6-O6	9.39	125.53	119.90
22	BA	1266	G	O5'-P-OP1	-9.33	97.30	105.70
22	BA	783	A	N3-C4-C5	9.31	133.31	126.80
22	BA	2744	G	C8-N9-C4	9.22	110.09	106.40
22	BA	2616	C	N1-C2-O2	-9.22	113.37	118.90
22	BA	1142	A	C5-N7-C8	-9.19	99.30	103.90
22	BA	2505	G	N1-C6-O6	-9.11	114.44	119.90
22	BA	747	U	N3-C4-O4	-9.09	113.04	119.40
22	BA	2036	C	C6-N1-C2	-9.06	116.67	120.30
22	BA	2698	U	C5-C4-O4	-9.06	120.46	125.90
22	BA	1332	G	C6-C5-N7	-9.06	124.97	130.40
22	BA	1191	G	O5'-P-OP2	-9.04	97.56	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	993	G	N1-C6-O6	-9.01	114.50	119.90
22	BA	2777	G	C8-N9-C4	-8.95	102.82	106.40
22	BA	1134	A	OP1-P-OP2	8.94	133.01	119.60
22	BA	2645	G	C6-C5-N7	-8.93	125.04	130.40
22	BA	841	G	N1-C6-O6	8.91	125.25	119.90
22	BA	984	A	C5-N7-C8	-8.91	99.45	103.90
22	BA	2049	G	C8-N9-C4	-8.87	102.85	106.40
22	BA	1676	A	O5'-P-OP2	-8.82	97.76	105.70
22	BA	2596	U	O5'-P-OP1	-8.81	97.77	105.70
22	BA	2606	C	C6-N1-C2	8.79	123.82	120.30
22	BA	1653	G	C8-N9-C4	-8.77	102.89	106.40
22	BA	1566	A	N1-C6-N6	8.76	123.86	118.60
23	BB	82	U	OP2-P-O3'	8.76	124.46	105.20
22	BA	2884	U	C5-C4-O4	8.75	131.15	125.90
22	BA	1240	U	O5'-P-OP2	-8.74	97.84	105.70
22	BA	658	U	O5'-P-OP1	-8.73	97.84	105.70
22	BA	2825	G	N3-C4-C5	-8.71	124.25	128.60
22	BA	2021	C	O5'-P-OP2	-8.64	97.92	105.70
22	BA	516	C	N1-C2-O2	-8.64	113.72	118.90
22	BA	831	G	N9-C4-C5	-8.63	101.95	105.40
22	BA	2250	G	N7-C8-N9	8.62	117.41	113.10
1	AA	279	A	N1-C6-N6	8.62	123.77	118.60
23	BB	83	G	O5'-P-OP1	8.60	121.02	110.70
1	CA	500	G	C8-N9-C1'	8.58	138.16	127.00
1	CA	26	A	O5'-P-OP2	-8.55	98.01	105.70
22	BA	2499	C	N1-C2-O2	-8.54	113.77	118.90
22	BA	2513	A	N1-C6-N6	-8.49	113.51	118.60
22	BA	2787	C	C2-N3-C4	-8.48	115.66	119.90
22	BA	745	G	C6-N1-C2	-8.47	120.02	125.10
22	BA	2569	G	N1-C6-O6	-8.44	114.84	119.90
22	BA	782	A	O5'-P-OP1	-8.43	98.11	105.70
22	BA	831	G	C5-C6-O6	-8.43	123.55	128.60
22	BA	783	A	N7-C8-N9	8.40	118.00	113.80
22	BA	2499	C	O5'-P-OP2	-8.38	98.15	105.70
1	AA	500	G	N7-C8-N9	8.38	117.29	113.10
22	BA	2041	U	O5'-P-OP2	-8.37	98.17	105.70
22	BA	1002	G	N1-C6-O6	-8.36	114.89	119.90
22	BA	528	A	C5-C6-N1	-8.35	113.52	117.70
22	BA	512	G	N7-C8-N9	8.35	117.27	113.10
22	BA	2383	G	C5-N7-C8	-8.35	100.13	104.30
22	BA	2425	A	P-O3'-C3'	8.33	129.69	119.70
22	BA	1452	G	C5-N7-C8	-8.32	100.14	104.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	2678	C	N3-C4-C5	8.32	125.23	121.90
22	BA	2005	A	C5-C6-N6	-8.29	117.07	123.70
22	BA	1663	G	N9-C4-C5	-8.29	102.08	105.40
22	BA	2490	G	C5-C6-O6	-8.26	123.64	128.60
22	BA	561	G	C5-C6-O6	-8.25	123.65	128.60
22	BA	2679	A	N1-C6-N6	8.24	123.55	118.60
22	BA	2627	G	O5'-P-OP2	-8.24	98.28	105.70
22	BA	984	A	C4-C5-N7	8.22	114.81	110.70
22	BA	1020	A	N1-C6-N6	8.22	123.53	118.60
22	BA	1618	A	C5-C6-N1	-8.21	113.59	117.70
22	BA	731	C	C2-N3-C4	-8.21	115.80	119.90
22	BA	752	A	C8-N9-C4	-8.20	102.52	105.80
22	BA	1147	A	O5'-P-OP2	-8.20	98.32	105.70
22	BA	2502	G	N9-C4-C5	8.20	108.68	105.40
22	BA	1303	G	N1-C6-O6	-8.18	114.99	119.90
22	BA	577	G	N1-C6-O6	8.17	124.80	119.90
22	BA	2837	A	O5'-P-OP1	-8.17	98.35	105.70
22	DA	1694	C	N1-C2-O2	8.16	123.80	118.90
22	BA	783	A	C6-C5-N7	-8.15	126.59	132.30
22	BA	1282	U	N3-C2-O2	8.15	127.91	122.20
22	BA	1617	C	N1-C2-O2	-8.15	114.01	118.90
23	BB	15	A	C4-C5-N7	8.13	114.77	110.70
22	BA	1293	C	N1-C2-O2	-8.11	114.03	118.90
22	BA	1677	A	C5-C6-N6	-8.11	117.21	123.70
22	BA	2001	C	C6-N1-C2	8.11	123.54	120.30
22	BA	446	G	C5-N7-C8	-8.10	100.25	104.30
22	BA	2626	C	C6-N1-C2	8.08	123.53	120.30
22	BA	1663	G	C8-N9-C4	8.03	109.61	106.40
22	BA	2814	A	N9-C4-C5	-8.04	102.59	105.80
22	BA	823	C	N1-C2-O2	-8.01	114.09	118.90
22	BA	2450	A	O5'-P-OP2	-8.01	98.49	105.70
22	BA	1653	G	N9-C4-C5	8.01	108.60	105.40
22	BA	101	A	C2-N3-C4	-8.00	106.60	110.60
22	BA	1229	C	N1-C2-O2	-7.99	114.11	118.90
22	BA	1426	G	C5-C6-O6	-7.98	123.81	128.60
22	BA	518	G	O5'-P-OP2	-7.97	98.52	105.70
22	BA	1357	C	N1-C2-O2	-7.97	114.12	118.90
1	AA	500	G	N1-C6-O6	7.96	124.67	119.90
22	BA	1134	A	O5'-P-OP1	-7.95	98.55	105.70
22	BA	861	A	OP1-P-O3'	7.92	122.62	105.20
22	BA	2070	A	C5-C6-N6	-7.90	117.38	123.70
22	BA	1779	U	N3-C4-O4	-7.89	113.88	119.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	1379	U	C5-C4-O4	-7.87	121.18	125.90
22	BA	2283	C	C6-N1-C2	7.87	123.45	120.30
22	BA	1040	A	N1-C6-N6	7.85	123.31	118.60
22	BA	1549	A	N1-C6-N6	7.84	123.31	118.60
22	BA	1293	C	N3-C2-O2	7.80	127.36	121.90
22	BA	747	U	C5-C4-O4	7.79	130.58	125.90
22	BA	2814	A	C2-N3-C4	-7.78	106.71	110.60
22	BA	1790	C	C2-N3-C4	-7.77	116.02	119.90
22	BA	784	G	P-O3'-C3'	7.77	129.02	119.70
25	BD	152	PRO	CA-N-CD	-7.76	100.63	111.50
22	BA	750	A	O5'-P-OP2	-7.76	98.72	105.70
22	BA	2083	G	N1-C6-O6	7.75	124.55	119.90
22	BA	261	G	O5'-P-OP2	-7.74	98.73	105.70
22	BA	1192	G	C5-C6-O6	7.74	133.25	128.60
22	BA	753	A	C5-C6-N1	7.74	121.57	117.70
22	BA	2442	C	N3-C4-C5	7.74	125.00	121.90
22	BA	831	G	C8-N9-C4	7.74	109.50	106.40
22	BA	2809	A	O5'-P-OP1	-7.71	98.76	105.70
22	BA	2387	U	O5'-P-OP2	-7.70	98.77	105.70
22	BA	520	G	C5-C6-O6	7.69	133.22	128.60
1	CA	919	A	O5'-P-OP2	-7.69	98.78	105.70
22	BA	2825	G	C8-N9-C4	-7.68	103.33	106.40
22	BA	577	G	C5-C6-O6	-7.64	124.02	128.60
22	BA	61	C	C2-N3-C4	-7.63	116.09	119.90
22	BA	1997	C	C6-N1-C2	7.63	123.35	120.30
22	BA	1287	A	C8-N9-C4	-7.62	102.75	105.80
22	BA	1271	G	C5-C6-N1	7.61	115.30	111.50
22	BA	1452	G	C2-N3-C4	-7.60	108.10	111.90
22	BA	1658	C	N3-C4-C5	7.59	124.94	121.90
22	BA	1614	A	C2-N3-C4	-7.59	106.80	110.60
22	BA	1394	U	O5'-P-OP1	-7.59	98.87	105.70
22	BA	2023	C	C6-N1-C2	7.59	123.34	120.30
22	BA	752	A	N1-C6-N6	7.58	123.15	118.60
22	BA	974	G	N7-C8-N9	7.56	116.88	113.10
28	BG	149	ARG	NE-CZ-NH1	7.56	124.08	120.30
22	BA	255	A	C2-N3-C4	-7.55	106.82	110.60
22	BA	1220	G	N3-C2-N2	7.55	125.19	119.90
22	BA	2884	U	N3-C4-O4	-7.54	114.12	119.40
22	BA	561	G	C4-C5-N7	7.51	113.81	110.80
22	BA	2744	G	N1-C6-O6	7.51	124.41	119.90
22	BA	1639	C	N1-C2-O2	-7.51	114.39	118.90
22	BA	528	A	C5-C6-N6	-7.51	117.69	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	CA	500	G	C4-C5-C6	-7.50	114.30	118.80
22	BA	2825	G	C4-N9-C1'	7.50	136.24	126.50
1	CA	500	G	C8-N9-C4	-7.49	103.40	106.40
22	BA	687	C	N1-C2-O2	-7.49	114.41	118.90
1	AA	500	G	C4-C5-N7	7.49	113.80	110.80
22	BA	2248	C	O5'-P-OP2	-7.49	98.96	105.70
22	BA	1663	G	N1-C6-O6	7.49	124.39	119.90
22	BA	1618	A	C5-C6-N6	7.48	129.69	123.70
22	BA	752	A	C4-C5-N7	7.48	114.44	110.70
22	BA	823	C	C2-N3-C4	-7.47	116.17	119.90
22	BA	1296	G	C8-N9-C4	7.47	109.39	106.40
22	BA	2091	C	N1-C2-O2	-7.46	114.42	118.90
23	BB	106	G	C5-C6-O6	-7.45	124.13	128.60
44	BW	39	ARG	NE-CZ-NH2	-7.43	116.58	120.30
23	BB	15	A	C5-C6-N6	-7.43	117.76	123.70
1	AA	500	G	C8-N9-C1'	7.42	136.65	127.00
22	BA	853	C	O5'-P-OP2	-7.42	99.03	105.70
22	BA	531	C	O4'-C1'-N1	-7.41	102.27	108.20
22	BA	1653	G	N3-C4-C5	-7.41	124.89	128.60
22	BA	2898	U	O5'-P-OP2	-7.41	99.03	105.70
22	BA	2030	A	C5-C6-N1	-7.40	114.00	117.70
22	BA	785	G	O5'-P-OP1	-7.40	99.04	105.70
22	BA	208	C	C6-N1-C2	7.38	123.25	120.30
22	BA	1679	A	N1-C6-N6	7.36	123.02	118.60
22	BA	579	G	N1-C6-O6	-7.36	115.48	119.90
22	BA	2324	U	C5-C4-O4	-7.36	121.49	125.90
22	BA	988	A	N1-C6-N6	7.35	123.01	118.60
22	BA	127	A	N1-C6-N6	7.34	123.00	118.60
22	BA	561	G	C6-C5-N7	-7.32	126.01	130.40
22	BA	2575	C	N1-C2-O2	7.32	123.29	118.90
22	BA	2633	G	C2-N3-C4	-7.31	108.25	111.90
22	BA	1227	G	C2-N3-C4	-7.30	108.25	111.90
22	BA	1271	G	N1-C6-O6	-7.30	115.52	119.90
22	BA	2689	U	N3-C4-O4	-7.28	114.30	119.40
22	BA	1669	A	C6-N1-C2	-7.28	114.23	118.60
22	BA	2049	G	N7-C8-N9	7.28	116.74	113.10
22	BA	528	A	N7-C8-N9	7.28	117.44	113.80
22	BA	914	G	C5-C6-O6	-7.28	124.23	128.60
22	BA	691	C	N1-C2-O2	-7.26	114.54	118.90
22	BA	2787	C	N1-C2-O2	-7.25	114.55	118.90
1	AA	365	U	C5-C4-O4	7.24	130.25	125.90
22	BA	2513	A	N9-C4-C5	7.24	108.70	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	2744	G	N9-C4-C5	-7.24	102.50	105.40
1	AA	365	U	C5-C6-N1	-7.23	119.08	122.70
22	BA	1303	G	C5-C6-O6	7.23	132.94	128.60
22	BA	534	U	N3-C4-O4	-7.23	114.34	119.40
22	BA	1784	A	C2-N3-C4	-7.23	106.99	110.60
23	BB	102	G	C5-C6-O6	-7.22	124.27	128.60
22	BA	1939	U	N3-C4-O4	-7.21	114.35	119.40
22	BA	2699	C	C6-N1-C2	7.20	123.18	120.30
22	BA	784	G	O4'-C1'-N9	-7.18	102.45	108.20
22	BA	2733	A	O5'-P-OP2	-7.18	99.23	105.70
22	BA	192	C	N1-C2-O2	-7.18	114.59	118.90
22	BA	1192	G	N1-C6-O6	-7.18	115.59	119.90
22	BA	213	A	N1-C6-N6	7.17	122.91	118.60
22	BA	2787	C	C5-C6-N1	-7.17	117.41	121.00
1	CA	35	G	O5'-P-OP1	-7.17	99.25	105.70
22	BA	1779	U	C5-C4-O4	7.16	130.19	125.90
22	BA	752	A	C5-C6-N1	-7.16	114.12	117.70
1	AA	365	U	C2-N1-C1'	-7.15	109.12	117.70
23	BB	70	C	N1-C2-O2	-7.15	114.61	118.90
22	BA	1294	U	O5'-P-OP2	-7.14	99.27	105.70
22	BA	2808	G	O5'-P-OP2	-7.14	99.27	105.70
22	BA	984	A	C5-C6-N1	-7.14	114.13	117.70
22	BA	2889	C	N3-C2-O2	7.13	126.89	121.90
22	BA	2505	G	C5-C6-O6	7.13	132.88	128.60
22	BA	961	C	C6-N1-C2	7.13	123.15	120.30
22	BA	1649	G	C8-N9-C4	-7.13	103.55	106.40
22	BA	1331	G	C8-N9-C4	-7.11	103.56	106.40
22	BA	249	C	N1-C2-O2	-7.10	114.64	118.90
22	BA	1220	G	N1-C2-N2	-7.10	109.81	116.20
22	BA	2211	A	P-O3'-C3'	7.09	128.21	119.70
22	BA	961	C	O5'-P-OP2	-7.09	99.32	105.70
22	BA	1452	G	C4-C5-N7	7.09	113.64	110.80
22	BA	1936	A	N3-C4-C5	7.09	131.76	126.80
22	BA	2250	G	C2-N3-C4	-7.08	108.36	111.90
22	BA	869	G	O5'-P-OP2	-7.08	99.33	105.70
51	B3	13	ARG	NE-CZ-NH2	7.08	123.84	120.30
23	BB	15	A	N9-C4-C5	-7.08	102.97	105.80
22	BA	1976	U	N1-C2-O2	-7.07	117.85	122.80
22	BA	1251	C	N3-C4-N4	7.07	122.95	118.00
22	BA	1296	G	O5'-P-OP2	-7.05	99.36	105.70
22	BA	801	G	N9-C4-C5	7.04	108.22	105.40
22	BA	984	A	O4'-C1'-N9	7.04	113.83	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	2053	G	N1-C6-O6	7.04	124.12	119.90
22	BA	1023	U	N1-C2-O2	-7.03	117.88	122.80
22	BA	1282	U	N1-C2-O2	-7.03	117.88	122.80
22	BA	1976	U	N3-C2-O2	7.03	127.12	122.20
41	BT	3	ARG	N-CA-C	-7.02	92.05	111.00
22	BA	312	G	N1-C6-O6	7.02	124.11	119.90
22	BA	758	C	C2-N3-C4	-7.01	116.39	119.90
22	BA	1296	G	O5'-P-OP1	7.01	119.12	110.70
22	BA	1319	C	N1-C2-O2	-7.00	114.70	118.90
22	BA	567	U	N1-C2-O2	-7.00	117.90	122.80
22	BA	2297	A	O5'-P-OP1	-7.00	99.41	105.70
22	BA	1656	C	N3-C4-C5	6.99	124.70	121.90
22	BA	745	G	N1-C2-N3	6.98	128.09	123.90
22	BA	2424	C	C2-N3-C4	-6.97	116.42	119.90
22	BA	1428	C	N1-C2-O2	-6.96	114.72	118.90
22	BA	2503	A	N1-C6-N6	6.96	122.78	118.60
22	BA	2820	A	C2-N3-C4	-6.96	107.12	110.60
22	BA	535	G	N1-C6-O6	-6.95	115.73	119.90
1	AA	124	C	N1-C2-O2	-6.94	114.73	118.90
22	BA	961	C	N3-C2-O2	6.94	126.76	121.90
22	BA	831	G	N1-C6-O6	6.93	124.06	119.90
22	BA	2814	A	C5-C6-N6	-6.93	118.16	123.70
22	BA	1936	A	N1-C2-N3	6.92	132.76	129.30
22	BA	2476	A	O5'-P-OP2	-6.92	99.47	105.70
22	BA	577	G	C6-C5-N7	-6.91	126.25	130.40
22	BA	32	C	C2-N1-C1'	-6.90	111.21	118.80
22	BA	1279	G	N1-C6-O6	-6.90	115.76	119.90
22	BA	956	G	N1-C6-O6	6.89	124.03	119.90
22	BA	61	C	N3-C4-C5	6.89	124.66	121.90
22	BA	672	C	N3-C4-C5	6.88	124.65	121.90
22	BA	698	C	C6-N1-C2	6.87	123.05	120.30
22	BA	944	C	O5'-P-OP1	6.87	118.94	110.70
22	BA	2679	A	C5-C6-N6	-6.86	118.21	123.70
22	BA	2529	G	C5-C6-O6	-6.86	124.49	128.60
22	BA	2352	A	O5'-P-OP1	-6.86	99.53	105.70
22	BA	1313	U	C2-N1-C1'	6.85	125.92	117.70
22	BA	975	A	O5'-P-OP1	-6.84	99.54	105.70
1	AA	919	A	O5'-P-OP2	-6.84	99.54	105.70
22	BA	16	C	O5'-P-OP2	-6.84	99.55	105.70
22	BA	1020	A	C4-C5-N7	6.83	114.12	110.70
22	BA	835	C	C6-N1-C2	6.83	123.03	120.30
22	BA	61	C	N1-C2-O2	-6.83	114.80	118.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	1387	A	N1-C6-N6	6.82	122.69	118.60
22	BA	1284	A	N1-C6-N6	6.82	122.69	118.60
1	AA	1509	C	C6-N1-C2	6.82	123.03	120.30
22	BA	2018	G	C4-C5-N7	-6.80	108.08	110.80
22	BA	520	G	N1-C6-O6	-6.79	115.83	119.90
34	BM	20	LEU	CA-CB-CG	6.79	130.92	115.30
1	AA	1071	C	N1-C2-O2	-6.79	114.83	118.90
22	BA	515	A	N1-C6-N6	-6.79	114.53	118.60
22	BA	454	A	O5'-P-OP2	-6.79	99.59	105.70
22	BA	1957	C	N3-C4-C5	6.79	124.61	121.90
22	BA	255	A	O5'-P-OP1	-6.78	99.59	105.70
22	BA	2608	G	C5-C6-N1	6.78	114.89	111.50
1	CA	575	G	N3-C4-C5	6.78	131.99	128.60
22	BA	1670	C	N1-C2-O2	-6.78	114.83	118.90
22	BA	567	U	N3-C2-O2	6.78	126.94	122.20
22	BA	581	C	N1-C2-O2	-6.77	114.84	118.90
22	BA	1332	G	C4-C5-N7	6.77	113.51	110.80
22	BA	945	A	C8-N9-C4	6.77	108.51	105.80
22	BA	1663	G	C4-C5-N7	6.77	113.51	110.80
22	BA	180	G	C4-N9-C1'	-6.76	117.71	126.50
22	BA	1569	A	O5'-P-OP1	-6.76	99.62	105.70
22	BA	717	C	N1-C2-O2	-6.75	114.85	118.90
22	BA	1658	C	N1-C2-O2	-6.75	114.85	118.90
22	BA	2520	C	O5'-P-OP1	-6.75	99.63	105.70
22	BA	521	U	N1-C2-N3	6.74	118.94	114.90
22	BA	2393	U	O5'-P-OP2	-6.74	99.64	105.70
22	BA	1663	G	N3-C4-C5	6.73	131.97	128.60
1	AA	1530	G	C4-N9-C1'	-6.73	117.75	126.50
22	BA	1296	G	N7-C8-N9	-6.73	109.74	113.10
22	BA	705	A	N1-C6-N6	6.73	122.64	118.60
22	BA	752	A	C6-C5-N7	-6.72	127.59	132.30
22	BA	1299	G	C6-C5-N7	-6.72	126.37	130.40
22	BA	1669	A	C5-C6-N1	6.72	121.06	117.70
22	BA	2009	A	C8-N9-C4	-6.71	103.11	105.80
22	BA	1595	C	N1-C2-O2	-6.71	114.87	118.90
22	BA	1609	A	C4-C5-C6	6.71	120.36	117.00
22	DA	741	U	N3-C2-O2	6.71	126.90	122.20
22	BA	1251	C	C6-N1-C1'	-6.71	112.75	120.80
22	BA	2824	C	N3-C4-C5	-6.71	119.22	121.90
22	BA	2480	C	N1-C2-O2	-6.70	114.88	118.90
22	BA	32	C	C6-N1-C1'	6.70	128.84	120.80
22	BA	1681	G	C5-C6-O6	-6.70	124.58	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	2635	A	C8-N9-C4	-6.69	103.12	105.80
22	BA	93	G	N3-C2-N2	-6.69	115.22	119.90
1	AA	500	G	C5-C6-O6	-6.68	124.59	128.60
22	BA	561	G	N3-C4-C5	6.68	131.94	128.60
22	BA	1326	U	N1-C2-O2	-6.68	118.13	122.80
22	BA	1985	C	C5-C6-N1	-6.67	117.66	121.00
22	BA	1301	A	N9-C4-C5	-6.67	103.13	105.80
22	BA	831	G	N3-C4-N9	6.67	130.00	126.00
22	BA	1783	A	C4-C5-C6	6.67	120.33	117.00
22	BA	729	G	O5'-P-OP1	-6.66	99.70	105.70
22	BA	1985	C	C2-N1-C1'	-6.66	111.47	118.80
22	BA	752	A	O4'-C1'-N9	6.66	113.53	108.20
1	CA	857	C	N1-C2-O2	-6.66	114.91	118.90
22	BA	1609	A	N1-C6-N6	6.65	122.59	118.60
22	BA	127	A	C5-C6-N6	-6.65	118.38	123.70
22	BA	1251	C	C2-N3-C4	-6.64	116.58	119.90
22	BA	2252	G	N1-C6-O6	6.63	123.88	119.90
22	BA	2019	A	O5'-P-OP2	-6.63	99.73	105.70
22	BA	1784	A	C8-N9-C4	6.62	108.45	105.80
22	BA	462	C	O5'-P-OP2	-6.62	99.74	105.70
22	BA	2787	C	N3-C4-C5	6.62	124.55	121.90
22	BA	1231	U	C5-C4-O4	-6.61	121.93	125.90
22	BA	2525	G	OP2-P-O3'	6.61	119.75	105.20
22	BA	2788	C	N1-C2-O2	-6.61	114.93	118.90
22	BA	1779	U	N3-C2-O2	-6.60	117.58	122.20
22	BA	1022	G	C6-N1-C2	-6.59	121.14	125.10
22	BA	1692	U	C6-N1-C2	6.59	124.95	121.00
1	AA	1509	C	N1-C2-O2	-6.58	114.95	118.90
22	BA	587	C	N1-C2-O2	-6.58	114.95	118.90
22	BA	1653	G	C5-C6-O6	6.58	132.55	128.60
22	BA	1790	C	N1-C2-O2	-6.58	114.95	118.90
22	BA	2572	A	OP1-P-OP2	6.58	129.47	119.60
22	BA	451	U	C2-N1-C1'	-6.58	109.81	117.70
22	BA	19	A	C4-C5-C6	6.58	120.29	117.00
22	BA	1659	G	N1-C6-O6	-6.57	115.96	119.90
22	BA	2894	G	C8-N9-C1'	6.57	135.53	127.00
22	BA	533	G	C4-C5-N7	6.55	113.42	110.80
22	BA	1708	C	C6-N1-C2	6.55	122.92	120.30
22	BA	974	G	O4'-C1'-N9	6.54	113.44	108.20
22	BA	1348	C	O5'-P-OP1	-6.54	99.82	105.70
22	BA	2376	A	C2-N3-C4	-6.53	107.33	110.60
22	BA	1676	A	N1-C6-N6	-6.53	114.68	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	2342	C	O5'-P-OP1	-6.53	99.82	105.70
22	BA	1223	G	O5'-P-OP1	-6.52	99.83	105.70
22	BA	1020	A	O5'-P-OP2	-6.52	99.83	105.70
22	BA	1679	A	C6-C5-N7	-6.51	127.74	132.30
22	BA	2825	G	N3-C4-N9	6.51	129.91	126.00
22	BA	752	A	N3-C4-C5	6.51	131.35	126.80
22	BA	2858	C	C2-N3-C4	-6.51	116.65	119.90
22	BA	1297	C	N1-C2-O2	-6.50	115.00	118.90
22	BA	2362	C	N1-C2-O2	-6.50	115.00	118.90
22	BA	2023	C	N1-C2-O2	-6.49	115.00	118.90
22	DA	776	G	C4-N9-C1'	6.49	134.94	126.50
22	BA	1128	G	O5'-P-OP1	-6.49	99.86	105.70
22	BA	1132	U	N1-C2-O2	-6.49	118.26	122.80
22	BA	536	G	C5-C6-O6	-6.48	124.71	128.60
22	BA	974	G	C8-N9-C4	-6.48	103.81	106.40
22	BA	567	U	C5-C4-O4	-6.48	122.01	125.90
22	BA	655	A	C5-C6-N6	6.48	128.88	123.70
22	BA	2286	G	C2-N3-C4	-6.48	108.66	111.90
22	BA	400	G	N1-C6-O6	6.47	123.78	119.90
22	BA	2283	C	C2-N1-C1'	-6.47	111.68	118.80
22	BA	2487	G	C5-C6-O6	-6.47	124.72	128.60
22	BA	535	G	O5'-P-OP1	6.47	118.47	110.70
22	DA	1819	A	O5'-P-OP2	-6.47	99.88	105.70
22	BA	786	C	N3-C4-C5	6.47	124.49	121.90
22	BA	1167	C	O5'-P-OP2	-6.47	99.88	105.70
1	CA	624	C	O5'-P-OP2	-6.46	99.88	105.70
22	BA	2787	C	C6-N1-C2	6.46	122.88	120.30
22	BA	1300	G	C5-C6-O6	-6.46	124.73	128.60
22	BA	731	C	N1-C2-O2	-6.46	115.03	118.90
1	CA	288	A	N1-C6-N6	6.45	122.47	118.60
22	BA	2218	G	N1-C6-O6	-6.45	116.03	119.90
22	DA	1983	G	C8-N9-C4	-6.44	103.82	106.40
40	BS	19	LEU	CB-CG-CD2	-6.44	100.05	111.00
22	BA	1695	G	N3-C4-N9	6.44	129.86	126.00
22	BA	2005	A	C4-C5-N7	6.43	113.92	110.70
22	BA	991	C	C6-N1-C2	-6.43	117.73	120.30
22	BA	1665	A	C2-N3-C4	-6.43	107.39	110.60
22	BA	26	G	N3-C4-N9	6.42	129.85	126.00
22	BA	2427	C	N1-C2-O2	-6.42	115.05	118.90
22	BA	1779	U	C2-N3-C4	-6.42	123.15	127.00
22	BA	1932	A	O5'-P-OP1	-6.42	99.92	105.70
23	BB	90	C	OP1-P-OP2	6.42	129.23	119.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	2033	A	O4'-C1'-N9	6.41	113.33	108.20
1	AA	11	G	O5'-P-OP1	-6.41	99.93	105.70
1	AA	500	G	C8-N9-C4	-6.41	103.84	106.40
22	BA	561	G	N9-C4-C5	-6.40	102.84	105.40
22	BA	2368	C	N1-C2-O2	-6.40	115.06	118.90
22	BA	2820	A	C8-N9-C4	6.40	108.36	105.80
22	BA	704	G	O4'-C1'-N9	6.40	113.32	108.20
22	BA	954	G	OP2-P-O3'	6.40	119.28	105.20
22	BA	1272	A	C2-N3-C4	-6.40	107.40	110.60
22	DA	1677	A	N1-C6-N6	6.40	122.44	118.60
22	BA	964	C	O5'-P-OP1	-6.39	99.95	105.70
22	BA	527	C	C2-N1-C1'	-6.39	111.77	118.80
22	BA	2049	G	C5-N7-C8	-6.39	101.11	104.30
22	BA	2286	G	N3-C4-C5	6.39	131.79	128.60
22	BA	2762	C	N1-C2-O2	-6.39	115.07	118.90
22	BA	2760	C	C5-C6-N1	-6.38	117.81	121.00
22	BA	2760	C	C2-N1-C1'	-6.38	111.78	118.80
23	BB	15	A	C6-C5-N7	-6.38	127.83	132.30
1	AA	571	U	N1-C2-O2	-6.38	118.34	122.80
22	BA	84	A	N1-C6-N6	-6.38	114.77	118.60
22	BA	980	A	C5-N7-C8	-6.38	100.71	103.90
22	BA	2000	C	OP2-P-O3'	6.38	119.22	105.20
22	BA	1323	C	N3-C4-C5	6.37	124.45	121.90
1	CA	1499	A	N1-C6-N6	6.37	122.42	118.60
22	BA	1121	C	C2-N3-C4	-6.37	116.72	119.90
22	BA	1319	C	N3-C2-O2	6.37	126.36	121.90
22	BA	1976	U	O5'-P-OP2	-6.37	99.97	105.70
22	BA	1775	U	C2-N3-C4	-6.36	123.18	127.00
22	BA	2250	G	N1-C6-O6	6.36	123.72	119.90
22	DA	2240	U	N3-C2-O2	-6.36	117.75	122.20
22	BA	1297	C	OP1-P-OP2	-6.35	110.07	119.60
22	BA	681	G	OP2-P-O3'	6.35	119.16	105.20
22	BA	1142	A	C8-N9-C1'	6.34	139.12	127.70
1	CA	500	G	C4-N9-C1'	-6.34	118.26	126.50
22	BA	1788	C	C5-C4-N4	-6.33	115.77	120.20
22	BA	2565	A	C8-N9-C4	6.33	108.33	105.80
22	BA	2265	U	O5'-P-OP1	-6.33	100.00	105.70
22	BA	1156	A	C5-C6-N6	-6.33	118.64	123.70
22	BA	2071	A	C2-N3-C4	-6.33	107.44	110.60
22	BA	781	A	C5-C6-N1	6.32	120.86	117.70
1	CA	869	G	O5'-P-OP1	-6.32	100.01	105.70
22	DA	798	G	C8-N9-C4	-6.32	103.87	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	1261	C	N3-C4-N4	-6.32	113.57	118.00
22	BA	2503	A	C5-C6-N6	-6.32	118.65	123.70
22	BA	783	A	N3-C4-N9	-6.31	122.35	127.40
22	DA	1779	U	C5-C4-O4	6.31	129.69	125.90
22	BA	1022	G	N3-C4-C5	-6.30	125.45	128.60
23	BB	92	C	C6-N1-C2	6.30	122.82	120.30
1	AA	279	A	C4-C5-N7	6.29	113.85	110.70
22	BA	1948	G	C5-C6-O6	6.29	132.37	128.60
22	BA	1543	G	C8-N9-C4	-6.28	103.89	106.40
22	BA	2505	G	C6-C5-N7	6.28	134.17	130.40
22	BA	2502	G	C4-C5-N7	-6.28	108.29	110.80
23	BB	93	C	O5'-P-OP1	6.28	118.23	110.70
22	BA	557	C	OP1-P-OP2	6.27	129.01	119.60
1	AA	815	A	O5'-P-OP2	-6.27	100.06	105.70
22	BA	527	C	C6-N1-C1'	6.27	128.32	120.80
22	BA	2814	A	C4-C5-N7	6.27	113.83	110.70
1	AA	1530	G	N3-C4-C5	6.26	131.73	128.60
1	AA	1523	G	C4-C5-N7	-6.25	108.30	110.80
22	BA	213	A	N9-C4-C5	-6.25	103.30	105.80
22	BA	823	C	N3-C4-C5	6.25	124.40	121.90
22	BA	817	C	N1-C2-O2	-6.24	115.15	118.90
22	BA	2839	G	OP2-P-O3'	6.23	118.91	105.20
1	AA	279	A	C5-N7-C8	-6.23	100.78	103.90
22	BA	684	G	C5-N7-C8	-6.23	101.19	104.30
22	BA	1394	U	O5'-P-OP2	6.23	118.17	110.70
22	BA	2820	A	N1-C6-N6	6.23	122.34	118.60
1	AA	365	U	N3-C4-O4	-6.22	115.04	119.40
22	BA	743	A	O5'-P-OP2	-6.22	100.10	105.70
1	AA	1279	G	C8-N9-C4	-6.22	103.91	106.40
22	BA	761	A	N1-C6-N6	-6.22	114.87	118.60
22	DA	776	G	N3-C4-N9	6.22	129.73	126.00
22	BA	2644	G	C5-C6-O6	-6.21	124.87	128.60
22	BA	2866	U	N3-C2-O2	-6.21	117.85	122.20
22	BA	489	G	N3-C2-N2	-6.21	115.55	119.90
22	BA	193	U	O5'-P-OP1	6.21	118.15	110.70
22	BA	575	A	O5'-P-OP1	-6.21	100.11	105.70
22	BA	867	C	N1-C2-O2	-6.21	115.18	118.90
22	BA	1311	G	C5-N7-C8	-6.21	101.20	104.30
22	BA	2875	C	N3-C4-N4	6.21	122.34	118.00
22	BA	1948	G	N1-C6-O6	-6.20	116.18	119.90
22	BA	2279	G	N1-C6-O6	-6.20	116.18	119.90
1	AA	889	A	O5'-P-OP2	-6.20	100.12	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	517	C	O5'-P-OP2	-6.19	100.13	105.70
1	AA	914	A	C8-N9-C4	-6.19	103.33	105.80
22	BA	2076	U	N3-C2-O2	-6.19	117.87	122.20
22	BA	665	U	C6-N1-C2	6.18	124.71	121.00
1	CA	575	G	C4-N9-C1'	-6.18	118.46	126.50
22	BA	1663	G	C6-C5-N7	-6.18	126.69	130.40
22	BA	1653	G	C4-C5-N7	-6.18	108.33	110.80
22	BA	673	C	N3-C2-O2	6.17	126.22	121.90
22	BA	752	A	N3-C4-N9	-6.17	122.46	127.40
1	CA	207	C	C2-N1-C1'	6.17	125.59	118.80
22	BA	575	A	O5'-P-OP2	6.17	118.11	110.70
22	BA	834	G	C5-C6-O6	-6.17	124.90	128.60
22	BA	1566	A	C6-C5-N7	-6.17	127.98	132.30
22	BA	1576	U	OP2-P-O3'	6.17	118.77	105.20
22	BA	512	G	C8-N9-C4	-6.17	103.93	106.40
23	BB	79	G	O5'-P-OP1	6.17	118.10	110.70
22	BA	2505	G	N9-C4-C5	6.16	107.87	105.40
22	BA	1649	G	N9-C4-C5	6.16	107.86	105.40
22	BA	1658	C	C2-N3-C4	-6.16	116.82	119.90
22	BA	1758	U	C5-C6-N1	-6.16	119.62	122.70
22	BA	630	G	C5-C6-O6	-6.16	124.91	128.60
22	BA	595	C	N1-C2-O2	-6.16	115.21	118.90
22	BA	2816	G	O5'-P-OP2	-6.15	100.16	105.70
22	BA	2777	G	C5-C6-O6	6.15	132.29	128.60
22	BA	474	G	N3-C4-C5	-6.15	125.53	128.60
22	BA	2385	C	C6-N1-C2	6.15	122.76	120.30
22	BA	728	G	N1-C6-O6	-6.15	116.21	119.90
22	BA	1934	C	N3-C4-N4	-6.14	113.70	118.00
22	BA	748	G	O4'-C1'-N9	6.14	113.11	108.20
1	CA	575	G	N3-C4-N9	-6.14	122.31	126.00
22	BA	1142	A	N1-C6-N6	6.14	122.28	118.60
22	BA	1758	U	N1-C2-N3	6.14	118.58	114.90
22	BA	515	A	N9-C4-C5	6.14	108.25	105.80
22	BA	853	C	O5'-P-OP1	6.14	118.06	110.70
1	CA	18	C	O5'-P-OP1	-6.14	100.18	105.70
22	BA	942	G	C5-C6-O6	-6.13	124.92	128.60
22	BA	203	A	N1-C6-N6	6.13	122.28	118.60
22	BA	1609	A	C6-C5-N7	-6.13	128.01	132.30
22	BA	2744	G	C4-C5-N7	6.13	113.25	110.80
22	BA	1027	A	C4-C5-C6	6.13	120.06	117.00
22	BA	2815	C	N3-C4-C5	6.13	124.35	121.90
22	BA	704	G	C5-C6-O6	-6.12	124.92	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	845	A	N1-C6-N6	6.12	122.28	118.60
22	BA	1635	A	O5'-P-OP2	-6.12	100.19	105.70
22	BA	993	G	C5-C6-O6	6.12	132.27	128.60
22	BA	1930	G	C4-N9-C1'	-6.12	118.55	126.50
22	BA	2196	C	N1-C2-O2	-6.12	115.23	118.90
22	BA	2208	C	O5'-P-OP2	6.12	118.04	110.70
22	BA	2046	G	N1-C6-O6	-6.11	116.23	119.90
22	BA	2588	G	C8-N9-C4	6.11	108.84	106.40
22	BA	2718	G	O5'-P-OP1	-6.11	100.20	105.70
22	DA	246	C	C6-N1-C2	6.10	122.74	120.30
22	BA	1309	G	N1-C6-O6	-6.10	116.24	119.90
22	BA	1692	U	N1-C2-O2	-6.10	118.53	122.80
22	DA	692	C	C6-N1-C2	-6.10	117.86	120.30
22	BA	752	A	N9-C1'-C2'	6.10	121.92	114.00
22	BA	990	A	N1-C2-N3	6.09	132.34	129.30
22	BA	1476	U	OP2-P-O3'	6.09	118.60	105.20
22	BA	2805	C	C2-N1-C1'	-6.09	112.10	118.80
22	BA	1274	A	N1-C6-N6	6.09	122.25	118.60
22	BA	2281	A	O5'-P-OP2	6.08	118.00	110.70
1	CA	244	U	N3-C2-O2	-6.08	117.94	122.20
22	BA	1326	U	C5-C4-O4	-6.08	122.25	125.90
22	BA	1253	A	O5'-P-OP2	-6.08	100.23	105.70
22	BA	2631	G	O5'-P-OP2	-6.08	100.23	105.70
22	BA	2832	U	N1-C2-O2	-6.08	118.55	122.80
22	DA	2455	G	C8-N9-C4	-6.08	103.97	106.40
22	BA	2595	G	OP1-P-O3'	6.07	118.56	105.20
22	BA	1114	C	C6-N1-C2	6.07	122.73	120.30
22	BA	2468	A	N1-C6-N6	6.07	122.24	118.60
22	BA	2811	G	C8-N9-C4	6.07	108.83	106.40
22	BA	1002	G	C4-C5-N7	-6.07	108.37	110.80
22	BA	942	G	C8-N9-C4	6.06	108.83	106.40
22	BA	1819	A	O5'-P-OP1	-6.06	100.24	105.70
22	BA	1771	C	O5'-P-OP1	-6.06	100.25	105.70
22	BA	2811	G	O5'-P-OP2	-6.06	100.25	105.70
22	BA	2831	G	N1-C6-O6	6.05	123.53	119.90
22	BA	2815	C	C6-N1-C2	6.05	122.72	120.30
22	BA	1263	U	N3-C4-C5	6.05	118.23	114.60
22	BA	1758	U	C2-N3-C4	-6.05	123.37	127.00
22	BA	2837	A	O5'-P-OP2	6.05	117.96	110.70
22	BA	1985	C	C6-N1-C2	6.04	122.72	120.30
22	BA	655	A	N1-C6-N6	-6.04	114.97	118.60
22	BA	1265	A	OP1-P-O3'	6.04	118.49	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	1667	G	C4-C5-N7	6.04	113.22	110.80
22	BA	1327	A	OP1-P-OP2	-6.04	110.54	119.60
22	BA	1692	U	C5-C4-O4	-6.04	122.28	125.90
22	BA	836	G	C5-C6-O6	-6.04	124.98	128.60
22	BA	1981	A	C6-N1-C2	6.04	122.22	118.60
22	BA	1548	A	N1-C6-N6	6.04	122.22	118.60
22	BA	786	C	C2-N3-C4	-6.03	116.88	119.90
22	BA	2760	C	C2-N3-C4	-6.03	116.88	119.90
22	BA	1469	A	N1-C6-N6	6.03	122.22	118.60
22	BA	2606	C	C5-C6-N1	-6.03	117.98	121.00
22	BA	2820	A	N9-C4-C5	-6.03	103.39	105.80
1	AA	716	A	OP2-P-O3'	6.03	118.47	105.20
22	BA	579	G	C5-C6-O6	6.03	132.22	128.60
22	BA	2270	A	OP2-P-O3'	6.03	118.47	105.20
22	BA	2572	A	O5'-P-OP2	-6.03	100.27	105.70
22	BA	561	G	C2-N3-C4	-6.03	108.89	111.90
22	BA	596	U	N1-C2-O2	-6.03	118.58	122.80
22	BA	1681	G	N1-C6-O6	6.03	123.52	119.90
22	BA	1332	G	C5-C6-O6	-6.03	124.98	128.60
22	BA	2799	A	N1-C6-N6	6.02	122.22	118.60
22	BA	984	A	C4-N9-C1'	-6.02	115.46	126.30
22	BA	783	A	C5-C6-N1	-6.02	114.69	117.70
22	BA	2870	C	OP2-P-O3'	6.02	118.44	105.20
22	BA	997	G	C5-C6-O6	6.01	132.21	128.60
22	BA	1022	G	N9-C4-C5	6.01	107.81	105.40
22	BA	1639	C	N3-C2-O2	6.01	126.11	121.90
22	BA	2283	C	N3-C2-O2	6.01	126.11	121.90
22	BA	905	A	N9-C4-C5	6.01	108.20	105.80
22	BA	2326	C	C6-N1-C2	-6.01	117.90	120.30
22	BA	203	A	C5-C6-N6	-6.01	118.89	123.70
22	BA	2757	A	N1-C6-N6	6.01	122.20	118.60
22	BA	1650	A	C8-N9-C4	-6.00	103.40	105.80
22	BA	1975	G	O5'-P-OP1	-6.00	100.30	105.70
1	CA	1531	A	N1-C6-N6	6.00	122.20	118.60
22	BA	2777	G	N9-C4-C5	6.00	107.80	105.40
22	BA	723	C	C6-N1-C2	6.00	122.70	120.30
22	BA	1437	C	N1-C2-O2	-6.00	115.30	118.90
22	BA	1976	U	N3-C4-C5	6.00	118.20	114.60
22	BA	2091	C	C2-N3-C4	-6.00	116.90	119.90
22	BA	2825	G	C6-C5-N7	-6.00	126.80	130.40
1	CA	769	G	O5'-P-OP2	-5.99	100.31	105.70
1	CA	1528	U	O5'-P-OP2	-5.99	100.31	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	1218	G	N3-C4-N9	-5.99	122.41	126.00
22	BA	2580	U	OP2-P-O3'	5.99	118.37	105.20
22	BA	1319	C	C6-N1-C2	5.98	122.69	120.30
22	BA	2824	C	C6-N1-C2	-5.98	117.91	120.30
22	BA	1946	U	N3-C2-O2	-5.98	118.02	122.20
22	BA	2070	A	N9-C4-C5	-5.97	103.41	105.80
22	BA	2420	C	N1-C2-O2	-5.97	115.31	118.90
22	BA	2393	U	N3-C4-O4	5.97	123.58	119.40
22	BA	839	U	O5'-P-OP2	-5.97	100.33	105.70
22	BA	1187	G	C6-C5-N7	-5.97	126.82	130.40
22	BA	2382	G	N3-C4-N9	5.96	129.58	126.00
22	BA	2389	G	C8-N9-C4	-5.96	104.01	106.40
22	BA	572	A	C5-N7-C8	-5.96	100.92	103.90
1	AA	1468	A	C8-N9-C4	5.96	108.19	105.80
22	BA	2499	C	N3-C2-O2	5.96	126.07	121.90
22	BA	180	G	N3-C4-C5	5.96	131.58	128.60
22	BA	1785	A	C4-C5-C6	5.95	119.98	117.00
22	BA	2427	C	O5'-P-OP1	-5.95	100.34	105.70
22	BA	970	U	OP2-P-O3'	5.95	118.29	105.20
22	BA	2522	U	O5'-P-OP2	-5.95	100.34	105.70
22	DA	1815	A	O5'-P-OP2	-5.95	100.34	105.70
22	BA	2449	U	C2-N3-C4	-5.95	123.43	127.00
22	BA	1227	G	N1-C2-N3	5.95	127.47	123.90
22	BA	1689	A	OP2-P-O3'	5.94	118.28	105.20
22	BA	2503	A	N9-C4-C5	-5.94	103.42	105.80
22	BA	2720	U	N1-C2-O2	-5.94	118.64	122.80
22	BA	2005	A	N9-C4-C5	-5.94	103.43	105.80
22	BA	1142	A	C5-C6-N1	-5.93	114.73	117.70
22	BA	328	U	O5'-P-OP1	-5.93	100.36	105.70
22	BA	862	G	OP1-P-O3'	5.93	118.25	105.20
22	BA	2038	G	N1-C6-O6	5.93	123.46	119.90
22	BA	2469	A	C8-N9-C4	5.93	108.17	105.80
1	CA	287	U	C6-N1-C2	5.93	124.56	121.00
22	BA	731	C	N1-C2-N3	5.92	123.35	119.20
22	BA	585	G	C5-C6-O6	-5.92	125.05	128.60
22	BA	1983	G	N1-C6-O6	-5.92	116.35	119.90
22	BA	2059	A	N1-C6-N6	5.92	122.15	118.60
23	BB	107	G	C6-C5-N7	-5.92	126.85	130.40
22	BA	1231	U	N3-C2-O2	5.92	126.34	122.20
22	BA	1900	A	O5'-P-OP1	-5.92	100.37	105.70
23	BB	106	G	C4-C5-N7	5.92	113.17	110.80
22	BA	1513	U	N3-C4-O4	-5.91	115.26	119.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	1776	G	C8-N9-C1'	-5.91	119.31	127.00
22	BA	1009	A	N1-C6-N6	5.91	122.15	118.60
22	BA	19	A	C6-C5-N7	-5.91	128.16	132.30
22	BA	31	C	O5'-P-OP2	-5.91	100.38	105.70
22	BA	474	G	N1-C6-O6	-5.90	116.36	119.90
22	BA	930	G	O5'-P-OP2	-5.90	100.39	105.70
22	BA	2250	G	O4'-C1'-N9	-5.90	103.48	108.20
22	BA	2590	A	OP1-P-OP2	-5.90	110.75	119.60
22	DA	1373	A	O5'-P-OP1	-5.90	100.39	105.70
1	AA	1080	A	N1-C6-N6	-5.90	115.06	118.60
22	BA	565	C	N3-C2-O2	5.90	126.03	121.90
22	BA	595	C	N3-C2-O2	5.90	126.03	121.90
22	BA	2870	C	N3-C4-C5	5.90	124.26	121.90
22	BA	312	G	C5-C6-O6	-5.90	125.06	128.60
22	BA	455	C	N3-C2-O2	-5.90	117.77	121.90
22	BA	1284	A	C5-C6-N6	-5.90	118.98	123.70
22	BA	2388	A	OP1-P-OP2	-5.90	110.75	119.60
22	BA	998	C	C6-N1-C2	-5.89	117.94	120.30
22	BA	1677	A	N9-C4-C5	-5.89	103.44	105.80
22	BA	1976	U	C2-N3-C4	-5.89	123.46	127.00
22	BA	554	U	O5'-P-OP1	5.89	117.77	110.70
22	BA	1348	C	C5-C4-N4	-5.89	116.08	120.20
22	BA	1339	G	C5-C6-O6	-5.89	125.06	128.60
22	BA	2442	C	C5-C4-N4	-5.89	116.08	120.20
1	CA	572	A	O5'-P-OP2	-5.89	100.40	105.70
22	BA	1379	U	C6-N1-C2	5.88	124.53	121.00
22	BA	1679	A	C4-C5-C6	5.88	119.94	117.00
22	BA	512	G	O4'-C1'-N9	5.88	112.90	108.20
22	BA	678	C	C5-C4-N4	-5.88	116.09	120.20
22	BA	2063	C	N3-C4-C5	5.88	124.25	121.90
22	BA	521	U	C6-N1-C2	-5.87	117.48	121.00
22	BA	2814	A	C6-C5-N7	-5.87	128.19	132.30
22	BA	1230	A	O5'-P-OP2	-5.87	100.42	105.70
22	BA	2383	G	C4-C5-C6	-5.87	115.28	118.80
1	AA	323	U	N3-C2-O2	5.87	126.31	122.20
22	BA	997	G	OP1-P-O3'	5.87	118.11	105.20
22	BA	1253	A	O4'-C1'-N9	-5.87	103.51	108.20
22	BA	2894	G	C4-N9-C1'	-5.87	118.87	126.50
22	BA	537	G	C5-C6-O6	-5.87	125.08	128.60
22	BA	589	U	C5-C4-O4	-5.86	122.38	125.90
22	BA	2012	G	C5-C6-O6	-5.86	125.08	128.60
22	BA	2630	G	O5'-P-OP2	-5.86	100.43	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	33	C	N1-C2-O2	-5.86	115.39	118.90
22	BA	1959	G	C4-C5-N7	-5.86	108.46	110.80
22	BA	2520	C	N1-C2-O2	-5.86	115.39	118.90
22	BA	2693	G	N3-C2-N2	5.85	124.00	119.90
22	BA	2836	U	OP1-P-O3'	5.85	118.08	105.20
22	BA	1319	C	O5'-P-OP1	-5.85	100.43	105.70
22	BA	2782	G	N3-C4-N9	5.85	129.51	126.00
22	BA	1138	G	C8-N9-C4	-5.84	104.06	106.40
22	BA	2333	A	O5'-P-OP2	-5.84	100.44	105.70
22	BA	2343	U	O5'-P-OP1	-5.84	100.44	105.70
22	BA	1466	U	O5'-P-OP2	-5.84	100.44	105.70
22	DA	2603	G	C5-C6-O6	-5.84	125.10	128.60
1	AA	124	C	N3-C2-O2	5.84	125.99	121.90
22	BA	2286	G	N3-C4-N9	-5.84	122.50	126.00
22	BA	116	C	O5'-P-OP2	-5.83	100.45	105.70
22	BA	726	G	C8-N9-C4	5.83	108.73	106.40
1	AA	124	C	C2-N1-C1'	-5.83	112.39	118.80
22	BA	814	C	N3-C4-N4	-5.83	113.92	118.00
22	BA	1992	G	N1-C2-N3	5.82	127.39	123.90
22	BA	790	U	P-O3'-C3'	-5.82	112.72	119.70
22	BA	2070	A	N1-C6-N6	5.82	122.09	118.60
22	BA	2253	G	C6-C5-N7	-5.81	126.91	130.40
23	BB	102	G	N1-C6-O6	5.81	123.39	119.90
22	BA	1779	U	N1-C2-N3	5.81	118.39	114.90
22	BA	595	C	C5-C4-N4	-5.81	116.13	120.20
22	BA	1294	U	OP1-P-OP2	5.81	128.31	119.60
1	AA	811	C	N1-C2-O2	-5.81	115.42	118.90
22	BA	2379	G	N1-C6-O6	5.81	123.38	119.90
22	BA	190	A	N1-C6-N6	5.80	122.08	118.60
22	BA	957	C	OP2-P-O3'	5.80	117.96	105.20
22	DA	1937	A	O4'-C1'-N9	5.80	112.84	108.20
22	BA	1514	G	C5-C6-O6	-5.80	125.12	128.60
22	BA	1676	A	N9-C4-C5	5.80	108.12	105.80
22	BA	2276	G	O5'-P-OP1	-5.80	100.48	105.70
1	CA	1406	U	O5'-P-OP1	5.80	117.66	110.70
1	CA	716	A	OP2-P-O3'	5.80	117.95	105.20
22	BA	2069	G	N3-C4-N9	-5.79	122.52	126.00
22	BA	128	C	O5'-P-OP2	-5.79	100.49	105.70
22	BA	789	A	C2-N3-C4	-5.79	107.71	110.60
22	BA	1266	G	C8-N9-C4	5.79	108.71	106.40
22	BA	1022	G	C8-N9-C4	-5.78	104.09	106.40
22	BA	2038	G	C6-C5-N7	-5.78	126.93	130.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	1034	G	N3-C4-C5	-5.78	125.71	128.60
22	BA	1637	A	O5'-P-OP2	5.78	117.63	110.70
22	BA	1940	U	O5'-P-OP2	-5.78	100.50	105.70
22	BA	850	U	O5'-P-OP1	-5.78	100.50	105.70
22	BA	916	G	C6-C5-N7	-5.78	126.94	130.40
22	BA	954	G	O5'-P-OP2	5.78	117.63	110.70
22	BA	1332	G	N3-C4-N9	5.78	129.47	126.00
1	AA	1080	A	C4-C5-N7	-5.77	107.81	110.70
22	BA	2083	G	C5-C6-O6	-5.77	125.14	128.60
22	BA	446	G	C4-C5-N7	5.77	113.11	110.80
22	BA	984	A	C8-N9-C1'	5.77	138.09	127.70
22	BA	1693	U	OP1-P-OP2	5.77	128.25	119.60
22	BA	11	C	N1-C2-O2	-5.77	115.44	118.90
22	BA	240	C	N1-C2-O2	-5.76	115.44	118.90
22	BA	2830	C	N3-C4-C5	5.76	124.21	121.90
22	DA	777	G	C8-N9-C4	5.76	108.71	106.40
1	CA	812	G	O4'-C1'-N9	5.76	112.81	108.20
22	BA	1032	A	N9-C4-C5	5.76	108.10	105.80
22	BA	1635	A	C2-N3-C4	-5.76	107.72	110.60
22	BA	1426	G	C6-N1-C2	-5.76	121.65	125.10
22	BA	1900	A	C8-N9-C4	-5.76	103.50	105.80
22	BA	2330	G	C5-C6-O6	-5.75	125.15	128.60
22	BA	1611	C	C2-N3-C4	-5.75	117.02	119.90
22	BA	2422	C	N1-C2-O2	-5.75	115.45	118.90
22	BA	152	A	N1-C6-N6	5.75	122.05	118.60
22	BA	1352	U	C2-N3-C4	-5.74	123.56	127.00
22	BA	1352	U	C2-N1-C1'	-5.74	110.81	117.70
22	BA	1679	A	C5-C6-N6	-5.74	119.11	123.70
22	BA	479	A	O4'-C1'-N9	5.74	112.79	108.20
22	DA	2603	G	N1-C6-O6	5.74	123.34	119.90
22	BA	945	A	C5-C6-N6	-5.74	119.11	123.70
22	BA	2524	G	OP2-P-O3'	5.74	117.82	105.20
22	BA	2645	G	C4-C5-N7	5.73	113.09	110.80
22	BA	1759	A	C6-N1-C2	-5.73	115.16	118.60
23	BB	15	A	C5-N7-C8	-5.73	101.03	103.90
22	BA	1352	U	N1-C2-O2	-5.73	118.79	122.80
22	BA	1640	A	N9-C4-C5	5.73	108.09	105.80
22	BA	1779	U	C4-C5-C6	5.73	123.14	119.70
22	BA	2254	C	N1-C2-O2	-5.73	115.46	118.90
22	BA	2497	A	OP1-P-O3'	5.73	117.81	105.20
22	BA	33	C	N3-C4-N4	5.73	122.01	118.00
22	BA	1374	G	C5-N7-C8	-5.73	101.44	104.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	248	G	C8-N9-C4	5.72	108.69	106.40
22	BA	512	G	C4-C5-N7	5.72	113.09	110.80
22	BA	530	G	C5-C6-O6	-5.72	125.17	128.60
22	BA	1429	G	C5-C6-O6	5.72	132.03	128.60
22	BA	1444	G	O5'-P-OP2	-5.72	100.55	105.70
22	BA	2762	C	N3-C2-O2	5.72	125.91	121.90
22	BA	665	U	C5-C6-N1	-5.72	119.84	122.70
22	BA	1265	A	N1-C2-N3	5.72	132.16	129.30
22	BA	2036	C	N3-C2-O2	-5.71	117.90	121.90
22	BA	757	G	N1-C6-O6	5.71	123.33	119.90
22	BA	1695	G	OP1-P-OP2	5.71	128.17	119.60
22	BA	1758	U	N3-C4-O4	-5.71	115.40	119.40
22	BA	2678	C	C2-N3-C4	-5.71	117.04	119.90
23	BB	39	A	N1-C6-N6	5.71	122.03	118.60
23	BB	77	U	N3-C2-O2	5.71	126.20	122.20
22	DA	2443	C	C6-N1-C2	-5.71	118.02	120.30
22	BA	557	C	N1-C2-O2	-5.71	115.47	118.90
22	BA	1225	G	C6-C5-N7	-5.71	126.97	130.40
22	BA	2023	C	N3-C2-O2	5.71	125.89	121.90
22	BA	2049	G	C5-C6-N1	5.71	114.35	111.50
22	BA	1756	G	C4-C5-C6	5.71	122.22	118.80
22	BA	1784	A	C5-C6-N1	-5.71	114.85	117.70
22	BA	569	U	N1-C2-O2	-5.70	118.81	122.80
22	BA	630	G	N1-C6-O6	5.70	123.32	119.90
22	BA	1343	G	C4-N9-C1'	5.70	133.91	126.50
22	DA	250	G	N3-C4-C5	-5.70	125.75	128.60
22	BA	956	G	C5-C6-O6	-5.70	125.18	128.60
22	BA	2813	A	C5-C6-N6	-5.70	119.14	123.70
23	BB	26	C	N1-C2-O2	-5.70	115.48	118.90
22	BA	1208	C	C5-C4-N4	-5.70	116.21	120.20
22	BA	2283	C	N1-C2-O2	-5.70	115.48	118.90
22	BA	2360	G	C6-C5-N7	-5.70	126.98	130.40
22	BA	2515	C	N1-C2-O2	-5.70	115.48	118.90
22	BA	2519	U	C5-C4-O4	-5.70	122.48	125.90
22	BA	2525	G	O5'-P-OP2	-5.70	100.58	105.70
22	BA	2645	G	N1-C6-O6	5.70	123.32	119.90
22	BA	528	A	O4'-C1'-N9	-5.69	103.64	108.20
22	BA	1977	A	C2-N3-C4	-5.69	107.75	110.60
22	BA	81	G	O5'-P-OP1	-5.69	100.58	105.70
22	BA	2383	G	N1-C6-O6	-5.69	116.48	119.90
22	BA	672	C	C2-N3-C4	-5.69	117.06	119.90
22	BA	554	U	O5'-P-OP2	-5.68	100.58	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	1020	A	C5-N7-C8	-5.68	101.06	103.90
23	BB	106	G	C6-C5-N7	-5.68	126.99	130.40
22	BA	213	A	C5-C6-N6	-5.68	119.16	123.70
22	BA	1566	A	C5-C6-N6	-5.68	119.16	123.70
22	BA	1282	U	C5-C4-O4	-5.68	122.49	125.90
22	BA	33	C	C5-C4-N4	-5.67	116.23	120.20
22	BA	1142	A	C4-N9-C1'	-5.67	116.08	126.30
22	BA	1265	A	O5'-P-OP2	-5.67	100.60	105.70
22	DA	776	G	C8-N9-C1'	-5.67	119.63	127.00
22	BA	2545	G	C5-C6-N1	5.67	114.33	111.50
1	AA	857	C	O5'-P-OP2	-5.67	100.60	105.70
1	AA	1080	A	C5-C6-N6	5.67	128.23	123.70
22	BA	531	C	N3-C4-N4	-5.67	114.03	118.00
22	BA	726	G	C5-C6-O6	-5.67	125.20	128.60
22	BA	1223	G	O5'-P-OP2	5.67	117.50	110.70
22	BA	2781	A	OP1-P-OP2	5.67	128.10	119.60
22	BA	1320	C	N3-C4-C5	-5.66	119.64	121.90
22	BA	1785	A	N1-C6-N6	5.66	122.00	118.60
22	BA	2340	A	C8-N9-C4	-5.66	103.53	105.80
22	BA	1357	C	N3-C2-O2	5.66	125.86	121.90
22	BA	2712	C	C2-N3-C4	-5.66	117.07	119.90
23	BB	71	C	C6-N1-C2	5.66	122.56	120.30
1	AA	279	A	C5-C6-N6	-5.66	119.17	123.70
1	AA	732	C	OP1-P-O3'	5.66	117.65	105.20
22	BA	1251	C	C2-N1-C1'	5.66	125.03	118.80
22	BA	1756	G	C6-C5-N7	-5.66	127.00	130.40
22	BA	1999	C	OP2-P-O3'	5.66	117.65	105.20
22	BA	578	G	N3-C4-N9	5.66	129.39	126.00
22	DA	2501	C	C6-N1-C1'	5.66	127.59	120.80
22	BA	1020	A	C5-C6-N6	-5.66	119.18	123.70
22	BA	1378	A	P-O3'-C3'	5.65	126.48	119.70
1	AA	819	A	O5'-P-OP1	-5.65	100.61	105.70
22	BA	945	A	N9-C4-C5	-5.65	103.54	105.80
22	BA	2070	A	C5-C6-N1	5.65	120.53	117.70
22	BA	2465	C	O5'-P-OP2	-5.65	100.61	105.70
1	AA	1523	G	C5-C6-O6	5.65	131.99	128.60
22	BA	1984	G	N1-C6-O6	-5.64	116.51	119.90
22	BA	26	G	N9-C4-C5	-5.64	103.14	105.40
22	BA	1262	A	C6-N1-C2	-5.64	115.22	118.60
23	BB	91	C	N1-C2-O2	-5.64	115.52	118.90
22	BA	1231	U	N1-C2-O2	-5.64	118.86	122.80
22	BA	565	C	C2-N3-C4	-5.63	117.08	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	575	A	O4'-C1'-N9	5.63	112.71	108.20
23	BB	15	A	C2-N3-C4	-5.63	107.78	110.60
1	CA	547	A	O5'-P-OP1	-5.63	100.63	105.70
1	CA	551	U	OP2-P-O3'	5.63	117.60	105.20
22	BA	726	G	N9-C4-C5	-5.63	103.15	105.40
22	BA	2460	U	N3-C2-O2	-5.63	118.26	122.20
1	AA	820	U	O5'-P-OP1	-5.63	100.64	105.70
22	BA	914	G	C6-C5-N7	-5.63	127.02	130.40
22	BA	733	G	C5-N7-C8	-5.62	101.49	104.30
1	AA	264	C	N3-C4-N4	5.62	121.94	118.00
22	BA	952	G	OP2-P-O3'	5.62	117.57	105.20
22	BA	2451	A	OP1-P-OP2	-5.62	111.17	119.60
22	BA	855	G	N9-C4-C5	5.62	107.65	105.40
22	BA	1989	G	N9-C4-C5	-5.62	103.15	105.40
22	BA	1549	A	N9-C4-C5	-5.62	103.55	105.80
22	BA	2226	C	C2-N1-C1'	5.62	124.98	118.80
22	BA	2678	C	C6-N1-C2	5.62	122.55	120.30
22	BA	1288	G	N3-C2-N2	5.62	123.83	119.90
22	DA	2501	C	C2-N1-C1'	-5.62	112.62	118.80
22	BA	61	C	C6-N1-C2	5.62	122.55	120.30
22	BA	2829	A	C2-N3-C4	-5.62	107.79	110.60
22	BA	2455	G	N3-C4-N9	5.61	129.37	126.00
22	BA	1343	G	C8-N9-C4	-5.61	104.16	106.40
1	AA	279	A	C6-C5-N7	-5.61	128.37	132.30
22	BA	799	G	C5-C6-N1	5.61	114.30	111.50
22	DA	1789	A	C8-N9-C4	5.61	108.04	105.80
1	AA	779	C	C6-N1-C2	-5.61	118.06	120.30
22	BA	532	A	OP1-P-OP2	5.61	128.01	119.60
22	BA	1002	G	N3-C4-C5	-5.61	125.80	128.60
22	BA	1773	A	C8-N9-C4	5.60	108.04	105.80
22	BA	2672	U	C5-C4-O4	-5.60	122.54	125.90
23	BB	103	U	C5-C4-O4	-5.60	122.54	125.90
22	BA	2073	C	N3-C4-N4	5.60	121.92	118.00
22	BA	462	C	N1-C2-O2	-5.60	115.54	118.90
22	DA	1798	U	C2-N1-C1'	-5.60	110.98	117.70
1	CA	1079	G	C8-N9-C4	-5.60	104.16	106.40
23	BB	71	C	C2-N3-C4	-5.60	117.10	119.90
22	BA	723	C	C5-C6-N1	-5.59	118.20	121.00
22	BA	1959	G	N9-C4-C5	5.59	107.64	105.40
22	BA	2827	C	N3-C4-C5	5.59	124.14	121.90
23	BB	99	A	C2-N3-C4	-5.59	107.81	110.60
22	BA	378	C	N3-C4-C5	5.59	124.13	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	1831	G	OP2-P-O3'	5.59	117.49	105.20
22	BA	257	C	N3-C4-C5	5.58	124.13	121.90
22	BA	451	U	N1-C2-O2	-5.58	118.89	122.80
1	AA	323	U	N1-C2-O2	-5.58	118.89	122.80
22	BA	2449	U	C5-C6-N1	-5.58	119.91	122.70
22	DA	1276	A	N1-C6-N6	5.58	121.95	118.60
22	BA	717	C	N3-C2-O2	5.58	125.80	121.90
22	BA	1450	G	N3-C2-N2	5.58	123.80	119.90
22	BA	1567	G	OP1-P-O3'	5.58	117.47	105.20
22	BA	2689	U	N3-C2-O2	-5.57	118.30	122.20
22	BA	2712	C	C4-C5-C6	5.57	120.19	117.40
22	BA	2444	G	OP2-P-O3'	5.57	117.46	105.20
22	BA	200	U	N3-C2-O2	5.57	126.10	122.20
22	BA	520	G	N3-C2-N2	5.57	123.80	119.90
22	BA	837	C	N1-C2-O2	-5.57	115.56	118.90
22	BA	2588	G	N9-C4-C5	-5.57	103.17	105.40
22	BA	2038	G	C5-C6-O6	-5.57	125.26	128.60
22	BA	596	U	C5-C4-O4	-5.57	122.56	125.90
22	BA	1273	U	C2-N3-C4	-5.57	123.66	127.00
22	BA	918	A	C2-N3-C4	-5.56	107.82	110.60
22	BA	825	A	N1-C6-N6	-5.56	115.27	118.60
22	BA	2838	G	OP2-P-O3'	5.56	117.43	105.20
22	BA	1642	G	N3-C4-C5	5.56	131.38	128.60
22	BA	2578	G	OP1-P-O3'	5.55	117.42	105.20
23	BB	92	C	N1-C2-O2	-5.55	115.57	118.90
1	AA	888	G	O5'-P-OP2	-5.55	100.70	105.70
22	BA	1774	C	N1-C2-O2	-5.55	115.57	118.90
22	BA	2051	A	N1-C6-N6	5.55	121.93	118.60
22	BA	2450	A	OP2-P-O3'	5.55	117.41	105.20
22	BA	585	G	N3-C4-C5	-5.55	125.83	128.60
23	BB	79	G	C2-N3-C4	5.55	114.67	111.90
22	BA	708	G	C6-C5-N7	-5.55	127.07	130.40
22	BA	1707	G	O5'-P-OP2	-5.55	100.71	105.70
22	BA	213	A	C8-N9-C4	5.54	108.02	105.80
22	BA	1311	G	N7-C8-N9	5.54	115.87	113.10
22	BA	841	G	C2-N3-C4	-5.54	109.13	111.90
22	BA	858	G	O5'-P-OP2	-5.54	100.71	105.70
22	BA	995	C	C5'-C4'-O4'	-5.54	102.45	109.10
22	BA	2024	G	N3-C2-N2	-5.54	116.02	119.90
22	BA	2442	C	C6-N1-C1'	-5.54	114.15	120.80
1	CA	27	G	N1-C6-O6	5.54	123.22	119.90
22	BA	2056	G	OP1-P-O3'	5.54	117.39	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	986	C	C6-N1-C2	-5.54	118.08	120.30
22	BA	1301	A	N1-C6-N6	5.54	121.92	118.60
22	BA	811	U	N3-C4-O4	-5.54	115.53	119.40
22	BA	2067	G	C5-C6-N1	5.54	114.27	111.50
22	BA	2383	G	C4-C5-N7	5.53	113.01	110.80
23	BB	93	C	C2-N3-C4	-5.53	117.14	119.90
22	BA	2800	A	O5'-P-OP2	-5.53	100.72	105.70
22	BA	565	C	N3-C4-N4	5.53	121.87	118.00
22	DA	2620	C	C6-N1-C2	5.53	122.51	120.30
22	BA	2375	G	N1-C6-O6	-5.53	116.58	119.90
22	BA	2350	C	C6-N1-C2	5.52	122.51	120.30
22	BA	535	G	N3-C2-N2	5.52	123.77	119.90
22	BA	771	G	OP1-P-O3'	5.52	117.35	105.20
22	BA	2407	A	C4-C5-C6	5.52	119.76	117.00
22	BA	190	A	C4-C5-N7	5.52	113.46	110.70
22	BA	1638	C	N1-C2-O2	-5.52	115.59	118.90
22	BA	1957	C	C6-N1-C2	5.52	122.51	120.30
1	AA	1510	C	C5-C4-N4	-5.52	116.34	120.20
22	BA	1831	G	C8-N9-C4	-5.52	104.19	106.40
22	BA	1940	U	N1-C2-O2	5.52	126.66	122.80
22	BA	697	G	C5-C6-O6	5.52	131.91	128.60
22	BA	1032	A	N1-C2-N3	5.52	132.06	129.30
22	BA	1792	G	OP2-P-O3'	5.52	117.33	105.20
22	BA	686	U	O5'-P-OP2	-5.51	100.74	105.70
22	BA	1829	A	N1-C6-N6	-5.51	115.29	118.60
22	DA	691	C	C6-N1-C2	-5.51	118.10	120.30
1	AA	264	C	C5-C4-N4	-5.51	116.34	120.20
22	BA	27	G	C5-C6-N1	5.51	114.25	111.50
22	BA	2000	C	C6-N1-C2	5.51	122.50	120.30
22	BA	2008	C	C4-C5-C6	5.51	120.15	117.40
22	BA	1250	G	C8-N9-C4	-5.51	104.20	106.40
22	BA	2503	A	N3-C4-N9	5.51	131.81	127.40
22	BA	2628	C	C6-N1-C2	5.51	122.50	120.30
22	BA	1020	A	N9-C4-C5	-5.50	103.60	105.80
1	CA	535	A	C8-N9-C4	5.50	108.00	105.80
1	AA	1504	G	O4'-C1'-N9	5.50	112.60	108.20
22	BA	680	C	N3-C2-O2	5.50	125.75	121.90
22	BA	636	G	N3-C2-N2	-5.50	116.05	119.90
22	BA	762	U	O5'-P-OP2	-5.50	100.75	105.70
22	BA	2606	C	C2-N1-C1'	-5.50	112.75	118.80
22	DA	2045	C	C5-C4-N4	-5.50	116.35	120.20
22	BA	474	G	C5-C6-O6	5.50	131.90	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	1225	G	C5-C6-O6	-5.50	125.30	128.60
22	BA	2025	C	O5'-P-OP2	-5.50	100.75	105.70
23	BB	71	C	N1-C2-O2	-5.50	115.60	118.90
22	BA	1666	G	O5'-P-OP2	-5.50	100.75	105.70
1	AA	365	U	C6-N1-C1'	5.50	128.89	121.20
22	BA	1384	A	O5'-P-OP1	-5.49	100.76	105.70
22	BA	519	U	N3-C2-O2	-5.49	118.36	122.20
22	BA	1035	U	C5-C6-N1	-5.49	119.95	122.70
22	BA	2815	C	C2-N3-C4	-5.49	117.16	119.90
22	DA	1819	A	N9-C4-C5	5.49	108.00	105.80
22	BA	732	C	N1-C2-O2	-5.49	115.61	118.90
22	BA	1514	G	N1-C6-O6	5.49	123.19	119.90
22	BA	1833	C	N1-C2-O2	-5.49	115.61	118.90
22	BA	670	A	O4'-C1'-N9	-5.49	103.81	108.20
22	BA	1687	G	N3-C4-C5	-5.49	125.86	128.60
22	BA	2502	G	O5'-P-OP2	-5.49	100.76	105.70
22	BA	2581	G	C4-C5-N7	5.49	112.99	110.80
1	CA	1395	C	N3-C4-N4	5.49	121.84	118.00
22	DA	946	C	C6-N1-C2	-5.49	118.11	120.30
22	BA	1239	G	OP2-P-O3'	5.48	117.27	105.20
1	AA	1094	G	O4'-C1'-N9	5.48	112.58	108.20
22	BA	867	C	O5'-P-OP2	5.48	117.28	110.70
22	BA	2030	A	O5'-P-OP1	-5.48	100.77	105.70
22	BA	2529	G	C6-C5-N7	-5.48	127.11	130.40
22	BA	1637	A	C4-C5-N7	5.48	113.44	110.70
22	BA	2425	A	OP2-P-O3'	5.48	117.25	105.20
22	BA	1193	G	N1-C6-O6	-5.47	116.62	119.90
22	BA	1276	A	O5'-P-OP2	-5.47	100.77	105.70
22	BA	692	C	OP2-P-O3'	5.47	117.24	105.20
22	BA	2052	A	C5-N7-C8	5.47	106.64	103.90
22	BA	2232	C	N3-C4-C5	5.47	124.09	121.90
1	CA	1517	G	O5'-P-OP2	-5.47	100.78	105.70
22	BA	1013	C	O5'-P-OP2	-5.47	100.78	105.70
22	BA	1779	U	OP1-P-O3'	5.47	117.23	105.20
22	BA	1563	U	C6-N1-C2	-5.47	117.72	121.00
22	BA	1969	A	OP1-P-O3'	5.47	117.23	105.20
22	BA	2785	C	OP1-P-OP2	5.47	127.80	119.60
1	AA	766	A	C8-N9-C4	5.46	107.99	105.80
22	BA	1428	C	N3-C2-O2	5.46	125.72	121.90
1	AA	793	U	OP2-P-O3'	5.46	117.21	105.20
22	BA	1182	G	OP1-P-O3'	5.46	117.21	105.20
22	BA	1142	A	C4-C5-N7	5.46	113.43	110.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	224	U	OP2-P-O3'	5.45	117.20	105.20
22	BA	740	C	O5'-P-OP2	5.45	117.25	110.70
22	BA	1917	U	C2-N1-C1'	5.45	124.24	117.70
22	BA	2250	G	N9-C4-C5	-5.45	103.22	105.40
22	BA	675	A	C8-N9-C4	5.45	107.98	105.80
22	BA	1976	U	C5-C4-O4	-5.45	122.63	125.90
22	BA	2825	G	C8-N9-C1'	-5.45	119.92	127.00
22	BA	2892	G	C2-N3-C4	-5.45	109.18	111.90
22	BA	1022	G	C5-C6-O6	-5.45	125.33	128.60
22	BA	35	G	N1-C6-O6	-5.44	116.63	119.90
22	BA	1934	C	C5-C4-N4	5.44	124.01	120.20
22	BA	2447	G	C8-N9-C4	5.44	108.58	106.40
22	BA	2768	U	N1-C2-O2	-5.44	118.99	122.80
1	AA	1511	G	C5-C6-O6	-5.44	125.34	128.60
22	BA	1295	C	OP2-P-O3'	5.44	117.17	105.20
1	AA	1487	G	OP2-P-O3'	5.44	117.16	105.20
22	BA	1227	G	N1-C6-O6	5.44	123.16	119.90
22	BA	400	G	C5-C6-O6	-5.43	125.34	128.60
22	BA	841	G	C6-C5-N7	-5.43	127.14	130.40
22	BA	318	C	N1-C2-O2	-5.43	115.64	118.90
22	BA	1283	G	N1-C6-O6	-5.43	116.64	119.90
22	BA	1349	C	O5'-P-OP2	-5.43	100.82	105.70
22	BA	2337	G	C8-N9-C4	-5.43	104.23	106.40
1	AA	254	G	O5'-P-OP1	-5.42	100.82	105.70
22	BA	2035	G	N1-C6-O6	-5.42	116.65	119.90
22	BA	1756	G	N3-C4-C5	-5.42	125.89	128.60
22	BA	2863	C	N1-C2-O2	-5.42	115.65	118.90
23	BB	17	C	N1-C2-O2	-5.42	115.65	118.90
22	BA	695	G	N1-C6-O6	-5.42	116.65	119.90
22	BA	32	C	N1-C2-O2	-5.42	115.65	118.90
22	BA	2569	G	C5-C6-O6	5.42	131.85	128.60
22	BA	674	G	C2-N3-C4	5.42	114.61	111.90
22	BA	2680	U	OP1-P-OP2	-5.42	111.47	119.60
22	BA	483	A	C8-N9-C4	5.42	107.97	105.80
22	BA	1829	A	N7-C8-N9	-5.42	111.09	113.80
1	CA	287	U	N3-C2-O2	5.42	125.99	122.20
1	CA	1522	U	O5'-P-OP2	-5.41	100.83	105.70
1	AA	1530	G	C8-N9-C1'	5.41	134.03	127.00
22	BA	851	C	N1-C2-O2	-5.41	115.65	118.90
22	BA	559	G	OP2-P-O3'	5.41	117.10	105.20
22	BA	645	C	N1-C2-O2	-5.41	115.66	118.90
22	BA	955	U	N3-C4-C5	5.41	117.84	114.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	2788	C	C6-N1-C2	5.41	122.46	120.30
22	BA	1230	A	O5'-P-OP1	5.41	117.19	110.70
22	BA	1387	A	C6-C5-N7	-5.41	128.52	132.30
22	BA	1562	U	N1-C2-O2	-5.41	119.02	122.80
1	AA	780	A	N9-C4-C5	-5.40	103.64	105.80
22	BA	1796	U	N1-C2-O2	-5.40	119.02	122.80
1	CA	1395	C	C5-C4-N4	-5.40	116.42	120.20
22	BA	451	U	N3-C2-O2	5.40	125.98	122.20
22	BA	565	C	C5-C4-N4	-5.40	116.42	120.20
22	BA	1607	C	C6-N1-C2	-5.40	118.14	120.30
22	BA	512	G	C4-C5-C6	-5.40	115.56	118.80
22	BA	2581	G	N9-C4-C5	-5.40	103.24	105.40
22	BA	184	C	N1-C2-O2	-5.40	115.66	118.90
22	BA	975	A	OP1-P-OP2	5.39	127.69	119.60
22	BA	1251	C	N3-C4-C5	5.39	124.06	121.90
22	BA	1651	G	N3-C4-C5	-5.39	125.90	128.60
22	BA	1694	C	N3-C2-O2	5.39	125.68	121.90
1	AA	1509	C	N3-C2-O2	5.39	125.67	121.90
22	BA	1225	G	N1-C6-O6	5.39	123.14	119.90
22	BA	1516	G	N1-C6-O6	5.39	123.14	119.90
22	BA	1930	G	O5'-P-OP2	-5.39	100.85	105.70
23	BB	81	G	N1-C2-N3	5.39	127.14	123.90
22	BA	753	A	OP1-P-OP2	5.39	127.69	119.60
22	BA	2395	C	C6-N1-C2	5.39	122.46	120.30
22	BA	1264	A	C5-C6-N1	5.39	120.39	117.70
22	BA	1370	C	N1-C2-O2	-5.39	115.67	118.90
22	BA	2042	A	C6-N1-C2	-5.39	115.37	118.60
1	AA	500	G	C4-N9-C1'	-5.39	119.50	126.50
22	BA	957	C	C2-N3-C4	5.39	122.59	119.90
22	BA	2457	U	C5-C6-N1	5.39	125.39	122.70
1	AA	1479	C	N1-C2-O2	-5.39	115.67	118.90
22	BA	486	C	N1-C2-O2	-5.39	115.67	118.90
22	BA	760	G	N1-C6-O6	5.39	123.13	119.90
22	BA	855	G	C5-C6-O6	5.39	131.83	128.60
22	BA	1025	G	C5-C6-O6	5.39	131.83	128.60
22	BA	2852	G	N3-C4-C5	-5.39	125.91	128.60
22	BA	981	A	N1-C6-N6	5.38	121.83	118.60
22	BA	2053	G	N9-C4-C5	-5.38	103.25	105.40
22	BA	2837	A	N1-C6-N6	5.38	121.83	118.60
22	BA	2633	G	N1-C2-N2	-5.38	111.36	116.20
22	BA	560	C	N1-C2-O2	5.38	122.13	118.90
22	BA	1034	G	C6-N1-C2	-5.38	121.87	125.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	795	C	N3-C4-C5	5.38	124.05	121.90
22	BA	1398	C	N1-C2-O2	-5.38	115.67	118.90
22	BA	1776	G	O4'-C1'-N9	-5.38	103.90	108.20
22	BA	1426	G	N3-C4-N9	5.38	129.23	126.00
22	DA	1779	U	O4'-C1'-N1	5.38	112.50	108.20
22	BA	513	A	N9-C4-C5	-5.38	103.65	105.80
22	BA	789	A	N9-C4-C5	5.38	107.95	105.80
22	BA	516	C	C2-N3-C4	-5.37	117.21	119.90
22	BA	1256	G	C8-N9-C1'	-5.37	120.02	127.00
22	BA	740	C	OP2-P-O3'	5.37	117.02	105.20
22	BA	2676	C	C2-N3-C4	-5.37	117.21	119.90
22	BA	2702	G	OP2-P-O3'	5.37	117.02	105.20
1	AA	1467	C	N1-C2-O2	-5.37	115.68	118.90
22	BA	1270	C	N1-C2-O2	-5.37	115.68	118.90
22	BA	271	G	P-O3'-C3'	5.37	126.14	119.70
22	BA	905	A	N1-C6-N6	-5.37	115.38	118.60
22	BA	968	C	O5'-P-OP1	5.37	117.14	110.70
22	BA	2422	C	N3-C2-O2	5.37	125.66	121.90
22	BA	789	A	N3-C4-N9	-5.37	123.11	127.40
22	BA	2807	U	OP2-P-O3'	5.37	117.00	105.20
22	BA	1123	C	N3-C4-C5	5.36	124.05	121.90
22	BA	1936	A	N9-C1'-C2'	-5.36	106.10	112.00
22	BA	908	C	C6-N1-C2	-5.36	118.16	120.30
22	BA	914	G	C4-C5-N7	5.36	112.94	110.80
22	BA	2030	A	N1-C2-N3	5.36	131.98	129.30
22	BA	845	A	C4-C5-C6	5.36	119.68	117.00
22	BA	816	C	C6-N1-C2	-5.36	118.16	120.30
22	BA	2819	G	O5'-P-OP2	5.36	117.13	110.70
22	BA	1006	C	N1-C2-O2	-5.36	115.69	118.90
22	BA	801	G	C8-N9-C4	-5.35	104.26	106.40
22	BA	835	C	N3-C4-C5	5.35	124.04	121.90
22	BA	2060	A	N9-C4-C5	5.35	107.94	105.80
22	BA	98	G	OP1-P-OP2	5.35	127.62	119.60
22	BA	1658	C	C5-C4-N4	-5.35	116.46	120.20
22	BA	2318	G	N3-C4-N9	5.35	129.21	126.00
22	BA	180	G	C8-N9-C1'	5.34	133.95	127.00
22	BA	1249	U	N1-C2-O2	-5.34	119.06	122.80
22	BA	1299	G	C4-C5-C6	5.34	122.01	118.80
22	BA	1677	A	C6-C5-N7	-5.34	128.56	132.30
22	BA	2639	A	N1-C6-N6	5.34	121.81	118.60
22	BA	993	G	C4-C5-N7	-5.34	108.66	110.80
22	BA	996	A	C4-C5-C6	5.34	119.67	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	2545	G	C2-N3-C4	5.34	114.57	111.90
1	AA	780	A	N1-C6-N6	5.34	121.80	118.60
1	AA	811	C	OP2-P-O3'	5.34	116.95	105.20
22	BA	1637	A	C5-N7-C8	-5.34	101.23	103.90
22	BA	979	A	C5-N7-C8	-5.34	101.23	103.90
22	BA	1757	A	N1-C2-N3	5.34	131.97	129.30
22	BA	2324	U	O4'-C1'-N1	-5.34	103.93	108.20
23	BB	8	C	N1-C2-O2	-5.34	115.70	118.90
1	AA	1279	G	N7-C8-N9	5.34	115.77	113.10
22	BA	2677	G	C8-N9-C4	5.34	108.53	106.40
22	BA	677	A	O5'-P-OP2	-5.33	100.90	105.70
22	BA	980	A	C4-C5-N7	5.33	113.37	110.70
22	BA	1799	G	OP2-P-O3'	5.33	116.92	105.20
22	BA	2889	C	C5-C6-N1	-5.33	118.33	121.00
23	BB	99	A	C8-N9-C4	5.33	107.93	105.80
22	BA	840	C	C5-C4-N4	-5.33	116.47	120.20
1	AA	1486	G	O5'-P-OP2	-5.33	100.91	105.70
22	BA	1027	A	N1-C6-N6	5.33	121.80	118.60
22	BA	1687	G	C8-N9-C4	-5.33	104.27	106.40
22	BA	1032	A	C8-N9-C4	-5.32	103.67	105.80
22	BA	1223	G	OP2-P-O3'	5.32	116.91	105.20
22	BA	2320	U	N3-C2-O2	5.32	125.92	122.20
22	DA	1993	U	N3-C2-O2	-5.32	118.48	122.20
22	BA	745	G	C5-C6-N1	5.32	114.16	111.50
22	BA	1632	A	C5-C6-N6	-5.32	119.44	123.70
22	BA	2587	A	C8-N9-C4	5.32	107.93	105.80
22	BA	1963	U	C2-N1-C1'	5.32	124.08	117.70
22	BA	2501	C	N3-C4-C5	5.32	124.03	121.90
22	BA	2729	G	N3-C4-N9	5.32	129.19	126.00
22	BA	1645	G	C4-C5-N7	5.31	112.92	110.80
22	BA	1695	G	N3-C2-N2	5.31	123.62	119.90
22	BA	2040	G	N3-C4-N9	5.31	129.19	126.00
25	BD	165	MET	CG-SD-CE	-5.31	91.71	100.20
22	BA	1145	C	N1-C2-O2	-5.31	115.72	118.90
22	BA	2001	C	N3-C2-O2	5.31	125.61	121.90
22	BA	2518	A	O4'-C1'-N9	-5.31	103.95	108.20
22	BA	2705	A	O5'-P-OP2	5.31	117.07	110.70
22	BA	8	C	N1-C2-O2	-5.30	115.72	118.90
22	BA	1651	G	N1-C6-O6	-5.30	116.72	119.90
22	BA	1819	A	N1-C6-N6	5.30	121.78	118.60
22	BA	1303	G	C6-C5-N7	5.30	133.58	130.40
22	DA	1928	A	N1-C6-N6	5.30	121.78	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	967	U	C5-C4-O4	5.30	129.08	125.90
22	BA	2001	C	N3-C4-C5	5.30	124.02	121.90
22	BA	1330	C	N3-C4-C5	-5.30	119.78	121.90
22	BA	2051	A	OP1-P-O3'	5.30	116.86	105.20
22	BA	2075	U	C2-N3-C4	-5.30	123.82	127.00
22	DA	776	G	N3-C4-C5	-5.30	125.95	128.60
1	AA	729	A	N1-C6-N6	5.30	121.78	118.60
22	BA	1981	A	N1-C2-N3	-5.30	126.65	129.30
22	BA	2741	A	N1-C6-N6	5.30	121.78	118.60
1	AA	834	U	O5'-P-OP2	-5.29	100.94	105.70
22	BA	446	G	N3-C4-C5	5.29	131.25	128.60
22	BA	561	G	C5-N7-C8	-5.29	101.65	104.30
22	BA	69	C	C5-C6-N1	-5.29	118.35	121.00
22	BA	2051	A	C4-C5-N7	5.29	113.34	110.70
22	BA	2846	G	C2-N3-C4	-5.29	109.25	111.90
22	BA	1208	C	N3-C4-N4	5.29	121.70	118.00
22	BA	2777	G	N3-C2-N2	5.29	123.60	119.90
22	BA	2076	U	N1-C2-N3	5.29	118.07	114.90
22	BA	726	G	N1-C6-O6	5.29	123.07	119.90
22	BA	2042	A	O5'-P-OP1	5.29	117.04	110.70
22	BA	2069	G	C4-C5-C6	-5.29	115.63	118.80
22	BA	974	G	C2-N3-C4	-5.28	109.26	111.90
22	BA	1262	A	OP1-P-O3'	5.28	116.82	105.20
22	BA	1609	A	C5-C6-N6	-5.28	119.47	123.70
1	AA	1523	G	N3-C4-N9	-5.28	122.83	126.00
22	BA	2777	G	N1-C2-N3	-5.28	120.73	123.90
22	BA	749	A	OP1-P-OP2	5.28	127.52	119.60
22	BA	19	A	N1-C6-N6	5.28	121.77	118.60
22	BA	1994	C	C2-N3-C4	-5.28	117.26	119.90
22	BA	523	C	N3-C4-C5	5.28	124.01	121.90
22	BA	998	C	N3-C2-O2	-5.28	118.21	121.90
22	BA	2073	C	N1-C2-O2	-5.28	115.73	118.90
22	BA	1584	U	C2-N1-C1'	5.27	124.03	117.70
22	BA	2517	C	O4'-C1'-N1	5.27	112.42	108.20
22	BA	980	A	N7-C8-N9	5.27	116.44	113.80
22	BA	747	U	N3-C2-O2	-5.27	118.51	122.20
22	BA	27	G	N1-C6-O6	-5.27	116.74	119.90
22	BA	1469	A	C5-C6-N6	-5.27	119.49	123.70
1	CA	428	G	N1-C6-O6	5.27	123.06	119.90
22	BA	1260	A	C8-N9-C4	5.26	107.91	105.80
22	BA	1703	G	O5'-P-OP2	-5.26	100.96	105.70
38	BQ	53	ARG	NE-CZ-NH1	5.26	122.93	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	723	C	C2-N1-C1'	-5.26	113.01	118.80
22	BA	922	C	C6-N1-C2	5.26	122.40	120.30
22	BA	2752	C	C5-C4-N4	-5.26	116.52	120.20
22	BA	2824	C	N3-C2-O2	5.26	125.58	121.90
22	DA	1824	G	O5'-P-OP2	5.26	117.02	110.70
22	BA	572	A	N1-C2-N3	5.26	131.93	129.30
22	BA	1606	C	P-O3'-C3'	5.26	126.01	119.70
22	BA	2642	G	N1-C6-O6	-5.26	116.74	119.90
22	BA	767	U	OP2-P-O3'	5.26	116.76	105.20
23	BB	90	C	C2-N3-C4	-5.26	117.27	119.90
1	AA	124	C	C6-N1-C2	5.25	122.40	120.30
22	BA	534	U	C5-C4-O4	5.25	129.05	125.90
1	AA	811	C	N3-C2-O2	5.25	125.58	121.90
1	AA	1483	A	C8-N9-C4	5.25	107.90	105.80
22	BA	455	C	N1-C2-O2	5.25	122.05	118.90
1	CA	770	C	C6-N1-C2	5.25	122.40	120.30
22	BA	1303	G	N9-C4-C5	5.25	107.50	105.40
22	BA	1565	C	O5'-P-OP2	-5.25	100.98	105.70
22	BA	1930	G	N1-C6-O6	-5.25	116.75	119.90
23	BB	106	G	N1-C6-O6	5.25	123.05	119.90
22	BA	948	C	C6-N1-C2	-5.25	118.20	120.30
22	DA	1900	A	O4'-C1'-N9	-5.25	104.00	108.20
22	BA	530	G	N1-C6-O6	5.24	123.05	119.90
22	BA	680	C	N1-C2-O2	-5.24	115.75	118.90
1	AA	1079	G	C8-N9-C4	-5.24	104.30	106.40
22	BA	298	G	N1-C6-O6	5.24	123.05	119.90
22	BA	2424	C	C5-C6-N1	-5.24	118.38	121.00
22	BA	2516	A	OP2-P-O3'	5.24	116.72	105.20
22	BA	1326	U	N3-C4-O4	5.24	123.06	119.40
22	BA	2718	G	N1-C6-O6	5.24	123.04	119.90
22	DA	2240	U	O5'-P-OP2	-5.24	100.99	105.70
22	BA	1798	U	C6-N1-C2	5.23	124.14	121.00
22	BA	2069	G	N3-C4-C5	5.23	131.22	128.60
22	BA	336	C	N1-C2-O2	-5.23	115.76	118.90
22	BA	678	C	N3-C4-C5	5.23	123.99	121.90
22	BA	1972	G	C8-N9-C4	-5.23	104.31	106.40
22	BA	2799	A	C5-N7-C8	-5.23	101.29	103.90
22	BA	218	A	N1-C6-N6	5.23	121.73	118.60
1	CA	467	U	C2-N1-C1'	5.23	123.97	117.70
22	BA	66	C	C6-N1-C2	-5.22	118.21	120.30
22	BA	599	A	N1-C6-N6	-5.22	115.47	118.60
22	BA	523	C	C6-N1-C2	5.22	122.39	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	974	G	OP2-P-O3'	5.22	116.68	105.20
22	BA	2382	G	C8-N9-C1'	-5.22	120.22	127.00
22	BA	2825	G	N7-C8-N9	5.22	115.71	113.10
22	DA	1666	G	C4-C5-N7	-5.22	108.71	110.80
1	AA	229	U	N3-C2-O2	5.22	125.85	122.20
1	AA	1279	G	C6-C5-N7	-5.22	127.27	130.40
22	BA	2788	C	C5-C6-N1	-5.22	118.39	121.00
22	BA	2373	G	N3-C4-N9	-5.21	122.87	126.00
22	BA	1674	G	O4'-C1'-N9	-5.21	104.03	108.20
1	CA	1530	G	O4'-C1'-N9	5.21	112.37	108.20
22	BA	536	G	C5-C6-N1	5.21	114.10	111.50
22	BA	665	U	N3-C2-O2	5.21	125.85	122.20
22	BA	952	G	N1-C6-O6	5.21	123.03	119.90
22	BA	1027	A	C6-C5-N7	-5.21	128.65	132.30
22	BA	2815	C	C5-C6-N1	-5.21	118.39	121.00
22	BA	2866	U	N1-C2-O2	5.21	126.45	122.80
22	BA	725	G	O5'-P-OP1	-5.21	101.02	105.70
22	BA	1283	G	C8-N9-C4	-5.21	104.32	106.40
22	BA	2393	U	C5-C4-O4	-5.21	122.78	125.90
22	BA	248	G	OP1-P-OP2	5.20	127.41	119.60
22	BA	749	A	O5'-P-OP2	-5.20	101.02	105.70
22	BA	1002	G	OP1-P-O3'	5.20	116.65	105.20
22	BA	1649	G	OP1-P-OP2	5.20	127.41	119.60
22	BA	1999	C	C2-N1-C1'	-5.20	113.08	118.80
23	BB	99	A	N3-C4-C5	5.20	130.44	126.80
22	BA	1977	A	N1-C6-N6	-5.20	115.48	118.60
1	CA	1521	C	OP2-P-O3'	5.20	116.64	105.20
22	BA	818	G	OP1-P-OP2	-5.20	111.80	119.60
22	BA	2643	G	C8-N9-C4	5.20	108.48	106.40
22	BA	2837	A	C4-C5-N7	5.20	113.30	110.70
22	DA	1958	C	N1-C2-O2	-5.20	115.78	118.90
22	BA	841	G	C5-C6-N1	-5.20	108.90	111.50
22	BA	580	U	N1-C2-N3	5.20	118.02	114.90
22	BA	2318	G	N3-C4-C5	-5.20	126.00	128.60
22	BA	1273	U	N3-C4-C5	5.19	117.72	114.60
22	BA	1273	U	C2-N1-C1'	-5.19	111.47	117.70
22	BA	1319	C	C5-C4-N4	-5.19	116.57	120.20
22	BA	2324	U	N3-C4-O4	5.19	123.03	119.40
22	BA	1009	A	C5-C6-N6	-5.19	119.55	123.70
22	BA	691	C	OP1-P-OP2	-5.18	111.82	119.60
22	BA	1682	G	N1-C2-N2	-5.18	111.53	116.20
22	BA	1917	U	C5-C6-N1	5.18	125.29	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	2644	G	N1-C6-O6	5.18	123.01	119.90
22	BA	48	G	N1-C6-O6	5.18	123.01	119.90
22	BA	2252	G	C5-C6-O6	-5.18	125.49	128.60
22	BA	2615	U	OP1-P-O3'	5.18	116.60	105.20
22	BA	772	C	O5'-P-OP1	-5.18	101.04	105.70
22	BA	2455	G	OP2-P-O3'	5.18	116.60	105.20
22	BA	2824	C	N3-C4-N4	5.18	121.62	118.00
22	BA	1040	A	C5-C6-N6	-5.18	119.56	123.70
22	BA	1313	U	N3-C4-O4	5.18	123.02	119.40
22	BA	1332	G	C6-N1-C2	-5.18	122.00	125.10
22	BA	1779	U	C2-N1-C1'	-5.18	111.49	117.70
22	BA	1926	U	C5-C4-O4	-5.18	122.79	125.90
22	DA	1993	U	N1-C2-O2	5.18	126.42	122.80
22	BA	2383	G	N7-C8-N9	5.17	115.69	113.10
22	BA	2484	G	C8-N9-C4	5.17	108.47	106.40
22	BA	2021	C	OP1-P-O3'	5.17	116.58	105.20
22	BA	2427	C	C2-N3-C4	-5.17	117.31	119.90
1	AA	814	A	OP2-P-O3'	5.17	116.57	105.20
1	AA	888	G	O5'-P-OP1	5.17	116.90	110.70
22	BA	771	G	C5-C6-O6	-5.17	125.50	128.60
22	BA	1265	A	C6-N1-C2	-5.17	115.50	118.60
22	BA	2624	G	C8-N9-C4	5.17	108.47	106.40
22	BA	2829	A	N1-C2-N3	5.17	131.88	129.30
1	AA	899	C	N3-C2-O2	5.17	125.52	121.90
1	CA	767	A	N1-C6-N6	5.17	121.70	118.60
22	BA	675	A	N9-C4-C5	-5.16	103.73	105.80
22	BA	2575	C	N3-C2-O2	-5.16	118.28	121.90
22	BA	2070	A	C8-N9-C4	5.16	107.86	105.80
22	BA	2509	G	C5-C6-N1	5.16	114.08	111.50
22	BA	555	G	N3-C2-N2	-5.16	116.29	119.90
22	BA	512	G	N3-C4-N9	-5.16	122.91	126.00
22	BA	938	G	C4-N9-C1'	-5.16	119.80	126.50
22	BA	1618	A	OP2-P-O3'	5.16	116.54	105.20
22	BA	2557	G	N3-C4-C5	-5.16	126.02	128.60
22	DA	847	U	C2-N1-C1'	5.15	123.89	117.70
22	BA	1936	A	N3-C4-N9	-5.15	123.28	127.40
22	BA	2010	G	OP1-P-OP2	-5.15	111.87	119.60
22	BA	1263	U	O5'-P-OP2	-5.15	101.07	105.70
22	BA	1129	A	OP1-P-O3'	5.15	116.53	105.20
22	BA	1326	U	N3-C2-O2	5.15	125.80	122.20
22	BA	1788	C	C2-N3-C4	-5.15	117.33	119.90
22	BA	2046	G	C5-C6-O6	5.15	131.69	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	560	C	OP1-P-OP2	-5.15	111.88	119.60
22	BA	2563	U	O5'-P-OP2	-5.15	101.07	105.70
23	BB	71	C	C5-C6-N1	-5.15	118.43	121.00
22	BA	799	G	N1-C6-O6	-5.14	116.81	119.90
22	BA	943	A	C2-N3-C4	-5.14	108.03	110.60
22	BA	1309	G	C5-N7-C8	5.14	106.87	104.30
22	BA	1570	A	C4-C5-C6	5.14	119.57	117.00
1	AA	50	A	N1-C6-N6	5.14	121.68	118.60
22	BA	573	U	OP1-P-OP2	-5.14	111.89	119.60
22	BA	941	A	N1-C6-N6	5.14	121.68	118.60
22	BA	1977	A	N1-C2-N3	5.14	131.87	129.30
22	BA	2240	U	C5-C4-O4	-5.14	122.82	125.90
1	AA	1500	A	C8-N9-C4	5.14	107.86	105.80
22	BA	759	G	C4-C5-N7	5.14	112.86	110.80
22	BA	759	G	N9-C4-C5	-5.14	103.34	105.40
22	BA	2001	C	O5'-P-OP2	-5.14	101.08	105.70
1	CA	917	G	O5'-P-OP2	5.14	116.86	110.70
22	BA	128	C	N3-C2-O2	5.13	125.49	121.90
22	BA	1450	G	N1-C2-N2	-5.13	111.58	116.20
1	CA	893	C	N3-C4-C5	5.13	123.95	121.90
22	BA	1340	U	C2-N1-C1'	-5.13	111.54	117.70
22	BA	143	C	N3-C4-N4	5.13	121.59	118.00
22	BA	1299	G	N3-C4-N9	5.13	129.08	126.00
22	BA	2814	A	OP1-P-OP2	5.13	127.30	119.60
1	CA	887	G	N1-C6-O6	5.13	122.98	119.90
22	BA	726	G	C2-N3-C4	-5.13	109.33	111.90
22	BA	1438	U	N1-C2-O2	-5.13	119.21	122.80
22	BA	2707	U	C5-C6-N1	-5.13	120.14	122.70
22	BA	1258	U	C5-C6-N1	5.13	125.26	122.70
22	BA	2252	G	C6-C5-N7	-5.13	127.32	130.40
22	BA	1249	U	N3-C2-O2	5.13	125.79	122.20
22	BA	1930	G	C6-C5-N7	5.13	133.48	130.40
22	BA	2097	A	N1-C6-N6	-5.13	115.52	118.60
22	BA	2281	A	O5'-P-OP1	-5.13	101.09	105.70
22	BA	1941	C	N1-C2-O2	-5.12	115.83	118.90
22	BA	1138	G	N7-C8-N9	5.12	115.66	113.10
22	BA	2498	C	C5-C4-N4	5.12	123.78	120.20
22	BA	203	A	C4-C5-N7	5.12	113.26	110.70
22	BA	1648	U	C2-N1-C1'	-5.12	111.56	117.70
22	BA	1707	G	N3-C2-N2	5.12	123.48	119.90
22	BA	2075	U	N1-C2-N3	5.12	117.97	114.90
22	BA	2360	G	N1-C6-O6	5.12	122.97	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	DA	757	G	N3-C4-C5	5.12	131.16	128.60
22	BA	19	A	N3-C4-C5	-5.12	123.22	126.80
22	BA	41	C	N1-C2-O2	-5.12	115.83	118.90
22	BA	2362	C	N3-C4-C5	5.12	123.95	121.90
22	BA	2626	C	C5-C6-N1	-5.12	118.44	121.00
22	BA	1618	A	C6-N1-C2	5.11	121.67	118.60
22	BA	1780	A	N1-C2-N3	5.11	131.86	129.30
22	BA	318	C	N3-C2-O2	5.11	125.48	121.90
22	BA	658	U	C5-C4-O4	-5.11	122.83	125.90
22	BA	1283	G	C5-C6-O6	5.11	131.67	128.60
22	BA	1930	G	C8-N9-C1'	5.11	133.64	127.00
22	BA	1271	G	N3-C2-N2	5.11	123.48	119.90
22	BA	527	C	O4'-C1'-N1	5.11	112.29	108.20
22	BA	1661	G	C4-C5-N7	-5.11	108.76	110.80
22	BA	2076	U	N3-C4-O4	-5.11	115.82	119.40
22	BA	2365	G	N1-C6-O6	5.11	122.97	119.90
22	BA	2496	C	C6-N1-C1'	5.11	126.93	120.80
22	BA	1006	C	OP2-P-O3'	5.11	116.43	105.20
22	BA	1260	A	O5'-P-OP2	-5.11	101.11	105.70
1	CA	729	A	OP1-P-O3'	5.11	116.43	105.20
22	BA	866	A	OP1-P-OP2	-5.10	111.94	119.60
22	BA	1278	C	C2-N1-C1'	-5.10	113.19	118.80
22	BA	2382	G	C6-C5-N7	-5.10	127.34	130.40
22	BA	143	C	C5-C4-N4	-5.10	116.63	120.20
22	BA	565	C	OP1-P-OP2	5.10	127.25	119.60
22	BA	2018	G	N3-C4-C5	-5.10	126.05	128.60
23	BB	6	G	C6-C5-N7	-5.10	127.34	130.40
22	DA	764	A	N1-C6-N6	-5.10	115.54	118.60
22	BA	696	G	O5'-P-OP1	-5.10	101.11	105.70
22	BA	1395	A	C4-N9-C1'	-5.10	117.12	126.30
22	BA	1406	U	N3-C4-O4	-5.10	115.83	119.40
22	BA	1981	A	C4-C5-C6	-5.10	114.45	117.00
22	BA	2251	G	C4-C5-N7	-5.10	108.76	110.80
22	BA	531	C	N3-C4-C5	5.10	123.94	121.90
22	BA	1566	A	C4-C5-C6	5.10	119.55	117.00
22	BA	597	G	C6-C5-N7	-5.09	127.34	130.40
22	BA	760	G	O5'-P-OP2	5.09	116.81	110.70
22	BA	1426	G	C6-C5-N7	-5.09	127.34	130.40
1	CA	207	C	C6-N1-C2	-5.09	118.26	120.30
22	BA	993	G	N9-C4-C5	5.09	107.44	105.40
22	BA	1940	U	N3-C2-O2	-5.09	118.64	122.20
22	BA	2487	G	N1-C6-O6	5.08	122.95	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	1438	U	N3-C2-O2	5.08	125.76	122.20
22	BA	2022	U	OP1-P-OP2	-5.08	111.98	119.60
22	BA	2262	U	N3-C4-O4	-5.08	115.84	119.40
1	CA	910	C	O5'-P-OP2	5.08	116.80	110.70
23	BB	92	C	C5-C6-N1	-5.08	118.46	121.00
22	BA	513	A	N1-C6-N6	5.08	121.65	118.60
22	BA	1989	G	N3-C4-N9	5.08	129.05	126.00
22	BA	2616	C	C6-N1-C1'	5.08	126.89	120.80
23	BB	72	G	OP2-P-O3'	5.08	116.38	105.20
22	BA	907	G	C8-N9-C4	-5.08	104.37	106.40
22	BA	2645	G	O4'-C1'-N9	5.08	112.26	108.20
22	BA	2788	C	C2-N3-C4	-5.08	117.36	119.90
22	BA	770	G	C8-N9-C4	-5.07	104.37	106.40
22	BA	1549	A	C4-C5-N7	5.07	113.24	110.70
22	BA	378	C	C2-N3-C4	-5.07	117.36	119.90
22	BA	913	U	C5-C4-O4	5.07	128.94	125.90
22	BA	1779	U	O4'-C1'-N1	5.07	112.26	108.20
22	BA	1669	A	C2-N3-C4	5.07	113.14	110.60
22	BA	2645	G	C4-C5-C6	5.07	121.84	118.80
23	BB	39	A	N9-C4-C5	-5.07	103.77	105.80
22	BA	1688	U	N1-C2-N3	5.07	117.94	114.90
22	DA	1786	A	O4'-C1'-N9	5.07	112.25	108.20
22	BA	410	G	C5-C6-O6	-5.07	125.56	128.60
22	BA	2557	G	C8-N9-C4	-5.07	104.37	106.40
1	AA	556	C	O5'-P-OP1	-5.07	101.14	105.70
22	BA	758	C	O5'-P-OP2	-5.07	101.14	105.70
22	BA	761	A	C8-N9-C4	-5.07	103.77	105.80
22	DA	1837	C	O5'-P-OP1	-5.07	101.14	105.70
22	BA	2621	G	C8-N9-C4	-5.06	104.37	106.40
1	AA	1484	C	C5-C4-N4	-5.06	116.66	120.20
22	BA	1009	A	N9-C4-C5	-5.06	103.78	105.80
22	BA	1154	G	C8-N9-C4	-5.06	104.38	106.40
22	BA	2018	G	N1-C6-O6	-5.06	116.86	119.90
22	BA	2707	U	N3-C4-O4	-5.06	115.86	119.40
22	BA	533	G	C5-N7-C8	-5.06	101.77	104.30
22	BA	2056	G	O5'-P-OP2	-5.06	101.15	105.70
22	BA	2851	A	N1-C6-N6	5.06	121.64	118.60
22	BA	794	A	N1-C6-N6	5.06	121.63	118.60
22	BA	2303	G	OP1-P-O3'	5.06	116.32	105.20
22	BA	1256	G	C4-N9-C1'	5.05	133.07	126.50
22	BA	1653	G	N1-C6-O6	-5.05	116.87	119.90
22	BA	1694	C	C2-N3-C4	-5.05	117.37	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	1826	G	N3-C4-N9	5.05	129.03	126.00
22	BA	2609	U	O4'-C1'-N1	5.05	112.24	108.20
22	BA	124	G	OP2-P-O3'	5.05	116.32	105.20
22	BA	1976	U	OP1-P-O3'	5.05	116.31	105.20
39	BR	51	VAL	N-CA-C	-5.05	97.36	111.00
22	BA	240	C	C5-C4-N4	-5.05	116.67	120.20
22	BA	1826	G	C6-C5-N7	-5.05	127.37	130.40
22	BA	2040	G	C5-C6-N1	5.05	114.03	111.50
22	DA	2465	C	C6-N1-C2	-5.05	118.28	120.30
22	BA	2625	G	N1-C2-N2	5.05	120.74	116.20
1	CA	575	G	C8-N9-C1'	5.05	133.56	127.00
22	BA	2407	A	C6-C5-N7	-5.04	128.77	132.30
1	AA	326	G	N3-C4-N9	5.04	129.03	126.00
22	BA	1775	U	C5-C4-O4	-5.04	122.87	125.90
22	BA	2870	C	C6-N1-C2	5.04	122.32	120.30
22	BA	697	G	O5'-P-OP1	-5.04	101.16	105.70
23	BB	92	C	C2-N3-C4	-5.04	117.38	119.90
22	BA	2689	U	N1-C2-N3	5.04	117.92	114.90
22	BA	2838	G	N1-C6-O6	5.04	122.92	119.90
22	DA	1257	C	C6-N1-C2	-5.04	118.28	120.30
22	BA	1004	U	C6-N1-C2	-5.04	117.98	121.00
22	BA	1218	G	C2-N3-C4	-5.04	109.38	111.90
22	BA	2858	C	N3-C4-C5	5.04	123.91	121.90
22	BA	682	G	C5-C6-N1	5.03	114.02	111.50
22	BA	2827	C	C2-N3-C4	-5.03	117.38	119.90
22	BA	814	C	C5-C4-N4	5.03	123.72	120.20
22	BA	1141	U	C5-C6-N1	5.03	125.22	122.70
22	BA	1343	G	N3-C4-C5	-5.03	126.08	128.60
22	BA	1900	A	N9-C4-C5	5.03	107.81	105.80
22	BA	2890	G	N3-C2-N2	5.03	123.42	119.90
1	CA	287	U	C2-N1-C1'	-5.03	111.66	117.70
1	AA	584	G	C6-C5-N7	-5.03	127.38	130.40
22	BA	2000	C	O5'-P-OP2	-5.03	101.17	105.70
22	BA	2030	A	C6-C5-N7	5.03	135.82	132.30
22	BA	2794	C	N3-C2-O2	5.03	125.42	121.90
22	BA	2320	U	N1-C2-O2	-5.02	119.28	122.80
22	BA	493	G	N3-C4-N9	-5.02	122.99	126.00
22	BA	515	A	O4'-C1'-N9	5.02	112.22	108.20
22	BA	1775	U	N1-C2-O2	-5.02	119.29	122.80
22	BA	1273	U	N1-C2-O2	-5.02	119.29	122.80
22	BA	381	G	C5-C6-O6	-5.02	125.59	128.60
22	BA	705	A	C2-N3-C4	-5.02	108.09	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BA	1650	A	N7-C8-N9	5.02	116.31	113.80
1	AA	634	C	N3-C2-O2	-5.02	118.39	121.90
22	BA	1378	A	C2-N3-C4	5.02	113.11	110.60
22	BA	2250	G	C6-N1-C2	5.02	128.11	125.10
22	BA	2610	C	N3-C2-O2	-5.02	118.39	121.90
22	BA	2036	C	C2-N1-C1'	5.01	124.32	118.80
23	BB	92	C	N3-C4-C5	5.01	123.91	121.90
22	BA	2012	G	N1-C6-O6	5.01	122.91	119.90
22	BA	2814	A	C8-N9-C4	5.01	107.81	105.80
22	BA	1665	A	OP2-P-O3'	5.01	116.22	105.20
22	BA	914	G	C5-N7-C8	-5.01	101.80	104.30
22	BA	1274	A	C5-C6-N6	-5.01	119.69	123.70
22	BA	1644	C	N3-C4-N4	-5.01	114.49	118.00
22	BA	1829	A	C4-C5-N7	-5.01	108.19	110.70
22	BA	1936	A	C5-N7-C8	-5.01	101.39	103.90
22	BA	2837	A	C5-C6-N6	-5.01	119.69	123.70
22	BA	1035	U	N3-C4-O4	-5.01	115.89	119.40
22	BA	1432	G	OP1-P-O3'	5.01	116.21	105.20
22	BA	1938	A	N9-C4-C5	-5.01	103.80	105.80
1	AA	115	G	P-O3'-C3'	5.00	125.70	119.70
22	BA	1695	G	C5-C6-N1	5.00	114.00	111.50
22	BA	1929	G	C8-N9-C1'	-5.00	120.50	127.00
1	AA	1511	G	N1-C6-O6	5.00	122.90	119.90
22	BA	820	A	C5-C6-N1	5.00	120.20	117.70
22	BA	993	G	N3-C4-C5	-5.00	126.10	128.60
22	BA	2051	A	C5-N7-C8	-5.00	101.40	103.90

There are no chirality outliers.

All (9) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
11	AK	126	LYS	Peptide
25	BD	132	ALA	Peptide
41	BT	2	ILE	Peptide
47	BZ	15	GLY	Peptide
5	CE	102	GLY	Peptide
5	CE	104	GLY	Peptide
11	CK	126	LYS	Peptide
12	CL	38	TYR	Peptide
25	DD	151	THR	Peptide

5.2 Too-close contacts ⓘ

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	AA	32995	0	16607	893	5
1	CA	33015	0	16617	1011	0
2	AB	1705	0	1732	112	0
2	CB	1705	0	1732	107	0
3	AC	1625	0	1696	57	0
3	CC	1625	0	1696	49	0
4	AD	1643	0	1707	117	0
4	CD	1643	0	1707	96	0
5	AE	1106	0	1148	81	0
5	CE	1106	0	1148	88	0
6	AF	818	0	808	47	0
6	CF	818	0	808	47	0
7	AG	1182	0	1238	47	0
7	CG	1182	0	1238	46	0
8	AH	979	0	1031	41	0
8	CH	979	0	1031	36	0
9	AI	1022	0	1070	55	0
9	CI	1022	0	1070	36	0
10	AJ	787	0	828	42	0
10	CJ	787	0	828	32	0
11	AK	877	0	887	59	0
11	CK	877	0	887	52	0
12	AL	955	0	1016	61	0
12	CL	955	0	1016	73	0
13	AM	884	0	941	51	0
13	CM	884	0	941	35	0
14	AN	774	0	824	44	0
14	CN	774	0	824	26	0
15	AO	714	0	734	28	0
15	CO	714	0	734	36	0
16	AP	649	0	666	27	0
16	CP	649	0	666	43	0
17	AQ	649	0	691	46	0
17	CQ	649	0	691	38	0
18	AR	456	0	478	9	0
18	CR	456	0	478	22	0
19	AS	638	0	665	33	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
19	CS	638	0	665	22	0
20	AT	665	0	714	41	0
20	CT	665	0	714	34	0
21	AU	426	0	449	59	0
21	CU	426	0	449	44	0
22	BA	62195	0	31278	1182	0
22	DA	62195	0	31280	2055	0
23	BB	2549	0	1291	36	0
23	DB	2529	0	1281	42	0
24	BC	2083	0	2154	81	0
24	DC	2083	0	2154	161	0
25	BD	1565	0	1616	59	0
25	DD	1565	0	1616	69	0
26	BE	1552	0	1619	57	0
26	DE	1552	0	1619	80	0
27	BF	1411	0	1444	66	0
27	DF	1411	0	1444	26	0
28	BG	1323	0	1371	44	0
28	DG	1323	0	1371	41	0
29	BH	1110	0	1148	166	0
29	DH	1110	0	1148	86	5
30	BI	1032	0	1085	54	0
30	DI	1032	0	1085	38	0
31	BJ	1129	0	1162	41	0
31	DJ	1129	0	1162	43	0
32	BK	939	0	1012	29	0
32	DK	939	0	1012	33	0
33	BL	1045	0	1117	54	0
33	DL	1045	0	1117	59	0
34	BM	1074	0	1157	40	0
34	DM	1074	0	1157	23	0
35	BN	961	0	1000	46	0
35	DN	961	0	1000	47	0
36	BO	892	0	923	48	0
36	DO	892	0	923	36	0
37	BP	917	0	962	24	0
37	DP	917	0	962	40	0
38	BQ	947	0	1019	49	0
38	DQ	947	0	1019	47	0
39	BR	816	0	839	59	0
39	DR	816	0	839	45	0
40	BS	857	0	922	29	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
40	DS	857	0	922	29	0
41	BT	739	0	807	49	0
41	DT	739	0	807	27	0
42	BU	780	0	831	33	0
42	DU	780	0	831	61	0
43	BV	753	0	780	21	0
43	DV	753	0	780	14	0
44	BW	580	0	594	22	0
44	DW	569	0	581	11	0
45	BX	625	0	652	21	0
45	DX	625	0	652	54	0
46	BY	509	0	543	30	0
46	DY	509	0	543	24	0
47	BZ	449	0	488	12	0
47	DZ	449	0	488	15	0
48	B0	444	0	458	24	0
48	D0	444	0	458	18	0
49	B1	410	0	440	15	0
49	D1	410	0	440	12	0
50	B2	377	0	418	14	0
50	D2	377	0	418	41	0
51	B3	504	0	572	23	0
51	D3	504	0	572	22	0
52	B4	302	0	340	9	0
52	D4	302	0	342	10	0
53	B5	1142	0	865	26	0
54	AA	71	0	0	0	0
54	AM	1	0	0	0	0
54	BA	194	0	0	0	0
54	BB	4	0	0	0	0
54	BQ	1	0	0	0	0
54	CA	56	0	0	0	0
54	CT	1	0	0	0	0
54	D2	1	0	0	0	0
54	DA	164	0	0	0	0
54	DB	3	0	0	0	0
54	DL	2	0	0	0	0
54	DQ	1	0	0	0	0
55	B4	1	0	0	0	0
55	D4	1	0	0	0	0
56	CA	17	0	19	1	0
57	AA	196	0	0	18	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
57	AE	1	0	0	0	0
57	AL	1	0	0	0	0
57	AN	3	0	0	1	0
57	AT	1	0	0	0	0
57	AU	1	0	0	0	0
57	B2	1	0	0	0	0
57	B3	3	0	0	0	0
57	B4	2	0	0	0	0
57	BA	620	0	0	82	0
57	BB	14	0	0	0	0
57	BC	10	0	0	1	0
57	BD	4	0	0	2	0
57	BF	1	0	0	0	0
57	BG	1	0	0	0	0
57	BL	6	0	0	2	0
57	BN	3	0	0	0	0
57	BS	1	0	0	0	0
57	CA	186	0	0	16	0
57	CL	1	0	0	0	0
57	CN	3	0	0	0	0
57	CT	3	0	0	1	0
57	CU	1	0	0	0	0
57	D2	1	0	0	1	0
57	D3	2	0	0	0	0
57	D4	1	0	0	1	0
57	DA	611	0	0	87	0
57	DB	13	0	0	1	0
57	DC	8	0	0	0	0
57	DD	3	0	0	1	0
57	DE	5	0	0	0	0
57	DJ	1	0	0	0	0
57	DL	4	0	0	0	0
57	DN	1	0	0	0	0
57	DS	1	0	0	0	0
57	DT	3	0	0	0	0
57	DU	1	0	0	0	0
57	DV	1	0	0	0	0
All	All	288204	0	192819	8827	5

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 19.

All (8827) 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)
29:BH:94:ILE:H	29:BH:122:LEU:CB	1.08	1.58
29:BH:94:ILE:O	29:BH:122:LEU:CD1	1.69	1.41
29:BH:94:ILE:N	29:BH:122:LEU:CB	1.88	1.34
29:BH:94:ILE:N	29:BH:122:LEU:HB3	1.41	1.28
29:BH:122:LEU:HD21	1:CA:368:U:OP2	1.09	1.25
29:BH:122:LEU:CD2	1:CA:368:U:OP2	1.88	1.20
29:BH:93:SER:CA	29:BH:122:LEU:HB2	1.74	1.17
22:BA:731:C:OP2	57:BA:3699:HOH:O	1.61	1.17
29:BH:94:ILE:N	29:BH:122:LEU:HG	1.60	1.15
1:CA:1500:A:OP2	57:CA:1877:HOH:O	1.66	1.14
29:BH:94:ILE:C	29:BH:122:LEU:HD12	1.68	1.13
22:BA:842:U:O4	57:BA:3592:HOH:O	1.69	1.09
22:BA:2499:C:OP2	57:BA:3691:HOH:O	1.68	1.08
6:CF:12:PRO:O	6:CF:15:SER:OG	1.70	1.08
25:DD:151:THR:O	25:DD:153:GLY:N	1.84	1.08
22:DA:1154:G:OP2	38:DQ:58:ARG:NH1	1.87	1.08
29:BH:94:ILE:H	29:BH:122:LEU:CG	1.66	1.08
22:BA:2575:C:OP2	57:BA:3717:HOH:O	1.67	1.08
29:BH:93:SER:HA	29:BH:122:LEU:HB2	1.29	1.07
29:BH:94:ILE:N	29:BH:122:LEU:CG	2.16	1.07
22:BA:1342:A:OP2	57:BA:3721:HOH:O	1.71	1.06
22:DA:1936:A:OP1	57:DA:3459:HOH:O	1.73	1.06
22:BA:2574:G:OP1	57:BA:3717:HOH:O	1.74	1.05
39:BR:49:ILE:HG22	39:BR:53:PHE:N	1.72	1.04
22:DA:1357:C:C5	57:DA:3401:HOH:O	2.10	1.04
1:CA:1181:G:O2'	1:CA:1182:G:N7	1.89	1.03
22:DA:756:A:N7	57:DA:3299:HOH:O	1.92	1.02
29:BH:93:SER:HB2	29:BH:122:LEU:HD23	1.08	1.02
22:BA:2269:G:OP1	57:BA:3517:HOH:O	1.77	1.02
22:DA:822:G:OP2	57:DA:3348:HOH:O	1.78	1.00
22:DA:370:G:N7	57:DA:3559:HOH:O	1.92	1.00
1:AA:533:A:OP1	57:AA:1850:HOH:O	1.78	0.99
22:BA:2243:U:OP1	57:BA:3747:HOH:O	1.81	0.98
29:BH:93:SER:HB2	29:BH:122:LEU:CD2	1.93	0.98
22:DA:1010:A:OP2	57:DA:3780:HOH:O	1.83	0.97
22:DA:211:C:OP1	50:D2:25:LYS:NZ	1.97	0.97
29:DH:40:THR:O	29:DH:42:LYS:N	1.98	0.96
22:DA:1789:A:OP2	24:DC:221:ARG:NH1	1.99	0.95
4:CD:192:SER:OG	4:CD:193:ALA:N	1.93	0.94
22:DA:761:A:N7	57:DA:3295:HOH:O	1.99	0.94
29:BH:83:LYS:HD2	1:CA:55:A:O2'	1.68	0.94
22:DA:602:A:O2'	22:DA:604:G:O2'	1.81	0.93

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AB:193:PRO:O	2:AB:195:GLY:N	2.01	0.93
14:CN:41:ARG:NH1	14:CN:42:TRP:O	2.01	0.93
22:DA:2006:C:OP1	57:DA:3380:HOH:O	1.86	0.93
22:BA:622:G:OP2	57:BA:3296:HOH:O	1.86	0.93
1:AA:702:A:N6	22:BA:1846:G:O2'	2.00	0.93
2:AB:82:ASP:O	2:AB:85:LEU:N	2.02	0.93
12:AL:24:LEU:O	12:AL:26:ALA:N	2.01	0.93
22:BA:2005:A:OP1	57:BA:3387:HOH:O	1.85	0.92
22:BA:2611:C:OP2	57:BA:3548:HOH:O	1.87	0.92
1:CA:1095:U:OP2	57:CA:1849:HOH:O	1.86	0.92
22:BA:1073:A:H3'	22:BA:1074:G:H5''	1.50	0.92
25:BD:140:HIS:NE2	57:BD:303:HOH:O	2.01	0.91
22:BA:1171:G:N2	22:BA:1178:C:O2	2.02	0.91
22:BA:1916:A:H2'	22:BA:1917:U:O4'	1.70	0.91
22:BA:192:C:OP1	57:BA:3747:HOH:O	1.89	0.91
22:DA:1607:C:N4	22:DA:1622:G:N7	2.19	0.90
2:CB:103:ASN:OD1	2:CB:106:THR:OG1	1.89	0.90
29:BH:94:ILE:O	29:BH:122:LEU:HD12	0.73	0.90
22:DA:1371:G:N7	57:DA:3400:HOH:O	2.05	0.90
22:DA:58:G:OP1	41:DT:78:SER:OG	1.90	0.90
28:BG:104:ASN:ND2	28:BG:114:ASP:OD1	2.03	0.89
1:AA:537:G:OP1	12:AL:110:ARG:NH2	2.04	0.89
1:CA:201:G:N2	1:CA:469:C:O2	2.04	0.89
22:DA:618:G:O6	57:DA:3289:HOH:O	1.89	0.89
22:DA:1439:A:OP2	57:DA:3631:HOH:O	1.91	0.88
1:AA:980:C:OP1	57:AA:1838:HOH:O	1.91	0.88
22:BA:301:G:OP2	42:BU:82:ARG:NH1	2.06	0.88
1:CA:890:G:O2'	1:CA:906:A:N6	2.06	0.88
28:BG:155:GLU:OE2	28:BG:158:LYS:N	2.06	0.88
27:DF:122:PHE:O	27:DF:124:GLY:N	2.06	0.88
22:DA:1187:G:N7	57:DA:3578:HOH:O	2.05	0.88
31:BJ:17:VAL:HG23	31:BJ:137:PRO:HB2	1.55	0.88
25:BD:140:HIS:CD2	57:BD:303:HOH:O	2.27	0.88
22:DA:2627:G:O2'	22:DA:2781:A:N1	2.05	0.88
24:BC:182:ARG:NH2	24:BC:183:LYS:O	2.07	0.88
22:DA:2262:U:OP1	44:DW:41:ARG:NH2	2.07	0.88
22:BA:1509:A:O2'	22:BA:1510:G:OP2	1.92	0.87
22:BA:512:G:C8	57:BA:3777:HOH:O	2.13	0.87
22:DA:1251:C:OP2	38:DQ:6:ARG:NH2	2.07	0.87
24:DC:157:SER:O	24:DC:160:THR:OG1	1.92	0.86
22:BA:1179:G:C5	22:BA:1180:U:H1'	2.11	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:2874:C:OP1	57:DA:3803:HOH:O	1.93	0.86
4:CD:100:ASN:OD1	4:CD:111:ARG:NH1	2.08	0.86
22:DA:1395:A:OP2	57:DA:3404:HOH:O	1.94	0.86
29:BH:147:VAL:HG12	29:BH:149:GLU:HG3	1.57	0.86
22:DA:1266:G:O2'	22:DA:2012:G:O6	1.94	0.85
22:BA:572:A:OP2	39:BR:80:ARG:NH2	2.08	0.85
22:DA:618:G:N7	57:DA:3287:HOH:O	2.09	0.85
1:CA:536:C:OP1	57:CA:1770:HOH:O	1.93	0.85
17:AQ:17:MET:N	17:AQ:17:MET:SD	2.48	0.85
22:BA:1925:C:H5''	22:BA:1926:U:O4	1.77	0.85
41:BT:2:ILE:HD12	41:BT:7:LEU:HD11	1.58	0.85
22:BA:1603:A:OP1	57:BA:3415:HOH:O	1.94	0.85
29:BH:121:VAL:O	29:BH:128:HIS:CD2	2.29	0.85
40:DS:28:LYS:O	40:DS:30:SER:N	2.10	0.85
1:CA:1124:G:O2'	1:CA:1145:A:N6	2.09	0.85
22:DA:450:G:O6	57:DA:3240:HOH:O	1.93	0.85
12:CL:75:GLN:O	12:CL:78:SER:OG	1.92	0.84
29:BH:94:ILE:H	29:BH:122:LEU:HB3	0.68	0.84
21:AU:44:GLU:OE2	21:AU:45:ARG:NH1	2.09	0.84
22:BA:273:G:N2	22:BA:365:U:O2	2.10	0.84
4:AD:163:GLU:OE2	4:AD:164:GLN:N	2.11	0.84
29:DH:82:SER:O	29:DH:84:ALA:N	2.10	0.84
39:BR:24:LYS:HA	39:BR:94:THR:HG23	1.60	0.84
22:DA:2234:G:C6	22:DA:2235:G:N7	2.46	0.84
22:BA:627:A:OP1	33:BL:78:ARG:NH1	2.10	0.83
1:AA:1225:A:H2'	1:AA:1226:C:C5	2.12	0.83
12:CL:34:CYS:HA	12:CL:55:VAL:HA	1.59	0.83
39:BR:49:ILE:HG22	39:BR:52:PRO:C	1.99	0.83
22:DA:2144:G:N2	22:DA:2148:G:O6	2.11	0.83
22:DA:247:G:H4'	22:DA:386:G:C5	2.12	0.83
22:DA:2711:A:OP2	57:DA:3549:HOH:O	1.96	0.83
34:DM:66:ARG:NH1	34:DM:104:GLU:OE1	2.11	0.83
22:DA:2057:G:OP1	57:DA:3671:HOH:O	1.96	0.83
22:DA:1826:G:C5	22:DA:1827:U:C5	2.67	0.83
29:DH:94:ILE:HB	29:DH:122:LEU:HD12	1.60	0.83
41:BT:2:ILE:HD12	41:BT:7:LEU:CD1	2.08	0.83
22:DA:1317:G:C2	22:DA:1336:A:C2	2.67	0.83
1:CA:1197:A:O3'	56:CA:1657:NEG:N4	2.11	0.83
22:DA:1378:A:O2'	22:DA:1380:G:N7	2.11	0.83
22:DA:192:C:OP1	57:DA:3739:HOH:O	1.96	0.83
18:CR:20:GLU:O	18:CR:22:ASP:N	2.11	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:2720:U:OP1	37:DP:53:ARG:NH2	2.11	0.82
29:BH:117:LEU:HD21	29:BH:122:LEU:CD1	2.09	0.82
35:BN:12:ARG:NE	35:BN:20:MET:HE3	1.94	0.82
22:DA:53:A:C2	22:DA:179:C:H4'	2.14	0.82
22:BA:1916:A:H2'	22:BA:1917:U:C1'	2.08	0.82
22:DA:2171:A:O2'	22:DA:2173:A:OP1	1.97	0.82
22:DA:2271:G:O6	57:DA:3510:HOH:O	1.97	0.82
22:BA:1924:C:O2	22:BA:1926:U:O4	1.97	0.82
22:DA:1359:A:C8	22:DA:1373:A:N1	2.48	0.82
50:D2:44:VAL:O	50:D2:45:SER:OG	1.97	0.81
22:BA:761:A:C8	57:BA:3298:HOH:O	2.32	0.81
34:BM:27:SER:N	34:BM:104:GLU:OE1	2.12	0.81
13:CM:13:LYS:O	13:CM:14:HIS:ND1	2.13	0.81
22:BA:1779:U:H5	22:BA:1784:A:N7	1.78	0.81
22:DA:201:C:C5	22:DA:202:U:C5	2.69	0.81
22:BA:1917:U:H2'	22:BA:1918:A:H5'	1.62	0.80
22:BA:1916:A:N3	22:BA:1917:U:H1'	1.96	0.80
31:BJ:81:ILE:HG23	31:BJ:82:GLY:H	1.45	0.80
1:AA:1397:C:O2'	1:AA:1398:A:OP1	1.99	0.80
22:BA:2191:A:C2	22:BA:2192:U:C2	2.69	0.80
1:AA:452:A:N6	1:AA:480:U:O2	2.14	0.80
29:BH:91:PHE:O	1:CA:55:A:C6	2.34	0.80
31:BJ:81:ILE:HG23	31:BJ:82:GLY:N	1.96	0.80
1:AA:939:G:N7	57:AA:1771:HOH:O	2.13	0.80
1:CA:373:A:C2	1:CA:374:A:C8	2.70	0.80
45:BX:2:SER:O	45:BX:4:VAL:N	2.15	0.80
42:BU:16:GLY:O	42:BU:18:ASP:N	2.15	0.80
22:DA:761:A:OP2	57:DA:3293:HOH:O	1.98	0.80
22:BA:1179:G:C6	22:BA:1180:U:H1'	2.17	0.80
4:AD:100:ASN:OD1	4:AD:111:ARG:NH1	2.14	0.80
7:AG:68:ASN:O	7:AG:138:ARG:NH1	2.14	0.80
1:AA:880:C:OP1	12:AL:9:ARG:NH1	2.13	0.80
16:CP:23:ASP:OD2	16:CP:25:ARG:NH2	2.15	0.80
22:DA:616:A:H4'	26:DE:101:TYR:CZ	2.16	0.80
22:DA:2243:U:OP1	57:DA:3742:HOH:O	1.98	0.80
11:AK:126:LYS:C	21:AU:34:ARG:CZ	2.51	0.79
3:AC:40:ARG:NH1	3:AC:55:ILE:O	2.15	0.79
22:BA:2637:U:C2'	22:BA:2638:G:H5'	2.12	0.79
29:BH:119:ASN:HB3	29:BH:123:ARG:HH12	1.47	0.79
1:CA:505:G:C6	1:CA:535:A:C2	2.70	0.79
22:DA:1344:U:O2'	22:DA:1345:C:OP2	1.99	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:1131:G:OP1	31:BJ:82:GLY:HA2	1.82	0.79
22:DA:132:G:N2	22:DA:148:U:C2	2.51	0.79
5:AE:65:GLU:OE2	5:AE:69:ARG:NH2	2.16	0.79
24:BC:247:PRO:HD2	24:BC:248:TRP:CZ3	2.18	0.79
1:CA:794:A:HO2'	1:CA:1521:C:HO2'	1.08	0.79
1:AA:1223:C:OP2	19:AS:78:ARG:NH1	2.16	0.79
22:DA:1050:A:N6	22:DA:1109:C:O2	2.16	0.79
17:AQ:52:GLU:N	17:AQ:52:GLU:OE1	2.14	0.79
22:BA:1998:A:OP2	25:BD:141:ARG:NH2	2.16	0.79
37:DP:65:SER:O	37:DP:67:GLY:N	2.15	0.79
22:BA:12:U:O2	22:BA:12:U:H2'	1.83	0.79
29:BH:122:LEU:O	29:BH:122:LEU:HD22	1.83	0.78
22:BA:2748:A:H1'	28:BG:67:THR:HG22	1.65	0.78
24:BC:168:ASP:OD1	24:BC:169:GLY:N	2.14	0.78
34:DM:76:LYS:NZ	34:DM:85:GLY:O	2.16	0.78
4:AD:32:CYS:SG	4:AD:33:LYS:N	2.55	0.78
22:DA:2883:A:OP1	48:D0:49:TYR:OH	2.01	0.78
5:CE:104:GLY:HA3	5:CE:122:ASN:HA	1.64	0.78
11:AK:69:ARG:HD2	22:BA:2146:C:N3	1.99	0.78
24:BC:204:VAL:O	24:BC:205:LEU:HB2	1.82	0.78
11:AK:127:ARG:N	21:AU:34:ARG:NH1	2.31	0.78
22:BA:2187:U:C4	22:BA:2188:U:C4	2.72	0.78
22:DA:1984:G:C5	22:DA:1985:C:C5	2.72	0.78
5:CE:157:ARG:O	5:CE:159:LYS:N	2.17	0.78
1:AA:516:U:O4	57:AA:1848:HOH:O	2.00	0.78
6:AF:16:GLU:OE2	4:CD:188:ARG:NH1	2.17	0.78
39:DR:60:LYS:NZ	39:DR:102:SER:OG	2.17	0.78
25:BD:103:ASP:O	25:BD:105:LYS:N	2.16	0.77
22:DA:1851:U:N3	22:DA:1891:G:O6	2.16	0.77
1:AA:1166:G:N1	1:AA:1169:A:OP2	2.17	0.77
22:DA:2550:G:O6	22:DA:2551:C:N4	2.17	0.77
29:DH:124:THR:OG1	29:DH:125:THR:N	2.17	0.77
29:BH:118:PRO:O	29:BH:120:GLY:N	2.18	0.77
29:BH:83:LYS:HG3	1:CA:55:A:N3	2.00	0.77
35:BN:73:ASN:HA	35:BN:76:VAL:HG12	1.67	0.77
1:CA:484:G:H4'	1:CA:485:U:O5'	1.85	0.77
22:DA:514:A:N3	22:DA:581:C:O2'	2.15	0.77
5:CE:98:PRO:O	5:CE:122:ASN:ND2	2.16	0.77
22:BA:2321:U:H5'	22:BA:2322:A:OP2	1.84	0.77
1:AA:1232:U:OP1	9:AI:126:GLN:NE2	2.18	0.77
1:CA:977:A:O2'	1:CA:1223:C:N4	2.18	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:1917:U:C4	22:BA:1918:A:C5	2.73	0.77
22:DA:2800:A:C2	22:DA:2895:G:H1'	2.20	0.77
22:DA:1847:A:HO2'	22:DA:1848:A:H8	1.32	0.76
1:AA:328:C:O2	1:AA:328:C:H2'	1.84	0.76
29:BH:83:LYS:CD	1:CA:55:A:O2'	2.33	0.76
22:DA:1791:A:N6	22:DA:1828:G:O2'	2.17	0.76
29:DH:1:MET:SD	29:DH:27:ARG:NH1	2.58	0.76
22:DA:250:G:OP2	51:D3:13:ARG:NH1	2.18	0.76
22:BA:481:G:C4	22:BA:507:A:C2	2.74	0.76
22:DA:1671:U:OP2	57:DA:3434:HOH:O	2.03	0.76
22:DA:2005:A:OP1	57:DA:3384:HOH:O	2.02	0.76
1:CA:527:G:C2	1:CA:528:C:C6	2.73	0.76
1:AA:701:U:O2	1:AA:703:G:N1	2.19	0.76
22:DA:2821:A:OP2	25:DD:115:GLY:N	2.18	0.76
29:DH:53:GLU:O	29:DH:55:GLU:N	2.19	0.76
22:DA:842:U:O4	57:DA:3581:HOH:O	2.02	0.76
29:DH:45:GLU:O	29:DH:49:ALA:N	2.19	0.76
22:BA:2683:C:OP1	37:BP:51:ARG:NH2	2.18	0.76
22:DA:466:A:N1	22:DA:795:C:O2'	2.19	0.76
38:BQ:87:SER:HB3	39:BR:51:VAL:HA	1.68	0.76
22:DA:1826:G:C6	22:DA:1827:U:C4	2.73	0.76
1:CA:62:U:O2'	1:CA:379:C:O2	2.03	0.76
29:BH:94:ILE:C	29:BH:122:LEU:CD1	2.40	0.76
22:DA:1209:U:O2	22:DA:1210:G:N2	2.19	0.76
22:DA:1995:U:OP1	57:DA:3808:HOH:O	2.03	0.76
27:BF:17:MET:HE1	27:BF:22:TYR:O	1.85	0.76
21:CU:44:GLU:OE1	21:CU:45:ARG:NH1	2.19	0.75
39:BR:49:ILE:CG2	39:BR:53:PHE:N	2.49	0.75
22:DA:512:G:N7	57:DA:3769:HOH:O	2.19	0.75
24:BC:237:GLY:O	57:BC:309:HOH:O	2.04	0.75
4:CD:30:THR:C	4:CD:31:LYS:HD3	2.06	0.75
22:DA:186:G:N2	22:DA:211:C:O2	2.19	0.75
22:DA:183:C:C5	22:DA:184:C:C5	2.74	0.75
22:DA:1316:U:C2	22:DA:1337:G:N2	2.54	0.75
1:AA:673:A:H2'	1:AA:674:G:C8	2.21	0.75
4:CD:25:VAL:O	4:CD:26:ARG:O	2.04	0.75
29:BH:117:LEU:HD21	29:BH:122:LEU:HD13	1.67	0.75
6:CF:91:ARG:O	6:CF:92:THR:OG1	2.04	0.75
1:AA:79:G:N2	1:AA:91:U:O4	2.19	0.75
22:BA:2757:A:N1	28:BG:67:THR:HG21	2.01	0.75
1:AA:108:G:C6	20:AT:10:ARG:HG2	2.22	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:154:U:C2	1:AA:168:G:N2	2.55	0.74
1:CA:818:G:O2'	1:CA:819:A:H5'	1.86	0.74
1:AA:1031:C:O2'	1:AA:1032:G:OP2	2.05	0.74
22:DA:724:U:H2'	22:DA:725:G:O4'	1.87	0.74
7:CG:24:ALA:O	7:CG:28:ASN:ND2	2.19	0.74
22:DA:1984:G:C6	22:DA:1985:C:C4	2.74	0.74
1:CA:803:G:OP1	57:CA:1800:HOH:O	2.04	0.74
8:AH:42:GLU:N	8:AH:42:GLU:OE1	2.20	0.74
1:AA:451:A:C8	1:AA:452:A:C2	2.75	0.74
22:DA:1339:G:O4'	22:DA:1393:A:C2	2.39	0.74
22:DA:782:A:O2'	24:DC:224:ALA:O	2.03	0.74
22:BA:2127:G:O2'	22:BA:2128:G:O5'	2.04	0.74
22:BA:792:A:N3	22:BA:2072:C:O2'	2.19	0.74
35:DN:90:ARG:CZ	35:DN:116:VAL:HG11	2.17	0.74
29:BH:88:GLY:O	29:BH:125:THR:OG1	2.04	0.74
13:AM:11:ASP:OD1	13:AM:12:HIS:N	2.20	0.74
5:CE:106:ILE:HD11	5:CE:124:LEU:HD23	1.68	0.74
1:AA:69:G:O6	1:AA:98:A:N6	2.20	0.74
22:BA:198:C:OP2	57:BA:3767:HOH:O	2.04	0.74
1:AA:685:G:N1	1:AA:686:U:O4	2.21	0.74
2:CB:131:LYS:O	2:CB:135:LEU:N	2.20	0.74
29:BH:91:PHE:HB3	1:CA:55:A:C4	2.23	0.74
27:BF:61:SER:O	27:BF:63:GLN:N	2.18	0.74
1:CA:17:U:H2'	1:CA:18:C:C6	2.23	0.74
31:DJ:41:LYS:O	31:DJ:44:TYR:N	2.20	0.74
22:DA:2407:A:OP2	57:DA:3562:HOH:O	2.05	0.74
22:BA:686:U:OP2	57:BA:3727:HOH:O	2.04	0.74
17:AQ:16:LYS:C	17:AQ:17:MET:SD	2.66	0.74
22:DA:2550:G:OP1	57:DA:3722:HOH:O	2.03	0.74
22:DA:249:C:O5'	22:DA:2394:C:O2'	2.06	0.73
26:DE:58:LYS:NZ	26:DE:70:SER:O	2.20	0.73
1:AA:131:A:H2'	1:AA:132:C:C6	2.23	0.73
22:BA:572:A:H5''	22:BA:573:U:OP2	1.89	0.73
41:BT:2:ILE:HG22	41:BT:3:ARG:N	2.03	0.73
22:DA:784:G:OP1	57:DA:3314:HOH:O	2.05	0.73
24:DC:141:VAL:HG11	24:DC:190:ALA:HB1	1.71	0.73
1:AA:1077:G:N7	57:AA:1793:HOH:O	2.20	0.73
1:CA:1521:C:C4	1:CA:1522:U:C5	2.75	0.73
1:CA:687:A:O2'	1:CA:701:U:O4	2.05	0.73
27:BF:73:SER:OG	27:BF:80:ARG:HA	1.88	0.73
1:AA:203:G:N2	1:AA:215:C:C2	2.56	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:1746:A:H2'	22:BA:1747:U:C6	2.22	0.73
22:DA:185:G:C6	22:DA:212:G:N2	2.56	0.73
2:AB:188:ASP:HB2	2:AB:204:ASP:HB3	1.71	0.73
11:AK:76:GLU:N	11:AK:76:GLU:OE2	2.22	0.73
22:BA:977:G:C6	57:BA:3598:HOH:O	2.41	0.73
22:DA:1359:A:C2	22:DA:1360:G:H1'	2.24	0.73
21:CU:18:ARG:O	21:CU:21:ARG:N	2.21	0.73
19:AS:29:LYS:HB3	19:AS:30:PRO:HD2	1.71	0.73
29:BH:93:SER:CB	29:BH:122:LEU:HB2	2.18	0.73
29:BH:93:SER:HA	29:BH:122:LEU:CB	2.14	0.73
22:BA:1917:U:C2'	22:BA:1918:A:H5'	2.17	0.73
1:CA:724:G:OP2	1:CA:833:G:O2'	2.06	0.73
24:DC:15:HIS:O	24:DC:204:VAL:HG21	1.89	0.73
22:DA:2209:G:C2	22:DA:2216:G:C2	2.75	0.73
22:DA:593:U:H2'	22:DA:594:U:C6	2.23	0.73
22:BA:997:G:OP1	38:BQ:92:ARG:HG2	1.89	0.73
22:BA:370:G:OP2	57:BA:3565:HOH:O	2.07	0.73
22:DA:1019:U:OP1	22:DA:1035:U:O2'	2.06	0.73
46:BY:56:LEU:O	46:BY:57:LEU:CB	2.37	0.73
1:AA:970:C:N4	9:AI:130:ARG:OXT	2.20	0.73
1:CA:409:U:OP1	4:CD:24:GLY:HA3	1.89	0.72
22:DA:674:G:H1'	26:DE:69:ARG:NE	2.05	0.72
22:BA:588:U:H2'	22:BA:589:U:C6	2.23	0.72
22:BA:1508:A:O2'	22:BA:1509:A:O4'	2.07	0.72
4:AD:3:ARG:CZ	4:AD:115:ARG:HD3	2.19	0.72
22:DA:377:G:C6	22:DA:378:C:C4	2.77	0.72
1:AA:1406:U:C5	1:AA:1407:C:C5	2.77	0.72
1:CA:31:G:N7	1:CA:306:A:H1'	2.04	0.72
22:DA:910:A:N3	22:DA:2264:C:O2'	2.22	0.72
27:BF:25:VAL:O	27:BF:28:VAL:HG12	1.89	0.72
22:DA:84:A:N1	22:DA:98:G:O2'	2.18	0.72
22:DA:161:A:H3'	22:DA:162:U:H5''	1.72	0.72
26:BE:108:ILE:HD11	26:BE:180:LEU:HB3	1.69	0.72
35:BN:103:ARG:HB2	35:BN:110:MET:HE3	1.70	0.72
22:DA:2032:G:N7	57:DA:3533:HOH:O	2.22	0.72
22:DA:370:G:O2'	22:DA:424:G:OP1	2.07	0.72
22:DA:1973:G:OP1	57:DA:3464:HOH:O	2.06	0.72
22:DA:374:A:C6	22:DA:401:A:C8	2.77	0.72
22:BA:1141:U:H4'	22:BA:1142:A:O4'	1.88	0.72
25:DD:12:THR:OG1	25:DD:13:ARG:N	2.18	0.72
24:DC:210:ALA:HA	24:DC:213:TRP:CE2	2.24	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AL:21:VAL:HG23	12:AL:95:TYR:CE2	2.24	0.72
1:AA:496:A:C2	1:AA:497:G:C5	2.77	0.72
22:DA:2248:C:OP2	57:DA:3505:HOH:O	2.05	0.72
22:DA:185:G:C6	22:DA:212:G:C2	2.78	0.72
1:CA:995:C:N3	1:CA:1046:A:O2'	2.23	0.72
22:DA:528:A:C2	22:DA:2043:C:H4'	2.23	0.72
22:DA:287:G:C2	22:DA:354:A:C2	2.78	0.72
22:BA:2355:G:O3'	44:BW:24:LYS:NZ	2.23	0.72
1:CA:1309:G:C6	1:CA:1329:A:C2	2.78	0.72
22:DA:995:C:O2	31:DJ:3:THR:OG1	2.07	0.72
22:DA:746:U:HO2'	22:DA:2611:C:HO2'	1.37	0.71
3:AC:7:PRO:HG2	3:AC:184:TYR:CD2	2.25	0.71
5:CE:137:VAL:HG22	5:CE:137:VAL:O	1.90	0.71
22:DA:2550:G:C6	22:DA:2551:C:N4	2.57	0.71
1:CA:1169:A:C2	1:CA:1170:A:C4	2.78	0.71
22:DA:570:G:OP1	57:DA:3773:HOH:O	2.08	0.71
22:BA:1394:U:P	57:BA:3413:HOH:O	2.48	0.71
23:DB:8:C:O2'	36:DO:25:ARG:NH1	2.23	0.71
41:BT:2:ILE:HD11	41:BT:45:ALA:HB1	1.70	0.71
22:DA:247:G:H4'	22:DA:386:G:C4	2.24	0.71
50:D2:43:THR:OG1	50:D2:44:VAL:N	2.22	0.71
29:BH:119:ASN:O	29:BH:123:ARG:NH1	2.23	0.71
29:DH:31:VAL:HB	29:DH:32:PRO:CD	2.20	0.71
22:BA:574:A:OP2	57:BA:3269:HOH:O	2.07	0.71
22:DA:1645:G:OP1	22:DA:1646:C:H5'	1.90	0.71
4:CD:32:CYS:SG	4:CD:33:LYS:N	2.63	0.71
22:BA:2498:C:OP2	57:BA:3690:HOH:O	2.08	0.71
1:AA:958:A:N6	1:AA:959:A:N1	2.37	0.71
22:DA:690:G:O2'	22:DA:780:G:OP1	2.07	0.71
29:BH:120:GLY:O	29:BH:122:LEU:N	2.23	0.71
22:DA:396:G:H4'	45:DX:29:PHE:O	1.90	0.71
22:DA:1805:A:N3	22:DA:1813:G:C2	2.58	0.71
22:DA:1544:A:N6	22:DA:1545:A:N1	2.39	0.71
4:CD:29:ASP:O	4:CD:31:LYS:NZ	2.23	0.71
22:DA:1973:G:C5	22:DA:1974:C:C5	2.79	0.71
4:CD:151:LYS:O	4:CD:152:GLN:OE1	2.08	0.71
22:DA:1821:A:OP1	24:DC:200:HIS:NE2	2.20	0.71
1:AA:532:A:O3'	57:AA:1849:HOH:O	2.07	0.71
17:AQ:17:MET:HG2	17:AQ:20:SER:HB3	1.73	0.71
22:DA:1336:A:H2'	22:DA:1337:G:C8	2.26	0.71
22:DA:1315:C:OP2	57:DA:3762:HOH:O	2.07	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:2720:U:OP1	37:BP:53:ARG:NH2	2.24	0.71
1:AA:914:A:C4	1:AA:915:A:C8	2.78	0.71
1:AA:468:A:C2	1:AA:469:C:C4	2.79	0.71
22:DA:1619:G:N7	57:DA:3644:HOH:O	2.22	0.71
22:BA:2049:G:N7	57:BA:3679:HOH:O	2.22	0.71
1:AA:9:G:OP2	5:AE:126:LYS:NZ	2.20	0.71
24:DC:247:PRO:HG2	24:DC:248:TRP:CZ3	2.25	0.71
16:AP:43:ALA:O	16:AP:46:LYS:HD2	1.90	0.71
22:BA:877:A:N6	22:BA:899:A:N6	2.39	0.71
11:AK:125:LYS:O	21:AU:34:ARG:NE	2.24	0.71
22:DA:511:U:OP2	57:DA:3769:HOH:O	2.08	0.71
22:DA:1450:G:C6	22:DA:1451:C:N4	2.59	0.71
22:DA:1652:A:OP1	35:DN:8:ARG:NH2	2.24	0.71
1:CA:109:A:O2'	1:CA:326:G:N2	2.24	0.71
22:BA:528:A:C8	22:BA:528:A:H3'	2.25	0.71
22:BA:1509:A:O2'	22:BA:1510:G:P	2.49	0.70
22:BA:1380:G:OP2	57:BA:3758:HOH:O	2.09	0.70
1:AA:1220:G:OP1	19:AS:37:ARG:NH2	2.24	0.70
22:BA:245:G:O6	51:B3:8:ARG:HD3	1.91	0.70
1:CA:552:U:C4	1:CA:553:A:N7	2.59	0.70
22:DA:1581:G:C6	22:DA:1582:C:N4	2.59	0.70
1:CA:978:A:OP2	1:CA:1362:A:N6	2.24	0.70
28:BG:80:THR:HG22	28:BG:81:GLU:N	2.06	0.70
1:CA:1001:C:H2'	1:CA:1002:G:C8	2.26	0.70
22:DA:449:A:OP2	57:DA:3241:HOH:O	2.09	0.70
22:DA:251:A:H4'	33:DL:47:ARG:NH2	2.06	0.70
22:DA:1196:C:H1'	22:DA:1226:A:C4	2.27	0.70
1:CA:207:C:O2'	1:CA:213:G:N2	2.24	0.70
12:CL:100:GLY:N	12:CL:105:SER:O	2.23	0.70
22:DA:591:U:C2	22:DA:592:A:C8	2.79	0.70
22:DA:370:G:OP2	57:DA:3560:HOH:O	2.10	0.70
17:AQ:69:LYS:O	17:AQ:70:THR:HB	1.91	0.70
22:DA:1208:C:C4	22:DA:1209:U:C5	2.78	0.70
2:CB:15:HIS:O	2:CB:17:GLY:N	2.24	0.70
1:CA:577:G:C2	1:CA:578:C:C6	2.80	0.70
1:CA:1107:C:C2	1:CA:1108:G:C8	2.80	0.70
1:AA:178:C:OP2	20:AT:60:ARG:NH2	2.24	0.70
27:BF:127:ASN:OD1	27:BF:157:THR:HA	1.91	0.70
1:CA:933:G:N7	7:CG:3:ARG:NH2	2.38	0.70
22:BA:2498:C:OP2	57:BA:3691:HOH:O	2.08	0.70
12:CL:68:GLY:O	12:CL:99:ARG:NH1	2.24	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:826:U:O2'	33:DL:53:GLY:HA3	1.92	0.70
5:AE:82:GLN:NE2	5:AE:150:PRO:HD3	2.07	0.70
22:BA:784:G:OP2	57:BA:3316:HOH:O	2.08	0.70
22:DA:2711:A:OP2	57:DA:3546:HOH:O	2.09	0.70
46:DY:56:LEU:O	46:DY:57:LEU:HB3	1.92	0.70
22:DA:684:G:C2	22:DA:794:A:C2	2.79	0.70
22:BA:1433:A:O2'	22:BA:1434:A:H5'	1.92	0.70
29:BH:94:ILE:HG22	29:BH:99:ILE:HG13	1.72	0.70
14:AN:61:ARG:O	14:AN:62:ASN:HB2	1.89	0.70
1:AA:322:C:O2'	20:AT:18:ARG:HG3	1.92	0.70
1:CA:374:A:O3'	16:CP:70:ARG:NH1	2.25	0.69
5:AE:101:GLU:HB3	5:AE:122:ASN:HB3	1.73	0.69
3:AC:25:ASN:O	3:AC:27:LYS:N	2.24	0.69
5:AE:157:ARG:HD2	8:AH:43:GLU:O	1.92	0.69
22:DA:1469:A:C2	22:DA:1470:A:C5	2.80	0.69
22:DA:564:C:O4'	38:DQ:37:GLN:NE2	2.24	0.69
1:CA:1027:C:N4	1:CA:1034:G:O6	2.25	0.69
6:CF:13:ASP:O	6:CF:15:SER:N	2.25	0.69
2:AB:82:ASP:O	2:AB:84:ALA:N	2.25	0.69
22:DA:447:A:H5'	22:DA:449:A:C5	2.26	0.69
1:CA:247:G:C6	1:CA:278:G:C2	2.80	0.69
22:DA:89:A:C2	22:DA:90:U:C2	2.80	0.69
22:DA:1823:G:N7	57:DA:3659:HOH:O	2.26	0.69
22:DA:1839:G:O6	57:DA:3810:HOH:O	2.06	0.69
22:DA:301:G:C2	22:DA:302:C:N3	2.60	0.69
1:AA:771:G:O2'	1:AA:772:U:H5'	1.92	0.69
29:DH:27:ARG:HE	45:DX:60:ASP:CG	1.95	0.69
5:AE:159:LYS:O	8:AH:64:LYS:NZ	2.25	0.69
22:DA:2467:C:N4	22:DA:2468:A:C6	2.60	0.69
22:DA:753:A:C2	22:DA:754:U:C2	2.81	0.69
8:CH:92:LEU:O	8:CH:117:ARG:NH1	2.26	0.69
22:DA:1364:G:N7	45:DX:2:SER:N	2.41	0.69
5:CE:15:LEU:HD12	5:CE:15:LEU:C	2.12	0.69
44:BW:49:ALA:O	44:BW:50:ASN:HB2	1.92	0.69
29:BH:94:ILE:CA	29:BH:122:LEU:HG	2.22	0.69
22:DA:2550:G:C6	22:DA:2551:C:C4	2.81	0.69
1:AA:1277:C:HO2'	1:AA:1279:G:H8	1.38	0.69
23:DB:43:C:O2	27:DF:92:ARG:NH2	2.25	0.69
1:CA:32:A:C2	1:CA:33:A:C5	2.80	0.69
2:CB:21:ARG:HA	2:CB:21:ARG:NH1	2.06	0.69
22:BA:2271:G:O6	57:BA:3518:HOH:O	2.09	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:DC:66:ASP:OD2	24:DC:102:ARG:NH1	2.25	0.69
29:BH:116:ARG:O	29:BH:118:PRO:HD3	1.92	0.69
29:BH:117:LEU:HD11	29:BH:121:VAL:N	2.05	0.69
29:BH:93:SER:C	29:BH:122:LEU:CB	2.60	0.69
22:BA:2192:U:C4	22:BA:2193:G:N7	2.60	0.69
29:BH:119:ASN:HB3	29:BH:123:ARG:NH1	2.06	0.69
1:AA:405:U:OP1	1:AA:406:G:O2'	2.09	0.69
22:BA:276:U:O2'	22:BA:278:A:N7	2.26	0.69
22:DA:2093:G:C5	22:DA:2225:A:C8	2.81	0.69
42:DU:82:ARG:HB2	42:DU:97:LYS:HG3	1.75	0.69
24:DC:108:LYS:N	24:DC:194:GLU:O	2.24	0.69
22:DA:1300:G:O6	22:DA:1626:A:O2'	2.11	0.69
29:BH:97:ARG:HD2	1:CA:369:G:O2'	1.93	0.69
22:BA:2502:G:OP2	57:BA:3496:HOH:O	2.10	0.69
22:BA:2196:C:O2'	22:BA:2197:U:H5'	1.93	0.69
17:AQ:15:ASP:C	17:AQ:17:MET:SD	2.71	0.69
1:CA:577:G:C4	1:CA:816:A:C2	2.80	0.69
1:CA:673:A:H2'	1:CA:674:G:C8	2.28	0.69
22:DA:658:U:N3	22:DA:659:G:N7	2.41	0.69
22:BA:2445:G:OP1	26:BE:69:ARG:NH2	2.26	0.69
33:BL:87:GLY:O	33:BL:89:VAL:N	2.26	0.69
22:BA:2151:U:H2'	22:BA:2152:G:C8	2.28	0.69
22:BA:2448:A:OP2	57:BA:3691:HOH:O	2.09	0.68
22:DA:1350:C:N3	22:DA:1382:G:C2	2.61	0.68
4:AD:59:GLN:O	4:AD:63:ARG:HG2	1.93	0.68
22:BA:686:U:H2'	22:BA:788:A:N1	2.09	0.68
1:CA:579:A:H2'	1:CA:580:C:C6	2.28	0.68
2:AB:73:LYS:O	2:AB:75:ALA:N	2.26	0.68
22:BA:2499:C:OP1	57:BA:3693:HOH:O	2.10	0.68
22:BA:977:G:O6	57:BA:3598:HOH:O	2.10	0.68
22:DA:789:A:N1	57:DA:3310:HOH:O	2.27	0.68
2:CB:193:PRO:O	2:CB:195:GLY:N	2.26	0.68
22:BA:2297:A:N1	22:BA:2321:U:H5	1.92	0.68
1:CA:1108:G:O6	57:CA:1851:HOH:O	2.10	0.68
22:BA:2305:U:C2	27:BF:151:GLY:HA3	2.29	0.68
1:AA:1323:G:H2'	1:AA:1324:A:C8	2.29	0.68
23:BB:8:C:O3'	36:BO:25:ARG:NH1	2.26	0.68
25:BD:2:ILE:HG13	25:BD:100:LEU:HD21	1.75	0.68
22:DA:2780:G:N1	31:DJ:102:GLU:OE2	2.26	0.68
38:BQ:112:LYS:NZ	39:BR:50:GLY:HA2	2.08	0.68
22:DA:1715:G:O2'	22:DA:1743:G:O6	2.09	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:CC:40:ARG:HG2	3:CC:55:ILE:HD11	1.74	0.68
1:CA:1178:G:N2	1:CA:1181:G:OP2	2.26	0.68
1:AA:254:G:OP1	17:AQ:70:THR:HG22	1.93	0.68
39:BR:66:HIS:CE1	39:BR:94:THR:HG22	2.29	0.68
24:BC:247:PRO:HD2	24:BC:248:TRP:CE3	2.28	0.68
20:CT:5:LYS:O	20:CT:7:ALA:N	2.25	0.68
6:CF:45:ARG:O	6:CF:56:LYS:HA	1.94	0.68
1:CA:811:C:O2'	1:CA:901:A:N1	2.24	0.68
22:BA:2192:U:C2	22:BA:2193:G:C8	2.82	0.68
22:BA:2502:G:H5'	22:BA:2503:A:H5''	1.76	0.68
40:BS:63:GLY:O	40:BS:64:ALA:CB	2.42	0.68
39:BR:51:VAL:HB	39:BR:52:PRO:CD	2.24	0.68
22:BA:1922:G:C2	22:BA:1923:U:C6	2.81	0.68
22:DA:2091:C:H3'	22:DA:2092:U:H5''	1.76	0.68
22:DA:777:G:N7	22:DA:793:A:H2	1.92	0.68
22:BA:181:A:H2'	22:BA:182:A:C8	2.29	0.68
39:BR:3:ALA:HB3	39:BR:59:ILE:HD11	1.75	0.68
22:DA:297:G:H5''	42:DU:85:PHE:HB2	1.75	0.68
22:DA:771:G:C2	22:DA:772:C:C6	2.81	0.68
17:AQ:14:SER:HB3	17:AQ:22:VAL:HG12	1.75	0.68
39:DR:58:VAL:HG13	39:DR:102:SER:HB2	1.76	0.68
2:AB:188:ASP:OD2	2:AB:204:ASP:CG	2.32	0.68
8:CH:114:ARG:O	8:CH:117:ARG:N	2.25	0.68
22:BA:2380:C:OP1	36:BO:17:LYS:NZ	2.24	0.68
22:DA:1938:A:C6	22:DA:2590:A:H1'	2.29	0.68
22:DA:201:C:C4	22:DA:202:U:C5	2.81	0.68
22:BA:2845:U:H5''	37:BP:52:ASN:O	1.94	0.68
1:CA:577:G:C8	1:CA:816:A:N1	2.62	0.68
22:DA:740:C:H5'	22:DA:1784:A:H3'	1.74	0.68
5:AE:90:THR:HG22	5:AE:91:GLY:N	2.09	0.68
22:DA:260:G:C6	22:DA:261:G:N7	2.62	0.68
1:CA:1377:A:C5	7:CG:7:ILE:CD1	2.77	0.68
22:DA:42:A:C2	22:DA:438:G:C2	2.82	0.68
22:BA:1288:G:C4	22:BA:1327:A:C2	2.82	0.68
28:BG:149:ARG:HG2	28:BG:149:ARG:HH11	1.59	0.68
2:CB:87:CYS:O	2:CB:89:GLN:N	2.26	0.68
29:BH:93:SER:C	29:BH:122:LEU:HB2	2.14	0.67
11:CK:126:LYS:O	21:CU:34:ARG:NE	2.27	0.67
22:BA:877:A:C6	22:BA:899:A:C6	2.82	0.67
22:DA:443:A:C8	26:DE:40:ARG:HD3	2.29	0.67
20:AT:5:LYS:O	20:AT:7:ALA:N	2.27	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:123:U:OP1	1:CA:311:C:O2'	2.11	0.67
22:DA:1737:G:C6	22:DA:1738:G:N1	2.62	0.67
22:DA:563:A:C4	22:DA:2018:G:C2	2.83	0.67
1:AA:119:A:C2	1:AA:240:G:C8	2.82	0.67
22:DA:1091:G:O2'	22:DA:1092:C:OP2	2.08	0.67
4:CD:35:GLU:O	4:CD:37:ALA:N	2.27	0.67
33:BL:82:LEU:HD23	33:BL:83:ALA:N	2.08	0.67
22:DA:2093:G:C6	22:DA:2225:A:C8	2.82	0.67
22:DA:749:A:C5	22:DA:750:A:N7	2.63	0.67
22:DA:669:G:N2	22:DA:670:A:N1	2.42	0.67
4:AD:191:LEU:O	4:AD:192:SER:HB3	1.93	0.67
22:DA:1805:A:C2	22:DA:1813:G:C2	2.83	0.67
12:CL:90:LEU:HB2	12:CL:93:VAL:HG21	1.75	0.67
4:AD:107:PHE:CG	4:AD:145:ILE:HD11	2.29	0.67
5:CE:56:VAL:N	5:CE:57:PRO:HD2	2.10	0.67
22:DA:398:C:OP1	45:DX:32:ASN:ND2	2.27	0.67
21:AU:37:PHE:HA	21:AU:40:LYS:HE3	1.76	0.67
22:DA:1627:G:C2	22:DA:1628:G:C8	2.82	0.67
2:CB:16:PHE:CE2	2:CB:18:HIS:CE1	2.82	0.67
1:CA:309:A:O2'	1:CA:607:A:N1	2.25	0.67
24:DC:70:ASN:O	24:DC:72:ASP:N	2.28	0.67
22:DA:1935:G:H1'	22:DA:1964:G:N2	2.10	0.67
22:DA:1427:A:N6	22:DA:1571:A:OP2	2.26	0.67
51:B3:31:HIS:CD2	51:B3:32:ILE:HG13	2.30	0.67
22:BA:142:A:C5	22:BA:143:C:C4	2.83	0.67
22:DA:2427:C:OP1	57:DA:3699:HOH:O	2.11	0.67
11:CK:127:ARG:N	21:CU:34:ARG:NH2	2.42	0.67
5:CE:102:GLY:O	5:CE:104:GLY:N	2.28	0.67
22:BA:2591:C:OP2	24:BC:237:GLY:O	2.13	0.67
22:DA:1773:A:N3	22:DA:1978:A:C2	2.62	0.67
1:CA:1266:G:N2	1:CA:1269:A:OP2	2.28	0.67
1:CA:1458:G:H5'	20:CT:27:MET:HB3	1.77	0.67
1:AA:858:G:N7	57:AA:1824:HOH:O	2.28	0.67
24:BC:15:HIS:O	24:BC:204:VAL:HG21	1.94	0.67
22:DA:14:A:C6	22:DA:526:A:C2	2.83	0.67
1:AA:572:A:H5'	1:AA:573:A:OP2	1.95	0.67
22:DA:1258:U:H2'	22:DA:1259:G:C8	2.30	0.67
22:DA:2016:U:O2	48:D0:4:GLN:NE2	2.27	0.67
22:DA:187:G:C2	22:DA:210:C:C2	2.82	0.67
24:DC:62:TYR:CE2	24:DC:63:ARG:O	2.48	0.67
22:DA:310:A:O2'	22:DA:311:A:OP2	2.09	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:DP:89:ARG:NH1	37:DP:115:ASN:OXT	2.27	0.67
45:BX:17:ASN:OD1	45:BX:27:ARG:HD2	1.94	0.67
32:DK:34:GLY:O	32:DK:36:GLY:N	2.27	0.67
22:DA:945:A:C8	22:DA:2448:A:C2	2.83	0.67
4:AD:130:VAL:HG11	4:AD:135:TYR:CD2	2.29	0.67
22:DA:2162:G:H4'	22:DA:2163:A:OP1	1.94	0.67
22:DA:2392:A:C8	22:DA:2429:G:C2	2.83	0.67
1:AA:872:A:C4	1:AA:874:G:N7	2.63	0.67
22:DA:334:C:OP1	22:DA:335:C:N4	2.26	0.66
28:BG:121:ILE:HD12	28:BG:141:ILE:HG22	1.77	0.66
47:DZ:6:LYS:HB2	47:DZ:58:GLU:HG3	1.75	0.66
22:BA:2192:U:N3	22:BA:2193:G:N7	2.44	0.66
17:AQ:48:ASP:OD2	17:AQ:52:GLU:OE1	2.13	0.66
22:DA:2093:G:C5	22:DA:2225:A:N7	2.63	0.66
22:DA:1654:A:P	35:DN:1:MET:HA	2.35	0.66
29:BH:27:ARG:O	29:BH:28:ASN:HB2	1.95	0.66
5:AE:25:VAL:O	5:AE:27:GLY:N	2.28	0.66
22:BA:1746:A:H2'	22:BA:1747:U:H6	1.59	0.66
22:BA:831:G:OP1	57:BA:3265:HOH:O	2.14	0.66
24:DC:45:ASN:OD1	24:DC:46:ASN:N	2.28	0.66
29:BH:94:ILE:CG2	29:BH:99:ILE:HG13	2.26	0.66
22:DA:2055:C:N3	57:DA:3532:HOH:O	2.28	0.66
25:DD:13:ARG:HD2	25:DD:15:PHE:CE2	2.30	0.66
40:DS:58:ALA:O	40:DS:64:ALA:N	2.29	0.66
22:BA:1385:A:H1'	22:BA:1386:C:C6	2.30	0.66
4:CD:168:PRO:HB2	4:CD:171:LEU:CD1	2.26	0.66
51:B3:27:ALA:O	51:B3:28:ASN:HB2	1.93	0.66
22:DA:1649:G:O6	22:DA:2009:A:N6	2.29	0.66
22:DA:696:G:C2	22:DA:767:U:O2	2.47	0.66
22:DA:450:G:N1	22:DA:454:A:OP2	2.27	0.66
1:AA:532:A:N6	3:AC:192:THR:OG1	2.28	0.66
22:BA:357:C:H2'	22:BA:358:U:C6	2.31	0.66
42:BU:12:ILE:HG21	42:BU:80:ALA:HB2	1.78	0.66
29:BH:121:VAL:O	29:BH:128:HIS:NE2	2.29	0.66
1:CA:374:A:H5''	1:CA:452:A:N1	2.10	0.66
1:CA:577:G:N3	1:CA:578:C:C6	2.64	0.66
2:CB:72:THR:HG22	2:CB:95:ARG:NH1	2.11	0.66
44:BW:19:LYS:O	44:BW:21:LEU:HG	1.94	0.66
2:AB:88:ASP:C	2:AB:89:GLN:HG3	2.16	0.66
1:CA:412:A:O2'	1:CA:413:G:H4'	1.95	0.66
22:DA:1380:G:OP2	57:DA:3751:HOH:O	2.14	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:2134:A:N6	22:DA:2157:G:O2'	2.28	0.66
45:BX:13:VAL:HG22	45:BX:29:PHE:HB2	1.78	0.66
1:CA:716:A:N3	11:CK:119:ASN:O	2.29	0.66
22:BA:798:G:O6	57:BA:3326:HOH:O	2.12	0.66
1:AA:16:A:O2'	1:AA:17:U:H5'	1.95	0.66
39:BR:66:HIS:ND1	39:BR:94:THR:HG22	2.11	0.66
22:DA:189:G:C4	22:DA:205:G:N2	2.63	0.66
22:BA:2190:G:C6	22:BA:2191:A:C6	2.83	0.66
25:BD:13:ARG:HD2	25:BD:15:PHE:CZ	2.31	0.66
22:DA:2520:C:HO2'	22:DA:2565:A:HO2'	1.39	0.66
1:AA:382:A:H2'	1:AA:383:A:C8	2.31	0.66
1:AA:721:G:H4'	1:AA:722:G:O4'	1.95	0.66
29:BH:86:ASP:HB2	1:CA:359:G:O2'	1.96	0.66
1:CA:728:A:H2'	1:CA:729:A:C8	2.31	0.66
22:DA:574:A:H4'	22:DA:575:A:C5'	2.25	0.66
16:AP:50:THR:O	16:AP:50:THR:HG22	1.96	0.66
22:BA:1915:U:C2	22:BA:1916:A:C8	2.84	0.65
1:CA:1071:C:O2	1:CA:1072:G:C8	2.49	0.65
22:BA:1779:U:C5	22:BA:1784:A:N7	2.62	0.65
5:CE:80:THR:OG1	5:CE:122:ASN:ND2	2.29	0.65
1:CA:31:G:O4'	1:CA:306:A:C2	2.49	0.65
50:D2:11:LYS:NZ	57:D2:201:HOH:O	2.29	0.65
14:AN:61:ARG:NH1	57:AN:202:HOH:O	2.25	0.65
22:DA:2131:U:O2	22:DA:2158:A:N6	2.29	0.65
38:BQ:79:PHE:CZ	38:BQ:83:LEU:HD11	2.31	0.65
22:DA:1530:G:N2	22:DA:1542:U:O2	2.28	0.65
22:DA:1181:U:H2'	22:DA:1182:G:C8	2.31	0.65
1:CA:1524:C:OP2	11:CK:125:LYS:HD2	1.95	0.65
22:BA:11:C:H2'	22:BA:12:U:H5'	1.79	0.65
22:BA:555:G:O2'	22:BA:556:A:OP2	2.14	0.65
22:DA:2004:G:OP2	57:DA:3801:HOH:O	2.13	0.65
24:DC:67:PHE:HB3	24:DC:151:GLY:O	1.96	0.65
29:BH:14:SER:O	29:BH:15:LEU:HB2	1.95	0.65
25:BD:149:ASN:OD1	25:BD:150:GLN:N	2.29	0.65
5:AE:114:VAL:HG21	5:AE:141:ILE:HD12	1.78	0.65
22:DA:2164:C:H2'	22:DA:2165:C:C6	2.32	0.65
22:DA:2074:U:H2'	22:DA:2075:U:C6	2.32	0.65
17:CQ:14:SER:O	17:CQ:17:MET:HE1	1.96	0.65
1:CA:821:G:H2'	1:CA:822:U:C6	2.31	0.65
29:BH:122:LEU:HD21	1:CA:368:U:P	2.33	0.65
22:DA:1317:G:H2'	22:DA:1318:U:O4'	1.97	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:2637:U:H2'	22:BA:2638:G:H5'	1.79	0.65
38:BQ:92:ARG:HA	38:BQ:95:LEU:HB2	1.79	0.65
22:DA:1973:G:C6	22:DA:1974:C:C4	2.85	0.65
22:DA:2429:G:OP2	22:DA:2430:A:OP2	2.15	0.65
22:DA:613:A:OP2	22:DA:614:A:N7	2.29	0.65
1:AA:71:A:O2'	1:AA:72:A:OP2	2.15	0.65
12:CL:16:VAL:O	12:CL:17:ALA:O	2.14	0.65
22:BA:933:A:H5'	22:BA:934:U:OP2	1.97	0.65
22:BA:1410:G:O6	57:BA:3628:HOH:O	2.10	0.65
1:AA:1180:A:OP2	9:AI:99:ARG:NH2	2.30	0.65
22:BA:2128:G:H2'	22:BA:2129:C:O4'	1.96	0.65
25:BD:12:THR:HG23	37:BP:9:GLU:OE2	1.96	0.65
1:AA:1054:C:C5	1:AA:1196:A:H2'	2.31	0.65
29:BH:139:PHE:O	29:BH:140:ALA:CB	2.44	0.65
22:DA:1359:A:C8	22:DA:1373:A:C2	2.84	0.65
5:CE:99:ALA:O	5:CE:101:GLU:N	2.29	0.65
29:BH:114:GLU:HB3	29:BH:133:GLN:O	1.97	0.65
22:DA:1843:C:H4'	24:DC:251:GLN:HG2	1.78	0.65
1:AA:8:A:C6	4:AD:206:LYS:HB3	2.31	0.65
22:DA:105:C:H2'	22:DA:106:C:C6	2.32	0.65
23:DB:4:C:C2	23:DB:117:G:N2	2.64	0.65
22:DA:118:A:C8	22:DA:119:A:C8	2.85	0.65
22:DA:1627:G:C2	22:DA:1628:G:N7	2.65	0.65
3:AC:36:ASP:OD1	3:AC:59:ARG:NH1	2.29	0.65
1:AA:484:G:H4'	1:AA:485:U:OP1	1.97	0.65
6:AF:14:GLN:OE1	6:AF:17:GLN:HB2	1.97	0.65
22:BA:2286:G:H4'	22:BA:2287:A:O5'	1.97	0.65
22:BA:761:A:OP1	57:BA:3701:HOH:O	2.14	0.65
1:CA:675:A:C4	1:CA:676:A:C8	2.85	0.65
22:DA:1638:C:O2	22:DA:2698:U:O2'	2.10	0.65
32:BK:107:LEU:O	32:BK:109:SER:N	2.30	0.65
1:AA:1255:G:O2'	1:AA:1258:G:N3	2.30	0.65
1:AA:933:G:OP2	7:AG:3:ARG:HB3	1.96	0.65
29:DH:149:GLU:O	29:DH:149:GLU:HG2	1.96	0.65
42:DU:77:THR:O	42:DU:79:LYS:N	2.30	0.65
17:CQ:19:LYS:NZ	17:CQ:49:GLU:OE1	2.17	0.65
5:AE:72:ILE:HD13	5:AE:145:GLU:CD	2.17	0.65
1:AA:1299:A:H2'	1:AA:1299:A:N3	2.11	0.65
42:BU:72:ILE:N	42:BU:72:ILE:HD13	2.11	0.65
22:BA:1224:U:H4'	39:BR:88:GLY:O	1.96	0.65
1:CA:1410:A:H2'	1:CA:1411:C:C6	2.32	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:771:G:C2'	1:AA:772:U:H5'	2.26	0.64
22:DA:1364:G:H1'	22:DA:1368:G:N2	2.12	0.64
22:DA:2757:A:N1	28:DG:67:THR:HG21	2.12	0.64
22:BA:1789:A:OP2	24:BC:221:ARG:NH1	2.30	0.64
22:BA:747:U:C5	22:BA:2613:U:C5	2.85	0.64
4:AD:125:VAL:O	4:AD:127:GLY:N	2.27	0.64
22:DA:1277:G:H5'	35:DN:20:MET:CE	2.27	0.64
22:BA:981:A:OP1	57:BA:3600:HOH:O	2.14	0.64
1:CA:1499:A:H3'	57:CA:1877:HOH:O	1.97	0.64
24:BC:231:PRO:HD2	24:BC:247:PRO:HA	1.79	0.64
26:DE:52:VAL:HG21	26:DE:81:GLY:HA2	1.79	0.64
7:AG:75:VAL:HG11	7:AG:144:MET:HG3	1.80	0.64
1:CA:642:A:C5	8:CH:107:SER:HA	2.32	0.64
22:BA:686:U:H2'	22:BA:788:A:C2	2.33	0.64
22:DA:1780:A:OP1	57:DA:3690:HOH:O	2.15	0.64
1:AA:188:C:N3	1:AA:189:A:C2	2.65	0.64
2:CB:104:TRP:CH2	2:CB:155:GLY:C	2.70	0.64
1:AA:1055:A:C6	1:AA:1206:G:C5	2.86	0.64
22:BA:96:C:H4'	46:BY:41:HIS:CD2	2.32	0.64
22:BA:2066:C:OP1	57:BA:3514:HOH:O	2.15	0.64
2:CB:73:LYS:O	2:CB:75:ALA:N	2.31	0.64
32:DK:105:ARG:NH2	37:DP:34:GLU:OE2	2.29	0.64
1:AA:262:A:H2'	1:AA:263:A:C8	2.32	0.64
33:BL:100:ILE:HG13	33:BL:101:ILE:HG23	1.80	0.64
1:CA:547:A:P	57:CA:1774:HOH:O	2.56	0.64
22:DA:1269:A:N7	57:DA:3383:HOH:O	2.30	0.64
22:DA:813:U:H2'	22:DA:814:C:C6	2.33	0.64
12:CL:28:PRO:HB2	12:CL:29:GLN:OE1	1.96	0.64
1:AA:111:G:H5"	1:AA:112:G:OP2	1.97	0.64
22:DA:1649:G:C6	22:DA:2009:A:N6	2.65	0.64
39:BR:47:VAL:HG11	39:BR:54:VAL:HG13	1.79	0.64
23:BB:47:C:P	36:BO:3:LYS:HD2	2.36	0.64
49:B1:23:THR:OG1	49:B1:24:THR:N	2.31	0.64
33:DL:93:ASN:O	33:DL:95:LEU:N	2.30	0.64
22:DA:1441:G:N2	22:DA:1442:U:C2	2.66	0.64
15:CO:87:LEU:O	15:CO:88:ARG:HB3	1.96	0.64
33:DL:61:LEU:O	51:D3:13:ARG:HD3	1.97	0.64
1:AA:202:G:O2'	1:AA:468:A:C8	2.48	0.64
2:AB:50:PHE:HA	2:AB:213:TYR:OH	1.98	0.64
22:DA:1824:G:N7	57:DA:3656:HOH:O	2.30	0.64
22:DA:388:G:N7	22:DA:390:U:H2'	2.13	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:DT:61:LEU:HD12	41:DT:62:VAL:N	2.13	0.64
22:BA:1509:A:HO2'	22:BA:1510:G:P	2.20	0.64
22:DA:686:U:OP2	57:DA:3717:HOH:O	2.15	0.64
30:BI:16:GLY:HA2	30:BI:51:LYS:HB3	1.79	0.64
31:DJ:4:PHE:O	38:DQ:64:ARG:NH2	2.26	0.64
22:BA:1921:G:C2	22:BA:1922:G:C8	2.86	0.64
28:DG:170:ARG:NH1	52:D4:29:ALA:O	2.30	0.64
22:BA:2315:G:H2'	22:BA:2316:G:H8	1.63	0.64
22:BA:930:G:H1'	47:BZ:25:LEU:HD21	1.80	0.64
22:BA:2554:U:C4	22:BA:2555:U:O4	2.51	0.64
22:BA:2187:U:C5	22:BA:2188:U:C4	2.86	0.64
4:CD:130:VAL:HG11	4:CD:135:TYR:CD2	2.33	0.64
36:DO:2:ASP:O	36:DO:6:ALA:HB2	1.97	0.64
22:BA:197:A:N6	22:BA:2430:A:H2'	2.13	0.64
22:BA:45:G:H5''	22:BA:46:G:OP1	1.97	0.64
22:DA:2286:G:H4'	22:DA:2287:A:O5'	1.98	0.64
1:CA:22:G:O2'	1:CA:913:A:N1	2.29	0.64
32:DK:76:VAL:HG12	37:DP:73:VAL:HG22	1.78	0.64
8:AH:18:GLN:O	8:AH:21:ASN:N	2.30	0.64
33:BL:68:SER:O	33:BL:69:ARG:HB2	1.97	0.64
22:BA:1925:C:H4'	22:BA:1926:U:C4	2.33	0.63
17:AQ:16:LYS:N	17:AQ:17:MET:HE1	2.13	0.63
1:CA:1296:C:H4'	1:CA:1302:C:N4	2.13	0.63
1:AA:1358:U:C5	1:AA:1359:C:C4	2.86	0.63
1:CA:576:C:H3'	1:CA:577:G:H5''	1.79	0.63
24:DC:212:ARG:NE	24:DC:216:VAL:O	2.31	0.63
33:DL:80:SER:O	33:DL:84:LYS:NZ	2.31	0.63
1:CA:966:G:O2'	9:CI:130:ARG:OXT	2.16	0.63
22:DA:2330:G:N2	22:DA:2386:A:C2	2.66	0.63
29:DH:117:LEU:CD1	29:DH:130:VAL:HG22	2.28	0.63
4:CD:26:ARG:O	4:CD:27:ALA:HB2	1.98	0.63
6:AF:3:HIS:O	6:AF:92:THR:HA	1.98	0.63
28:DG:24:ILE:HD11	28:DG:43:VAL:HG11	1.81	0.63
2:AB:67:ILE:HG21	2:AB:69:PHE:CE1	2.33	0.63
22:BA:591:U:HO2'	51:B3:2:PRO:N	1.95	0.63
22:DA:781:A:H2'	22:DA:1777:U:O2'	1.98	0.63
22:DA:1141:U:H4'	22:DA:1142:A:O4'	1.98	0.63
22:DA:1973:G:C5	22:DA:1974:C:C4	2.87	0.63
1:CA:322:C:O2	1:CA:332:G:N2	2.30	0.63
1:CA:577:G:C8	1:CA:816:A:C2	2.87	0.63
22:DA:1469:A:H2'	22:DA:1470:A:C8	2.33	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CB:86:SER:O	2:CB:87:CYS:O	2.17	0.63
22:DA:235:U:C4	22:DA:236:C:C5	2.86	0.63
22:BA:319:G:C4	22:BA:333:G:N2	2.66	0.63
22:BA:2516:A:O2'	22:BA:2517:C:H5'	1.98	0.63
1:AA:682:G:N2	1:AA:709:U:C2	2.67	0.63
1:CA:134:G:H2'	1:CA:135:C:O4'	1.99	0.63
22:DA:1826:G:O2'	22:DA:1971:U:OP2	2.17	0.63
45:DX:28:ARG:CZ	45:DX:30:LEU:HD21	2.29	0.63
22:BA:475:C:C4	22:BA:481:G:O6	2.51	0.63
22:DA:684:G:OP1	50:D2:16:HIS:ND1	2.31	0.63
22:DA:684:G:N2	22:DA:788:A:OP2	2.30	0.63
22:DA:749:A:C6	22:DA:750:A:N7	2.66	0.63
24:DC:67:PHE:CE1	24:DC:156:ARG:CZ	2.81	0.63
11:CK:23:ILE:HG21	11:CK:96:THR:HG21	1.79	0.63
36:BO:51:ALA:HB3	36:BO:78:VAL:HG13	1.79	0.63
5:CE:77:ASN:HB2	5:CE:82:GLN:HG2	1.80	0.63
22:DA:2189:U:H2'	22:DA:2190:G:H5''	1.80	0.63
1:AA:1121:U:N3	1:AA:1122:U:C5	2.67	0.63
22:BA:324:A:N6	22:BA:338:G:O2'	2.30	0.63
4:CD:107:PHE:CG	4:CD:145:ILE:HD11	2.33	0.63
22:DA:2707:U:H2'	22:DA:2708:G:C8	2.34	0.63
22:DA:2125:G:N1	22:DA:2171:A:OP1	2.32	0.63
46:BY:22:LEU:O	46:BY:23:ARG:O	2.16	0.63
22:DA:1364:G:N2	22:DA:1367:A:OP2	2.29	0.63
29:DH:117:LEU:HG	29:DH:120:GLY:O	1.98	0.63
11:AK:29:ASN:OD1	11:AK:30:THR:N	2.31	0.63
1:CA:623:C:C4	1:CA:624:C:C5	2.86	0.63
40:BS:43:ALA:HA	40:BS:46:LEU:HD12	1.80	0.63
7:AG:146:GLU:HA	7:AG:149:LYS:HB2	1.80	0.63
50:B2:43:THR:O	50:B2:44:VAL:HG12	1.98	0.63
39:BR:49:ILE:CG2	39:BR:53:PHE:H	2.11	0.63
17:AQ:69:LYS:O	17:AQ:70:THR:CB	2.46	0.63
22:BA:2296:U:H4'	22:BA:2297:A:OP1	1.99	0.63
22:DA:1847:A:O2'	22:DA:1848:A:H8	1.80	0.63
22:DA:106:C:O2'	22:DA:294:A:O2'	2.07	0.63
38:DQ:25:TYR:CD2	38:DQ:26:GLY:N	2.66	0.63
1:CA:568:G:O6	12:CL:2:ALA:HB2	1.98	0.63
23:DB:23:G:O6	57:DB:304:HOH:O	2.13	0.63
30:BI:131:GLY:O	30:BI:134:ARG:N	2.31	0.63
22:BA:2062:A:OP1	57:BA:3502:HOH:O	2.15	0.63
22:DA:1835:G:N3	22:DA:1836:C:C6	2.67	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:AF:47:LEU:HD13	6:AF:51:ILE:HG23	1.79	0.63
22:DA:749:A:C4	22:DA:750:A:C8	2.87	0.63
37:DP:29:LYS:HB3	37:DP:40:LEU:HD21	1.80	0.63
1:CA:773:G:C2	1:CA:807:A:C2	2.86	0.63
1:CA:131:A:O2'	1:CA:262:A:N3	2.31	0.63
1:AA:1217:C:OP1	14:AN:9:ARG:NE	2.32	0.63
15:AO:19:ALA:O	15:AO:20:ASN:HB2	1.99	0.63
22:DA:1825:U:O4	57:DA:3783:HOH:O	2.14	0.63
2:CB:210:VAL:O	2:CB:214:LEU:HB2	1.98	0.63
13:AM:80:LEU:HD22	13:AM:87:ARG:HB2	1.81	0.63
1:AA:1269:A:N1	1:AA:1312:G:O2'	2.19	0.63
22:BA:1915:U:C2'	22:BA:1916:A:H5'	2.29	0.63
22:DA:1395:A:O2'	22:DA:1397:U:C6	2.52	0.63
22:DA:1798:U:O2'	22:DA:1802:A:N3	2.32	0.63
42:BU:72:ILE:H	42:BU:72:ILE:HD13	1.64	0.63
48:B0:15:MET:O	48:B0:18:SER:HB3	1.98	0.63
1:CA:1077:G:N2	1:CA:1080:A:OP2	2.31	0.63
31:BJ:42:ALA:O	38:BQ:64:ARG:HG2	1.98	0.63
22:DA:1432:G:H2'	22:DA:1433:A:C8	2.33	0.63
25:DD:140:HIS:NE2	57:DD:302:HOH:O	2.14	0.63
13:AM:46:SER:O	13:AM:47:GLU:HB3	1.99	0.63
22:BA:2887:A:H5'	22:BA:2888:C:OP2	1.99	0.63
1:CA:1417:G:N2	1:CA:1484:C:C4	2.67	0.63
22:DA:1833:C:C2	22:DA:1834:U:C6	2.86	0.63
1:CA:784:A:C2	1:CA:785:G:C4	2.86	0.63
35:BN:117:ASP:O	35:BN:119:SER:N	2.32	0.63
22:BA:1916:A:H2'	22:BA:1917:U:H1'	1.79	0.63
17:AQ:12:VAL:O	17:AQ:22:VAL:O	2.16	0.63
38:BQ:112:LYS:HZ1	39:BR:50:GLY:HA2	1.64	0.63
22:DA:1649:G:C6	22:DA:2009:A:C6	2.87	0.63
22:DA:2349:G:OP1	51:D3:45:ARG:NH2	2.31	0.63
1:AA:655:A:C2	1:AA:656:G:C4	2.87	0.63
22:BA:2025:C:OP2	57:BA:3479:HOH:O	2.16	0.63
22:DA:587:C:OP2	33:DL:21:ARG:NH1	2.32	0.63
35:DN:55:ALA:HA	35:DN:80:PHE:CE1	2.34	0.63
1:CA:223:A:C5	1:CA:224:U:C5	2.87	0.63
24:BC:232:HIS:O	24:BC:233:GLY:O	2.17	0.63
17:AQ:14:SER:OG	17:AQ:17:MET:CE	2.47	0.62
4:AD:192:SER:OG	4:AD:193:ALA:N	2.28	0.62
1:AA:1077:G:C5	57:AA:1793:HOH:O	2.52	0.62
22:DA:1973:G:N7	22:DA:1974:C:C5	2.66	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:DY:56:LEU:O	46:DY:57:LEU:CB	2.46	0.62
1:CA:1240:U:H5'	1:CA:1241:G:C8	2.34	0.62
22:BA:1071:G:C8	22:BA:1089:A:N6	2.66	0.62
22:BA:1917:U:O4	22:BA:1918:A:C6	2.52	0.62
1:AA:108:G:N3	1:AA:108:G:H5'	2.14	0.62
46:BY:17:GLU:HG3	46:BY:18:LEU:N	2.12	0.62
22:DA:1715:G:N2	22:DA:1744:A:OP2	2.32	0.62
42:BU:39:ILE:HG22	42:BU:40:ASN:N	2.14	0.62
1:AA:1317:C:OP1	14:AN:56:SER:OG	2.12	0.62
28:DG:11:VAL:O	28:DG:48:ASN:ND2	2.32	0.62
1:CA:1296:C:N4	1:CA:1297:G:O6	2.32	0.62
13:CM:14:HIS:HB2	13:CM:17:ILE:HD12	1.81	0.62
1:AA:68:G:C5	1:AA:69:G:H1'	2.34	0.62
22:DA:2214:C:C2	22:DA:2215:C:C6	2.87	0.62
22:DA:686:U:H6	22:DA:788:A:N1	1.96	0.62
22:DA:668:A:C2	22:DA:670:A:C5	2.88	0.62
44:BW:19:LYS:O	44:BW:21:LEU:N	2.32	0.62
22:DA:319:G:H2'	22:DA:320:A:O4'	1.99	0.62
1:AA:114:U:O2'	1:AA:115:G:H5'	1.99	0.62
20:CT:25:ARG:O	20:CT:29:ARG:HG2	1.99	0.62
22:BA:2455:G:O6	57:BA:3534:HOH:O	2.14	0.62
22:DA:308:G:C6	22:DA:309:A:C6	2.88	0.62
1:AA:1407:C:O2'	22:BA:1912:A:N1	2.24	0.62
22:DA:514:A:N1	22:DA:515:A:C2	2.67	0.62
1:AA:597:G:C2	1:AA:644:U:C2	2.87	0.62
33:BL:62:PRO:CG	51:B3:25:LYS:HD3	2.30	0.62
41:BT:61:LEU:C	41:BT:61:LEU:HD12	2.19	0.62
1:CA:1219:A:N6	1:CA:1220:G:O6	2.32	0.62
28:BG:38:ASN:O	28:BG:39:ASP:HB2	1.99	0.62
20:CT:78:ASN:ND2	57:CT:202:HOH:O	2.33	0.62
22:DA:1308:A:N6	22:DA:1309:G:C2	2.68	0.62
2:CB:99:GLY:O	2:CB:103:ASN:ND2	2.32	0.62
22:DA:2054:A:OP1	22:DA:2055:C:O2'	2.18	0.62
22:DA:53:A:N3	22:DA:179:C:H4'	2.14	0.62
22:BA:998:C:C6	57:BA:3364:HOH:O	2.51	0.62
4:CD:32:CYS:O	4:CD:33:LYS:CB	2.47	0.62
1:CA:577:G:N9	1:CA:816:A:C2	2.67	0.62
15:CO:58:ARG:O	15:CO:62:GLN:HB2	2.00	0.62
22:BA:1817:G:OP1	24:BC:87:ARG:NH2	2.30	0.62
7:CG:65:ALA:O	7:CG:127:ALA:HB1	2.00	0.62
24:BC:70:ASN:O	24:BC:72:ASP:N	2.32	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:AC:141:ALA:O	3:AC:146:ALA:HB3	2.00	0.62
25:DD:151:THR:O	25:DD:152:PRO:C	2.38	0.62
41:BT:2:ILE:CD1	41:BT:7:LEU:HD11	2.27	0.62
22:DA:118:A:N3	22:DA:178:G:H1'	2.15	0.62
22:DA:1693:U:O4	22:DA:1976:U:O2'	2.16	0.62
22:BA:528:A:H3'	22:BA:528:A:H8	1.62	0.62
1:CA:33:A:H2'	1:CA:34:C:C6	2.34	0.62
22:DA:2521:C:C2	22:DA:2545:G:N2	2.68	0.62
1:AA:1253:G:H2'	1:AA:1254:A:H8	1.64	0.62
1:CA:782:A:N7	1:CA:783:C:C5	2.68	0.62
22:BA:1156:A:C8	38:BQ:51:ARG:HG2	2.34	0.62
9:AI:57:MET:SD	9:AI:58:VAL:N	2.70	0.62
53:B5:50:ILE:C	53:B5:52:PRO:HD3	2.19	0.62
24:BC:260:ASN:O	24:BC:262:ARG:N	2.30	0.62
22:BA:2314:A:OP1	27:BF:88:LYS:NZ	2.33	0.62
1:AA:995:C:N3	1:AA:1046:A:O2'	2.30	0.62
39:BR:49:ILE:CG2	39:BR:52:PRO:HA	2.29	0.62
22:DA:1440:U:O4	57:DA:3631:HOH:O	2.16	0.62
22:DA:1250:G:C5'	38:DQ:6:ARG:HD2	2.29	0.62
22:DA:248:G:H5'	22:DA:250:G:N7	2.15	0.62
31:DJ:41:LYS:O	31:DJ:43:GLU:N	2.33	0.62
22:BA:276:U:O2	22:BA:276:U:H2'	2.00	0.62
1:AA:64:G:C8	1:AA:99:C:N4	2.68	0.62
11:AK:67:ALA:HB1	11:AK:100:LEU:HD13	1.81	0.62
22:DA:1304:A:C6	22:DA:1305:C:C4	2.87	0.62
22:DA:1360:G:C6	22:DA:1361:G:H1'	2.34	0.62
5:CE:154:ALA:HA	5:CE:157:ARG:HB3	1.82	0.62
1:CA:429:U:H3'	4:CD:9:LEU:HD22	1.80	0.62
4:CD:35:GLU:O	4:CD:38:PRO:HD3	2.00	0.62
14:CN:91:GLY:O	14:CN:93:ILE:N	2.33	0.62
22:BA:1057:A:C2	22:BA:1086:A:C2	2.87	0.62
2:CB:167:ASP:OD2	2:CB:191:SER:HA	1.99	0.62
22:BA:58:G:OP1	41:BT:78:SER:CB	2.48	0.62
22:BA:1926:U:H2'	22:BA:1926:U:O2	1.99	0.62
22:DA:2056:G:C2	22:DA:2057:G:C8	2.88	0.62
25:BD:104:VAL:O	25:BD:105:LYS:HB2	1.99	0.62
22:DA:311:A:C2	22:DA:328:U:O4	2.52	0.62
3:AC:139:GLN:O	3:AC:141:ALA:N	2.33	0.62
1:CA:257:G:C6	57:CA:1718:HOH:O	2.51	0.62
34:DM:136:MET:O	43:DV:79:ARG:NH2	2.32	0.62
22:BA:2334:U:C4	36:BO:16:ARG:HD3	2.35	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:858:G:O6	1:CA:869:G:H3'	2.00	0.62
22:DA:1088:A:N6	30:DI:135:SER:OG	2.33	0.62
29:BH:94:ILE:C	29:BH:122:LEU:HG	2.21	0.62
22:DA:192:C:O2'	22:DA:802:A:N3	2.33	0.62
33:BL:77:ILE:HG23	33:BL:100:ILE:HD11	1.82	0.62
24:BC:107:PRO:HB3	24:BC:142:HIS:CE1	2.34	0.62
22:DA:1240:U:O2'	22:DA:1241:A:O5'	2.17	0.62
1:AA:1212:U:H4'	1:AA:1213:A:C8	2.35	0.62
23:DB:58:A:H2'	23:DB:59:A:O4'	2.00	0.62
5:CE:24:THR:HA	5:CE:29:ARG:HA	1.82	0.62
22:DA:1676:A:H2'	22:DA:1677:A:O4'	2.00	0.62
22:BA:2499:C:P	57:BA:3691:HOH:O	2.48	0.61
22:DA:1309:G:H4'	50:D2:7:PRO:HB2	1.82	0.61
22:DA:1345:C:H5'	22:DA:1396:U:O4	2.00	0.61
1:AA:976:G:OP2	1:AA:1358:U:O2'	2.18	0.61
5:CE:36:LEU:HD21	5:CE:137:VAL:HG11	1.82	0.61
22:BA:528:A:C8	22:BA:528:A:C3'	2.82	0.61
44:BW:37:ILE:HG21	44:BW:80:ILE:HG21	1.82	0.61
22:BA:1376:C:OP2	57:BA:3406:HOH:O	2.16	0.61
1:CA:1452:C:H5'	1:CA:1453:G:C4	2.35	0.61
1:CA:920:U:H2'	1:CA:921:U:C6	2.34	0.61
22:BA:2800:A:C2	22:BA:2895:G:H1'	2.34	0.61
22:BA:1588:G:C2	22:BA:1589:U:C6	2.88	0.61
22:BA:1589:U:C2	22:BA:1590:A:C8	2.88	0.61
22:BA:2707:U:O2	35:BN:71:ARG:NH1	2.33	0.61
1:CA:582:C:O2	1:CA:760:G:N2	2.33	0.61
29:DH:32:PRO:O	29:DH:33:GLN:CB	2.48	0.61
11:AK:88:GLY:H	11:AK:114:THR:HG22	1.64	0.61
22:DA:2300:C:C2	22:DA:2317:A:C2	2.88	0.61
22:BA:636:G:C6	33:BL:111:ILE:HD11	2.35	0.61
2:CB:24:ASN:O	2:CB:26:LYS:N	2.32	0.61
1:CA:386:C:N4	1:CA:387:U:O4	2.33	0.61
22:BA:580:U:H2'	22:BA:581:C:C6	2.33	0.61
13:AM:29:ARG:NH2	13:AM:63:PHE:HB2	2.15	0.61
23:DB:7:G:O2'	36:DO:38:GLN:OE1	2.17	0.61
5:AE:115:LEU:HG	5:AE:120:VAL:HG21	1.81	0.61
1:AA:518:C:H2'	1:AA:530:G:C8	2.35	0.61
22:BA:1910:G:H2'	22:BA:1911:U:O4'	2.00	0.61
1:CA:373:A:N3	1:CA:374:A:C8	2.68	0.61
1:CA:1518:A:C2	1:CA:1519:A:C2	2.88	0.61
22:DA:828:U:O2'	22:DA:829:A:O5'	2.19	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1417:G:N2	1:AA:1482:G:H2'	2.16	0.61
1:AA:408:A:C2	1:AA:435:A:C2	2.88	0.61
11:CK:112:ASP:CB	21:CU:20:LYS:HE3	2.30	0.61
1:AA:955:U:O4'	1:AA:1227:A:N6	2.33	0.61
6:AF:16:GLU:OE2	4:CD:188:ARG:CZ	2.47	0.61
22:BA:2683:C:O2	32:BK:70:ARG:NH2	2.33	0.61
12:CL:24:LEU:O	12:CL:26:ALA:N	2.34	0.61
22:DA:830:G:C4	22:DA:2448:A:C5	2.88	0.61
22:BA:830:G:H4'	22:BA:831:G:OP2	1.99	0.61
22:DA:2300:C:N3	22:DA:2317:A:C2	2.68	0.61
26:DE:15:SER:OG	26:DE:197:GLU:OE2	2.17	0.61
22:BA:861:A:C2	22:BA:917:A:C4	2.89	0.61
12:AL:114:ARG:O	12:AL:116:LYS:O	2.19	0.61
53:B5:40:GLU:HA	53:B5:181:PHE:HA	1.82	0.61
6:AF:68:GLN:HA	6:AF:71:ILE:CG2	2.29	0.61
16:CP:6:LEU:CD1	16:CP:71:VAL:HG23	2.31	0.61
22:DA:27:G:N2	22:DA:512:G:H1'	2.15	0.61
1:AA:872:A:C5	1:AA:874:G:C8	2.89	0.61
39:BR:47:VAL:CG1	39:BR:54:VAL:HG13	2.30	0.61
22:BA:580:U:H2'	22:BA:581:C:H6	1.66	0.61
1:CA:72:A:C6	1:CA:73:C:N4	2.68	0.61
13:CM:10:PRO:O	13:CM:11:ASP:CB	2.49	0.61
22:DA:2204:G:C2	22:DA:2205:A:C8	2.88	0.61
35:DN:87:PHE:O	35:DN:89:SER:N	2.34	0.61
1:AA:1059:C:H2'	1:AA:1060:U:H6	1.64	0.61
22:BA:1584:U:H2'	22:BA:1584:U:O2	1.99	0.61
1:CA:790:A:C6	1:CA:791:G:C6	2.88	0.61
39:BR:66:HIS:CE1	39:BR:94:THR:CG2	2.83	0.61
4:AD:32:CYS:O	4:AD:33:LYS:HB2	2.00	0.61
46:BY:22:LEU:O	46:BY:23:ARG:C	2.38	0.61
1:CA:765:G:C6	1:CA:812:G:C4	2.89	0.61
9:AI:35:LEU:HD11	9:AI:48:VAL:HG21	1.82	0.61
15:CO:16:GLY:O	15:CO:18:ASP:N	2.33	0.61
22:BA:281:C:H2'	22:BA:282:A:C8	2.35	0.61
8:AH:10:MET:HE2	8:AH:33:LYS:HG2	1.82	0.61
10:AJ:11:LYS:HG3	10:AJ:97:ASP:HB3	1.81	0.61
11:AK:126:LYS:O	21:AU:34:ARG:CZ	2.49	0.61
1:CA:938:A:O3'	7:CG:95:ARG:NH2	2.34	0.61
22:DA:1831:G:H2'	22:DA:1832:C:C6	2.36	0.61
22:BA:1588:G:N3	22:BA:1589:U:C6	2.68	0.61
23:DB:81:G:C5	23:DB:82:U:C5	2.88	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:BI:39:CYS:HA	30:BI:42:PHE:HB3	1.83	0.61
44:BW:10:THR:O	44:BW:11:ARG:HB2	2.00	0.61
1:CA:455:G:N2	1:CA:478:A:C2	2.68	0.61
1:CA:562:U:H1'	12:CL:12:ARG:CG	2.30	0.61
2:AB:10:LEU:HD23	2:AB:10:LEU:C	2.21	0.61
1:AA:769:G:H4'	1:AA:1513:A:H4'	1.82	0.61
1:CA:33:A:H2'	1:CA:34:C:H6	1.65	0.61
22:DA:2291:U:H2'	22:DA:2292:U:C6	2.35	0.61
1:CA:1202:U:H2'	1:CA:1203:C:O4'	2.00	0.61
22:DA:2339:C:H2'	22:DA:2340:A:C8	2.36	0.61
5:AE:137:VAL:O	5:AE:138:ARG:HB2	2.01	0.61
1:AA:328:C:C2'	1:AA:328:C:O2	2.49	0.61
1:CA:369:G:OP2	1:CA:388:G:N1	2.30	0.61
22:DA:659:G:H2'	22:DA:660:C:C6	2.36	0.61
53:B5:52:PRO:O	53:B5:53:ARG:HB2	2.01	0.61
27:BF:146:VAL:O	27:BF:147:ASP:C	2.37	0.61
1:CA:1191:A:H5''	3:CC:4:LYS:HE3	1.82	0.61
22:DA:2573:C:OP1	22:DA:2574:G:H5''	2.01	0.61
1:CA:1386:G:O2'	1:CA:1387:G:H5'	2.00	0.61
40:BS:84:ARG:HB2	40:BS:96:ILE:CG1	2.31	0.61
1:AA:1504:G:O3'	57:AA:1871:HOH:O	2.16	0.61
22:DA:2502:G:H5'	22:DA:2503:A:O5'	2.01	0.61
22:DA:56:A:C2	22:DA:57:C:C2	2.89	0.61
22:BA:272:A:H2'	22:BA:273:G:O4'	2.00	0.61
22:DA:1790:C:O2'	24:DC:208:ALA:HB2	2.00	0.61
18:CR:35:GLU:HB2	21:CU:19:PHE:CZ	2.36	0.61
1:CA:552:U:C2	1:CA:553:A:C8	2.89	0.61
1:AA:1151:A:O2'	1:AA:1152:A:O5'	2.18	0.61
22:BA:846:U:O2'	22:BA:847:U:P	2.59	0.61
1:AA:457:G:C6	1:AA:458:U:N3	2.69	0.61
22:BA:2448:A:OP2	57:BA:3690:HOH:O	2.15	0.60
41:BT:2:ILE:O	41:BT:3:ARG:HB2	1.99	0.60
45:DX:30:LEU:HB3	45:DX:31:PRO:CD	2.31	0.60
22:DA:2091:C:H1'	45:DX:34:HIS:NE2	2.16	0.60
22:DA:1109:C:C4	22:DA:1110:G:C6	2.89	0.60
4:CD:202:GLU:OE1	5:CE:105:ILE:HG23	2.01	0.60
22:DA:2093:G:O2'	22:DA:2094:A:H5'	2.01	0.60
12:CL:39:THR:O	12:CL:40:THR:OG1	2.13	0.60
22:DA:306:U:O2	22:DA:312:G:N2	2.34	0.60
22:DA:2156:G:C6	22:DA:2157:G:C2	2.89	0.60
22:DA:1638:C:H5''	22:DA:2710:C:O2'	2.01	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:DH:81:ALA:O	29:DH:149:GLU:OE1	2.18	0.60
22:DA:1515:A:HO2'	22:DA:1556:C:HO2'	1.38	0.60
1:AA:448:A:C4	1:AA:487:A:C2	2.89	0.60
1:AA:438:U:C2	1:AA:494:G:C6	2.89	0.60
22:BA:790:U:O2'	22:BA:791:C:OP2	2.17	0.60
29:BH:100:ALA:HB1	29:BH:112:LYS:HA	1.83	0.60
22:BA:1078:U:H1'	22:BA:1088:A:N1	2.16	0.60
22:DA:2262:U:H1'	22:DA:2328:A:H1'	1.83	0.60
35:BN:12:ARG:NE	35:BN:20:MET:CE	2.63	0.60
22:DA:120:U:O4	22:DA:177:G:C8	2.54	0.60
1:CA:375:U:OP1	16:CP:70:ARG:HD3	2.00	0.60
22:DA:1805:A:H1'	24:DC:50:THR:O	2.01	0.60
1:AA:1319:A:C8	1:AA:1323:G:C5	2.89	0.60
2:AB:47:VAL:HB	2:AB:48:PRO:HD3	1.82	0.60
30:BI:127:ARG:HA	30:BI:130:GLU:HG3	1.83	0.60
13:AM:29:ARG:O	13:AM:33:ILE:HG12	2.01	0.60
22:DA:142:A:C6	22:DA:143:C:N4	2.69	0.60
22:BA:1138:G:O2'	31:BJ:107:GLY:HA3	2.00	0.60
22:BA:1853:A:N1	22:BA:2087:G:H1'	2.16	0.60
2:AB:91:PHE:CD1	2:AB:150:GLY:HA3	2.36	0.60
52:D4:20:ASP:OD1	57:D4:201:HOH:O	2.15	0.60
4:CD:4:TYR:O	4:CD:5:LEU:HB2	2.01	0.60
17:AQ:17:MET:HG2	17:AQ:20:SER:CB	2.31	0.60
22:DA:2053:G:H2'	22:DA:2054:A:O4'	2.02	0.60
22:DA:2128:G:N3	22:DA:2173:A:O2'	2.35	0.60
22:DA:1767:G:N1	22:DA:1986:C:C4	2.69	0.60
22:DA:1773:A:N7	22:DA:1829:A:H1'	2.16	0.60
4:AD:22:LYS:O	4:AD:24:GLY:N	2.34	0.60
1:CA:577:G:C2	1:CA:578:C:C5	2.89	0.60
22:DA:443:A:N6	26:DE:36:ALA:O	2.34	0.60
22:DA:669:G:N2	22:DA:670:A:C2	2.69	0.60
9:AI:46:MET:N	9:AI:46:MET:SD	2.66	0.60
25:DD:104:VAL:O	25:DD:105:LYS:HB3	1.99	0.60
1:AA:1461:G:C5	1:AA:1462:C:C5	2.89	0.60
22:BA:1738:G:O2'	22:BA:1739:A:O5'	2.18	0.60
1:AA:1338:G:H2'	1:AA:1339:A:C8	2.36	0.60
22:DA:1905:C:O4'	22:DA:1928:A:C2	2.54	0.60
32:BK:70:ARG:HD3	32:BK:76:VAL:HG22	1.83	0.60
13:AM:29:ARG:CZ	13:AM:63:PHE:HB2	2.31	0.60
22:BA:1993:U:H4'	25:BD:133:THR:HG21	1.82	0.60
36:BO:55:GLU:OE1	36:BO:81:ARG:NH1	2.32	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:2589:A:C8	57:BA:3551:HOH:O	2.54	0.60
1:AA:1101:A:H4'	1:AA:1102:A:O5'	2.02	0.60
22:BA:194:G:N7	57:BA:3765:HOH:O	2.31	0.60
48:B0:34:SER:OG	48:B0:36:GLU:HG2	2.00	0.60
22:BA:1559:U:H4'	22:BA:1560:G:OP2	2.01	0.60
48:D0:25:VAL:O	48:D0:26:THR:OG1	2.15	0.60
29:DH:126:GLY:O	29:DH:146:VAL:HG23	2.00	0.60
40:DS:79:GLY:N	40:DS:100:THR:O	2.35	0.60
22:DA:1936:A:C8	22:DA:1945:G:C6	2.89	0.60
3:AC:7:PRO:HD2	3:AC:184:TYR:CD1	2.37	0.60
22:BA:2847:U:C2'	22:BA:2848:G:H5'	2.32	0.60
41:DT:34:VAL:HG21	41:DT:43:ILE:HD11	1.82	0.60
1:CA:502:A:H2'	1:CA:503:C:O4'	2.01	0.60
22:DA:561:G:O2'	38:DQ:45:TYR:OH	2.14	0.60
26:BE:12:LEU:HD23	26:BE:13:THR:N	2.16	0.60
22:DA:1127:A:H2'	22:DA:1128:G:H5'	1.84	0.60
17:AQ:12:VAL:O	17:AQ:13:VAL:HG12	2.02	0.60
22:DA:1207:C:N3	22:DA:1208:C:C5	2.69	0.60
50:D2:15:SER:HG	50:D2:16:HIS:CD2	2.20	0.60
1:CA:642:A:N3	8:CH:105:SER:OG	2.32	0.60
9:AI:44:ALA:HA	9:AI:46:MET:HE2	1.84	0.60
22:DA:1313:U:H4'	22:DA:1332:G:H4'	1.84	0.60
32:BK:4:GLU:O	32:BK:5:GLN:HB2	2.02	0.60
22:BA:1585:C:H2'	22:BA:1586:A:O4'	2.02	0.60
19:CS:80:TYR:CG	19:CS:81:ARG:N	2.69	0.60
22:DA:1855:U:C5	22:DA:1856:U:C5	2.88	0.60
1:AA:901:A:N7	1:AA:902:G:H1'	2.17	0.60
22:BA:2128:G:H5'	53:B5:36:ALA:HA	1.83	0.60
22:DA:575:A:C2	22:DA:576:U:C6	2.90	0.60
2:AB:213:TYR:O	2:AB:217:VAL:HG23	2.01	0.60
22:BA:2406:A:C2	33:BL:69:ARG:NH2	2.69	0.60
22:DA:2330:G:N2	22:DA:2386:A:C4	2.70	0.60
22:DA:1835:G:C4	22:DA:1836:C:C5	2.89	0.60
36:BO:64:TYR:HB3	36:BO:67:ASN:ND2	2.17	0.60
32:BK:21:CYS:HA	32:BK:41:ILE:HG22	1.83	0.60
22:DA:2373:G:H2'	22:DA:2374:C:C6	2.35	0.60
22:DA:635:C:O2'	22:DA:639:U:H5''	2.01	0.60
32:BK:116:ILE:HD12	32:BK:117:SER:N	2.16	0.60
1:CA:1053:G:O5'	1:CA:1054:C:H3'	2.02	0.60
41:BT:2:ILE:HD12	41:BT:7:LEU:CG	2.32	0.60
12:CL:71:GLY:O	12:CL:99:ARG:NH2	2.35	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1107:C:C4	1:AA:1108:G:N7	2.70	0.60
2:CB:85:LEU:O	2:CB:85:LEU:HG	2.01	0.60
22:BA:1794:A:H2'	22:BA:1795:C:C6	2.37	0.60
22:BA:271:G:C4	22:BA:367:G:N2	2.70	0.60
22:DA:1826:G:C4	22:DA:1827:U:C6	2.90	0.60
1:AA:1343:G:H2'	1:AA:1344:C:C6	2.36	0.60
1:CA:1521:C:N3	1:CA:1522:U:C5	2.70	0.60
46:BY:56:LEU:O	46:BY:57:LEU:HB2	2.02	0.60
1:CA:1480:A:C5	1:CA:1481:U:C5	2.90	0.60
1:CA:899:C:O2'	22:DA:1832:C:OP1	2.19	0.60
22:BA:1817:G:H2'	22:BA:1818:U:H5'	1.84	0.60
22:BA:1450:G:C6	22:BA:1451:C:N4	2.70	0.60
22:DA:608:A:H2'	22:DA:609:A:C8	2.37	0.60
22:BA:289:G:H2'	22:BA:290:U:O4'	2.02	0.60
22:BA:1832:C:N4	22:BA:1833:C:C4	2.70	0.60
22:DA:483:A:C8	22:DA:484:C:C5	2.90	0.60
1:AA:667:G:H4'	15:AO:51:HIS:ND1	2.17	0.60
1:CA:840:C:N3	1:CA:842:U:H4'	2.17	0.60
29:BH:117:LEU:CD2	29:BH:122:LEU:CD1	2.80	0.60
39:BR:49:ILE:C	39:BR:51:VAL:O	2.40	0.60
22:BA:1910:G:H2'	22:BA:1911:U:C6	2.37	0.60
22:DA:2053:G:N2	22:DA:2054:A:H1'	2.17	0.60
1:CA:728:A:N1	1:CA:729:A:C6	2.70	0.60
22:DA:1818:U:OP2	24:DC:156:ARG:NH1	2.35	0.60
22:DA:1835:G:H2'	22:DA:1836:C:H6	1.67	0.60
22:BA:1587:G:C5	22:BA:1588:G:N7	2.70	0.60
1:AA:1059:C:C2	1:AA:1060:U:C5	2.90	0.60
22:DA:1140:C:O4'	22:DA:1143:A:C2	2.55	0.60
22:DA:527:C:OP2	22:DA:2779:U:N3	2.34	0.60
22:DA:1252:G:H5''	57:DA:3283:HOH:O	2.00	0.60
1:CA:1087:G:N2	1:CA:1099:G:H1'	2.17	0.60
22:DA:2864:G:H2'	22:DA:2865:U:O4'	2.02	0.60
29:BH:99:ILE:HB	29:BH:115:VAL:HG11	1.84	0.59
22:BA:1060:U:O4'	22:BA:1062:G:H5'	2.02	0.59
22:DA:24:G:N2	22:DA:517:C:C2	2.70	0.59
22:DA:842:U:N3	22:DA:843:G:N7	2.50	0.59
13:CM:10:PRO:O	13:CM:11:ASP:HB2	2.02	0.59
1:CA:475:C:H2'	1:CA:476:U:C6	2.36	0.59
22:DA:629:G:N3	22:DA:639:U:O2'	2.31	0.59
22:DA:135:U:H2'	22:DA:136:G:C8	2.37	0.59
22:BA:2278:A:OP1	34:BM:10:ARG:NH2	2.35	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:CJ:52:LEU:HB2	14:CN:81:ARG:HD2	1.84	0.59
1:AA:951:G:C2	1:AA:952:U:C2	2.90	0.59
13:CM:40:ALA:O	13:CM:42:ASP:N	2.34	0.59
14:CN:51:LEU:O	14:CN:53:ARG:N	2.34	0.59
22:BA:1867:G:O2'	22:BA:1868:C:H5'	2.02	0.59
26:DE:149:ILE:HG23	26:DE:188:MET:HG2	1.83	0.59
17:AQ:14:SER:HB3	17:AQ:22:VAL:CG1	2.31	0.59
22:DA:1827:U:H2'	22:DA:1828:G:C8	2.37	0.59
22:DA:397:U:OP1	45:DX:31:PRO:HA	2.02	0.59
22:DA:727:A:H2'	22:DA:728:G:C8	2.37	0.59
11:AK:13:ARG:N	22:BA:2141:G:H4'	2.16	0.59
22:BA:2287:A:OP1	49:B1:30:LYS:NZ	2.35	0.59
1:AA:988:G:N2	1:AA:1217:C:O2	2.35	0.59
2:AB:184:PHE:CZ	2:AB:198:PHE:CD2	2.89	0.59
22:DA:2046:G:OP1	48:D0:12:LYS:NZ	2.35	0.59
1:AA:1142:G:H2'	1:AA:1143:G:O4'	2.02	0.59
22:BA:1869:G:C2	22:BA:1873:G:C6	2.90	0.59
22:DA:2415:G:C6	22:DA:2416:C:C4	2.89	0.59
1:CA:581:G:OP1	15:CO:65:LYS:NZ	2.32	0.59
22:BA:278:A:C2	22:BA:362:A:C8	2.90	0.59
22:BA:1588:G:C4	22:BA:1589:U:C5	2.91	0.59
4:AD:78:GLU:OE1	4:AD:81:ARG:NH1	2.35	0.59
22:BA:1936:A:H2	22:BA:1943:U:N3	1.99	0.59
22:DA:2111:U:C5	22:DA:2145:C:H2'	2.38	0.59
1:AA:691:G:O6	11:AK:57:LYS:NZ	2.28	0.59
1:CA:1486:G:H2'	1:CA:1487:G:O4'	2.01	0.59
2:CB:119:THR:O	2:CB:120:GLN:HB2	2.01	0.59
3:AC:148:GLY:HA3	3:AC:172:ARG:O	2.03	0.59
23:DB:66:A:N6	23:DB:107:G:H2'	2.17	0.59
38:DQ:58:ARG:NH2	38:DQ:92:ARG:NH1	2.50	0.59
2:CB:102:THR:O	2:CB:103:ASN:HB3	2.02	0.59
22:BA:139:U:O4	41:BT:2:ILE:HG13	2.02	0.59
22:DA:1826:G:C5	22:DA:1827:U:C4	2.90	0.59
22:DA:1131:G:O6	22:DA:2024:G:O2'	2.16	0.59
5:AE:15:LEU:HB3	5:AE:37:THR:HG22	1.83	0.59
22:BA:2339:C:H2'	22:BA:2340:A:C8	2.37	0.59
22:DA:71:A:OP2	22:DA:113:U:H5'	2.03	0.59
22:DA:2232:C:P	45:DX:27:ARG:HH12	2.25	0.59
31:DJ:7:LYS:O	31:DJ:11:VAL:HG23	2.03	0.59
29:BH:94:ILE:HG22	29:BH:99:ILE:CG1	2.32	0.59
22:DA:1358:G:O2'	22:DA:1359:A:H5'	2.01	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:976:G:C4	1:AA:1363:A:N6	2.70	0.59
1:CA:207:C:O2	1:CA:207:C:H2'	2.02	0.59
11:CK:112:ASP:HB3	21:CU:20:LYS:HE3	1.83	0.59
22:BA:2589:A:N7	57:BA:3551:HOH:O	2.32	0.59
29:DH:126:GLY:O	29:DH:146:VAL:N	2.35	0.59
22:BA:1132:U:H3'	22:BA:1133:A:H5''	1.85	0.59
22:DA:2069:G:N2	22:DA:2443:C:C2	2.71	0.59
1:CA:1348:U:H4'	9:CI:122:ARG:HG3	1.83	0.59
1:AA:1387:G:H2'	1:AA:1388:C:C6	2.36	0.59
2:CB:64:LYS:HD3	2:CB:64:LYS:C	2.23	0.59
22:BA:1223:G:OP2	39:BR:68:ARG:NH1	2.35	0.59
30:DI:58:VAL:O	30:DI:69:PHE:HB3	2.02	0.59
22:BA:2884:U:O4'	22:BA:2884:U:O2	2.14	0.59
23:DB:61:G:C6	23:DB:62:C:C4	2.90	0.59
22:BA:2114:A:N3	22:BA:2114:A:H2'	2.18	0.59
7:AG:99:LEU:O	7:AG:102:ARG:N	2.35	0.59
22:BA:1792:G:OP1	24:BC:204:VAL:O	2.20	0.59
24:DC:204:VAL:O	24:DC:206:GLY:N	2.35	0.59
1:AA:1121:U:C2	1:AA:1122:U:C6	2.91	0.59
9:AI:57:MET:O	9:AI:59:GLU:N	2.36	0.59
1:CA:1055:A:O2'	3:CC:161:GLU:O	2.20	0.59
29:BH:1:MET:O	29:BH:20:ASN:ND2	2.35	0.59
12:AL:56:ARG:NH1	12:AL:62:GLU:HG3	2.18	0.59
4:CD:174:ASP:O	4:CD:175:ALA:CB	2.51	0.59
12:AL:76:GLU:O	12:AL:77:HIS:HB2	2.02	0.59
22:DA:2886:A:C2	22:DA:2887:A:H1'	2.37	0.59
22:BA:1091:G:O2'	22:BA:1092:C:OP2	2.14	0.59
1:CA:1124:G:C2	1:CA:1127:G:N2	2.71	0.59
22:DA:2234:G:C5	22:DA:2235:G:N7	2.70	0.59
22:BA:2191:A:N1	22:BA:2192:U:N3	2.51	0.59
21:AU:34:ARG:O	21:AU:35:ARG:O	2.20	0.59
22:BA:977:G:C5	57:BA:3598:HOH:O	2.56	0.59
16:AP:47:GLU:O	16:AP:48:GLU:O	2.20	0.59
1:CA:1417:G:C6	1:CA:1482:G:C6	2.91	0.59
10:AJ:15:HIS:CG	10:AJ:16:ARG:N	2.70	0.59
1:AA:194:C:O2'	1:AA:195:A:H5'	2.02	0.59
22:DA:158:U:O2	22:DA:169:G:C2	2.56	0.59
22:DA:11:C:C2'	22:DA:12:U:H5'	2.33	0.59
15:AO:27:VAL:O	15:AO:31:LEU:HG	2.03	0.59
34:DM:70:ASP:OD1	34:DM:70:ASP:C	2.41	0.59
22:DA:2811:G:H2'	22:DA:2812:G:O4'	2.03	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
45:BX:64:ILE:HG23	45:BX:65:ASP:N	2.17	0.59
22:DA:185:G:C5	22:DA:212:G:N2	2.71	0.59
35:DN:90:ARG:NH2	35:DN:116:VAL:HG21	2.18	0.59
22:DA:1651:G:N2	22:DA:2007:U:O2	2.35	0.59
22:DA:658:U:C2	22:DA:659:G:C8	2.91	0.59
22:DA:1204:A:C2	22:DA:1240:U:N3	2.70	0.59
22:DA:2345:G:C6	22:DA:2347:C:N4	2.70	0.59
30:DI:28:LEU:HD13	30:DI:38:PHE:CD1	2.37	0.59
22:DA:2230:G:O3'	45:DX:30:LEU:HB2	2.02	0.59
22:DA:200:U:C4	22:DA:248:G:N2	2.71	0.59
1:AA:1029:U:O2'	1:AA:1032:G:O6	2.21	0.59
22:DA:1773:A:C2	22:DA:1978:A:C2	2.91	0.59
2:CB:23:TRP:O	2:CB:23:TRP:CD1	2.56	0.59
4:CD:35:GLU:HG3	4:CD:36:GLN:N	2.18	0.59
24:DC:67:PHE:CE1	24:DC:156:ARG:NH2	2.71	0.59
14:CN:51:LEU:HB3	14:CN:52:PRO:HD2	1.85	0.59
13:AM:73:ILE:O	13:AM:76:SER:OG	2.21	0.59
22:DA:2461:A:N1	22:DA:2490:G:N2	2.50	0.59
10:AJ:32:THR:OG1	10:AJ:33:GLY:N	2.36	0.59
22:DA:204:A:H5'	22:DA:206:U:O4'	2.03	0.59
22:DA:214:G:O2'	22:DA:216:A:O3'	2.20	0.59
22:DA:1596:A:N1	22:DA:1597:A:C2	2.71	0.59
22:DA:186:G:C2	22:DA:211:C:O2	2.56	0.59
22:DA:2328:A:H2'	22:DA:2329:U:C6	2.38	0.59
22:DA:2056:G:N3	22:DA:2056:G:H2'	2.18	0.59
1:CA:374:A:H5''	1:CA:452:A:C2	2.37	0.59
1:CA:505:G:C5	1:CA:535:A:C2	2.91	0.59
29:DH:81:ALA:C	29:DH:149:GLU:HB2	2.23	0.59
1:AA:1505:G:P	57:AA:1871:HOH:O	2.60	0.59
16:CP:5:ARG:O	16:CP:19:VAL:HA	2.01	0.59
22:BA:2856:A:N6	22:BA:2857:G:C6	2.71	0.59
23:DB:48:U:H2'	23:DB:49:C:C6	2.38	0.59
20:AT:58:VAL:HG12	20:AT:59:ASP:N	2.18	0.59
12:AL:44:LYS:HB2	12:AL:45:PRO:CD	2.33	0.59
1:CA:1463:U:H2'	1:CA:1464:U:C6	2.38	0.59
1:AA:209:U:HO2'	1:AA:210:C:P	2.26	0.59
22:BA:2615:U:C2	48:B0:4:GLN:HA	2.38	0.59
13:CM:6:GLY:O	13:CM:8:ASN:N	2.35	0.59
22:DA:188:G:HO2'	22:DA:1365:A:N6	2.01	0.58
22:DA:201:C:C4	22:DA:202:U:C4	2.91	0.58
22:DA:1951:U:H2'	22:DA:1953:A:OP2	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CB:21:ARG:HA	2:CB:21:ARG:CZ	2.33	0.58
17:CQ:14:SER:OG	17:CQ:22:VAL:HG12	2.03	0.58
22:BA:58:G:OP1	41:BT:78:SER:HB3	2.02	0.58
22:DA:2024:G:OP2	22:DA:2034:U:H4'	2.02	0.58
22:BA:2665:A:C2	22:BA:2666:C:C6	2.90	0.58
22:DA:2350:C:H2'	22:DA:2351:G:O4'	2.02	0.58
36:BO:31:THR:O	36:BO:102:ARG:NH1	2.35	0.58
22:DA:2563:U:H1'	22:DA:2566:A:C6	2.38	0.58
9:AI:10:GLY:HA2	9:AI:81:HIS:ND1	2.18	0.58
22:BA:2233:U:H2'	22:BA:2234:G:C8	2.38	0.58
22:BA:70:G:H4'	22:BA:71:A:OP1	2.01	0.58
22:DA:1275:A:C8	35:DN:16:HIS:ND1	2.71	0.58
27:BF:171:ALA:O	27:BF:174:ASP:N	2.36	0.58
22:BA:2346:A:H4'	22:BA:2347:C:OP2	2.02	0.58
1:CA:404:G:N7	4:CD:2:ALA:HB3	2.18	0.58
22:DA:1605:C:H2'	22:DA:1606:C:H5'	1.84	0.58
24:DC:160:THR:HG22	24:DC:177:ARG:HG2	1.85	0.58
22:BA:2187:U:H2'	22:BA:2188:U:O4'	2.03	0.58
38:BQ:76:TYR:CZ	38:BQ:80:ILE:HG13	2.37	0.58
50:D2:15:SER:OG	50:D2:16:HIS:CD2	2.56	0.58
1:CA:513:C:H2'	1:CA:514:C:C6	2.38	0.58
4:AD:150:LYS:O	4:AD:151:LYS:C	2.41	0.58
24:DC:136:PRO:O	24:DC:139:SER:OG	2.22	0.58
33:BL:26:GLY:O	33:BL:27:LEU:HD23	2.02	0.58
1:CA:1125:U:C6	10:CJ:40:ILE:HD13	2.39	0.58
40:DS:73:LYS:HB2	40:DS:106:VAL:HB	1.84	0.58
24:DC:108:LYS:HA	24:DC:196:GLY:HA2	1.84	0.58
22:DA:2259:U:H1'	22:DA:2427:C:C2	2.38	0.58
1:CA:252:U:O4	1:CA:253:A:N6	2.35	0.58
5:CE:82:GLN:N	5:CE:147:MET:HE3	2.18	0.58
1:CA:72:A:C5	1:CA:73:C:C4	2.91	0.58
22:DA:2566:A:N1	32:DK:28:SER:HB2	2.17	0.58
1:AA:1001:C:H2'	1:AA:1002:G:C8	2.38	0.58
21:AU:25:LYS:HD2	21:AU:26:ALA:H	1.67	0.58
1:CA:801:U:H2'	1:CA:802:A:H8	1.68	0.58
22:DA:1060:U:H4'	22:DA:1061:U:H5'	1.85	0.58
1:CA:195:A:OP1	20:CT:60:ARG:NH1	2.37	0.58
22:DA:2576:G:C8	22:DA:2580:U:O4	2.56	0.58
22:DA:30:G:C6	22:DA:31:C:N3	2.71	0.58
22:DA:2235:G:C5	22:DA:2236:U:C5	2.91	0.58
1:AA:496:A:C2	1:AA:497:G:C6	2.91	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:792:A:H1'	22:DA:2072:C:O2'	2.03	0.58
22:DA:749:A:C2	22:DA:750:A:C8	2.91	0.58
22:DA:1833:C:N3	22:DA:1834:U:C5	2.72	0.58
22:DA:1042:G:C6	22:DA:1043:C:C4	2.92	0.58
1:AA:1288:A:C6	1:AA:1289:A:C5	2.90	0.58
1:AA:1063:C:H2'	1:AA:1064:G:C8	2.39	0.58
22:DA:933:A:H5'	22:DA:934:U:OP2	2.03	0.58
17:CQ:70:THR:HG22	17:CQ:71:LYS:N	2.18	0.58
8:AH:11:LEU:HD11	8:AH:127:CYS:HB3	1.85	0.58
28:BG:20:ASN:ND2	28:BG:20:ASN:O	2.35	0.58
22:BA:1653:G:H3'	35:BN:2:ARG:HG3	1.84	0.58
22:BA:1435:G:O2'	22:BA:1436:G:H5'	2.03	0.58
22:BA:731:C:P	57:BA:3699:HOH:O	2.42	0.58
22:BA:1916:A:C2'	22:BA:1917:U:O4'	2.47	0.58
22:DA:1358:G:O6	22:DA:1371:G:C8	2.56	0.58
22:BA:1078:U:H1'	22:BA:1088:A:C2	2.38	0.58
33:DL:59:ARG:CZ	33:DL:59:ARG:HB3	2.33	0.58
22:DA:1208:C:C4	22:DA:1209:U:C4	2.91	0.58
22:DA:2093:G:C2	22:DA:2094:A:C5	2.91	0.58
22:DA:1351:C:HO2'	22:DA:1571:A:H1'	1.68	0.58
25:DD:103:ASP:O	25:DD:105:LYS:N	2.37	0.58
1:CA:1317:C:OP1	14:CN:57:PRO:HD2	2.04	0.58
36:BO:79:ALA:HB2	36:BO:110:ALA:HA	1.85	0.58
48:B0:55:ILE:HG22	48:B0:56:ALA:N	2.16	0.58
22:DA:1097:U:C5	22:DA:1098:A:H1'	2.38	0.58
1:CA:64:G:C8	1:CA:99:C:N4	2.71	0.58
45:BX:21:ALA:O	45:BX:22:LEU:HB2	2.03	0.58
1:AA:1521:C:C2	1:AA:1522:U:C5	2.91	0.58
5:AE:109:GLY:O	5:AE:110:ALA:HB3	2.04	0.58
22:DA:1245:G:H4'	26:DE:33:VAL:HG11	1.86	0.58
29:BH:31:VAL:N	29:BH:32:PRO:HD2	2.18	0.58
39:BR:49:ILE:HG22	39:BR:52:PRO:CA	2.34	0.58
22:DA:1378:A:O2'	57:DA:3753:HOH:O	2.17	0.58
22:DA:2115:G:N3	22:DA:2117:A:N7	2.51	0.58
22:BA:2298:A:C6	22:BA:2321:U:O4	2.56	0.58
21:AU:40:LYS:HA	21:AU:43:THR:HG23	1.86	0.58
22:DA:2134:A:OP2	22:DA:2157:G:N2	2.36	0.58
17:CQ:70:THR:HG22	17:CQ:71:LYS:H	1.67	0.58
40:DS:29:VAL:CG1	40:DS:55:ILE:HD11	2.34	0.58
22:BA:1359:A:C8	22:BA:1373:A:C2	2.92	0.58
1:CA:1084:G:C5	1:CA:1085:U:C4	2.92	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:DH:34:GLY:O	29:DH:35:LYS:CB	2.51	0.58
1:CA:604:G:H2'	1:CA:605:U:O4'	2.03	0.58
1:AA:1313:U:P	19:AS:6:LYS:HB3	2.44	0.58
22:DA:1199:U:H2'	22:DA:1200:C:C6	2.38	0.58
28:DG:166:ASP:OD1	28:DG:166:ASP:N	2.37	0.58
8:CH:3:MET:SD	8:CH:3:MET:N	2.76	0.58
51:D3:26:HIS:NE2	51:D3:48:ALA:HB2	2.19	0.58
22:DA:1389:G:N2	22:DA:1390:U:O2	2.36	0.58
1:AA:705:G:C5	1:AA:706:A:C8	2.92	0.58
22:DA:82:U:C2	22:DA:83:A:C8	2.92	0.58
22:DA:1805:A:N3	22:DA:1813:G:N2	2.52	0.58
1:CA:1411:C:H2'	1:CA:1412:C:H6	1.69	0.58
22:DA:587:C:N3	33:DL:33:ARG:NH2	2.51	0.58
2:AB:23:TRP:CZ3	2:AB:25:PRO:HA	2.38	0.58
16:CP:20:VAL:HG21	16:CP:32:PHE:HB2	1.86	0.58
1:AA:365:U:H5''	1:AA:366:A:OP1	2.02	0.58
38:BQ:89:GLU:H	39:BR:49:ILE:HD12	1.67	0.58
50:D2:10:LEU:HD11	50:D2:14:ARG:CZ	2.33	0.58
22:BA:1588:G:C2	22:BA:1589:U:C5	2.92	0.58
22:DA:55:G:C2	22:DA:56:A:C8	2.91	0.58
22:DA:1179:G:C5	22:DA:1180:U:H1'	2.38	0.58
22:DA:732:C:N4	22:DA:733:G:C5	2.72	0.58
24:BC:244:PRO:O	24:BC:251:GLN:OE1	2.21	0.58
16:CP:52:LEU:HD23	16:CP:53:ASP:N	2.19	0.58
5:CE:72:ILE:HD13	5:CE:145:GLU:CD	2.24	0.58
36:DO:79:ALA:O	36:DO:83:LEU:HG	2.03	0.58
22:DA:2677:G:C2	22:DA:2731:G:C2	2.92	0.58
22:DA:747:U:O2	22:DA:2014:A:H1'	2.03	0.58
22:BA:975:A:C2	22:BA:990:A:C8	2.91	0.58
2:CB:90:PHE:CD1	2:CB:150:GLY:O	2.56	0.58
42:DU:54:GLN:N	42:DU:55:PRO:HD3	2.19	0.58
52:D4:16:ILE:HD13	52:D4:25:VAL:HG22	1.85	0.58
22:DA:2143:C:H2'	22:DA:2144:G:O4'	2.03	0.58
1:CA:833:G:C6	1:CA:834:U:C4	2.91	0.58
22:DA:1525:A:C6	22:DA:1526:C:N3	2.72	0.58
50:D2:10:LEU:HD11	50:D2:14:ARG:NE	2.19	0.58
1:AA:1319:A:C8	1:AA:1323:G:C6	2.91	0.58
17:CQ:17:MET:HE2	17:CQ:20:SER:O	2.02	0.58
2:AB:49:MET:O	2:AB:53:ALA:HB2	2.03	0.58
22:DA:2204:G:C5	22:DA:2221:G:C2	2.92	0.58
1:CA:1201:A:H1'	1:CA:1202:U:OP2	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1241:G:C2	1:AA:1242:G:C5	2.92	0.58
22:DA:1324:G:N2	22:DA:1328:A:N1	2.51	0.58
22:DA:2817:U:O2'	22:DA:2836:U:O2	2.14	0.58
24:DC:34:LEU:O	24:DC:35:GLU:HB3	2.03	0.58
1:AA:73:C:O2'	1:AA:74:A:H5''	2.04	0.58
1:AA:212:G:N2	1:AA:213:G:C4	2.71	0.58
1:CA:1291:U:OP1	7:CG:37:SER:HB3	2.03	0.58
13:AM:114:LYS:HB2	13:AM:115:PRO:HD3	1.85	0.58
22:BA:1300:G:H4'	22:BA:1301:A:H5'	1.86	0.58
22:DA:834:G:H1'	22:DA:2358:A:N3	2.18	0.58
22:BA:1045:C:C3'	22:BA:1046:A:H5'	2.33	0.58
1:AA:109:A:H2'	1:AA:326:G:N2	2.19	0.58
1:CA:687:A:N3	1:CA:688:G:H1'	2.19	0.58
1:AA:144:G:C4	1:AA:179:A:C2	2.92	0.58
22:DA:684:G:H5'	50:D2:16:HIS:CE1	2.39	0.58
22:DA:2019:A:H4'	38:DQ:34:VAL:CG2	2.34	0.58
1:AA:1053:G:H4'	1:AA:1054:C:H5'	1.86	0.58
22:DA:2330:G:C2	22:DA:2386:A:C2	2.91	0.58
6:AF:5:GLU:O	6:AF:6:ILE:HB	2.04	0.58
1:CA:761:G:C2	1:CA:762:U:C2	2.91	0.58
36:DO:33:ARG:O	36:DO:34:HIS:HB2	2.04	0.58
1:CA:243:A:H4'	1:CA:244:U:H5''	1.86	0.58
22:DA:630:G:N2	22:DA:633:A:OP2	2.37	0.58
22:BA:2052:A:H4'	25:BD:148:GLN:O	2.04	0.58
1:AA:316:C:C2	1:AA:317:U:C5	2.91	0.58
1:AA:983:A:H2'	1:AA:983:A:N3	2.18	0.58
22:BA:1847:A:C8	22:BA:1847:A:OP2	2.57	0.58
31:BJ:125:TYR:OH	31:BJ:132:HIS:NE2	2.34	0.58
22:DA:681:G:C4	22:DA:682:G:C8	2.92	0.58
4:CD:30:THR:HG22	4:CD:30:THR:O	2.02	0.57
22:DA:1315:C:N3	22:DA:1338:G:C2	2.72	0.57
35:DN:90:ARG:NH1	35:DN:116:VAL:HG11	2.18	0.57
22:DA:197:A:C8	22:DA:2430:A:C8	2.92	0.57
1:CA:729:A:C4	1:CA:730:G:C8	2.92	0.57
1:AA:1299:A:C6	1:AA:1301:U:O2	2.57	0.57
35:DN:54:LEU:HD23	35:DN:66:ALA:HB2	1.85	0.57
26:DE:21:ARG:O	26:DE:114:ARG:NH2	2.37	0.57
1:CA:38:G:C2	1:CA:397:A:C2	2.92	0.57
22:BA:1319:C:O2'	22:BA:1320:C:H5'	2.03	0.57
29:DH:108:VAL:O	29:DH:110:VAL:N	2.36	0.57
20:AT:81:ALA:O	20:AT:85:LYS:HG2	2.03	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:1421:G:C2	22:BA:1422:G:C8	2.92	0.57
22:BA:1985:C:O2	22:BA:1985:C:H2'	2.03	0.57
1:CA:495:A:C2	1:CA:496:A:C6	2.92	0.57
1:CA:1222:G:O6	57:CA:1858:HOH:O	2.17	0.57
1:CA:31:G:H5'	1:CA:306:A:N1	2.19	0.57
1:CA:1022:A:C5	1:CA:1023:U:C4	2.92	0.57
12:CL:38:TYR:HB3	12:CL:39:THR:O	2.03	0.57
22:DA:1555:G:C2	22:DA:1556:C:C2	2.92	0.57
22:DA:11:C:H2'	22:DA:12:U:H5'	1.85	0.57
2:CB:91:PHE:O	2:CB:150:GLY:HA3	2.04	0.57
2:AB:96:TRP:CH2	2:AB:100:MET:HB3	2.38	0.57
1:AA:473:U:C2	1:AA:474:G:C8	2.91	0.57
22:DA:2183:A:H2'	22:DA:2184:A:C8	2.38	0.57
22:DA:936:A:C6	22:DA:937:C:C4	2.92	0.57
30:DI:20:PRO:HB2	30:DI:23:PRO:HD2	1.86	0.57
1:AA:1118:U:O4'	1:AA:1179:A:H1'	2.05	0.57
29:DH:62:LEU:HD13	29:DH:62:LEU:C	2.25	0.57
25:DD:148:GLN:HB2	25:DD:152:PRO:HG2	1.86	0.57
11:CK:125:LYS:O	21:CU:34:ARG:NE	2.37	0.57
22:BA:11:C:C2'	22:BA:12:U:H5'	2.33	0.57
1:AA:971:G:C8	1:AA:1365:G:H4'	2.40	0.57
1:CA:109:A:C6	1:CA:327:A:C6	2.91	0.57
22:DA:1240:U:HO2'	22:DA:1241:A:P	2.27	0.57
22:DA:1855:U:C5	22:DA:1856:U:C4	2.92	0.57
40:BS:29:VAL:CG1	40:BS:55:ILE:HD11	2.34	0.57
30:DI:103:ARG:O	30:DI:107:GLN:HB2	2.04	0.57
5:CE:41:ASP:OD1	5:CE:42:GLY:N	2.37	0.57
8:AH:2:SER:N	8:AH:4:GLN:HG3	2.19	0.57
22:DA:1361:G:C5	22:DA:1371:G:N2	2.72	0.57
37:DP:53:ARG:O	37:DP:56:HIS:N	2.36	0.57
35:BN:12:ARG:CZ	35:BN:20:MET:HE1	2.34	0.57
22:DA:1984:G:C6	22:DA:1985:C:C5	2.92	0.57
1:CA:409:U:H2'	1:CA:410:G:O4'	2.05	0.57
22:DA:2093:G:N2	22:DA:2094:A:C4	2.73	0.57
1:CA:671:G:C6	1:CA:672:U:N3	2.72	0.57
1:AA:1319:A:C5	1:AA:1323:G:C4	2.91	0.57
33:BL:68:SER:O	33:BL:69:ARG:CB	2.53	0.57
22:BA:1869:G:H3'	22:BA:1870:C:H5'	1.85	0.57
1:AA:208:U:C5	1:AA:210:C:C4	2.91	0.57
15:CO:19:ALA:O	15:CO:20:ASN:HB2	2.03	0.57
22:BA:1185:G:H4'	22:BA:1186:G:OP1	2.04	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:AN:91:GLY:O	14:AN:93:ILE:N	2.37	0.57
22:DA:2484:G:OP1	34:DM:44:ARG:NH2	2.37	0.57
7:CG:70:ARG:HD2	7:CG:97:ASN:HB3	1.86	0.57
14:AN:54:ASP:OD1	14:AN:59:ARG:NH1	2.37	0.57
22:DA:1584:U:O2	22:DA:1584:U:H3'	2.03	0.57
1:CA:1086:U:OP1	1:CA:1086:U:H4'	2.05	0.57
22:BA:1073:A:H3'	22:BA:1074:G:C5'	2.29	0.57
41:BT:3:ARG:HD2	41:BT:5:GLU:HB2	1.86	0.57
22:DA:1317:G:N2	22:DA:1336:A:N3	2.53	0.57
22:DA:374:A:C2	22:DA:401:A:C4	2.92	0.57
1:CA:328:C:O2	1:CA:328:C:C2'	2.51	0.57
22:DA:687:C:H1'	50:D2:4:THR:HG23	1.86	0.57
1:CA:728:A:C8	15:CO:54:ARG:NH1	2.73	0.57
22:DA:2133:G:N2	22:DA:2158:A:C6	2.72	0.57
29:DH:117:LEU:HB3	29:DH:120:GLY:O	2.05	0.57
1:CA:72:A:C5	1:CA:73:C:N4	2.72	0.57
22:DA:1343:G:H1'	22:DA:1597:A:C4	2.39	0.57
1:AA:209:U:O2'	1:AA:210:C:OP1	2.13	0.57
1:CA:1237:C:C5	1:CA:1336:C:C4	2.92	0.57
21:AU:4:ILE:HA	21:AU:20:LYS:CE	2.34	0.57
22:BA:740:C:OP2	57:BA:3703:HOH:O	2.17	0.57
49:B1:40:ASP:C	49:B1:40:ASP:OD1	2.43	0.57
1:CA:774:G:C6	1:CA:775:G:C5	2.92	0.57
1:AA:1124:G:H2'	1:AA:1145:A:C6	2.40	0.57
1:AA:1124:G:H2'	1:AA:1145:A:N6	2.18	0.57
22:BA:137:U:H2'	22:BA:140:C:C2	2.39	0.57
3:CC:7:PRO:HD2	3:CC:184:TYR:CD2	2.39	0.57
12:CL:114:ARG:HD2	12:CL:119:VAL:HG12	1.86	0.57
22:DA:2407:A:OP1	57:DA:3564:HOH:O	2.17	0.57
1:AA:131:A:H2'	1:AA:132:C:H6	1.68	0.57
1:AA:976:G:N9	1:AA:1363:A:N6	2.52	0.57
22:DA:197:A:H62	22:DA:2430:A:H2'	1.68	0.57
1:CA:671:G:C2	1:CA:672:U:C2	2.93	0.57
22:DA:600:G:OP1	26:DE:24:ASN:ND2	2.38	0.57
5:CE:57:PRO:O	5:CE:60:ILE:HG13	2.05	0.57
39:BR:47:VAL:HG11	39:BR:54:VAL:CG1	2.33	0.57
1:CA:135:C:O2	16:CP:1:MET:HB2	2.04	0.57
22:DA:1833:C:C2	22:DA:1834:U:C5	2.92	0.57
7:AG:120:LEU:HD22	7:AG:124:LEU:HD23	1.86	0.57
28:BG:74:SER:HA	28:BG:77:ILE:HG13	1.87	0.57
22:DA:938:G:C2	22:DA:939:G:N7	2.72	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:1998:A:OP2	25:DD:141:ARG:NH2	2.37	0.57
22:DA:269:C:N3	22:DA:270:A:C8	2.72	0.57
1:CA:692:U:O2'	1:CA:694:A:N7	2.32	0.57
15:CO:37:ASN:O	15:CO:40:GLN:HB2	2.03	0.57
22:DA:1031:G:H4'	52:D4:6:SER:HB2	1.86	0.57
29:BH:95:GLY:HA2	29:BH:117:LEU:HD22	1.87	0.57
1:CA:505:G:H2'	1:CA:506:G:C8	2.39	0.57
29:DH:21:VAL:HG22	29:DH:22:LYS:N	2.19	0.57
2:AB:188:ASP:OD2	2:AB:204:ASP:OD1	2.23	0.57
1:AA:409:U:OP1	4:AD:24:GLY:HA3	2.04	0.57
22:DA:528:A:H2'	22:DA:529:A:H5''	1.87	0.57
19:AS:37:ARG:O	19:AS:70:LYS:HD2	2.05	0.57
42:DU:2:ALA:HA	42:DU:85:PHE:CE1	2.40	0.57
20:AT:3:ASN:O	20:AT:4:ILE:C	2.43	0.57
22:DA:668:A:C4	22:DA:670:A:N7	2.73	0.57
22:DA:1654:A:OP1	35:DN:1:MET:HA	2.05	0.57
22:DA:391:A:C8	22:DA:392:U:C5	2.93	0.57
6:AF:6:ILE:HG12	6:AF:89:VAL:HG12	1.87	0.57
11:AK:25:ALA:HA	11:AK:30:THR:HG22	1.86	0.57
22:DA:1832:C:C4	22:DA:1833:C:C6	2.92	0.57
1:AA:652:U:C2	1:AA:752:G:N2	2.73	0.57
39:BR:68:ARG:HD3	39:BR:92:TRP:CZ2	2.38	0.57
10:AJ:32:THR:HG21	10:AJ:86:ALA:CB	2.34	0.57
1:CA:1084:G:C8	1:CA:1085:U:C6	2.92	0.57
2:AB:23:TRP:CH2	2:AB:25:PRO:HA	2.39	0.57
22:DA:733:G:OP2	57:DA:3294:HOH:O	2.18	0.57
6:CF:16:GLU:O	6:CF:19:PRO:HD2	2.05	0.57
22:DA:2813:A:H2'	22:DA:2814:A:C8	2.40	0.57
38:DQ:47:TYR:C	38:DQ:47:TYR:CD1	2.78	0.57
22:DA:109:C:H4'	22:DA:348:A:H4'	1.86	0.57
19:CS:4:SER:O	19:CS:5:LEU:HB2	2.03	0.57
22:DA:647:G:C5	22:DA:648:G:N7	2.73	0.57
1:AA:430:A:OP1	4:AD:9:LEU:HB2	2.05	0.57
22:DA:2772:C:H2'	22:DA:2773:C:C6	2.39	0.57
29:BH:132:PHE:CE2	29:BH:142:VAL:HG21	2.40	0.57
1:AA:71:A:O2'	1:AA:72:A:P	2.63	0.57
33:BL:62:PRO:HG2	51:B3:25:LYS:HD3	1.85	0.57
1:CA:72:A:N6	1:CA:73:C:N4	2.52	0.57
1:AA:457:G:C6	1:AA:458:U:C2	2.93	0.57
6:CF:64:VAL:HG12	6:CF:65:GLU:N	2.19	0.57
39:DR:16:GLU:OE1	39:DR:100:GLY:HA2	2.04	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:BG:84:THR:OG1	28:BG:134:LYS:HG2	2.05	0.57
33:BL:59:ARG:HD2	57:BL:306:HOH:O	2.04	0.57
1:CA:1314:C:H2'	1:CA:1315:U:C6	2.40	0.57
22:DA:697:G:C2	22:DA:766:U:O2	2.57	0.57
30:DI:89:GLY:HA3	30:DI:136:MET:HE3	1.86	0.57
29:BH:99:ILE:CD1	29:BH:121:VAL:HG21	2.35	0.57
24:DC:107:PRO:CD	24:DC:110:LEU:HD22	2.34	0.57
22:DA:120:U:C2	22:DA:149:A:C6	2.92	0.57
22:DA:202:U:H2'	22:DA:203:A:C8	2.40	0.57
1:AA:923:A:O4'	1:AA:1398:A:C2	2.58	0.57
24:BC:17:VAL:N	24:BC:204:VAL:HG22	2.19	0.57
1:AA:914:A:C5	1:AA:915:A:N7	2.73	0.57
5:AE:80:THR:OG1	5:AE:81:LEU:N	2.38	0.57
22:BA:790:U:O2'	22:BA:791:C:P	2.63	0.57
1:CA:35:G:N3	12:CL:115:SER:OG	2.38	0.57
4:AD:147:GLU:O	4:AD:150:LYS:HB2	2.05	0.57
22:BA:265:A:H4'	22:BA:266:G:OP1	2.02	0.57
36:DO:64:TYR:O	36:DO:67:ASN:ND2	2.37	0.57
5:CE:95:PHE:O	5:CE:125:ALA:O	2.23	0.57
22:DA:734:A:C5	22:DA:735:A:C8	2.92	0.57
22:DA:1914:C:C5	22:DA:1915:U:C2	2.92	0.57
16:AP:42:ILE:HG22	16:AP:42:ILE:O	2.05	0.57
5:CE:76:LEU:N	5:CE:76:LEU:HD12	2.19	0.57
6:AF:64:VAL:HG12	6:AF:65:GLU:N	2.19	0.57
34:BM:77:PRO:HG2	34:BM:80:VAL:CG2	2.35	0.57
1:CA:971:G:OP1	1:CA:972:C:H5''	2.05	0.57
22:BA:1915:U:O2'	22:BA:1916:A:H5'	2.05	0.57
22:BA:1059:G:H5''	22:BA:1060:U:H2'	1.87	0.57
18:CR:35:GLU:HB2	21:CU:19:PHE:HZ	1.70	0.57
22:DA:377:G:C5	22:DA:378:C:C5	2.93	0.57
3:AC:7:PRO:HG2	3:AC:184:TYR:CG	2.40	0.57
4:CD:105:MET:SD	4:CD:143:VAL:HG13	2.45	0.57
4:CD:146:ARG:O	4:CD:150:LYS:HB2	2.05	0.57
2:CB:87:CYS:HB2	2:CB:89:GLN:NE2	2.18	0.57
1:CA:804:U:H5''	1:CA:805:C:OP2	2.05	0.57
14:AN:9:ARG:O	14:AN:13:ARG:HG2	2.04	0.57
22:BA:1084:A:C5	22:BA:1085:A:C6	2.93	0.57
23:DB:7:G:H5'	36:DO:29:HIS:CE1	2.40	0.57
22:BA:1936:A:C2	22:BA:1945:G:C8	2.92	0.57
22:DA:732:C:C4	22:DA:733:G:N7	2.73	0.57
16:CP:43:ALA:O	16:CP:44:SER:OG	2.20	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1296:C:H4'	1:AA:1302:C:C5	2.40	0.57
1:CA:711:G:O2'	1:CA:712:A:H5'	2.05	0.57
22:DA:1012:U:O4	31:DJ:30:THR:HG21	2.04	0.57
22:DA:1462:C:C2	22:DA:1463:C:C5	2.93	0.57
22:BA:1880:U:H2'	22:BA:1881:C:C6	2.39	0.57
28:BG:30:ASN:O	28:BG:30:ASN:OD1	2.23	0.57
1:CA:570:G:C6	1:CA:873:A:C2	2.93	0.57
23:DB:39:A:H2'	23:DB:40:U:C6	2.40	0.57
22:BA:570:G:H2'	22:BA:2030:A:N7	2.19	0.56
22:BA:627:A:C6	22:BA:637:A:C8	2.92	0.56
1:AA:1226:C:O2'	13:AM:110:LYS:NZ	2.37	0.56
45:DX:3:ARG:HG2	45:DX:33:LEU:HD22	1.87	0.56
22:DA:27:G:O2'	22:DA:28:A:OP2	2.17	0.56
1:CA:920:U:C2	1:CA:921:U:C5	2.93	0.56
10:AJ:44:THR:HG22	10:AJ:70:HIS:HA	1.86	0.56
22:DA:30:G:C2	22:DA:31:C:C2	2.93	0.56
1:AA:1286:U:H2'	1:AA:1286:U:O2	2.05	0.56
13:CM:91:HIS:HA	13:CM:109:ARG:NH2	2.20	0.56
8:AH:89:LYS:HG3	8:AH:90:ASP:N	2.20	0.56
42:BU:49:VAL:O	42:BU:51:ALA:N	2.38	0.56
22:BA:702:U:O2	22:BA:702:U:H2'	2.04	0.56
31:DJ:77:HIS:HA	31:DJ:83:GLY:O	2.05	0.56
22:BA:1912:A:C2	22:BA:1919:A:C5	2.92	0.56
22:DA:1394:U:H4'	22:DA:1603:A:H4'	1.88	0.56
21:CU:34:ARG:NE	21:CU:35:ARG:HB2	2.20	0.56
1:CA:552:U:N3	1:CA:553:A:N7	2.52	0.56
4:CD:34:ILE:O	4:CD:35:GLU:HB3	2.05	0.56
2:CB:94:HIS:CD2	2:CB:146:ASN:HB2	2.40	0.56
1:AA:262:A:C6	1:AA:263:A:C6	2.92	0.56
7:AG:145:ALA:O	7:AG:146:GLU:HB2	2.04	0.56
1:AA:1201:A:H1'	1:AA:1202:U:OP2	2.05	0.56
22:DA:1126:A:H4'	22:DA:1127:A:O5'	2.05	0.56
22:DA:699:A:C2'	22:DA:700:G:H5'	2.35	0.56
22:BA:2529:G:O6	52:B4:32:LYS:NZ	2.37	0.56
6:CF:66:ALA:HB1	6:CF:67:PRO:HD2	1.87	0.56
30:BI:117:MET:SD	30:BI:129:ILE:HD11	2.45	0.56
22:DA:583:G:C6	22:DA:584:C:C4	2.93	0.56
46:BY:45:GLN:O	46:BY:46:VAL:HB	2.03	0.56
22:BA:1917:U:O2	22:BA:1918:A:O4'	2.23	0.56
22:BA:1924:C:C2	22:BA:1926:U:O4	2.59	0.56
22:DA:1358:G:N2	22:DA:1374:G:C6	2.73	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:1124:G:N2	1:CA:1127:G:C2	2.73	0.56
22:DA:191:A:C6	22:DA:192:C:N4	2.74	0.56
22:DA:1809:A:C6	22:DA:1810:A:C6	2.94	0.56
4:CD:26:ARG:HG3	4:CD:27:ALA:N	2.19	0.56
1:AA:91:U:H2'	1:AA:92:U:O4'	2.05	0.56
22:BA:1745:A:C2	22:BA:1746:A:C8	2.94	0.56
1:CA:833:G:C5	1:CA:834:U:C5	2.93	0.56
22:DA:372:G:N2	22:DA:401:A:OP2	2.34	0.56
22:DA:352:A:H2'	22:DA:353:C:O4'	2.05	0.56
22:DA:301:G:C2	22:DA:302:C:C2	2.93	0.56
1:CA:32:A:H2'	1:CA:33:A:C8	2.40	0.56
22:DA:2093:G:N7	22:DA:2225:A:C8	2.73	0.56
22:DA:1379:U:H2'	22:DA:1379:U:O2	2.04	0.56
24:DC:72:ASP:HA	24:DC:118:SER:O	2.06	0.56
1:AA:315:A:O2'	1:AA:330:C:H4'	2.05	0.56
14:AN:14:VAL:HA	14:AN:60:GLN:OE1	2.04	0.56
1:CA:783:C:C2	1:CA:784:A:C8	2.93	0.56
53:B5:204:GLY:O	53:B5:205:ALA:HB3	2.05	0.56
22:BA:1738:G:HO2'	22:BA:1739:A:P	2.28	0.56
1:CA:610:U:C4	1:CA:611:C:C5	2.93	0.56
33:DL:108:ALA:HB3	33:DL:125:LEU:HG	1.87	0.56
22:DA:989:G:OP2	47:DZ:12:SER:OG	2.12	0.56
22:DA:2642:G:OP1	31:DJ:78:THR:OG1	2.24	0.56
28:BG:124:GLU:OE2	28:BG:125:CYS:N	2.37	0.56
23:BB:37:C:C5	23:BB:38:C:C4	2.93	0.56
22:DA:2297:A:C2	22:DA:2298:A:C8	2.92	0.56
20:AT:67:ILE:HD11	20:AT:71:LYS:CE	2.34	0.56
22:DA:65:U:H2'	22:DA:66:C:C6	2.38	0.56
15:AO:63:ARG:HG2	15:AO:67:LEU:HD12	1.88	0.56
1:CA:1239:A:H2'	1:CA:1298:U:O4	2.05	0.56
1:CA:1147:C:O2	9:CI:18:ARG:NH2	2.38	0.56
22:BA:695:G:C2	22:BA:696:G:C8	2.93	0.56
43:DV:9:ARG:CG	43:DV:41:GLU:HB3	2.35	0.56
9:CI:106:ARG:NH1	9:CI:107:ASP:O	2.38	0.56
1:AA:1236:A:H2'	1:AA:1237:C:C6	2.41	0.56
1:CA:682:G:C2	1:CA:683:G:C8	2.94	0.56
50:D2:18:PHE:O	50:D2:19:ARG:C	2.44	0.56
22:DA:305:C:H1'	22:DA:313:G:N2	2.20	0.56
35:BN:58:ASP:OD1	35:BN:63:ARG:HD2	2.06	0.56
41:BT:2:ILE:HG22	41:BT:3:ARG:H	1.68	0.56
1:CA:451:A:OP2	16:CP:70:ARG:NH2	2.36	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:728:G:C2	22:DA:730:A:C4	2.94	0.56
21:CU:41:PRO:O	21:CU:43:THR:N	2.39	0.56
22:DA:2199:A:C5	22:DA:2225:A:N1	2.74	0.56
1:AA:115:G:H4'	1:AA:116:A:O5'	2.04	0.56
33:DL:29:LYS:HG2	33:DL:29:LYS:O	2.03	0.56
22:BA:280:U:H2'	22:BA:281:C:C6	2.39	0.56
22:DA:1104:C:N4	22:DA:1105:U:O4	2.38	0.56
1:CA:445:G:C2	1:CA:446:G:C8	2.93	0.56
22:BA:2715:C:C4	22:BA:2716:C:C5	2.93	0.56
28:DG:40:ALA:HA	28:DG:58:TYR:CD1	2.40	0.56
5:CE:25:VAL:N	5:CE:28:GLY:O	2.34	0.56
2:AB:149:GLY:O	2:AB:151:ILE:N	2.38	0.56
30:BI:11:LEU:HD11	30:BI:27:ALA:O	2.05	0.56
22:DA:2516:A:N6	22:DA:2517:C:N4	2.53	0.56
22:BA:1344:U:O2'	22:BA:1345:C:P	2.64	0.56
1:CA:1181:G:O2'	1:CA:1182:G:C8	2.49	0.56
45:DX:33:LEU:O	45:DX:34:HIS:ND1	2.38	0.56
1:AA:1397:C:HO2'	1:AA:1398:A:P	2.23	0.56
1:AA:80:A:C2	1:AA:90:C:N3	2.73	0.56
50:D2:15:SER:OG	50:D2:16:HIS:NE2	2.38	0.56
22:DA:2018:G:H2'	22:DA:2019:A:O4'	2.05	0.56
42:DU:82:ARG:CB	42:DU:97:LYS:HG3	2.35	0.56
1:AA:502:A:OP1	12:AL:115:SER:CB	2.54	0.56
22:DA:749:A:C6	22:DA:1618:A:C2	2.93	0.56
12:CL:66:TYR:O	12:CL:97:THR:OG1	2.23	0.56
1:CA:378:G:C2	1:CA:386:C:O2	2.58	0.56
1:CA:562:U:O2'	12:CL:13:ALA:O	2.14	0.56
39:BR:68:ARG:HD3	39:BR:92:TRP:CE2	2.40	0.56
1:CA:1298:U:O2	1:CA:1298:U:C2'	2.53	0.56
1:CA:159:G:N2	1:CA:161:A:H3'	2.20	0.56
36:BO:116:GLN:O	36:BO:117:PHE:HB3	2.06	0.56
21:CU:53:VAL:HG13	21:CU:54:LYS:N	2.21	0.56
19:CS:48:THR:HG22	19:CS:61:PHE:CD2	2.41	0.56
38:DQ:78:LYS:HE2	38:DQ:117:LEU:HD21	1.88	0.56
34:BM:22:GLN:HA	34:BM:22:GLN:OE1	2.05	0.56
34:BM:62:LYS:HD3	34:BM:64:TRP:CZ2	2.40	0.56
57:BA:3296:HOH:O	33:BL:99:ASN:ND2	2.29	0.56
35:BN:103:ARG:HD3	35:BN:110:MET:HE3	1.86	0.56
5:CE:13:GLU:HB2	5:CE:39:VAL:HG12	1.87	0.56
1:CA:1053:G:N7	1:CA:1200:C:H5''	2.20	0.56
1:CA:1461:G:C5	1:CA:1462:C:C4	2.94	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:731:G:OP1	1:AA:766:A:H1'	2.06	0.56
1:AA:669:G:O2'	1:AA:670:G:H5'	2.06	0.56
1:CA:283:U:C4	1:CA:284:C:C4	2.94	0.56
2:CB:47:VAL:HB	2:CB:48:PRO:HD3	1.88	0.56
22:BA:694:U:OP1	24:BC:59:LYS:NZ	2.39	0.56
22:DA:952:G:C2	22:DA:966:G:C2	2.93	0.56
45:DX:21:ALA:O	45:DX:22:LEU:HB2	2.05	0.56
1:CA:375:U:P	16:CP:70:ARG:HH11	2.29	0.56
1:CA:429:U:O3'	4:CD:22:LYS:HE3	2.06	0.56
22:DA:1830:C:H5'	24:DC:15:HIS:CE1	2.41	0.56
22:DA:1651:G:C2	22:DA:2007:U:O2	2.59	0.56
8:CH:114:ARG:O	8:CH:115:ALA:C	2.44	0.56
2:CB:21:ARG:O	2:CB:23:TRP:N	2.38	0.56
1:AA:1157:A:C6	1:AA:1180:A:C5	2.93	0.56
12:CL:29:GLN:O	12:CL:30:LYS:HG2	2.06	0.56
22:BA:2315:G:H2'	22:BA:2316:G:C8	2.40	0.56
6:AF:91:ARG:O	6:AF:92:THR:OG1	2.22	0.56
22:DA:503:A:C2	22:DA:506:G:C4	2.94	0.56
22:DA:2351:G:O2'	22:DA:2366:A:N6	2.39	0.56
2:CB:140:GLU:O	2:CB:144:LEU:HG	2.06	0.56
22:DA:1980:G:O2'	22:DA:1982:U:OP2	2.21	0.56
19:CS:69:HIS:ND1	19:CS:73:GLU:OE2	2.38	0.56
52:B4:26:ILE:HD13	52:B4:26:ILE:N	2.21	0.56
27:BF:41:GLY:O	27:BF:43:ALA:N	2.39	0.56
1:CA:1302:C:C4	13:CM:17:ILE:HD11	2.40	0.56
22:BA:2190:G:OP2	22:BA:2190:G:H8	1.88	0.56
6:CF:92:THR:HG22	6:CF:93:LYS:N	2.20	0.56
4:AD:3:ARG:NH2	4:AD:115:ARG:HD3	2.20	0.56
1:CA:247:G:O6	1:CA:278:G:N1	2.38	0.56
22:DA:668:A:C2	22:DA:670:A:C6	2.94	0.56
22:BA:2517:C:C6	22:BA:2542:A:N7	2.73	0.56
11:AK:88:GLY:N	11:AK:114:THR:HG22	2.21	0.56
22:DA:577:G:O2'	22:DA:1254:A:OP1	2.23	0.56
25:BD:133:THR:HG23	25:BD:134:HIS:N	2.21	0.56
22:DA:572:A:C2	22:DA:2033:A:C2	2.94	0.56
4:CD:174:ASP:OD2	4:CD:176:GLY:N	2.39	0.56
22:DA:765:C:H2'	22:DA:766:U:C6	2.41	0.56
41:DT:64:LYS:HA	41:DT:79:ASP:OD2	2.06	0.56
29:BH:40:THR:OG1	29:BH:43:ASN:OD1	2.24	0.56
1:CA:1306:A:C6	1:CA:1307:U:N3	2.72	0.56
11:CK:67:ALA:O	11:CK:71:ALA:N	2.39	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:250:G:OP2	51:B3:13:ARG:NH1	2.38	0.56
22:DA:545:U:O2'	22:DA:547:A:OP1	2.23	0.56
22:BA:508:A:H4'	22:BA:509:C:OP2	2.06	0.56
22:BA:2520:C:C6	22:BA:2567:G:H1'	2.41	0.56
15:AO:45:GLU:HG2	15:AO:46:HIS:N	2.20	0.56
1:AA:810:C:H2'	1:AA:810:C:O2	2.05	0.56
2:CB:117:LEU:O	2:CB:121:SER:N	2.39	0.56
1:AA:813:U:H2'	1:AA:814:A:H5''	1.87	0.56
1:CA:1521:C:C2	1:CA:1522:U:C6	2.93	0.56
4:AD:32:CYS:SG	4:AD:34:ILE:N	2.77	0.56
11:AK:69:ARG:CD	22:BA:2146:C:N3	2.67	0.56
1:CA:484:G:C5	1:CA:486:U:H1'	2.40	0.56
22:DA:1208:C:C2	22:DA:1209:U:C6	2.93	0.56
35:BN:103:ARG:CD	35:BN:110:MET:HE3	2.36	0.56
2:CB:20:THR:O	2:CB:21:ARG:NH2	2.39	0.56
42:DU:98:SER:O	42:DU:99:ASN:HB3	2.06	0.56
22:BA:674:G:H1'	26:BE:69:ARG:HD3	1.88	0.56
22:DA:362:A:C4	22:DA:363:G:C8	2.94	0.56
12:CL:74:LEU:HD11	12:CL:80:ILE:HG21	1.88	0.56
22:BA:1587:G:C4	22:BA:1588:G:C8	2.93	0.56
22:DA:503:A:N3	22:DA:506:G:C8	2.73	0.56
1:AA:1033:G:H2'	1:AA:1034:G:H5'	1.88	0.56
30:BI:116:ASP:O	30:BI:117:MET:HB2	2.05	0.56
22:DA:126:A:OP2	50:D2:19:ARG:HG3	2.06	0.56
1:AA:184:G:H2'	1:AA:185:U:C6	2.39	0.56
33:DL:56:PRO:O	33:DL:60:ARG:CB	2.53	0.56
22:BA:947:A:HO2'	22:BA:984:A:H2	1.49	0.56
1:CA:1036:A:H3'	1:CA:1037:C:C6	2.41	0.56
8:CH:88:ARG:O	8:CH:122:GLY:HA3	2.05	0.56
19:CS:50:ALA:HB1	19:CS:57:HIS:HB3	1.87	0.56
26:BE:181:ILE:CG2	33:BL:2:ARG:HB3	2.36	0.56
1:CA:130:A:N3	1:CA:263:A:O2'	2.30	0.56
22:DA:1802:A:OP2	22:DA:1815:A:N6	2.39	0.56
39:BR:3:ALA:CB	39:BR:59:ILE:HD11	2.36	0.56
22:BA:1288:G:C5	22:BA:1327:A:C2	2.94	0.56
22:DA:1441:G:H2'	22:DA:1442:U:C6	2.41	0.56
1:CA:624:C:H2'	1:CA:625:U:O4'	2.05	0.56
22:BA:2887:A:H2'	22:BA:2887:A:N3	2.19	0.56
1:CA:782:A:C8	1:CA:783:C:C5	2.94	0.56
22:DA:225:C:H2'	22:DA:226:A:O4'	2.06	0.56
49:B1:17:THR:HG21	49:B1:42:VAL:HB	1.87	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:AH:78:VAL:HG11	8:AH:125:ILE:CD1	2.36	0.56
1:AA:660:C:P	15:AO:5:THR:HG21	2.45	0.56
1:AA:205:A:H2'	1:AA:205:A:N3	2.21	0.56
1:CA:955:U:H2'	1:CA:956:U:O4'	2.06	0.56
22:DA:1731:G:C2	22:DA:1733:G:C4	2.94	0.56
22:BA:2080:A:O5'	45:BX:19:SER:HB2	2.05	0.56
31:BJ:24:THR:HG22	31:BJ:24:THR:O	2.06	0.56
22:DA:2796:U:C4	22:DA:2798:U:C4	2.94	0.56
20:CT:67:ILE:HG13	20:CT:71:LYS:HD3	1.88	0.56
35:BN:45:ARG:HG2	35:BN:95:THR:HG21	1.86	0.56
1:CA:1092:A:N6	1:CA:1093:A:N6	2.54	0.55
22:DA:187:G:N2	22:DA:210:C:H1'	2.21	0.55
11:AK:76:GLU:HA	22:BA:2141:G:P	2.46	0.55
12:CL:22:PRO:O	12:CL:24:LEU:N	2.35	0.55
1:CA:978:A:HO2'	1:CA:1322:C:H5	1.53	0.55
1:CA:32:A:N3	1:CA:33:A:C8	2.74	0.55
1:CA:938:A:N6	1:CA:939:G:C6	2.75	0.55
12:CL:90:LEU:HB2	12:CL:93:VAL:CG2	2.35	0.55
1:AA:1074:G:C4	1:AA:1102:A:C2	2.94	0.55
22:BA:1866:A:N1	22:BA:1876:A:C8	2.74	0.55
23:BB:28:C:OP1	36:BO:31:THR:HG21	2.05	0.55
22:DA:1095:A:C6	22:DA:1096:A:C2	2.94	0.55
22:DA:2223:G:C6	22:DA:2224:G:C4	2.93	0.55
4:CD:126:ASN:OD1	4:CD:142:VAL:HG22	2.06	0.55
15:CO:53:ARG:O	15:CO:56:LEU:HB3	2.06	0.55
22:BA:455:C:N3	22:BA:472:A:H2'	2.20	0.55
22:DA:533:G:H5'	38:DQ:24:TYR:CE1	2.41	0.55
22:DA:1280:G:C6	22:DA:1281:G:C5	2.94	0.55
39:DR:49:ILE:HD12	39:DR:52:PRO:HA	1.87	0.55
2:AB:9:MET:SD	2:AB:9:MET:N	2.78	0.55
42:DU:59:VAL:CG1	42:DU:61:LYS:HD3	2.37	0.55
12:AL:24:LEU:HB2	12:AL:59:ASN:ND2	2.20	0.55
22:DA:36:G:C2'	22:DA:450:G:HO2'	2.19	0.55
11:AK:126:LYS:O	21:AU:34:ARG:NE	2.39	0.55
1:AA:1357:A:C5	1:AA:1358:U:C4	2.94	0.55
22:DA:674:G:O2'	26:DE:69:ARG:HD3	2.06	0.55
22:DA:674:G:O2'	26:DE:69:ARG:NE	2.39	0.55
28:BG:149:ARG:HH11	28:BG:149:ARG:CG	2.19	0.55
1:CA:786:G:C2	1:CA:797:C:O2	2.60	0.55
1:CA:386:C:N4	1:CA:387:U:C4	2.75	0.55
9:AI:43:THR:O	9:AI:44:ALA:HB3	2.05	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:AE:137:VAL:HG13	5:AE:137:VAL:O	2.05	0.55
22:BA:1435:G:C2'	22:BA:1436:G:H5'	2.36	0.55
22:DA:1096:A:H2'	22:DA:1097:U:O4'	2.07	0.55
40:BS:29:VAL:HG13	40:BS:55:ILE:HD11	1.88	0.55
22:DA:228:C:H5''	22:DA:229:C:C6	2.41	0.55
26:DE:41:GLN:O	26:DE:43:THR:N	2.36	0.55
11:AK:26:SER:O	11:AK:28:ASN:N	2.39	0.55
1:AA:1328:C:H5''	13:AM:28:THR:HG21	1.88	0.55
1:CA:668:G:N2	1:CA:739:C:C2	2.74	0.55
2:CB:203:ASN:OD1	2:CB:204:ASP:N	2.38	0.55
22:DA:151:C:H2'	22:DA:152:A:C8	2.41	0.55
1:AA:412:A:H4'	1:AA:413:G:OP1	2.06	0.55
1:AA:1221:G:OP1	1:AA:1320:C:N4	2.40	0.55
22:DA:1832:C:N4	22:DA:1833:C:C5	2.74	0.55
12:AL:114:ARG:HB2	12:AL:114:ARG:CZ	2.36	0.55
1:AA:1313:U:O4	19:AS:4:SER:HA	2.05	0.55
16:CP:42:ILE:O	16:CP:43:ALA:HB3	2.04	0.55
22:DA:2223:G:H2'	22:DA:2224:G:H5'	1.87	0.55
1:AA:1098:C:OP1	2:AB:143:LYS:NZ	2.30	0.55
22:BA:65:U:H2'	22:BA:66:C:C6	2.41	0.55
22:DA:40:U:C4	22:DA:41:C:N4	2.75	0.55
49:B1:34:LEU:H	49:B1:52:ALA:CB	2.19	0.55
32:DK:63:VAL:HB	32:DK:103:VAL:HG12	1.89	0.55
26:BE:106:LYS:HG3	26:BE:200:LEU:HG	1.86	0.55
22:DA:1063:G:N2	22:DA:1076:C:O2	2.39	0.55
29:BH:86:ASP:O	29:BH:87:GLU:CB	2.53	0.55
21:CU:41:PRO:O	21:CU:44:GLU:N	2.39	0.55
22:DA:674:G:N2	22:DA:2445:G:OP1	2.39	0.55
22:DA:1692:U:O2'	22:DA:1693:U:H2'	2.07	0.55
1:CA:672:U:H2'	1:CA:673:A:H8	1.71	0.55
22:DA:443:A:N7	26:DE:40:ARG:HG3	2.22	0.55
24:DC:67:PHE:CZ	24:DC:156:ARG:NH2	2.74	0.55
5:AE:114:VAL:HG21	5:AE:141:ILE:CD1	2.36	0.55
33:BL:76:GLU:HB2	33:BL:111:ILE:HD13	1.89	0.55
22:DA:2345:G:C5	22:DA:2381:A:C2	2.94	0.55
1:AA:1288:A:C6	1:AA:1289:A:N7	2.74	0.55
22:DA:1327:A:N6	22:DA:1328:A:C2	2.73	0.55
22:DA:2716:C:H2'	22:DA:2717:C:H6	1.71	0.55
37:DP:21:ARG:HB3	37:DP:22:PRO:HD2	1.88	0.55
26:DE:75:SER:O	26:DE:78:TRP:HB2	2.06	0.55
2:CB:28:LYS:N	2:CB:29:PRO:CD	2.69	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:102:U:C2	46:DY:2:LYS:HE2	2.41	0.55
43:BV:20:LEU:HD23	43:BV:25:LYS:HB2	1.87	0.55
32:DK:90:ASN:OD1	32:DK:90:ASN:N	2.40	0.55
22:BA:226:A:C6	22:BA:227:A:C6	2.95	0.55
22:BA:870:U:H2'	22:BA:871:U:H5'	1.88	0.55
22:DA:677:A:C2	22:DA:678:C:C2	2.94	0.55
22:DA:787:C:OP1	57:DA:3755:HOH:O	2.18	0.55
10:AJ:54:SER:O	14:AN:81:ARG:NH2	2.40	0.55
29:BH:98:ASP:O	29:BH:102:ALA:HB3	2.07	0.55
22:BA:192:C:P	57:BA:3749:HOH:O	2.64	0.55
31:BJ:81:ILE:CG2	31:BJ:82:GLY:N	2.64	0.55
22:DA:1984:G:O6	22:DA:1985:C:N4	2.39	0.55
1:CA:411:A:C5	1:CA:429:U:C5	2.95	0.55
46:BY:18:LEU:O	46:BY:22:LEU:HB2	2.06	0.55
22:DA:804:A:H2'	22:DA:806:C:C4	2.41	0.55
22:DA:2094:A:C2	22:DA:2196:C:C2	2.95	0.55
22:DA:2200:C:OP2	45:DX:37:ARG:NH2	2.39	0.55
26:DE:52:VAL:HG21	26:DE:81:GLY:CA	2.35	0.55
20:CT:78:ASN:O	20:CT:82:GLN:HG2	2.06	0.55
14:CN:52:PRO:O	14:CN:53:ARG:HB3	2.06	0.55
17:CQ:69:LYS:O	17:CQ:70:THR:CB	2.55	0.55
2:AB:147:SER:O	2:AB:148:LEU:HB2	2.07	0.55
22:BA:118:A:C8	22:BA:119:A:C8	2.94	0.55
32:BK:7:MET:SD	32:BK:20:MET:HB2	2.47	0.55
28:DG:105:LEU:HB2	28:DG:113:VAL:HB	1.88	0.55
30:BI:80:LEU:HD11	30:BI:133:ALA:HA	1.88	0.55
1:AA:417:G:C6	1:AA:418:C:C4	2.95	0.55
22:DA:2681:C:C2	22:DA:2724:U:O4	2.59	0.55
22:DA:2513:A:OP1	25:DD:160:LYS:NZ	2.39	0.55
22:DA:1985:C:C2	22:DA:1986:C:C5	2.95	0.55
1:AA:145:G:N2	1:AA:178:C:C2	2.75	0.55
2:AB:72:THR:O	2:AB:73:LYS:HG2	2.06	0.55
21:AU:36:GLU:O	21:AU:37:PHE:HB2	2.07	0.55
41:DT:62:VAL:HG12	41:DT:63:VAL:N	2.21	0.55
22:BA:846:U:O2'	22:BA:847:U:OP2	2.22	0.55
1:AA:1100:C:O2'	1:AA:1102:A:OP1	2.25	0.55
41:BT:16:VAL:O	41:BT:17:SER:HB3	2.07	0.55
22:DA:2788:C:H2'	22:DA:2789:C:C6	2.42	0.55
24:BC:29:PRO:HG2	24:BC:34:LEU:HD11	1.86	0.55
22:DA:622:G:H2'	22:DA:623:C:C6	2.40	0.55
22:DA:1082:U:H5'	30:DI:119:GLY:HA2	1.88	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AB:118:GLU:HA	2:AB:121:SER:HB2	1.89	0.55
3:CC:59:ARG:HB2	3:CC:63:SER:O	2.07	0.55
22:DA:1912:A:C8	22:DA:1918:A:C2	2.95	0.55
1:CA:1330:U:H4'	13:CM:23:TYR:CE2	2.41	0.55
10:CJ:81:GLU:HA	10:CJ:84:VAL:HG12	1.89	0.55
22:DA:1682:G:H2'	22:DA:1683:U:C6	2.41	0.55
22:DA:1807:G:C2	22:DA:1809:A:OP2	2.60	0.55
1:AA:934:C:H5''	57:AA:1767:HOH:O	2.06	0.55
29:DH:27:ARG:HE	45:DX:60:ASP:CB	2.19	0.55
4:AD:56:ARG:NH2	4:AD:59:GLN:HG2	2.22	0.55
12:CL:65:SER:HB2	12:CL:82:ILE:HD11	1.89	0.55
48:B0:11:SER:O	48:B0:15:MET:HG3	2.06	0.55
1:AA:654:G:H2'	1:AA:655:A:H5'	1.88	0.55
1:CA:984:C:N4	57:CA:1859:HOH:O	2.40	0.55
22:BA:1441:G:H2'	22:BA:1442:U:C6	2.41	0.55
1:CA:892:A:O2'	1:CA:1415:G:H4'	2.07	0.55
33:DL:116:VAL:HG21	33:DL:134:ALA:O	2.07	0.55
1:CA:157:U:H1'	1:CA:165:G:N2	2.21	0.55
3:AC:156:ARG:HD3	3:AC:193:TYR:O	2.07	0.55
1:AA:307:C:H5''	1:AA:308:C:OP2	2.07	0.55
17:CQ:62:ARG:C	17:CQ:73:TRP:CE3	2.80	0.55
22:DA:1476:U:C5	22:DA:1514:G:N2	2.74	0.55
41:DT:37:ASP:O	41:DT:38:ALA:HB3	2.06	0.55
24:DC:114:ASP:OD1	24:DC:114:ASP:N	2.38	0.55
28:BG:11:VAL:O	28:BG:11:VAL:HG23	2.06	0.55
5:CE:50:TYR:O	5:CE:51:GLY:O	2.25	0.55
22:BA:2786:U:O2'	25:BD:66:GLY:HA3	2.06	0.55
7:CG:25:LYS:O	7:CG:29:ILE:HG12	2.07	0.55
1:CA:36:C:OP1	12:CL:120:LYS:HE3	2.07	0.55
1:AA:685:G:C6	1:AA:686:U:O4	2.60	0.55
1:CA:577:G:C4	1:CA:578:C:C5	2.95	0.55
1:CA:734:G:C5	1:CA:735:C:C5	2.94	0.55
22:DA:1239:G:C6	22:DA:1240:U:C4	2.94	0.55
22:DA:1064:C:N3	22:DA:1074:G:N2	2.54	0.55
1:AA:737:C:H2'	1:AA:738:C:H6	1.72	0.55
2:CB:100:MET:HA	2:CB:107:VAL:HG21	1.89	0.55
22:DA:2819:G:H5''	57:DA:3805:HOH:O	2.07	0.55
22:BA:400:G:N7	45:BX:57:ARG:NH1	2.51	0.55
40:DS:14:ALA:HB1	40:DS:18:ARG:CZ	2.37	0.55
22:DA:1665:A:OP1	32:DK:66:LYS:CE	2.55	0.55
22:DA:2657:A:H1'	22:DA:2665:A:N6	2.22	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AL:74:LEU:HD21	12:AL:104:CYS:SG	2.46	0.55
20:CT:43:ASP:HB3	20:CT:46:ALA:HB3	1.88	0.55
35:BN:100:CYS:O	35:BN:100:CYS:SG	2.65	0.55
31:BJ:140:LEU:HD11	31:BJ:142:ILE:HD13	1.88	0.55
24:BC:125:LYS:HG2	24:BC:128:ASN:ND2	2.22	0.55
1:AA:1446:A:H2'	1:AA:1447:A:H5'	1.89	0.55
44:BW:68:LYS:HD3	44:BW:83:GLU:OE2	2.07	0.55
1:AA:455:G:N2	1:AA:478:A:C2	2.75	0.55
22:BA:1915:U:H2'	22:BA:1916:A:O4'	2.07	0.55
1:CA:1070:U:H2'	1:CA:1071:C:H6	1.72	0.55
22:DA:2234:G:C6	22:DA:2235:G:C5	2.95	0.55
11:CK:126:LYS:C	21:CU:34:ARG:NH2	2.60	0.55
38:BQ:76:TYR:OH	38:BQ:92:ARG:NH1	2.39	0.55
1:AA:202:G:N2	1:AA:216:U:O2	2.39	0.55
1:CA:1027:C:N4	1:CA:1034:G:C6	2.75	0.55
22:DA:792:A:H5''	22:DA:793:A:H5'	1.87	0.55
22:DA:2818:U:OP2	35:DN:42:LYS:NZ	2.36	0.55
27:DF:108:VAL:HG11	27:DF:176:PRO:HG2	1.88	0.55
16:AP:10:GLY:O	16:AP:11:ALA:HB2	2.07	0.55
29:BH:10:ALA:O	29:BH:12:LEU:N	2.40	0.55
19:AS:65:GLU:CD	19:AS:65:GLU:N	2.60	0.55
46:BY:13:GLU:HA	46:BY:13:GLU:OE2	2.06	0.55
22:BA:1314:C:H2'	22:BA:1314:C:O2	2.06	0.55
25:BD:4:LEU:HD23	25:BD:101:PHE:CE1	2.41	0.55
22:BA:1179:G:H3'	22:BA:1180:U:H4'	1.89	0.55
22:DA:2199:A:C5	22:DA:2225:A:C6	2.95	0.55
1:CA:676:A:C2	1:CA:677:U:C4	2.95	0.55
40:BS:63:GLY:O	40:BS:64:ALA:HB2	2.07	0.55
25:DD:101:PHE:HB3	25:DD:104:VAL:HG21	1.89	0.55
22:BA:1985:C:O2	22:BA:1985:C:C2'	2.49	0.55
1:AA:1144:G:N1	1:AA:1145:A:C2	2.75	0.55
1:AA:1145:A:O2'	1:AA:1146:A:O5'	2.22	0.55
46:BY:45:GLN:O	46:BY:46:VAL:CB	2.55	0.55
22:DA:305:C:C2	22:DA:313:G:N1	2.75	0.55
5:AE:98:PRO:O	5:AE:99:ALA:HB3	2.07	0.55
32:DK:92:GLU:O	32:DK:93:GLN:HB2	2.07	0.55
38:BQ:9:ILE:HG13	38:BQ:10:ALA:N	2.22	0.55
22:DA:1769:U:O2'	22:DA:1958:C:OP1	2.25	0.55
22:BA:2454:G:N7	57:BA:3529:HOH:O	2.33	0.55
7:CG:23:LEU:HD23	7:CG:26:PHE:HB3	1.87	0.55
9:CI:50:GLN:N	9:CI:51:PRO:HD2	2.22	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AL:25:GLU:O	12:AL:26:ALA:C	2.44	0.54
22:BA:364:C:H2'	22:BA:365:U:C6	2.41	0.54
22:DA:2235:G:C4	22:DA:2236:U:C6	2.95	0.54
22:DA:1335:C:N4	57:DA:3393:HOH:O	2.39	0.54
18:CR:20:GLU:HG3	18:CR:55:LEU:HD13	1.89	0.54
29:DH:31:VAL:HB	29:DH:32:PRO:HD3	1.89	0.54
1:CA:562:U:H1'	12:CL:12:ARG:HG3	1.88	0.54
1:CA:1201:A:H4'	1:CA:1202:U:O5'	2.07	0.54
31:DJ:6:ALA:O	31:DJ:7:LYS:HG3	2.07	0.54
35:BN:33:ILE:HG23	35:BN:33:ILE:O	2.07	0.54
3:CC:150:LYS:HE3	3:CC:201:TRP:CE3	2.42	0.54
1:AA:205:A:OP1	1:AA:205:A:H4'	2.07	0.54
22:BA:870:U:C2'	22:BA:871:U:H5'	2.37	0.54
22:DA:1665:A:H5''	32:DK:66:LYS:HG3	1.90	0.54
9:CI:30:ILE:HA	9:CI:65:ILE:O	2.07	0.54
19:AS:50:ALA:HB1	19:AS:57:HIS:HB3	1.88	0.54
13:AM:15:ALA:HB3	13:AM:34:LEU:HD21	1.89	0.54
39:DR:8:GLY:O	39:DR:10:LYS:NZ	2.40	0.54
49:D1:23:THR:OG1	49:D1:24:THR:N	2.40	0.54
22:BA:1973:G:OP1	57:BA:3471:HOH:O	2.18	0.54
31:BJ:84:ILE:HG23	31:BJ:84:ILE:O	2.07	0.54
22:BA:2728:U:O2'	22:BA:2729:G:OP2	2.20	0.54
16:CP:6:LEU:CD1	16:CP:71:VAL:CG2	2.85	0.54
4:AD:95:GLU:OE2	4:AD:100:ASN:ND2	2.40	0.54
22:DA:1993:U:H4'	25:DD:133:THR:HG22	1.90	0.54
1:CA:1169:A:H2'	1:CA:1170:A:C8	2.42	0.54
22:BA:2502:G:C5'	22:BA:2503:A:H5''	2.36	0.54
33:BL:87:GLY:O	33:BL:89:VAL:HG12	2.06	0.54
2:AB:72:THR:O	2:AB:73:LYS:CB	2.55	0.54
22:DA:443:A:N7	26:DE:40:ARG:CG	2.70	0.54
22:BA:142:A:H2'	22:BA:143:C:O4'	2.06	0.54
4:CD:168:PRO:HB3	4:CD:170:TRP:CH2	2.42	0.54
13:AM:20:THR:HA	13:AM:25:VAL:HG23	1.88	0.54
22:DA:2343:U:HO2'	22:DA:2373:G:HO2'	1.55	0.54
32:BK:113:MET:O	32:BK:116:ILE:HG13	2.06	0.54
22:DA:1917:U:H2'	22:DA:1918:A:H5'	1.90	0.54
21:AU:10:GLU:CG	21:AU:11:PRO:HD3	2.38	0.54
22:BA:1080:A:C2	22:BA:1081:U:C5	2.96	0.54
1:AA:895:G:H2'	1:AA:896:C:C6	2.43	0.54
22:DA:2834:G:H2'	22:DA:2879:A:N6	2.22	0.54
25:BD:151:THR:HG22	25:BD:152:PRO:HD3	1.89	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:1438:U:C5	22:DA:1552:A:C2	2.95	0.54
28:DG:98:VAL:HG23	28:DG:125:CYS:SG	2.47	0.54
11:AK:23:ILE:HD11	11:AK:86:VAL:HG13	1.87	0.54
24:DC:260:ASN:O	24:DC:261:LYS:HB2	2.06	0.54
22:BA:374:A:C2	22:BA:401:A:C4	2.95	0.54
22:BA:1812:U:O2	22:BA:1812:U:H2'	2.06	0.54
1:AA:257:G:C2	1:AA:258:G:C5	2.94	0.54
8:CH:9:ASP:OD2	8:CH:13:ARG:NH1	2.40	0.54
29:BH:90:LEU:CD2	29:BH:93:SER:HA	2.37	0.54
22:DA:1606:C:O2'	22:DA:1607:C:P	2.65	0.54
22:DA:192:C:C5	22:DA:193:U:C2	2.95	0.54
45:DX:31:PRO:HB2	45:DX:33:LEU:HD13	1.89	0.54
1:AA:858:G:O2'	1:AA:859:G:H5'	2.06	0.54
4:AD:124:MET:HG3	4:AD:146:ARG:HG2	1.90	0.54
22:DA:699:A:H2'	22:DA:700:G:H5'	1.89	0.54
42:DU:59:VAL:HG12	42:DU:61:LYS:HD3	1.90	0.54
22:BA:2291:U:H2'	22:BA:2292:U:C6	2.42	0.54
22:DA:969:G:H2'	22:DA:970:U:C6	2.42	0.54
22:BA:31:C:O3'	22:BA:1238:G:H5''	2.08	0.54
17:AQ:81:LYS:HD3	17:AQ:81:LYS:N	2.22	0.54
26:DE:128:ALA:O	26:DE:130:LYS:N	2.40	0.54
10:AJ:8:ILE:HA	10:AJ:99:GLN:O	2.07	0.54
1:AA:1418:A:C2	1:AA:1483:A:C2	2.96	0.54
30:DI:101:ILE:HG22	30:DI:105:GLN:HB2	1.88	0.54
7:CG:145:ALA:O	7:CG:146:GLU:HB2	2.07	0.54
16:AP:78:VAL:O	16:AP:78:VAL:HG13	2.06	0.54
14:AN:49:GLN:OE1	14:AN:49:GLN:HA	2.07	0.54
34:BM:55:ARG:CZ	34:BM:55:ARG:HB3	2.36	0.54
27:BF:108:VAL:CG1	27:BF:114:PHE:CZ	2.90	0.54
1:AA:397:A:C6	1:AA:548:G:N7	2.75	0.54
29:BH:90:LEU:HA	29:BH:125:THR:HG23	1.90	0.54
29:BH:83:LYS:CE	1:CA:55:A:O2'	2.56	0.54
22:BA:1340:U:OP1	41:BT:19:LYS:NZ	2.34	0.54
22:BA:686:U:O4	50:B2:12:ARG:HB2	2.07	0.54
22:DA:2164:C:H2'	22:DA:2165:C:C5	2.43	0.54
22:DA:2232:C:OP1	45:DX:27:ARG:NH1	2.40	0.54
7:CG:93:PRO:O	7:CG:97:ASN:ND2	2.40	0.54
22:DA:2773:C:H2'	22:DA:2774:C:C6	2.42	0.54
22:DA:1914:C:C6	22:DA:1915:U:C6	2.96	0.54
34:BM:77:PRO:HG2	34:BM:80:VAL:HG21	1.89	0.54
20:AT:25:ARG:HG2	20:AT:29:ARG:NH1	2.23	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:AU:14:VAL:HG13	21:AU:16:LEU:HG	1.88	0.54
22:DA:13:A:C6	22:DA:525:U:C2	2.96	0.54
22:DA:801:G:C8	26:DE:49:ARG:HG3	2.43	0.54
22:DA:1713:A:C6	22:DA:1716:U:H1'	2.42	0.54
22:DA:636:G:N1	33:DL:76:GLU:OE2	2.37	0.54
6:CF:99:ALA:O	6:CF:100:SER:CB	2.55	0.54
4:CD:166:GLU:O	4:CD:167:LYS:HB2	2.06	0.54
5:AE:50:TYR:CE1	5:AE:134:ILE:HD11	2.42	0.54
29:DH:79:THR:HA	29:DH:145:ASN:HB2	1.89	0.54
22:BA:345:A:N3	22:BA:346:A:N6	2.52	0.54
29:BH:103:VAL:HG21	29:BH:132:PHE:CE1	2.42	0.54
22:DA:1208:C:N3	22:DA:1209:U:C5	2.75	0.54
29:BH:14:SER:OG	29:BH:17:ASP:CG	2.46	0.54
13:AM:25:VAL:HG12	13:AM:29:ARG:HB3	1.90	0.54
22:DA:1027:A:C6	22:DA:1126:A:C4	2.94	0.54
15:CO:37:ASN:O	15:CO:40:GLN:CB	2.56	0.54
24:BC:123:ALA:O	24:BC:128:ASN:ND2	2.40	0.54
34:BM:132:THR:HG22	34:BM:133:LYS:N	2.23	0.54
36:BO:10:ARG:NH2	36:BO:96:GLY:O	2.41	0.54
1:CA:41:G:H2'	1:CA:42:G:C8	2.43	0.54
22:DA:2369:A:N6	22:DA:2382:G:O6	2.40	0.54
5:CE:12:GLN:HB3	5:CE:40:GLY:O	2.08	0.54
22:BA:682:G:H5'	50:B2:26:ASN:OD1	2.06	0.54
10:CJ:63:ASP:OD2	14:CN:85:ARG:NH1	2.40	0.54
27:DF:64:LYS:HG2	27:DF:64:LYS:O	2.07	0.54
11:AK:17:SER:HA	11:AK:79:ILE:HA	1.90	0.54
1:AA:666:G:C5	1:AA:741:G:C6	2.96	0.54
22:DA:2056:G:OP2	57:DA:3487:HOH:O	2.18	0.54
22:DA:1809:A:H2'	22:DA:1810:A:C8	2.43	0.54
22:BA:2191:A:C6	22:BA:2192:U:C4	2.96	0.54
42:BU:16:GLY:O	42:BU:17:LYS:C	2.45	0.54
22:DA:2550:G:C5	22:DA:2551:C:C5	2.95	0.54
12:CL:65:SER:OG	12:CL:97:THR:HG23	2.07	0.54
22:DA:1259:G:H2'	22:DA:1260:A:C8	2.43	0.54
1:AA:872:A:C8	1:AA:874:G:C8	2.96	0.54
7:AG:75:VAL:HG21	7:AG:144:MET:HG2	1.88	0.54
1:CA:1125:U:C5	10:CJ:40:ILE:HG21	2.43	0.54
48:B0:54:VAL:O	48:B0:56:ALA:N	2.40	0.54
22:BA:1373:A:H5''	22:BA:1374:G:OP2	2.08	0.54
22:BA:693:A:O2'	22:BA:694:U:H5'	2.08	0.54
22:DA:1940:U:C2	22:DA:1965:C:OP2	2.61	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:DE:5:LEU:HD23	26:DE:122:GLU:CD	2.27	0.54
22:BA:1002:G:N7	57:BA:3746:HOH:O	2.33	0.54
1:AA:949:A:C6	1:AA:950:U:N3	2.76	0.54
46:DY:1:MET:N	46:DY:4:LYS:HD3	2.23	0.54
4:AD:10:LYS:HA	4:AD:13:ARG:HG3	1.89	0.54
23:BB:51:G:H2'	23:BB:52:A:C8	2.42	0.54
1:AA:81:A:H2'	1:AA:82:G:H5'	1.89	0.54
22:BA:1114:C:O2'	22:BA:1115:G:H5'	2.08	0.54
22:BA:1411:U:H2'	22:BA:1412:U:O4'	2.08	0.54
35:BN:24:MET:HG2	35:BN:44:LEU:HD22	1.89	0.54
29:BH:77:THR:CG2	29:BH:77:THR:O	2.56	0.54
22:BA:995:C:H5'	22:BA:995:C:H6	1.73	0.54
1:AA:1493:A:OP2	1:AA:1493:A:C8	2.61	0.54
1:AA:990:C:C4	1:AA:991:U:O4	2.61	0.54
51:D3:16:LYS:HE2	51:D3:20:GLY:HA2	1.89	0.54
22:DA:856:G:N2	22:DA:922:C:C2	2.76	0.54
22:DA:2603:G:C6	22:DA:2604:U:C4	2.96	0.54
22:DA:1869:G:C3'	22:DA:1870:C:H5'	2.37	0.54
2:AB:167:ASP:OD1	2:AB:168:HIS:N	2.41	0.54
25:DD:149:ASN:OD1	25:DD:150:GLN:N	2.41	0.54
12:AL:22:PRO:C	12:AL:24:LEU:H	2.11	0.54
22:BA:1917:U:N3	22:BA:1918:A:C4	2.76	0.54
1:AA:451:A:N6	1:AA:481:G:H5''	2.23	0.54
1:AA:451:A:H61	1:AA:481:G:H5''	1.73	0.54
38:DQ:27:ALA:HB1	38:DQ:31:VAL:CG2	2.37	0.54
22:DA:2209:G:N2	22:DA:2216:G:N3	2.55	0.54
24:BC:230:HIS:C	24:BC:232:HIS:H	2.11	0.54
12:AL:114:ARG:HB3	12:AL:119:VAL:HB	1.89	0.54
22:DA:2346:A:H3'	22:DA:2347:C:H5'	1.89	0.54
1:CA:682:G:C2	1:CA:683:G:N7	2.75	0.54
33:DL:77:ILE:O	33:DL:110:VAL:O	2.26	0.54
1:CA:992:U:C4	1:CA:1043:G:C8	2.96	0.54
22:BA:639:U:H2'	22:BA:640:C:C6	2.43	0.54
1:CA:1474:U:H2'	1:CA:1475:G:H5''	1.90	0.54
22:DA:1000:A:N1	22:DA:1001:A:C2	2.76	0.54
44:BW:41:ARG:NH1	44:BW:41:ARG:HG3	2.22	0.54
24:DC:51:THR:CG2	24:DC:54:ILE:HD11	2.38	0.54
3:CC:77:ILE:HA	3:CC:84:VAL:HG23	1.88	0.54
14:CN:54:ASP:OD1	14:CN:59:ARG:NH1	2.40	0.54
22:DA:1809:A:N6	22:DA:1810:A:C6	2.76	0.54
22:DA:1808:A:N1	45:DX:28:ARG:HD2	2.23	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AK:127:ARG:N	21:AU:34:ARG:CZ	2.69	0.54
4:CD:30:THR:O	4:CD:31:LYS:HD3	2.06	0.54
33:BL:81:ASP:HB3	33:BL:100:ILE:HD13	1.89	0.54
1:AA:872:A:C4	1:AA:874:G:C8	2.96	0.54
1:CA:718:A:H5'	11:CK:119:ASN:CG	2.28	0.54
1:AA:71:A:H3'	1:AA:71:A:OP2	2.07	0.54
1:CA:1068:G:C2'	1:CA:1069:C:H5'	2.37	0.54
22:DA:1856:U:O4	22:DA:1857:G:N1	2.40	0.54
35:DN:12:ARG:O	35:DN:17:ARG:NH2	2.41	0.54
16:CP:20:VAL:HG21	16:CP:32:PHE:CB	2.37	0.54
22:BA:2014:A:H2'	22:BA:2015:A:C8	2.43	0.54
1:CA:76:G:N2	1:CA:95:C:N3	2.55	0.54
51:D3:6:THR:O	51:D3:8:ARG:HG2	2.08	0.54
1:CA:298:A:H2'	1:CA:299:G:O4'	2.08	0.54
22:BA:84:A:H4'	22:BA:85:G:O5'	2.08	0.54
1:CA:463:U:H3'	1:CA:464:U:C6	2.42	0.54
2:AB:21:ARG:NH1	2:AB:21:ARG:HA	2.23	0.54
1:AA:1526:G:OP1	21:AU:39:GLU:HB2	2.08	0.54
1:CA:1513:A:H2'	1:CA:1514:G:C8	2.43	0.54
22:BA:613:A:O2'	22:BA:614:A:OP1	2.26	0.54
2:AB:83:ALA:HA	2:AB:86:SER:OG	2.08	0.54
1:AA:1377:A:N3	7:AG:2:PRO:HG3	2.23	0.54
22:BA:480:A:OP2	42:BU:44:LYS:HD2	2.08	0.54
1:CA:686:U:O2'	1:CA:687:A:OP2	2.20	0.54
1:CA:327:A:C2	1:CA:329:A:C4	2.95	0.54
1:CA:978:A:H4'	1:CA:1322:C:C5	2.43	0.54
22:DA:1571:A:H2'	22:DA:1572:A:C8	2.43	0.54
1:CA:511:C:C2	1:CA:512:U:C6	2.96	0.54
1:AA:96:U:O2'	1:AA:97:G:P	2.65	0.54
30:DI:22:PRO:HB2	30:DI:23:PRO:HD3	1.90	0.54
30:BI:11:LEU:HD12	30:BI:24:VAL:HG12	1.90	0.54
21:AU:10:GLU:CD	21:AU:11:PRO:HD3	2.28	0.54
36:BO:7:ARG:CG	36:BO:96:GLY:HA3	2.37	0.54
1:AA:949:A:C6	1:AA:950:U:C2	2.96	0.54
22:DA:500:G:N2	22:DA:502:A:C8	2.75	0.54
22:DA:2406:A:H2'	22:DA:2406:A:OP2	2.07	0.54
9:AI:50:GLN:N	9:AI:51:PRO:HD2	2.23	0.54
41:DT:44:LYS:O	41:DT:48:GLN:HG2	2.08	0.54
22:BA:585:G:N7	38:BQ:6:ARG:NH1	2.55	0.54
22:DA:532:A:N7	22:DA:2021:C:H2'	2.23	0.54
22:BA:666:A:H4'	33:BL:48:ARG:HD3	1.90	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:AT:80:THR:O	20:AT:83:ILE:HG13	2.07	0.54
4:AD:90:LEU:HD23	4:AD:200:ILE:HD11	1.89	0.54
7:CG:5:ARG:HA	7:CG:5:ARG:NE	2.22	0.54
22:DA:2531:A:H5'	28:DG:157:TYR:CZ	2.43	0.54
22:BA:570:G:O6	57:BA:3693:HOH:O	2.17	0.54
39:BR:51:VAL:HB	39:BR:52:PRO:HD2	1.90	0.54
22:BA:1090:A:H2'	22:BA:1091:G:H5'	1.89	0.54
1:CA:577:G:C8	1:CA:816:A:C6	2.95	0.54
1:CA:1106:G:C4	1:CA:1107:C:C5	2.96	0.54
22:DA:1469:A:C2	22:DA:1470:A:C6	2.96	0.54
22:DA:2199:A:N7	22:DA:2225:A:C6	2.76	0.54
22:BA:142:A:H2'	22:BA:143:C:C6	2.43	0.54
14:AN:7:LYS:O	14:AN:10:GLU:N	2.40	0.54
1:AA:49:U:O4	1:AA:365:U:H5	1.91	0.54
1:CA:1298:U:O2	1:CA:1298:U:H2'	2.06	0.54
22:DA:223:A:C4	22:DA:408:G:H1'	2.43	0.54
42:DU:34:VAL:HG22	42:DU:65:ILE:O	2.08	0.54
24:BC:33:LEU:O	24:BC:64:ILE:HG13	2.08	0.54
5:AE:149:SER:OG	5:AE:152:MET:HB2	2.07	0.54
12:AL:86:ARG:NE	12:AL:88:LYS:HB3	2.23	0.54
22:BA:1824:G:C5	22:BA:1825:U:C5	2.96	0.54
22:DA:2370:G:C6	22:DA:2371:G:C5	2.96	0.54
10:CJ:26:VAL:HG13	10:CJ:36:VAL:HG11	1.90	0.54
6:CF:97:THR:O	6:CF:98:GLU:HB3	2.08	0.54
13:AM:3:ARG:O	13:AM:4:ILE:HG12	2.08	0.54
22:DA:45:G:H2'	22:DA:215:G:N7	2.22	0.54
1:AA:232:G:H2'	1:AA:233:C:O4'	2.07	0.54
2:AB:126:PHE:N	2:AB:126:PHE:HD1	2.06	0.54
22:DA:1867:G:O6	22:DA:1875:G:N2	2.40	0.54
1:CA:227:G:H2'	1:CA:228:A:O4'	2.07	0.54
1:CA:951:G:C2	1:CA:1231:G:C2	2.96	0.54
1:AA:728:A:C6	1:AA:729:A:C6	2.96	0.54
42:BU:99:ASN:O	42:BU:101:GLU:N	2.41	0.54
22:DA:1389:G:C2	22:DA:1390:U:C2	2.96	0.53
22:BA:2748:A:C1'	28:BG:67:THR:HG22	2.36	0.53
22:DA:1034:G:C6	22:DA:1035:U:C2	2.97	0.53
5:AE:82:GLN:HG2	5:AE:150:PRO:HB3	1.89	0.53
22:DA:301:G:N2	22:DA:302:C:C2	2.76	0.53
22:DA:741:U:O2'	22:DA:1676:A:OP1	2.18	0.53
1:AA:1195:C:H2'	1:AA:1197:A:O4'	2.07	0.53
1:CA:679:C:O2	1:CA:712:A:C2	2.61	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:AU:12:PHE:HD1	21:AU:12:PHE:N	2.05	0.53
22:DA:222:A:H3'	22:DA:421:C:H5'	1.90	0.53
1:CA:1320:C:N3	19:CS:36:ARG:NH1	2.56	0.53
6:CF:88:MET:SD	6:CF:90:MET:SD	3.06	0.53
22:BA:492:A:H2'	22:BA:493:G:O4'	2.08	0.53
1:CA:102:G:C2	1:CA:103:U:C4	2.95	0.53
1:AA:792:A:H1'	1:AA:794:A:N7	2.24	0.53
26:BE:77:ILE:HG22	26:BE:77:ILE:O	2.08	0.53
1:AA:624:C:C5	1:AA:625:U:C5	2.95	0.53
22:BA:2557:G:H2'	22:BA:2558:C:C6	2.43	0.53
22:DA:2032:G:H1'	25:DD:150:GLN:OE1	2.08	0.53
22:BA:1922:G:N2	22:BA:1923:U:C1'	2.71	0.53
22:DA:1340:U:C4	22:DA:1603:A:C8	2.96	0.53
22:DA:1345:C:C2	22:DA:1346:G:C8	2.96	0.53
22:DA:2816:G:O3'	35:DN:99:LYS:HE2	2.08	0.53
5:CE:82:GLN:H	5:CE:147:MET:HE3	1.73	0.53
22:DA:1431:A:C2	22:DA:1432:G:C4	2.96	0.53
1:CA:257:G:C2	1:CA:270:A:N1	2.76	0.53
1:AA:792:A:H4'	1:AA:793:U:O5'	2.08	0.53
22:BA:533:G:H5'	38:BQ:24:TYR:CD1	2.43	0.53
1:CA:1277:C:H2'	1:CA:1278:G:H5''	1.90	0.53
51:D3:62:LEU:HB3	51:D3:65:ALA:HB2	1.89	0.53
1:AA:579:A:H2'	1:AA:580:C:H6	1.74	0.53
1:CA:427:U:OP1	4:CD:13:ARG:NH2	2.41	0.53
7:AG:15:ASP:OD1	7:AG:44:TYR:OH	2.26	0.53
22:DA:2268:A:OP1	57:DA:3509:HOH:O	2.18	0.53
36:DO:92:PHE:HB2	36:DO:117:PHE:CD1	2.43	0.53
22:DA:289:G:H2'	22:DA:290:U:O4'	2.08	0.53
2:CB:54:LEU:HA	2:CB:57:LEU:HB3	1.90	0.53
4:AD:148:LYS:O	4:AD:149:ALA:C	2.46	0.53
29:BH:94:ILE:CA	29:BH:122:LEU:HB3	2.35	0.53
50:D2:25:LYS:O	50:D2:29:GLN:HG3	2.08	0.53
1:AA:188:C:O2	1:AA:188:C:H2'	2.07	0.53
1:CA:782:A:H2'	1:CA:783:C:H5'	1.89	0.53
1:AA:448:A:C5	1:AA:487:A:C2	2.96	0.53
22:DA:1139:G:O2'	22:DA:1140:C:H5'	2.09	0.53
8:AH:113:ASP:OD2	8:AH:117:ARG:NH2	2.41	0.53
11:AK:34:ILE:HG12	11:AK:70:CYS:SG	2.48	0.53
1:CA:620:C:H2'	1:CA:621:A:O4'	2.09	0.53
1:CA:976:G:OP2	1:CA:1358:U:O2'	2.24	0.53
2:CB:163:VAL:HG23	2:CB:185:ALA:HB2	1.89	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:1001:A:OP2	57:BA:3743:HOH:O	2.18	0.53
1:CA:706:A:N7	1:CA:707:U:C5	2.76	0.53
12:CL:88:LYS:O	12:CL:88:LYS:HG3	2.07	0.53
5:AE:78:ASN:OD1	5:AE:79:GLY:N	2.40	0.53
2:CB:126:PHE:CZ	2:CB:127:ASP:OD1	2.60	0.53
22:BA:1917:U:C4	22:BA:1918:A:C4	2.97	0.53
22:BA:1482:G:C6	22:BA:1508:A:N1	2.76	0.53
22:DA:1826:G:C4	22:DA:1827:U:C5	2.96	0.53
22:DA:188:G:O2'	22:DA:1365:A:N6	2.41	0.53
1:AA:939:G:H4'	7:AG:102:ARG:NH2	2.24	0.53
1:CA:579:A:H2'	1:CA:580:C:H6	1.74	0.53
1:CA:580:C:H2'	1:CA:581:G:C8	2.44	0.53
22:DA:685:A:C8	22:DA:774:G:C6	2.96	0.53
1:CA:734:G:C4	1:CA:735:C:C5	2.96	0.53
1:CA:728:A:C2	1:CA:729:A:C5	2.96	0.53
1:AA:1157:A:H5'	1:AA:1158:C:C6	2.43	0.53
30:DI:58:VAL:HG12	30:DI:59:ILE:N	2.22	0.53
10:AJ:28:THR:HG22	10:AJ:86:ALA:HB1	1.90	0.53
22:DA:920:A:OP1	47:DZ:19:LYS:HE3	2.09	0.53
22:BA:1907:G:C6	22:BA:1908:C:N3	2.77	0.53
22:DA:522:A:C6	22:DA:523:C:N3	2.76	0.53
22:DA:2567:G:H2'	22:DA:2568:U:C6	2.43	0.53
13:AM:40:ALA:HB3	13:AM:43:VAL:HG13	1.89	0.53
28:BG:154:PRO:HD3	28:BG:162:VAL:O	2.09	0.53
22:BA:1971:U:OP2	22:BA:1971:U:H4'	2.09	0.53
22:BA:2193:G:O2'	22:BA:2194:U:H5'	2.09	0.53
24:BC:17:VAL:H	24:BC:204:VAL:HG22	1.72	0.53
22:BA:2129:C:OP1	53:B5:35:THR:HG23	2.08	0.53
1:AA:1358:U:C6	1:AA:1359:C:C5	2.97	0.53
29:DH:32:PRO:O	29:DH:33:GLN:HB2	2.09	0.53
1:AA:201:G:C2	1:AA:217:C:O2	2.62	0.53
1:CA:1190:G:H5'	3:CC:176:HIS:CE1	2.43	0.53
22:DA:575:A:C2	22:DA:576:U:C5	2.96	0.53
1:AA:1055:A:N6	1:AA:1206:G:C5	2.76	0.53
22:DA:1833:C:O2	22:DA:1833:C:H2'	2.07	0.53
22:BA:1056:G:O2'	22:BA:1086:A:H1'	2.08	0.53
22:DA:828:U:HO2'	22:DA:829:A:C5'	2.21	0.53
1:AA:1339:A:H2'	1:AA:1340:A:O4'	2.08	0.53
22:BA:1185:G:H5''	22:BA:1186:G:P	2.49	0.53
1:AA:792:A:N3	1:AA:794:A:C5	2.76	0.53
16:CP:16:PHE:CE1	16:CP:38:PHE:HB2	2.44	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CF:52:ASN:O	6:CF:53:LYS:HB2	2.08	0.53
22:BA:1095:A:C6	22:BA:1096:A:N6	2.76	0.53
34:BM:12:MET:HE3	34:BM:71:LYS:HG3	1.90	0.53
21:CU:12:PHE:O	21:CU:13:ASP:CB	2.57	0.53
4:AD:50:ASP:OD1	4:AD:54:GLN:NE2	2.41	0.53
31:DJ:40:HIS:O	38:DQ:67:ALA:HB1	2.08	0.53
1:CA:68:G:C5	1:CA:69:G:H1'	2.43	0.53
24:DC:159:GLY:N	24:DC:197:ASN:O	2.41	0.53
26:DE:170:ARG:CG	26:DE:174:GLY:O	2.56	0.53
22:DA:953:G:O2'	22:DA:954:G:H5'	2.09	0.53
2:AB:15:HIS:O	2:AB:16:PHE:C	2.47	0.53
17:AQ:51:ASN:N	17:AQ:51:ASN:OD1	2.42	0.53
32:DK:71:ARG:HB3	32:DK:72:PRO:HD2	1.91	0.53
29:BH:147:VAL:CG1	29:BH:149:GLU:HG3	2.36	0.53
22:BA:192:C:OP1	57:BA:3749:HOH:O	2.19	0.53
22:BA:621:A:OP2	33:BL:99:ASN:ND2	2.41	0.53
22:BA:1925:C:H5''	22:BA:1926:U:C4	2.41	0.53
22:BA:1925:C:H4'	22:BA:1926:U:C5	2.44	0.53
22:BA:1070:A:O2'	22:BA:1097:U:OP1	2.17	0.53
22:DA:2148:G:C2	22:DA:2149:U:C4	2.96	0.53
22:DA:1809:A:C5	22:DA:1810:A:C5	2.97	0.53
1:AA:451:A:C8	1:AA:452:A:N1	2.76	0.53
11:AK:76:GLU:HA	22:BA:2141:G:OP2	2.08	0.53
1:AA:958:A:C6	1:AA:959:A:N1	2.77	0.53
45:DX:2:SER:O	45:DX:4:VAL:N	2.40	0.53
22:DA:1352:U:C5	22:DA:1377:G:C6	2.97	0.53
26:DE:52:VAL:HB	26:DE:74:LYS:HD3	1.91	0.53
27:BF:173:PHE:O	27:BF:174:ASP:HB3	2.09	0.53
35:BN:2:ARG:HA	35:BN:5:LYS:HD2	1.90	0.53
22:DA:1245:G:H4'	26:DE:33:VAL:CG1	2.39	0.53
1:CA:1237:C:C5	1:CA:1336:C:N4	2.77	0.53
22:BA:2469:A:H4'	34:BM:55:ARG:HH12	1.73	0.53
1:AA:1149:C:OP2	9:AI:11:ARG:NH2	2.42	0.53
31:BJ:30:THR:HG22	31:BJ:31:GLU:N	2.24	0.53
33:DL:82:LEU:HA	33:DL:85:VAL:HG13	1.90	0.53
33:BL:14:LYS:HD3	33:BL:15:ALA:HB3	1.89	0.53
6:AF:12:PRO:O	6:AF:15:SER:HB2	2.09	0.53
22:BA:1487:U:C2	22:BA:1503:A:C2	2.97	0.53
2:CB:183:VAL:N	2:CB:197:ASP:OD2	2.41	0.53
29:DH:2:GLN:O	29:DH:3:VAL:HG22	2.09	0.53
29:DH:103:VAL:HA	29:DH:106:ALA:HB3	1.89	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:CK:43:GLY:HA3	11:CK:74:VAL:HG12	1.91	0.53
31:DJ:15:TRP:O	31:DJ:137:PRO:HA	2.09	0.53
43:DV:42:LEU:HD12	43:DV:47:VAL:HG21	1.89	0.53
22:DA:2454:G:H1'	57:DA:3533:HOH:O	2.08	0.53
1:AA:451:A:H4'	1:AA:452:A:O5'	2.08	0.53
22:DA:705:A:C2	22:DA:727:A:H1'	2.43	0.53
46:BY:6:LEU:HD13	46:BY:56:LEU:HD22	1.90	0.53
24:DC:258:ARG:NH2	24:DC:264:ASP:OD1	2.41	0.53
22:DA:785:G:O2'	22:DA:1779:U:H5'	2.09	0.53
1:AA:1053:G:N7	1:AA:1200:C:H5''	2.24	0.53
1:AA:1198:G:H5''	57:AA:1835:HOH:O	2.08	0.53
21:CU:4:ILE:HG22	21:CU:4:ILE:O	2.09	0.53
28:DG:39:ASP:HB3	28:DG:58:TYR:OH	2.07	0.53
22:BA:804:A:H5''	22:BA:805:G:OP1	2.09	0.53
27:BF:121:SER:HB2	27:BF:128:TYR:CE1	2.44	0.53
5:CE:96:MET:HE3	5:CE:111:MET:CE	2.39	0.53
37:DP:103:ARG:HB3	37:DP:108:ALA:HB2	1.90	0.53
51:B3:15:LYS:HD2	51:B3:23:LYS:HE2	1.90	0.53
1:CA:1081:A:H5'	5:CE:23:LYS:HG3	1.89	0.53
1:AA:994:A:N3	1:AA:994:A:H2'	2.24	0.53
1:AA:439:U:C5	1:AA:440:C:C5	2.97	0.53
47:BZ:35:THR:CG2	47:BZ:36:VAL:N	2.71	0.53
2:AB:76:ALA:O	2:AB:80:VAL:HG23	2.09	0.53
11:AK:76:GLU:C	22:BA:2141:G:OP1	2.47	0.53
22:BA:574:A:H4'	22:BA:575:A:O5'	2.07	0.53
22:DA:1645:G:H4'	22:DA:1646:C:C6	2.44	0.53
22:DA:690:G:H1'	22:DA:779:U:O3'	2.09	0.53
5:AE:81:LEU:CD2	5:AE:123:VAL:HG12	2.39	0.53
1:CA:938:A:C2	1:CA:1345:U:O4	2.62	0.53
1:AA:188:C:O2	1:AA:189:A:C4	2.62	0.53
22:BA:2065:C:H2'	22:BA:2066:C:H6	1.74	0.53
1:CA:899:C:OP1	1:CA:899:C:H6	1.91	0.53
42:DU:47:LYS:HG3	42:DU:48:PRO:HD2	1.91	0.53
1:AA:1062:U:H2'	1:AA:1063:C:C6	2.44	0.53
16:CP:20:VAL:CG2	16:CP:32:PHE:HB2	2.39	0.53
6:AF:29:ILE:HD13	6:AF:64:VAL:HG11	1.91	0.53
27:DF:108:VAL:N	27:DF:109:PRO:CD	2.72	0.53
22:DA:532:A:N3	22:DA:532:A:H2'	2.24	0.53
22:DA:224:U:OP2	22:DA:408:G:N2	2.42	0.53
1:CA:228:A:H4'	16:CP:63:GLN:HG2	1.91	0.53
22:DA:919:U:H2'	22:DA:920:A:O4'	2.09	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:2079:U:H2'	22:DA:2080:A:O4'	2.09	0.53
1:CA:518:C:H4'	1:CA:519:C:O5'	2.08	0.53
47:DZ:24:LEU:HD22	47:DZ:29:LEU:HD12	1.91	0.53
22:BA:1439:A:C2	22:BA:1553:A:C5	2.97	0.53
1:AA:938:A:N3	1:AA:1376:U:O2'	2.39	0.53
29:DH:37:VAL:CG2	29:DH:38:PRO:HD2	2.39	0.53
22:DA:1728:C:C2	22:DA:1730:C:O2	2.62	0.53
22:BA:2883:A:OP2	48:B0:50:ARG:NH1	2.42	0.53
11:CK:52:PHE:CE1	11:CK:62:ALA:HB1	2.44	0.53
25:DD:176:ASP:O	25:DD:190:LYS:N	2.39	0.53
26:DE:109:LEU:O	26:DE:112:LEU:HB2	2.08	0.53
1:CA:1070:U:C2	1:CA:1071:C:C5	2.97	0.53
22:DA:2147:A:C8	22:DA:2148:G:C8	2.97	0.53
50:D2:44:VAL:O	50:D2:45:SER:CB	2.56	0.53
16:CP:6:LEU:HD12	16:CP:71:VAL:CG2	2.38	0.53
29:DH:32:PRO:HB3	45:DX:39:TRP:HB3	1.91	0.53
12:CL:116:LYS:O	12:CL:117:TYR:CB	2.57	0.53
22:DA:563:A:C6	22:DA:2018:G:C4	2.97	0.53
22:BA:674:G:N3	26:BE:69:ARG:NH1	2.57	0.53
22:DA:526:A:OP1	57:DA:3244:HOH:O	2.18	0.53
24:DC:61:ALA:O	24:DC:63:ARG:NH2	2.42	0.53
22:DA:696:G:C6	22:DA:767:U:O2	2.62	0.53
1:AA:1417:G:C6	1:AA:1482:G:C6	2.96	0.53
22:DA:2502:G:H5'	22:DA:2503:A:C5'	2.39	0.53
22:DA:2415:G:C2	22:DA:2416:C:C2	2.97	0.53
1:AA:1285:A:O2'	1:AA:1286:U:OP2	2.26	0.53
16:AP:72:ALA:O	16:AP:75:ILE:HG13	2.08	0.53
1:AA:557:G:C6	1:AA:558:G:N1	2.77	0.53
22:BA:2750:A:H1'	22:BA:2752:C:N4	2.23	0.53
25:DD:51:THR:OG1	25:DD:76:GLY:HA3	2.09	0.53
7:CG:66:LEU:O	7:CG:66:LEU:HG	2.09	0.53
42:DU:74:ASN:HA	42:DU:96:PHE:CZ	2.43	0.53
22:BA:1205:A:H4'	22:BA:1206:G:OP2	2.09	0.53
22:BA:1206:G:C6	22:BA:1207:C:C4	2.97	0.53
22:BA:852:U:H2'	22:BA:853:C:C6	2.44	0.53
22:BA:2425:A:H5''	22:BA:2427:C:O4'	2.09	0.53
1:CA:1012:A:N6	1:CA:1013:G:O6	2.42	0.53
41:BT:64:LYS:HA	41:BT:79:ASP:OD1	2.09	0.53
12:AL:66:TYR:CD2	12:AL:87:VAL:HG21	2.44	0.53
29:DH:40:THR:O	29:DH:41:LYS:C	2.48	0.53
1:AA:554:A:C5'	12:AL:26:ALA:HB1	2.39	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:187:G:C2	22:DA:210:C:O2	2.61	0.53
22:DA:910:A:C2	34:DM:13:HIS:CE1	2.97	0.53
25:BD:99:GLU:HG2	25:BD:182:ALA:HB2	1.90	0.53
42:DU:33:LYS:HE2	42:DU:66:GLN:NE2	2.24	0.53
22:DA:696:G:N1	22:DA:767:U:C2	2.77	0.53
39:BR:47:VAL:CG1	39:BR:54:VAL:CG1	2.87	0.53
22:DA:142:A:H2'	22:DA:143:C:C6	2.44	0.53
28:DG:98:VAL:HG21	28:DG:124:GLU:HA	1.91	0.53
22:DA:2438:U:O2'	22:DA:2440:C:OP1	2.24	0.53
22:DA:2441:U:OP1	22:DA:2441:U:H4'	2.09	0.53
22:BA:2480:C:C2'	22:BA:2481:G:H5'	2.39	0.53
22:DA:2365:G:H4'	44:DW:60:PHE:CE2	2.44	0.53
43:BV:10:LYS:CE	43:BV:10:LYS:H	2.21	0.53
22:DA:476:G:N2	22:DA:479:A:C8	2.77	0.53
22:DA:1483:G:C6	22:DA:1484:U:C4	2.96	0.53
22:BA:497:A:C4	22:BA:498:G:C8	2.97	0.53
42:DU:13:VAL:CG1	42:DU:18:ASP:O	2.57	0.53
22:DA:2734:A:N6	22:DA:2770:G:O2'	2.42	0.53
22:DA:1747:U:H2'	22:DA:1748:C:C6	2.44	0.53
9:AI:97:GLU:OE2	9:AI:97:GLU:N	2.42	0.53
27:BF:40:VAL:HG11	27:BF:50:LEU:HD13	1.91	0.53
22:BA:608:A:C6	22:BA:609:A:C6	2.96	0.53
1:AA:173:U:C6	1:AA:197:A:C2	2.97	0.53
22:DA:2750:A:O2'	22:DA:2752:C:N4	2.42	0.53
22:DA:1680:U:H2'	22:DA:1681:G:O4'	2.08	0.53
8:AH:8:ALA:HB2	8:AH:77:ARG:CD	2.38	0.53
1:CA:1160:G:O6	1:CA:1181:G:C6	2.62	0.52
22:BA:1069:A:O2'	22:BA:1070:A:H5''	2.08	0.52
29:DH:31:VAL:CB	29:DH:32:PRO:CD	2.86	0.52
22:DA:1651:G:N2	22:DA:2007:U:C2	2.77	0.52
22:DA:79:C:C4	22:DA:80:G:N7	2.77	0.52
42:BU:72:ILE:HD12	42:BU:83:VAL:HG23	1.91	0.52
22:BA:1870:C:H2'	22:BA:1871:A:C2	2.44	0.52
23:DB:39:A:H2'	23:DB:40:U:C5	2.43	0.52
5:AE:151:GLU:HG3	5:AE:152:MET:SD	2.49	0.52
10:CJ:35:GLN:O	10:CJ:36:VAL:HB	2.09	0.52
1:CA:519:C:H2'	1:CA:520:A:O4'	2.08	0.52
1:CA:154:U:C2'	1:CA:155:A:H5'	2.39	0.52
9:AI:13:LYS:HG2	9:AI:13:LYS:O	2.08	0.52
9:AI:16:ALA:O	9:AI:67:VAL:HA	2.09	0.52
30:DI:10:LYS:HB2	30:DI:56:PRO:HB3	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:DN:22:ARG:HG3	35:DN:70:THR:HA	1.90	0.52
14:AN:21:PHE:HA	14:AN:25:ALA:HB3	1.91	0.52
20:CT:58:VAL:HG13	20:CT:72:ALA:HB1	1.92	0.52
12:CL:51:LYS:HD2	12:CL:51:LYS:N	2.24	0.52
22:DA:1497:U:C5	22:DA:1578:U:O5'	2.62	0.52
22:DA:2899:A:H2'	22:DA:2900:A:C8	2.44	0.52
25:DD:112:THR:O	25:DD:195:GLY:HA2	2.09	0.52
26:DE:32:VAL:HG21	33:DL:6:LEU:HD13	1.91	0.52
1:CA:1328:C:H5''	13:CM:28:THR:HG21	1.92	0.52
1:AA:903:G:C5	1:AA:904:U:C5	2.97	0.52
43:BV:80:HIS:CE1	43:BV:83:LYS:HD2	2.43	0.52
38:BQ:88:VAL:HG13	39:BR:49:ILE:HD11	1.92	0.52
22:BA:1098:A:H5'	22:BA:1099:G:OP2	2.09	0.52
1:AA:970:C:H5''	1:AA:971:G:OP1	2.09	0.52
1:CA:1211:U:C2'	1:CA:1212:U:OP2	2.56	0.52
22:DA:1544:A:C6	22:DA:1545:A:N1	2.77	0.52
5:CE:15:LEU:C	5:CE:16:ILE:HD12	2.28	0.52
22:BA:674:G:O2'	26:BE:69:ARG:HB3	2.09	0.52
22:DA:777:G:C2	22:DA:778:G:C8	2.97	0.52
12:CL:80:ILE:HD12	12:CL:97:THR:HG21	1.91	0.52
22:DA:1835:G:C5	22:DA:1836:C:C5	2.96	0.52
22:DA:1835:G:C2	22:DA:1836:C:C6	2.97	0.52
22:DA:1432:G:C2	22:DA:1433:A:C4	2.98	0.52
11:CK:87:LYS:HA	11:CK:114:THR:HG22	1.91	0.52
13:CM:40:ALA:C	13:CM:41:GLU:HG3	2.30	0.52
19:CS:4:SER:O	19:CS:5:LEU:CB	2.57	0.52
22:DA:2772:C:H2'	22:DA:2773:C:H6	1.74	0.52
15:CO:56:LEU:O	15:CO:59:MET:N	2.42	0.52
21:AU:12:PHE:N	21:AU:12:PHE:CD1	2.75	0.52
36:BO:7:ARG:HG3	36:BO:96:GLY:HA3	1.91	0.52
29:BH:77:THR:HA	29:BH:143:ILE:O	2.09	0.52
5:CE:96:MET:HE3	5:CE:111:MET:HE3	1.91	0.52
25:DD:7:LYS:HG3	25:DD:198:GLY:O	2.09	0.52
7:CG:18:PHE:CE1	7:CG:58:GLU:HG2	2.44	0.52
2:AB:54:LEU:HD12	2:AB:220:THR:HG21	1.91	0.52
3:AC:206:GLU:O	3:AC:207:ILE:O	2.28	0.52
7:AG:31:MET:CG	7:AG:32:VAL:N	2.71	0.52
22:BA:2811:G:OP1	25:BD:62:LYS:HB2	2.09	0.52
25:DD:35:THR:HG22	25:DD:73:VAL:HG21	1.91	0.52
1:AA:269:C:H2'	1:AA:270:A:C8	2.44	0.52
1:AA:21:G:N2	1:AA:22:G:C6	2.77	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:BH:89:LYS:HG2	1:CA:359:G:OP1	2.09	0.52
22:DA:370:G:C6	22:DA:424:G:N7	2.77	0.52
22:BA:301:G:C4	22:BA:302:C:C5	2.97	0.52
22:DA:2234:G:C4	22:DA:2235:G:C8	2.96	0.52
5:AE:25:VAL:O	5:AE:28:GLY:N	2.32	0.52
22:DA:25:U:O2	22:DA:515:A:N6	2.42	0.52
1:AA:971:G:O6	1:AA:1364:U:O2'	2.27	0.52
1:AA:827:U:C4	1:AA:870:U:C2	2.98	0.52
39:BR:47:VAL:HG12	39:BR:54:VAL:CG2	2.40	0.52
36:BO:51:ALA:O	36:BO:74:VAL:HG13	2.10	0.52
1:CA:785:G:C2	1:CA:786:G:C8	2.98	0.52
22:DA:2343:U:H2'	22:DA:2344:U:C6	2.44	0.52
12:AL:44:LYS:CB	12:AL:45:PRO:CD	2.87	0.52
22:DA:125:A:H3'	50:D2:19:ARG:HG3	1.92	0.52
15:CO:42:HIS:O	15:CO:45:GLU:O	2.27	0.52
1:AA:36:C:OP1	12:AL:120:LYS:HE3	2.09	0.52
22:DA:2273:A:H2'	22:DA:2274:A:C8	2.45	0.52
1:CA:151:A:H2'	1:CA:152:A:O4'	2.10	0.52
1:AA:595:A:C5	1:AA:641:U:C5	2.97	0.52
46:DY:18:LEU:O	46:DY:22:LEU:CB	2.56	0.52
47:DZ:14:ILE:HG22	47:DZ:15:GLY:N	2.23	0.52
22:DA:931:U:O4	22:DA:1166:G:N2	2.42	0.52
22:DA:1534:U:O2'	22:DA:1537:G:O6	2.28	0.52
1:AA:76:G:N2	1:AA:95:C:C2	2.77	0.52
1:AA:125:U:H2'	1:AA:126:G:O4'	2.08	0.52
45:DX:74:ARG:NH2	45:DX:76:GLU:OE2	2.40	0.52
22:BA:2325:G:C6	22:BA:2326:C:N4	2.77	0.52
22:BA:675:A:OP1	26:BE:58:LYS:HE2	2.08	0.52
28:DG:123:ALA:HB2	28:DG:133:LEU:HB3	1.91	0.52
24:BC:91:ILE:HD12	24:BC:103:TYR:CD1	2.45	0.52
1:AA:11:G:C6	1:AA:12:U:C4	2.98	0.52
17:CQ:12:VAL:HG12	17:CQ:13:VAL:N	2.23	0.52
22:DA:1809:A:C6	22:DA:1810:A:C5	2.98	0.52
25:DD:13:ARG:HD3	25:DD:21:SER:OG	2.08	0.52
22:BA:2097:A:C2	22:BA:2193:G:C6	2.97	0.52
1:CA:384:G:H2'	1:CA:385:C:C6	2.45	0.52
1:AA:914:A:N3	1:AA:915:A:C8	2.78	0.52
22:DA:89:A:C6	22:DA:90:U:C4	2.97	0.52
22:DA:2226:C:H2'	22:DA:2227:A:O4'	2.09	0.52
1:CA:671:G:N1	1:CA:672:U:C2	2.77	0.52
50:B2:43:THR:O	50:B2:44:VAL:CB	2.57	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:1417:C:H2'	22:BA:1418:G:O4'	2.09	0.52
22:BA:1794:A:H2'	22:BA:1795:C:H6	1.73	0.52
22:DA:1275:A:N7	35:DN:16:HIS:CG	2.78	0.52
22:DA:649:G:H2'	22:DA:650:C:C6	2.44	0.52
2:AB:147:SER:O	2:AB:148:LEU:CB	2.56	0.52
1:AA:731:G:O2'	1:AA:732:C:H5'	2.08	0.52
22:DA:1917:U:C2'	22:DA:1918:A:H5'	2.40	0.52
22:DA:987:C:O2'	22:DA:1000:A:N3	2.39	0.52
22:BA:665:U:O2'	22:BA:666:A:H5'	2.09	0.52
5:AE:74:VAL:O	5:AE:74:VAL:HG23	2.09	0.52
7:AG:27:VAL:O	7:AG:31:MET:N	2.40	0.52
35:BN:78:LYS:C	35:BN:79:LEU:O	2.45	0.52
1:AA:157:U:O2'	1:AA:158:G:H5'	2.09	0.52
22:DA:971:G:O2'	22:DA:983:A:N3	2.39	0.52
12:AL:38:TYR:O	12:AL:39:THR:HG22	2.09	0.52
19:CS:58:VAL:HG11	19:CS:75:ALA:HA	1.90	0.52
12:AL:41:THR:HG22	12:AL:49:LEU:HD12	1.92	0.52
1:AA:1306:A:C4	1:AA:1307:U:C6	2.98	0.52
24:DC:252:THR:O	24:DC:253:LYS:HB2	2.09	0.52
43:DV:30:ILE:HD11	43:DV:63:ILE:HD12	1.92	0.52
4:AD:98:LEU:HD23	4:AD:118:VAL:CG1	2.38	0.52
1:AA:443:C:C2	1:AA:444:G:C8	2.98	0.52
22:BA:1161:C:H1'	39:BR:8:GLY:O	2.09	0.52
22:DA:1344:U:O2'	22:DA:1345:C:P	2.68	0.52
1:AA:701:U:O2	1:AA:703:G:C2	2.63	0.52
22:DA:410:G:C2	22:DA:2407:A:C5	2.98	0.52
1:CA:1007:U:H2'	1:CA:1008:U:H5''	1.91	0.52
5:AE:101:GLU:HB3	5:AE:122:ASN:CB	2.38	0.52
1:CA:764:C:C2	1:CA:765:G:C8	2.98	0.52
1:AA:1053:G:C4'	1:AA:1054:C:H5'	2.40	0.52
1:AA:1101:A:N7	2:AB:171:ILE:HG23	2.24	0.52
21:AU:25:LYS:O	21:AU:29:LEU:HB2	2.09	0.52
29:DH:34:GLY:O	29:DH:35:LYS:HD2	2.10	0.52
1:AA:1124:G:C2'	1:AA:1145:A:N6	2.73	0.52
46:BY:42:LEU:O	46:BY:45:GLN:O	2.28	0.52
30:DI:90:SER:HB3	30:DI:93:PRO:HG3	1.92	0.52
5:CE:35:ALA:O	5:CE:50:TYR:O	2.27	0.52
20:CT:43:ASP:HB3	20:CT:46:ALA:CB	2.40	0.52
2:AB:126:PHE:N	2:AB:126:PHE:CD1	2.76	0.52
42:DU:13:VAL:HG21	42:DU:39:ILE:CG2	2.40	0.52
22:DA:819:A:C8	22:DA:1188:U:O4	2.62	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AL:41:THR:HG22	12:AL:48:ALA:O	2.10	0.52
6:CF:6:ILE:HG22	6:CF:7:VAL:N	2.25	0.52
29:BH:51:ARG:NH1	29:BH:55:GLU:OE1	2.43	0.52
1:AA:657:U:O2'	1:AA:658:C:H5'	2.09	0.52
32:BK:78:ARG:NH1	37:BP:71:GLU:OE2	2.40	0.52
13:AM:75:MET:SD	27:BF:112:ARG:HD3	2.50	0.52
19:AS:36:ARG:NH2	19:AS:75:ALA:O	2.42	0.52
2:AB:113:ARG:O	2:AB:117:LEU:HB2	2.10	0.52
3:CC:177:THR:HG22	3:CC:179:ARG:HG3	1.92	0.52
1:CA:371:A:H1'	1:CA:482:A:H1'	1.91	0.52
41:DT:51:PHE:C	41:DT:52:GLU:HG2	2.30	0.52
14:AN:79:LEU:HB2	14:AN:84:VAL:HG23	1.91	0.52
22:DA:77:G:OP1	46:DY:52:ARG:HD3	2.07	0.52
29:DH:72:ILE:HG22	29:DH:72:ILE:O	2.09	0.52
29:BH:94:ILE:C	29:BH:122:LEU:CG	2.77	0.52
39:BR:49:ILE:HG22	39:BR:52:PRO:HA	1.90	0.52
22:BA:1917:U:C4	22:BA:1918:A:C6	2.97	0.52
22:DA:1361:G:C6	22:DA:1371:G:C2	2.97	0.52
22:DA:811:U:O2	22:DA:1251:C:C5	2.63	0.52
22:DA:445:C:O2'	22:DA:449:A:N3	2.41	0.52
22:BA:2191:A:C2	22:BA:2192:U:N3	2.77	0.52
1:CA:373:A:O2'	1:CA:374:A:H5'	2.09	0.52
1:CA:448:A:N6	1:CA:449:G:C2	2.78	0.52
22:DA:82:U:H5'	22:DA:296:U:H5''	1.90	0.52
22:DA:197:A:C8	22:DA:2430:A:N7	2.77	0.52
5:CE:15:LEU:C	5:CE:15:LEU:CD1	2.78	0.52
22:DA:2093:G:C6	22:DA:2225:A:N7	2.78	0.52
22:DA:2225:A:H4'	22:DA:2226:C:O5'	2.09	0.52
1:AA:502:A:OP1	12:AL:115:SER:HB3	2.09	0.52
22:BA:142:A:N7	22:BA:143:C:C4	2.77	0.52
1:AA:596:A:C6	1:AA:645:G:C2	2.98	0.52
22:DA:734:A:N7	22:DA:735:A:N7	2.58	0.52
1:CA:610:U:C4	1:CA:611:C:C4	2.97	0.52
13:AM:15:ALA:CB	13:AM:34:LEU:HD21	2.39	0.52
21:AU:14:VAL:HG13	21:AU:16:LEU:CD2	2.38	0.52
1:CA:1513:A:H2'	1:CA:1514:G:H8	1.75	0.52
12:AL:39:THR:OG1	12:AL:39:THR:O	2.28	0.52
13:CM:107:ARG:NH2	13:CM:111:GLY:O	2.42	0.52
22:BA:1696:G:C6	22:BA:1697:G:C4	2.98	0.52
12:AL:29:GLN:HB2	12:AL:82:ILE:O	2.09	0.52
22:DA:1754:A:N6	22:DA:1755:A:C6	2.78	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:947:G:C2	1:AA:948:C:C2	2.97	0.52
22:BA:858:G:H3'	22:BA:859:G:C8	2.44	0.52
34:BM:17:ASN:O	34:BM:38:ARG:HD3	2.10	0.52
22:DA:1464:G:H2'	22:DA:1465:G:C8	2.44	0.52
22:DA:1190:G:OP1	33:DL:32:GLY:HA2	2.09	0.52
22:DA:46:G:C2	22:DA:47:C:C5	2.97	0.52
22:BA:864:G:C6	22:BA:865:C:N4	2.77	0.52
22:BA:1680:U:H2'	22:BA:1681:G:O4'	2.09	0.52
22:DA:2596:U:C5	22:DA:2597:G:C5	2.98	0.52
1:AA:617:G:C2	1:AA:618:C:C5	2.98	0.52
1:AA:774:G:N2	1:AA:806:C:C2	2.78	0.52
29:BH:94:ILE:HD12	29:BH:98:ASP:HB3	1.92	0.52
22:DA:186:G:N1	22:DA:211:C:N3	2.58	0.52
22:DA:190:A:N6	22:DA:191:A:N1	2.58	0.52
1:CA:527:G:N1	1:CA:528:C:C5	2.78	0.52
1:AA:957:U:C2	1:AA:959:A:OP2	2.63	0.52
24:DC:230:HIS:CD2	24:DC:247:PRO:HB3	2.45	0.52
22:DA:1580:A:H3'	22:DA:1581:G:O4'	2.10	0.52
1:CA:717:U:O2'	1:CA:734:G:O4'	2.26	0.52
11:AK:25:ALA:O	11:AK:88:GLY:HA3	2.10	0.52
22:DA:1835:G:C4	22:DA:1836:C:C6	2.98	0.52
53:B5:52:PRO:O	53:B5:53:ARG:CB	2.58	0.52
23:DB:29:A:H2'	23:DB:30:C:C6	2.44	0.52
11:CK:112:ASP:HB3	21:CU:4:ILE:CG2	2.40	0.52
1:AA:690:G:H2'	1:AA:691:G:C8	2.45	0.52
1:AA:1492:A:OP1	12:AL:44:LYS:HA	2.10	0.52
29:DH:34:GLY:O	29:DH:35:LYS:HB2	2.08	0.52
1:CA:604:G:C2	1:CA:635:A:C2	2.98	0.52
1:AA:429:U:H4'	1:AA:430:A:OP1	2.08	0.52
1:CA:160:A:H1'	1:CA:344:A:N7	2.25	0.52
22:BA:65:U:H2'	22:BA:66:C:H6	1.73	0.52
10:AJ:52:LEU:HB3	14:AN:81:ARG:NE	2.24	0.52
25:BD:151:THR:HG22	25:BD:152:PRO:CD	2.39	0.52
37:DP:103:ARG:HG2	37:DP:107:ALA:HB1	1.91	0.52
22:BA:864:G:O2'	22:BA:865:C:H5'	2.10	0.52
1:AA:463:U:H3'	1:AA:464:U:C6	2.44	0.52
41:DT:82:LYS:HG2	41:DT:83:ALA:N	2.25	0.52
1:AA:394:G:C5	1:AA:395:C:C5	2.98	0.52
22:DA:2624:G:H1'	48:D0:19:HIS:CE1	2.44	0.52
11:AK:63:ALA:CB	11:AK:92:GLY:HA3	2.40	0.52
9:CI:95:ARG:HA	9:CI:98:LEU:HB3	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:DT:6:ARG:O	41:DT:10:VAL:HG23	2.09	0.52
24:DC:2:ALA:N	24:DC:199:GLU:OE2	2.42	0.52
1:CA:1499:A:OP2	57:CA:1877:HOH:O	2.18	0.52
1:AA:254:G:OP1	17:AQ:70:THR:CG2	2.57	0.52
7:AG:135:VAL:HB	7:AG:138:ARG:NH2	2.24	0.52
46:BY:56:LEU:O	46:BY:57:LEU:HB3	2.09	0.52
4:CD:32:CYS:O	4:CD:33:LYS:HB2	2.10	0.52
22:BA:182:A:C6	22:BA:183:C:C4	2.98	0.52
32:BK:113:MET:SD	32:BK:116:ILE:HD11	2.49	0.52
22:DA:2357:G:H5'	22:DA:2358:A:OP2	2.09	0.52
1:AA:204:G:H2'	1:AA:205:A:O4'	2.10	0.52
22:DA:37:C:H2'	22:DA:38:A:C8	2.44	0.52
38:BQ:9:ILE:O	38:BQ:13:ARG:HG3	2.09	0.52
11:CK:51:GLY:O	11:CK:52:PHE:O	2.28	0.52
34:DM:56:ALA:C	34:DM:58:LYS:H	2.13	0.52
1:AA:1372:U:H2'	1:AA:1373:G:O4'	2.10	0.52
9:CI:26:GLY:H	9:CI:59:GLU:HA	1.75	0.52
3:CC:141:ALA:O	3:CC:146:ALA:HB3	2.09	0.52
22:BA:1731:G:C6	22:BA:1733:G:C5	2.98	0.52
29:DH:25:TYR:CZ	29:DH:30:LEU:HD21	2.45	0.52
1:CA:1364:U:O2	1:CA:1364:U:C2'	2.57	0.52
22:DA:1087:G:N1	22:DA:1089:A:C2	2.77	0.52
1:AA:1129:C:O2	1:AA:1130:A:N6	2.42	0.52
1:AA:697:U:C5	1:AA:698:G:C8	2.97	0.52
15:AO:3:LEU:HB2	15:AO:35:GLN:HG2	1.91	0.52
22:DA:1559:U:H4'	22:DA:1560:G:OP2	2.10	0.52
22:DA:2790:U:H5'	22:DA:2893:A:N7	2.25	0.52
22:BA:1667:G:O2'	22:BA:1991:U:O4	2.22	0.52
38:BQ:88:VAL:HG13	39:BR:49:ILE:CD1	2.40	0.52
4:AD:95:GLU:HG2	4:AD:186:PRO:HG2	1.92	0.52
22:DA:582:A:N7	57:DA:3284:HOH:O	2.34	0.52
21:CU:19:PHE:HD1	21:CU:19:PHE:O	1.93	0.52
22:DA:1969:A:O2'	22:DA:1972:G:N3	2.39	0.52
1:CA:1362:A:H4'	1:CA:1362:A:OP1	2.08	0.52
22:DA:590:A:C6	22:DA:591:U:C4	2.98	0.52
2:CB:20:THR:O	2:CB:21:ARG:CZ	2.57	0.52
17:CQ:16:LYS:C	17:CQ:17:MET:SD	2.88	0.52
1:CA:755:G:C2	1:CA:756:C:C6	2.98	0.52
6:AF:93:LYS:O	6:AF:94:HIS:HB2	2.09	0.52
3:AC:156:ARG:NH1	3:AC:193:TYR:O	2.43	0.52
1:CA:154:U:H2'	1:CA:155:A:H5'	1.92	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:AD:118:VAL:HA	4:AD:123:ILE:HD12	1.91	0.52
1:AA:1095:U:H2'	1:AA:1096:C:O4'	2.10	0.52
3:CC:42:TYR:CE2	3:CC:90:VAL:HG21	2.45	0.52
29:BH:2:GLN:O	29:BH:3:VAL:HG22	2.10	0.52
4:AD:17:THR:CG2	4:AD:18:ASP:N	2.73	0.52
33:BL:9:ALA:O	33:BL:12:SER:OG	2.28	0.52
22:BA:1283:G:N2	22:BA:1285:A:H3'	2.25	0.52
22:DA:2557:G:H2'	22:DA:2558:C:C6	2.45	0.52
14:AN:20:TYR:CE1	14:AN:52:PRO:HG2	2.45	0.52
22:DA:664:G:O2'	22:DA:941:A:OP1	2.19	0.52
32:DK:21:CYS:SG	32:DK:39:ILE:HB	2.49	0.52
22:BA:1028:A:N6	22:BA:1125:G:H2'	2.24	0.52
16:AP:22:ALA:HB2	16:AP:32:PHE:HA	1.91	0.52
43:BV:2:PHE:HB2	43:BV:61:LEU:HD12	1.92	0.52
35:BN:65:LEU:HD11	35:BN:69:ARG:HH21	1.74	0.52
1:AA:977:A:H8	1:AA:1223:C:C2	2.28	0.52
31:BJ:55:ILE:HG21	31:BJ:130:HIS:CD2	2.45	0.52
22:DA:1378:A:C2'	22:DA:1380:G:N7	2.73	0.52
5:AE:25:VAL:O	5:AE:26:LYS:C	2.48	0.52
22:DA:132:G:N2	22:DA:148:U:O2	2.43	0.52
22:DA:183:C:C4	22:DA:184:C:C5	2.98	0.52
22:DA:2043:C:O2	22:DA:2043:C:H2'	2.10	0.52
1:CA:32:A:OP1	1:CA:398:U:H1'	2.09	0.52
22:DA:2199:A:C4	22:DA:2225:A:C2	2.98	0.52
20:AT:6:SER:OG	20:AT:7:ALA:N	2.43	0.52
1:CA:734:G:C6	1:CA:735:C:C4	2.98	0.52
1:CA:1411:C:H2'	1:CA:1412:C:C6	2.45	0.52
37:DP:60:THR:HG23	37:DP:73:VAL:HG12	1.91	0.52
1:AA:1285:A:H5''	1:AA:1286:U:C4	2.45	0.52
27:BF:107:ALA:O	27:BF:110:ARG:N	2.43	0.52
22:DA:2037:A:C6	22:DA:2038:G:C6	2.98	0.52
39:DR:81:LYS:O	39:DR:82:HIS:C	2.49	0.52
1:CA:1149:C:N4	1:CA:1150:A:C6	2.77	0.52
22:DA:2805:C:H2'	22:DA:2806:C:C6	2.45	0.52
2:AB:181:ILE:O	2:AB:183:VAL:HG23	2.10	0.52
48:B0:43:ILE:HG22	48:B0:49:TYR:HB2	1.92	0.52
24:BC:258:ARG:HD2	24:BC:270:ARG:NH2	2.25	0.52
1:CA:49:U:O4	1:CA:365:U:H5	1.93	0.52
1:CA:748:G:C2	1:CA:749:A:C5	2.98	0.52
9:CI:25:ASN:O	9:CI:62:ASP:HA	2.10	0.52
12:CL:3:THR:HB	12:CL:6:GLN:HG3	1.92	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
50:B2:6:GLN:HA	50:B2:6:GLN:OE1	2.09	0.52
1:AA:137:U:H2'	1:AA:137:U:O2	2.09	0.52
23:DB:42:C:C5	27:DF:66:LEU:HD22	2.45	0.52
29:BH:91:PHE:HB3	1:CA:55:A:N3	2.25	0.51
22:DA:58:G:N3	22:DA:70:G:N2	2.58	0.51
1:CA:505:G:H2'	1:CA:506:G:H8	1.74	0.51
4:CD:31:LYS:HD3	4:CD:31:LYS:N	2.26	0.51
38:BQ:115:ALA:O	38:BQ:117:LEU:N	2.43	0.51
1:CA:1377:A:C5	7:CG:7:ILE:HD12	2.44	0.51
22:BA:1386:C:H2'	22:BA:1387:A:C8	2.45	0.51
22:DA:1304:A:N6	22:DA:1305:C:N4	2.58	0.51
44:BW:69:PHE:CE2	44:BW:80:ILE:HD11	2.45	0.51
1:CA:1387:G:H2'	1:CA:1388:C:C6	2.45	0.51
32:BK:4:GLU:O	32:BK:5:GLN:CB	2.58	0.51
22:DA:1275:A:C8	35:DN:16:HIS:CG	2.98	0.51
28:BG:30:ASN:CG	28:BG:30:ASN:O	2.49	0.51
34:DM:2:LEU:O	34:DM:3:GLN:HB3	2.09	0.51
6:CF:2:ARG:HB3	6:CF:4:TYR:CZ	2.45	0.51
2:CB:154:MET:SD	2:CB:158:PRO:HD3	2.50	0.51
52:B4:36:ARG:HG2	52:B4:37:GLN:N	2.24	0.51
53:B5:42:VAL:HG12	53:B5:214:TYR:HA	1.92	0.51
30:BI:101:ILE:CG1	30:BI:138:LEU:HD13	2.40	0.51
22:BA:2223:G:OP1	24:BC:171:TYR:OH	2.22	0.51
17:CQ:46:VAL:HG11	17:CQ:61:ILE:HG13	1.93	0.51
7:CG:75:VAL:HG11	7:CG:144:MET:HG3	1.92	0.51
22:DA:1490:A:H2'	22:DA:1490:A:N3	2.24	0.51
12:AL:51:LYS:HD2	12:AL:51:LYS:N	2.25	0.51
22:DA:1494:A:H2'	22:DA:1495:A:C8	2.44	0.51
4:AD:138:SER:HB2	4:AD:139:PRO:HD2	1.92	0.51
1:AA:1405:G:H1'	1:AA:1519:A:O4'	2.10	0.51
17:AQ:4:LYS:HG3	17:AQ:7:THR:HG23	1.91	0.51
39:BR:53:PHE:CD1	39:BR:53:PHE:N	2.72	0.51
29:BH:83:LYS:HE2	1:CA:55:A:H2'	1.93	0.51
1:CA:780:A:C2	1:CA:803:G:N1	2.78	0.51
22:BA:198:C:P	57:BA:3767:HOH:O	2.67	0.51
22:DA:2043:C:C2	22:DA:2044:C:C5	2.98	0.51
12:CL:116:LYS:O	12:CL:117:TYR:HB2	2.10	0.51
1:CA:32:A:N3	1:CA:32:A:H2'	2.24	0.51
22:DA:600:G:C5'	26:DE:27:LEU:HD22	2.40	0.51
22:DA:1351:C:OP2	57:DA:3398:HOH:O	2.19	0.51
17:CQ:17:MET:CE	17:CQ:20:SER:O	2.58	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:294:A:C5	22:DA:345:A:C2	2.97	0.51
23:BB:47:C:OP2	36:BO:3:LYS:HE3	2.11	0.51
35:BN:117:ASP:O	35:BN:118:ARG:C	2.49	0.51
22:DA:586:A:C2	22:DA:810:U:O4'	2.63	0.51
6:AF:68:GLN:HA	6:AF:71:ILE:HG21	1.92	0.51
17:CQ:45:HIS:HB2	17:CQ:70:THR:HG22	1.93	0.51
6:CF:1:MET:SD	6:CF:67:PRO:HD3	2.50	0.51
22:BA:569:U:H1'	22:BA:947:A:O4'	2.10	0.51
39:DR:52:PRO:O	39:DR:53:PHE:CB	2.57	0.51
22:BA:1441:G:H2'	22:BA:1442:U:H6	1.75	0.51
4:AD:188:ARG:NH2	4:AD:197:GLU:OE1	2.43	0.51
5:CE:96:MET:CE	5:CE:111:MET:CE	2.89	0.51
27:BF:40:VAL:O	27:BF:42:GLU:N	2.44	0.51
4:AD:138:SER:HB2	4:AD:139:PRO:CD	2.39	0.51
22:DA:1786:A:OP1	57:DA:3443:HOH:O	2.19	0.51
22:DA:2722:G:C2	22:DA:2723:C:O2	2.63	0.51
1:CA:1090:U:H2'	1:CA:1091:U:C6	2.45	0.51
12:AL:3:THR:HB	12:AL:6:GLN:HG3	1.92	0.51
35:BN:14:SER:HA	35:BN:17:ARG:NH1	2.25	0.51
31:BJ:118:MET:HA	31:BJ:121:LYS:NZ	2.24	0.51
8:CH:126:ILE:HG22	8:CH:127:CYS:SG	2.50	0.51
28:DG:111:HIS:O	28:DG:111:HIS:ND1	2.44	0.51
57:DA:3249:HOH:O	31:DJ:111:LYS:HE2	2.10	0.51
6:CF:43:GLY:HA2	6:CF:58:HIS:CD2	2.45	0.51
29:BH:99:ILE:O	29:BH:103:VAL:CG2	2.58	0.51
22:DA:1372:U:H2'	22:DA:1373:A:C8	2.46	0.51
1:CA:1124:G:O2'	1:CA:1127:G:O6	2.29	0.51
22:DA:2697:G:C2	22:DA:2711:A:C2	2.98	0.51
35:BN:12:ARG:HE	35:BN:20:MET:HE3	1.75	0.51
22:DA:2217:G:C5	22:DA:2218:G:N7	2.79	0.51
4:AD:3:ARG:CZ	4:AD:115:ARG:CD	2.86	0.51
1:CA:1361:G:H2'	1:CA:1362:A:H5''	1.92	0.51
1:AA:1277:C:O2'	1:AA:1279:G:H8	1.91	0.51
22:DA:740:C:H5'	22:DA:1784:A:C3'	2.39	0.51
22:DA:1737:G:C6	22:DA:1738:G:C6	2.98	0.51
29:BH:110:VAL:HG22	29:BH:114:GLU:HB2	1.90	0.51
28:BG:38:ASN:O	28:BG:39:ASP:CB	2.58	0.51
21:AU:29:LEU:HD23	21:AU:29:LEU:O	2.10	0.51
3:CC:184:TYR:HE1	3:CC:201:TRP:CZ2	2.29	0.51
1:CA:680:C:C2	1:CA:711:G:N2	2.78	0.51
42:BU:99:ASN:OD1	42:BU:99:ASN:C	2.48	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:CK:31:ILE:O	11:CK:31:ILE:HG12	2.09	0.51
1:AA:1091:U:O2	1:AA:1095:U:C2	2.64	0.51
1:CA:1089:G:C5	1:CA:1090:U:C5	2.98	0.51
22:DA:134:G:C2	22:DA:146:A:C2	2.99	0.51
8:AH:66:PHE:CD2	8:AH:67:GLN:HG2	2.46	0.51
7:AG:89:VAL:HG22	7:AG:90:GLU:N	2.26	0.51
1:CA:1271:A:H2'	1:CA:1272:G:C8	2.46	0.51
36:BO:87:ILE:HG22	36:BO:88:LYS:N	2.24	0.51
1:CA:66:A:H4'	1:CA:173:U:C5	2.45	0.51
4:AD:168:PRO:HG2	4:AD:171:LEU:HD11	1.91	0.51
44:DW:34:GLY:N	44:DW:61:ALA:O	2.36	0.51
1:AA:844:G:N2	1:AA:846:G:H4'	2.25	0.51
22:DA:2250:G:N2	34:DM:82:MET:HG3	2.25	0.51
9:AI:114:LYS:NZ	9:AI:118:LEU:O	2.43	0.51
2:AB:64:LYS:HA	2:AB:64:LYS:HE2	1.92	0.51
2:AB:108:ARG:O	2:AB:111:ILE:HB	2.10	0.51
22:DA:2493:U:O2'	34:DM:78:LEU:HD21	2.11	0.51
22:BA:2331:G:O2'	22:BA:2336:A:N1	2.32	0.51
16:AP:3:THR:HG22	16:AP:4:ILE:N	2.25	0.51
22:DA:2594:C:N4	22:DA:2595:G:O6	2.42	0.51
26:BE:119:ILE:CG2	26:BE:187:VAL:HG22	2.40	0.51
1:AA:1321:U:C4	1:AA:1322:C:N4	2.79	0.51
29:DH:121:VAL:O	29:DH:122:LEU:HB2	2.11	0.51
22:DA:152:A:C2	22:DA:175:G:C2	2.99	0.51
22:BA:2192:U:N3	22:BA:2193:G:C8	2.79	0.51
1:CA:374:A:C2	1:CA:375:U:C6	2.98	0.51
4:AD:95:GLU:OE2	4:AD:104:ARG:NH1	2.43	0.51
29:DH:23:ALA:O	29:DH:27:ARG:N	2.38	0.51
2:CB:131:LYS:HA	2:CB:134:ALA:HB3	1.92	0.51
1:AA:144:G:C5	1:AA:179:A:C2	2.99	0.51
22:DA:771:G:C6	22:DA:772:C:C5	2.99	0.51
33:DL:93:ASN:OD1	33:DL:94:THR:N	2.44	0.51
1:AA:1337:G:H5''	1:AA:1338:G:OP1	2.11	0.51
1:CA:243:A:C5	1:CA:281:G:N2	2.79	0.51
1:AA:455:G:C2	1:AA:478:A:N1	2.78	0.51
7:CG:145:ALA:O	7:CG:146:GLU:CB	2.59	0.51
22:DA:1754:A:C6	22:DA:1755:A:C6	2.98	0.51
1:CA:1263:C:H2'	1:CA:1264:U:C6	2.46	0.51
1:CA:1166:G:C6	1:CA:1168:U:H5''	2.45	0.51
22:BA:277:G:H1'	22:BA:361:G:O6	2.10	0.51
12:AL:27:CYS:SG	12:AL:30:LYS:HE3	2.50	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:991:C:H5'	22:DA:1186:G:H5'	1.92	0.51
1:AA:1348:U:H4'	9:AI:122:ARG:HG3	1.92	0.51
22:BA:495:G:C1'	40:BS:57:ASN:HD21	2.23	0.51
28:DG:107:LEU:O	28:DG:152:ARG:NH2	2.41	0.51
22:DA:380:G:O3'	45:DX:16:ASN:HB2	2.11	0.51
8:AH:14:ILE:O	8:AH:16:ASN:N	2.44	0.51
50:D2:34:ARG:HB2	50:D2:42:LEU:HD13	1.92	0.51
22:DA:1401:G:C5	22:DA:1402:U:C5	2.98	0.51
29:BH:132:PHE:O	29:BH:139:PHE:HB3	2.11	0.51
22:DA:187:G:N2	22:DA:210:C:C2	2.78	0.51
24:BC:167:ARG:O	24:BC:168:ASP:CB	2.59	0.51
29:DH:53:GLU:O	29:DH:54:LEU:C	2.49	0.51
22:DA:1315:C:C2	22:DA:1338:G:N2	2.79	0.51
5:AE:104:GLY:O	5:AE:105:ILE:HB	2.11	0.51
22:DA:2136:G:N1	22:DA:2156:G:H1'	2.25	0.51
24:DC:65:VAL:CG1	24:DC:67:PHE:CE2	2.94	0.51
17:CQ:48:ASP:OD1	17:CQ:48:ASP:N	2.43	0.51
1:AA:263:A:H2'	1:AA:264:C:C5	2.45	0.51
1:CA:1481:U:H2'	1:CA:1482:G:C8	2.46	0.51
1:CA:786:G:N2	1:CA:787:A:H1'	2.26	0.51
1:CA:1068:G:N3	1:CA:1191:A:C2	2.79	0.51
22:DA:1857:G:C4	22:DA:1884:G:C2	2.98	0.51
22:DA:2345:G:C4	22:DA:2381:A:C2	2.98	0.51
1:AA:257:G:C2	1:AA:258:G:N7	2.79	0.51
1:AA:1149:C:OP1	9:AI:11:ARG:NE	2.44	0.51
1:AA:173:U:C2	1:AA:197:A:N1	2.79	0.51
22:BA:1536:C:O4'	22:BA:1537:G:C2	2.63	0.51
26:BE:147:LEU:HB3	26:BE:186:VAL:HG13	1.92	0.51
22:BA:2228:G:H2'	22:BA:2229:U:C6	2.46	0.51
7:CG:78:ARG:O	7:CG:79:ARG:HB2	2.11	0.51
16:CP:75:ILE:C	16:CP:77:GLU:H	2.13	0.51
22:DA:2575:C:OP2	57:DA:3710:HOH:O	2.18	0.51
37:DP:98:TYR:CD1	37:DP:101:ARG:NH2	2.79	0.51
22:BA:2056:G:C2	22:BA:2057:G:C8	2.99	0.51
22:DA:2201:G:H2'	22:DA:2202:U:H6	1.74	0.51
44:BW:38:VAL:CG1	44:BW:39:ARG:N	2.73	0.51
1:AA:358:U:H2'	1:AA:359:G:C8	2.46	0.51
29:BH:100:ALA:CB	29:BH:112:LYS:HA	2.41	0.51
22:BA:570:G:H2'	22:BA:2030:A:C8	2.46	0.51
29:BH:85:GLY:HA2	29:BH:91:PHE:CE2	2.46	0.51
12:AL:24:LEU:HB2	12:AL:59:ASN:HD22	1.75	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:1922:G:N2	22:BA:1923:U:H1'	2.25	0.51
41:BT:1:MET:O	41:BT:2:ILE:HD13	2.10	0.51
41:BT:6:ARG:O	41:BT:7:LEU:C	2.48	0.51
22:DA:2148:G:N2	22:DA:2149:U:C4	2.78	0.51
22:DA:49:A:H1'	22:DA:51:G:C6	2.46	0.51
1:AA:1525:G:OP1	11:AK:122:ARG:NH2	2.42	0.51
25:BD:13:ARG:HD2	25:BD:15:PHE:CE2	2.46	0.51
1:AA:1031:C:H4'	1:AA:1032:G:O5'	2.10	0.51
12:AL:21:VAL:HG23	12:AL:95:TYR:HE2	1.74	0.51
1:AA:1217:C:H2'	1:AA:1218:C:C6	2.46	0.51
22:BA:1416:G:HO2'	22:BA:1417:C:H6	1.56	0.51
8:AH:10:MET:HE1	8:AH:33:LYS:HA	1.93	0.51
22:BA:2847:U:H2'	22:BA:2848:G:H5'	1.92	0.51
3:CC:184:TYR:CE1	3:CC:201:TRP:CE2	2.99	0.51
1:CA:1313:U:P	19:CS:6:LYS:HB3	2.51	0.51
1:CA:1310:G:N2	1:CA:1328:C:O2	2.43	0.51
22:DA:982:C:C5'	22:DA:983:A:P	2.99	0.51
1:AA:657:U:O2	15:AO:22:THR:HG23	2.11	0.51
29:DH:26:ALA:HA	29:DH:30:LEU:HB2	1.92	0.51
22:BA:1266:G:N7	40:BS:16:LYS:HE3	2.25	0.51
22:BA:2656:U:C5	22:BA:2664:G:N2	2.78	0.51
45:DX:58:VAL:HG12	45:DX:59:ILE:N	2.24	0.51
5:CE:90:THR:HG22	5:CE:91:GLY:N	2.25	0.51
1:CA:965:U:OP1	1:CA:1198:G:H5''	2.10	0.51
22:DA:2852:G:H2'	22:DA:2853:C:O4'	2.11	0.51
22:DA:2307:G:N2	22:DA:2312:U:C2	2.79	0.51
22:DA:1924:C:H2'	22:DA:1925:C:O4'	2.11	0.51
27:BF:20:PHE:O	27:BF:21:ASN:ND2	2.43	0.51
1:CA:1494:G:C2	1:CA:1495:U:C5	2.98	0.51
29:BH:94:ILE:CG2	29:BH:99:ILE:CG1	2.88	0.51
22:BA:622:G:H2'	22:BA:623:C:C6	2.46	0.51
1:CA:1097:C:H2'	1:CA:1098:C:C6	2.46	0.51
22:BA:1918:A:O2'	22:BA:1920:C:C5	2.64	0.51
22:DA:1267:U:O3'	57:DA:3379:HOH:O	2.18	0.51
22:DA:189:G:C2'	22:DA:190:A:O5'	2.58	0.51
1:AA:1167:A:N7	1:AA:1169:A:C6	2.78	0.51
1:CA:106:C:O2	1:CA:379:C:H4'	2.10	0.51
46:BY:53:VAL:O	46:BY:56:LEU:O	2.28	0.51
1:CA:1107:C:N3	1:CA:1108:G:C8	2.79	0.51
1:CA:1108:G:OP1	3:CC:176:HIS:CD2	2.64	0.51
1:AA:858:G:C2'	1:AA:859:G:H5'	2.41	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:1800:C:HO2'	22:BA:1818:U:H3	1.59	0.51
22:DA:39:G:N2	22:DA:441:U:C2	2.79	0.51
22:DA:1063:G:H2'	22:DA:1064:C:O4'	2.10	0.51
22:DA:621:A:C5	22:DA:622:G:H1'	2.46	0.51
22:DA:1866:A:C8	22:DA:1867:G:C8	2.99	0.51
22:DA:2037:A:N6	22:DA:2038:G:C6	2.78	0.51
1:CA:748:G:N2	1:CA:749:A:C4	2.79	0.51
22:DA:2722:G:C2	22:DA:2723:C:C2	2.98	0.51
1:AA:843:U:OP1	1:AA:846:G:N2	2.44	0.51
53:B5:99:GLU:O	53:B5:103:LYS:CB	2.59	0.51
22:BA:1176:U:C4	22:BA:1177:G:O6	2.64	0.51
22:BA:1182:G:H2'	22:BA:1183:U:O4'	2.10	0.51
22:BA:28:A:C4	22:BA:29:U:C6	2.98	0.51
22:DA:2283:C:C2	22:DA:2389:G:C2	2.98	0.51
30:BI:108:GLU:HA	30:BI:111:GLN:HB3	1.91	0.51
22:BA:1039:A:H2'	22:BA:1040:A:O4'	2.10	0.51
22:DA:21:A:C2	22:DA:520:G:C2	2.99	0.51
1:CA:265:G:O3'	17:CQ:68:SER:HA	2.10	0.51
22:BA:62:U:O2	22:BA:63:A:N6	2.44	0.51
22:BA:1496:A:N3	22:BA:1577:C:O2'	2.38	0.51
10:CJ:77:VAL:O	10:CJ:79:PRO:HD3	2.11	0.51
2:AB:120:GLN:HG2	2:AB:125:THR:O	2.11	0.51
2:CB:43:LEU:HG	2:CB:44:GLU:N	2.25	0.51
3:AC:114:LYS:HD3	3:AC:185:ASN:OD1	2.11	0.51
1:CA:777:A:C2	1:CA:778:G:H1'	2.45	0.51
22:BA:2547:A:H2'	22:BA:2548:U:C6	2.45	0.51
22:DA:485:C:C2	22:DA:496:G:N2	2.79	0.51
27:BF:8:TYR:HA	27:BF:12:VAL:CG2	2.41	0.51
25:BD:55:LYS:HD2	25:BD:60:VAL:HG22	1.93	0.51
39:BR:49:ILE:HG22	39:BR:53:PHE:H	1.63	0.51
22:BA:2243:U:H2'	22:BA:2244:U:C6	2.46	0.51
29:BH:83:LYS:HA	29:BH:148:ALA:HA	1.93	0.51
2:AB:27:MET:SD	2:AB:193:PRO:HD3	2.51	0.51
22:DA:2683:C:H4'	25:DD:13:ARG:NH1	2.26	0.51
16:CP:70:ARG:O	16:CP:74:LEU:HG	2.11	0.51
1:AA:796:C:OP1	11:AK:126:LYS:HB2	2.10	0.51
22:DA:830:G:C6	22:DA:2448:A:C8	2.99	0.51
17:CQ:48:ASP:HB2	17:CQ:52:GLU:OE1	2.11	0.51
1:CA:784:A:C2	1:CA:785:G:C5	2.98	0.51
44:BW:10:THR:O	44:BW:11:ARG:CB	2.59	0.51
22:BA:1795:C:C4	22:BA:1796:U:C5	2.99	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:565:C:H4'	22:DA:1253:A:N6	2.26	0.51
1:AA:1033:G:C2'	1:AA:1034:G:H5'	2.41	0.51
22:DA:2621:G:H2'	22:DA:2622:U:O4'	2.11	0.51
22:DA:206:U:H2'	22:DA:207:A:H8	1.76	0.51
4:AD:151:LYS:HB2	4:AD:156:LYS:HE3	1.93	0.51
22:DA:2297:A:N7	22:DA:2320:U:C4	2.79	0.51
22:DA:2223:G:C2'	22:DA:2224:G:H5'	2.41	0.51
32:DK:107:LEU:O	32:DK:109:SER:N	2.44	0.51
22:DA:1665:A:H2'	22:DA:1666:G:O4'	2.11	0.51
5:CE:11:LEU:HG	5:CE:12:GLN:H	1.76	0.51
4:AD:13:ARG:NH2	4:AD:37:ALA:O	2.43	0.51
24:DC:51:THR:HG22	24:DC:54:ILE:HD11	1.93	0.51
42:DU:13:VAL:HG21	42:DU:39:ILE:HD12	1.92	0.51
1:AA:466:A:H5'	1:AA:467:U:OP2	2.10	0.51
27:DF:44:ILE:HG21	27:DF:79:ILE:HG22	1.91	0.51
10:CJ:15:HIS:HA	10:CJ:18:ILE:HG22	1.93	0.51
47:BZ:24:LEU:HD11	47:BZ:54:MET:CE	2.41	0.51
22:BA:973:A:O4'	22:BA:1188:U:C6	2.64	0.51
1:CA:667:G:OP1	1:CA:732:C:O2'	2.21	0.51
1:AA:1270:G:O3'	1:AA:1314:C:H5'	2.10	0.51
48:D0:53:LYS:HE3	48:D0:56:ALA:HA	1.92	0.51
22:BA:321:U:H5"	26:BE:131:THR:HG23	1.91	0.51
40:DS:47:VAL:HG23	40:DS:103:ILE:HG21	1.92	0.51
30:BI:97:LYS:CG	30:BI:139:VAL:HG22	2.41	0.51
1:CA:288:A:H2'	1:CA:289:G:H4'	1.93	0.51
41:BT:88:LYS:O	41:BT:89:GLU:HG2	2.11	0.51
27:BF:169:LEU:HD12	27:BF:169:LEU:O	2.11	0.51
22:DA:864:G:O2'	22:DA:914:G:O6	2.26	0.51
1:CA:1324:A:C6	1:CA:1325:C:N3	2.79	0.51
29:BH:121:VAL:CG2	29:BH:130:VAL:HG22	2.41	0.51
1:AA:1406:U:C6	1:AA:1407:C:C5	2.99	0.51
22:DA:811:U:C2	22:DA:1251:C:C4	2.99	0.51
1:CA:1144:G:C2	1:CA:1145:A:C2	2.99	0.51
22:DA:52:A:C5	22:DA:118:A:C2	2.99	0.51
1:CA:1296:C:N4	1:CA:1297:G:C6	2.79	0.51
11:AK:125:LYS:O	21:AU:34:ARG:NH2	2.44	0.51
22:DA:1045:C:O2	22:DA:1047:G:N1	2.44	0.51
22:DA:185:G:N1	22:DA:212:G:C2	2.79	0.51
35:DN:10:LEU:HD13	35:DN:40:LYS:HG3	1.92	0.51
5:AE:82:GLN:H	5:AE:147:MET:HE3	1.76	0.51
22:DA:2093:G:N7	22:DA:2225:A:C4	2.79	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:675:A:C5	1:CA:676:A:N7	2.79	0.51
26:DE:24:ASN:ND2	26:DE:27:LEU:CB	2.74	0.51
12:CL:40:THR:CG2	12:CL:90:LEU:HD12	2.41	0.51
22:DA:310:A:HO2'	22:DA:311:A:P	2.30	0.51
22:DA:311:A:H5'	22:DA:332:A:C2	2.46	0.51
1:AA:484:G:C5	1:AA:486:U:H1'	2.46	0.51
11:CK:23:ILE:HD11	11:CK:86:VAL:HG22	1.93	0.51
1:AA:1312:G:N7	19:AS:3:ARG:N	2.59	0.51
40:BS:84:ARG:HB2	40:BS:96:ILE:HG13	1.93	0.51
1:CA:35:G:O2'	12:CL:115:SER:O	2.29	0.51
22:DA:484:C:P	42:DU:48:PRO:HG3	2.51	0.51
1:CA:1461:G:H2'	1:CA:1462:C:C6	2.46	0.51
1:CA:1084:G:C8	1:CA:1085:U:C5	2.99	0.51
24:BC:251:GLN:HG3	24:BC:252:THR:O	2.11	0.51
1:AA:96:U:HO2'	1:AA:97:G:P	2.33	0.51
1:AA:1285:A:C5'	1:AA:1286:U:C4	2.94	0.51
20:AT:67:ILE:CD1	20:AT:71:LYS:HD3	2.41	0.51
1:AA:1335:U:H5''	1:AA:1336:C:OP1	2.11	0.51
21:AU:11:PRO:O	21:AU:12:PHE:HB3	2.11	0.51
22:BA:1002:G:C6	57:BA:3746:HOH:O	2.62	0.51
22:DA:1219:U:H2'	22:DA:1220:G:C8	2.46	0.51
45:DX:36:HIS:O	45:DX:48:THR:HA	2.11	0.51
29:DH:5:LEU:HA	29:DH:36:ALA:HA	1.93	0.51
32:BK:77:ILE:CD1	37:BP:72:ARG:HD2	2.41	0.51
22:BA:1850:G:C5	22:BA:1851:U:C5	2.98	0.51
9:CI:114:LYS:HG3	9:CI:120:LYS:HA	1.93	0.51
22:BA:2619:C:OP1	25:BD:157:LYS:HE2	2.10	0.51
42:BU:88:GLU:O	42:BU:89:ASP:HB3	2.11	0.51
13:AM:95:LEU:HB3	13:AM:96:PRO:HD2	1.93	0.51
22:DA:85:G:OP2	42:DU:7:ARG:HG2	2.11	0.51
22:DA:52:A:C6	22:DA:118:A:C2	2.99	0.51
16:CP:6:LEU:HD12	16:CP:71:VAL:HG23	1.93	0.51
22:DA:705:A:C2	22:DA:727:A:C1'	2.94	0.51
5:AE:126:LYS:HD3	5:AE:128:TYR:CE1	2.47	0.51
22:BA:784:G:H5''	24:BC:226:ASN:OD1	2.11	0.51
29:BH:14:SER:OG	29:BH:17:ASP:OD1	2.29	0.51
1:CA:575:G:C6	1:CA:821:G:N7	2.79	0.51
1:CA:372:C:N4	1:CA:387:U:H2'	2.26	0.51
22:BA:1735:A:C2	22:BA:1736:U:H1'	2.46	0.51
22:DA:1095:A:C6	22:DA:1096:A:N1	2.79	0.51
1:CA:635:A:C6	1:CA:636:U:C4	2.98	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:732:C:C4	22:DA:733:G:C8	2.99	0.51
1:CA:774:G:C5	1:CA:775:G:N7	2.79	0.51
20:AT:67:ILE:HD11	20:AT:71:LYS:HD3	1.92	0.51
22:DA:677:A:O2'	22:DA:2071:A:H5'	2.11	0.51
4:AD:37:ALA:N	4:AD:38:PRO:HD3	2.26	0.51
8:AH:8:ALA:HB2	8:AH:77:ARG:HD2	1.92	0.51
36:BO:88:LYS:O	36:BO:89:ASP:HB2	2.10	0.51
33:DL:68:SER:O	33:DL:69:ARG:CB	2.59	0.51
22:DA:1799:G:C8	24:DC:180:GLU:OE2	2.63	0.51
22:DA:325:G:O6	22:DA:338:G:C2	2.64	0.51
22:DA:2466:C:OP1	52:D4:4:ARG:HB3	2.10	0.51
27:BF:135:GLN:O	27:BF:135:GLN:HG2	2.10	0.51
1:CA:301:G:H2'	1:CA:302:G:C8	2.46	0.51
22:DA:1803:A:H2'	22:DA:1804:C:O4'	2.11	0.51
25:BD:26:VAL:HG22	25:BD:188:LEU:CD2	2.41	0.51
37:DP:92:VAL:HG21	37:DP:97:LEU:HD11	1.92	0.51
22:BA:2394:C:OP2	51:B3:30:ARG:HD3	2.11	0.51
42:BU:54:GLN:N	42:BU:55:PRO:CD	2.73	0.51
45:BX:18:ARG:CZ	45:BX:24:ALA:HB2	2.40	0.51
32:BK:8:LEU:HD12	32:BK:8:LEU:N	2.25	0.51
35:DN:28:LEU:O	35:DN:32:GLU:HA	2.11	0.51
22:DA:2612:C:H5''	22:DA:2613:U:OP1	2.11	0.51
4:CD:72:PHE:O	4:CD:75:TYR:HB2	2.10	0.51
22:BA:18:U:OP1	38:BQ:30:ARG:NH2	2.44	0.51
22:DA:1788:C:H2'	22:DA:1789:A:O4'	2.11	0.50
22:BA:2190:G:C6	22:BA:2191:A:C5	2.99	0.50
1:AA:1233:G:OP1	9:AI:126:GLN:HB3	2.11	0.50
1:AA:154:U:O2	1:AA:168:G:N2	2.43	0.50
1:CA:247:G:C5	1:CA:278:G:N2	2.79	0.50
22:DA:2780:G:P	31:DJ:120:ARG:HE	2.34	0.50
22:DA:2330:G:N2	22:DA:2386:A:N3	2.60	0.50
1:AA:1212:U:H5'	1:AA:1213:A:C5	2.45	0.50
1:CA:1384:C:N4	1:CA:1385:G:O6	2.43	0.50
12:AL:44:LYS:HB2	12:AL:45:PRO:HD3	1.92	0.50
22:DA:845:A:C6	22:DA:847:U:C6	2.98	0.50
22:DA:2771:C:H2'	22:DA:2772:C:C6	2.46	0.50
22:DA:2516:A:O2'	22:DA:2517:C:H5'	2.11	0.50
39:DR:39:LEU:O	39:DR:49:ILE:HG23	2.11	0.50
22:DA:457:A:N1	22:DA:470:A:H5''	2.26	0.50
22:DA:2624:G:H1'	48:D0:19:HIS:HE1	1.76	0.50
22:DA:1087:G:C4	22:DA:1089:A:H1'	2.46	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:AR:33:ILE:HA	18:AR:40:VAL:HG23	1.92	0.50
7:AG:70:ARG:HG3	7:AG:96:ARG:HG2	1.92	0.50
24:DC:24:LEU:HD11	24:DC:90:ASN:ND2	2.27	0.50
22:BA:2773:C:H5''	25:BD:169:ARG:HG2	1.92	0.50
1:CA:456:A:H2'	1:CA:457:G:O4'	2.11	0.50
22:DA:2289:G:HO2'	22:DA:2383:G:HO2'	1.57	0.50
37:DP:55:LEU:HA	37:DP:77:HIS:CD2	2.46	0.50
26:DE:23:PHE:HB2	26:DE:111:GLU:OE2	2.11	0.50
1:CA:690:G:H2'	1:CA:691:G:O4'	2.10	0.50
35:DN:30:ARG:HD2	35:DN:31:HIS:NE2	2.26	0.50
8:AH:54:ASP:OD1	8:AH:55:THR:N	2.44	0.50
22:DA:1373:A:C5	22:DA:1374:G:H1'	2.46	0.50
1:AA:977:A:O2'	1:AA:979:C:OP2	2.28	0.50
22:DA:49:A:C8	22:DA:51:G:N1	2.79	0.50
1:CA:411:A:C6	1:CA:429:U:C5	2.99	0.50
22:DA:1829:A:O2'	24:DC:15:HIS:NE2	2.39	0.50
22:DA:195:A:C6	22:DA:198:C:C5	3.00	0.50
22:DA:658:U:C4	22:DA:659:G:N7	2.78	0.50
22:DA:306:U:O4	22:DA:307:G:C6	2.64	0.50
1:AA:1253:G:H2'	1:AA:1254:A:C8	2.46	0.50
22:DA:1027:A:N6	22:DA:1126:A:H1'	2.26	0.50
22:DA:2342:C:O2'	22:DA:2374:C:H5''	2.11	0.50
1:CA:510:A:H5''	1:CA:511:C:P	2.50	0.50
22:DA:66:C:C4	22:DA:67:U:C5	2.99	0.50
1:AA:1237:C:C4	1:AA:1336:C:N3	2.78	0.50
49:B1:34:LEU:H	49:B1:52:ALA:HB3	1.76	0.50
1:CA:157:U:O2	1:CA:165:G:C2	2.65	0.50
1:CA:706:A:C1'	11:CK:31:ILE:HD11	2.41	0.50
22:DA:1188:U:O2'	22:DA:1189:A:H5'	2.12	0.50
2:AB:219:ALA:O	2:AB:220:THR:HB	2.11	0.50
22:DA:2250:G:C2	34:DM:82:MET:HB2	2.46	0.50
1:CA:401:C:OP2	4:CD:70:ARG:HD3	2.11	0.50
24:DC:185:GLU:HB2	24:DC:188:CYS:SG	2.51	0.50
1:CA:681:A:N1	1:CA:710:G:C6	2.80	0.50
44:BW:46:HIS:CE1	44:BW:77:ARG:HD3	2.47	0.50
38:BQ:68:ALA:O	38:BQ:71:GLN:HB3	2.11	0.50
2:CB:96:TRP:CE3	2:CB:97:LEU:O	2.64	0.50
19:CS:18:LYS:HE3	19:CS:33:THR:CG2	2.41	0.50
22:BA:1105:U:H2'	22:BA:1106:G:H8	1.76	0.50
7:CG:46:ALA:HA	7:CG:121:ALA:HB2	1.94	0.50
1:AA:1125:U:C5	1:AA:1127:G:C5	3.00	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:51:A:H4'	1:CA:52:C:C5'	2.41	0.50
22:BA:215:G:H4'	22:BA:216:A:H4'	1.93	0.50
2:AB:133:GLU:O	2:AB:137:ARG:N	2.43	0.50
22:DA:1068:G:N3	22:DA:1068:G:H2'	2.25	0.50
15:CO:73:LYS:HE2	15:CO:73:LYS:HA	1.93	0.50
17:AQ:36:LYS:HG2	17:AQ:37:PHE:N	2.27	0.50
1:AA:1537:U:C4	1:AA:1538:C:C4	3.00	0.50
22:DA:219:A:C6	22:DA:220:G:C6	2.99	0.50
22:DA:36:G:H4'	22:DA:451:U:C2	2.46	0.50
5:CE:98:PRO:O	5:CE:99:ALA:HB3	2.11	0.50
22:DA:1993:U:H4'	25:DD:133:THR:CG2	2.41	0.50
24:DC:189:ARG:O	24:DC:190:ALA:HB2	2.10	0.50
2:AB:186:ILE:HG12	2:AB:186:ILE:O	2.12	0.50
4:AD:109:ALA:N	4:AD:113:GLU:OE2	2.40	0.50
1:CA:327:A:C2	1:CA:329:A:N3	2.79	0.50
22:BA:2305:U:O2	27:BF:151:GLY:HA3	2.11	0.50
22:DA:685:A:H5''	22:DA:774:G:O6	2.11	0.50
40:DS:59:GLU:OE2	40:DS:66:ILE:HD11	2.11	0.50
22:DA:1182:G:H2'	22:DA:1183:U:O4'	2.12	0.50
5:AE:115:LEU:O	5:AE:120:VAL:HG22	2.11	0.50
1:CA:243:A:C2	1:CA:245:U:C2	2.99	0.50
22:BA:1319:C:C2'	22:BA:1320:C:H5'	2.42	0.50
6:CF:64:VAL:HG12	6:CF:65:GLU:H	1.76	0.50
23:BB:37:C:C5	23:BB:38:C:C5	2.99	0.50
48:D0:54:VAL:O	48:D0:56:ALA:N	2.44	0.50
1:CA:1244:G:C2	1:CA:1294:G:C2	2.99	0.50
29:BH:66:ASN:OD1	29:BH:138:VAL:HG21	2.11	0.50
1:CA:542:G:C2	1:CA:543:U:C5	3.00	0.50
22:BA:855:G:H2'	22:BA:856:G:O5'	2.12	0.50
22:BA:78:U:H2'	22:BA:79:C:C6	2.46	0.50
22:BA:958:U:H2'	23:BB:89:U:C2	2.47	0.50
57:BA:3790:HOH:O	24:BC:219:THR:HB	2.10	0.50
34:BM:41:LEU:CD2	34:BM:125:PRO:HD2	2.42	0.50
29:BH:80:ILE:O	29:BH:147:VAL:N	2.44	0.50
29:DH:44:ILE:O	29:DH:48:GLU:HB2	2.11	0.50
22:DA:2142:A:C5	22:DA:2143:C:C5	2.99	0.50
22:DA:2821:A:OP2	25:DD:115:GLY:CA	2.58	0.50
1:CA:1169:A:C2	1:CA:1170:A:C5	3.00	0.50
20:CT:4:ILE:O	20:CT:5:LYS:HB2	2.11	0.50
24:DC:43:ARG:NH2	24:DC:49:ILE:HD11	2.26	0.50
2:CB:93:ASN:OD1	2:CB:94:HIS:ND1	2.44	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:736:C:H2'	1:CA:737:C:C6	2.47	0.50
22:DA:2574:G:N3	22:DA:2574:G:H2'	2.26	0.50
4:CD:174:ASP:O	4:CD:175:ALA:HB3	2.11	0.50
22:DA:30:G:C5	22:DA:31:C:C4	2.99	0.50
30:BI:110:ALA:HB1	30:BI:129:ILE:HG13	1.93	0.50
22:BA:947:A:O2'	22:BA:984:A:C2	2.59	0.50
22:BA:1392:A:C6	22:BA:1393:A:C6	2.99	0.50
4:AD:62:ARG:NH1	4:AD:69:GLU:HG2	2.26	0.50
4:AD:58:LYS:NZ	4:AD:69:GLU:OE2	2.40	0.50
6:CF:51:ILE:HG12	6:CF:51:ILE:O	2.12	0.50
22:BA:498:G:O2'	42:BU:45:HIS:NE2	2.43	0.50
25:DD:7:LYS:O	25:DD:27:ILE:HA	2.11	0.50
22:BA:2810:A:H2'	22:BA:2811:G:O4'	2.10	0.50
22:DA:981:A:OP2	22:DA:982:C:N4	2.41	0.50
1:CA:747:A:N6	1:CA:748:G:C6	2.79	0.50
7:CG:33:ASP:HB3	7:CG:35:LYS:HE3	1.92	0.50
34:BM:2:LEU:O	34:BM:3:GLN:HB3	2.11	0.50
3:CC:130:PHE:CE2	3:CC:157:LEU:HB3	2.46	0.50
22:DA:1149:G:H2'	22:DA:1150:C:C6	2.46	0.50
1:AA:986:U:H1'	19:AS:54:GLY:O	2.11	0.50
1:AA:723:U:H5''	21:AU:49:LYS:HG2	1.92	0.50
1:AA:1048:G:N3	1:AA:1050:G:N7	2.60	0.50
3:CC:117:ALA:HB1	3:CC:187:SER:HB3	1.91	0.50
1:CA:1391:U:H2'	1:CA:1392:G:C8	2.46	0.50
22:DA:927:A:H2'	22:DA:928:A:C8	2.46	0.50
22:DA:2686:G:H2'	22:DA:2687:U:C6	2.46	0.50
20:CT:55:GLN:N	20:CT:56:PRO:CD	2.74	0.50
1:CA:597:G:H2'	1:CA:598:U:H5'	1.92	0.50
22:DA:35:G:C4	22:DA:454:A:C2	3.00	0.50
1:AA:452:A:N6	1:AA:480:U:C2	2.79	0.50
1:CA:449:G:H2'	1:CA:450:G:C8	2.46	0.50
22:DA:536:G:C6	22:DA:537:G:C4	3.00	0.50
1:CA:1106:G:N3	1:CA:1107:C:C6	2.79	0.50
22:DA:1718:G:O6	22:DA:1743:G:C2	2.64	0.50
1:CA:938:A:N6	1:CA:939:G:C5	2.80	0.50
5:CE:56:VAL:N	5:CE:57:PRO:CD	2.75	0.50
17:CQ:19:LYS:HD2	17:CQ:49:GLU:HA	1.93	0.50
5:AE:45:ARG:HG2	5:AE:73:ASN:HB3	1.94	0.50
22:BA:1085:A:C6	22:BA:1086:A:N6	2.79	0.50
23:DB:29:A:O2'	23:DB:58:A:N1	2.38	0.50
1:CA:72:A:C6	1:CA:73:C:C4	3.00	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:2204:G:N3	22:DA:2205:A:C8	2.80	0.50
22:DA:2563:U:H1'	22:DA:2566:A:N6	2.27	0.50
1:AA:73:C:HO2'	1:AA:74:A:H5''	1.75	0.50
15:CO:37:ASN:O	15:CO:38:HIS:C	2.50	0.50
39:DR:52:PRO:O	39:DR:53:PHE:CG	2.64	0.50
39:DR:52:PRO:O	39:DR:53:PHE:CD2	2.64	0.50
1:AA:1182:G:H4'	1:AA:1183:U:H5'	1.94	0.50
1:CA:706:A:C5	1:CA:707:U:C5	3.00	0.50
15:AO:2:SER:O	15:AO:3:LEU:CB	2.58	0.50
22:BA:277:G:O2'	22:BA:361:G:N1	2.44	0.50
22:BA:445:C:H5''	38:BQ:3:ARG:HB2	1.92	0.50
26:BE:48:THR:HG22	26:BE:86:ALA:HB3	1.92	0.50
19:CS:16:LEU:O	19:CS:20:GLU:HG2	2.11	0.50
1:AA:942:G:H2'	1:AA:942:G:N3	2.25	0.50
4:CD:161:LEU:H	4:CD:161:LEU:HD22	1.77	0.50
1:AA:862:C:O2'	1:AA:863:U:H5'	2.11	0.50
30:BI:125:MET:O	30:BI:128:SER:OG	2.25	0.50
28:BG:89:LEU:CD1	28:BG:89:LEU:N	2.74	0.50
18:AR:37:GLY:O	18:AR:63:ARG:NH2	2.45	0.50
22:DA:1370:C:H2'	22:DA:1371:G:C8	2.47	0.50
22:BA:1482:G:C2	22:BA:1483:G:C8	3.00	0.50
22:DA:192:C:H2'	22:DA:193:U:O4'	2.11	0.50
22:DA:397:U:N3	22:DA:398:C:C5	2.80	0.50
27:BF:63:GLN:OE1	27:BF:95:ARG:HD3	2.12	0.50
22:BA:1022:G:N7	31:BJ:68:LYS:HE2	2.26	0.50
22:DA:161:A:C3'	22:DA:162:U:H5''	2.39	0.50
22:DA:1798:U:H5''	24:DC:258:ARG:HB2	1.93	0.50
22:DA:1833:C:C4	22:DA:1834:U:C5	2.99	0.50
22:BA:1138:G:H5''	22:BA:1139:G:OP2	2.12	0.50
26:DE:149:ILE:CG2	26:DE:188:MET:HG2	2.42	0.50
22:DA:1060:U:O4'	22:DA:1062:G:H5'	2.12	0.50
22:BA:1984:G:C6	22:BA:1985:C:C5	3.00	0.50
3:CC:184:TYR:CE1	3:CC:201:TRP:CZ2	3.00	0.50
22:DA:533:G:H5'	38:DQ:24:TYR:CD1	2.47	0.50
19:AS:32:ARG:HA	19:AS:50:ALA:HB3	1.93	0.50
22:DA:411:G:OP2	22:DA:2406:A:O2'	2.28	0.50
2:CB:54:LEU:HD12	2:CB:220:THR:HG21	1.93	0.50
4:AD:123:ILE:O	4:AD:123:ILE:HD13	2.12	0.50
29:DH:5:LEU:HD11	29:DH:13:GLY:HA2	1.93	0.50
27:BF:135:GLN:HG2	27:BF:141:ILE:HG12	1.94	0.50
22:DA:2714:G:C5	22:DA:2715:C:C5	2.98	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:722:A:H2'	22:BA:723:C:O4'	2.12	0.50
46:BY:61:ALA:O	46:BY:63:ALA:N	2.37	0.50
22:BA:962:G:H21	22:BA:2250:G:H1	1.59	0.50
1:CA:307:C:H5''	1:CA:308:C:OP2	2.12	0.50
1:CA:629:A:H2'	1:CA:630:A:O4'	2.11	0.50
28:DG:35:ARG:NE	28:DG:71:LEU:HD22	2.26	0.50
22:BA:2703:C:H2'	22:BA:2704:C:H6	1.76	0.50
22:BA:2023:C:H2'	22:BA:2023:C:O2	2.11	0.50
13:AM:85:CYS:O	13:AM:89:LEU:HG	2.12	0.50
22:DA:1310:G:C2'	22:DA:1311:G:H5'	2.42	0.50
22:DA:1359:A:N1	22:DA:1360:G:H1'	2.27	0.50
22:DA:58:G:C4	22:DA:70:G:N2	2.79	0.50
22:DA:2683:C:OP1	37:DP:56:HIS:HB3	2.12	0.50
1:AA:374:A:C5	1:AA:375:U:C5	2.99	0.50
5:CE:122:ASN:O	5:CE:123:VAL:O	2.30	0.50
1:AA:1077:G:N2	1:AA:1080:A:OP2	2.43	0.50
1:AA:203:G:O2'	1:AA:465:A:N1	2.40	0.50
22:DA:2217:G:C6	22:DA:2218:G:N7	2.79	0.50
22:DA:570:G:C4	22:DA:2030:A:N7	2.80	0.50
20:CT:79:LEU:HA	20:CT:82:GLN:HB2	1.94	0.50
1:AA:428:G:O4'	1:AA:430:A:C8	2.65	0.50
6:CF:1:MET:HG3	6:CF:66:ALA:O	2.12	0.50
23:DB:76:G:OP1	43:DV:9:ARG:NH2	2.39	0.50
1:CA:892:A:C5	1:CA:893:C:C5	3.00	0.50
22:DA:1865:U:C5	22:DA:1875:G:C2	3.00	0.50
5:CE:96:MET:CE	5:CE:111:MET:HE3	2.42	0.50
22:BA:973:A:H5'	22:BA:1188:U:H1'	1.93	0.50
40:BS:17:VAL:HG12	40:BS:76:VAL:HG21	1.94	0.50
1:AA:683:G:N2	11:AK:40:ASN:HA	2.27	0.50
22:DA:883:G:N2	22:DA:894:U:O2	2.45	0.50
22:DA:751:A:H5''	22:DA:752:A:OP1	2.12	0.50
22:BA:2771:C:H2'	22:BA:2772:C:H6	1.77	0.50
22:DA:2652:C:C4	22:DA:2653:U:C4	3.00	0.50
22:BA:826:U:O2'	33:BL:53:GLY:HA3	2.10	0.50
4:AD:101:VAL:HG12	4:AD:101:VAL:O	2.12	0.50
1:AA:512:U:H2'	1:AA:513:C:C6	2.47	0.50
1:AA:260:G:H2'	1:AA:261:U:C6	2.46	0.50
4:AD:91:LEU:HD21	4:AD:195:ILE:HD12	1.93	0.50
3:AC:154:SER:O	3:AC:196:ILE:HG23	2.10	0.50
22:DA:997:G:OP1	38:DQ:92:ARG:HG2	2.11	0.50
45:DX:33:LEU:HD12	45:DX:51:VAL:O	2.12	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:452:A:C8	1:AA:453:G:C8	2.99	0.50
4:AD:170:TRP:CE2	4:AD:186:PRO:HG3	2.47	0.50
22:BA:588:U:H2'	22:BA:589:U:H6	1.73	0.50
1:AA:495:A:C2	1:AA:496:A:N6	2.80	0.50
5:CE:133:PRO:O	5:CE:137:VAL:HG12	2.11	0.50
16:AP:46:LYS:HD3	16:AP:47:GLU:N	2.27	0.50
1:AA:145:G:N2	1:AA:178:C:N3	2.59	0.50
22:DA:753:A:H2'	22:DA:754:U:C6	2.47	0.50
22:DA:2196:C:O2'	22:DA:2197:U:H5'	2.12	0.50
22:DA:599:A:N3	22:DA:659:G:C2	2.80	0.50
4:CD:38:PRO:HD2	4:CD:42:GLY:CA	2.42	0.50
22:DA:107:G:O3'	22:DA:293:U:O2'	2.25	0.50
23:BB:46:A:C5	23:BB:47:C:C5	3.00	0.50
24:DC:212:ARG:HD2	24:DC:216:VAL:O	2.11	0.50
4:CD:145:ILE:HD13	4:CD:178:MET:HB3	1.94	0.50
22:DA:1241:A:C2	22:DA:1242:U:H1'	2.47	0.50
1:CA:1518:A:N1	1:CA:1519:A:C6	2.80	0.50
22:DA:2345:G:C5	22:DA:2347:C:N4	2.80	0.50
1:AA:1062:U:H2'	1:AA:1063:C:C5	2.47	0.50
13:AM:114:LYS:CB	13:AM:115:PRO:HD3	2.41	0.50
27:BF:41:GLY:C	27:BF:43:ALA:H	2.14	0.50
33:DL:56:PRO:O	33:DL:60:ARG:HB2	2.11	0.50
22:DA:1865:U:C4	22:DA:1875:G:C2	2.98	0.50
20:CT:55:GLN:N	20:CT:56:PRO:HD2	2.26	0.50
22:DA:2808:G:H4'	22:DA:2809:A:O5'	2.12	0.50
1:CA:1434:A:N6	1:CA:1435:G:C6	2.80	0.50
1:CA:251:G:N1	1:CA:266:G:C6	2.80	0.50
22:BA:2579:C:O2'	22:BA:2580:U:H5'	2.11	0.50
1:CA:209:U:H4'	1:CA:210:C:OP2	2.12	0.50
31:DJ:89:PHE:CE2	31:DJ:100:VAL:HG11	2.47	0.50
1:CA:1072:G:C5	1:CA:1073:U:C4	3.00	0.50
41:BT:2:ILE:HD12	41:BT:7:LEU:HG	1.93	0.50
35:BN:12:ARG:NH1	35:BN:20:MET:HE1	2.27	0.50
22:DA:2823:A:C2	22:DA:2824:C:C6	3.00	0.50
22:DA:528:A:N1	22:DA:2043:C:C5'	2.75	0.50
57:DA:3717:HOH:O	50:D2:11:LYS:NZ	2.44	0.50
17:CQ:14:SER:C	17:CQ:17:MET:HE1	2.33	0.50
12:CL:27:CYS:HB3	12:CL:29:GLN:O	2.12	0.50
8:AH:18:GLN:O	8:AH:20:ALA:N	2.44	0.50
22:DA:1832:C:C4	22:DA:1833:C:C5	3.00	0.50
2:AB:96:TRP:HZ3	2:AB:175:GLU:OE2	1.95	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:126:A:P	50:D2:19:ARG:HG3	2.52	0.50
2:AB:151:ILE:O	2:AB:152:LYS:C	2.50	0.50
1:CA:158:G:C4	1:CA:159:G:C8	3.00	0.50
22:BA:371:A:O2'	45:BX:61:LYS:NZ	2.41	0.50
22:DA:1801:A:C5	24:DC:262:ARG:NH2	2.80	0.50
27:BF:108:VAL:HG11	27:BF:176:PRO:HG2	1.93	0.50
21:AU:39:GLU:OE2	21:AU:42:THR:HG21	2.12	0.50
16:AP:4:ILE:HG12	16:AP:21:VAL:HG22	1.94	0.50
1:AA:1209:C:O2'	1:AA:1210:C:H5'	2.11	0.50
20:AT:68:HIS:O	20:AT:69:LYS:NZ	2.44	0.50
22:BA:1488:C:O2	22:BA:1502:A:C2	2.65	0.50
22:DA:2753:A:C6	22:DA:2754:U:C4	2.99	0.50
11:AK:72:ASP:O	11:AK:73:ALA:HB2	2.12	0.50
22:BA:2258:C:H4'	22:BA:2259:U:OP2	2.12	0.50
22:BA:1405:U:H2'	22:BA:1406:U:C6	2.46	0.50
22:DA:2050:C:N4	22:DA:2051:A:C6	2.80	0.50
39:DR:68:ARG:HD2	39:DR:90:ARG:HB3	1.92	0.50
8:CH:18:GLN:NE2	8:CH:70:ALA:HB1	2.26	0.50
1:AA:818:G:O2'	1:AA:819:A:H5'	2.12	0.50
22:BA:1916:A:C5	22:BA:1917:U:C6	3.00	0.49
22:DA:49:A:C8	22:DA:51:G:C2	3.00	0.49
1:CA:106:C:H2'	1:CA:107:G:O4'	2.12	0.49
22:DA:2216:G:H2'	22:DA:2217:G:C8	2.47	0.49
1:AA:1178:G:C5	9:AI:99:ARG:NH2	2.80	0.49
1:AA:8:A:N6	4:AD:202:GLU:O	2.44	0.49
6:AF:3:HIS:H	6:AF:92:THR:HG23	1.77	0.49
1:CA:1219:A:C5	1:CA:1220:G:N7	2.80	0.49
22:BA:1414:C:C4	22:BA:1415:U:C5	2.99	0.49
22:BA:1589:U:N3	22:BA:1590:A:N7	2.59	0.49
22:DA:810:U:C6	33:DL:29:LYS:HA	2.47	0.49
1:AA:1152:A:OP1	10:AJ:15:HIS:HB2	2.11	0.49
22:DA:639:U:H2'	22:DA:640:C:C6	2.46	0.49
29:BH:43:ASN:O	29:BH:46:PHE:HB3	2.12	0.49
8:AH:125:ILE:O	8:AH:125:ILE:HG13	2.12	0.49
29:DH:127:GLU:HG3	29:DH:145:ASN:HA	1.93	0.49
11:AK:34:ILE:HB	11:AK:74:VAL:HG11	1.93	0.49
1:CA:1525:G:O6	57:CA:1885:HOH:O	2.20	0.49
41:DT:7:LEU:HD22	41:DT:46:ALA:HB2	1.94	0.49
1:CA:1366:C:O2'	10:CJ:62:ARG:NH2	2.45	0.49
22:BA:741:U:P	57:BA:3702:HOH:O	2.70	0.49
1:CA:1122:U:C4	1:CA:1123:U:C4	3.00	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1304:G:N1	1:AA:1305:G:N2	2.60	0.49
4:AD:46:PRO:O	4:AD:48:LEU:N	2.45	0.49
2:CB:35:ARG:O	2:CB:38:VAL:N	2.42	0.49
22:BA:2352:A:H5''	22:BA:2353:G:OP2	2.12	0.49
31:BJ:56:VAL:HB	31:BJ:124:VAL:HB	1.94	0.49
1:CA:211:G:N3	1:CA:211:G:H2'	2.27	0.49
22:DA:2326:C:H1'	22:DA:2327:A:OP1	2.11	0.49
28:BG:25:THR:HG23	28:BG:34:THR:OG1	2.11	0.49
5:AE:132:ASN:O	5:AE:136:VAL:HG12	2.12	0.49
29:BH:99:ILE:O	29:BH:99:ILE:HG22	2.12	0.49
1:CA:58:C:H2'	1:CA:59:A:O5'	2.12	0.49
14:CN:41:ARG:HG2	14:CN:42:TRP:N	2.27	0.49
1:AA:960:U:H2'	1:AA:1225:A:H62	1.77	0.49
22:DA:1807:G:H2'	22:DA:1808:A:O5'	2.12	0.49
31:BJ:81:ILE:CG2	31:BJ:82:GLY:H	2.15	0.49
5:CE:81:LEU:HD22	5:CE:123:VAL:HG12	1.93	0.49
22:DA:1020:A:C2	22:DA:1141:U:C2	3.00	0.49
22:DA:160:A:C6	22:DA:161:A:C6	3.00	0.49
22:BA:2846:G:OP2	37:BP:52:ASN:HB2	2.12	0.49
1:AA:468:A:C2	1:AA:469:C:C5	3.00	0.49
22:DA:1195:G:O2'	22:DA:1226:A:N1	2.32	0.49
1:AA:544:G:OP1	4:AD:59:GLN:NE2	2.46	0.49
22:DA:362:A:C5	22:DA:363:G:C8	3.00	0.49
5:AE:45:ARG:HA	5:AE:72:ILE:O	2.12	0.49
1:AA:188:C:O2	1:AA:188:C:C2'	2.60	0.49
1:CA:1480:A:C4	1:CA:1481:U:C6	3.00	0.49
30:DI:69:PHE:N	30:DI:69:PHE:CD1	2.80	0.49
23:DB:48:U:H4'	36:DO:100:HIS:CD2	2.47	0.49
21:AU:17:ARG:NH1	21:AU:20:LYS:HG3	2.27	0.49
22:DA:1476:U:H1'	22:DA:1732:C:O2	2.12	0.49
1:AA:1539:C:O3'	21:AU:18:ARG:HB3	2.12	0.49
1:AA:1129:C:H5'	9:AI:18:ARG:HH22	1.77	0.49
22:BA:28:A:C5	22:BA:29:U:C5	3.00	0.49
50:D2:22:MET:SD	50:D2:28:ARG:HG2	2.52	0.49
22:DA:2282:G:N3	22:DA:2425:A:N6	2.61	0.49
22:DA:167:A:H2'	22:DA:168:G:O4'	2.12	0.49
1:CA:558:G:C4	1:CA:559:A:C2	3.00	0.49
24:BC:43:ARG:HA	24:BC:48:ARG:O	2.12	0.49
30:DI:54:PRO:O	30:DI:75:PRO:HD2	2.11	0.49
1:CA:571:U:H5''	1:CA:572:A:OP2	2.12	0.49
22:BA:827:U:H2'	22:BA:2068:U:C2	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1508:A:H2'	1:AA:1509:C:O4'	2.12	0.49
22:DA:1293:C:H2'	22:DA:1294:U:O4'	2.12	0.49
24:BC:187:ASP:N	24:BC:187:ASP:OD1	2.44	0.49
22:DA:1:G:H2'	22:DA:2:G:C8	2.47	0.49
22:DA:389:G:C2	22:DA:2413:G:H1'	2.47	0.49
18:AR:26:ILE:O	18:AR:30:LYS:HG3	2.12	0.49
22:BA:967:U:H2'	22:BA:968:C:C6	2.47	0.49
22:BA:994:C:H3'	38:BQ:54:LYS:HE3	1.94	0.49
6:CF:38:ARG:HG2	6:CF:63:ASN:HB3	1.94	0.49
1:CA:577:G:H1'	1:CA:816:A:N3	2.27	0.49
27:DF:33:LYS:HD3	27:DF:92:ARG:NH1	2.27	0.49
5:CE:18:VAL:HG21	5:CE:56:VAL:HG13	1.94	0.49
4:CD:168:PRO:HB2	4:CD:171:LEU:HD13	1.94	0.49
22:DA:2747:G:C2	22:DA:2756:U:C5	3.00	0.49
1:CA:568:G:N2	1:CA:883:C:C2	2.80	0.49
8:AH:10:MET:HG3	8:AH:27:MET:SD	2.52	0.49
1:CA:64:G:H4'	1:CA:65:A:H5''	1.94	0.49
1:AA:74:A:C2	1:AA:97:G:C4	3.00	0.49
22:DA:2370:G:C6	22:DA:2371:G:C6	3.01	0.49
1:AA:439:U:H1'	4:AD:119:SER:O	2.12	0.49
1:CA:154:U:C2	1:CA:168:G:N2	2.80	0.49
1:CA:1408:A:C2	1:CA:1494:G:C5	3.00	0.49
1:CA:264:C:H2'	1:CA:265:G:O4'	2.13	0.49
22:BA:17:G:H2'	22:BA:18:U:C6	2.47	0.49
22:DA:43:G:C2	22:DA:437:U:C2	3.00	0.49
1:AA:866:C:C4	1:AA:867:G:H1'	2.48	0.49
17:CQ:79:VAL:O	17:CQ:80:GLU:HB2	2.12	0.49
13:CM:20:THR:HG22	13:CM:26:GLY:O	2.10	0.49
26:BE:189:THR:HG22	26:BE:191:ASP:N	2.27	0.49
36:BO:24:THR:HG22	36:BO:41:ALA:HA	1.95	0.49
22:DA:569:U:H5''	22:DA:821:A:C2	2.48	0.49
22:DA:1274:A:N3	22:DA:1297:C:H1'	2.28	0.49
11:CK:33:THR:O	11:CK:33:THR:HG22	2.13	0.49
29:BH:86:ASP:O	29:BH:87:GLU:HB2	2.11	0.49
22:BA:1073:A:C8	22:BA:1074:G:O4'	2.66	0.49
1:CA:1070:U:O2'	1:CA:1071:C:H5'	2.12	0.49
2:CB:99:GLY:HA2	2:CB:102:THR:HG22	1.94	0.49
22:DA:2127:G:O2'	22:DA:2173:A:C2	2.65	0.49
1:AA:373:A:C2	1:AA:374:A:C8	3.01	0.49
11:CK:124:PRO:HB2	11:CK:126:LYS:HE3	1.93	0.49
22:BA:362:A:C8	22:BA:362:A:OP2	2.66	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:599:A:C2	22:DA:659:G:C6	3.00	0.49
22:DA:789:A:C2	57:DA:3310:HOH:O	2.64	0.49
22:DA:1272:A:C5	22:DA:1618:A:H1'	2.47	0.49
22:DA:668:A:N6	22:DA:670:A:O2'	2.46	0.49
22:DA:1441:G:C2	22:DA:1442:U:C4	2.99	0.49
1:CA:583:A:C2	1:CA:759:A:C5	3.01	0.49
22:DA:2221:G:C2'	22:DA:2222:C:H5'	2.42	0.49
22:DA:2346:A:H4'	22:DA:2347:C:OP2	2.12	0.49
22:BA:1796:U:O2	22:BA:1796:U:H2'	2.12	0.49
4:AD:147:GLU:HA	4:AD:150:LYS:HD2	1.93	0.49
22:BA:1045:C:H3'	22:BA:1046:A:H5'	1.95	0.49
1:AA:1349:A:H2'	1:AA:1350:A:O4'	2.13	0.49
22:DA:1665:A:OP1	32:DK:66:LYS:HE3	2.11	0.49
22:DA:420:C:H2'	22:DA:421:C:H6	1.78	0.49
1:AA:557:G:C6	1:AA:558:G:C6	2.99	0.49
1:CA:1310:G:N2	1:CA:1328:C:C2	2.81	0.49
26:BE:119:ILE:HB	26:BE:187:VAL:CG2	2.42	0.49
20:AT:69:LYS:HB2	20:AT:69:LYS:NZ	2.27	0.49
6:CF:38:ARG:NH1	6:CF:61:LEU:HD21	2.27	0.49
22:BA:440:C:O2'	22:BA:441:U:H5'	2.12	0.49
22:DA:1506:U:H2'	22:DA:1507:C:C6	2.47	0.49
22:BA:1540:G:H2'	22:BA:1541:C:O4'	2.12	0.49
7:CG:57:SER:HB3	7:CG:60:GLU:HG2	1.93	0.49
22:DA:839:U:H2'	22:DA:840:C:C6	2.47	0.49
22:BA:2489:U:O2	22:BA:2491:U:C4	2.66	0.49
19:AS:64:ASP:HB2	27:BF:115:ARG:NH2	2.27	0.49
3:AC:42:TYR:CZ	3:AC:90:VAL:HG21	2.47	0.49
22:BA:868:U:C4	22:BA:869:G:N7	2.80	0.49
42:BU:102:THR:HG22	42:BU:103:ILE:N	2.27	0.49
35:BN:106:ASP:OD1	35:BN:106:ASP:C	2.48	0.49
34:BM:110:GLU:OE2	34:BM:114:ARG:NH2	2.45	0.49
22:DA:1384:A:O2'	22:DA:1404:C:O2	2.29	0.49
1:CA:57:G:H2'	1:CA:58:C:C6	2.48	0.49
22:DA:449:A:C5	22:DA:450:G:C8	3.01	0.49
18:CR:32:TYR:CG	18:CR:55:LEU:HD21	2.47	0.49
22:BA:2190:G:C2	22:BA:2191:A:C4	3.00	0.49
1:AA:452:A:N7	1:AA:453:G:C4	2.80	0.49
1:AA:450:G:N7	1:AA:481:G:C6	2.81	0.49
22:BA:2297:A:N1	22:BA:2321:U:C5	2.78	0.49
22:DA:2217:G:C4	22:DA:2218:G:C8	3.01	0.49
22:DA:674:G:O6	22:DA:806:C:N3	2.45	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:301:G:N3	22:DA:302:C:C2	2.81	0.49
1:CA:369:G:OP2	1:CA:388:G:N2	2.45	0.49
2:AB:72:THR:O	2:AB:73:LYS:HB3	2.12	0.49
4:CD:107:PHE:CD2	4:CD:145:ILE:HD11	2.47	0.49
22:BA:1322:A:O3'	40:BS:84:ARG:NH1	2.42	0.49
1:AA:1153:G:C2	1:AA:1154:G:C8	3.00	0.49
36:BO:76:LYS:HE3	36:BO:80:GLU:OE2	2.11	0.49
22:BA:1866:A:H2'	22:BA:1867:G:H5'	1.95	0.49
1:AA:972:C:H4'	10:AJ:59:LYS:CE	2.42	0.49
27:BF:108:VAL:HG13	27:BF:114:PHE:CZ	2.47	0.49
8:AH:5:ASP:OD1	8:AH:77:ARG:NH1	2.45	0.49
22:DA:983:A:N6	22:DA:984:A:C2	2.80	0.49
1:AA:843:U:OP1	1:AA:846:G:N1	2.46	0.49
34:BM:135:VAL:O	34:BM:136:MET:HB3	2.12	0.49
22:DA:2085:U:C2	22:DA:2086:U:C5	3.00	0.49
22:BA:1720:U:H2'	22:BA:1721:G:O4'	2.11	0.49
51:D3:7:VAL:O	51:D3:10:ALA:HB3	2.12	0.49
22:BA:2307:G:N3	22:BA:2308:G:O6	2.45	0.49
22:BA:923:G:H4'	44:BW:29:GLU:HG3	1.95	0.49
24:DC:145:GLU:HG2	24:DC:152:GLY:N	2.27	0.49
1:CA:926:G:C6	1:CA:1505:G:C6	2.99	0.49
1:AA:965:U:OP2	57:AA:1833:HOH:O	2.19	0.49
29:DH:112:LYS:CG	29:DH:113:SER:N	2.76	0.49
49:D1:35:GLU:HG2	49:D1:50:LYS:HG3	1.95	0.49
33:BL:132:ARG:HG3	33:BL:142:ILE:HD13	1.93	0.49
28:DG:89:LEU:CD1	28:DG:162:VAL:HG22	2.42	0.49
22:BA:1483:G:C2	22:BA:1484:U:C2	3.01	0.49
41:BT:7:LEU:HD22	41:BT:46:ALA:HB2	1.94	0.49
22:DA:1810:A:H2'	22:DA:1811:G:O4'	2.12	0.49
45:DX:28:ARG:O	45:DX:29:PHE:CD1	2.66	0.49
34:BM:26:VAL:HA	34:BM:104:GLU:OE1	2.13	0.49
24:BC:167:ARG:O	24:BC:168:ASP:HB3	2.13	0.49
1:AA:672:U:H2'	1:AA:673:A:H8	1.77	0.49
46:BY:23:ARG:O	46:BY:24:GLU:C	2.49	0.49
22:DA:1973:G:C8	22:DA:1974:C:C5	3.00	0.49
22:DA:1544:A:C6	22:DA:1545:A:C2	3.00	0.49
24:DC:101:ARG:O	24:DC:102:ARG:HG3	2.13	0.49
1:AA:502:A:H2'	1:AA:503:C:O4'	2.13	0.49
1:CA:1480:A:H2'	1:CA:1481:U:O4'	2.13	0.49
1:CA:223:A:C6	1:CA:224:U:C4	3.00	0.49
1:AA:1072:G:H2'	1:AA:1073:U:C6	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:2810:A:H2'	22:DA:2811:G:O4'	2.13	0.49
45:BX:42:SER:OG	45:BX:65:ASP:OD1	2.30	0.49
1:CA:511:C:C2	1:CA:512:U:C5	3.00	0.49
2:CB:69:PHE:O	2:CB:91:PHE:HA	2.12	0.49
2:AB:151:ILE:HD12	2:AB:154:MET:SD	2.52	0.49
20:CT:67:ILE:HD11	20:CT:71:LYS:HE2	1.94	0.49
27:BF:42:GLU:O	27:BF:42:GLU:HG2	2.12	0.49
1:CA:301:G:C4	1:CA:302:G:C8	3.01	0.49
22:DA:2712:C:OP1	22:DA:2714:G:H4'	2.12	0.49
48:B0:10:ARG:HB2	48:B0:13:ARG:NH2	2.27	0.49
22:DA:2261:C:C2	22:DA:2280:G:C2	3.01	0.49
1:CA:525:C:C2	1:CA:526:C:C5	3.01	0.49
1:CA:1343:G:H2'	1:CA:1344:C:C6	2.47	0.49
44:DW:23:VAL:HG22	44:DW:38:VAL:CG1	2.42	0.49
4:AD:30:THR:O	4:AD:31:LYS:C	2.51	0.49
22:BA:2869:G:H2'	22:BA:2870:C:O4'	2.11	0.49
24:DC:232:HIS:NE2	24:DC:244:PRO:HA	2.27	0.49
53:B5:212:SER:HA	53:B5:221:PRO:CB	2.43	0.49
1:CA:465:A:C2	1:CA:466:A:C4	3.01	0.49
3:AC:58:GLU:HG3	3:AC:65:ARG:HB3	1.95	0.49
49:D1:25:LYS:NZ	49:D1:32:GLU:O	2.45	0.49
36:BO:36:TYR:N	36:BO:36:TYR:CD1	2.81	0.49
22:DA:990:A:N3	39:DR:76:LYS:NZ	2.60	0.49
1:CA:872:A:C4	1:CA:874:G:C8	3.01	0.49
26:BE:61:ARG:NH2	26:BE:64:GLY:HA3	2.27	0.49
2:AB:78:GLU:C	2:AB:80:VAL:H	2.16	0.49
1:CA:1074:G:H4'	2:CB:103:ASN:HB3	1.95	0.49
22:DA:1267:U:C5	22:DA:2012:G:N2	2.80	0.49
22:DA:1791:A:H5'	24:DC:207:LYS:O	2.13	0.49
22:DA:2091:C:H1'	45:DX:34:HIS:CD2	2.47	0.49
22:DA:176:A:N6	22:DA:177:G:O6	2.45	0.49
50:D2:12:ARG:NH2	50:D2:44:VAL:HG13	2.28	0.49
1:AA:374:A:C6	1:AA:375:U:C4	3.01	0.49
5:CE:99:ALA:O	5:CE:122:ASN:ND2	2.42	0.49
22:DA:250:G:H4'	33:DL:59:ARG:HD3	1.93	0.49
22:DA:1829:A:HO2'	24:DC:15:HIS:CD2	2.29	0.49
22:DA:1566:A:N1	24:DC:213:TRP:CE3	2.80	0.49
1:AA:956:U:C4	1:AA:957:U:C5	3.01	0.49
1:CA:1106:G:C2	1:CA:1107:C:C6	3.00	0.49
1:CA:1107:C:N3	1:CA:1108:G:N7	2.60	0.49
42:DU:82:ARG:HB2	42:DU:97:LYS:CG	2.39	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:770:G:H1'	22:DA:1379:U:C4	2.48	0.49
22:BA:1327:A:N6	22:BA:1328:A:C2	2.80	0.49
22:DA:669:G:N3	22:DA:669:G:C2'	2.75	0.49
22:DA:1351:C:O2'	22:DA:1571:A:H1'	2.11	0.49
1:CA:820:U:H4'	1:CA:821:G:OP2	2.13	0.49
22:DA:320:A:H4'	22:DA:322:A:N7	2.27	0.49
1:CA:386:C:C4	1:CA:387:U:C4	3.01	0.49
23:DB:81:G:C6	23:DB:82:U:C4	3.01	0.49
22:DA:2443:C:H2'	22:DA:2444:G:O4'	2.12	0.49
22:DA:9:G:C6	22:DA:2629:U:C5	3.01	0.49
19:AS:5:LEU:C	19:AS:6:LYS:HG3	2.33	0.49
26:BE:18:THR:HA	26:BE:106:LYS:HG2	1.95	0.49
21:AU:10:GLU:CB	21:AU:11:PRO:HD3	2.43	0.49
5:AE:50:TYR:CD1	5:AE:134:ILE:HD11	2.47	0.49
1:CA:102:G:N1	1:CA:103:U:C4	2.80	0.49
4:AD:53:VAL:CG2	4:AD:54:GLN:N	2.76	0.49
1:CA:1150:A:N6	1:CA:1151:A:H62	2.11	0.49
1:CA:1408:A:N1	1:CA:1494:G:C6	2.81	0.49
1:CA:1368:A:C8	9:CI:114:LYS:HD3	2.47	0.49
3:CC:130:PHE:CE1	3:CC:131:ARG:HD3	2.47	0.49
24:BC:144:VAL:HG12	24:BC:145:GLU:O	2.11	0.49
37:BP:2:SER:O	37:BP:6:LYS:HG2	2.13	0.49
22:DA:2636:C:H4'	25:DD:81:GLU:CD	2.33	0.49
28:DG:93:GLY:HA2	28:DG:95:ARG:NH2	2.28	0.49
45:DX:41:GLU:O	45:DX:44:LYS:HD2	2.12	0.49
22:DA:2845:U:H5''	37:DP:52:ASN:O	2.12	0.49
22:BA:340:A:H2'	22:BA:341:C:O4'	2.13	0.49
42:BU:6:ARG:HG3	42:BU:94:ARG:NH2	2.28	0.49
3:CC:192:THR:HG23	3:CC:193:TYR:CD1	2.46	0.49
2:AB:123:ASP:OD1	2:AB:123:ASP:N	2.46	0.49
1:AA:160:A:HO2'	1:AA:344:A:N6	2.10	0.49
40:DS:70:LYS:O	40:DS:107:VAL:HG23	2.13	0.49
22:DA:1359:A:N7	22:DA:1373:A:C2	2.80	0.49
39:BR:66:HIS:HE1	39:BR:94:THR:CG2	2.26	0.49
22:DA:2053:G:H21	22:DA:2054:A:H1'	1.78	0.49
1:AA:920:U:O2'	1:AA:921:U:H5'	2.13	0.49
22:DA:2816:G:N3	22:DA:2883:A:O2'	2.42	0.49
22:DA:2823:A:H2'	22:DA:2824:C:H5'	1.95	0.49
22:DA:784:G:N1	24:DC:228:VAL:HG21	2.27	0.49
12:CL:25:GLU:O	12:CL:26:ALA:C	2.49	0.49
22:DA:105:C:H2'	22:DA:106:C:H6	1.76	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:784:A:H2'	1:CA:785:G:C8	2.48	0.49
1:AA:652:U:O2'	1:AA:653:U:OP2	2.23	0.49
22:DA:1515:A:H5'	22:DA:1516:G:OP2	2.12	0.49
1:CA:1055:A:C6	1:CA:1206:G:C5	3.01	0.49
22:DA:2033:A:H4'	22:DA:2034:U:OP1	2.13	0.49
31:DJ:80:HIS:O	31:DJ:82:GLY:N	2.46	0.49
20:AT:54:MET:CE	20:AT:58:VAL:HG21	2.43	0.49
22:BA:1358:G:N2	22:BA:1374:G:C5	2.80	0.49
22:BA:251:A:OP1	33:BL:58:TYR:OH	2.16	0.49
9:CI:51:PRO:HD3	9:CI:80:ARG:HG2	1.95	0.49
1:CA:202:G:N2	1:CA:216:U:O2	2.46	0.49
22:DA:2267:A:H5''	22:DA:2268:A:H5'	1.94	0.49
22:DA:7:G:C5	22:DA:8:C:C5	3.01	0.49
47:BZ:35:THR:HG22	47:BZ:36:VAL:N	2.27	0.49
29:DH:72:ILE:O	29:DH:141:LYS:O	2.30	0.49
22:DA:2722:G:H8	22:DA:2722:G:O5'	1.96	0.49
1:CA:1365:G:H2'	1:CA:1366:C:O4'	2.13	0.49
22:BA:2749:A:OP1	28:BG:2:SER:N	2.45	0.49
34:BM:6:ARG:O	34:BM:7:THR:CG2	2.61	0.49
16:CP:39:PHE:HD1	16:CP:50:THR:HG1	1.61	0.49
22:DA:2379:G:H4'	36:DO:21:LEU:HD11	1.93	0.49
1:CA:182:A:C4	1:CA:184:G:C8	3.01	0.49
22:BA:753:A:H2'	22:BA:754:U:H6	1.77	0.49
1:CA:656:G:N2	1:CA:751:U:C2	2.81	0.49
22:BA:558:U:OP2	31:BJ:113:PRO:HG2	2.13	0.49
3:AC:150:LYS:HG3	3:AC:201:TRP:CE3	2.48	0.49
22:DA:892:A:C8	22:DA:892:A:P	3.06	0.49
42:DU:95:PHE:O	42:DU:95:PHE:CG	2.66	0.49
22:DA:1361:G:C2	22:DA:1362:C:C6	3.01	0.49
22:DA:2234:G:C5	22:DA:2235:G:C8	3.01	0.49
22:DA:1791:A:C5'	24:DC:207:LYS:O	2.60	0.49
22:DA:1790:C:H4'	24:DC:208:ALA:CB	2.42	0.49
22:DA:2683:C:C5	22:DA:2684:U:C5	3.00	0.49
22:BA:1656:C:OP1	25:BD:141:ARG:HD3	2.13	0.49
1:AA:79:G:N2	1:AA:91:U:C4	2.81	0.49
22:DA:83:A:H5''	22:DA:84:A:P	2.53	0.49
22:DA:353:C:H2'	22:DA:354:A:C8	2.47	0.49
4:CD:150:LYS:O	4:CD:151:LYS:C	2.50	0.49
5:CE:16:ILE:HD11	5:CE:38:VAL:HG23	1.93	0.49
1:AA:502:A:C2	1:AA:544:G:C2	3.01	0.49
1:CA:1413:A:N1	1:CA:1488:G:C2	2.81	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1055:A:C6	1:AA:1206:G:C4	3.01	0.49
1:AA:682:G:C2	1:AA:709:U:C2	3.00	0.49
14:AN:13:ARG:HB3	14:AN:60:GLN:HG2	1.95	0.49
22:BA:1083:U:O2	22:BA:1086:A:N1	2.46	0.49
1:CA:790:A:N6	1:CA:791:G:C6	2.81	0.49
9:AI:45:ARG:HG2	9:AI:46:MET:SD	2.53	0.49
9:AI:45:ARG:O	9:AI:48:VAL:HG22	2.12	0.49
1:AA:438:U:N3	1:AA:494:G:C6	2.81	0.49
22:DA:2817:U:O2	22:DA:2836:U:H1'	2.13	0.49
26:BE:200:LEU:O	26:BE:201:ALA:HB3	2.12	0.49
1:AA:1160:G:O2'	1:AA:1161:C:P	2.71	0.49
26:DE:5:LEU:HA	26:DE:120:VAL:O	2.12	0.49
2:CB:54:LEU:HD11	2:CB:217:VAL:HG22	1.94	0.49
1:CA:1358:U:H5''	14:CN:73:PHE:O	2.13	0.49
41:DT:82:LYS:HG2	41:DT:83:ALA:H	1.78	0.49
1:CA:1150:A:N6	1:CA:1151:A:N6	2.61	0.49
36:BO:41:ALA:HB2	36:BO:48:LEU:HD21	1.95	0.49
1:CA:872:A:C5	1:CA:874:G:C8	3.01	0.49
22:BA:754:U:H2'	22:BA:755:U:C6	2.47	0.49
22:DA:1320:C:N4	22:DA:1333:G:C6	2.81	0.49
1:AA:293:G:C4	1:AA:294:U:C5	3.01	0.49
22:DA:428:A:H2'	22:DA:429:A:O4'	2.13	0.49
24:DC:84:ASP:OD2	24:DC:87:ARG:NE	2.45	0.49
24:DC:226:ASN:HB3	24:DC:227:PRO:HD2	1.94	0.49
33:BL:30:THR:O	33:BL:33:ARG:HG2	2.13	0.49
22:DA:321:U:H5''	26:DE:131:THR:HG23	1.95	0.49
22:BA:2125:G:N2	22:BA:2171:A:O5'	2.46	0.49
3:AC:79:LYS:O	3:AC:82:GLU:HG3	2.12	0.49
46:BY:11:VAL:O	46:BY:15:ASN:ND2	2.46	0.49
22:DA:1812:U:H2'	22:DA:1812:U:O2	2.13	0.49
1:AA:1164:G:C2	1:AA:1173:U:O2	2.65	0.49
22:BA:1700:A:H5'	22:BA:1701:A:OP2	2.13	0.49
1:CA:57:G:C6	1:CA:58:C:N4	2.81	0.49
1:AA:553:A:O2'	1:AA:554:A:H5'	2.12	0.49
22:DA:1606:C:C2'	22:DA:1607:C:OP2	2.60	0.49
28:BG:67:THR:O	28:BG:71:LEU:HG	2.13	0.49
24:DC:141:VAL:HG13	24:DC:191:THR:O	2.13	0.49
22:BA:1747:U:H2'	22:BA:1748:C:C6	2.48	0.49
22:BA:1287:A:H5'	35:BN:103:ARG:HD2	1.94	0.49
22:DA:2247:A:H3'	57:DA:3505:HOH:O	2.13	0.49
22:DA:1651:G:C2	22:DA:2007:U:C2	3.01	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:88:G:C2	22:DA:89:A:C8	3.01	0.49
1:CA:756:C:O2'	1:CA:757:U:H5'	2.13	0.49
1:AA:709:U:H2'	1:AA:710:G:C8	2.47	0.49
24:BC:230:HIS:O	24:BC:232:HIS:N	2.45	0.49
9:AI:44:ALA:O	9:AI:47:VAL:HG22	2.13	0.49
22:DA:1856:U:O4	22:DA:1857:G:C6	2.66	0.49
16:CP:20:VAL:HG21	16:CP:32:PHE:CG	2.48	0.49
22:DA:680:C:C2	22:DA:681:G:N7	2.81	0.49
1:CA:1314:C:OP2	19:CS:6:LYS:HG2	2.13	0.49
1:CA:158:G:C5	1:CA:159:G:N7	2.81	0.49
49:B1:25:LYS:HD2	49:B1:52:ALA:HB1	1.95	0.49
28:BG:11:VAL:CG2	28:BG:11:VAL:O	2.60	0.49
22:BA:613:A:H2'	22:BA:614:A:H5''	1.94	0.49
25:DD:7:LYS:HG2	25:DD:8:LYS:N	2.28	0.49
1:CA:1392:G:C2'	1:CA:1393:U:H5'	2.42	0.49
42:BU:5:ILE:C	42:BU:6:ARG:HG2	2.33	0.49
22:DA:231:A:N6	22:DA:232:G:C2	2.80	0.49
1:CA:1250:A:N6	1:CA:1251:A:C6	2.81	0.49
36:DO:53:THR:O	36:DO:59:ALA:HB2	2.13	0.49
46:DY:42:LEU:O	46:DY:46:VAL:HG23	2.13	0.49
1:AA:32:A:OP1	1:AA:398:U:H1'	2.13	0.49
18:CR:33:ILE:HA	18:CR:40:VAL:HG23	1.95	0.49
22:DA:2001:C:H4'	22:DA:2689:U:H2'	1.94	0.49
39:BR:76:LYS:O	39:BR:84:ARG:HA	2.13	0.49
22:BA:2323:G:C2'	22:BA:2324:U:H5'	2.43	0.49
43:BV:6:ALA:HB1	43:BV:40:ILE:HG23	1.94	0.49
22:DA:655:A:H4'	22:DA:656:G:OP1	2.13	0.49
22:DA:672:C:C2	22:DA:809:G:N2	2.81	0.49
39:DR:66:HIS:CE1	39:DR:94:THR:HG21	2.47	0.49
1:AA:1422:G:O3'	32:BK:49:ARG:NH2	2.46	0.49
22:BA:2360:G:H1'	33:BL:60:ARG:HD3	1.95	0.49
22:DA:2704:C:H2'	22:DA:2705:A:O4'	2.13	0.49
22:DA:1009:A:N3	22:DA:1153:C:O2'	2.36	0.48
1:CA:56:U:H2'	1:CA:57:G:C8	2.48	0.48
1:CA:1093:A:O2'	1:CA:1095:U:OP1	2.19	0.48
1:CA:1096:C:H2'	1:CA:1097:C:C6	2.48	0.48
1:CA:505:G:H5'	1:CA:534:U:C2	2.48	0.48
29:DH:21:VAL:CG2	29:DH:22:LYS:N	2.76	0.48
22:DA:1208:C:C2	22:DA:1209:U:C5	3.01	0.48
21:CU:39:GLU:HA	21:CU:42:THR:OG1	2.13	0.48
22:DA:1315:C:O2'	22:DA:1392:A:N3	2.41	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:1364:G:C8	45:DX:2:SER:N	2.81	0.48
24:DC:48:ARG:HG3	24:DC:48:ARG:HH11	1.78	0.48
24:DC:45:ASN:O	24:DC:47:GLY:N	2.46	0.48
1:AA:1121:U:C2	1:AA:1122:U:C5	3.01	0.48
36:DO:80:GLU:HA	36:DO:83:LEU:HD12	1.95	0.48
1:AA:1145:A:HO2'	1:AA:1146:A:P	2.36	0.48
42:DU:71:ALA:HB1	42:DU:81:ASP:O	2.12	0.48
35:BN:77:ALA:O	35:BN:79:LEU:O	2.31	0.48
22:DA:46:G:C2	22:DA:47:C:C6	3.00	0.48
12:CL:3:THR:O	12:CL:4:VAL:C	2.51	0.48
12:AL:3:THR:HG22	12:AL:4:VAL:N	2.28	0.48
3:AC:113:ALA:O	3:AC:116:VAL:HB	2.13	0.48
8:AH:106:THR:HG22	8:AH:122:GLY:O	2.13	0.48
22:DA:1862:G:C2	22:DA:1881:C:C2	3.01	0.48
8:AH:111:MET:HE2	8:AH:116:ALA:N	2.28	0.48
15:CO:64:ARG:NH1	15:CO:68:ASP:OD1	2.44	0.48
1:CA:537:G:H2'	1:CA:538:G:C8	2.48	0.48
19:CS:29:LYS:CB	19:CS:30:PRO:HD2	2.43	0.48
46:DY:60:LYS:HA	46:DY:63:ALA:OXT	2.13	0.48
40:BS:73:LYS:HB2	40:BS:106:VAL:HB	1.95	0.48
12:CL:44:LYS:HB2	12:CL:45:PRO:HD3	1.94	0.48
31:DJ:138:GLN:HG3	31:DJ:138:GLN:O	2.13	0.48
22:DA:1029:A:N1	22:DA:2465:C:O2'	2.35	0.48
1:AA:627:G:OP1	16:AP:51:ARG:NH2	2.46	0.48
36:DO:18:LEU:O	36:DO:22:GLY:N	2.46	0.48
22:DA:2032:G:C2	25:DD:150:GLN:HG2	2.48	0.48
39:BR:49:ILE:CA	39:BR:51:VAL:O	2.61	0.48
22:BA:1079:C:C5	22:BA:1088:A:C2	3.01	0.48
1:CA:685:G:C2	1:CA:686:U:C4	3.01	0.48
1:AA:914:A:C2	1:AA:915:A:C8	3.00	0.48
1:AA:914:A:C6	1:AA:915:A:N7	2.81	0.48
5:AE:81:LEU:HB3	5:AE:147:MET:HE3	1.95	0.48
3:CC:36:ASP:O	3:CC:40:ARG:HG3	2.13	0.48
1:CA:123:U:H2'	1:CA:124:C:C6	2.48	0.48
33:BL:81:ASP:O	33:BL:83:ALA:N	2.43	0.48
2:AB:67:ILE:O	2:AB:68:LEU:CB	2.60	0.48
53:B5:52:PRO:HB2	53:B5:204:GLY:O	2.13	0.48
24:BC:143:ASN:OD1	24:BC:152:GLY:HA3	2.13	0.48
22:DA:2773:C:H2'	22:DA:2774:C:H6	1.78	0.48
22:DA:1462:C:N3	22:DA:1463:C:C5	2.80	0.48
21:AU:16:LEU:C	21:AU:18:ARG:HE	2.16	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:76:G:C2	1:CA:95:C:N3	2.82	0.48
22:BA:1250:G:OP2	33:BL:21:ARG:NH2	2.46	0.48
1:AA:1347:G:N2	1:AA:1373:G:H2'	2.28	0.48
9:AI:114:LYS:HG2	9:AI:120:LYS:HA	1.95	0.48
40:DS:47:VAL:HG23	40:DS:103:ILE:CG2	2.42	0.48
22:DA:632:A:H4'	33:DL:68:SER:HB2	1.94	0.48
1:CA:681:A:C6	1:CA:710:G:C6	3.01	0.48
39:DR:66:HIS:CD2	39:DR:94:THR:HG22	2.48	0.48
15:AO:88:ARG:O	15:AO:89:ARG:HB2	2.13	0.48
1:AA:830:G:H2'	1:AA:831:A:C8	2.49	0.48
22:BA:1823:G:N7	57:BA:3663:HOH:O	2.34	0.48
22:DA:2736:A:C2	22:DA:2769:U:O2	2.66	0.48
1:CA:1300:G:C6	1:CA:1335:U:C6	3.02	0.48
22:BA:1820:U:OP1	24:BC:177:ARG:NH2	2.45	0.48
27:DF:111:ILE:HB	27:DF:114:PHE:HB2	1.94	0.48
22:BA:2897:U:H2'	22:BA:2898:U:C6	2.48	0.48
29:BH:139:PHE:O	29:BH:140:ALA:HB2	2.14	0.48
1:AA:1226:C:H4'	1:AA:1227:A:OP1	2.12	0.48
1:AA:1226:C:H2'	13:AM:102:THR:HB	1.95	0.48
22:DA:2823:A:C5	22:DA:2824:C:C5	3.01	0.48
32:BK:70:ARG:HD3	32:BK:76:VAL:CG2	2.43	0.48
22:DA:183:C:N4	22:DA:184:C:C4	2.81	0.48
31:DJ:41:LYS:CE	31:DJ:52:ASP:OD1	2.62	0.48
22:BA:1379:U:C6	22:BA:1379:U:OP1	2.67	0.48
50:D2:11:LYS:O	50:D2:14:ARG:N	2.46	0.48
37:DP:113:ARG:O	37:DP:114:LEU:C	2.52	0.48
1:CA:821:G:H2'	1:CA:822:U:H6	1.76	0.48
1:AA:1157:A:C6	1:AA:1180:A:C6	3.01	0.48
7:AG:146:GLU:O	7:AG:149:LYS:CB	2.61	0.48
1:CA:1479:C:C2	1:CA:1480:A:C8	3.01	0.48
22:BA:1007:C:OP1	31:BJ:39:LYS:HE2	2.14	0.48
22:DA:1856:U:C4	22:DA:1857:G:C6	3.01	0.48
22:BA:1795:C:C2	22:BA:1796:U:C6	3.01	0.48
22:DA:136:G:N2	22:DA:144:A:C5	2.81	0.48
22:DA:630:G:H5''	22:DA:631:A:OP2	2.12	0.48
22:DA:1155:A:H5''	38:DQ:55:ARG:HD3	1.95	0.48
33:BL:59:ARG:NH1	57:BL:306:HOH:O	2.38	0.48
22:DA:1665:A:OP1	32:DK:66:LYS:HE2	2.13	0.48
31:BJ:23:LYS:CE	31:BJ:142:ILE:OXT	2.62	0.48
20:AT:83:ILE:O	20:AT:87:ALA:HB3	2.13	0.48
1:CA:102:G:N2	1:CA:103:U:C2	2.82	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:AM:40:ALA:HB3	13:AM:43:VAL:CG1	2.43	0.48
1:CA:1328:C:OP1	13:CM:28:THR:HG21	2.13	0.48
22:DA:2852:G:C2	22:DA:2853:C:C2	3.02	0.48
1:CA:740:U:O2'	1:CA:741:G:H5'	2.13	0.48
42:BU:89:ASP:CG	42:BU:90:GLY:H	2.15	0.48
22:DA:2539:C:H4'	52:D4:3:VAL:HG11	1.95	0.48
28:BG:94:TYR:HA	28:BG:106:SER:O	2.14	0.48
6:AF:97:THR:O	6:AF:98:GLU:CG	2.61	0.48
22:DA:1509:A:O2'	22:DA:1510:G:OP2	2.26	0.48
22:DA:1512:C:C4	22:DA:1513:U:C4	3.00	0.48
1:CA:865:A:H2	1:CA:918:A:H4'	1.79	0.48
12:AL:79:VAL:O	12:AL:102:LEU:HB3	2.13	0.48
1:CA:127:G:N2	1:CA:235:C:C2	2.80	0.48
22:DA:2119:A:N6	22:DA:2167:U:O2'	2.46	0.48
1:AA:1410:A:H2'	1:AA:1411:C:C6	2.48	0.48
22:BA:1714:U:O2	22:BA:1714:U:H2'	2.13	0.48
22:BA:464:U:H5'	50:B2:5:PHE:CD1	2.48	0.48
4:CD:51:TYR:O	4:CD:52:GLY:C	2.52	0.48
1:AA:620:C:H2'	1:AA:621:A:O4'	2.13	0.48
35:DN:34:ILE:HD11	35:DN:44:LEU:CD2	2.43	0.48
24:BC:76:ALA:HB2	24:BC:96:TYR:CD2	2.49	0.48
22:BA:1169:A:H2'	22:BA:1170:C:O4'	2.13	0.48
1:AA:1225:A:H2'	1:AA:1226:C:C6	2.48	0.48
35:BN:12:ARG:CZ	35:BN:20:MET:CE	2.91	0.48
4:AD:95:GLU:OE1	4:AD:191:LEU:HD22	2.13	0.48
4:AD:34:ILE:O	4:AD:35:GLU:CB	2.60	0.48
22:DA:1805:A:C2	22:DA:1813:G:N1	2.81	0.48
4:CD:124:MET:O	4:CD:143:VAL:HA	2.13	0.48
50:B2:43:THR:O	50:B2:44:VAL:CG1	2.61	0.48
4:CD:3:ARG:O	4:CD:5:LEU:HD13	2.13	0.48
48:B0:34:SER:OG	48:B0:36:GLU:CG	2.61	0.48
22:DA:527:C:H2'	22:DA:2779:U:O2	2.13	0.48
22:DA:1252:G:C4	22:DA:1253:A:C2	3.01	0.48
1:AA:951:G:H1'	1:AA:1231:G:N2	2.27	0.48
27:BF:171:ALA:O	27:BF:173:PHE:N	2.46	0.48
1:CA:801:U:C2'	1:CA:802:A:O5'	2.61	0.48
5:CE:25:VAL:O	5:CE:26:LYS:C	2.51	0.48
1:CA:1306:A:C5	1:CA:1307:U:C4	3.01	0.48
1:AA:737:C:H2'	1:AA:738:C:C6	2.49	0.48
29:DH:127:GLU:HG3	29:DH:144:VAL:O	2.13	0.48
1:CA:216:U:H4'	1:CA:464:U:H4'	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:1012:A:C6	1:CA:1013:G:C6	3.02	0.48
1:AA:36:C:H2'	1:AA:37:U:O4'	2.14	0.48
1:AA:575:G:C6	1:AA:821:G:C8	3.01	0.48
48:B0:25:VAL:O	48:B0:27:SER:N	2.47	0.48
22:BA:1644:C:O2	22:BA:1644:C:H2'	2.13	0.48
32:BK:34:GLY:O	32:BK:35:VAL:C	2.51	0.48
15:CO:76:ALA:O	15:CO:80:GLN:HG3	2.14	0.48
9:CI:49:ARG:CZ	9:CI:53:GLU:HG2	2.43	0.48
14:AN:46:LEU:O	14:AN:48:LEU:N	2.46	0.48
6:CF:50:PRO:HD2	18:CR:74:HIS:HB3	1.95	0.48
22:BA:1967:C:H2'	22:BA:1968:G:H5'	1.96	0.48
22:DA:1931:U:O2	22:DA:1931:U:H2'	2.14	0.48
1:CA:878:A:C5	1:CA:879:C:C5	3.01	0.48
1:AA:1086:U:O2'	1:AA:1087:G:H5'	2.13	0.48
1:CA:1140:C:O2'	1:CA:1141:C:O5'	2.31	0.48
46:DY:9:LYS:HG2	46:DY:10:SER:N	2.29	0.48
22:DA:1153:C:C4	22:DA:1154:G:C5	3.00	0.48
22:BA:1073:A:OP1	22:BA:1073:A:C8	2.67	0.48
1:CA:1071:C:H2'	1:CA:1072:G:H8	1.79	0.48
22:DA:1360:G:N1	22:DA:1361:G:H1'	2.29	0.48
45:DX:28:ARG:NH1	45:DX:30:LEU:HD21	2.29	0.48
1:CA:1296:C:H4'	1:CA:1302:C:C4	2.49	0.48
22:BA:1657:U:OP2	25:BD:141:ARG:HG3	2.14	0.48
1:AA:203:G:C2	1:AA:215:C:C2	3.01	0.48
22:BA:996:A:N6	22:BA:1160:G:C6	2.81	0.48
22:DA:2064:C:H2'	22:DA:2065:C:C6	2.48	0.48
22:DA:2199:A:C6	22:DA:2200:C:N3	2.82	0.48
22:BA:2502:G:H5'	22:BA:2503:A:C5'	2.43	0.48
51:B3:27:ALA:O	51:B3:28:ASN:CB	2.61	0.48
15:CO:88:ARG:O	15:CO:89:ARG:HB2	2.13	0.48
24:DC:212:ARG:CD	24:DC:216:VAL:O	2.62	0.48
33:BL:62:PRO:HG2	51:B3:25:LYS:CD	2.42	0.48
1:CA:760:G:N7	1:CA:761:G:C8	2.81	0.48
24:DC:260:ASN:OD1	24:DC:262:ARG:HB3	2.14	0.48
22:DA:636:G:N2	33:DL:76:GLU:OE2	2.46	0.48
22:DA:2370:G:O6	22:DA:2371:G:C6	2.67	0.48
8:AH:14:ILE:O	8:AH:15:ARG:C	2.51	0.48
22:DA:1907:G:C2	22:DA:1924:C:C2	3.01	0.48
22:DA:1403:A:C2	22:DA:1404:C:C2	3.02	0.48
1:AA:41:G:H2'	1:AA:42:G:H8	1.78	0.48
3:CC:148:GLY:O	3:CC:203:PHE:N	2.43	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:CD:119:SER:O	4:CD:131:ASN:OD1	2.31	0.48
22:DA:1659:G:C6	22:DA:1660:G:N7	2.82	0.48
22:DA:993:G:H1'	39:DR:91:GLN:OE1	2.13	0.48
22:DA:1446:C:N4	22:DA:1447:C:N4	2.61	0.48
25:BD:33:ARG:NH2	25:BD:74:GLU:O	2.42	0.48
35:BN:28:LEU:O	35:BN:32:GLU:N	2.43	0.48
1:CA:389:A:C6	1:CA:390:U:H1'	2.48	0.48
33:DL:20:GLY:N	33:DL:27:LEU:O	2.45	0.48
1:AA:1275:A:H2'	1:AA:1276:G:O4'	2.13	0.48
22:DA:1153:C:C4	22:DA:1154:G:C6	3.02	0.48
39:BR:49:ILE:HB	39:BR:51:VAL:O	2.13	0.48
2:AB:82:ASP:C	2:AB:84:ALA:N	2.64	0.48
2:CB:102:THR:HG23	2:CB:102:THR:O	2.13	0.48
1:AA:701:U:C2	1:AA:703:G:C2	3.01	0.48
21:CU:37:PHE:HA	21:CU:40:LYS:HE3	1.96	0.48
24:DC:228:VAL:CG1	24:DC:229:ASP:N	2.76	0.48
1:CA:325:A:N6	1:CA:326:G:C6	2.81	0.48
22:DA:591:U:H2'	22:DA:592:A:O4'	2.14	0.48
22:DA:90:U:H5	22:DA:91:A:HO2'	1.60	0.48
26:DE:24:ASN:ND2	26:DE:27:LEU:HB3	2.28	0.48
22:DA:1425:G:H2'	22:DA:1426:G:O4'	2.13	0.48
24:DC:65:VAL:HG12	24:DC:67:PHE:CD2	2.49	0.48
1:CA:1484:C:H2'	1:CA:1485:U:O4'	2.13	0.48
22:DA:1855:U:C4	22:DA:1856:U:C4	3.01	0.48
1:CA:634:C:N4	1:CA:635:A:N6	2.61	0.48
22:DA:734:A:C4	22:DA:735:A:C8	3.01	0.48
22:DA:1914:C:H5	22:DA:1915:U:C2	2.31	0.48
30:BI:76:ALA:HB1	30:BI:129:ILE:HG23	1.95	0.48
22:DA:2359:C:O2'	51:D3:54:ASP:OD2	2.19	0.48
22:BA:372:G:O2'	22:BA:400:G:O6	2.25	0.48
22:BA:1439:A:C2	22:BA:1553:A:C4	3.02	0.48
2:CB:68:LEU:HD12	2:CB:158:PRO:HG2	1.94	0.48
2:CB:35:ARG:O	2:CB:37:LYS:N	2.47	0.48
1:CA:525:C:N3	1:CA:526:C:C5	2.82	0.48
9:CI:28:ILE:HB	9:CI:35:LEU:HB2	1.93	0.48
24:DC:94:VAL:HG11	24:DC:104:ILE:HD11	1.95	0.48
24:BC:65:VAL:HG11	24:BC:67:PHE:CZ	2.49	0.48
22:DA:1709:U:H2'	22:DA:1710:G:C8	2.48	0.48
22:DA:262:A:H4'	22:DA:611:C:OP1	2.13	0.48
27:BF:57:LEU:HD21	27:BF:152:LEU:HD11	1.95	0.48
22:BA:1187:G:H5"	39:BR:83:TYR:CE1	2.49	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:AO:30:ALA:HA	15:AO:85:LEU:HD21	1.95	0.48
13:AM:104:THR:O	13:AM:105:ASN:C	2.52	0.48
22:BA:595:C:H2'	22:BA:596:U:C6	2.49	0.48
22:DA:1567:G:H2'	24:DC:85:PRO:HG3	1.96	0.48
22:BA:2344:U:H4'	22:BA:2345:G:OP1	2.13	0.48
22:BA:1619:G:C8	22:BA:1619:G:OP2	2.66	0.48
3:AC:97:VAL:HB	3:AC:98:PRO:HD2	1.95	0.48
24:BC:118:SER:HB2	24:BC:129:THR:HB	1.94	0.48
1:AA:781:A:C5	1:AA:802:A:C2	3.01	0.48
22:BA:1246:A:C2'	22:BA:1247:A:O5'	2.61	0.48
29:BH:103:VAL:HG21	29:BH:132:PHE:CZ	2.49	0.48
22:DA:996:A:C2	22:DA:997:G:C8	3.01	0.48
1:CA:1177:G:O6	1:CA:1178:G:C6	2.67	0.48
4:AD:190:ASP:O	4:AD:191:LEU:O	2.31	0.48
11:AK:126:LYS:N	11:AK:126:LYS:HD3	2.29	0.48
24:DC:162:VAL:HG12	24:DC:163:GLN:N	2.28	0.48
2:AB:71:GLY:HA2	2:AB:164:ILE:HG22	1.95	0.48
22:DA:1784:A:H4'	22:DA:1785:A:O5'	2.13	0.48
22:DA:1270:C:N4	22:DA:1648:U:O4	2.46	0.48
24:DC:212:ARG:HD3	24:DC:218:PRO:HD3	1.94	0.48
22:BA:2313:C:H5"	27:BF:88:LYS:HD3	1.96	0.48
13:AM:33:ILE:HD12	13:AM:59:GLU:HB3	1.96	0.48
2:AB:90:PHE:HB3	2:AB:150:GLY:O	2.13	0.48
1:AA:945:G:C2	1:AA:946:A:C8	3.01	0.48
40:DS:79:GLY:CA	40:DS:100:THR:O	2.61	0.48
22:DA:2461:A:C2	22:DA:2490:G:N2	2.81	0.48
36:BO:31:THR:HG23	36:BO:34:HIS:H	1.79	0.48
2:AB:139:ARG:HG3	2:AB:140:GLU:N	2.29	0.48
1:CA:1227:A:OP2	13:CM:110:LYS:HE3	2.13	0.48
42:DU:74:ASN:HB2	42:DU:81:ASP:OD2	2.13	0.48
1:AA:594:U:C4	1:AA:595:A:C6	3.01	0.48
10:CJ:5:ARG:HG3	10:CJ:6:ILE:HG13	1.96	0.48
33:DL:68:SER:O	33:DL:69:ARG:HB2	2.13	0.48
20:AT:68:HIS:C	20:AT:69:LYS:HZ2	2.16	0.48
1:CA:572:A:H5'	1:CA:573:A:OP2	2.14	0.48
44:DW:38:VAL:HG23	44:DW:59:LEU:HB2	1.95	0.48
22:DA:807:U:OP2	33:DL:41:ARG:NH1	2.47	0.48
22:DA:2469:A:O2'	34:DM:55:ARG:NH2	2.46	0.48
22:DA:1838:C:C6	22:DA:1899:A:C6	3.02	0.48
1:CA:1118:U:H1'	1:CA:1179:A:C4	2.48	0.48
40:BS:41:LYS:HE3	48:B0:22:LEU:HD21	1.96	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:BI:122:ILE:O	30:BI:122:ILE:HG22	2.14	0.48
1:CA:295:C:C4	1:CA:296:U:C5	3.02	0.48
25:BD:8:LYS:NZ	25:BD:195:GLY:O	2.32	0.48
8:CH:86:TYR:CE1	8:CH:124:GLU:HB2	2.48	0.48
22:BA:593:U:H2'	22:BA:594:U:C6	2.49	0.48
39:DR:101:ILE:O	39:DR:103:ALA:N	2.47	0.48
1:AA:4:U:O2	1:AA:4:U:H2'	2.13	0.48
10:AJ:10:LEU:HG	10:AJ:98:VAL:HG12	1.96	0.48
1:AA:1322:C:OP1	19:AS:78:ARG:NH2	2.47	0.48
22:DA:246:C:H2'	22:DA:247:G:H5'	1.96	0.48
22:DA:192:C:P	57:DA:3739:HOH:O	2.68	0.48
22:DA:511:U:C5	22:DA:512:G:C5	3.02	0.48
12:CL:107:VAL:HG23	12:CL:117:TYR:HB3	1.96	0.48
22:DA:197:A:N6	22:DA:2430:A:H2'	2.28	0.48
22:DA:825:A:C6	22:DA:826:U:C4	3.01	0.48
1:AA:858:G:O6	1:AA:869:G:C8	2.66	0.48
37:DP:91:ALA:HB2	37:DP:113:ARG:HA	1.95	0.48
22:BA:2356:U:O3'	44:BW:20:ARG:HD3	2.13	0.48
1:CA:728:A:C6	1:CA:729:A:N6	2.82	0.48
22:DA:2131:U:H5'	22:DA:2132:U:H5''	1.95	0.48
1:CA:782:A:C2'	1:CA:783:C:H5'	2.43	0.48
1:AA:345:C:N3	32:BK:117:SER:OG	2.44	0.48
22:BA:1359:A:P	57:BA:3622:HOH:O	2.72	0.48
1:AA:1228:C:OP2	13:AM:107:ARG:NH2	2.42	0.48
1:CA:775:G:C4	1:CA:776:G:C8	3.01	0.48
22:BA:64:A:H2'	22:BA:65:U:C6	2.49	0.48
22:DA:1866:A:N7	22:DA:1867:G:C8	2.81	0.48
1:CA:1327:C:C4	1:CA:1328:C:N4	2.82	0.48
1:AA:617:G:C2	1:AA:618:C:C6	3.02	0.48
1:AA:1405:G:O4'	1:AA:1519:A:H4'	2.13	0.48
1:CA:456:A:C6	1:CA:457:G:C5	3.01	0.48
20:AT:68:HIS:HB3	20:AT:69:LYS:HE3	1.94	0.48
22:DA:2050:C:N4	22:DA:2051:A:N1	2.62	0.48
22:DA:2447:G:C8	22:DA:2500:U:C6	3.02	0.48
1:AA:575:G:C6	1:AA:821:G:N7	2.82	0.48
24:BC:141:VAL:N	24:BC:162:VAL:O	2.46	0.48
1:CA:458:U:H2'	1:CA:459:A:C8	2.48	0.48
22:DA:448:U:H5''	57:DA:3242:HOH:O	2.12	0.48
22:BA:1489:C:C2	22:BA:1501:G:C2	3.01	0.48
29:BH:135:HIS:CD2	29:BH:137:GLU:HG3	2.48	0.48
12:CL:58:THR:HG22	12:CL:59:ASN:N	2.28	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:BT:12:ARG:HB2	41:BT:33:LYS:O	2.13	0.48
11:CK:107:ILE:O	11:CK:107:ILE:HG23	2.13	0.48
4:AD:161:LEU:HD22	4:AD:161:LEU:N	2.29	0.48
9:CI:118:LEU:HD12	9:CI:118:LEU:N	2.29	0.48
1:CA:313:A:H2'	1:CA:314:C:C6	2.49	0.48
46:BY:32:ALA:HB2	46:BY:37:LEU:CD2	2.44	0.48
22:BA:332:A:C2	22:BA:335:C:C5	3.02	0.48
1:AA:716:A:H2'	1:AA:717:U:O5'	2.14	0.48
1:AA:539:A:N6	1:AA:540:G:O6	2.47	0.48
22:DA:1009:A:OP2	31:DJ:39:LYS:HE3	2.14	0.48
39:BR:49:ILE:CB	39:BR:51:VAL:O	2.62	0.48
22:DA:1358:G:H2'	22:DA:1359:A:OP2	2.14	0.48
22:DA:842:U:C2	22:DA:843:G:C8	3.02	0.48
4:CD:147:GLU:O	4:CD:150:LYS:HB3	2.13	0.48
22:DA:1351:C:H2'	22:DA:1352:U:O4'	2.14	0.48
1:CA:756:C:C4	1:CA:757:U:C5	3.01	0.48
1:AA:64:G:C2	1:AA:67:C:N4	2.82	0.48
15:AO:19:ALA:O	15:AO:20:ASN:CB	2.62	0.48
22:BA:2886:A:C4	22:BA:2887:A:C8	3.02	0.48
1:CA:954:G:C5	1:CA:955:U:C4	3.02	0.48
1:CA:954:G:C6	1:CA:955:U:C4	3.02	0.48
22:DA:1076:C:H1'	30:DI:93:PRO:HG2	1.95	0.48
13:CM:22:ILE:HG23	13:CM:66:GLU:HG2	1.96	0.48
33:DL:112:LEU:CD1	33:DL:134:ALA:HB2	2.44	0.48
22:DA:920:A:C6	22:DA:921:C:C4	3.02	0.48
22:DA:2272:U:H5''	22:DA:2273:A:OP1	2.14	0.48
45:DX:7:VAL:HG21	45:DX:59:ILE:HD11	1.95	0.48
13:AM:88:GLY:C	13:AM:90:ARG:N	2.66	0.48
4:CD:72:PHE:CE2	4:CD:200:ILE:HD11	2.49	0.48
22:DA:2651:C:O2'	22:DA:2652:C:H5'	2.14	0.48
22:BA:2492:U:H2'	22:BA:2493:U:H6	1.77	0.48
39:DR:76:LYS:HB2	39:DR:85:LYS:HB2	1.96	0.48
35:BN:28:LEU:HD23	35:BN:48:VAL:HG11	1.96	0.48
13:CM:114:LYS:HB2	13:CM:115:PRO:HD3	1.96	0.48
22:DA:1231:U:H2'	22:DA:1232:G:C8	2.48	0.48
5:AE:94:VAL:HG21	5:AE:140:THR:HG22	1.95	0.48
47:DZ:16:ARG:O	47:DZ:21:LYS:NZ	2.31	0.48
22:DA:1428:C:C4	22:DA:1569:A:H5''	2.48	0.48
22:DA:323:C:H2'	26:DE:163:ASN:OD1	2.13	0.48
2:CB:164:ILE:O	2:CB:186:ILE:O	2.32	0.48
7:CG:88:PRO:HD2	7:CG:151:PHE:O	2.14	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:AF:52:ASN:O	6:AF:53:LYS:CB	2.61	0.48
38:DQ:76:TYR:CZ	38:DQ:80:ILE:HG13	2.48	0.48
22:DA:280:U:H2'	22:DA:281:C:C6	2.49	0.48
6:AF:38:ARG:HB3	6:AF:63:ASN:HB2	1.96	0.48
22:BA:74:A:H5''	22:BA:74:A:N3	2.28	0.48
48:D0:38:HIS:CD2	48:D0:44:THR:HG22	2.48	0.48
13:CM:55:THR:O	13:CM:59:GLU:OE2	2.31	0.48
39:BR:51:VAL:CB	39:BR:52:PRO:CD	2.91	0.48
22:DA:370:G:C6	22:DA:424:G:C5	3.02	0.48
22:DA:1250:G:H5'	38:DQ:6:ARG:HD2	1.95	0.48
24:DC:107:PRO:HD2	24:DC:110:LEU:HD22	1.94	0.48
22:DA:177:G:H3'	22:DA:178:G:C8	2.49	0.48
22:BA:2129:C:H2'	22:BA:2130:U:C6	2.49	0.48
1:AA:1077:G:C6	57:AA:1793:HOH:O	2.59	0.48
22:DA:1566:A:C2	24:DC:213:TRP:CE3	3.02	0.48
22:DA:197:A:C8	22:DA:2430:A:C5	3.02	0.48
5:AE:122:ASN:C	5:AE:122:ASN:HD22	2.17	0.48
8:CH:114:ARG:HG2	8:CH:115:ALA:N	2.28	0.48
1:CA:716:A:C6	1:CA:717:U:N3	2.82	0.48
22:BA:2554:U:H2'	22:BA:2555:U:C6	2.49	0.48
22:DA:2286:G:H5''	22:DA:2287:A:OP1	2.14	0.48
7:AG:146:GLU:O	7:AG:149:LYS:HB3	2.13	0.48
22:DA:1430:G:C4	22:DA:1431:A:C8	3.02	0.48
20:CT:83:ILE:HD12	20:CT:84:ASN:N	2.29	0.48
1:CA:72:A:N6	1:CA:73:C:H42	2.12	0.48
22:DA:2854:G:C2	22:DA:2864:G:C2	3.02	0.48
1:AA:689:C:H2'	1:AA:690:G:O4'	2.14	0.48
22:DA:649:G:H2'	22:DA:650:C:O4'	2.14	0.48
1:AA:184:G:C6	1:AA:185:U:O4	2.66	0.48
22:BA:2294:G:H5''	36:BO:10:ARG:HD3	1.96	0.48
42:DU:28:VAL:HG23	42:DU:34:VAL:HG12	1.95	0.48
5:AE:74:VAL:O	5:AE:76:LEU:HD12	2.14	0.48
1:AA:1371:G:P	9:AI:13:LYS:HD3	2.54	0.48
12:AL:35:THR:HG22	12:AL:36:ARG:HD2	1.95	0.48
22:DA:325:G:O2'	22:DA:326:G:H5'	2.14	0.48
22:DA:752:A:N3	22:DA:752:A:H2'	2.28	0.48
2:CB:200:ILE:N	2:CB:200:ILE:HD12	2.29	0.48
22:DA:1960:A:O2'	22:DA:1961:C:H5'	2.14	0.48
26:BE:15:SER:N	26:BE:197:GLU:OE2	2.47	0.48
17:CQ:24:ALA:HA	17:CQ:43:LYS:HA	1.96	0.48
42:DU:72:ILE:CD1	42:DU:83:VAL:HG23	2.43	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:363:A:OP1	12:CL:31:ARG:N	2.43	0.48
1:CA:1288:A:N1	1:CA:1371:G:H1'	2.28	0.48
24:DC:154:LEU:HD13	24:DC:176:LEU:HD21	1.95	0.48
35:BN:8:ARG:HB3	35:BN:10:LEU:HG	1.95	0.48
1:AA:129:A:O2'	1:AA:130:A:H5''	2.14	0.48
49:D1:26:ASN:O	49:D1:27:LYS:C	2.53	0.48
1:CA:110:C:H2'	1:CA:111:G:O4'	2.14	0.48
24:BC:209:GLY:O	24:BC:210:ALA:C	2.52	0.48
22:BA:1605:C:C2'	22:BA:1606:C:H5'	2.44	0.48
22:BA:1317:G:C2	22:BA:1336:A:C2	3.02	0.48
39:DR:87:GLN:HG2	39:DR:88:GLY:N	2.29	0.48
17:AQ:15:ASP:HA	17:AQ:21:ILE:CD1	2.42	0.47
22:DA:1335:C:H2'	22:DA:1336:A:C8	2.49	0.47
22:DA:188:G:C6	22:DA:189:G:C4	3.02	0.47
22:BA:278:A:N1	22:BA:362:A:C8	2.82	0.47
22:DA:668:A:C5	22:DA:670:A:C8	3.02	0.47
22:DA:310:A:O2'	22:DA:311:A:P	2.72	0.47
1:CA:729:A:C5	1:CA:730:G:N7	2.82	0.47
1:AA:1053:G:C3'	1:AA:1054:C:H5'	2.44	0.47
29:DH:117:LEU:HD11	29:DH:130:VAL:HG22	1.95	0.47
1:CA:789:U:O2'	1:CA:791:G:N7	2.38	0.47
5:AE:36:LEU:HD21	5:AE:137:VAL:HG11	1.95	0.47
22:BA:2856:A:N6	22:BA:2857:G:O6	2.47	0.47
1:CA:510:A:H5''	1:CA:511:C:OP2	2.14	0.47
1:AA:1521:C:N3	1:AA:1522:U:C5	2.82	0.47
16:CP:52:LEU:HD21	16:CP:57:ILE:HD12	1.95	0.47
22:DA:681:G:N3	22:DA:682:G:C8	2.82	0.47
27:BF:108:VAL:N	27:BF:109:PRO:CD	2.77	0.47
22:BA:345:A:C2	22:BA:346:A:N6	2.82	0.47
22:DA:2405:G:O2'	22:DA:2411:A:N6	2.47	0.47
1:CA:137:U:H1'	1:CA:227:G:N2	2.29	0.47
11:CK:24:HIS:HB3	11:CK:31:ILE:HG23	1.95	0.47
16:AP:79:ASN:O	16:AP:80:LYS:HE3	2.14	0.47
22:DA:167:A:C2	22:DA:168:G:H1'	2.49	0.47
22:DA:323:C:H6	22:DA:1205:A:N1	2.12	0.47
40:BS:20:VAL:O	40:BS:23:LEU:HB2	2.13	0.47
22:BA:813:U:H2'	22:BA:814:C:C6	2.48	0.47
3:CC:92:ALA:O	3:CC:96:GLY:N	2.47	0.47
39:DR:19:THR:CG2	39:DR:95:ASP:HB3	2.44	0.47
19:AS:51:VAL:HG22	19:AS:71:LEU:HD13	1.96	0.47
27:BF:158:THR:HG22	27:BF:160:ALA:H	1.78	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:967:C:H4'	9:AI:127:PHE:HE2	1.78	0.47
26:DE:146:VAL:O	26:DE:167:VAL:HA	2.13	0.47
22:BA:1304:A:O2'	22:BA:1305:C:H5'	2.13	0.47
25:BD:39:ASP:CG	25:BD:40:LEU:N	2.67	0.47
24:DC:174:LEU:O	24:DC:181:MET:HA	2.14	0.47
31:DJ:116:ARG:NH1	31:DJ:116:ARG:HB2	2.29	0.47
4:CD:206:LYS:OXT	4:CD:206:LYS:HG3	2.14	0.47
26:BE:134:LEU:O	26:BE:134:LEU:HD12	2.14	0.47
43:BV:66:ASP:O	43:BV:67:GLY:C	2.53	0.47
53:B5:48:LEU:HA	53:B5:208:THR:CB	2.44	0.47
22:BA:1060:U:O4	22:BA:1088:A:C8	2.68	0.47
1:AA:1222:G:C6	1:AA:1223:C:C4	3.02	0.47
1:AA:1322:C:O4'	1:AA:1322:C:O2	2.30	0.47
22:BA:1340:U:C5	22:BA:1603:A:C8	3.01	0.47
42:BU:16:GLY:O	42:BU:19:LYS:N	2.40	0.47
1:AA:89:U:O2'	1:AA:90:C:H5''	2.14	0.47
22:DA:1338:G:O2'	22:DA:1393:A:N1	2.31	0.47
2:CB:134:ALA:O	2:CB:138:THR:N	2.43	0.47
21:AU:37:PHE:HB3	21:AU:41:PRO:CG	2.43	0.47
2:CB:94:HIS:HB2	2:CB:146:ASN:O	2.14	0.47
1:CA:728:A:C6	1:CA:729:A:C6	3.02	0.47
22:BA:1588:G:C5	22:BA:1589:U:C5	3.02	0.47
35:BN:71:ARG:CG	35:BN:71:ARG:HH21	2.26	0.47
22:BA:1006:C:C2	22:BA:1138:G:N2	2.81	0.47
1:CA:1439:G:C2	1:CA:1463:U:O2	2.67	0.47
1:CA:1125:U:H4'	10:CJ:7:ARG:NH1	2.29	0.47
22:DA:1288:G:C4	22:DA:1327:A:C2	3.02	0.47
1:CA:280:C:H4'	1:CA:281:G:OP2	2.14	0.47
7:CG:92:ARG:HB3	7:CG:93:PRO:HD2	1.96	0.47
16:AP:10:GLY:O	16:AP:11:ALA:CB	2.62	0.47
21:AU:16:LEU:C	21:AU:18:ARG:NE	2.68	0.47
22:DA:980:A:C4	22:DA:1136:G:O4'	2.67	0.47
1:CA:749:A:C6	1:CA:750:C:C4	3.02	0.47
1:CA:927:G:N2	1:CA:1391:U:H1'	2.29	0.47
9:CI:49:ARG:NH1	9:CI:53:GLU:HG2	2.29	0.47
22:BA:1967:C:C2'	22:BA:1968:G:H5'	2.44	0.47
21:AU:22:SER:C	21:AU:23:CYS:SG	2.92	0.47
19:CS:17:LYS:O	19:CS:21:LYS:HB2	2.13	0.47
37:DP:30:VAL:HG12	37:DP:31:TRP:O	2.14	0.47
9:AI:23:PRO:HA	9:AI:61:LEU:HA	1.96	0.47
24:BC:11:PRO:C	24:BC:13:ARG:H	2.18	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:DU:88:GLU:O	42:DU:89:ASP:HB3	2.14	0.47
35:DN:69:ARG:O	35:DN:71:ARG:N	2.41	0.47
22:DA:1434:A:H2'	22:DA:1435:G:C8	2.48	0.47
3:AC:130:PHE:CE1	3:AC:131:ARG:HD2	2.49	0.47
3:CC:47:LEU:HB3	3:CC:50:ALA:HB3	1.97	0.47
22:BA:391:A:H1'	22:BA:411:G:O4'	2.13	0.47
26:BE:111:GLU:HG2	26:BE:114:ARG:NH1	2.29	0.47
6:AF:55:HIS:O	6:AF:56:LYS:HB2	2.14	0.47
1:AA:785:G:C2'	1:AA:786:G:H5'	2.44	0.47
1:AA:1419:G:C5	1:AA:1420:U:C5	3.02	0.47
1:AA:237:G:OP1	17:AQ:42:THR:OG1	2.29	0.47
46:BY:26:PHE:CD2	46:BY:26:PHE:C	2.87	0.47
51:D3:32:ILE:HG22	51:D3:32:ILE:O	2.14	0.47
17:AQ:5:ILE:N	17:AQ:5:ILE:HD12	2.29	0.47
7:AG:55:GLY:O	7:AG:57:SER:N	2.42	0.47
18:AR:34:THR:OG1	18:AR:35:GLU:N	2.47	0.47
25:BD:5:VAL:HG21	25:BD:80:TRP:CE3	2.49	0.47
22:DA:1187:G:C5	57:DA:3578:HOH:O	2.60	0.47
22:BA:2296:U:C4'	22:BA:2297:A:OP1	2.62	0.47
1:AA:684:U:O4	1:AA:685:G:C6	2.67	0.47
22:DA:2215:C:O2'	22:DA:2216:G:H5'	2.14	0.47
22:DA:1805:A:C4	22:DA:1813:G:N2	2.82	0.47
22:DA:1350:C:C2	22:DA:1382:G:C2	3.01	0.47
1:AA:119:A:C4	1:AA:240:G:N7	2.82	0.47
22:BA:1385:A:C4	22:BA:1386:C:C5	3.03	0.47
1:CA:718:A:H5'	11:CK:119:ASN:ND2	2.29	0.47
22:BA:934:U:H2'	22:BA:935:C:C6	2.49	0.47
25:BD:12:THR:HG21	37:BP:9:GLU:CG	2.44	0.47
1:AA:1254:A:H2'	1:AA:1255:G:C8	2.49	0.47
36:BO:53:THR:HB	36:BO:65:THR:HG22	1.96	0.47
2:AB:91:PHE:CD1	2:AB:150:GLY:CA	2.97	0.47
22:DA:609:A:H2'	22:DA:610:C:O4'	2.13	0.47
9:AI:78:ALA:O	9:AI:81:HIS:HB3	2.14	0.47
1:AA:411:A:C5	1:AA:429:U:C5	3.02	0.47
5:CE:25:VAL:O	5:CE:28:GLY:N	2.45	0.47
2:AB:154:MET:CE	2:AB:158:PRO:HG3	2.44	0.47
27:BF:41:GLY:HA2	27:BF:85:ILE:HG13	1.96	0.47
22:DA:621:A:H2'	22:DA:622:G:O4'	2.15	0.47
1:CA:951:G:H2'	1:CA:952:U:C6	2.49	0.47
26:DE:170:ARG:CZ	26:DE:176:ASP:OD1	2.63	0.47
15:CO:42:HIS:CE1	15:CO:46:HIS:CD2	3.02	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:DR:81:LYS:O	39:DR:83:TYR:N	2.47	0.47
1:AA:444:G:C6	1:AA:445:G:N7	2.82	0.47
1:AA:1130:A:O2'	9:AI:5:GLN:HG3	2.14	0.47
12:AL:5:ASN:O	12:AL:8:VAL:N	2.46	0.47
2:AB:119:THR:O	2:AB:120:GLN:HB2	2.13	0.47
42:DU:7:ARG:O	42:DU:25:VAL:HB	2.14	0.47
14:AN:2:ALA:O	14:AN:3:LYS:CB	2.62	0.47
4:AD:25:VAL:O	4:AD:26:ARG:O	2.32	0.47
22:DA:807:U:H4'	22:DA:2446:G:OP1	2.14	0.47
22:DA:1838:C:N3	22:DA:1899:A:C2	2.82	0.47
1:CA:1253:G:N1	1:CA:1285:A:N6	2.63	0.47
22:DA:242:G:N7	51:D3:5:LYS:HG2	2.29	0.47
22:BA:1199:U:H2'	22:BA:1200:C:C6	2.49	0.47
23:BB:91:C:OP2	34:BM:18:ARG:HG2	2.13	0.47
34:BM:97:GLN:N	34:BM:97:GLN:OE1	2.47	0.47
8:CH:64:LYS:HE2	8:CH:71:VAL:HG21	1.95	0.47
53:B5:69:LEU:HD12	53:B5:163:GLU:O	2.14	0.47
20:AT:70:ASN:N	20:AT:70:ASN:OD1	2.46	0.47
22:DA:949:G:C6	22:DA:950:G:N7	2.83	0.47
22:DA:2334:U:C4	36:DO:16:ARG:HD3	2.49	0.47
22:BA:1179:G:N7	22:BA:1180:U:H1'	2.29	0.47
13:CM:14:HIS:HB2	13:CM:17:ILE:CD1	2.44	0.47
22:BA:2298:A:N1	22:BA:2321:U:C4	2.81	0.47
1:AA:974:A:H4'	1:AA:975:A:H3'	1.95	0.47
22:DA:1525:A:C5	22:DA:1526:C:C4	3.02	0.47
24:DC:178:SER:O	24:DC:271:ARG:HB2	2.14	0.47
22:DA:197:A:C2	22:DA:198:C:H1'	2.49	0.47
5:AE:81:LEU:HA	5:AE:147:MET:HE1	1.97	0.47
29:BH:111:ALA:O	29:BH:114:GLU:HB2	2.14	0.47
1:AA:8:A:C5	4:AD:206:LYS:HB3	2.50	0.47
1:AA:1123:U:O2'	10:AJ:39:PRO:O	2.30	0.47
1:AA:1462:C:C2	1:AA:1463:U:C6	3.02	0.47
22:BA:1449:G:C6	22:BA:1450:G:N7	2.82	0.47
1:AA:192:A:C5	1:AA:193:C:C5	3.03	0.47
22:DA:1324:G:C2	22:DA:1328:A:C6	3.01	0.47
29:DH:62:LEU:HD22	29:DH:62:LEU:O	2.14	0.47
9:CI:12:ARG:HD2	9:CI:107:ASP:HB3	1.96	0.47
35:BN:95:THR:HG21	35:BN:113:ILE:HD11	1.97	0.47
22:DA:39:G:C6	22:DA:40:U:C4	3.02	0.47
22:BA:372:G:OP2	45:BX:61:LYS:HD3	2.13	0.47
4:AD:197:GLU:O	4:AD:200:ILE:N	2.47	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:479:A:H4'	22:DA:480:A:OP1	2.14	0.47
14:AN:21:PHE:O	14:AN:22:ALA:HB3	2.15	0.47
22:DA:2283:C:H2'	22:DA:2284:A:O4'	2.14	0.47
13:AM:88:GLY:O	13:AM:91:HIS:N	2.47	0.47
1:CA:182:A:C8	1:CA:184:G:N7	2.82	0.47
22:DA:1509:A:O2'	22:DA:1510:G:P	2.72	0.47
9:CI:49:ARG:NH2	9:CI:53:GLU:HA	2.29	0.47
10:AJ:26:VAL:HG12	10:AJ:30:LYS:HD3	1.97	0.47
22:BA:907:G:C2'	22:BA:908:C:H5'	2.45	0.47
25:DD:66:GLY:O	25:DD:69:ALA:HB3	2.14	0.47
22:DA:2452:C:N4	22:DA:2453:A:N6	2.62	0.47
22:BA:1672:A:C2	22:BA:2582:G:H5'	2.49	0.47
1:CA:815:A:N7	1:CA:1509:C:O2'	2.31	0.47
1:CA:632:U:H3'	1:CA:633:G:H5'	1.97	0.47
20:CT:65:GLY:HA2	20:CT:68:HIS:CD2	2.49	0.47
6:AF:18:VAL:N	6:AF:19:PRO:HD2	2.30	0.47
11:AK:110:ILE:HB	21:AU:6:VAL:CG2	2.44	0.47
24:DC:168:ASP:O	24:DC:171:TYR:O	2.31	0.47
24:BC:92:ALA:HB3	24:BC:104:ILE:HG13	1.97	0.47
23:BB:7:G:H5'	36:BO:29:HIS:CD2	2.49	0.47
4:CD:48:LEU:HD23	4:CD:53:VAL:HA	1.96	0.47
22:DA:1028:A:N6	22:DA:1125:G:H2'	2.29	0.47
35:DN:38:LEU:HB3	35:DN:39:PRO:HD3	1.97	0.47
22:BA:164:C:H2'	22:BA:165:A:O4'	2.13	0.47
22:BA:1483:G:N2	22:BA:1484:U:C2	2.83	0.47
22:DA:1794:A:H2'	22:DA:1795:C:C6	2.49	0.47
22:DA:396:G:O4'	45:DX:29:PHE:HB3	2.14	0.47
1:AA:934:C:N3	1:AA:1345:U:C5	2.83	0.47
21:AU:34:ARG:NH2	21:AU:35:ARG:HD2	2.30	0.47
24:DC:17:VAL:HB	24:DC:204:VAL:HG22	1.97	0.47
22:DA:2276:G:O2'	22:DA:2277:G:H5'	2.14	0.47
1:CA:1309:G:O6	1:CA:1329:A:N1	2.47	0.47
22:DA:536:G:C6	22:DA:537:G:C5	3.02	0.47
24:DC:200:HIS:O	24:DC:203:ARG:HG2	2.14	0.47
22:DA:776:G:N7	22:DA:793:A:C4	2.82	0.47
22:DA:1736:U:H2'	22:DA:1737:G:O4'	2.14	0.47
1:AA:381:C:H2'	1:AA:382:A:O4'	2.15	0.47
11:CK:112:ASP:HB2	21:CU:20:LYS:HE3	1.95	0.47
11:CK:87:LYS:HG3	11:CK:114:THR:HA	1.95	0.47
25:DD:4:LEU:HD22	25:DD:101:PHE:CZ	2.49	0.47
22:DA:1313:U:H2'	22:DA:1313:U:O2	2.13	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:2346:A:H3'	22:DA:2347:C:C5'	2.45	0.47
22:BA:2339:C:H2'	22:BA:2340:A:H8	1.79	0.47
13:AM:66:GLU:O	13:AM:69:LEU:N	2.48	0.47
36:BO:30:ARG:HG2	36:BO:31:THR:N	2.29	0.47
22:DA:732:C:H2'	22:DA:733:G:O4'	2.15	0.47
27:BF:85:ILE:HG23	27:BF:85:ILE:O	2.15	0.47
10:AJ:51:VAL:HB	14:AN:81:ARG:CB	2.44	0.47
33:DL:76:GLU:HG3	33:DL:76:GLU:O	2.14	0.47
14:AN:26:GLU:O	14:AN:28:LYS:N	2.47	0.47
1:AA:1306:A:C5	1:AA:1307:U:C5	3.01	0.47
6:CF:6:ILE:HD12	6:CF:6:ILE:N	2.29	0.47
1:CA:748:G:H2'	1:CA:749:A:H8	1.80	0.47
26:BE:119:ILE:HB	26:BE:187:VAL:HG22	1.97	0.47
45:DX:54:LYS:O	45:DX:58:VAL:N	2.41	0.47
22:DA:219:A:N6	22:DA:220:G:C6	2.83	0.47
3:CC:153:VAL:HG23	3:CC:157:LEU:HD21	1.95	0.47
22:BA:2258:C:C2	22:BA:2426:A:H4'	2.49	0.47
22:DA:874:G:C2	22:DA:904:G:C2	3.01	0.47
1:AA:152:A:N6	1:AA:170:U:C2	2.82	0.47
22:DA:1861:G:N2	22:DA:1882:U:H1'	2.29	0.47
22:BA:2805:C:C4	22:BA:2806:C:C4	3.03	0.47
27:BF:134:GLU:HB3	27:BF:136:ILE:HD12	1.96	0.47
1:AA:760:G:C5	1:AA:761:G:C8	3.03	0.47
24:BC:36:LYS:O	24:BC:37:ASN:CB	2.61	0.47
22:BA:807:U:P	33:BL:36:LYS:HE2	2.55	0.47
22:DA:1722:A:C2	22:DA:1739:A:H1'	2.50	0.47
1:CA:929:G:H5''	1:CA:1535:C:H5''	1.96	0.47
22:BA:2327:A:H2'	22:BA:2328:A:C8	2.50	0.47
22:BA:2118:U:O4	22:BA:2149:U:H5'	2.15	0.47
43:BV:82:TYR:O	43:BV:82:TYR:CD2	2.67	0.47
2:CB:32:PHE:N	2:CB:40:ILE:O	2.46	0.47
22:BA:2117:A:N1	22:BA:2170:A:N1	2.63	0.47
23:BB:109:A:C5	23:BB:110:C:C4	3.02	0.47
35:DN:18:GLN:HG2	35:DN:18:GLN:O	2.13	0.47
22:DA:295:G:N3	22:DA:295:G:H2'	2.28	0.47
38:DQ:98:ILE:HG22	38:DQ:106:PHE:HB2	1.97	0.47
41:DT:69:ARG:HA	41:DT:74:ILE:HG22	1.95	0.47
22:DA:2572:A:N7	25:DD:150:GLN:HB3	2.29	0.47
22:BA:622:G:O5'	57:BA:3297:HOH:O	2.20	0.47
22:DA:247:G:OP2	22:DA:249:C:N4	2.48	0.47
1:CA:1302:C:C5	13:CM:17:ILE:CD1	2.98	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:AQ:48:ASP:HB2	17:AQ:52:GLU:OE2	2.14	0.47
22:BA:2187:U:C4	22:BA:2188:U:N3	2.83	0.47
22:BA:2298:A:C2	22:BA:2321:U:C4	3.03	0.47
22:DA:674:G:O2'	26:DE:69:ARG:CD	2.62	0.47
1:CA:33:A:N3	1:CA:34:C:C6	2.83	0.47
22:BA:362:A:H2'	22:BA:362:A:N3	2.29	0.47
22:BA:981:A:C5'	57:BA:3600:HOH:O	2.61	0.47
2:CB:207:ILE:O	2:CB:210:VAL:HG22	2.15	0.47
35:DN:87:PHE:CZ	35:DN:94:TYR:HB3	2.49	0.47
1:CA:1387:G:H2'	1:CA:1388:C:H6	1.80	0.47
20:AT:54:MET:HE1	20:AT:58:VAL:HG21	1.96	0.47
22:BA:1358:G:C8	22:BA:1371:G:O6	2.68	0.47
22:DA:1323:C:C4	22:DA:1324:G:N7	2.83	0.47
22:DA:833:A:H2'	22:DA:834:G:C8	2.50	0.47
15:CO:53:ARG:O	15:CO:56:LEU:N	2.47	0.47
44:BW:41:ARG:HG3	44:BW:41:ARG:HH11	1.79	0.47
1:CA:461:A:H2'	1:CA:462:G:O4'	2.15	0.47
21:CU:8:GLU:HB3	21:CU:12:PHE:CD2	2.49	0.47
33:DL:81:ASP:O	33:DL:82:LEU:HB3	2.13	0.47
39:DR:82:HIS:O	39:DR:82:HIS:CG	2.67	0.47
2:AB:179:LEU:O	2:AB:181:ILE:HG13	2.14	0.47
6:CF:41:ASP:OD1	6:CF:58:HIS:NE2	2.47	0.47
22:BA:1176:U:C4	22:BA:1177:G:C6	3.02	0.47
29:DH:5:LEU:HD13	29:DH:13:GLY:HA3	1.96	0.47
42:BU:89:ASP:CG	42:BU:90:GLY:N	2.67	0.47
13:AM:90:ARG:HD2	13:AM:96:PRO:O	2.15	0.47
24:DC:232:HIS:CE1	24:DC:244:PRO:HA	2.50	0.47
22:DA:2636:C:H4'	25:DD:81:GLU:OE1	2.14	0.47
22:DA:1660:G:C2	22:DA:1661:G:C8	3.03	0.47
22:BA:1246:A:H2'	22:BA:1247:A:O5'	2.15	0.47
22:DA:1428:C:C5	22:DA:1569:A:H5''	2.48	0.47
1:CA:1423:G:OP1	32:DK:49:ARG:NH1	2.45	0.47
3:AC:152:GLU:HB3	3:AC:167:TRP:HB3	1.96	0.47
22:BA:1360:G:C6	22:BA:1372:U:C2	3.02	0.47
1:AA:1381:U:C6	7:AG:78:ARG:NH2	2.83	0.47
8:AH:73:GLU:N	8:AH:130:ALA:O	2.48	0.47
22:DA:605:G:N7	22:DA:606:U:C5	2.82	0.47
7:AG:92:ARG:HB3	7:AG:93:PRO:HD2	1.97	0.47
22:DA:805:G:H5''	33:DL:38:GLN:HG3	1.95	0.47
24:DC:237:GLY:O	24:DC:239:ASN:N	2.47	0.47
3:CC:169:ARG:HD2	3:CC:170:GLU:N	2.29	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:DI:39:CYS:HA	30:DI:42:PHE:HB3	1.95	0.47
52:B4:10:LEU:HD12	52:B4:33:HIS:CD2	2.49	0.47
22:DA:282:A:C6	22:DA:359:G:N1	2.82	0.47
1:AA:134:G:H1'	1:AA:325:A:C5	2.50	0.47
3:AC:106:VAL:HG12	3:AC:106:VAL:O	2.14	0.47
1:CA:1299:A:N3	1:CA:1299:A:H2'	2.30	0.47
41:BT:73:ARG:NH2	41:BT:73:ARG:HB3	2.29	0.47
34:BM:74:THR:HG22	34:BM:88:ASN:C	2.35	0.47
22:DA:1213:A:H2'	22:DA:1214:A:O4'	2.14	0.47
22:BA:2198:A:C2	29:BH:29:PHE:HB2	2.49	0.47
29:BH:86:ASP:CB	1:CA:359:G:O2'	2.62	0.47
22:BA:1178:C:H2'	22:BA:1179:G:N7	2.30	0.47
22:BA:1926:U:H2'	22:BA:1927:A:OP2	2.14	0.47
1:CA:906:A:N6	57:CA:1824:HOH:O	2.47	0.47
22:DA:2874:C:H2'	22:DA:2875:C:C6	2.49	0.47
1:AA:254:G:N2	1:AA:273:U:C2	2.83	0.47
22:DA:1794:A:H1'	22:DA:1900:A:C2	2.49	0.47
18:CR:24:LYS:O	18:CR:26:ILE:N	2.48	0.47
5:AE:65:GLU:HA	5:AE:68:ARG:HG3	1.96	0.47
5:CE:105:ILE:HG13	5:CE:112:ARG:HD3	1.96	0.47
22:DA:2550:G:C2	22:DA:2559:C:O2	2.68	0.47
31:BJ:64:VAL:HG13	31:BJ:68:LYS:HB2	1.97	0.47
22:BA:1378:A:O2'	22:BA:1380:G:N7	2.45	0.47
12:CL:24:LEU:HG	12:CL:25:GLU:HG3	1.95	0.47
2:CB:17:GLY:O	2:CB:39:HIS:O	2.32	0.47
22:DA:749:A:C4	22:DA:750:A:H8	2.31	0.47
29:BH:14:SER:O	29:BH:15:LEU:CB	2.61	0.47
1:CA:254:G:H4'	17:CQ:20:SER:OG	2.15	0.47
22:DA:1265:A:OP2	57:DA:3747:HOH:O	2.19	0.47
1:CA:1416:G:C2	1:CA:1485:U:O2	2.67	0.47
1:AA:654:G:C5	1:AA:655:A:N7	2.82	0.47
2:CB:187:VAL:HB	2:CB:191:SER:HB2	1.96	0.47
22:BA:58:G:OP1	41:BT:78:SER:HB2	2.15	0.47
33:BL:111:ILE:HD12	33:BL:111:ILE:N	2.30	0.47
22:DA:2621:G:OP1	25:DD:124:ARG:NH2	2.47	0.47
1:AA:1133:G:N1	1:AA:1142:G:C6	2.82	0.47
22:DA:2886:A:C2	48:D0:29:SER:HB3	2.49	0.47
1:AA:181:A:N6	1:AA:195:A:C8	2.83	0.47
21:AU:25:LYS:HD2	21:AU:26:ALA:N	2.30	0.47
1:AA:1288:A:C5	1:AA:1289:A:C8	3.03	0.47
22:DA:1179:G:C6	22:DA:1180:U:H1'	2.49	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:681:G:H2'	22:DA:682:G:O4'	2.15	0.47
22:DA:271:G:H1'	22:DA:272:A:O5'	2.14	0.47
6:AF:64:VAL:CG1	6:AF:65:GLU:N	2.77	0.47
1:CA:679:C:C2	1:CA:712:A:C2	3.03	0.47
2:AB:151:ILE:HG23	2:AB:152:LYS:N	2.30	0.47
22:BA:250:G:C6	22:BA:251:A:C6	3.03	0.47
22:BA:118:A:N3	22:BA:178:G:H1'	2.30	0.47
4:AD:38:PRO:HD2	4:AD:42:GLY:CA	2.45	0.47
1:CA:992:U:C5	1:CA:1043:G:C8	3.03	0.47
1:AA:728:A:OP1	15:AO:54:ARG:NH2	2.47	0.47
26:DE:181:ILE:HG23	33:DL:2:ARG:CZ	2.45	0.47
26:DE:28:VAL:O	26:DE:32:VAL:HG23	2.14	0.47
35:BN:65:LEU:HD11	35:BN:69:ARG:NH2	2.29	0.47
17:CQ:46:VAL:HG22	17:CQ:61:ILE:HD11	1.96	0.47
22:DA:2201:G:N3	22:DA:2202:U:C6	2.83	0.47
22:BA:29:U:H2'	22:BA:30:G:C8	2.50	0.47
22:BA:1106:G:N2	22:BA:1107:G:H1'	2.29	0.47
1:CA:1243:C:N4	1:CA:1244:G:O6	2.48	0.47
1:CA:1243:C:H2'	1:CA:1244:G:C8	2.49	0.47
34:BM:136:MET:HE2	43:BV:57:TYR:CE1	2.50	0.47
22:DA:1686:C:H2'	22:DA:1687:G:O4'	2.14	0.47
19:AS:51:VAL:HG22	19:AS:71:LEU:CD1	2.44	0.47
22:DA:1436:G:C2	22:DA:1437:C:H1'	2.49	0.47
24:BC:36:LYS:O	24:BC:37:ASN:HB2	2.14	0.47
4:CD:116:GLN:HG3	4:CD:120:HIS:ND1	2.30	0.47
17:AQ:59:VAL:HG22	17:AQ:61:ILE:HD12	1.97	0.47
12:CL:94:ARG:O	12:CL:95:TYR:CD1	2.67	0.47
36:DO:7:ARG:NH1	36:DO:97:PHE:CE2	2.83	0.47
22:DA:2293:G:H2'	22:DA:2294:G:O4'	2.14	0.47
22:DA:1819:A:H4'	22:DA:1820:U:H5''	1.97	0.47
3:CC:64:ILE:HG12	3:CC:66:VAL:HG23	1.96	0.47
49:B1:47:VAL:HG13	49:B1:48:ILE:N	2.28	0.47
30:DI:53:LEU:HD21	30:DI:82:LYS:HE2	1.96	0.47
22:DA:2518:A:H2'	22:DA:2518:A:N3	2.29	0.47
25:DD:56:LYS:O	25:DD:58:ASN:N	2.47	0.47
26:BE:37:ALA:O	26:BE:40:ARG:HB2	2.14	0.47
22:DA:508:A:N6	40:DS:9:HIS:CE1	2.82	0.47
22:BA:547:A:H8	22:BA:548:G:N3	2.12	0.47
5:CE:69:ARG:O	5:CE:70:ASN:HB2	2.14	0.47
26:DE:97:ASN:HB2	26:DE:100:MET:SD	2.55	0.47
25:DD:74:GLU:HG2	25:DD:75:ALA:N	2.30	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:CG:130:ASN:HB2	7:CG:135:VAL:HG21	1.97	0.47
28:BG:24:ILE:HG22	28:BG:26:ILE:HD11	1.96	0.47
22:DA:2851:A:O2'	35:DN:64:ARG:NH2	2.47	0.47
22:BA:1230:A:H2'	22:BA:1231:U:O4'	2.15	0.47
4:AD:126:ASN:HA	4:AD:142:VAL:HG23	1.96	0.47
1:AA:1434:A:H2'	1:AA:1435:G:O4'	2.14	0.47
35:BN:55:ALA:HB1	35:BN:80:PHE:H	1.80	0.47
1:CA:1041:G:H2'	1:CA:1042:A:C8	2.50	0.47
48:D0:36:GLU:OE2	48:D0:46:ASP:HB2	2.15	0.47
22:DA:2784:U:H4'	25:DD:42:ASN:O	2.14	0.47
22:DA:2522:U:C2'	22:DA:2523:G:H5'	2.45	0.47
6:CF:13:ASP:C	6:CF:15:SER:H	2.13	0.47
31:BJ:17:VAL:HG23	31:BJ:137:PRO:CB	2.35	0.47
22:BA:572:A:C2	22:BA:2033:A:C2	3.03	0.47
1:AA:1524:C:OP2	11:AK:125:LYS:NZ	2.48	0.47
11:AK:125:LYS:O	21:AU:34:ARG:CZ	2.63	0.47
22:BA:480:A:H2'	22:BA:481:G:OP1	2.14	0.47
1:CA:780:A:H2	1:CA:803:G:C6	2.33	0.47
22:DA:2264:C:C2	22:DA:2277:G:N2	2.82	0.47
1:CA:1107:C:C4	1:CA:1108:G:N7	2.83	0.47
1:AA:769:G:C2'	1:AA:770:C:H5'	2.45	0.47
22:BA:2196:C:C2'	22:BA:2197:U:H5'	2.45	0.47
22:DA:776:G:C8	22:DA:793:A:C4	3.02	0.47
22:DA:669:G:H2'	22:DA:669:G:N3	2.30	0.47
5:CE:18:VAL:CG2	5:CE:56:VAL:HG13	2.45	0.47
14:CN:49:GLN:O	14:CN:51:LEU:N	2.48	0.47
22:DA:732:C:C4	22:DA:733:G:C5	3.03	0.47
1:AA:1402:C:H2'	1:AA:1403:C:O4'	2.15	0.47
22:DA:938:G:C2	22:DA:939:G:C8	3.03	0.47
41:BT:16:VAL:O	41:BT:17:SER:CB	2.62	0.47
22:DA:13:A:C5	22:DA:525:U:N3	2.83	0.47
22:DA:85:G:OP1	42:DU:6:ARG:HB2	2.15	0.47
22:DA:2652:C:C4	22:DA:2653:U:C5	3.03	0.47
22:DA:2050:C:C4	22:DA:2051:A:C6	3.02	0.47
22:DA:566:U:H2'	22:DA:567:U:O4'	2.14	0.47
22:DA:1838:C:C5	22:DA:1899:A:C6	3.03	0.47
22:BA:2329:U:H2'	22:BA:2330:G:C8	2.50	0.47
22:DA:2077:A:C2	22:DA:2078:C:C5	3.02	0.47
22:DA:2435:A:H2'	22:DA:2436:G:O4'	2.14	0.47
9:CI:83:ILE:O	9:CI:87:LEU:HG	2.15	0.47
22:BA:26:G:H1'	22:BA:514:A:N6	2.30	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AB:166:ALA:HB2	2:AB:187:VAL:HG12	1.96	0.47
6:AF:99:ALA:O	6:AF:100:SER:HB2	2.15	0.47
1:AA:1392:G:C2'	1:AA:1393:U:H5'	2.45	0.47
1:AA:570:G:H2'	1:AA:571:U:C6	2.50	0.47
30:BI:59:ILE:HG22	30:BI:61:VAL:HG23	1.97	0.47
22:BA:2214:C:C5	22:BA:2215:C:C5	3.02	0.47
1:CA:1187:G:C4	1:CA:1188:A:C8	3.02	0.47
22:DA:624:C:O2'	22:DA:657:U:H5''	2.15	0.47
1:CA:937:A:C2	1:CA:1379:G:O6	2.67	0.47
22:DA:836:G:C5	22:DA:837:C:C4	3.03	0.47
1:CA:881:G:C6	1:CA:882:C:C4	3.02	0.47
22:DA:1153:C:H3'	22:DA:1154:G:H8	1.80	0.47
22:DA:2055:C:H5'	22:DA:2056:G:OP1	2.15	0.47
22:BA:11:C:C3'	22:BA:12:U:H5'	2.44	0.47
27:BF:73:SER:HB2	27:BF:81:GLN:N	2.30	0.47
22:DA:571:U:H4'	22:DA:573:U:H5	1.80	0.47
1:AA:469:C:C4	1:AA:470:C:C4	3.02	0.47
35:DN:8:ARG:HB3	35:DN:10:LEU:HG	1.96	0.47
1:CA:1359:C:H2'	1:CA:1361:G:OP2	2.15	0.47
22:DA:2019:A:H4'	38:DQ:34:VAL:HG21	1.97	0.47
22:DA:2195:U:H2'	22:DA:2196:C:C6	2.50	0.47
12:CL:64:THR:HG23	12:CL:93:VAL:HA	1.96	0.47
22:DA:1352:U:C5	22:DA:1377:G:O6	2.68	0.47
22:BA:1789:A:P	24:BC:221:ARG:HH11	2.37	0.47
22:BA:747:U:C4	22:BA:2613:U:C4	3.03	0.47
1:AA:1268:G:C6	1:AA:1269:A:N6	2.82	0.47
22:DA:308:G:N1	22:DA:309:A:C2	2.83	0.47
1:CA:1460:C:N4	1:CA:1461:G:C6	2.82	0.47
36:BO:102:ARG:O	36:BO:105:ALA:HB3	2.15	0.47
22:BA:1359:A:N7	22:BA:1373:A:C2	2.83	0.47
22:DA:682:G:H2'	22:DA:682:G:N3	2.30	0.47
38:DQ:47:TYR:CD1	38:DQ:47:TYR:O	2.67	0.47
1:AA:427:U:H3'	1:AA:428:G:H2'	1.97	0.47
22:BA:702:U:C2'	22:BA:702:U:O2	2.63	0.47
2:AB:140:GLU:O	2:AB:144:LEU:HG	2.15	0.47
32:DK:63:VAL:HG12	32:DK:107:LEU:HD21	1.96	0.47
22:BA:2728:U:O2'	22:BA:2729:G:P	2.73	0.47
13:AM:3:ARG:HA	13:AM:9:ILE:HA	1.97	0.47
6:CF:51:ILE:O	6:CF:52:ASN:HB2	2.15	0.47
7:AG:43:VAL:O	7:AG:47:LEU:HB2	2.15	0.47
17:AQ:4:LYS:O	17:AQ:4:LYS:HD2	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:DP:54:GLY:O	37:DP:77:HIS:NE2	2.47	0.47
1:CA:4:U:O4	1:CA:630:A:O2'	2.32	0.47
34:BM:110:GLU:O	34:BM:111:GLU:C	2.54	0.47
40:DS:107:VAL:HG13	40:DS:107:VAL:O	2.15	0.47
1:CA:865:A:C2	1:CA:918:A:H4'	2.49	0.47
3:CC:72:ARG:HB3	3:CC:75:ILE:HG22	1.96	0.47
23:BB:109:A:H2'	23:BB:110:C:C6	2.50	0.47
1:CA:9:G:H5'	5:CE:108:GLY:HA3	1.95	0.47
44:DW:49:ALA:O	44:DW:50:ASN:HB2	2.15	0.47
22:DA:2285:C:H5'	22:DA:2288:A:N6	2.30	0.47
22:DA:848:C:H2'	22:DA:849:A:C8	2.50	0.47
11:AK:102:ALA:C	11:AK:104:GLY:H	2.18	0.47
1:CA:1517:G:C8	22:DA:1920:C:OP1	2.68	0.47
1:CA:723:U:O5'	21:CU:49:LYS:HG2	2.15	0.47
4:CD:148:LYS:CD	4:CD:148:LYS:H	2.28	0.47
24:BC:197:ASN:CG	24:BC:197:ASN:O	2.53	0.47
9:AI:25:ASN:HB2	9:AI:27:LYS:HE3	1.96	0.47
52:D4:27:CYS:SG	52:D4:30:GLU:N	2.88	0.47
22:DA:1355:G:O2'	22:DA:1356:G:H5'	2.14	0.47
22:DA:1374:G:H5'	22:DA:1375:U:OP2	2.15	0.47
22:DA:396:G:C1'	45:DX:29:PHE:HB3	2.45	0.47
22:DA:149:A:C5	22:DA:150:U:C5	3.03	0.47
1:AA:453:G:H2'	1:AA:454:G:C8	2.50	0.47
4:AD:100:ASN:O	4:AD:104:ARG:HB2	2.15	0.47
4:CD:29:ASP:C	4:CD:31:LYS:H	2.18	0.47
12:AL:63:VAL:HG21	12:AL:95:TYR:CE1	2.50	0.47
1:AA:495:A:O4'	1:AA:496:A:C8	2.68	0.47
22:BA:528:A:H2	22:BA:2043:C:H5'	1.80	0.47
22:BA:278:A:C6	22:BA:362:A:N7	2.83	0.47
1:AA:1180:A:P	9:AI:99:ARG:HH22	2.37	0.47
22:DA:293:U:C5'	22:DA:294:A:OP2	2.63	0.47
23:BB:30:C:H2'	23:BB:31:C:H5'	1.96	0.47
24:BC:260:ASN:O	24:BC:261:LYS:HB2	2.15	0.47
1:CA:268:U:H2'	1:CA:269:C:C6	2.50	0.47
22:DA:1906:G:H5''	22:DA:1929:G:O2'	2.14	0.47
1:AA:192:A:C6	1:AA:193:C:C4	3.03	0.47
22:DA:2563:U:C1'	22:DA:2566:A:N6	2.78	0.47
1:CA:495:A:C2	1:CA:496:A:N6	2.83	0.47
1:AA:1374:A:C2	1:AA:1375:A:C8	3.03	0.47
10:CJ:35:GLN:O	10:CJ:36:VAL:CB	2.62	0.47
22:DA:45:G:N2	22:DA:434:U:C2	2.82	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:478:A:C2	22:DA:480:A:C4	3.03	0.47
4:AD:29:ASP:C	4:AD:30:THR:O	2.51	0.47
11:AK:102:ALA:O	11:AK:104:GLY:N	2.48	0.47
1:AA:284:C:H2'	1:AA:285:C:C6	2.49	0.47
22:DA:2785:C:H2'	22:DA:2786:U:O4'	2.15	0.47
22:BA:644:A:H2'	22:BA:645:C:O4'	2.15	0.47
35:DN:114:GLU:OE2	35:DN:118:ARG:HD3	2.15	0.47
1:CA:540:G:C6	1:CA:541:G:C5	3.02	0.47
22:BA:1579:A:H2'	22:BA:1580:A:C8	2.50	0.47
47:DZ:5:ILE:HD11	47:DZ:57:VAL:HG21	1.97	0.47
42:DU:12:ILE:HG13	42:DU:21:LYS:O	2.14	0.47
1:AA:505:G:OP2	1:AA:534:U:H2'	2.15	0.47
1:CA:366:A:H1'	1:CA:395:C:O2	2.14	0.47
22:BA:2076:U:O4'	22:BA:2076:U:O2	2.33	0.47
48:D0:39:LEU:O	48:D0:42:HIS:HB2	2.15	0.47
31:DJ:106:LYS:HD2	31:DJ:119:PHE:CZ	2.50	0.47
22:BA:601:C:O2	22:BA:605:G:H4'	2.15	0.47
2:AB:99:GLY:O	2:AB:103:ASN:N	2.42	0.47
22:BA:1925:C:C4'	22:BA:1926:U:C4	2.98	0.46
1:CA:1072:G:C6	1:CA:1073:U:C4	3.03	0.46
22:BA:1064:C:N4	22:BA:1070:A:OP2	2.48	0.46
22:DA:1394:U:H6	22:DA:1394:U:H3'	1.80	0.46
41:BT:2:ILE:HG22	41:BT:3:ARG:C	2.36	0.46
22:DA:118:A:N7	22:DA:119:A:C8	2.82	0.46
1:CA:792:A:H1'	1:CA:794:A:N7	2.29	0.46
22:BA:2683:C:H4'	25:BD:13:ARG:NH1	2.30	0.46
1:CA:429:U:H3'	4:CD:9:LEU:CD2	2.44	0.46
1:CA:724:G:C2	1:CA:725:G:C8	3.02	0.46
22:DA:2214:C:H2'	22:DA:2214:C:O2	2.14	0.46
22:BA:527:C:H4'	22:BA:528:A:O5'	2.15	0.46
1:CA:213:G:N7	1:CA:214:C:C2	2.83	0.46
40:DS:59:GLU:HA	40:DS:64:ALA:HA	1.97	0.46
22:DA:1649:G:N1	22:DA:2009:A:C6	2.83	0.46
1:CA:756:C:C2'	1:CA:757:U:H5'	2.45	0.46
37:DP:37:LYS:HD2	37:DP:39:ARG:HD2	1.97	0.46
22:BA:1818:U:OP2	24:BC:156:ARG:NH1	2.48	0.46
22:DA:460:A:H2'	22:DA:461:C:O4'	2.15	0.46
22:DA:1883:U:H2'	22:DA:1884:G:H5'	1.97	0.46
1:CA:1461:G:C6	1:CA:1462:C:C4	3.03	0.46
2:CB:114:LEU:HD13	2:CB:144:LEU:HB2	1.97	0.46
1:CA:956:U:C4	1:CA:957:U:C5	3.03	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:995:C:H5'	22:BA:995:C:C6	2.51	0.46
31:DJ:130:HIS:CE1	31:DJ:137:PRO:HG3	2.50	0.46
45:DX:74:ARG:HG3	45:DX:76:GLU:HG3	1.97	0.46
12:AL:4:VAL:O	12:AL:8:VAL:HG23	2.15	0.46
22:DA:2308:G:O6	22:DA:2311:A:N7	2.48	0.46
10:CJ:12:ALA:HB3	10:CJ:18:ILE:HB	1.96	0.46
22:BA:78:U:H2'	22:BA:79:C:H6	1.79	0.46
3:CC:130:PHE:CZ	3:CC:131:ARG:HD3	2.50	0.46
3:AC:42:TYR:CE2	3:AC:90:VAL:HG21	2.50	0.46
28:BG:2:SER:C	28:BG:4:VAL:N	2.68	0.46
22:DA:1708:C:H2'	22:DA:1709:U:C6	2.50	0.46
22:DA:1838:C:C4	22:DA:1899:A:C2	3.04	0.46
1:CA:1039:G:C6	1:CA:1040:U:C4	3.03	0.46
2:AB:169:GLU:O	2:AB:170:HIS:C	2.54	0.46
22:BA:1713:A:C2	22:BA:1716:U:C6	3.03	0.46
22:DA:693:A:C6	22:DA:694:U:C4	3.03	0.46
26:DE:189:THR:O	26:DE:193:VAL:HG23	2.16	0.46
32:BK:121:GLU:O	32:BK:122:VAL:OXT	2.33	0.46
28:BG:105:LEU:HB2	28:BG:113:VAL:HB	1.98	0.46
30:DI:33:VAL:HG22	30:DI:67:PHE:CD1	2.50	0.46
33:BL:79:LEU:HB2	33:BL:114:GLY:O	2.14	0.46
22:BA:465:G:H2'	22:BA:466:A:C8	2.50	0.46
22:BA:1857:G:N2	22:BA:1884:G:H1'	2.30	0.46
22:BA:2241:A:N7	57:BA:3510:HOH:O	2.36	0.46
1:AA:341:C:H2'	1:AA:342:C:C6	2.50	0.46
1:CA:132:C:C4	1:CA:133:U:C5	3.03	0.46
22:BA:1515:A:H2'	22:BA:1516:G:O4'	2.15	0.46
22:BA:1770:G:C5	22:BA:1983:G:C6	3.04	0.46
22:BA:2575:C:O2'	25:BD:145:SER:HB2	2.15	0.46
22:DA:756:A:H2'	22:DA:757:G:O4'	2.16	0.46
29:DH:41:LYS:O	29:DH:44:ILE:HG12	2.15	0.46
1:CA:57:G:C6	1:CA:58:C:C4	3.03	0.46
1:AA:374:A:C4	1:AA:375:U:C5	3.04	0.46
16:CP:6:LEU:HD11	16:CP:71:VAL:CG2	2.46	0.46
5:CE:105:ILE:HG13	5:CE:105:ILE:O	2.16	0.46
22:DA:593:U:C2	22:DA:594:U:C5	3.04	0.46
22:DA:1645:G:H5''	22:DA:1646:C:O5'	2.15	0.46
22:DA:2094:A:C2	22:DA:2196:C:O2	2.69	0.46
22:DA:2093:G:N1	22:DA:2094:A:C5	2.83	0.46
1:CA:672:U:H2'	1:CA:673:A:C8	2.50	0.46
6:CF:55:HIS:O	6:CF:56:LYS:O	2.33	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:CL:40:THR:HG21	12:CL:90:LEU:HD12	1.97	0.46
1:CA:254:G:N2	1:CA:273:U:C2	2.83	0.46
49:B1:29:THR:O	49:B1:30:LYS:HD3	2.15	0.46
26:DE:48:THR:O	26:DE:52:VAL:HG23	2.15	0.46
23:BB:29:A:H2'	23:BB:30:C:O4'	2.15	0.46
22:BA:2886:A:C5	22:BA:2887:A:C8	3.03	0.46
20:CT:25:ARG:HD2	20:CT:29:ARG:NH2	2.30	0.46
22:DA:503:A:C4	22:DA:506:G:N7	2.83	0.46
19:AS:5:LEU:O	19:AS:7:LYS:N	2.47	0.46
22:DA:2361:G:OP1	51:D3:26:HIS:HA	2.15	0.46
30:DI:80:LEU:HD23	30:DI:84:ALA:HB2	1.97	0.46
22:BA:2567:G:H2'	22:BA:2568:U:C6	2.51	0.46
1:CA:956:U:C5	1:CA:957:U:C5	3.03	0.46
39:DR:49:ILE:CG2	39:DR:54:VAL:HG23	2.46	0.46
1:AA:972:C:H4'	10:AJ:59:LYS:HE2	1.96	0.46
22:BA:84:A:H62	22:BA:101:A:H2	1.63	0.46
12:AL:90:LEU:HB3	12:AL:93:VAL:CG2	2.45	0.46
1:AA:1091:U:O2	1:AA:1095:U:N3	2.48	0.46
1:CA:708:C:H2'	1:CA:709:U:C6	2.51	0.46
1:CA:1244:G:C6	1:CA:1245:C:N4	2.83	0.46
6:CF:38:ARG:HG2	6:CF:63:ASN:CB	2.45	0.46
1:AA:160:A:H2'	1:AA:161:A:O4'	2.15	0.46
40:DS:69:LEU:HG	40:DS:107:VAL:CG2	2.45	0.46
1:CA:312:C:H2'	1:CA:313:A:O4'	2.16	0.46
2:CB:164:ILE:O	2:CB:186:ILE:HG23	2.14	0.46
22:DA:605:G:C5	22:DA:606:U:C5	3.03	0.46
22:BA:170:U:H2'	22:BA:171:U:H6	1.80	0.46
22:BA:368:A:N6	22:BA:369:U:O4	2.47	0.46
26:BE:149:ILE:CD1	26:BE:172:ALA:HA	2.46	0.46
1:CA:1273:C:H2'	1:CA:1274:A:O4'	2.15	0.46
48:D0:13:ARG:HG2	48:D0:17:ARG:HD2	1.96	0.46
23:DB:25:U:C4	23:DB:26:C:C4	3.03	0.46
1:AA:224:U:H2'	1:AA:225:C:C6	2.49	0.46
1:CA:544:G:OP1	4:CD:59:GLN:HG2	2.15	0.46
51:B3:62:LEU:HB3	51:B3:65:ALA:HB2	1.96	0.46
22:DA:1599:U:O4	22:DA:1600:C:N4	2.47	0.46
2:CB:133:GLU:O	2:CB:137:ARG:HB3	2.15	0.46
38:BQ:90:ILE:HG13	39:BR:49:ILE:HD11	1.96	0.46
22:BA:622:G:H2'	22:BA:623:C:H6	1.79	0.46
22:DA:1365:A:O3'	45:DX:11:ARG:NH1	2.47	0.46
22:BA:1131:G:O2'	22:BA:2026:U:H5'	2.14	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:2821:A:H2'	22:DA:2822:G:O4'	2.15	0.46
11:CK:122:ARG:CZ	21:CU:36:GLU:HG3	2.45	0.46
1:CA:332:G:OP2	20:CT:5:LYS:HB3	2.15	0.46
1:CA:1106:G:H2'	1:CA:1107:C:H6	1.80	0.46
22:DA:2093:G:C8	22:DA:2225:A:C4	3.03	0.46
38:BQ:115:ALA:C	38:BQ:117:LEU:H	2.19	0.46
22:DA:669:G:N2	22:DA:670:A:C6	2.82	0.46
22:DA:574:A:P	57:DA:3263:HOH:O	2.74	0.46
22:DA:1264:A:C8	22:DA:1265:A:C8	3.04	0.46
22:DA:1341:G:C2	41:DT:84:TYR:CD2	3.03	0.46
12:CL:7:LEU:HD22	12:CL:12:ARG:HD2	1.96	0.46
22:BA:1866:A:C6	22:BA:1876:A:N7	2.83	0.46
22:DA:2066:C:N3	22:DA:2067:G:N7	2.62	0.46
21:AU:14:VAL:CG1	21:AU:16:LEU:HD21	2.45	0.46
22:DA:13:A:N1	22:DA:525:U:H2'	2.29	0.46
1:CA:101:A:C5	1:CA:102:G:N7	2.83	0.46
21:CU:12:PHE:O	21:CU:13:ASP:HB2	2.14	0.46
22:DA:982:C:H5''	22:DA:983:A:OP2	2.15	0.46
22:BA:1176:U:H2'	22:BA:1177:G:N9	2.31	0.46
1:CA:301:G:H2'	1:CA:302:G:H8	1.79	0.46
28:DG:89:LEU:HD12	28:DG:162:VAL:HG22	1.97	0.46
15:AO:89:ARG:NH2	22:BA:714:U:OP2	2.48	0.46
46:DY:9:LYS:HG2	46:DY:10:SER:H	1.80	0.46
1:AA:929:G:C6	1:AA:930:C:C4	3.04	0.46
1:CA:391:G:H5''	16:CP:8:ARG:CD	2.45	0.46
22:BA:2861:U:C2	22:BA:2862:G:C8	3.04	0.46
1:CA:187:G:H5'	20:CT:80:THR:HG21	1.97	0.46
11:CK:61:PHE:O	11:CK:65:VAL:HG12	2.15	0.46
22:DA:2658:C:OP1	28:DG:158:LYS:NZ	2.49	0.46
26:DE:83:VAL:CG1	26:DE:86:ALA:HA	2.45	0.46
1:AA:614:C:H2'	1:AA:615:G:O4'	2.16	0.46
1:AA:637:C:C2'	1:AA:638:U:H5'	2.45	0.46
29:BH:132:PHE:CD2	29:BH:142:VAL:CG2	2.99	0.46
29:BH:80:ILE:HG21	29:BH:94:ILE:CG1	2.45	0.46
1:CA:54:C:N4	1:CA:352:C:H2'	2.30	0.46
12:AL:22:PRO:C	12:AL:24:LEU:N	2.69	0.46
22:BA:1074:G:H2'	22:BA:1074:G:N3	2.31	0.46
17:AQ:16:LYS:N	17:AQ:17:MET:SD	2.88	0.46
17:AQ:17:MET:CG	17:AQ:20:SER:HB3	2.43	0.46
22:DA:2091:C:C3'	22:DA:2092:U:H5''	2.43	0.46
22:DA:150:U:H2'	22:DA:151:C:C6	2.50	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:451:A:H61	1:AA:481:G:C5'	2.28	0.46
22:DA:26:G:C6	22:DA:27:G:N1	2.84	0.46
19:AS:29:LYS:HB3	19:AS:30:PRO:CD	2.44	0.46
1:CA:328:C:O2	1:CA:328:C:O2'	2.32	0.46
5:CE:13:GLU:O	5:CE:13:GLU:HG3	2.15	0.46
22:DA:1426:G:N2	22:DA:1571:A:N7	2.63	0.46
39:BR:47:VAL:HG12	39:BR:54:VAL:HG22	1.96	0.46
1:CA:1416:G:N2	1:CA:1485:U:H1'	2.30	0.46
12:CL:7:LEU:HD22	12:CL:12:ARG:CD	2.46	0.46
1:AA:456:A:H5'	1:AA:457:G:OP2	2.16	0.46
22:DA:847:U:O2	22:DA:847:U:H2'	2.14	0.46
22:BA:974:G:C4	22:BA:1186:G:C2	3.03	0.46
33:BL:63:LYS:HA	51:B3:13:ARG:HB3	1.98	0.46
22:DA:987:C:N4	22:DA:988:A:C6	2.83	0.46
22:BA:1250:G:C5'	38:BQ:6:ARG:HD3	2.45	0.46
39:DR:82:HIS:O	39:DR:82:HIS:ND1	2.49	0.46
41:BT:89:GLU:OE2	41:BT:89:GLU:HA	2.15	0.46
1:AA:1049:U:O4	14:AN:2:ALA:HB1	2.16	0.46
22:DA:748:G:O6	22:DA:751:A:H5'	2.16	0.46
31:DJ:25:LEU:CD1	31:DJ:100:VAL:HG12	2.45	0.46
45:DX:40:VAL:O	45:DX:41:GLU:C	2.53	0.46
24:DC:87:ARG:NH1	24:DC:87:ARG:HB3	2.29	0.46
24:DC:173:THR:O	24:DC:174:LEU:HD12	2.15	0.46
1:AA:927:G:N1	1:AA:1391:U:C2	2.84	0.46
30:DI:29:GLY:HA2	30:DI:33:VAL:HB	1.97	0.46
8:CH:20:ALA:O	8:CH:21:ASN:HB2	2.16	0.46
45:BX:68:LEU:HD13	45:BX:78:TYR:CE2	2.51	0.46
1:AA:1486:G:H2'	1:AA:1487:G:O4'	2.14	0.46
22:DA:1255:U:C5	26:DE:68:ALA:HA	2.50	0.46
22:DA:1616:A:H2	22:DA:1647:U:C5	2.33	0.46
33:DL:10:GLU:HA	33:DL:10:GLU:OE1	2.16	0.46
22:BA:1808:A:O2'	45:BX:3:ARG:NH1	2.49	0.46
22:DA:370:G:N1	22:DA:424:G:C5	2.83	0.46
2:AB:85:LEU:HG	2:AB:86:SER:N	2.31	0.46
22:BA:1918:A:O2'	22:BA:1920:C:C4	2.68	0.46
24:DC:107:PRO:HG2	24:DC:110:LEU:HD13	1.97	0.46
1:CA:1280:A:C8	10:CJ:42:LEU:HD23	2.51	0.46
3:AC:40:ARG:HG2	3:AC:55:ILE:HG12	1.98	0.46
5:AE:68:ARG:O	5:AE:71:MET:HE3	2.15	0.46
22:DA:1953:A:C6	22:DA:2550:G:O4'	2.69	0.46
1:AA:154:U:C2	1:AA:168:G:C2	3.03	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:407:U:O2'	4:AD:113:GLU:HG3	2.15	0.46
22:BA:2305:U:H5''	27:BF:131:GLY:HA3	1.96	0.46
1:AA:1319:A:N7	1:AA:1323:G:C5	2.84	0.46
1:CA:253:A:C2	1:CA:254:G:C5	3.03	0.46
22:DA:1432:G:N2	22:DA:1433:A:N3	2.63	0.46
22:DA:2291:U:H2'	22:DA:2292:U:H6	1.80	0.46
23:BB:37:C:C6	23:BB:38:C:C5	3.04	0.46
9:CI:12:ARG:CD	9:CI:107:ASP:HB3	2.46	0.46
1:AA:1446:A:C2'	1:AA:1447:A:H5'	2.45	0.46
17:AQ:81:LYS:N	17:AQ:81:LYS:CD	2.78	0.46
34:BM:55:ARG:CZ	34:BM:55:ARG:CB	2.93	0.46
27:BF:106:ILE:C	27:BF:109:PRO:HD2	2.34	0.46
21:AU:14:VAL:O	21:AU:16:LEU:HG	2.15	0.46
12:AL:88:LYS:HG3	12:AL:88:LYS:O	2.15	0.46
4:AD:49:SER:O	4:AD:53:VAL:HG13	2.15	0.46
7:AG:32:VAL:O	7:AG:34:GLY:N	2.48	0.46
22:BA:28:A:C2'	22:BA:29:U:H5'	2.46	0.46
24:BC:162:VAL:HG22	24:BC:176:LEU:HA	1.98	0.46
42:DU:88:GLU:O	42:DU:89:ASP:CB	2.62	0.46
1:AA:151:A:H2'	1:AA:152:A:O4'	2.14	0.46
17:AQ:8:LEU:HD23	17:AQ:25:ILE:HG13	1.97	0.46
53:B5:101:ILE:HG22	53:B5:102:GLN:HG3	1.98	0.46
34:BM:95:LEU:O	34:BM:96:ILE:HD13	2.15	0.46
1:CA:436:C:C2	1:CA:437:U:C5	3.04	0.46
1:CA:439:U:C5	1:CA:440:C:C5	3.04	0.46
30:BI:106:LEU:HA	30:BI:109:ILE:HB	1.97	0.46
11:AK:51:GLY:O	11:AK:52:PHE:O	2.33	0.46
1:AA:1114:C:C5	1:AA:1115:U:C5	3.04	0.46
22:DA:1623:G:C2	22:DA:1624:U:C6	3.04	0.46
1:AA:587:G:C2	1:AA:755:G:C5	3.03	0.46
47:DZ:40:ASP:OD2	47:DZ:45:ARG:NE	2.48	0.46
22:BA:1840:G:C6	22:BA:1841:U:C4	3.03	0.46
1:AA:1326:U:H2'	1:AA:1327:C:C6	2.50	0.46
16:AP:19:VAL:HG13	16:AP:37:GLY:C	2.35	0.46
22:DA:2207:C:O2'	22:DA:2208:C:H5'	2.15	0.46
26:BE:29:HIS:CE1	33:BL:8:PRO:HB3	2.51	0.46
9:CI:67:VAL:HG11	9:CI:79:ILE:HD11	1.97	0.46
50:D2:6:GLN:HA	50:D2:6:GLN:OE1	2.16	0.46
16:AP:29:ASN:N	16:AP:29:ASN:OD1	2.48	0.46
16:CP:10:GLY:HA3	16:CP:15:PRO:HA	1.98	0.46
41:DT:8:LEU:HD23	41:DT:50:LEU:HD21	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:BE:115:GLN:HA	26:BE:115:GLN:OE1	2.16	0.46
38:BQ:88:VAL:HA	39:BR:49:ILE:HD12	1.97	0.46
22:DA:1775:U:O4	22:DA:1789:A:H2	1.98	0.46
21:CU:34:ARG:HG2	21:CU:35:ARG:N	2.31	0.46
22:DA:781:A:O2'	22:DA:782:A:OP2	2.29	0.46
22:BA:2127:G:H4'	22:BA:2128:G:OP1	2.15	0.46
22:DA:537:G:OP1	22:DA:995:C:N4	2.48	0.46
1:CA:1022:A:C6	1:CA:1023:U:C4	3.04	0.46
1:CA:978:A:O2'	1:CA:1322:C:H5	1.97	0.46
5:AE:157:ARG:O	5:AE:159:LYS:N	2.46	0.46
22:DA:1257:C:N4	22:DA:1258:U:O4	2.49	0.46
22:DA:1817:G:C2'	22:DA:1818:U:H5'	2.46	0.46
11:AK:29:ASN:OD1	11:AK:47:ALA:HB3	2.15	0.46
1:AA:654:G:C2'	1:AA:655:A:H5'	2.44	0.46
22:BA:140:C:O2	22:BA:140:C:O4'	2.34	0.46
22:DA:228:C:O2	22:DA:418:C:H4'	2.15	0.46
13:AM:26:GLY:O	13:AM:28:THR:N	2.42	0.46
10:AJ:52:LEU:HD22	10:AJ:62:ARG:HG2	1.98	0.46
1:AA:455:G:C2	1:AA:478:A:C2	3.04	0.46
4:AD:38:PRO:HD2	4:AD:42:GLY:HA2	1.98	0.46
33:DL:110:VAL:C	33:DL:111:ILE:HD12	2.35	0.46
5:AE:77:ASN:O	5:AE:78:ASN:HB3	2.16	0.46
26:DE:181:ILE:HG23	33:DL:2:ARG:NH1	2.31	0.46
22:DA:818:G:C2'	22:DA:819:A:H5''	2.45	0.46
10:CJ:15:HIS:CG	10:CJ:16:ARG:N	2.83	0.46
22:DA:2538:C:H2'	22:DA:2539:C:C6	2.50	0.46
17:CQ:28:PHE:HE2	17:CQ:37:PHE:O	1.99	0.46
19:CS:11:ILE:HG13	19:CS:12:ASP:N	2.31	0.46
22:BA:927:A:H2'	22:BA:928:A:C8	2.51	0.46
22:BA:478:A:N1	22:BA:500:G:H4'	2.31	0.46
1:CA:909:A:H2'	1:CA:910:C:O4'	2.16	0.46
22:DA:1262:A:N1	22:DA:1263:U:C2	2.83	0.46
32:BK:15:GLY:HA2	32:BK:47:ILE:HD13	1.98	0.46
22:DA:2848:G:C8	37:DP:95:ALA:HB2	2.50	0.46
45:BX:11:ARG:HB2	45:BX:12:PRO:CD	2.46	0.46
28:DG:96:ALA:N	28:DG:128:GLN:O	2.47	0.46
8:AH:25:VAL:HG13	8:AH:25:VAL:O	2.15	0.46
22:BA:2518:A:H2'	22:BA:2518:A:N3	2.31	0.46
1:CA:1496:C:H2'	1:CA:1497:G:O4'	2.15	0.46
22:DA:1946:U:H2'	22:DA:1947:C:C6	2.51	0.46
22:DA:851:C:C2	22:DA:852:U:C5	3.03	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:AS:15:LEU:O	19:AS:19:VAL:HG23	2.14	0.46
22:BA:988:A:O5'	47:BZ:12:SER:HB2	2.16	0.46
1:AA:1353:G:C4	1:AA:1354:U:C5	3.03	0.46
49:D1:33:LYS:HA	49:D1:52:ALA:HB3	1.97	0.46
29:BH:37:VAL:CG2	29:BH:38:PRO:HD2	2.45	0.46
29:BH:94:ILE:O	29:BH:122:LEU:CG	2.57	0.46
29:BH:94:ILE:HG23	29:BH:98:ASP:HB2	1.98	0.46
22:BA:1171:G:C2	22:BA:1178:C:O2	2.68	0.46
22:DA:2142:A:C6	22:DA:2143:C:C4	3.04	0.46
22:DA:1792:G:C5	22:DA:1793:C:C5	3.04	0.46
22:DA:187:G:O2'	22:DA:1365:A:C2	2.64	0.46
22:DA:201:C:N4	22:DA:202:U:C4	2.83	0.46
16:AP:17:TYR:N	16:AP:17:TYR:CD1	2.84	0.46
1:CA:1346:A:N6	1:CA:1374:A:C8	2.84	0.46
24:BC:17:VAL:HB	24:BC:204:VAL:HG22	1.96	0.46
22:BA:528:A:C2	22:BA:2042:A:H2'	2.51	0.46
22:DA:301:G:H1'	22:DA:302:C:C6	2.50	0.46
1:AA:1256:A:N6	1:AA:1277:C:C2	2.83	0.46
22:DA:2199:A:C6	22:DA:2200:C:C2	3.03	0.46
22:DA:2199:A:C5	22:DA:2200:C:C4	3.04	0.46
22:DA:613:A:OP2	22:DA:614:A:C8	2.69	0.46
22:DA:1268:A:H2'	22:DA:1269:A:O4'	2.16	0.46
22:DA:2189:U:C2'	22:DA:2190:G:H5''	2.44	0.46
1:AA:1417:G:H22	1:AA:1482:G:H2'	1.79	0.46
1:CA:563:A:N7	1:CA:567:G:H1'	2.31	0.46
1:AA:1491:G:H5''	12:AL:43:LYS:HG3	1.97	0.46
29:DH:34:GLY:O	29:DH:35:LYS:CG	2.64	0.46
22:DA:1635:A:H2'	22:DA:1636:U:O4'	2.16	0.46
1:AA:1374:A:N3	1:AA:1375:A:C8	2.83	0.46
39:DR:49:ILE:HG22	39:DR:54:VAL:N	2.31	0.46
10:AJ:51:VAL:HB	14:AN:81:ARG:HB3	1.98	0.46
7:CG:146:GLU:HA	7:CG:149:LYS:HB2	1.98	0.46
20:AT:44:LYS:HE2	20:AT:86:LEU:O	2.16	0.46
1:CA:102:G:C2	1:CA:103:U:C5	3.04	0.46
22:DA:893:C:H2'	22:DA:894:U:O4'	2.15	0.46
30:BI:122:ILE:O	30:BI:126:THR:OG1	2.33	0.46
23:BB:7:G:O2'	36:BO:38:GLN:OE1	2.09	0.46
7:CG:68:ASN:HB3	7:CG:130:ASN:HB3	1.96	0.46
22:BA:270:A:N6	22:BA:369:U:O2	2.49	0.46
24:DC:116:ILE:HB	24:DC:127:GLY:O	2.16	0.46
46:DY:23:ARG:NH1	46:DY:27:ASN:OD1	2.48	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AB:31:ILE:HD11	2:AB:39:HIS:ND1	2.31	0.46
3:AC:181:ASP:HB3	3:AC:204:LYS:HB2	1.98	0.46
11:AK:31:ILE:HB	11:AK:46:THR:HG22	1.98	0.46
2:CB:184:PHE:CD2	2:CB:198:PHE:HB2	2.50	0.46
2:CB:165:ASP:O	2:CB:169:GLU:HG2	2.16	0.46
22:BA:7:G:H2'	22:BA:8:C:C6	2.51	0.46
22:BA:1935:G:C6	22:BA:1962:C:C5	3.04	0.46
18:CR:58:ALA:O	18:CR:59:ILE:C	2.54	0.46
1:CA:121:U:C4'	1:CA:121:U:OP2	2.64	0.46
22:DA:712:G:C2	22:DA:720:U:O2	2.69	0.46
1:CA:487:A:H2'	1:CA:488:C:O4'	2.16	0.46
22:DA:868:U:C4	22:DA:869:G:N7	2.84	0.46
2:AB:18:HIS:O	2:AB:19:GLN:HB2	2.16	0.46
2:AB:80:VAL:O	2:AB:84:ALA:HB3	2.15	0.46
22:DA:119:A:H5'	57:DA:3220:HOH:O	2.14	0.46
1:CA:795:C:H5'	1:CA:1522:U:OP1	2.15	0.46
22:BA:2321:U:C5'	22:BA:2322:A:OP2	2.60	0.46
22:BA:1020:A:C2	22:BA:1141:U:C2	3.03	0.46
2:AB:186:ILE:HA	2:AB:200:ILE:HB	1.97	0.46
22:DA:1645:G:H4'	22:DA:1646:C:C5	2.50	0.46
1:CA:676:A:C2	1:CA:677:U:C5	3.03	0.46
22:BA:674:G:O2'	26:BE:69:ARG:HD3	2.16	0.46
22:DA:785:G:O2'	22:DA:1779:U:C5'	2.64	0.46
22:DA:750:A:N3	22:DA:750:A:H2'	2.30	0.46
12:CL:93:VAL:O	12:CL:93:VAL:HG23	2.16	0.46
1:AA:1238:A:C2	1:AA:1303:C:H4'	2.51	0.46
1:AA:408:A:N3	1:AA:435:A:C2	2.84	0.46
1:AA:1152:A:C5	1:AA:1153:G:N7	2.84	0.46
22:DA:141:G:H3'	22:DA:142:A:C8	2.50	0.46
1:CA:502:A:OP1	12:CL:115:SER:CB	2.64	0.46
22:DA:2373:G:C6	22:DA:2374:C:N4	2.83	0.46
22:DA:482:A:O2'	22:DA:497:A:N1	2.45	0.46
1:AA:193:C:O2'	1:AA:194:C:H5'	2.14	0.46
1:AA:1521:C:C2	1:AA:1522:U:C6	3.04	0.46
22:DA:680:C:H2'	22:DA:681:G:C8	2.51	0.46
22:DA:647:G:OP1	22:DA:647:G:H4'	2.15	0.46
28:DG:40:ALA:HA	28:DG:58:TYR:CE1	2.50	0.46
1:AA:660:C:OP1	15:AO:5:THR:HG21	2.15	0.46
20:CT:70:ASN:O	20:CT:73:ALA:N	2.49	0.46
22:DA:228:C:C5'	22:DA:229:C:C6	2.99	0.46
11:CK:51:GLY:O	11:CK:52:PHE:CD1	2.69	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:479:A:N3	22:DA:481:G:H5''	2.31	0.46
22:DA:2898:U:H2'	22:DA:2899:A:C8	2.51	0.46
1:CA:666:G:C6	1:CA:741:G:C6	3.04	0.46
1:CA:927:G:H4'	1:CA:1503:A:N7	2.31	0.46
20:AT:68:HIS:HB3	20:AT:69:LYS:CE	2.46	0.46
22:BA:1384:A:H1'	22:BA:1405:U:H1'	1.98	0.46
22:DA:2326:C:C1'	22:DA:2327:A:OP1	2.64	0.46
24:DC:226:ASN:OD1	24:DC:226:ASN:N	2.49	0.46
22:DA:2518:A:P	57:DA:3534:HOH:O	2.73	0.46
22:DA:2436:G:C2	22:DA:2437:G:C8	3.04	0.46
1:CA:881:G:H2'	1:CA:882:C:O4'	2.16	0.46
22:DA:1262:A:H2'	22:DA:1262:A:N3	2.30	0.46
22:DA:1319:C:O2	22:DA:1334:G:C2	2.68	0.46
17:CQ:26:GLU:HA	17:CQ:40:ARG:O	2.14	0.46
4:CD:81:ARG:NH2	4:CD:82:LEU:HD23	2.30	0.46
41:DT:45:ALA:O	41:DT:49:LYS:HE3	2.16	0.46
40:BS:50:VAL:O	40:BS:53:SER:N	2.49	0.46
32:DK:18:ARG:HB2	32:DK:45:GLU:HB3	1.98	0.46
14:AN:68:GLY:O	14:AN:69:ARG:C	2.54	0.46
22:DA:2031:A:C6	22:DA:2498:C:H1'	2.51	0.46
28:DG:86:LYS:O	28:DG:165:ALA:HB2	2.15	0.46
1:AA:279:A:H8	1:AA:279:A:H5'	1.80	0.46
31:DJ:109:LEU:HD22	31:DJ:118:MET:HG3	1.98	0.46
10:CJ:91:ASP:O	10:CJ:92:LEU:HG	2.15	0.46
1:CA:1501:C:OP2	1:CA:1504:G:O2'	2.27	0.46
1:CA:1158:C:N3	1:CA:1160:G:N7	2.64	0.46
22:DA:1790:C:C5	22:DA:1828:G:C2	3.04	0.46
22:DA:149:A:C8	22:DA:150:U:C5	3.04	0.46
22:DA:182:A:H2'	22:DA:183:C:C6	2.51	0.46
13:AM:11:ASP:O	13:AM:12:HIS:HB2	2.16	0.46
22:DA:410:G:H2'	22:DA:2407:A:N7	2.31	0.46
22:DA:1814:G:OP1	24:DC:40:SER:OG	2.33	0.46
1:CA:322:C:H5	1:CA:328:C:C5	2.33	0.46
1:AA:1220:G:H2'	1:AA:1221:G:O4'	2.16	0.46
5:CE:16:ILE:HD12	5:CE:16:ILE:N	2.30	0.46
2:CB:83:ALA:O	2:CB:86:SER:OG	2.33	0.46
1:CA:822:U:H2'	1:CA:823:C:H6	1.80	0.46
1:AA:1157:A:C5	1:AA:1180:A:C6	3.04	0.46
7:AG:86:GLN:HB3	7:AG:148:ASN:ND2	2.31	0.46
1:AA:189:A:O2'	1:AA:190:A:H5'	2.15	0.46
4:CD:107:PHE:CD1	4:CD:107:PHE:N	2.82	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:DB:78:A:C6	23:DB:99:A:C8	3.04	0.46
1:CA:858:G:C5	57:CA:1818:HOH:O	2.62	0.46
22:BA:1376:C:H2'	22:BA:1377:G:O4'	2.15	0.46
10:AJ:17:LEU:HD23	10:AJ:18:ILE:N	2.31	0.46
22:BA:1734:G:C4	22:BA:1735:A:C8	3.03	0.46
22:BA:1735:A:C2	22:BA:1736:U:C1'	2.98	0.46
22:DA:1855:U:C6	22:DA:1856:U:C5	3.03	0.46
22:DA:2232:C:OP2	45:DX:27:ARG:NH2	2.44	0.46
22:DA:158:U:O2	22:DA:169:G:N1	2.49	0.46
1:AA:1145:A:O2'	1:AA:1146:A:P	2.74	0.46
17:AQ:79:VAL:HG12	17:AQ:80:GLU:HG3	1.97	0.46
17:AQ:50:ASN:O	17:AQ:51:ASN:C	2.53	0.46
43:DV:42:LEU:CD1	43:DV:47:VAL:HG21	2.46	0.46
26:DE:108:ILE:O	26:DE:112:LEU:HG	2.16	0.46
16:AP:75:ILE:HG22	16:AP:80:LYS:HE2	1.98	0.46
1:CA:1150:A:C6	1:CA:1151:A:N6	2.83	0.46
44:BW:38:VAL:HG12	44:BW:39:ARG:N	2.31	0.46
10:CJ:5:ARG:HD2	10:CJ:79:PRO:HB3	1.98	0.46
1:CA:667:G:C2	1:CA:740:U:O2	2.69	0.46
19:CS:15:LEU:HD13	19:CS:33:THR:HG21	1.98	0.46
6:AF:97:THR:O	6:AF:98:GLU:HG2	2.15	0.46
22:DA:1512:C:C4	22:DA:1513:U:C5	3.04	0.46
6:AF:54:LEU:HD22	6:AF:55:HIS:O	2.16	0.46
47:DZ:42:PRO:HA	47:DZ:45:ARG:HB2	1.98	0.46
1:AA:1311:A:C2	1:AA:1327:C:N3	2.84	0.46
1:CA:1431:A:C6	1:CA:1432:G:C6	3.03	0.46
4:AD:166:GLU:O	4:AD:167:LYS:HB2	2.16	0.46
16:CP:7:ALA:HB1	16:CP:29:ASN:OD1	2.16	0.46
8:CH:43:GLU:OE1	8:CH:112:THR:HG21	2.16	0.46
22:BA:1676:A:H2'	22:BA:1677:A:O4'	2.16	0.46
24:DC:167:ARG:O	24:DC:167:ARG:CG	2.63	0.46
1:AA:695:A:H2'	1:AA:696:A:O4'	2.16	0.46
32:BK:36:GLY:HA2	32:BK:62:VAL:O	2.15	0.46
40:DS:36:LEU:HD13	40:DS:48:LYS:HB2	1.96	0.46
1:AA:882:C:O2'	1:AA:883:C:H5'	2.15	0.46
22:BA:1174:U:O2	22:BA:1174:U:O4'	2.32	0.46
30:BI:69:PHE:CD1	30:BI:69:PHE:O	2.68	0.46
29:DH:60:GLU:HA	29:DH:60:GLU:OE2	2.15	0.46
31:DJ:49:ASP:OD2	31:DJ:121:LYS:HD3	2.16	0.46
29:BH:90:LEU:HD21	29:BH:93:SER:HA	1.97	0.46
29:BH:90:LEU:HD23	29:BH:93:SER:HA	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:1374:G:H3'	22:DA:1375:U:C6	2.51	0.46
22:DA:118:A:OP2	22:DA:119:A:H5''	2.15	0.46
22:DA:2117:A:N1	22:DA:2171:A:N1	2.64	0.46
22:BA:2637:U:C3'	22:BA:2638:G:H5'	2.44	0.46
24:BC:246:THR:HB	24:BC:248:TRP:CE3	2.51	0.46
11:CK:126:LYS:O	11:CK:127:ARG:HB2	2.16	0.46
1:CA:552:U:H5'	12:CL:83:ARG:HD2	1.98	0.46
1:CA:1345:U:C2	1:CA:1377:A:C2	3.04	0.46
24:DC:65:VAL:HG12	24:DC:67:PHE:CE2	2.51	0.46
22:DA:1638:C:O2'	22:DA:2698:U:O2	2.33	0.46
17:CQ:48:ASP:O	17:CQ:49:GLU:C	2.54	0.46
22:BA:1224:U:C4	22:BA:1225:G:C6	3.04	0.46
1:CA:1412:C:H2'	1:CA:1413:A:C8	2.51	0.46
1:CA:21:G:H1'	1:CA:914:A:N6	2.30	0.46
30:BI:130:GLU:HB3	30:BI:134:ARG:NH2	2.31	0.46
1:CA:1202:U:N3	14:CN:82:ILE:HG21	2.30	0.46
22:DA:2502:G:H5'	22:DA:2503:A:H5''	1.98	0.46
52:D4:19:ARG:O	52:D4:20:ASP:HB2	2.16	0.46
22:BA:1736:U:H2'	22:BA:1737:G:O4'	2.16	0.46
22:DA:2347:C:H2'	22:DA:2348:U:C6	2.51	0.46
32:BK:113:MET:O	32:BK:114:LYS:C	2.53	0.46
48:B0:53:LYS:HE2	48:B0:56:ALA:HA	1.98	0.46
49:B1:40:ASP:HB2	49:B1:49:TYR:OH	2.16	0.46
22:BA:2292:U:H2'	22:BA:2293:G:C8	2.51	0.46
21:AU:16:LEU:HA	21:AU:18:ARG:CZ	2.46	0.46
22:DA:2440:C:C4	22:DA:2441:U:H1'	2.50	0.46
22:BA:1991:U:H2'	22:BA:1992:G:H5'	1.98	0.46
22:DA:2250:G:H8	22:DA:2250:G:O5'	1.99	0.46
25:BD:55:LYS:CD	25:BD:60:VAL:HG22	2.45	0.46
1:CA:663:A:O2'	1:CA:664:G:H5'	2.16	0.46
30:BI:97:LYS:HG3	30:BI:139:VAL:HG22	1.97	0.46
7:CG:60:GLU:HA	7:CG:63:GLU:HB3	1.98	0.46
22:BA:2488:G:O2'	22:BA:2489:U:H5'	2.16	0.46
24:DC:84:ASP:O	24:DC:86:ASN:N	2.49	0.46
19:AS:51:VAL:O	19:AS:58:VAL:HG13	2.15	0.46
26:DE:31:VAL:HG21	26:DE:100:MET:O	2.16	0.46
1:CA:1337:G:H5''	1:CA:1338:G:OP1	2.16	0.46
14:AN:87:ALA:O	14:AN:92:GLU:HB2	2.16	0.46
22:BA:634:C:H2'	22:BA:635:C:C6	2.51	0.46
22:DA:2138:G:N2	22:DA:2154:A:H1'	2.30	0.46
22:BA:734:A:C4	22:BA:735:A:C8	3.04	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:582:C:C4	1:AA:583:A:N7	2.84	0.46
31:BJ:21:THR:HA	31:BJ:61:LYS:HB3	1.98	0.46
22:DA:2744:G:N2	22:DA:2761:A:C4	2.84	0.46
22:DA:1114:C:H2'	22:DA:1115:G:C8	2.50	0.46
1:AA:605:U:O2'	1:AA:606:G:H5'	2.16	0.46
13:AM:79:ARG:O	13:AM:83:LEU:HG	2.16	0.46
22:BA:2273:A:H2'	22:BA:2274:A:C8	2.51	0.46
22:BA:151:C:H2'	22:BA:152:A:C8	2.51	0.46
39:DR:69:GLY:O	39:DR:70:GLU:C	2.54	0.46
22:BA:1816:C:C5	24:BC:62:TYR:CE2	3.03	0.46
29:BH:79:THR:HG23	29:BH:147:VAL:HB	1.98	0.45
22:DA:998:C:OP2	38:DQ:58:ARG:NH2	2.48	0.45
29:DH:39:ALA:O	29:DH:41:LYS:N	2.47	0.45
1:AA:554:A:H5'	12:AL:26:ALA:HB1	1.99	0.45
22:DA:2262:U:OP2	44:DW:16:SER:HB3	2.16	0.45
41:BT:3:ARG:CD	41:BT:5:GLU:HB2	2.45	0.45
41:BT:3:ARG:NE	41:BT:5:GLU:HB2	2.31	0.45
34:DM:66:ARG:HB2	34:DM:101:VAL:O	2.15	0.45
34:DM:76:LYS:HE2	34:DM:83:GLY:O	2.16	0.45
22:DA:2822:G:H2'	22:DA:2823:A:H5''	1.97	0.45
22:DA:1210:G:P	22:DA:1212:G:H5'	2.56	0.45
1:AA:1363:A:C4	1:AA:1365:G:C6	3.03	0.45
22:DA:1645:G:H5''	22:DA:1646:C:C5'	2.46	0.45
24:DC:230:HIS:NE2	24:DC:247:PRO:HA	2.31	0.45
1:CA:552:U:O2'	12:CL:83:ARG:O	2.29	0.45
22:DA:1585:C:C5	22:DA:1586:A:C5	3.05	0.45
22:DA:590:A:H2'	22:DA:591:U:C6	2.52	0.45
22:DA:2421:G:O6	51:D3:31:HIS:CD2	2.70	0.45
22:DA:2199:A:C5	22:DA:2225:A:C2	3.03	0.45
1:AA:987:G:H2'	1:AA:987:G:N3	2.31	0.45
1:AA:1150:A:N6	1:AA:1151:A:N6	2.65	0.45
22:BA:1737:G:C6	22:BA:1738:G:N1	2.84	0.45
22:DA:1331:G:O2'	22:DA:1332:G:H5'	2.16	0.45
22:DA:1275:A:C8	35:DN:16:HIS:CE1	3.03	0.45
2:AB:148:LEU:HA	2:AB:151:ILE:HG22	1.97	0.45
1:CA:955:U:O2'	1:CA:1227:A:N6	2.50	0.45
1:CA:957:U:H1'	1:CA:960:U:C4	2.51	0.45
22:DA:2867:G:C5	37:DP:21:ARG:NH2	2.85	0.45
22:DA:1438:U:C5	22:DA:1552:A:N1	2.84	0.45
24:DC:260:ASN:O	24:DC:261:LYS:CB	2.64	0.45
27:BF:105:THR:HG23	27:BF:106:ILE:HG23	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CB:126:PHE:CD1	2:CB:127:ASP:N	2.84	0.45
2:AB:16:PHE:O	2:AB:41:ILE:HD12	2.16	0.45
1:AA:1149:C:P	9:AI:11:ARG:HH21	2.39	0.45
22:BA:1176:U:N3	22:BA:1177:G:C6	2.84	0.45
1:CA:664:G:H2'	1:CA:666:G:OP1	2.17	0.45
22:DA:325:G:H2'	22:DA:326:G:O4'	2.16	0.45
22:BA:994:C:H1'	39:BR:10:LYS:HE3	1.98	0.45
29:DH:112:LYS:HG2	29:DH:113:SER:N	2.32	0.45
2:CB:169:GLU:O	2:CB:171:ILE:N	2.48	0.45
1:CA:121:U:H3'	1:CA:122:G:H5'	1.97	0.45
35:DN:49:GLU:N	35:DN:50:PRO:CD	2.79	0.45
1:AA:1109:C:OP2	3:AC:176:HIS:ND1	2.50	0.45
22:BA:1754:A:C6	22:BA:1755:A:C6	3.04	0.45
22:BA:2791:G:H2'	22:BA:2792:A:O4'	2.16	0.45
43:DV:14:LYS:HD3	43:DV:18:ARG:NH2	2.31	0.45
37:BP:4:ILE:H	37:BP:4:ILE:HD12	1.81	0.45
36:BO:100:HIS:CG	36:BO:101:GLY:N	2.83	0.45
1:AA:1456:A:H2'	1:AA:1457:G:O4'	2.16	0.45
29:BH:94:ILE:HG23	29:BH:98:ASP:CB	2.47	0.45
22:DA:1789:A:OP1	24:DC:220:VAL:HA	2.16	0.45
22:DA:2118:U:O4	22:DA:2149:U:H1'	2.16	0.45
22:DA:1317:G:N2	22:DA:1336:A:C2	2.83	0.45
22:BA:1131:G:C4	31:BJ:77:HIS:CD2	3.04	0.45
22:DA:410:G:C2	22:DA:2407:A:C6	3.04	0.45
46:BY:18:LEU:O	46:BY:22:LEU:CB	2.64	0.45
1:CA:578:C:O2	1:CA:579:A:C8	2.68	0.45
1:CA:32:A:C3'	1:CA:33:A:H8	2.29	0.45
22:DA:260:G:C6	22:DA:261:G:C5	3.03	0.45
12:CL:40:THR:HG22	12:CL:41:THR:N	2.31	0.45
1:AA:1198:G:H2'	1:AA:1199:U:H6	1.80	0.45
2:CB:211:THR:HA	2:CB:214:LEU:HB3	1.98	0.45
1:AA:1061:G:C6	1:AA:1197:A:C2	3.04	0.45
2:AB:96:TRP:CZ3	2:AB:175:GLU:OE2	2.69	0.45
3:CC:6:HIS:NE2	3:CC:184:TYR:HE2	2.14	0.45
22:BA:2520:C:O2'	22:BA:2521:C:H5'	2.16	0.45
22:DA:1076:C:H2'	22:DA:1077:A:O4'	2.17	0.45
22:DA:1666:G:O3'	32:DK:6:THR:HG23	2.16	0.45
36:DO:111:ARG:NH2	36:DO:117:PHE:OXT	2.48	0.45
8:CH:126:ILE:HG22	8:CH:127:CYS:CB	2.46	0.45
1:CA:115:G:C2	1:CA:289:G:N7	2.84	0.45
1:CA:115:G:H1'	1:CA:116:A:N7	2.32	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:597:G:N7	1:CA:598:U:C5	2.84	0.45
1:CA:1253:G:C2	1:CA:1285:A:N6	2.84	0.45
41:BT:71:GLY:O	41:BT:73:ARG:N	2.49	0.45
30:DI:61:VAL:HG22	30:DI:67:PHE:HB3	1.97	0.45
51:B3:61:CYS:O	51:B3:62:LEU:HD23	2.16	0.45
3:AC:14:ILE:O	3:AC:15:VAL:HG22	2.16	0.45
1:AA:246:A:C2	1:AA:282:A:C5	3.05	0.45
22:BA:2010:G:N7	57:BA:3381:HOH:O	2.36	0.45
1:CA:346:G:H2'	1:CA:347:G:O4'	2.15	0.45
22:BA:451:U:C2	22:BA:453:A:N7	2.84	0.45
1:AA:105:G:H2'	1:AA:106:C:C6	2.51	0.45
14:AN:15:ALA:O	14:AN:18:ASP:OD1	2.34	0.45
22:BA:2865:U:C4	22:BA:2866:U:C4	3.05	0.45
22:DA:912:C:N4	22:DA:913:U:O4	2.49	0.45
17:CQ:38:ILE:HD12	17:CQ:38:ILE:N	2.30	0.45
14:AN:100:SER:O	14:AN:101:TRP:HB3	2.16	0.45
16:AP:5:ARG:HA	16:AP:68:SER:OG	2.16	0.45
22:DA:2244:U:H2'	22:DA:2245:U:O4'	2.16	0.45
22:DA:1308:A:C6	22:DA:1309:G:C2	3.05	0.45
42:BU:82:ARG:O	42:BU:97:LYS:HG2	2.16	0.45
5:CE:115:LEU:HD12	5:CE:115:LEU:HA	1.87	0.45
1:CA:527:G:N2	1:CA:528:C:N1	2.64	0.45
1:CA:429:U:H5'	4:CD:9:LEU:HD23	1.97	0.45
38:DQ:28:ARG:HG2	38:DQ:34:VAL:HG12	1.98	0.45
5:CE:16:ILE:CD1	5:CE:38:VAL:HG23	2.46	0.45
5:CE:39:VAL:CG2	5:CE:71:MET:HE2	2.46	0.45
22:DA:526:A:C6	22:DA:2626:C:H4'	2.52	0.45
1:CA:269:C:H2'	1:CA:270:A:C8	2.52	0.45
1:CA:477:C:H2'	1:CA:478:A:C8	2.51	0.45
1:AA:457:G:O6	1:AA:458:U:N3	2.50	0.45
22:BA:1735:A:C2	22:BA:1736:U:N1	2.85	0.45
22:DA:482:A:C8	22:DA:506:G:C2	3.04	0.45
1:AA:1140:C:O2	1:AA:1141:C:C5	2.70	0.45
26:DE:61:ARG:HB3	26:DE:63:LYS:O	2.16	0.45
29:DH:34:GLY:O	29:DH:35:LYS:CD	2.65	0.45
22:BA:1045:C:H3'	22:BA:1046:A:C5'	2.46	0.45
1:AA:428:G:C5	1:AA:430:A:C6	3.04	0.45
1:CA:158:G:C5	1:CA:159:G:C8	3.04	0.45
1:AA:579:A:H2'	1:AA:580:C:C6	2.51	0.45
43:BV:41:GLU:O	43:BV:41:GLU:HG2	2.15	0.45
22:DA:1464:G:H2'	22:DA:1465:G:H8	1.80	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:BF:5:HIS:O	27:BF:8:TYR:HB3	2.16	0.45
22:DA:2652:C:N4	22:DA:2653:U:C4	2.84	0.45
2:CB:19:GLN:O	2:CB:38:VAL:CG2	2.64	0.45
39:DR:66:HIS:CE1	39:DR:94:THR:CG2	2.99	0.45
22:BA:1199:U:H1'	38:BQ:4:VAL:HG22	1.98	0.45
22:BA:2582:G:C2	22:BA:2583:G:C8	3.05	0.45
22:DA:2077:A:N1	22:DA:2078:C:C4	2.85	0.45
1:AA:284:C:H2'	1:AA:285:C:H6	1.81	0.45
22:DA:1262:A:C6	22:DA:1263:U:N3	2.85	0.45
3:AC:162:ILE:HG13	3:AC:163:ALA:N	2.30	0.45
1:AA:918:A:C6	1:AA:919:A:C6	3.05	0.45
22:BA:1638:C:H5''	22:BA:2710:C:O2'	2.16	0.45
22:DA:1422:G:N2	22:DA:1577:C:H1'	2.30	0.45
1:CA:1076:U:N3	1:CA:1082:A:C2	2.85	0.45
10:CJ:29:ALA:HB1	10:CJ:76:ILE:HD13	1.99	0.45
19:AS:62:VAL:HG12	19:AS:63:THR:N	2.32	0.45
22:BA:657:U:O5'	22:BA:657:U:H6	1.99	0.45
20:CT:22:ALA:O	20:CT:26:SER:HB2	2.15	0.45
6:AF:10:VAL:HG12	6:AF:11:HIS:N	2.31	0.45
21:CU:14:VAL:C	21:CU:16:LEU:HG	2.37	0.45
24:BC:199:GLU:O	24:BC:202:LEU:HB2	2.16	0.45
22:DA:1797:G:H4'	24:DC:255:LYS:O	2.15	0.45
1:AA:1412:C:H2'	1:AA:1413:A:C8	2.52	0.45
1:CA:980:C:O3'	14:CN:13:ARG:NH2	2.50	0.45
8:AH:9:ASP:O	8:AH:13:ARG:HB2	2.17	0.45
39:BR:98:ILE:HG21	39:BR:101:ILE:HD11	1.98	0.45
47:BZ:11:ARG:NH1	47:BZ:53:PHE:O	2.50	0.45
25:DD:183:GLU:OE1	25:DD:183:GLU:N	2.48	0.45
12:CL:18:LYS:HD2	12:CL:18:LYS:C	2.36	0.45
4:CD:20:PHE:N	4:CD:20:PHE:CD1	2.84	0.45
37:BP:33:VAL:HA	37:BP:37:LYS:O	2.16	0.45
50:B2:8:SER:OG	50:B2:11:LYS:HG3	2.17	0.45
1:AA:1427:C:C2'	1:AA:1428:A:H5'	2.46	0.45
22:DA:996:A:N6	22:DA:1160:G:C6	2.85	0.45
22:BA:1910:G:N2	22:BA:1921:G:H1'	2.31	0.45
22:DA:450:G:H2'	22:DA:451:U:H5''	1.97	0.45
1:AA:375:U:H4'	16:AP:17:TYR:CE2	2.52	0.45
22:DA:1985:C:H2'	22:DA:1986:C:H6	1.80	0.45
25:BD:104:VAL:HG23	25:BD:105:LYS:N	2.32	0.45
22:BA:2128:G:N2	22:BA:2173:A:O2'	2.49	0.45
1:CA:17:U:C2	1:CA:18:C:C5	3.04	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:CD:23:SER:HB2	4:CD:109:ALA:O	2.15	0.45
35:BN:103:ARG:CZ	35:BN:110:MET:HE1	2.47	0.45
5:AE:144:LEU:O	5:AE:147:MET:HB3	2.16	0.45
22:DA:297:G:H5''	42:DU:85:PHE:CB	2.43	0.45
33:BL:81:ASP:HA	33:BL:84:LYS:HE3	1.99	0.45
22:DA:1257:C:C4	22:DA:1258:U:O4	2.69	0.45
4:CD:168:PRO:HB3	4:CD:170:TRP:CZ3	2.52	0.45
1:CA:254:G:C2	1:CA:273:U:C2	3.04	0.45
22:DA:1277:G:H5'	35:DN:20:MET:HE2	1.96	0.45
1:AA:263:A:H2'	1:AA:264:C:C6	2.52	0.45
33:DL:90:VAL:HG13	33:DL:95:LEU:HD11	1.98	0.45
15:CO:88:ARG:HG3	15:CO:89:ARG:OXT	2.17	0.45
6:AF:92:THR:HG22	6:AF:93:LYS:N	2.31	0.45
14:AN:10:GLU:HA	14:AN:13:ARG:HG3	1.97	0.45
1:CA:223:A:C4	1:CA:224:U:C5	3.05	0.45
20:CT:82:GLN:O	20:CT:85:LYS:HB2	2.16	0.45
22:DA:2341:G:C2	22:DA:2342:C:C2	3.05	0.45
1:AA:1491:G:H2'	1:AA:1492:A:O4'	2.17	0.45
21:AU:17:ARG:NH1	21:AU:20:LYS:CG	2.79	0.45
31:DJ:30:THR:CG2	31:DJ:31:GLU:N	2.80	0.45
30:BI:76:ALA:CB	30:BI:129:ILE:HG23	2.45	0.45
22:BA:1344:U:C2'	22:BA:1345:C:OP1	2.64	0.45
1:CA:49:U:O4	1:CA:362:G:N2	2.47	0.45
26:BE:119:ILE:O	26:BE:187:VAL:HA	2.16	0.45
1:AA:1048:G:N2	1:AA:1050:G:C5	2.85	0.45
4:AD:25:VAL:O	4:AD:26:ARG:C	2.55	0.45
22:DA:254:G:OP2	51:D3:5:LYS:NZ	2.49	0.45
22:DA:1861:G:H22	22:DA:1882:U:H1'	1.80	0.45
1:AA:341:C:H2'	1:AA:342:C:H6	1.81	0.45
22:DA:1623:G:C6	22:DA:1624:U:C5	3.04	0.45
22:BA:500:G:N2	22:BA:502:A:H3'	2.31	0.45
22:BA:657:U:H2'	22:BA:658:U:C6	2.51	0.45
18:AR:42:SER:OG	18:AR:47:THR:HG23	2.17	0.45
4:AD:152:GLN:O	4:AD:155:VAL:HG12	2.16	0.45
24:DC:77:VAL:HG22	24:DC:78:VAL:N	2.32	0.45
22:DA:974:G:H1'	22:DA:975:A:C8	2.50	0.45
22:BA:1635:A:C6	22:BA:1636:U:C2	3.05	0.45
28:BG:69:ARG:C	28:BG:69:ARG:HD3	2.37	0.45
44:DW:56:ASP:OD2	44:DW:58:THR:OG1	2.35	0.45
36:DO:49:VAL:HG21	36:DO:82:ALA:HA	1.97	0.45
1:AA:1476:A:H2'	1:AA:1477:U:O4'	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:1248:G:O2'	38:DQ:3:ARG:HA	2.16	0.45
24:DC:120:VAL:HG13	24:DC:134:ASN:ND2	2.32	0.45
22:DA:1083:U:O2	22:DA:1086:A:N1	2.50	0.45
22:BA:839:U:H2'	22:BA:840:C:C6	2.51	0.45
43:BV:48:MET:SD	43:BV:86:LEU:HG	2.56	0.45
45:DX:64:ILE:O	45:DX:64:ILE:HD12	2.17	0.45
31:BJ:4:PHE:CD1	31:BJ:4:PHE:C	2.90	0.45
15:CO:25:THR:HG23	15:CO:66:LEU:HD12	1.99	0.45
6:AF:76:THR:O	6:AF:79:ARG:N	2.45	0.45
22:BA:1916:A:C2	22:BA:1917:U:C2	3.04	0.45
22:BA:1917:U:C3'	22:BA:1918:A:H5'	2.45	0.45
1:CA:1073:U:H5'	1:CA:1074:G:OP2	2.16	0.45
27:DF:122:PHE:O	27:DF:123:ASP:C	2.55	0.45
16:CP:6:LEU:HD11	16:CP:71:VAL:HG23	1.99	0.45
4:CD:202:GLU:OE1	5:CE:105:ILE:CG2	2.64	0.45
1:AA:1167:A:N7	1:AA:1169:A:C5	2.85	0.45
22:DA:2248:C:C2'	22:DA:2249:U:H5'	2.47	0.45
5:CE:68:ARG:O	5:CE:71:MET:HE3	2.17	0.45
1:AA:1278:G:H4'	1:AA:1279:G:C8	2.51	0.45
22:BA:2140:G:C2	22:BA:2152:G:N1	2.85	0.45
33:BL:95:LEU:HD22	33:BL:100:ILE:CD1	2.46	0.45
32:DK:34:GLY:O	32:DK:35:VAL:C	2.53	0.45
22:BA:2286:G:C2	49:B1:36:LEU:HD11	2.52	0.45
50:B2:43:THR:O	50:B2:44:VAL:HB	2.17	0.45
28:DG:11:VAL:HG23	28:DG:15:VAL:HB	1.98	0.45
3:AC:141:ALA:O	3:AC:146:ALA:CB	2.63	0.45
22:DA:136:G:C2	22:DA:144:A:C6	3.04	0.45
36:DO:33:ARG:O	36:DO:34:HIS:CB	2.64	0.45
14:AN:93:ILE:HG21	14:AN:96:LEU:HD22	1.98	0.45
22:DA:2297:A:H2'	22:DA:2297:A:N3	2.32	0.45
34:BM:62:LYS:HD3	34:BM:64:TRP:CH2	2.51	0.45
29:BH:40:THR:O	29:BH:42:LYS:N	2.48	0.45
22:DA:547:A:H3'	22:DA:548:G:C5'	2.47	0.45
15:AO:46:HIS:C	15:AO:48:LYS:H	2.20	0.45
22:DA:40:U:C4	22:DA:41:C:C4	3.05	0.45
10:AJ:53:ILE:HG22	10:AJ:61:ALA:CB	2.47	0.45
6:CF:51:ILE:CG1	6:CF:51:ILE:O	2.65	0.45
22:DA:7:G:H2'	22:DA:8:C:O4'	2.17	0.45
22:DA:980:A:C6	22:DA:981:A:N1	2.84	0.45
24:BC:258:ARG:NH1	24:BC:264:ASP:OD1	2.49	0.45
22:DA:324:A:N6	22:DA:338:G:O2'	2.50	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:449:A:H4'	38:BQ:3:ARG:NH1	2.32	0.45
22:BA:2307:G:H4'	22:BA:2308:G:O5'	2.17	0.45
1:AA:41:G:H2'	1:AA:42:G:C8	2.51	0.45
22:BA:2345:G:N3	22:BA:2381:A:H2'	2.32	0.45
42:DU:72:ILE:HD11	42:DU:83:VAL:HG23	1.98	0.45
46:DY:17:GLU:HA	46:DY:20:ASN:HB2	1.98	0.45
43:BV:8:VAL:HG23	43:BV:9:ARG:N	2.31	0.45
1:AA:1014:A:H2'	1:AA:1015:G:O4'	2.16	0.45
7:AG:98:ALA:HA	7:AG:101:MET:HE3	1.98	0.45
22:DA:341:C:H2'	22:DA:342:A:C8	2.50	0.45
24:DC:142:HIS:O	24:DC:143:ASN:CB	2.63	0.45
9:AI:128:SER:O	9:AI:129:LYS:C	2.55	0.45
1:CA:39:G:N2	1:CA:40:C:C2	2.85	0.45
22:DA:1073:A:O2'	22:DA:2474:U:H5'	2.17	0.45
22:DA:627:A:C2	22:DA:637:A:C4	3.04	0.45
1:AA:200:G:C2	1:AA:218:U:O2	2.70	0.45
39:DR:1:MET:HA	39:DR:42:ALA:O	2.16	0.45
22:BA:1956:U:H2'	22:BA:1957:C:H5'	1.98	0.45
1:AA:510:A:H5''	1:AA:511:C:P	2.57	0.45
24:BC:19:VAL:HG12	24:BC:19:VAL:O	2.16	0.45
13:CM:34:LEU:HB3	13:CM:39:ILE:HB	1.98	0.45
29:DH:147:VAL:HG12	29:DH:148:ALA:N	2.32	0.45
29:BH:72:ILE:HG23	29:BH:142:VAL:HG22	1.99	0.45
1:AA:533:A:P	57:AA:1850:HOH:O	2.62	0.45
22:BA:1088:A:H5''	22:BA:1088:A:N3	2.32	0.45
22:DA:2841:C:H2'	22:DA:2842:G:C8	2.52	0.45
4:CD:100:ASN:O	4:CD:104:ARG:HG2	2.17	0.45
33:DL:59:ARG:NH1	33:DL:59:ARG:HB3	2.31	0.45
22:DA:1314:C:C2	22:DA:1339:G:N2	2.84	0.45
2:AB:186:ILE:HD11	2:AB:204:ASP:HA	1.99	0.45
22:DA:1974:C:C2	22:DA:1975:G:C8	3.05	0.45
22:BA:1433:A:C2'	22:BA:1434:A:H5'	2.46	0.45
8:CH:114:ARG:O	8:CH:116:ALA:N	2.50	0.45
22:BA:674:G:H1'	26:BE:69:ARG:CD	2.46	0.45
24:DC:245:VAL:HG12	24:DC:251:GLN:HA	1.99	0.45
1:AA:1154:G:N3	1:AA:1154:G:H2'	2.30	0.45
22:BA:2847:U:O2'	22:BA:2848:G:H5'	2.17	0.45
1:AA:207:C:H2'	1:AA:208:U:C2	2.51	0.45
40:DS:29:VAL:HG11	40:DS:55:ILE:HD11	1.98	0.45
20:AT:67:ILE:HG13	20:AT:71:LYS:CG	2.46	0.45
32:DK:103:VAL:O	32:DK:122:VAL:HB	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BJ:139:VAL:CG1	31:BJ:140:LEU:N	2.76	0.45
51:B3:23:LYS:HA	51:B3:48:ALA:O	2.17	0.45
7:CG:66:LEU:HB2	7:CG:104:ILE:HD11	1.99	0.45
7:CG:75:VAL:HG21	7:CG:144:MET:HG2	1.99	0.45
22:DA:866:A:O4'	22:DA:914:G:N2	2.49	0.45
22:BA:855:G:C2'	22:BA:856:G:O5'	2.65	0.45
22:BA:826:U:H2'	22:BA:828:U:O4'	2.17	0.45
1:CA:878:A:C6	1:CA:879:C:C4	3.05	0.45
3:CC:57:ILE:HG13	3:CC:66:VAL:HG22	1.97	0.45
46:DY:23:ARG:NH1	46:DY:23:ARG:HA	2.32	0.45
31:DJ:37:ARG:HA	31:DJ:118:MET:SD	2.56	0.45
4:AD:152:GLN:O	4:AD:153:SER:C	2.53	0.45
34:BM:73:ILE:HG21	34:BM:91:TYR:CZ	2.51	0.45
34:DM:110:GLU:O	34:DM:114:ARG:HG3	2.16	0.45
13:CM:90:ARG:HB3	13:CM:97:VAL:HG22	1.99	0.45
13:AM:5:ALA:CB	13:AM:22:ILE:HG12	2.46	0.45
27:DF:63:GLN:HG3	27:DF:95:ARG:HD2	1.99	0.45
22:BA:2178:C:H2'	22:BA:2179:C:C6	2.52	0.45
12:CL:61:PHE:CD1	12:CL:61:PHE:N	2.84	0.45
11:CK:72:ASP:OD2	11:CK:73:ALA:N	2.49	0.45
30:BI:22:PRO:HB2	30:BI:23:PRO:HD3	1.99	0.45
1:AA:1360:A:H2'	1:AA:1361:G:O4'	2.16	0.45
1:AA:1272:G:H2'	1:AA:1273:C:O4'	2.16	0.45
22:BA:300:A:OP2	42:BU:97:LYS:NZ	2.50	0.45
22:DA:1793:C:C2	22:DA:1794:A:C8	3.05	0.45
22:DA:118:A:H1'	22:DA:178:G:O4'	2.16	0.45
11:AK:125:LYS:HE3	21:AU:35:ARG:NH2	2.32	0.45
25:BD:103:ASP:CG	25:BD:104:VAL:N	2.69	0.45
22:BA:2318:G:C6	22:BA:2319:G:N1	2.85	0.45
4:CD:26:ARG:O	4:CD:27:ALA:CB	2.63	0.45
1:AA:90:C:C2	1:AA:91:U:C5	3.05	0.45
1:CA:780:A:C2	1:CA:803:G:C6	3.05	0.45
1:AA:1358:U:H5	1:AA:1359:C:C4	2.35	0.45
4:AD:23:SER:O	4:AD:24:GLY:O	2.33	0.45
22:DA:528:A:N1	22:DA:2043:C:H4'	2.32	0.45
29:DH:31:VAL:CG1	29:DH:32:PRO:HD3	2.47	0.45
22:DA:1651:G:N2	22:DA:1652:A:H1'	2.31	0.45
1:CA:321:A:C8	1:CA:328:C:C2	3.04	0.45
42:DU:98:SER:O	42:DU:99:ASN:CB	2.65	0.45
12:CL:52:VAL:HG23	12:CL:65:SER:O	2.17	0.45
5:CE:18:VAL:HG21	5:CE:56:VAL:CG1	2.47	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1463:U:H2'	1:AA:1464:U:C6	2.52	0.45
22:DA:204:A:O4'	22:DA:206:U:C6	2.69	0.45
26:DE:106:LYS:HG3	26:DE:200:LEU:HD23	1.98	0.45
22:BA:1984:G:C2	22:BA:1985:C:C6	3.05	0.45
23:BB:104:A:H2'	23:BB:105:G:O4'	2.17	0.45
33:DL:77:ILE:HB	33:DL:109:LYS:O	2.17	0.45
13:AM:3:ARG:O	13:AM:8:ASN:O	2.35	0.45
22:DA:19:A:C2	22:DA:522:A:C2	3.04	0.45
42:DU:13:VAL:CG2	42:DU:39:ILE:HG21	2.47	0.45
1:AA:617:G:N1	1:AA:618:C:C5	2.84	0.45
1:CA:747:A:C6	1:CA:748:G:C6	3.05	0.45
1:CA:67:C:O2'	1:CA:171:A:N3	2.49	0.45
1:AA:1368:A:P	9:AI:114:LYS:O	2.75	0.45
45:DX:54:LYS:HA	45:DX:57:ARG:HB2	1.99	0.45
22:BA:60:G:C8	22:BA:62:U:C6	3.05	0.45
22:BA:1496:A:H2'	22:BA:1498:C:C5	2.52	0.45
3:CC:117:ALA:HB2	3:CC:200:VAL:HG12	1.99	0.45
19:AS:64:ASP:CB	27:BF:115:ARG:NH2	2.80	0.45
39:DR:66:HIS:CG	39:DR:94:THR:HG22	2.52	0.45
22:BA:1820:U:H4'	22:BA:1821:A:OP2	2.16	0.45
27:BF:57:LEU:CD2	27:BF:152:LEU:HD11	2.47	0.45
1:CA:296:U:O2'	1:CA:556:C:O2	2.29	0.45
3:AC:11:ARG:O	3:AC:14:ILE:O	2.35	0.45
22:DA:1248:G:C2	38:DQ:3:ARG:HD2	2.52	0.45
27:DF:36:LEU:HB2	27:DF:89:VAL:HB	1.98	0.45
22:DA:92:U:C6	22:DA:93:G:C8	3.05	0.45
22:DA:797:G:H5''	26:DE:55:SER:HB2	1.98	0.45
1:AA:380:G:C2	1:AA:384:G:C6	3.04	0.45
1:CA:260:G:H2'	1:CA:261:U:C6	2.52	0.45
31:DJ:13:ARG:HD3	31:DJ:51:GLY:O	2.17	0.45
25:DD:29:VAL:HB	25:DD:98:VAL:CG2	2.46	0.45
11:AK:61:PHE:O	11:AK:64:GLN:HB3	2.17	0.45
40:BS:59:GLU:HG3	40:BS:66:ILE:HD11	1.99	0.45
22:DA:2022:U:P	57:DA:3661:HOH:O	2.75	0.45
22:BA:2527:C:C2'	22:BA:2528:U:H5'	2.47	0.45
1:AA:1250:A:H2'	1:AA:1251:A:C8	2.52	0.45
32:DK:64:ARG:HD3	32:DK:102:PRO:O	2.16	0.45
29:DH:93:SER:HB3	29:DH:123:ARG:HG3	1.99	0.45
29:BH:76:GLU:HA	29:BH:142:VAL:CG1	2.46	0.45
25:DD:151:THR:HG22	25:DD:152:PRO:N	2.32	0.45
22:BA:1916:A:N1	22:BA:1917:U:C2	2.85	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:1071:C:C2	1:CA:1072:G:C8	3.04	0.45
24:DC:160:THR:H	24:DC:195:VAL:HG13	1.81	0.45
22:BA:273:G:C2	22:BA:365:U:O2	2.69	0.45
22:DA:1794:A:H2'	22:DA:1795:C:H6	1.81	0.45
22:DA:1791:A:C4'	24:DC:207:LYS:O	2.65	0.45
22:DA:1809:A:C5	22:DA:1810:A:N7	2.84	0.45
22:DA:118:A:N7	22:DA:119:A:N7	2.65	0.45
22:DA:149:A:N7	22:DA:150:U:C5	2.84	0.45
1:AA:797:C:OP1	11:AK:126:LYS:NZ	2.33	0.45
22:DA:1767:G:O6	22:DA:1986:C:N4	2.50	0.45
4:CD:25:VAL:HG12	4:CD:26:ARG:N	2.31	0.45
38:BQ:58:ARG:HA	38:BQ:61:TRP:CE3	2.51	0.45
22:DA:89:A:C2	22:DA:90:U:N3	2.85	0.45
1:AA:1206:G:C6	1:AA:1207:G:C5	3.05	0.45
22:BA:2515:C:O2	22:BA:2570:G:C2	2.70	0.45
19:CS:34:TRP:HA	19:CS:52:HIS:HB2	1.99	0.45
22:BA:2086:U:H2'	22:BA:2087:G:C8	2.52	0.45
25:BD:133:THR:HG23	25:BD:134:HIS:CG	2.52	0.45
30:DI:57:VAL:CG2	30:DI:69:PHE:HB2	2.46	0.45
7:CG:37:SER:OG	9:CI:43:THR:OG1	2.30	0.45
2:AB:148:LEU:HA	2:AB:151:ILE:CG2	2.47	0.45
22:DA:2359:C:N4	22:DA:2360:G:C6	2.85	0.45
29:BH:12:LEU:HG	29:BH:13:GLY:N	2.31	0.45
13:AM:40:ALA:O	13:AM:43:VAL:HG22	2.17	0.45
22:BA:1971:U:H5'	22:BA:1972:G:H5''	1.98	0.45
25:DD:176:ASP:O	25:DD:189:VAL:HA	2.16	0.45
14:AN:27:LEU:O	14:AN:28:LYS:HB3	2.17	0.45
22:BA:2056:G:H2'	22:BA:2056:G:N3	2.32	0.45
22:DA:2201:G:C4	22:DA:2202:U:C6	3.04	0.45
4:AD:30:THR:HG22	4:AD:31:LYS:H	1.81	0.45
22:DA:807:U:H2'	22:DA:808:G:O4'	2.17	0.45
1:CA:537:G:OP1	12:CL:110:ARG:NH2	2.47	0.45
46:DY:9:LYS:N	46:DY:12:GLU:HG3	2.32	0.45
22:DA:2077:A:C2	22:DA:2078:C:C6	3.05	0.45
26:DE:46:GLN:O	26:DE:88:ARG:NH1	2.49	0.45
3:AC:87:LEU:O	3:AC:91:VAL:HG23	2.17	0.45
22:DA:266:G:C2	22:DA:427:U:O2	2.70	0.45
1:CA:1356:G:H2'	1:CA:1357:A:C8	2.51	0.45
14:CN:64:CYS:SG	14:CN:83:LYS:HG3	2.57	0.45
48:B0:39:LEU:O	48:B0:40:ARG:C	2.55	0.45
11:AK:109:ASN:HB3	21:AU:7:ARG:HA	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:828:U:H2'	1:CA:829:G:O5'	2.16	0.45
41:DT:13:ALA:HB1	41:DT:14:PRO:HD2	1.99	0.45
22:DA:873:C:N3	22:DA:905:A:C2	2.84	0.45
32:BK:118:LEU:O	32:BK:119:ALA:HB3	2.16	0.45
22:DA:2868:A:C6	22:DA:2869:G:C6	3.04	0.45
22:BA:892:A:N3	22:BA:892:A:H2'	2.30	0.45
26:DE:77:ILE:O	26:DE:77:ILE:HG13	2.17	0.45
6:AF:67:PRO:O	6:AF:69:GLU:N	2.42	0.45
2:CB:123:ASP:O	2:CB:124:GLY:C	2.55	0.45
1:AA:714:G:N3	1:AA:777:A:H1'	2.32	0.45
4:CD:97:ARG:O	4:CD:101:VAL:HG23	2.17	0.45
2:AB:78:GLU:C	2:AB:80:VAL:N	2.70	0.45
31:BJ:17:VAL:HG22	31:BJ:55:ILE:HB	1.99	0.45
22:DA:187:G:N1	22:DA:210:C:C2	2.85	0.45
22:BA:977:G:N7	57:BA:3598:HOH:O	2.50	0.45
22:DA:1830:C:H5'	24:DC:15:HIS:HE1	1.82	0.45
22:DA:2064:C:H2'	22:DA:2065:C:H6	1.82	0.45
22:DA:806:C:O2'	22:DA:2445:G:H4'	2.17	0.45
22:DA:377:G:C5	22:DA:378:C:C4	3.05	0.45
1:CA:320:A:H2'	1:CA:321:A:O4'	2.16	0.45
22:BA:1378:A:H4'	22:BA:1379:U:OP1	2.16	0.45
22:DA:777:G:N7	22:DA:793:A:C2	2.80	0.45
22:BA:1327:A:H2'	22:BA:1328:A:O4'	2.17	0.45
22:DA:362:A:N7	22:DA:363:G:N7	2.65	0.45
22:BA:141:G:H3'	22:BA:142:A:C8	2.52	0.45
24:DC:153:GLN:C	24:DC:156:ARG:HD2	2.37	0.45
22:DA:391:A:N7	22:DA:392:U:C5	2.85	0.45
22:BA:2887:A:C5'	22:BA:2888:C:OP2	2.64	0.45
22:DA:1833:C:C4	22:DA:1834:U:C4	3.05	0.45
7:CG:65:ALA:HA	7:CG:128:ALA:HA	1.98	0.45
1:CA:73:C:C2	1:CA:74:A:C8	3.05	0.45
1:CA:476:U:O2'	1:CA:477:C:H5'	2.16	0.45
1:AA:1152:A:C6	1:AA:1153:G:C6	3.05	0.45
22:DA:141:G:N2	22:DA:142:A:C2	2.85	0.45
23:DB:27:C:OP1	36:DO:34:HIS:NE2	2.50	0.45
1:CA:775:G:C5	1:CA:776:G:N7	2.85	0.45
22:DA:68:G:N2	22:DA:69:C:H1'	2.31	0.45
22:DA:125:A:H5''	50:D2:19:ARG:HD3	1.99	0.45
22:DA:2516:A:C6	22:DA:2517:C:C4	3.04	0.45
9:CI:19:VAL:HG21	9:CI:82:GLY:HA3	1.98	0.45
1:AA:557:G:N1	1:AA:558:G:C2	2.85	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:851:C:H2'	22:BA:852:U:C6	2.51	0.45
1:AA:443:C:O2'	1:AA:444:G:H5'	2.17	0.45
34:DM:56:ALA:O	34:DM:58:LYS:N	2.48	0.45
1:AA:1048:G:H5''	14:AN:3:LYS:HG3	1.98	0.45
22:BA:2491:U:HO2'	22:BA:2492:U:H5	1.65	0.45
1:CA:1138:G:H2'	1:CA:1140:C:H5'	1.98	0.45
22:DA:1446:C:C4	22:DA:1447:C:C4	3.05	0.45
22:DA:1230:A:H2'	22:DA:1231:U:C6	2.52	0.45
2:AB:187:VAL:HG23	2:AB:187:VAL:O	2.17	0.45
28:DG:86:LYS:HB3	28:DG:165:ALA:CB	2.47	0.45
42:DU:43:LYS:HA	42:DU:60:GLU:HA	1.98	0.45
11:CK:91:PRO:O	11:CK:92:GLY:O	2.35	0.45
38:DQ:9:ILE:HG13	38:DQ:10:ALA:N	2.32	0.45
2:CB:152:LYS:HG3	2:CB:153:ASP:OD2	2.17	0.45
22:DA:2728:U:O2'	22:DA:2729:G:H5''	2.17	0.45
22:DA:2478:A:N7	22:DA:2529:G:C6	2.85	0.45
37:DP:99:TYR:CE2	37:DP:100:LEU:HD21	2.52	0.45
37:BP:25:THR:HB	37:BP:88:ARG:HB3	1.99	0.45
49:D1:11:LEU:HB2	49:D1:21:TYR:HB2	1.98	0.45
22:DA:2110:G:C6	22:DA:2120:G:C8	3.05	0.45
23:DB:28:C:OP1	36:DO:36:TYR:OH	2.17	0.45
29:DH:86:ASP:C	29:DH:88:GLY:H	2.19	0.45
38:DQ:72:ASN:HB3	38:DQ:110:VAL:HG11	1.98	0.45
22:BA:92:U:H2'	22:BA:92:U:O2	2.17	0.45
2:AB:24:ASN:OD1	2:AB:24:ASN:C	2.55	0.45
30:BI:67:PHE:N	30:BI:67:PHE:CD1	2.85	0.45
6:AF:49:TYR:O	6:AF:49:TYR:CD1	2.70	0.45
22:BA:999:U:C5	22:BA:1154:G:C5	3.05	0.45
1:CA:1225:A:H2'	1:CA:1226:C:C5	2.52	0.45
51:B3:36:LYS:O	51:B3:41:LYS:HE2	2.16	0.45
1:CA:142:G:C2	1:CA:143:A:H1'	2.52	0.45
12:AL:24:LEU:HG	12:AL:25:GLU:H	1.82	0.45
22:BA:1097:U:H3'	22:BA:1098:A:H4'	1.99	0.45
17:AQ:16:LYS:H	17:AQ:17:MET:HE1	1.82	0.45
22:BA:626:A:H2'	33:BL:78:ARG:NH1	2.31	0.45
1:AA:1226:C:C4	13:AM:103:LYS:HA	2.52	0.45
22:DA:1791:A:C8	22:DA:1792:G:C8	3.05	0.45
1:CA:1522:U:O2	1:CA:1523:G:C8	2.70	0.45
22:DA:27:G:HO2'	22:DA:28:A:P	2.38	0.45
22:DA:1773:A:N3	22:DA:1978:A:H2	2.14	0.45
22:DA:82:U:N3	22:DA:83:A:N7	2.65	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:CD:124:MET:HG3	4:CD:146:ARG:HG2	1.99	0.45
22:BA:2844:G:H2'	22:BA:2845:U:O4'	2.18	0.45
1:CA:553:A:O2'	12:CL:26:ALA:O	2.32	0.45
1:CA:577:G:N7	1:CA:816:A:N1	2.65	0.45
22:DA:2019:A:H4'	38:DQ:34:VAL:HG22	1.98	0.45
22:DA:1363:C:H2'	22:DA:1364:G:C8	2.52	0.45
22:BA:1109:C:C5	22:BA:1110:G:C6	3.05	0.45
22:DA:600:G:H2'	22:DA:601:C:C6	2.52	0.45
22:BA:1324:G:C4	22:BA:1328:A:N6	2.85	0.45
1:CA:1079:G:H2'	1:CA:1080:A:C8	2.52	0.45
1:AA:654:G:C4	1:AA:655:A:C8	3.05	0.45
22:DA:1239:G:C5	22:DA:1240:U:C5	3.05	0.45
22:DA:1242:U:H2'	22:DA:1243:C:O4'	2.17	0.45
25:DD:101:PHE:O	25:DD:104:VAL:HG22	2.17	0.45
1:AA:945:G:C2	1:AA:1337:G:C2	3.05	0.45
14:CN:46:LEU:HD21	19:CS:10:PHE:CD1	2.52	0.45
22:BA:1936:A:H2	22:BA:1943:U:H3	1.63	0.45
2:CB:64:LYS:HD3	2:CB:65:GLY:N	2.32	0.45
2:CB:81:LYS:HG3	2:CB:91:PHE:CE2	2.52	0.45
22:DA:1324:G:C2	22:DA:1328:A:N6	2.85	0.45
1:CA:774:G:C5	1:CA:775:G:C8	3.04	0.45
22:DA:271:G:C2	22:DA:367:G:C2	3.05	0.45
43:DV:9:ARG:HG3	43:DV:41:GLU:HB3	1.97	0.45
22:BA:248:G:H5'	22:BA:250:G:N7	2.32	0.45
1:CA:463:U:H5'	1:CA:464:U:OP2	2.17	0.45
22:DA:858:G:N2	22:DA:919:U:O4	2.48	0.45
14:AN:25:ALA:O	14:AN:28:LYS:HG2	2.17	0.45
6:CF:4:TYR:CD2	6:CF:71:ILE:HG12	2.52	0.45
1:AA:358:U:H2'	1:AA:359:G:H8	1.81	0.45
22:DA:43:G:N2	22:DA:437:U:C2	2.85	0.45
45:DX:40:VAL:HG23	45:DX:45:ARG:O	2.17	0.45
19:AS:58:VAL:O	19:AS:58:VAL:CG2	2.65	0.45
22:BA:476:G:H4'	22:BA:502:A:N1	2.32	0.45
1:AA:200:G:N2	1:AA:218:U:O2	2.49	0.45
1:CA:1226:C:C4	13:CM:103:LYS:HA	2.52	0.45
22:DA:1783:A:H5'	22:DA:2608:G:H4'	1.99	0.45
22:BA:2484:G:OP1	34:BM:44:ARG:HD3	2.17	0.45
22:BA:2024:G:OP2	22:BA:2034:U:H4'	2.17	0.45
43:DV:75:GLN:HB2	43:DV:92:VAL:HG23	1.99	0.45
22:BA:468:G:N7	50:B2:39:ARG:NH2	2.65	0.45
5:AE:57:PRO:HA	5:AE:60:ILE:HG12	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:BC:30:PHE:CZ	24:BC:32:PRO:HG2	2.52	0.45
22:DA:2107:G:H2'	22:DA:2108:A:O4'	2.16	0.45
22:DA:277:G:H3'	22:DA:277:G:N3	2.32	0.45
10:AJ:81:GLU:O	10:AJ:84:VAL:HG12	2.17	0.45
40:DS:32:ALA:O	40:DS:35:ILE:HB	2.17	0.45
35:BN:38:LEU:HB3	35:BN:39:PRO:HD3	1.98	0.45
22:BA:2341:G:H2'	22:BA:2342:C:O4'	2.16	0.45
22:DA:1932:A:N3	22:DA:1932:A:H2'	2.31	0.45
22:DA:60:G:HO2'	22:DA:62:U:P	2.39	0.45
29:DH:15:LEU:HD22	29:DH:15:LEU:N	2.32	0.45
41:BT:26:LYS:O	41:BT:26:LYS:HG2	2.17	0.45
27:BF:6:ASP:O	27:BF:7:TYR:C	2.55	0.45
1:CA:238:A:C5	1:CA:239:U:C5	3.05	0.45
22:BA:1431:A:H2'	22:BA:1432:G:C8	2.52	0.45
22:BA:1912:A:C2	22:BA:1919:A:C4	3.04	0.44
22:BA:1921:G:C2	22:BA:1922:G:N7	2.85	0.44
31:BJ:17:VAL:CG1	31:BJ:57:LEU:HD12	2.47	0.44
1:CA:1124:G:C2	1:CA:1127:G:C2	3.05	0.44
1:AA:1344:C:O2'	1:AA:1345:U:H5'	2.17	0.44
37:DP:65:SER:O	37:DP:66:ASN:C	2.54	0.44
22:DA:705:A:N3	22:DA:727:A:H1'	2.32	0.44
1:CA:107:G:H2'	1:CA:108:G:H5''	1.98	0.44
1:CA:31:G:C8	1:CA:306:A:H1'	2.51	0.44
22:DA:2248:C:H2'	22:DA:2249:U:H5'	1.99	0.44
22:DA:2093:G:O6	22:DA:2225:A:C8	2.70	0.44
22:DA:770:G:C4	22:DA:771:G:C8	3.05	0.44
22:DA:740:C:H5''	22:DA:1784:A:OP1	2.17	0.44
1:CA:756:C:N3	1:CA:757:U:C6	2.85	0.44
22:DA:1638:C:H4'	22:DA:2710:C:O2	2.17	0.44
22:BA:2315:G:C2	22:BA:2316:G:C4	3.05	0.44
1:AA:113:G:H2'	1:AA:114:U:C6	2.52	0.44
9:AI:52:LEU:HB3	9:AI:57:MET:CG	2.47	0.44
22:DA:460:A:OP2	50:D2:41:ARG:NH1	2.50	0.44
22:BA:1057:A:N6	22:BA:1087:G:OP2	2.50	0.44
22:DA:1930:G:H1'	22:DA:1968:G:N1	2.32	0.44
22:DA:1131:G:OP1	31:DJ:82:GLY:HA2	2.16	0.44
22:DA:1596:A:C6	22:DA:1597:A:C6	3.05	0.44
1:CA:512:U:H2'	1:CA:513:C:C6	2.52	0.44
17:CQ:45:HIS:ND1	17:CQ:70:THR:HG21	2.32	0.44
48:B0:55:ILE:O	48:B0:56:ALA:HB3	2.17	0.44
22:DA:1327:A:C6	22:DA:1328:A:C4	3.05	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:DC:34:LEU:O	24:DC:35:GLU:CB	2.65	0.44
22:DA:681:G:C2	22:DA:682:G:C8	3.05	0.44
20:AT:67:ILE:HA	20:AT:67:ILE:HD12	1.84	0.44
22:BA:1061:U:O4	30:BI:11:LEU:HA	2.17	0.44
22:BA:947:A:O2'	22:BA:984:A:H2	1.97	0.44
1:CA:129:A:H1'	1:CA:130:A:C8	2.52	0.44
22:BA:226:A:N6	22:BA:227:A:C6	2.86	0.44
22:DA:2512:C:H2'	22:DA:2513:A:O4'	2.17	0.44
20:AT:44:LYS:HB3	20:AT:87:ALA:HB1	2.00	0.44
47:BZ:7:ILE:HB	47:BZ:36:VAL:HG12	1.98	0.44
17:CQ:11:ARG:NE	17:CQ:12:VAL:O	2.49	0.44
1:AA:1037:C:H2'	1:AA:1038:C:C6	2.53	0.44
7:CG:78:ARG:HB2	7:CG:85:TYR:HB2	1.99	0.44
29:DH:5:LEU:CD1	29:DH:13:GLY:CA	2.95	0.44
15:AO:82:ILE:HB	15:AO:87:LEU:HD23	1.99	0.44
46:BY:36:GLN:O	46:BY:37:LEU:HB3	2.18	0.44
24:DC:168:ASP:OD1	24:DC:168:ASP:N	2.45	0.44
6:AF:99:ALA:O	6:AF:100:SER:CB	2.65	0.44
22:BA:1857:G:C2	22:BA:1884:G:N3	2.85	0.44
11:AK:52:PHE:HB3	11:AK:56:ARG:HB3	2.00	0.44
41:DT:49:LYS:HD3	41:DT:49:LYS:N	2.31	0.44
22:BA:2386:A:H2'	22:BA:2387:U:O4'	2.17	0.44
22:BA:350:G:H2'	22:BA:351:C:O4'	2.17	0.44
2:AB:33:GLY:O	2:AB:34:ALA:HB2	2.17	0.44
22:DA:2457:U:C4	22:DA:2458:G:C6	3.05	0.44
22:BA:2115:G:HO2'	22:BA:2116:G:H8	1.65	0.44
22:BA:2038:G:H2'	22:BA:2039:U:O4'	2.18	0.44
32:DK:26:GLY:O	32:DK:30:ARG:HD2	2.18	0.44
37:BP:10:GLN:HA	37:BP:13:MET:HG3	1.99	0.44
38:BQ:41:LYS:HB2	38:BQ:41:LYS:HE3	1.81	0.44
22:DA:2647:U:O2	22:DA:2647:U:H2'	2.17	0.44
38:BQ:72:ASN:HD22	38:BQ:72:ASN:N	2.15	0.44
22:DA:2581:G:H2'	22:DA:2581:G:N3	2.31	0.44
24:DC:125:LYS:O	24:DC:192:LEU:HB2	2.18	0.44
3:CC:53:SER:HB3	3:CC:115:LEU:HD21	1.98	0.44
45:BX:32:ASN:O	45:BX:52:SER:HA	2.17	0.44
26:BE:113:VAL:HG22	26:BE:118:LEU:HD23	2.00	0.44
1:CA:356:A:H2'	1:CA:357:G:O4'	2.17	0.44
22:BA:1910:G:C5	22:BA:1911:U:C4	3.05	0.44
22:DA:1606:C:O2'	22:DA:1607:C:OP2	2.35	0.44
2:CB:103:ASN:C	2:CB:105:LYS:N	2.70	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:1088:A:N3	22:BA:1088:A:H3'	2.32	0.44
22:BA:1092:C:H2'	22:BA:1093:G:O4'	2.16	0.44
22:DA:2127:G:H4'	22:DA:2128:G:OP1	2.17	0.44
33:DL:63:LYS:HA	51:D3:13:ARG:HG3	1.99	0.44
22:DA:1992:G:H4'	22:DA:1993:U:OP1	2.17	0.44
22:DA:1034:G:C6	22:DA:1035:U:N3	2.85	0.44
1:AA:406:G:C6	1:AA:495:A:C8	3.05	0.44
12:CL:99:ARG:HB2	12:CL:117:TYR:HA	1.99	0.44
29:BH:97:ARG:NH1	1:CA:369:G:O2'	2.51	0.44
1:CA:671:G:C6	1:CA:672:U:C4	3.05	0.44
22:DA:1090:A:C6	22:DA:1091:G:N7	2.86	0.44
22:BA:142:A:C6	22:BA:143:C:N3	2.85	0.44
7:AG:75:VAL:HB	7:AG:86:GLN:HG3	1.98	0.44
15:CO:63:ARG:NH1	15:CO:87:LEU:CD1	2.80	0.44
11:AK:24:HIS:O	11:AK:30:THR:HA	2.18	0.44
37:DP:40:LEU:HD23	37:DP:40:LEU:C	2.38	0.44
1:CA:563:A:H2'	1:CA:567:G:C8	2.53	0.44
1:AA:1078:U:O4	1:AA:1079:G:C6	2.71	0.44
22:DA:144:A:N3	22:DA:144:A:H2'	2.32	0.44
1:AA:1141:C:C2	1:AA:1142:G:C8	3.05	0.44
31:DJ:80:HIS:O	31:DJ:81:ILE:C	2.56	0.44
22:BA:1223:G:N7	39:BR:71:LYS:NZ	2.65	0.44
22:BA:1984:G:C5	22:BA:1985:C:C5	3.05	0.44
30:DI:136:MET:HG3	30:DI:138:LEU:HD11	1.99	0.44
22:DA:734:A:C8	22:DA:735:A:C8	3.05	0.44
20:AT:67:ILE:HG13	20:AT:71:LYS:HD3	1.98	0.44
1:CA:683:G:H2'	1:CA:684:U:C6	2.52	0.44
1:CA:159:G:H22	1:CA:161:A:H3'	1.82	0.44
22:DA:858:G:C4	22:DA:2268:A:C2	3.05	0.44
4:AD:98:LEU:HD23	4:AD:118:VAL:HG11	1.98	0.44
17:CQ:8:LEU:HB2	17:CQ:61:ILE:CG2	2.47	0.44
1:CA:66:A:N6	1:CA:67:C:C4	2.85	0.44
1:CA:663:A:H5'	1:CA:836:G:OP1	2.17	0.44
34:BM:7:THR:O	34:BM:8:LYS:C	2.54	0.44
22:DA:993:G:O2'	22:DA:994:C:H5'	2.17	0.44
1:AA:716:A:C6	1:AA:717:U:N3	2.85	0.44
30:DI:53:LEU:HD11	30:DI:82:LYS:HE2	1.98	0.44
18:CR:23:TYR:HA	18:CR:58:ALA:HB1	1.98	0.44
22:DA:264:C:O3'	22:DA:265:A:H2'	2.17	0.44
22:DA:2478:A:C8	22:DA:2529:G:C5	3.05	0.44
7:AG:115:SER:HB3	7:AG:118:LEU:HG	1.98	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:AT:19:LYS:O	20:AT:22:ALA:HB3	2.17	0.44
22:BA:2388:A:H5'	22:BA:2389:G:OP2	2.18	0.44
35:DN:2:ARG:O	35:DN:3:HIS:C	2.55	0.44
10:CJ:19:ASP:HA	10:CJ:22:THR:HB	1.99	0.44
9:AI:87:LEU:O	9:AI:94:LEU:HD11	2.18	0.44
8:CH:89:LYS:HG3	8:CH:90:ASP:N	2.32	0.44
22:DA:2104:C:H2'	22:DA:2105:U:O4'	2.17	0.44
26:DE:136:GLN:NE2	26:DE:140:ASP:OD1	2.50	0.44
25:BD:38:LYS:O	25:BD:46:ARG:HA	2.17	0.44
3:AC:147:LYS:HB2	3:AC:203:PHE:CD2	2.52	0.44
22:DA:61:C:OP1	46:DY:44:LYS:HD3	2.17	0.44
7:CG:36:LYS:O	7:CG:36:LYS:HG3	2.17	0.44
44:BW:56:ASP:O	44:BW:57:HIS:HB2	2.17	0.44
1:AA:622:A:C8	1:AA:623:C:C6	3.05	0.44
1:AA:1533:C:H5'	1:AA:1534:A:OP1	2.17	0.44
22:BA:1692:U:H2'	22:BA:1694:C:C5	2.52	0.44
22:DA:2314:A:C2	22:DA:2315:G:C5	3.05	0.44
29:BH:99:ILE:HD11	29:BH:121:VAL:HG11	2.00	0.44
22:DA:1153:C:H2'	22:DA:1154:G:C8	2.52	0.44
22:BA:1180:U:C2'	22:BA:1181:U:H5'	2.48	0.44
22:DA:1308:A:N6	22:DA:1309:G:N1	2.65	0.44
22:BA:761:A:P	57:BA:3701:HOH:O	2.74	0.44
31:BJ:80:HIS:O	31:BJ:83:GLY:N	2.47	0.44
1:AA:1377:A:O2'	7:AG:2:PRO:HB3	2.18	0.44
22:DA:705:A:C2	22:DA:727:A:O4'	2.70	0.44
32:BK:70:ARG:NH1	32:BK:74:GLY:O	2.50	0.44
22:DA:1773:A:H2'	22:DA:1774:C:H5'	2.00	0.44
22:DA:2063:C:O2	22:DA:2450:A:N1	2.51	0.44
1:AA:9:G:C6	1:AA:26:A:N6	2.86	0.44
22:BA:527:C:N3	22:BA:2779:U:H2'	2.32	0.44
50:D2:10:LEU:CD1	50:D2:14:ARG:NE	2.80	0.44
33:BL:77:ILE:CD1	33:BL:95:LEU:HD13	2.48	0.44
37:DP:73:VAL:HG23	37:DP:73:VAL:O	2.16	0.44
24:DC:212:ARG:CD	24:DC:218:PRO:HD3	2.47	0.44
1:CA:257:G:C5	57:CA:1718:HOH:O	2.69	0.44
31:DJ:5:THR:HB	31:DJ:7:LYS:NZ	2.32	0.44
10:AJ:33:GLY:O	10:AJ:34:ALA:HB2	2.17	0.44
22:DA:2351:G:H1'	22:DA:2367:G:N2	2.33	0.44
22:DA:1170:C:H2'	22:DA:1171:G:C8	2.52	0.44
26:DE:21:ARG:HD3	26:DE:106:LYS:HB3	2.00	0.44
1:CA:922:G:H4'	5:CE:25:VAL:HA	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:2849:U:O4	37:DP:21:ARG:NH2	2.49	0.44
27:DF:176:PRO:O	27:DF:177:PHE:HB2	2.18	0.44
9:AI:80:ARG:NH1	9:AI:103:PHE:CE1	2.84	0.44
7:AG:15:ASP:OD1	7:AG:16:PRO:HD2	2.18	0.44
29:DH:37:VAL:HG22	29:DH:38:PRO:HD2	1.98	0.44
25:DD:35:THR:O	25:DD:36:GLN:HB2	2.17	0.44
1:AA:11:G:C5	1:AA:12:U:C5	3.06	0.44
22:DA:948:C:O2	22:DA:984:A:O2'	2.34	0.44
1:AA:463:U:H5'	1:AA:464:U:OP2	2.17	0.44
22:DA:2556:C:H2'	22:DA:2557:G:O4'	2.17	0.44
8:CH:126:ILE:HG22	8:CH:127:CYS:N	2.33	0.44
22:DA:485:C:C2	22:DA:496:G:C2	3.05	0.44
1:CA:1103:C:H5''	2:CB:97:LEU:HD22	1.98	0.44
33:BL:53:GLY:O	33:BL:54:GLN:C	2.56	0.44
1:CA:926:G:C5	1:CA:1505:G:C6	3.05	0.44
22:DA:566:U:O2'	22:DA:809:G:OP2	2.28	0.44
6:AF:45:ARG:O	6:AF:56:LYS:HA	2.18	0.44
22:BA:26:G:H1'	22:BA:514:A:H61	1.82	0.44
22:BA:170:U:H2'	22:BA:171:U:C6	2.52	0.44
22:BA:1964:G:H4'	22:BA:1965:C:OP2	2.17	0.44
6:AF:85:ILE:O	6:AF:86:ARG:HG2	2.18	0.44
12:AL:85:GLY:O	12:AL:96:HIS:ND1	2.47	0.44
29:BH:57:LYS:CG	29:BH:58:LEU:N	2.81	0.44
22:BA:39:G:H2'	22:BA:40:U:C6	2.53	0.44
49:B1:10:LYS:O	49:B1:51:GLU:HG3	2.18	0.44
1:CA:1157:A:C6	1:CA:1180:A:C5	3.05	0.44
8:CH:106:THR:HG21	8:CH:121:LEU:HD13	1.99	0.44
28:DG:9:VAL:HG23	28:DG:69:ARG:CD	2.47	0.44
26:DE:79:ARG:O	26:DE:80:SER:HB2	2.18	0.44
27:BF:122:PHE:HB3	27:BF:163:ASP:OD2	2.18	0.44
3:CC:80:LYS:HE3	3:CC:80:LYS:HA	1.99	0.44
22:DA:2114:A:N3	22:DA:2114:A:H2'	2.32	0.44
25:BD:68:PHE:CE1	25:BD:75:ALA:HA	2.53	0.44
22:DA:1760:C:C6	22:DA:1761:C:C5	3.06	0.44
14:CN:21:PHE:O	14:CN:23:LYS:N	2.50	0.44
28:DG:83:PHE:CE2	28:DG:138:LYS:HB2	2.52	0.44
10:AJ:49:PHE:CD2	14:AN:77:PHE:CE2	3.05	0.44
13:CM:54:ASP:HA	13:CM:57:ARG:HB3	1.99	0.44
23:DB:68:C:H2'	23:DB:69:G:O4'	2.17	0.44
1:CA:355:C:H2'	1:CA:356:A:O4'	2.17	0.44
4:CD:104:ARG:HA	4:CD:104:ARG:HD2	1.84	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:CE:101:GLU:OE2	5:CE:103:THR:HA	2.17	0.44
5:CE:157:ARG:O	5:CE:159:LYS:HG2	2.17	0.44
29:DH:1:MET:CE	29:DH:27:ARG:NH1	2.80	0.44
22:DA:2821:A:OP2	25:DD:115:GLY:HA3	2.17	0.44
24:DC:204:VAL:O	24:DC:205:LEU:HB2	2.16	0.44
1:AA:975:A:H8	1:AA:1357:A:HO2'	1.61	0.44
22:DA:528:A:C2	22:DA:2043:C:C4'	2.99	0.44
1:CA:1170:A:H2'	1:CA:1171:A:O4'	2.17	0.44
22:DA:563:A:C2	22:DA:564:C:C2	3.05	0.44
22:BA:1050:A:C2	22:BA:2751:G:C5	3.05	0.44
22:DA:2134:A:C6	22:DA:2135:A:C5	3.06	0.44
1:CA:587:G:N2	1:CA:755:G:C4	2.85	0.44
1:CA:1413:A:C2	1:CA:1488:G:N2	2.86	0.44
15:CO:88:ARG:O	15:CO:89:ARG:CB	2.65	0.44
22:BA:2554:U:C4	22:BA:2555:U:C4	3.05	0.44
1:CA:806:C:O2'	1:CA:807:A:H5'	2.18	0.44
1:AA:113:G:H2'	1:AA:114:U:H6	1.83	0.44
33:DL:29:LYS:O	33:DL:30:THR:OG1	2.24	0.44
1:AA:1078:U:O4'	5:AE:89:HIS:HE1	2.00	0.44
5:AE:137:VAL:O	5:AE:138:ARG:CB	2.65	0.44
22:DA:2571:U:C4	22:DA:2574:G:C8	3.06	0.44
22:BA:1795:C:H2'	22:BA:1796:U:H6	1.81	0.44
23:DB:66:A:H61	23:DB:107:G:H2'	1.83	0.44
1:CA:1461:G:C5	1:CA:1462:C:C5	3.06	0.44
29:BH:31:VAL:N	29:BH:32:PRO:CD	2.80	0.44
22:BA:1373:A:C5	22:BA:1374:G:H1'	2.53	0.44
1:CA:243:A:H4'	1:CA:244:U:C5'	2.48	0.44
1:CA:496:A:H2'	1:CA:497:G:N7	2.32	0.44
30:DI:80:LEU:HD11	30:DI:133:ALA:HA	1.98	0.44
22:DA:305:C:O2	22:DA:313:G:C2	2.70	0.44
22:BA:2566:A:H4'	22:BA:2567:G:H5''	1.99	0.44
1:AA:184:G:C6	1:AA:185:U:C4	3.05	0.44
22:DA:1731:G:N1	22:DA:1733:G:C5	2.86	0.44
1:AA:417:G:C5	1:AA:418:C:C4	3.05	0.44
9:CI:30:ILE:HA	9:CI:65:ILE:HG13	1.99	0.44
11:AK:23:ILE:O	11:AK:23:ILE:HG13	2.18	0.44
42:DU:81:ASP:OD1	42:DU:96:PHE:HB3	2.17	0.44
22:DA:480:A:H5''	42:DU:44:LYS:HD2	1.98	0.44
29:DH:25:TYR:O	29:DH:29:PHE:HB3	2.18	0.44
1:CA:709:U:H2'	1:CA:710:G:H8	1.82	0.44
22:DA:2753:A:C6	22:DA:2754:U:N3	2.85	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:1123:U:O3'	10:CJ:38:GLY:HA3	2.17	0.44
22:BA:517:C:OP2	48:B0:10:ARG:NH2	2.50	0.44
12:CL:59:ASN:HD22	12:CL:59:ASN:N	2.16	0.44
1:CA:312:C:C5	1:CA:313:A:N7	2.86	0.44
22:BA:1360:G:O6	22:BA:1372:U:C2	2.71	0.44
1:AA:1353:G:C2	1:AA:1370:G:C2	3.05	0.44
35:BN:38:LEU:O	35:BN:38:LEU:HD12	2.18	0.44
22:BA:1903:G:O2'	22:BA:1904:G:H5'	2.16	0.44
25:DD:62:LYS:HB2	25:DD:63:PRO:HD3	1.99	0.44
24:DC:31:ALA:HB3	24:DC:32:PRO:HD3	2.00	0.44
7:CG:105:VAL:O	7:CG:109:ARG:HG2	2.17	0.44
27:DF:8:TYR:HB2	27:DF:173:PHE:CZ	2.53	0.44
22:DA:2561:U:H2'	22:DA:2562:U:O5'	2.17	0.44
26:DE:132:LYS:HA	26:DE:135:ALA:HB3	1.99	0.44
6:AF:40:GLU:HB2	6:AF:61:LEU:HB3	2.00	0.44
29:BH:62:LEU:O	29:BH:62:LEU:HD12	2.17	0.44
16:CP:11:ALA:O	16:CP:14:ARG:N	2.45	0.44
3:CC:101:ILE:HG23	3:CC:101:ILE:O	2.18	0.44
45:BX:35:SER:HA	45:BX:49:LEU:O	2.17	0.44
1:AA:147:G:N2	1:AA:176:C:C2	2.86	0.44
14:AN:83:LYS:HD3	14:AN:86:GLU:OE1	2.17	0.44
14:CN:3:LYS:HB3	14:CN:6:MET:HG2	2.00	0.44
29:BH:100:ALA:HB2	29:BH:115:VAL:HG21	1.98	0.44
22:DA:1154:G:P	38:DQ:58:ARG:HH11	2.40	0.44
22:BA:1068:G:H2'	22:BA:1069:A:H5'	2.00	0.44
22:DA:2843:G:N2	22:DA:2875:C:C2	2.85	0.44
22:DA:1790:C:H4'	24:DC:208:ALA:HB2	2.00	0.44
4:AD:170:TRP:O	4:AD:183:LYS:HB3	2.16	0.44
1:CA:62:U:H5''	1:CA:385:C:O2'	2.17	0.44
4:CD:9:LEU:HD12	4:CD:9:LEU:HA	1.74	0.44
1:AA:108:G:O6	20:AT:10:ARG:HG2	2.16	0.44
22:DA:1392:A:N6	22:DA:1393:A:N6	2.66	0.44
1:AA:1080:A:H5''	5:AE:21:VAL:HG11	1.99	0.44
22:DA:82:U:O2	22:DA:83:A:C8	2.71	0.44
1:CA:327:A:O2'	1:CA:328:C:O4'	2.29	0.44
22:DA:687:C:H5''	50:D2:2:LYS:HE2	1.99	0.44
23:DB:45:A:OP1	27:DF:92:ARG:HD3	2.18	0.44
21:AU:40:LYS:HG2	21:AU:41:PRO:HD3	2.00	0.44
22:DA:14:A:N6	22:DA:526:A:C2	2.85	0.44
22:DA:830:G:C2	22:DA:2448:A:N7	2.86	0.44
1:CA:273:U:C2'	1:CA:274:A:H5'	2.48	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:BO:2:ASP:OD1	36:BO:3:LYS:N	2.51	0.44
37:DP:39:ARG:HG3	37:DP:40:LEU:H	1.82	0.44
37:DP:39:ARG:HA	37:DP:39:ARG:NE	2.33	0.44
1:AA:1269:A:H2	1:AA:1312:G:N3	2.15	0.44
23:DB:29:A:OP2	36:DO:32:PRO:HD2	2.17	0.44
1:AA:1394:A:C5	1:AA:1501:C:H4'	2.53	0.44
22:DA:2347:C:O2'	49:D1:39:PHE:HB3	2.18	0.44
1:CA:1318:A:H4'	19:CS:10:PHE:CE2	2.53	0.44
30:DI:57:VAL:HG22	30:DI:58:VAL:N	2.33	0.44
22:DA:270:A:OP1	22:DA:271:G:H2'	2.18	0.44
2:AB:157:LEU:O	2:AB:158:PRO:C	2.52	0.44
39:DR:39:LEU:HA	39:DR:49:ILE:HG21	2.00	0.44
1:AA:1329:A:H5''	13:AM:26:GLY:N	2.33	0.44
22:DA:1869:G:C2	22:DA:1873:G:N1	2.86	0.44
22:BA:675:A:H4'	26:BE:62:GLN:OE1	2.18	0.44
1:AA:137:U:C2'	1:AA:137:U:O2	2.65	0.44
1:CA:1408:A:C2	1:CA:1494:G:C6	3.05	0.44
22:DA:2389:G:H5''	22:DA:2390:U:H5'	1.99	0.44
41:BT:33:LYS:HG3	41:BT:80:TRP:CE3	2.52	0.44
22:DA:1623:G:C5	22:DA:1624:U:C5	3.05	0.44
28:BG:52:PHE:CE1	28:BG:69:ARG:HA	2.53	0.44
11:CK:92:GLY:O	11:CK:93:ARG:HB3	2.17	0.44
14:CN:61:ARG:O	14:CN:62:ASN:HB2	2.16	0.44
22:DA:96:C:OP1	46:DY:41:HIS:CE1	2.70	0.44
1:CA:942:G:N2	1:CA:1342:C:C2	2.85	0.44
22:BA:2122:U:C4	22:BA:2123:G:N7	2.85	0.44
22:BA:1566:A:O2'	22:BA:1567:G:H5'	2.18	0.44
50:B2:21:ARG:HB2	50:B2:31:LEU:HD11	1.99	0.44
1:CA:1126:U:C2	1:CA:1281:C:C5	3.05	0.44
1:AA:591:U:H2'	1:AA:592:G:C8	2.53	0.44
48:D0:33:THR:HG22	48:D0:34:SER:N	2.32	0.44
3:AC:140:ASN:O	3:AC:140:ASN:ND2	2.48	0.44
24:BC:84:ASP:C	24:BC:84:ASP:OD1	2.56	0.44
9:AI:49:ARG:HD3	9:AI:49:ARG:C	2.37	0.44
4:CD:57:GLU:OE2	4:CD:196:ASN:N	2.45	0.44
22:DA:1299:G:H5'	22:DA:1301:A:O4'	2.17	0.44
22:DA:1215:G:H2'	22:DA:1216:G:O4'	2.17	0.44
5:AE:83:HIS:HB2	5:AE:84:PRO:HD2	1.99	0.44
1:AA:423:G:H2'	1:AA:424:G:O4'	2.18	0.44
1:AA:604:G:C2	1:AA:635:A:C2	3.06	0.44
22:DA:1132:U:H4'	31:DJ:75:TYR:CE2	2.52	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:BH:89:LYS:CE	29:BH:124:THR:HG22	2.48	0.44
22:BA:1923:U:O2'	22:BA:1924:C:H5'	2.17	0.44
17:AQ:16:LYS:N	17:AQ:17:MET:CE	2.81	0.44
1:CA:1144:G:N2	1:CA:1145:A:C2	2.86	0.44
22:DA:246:C:C2'	22:DA:247:G:H5'	2.47	0.44
22:DA:189:G:C5	22:DA:205:G:C2	3.05	0.44
31:BJ:80:HIS:O	31:BJ:81:ILE:C	2.56	0.44
31:BJ:77:HIS:HA	31:BJ:83:GLY:O	2.17	0.44
22:BA:2747:G:O2'	28:BG:67:THR:HB	2.17	0.44
28:BG:35:ARG:HD3	28:BG:71:LEU:HD13	2.00	0.44
24:DC:228:VAL:HG13	24:DC:229:ASP:N	2.33	0.44
24:DC:141:VAL:O	24:DC:162:VAL:N	2.47	0.44
35:BN:37:THR:HG22	35:BN:110:MET:CE	2.48	0.44
5:AE:81:LEU:HB3	5:AE:147:MET:CE	2.48	0.44
50:D2:10:LEU:HD11	50:D2:14:ARG:NH1	2.32	0.44
22:DA:89:A:N1	22:DA:90:U:N3	2.66	0.44
22:DA:1300:G:C6	22:DA:1626:A:O2'	2.66	0.44
22:DA:260:G:C6	22:DA:261:G:C8	3.06	0.44
1:CA:1458:G:O2'	20:CT:23:SER:HB3	2.18	0.44
1:AA:859:G:H2'	1:AA:860:A:C8	2.52	0.44
22:DA:307:G:N2	22:DA:310:A:C8	2.85	0.44
1:CA:1404:C:O2	1:CA:1519:A:O2'	2.34	0.44
1:AA:1059:C:N3	1:AA:1060:U:C5	2.86	0.44
1:CA:562:U:H4'	1:CA:563:A:O5'	2.18	0.44
22:DA:1178:C:H2'	22:DA:1179:G:C8	2.52	0.44
22:DA:269:C:C4	22:DA:270:A:N7	2.85	0.44
1:CA:682:G:N3	1:CA:683:G:C8	2.86	0.44
22:DA:126:A:C5	22:DA:127:A:C2	3.06	0.44
10:AJ:53:ILE:CG2	10:AJ:61:ALA:HB1	2.48	0.44
41:BT:17:SER:O	41:BT:18:GLU:C	2.56	0.44
2:CB:53:ALA:O	2:CB:57:LEU:HB2	2.18	0.44
1:CA:519:C:N4	1:CA:520:A:C6	2.86	0.44
27:BF:40:VAL:CG1	27:BF:50:LEU:HD13	2.47	0.44
26:BE:57:LYS:HG3	26:BE:58:LYS:N	2.30	0.44
41:BT:87:LEU:O	41:BT:88:LYS:C	2.55	0.44
1:CA:632:U:H2'	1:CA:633:G:OP1	2.18	0.44
22:DA:60:G:C4	22:DA:74:A:C2	3.06	0.44
1:CA:1066:C:H3'	1:CA:1067:A:C8	2.52	0.44
1:CA:1066:C:H3'	1:CA:1067:A:H8	1.83	0.44
46:DY:21:LEU:HA	46:DY:25:GLN:HB3	2.00	0.44
26:BE:79:ARG:O	26:BE:80:SER:HB2	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:AI:9:THR:O	9:AI:17:ALA:O	2.36	0.44
22:DA:2631:G:C6	22:DA:2632:A:N7	2.86	0.44
1:CA:144:G:C2	1:CA:145:G:C4	3.06	0.44
22:BA:1654:A:H1'	22:BA:2823:A:H5'	1.99	0.44
22:DA:1669:A:O4'	32:DK:5:GLN:HG3	2.17	0.44
22:DA:2663:G:H2'	22:DA:2664:G:O4'	2.17	0.44
1:CA:560:A:H4'	1:CA:561:U:H5''	1.99	0.44
9:CI:60:LYS:HD2	9:CI:61:LEU:HD22	2.00	0.44
18:CR:52:GLN:O	18:CR:52:GLN:HG3	2.17	0.44
29:BH:122:LEU:HD23	1:CA:368:U:OP2	2.03	0.44
22:DA:1356:G:C2	22:DA:1357:C:H1'	2.53	0.44
22:BA:1269:A:H2'	22:BA:1270:C:C6	2.53	0.44
1:CA:1095:U:H2'	1:CA:1096:C:C6	2.52	0.44
22:BA:1911:U:H2'	22:BA:1918:A:N1	2.32	0.44
39:BR:79:ARG:O	39:BR:80:ARG:HB2	2.18	0.44
22:DA:1853:A:H1'	22:DA:2234:G:H5'	2.00	0.44
50:D2:44:VAL:HG22	50:D2:45:SER:H	1.83	0.44
1:AA:922:G:C6	1:AA:923:A:C6	3.06	0.44
7:AG:69:VAL:HG12	7:AG:135:VAL:HA	1.99	0.44
22:DA:726:G:O2'	22:DA:727:A:OP2	2.31	0.44
22:DA:1767:G:C6	22:DA:1986:C:N4	2.86	0.44
22:DA:1671:U:P	57:DA:3434:HOH:O	2.73	0.44
1:CA:687:A:C2'	1:CA:701:U:O4	2.66	0.44
22:DA:1022:G:N2	22:DA:1142:A:C2	2.86	0.44
46:BY:6:LEU:O	46:BY:60:LYS:NZ	2.48	0.44
27:BF:28:VAL:O	27:BF:28:VAL:HG13	2.18	0.44
1:CA:1106:G:C5	1:CA:1107:C:C5	3.05	0.44
29:BH:97:ARG:O	29:BH:101:ASP:HB2	2.17	0.44
22:DA:749:A:C5	22:DA:750:A:C8	3.06	0.44
12:CL:38:TYR:HB2	12:CL:52:VAL:HG13	2.00	0.44
1:AA:15:G:C4	1:AA:16:A:C8	3.06	0.44
1:CA:728:A:C2	1:CA:729:A:C4	3.05	0.44
22:DA:1341:G:H5'	41:DT:61:LEU:HB3	1.99	0.44
1:CA:1416:G:H2'	1:CA:1417:G:O4'	2.18	0.44
22:DA:1831:G:C5	22:DA:1832:C:C4	3.06	0.44
40:BS:95:ARG:HG2	40:BS:96:ILE:N	2.32	0.44
22:DA:1906:G:OP1	22:DA:1930:G:N7	2.51	0.44
22:DA:136:G:N1	22:DA:144:A:N6	2.65	0.44
22:DA:2444:G:P	26:DE:63:LYS:HD2	2.57	0.44
1:AA:1313:U:OP2	19:AS:6:LYS:HB3	2.18	0.44
9:CI:13:LYS:O	9:CI:14:SER:HB3	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:545:U:O2	22:DA:545:U:O5'	2.36	0.44
10:AJ:52:LEU:HD21	10:AJ:59:LYS:HA	2.00	0.44
25:DD:125:TRP:CG	25:DD:160:LYS:HB3	2.53	0.44
24:BC:125:LYS:HB2	24:BC:126:PRO:HD2	1.99	0.44
16:AP:10:GLY:HA3	16:AP:15:PRO:HA	1.98	0.44
27:BF:108:VAL:N	27:BF:109:PRO:HD2	2.32	0.44
22:DA:1869:G:H3'	22:DA:1870:C:H5'	2.00	0.44
9:AI:84:THR:HG21	9:AI:103:PHE:CB	2.46	0.44
42:DU:34:VAL:O	42:DU:64:ALA:HA	2.18	0.44
1:AA:1094:G:O2'	1:AA:1095:U:OP2	2.36	0.44
1:AA:1095:U:P	57:AA:1865:HOH:O	2.75	0.44
52:B4:37:GLN:CG	52:B4:37:GLN:O	2.65	0.44
1:CA:1392:G:O2'	1:CA:1393:U:H5'	2.18	0.44
45:DX:68:LEU:HD22	45:DX:78:TYR:CZ	2.52	0.44
1:CA:1250:A:C6	1:CA:1251:A:C6	3.05	0.44
15:AO:87:LEU:O	15:AO:88:ARG:HB3	2.17	0.44
39:BR:83:TYR:CD2	39:BR:83:TYR:C	2.90	0.44
26:DE:146:VAL:HA	26:DE:185:LYS:O	2.17	0.44
22:DA:2452:C:C4	22:DA:2453:A:C6	3.06	0.44
23:BB:7:G:C5'	36:BO:29:HIS:CD2	3.01	0.44
1:AA:1435:G:H2'	1:AA:1436:U:C6	2.53	0.44
1:AA:104:G:C2	1:AA:105:G:C8	3.06	0.44
29:BH:57:LYS:HG3	29:BH:58:LEU:N	2.33	0.44
1:AA:186:C:O4'	20:AT:76:LYS:HD2	2.17	0.44
1:CA:28:A:OP1	4:CD:73:ARG:NH2	2.51	0.44
38:BQ:49:ASP:HA	38:BQ:52:GLN:HB2	1.99	0.44
8:CH:30:SER:O	8:CH:34:VAL:HG23	2.17	0.44
2:AB:35:ARG:O	2:AB:38:VAL:HG13	2.18	0.44
22:BA:2209:G:C6	22:BA:2210:U:C4	3.05	0.44
24:BC:166:ALA:HB3	24:BC:173:THR:HB	2.00	0.44
1:CA:490:C:H2'	1:CA:491:G:C8	2.53	0.44
22:BA:2639:A:C2	22:BA:2778:A:C8	3.06	0.44
36:DO:70:ALA:O	36:DO:74:VAL:HB	2.17	0.44
22:BA:2419:U:O2'	22:BA:2420:C:H5'	2.17	0.44
24:BC:7:LYS:HB3	24:BC:8:PRO:HD2	2.00	0.44
22:BA:2267:A:H5''	22:BA:2268:A:H5'	1.99	0.44
3:CC:43:LEU:HD21	3:CC:68:ILE:HD11	1.99	0.44
22:DA:1290:C:N3	22:DA:1291:C:C5	2.86	0.44
6:AF:13:ASP:OD1	6:AF:13:ASP:N	2.50	0.44
1:AA:992:U:C2	1:AA:1043:G:N7	2.86	0.44
41:DT:29:THR:OG1	41:DT:86:THR:HG22	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:875:G:N2	22:BA:903:C:C2	2.85	0.44
1:CA:1229:A:C2	1:CA:1230:C:C2	3.05	0.44
31:BJ:11:VAL:HG21	31:BJ:50:THR:HG22	2.00	0.44
29:BH:99:ILE:CD1	29:BH:117:LEU:HD13	2.48	0.44
1:CA:1504:G:OP2	1:CA:1507:A:O2'	2.34	0.44
1:AA:1343:G:C5	1:AA:1344:C:C4	3.05	0.44
31:DJ:41:LYS:NZ	31:DJ:52:ASP:OD1	2.51	0.44
24:DC:141:VAL:CG1	24:DC:190:ALA:HB1	2.44	0.44
22:BA:423:A:H5''	22:BA:424:G:H5'	2.00	0.44
4:AD:23:SER:O	4:AD:24:GLY:C	2.55	0.44
22:DA:89:A:N1	22:DA:90:U:C4	2.85	0.44
22:DA:1363:C:H2'	22:DA:1364:G:H8	1.83	0.44
24:DC:66:ASP:OD2	24:DC:102:ARG:HD3	2.17	0.44
24:DC:43:ARG:HG2	24:DC:48:ARG:O	2.18	0.44
22:BA:831:G:C6	22:BA:832:U:C4	3.06	0.44
22:DA:575:A:N3	22:DA:576:U:C6	2.86	0.44
29:BH:4:ILE:HG23	29:BH:17:ASP:O	2.17	0.44
5:AE:72:ILE:CD1	5:AE:145:GLU:OE1	2.65	0.44
2:AB:68:LEU:HD22	2:AB:70:VAL:HG23	2.00	0.44
22:BA:1584:U:C2'	22:BA:1584:U:O2	2.66	0.44
22:BA:1796:U:C2'	22:BA:1796:U:O2	2.64	0.44
45:BX:22:LEU:HD23	45:BX:22:LEU:HA	1.73	0.44
5:CE:42:GLY:O	5:CE:119:GLY:HA3	2.17	0.44
10:AJ:65:TYR:HB3	14:AN:96:LEU:HD11	1.99	0.44
1:CA:1237:C:H3'	1:CA:1238:A:H5'	1.99	0.44
1:CA:610:U:O4	1:CA:611:C:N4	2.51	0.44
10:AJ:53:ILE:HG22	10:AJ:61:ALA:HB1	1.99	0.44
6:CF:99:ALA:O	6:CF:100:SER:HB3	2.18	0.44
1:AA:1159:U:N3	1:AA:1182:G:C5	2.85	0.44
22:DA:245:G:O6	51:D3:8:ARG:HD3	2.18	0.44
1:CA:769:G:H4'	1:CA:1513:A:H4'	1.98	0.44
22:DA:222:A:H3'	22:DA:421:C:C5'	2.48	0.44
42:DU:13:VAL:HG21	42:DU:39:ILE:HG21	2.00	0.44
22:DA:1465:G:C5	22:DA:1466:U:C4	3.06	0.44
35:DN:98:LEU:HD13	48:D0:54:VAL:HG21	2.00	0.44
1:CA:45:G:H5''	1:CA:307:C:O2'	2.18	0.44
1:AA:1409:C:H2'	1:AA:1410:A:C8	2.53	0.44
46:BY:32:ALA:HB2	46:BY:37:LEU:HD23	1.99	0.44
1:CA:1370:G:O2'	1:CA:1371:G:H5'	2.17	0.44
22:DA:2453:A:N7	57:DA:3527:HOH:O	2.35	0.44
22:DA:2607:G:H2'	22:DA:2608:G:O4'	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:1802:A:N1	22:BA:1822:C:H1'	2.33	0.44
5:CE:53:ALA:HB2	5:CE:62:LYS:NZ	2.33	0.44
22:BA:900:A:C2	22:BA:901:C:H1'	2.53	0.44
22:BA:544:C:H2'	22:BA:545:U:O4'	2.18	0.44
22:DA:1249:U:H4'	38:DQ:4:VAL:HB	2.00	0.44
41:BT:44:LYS:O	41:BT:48:GLN:HG3	2.18	0.44
1:CA:949:A:H2'	1:CA:950:U:O4'	2.18	0.44
42:DU:45:HIS:HB3	42:DU:58:ILE:HG12	2.00	0.44
1:CA:230:G:H2'	1:CA:231:U:O4'	2.18	0.44
1:AA:636:U:H5''	17:AQ:6:ARG:HG2	1.99	0.44
29:BH:121:VAL:HA	29:BH:128:HIS:CG	2.53	0.44
29:BH:89:LYS:HE3	29:BH:124:THR:HG22	1.99	0.44
22:DA:996:A:C5	22:DA:1160:G:C2	3.06	0.44
39:BR:39:LEU:HA	39:BR:49:ILE:HG23	2.00	0.44
1:CA:1160:G:O2'	1:CA:1161:C:P	2.76	0.44
2:AB:82:ASP:OD1	2:AB:82:ASP:C	2.56	0.44
1:AA:920:U:H2'	1:AA:921:U:C6	2.53	0.44
24:BC:246:THR:HB	24:BC:247:PRO:HD2	2.00	0.44
5:CE:105:ILE:HD11	5:CE:112:ARG:HA	2.00	0.44
25:BD:104:VAL:HG23	25:BD:177:VAL:HG11	2.00	0.44
24:DC:162:VAL:HG13	24:DC:175:ARG:O	2.18	0.44
29:DH:31:VAL:CB	29:DH:32:PRO:HD3	2.47	0.44
22:BA:2846:G:OP1	37:BP:53:ARG:NH1	2.51	0.44
22:BA:877:A:N6	22:BA:899:A:C6	2.84	0.44
22:DA:563:A:C5	22:DA:2018:G:C2	3.06	0.44
23:DB:43:C:N4	23:DB:45:A:N1	2.65	0.44
40:DS:59:GLU:HG2	40:DS:59:GLU:O	2.18	0.44
2:AB:87:CYS:HB2	2:AB:89:GLN:CD	2.38	0.44
1:CA:757:U:OP1	1:CA:822:U:O2'	2.31	0.44
53:B5:50:ILE:O	53:B5:52:PRO:HD3	2.16	0.44
22:DA:1905:C:O4'	22:DA:1928:A:H2	1.99	0.44
22:BA:1795:C:H2'	22:BA:1796:U:C6	2.53	0.44
2:AB:162:PHE:HA	2:AB:184:PHE:O	2.18	0.44
23:DB:66:A:H4'	23:DB:67:G:OP1	2.18	0.44
5:AE:15:LEU:O	5:AE:15:LEU:CD1	2.65	0.44
2:CB:81:LYS:HG3	2:CB:91:PHE:CZ	2.53	0.44
22:DA:2297:A:C8	22:DA:2320:U:C4	3.05	0.44
26:BE:18:THR:O	26:BE:18:THR:HG22	2.18	0.44
20:CT:43:ASP:CB	20:CT:46:ALA:HB3	2.48	0.44
21:AU:14:VAL:HG13	21:AU:16:LEU:CG	2.48	0.44
22:BA:1002:G:C5	57:BA:3746:HOH:O	2.69	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:856:G:C2	22:DA:922:C:C2	3.06	0.44
1:AA:76:G:H2'	1:AA:76:G:N3	2.32	0.44
4:AD:122:ALA:O	4:AD:123:ILE:HG22	2.18	0.44
22:DA:2556:C:C4	22:DA:2557:G:C5	3.06	0.44
4:AD:138:SER:N	4:AD:141:ASP:OD2	2.50	0.44
1:AA:1516:G:N2	1:AA:1519:A:OP2	2.51	0.44
22:BA:958:U:C2	23:BB:89:U:H1'	2.53	0.44
11:AK:35:THR:HG1	11:AK:40:ASN:C	2.21	0.44
22:DA:43:G:C2	22:DA:437:U:N3	2.86	0.44
42:BU:102:THR:CG2	42:BU:103:ILE:N	2.81	0.44
22:DA:1862:G:C2	22:DA:1881:C:O2	2.71	0.44
22:BA:1644:C:C2'	22:BA:1645:G:H5'	2.48	0.44
11:CK:107:ILE:HD13	11:CK:107:ILE:C	2.38	0.44
1:CA:121:U:OP2	1:CA:121:U:H4'	2.17	0.44
22:DA:60:G:C5	22:DA:62:U:C4	3.06	0.44
22:DA:875:G:C2	22:DA:876:C:O2	2.70	0.44
1:AA:1191:A:C6	1:AA:1192:C:N4	2.86	0.44
10:AJ:42:LEU:HA	10:AJ:43:PRO:HD2	1.83	0.44
1:AA:6:G:O2'	1:AA:7:A:H5''	2.18	0.44
22:BA:1438:U:C5	22:BA:1552:A:C2	3.06	0.44
45:DX:5:CYS:O	45:DX:9:GLY:N	2.49	0.44
4:CD:139:PRO:O	4:CD:140:ASN:HB2	2.18	0.44
22:DA:607:U:O4	22:DA:619:G:H2'	2.18	0.44
38:DQ:17:ILE:HG23	38:DQ:39:VAL:HG21	2.00	0.44
25:BD:125:TRP:CE3	25:BD:160:LYS:HE2	2.53	0.44
10:AJ:71:LEU:O	10:AJ:72:ARG:NE	2.51	0.44
22:BA:1664:A:H1'	22:BA:2726:A:N1	2.33	0.44
22:BA:483:A:H2'	22:BA:484:C:H5'	1.99	0.44
36:DO:89:ASP:O	36:DO:90:VAL:HG13	2.18	0.44
22:DA:2591:C:OP1	24:DC:238:ARG:NH1	2.51	0.44
1:AA:75:G:N3	1:AA:75:G:H2'	2.33	0.44
11:AK:128:ARG:HH11	11:AK:128:ARG:HG2	1.83	0.44
1:AA:628:G:H2'	1:AA:629:A:C8	2.53	0.44
1:AA:815:A:H4'	1:AA:817:C:C4	2.53	0.44
1:AA:1406:U:C5	1:AA:1407:C:C4	3.06	0.43
28:BG:104:ASN:ND2	28:BG:114:ASP:CG	2.72	0.43
22:BA:273:G:O6	22:BA:364:C:N4	2.42	0.43
22:DA:1316:U:C2	22:DA:1337:G:C2	3.05	0.43
22:DA:189:G:H2'	22:DA:190:A:O5'	2.18	0.43
22:DA:2271:G:H5'	44:DW:20:ARG:HG3	2.00	0.43
7:AG:99:LEU:O	7:AG:100:ALA:C	2.57	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:CP:6:LEU:N	16:CP:6:LEU:HD12	2.33	0.43
22:BA:10:A:C5	22:BA:11:C:C5	3.06	0.43
24:DC:155:ALA:HB2	24:DC:162:VAL:HG23	2.01	0.43
22:DA:2009:A:N6	57:DA:3375:HOH:O	2.30	0.43
22:DA:696:G:C6	22:DA:767:U:C2	3.06	0.43
16:AP:50:THR:O	16:AP:50:THR:CG2	2.64	0.43
1:CA:642:A:N7	8:CH:107:SER:HA	2.33	0.43
12:CL:30:LYS:O	12:CL:81:LEU:HD12	2.17	0.43
53:B5:204:GLY:O	53:B5:205:ALA:CB	2.66	0.43
22:DA:2204:G:C6	22:DA:2221:G:C2	3.06	0.43
1:CA:502:A:O2'	1:CA:503:C:H5'	2.18	0.43
1:AA:1197:A:P	1:AA:1197:A:H3'	2.58	0.43
1:CA:158:G:C5	1:CA:164:G:C6	3.06	0.43
27:BF:41:GLY:C	27:BF:43:ALA:N	2.71	0.43
1:AA:1350:A:C6	1:AA:1351:U:C4	3.06	0.43
29:DH:127:GLU:CG	29:DH:144:VAL:O	2.65	0.43
1:CA:202:G:C2	1:CA:216:U:O2	2.71	0.43
1:CA:1514:G:H2'	1:CA:1515:G:C8	2.52	0.43
22:BA:1001:A:P	57:BA:3743:HOH:O	2.75	0.43
22:DA:2037:A:C6	22:DA:2038:G:C5	3.05	0.43
12:AL:47:SER:O	12:AL:48:ALA:HB2	2.18	0.43
22:BA:962:G:N2	22:BA:2250:G:H1	2.16	0.43
22:BA:2579:C:C2'	22:BA:2580:U:H5'	2.48	0.43
1:AA:865:A:O2'	1:AA:866:C:H5'	2.18	0.43
27:BF:57:LEU:HD21	27:BF:152:LEU:CD1	2.48	0.43
35:DN:114:GLU:OE2	35:DN:118:ARG:CD	2.66	0.43
3:AC:15:VAL:HG11	3:AC:179:ARG:O	2.18	0.43
30:BI:29:GLY:O	30:BI:35:ILE:HD11	2.18	0.43
37:DP:26:VAL:HG12	37:DP:28:VAL:HG23	1.99	0.43
27:BF:129:SER:HA	27:BF:154:ILE:O	2.18	0.43
37:BP:114:LEU:O	37:BP:115:ASN:HB3	2.17	0.43
9:AI:29:VAL:O	9:AI:65:ILE:HG12	2.18	0.43
22:BA:90:U:H2'	22:BA:91:A:C8	2.53	0.43
22:BA:563:A:C2	22:BA:564:C:C2	3.06	0.43
24:DC:147:LYS:HB2	24:DC:150:LYS:HB2	2.00	0.43
22:DA:273:G:N2	22:DA:365:U:O2	2.51	0.43
22:BA:1768:C:C2	22:BA:1769:U:C6	3.06	0.43
39:DR:48:LYS:HG2	39:DR:48:LYS:O	2.17	0.43
4:AD:2:ALA:O	4:AD:68:LEU:HD23	2.18	0.43
36:BO:115:LEU:HA	36:BO:115:LEU:HD12	1.89	0.43
40:DS:22:ASP:OD1	40:DS:22:ASP:N	2.51	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:DT:32:LEU:O	41:DT:32:LEU:HD12	2.18	0.43
22:BA:1429:G:C5	22:BA:1568:G:C6	3.06	0.43
39:DR:21:ARG:HG3	39:DR:93:PHE:HD2	1.82	0.43
7:AG:62:PHE:CE2	7:AG:66:LEU:HD22	2.53	0.43
29:BH:96:THR:O	29:BH:100:ALA:N	2.50	0.43
1:CA:356:A:C5	1:CA:357:G:C8	3.06	0.43
6:CF:11:HIS:ND1	6:CF:12:PRO:HD2	2.33	0.43
22:DA:1789:A:C5'	24:DC:219:THR:O	2.66	0.43
1:AA:553:A:C2'	1:AA:554:A:H5'	2.47	0.43
1:AA:452:A:N7	1:AA:453:G:N9	2.67	0.43
1:CA:374:A:C5	1:CA:375:U:C5	3.06	0.43
11:AK:126:LYS:C	21:AU:34:ARG:NH2	2.71	0.43
1:CA:1349:A:H1'	1:CA:1374:A:N6	2.33	0.43
22:DA:24:G:C6	22:DA:25:U:C4	3.06	0.43
27:BF:61:SER:O	27:BF:61:SER:OG	2.35	0.43
22:DA:1650:A:C2	22:DA:1651:G:C4	3.06	0.43
2:CB:20:THR:OG1	2:CB:21:ARG:N	2.51	0.43
22:DA:1806:C:O2'	24:DC:48:ARG:HG3	2.18	0.43
2:AB:88:ASP:HB2	2:AB:221:VAL:HG12	1.99	0.43
1:CA:729:A:H2'	1:CA:730:G:H8	1.82	0.43
26:DE:52:VAL:CG2	26:DE:81:GLY:HA2	2.45	0.43
22:BA:2516:A:C2'	22:BA:2517:C:H5'	2.49	0.43
20:CT:79:LEU:O	20:CT:83:ILE:HG23	2.17	0.43
22:DA:1131:G:N7	22:DA:2025:C:H4'	2.33	0.43
22:BA:1358:G:N2	22:BA:1374:G:C6	2.86	0.43
22:DA:976:G:H4'	22:DA:1156:A:N7	2.33	0.43
22:BA:1607:C:N4	22:BA:1622:G:C5	2.85	0.43
2:AB:144:LEU:O	2:AB:148:LEU:HB2	2.19	0.43
32:DK:38:ILE:CD1	32:DK:61:VAL:HB	2.48	0.43
33:BL:48:ARG:HG2	51:B3:60:ALA:HB1	1.99	0.43
20:AT:83:ILE:HD12	20:AT:84:ASN:N	2.33	0.43
22:DA:2267:A:H5''	22:DA:2268:A:C5'	2.48	0.43
22:DA:478:A:C6	22:DA:480:A:C6	3.06	0.43
16:AP:22:ALA:CB	16:AP:32:PHE:HA	2.48	0.43
35:DN:98:LEU:CD1	48:D0:54:VAL:HG21	2.48	0.43
4:CD:157:ALA:O	4:CD:161:LEU:HD22	2.17	0.43
34:BM:6:ARG:C	34:BM:7:THR:HG23	2.39	0.43
1:AA:802:A:H5''	1:AA:803:G:OP2	2.17	0.43
17:CQ:24:ALA:HB1	17:CQ:41:THR:HG23	2.00	0.43
1:AA:236:A:H2'	1:AA:237:G:C8	2.53	0.43
34:BM:19:GLY:O	34:BM:97:GLN:HG2	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:BL:36:LYS:O	33:BL:40:SER:HB3	2.18	0.43
1:CA:1039:G:H2'	1:CA:1040:U:O4'	2.19	0.43
26:BE:79:ARG:O	26:BE:80:SER:CB	2.65	0.43
39:DR:61:ALA:HB2	39:DR:98:ILE:HD13	2.01	0.43
22:BA:1356:G:C6	22:BA:1357:C:C4	3.06	0.43
7:AG:133:THR:O	7:AG:136:LYS:HB3	2.18	0.43
1:AA:633:G:OP2	8:AH:88:ARG:NH2	2.50	0.43
38:DQ:36:PHE:CE2	38:DQ:40:ILE:CD1	3.01	0.43
22:BA:769:U:C2	22:BA:770:G:C8	3.07	0.43
3:AC:66:VAL:HG12	3:AC:66:VAL:O	2.16	0.43
19:AS:79:THR:O	19:AS:79:THR:OG1	2.30	0.43
24:DC:161:TYR:C	24:DC:161:TYR:CD1	2.92	0.43
1:CA:754:C:O2	1:CA:754:C:H3'	2.18	0.43
1:AA:250:A:H4'	1:AA:251:G:O5'	2.17	0.43
22:DA:2141:G:N2	22:DA:2151:U:O2	2.51	0.43
41:BT:13:ALA:HB1	46:BY:33:ALA:HB1	1.99	0.43
22:BA:167:A:H2'	22:BA:168:G:O4'	2.18	0.43
22:DA:1355:G:C2'	22:DA:1356:G:H5'	2.47	0.43
1:CA:1124:G:N2	1:CA:1127:G:N2	2.66	0.43
18:CR:20:GLU:C	18:CR:22:ASP:N	2.71	0.43
5:AE:25:VAL:N	5:AE:28:GLY:O	2.46	0.43
22:BA:2756:U:H1'	22:BA:2757:A:H5''	2.00	0.43
4:CD:9:LEU:HD11	4:CD:29:ASP:OD1	2.18	0.43
22:BA:1142:A:C4	22:BA:1144:A:C8	3.06	0.43
22:DA:995:C:C4	38:DQ:57:PHE:CZ	3.06	0.43
12:CL:116:LYS:O	12:CL:117:TYR:CD2	2.72	0.43
42:BU:72:ILE:H	42:BU:72:ILE:CD1	2.31	0.43
22:BA:981:A:H5'	57:BA:3600:HOH:O	2.17	0.43
22:DA:1265:A:N1	22:DA:2013:A:H5''	2.32	0.43
41:DT:62:VAL:CG1	41:DT:63:VAL:N	2.81	0.43
1:CA:256:U:H2'	1:CA:257:G:C8	2.54	0.43
1:CA:761:G:O2'	1:CA:762:U:H5'	2.19	0.43
1:CA:71:A:C2	1:CA:72:A:C8	3.07	0.43
1:AA:1058:G:C6	1:AA:1059:C:C4	3.07	0.43
22:DA:2290:G:C6	22:DA:2291:U:C4	3.06	0.43
1:CA:1385:G:H2'	1:CA:1386:G:O4'	2.18	0.43
1:AA:1394:A:N1	1:AA:1500:A:O2'	2.46	0.43
1:CA:1133:G:C2	1:CA:1142:G:C2	3.07	0.43
5:AE:15:LEU:CB	5:AE:37:THR:HG22	2.48	0.43
16:CP:5:ARG:O	16:CP:19:VAL:HG23	2.18	0.43
22:DA:1275:A:H1'	22:DA:1276:A:O4'	2.17	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:AU:4:ILE:HA	21:AU:20:LYS:HE3	1.98	0.43
22:DA:126:A:O5'	50:D2:19:ARG:HG3	2.17	0.43
5:CE:26:LYS:HA	5:CE:26:LYS:HE2	1.98	0.43
1:CA:1227:A:H3'	1:CA:1227:A:N3	2.32	0.43
2:CB:188:ASP:OD2	2:CB:204:ASP:OD1	2.35	0.43
9:CI:19:VAL:HG22	9:CI:65:ILE:HG22	2.00	0.43
1:CA:215:C:H2'	1:CA:216:U:O4'	2.18	0.43
22:BA:613:A:O2'	22:BA:614:A:P	2.77	0.43
22:DA:965:C:H4'	22:DA:2273:A:H1'	2.00	0.43
22:DA:982:C:H5'	22:DA:983:A:P	2.58	0.43
22:DA:1464:G:C2	22:DA:1465:G:C4	3.06	0.43
12:AL:5:ASN:O	12:AL:6:GLN:C	2.56	0.43
1:CA:251:G:C6	1:CA:266:G:O6	2.71	0.43
1:AA:1409:C:H2'	1:AA:1410:A:H8	1.83	0.43
1:CA:295:C:C4	1:CA:296:U:C4	3.06	0.43
34:BM:36:VAL:HG22	43:BV:82:TYR:CD2	2.52	0.43
22:DA:1599:U:C4	22:DA:1600:C:N4	2.86	0.43
22:DA:1326:U:O4	22:DA:1647:U:O2	2.36	0.43
22:DA:852:U:H2'	22:DA:853:C:O4'	2.18	0.43
10:CJ:91:ASP:OD1	10:CJ:92:LEU:N	2.51	0.43
1:CA:1076:U:C2	1:CA:1082:A:C2	3.07	0.43
22:BA:2267:A:OP2	57:BA:3516:HOH:O	2.21	0.43
23:DB:46:A:C5	23:DB:47:C:C5	3.05	0.43
22:BA:1265:A:P	57:BA:3754:HOH:O	2.75	0.43
1:CA:584:G:H2'	1:CA:585:G:C8	2.54	0.43
22:BA:2849:U:N3	22:BA:2867:G:O4'	2.48	0.43
20:AT:55:GLN:N	20:AT:56:PRO:HD2	2.33	0.43
22:DA:1532:A:C2	22:DA:1540:G:C6	3.06	0.43
22:DA:459:U:C5	22:DA:469:G:N2	2.87	0.43
49:B1:4:GLY:O	49:B1:5:ILE:HB	2.18	0.43
1:AA:1128:C:O2'	1:AA:1147:C:N3	2.48	0.43
30:BI:90:SER:HB3	30:BI:93:PRO:HG3	1.99	0.43
4:AD:102:VAL:HG12	4:AD:114:ALA:HB1	1.99	0.43
2:CB:181:ILE:HD13	2:CB:181:ILE:N	2.33	0.43
1:AA:19:A:C2	1:AA:917:G:C5	3.07	0.43
22:DA:491:G:C6	22:DA:492:A:C6	3.06	0.43
38:DQ:94:ILE:HG22	38:DQ:95:LEU:N	2.33	0.43
29:BH:80:ILE:HG21	29:BH:94:ILE:HG13	2.00	0.43
22:BA:1073:A:N7	22:BA:1074:G:C8	2.86	0.43
22:DA:2125:G:N2	22:DA:2171:A:O5'	2.51	0.43
5:CE:102:GLY:C	5:CE:104:GLY:N	2.71	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:CE:105:ILE:H	5:CE:122:ASN:CA	2.31	0.43
22:DA:2823:A:C6	22:DA:2824:C:C4	3.06	0.43
1:CA:17:U:N3	1:CA:18:C:C4	2.86	0.43
22:DA:1969:A:H1'	22:DA:1973:G:H1'	1.99	0.43
22:BA:1378:A:O2'	22:BA:1380:G:OP2	2.36	0.43
1:CA:1105:A:C2	1:CA:1106:G:C8	3.06	0.43
22:DA:1742:U:O4	22:DA:1743:G:C6	2.71	0.43
22:DA:1779:U:H5	22:DA:1784:A:N7	2.17	0.43
22:DA:362:A:C5	22:DA:363:G:N7	2.86	0.43
22:DA:311:A:C6	22:DA:328:U:C4	3.06	0.43
29:BH:27:ARG:O	29:BH:28:ASN:CB	2.66	0.43
22:BA:832:U:H2'	22:BA:833:A:C8	2.54	0.43
1:CA:735:C:H5'	18:CR:60:LYS:HD3	1.99	0.43
22:BA:747:U:C4	22:BA:2613:U:C5	3.06	0.43
22:DA:387:U:O2	22:DA:388:G:N7	2.52	0.43
22:DA:1429:G:N3	22:DA:1568:G:C2	2.86	0.43
22:BA:1588:G:N1	22:BA:1589:U:C4	2.87	0.43
25:BD:133:THR:HG23	25:BD:134:HIS:CD2	2.52	0.43
24:DC:35:GLU:O	24:DC:35:GLU:HG3	2.19	0.43
2:AB:154:MET:O	2:AB:156:GLY:N	2.51	0.43
22:BA:225:C:H2'	22:BA:226:A:O4'	2.19	0.43
22:DA:1477:A:N6	22:DA:1514:G:O2'	2.51	0.43
21:AU:8:GLU:HB3	21:AU:12:PHE:CZ	2.53	0.43
22:DA:2531:A:C4	22:DA:2532:G:C8	3.06	0.43
22:DA:408:G:C6	22:DA:409:G:C5	3.05	0.43
1:CA:1256:A:N1	1:CA:1277:C:C4	2.87	0.43
47:BZ:6:LYS:HA	47:BZ:36:VAL:O	2.19	0.43
9:AI:67:VAL:HG11	9:AI:79:ILE:HD11	2.00	0.43
1:AA:903:G:C4	1:AA:904:U:C6	3.06	0.43
39:DR:78:ARG:HB3	39:DR:83:TYR:HD2	1.83	0.43
1:AA:619:U:N3	4:AD:131:ASN:OD1	2.44	0.43
22:DA:2753:A:N1	22:DA:2754:U:C2	2.86	0.43
4:AD:44:ARG:O	4:AD:46:PRO:HD3	2.19	0.43
16:CP:39:PHE:O	16:CP:41:PRO:HD3	2.18	0.43
22:DA:1435:G:C2'	22:DA:1436:G:H5'	2.48	0.43
22:DA:1290:C:C4	22:DA:1291:C:C5	3.06	0.43
5:CE:58:ALA:O	5:CE:62:LYS:HB2	2.19	0.43
16:CP:2:VAL:HG23	16:CP:65:ALA:HA	2.00	0.43
1:AA:855:U:H2'	1:AA:856:C:C6	2.54	0.43
3:CC:10:ILE:HD12	14:CN:98:LYS:HG3	1.99	0.43
11:CK:89:PRO:HD3	21:CU:29:LEU:HD11	2.01	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:2740:A:C6	22:DA:2764:A:C8	3.06	0.43
22:DA:2301:C:C2	22:DA:2316:G:N2	2.87	0.43
20:AT:79:LEU:O	20:AT:82:GLN:HB2	2.18	0.43
22:BA:914:G:C8	22:BA:914:G:H3'	2.53	0.43
25:DD:39:ASP:CG	25:DD:40:LEU:N	2.71	0.43
22:BA:918:A:H4'	23:BB:97:C:O2	2.18	0.43
22:BA:1334:G:C6	22:BA:1335:C:C4	3.06	0.43
15:AO:64:ARG:NH2	15:AO:68:ASP:OD1	2.51	0.43
22:BA:2094:A:H4'	29:BH:25:TYR:CZ	2.54	0.43
2:AB:94:HIS:O	2:AB:95:ARG:C	2.57	0.43
53:B5:45:HIS:CD2	53:B5:176:VAL:HA	2.53	0.43
1:AA:1458:G:OP1	20:AT:30:THR:OG1	2.34	0.43
29:BH:76:GLU:HA	29:BH:142:VAL:HG12	2.00	0.43
22:DA:1936:A:H3'	22:DA:1937:A:H5'	2.00	0.43
22:DA:1358:G:H1'	22:DA:1374:G:N2	2.33	0.43
22:BA:1059:G:H5''	22:BA:1060:U:C2'	2.48	0.43
1:AA:254:G:H4'	17:AQ:20:SER:HB2	2.00	0.43
22:DA:247:G:C4'	22:DA:386:G:C5	2.94	0.43
45:DX:32:ASN:ND2	45:DX:53:ALA:HB2	2.34	0.43
22:DA:176:A:C5	22:DA:177:G:C6	3.06	0.43
16:CP:71:VAL:O	16:CP:74:LEU:N	2.52	0.43
22:BA:480:A:C2'	22:BA:481:G:OP1	2.66	0.43
2:AB:204:ASP:OD1	2:AB:205:ASP:OD1	2.35	0.43
22:BA:996:A:C2	22:BA:997:G:C8	3.07	0.43
46:BY:17:GLU:HB2	46:BY:53:VAL:HG11	2.01	0.43
22:DA:2064:C:H1'	22:DA:2450:A:C6	2.53	0.43
22:BA:245:G:N7	51:B3:8:ARG:NH1	2.65	0.43
1:CA:207:C:C2'	1:CA:207:C:O2	2.65	0.43
5:AE:104:GLY:HA3	5:AE:122:ASN:HA	1.99	0.43
50:D2:10:LEU:O	50:D2:14:ARG:HG3	2.17	0.43
22:DA:684:G:C5'	50:D2:16:HIS:CE1	3.00	0.43
22:DA:599:A:C2	22:DA:659:G:C5	3.06	0.43
38:BQ:112:LYS:O	38:BQ:115:ALA:HB3	2.18	0.43
22:DA:1648:U:H2'	22:DA:1649:G:O4'	2.18	0.43
2:CB:94:HIS:HA	2:CB:95:ARG:NH2	2.34	0.43
22:DA:2545:G:N3	22:DA:2565:A:H2	2.16	0.43
22:BA:2517:C:C5	22:BA:2542:A:C5	3.06	0.43
11:AK:112:ASP:OD2	11:AK:114:THR:HG23	2.17	0.43
22:BA:1071:G:P	22:BA:1071:G:H8	2.42	0.43
22:BA:1056:G:C2	22:BA:1102:C:C5	3.07	0.43
8:AH:7:ILE:O	8:AH:11:LEU:HG	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:AH:2:SER:N	8:AH:4:GLN:HE21	2.17	0.43
14:AN:54:ASP:HA	14:AN:59:ARG:HD3	2.00	0.43
22:DA:2360:G:H1'	33:DL:60:ARG:HD3	2.00	0.43
10:CJ:89:ARG:O	10:CJ:90:LEU:CB	2.66	0.43
22:DA:2818:U:H2'	22:DA:2819:G:C8	2.54	0.43
29:BH:9:VAL:O	29:BH:10:ALA:O	2.36	0.43
32:DK:91:SER:O	32:DK:92:GLU:O	2.36	0.43
1:AA:949:A:N1	1:AA:950:U:C2	2.86	0.43
1:CA:202:G:H2'	1:CA:203:G:O4'	2.17	0.43
7:AG:27:VAL:HG12	7:AG:43:VAL:HG21	2.00	0.43
4:AD:17:THR:HG23	4:AD:18:ASP:N	2.33	0.43
39:DR:68:ARG:HD3	39:DR:92:TRP:CE2	2.53	0.43
22:DA:2325:G:C6	22:DA:2326:C:N4	2.86	0.43
43:BV:51:GLN:HB2	43:BV:57:TYR:OH	2.18	0.43
22:BA:322:A:C5	22:BA:340:A:C2	3.07	0.43
22:DA:2847:U:O4	22:DA:2848:G:C6	2.72	0.43
31:BJ:59:ALA:C	31:BJ:61:LYS:H	2.20	0.43
22:DA:2109:U:H4'	22:DA:2110:G:OP1	2.17	0.43
22:DA:277:G:H1'	22:DA:361:G:O6	2.18	0.43
25:DD:61:THR:HB	25:DD:63:PRO:HD2	2.00	0.43
3:AC:64:ILE:HG12	3:AC:66:VAL:HG23	2.01	0.43
1:CA:337:G:H2'	1:CA:338:A:C8	2.52	0.43
22:DA:425:G:N2	22:DA:426:C:C2	2.87	0.43
22:BA:659:G:H4'	26:BE:95:LYS:HD2	2.00	0.43
33:DL:136:GLU:HA	33:DL:140:GLY:HA3	2.00	0.43
5:AE:23:LYS:HB3	5:AE:30:ILE:HG23	2.00	0.43
22:BA:668:A:H2'	22:BA:670:A:H62	1.84	0.43
36:BO:35:ILE:HG23	36:BO:35:ILE:O	2.18	0.43
7:AG:50:LEU:O	7:AG:50:LEU:HD13	2.19	0.43
26:DE:187:VAL:O	26:DE:187:VAL:HG13	2.18	0.43
23:BB:73:A:H2'	23:BB:73:A:N3	2.34	0.43
23:DB:100:G:H2'	23:DB:101:A:O4'	2.18	0.43
1:AA:1171:A:H2'	1:AA:1172:C:C6	2.53	0.43
22:DA:1789:A:H5''	24:DC:219:THR:O	2.18	0.43
22:DA:1439:A:C8	22:DA:1440:U:C6	3.06	0.43
1:AA:979:C:OP1	1:AA:981:U:O4	2.37	0.43
22:DA:52:A:C2	22:DA:178:G:N3	2.87	0.43
50:D2:31:LEU:HD21	50:D2:43:THR:CG2	2.49	0.43
1:AA:1524:C:H2'	1:AA:1525:G:C8	2.54	0.43
22:DA:513:A:C2	22:DA:514:A:C5	3.06	0.43
1:CA:108:G:N3	1:CA:108:G:H5'	2.34	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:706:A:C6	1:AA:707:U:C4	3.06	0.43
22:DA:1829:A:O2'	24:DC:15:HIS:CE1	2.71	0.43
1:AA:1357:A:C6	1:AA:1358:U:C4	3.07	0.43
12:AL:94:ARG:HB2	12:AL:95:TYR:CE1	2.54	0.43
3:AC:7:PRO:HD2	3:AC:184:TYR:CG	2.53	0.43
22:DA:686:U:C6	22:DA:788:A:N1	2.84	0.43
2:CB:23:TRP:O	2:CB:23:TRP:CG	2.71	0.43
22:DA:2093:G:C2	22:DA:2094:A:C4	3.06	0.43
22:DA:2200:C:O2	22:DA:2226:C:N4	2.48	0.43
38:BQ:112:LYS:HD3	39:BR:48:LYS:HD2	2.00	0.43
22:DA:1935:G:H1'	22:DA:1964:G:H21	1.80	0.43
42:DU:77:THR:C	42:DU:79:LYS:H	2.22	0.43
22:DA:813:U:H2'	22:DA:814:C:H6	1.81	0.43
6:AF:92:THR:C	6:AF:93:LYS:HG2	2.38	0.43
1:CA:1068:G:H2'	1:CA:1069:C:H5'	2.00	0.43
22:DA:2502:G:C5'	22:DA:2503:A:O5'	2.66	0.43
1:CA:1317:C:O2'	14:CN:49:GLN:HG2	2.18	0.43
22:DA:1407:G:N2	22:DA:1596:A:N3	2.67	0.43
1:AA:208:U:C5	1:AA:210:C:N3	2.86	0.43
21:AU:26:ALA:O	21:AU:27:GLY:C	2.56	0.43
5:CE:45:ARG:HA	5:CE:72:ILE:O	2.19	0.43
1:CA:774:G:C4	1:CA:775:G:C8	3.06	0.43
22:DA:270:A:C2	22:DA:369:U:H4'	2.54	0.43
10:AJ:61:ALA:O	10:AJ:62:ARG:HB2	2.18	0.43
22:BA:1392:A:H61	41:BT:18:GLU:CD	2.22	0.43
10:CJ:80:THR:O	10:CJ:84:VAL:HB	2.17	0.43
22:DA:1476:U:H1'	22:DA:1732:C:C2	2.54	0.43
26:DE:5:LEU:O	26:DE:6:LYS:C	2.55	0.43
33:BL:21:ARG:HA	33:BL:21:ARG:HD3	1.77	0.43
1:CA:519:C:OP2	12:CL:47:SER:OG	2.36	0.43
14:AN:26:GLU:HG2	14:AN:27:LEU:N	2.33	0.43
36:BO:87:ILE:O	36:BO:88:LYS:O	2.37	0.43
22:DA:20:C:H2'	22:DA:21:A:C8	2.53	0.43
7:CG:46:ALA:CB	7:CG:120:LEU:HB3	2.48	0.43
3:CC:130:PHE:CE1	3:CC:157:LEU:HD23	2.53	0.43
1:CA:525:C:N4	1:CA:526:C:N4	2.67	0.43
22:DA:1838:C:C2	22:DA:1899:A:C2	3.07	0.43
42:DU:72:ILE:HG12	42:DU:83:VAL:HG23	2.00	0.43
1:AA:509:A:C6	1:AA:510:A:N1	2.86	0.43
1:AA:1251:A:H2'	1:AA:1252:A:C8	2.53	0.43
24:DC:31:ALA:N	24:DC:32:PRO:HD2	2.34	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:1429:G:O2'	22:BA:1430:G:H5'	2.19	0.43
1:AA:19:A:C2	1:AA:917:G:C4	3.06	0.43
29:BH:41:LYS:HA	29:BH:44:ILE:HG12	2.01	0.43
1:AA:350:G:O2'	1:AA:351:G:H5'	2.18	0.43
22:DA:2176:A:H2'	22:DA:2177:C:C6	2.54	0.43
1:AA:663:A:H5'	1:AA:836:G:OP1	2.18	0.43
1:AA:745:G:H2'	1:AA:746:A:C8	2.54	0.43
1:AA:749:A:C5	1:AA:750:C:C5	3.07	0.43
22:DA:1609:A:O2'	22:DA:1610:A:H5'	2.19	0.43
22:BA:1:G:H2'	22:BA:2:G:H8	1.83	0.43
36:DO:51:ALA:HB3	36:DO:78:VAL:HG22	1.99	0.43
1:CA:859:G:H2'	1:CA:860:A:C8	2.54	0.43
1:CA:392:C:H2'	1:CA:393:A:C8	2.53	0.43
1:AA:1431:A:C6	1:AA:1432:G:C6	3.06	0.43
32:BK:91:SER:O	32:BK:92:GLU:C	2.57	0.43
22:DA:1120:G:C6	22:DA:1121:C:C4	3.06	0.43
29:BH:103:VAL:O	29:BH:108:VAL:O	2.37	0.43
29:BH:93:SER:HB2	29:BH:122:LEU:HB2	1.99	0.43
29:BH:94:ILE:CD1	29:BH:98:ASP:HB3	2.48	0.43
1:CA:1394:A:H4'	1:CA:1395:C:OP2	2.18	0.43
2:AB:78:GLU:O	2:AB:80:VAL:N	2.52	0.43
22:DA:1388:G:N2	22:DA:1389:G:H1'	2.34	0.43
17:AQ:16:LYS:HA	17:AQ:16:LYS:HD2	1.91	0.43
1:CA:505:G:C2	1:CA:506:G:C5	3.06	0.43
5:CE:81:LEU:N	5:CE:81:LEU:CD1	2.82	0.43
4:CD:23:SER:O	4:CD:24:GLY:O	2.37	0.43
1:CA:109:A:C6	1:CA:327:A:C5	3.07	0.43
22:DA:2093:G:N7	22:DA:2225:A:N9	2.67	0.43
12:CL:74:LEU:HD11	12:CL:80:ILE:CG2	2.49	0.43
37:DP:89:ARG:HD2	37:DP:113:ARG:NH2	2.33	0.43
23:BB:30:C:OP1	36:BO:3:LYS:NZ	2.52	0.43
4:CD:130:VAL:CG1	4:CD:135:TYR:CD2	3.01	0.43
22:DA:322:A:O4'	22:DA:340:A:H1'	2.18	0.43
22:DA:1241:A:C2	22:DA:1242:U:C1'	3.02	0.43
22:DA:585:G:H2'	22:DA:586:A:N7	2.33	0.43
1:AA:1202:U:H2'	1:AA:1203:C:O4'	2.17	0.43
1:AA:1394:A:H4'	1:AA:1395:C:OP2	2.19	0.43
22:BA:846:U:C2'	22:BA:847:U:OP2	2.66	0.43
22:DA:142:A:C5	22:DA:143:C:N4	2.87	0.43
1:AA:1288:A:N1	1:AA:1289:A:C5	2.86	0.43
6:AF:29:ILE:HG23	6:AF:66:ALA:HB2	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:1713:A:C5	22:DA:1716:U:H1'	2.53	0.43
26:DE:5:LEU:CD2	26:DE:122:GLU:CD	2.86	0.43
11:CK:31:ILE:HB	11:CK:46:THR:HG22	2.01	0.43
21:CU:8:GLU:HB3	21:CU:12:PHE:CE2	2.52	0.43
22:DA:481:G:C4	22:DA:507:A:C2	3.07	0.43
42:DU:13:VAL:HB	42:DU:18:ASP:O	2.18	0.43
22:DA:2896:C:C4	22:DA:2897:U:C5	3.07	0.43
7:AG:31:MET:HG2	7:AG:32:VAL:N	2.33	0.43
46:DY:18:LEU:O	46:DY:22:LEU:HB3	2.18	0.43
50:D2:34:ARG:CB	50:D2:42:LEU:HD13	2.49	0.43
22:DA:1799:G:H8	24:DC:180:GLU:OE2	2.00	0.43
1:AA:1210:C:N4	1:AA:1211:U:C4	2.87	0.43
22:DA:1569:A:C6	22:DA:1570:A:C2	3.06	0.43
22:DA:1213:A:O2'	22:DA:1214:A:H5'	2.18	0.43
1:AA:1353:G:C2	1:AA:1354:U:C6	3.07	0.43
22:DA:2497:A:N3	22:DA:2498:C:N4	2.66	0.43
4:AD:165:ARG:O	4:AD:166:GLU:C	2.56	0.43
23:BB:48:U:H4'	36:BO:100:HIS:CD2	2.54	0.43
1:AA:504:C:H1'	1:AA:510:A:C4	2.53	0.43
10:AJ:80:THR:O	10:AJ:84:VAL:N	2.51	0.43
22:BA:1768:C:N3	22:BA:1769:U:C5	2.86	0.43
1:CA:517:G:C8	1:CA:531:U:C5	3.06	0.43
30:BI:34:ASN:CB	30:BI:37:GLU:HB2	2.49	0.43
1:AA:376:G:C2	1:AA:389:A:C2	3.07	0.43
43:DV:44:HIS:NE2	43:DV:85:LYS:HB2	2.33	0.43
22:BA:1930:G:HO2'	22:BA:1931:U:P	2.41	0.43
3:CC:22:TRP:CZ3	14:CN:94:PRO:HG2	2.53	0.43
1:AA:198:G:C5	1:AA:220:G:C2	3.06	0.43
22:BA:1966:A:N3	22:BA:2592:G:O2'	2.44	0.43
22:DA:1474:U:O4	22:DA:1475:G:N1	2.51	0.43
43:DV:35:GLU:N	43:DV:35:GLU:CD	2.71	0.43
40:DS:17:VAL:HB	40:DS:76:VAL:HG11	1.99	0.43
47:BZ:38:ARG:HB3	47:BZ:44:ILE:HD12	2.00	0.43
6:CF:47:LEU:CD2	6:CF:59:TYR:OH	2.66	0.43
1:CA:358:U:H2'	1:CA:358:U:O2	2.18	0.43
38:BQ:89:GLU:N	39:BR:49:ILE:HD12	2.32	0.43
24:DC:107:PRO:HA	24:DC:195:VAL:HA	1.99	0.43
1:CA:1127:G:H5'	1:CA:1280:A:O2'	2.19	0.43
29:DH:82:SER:O	29:DH:83:LYS:C	2.57	0.43
22:DA:396:G:O2'	22:DA:397:U:H5'	2.18	0.43
22:BA:760:G:C2'	22:BA:761:A:H5'	2.49	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:374:A:N3	1:CA:375:U:C6	2.86	0.43
22:DA:2212:A:C2	22:DA:2214:C:C4	3.07	0.43
1:CA:1309:G:O6	1:CA:1329:A:C2	2.71	0.43
22:DA:197:A:N3	22:DA:197:A:H2'	2.33	0.43
22:DA:488:G:C2	22:DA:493:G:O6	2.72	0.43
1:AA:1253:G:N3	1:AA:1254:A:C8	2.86	0.43
22:DA:2756:U:H4'	22:DA:2757:A:OP1	2.19	0.43
22:BA:2065:C:H2'	22:BA:2066:C:C6	2.52	0.43
22:BA:2886:A:C2	22:BA:2887:A:H1'	2.54	0.43
24:BC:105:LEU:O	24:BC:107:PRO:HD3	2.18	0.43
1:AA:74:A:N3	1:AA:97:G:C2	2.87	0.43
1:AA:427:U:C4	1:AA:428:G:C6	3.06	0.43
1:CA:159:G:H5'	1:CA:160:A:OP2	2.19	0.43
2:CB:188:ASP:OD2	2:CB:204:ASP:CG	2.57	0.43
10:CJ:88:MET:O	10:CJ:89:ARG:HB2	2.19	0.43
22:DA:2410:G:H2'	22:DA:2411:A:O4'	2.19	0.43
22:DA:1184:U:OP1	47:DZ:30:ARG:HD3	2.19	0.43
22:DA:2037:A:N6	22:DA:2038:G:O6	2.52	0.43
1:AA:464:U:H2'	1:AA:466:A:OP2	2.18	0.43
53:B5:213:VAL:O	53:B5:214:TYR:CB	2.67	0.43
1:AA:842:U:H3'	1:AA:843:U:C5'	2.48	0.43
22:BA:1176:U:OP1	22:BA:1176:U:H4'	2.19	0.43
22:BA:854:C:C2'	22:BA:855:G:H5'	2.49	0.43
42:BU:7:ARG:HD2	42:BU:26:LYS:O	2.19	0.43
22:DA:2378:A:N7	22:DA:2379:G:H1'	2.34	0.43
42:DU:95:PHE:O	42:DU:95:PHE:CD2	2.72	0.43
22:DA:567:U:H4'	22:DA:808:G:OP1	2.19	0.43
35:DN:34:ILE:HD11	35:DN:44:LEU:HD21	2.00	0.43
1:AA:1392:G:C5	1:AA:1393:U:C5	3.07	0.43
22:BA:1935:G:N2	22:BA:1964:G:C8	2.87	0.43
28:DG:86:LYS:HB3	28:DG:165:ALA:HB3	2.00	0.43
30:BI:67:PHE:N	30:BI:67:PHE:HD1	2.15	0.43
24:DC:126:PRO:HA	24:DC:192:LEU:O	2.18	0.43
23:DB:69:G:C2	23:DB:70:C:H1'	2.54	0.43
22:BA:768:G:C5	22:BA:769:U:C5	3.07	0.43
1:AA:338:A:H2'	1:AA:339:C:O4'	2.18	0.43
1:CA:1511:G:C5	1:CA:1512:U:C5	3.07	0.43
22:DA:2623:G:H4'	22:DA:2825:G:C8	2.54	0.43
1:AA:39:G:O2'	1:AA:40:C:H5'	2.19	0.43
33:DL:73:ILE:HA	33:DL:105:ILE:HD13	1.99	0.43
1:CA:678:U:N3	1:CA:713:G:N2	2.67	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:DG:8:PRO:HG3	28:DG:51:THR:HG22	2.01	0.43
22:BA:2309:A:N6	22:BA:2310:C:N4	2.67	0.43
47:DZ:27:LEU:O	47:DZ:36:VAL:HG11	2.19	0.43
1:CA:637:C:H2'	1:CA:638:U:C6	2.53	0.43
38:BQ:74:ILE:HG21	38:BQ:110:VAL:HG13	2.01	0.43
22:DA:442:G:N2	22:DA:444:C:C2	2.87	0.43
22:BA:2780:G:H4'	22:BA:2781:A:OP2	2.18	0.43
22:BA:393:C:C2	22:BA:394:C:C5	3.06	0.43
38:BQ:81:ASN:O	38:BQ:84:LYS:HB3	2.17	0.43
1:AA:775:G:O2'	1:AA:776:G:H5'	2.19	0.43
22:DA:1844:C:H5'	24:DC:254:GLY:O	2.19	0.43
16:AP:38:PHE:C	16:AP:38:PHE:CD1	2.92	0.43
5:CE:33:PHE:CD1	5:CE:33:PHE:N	2.87	0.43
31:DJ:84:ILE:HG13	31:DJ:84:ILE:O	2.17	0.43
22:DA:546:U:O2	22:DA:546:U:H3'	2.19	0.43
1:AA:1069:C:H2'	1:AA:1070:U:O5'	2.18	0.43
22:DA:1936:A:N7	22:DA:1945:G:C6	2.87	0.43
22:BA:1268:A:H2'	22:BA:1269:A:O4'	2.18	0.43
22:BA:1268:A:C2	22:BA:2013:A:C4	3.07	0.43
1:CA:1092:A:H62	1:CA:1093:A:N6	2.17	0.43
22:DA:70:G:H5''	22:DA:112:U:O2	2.19	0.43
22:DA:2147:A:N7	22:DA:2148:G:N7	2.67	0.43
1:AA:923:A:OP1	5:AE:26:LYS:HG2	2.19	0.43
1:AA:374:A:C4	1:AA:375:U:C6	3.06	0.43
1:AA:481:G:O2'	1:AA:482:A:P	2.77	0.43
22:DA:727:A:C6	22:DA:728:G:C6	3.06	0.43
22:DA:250:G:H4'	33:DL:59:ARG:CD	2.49	0.43
1:CA:322:C:OP2	1:CA:328:C:N4	2.52	0.43
2:CB:15:HIS:O	2:CB:16:PHE:C	2.57	0.43
5:AE:81:LEU:HD21	5:AE:123:VAL:CG1	2.49	0.43
22:DA:684:G:C5'	50:D2:16:HIS:HE1	2.32	0.43
22:DA:788:A:H1'	50:D2:4:THR:CG2	2.48	0.43
24:DC:100:GLU:HG2	24:DC:101:ARG:N	2.33	0.43
22:DA:1090:A:N6	22:DA:1091:G:O6	2.52	0.43
22:DA:574:A:H4'	22:DA:575:A:O5'	2.19	0.43
25:BD:12:THR:HG21	37:BP:9:GLU:HG2	2.01	0.43
22:BA:980:A:C6	22:BA:981:A:N1	2.86	0.43
22:DA:1833:C:O2	22:DA:1833:C:C2'	2.66	0.43
35:DN:79:LEU:O	35:DN:80:PHE:HB2	2.19	0.43
24:BC:107:PRO:HB3	24:BC:142:HIS:HE1	1.82	0.43
1:CA:920:U:H2'	1:CA:921:U:H6	1.78	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:CO:17:ARG:O	15:CO:18:ASP:HB3	2.19	0.43
22:BA:281:C:H2'	22:BA:282:A:H8	1.82	0.43
22:BA:1795:C:C4	22:BA:1796:U:C4	3.07	0.43
22:DA:484:C:N4	22:DA:497:A:C2	2.86	0.43
22:BA:2856:A:C6	22:BA:2857:G:C6	3.07	0.43
1:AA:1228:C:H2'	1:AA:1229:A:C8	2.54	0.43
29:DH:62:LEU:HD13	29:DH:63:ALA:N	2.34	0.43
36:DO:67:ASN:OD1	36:DO:69:ASP:HB2	2.19	0.43
22:DA:66:C:C4	22:DA:67:U:C4	3.07	0.43
2:CB:117:LEU:HB3	2:CB:141:LEU:HD11	2.00	0.43
27:BF:111:ILE:O	27:BF:114:PHE:HB2	2.19	0.43
20:AT:44:LYS:CD	20:AT:87:ALA:HA	2.49	0.43
5:AE:76:LEU:HB3	5:AE:77:ASN:H	1.68	0.43
1:AA:1003:G:H21	1:AA:1005:A:H5'	1.84	0.43
22:BA:1494:A:C2	22:BA:1495:A:C4	3.07	0.43
1:CA:778:G:C6	1:CA:779:C:N3	2.87	0.43
1:CA:777:A:C2'	1:CA:778:G:O5'	2.67	0.43
7:AG:71:PRO:O	7:AG:96:ARG:HG3	2.18	0.43
1:CA:926:G:C6	1:CA:1505:G:C5	3.07	0.43
28:DG:148:LEU:O	28:DG:162:VAL:HG11	2.19	0.43
22:DA:323:C:H2'	26:DE:163:ASN:CG	2.39	0.43
49:D1:26:ASN:CG	49:D1:29:THR:OG1	2.57	0.43
26:BE:115:GLN:O	26:BE:116:ASP:HB2	2.19	0.43
24:DC:69:ARG:NH2	24:DC:116:ILE:HD12	2.33	0.43
1:CA:487:A:H3'	1:CA:488:C:C6	2.53	0.43
43:DV:14:LYS:HG3	43:DV:15:GLY:N	2.34	0.43
22:BA:1638:C:O2	22:BA:2698:U:O2'	2.35	0.43
22:BA:2484:G:OP1	34:BM:44:ARG:HG2	2.18	0.43
6:AF:86:ARG:HG2	6:AF:86:ARG:HH11	1.83	0.43
22:BA:1664:A:C2	22:BA:2726:A:C8	3.07	0.43
3:AC:164:ARG:NH1	3:AC:166:GLU:OE1	2.52	0.43
29:BH:45:GLU:HA	29:BH:48:GLU:HB2	2.01	0.43
1:AA:1040:U:H2'	1:AA:1041:G:C8	2.54	0.43
7:CG:111:ARG:CZ	7:CG:122:ASN:HB3	2.49	0.43
22:BA:103:A:H2'	22:BA:104:A:O4'	2.19	0.43
15:CO:39:LEU:HA	15:CO:39:LEU:HD12	1.88	0.43
24:BC:20:VAL:O	24:BC:20:VAL:HG23	2.19	0.43
24:DC:266:PHE:CD1	24:DC:266:PHE:N	2.86	0.43
22:BA:158:U:O4	22:BA:159:G:C6	2.72	0.43
14:CN:16:LEU:HB3	14:CN:55:SER:HA	2.00	0.43
1:CA:54:C:H2'	1:CA:352:C:H41	1.84	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:BH:89:LYS:HB3	1:CA:359:G:H5''	2.00	0.43
1:CA:1182:G:H5'	1:CA:1184:G:H5''	2.00	0.43
22:BA:1917:U:O4	22:BA:1918:A:N1	2.52	0.43
28:BG:155:GLU:OE2	28:BG:158:LYS:HB2	2.19	0.43
1:AA:267:C:OP1	17:AQ:69:LYS:HB2	2.18	0.43
22:DA:179:C:C2	22:DA:180:G:C8	3.06	0.43
42:BU:13:VAL:HG12	42:BU:19:LYS:HA	2.00	0.43
22:DA:1045:C:N4	22:DA:1111:A:H2'	2.34	0.43
1:CA:977:A:C2'	1:CA:1223:C:N4	2.81	0.43
1:CA:977:A:N3	1:CA:977:A:H3'	2.34	0.43
29:DH:27:ARG:NE	45:DX:60:ASP:CG	2.68	0.43
4:AD:99:ASP:OD2	4:AD:115:ARG:NH2	2.52	0.43
1:AA:406:G:C5	1:AA:495:A:C8	3.07	0.43
4:AD:110:THR:O	4:AD:113:GLU:N	2.51	0.43
22:DA:536:G:O6	22:DA:537:G:C6	2.72	0.43
22:DA:753:A:C2	22:DA:754:U:N3	2.87	0.43
5:CE:38:VAL:HG12	5:CE:39:VAL:N	2.33	0.43
1:CA:674:G:H2'	1:CA:675:A:H8	1.84	0.43
1:AA:1319:A:C4	1:AA:1323:G:C8	3.06	0.43
20:AT:3:ASN:O	20:AT:5:LYS:N	2.52	0.43
22:DA:749:A:N3	22:DA:750:A:C8	2.87	0.43
22:BA:833:A:H2'	22:BA:834:G:C8	2.54	0.43
1:CA:718:A:C8	1:CA:719:C:C5	3.07	0.43
1:CA:735:C:H2'	1:CA:736:C:H6	1.82	0.43
22:BA:319:G:C5	22:BA:333:G:C2	3.07	0.43
1:AA:1122:U:C4	1:AA:1123:U:C5	3.06	0.43
22:DA:319:G:C5	22:DA:333:G:C2	3.06	0.43
2:CB:187:VAL:HG23	2:CB:187:VAL:O	2.19	0.43
33:DL:29:LYS:O	33:DL:30:THR:CB	2.67	0.43
25:DD:105:LYS:O	25:DD:177:VAL:HG12	2.19	0.43
30:DI:57:VAL:HG22	30:DI:69:PHE:HB2	2.01	0.43
22:DA:9:G:C6	22:DA:2629:U:C6	3.07	0.43
1:CA:801:U:C2	1:CA:802:A:C8	3.07	0.43
1:AA:1195:C:O2	1:AA:1197:A:H1'	2.19	0.43
1:AA:1521:C:H2'	1:AA:1522:U:H6	1.84	0.43
22:DA:648:G:C2	22:DA:649:G:C5	3.07	0.43
22:BA:1879:C:C5	22:BA:1880:U:C5	3.07	0.43
9:CI:12:ARG:NH1	9:CI:13:LYS:HB2	2.34	0.43
1:CA:1306:A:H1'	1:CA:1332:A:N7	2.33	0.43
33:DL:57:LEU:HA	33:DL:60:ARG:HB3	2.01	0.43
22:DA:38:A:H2'	22:DA:39:G:O4'	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:BQ:21:ALA:HA	38:BQ:24:TYR:CE2	2.54	0.43
12:AL:87:VAL:HG11	12:AL:90:LEU:HD22	2.01	0.43
22:DA:5:A:C2	22:DA:2899:A:C2	3.07	0.43
15:CO:41:GLY:O	15:CO:42:HIS:C	2.57	0.43
1:AA:595:A:C6	1:AA:641:U:C6	3.06	0.43
12:AL:38:TYR:O	12:AL:39:THR:CG2	2.66	0.43
2:AB:115:LYS:O	2:AB:117:LEU:N	2.49	0.43
1:AA:33:A:O2'	12:AL:29:GLN:OE1	2.19	0.43
1:AA:1129:C:C6	1:AA:1139:G:N7	2.87	0.43
37:DP:98:TYR:HD1	37:DP:101:ARG:NH2	2.16	0.43
1:CA:1408:A:C2	1:CA:1494:G:C4	3.07	0.43
26:BE:48:THR:C	26:BE:50:ALA:N	2.69	0.43
31:DJ:25:LEU:HD13	31:DJ:100:VAL:HG12	2.00	0.43
22:BA:2176:A:C6	22:BA:2177:C:N4	2.86	0.43
22:DA:321:U:H4'	26:DE:159:LEU:O	2.18	0.43
22:BA:716:A:N6	22:BA:717:C:C4	2.87	0.43
2:CB:162:PHE:HA	2:CB:184:PHE:O	2.19	0.43
22:BA:158:U:C4	22:BA:159:G:C5	3.07	0.43
22:BA:21:A:O2'	22:BA:22:C:H5'	2.19	0.43
28:BG:109:PHE:CE1	28:BG:152:ARG:NH2	2.87	0.43
9:AI:56:ASP:O	9:AI:60:LYS:NZ	2.28	0.43
22:DA:2552:U:C2	22:DA:2554:U:C5'	3.01	0.43
22:BA:47:C:C2'	22:BA:48:G:H5'	2.49	0.43
22:BA:742:A:H2'	22:BA:743:A:C8	2.54	0.43
22:BA:1562:U:H2'	22:BA:1563:U:O4'	2.19	0.43
3:AC:53:SER:O	3:AC:54:ARG:HB2	2.18	0.43
22:DA:2569:G:C2	22:DA:2570:G:C8	3.06	0.43
47:DZ:7:ILE:O	47:DZ:35:THR:HA	2.18	0.43
22:BA:204:A:O4'	22:BA:206:U:C6	2.72	0.43
41:BT:57:VAL:HG22	41:BT:58:VAL:N	2.33	0.43
16:AP:20:VAL:HG23	16:AP:35:ARG:HA	2.01	0.43
30:BI:98:VAL:HG12	30:BI:99:GLY:N	2.34	0.43
5:AE:33:PHE:CD2	5:AE:56:VAL:HG22	2.54	0.43
3:CC:175:LEU:O	3:CC:175:LEU:HG	2.19	0.43
29:BH:79:THR:CG2	29:BH:147:VAL:CG2	2.97	0.42
22:BA:1179:G:N7	22:BA:1180:U:C1'	2.82	0.42
22:DA:2303:G:O4'	27:DF:123:ASP:HA	2.19	0.42
22:DA:1794:A:C6	22:DA:1795:C:C4	3.07	0.42
22:DA:2127:G:C2	22:DA:2162:G:C8	3.07	0.42
22:DA:1953:A:C6	22:DA:2550:G:C4'	3.02	0.42
22:DA:1992:G:N2	22:DA:1995:U:C5	2.87	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AB:164:ILE:O	2:AB:186:ILE:HG12	2.19	0.42
22:DA:910:A:N3	34:DM:13:HIS:CE1	2.87	0.42
1:AA:958:A:C6	1:AA:959:A:C2	3.07	0.42
1:CA:577:G:O2'	1:CA:578:C:H5'	2.19	0.42
5:AE:101:GLU:OE2	5:AE:101:GLU:O	2.37	0.42
22:DA:1363:C:O2	22:DA:1369:G:C2	2.72	0.42
22:DA:2093:G:C5	22:DA:2225:A:C5	3.07	0.42
42:DU:97:LYS:O	42:DU:98:SER:OG	2.29	0.42
24:DC:48:ARG:HG3	24:DC:48:ARG:NH1	2.34	0.42
22:DA:2133:G:N3	22:DA:2158:A:N1	2.67	0.42
1:CA:273:U:H2'	1:CA:274:A:H5'	2.01	0.42
1:CA:756:C:C2	1:CA:757:U:C6	3.07	0.42
22:BA:2064:C:H2'	22:BA:2065:C:C6	2.54	0.42
33:DL:94:THR:O	33:DL:98:ALA:N	2.50	0.42
1:CA:622:A:H5''	1:CA:623:C:OP2	2.19	0.42
1:CA:1478:U:N3	1:CA:1479:C:C5	2.87	0.42
22:BA:1006:C:C2'	22:BA:1007:C:H5'	2.49	0.42
48:B0:4:GLN:HG3	48:B0:4:GLN:O	2.18	0.42
6:CF:18:VAL:HG12	6:CF:19:PRO:N	2.33	0.42
42:DU:57:GLY:O	42:DU:59:VAL:HG23	2.20	0.42
30:BI:77:ALA:HA	30:BI:80:LEU:HD12	2.01	0.42
1:AA:1539:C:OP1	21:AU:18:ARG:HG3	2.19	0.42
22:BA:1113:U:H2'	22:BA:1114:C:C6	2.53	0.42
22:DA:45:G:H2'	22:DA:215:G:C5	2.54	0.42
22:DA:420:C:C2	22:DA:421:C:C5	3.07	0.42
27:BF:42:GLU:OE2	27:BF:49:LEU:CD2	2.67	0.42
22:DA:1166:G:H2'	22:DA:1167:C:O4'	2.18	0.42
12:AL:36:ARG:HB3	12:AL:38:TYR:CE1	2.54	0.42
19:AS:36:ARG:NE	19:AS:52:HIS:O	2.47	0.42
1:CA:1234:C:HO2'	1:CA:1364:U:H6	1.66	0.42
1:CA:1264:U:O2	1:CA:1272:G:N2	2.52	0.42
45:DX:7:VAL:HG12	45:DX:8:THR:HG23	2.01	0.42
41:BT:54:GLU:HB3	41:BT:88:LYS:HG3	2.00	0.42
1:AA:1048:G:C2	1:AA:1050:G:C5	3.07	0.42
15:AO:82:ILE:HA	15:AO:87:LEU:CD2	2.49	0.42
33:DL:20:GLY:HA2	33:DL:28:GLY:HA2	2.00	0.42
22:BA:1605:C:H2'	22:BA:1606:C:H5'	2.01	0.42
3:AC:167:TRP:CE3	3:AC:167:TRP:N	2.86	0.42
1:CA:723:U:O2	1:CA:855:U:H4'	2.19	0.42
35:DN:117:ASP:O	35:DN:118:ARG:HB2	2.19	0.42
2:CB:165:ASP:OD2	2:CB:168:HIS:HB2	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BJ:59:ALA:C	31:BJ:61:LYS:N	2.73	0.42
22:BA:1638:C:H1'	22:BA:2698:U:O2'	2.19	0.42
22:DA:1796:U:H2'	22:DA:1797:G:C8	2.54	0.42
37:BP:13:MET:HB3	37:BP:13:MET:HE2	1.81	0.42
22:DA:2547:A:C2	22:DA:2562:U:C2	3.07	0.42
22:DA:464:U:C4	22:DA:465:G:C6	3.07	0.42
22:BA:2284:A:OP1	49:B1:4:GLY:O	2.37	0.42
7:AG:106:GLU:O	7:AG:110:LYS:HG2	2.19	0.42
2:AB:173:ILE:HG22	2:AB:177:ASN:ND2	2.34	0.42
25:BD:136:ASN:HD21	25:BD:139:SER:H	1.66	0.42
15:CO:57:LEU:O	15:CO:60:VAL:HB	2.19	0.42
25:DD:110:THR:HG22	25:DD:111:GLY:N	2.34	0.42
1:CA:1350:A:N6	1:CA:1373:G:N2	2.66	0.42
28:BG:75:MET:O	28:BG:78:GLY:N	2.52	0.42
11:CK:14:LYS:O	11:CK:15:GLN:HB3	2.17	0.42
11:CK:17:SER:O	11:CK:80:LYS:N	2.48	0.42
9:CI:44:ALA:HB1	9:CI:47:VAL:HG21	2.00	0.42
22:BA:1428:C:N4	22:BA:1570:A:OP2	2.41	0.42
31:BJ:104:ALA:O	31:BJ:108:MET:HG3	2.18	0.42
40:BS:109:ASP:OD1	40:BS:110:ARG:N	2.50	0.42
22:BA:2350:C:H2'	22:BA:2351:G:O4'	2.18	0.42
22:BA:946:C:P	57:BA:3353:HOH:O	2.77	0.42
8:CH:8:ALA:HB2	8:CH:77:ARG:HG3	2.00	0.42
34:BM:42:THR:O	34:BM:43:ALA:C	2.56	0.42
33:BL:116:VAL:O	33:BL:116:VAL:HG13	2.19	0.42
18:AR:52:GLN:HA	18:AR:52:GLN:OE1	2.19	0.42
2:CB:10:LEU:HD23	2:CB:10:LEU:O	2.19	0.42
40:DS:25:ARG:CZ	40:DS:25:ARG:HB2	2.49	0.42
1:AA:1084:G:C5	1:AA:1085:U:C4	3.07	0.42
22:DA:1594:U:H2'	22:DA:1595:C:C6	2.54	0.42
22:BA:2648:G:H2'	22:BA:2649:C:C6	2.54	0.42
22:BA:2280:G:O6	44:BW:14:ARG:HD2	2.19	0.42
22:DA:1607:C:O2	22:DA:1621:U:C5	2.72	0.42
22:DA:189:G:C5	22:DA:205:G:N2	2.87	0.42
1:CA:1521:C:H2'	1:CA:1522:U:O5'	2.20	0.42
5:CE:122:ASN:CG	5:CE:123:VAL:H	2.23	0.42
21:CU:36:GLU:OE2	21:CU:38:TYR:HD2	2.02	0.42
21:CU:38:TYR:O	21:CU:39:GLU:HB2	2.19	0.42
21:CU:41:PRO:C	21:CU:43:THR:N	2.70	0.42
1:AA:109:A:C6	1:AA:326:G:C6	3.07	0.42
22:DA:674:G:H1'	26:DE:69:ARG:CD	2.48	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:1692:U:H2'	22:DA:1694:C:C4	2.54	0.42
5:CE:133:PRO:HA	5:CE:136:VAL:HG12	2.01	0.42
1:CA:1105:A:N3	1:CA:1106:G:C8	2.88	0.42
5:AE:90:THR:CG2	5:AE:91:GLY:N	2.77	0.42
4:CD:38:PRO:HD2	4:CD:42:GLY:HA3	2.01	0.42
33:BL:82:LEU:C	33:BL:84:LYS:H	2.22	0.42
1:AA:872:A:C2	1:AA:874:G:C6	3.07	0.42
2:AB:50:PHE:HB2	2:AB:213:TYR:OH	2.19	0.42
6:AF:5:GLU:OE2	18:AR:24:LYS:HE3	2.20	0.42
22:DA:430:A:H2'	22:DA:431:U:H5'	2.01	0.42
22:BA:2888:C:O2	22:BA:2888:C:H2'	2.18	0.42
1:CA:502:A:OP1	12:CL:115:SER:HB2	2.19	0.42
22:BA:1833:C:H2'	22:BA:1834:U:H6	1.84	0.42
1:AA:1061:G:C4	1:AA:1062:U:C6	3.07	0.42
22:DA:1097:U:O2	30:DI:9:VAL:HG11	2.19	0.42
1:AA:1228:C:OP1	13:AM:115:PRO:HD2	2.19	0.42
23:DB:52:A:C5	36:DO:33:ARG:NH2	2.87	0.42
22:DA:697:G:C2	22:DA:766:U:C2	3.07	0.42
2:CB:141:LEU:O	2:CB:145:GLU:N	2.50	0.42
15:CO:47:LYS:O	15:CO:53:ARG:NH2	2.53	0.42
32:DK:6:THR:O	32:DK:8:LEU:HD12	2.19	0.42
19:AS:32:ARG:HD3	19:AS:57:HIS:NE2	2.34	0.42
22:DA:415:A:C2	22:DA:2409:G:C2	3.07	0.42
1:AA:66:A:H4'	1:AA:173:U:C5	2.54	0.42
22:DA:1087:G:C6	22:DA:1089:A:C2	3.07	0.42
22:BA:973:A:H5''	39:BR:81:LYS:HG3	2.01	0.42
1:CA:666:G:C4	1:CA:667:G:C8	3.07	0.42
26:BE:131:THR:HG22	26:BE:160:ALA:HA	2.00	0.42
3:CC:134:MET:SD	3:CC:153:VAL:HG22	2.60	0.42
22:BA:2492:U:H2'	22:BA:2493:U:C6	2.54	0.42
18:CR:33:ILE:O	18:CR:33:ILE:HG12	2.18	0.42
40:BS:20:VAL:HG11	40:BS:44:ALA:HA	2.00	0.42
1:CA:1534:A:H4'	1:CA:1535:C:H2'	2.01	0.42
26:DE:97:ASN:HB2	26:DE:100:MET:CG	2.49	0.42
30:DI:33:VAL:HG22	30:DI:67:PHE:CE1	2.53	0.42
3:AC:155:GLY:HA2	3:AC:163:ALA:HB1	2.01	0.42
21:CU:14:VAL:HG12	21:CU:16:LEU:HG	2.01	0.42
36:DO:36:TYR:CD1	36:DO:36:TYR:N	2.86	0.42
6:AF:49:TYR:C	6:AF:49:TYR:CD1	2.92	0.42
5:AE:60:ILE:O	5:AE:64:MET:HG2	2.19	0.42
22:BA:2838:G:C6	22:BA:2839:G:C5	3.08	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:DW:46:HIS:CD2	44:DW:77:ARG:HB3	2.54	0.42
4:AD:173:VAL:HG22	4:AD:174:ASP:N	2.34	0.42
8:AH:79:SER:OG	8:AH:84:ARG:HA	2.19	0.42
9:AI:113:ARG:NH2	10:AJ:64:GLN:OE1	2.48	0.42
53:B5:74:ARG:HA	53:B5:93:ASP:OD1	2.20	0.42
8:CH:67:GLN:C	8:CH:69:LYS:N	2.72	0.42
22:BA:2441:U:OP2	22:BA:2586:U:O2'	2.32	0.42
38:BQ:97:ASP:OD1	38:BQ:101:PHE:HD2	2.02	0.42
10:CJ:8:ILE:HG22	10:CJ:10:LEU:HD11	2.01	0.42
11:CK:40:ASN:O	11:CK:41:ALA:HB3	2.18	0.42
25:DD:16:THR:OG1	25:DD:18:ASP:OD1	2.33	0.42
24:BC:83:TYR:CD1	24:BC:83:TYR:C	2.93	0.42
27:DF:46:ASP:N	27:DF:46:ASP:OD1	2.52	0.42
2:CB:41:ILE:HD12	2:CB:41:ILE:C	2.40	0.42
15:CO:24:SER:O	15:CO:27:VAL:HB	2.19	0.42
25:DD:187:LEU:HD21	25:DD:203:VAL:HG11	2.00	0.42
22:BA:2458:G:C2	22:BA:2490:G:N2	2.87	0.42
22:BA:2377:A:O2'	22:BA:2378:A:H5'	2.18	0.42
1:AA:244:U:O4	1:AA:906:A:H1'	2.19	0.42
22:DA:996:A:C2	22:DA:997:G:N9	2.87	0.42
1:CA:1161:C:H2'	1:CA:1162:C:C6	2.53	0.42
22:BA:2243:U:O2'	22:BA:2244:U:H5'	2.19	0.42
22:BA:2659:G:OP2	28:BG:158:LYS:NZ	2.48	0.42
41:BT:2:ILE:CD1	41:BT:45:ALA:HB1	2.47	0.42
22:DA:447:A:H5'	22:DA:449:A:C4	2.53	0.42
38:DQ:27:ALA:HB1	38:DQ:31:VAL:HG23	2.01	0.42
22:BA:2298:A:N6	22:BA:2318:G:H1'	2.34	0.42
1:AA:324:G:N2	1:AA:326:G:H3'	2.35	0.42
1:CA:687:A:C8	1:CA:701:U:C5	3.07	0.42
22:BA:998:C:OP2	38:BQ:58:ARG:NH2	2.51	0.42
1:CA:1211:U:O2'	1:CA:1212:U:P	2.77	0.42
1:CA:991:U:N3	1:CA:1212:U:O4'	2.53	0.42
26:BE:108:ILE:HD12	26:BE:108:ILE:C	2.40	0.42
1:AA:406:G:N2	1:AA:407:U:C2	2.87	0.42
1:AA:495:A:H4'	1:AA:496:A:O4'	2.19	0.42
37:BP:52:ASN:C	37:BP:53:ARG:HG2	2.39	0.42
22:DA:1581:G:C6	22:DA:1582:C:C4	3.07	0.42
22:BA:181:A:C2	22:BA:182:A:C4	3.06	0.42
22:DA:2131:U:H4'	22:DA:2133:G:O4'	2.18	0.42
22:BA:2552:U:C2	22:BA:2554:U:C5'	3.02	0.42
4:CD:130:VAL:HG11	4:CD:135:TYR:CG	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:2341:G:C6	22:DA:2342:C:C4	3.08	0.42
22:DA:2347:C:H2'	22:DA:2348:U:C5	2.54	0.42
22:DA:2111:U:C4	22:DA:2145:C:H2'	2.54	0.42
22:DA:2366:A:C2	22:DA:2367:G:H1'	2.55	0.42
22:BA:2820:A:OP1	35:BN:2:ARG:NH2	2.42	0.42
22:DA:1171:G:C2	22:DA:1179:G:O6	2.72	0.42
22:BA:1901:A:O2'	24:BC:244:PRO:HG2	2.18	0.42
22:BA:136:G:H2'	22:BA:137:U:C6	2.54	0.42
6:CF:18:VAL:O	6:CF:21:MET:HB2	2.18	0.42
30:BI:113:LYS:HE3	30:BI:116:ASP:HB3	2.02	0.42
22:DA:2297:A:N1	22:DA:2321:U:H5	2.17	0.42
1:CA:922:G:H2'	1:CA:923:A:C8	2.54	0.42
1:AA:810:C:C2'	1:AA:810:C:O2	2.64	0.42
2:AB:17:GLY:HA2	2:AB:41:ILE:HG23	2.02	0.42
22:DA:2897:U:H2'	22:DA:2898:U:C6	2.54	0.42
25:BD:62:LYS:CB	25:BD:63:PRO:HD3	2.49	0.42
17:CQ:11:ARG:HA	17:CQ:58:VAL:HA	2.01	0.42
39:DR:78:ARG:CB	39:DR:83:TYR:HD2	2.32	0.42
1:AA:842:U:H3'	1:AA:843:U:H4'	2.01	0.42
22:BA:1578:U:H2'	22:BA:1578:U:O2	2.19	0.42
1:AA:716:A:C2'	1:AA:717:U:O5'	2.68	0.42
1:AA:539:A:C6	1:AA:540:G:C6	3.08	0.42
7:CG:68:ASN:O	7:CG:138:ARG:NH2	2.51	0.42
22:BA:1935:G:C6	22:BA:1962:C:C6	3.06	0.42
1:AA:1016:A:C5	1:AA:1017:U:H1'	2.53	0.42
13:CM:39:ILE:HG13	13:CM:56:LEU:HD21	2.01	0.42
22:DA:2456:C:C4	22:DA:2457:U:C5	3.07	0.42
29:BH:45:GLU:C	29:BH:47:PHE:N	2.72	0.42
22:DA:2070:A:C2	22:DA:2442:C:C2	3.07	0.42
8:CH:27:MET:HB2	8:CH:28:PRO:HD2	2.00	0.42
1:CA:1153:G:H2'	1:CA:1154:G:O4'	2.19	0.42
53:B5:64:SER:O	53:B5:65:LEU:CB	2.67	0.42
22:BA:630:G:H5''	22:BA:631:A:OP2	2.20	0.42
1:AA:560:A:H5'	1:AA:566:G:N2	2.34	0.42
1:AA:1290:G:N2	1:AA:1291:U:H1'	2.34	0.42
24:BC:119:GLY:O	24:BC:121:ASP:N	2.52	0.42
22:BA:1467:U:C4	22:BA:1546:G:C2	3.08	0.42
1:CA:1303:C:N4	1:CA:1304:G:C6	2.88	0.42
40:DS:6:LYS:HE2	40:DS:104:THR:HG23	2.01	0.42
22:DA:2100:G:C6	22:DA:2101:A:C5	3.07	0.42
1:CA:47:C:H4'	1:CA:48:C:OP1	2.18	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:DE:3:LEU:HD11	26:DE:19:PHE:CD2	2.54	0.42
1:CA:983:A:N3	1:CA:983:A:C2'	2.82	0.42
26:DE:105:LEU:HA	26:DE:105:LEU:HD23	1.94	0.42
50:B2:9:VAL:HG12	50:B2:13:ASN:ND2	2.34	0.42
1:AA:28:A:C2	1:AA:29:U:C2	3.07	0.42
4:CD:62:ARG:NH1	4:CD:69:GLU:HG2	2.34	0.42
1:CA:58:C:C2'	1:CA:59:A:O5'	2.66	0.42
22:BA:1918:A:H4'	22:BA:1919:A:OP1	2.19	0.42
22:DA:1361:G:N3	22:DA:1362:C:C6	2.86	0.42
11:AK:126:LYS:HA	21:AU:34:ARG:NH2	2.34	0.42
22:DA:1047:G:N2	22:DA:1110:G:O2'	2.52	0.42
22:DA:517:C:O4'	40:DS:78:GLU:OE2	2.36	0.42
25:BD:13:ARG:HD3	25:BD:21:SER:OG	2.19	0.42
1:AA:1077:G:C6	1:AA:1081:A:N6	2.87	0.42
1:CA:685:G:C2	1:CA:686:U:O4	2.73	0.42
4:AD:11:LEU:HD22	4:AD:63:ARG:HD3	2.01	0.42
22:BA:2380:C:H5'	36:BO:17:LYS:NZ	2.34	0.42
22:DA:328:U:O3'	42:DU:66:GLN:NE2	2.52	0.42
4:AD:130:VAL:CG1	4:AD:135:TYR:CD2	3.00	0.42
1:AA:1053:G:C4	1:AA:1199:U:C4	3.08	0.42
29:BH:114:GLU:CB	29:BH:133:GLN:O	2.66	0.42
1:CA:783:C:N3	1:CA:784:A:N7	2.68	0.42
1:CA:1220:G:H4'	19:CS:34:TRP:O	2.18	0.42
1:CA:582:C:O2	1:CA:760:G:C2	2.72	0.42
1:CA:1518:A:C2	1:CA:1519:A:C4	3.08	0.42
10:AJ:11:LYS:HA	10:AJ:70:HIS:O	2.19	0.42
1:AA:1074:G:O2'	1:AA:1101:A:N1	2.45	0.42
26:DE:149:ILE:CG1	26:DE:188:MET:HE3	2.49	0.42
22:DA:2067:G:C6	22:DA:2444:G:N1	2.87	0.42
1:AA:194:C:C2'	1:AA:195:A:H5'	2.49	0.42
22:DA:1094:U:H2'	22:DA:1096:A:OP2	2.19	0.42
22:DA:2677:G:C4	22:DA:2731:G:N2	2.87	0.42
22:BA:1045:C:C4'	22:BA:1046:A:H5'	2.49	0.42
1:CA:692:U:H1'	1:CA:695:A:N7	2.35	0.42
22:DA:1156:A:C2	38:DQ:47:TYR:HE1	2.37	0.42
30:BI:113:LYS:HE2	30:BI:116:ASP:OD2	2.20	0.42
39:DR:49:ILE:HG23	39:DR:54:VAL:HG23	1.99	0.42
32:DK:87:LEU:HD22	32:DK:92:GLU:HA	2.01	0.42
21:AU:14:VAL:O	21:AU:16:LEU:CD1	2.68	0.42
1:AA:82:G:N2	1:AA:84:U:O4	2.52	0.42
1:CA:101:A:C4	1:CA:102:G:C8	3.07	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:AD:50:ASP:O	4:AD:53:VAL:HG22	2.20	0.42
25:DD:176:ASP:HB2	25:DD:190:LYS:HB3	2.02	0.42
16:AP:77:GLU:C	16:AP:79:ASN:H	2.23	0.42
1:CA:1328:C:H5''	13:CM:28:THR:CG2	2.48	0.42
22:DA:931:U:O4	22:DA:1184:U:O4'	2.37	0.42
17:CQ:12:VAL:HG11	17:CQ:55:ILE:HA	2.01	0.42
35:BN:79:LEU:O	35:BN:81:ASN:N	2.49	0.42
29:DH:72:ILE:O	29:DH:72:ILE:CG2	2.67	0.42
1:AA:1516:G:H2'	1:AA:1518:A:OP2	2.19	0.42
13:AM:90:ARG:HB3	13:AM:97:VAL:HG22	2.00	0.42
22:BA:1406:U:H2'	22:BA:1407:G:H8	1.84	0.42
2:CB:19:GLN:O	2:CB:38:VAL:HG22	2.20	0.42
22:DA:2261:C:C2	22:DA:2280:G:N2	2.87	0.42
34:BM:6:ARG:O	34:BM:7:THR:HG23	2.19	0.42
25:BD:39:ASP:OD1	25:BD:40:LEU:N	2.53	0.42
3:CC:50:ALA:HA	3:CC:75:ILE:HG21	2.00	0.42
17:AQ:59:VAL:CG2	17:AQ:60:GLU:N	2.83	0.42
22:DA:1820:U:O2'	24:DC:158:ALA:HB3	2.19	0.42
22:DA:2077:A:C8	22:DA:2435:A:C4	3.07	0.42
4:CD:148:LYS:H	4:CD:148:LYS:HD3	1.85	0.42
1:AA:614:C:H2'	1:AA:615:G:O5'	2.20	0.42
36:DO:49:VAL:CG2	36:DO:85:LYS:HG3	2.49	0.42
1:CA:144:G:C5	1:CA:179:A:C2	3.07	0.42
22:DA:607:U:H5	22:DA:619:G:C4	2.38	0.42
4:AD:102:VAL:CG1	4:AD:114:ALA:HB1	2.49	0.42
1:AA:19:A:N3	1:AA:917:G:C2	2.87	0.42
25:BD:142:VAL:HB	25:BD:143:PRO:CD	2.49	0.42
22:BA:739:A:N1	57:BA:3305:HOH:O	2.37	0.42
22:DA:2732:G:O2'	22:DA:2733:A:H5'	2.19	0.42
34:BM:30:SER:H	34:BM:106:ASP:HB2	1.84	0.42
1:CA:866:C:C4	1:CA:867:G:H1'	2.54	0.42
52:B4:19:ARG:HB2	52:B4:24:ARG:HD2	2.00	0.42
30:BI:45:LYS:HE2	30:BI:45:LYS:HB2	1.87	0.42
2:AB:207:ILE:HD13	2:AB:207:ILE:N	2.34	0.42
18:CR:67:LEU:HD23	18:CR:67:LEU:N	2.34	0.42
23:DB:5:U:C2	23:DB:116:G:N2	2.87	0.42
39:BR:15:SER:O	39:BR:18:GLN:HB3	2.19	0.42
42:DU:8:ASP:O	42:DU:9:ASP:HB2	2.19	0.42
1:CA:147:G:N2	1:CA:148:G:C2	2.87	0.42
22:BA:1078:U:H5''	22:BA:1079:C:OP1	2.19	0.42
1:AA:979:C:OP1	1:AA:1223:C:N4	2.53	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:2394:C:OP2	51:D3:30:ARG:NH2	2.51	0.42
22:DA:51:G:H4'	22:DA:52:A:H5'	2.02	0.42
22:DA:201:C:C6	22:DA:202:U:C5	3.05	0.42
11:CK:123:PRO:CB	11:CK:124:PRO:HD2	2.49	0.42
5:CE:112:ARG:O	5:CE:115:LEU:N	2.53	0.42
22:BA:2187:U:C4	22:BA:2188:U:O4	2.72	0.42
1:AA:685:G:N2	1:AA:706:A:N6	2.67	0.42
22:DA:1774:C:O2	24:DC:11:PRO:HB2	2.20	0.42
22:DA:1975:G:C4	22:DA:1976:U:C6	3.08	0.42
29:DH:31:VAL:HG12	29:DH:32:PRO:HD3	2.02	0.42
1:CA:581:G:H3'	1:CA:758:C:H42	1.83	0.42
22:BA:1385:A:C2	22:BA:1386:C:N3	2.87	0.42
1:CA:273:U:C4	1:CA:274:A:N7	2.88	0.42
22:BA:2847:U:H2'	22:BA:2848:G:C5'	2.50	0.42
22:DA:1027:A:H61	22:DA:1126:A:H1'	1.84	0.42
22:DA:2810:A:C8	22:DA:2811:G:C8	3.08	0.42
22:BA:693:A:C2'	22:BA:694:U:H5'	2.49	0.42
5:AE:97:GLN:HG2	5:AE:98:PRO:HD2	2.02	0.42
1:CA:216:U:H2'	1:CA:217:C:C6	2.55	0.42
5:AE:151:GLU:O	5:AE:153:VAL:N	2.53	0.42
22:DA:1681:G:H2'	22:DA:1757:A:N1	2.34	0.42
22:BA:958:U:C6	34:BM:14:LYS:HD3	2.55	0.42
1:AA:942:G:N2	1:AA:943:U:C2	2.87	0.42
15:AO:88:ARG:O	15:AO:89:ARG:CB	2.67	0.42
25:BD:33:ARG:NH1	25:BD:53:GLY:O	2.53	0.42
20:CT:62:ALA:HA	20:CT:68:HIS:H	1.84	0.42
20:CT:62:ALA:HA	20:CT:68:HIS:N	2.35	0.42
1:CA:86:G:H1'	1:CA:87:C:O4'	2.20	0.42
22:BA:368:A:C6	22:BA:369:U:C4	3.06	0.42
1:AA:1000:A:C2	1:AA:1041:G:C2	3.07	0.42
22:BA:2000:C:O2'	22:BA:2001:C:H5'	2.20	0.42
23:DB:85:G:N2	23:DB:92:C:C2	2.87	0.42
1:CA:37:U:OP1	12:CL:121:ARG:HB2	2.19	0.42
22:BA:2443:C:O2'	22:BA:2444:G:H5'	2.20	0.42
22:DA:641:U:C5	22:DA:642:U:C4	3.08	0.42
7:AG:22:LEU:HD21	7:AG:97:ASN:ND2	2.35	0.42
26:DE:150:THR:OG1	26:DE:151:GLY:N	2.53	0.42
22:BA:2438:U:O2'	22:BA:2439:A:H5''	2.20	0.42
1:CA:853:C:H2'	1:CA:854:U:H6	1.84	0.42
1:CA:468:A:N3	1:CA:468:A:O4'	2.52	0.42
1:AA:1260:G:H4'	1:AA:1283:U:O2'	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:BH:90:LEU:CD2	29:BH:122:LEU:HA	2.49	0.42
39:BR:51:VAL:CB	39:BR:52:PRO:HD2	2.49	0.42
29:BH:91:PHE:CD1	1:CA:55:A:C8	3.07	0.42
22:DA:2234:G:N1	22:DA:2235:G:C5	2.87	0.42
22:DA:1345:C:H5'	22:DA:1396:U:C4	2.54	0.42
11:CK:126:LYS:C	21:CU:34:ARG:CZ	2.88	0.42
22:DA:28:A:C2	22:DA:513:A:C8	3.08	0.42
22:DA:250:G:H2'	22:DA:251:A:C8	2.54	0.42
21:CU:40:LYS:N	21:CU:41:PRO:CD	2.83	0.42
21:CU:41:PRO:O	21:CU:42:THR:C	2.57	0.42
22:DA:724:U:C4	22:DA:725:G:C5	3.08	0.42
26:BE:180:LEU:HD23	26:BE:180:LEU:HA	1.92	0.42
22:DA:571:U:C4	22:DA:2030:A:C6	3.08	0.42
4:CD:150:LYS:HG2	4:CD:151:LYS:N	2.34	0.42
8:AH:64:LYS:HB3	8:AH:64:LYS:HE2	1.87	0.42
24:DC:45:ASN:C	24:DC:47:GLY:N	2.72	0.42
1:AA:71:A:H2'	1:AA:71:A:OP2	2.20	0.42
22:BA:58:G:O2'	22:BA:73:A:N1	2.44	0.42
24:BC:105:LEU:HD22	24:BC:143:ASN:ND2	2.34	0.42
40:BS:96:ILE:HD12	40:BS:98:LYS:HG3	2.01	0.42
22:BA:1734:G:H2'	22:BA:1735:A:H8	1.84	0.42
22:DA:1857:G:C2	22:DA:1884:G:C4	3.08	0.42
22:DA:2344:U:OP1	49:D1:37:LYS:HD2	2.19	0.42
49:D1:39:PHE:CE2	49:D1:40:ASP:O	2.73	0.42
1:CA:1053:G:H4'	1:CA:1055:A:OP1	2.20	0.42
22:DA:2069:G:C2	22:DA:2443:C:C2	3.07	0.42
22:DA:204:A:C8	22:DA:206:U:C2	3.08	0.42
1:CA:775:G:C2'	1:CA:776:G:H5'	2.50	0.42
22:BA:1343:G:C4	22:BA:1344:U:C5	3.07	0.42
15:CO:55:GLY:O	15:CO:59:MET:HG3	2.20	0.42
22:DA:78:U:OP2	46:DY:2:LYS:HD2	2.19	0.42
30:BI:80:LEU:HA	30:BI:84:ALA:HB3	2.02	0.42
32:DK:61:VAL:O	32:DK:61:VAL:HG13	2.19	0.42
4:AD:37:ALA:HA	4:AD:42:GLY:HA3	2.01	0.42
4:AD:58:LYS:HG2	4:AD:203:LEU:CD2	2.49	0.42
22:DA:1864:U:H2'	22:DA:1865:U:H5'	2.00	0.42
22:BA:493:G:H2'	22:BA:494:G:O4'	2.20	0.42
26:DE:108:ILE:HB	33:DL:2:ARG:HH22	1.84	0.42
42:DU:14:LEU:HD11	42:DU:71:ALA:HB2	2.01	0.42
1:CA:1311:A:C2	1:CA:1327:C:N3	2.87	0.42
1:CA:115:G:H4'	1:CA:116:A:O5'	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:AM:88:GLY:O	13:AM:90:ARG:N	2.52	0.42
26:DE:23:PHE:CG	26:DE:111:GLU:HG3	2.54	0.42
22:DA:2446:G:C6	22:DA:2501:C:H2'	2.55	0.42
39:DR:25:LEU:O	39:DR:66:HIS:HE1	2.03	0.42
1:AA:801:U:H2'	1:AA:802:A:C8	2.55	0.42
16:CP:2:VAL:CG2	16:CP:65:ALA:HA	2.49	0.42
22:BA:1:G:H2'	22:BA:2:G:C8	2.54	0.42
1:CA:1228:C:H5'	13:CM:113:ARG:HB2	2.01	0.42
22:BA:447:A:OP2	57:BA:3210:HOH:O	2.21	0.42
9:CI:88:MET:HB2	9:CI:92:GLU:OE2	2.19	0.42
9:CI:88:MET:HG2	9:CI:88:MET:O	2.19	0.42
22:DA:1411:U:H2'	22:DA:1412:U:O4'	2.19	0.42
26:BE:83:VAL:O	26:BE:84:THR:C	2.58	0.42
1:CA:78:A:N6	1:CA:79:G:C6	2.88	0.42
22:DA:2548:U:O2'	32:DK:4:GLU:OE1	2.38	0.42
25:DD:22:ILE:HA	25:DD:23:PRO:HD3	1.91	0.42
46:DY:7:ARG:O	46:DY:8:GLU:HG3	2.19	0.42
22:BA:2533:U:H2'	22:BA:2534:A:H5'	2.02	0.42
26:DE:12:LEU:HD23	26:DE:13:THR:N	2.35	0.42
22:BA:115:C:O2'	22:BA:116:C:H5'	2.19	0.42
24:DC:6:CYS:SG	24:DC:16:VAL:HG12	2.59	0.42
8:CH:87:LYS:HG3	8:CH:91:GLU:HB3	2.00	0.42
29:DH:69:ALA:HB2	29:DH:138:VAL:HG12	2.02	0.42
23:BB:116:G:H4'	36:BO:54:VAL:HG13	2.01	0.42
1:AA:146:G:C2	1:AA:177:G:N7	2.88	0.42
22:BA:1403:A:O2'	22:BA:1404:C:H5'	2.19	0.42
22:BA:2082:A:H2'	22:BA:2083:G:O4'	2.19	0.42
24:BC:79:GLU:OE1	24:BC:101:ARG:NE	2.42	0.42
22:DA:2794:C:H2'	22:DA:2795:C:O4'	2.19	0.42
51:B3:4:ILE:CG2	51:B3:63:PRO:HG3	2.50	0.42
1:AA:1366:C:H2'	1:AA:1367:C:C6	2.55	0.42
29:BH:116:ARG:HB3	29:BH:131:SER:O	2.20	0.42
29:BH:89:LYS:O	29:BH:90:LEU:C	2.58	0.42
22:BA:842:U:H2'	22:BA:843:G:O4'	2.19	0.42
22:DA:1606:C:H4'	22:DA:1607:C:H5'	2.02	0.42
1:AA:979:C:C5	1:AA:980:C:C5	3.08	0.42
41:BT:6:ARG:O	41:BT:8:LEU:N	2.53	0.42
1:AA:1225:A:C2	1:AA:1226:C:C4	3.08	0.42
29:DH:121:VAL:O	29:DH:122:LEU:CB	2.67	0.42
29:BH:119:ASN:CB	29:BH:123:ARG:NH1	2.81	0.42
33:DL:63:LYS:CA	51:D3:13:ARG:HG3	2.50	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:1566:A:H5'	24:DC:214:ARG:CZ	2.50	0.42
1:CA:1007:U:H3'	1:CA:1008:U:H5''	2.00	0.42
5:AE:101:GLU:CD	5:AE:101:GLU:O	2.58	0.42
25:BD:99:GLU:O	25:BD:100:LEU:C	2.55	0.42
22:DA:770:G:O4'	22:DA:1379:U:C5	2.72	0.42
22:DA:773:U:O2'	24:DC:48:ARG:HD3	2.19	0.42
1:AA:828:U:C4	1:AA:859:G:C4	3.07	0.42
22:DA:328:U:O3'	42:DU:66:GLN:HG3	2.19	0.42
1:AA:17:U:H2'	1:AA:18:C:C6	2.55	0.42
22:BA:2287:A:C8	22:BA:2289:G:C8	3.07	0.42
22:DA:1441:G:C2	22:DA:1442:U:C5	3.07	0.42
22:BA:2555:U:C5	22:BA:2556:C:C2	3.08	0.42
22:DA:234:U:H2'	22:DA:235:U:H6	1.84	0.42
1:CA:1478:U:C2	1:CA:1479:C:C5	3.07	0.42
1:AA:206:C:H2'	1:AA:207:C:O4'	2.19	0.42
1:CA:1125:U:C5	10:CJ:40:ILE:HD13	2.55	0.42
22:BA:1984:G:C4	22:BA:1985:C:C6	3.08	0.42
22:BA:250:G:P	51:B3:13:ARG:HH12	2.43	0.42
22:DA:775:G:O6	22:DA:787:C:H2'	2.20	0.42
22:DA:1916:A:H2'	22:DA:1917:U:O4'	2.19	0.42
19:AS:65:GLU:H	19:AS:65:GLU:CD	2.22	0.42
21:AU:14:VAL:HG13	21:AU:16:LEU:HD21	2.01	0.42
9:AI:51:PRO:HB3	9:AI:84:THR:HG23	2.00	0.42
22:DA:45:G:N2	22:DA:434:U:N3	2.67	0.42
22:DA:920:A:C5	22:DA:921:C:C5	3.07	0.42
22:BA:196:A:O2'	22:BA:805:G:O6	2.30	0.42
22:BA:2480:C:H2'	22:BA:2481:G:H5'	2.02	0.42
22:DA:1746:A:H2'	22:DA:1747:U:C6	2.55	0.42
22:DA:2852:G:H2'	22:DA:2853:C:C6	2.53	0.42
3:AC:113:ALA:O	3:AC:114:LYS:C	2.57	0.42
27:BF:134:GLU:CB	27:BF:136:ILE:HD12	2.50	0.42
22:BA:487:C:O2	40:BS:53:SER:OG	2.38	0.42
22:DA:265:A:H4'	22:DA:266:G:OP1	2.19	0.42
1:CA:238:A:C2'	1:CA:239:U:H5'	2.48	0.42
38:DQ:39:VAL:O	38:DQ:42:ALA:HB3	2.19	0.42
22:DA:425:G:C6	22:DA:426:C:N4	2.88	0.42
33:DL:135:ILE:HG22	33:DL:140:GLY:HA2	2.00	0.42
23:DB:34:A:N6	23:DB:44:G:O2'	2.49	0.42
22:BA:817:C:H2'	22:BA:818:G:O4'	2.19	0.42
40:BS:38:TYR:CE1	48:B0:28:LEU:HD21	2.54	0.42
22:BA:920:A:C6	22:BA:921:C:C4	3.08	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:AJ:7:ARG:HB2	10:AJ:75:ASP:OD1	2.20	0.42
22:DA:1452:G:H2'	22:DA:1457:U:O4	2.20	0.42
38:DQ:61:TRP:CH2	38:DQ:93:LYS:HB2	2.55	0.42
3:AC:77:ILE:HA	3:AC:84:VAL:CG2	2.50	0.42
12:AL:81:LEU:HD12	12:AL:81:LEU:HA	1.88	0.42
42:DU:51:ALA:O	42:DU:52:LEU:HB2	2.18	0.42
40:DS:33:LEU:CD2	40:DS:51:LEU:HD23	2.50	0.42
22:BA:749:A:C6	22:BA:1618:A:C2	3.07	0.42
22:DA:2618:G:C5	22:DA:2619:C:C5	3.08	0.42
36:BO:109:ALA:O	36:BO:112:GLU:HB2	2.20	0.42
1:CA:357:G:C2	1:CA:358:U:C6	3.08	0.42
1:CA:1162:C:C2	1:CA:1175:G:N2	2.88	0.42
29:DH:41:LYS:HE2	29:DH:44:ILE:CD1	2.50	0.42
22:BA:1483:G:C6	22:BA:1484:U:C4	3.07	0.42
1:CA:1128:C:N3	1:CA:1145:A:N1	2.67	0.42
25:DD:12:THR:HG21	37:DP:5:ILE:HG23	2.02	0.42
22:BA:1778:U:H2'	22:BA:1784:A:N6	2.34	0.42
1:AA:374:A:N3	1:AA:375:U:C6	2.88	0.42
22:DA:2883:A:P	48:D0:49:TYR:HH	2.38	0.42
22:DA:1851:U:C2	22:DA:1891:G:O6	2.72	0.42
46:BY:21:LEU:O	46:BY:22:LEU:O	2.37	0.42
22:BA:589:U:H2'	22:BA:590:A:C8	2.55	0.42
4:AD:23:SER:HB2	4:AD:110:THR:HB	2.01	0.42
29:DH:31:VAL:HB	29:DH:32:PRO:HD2	2.00	0.42
22:DA:1742:U:C4	22:DA:1743:G:C6	3.08	0.42
33:BL:82:LEU:CG	33:BL:90:VAL:HG21	2.49	0.42
1:AA:933:G:OP2	7:AG:3:ARG:CB	2.67	0.42
22:DA:2756:U:H1'	22:DA:2757:A:H5''	2.01	0.42
11:AK:67:ALA:HB3	11:AK:99:ALA:HB3	2.02	0.42
22:DA:2204:G:C5	22:DA:2221:G:N2	2.88	0.42
5:AE:133:PRO:O	5:AE:137:VAL:HG12	2.19	0.42
22:BA:1876:A:C2	22:BA:1877:A:C4	3.08	0.42
22:DA:11:C:C3'	22:DA:12:U:H5'	2.49	0.42
22:DA:1042:G:C5	22:DA:1043:C:C4	3.07	0.42
22:DA:699:A:H2'	22:DA:700:G:C5'	2.49	0.42
1:AA:1124:G:H3'	1:AA:1145:A:N6	2.35	0.42
30:DI:80:LEU:HD13	30:DI:136:MET:SD	2.60	0.42
1:CA:445:G:H2'	1:CA:445:G:N3	2.34	0.42
45:DX:21:ALA:HB3	45:DX:23:ASN:OD1	2.20	0.42
41:BT:17:SER:O	41:BT:20:ALA:N	2.52	0.42
10:CJ:84:VAL:O	10:CJ:88:MET:HG2	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:1442:U:H2'	22:BA:1443:U:C6	2.54	0.42
22:DA:2370:G:H4'	49:D1:44:ARG:NH1	2.35	0.42
22:DA:2899:A:H2'	22:DA:2900:A:H8	1.83	0.42
22:DA:2250:G:C2	34:DM:82:MET:CB	3.02	0.42
22:DA:2311:A:C2	27:DF:85:ILE:HD11	2.54	0.42
22:DA:2712:C:C2	22:DA:2715:C:OP1	2.73	0.42
1:AA:620:C:H1'	4:AD:132:ILE:HD11	2.01	0.42
26:BE:23:PHE:CD1	26:BE:111:GLU:HG3	2.55	0.42
1:CA:815:A:H4'	1:CA:817:C:C4	2.54	0.42
31:DJ:36:LEU:O	31:DJ:121:LYS:NZ	2.53	0.42
3:AC:14:ILE:O	3:AC:16:LYS:N	2.50	0.42
1:AA:1477:U:H2'	1:AA:1478:U:C6	2.54	0.42
22:DA:2528:U:O2'	22:DA:2529:G:H3'	2.19	0.42
22:BA:2093:G:O2'	22:BA:2094:A:H5'	2.19	0.42
22:DA:2100:G:O6	22:DA:2101:A:C6	2.72	0.42
2:AB:28:LYS:N	2:AB:29:PRO:CD	2.82	0.42
22:BA:1352:U:H2'	22:BA:1353:A:H5'	2.02	0.42
22:DA:416:U:H2'	22:DA:417:C:C6	2.55	0.42
37:DP:48:ILE:HD13	37:DP:62:ARG:HB2	2.01	0.42
5:CE:92:SER:OG	5:CE:130:SER:O	2.33	0.42
3:CC:16:LYS:HG3	3:CC:17:PRO:HD2	2.00	0.42
22:BA:1381:G:H1'	22:BA:1571:A:N1	2.34	0.42
5:CE:155:ALA:HB1	8:CH:66:PHE:CE2	2.54	0.42
41:DT:21:SER:O	41:DT:23:ALA:N	2.52	0.42
22:DA:738:G:O2'	22:DA:739:A:H5'	2.20	0.42
1:AA:687:A:C2	1:AA:700:G:N3	2.88	0.42
11:CK:113:VAL:HB	18:CR:73:ARG:NH2	2.35	0.42
8:CH:125:ILE:O	8:CH:125:ILE:HG12	2.18	0.42
40:DS:45:VAL:HG12	40:DS:45:VAL:O	2.20	0.42
22:BA:2899:A:H2'	22:BA:2900:A:C8	2.54	0.42
1:CA:1406:U:H2'	1:CA:1407:C:O4'	2.19	0.42
52:B4:7:VAL:O	52:B4:8:LYS:HG3	2.20	0.42
17:CQ:15:ASP:HA	17:CQ:21:ILE:HD12	2.01	0.42
1:CA:358:U:H2'	1:CA:359:G:C8	2.54	0.42
1:AA:980:C:H2'	1:AA:981:U:H5'	2.02	0.42
22:DA:1389:G:C2	22:DA:1390:U:O2	2.72	0.42
22:DA:451:U:C2	22:DA:453:A:N7	2.88	0.42
22:DA:1809:A:H2'	22:DA:1810:A:H8	1.83	0.42
50:D2:43:THR:O	50:D2:44:VAL:HB	2.20	0.42
1:CA:505:G:C5'	1:CA:534:U:C2	3.03	0.42
22:DA:511:U:OP2	22:DA:512:G:N7	2.53	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:DL:54:GLN:HG2	33:DL:55:MET:N	2.35	0.42
1:CA:527:G:C2	1:CA:528:C:C5	3.07	0.42
1:CA:688:G:C5	1:CA:700:G:C2	3.08	0.42
22:DA:1974:C:H2'	22:DA:1975:G:H8	1.85	0.42
22:DA:537:G:N1	22:DA:555:G:C2	2.88	0.42
1:CA:1105:A:C2	1:CA:1106:G:N7	2.88	0.42
22:DA:599:A:N1	22:DA:600:G:C6	2.88	0.42
4:CD:36:GLN:O	4:CD:37:ALA:HB2	2.20	0.42
51:B3:31:HIS:CD2	51:B3:31:HIS:C	2.93	0.42
1:AA:827:U:H5''	1:AA:828:U:OP2	2.20	0.42
22:DA:1817:G:O2'	22:DA:1818:U:H5'	2.20	0.42
1:CA:756:C:H2'	1:CA:757:U:C5'	2.50	0.42
22:BA:2286:G:C4'	22:BA:2287:A:O5'	2.67	0.42
42:BU:72:ILE:N	42:BU:72:ILE:CD1	2.81	0.42
1:AA:709:U:H2'	1:AA:710:G:H8	1.85	0.42
35:BN:114:GLU:OE2	35:BN:118:ARG:NH2	2.49	0.42
9:AI:57:MET:SD	9:AI:57:MET:N	2.93	0.42
22:DA:1304:A:C5	22:DA:1305:C:C5	3.08	0.42
1:CA:15:G:O4'	5:CE:29:ARG:NH2	2.53	0.42
5:AE:89:HIS:CE1	5:AE:138:ARG:HD3	2.55	0.42
13:AM:107:ARG:HH21	13:AM:113:ARG:HB3	1.84	0.42
1:AA:473:U:N3	1:AA:474:G:N7	2.67	0.42
1:AA:1296:C:H5''	1:AA:1297:G:OP2	2.19	0.42
22:DA:2516:A:N6	22:DA:2517:C:H42	2.16	0.42
32:DK:38:ILE:HD12	32:DK:87:LEU:CD1	2.50	0.42
29:DH:127:GLU:HA	29:DH:144:VAL:O	2.19	0.42
1:AA:81:A:C2'	1:AA:82:G:H5'	2.50	0.42
4:AD:58:LYS:HB3	4:AD:200:ILE:HG22	2.02	0.42
22:DA:415:A:C2	22:DA:2409:G:C6	3.07	0.42
22:DA:415:A:C2	22:DA:2409:G:N1	2.87	0.42
16:CP:16:PHE:HE1	16:CP:38:PHE:HB2	1.84	0.42
33:DL:82:LEU:O	33:DL:120:VAL:HG21	2.20	0.42
8:AH:5:ASP:OD2	8:AH:8:ALA:HB2	2.20	0.42
25:BD:62:LYS:HB2	25:BD:63:PRO:HD3	2.02	0.42
1:AA:1003:G:N2	1:AA:1005:A:H5'	2.35	0.42
22:DA:1220:G:H2'	22:DA:1221:C:C6	2.54	0.42
22:BA:189:G:O6	22:BA:205:G:O2'	2.26	0.42
22:BA:322:A:H5'	22:BA:340:A:H1'	2.01	0.42
28:BG:2:SER:C	28:BG:4:VAL:H	2.22	0.42
1:CA:235:C:H2'	1:CA:236:A:C8	2.55	0.42
39:DR:3:ALA:HB3	39:DR:101:ILE:HD12	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:DU:83:VAL:HG11	42:DU:94:ARG:HD2	2.00	0.42
10:AJ:26:VAL:HG12	10:AJ:30:LYS:CE	2.50	0.42
1:CA:1040:U:H2'	1:CA:1041:G:C8	2.54	0.42
22:DA:1599:U:C4	22:DA:1600:C:C4	3.08	0.42
1:AA:929:G:H2'	1:AA:930:C:C6	2.55	0.42
30:BI:83:ALA:HB1	30:BI:109:ILE:HD13	2.01	0.42
1:AA:346:G:C8	37:BP:37:LYS:HE2	2.55	0.42
22:BA:545:U:O2	22:BA:545:U:O5'	2.36	0.42
8:CH:77:ARG:NE	8:CH:79:SER:O	2.52	0.42
34:DM:106:ASP:OD1	34:DM:107:GLY:N	2.52	0.42
1:CA:522:C:H41	12:CL:50:ARG:NH2	2.18	0.42
22:BA:2018:G:O2'	22:BA:2019:A:H5'	2.20	0.42
41:BT:67:VAL:HG22	41:BT:76:ARG:HG3	2.01	0.42
17:AQ:46:VAL:HG13	17:AQ:73:TRP:O	2.20	0.42
24:DC:93:LEU:HD13	24:DC:103:TYR:CE1	2.54	0.42
1:CA:1279:G:O2'	1:CA:1282:C:N4	2.53	0.42
22:BA:1842:G:C6	22:BA:1843:C:C4	3.08	0.42
22:BA:2795:C:H2'	22:BA:2796:U:C6	2.55	0.42
3:AC:60:PRO:O	3:AC:61:ALA:O	2.37	0.42
13:AM:44:LYS:HE2	13:AM:44:LYS:HB3	1.94	0.42
22:BA:796:C:H2'	22:BA:797:G:C8	2.55	0.42
22:BA:989:G:C8	47:BZ:14:ILE:HD11	2.55	0.42
24:DC:9:THR:O	24:DC:10:SER:HB3	2.19	0.42
22:BA:1610:A:C4'	22:BA:1611:C:OP2	2.68	0.42
22:BA:15:G:O2'	22:BA:16:C:H5'	2.20	0.42
29:BH:82:SER:HG	29:BH:90:LEU:HG	1.85	0.42
1:CA:1097:C:OP1	2:CB:139:ARG:NH2	2.53	0.42
22:BA:1924:C:H2'	22:BA:1926:U:O4	2.19	0.42
31:BJ:55:ILE:HG21	31:BJ:130:HIS:HD2	1.83	0.42
22:DA:811:U:O2	22:DA:1251:C:C6	2.73	0.42
22:DA:1340:U:C5	22:DA:1603:A:C8	3.07	0.42
7:AG:135:VAL:HB	7:AG:138:ARG:HH21	1.85	0.42
34:DM:76:LYS:O	34:DM:77:PRO:O	2.37	0.42
5:CE:154:ALA:C	5:CE:156:LYS:N	2.73	0.42
1:AA:108:G:N2	1:AA:109:A:C2	2.88	0.42
13:AM:11:ASP:CG	13:AM:12:HIS:N	2.74	0.42
22:DA:804:A:C2'	22:DA:806:C:C4	3.03	0.42
26:BE:108:ILE:CD1	26:BE:180:LEU:HB3	2.44	0.42
4:CD:123:ILE:HG22	4:CD:124:MET:N	2.34	0.42
22:DA:2428:G:H5''	22:DA:2429:G:OP1	2.20	0.42
2:CB:21:ARG:C	2:CB:23:TRP:N	2.73	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:674:G:C2	1:CA:675:A:C5	3.08	0.42
20:AT:5:LYS:O	20:AT:6:SER:C	2.57	0.42
1:AA:868:C:H2'	1:AA:869:G:C5'	2.50	0.42
1:CA:728:A:C8	15:CO:54:ARG:CZ	3.03	0.42
29:BH:104:THR:CG2	29:BH:110:VAL:O	2.68	0.42
1:AA:1239:A:H62	1:AA:1299:A:N6	2.18	0.42
22:BA:2553:G:H5''	22:BA:2554:U:OP2	2.20	0.42
18:AR:22:ASP:OD2	18:AR:24:LYS:N	2.50	0.42
1:CA:223:A:C4	1:CA:224:U:C6	3.08	0.42
9:AI:43:THR:O	9:AI:44:ALA:CB	2.67	0.42
22:DA:2290:G:C5	22:DA:2291:U:C4	3.08	0.42
1:CA:35:G:N2	12:CL:115:SER:OG	2.53	0.42
1:AA:1061:G:C5	1:AA:1062:U:C5	3.08	0.42
22:DA:732:C:C5	22:DA:733:G:N7	2.88	0.42
39:DR:52:PRO:C	39:DR:53:PHE:CG	2.90	0.42
22:DA:40:U:H2'	22:DA:41:C:C6	2.55	0.42
34:BM:132:THR:HG22	34:BM:133:LYS:H	1.83	0.42
22:BA:1002:G:O6	57:BA:3746:HOH:O	2.22	0.42
22:BA:83:A:N6	22:BA:101:A:C5	2.88	0.42
5:AE:151:GLU:HG2	5:AE:152:MET:H	1.85	0.42
22:DA:2364:C:H2'	22:DA:2365:G:O4'	2.20	0.42
43:BV:10:LYS:NZ	43:BV:41:GLU:OE2	2.52	0.42
1:CA:714:G:H21	1:CA:777:A:H1'	1.84	0.42
41:BT:89:GLU:O	41:BT:91:GLN:HG2	2.19	0.42
1:CA:681:A:C2	1:CA:710:G:C2	3.08	0.42
7:CG:46:ALA:CA	7:CG:121:ALA:HB2	2.50	0.42
17:CQ:79:VAL:O	17:CQ:80:GLU:CB	2.67	0.42
45:DX:40:VAL:HG11	45:DX:68:LEU:CD1	2.49	0.42
23:BB:110:C:C4	23:BB:111:U:C5	3.08	0.42
1:CA:855:U:H2'	1:CA:856:C:C6	2.55	0.42
22:DA:1624:U:N3	22:DA:1625:C:C5	2.88	0.42
21:CU:14:VAL:O	21:CU:16:LEU:HG	2.19	0.42
39:BR:61:ALA:HB2	39:BR:98:ILE:HD13	2.02	0.42
22:DA:2104:C:H2'	22:DA:2105:U:C6	2.55	0.42
22:BA:901:C:C4	22:BA:902:C:C5	3.08	0.42
22:BA:1265:A:O4'	22:BA:1267:U:C6	2.73	0.42
53:B5:59:VAL:HG21	53:B5:167:ASP:C	2.40	0.42
1:AA:999:C:N4	1:AA:1000:A:N6	2.68	0.42
40:BS:38:TYR:CD1	48:B0:28:LEU:HD21	2.55	0.42
22:DA:495:G:H4'	40:DS:4:ILE:O	2.20	0.42
46:BY:5:GLU:HA	46:BY:8:GLU:HG3	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1247:U:O2'	1:AA:1248:A:H5'	2.20	0.42
22:BA:1296:G:OP1	22:BA:2709:G:O2'	2.32	0.42
1:AA:1496:C:H2'	1:AA:1497:G:O4'	2.20	0.42
40:BS:8:ARG:O	40:BS:9:HIS:HB2	2.20	0.42
1:CA:643:C:H5'	8:CH:32:LEU:HD13	2.02	0.42
2:AB:206:ALA:O	2:AB:210:VAL:HG22	2.19	0.42
22:BA:1263:U:O2'	48:B0:8:PRO:HD2	2.20	0.42
49:B1:39:PHE:HB2	49:B1:46:HIS:CE1	2.55	0.42
22:BA:2142:A:H2'	22:BA:2143:C:C6	2.55	0.42
1:CA:380:G:N2	1:CA:383:A:OP2	2.48	0.42
23:BB:59:A:H2'	23:BB:60:C:O4'	2.20	0.42
22:BA:2153:C:H2'	22:BA:2154:A:O4'	2.19	0.42
42:BU:73:PHE:CZ	42:BU:78:GLY:HA2	2.55	0.42
41:DT:2:ILE:HA	41:DT:3:ARG:CB	2.49	0.42
22:DA:1025:G:H4'	22:DA:1026:G:OP2	2.20	0.42
22:BA:246:C:C2'	22:BA:247:G:H5'	2.50	0.42
6:CF:29:ILE:CG2	6:CF:34:GLY:O	2.68	0.42
2:AB:161:LEU:CD1	2:AB:176:ALA:HB2	2.50	0.42
2:CB:222:ARG:NE	2:CB:223:GLU:HB2	2.35	0.42
29:BH:139:PHE:O	29:BH:140:ALA:HB3	2.20	0.41
12:AL:23:ALA:O	12:AL:24:LEU:O	2.38	0.41
22:BA:1269:A:OP2	57:BA:3387:HOH:O	2.21	0.41
2:CB:103:ASN:HD22	2:CB:103:ASN:N	2.16	0.41
22:BA:1090:A:C2'	22:BA:1091:G:H5'	2.50	0.41
22:BA:2659:G:OP1	28:BG:158:LYS:HE3	2.20	0.41
17:AQ:12:VAL:O	17:AQ:13:VAL:CB	2.68	0.41
22:BA:271:G:H4'	22:BA:272:A:OP1	2.19	0.41
22:DA:191:A:N6	22:DA:192:C:N4	2.67	0.41
45:BX:4:VAL:N	45:BX:33:LEU:HD11	2.35	0.41
4:AD:192:SER:O	4:AD:194:ASP:N	2.46	0.41
21:CU:34:ARG:CD	21:CU:35:ARG:HB2	2.49	0.41
1:CA:1222:G:C6	1:CA:1223:C:C4	3.08	0.41
8:AH:42:GLU:OE1	8:AH:42:GLU:CA	2.68	0.41
13:AM:11:ASP:O	13:AM:12:HIS:CB	2.68	0.41
22:DA:410:G:H2'	22:DA:2407:A:C8	2.55	0.41
22:BA:1022:G:N2	22:BA:1142:A:C2	2.77	0.41
22:DA:2392:A:OP2	51:D3:31:HIS:CE1	2.73	0.41
42:DU:82:ARG:HB2	42:DU:97:LYS:HB2	2.02	0.41
22:DA:599:A:H1'	22:DA:659:G:N2	2.35	0.41
4:AD:11:LEU:HD13	4:AD:63:ARG:HD3	2.02	0.41
1:CA:735:C:H2'	1:CA:736:C:C6	2.54	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:AD:124:MET:SD	4:AD:127:GLY:O	2.78	0.41
2:AB:50:PHE:HA	2:AB:53:ALA:HB3	2.02	0.41
22:BA:44:A:H2'	22:BA:45:G:O4'	2.19	0.41
36:BO:53:THR:HG23	36:BO:74:VAL:HG21	2.02	0.41
4:CD:102:VAL:HG13	4:CD:107:PHE:HB2	2.01	0.41
23:DB:23:G:C2	23:DB:24:G:O6	2.73	0.41
2:CB:207:ILE:HG12	2:CB:208:ARG:N	2.35	0.41
22:DA:1430:G:H2'	22:DA:1431:A:O4'	2.19	0.41
1:CA:1219:A:N6	1:CA:1220:G:C6	2.88	0.41
15:CO:17:ARG:O	15:CO:18:ASP:CB	2.68	0.41
1:AA:1500:A:OP2	1:AA:1505:G:OP1	2.38	0.41
1:CA:1462:C:H2'	1:CA:1463:U:C6	2.55	0.41
5:AE:109:GLY:O	5:AE:110:ALA:CB	2.66	0.41
36:DO:33:ARG:HG2	36:DO:34:HIS:CD2	2.55	0.41
1:CA:775:G:C6	1:CA:776:G:N7	2.88	0.41
22:DA:1462:C:H2'	22:DA:1463:C:H6	1.84	0.41
22:BA:2478:A:C2'	22:BA:2479:U:H5'	2.50	0.41
1:AA:668:G:O2'	15:AO:46:HIS:CD2	2.73	0.41
1:CA:892:A:C6	1:CA:893:C:C4	3.08	0.41
22:BA:2292:U:H4'	22:BA:2375:G:H4'	2.02	0.41
1:AA:666:G:H5'	1:AA:726:C:H1'	2.02	0.41
1:AA:545:C:H5'	4:AD:69:GLU:HG3	2.00	0.41
22:DA:415:A:O2'	22:DA:1865:U:H5''	2.20	0.41
1:AA:624:C:C4	1:AA:625:U:C5	3.08	0.41
1:AA:66:A:O4'	1:AA:173:U:C4	2.73	0.41
9:AI:12:ARG:O	9:AI:13:LYS:C	2.58	0.41
30:BI:101:ILE:HD11	30:BI:138:LEU:HD13	2.02	0.41
13:AM:91:HIS:HA	13:AM:109:ARG:NH2	2.34	0.41
4:AD:88:GLU:O	4:AD:91:LEU:N	2.52	0.41
1:CA:1343:G:C6	1:CA:1344:C:N4	2.88	0.41
27:BF:136:ILE:HG21	27:BF:143:TYR:CD1	2.54	0.41
22:BA:547:A:C8	22:BA:548:G:N3	2.88	0.41
35:DN:33:ILE:HD12	35:DN:118:ARG:CZ	2.49	0.41
22:DA:2527:C:C4	22:DA:2528:U:C5	3.08	0.41
22:BA:1438:U:C4	22:BA:1552:A:C2	3.08	0.41
28:BG:109:PHE:CE1	28:BG:152:ARG:CZ	3.03	0.41
43:DV:52:ALA:HB3	43:DV:53:LYS:HE2	2.01	0.41
23:BB:33:G:O2'	23:BB:34:A:H5'	2.20	0.41
22:BA:2221:G:O2'	22:BA:2222:C:H5'	2.20	0.41
4:AD:4:TYR:CZ	4:AD:6:GLY:HA3	2.54	0.41
22:BA:307:G:N2	22:BA:309:A:H3'	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:BD:48:ILE:HG23	25:BD:84:LEU:HD21	2.01	0.41
41:BT:50:LEU:HD12	41:BT:50:LEU:HA	1.85	0.41
22:BA:1309:G:H4'	50:B2:7:PRO:HG2	2.02	0.41
22:BA:257:C:H2'	22:BA:258:G:O4'	2.20	0.41
22:BA:2030:A:C2	22:BA:2499:C:H5''	2.55	0.41
22:DA:1009:A:O2'	22:DA:1153:C:H4'	2.20	0.41
29:DH:40:THR:OG1	29:DH:43:ASN:ND2	2.53	0.41
22:DA:1809:A:C4	22:DA:1810:A:C8	3.07	0.41
1:AA:374:A:H5''	1:AA:452:A:C2	2.55	0.41
22:DA:1663:G:C6	22:DA:1992:G:C8	3.07	0.41
1:CA:833:G:C4	1:CA:834:U:C6	3.08	0.41
22:BA:998:C:P	38:BQ:92:ARG:NH2	2.93	0.41
22:BA:370:G:O2'	22:BA:424:G:OP1	2.28	0.41
1:AA:975:A:O4'	1:AA:1358:U:H1'	2.20	0.41
26:BE:104:ALA:O	26:BE:108:ILE:HG23	2.20	0.41
22:DA:1975:G:C5	22:DA:1976:U:C5	3.08	0.41
5:AE:81:LEU:HD21	5:AE:123:VAL:HG12	2.01	0.41
1:CA:247:G:O6	1:CA:278:G:C6	2.73	0.41
1:AA:1277:C:O2'	1:AA:1279:G:C8	2.70	0.41
22:DA:658:U:N3	22:DA:659:G:C8	2.87	0.41
22:DA:659:G:H4'	26:DE:95:LYS:HD3	2.01	0.41
22:DA:772:C:C2	22:DA:773:U:C5	3.08	0.41
22:BA:1326:U:C2'	22:BA:1327:A:H5'	2.50	0.41
33:BL:77:ILE:HD11	33:BL:101:ILE:HG21	2.02	0.41
44:BW:18:ALA:HB3	44:BW:20:ARG:HH21	1.85	0.41
1:CA:734:G:C4	1:CA:735:C:C6	3.08	0.41
22:DA:2164:C:H2'	22:DA:2165:C:H6	1.83	0.41
22:DA:80:G:H4'	22:DA:346:A:H1'	2.02	0.41
22:BA:2406:A:N1	33:BL:69:ARG:NH2	2.67	0.41
1:AA:644:U:O2'	1:AA:645:G:H5'	2.20	0.41
11:CK:112:ASP:HB3	21:CU:4:ILE:HG22	2.00	0.41
1:CA:840:C:C4	1:CA:842:U:H4'	2.55	0.41
22:DA:1140:C:H1'	22:DA:1143:A:N3	2.35	0.41
22:DA:2067:G:C5	22:DA:2444:G:C2	3.08	0.41
22:BA:2857:G:N2	22:BA:2860:A:OP2	2.41	0.41
22:DA:2677:G:C2	22:DA:2731:G:N3	2.88	0.41
30:BI:77:ALA:HB2	30:BI:132:THR:CG2	2.49	0.41
10:CJ:80:THR:O	10:CJ:84:VAL:N	2.52	0.41
1:CA:892:A:C2	1:CA:907:A:C4	3.08	0.41
3:AC:193:TYR:CD2	3:AC:193:TYR:N	2.85	0.41
6:CF:97:THR:O	6:CF:98:GLU:CB	2.68	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:952:U:H2'	1:CA:953:G:C8	2.55	0.41
1:CA:1326:U:H2'	1:CA:1327:C:C6	2.55	0.41
1:AA:268:U:H2'	1:AA:269:C:C6	2.55	0.41
22:DA:971:G:H2'	22:DA:972:A:O4'	2.20	0.41
4:AD:118:VAL:O	4:AD:131:ASN:HA	2.20	0.41
22:BA:1495:A:O2'	22:BA:1496:A:H5'	2.20	0.41
1:CA:1118:U:H1'	1:CA:1179:A:C5	2.55	0.41
22:BA:2328:A:H2'	22:BA:2329:U:C6	2.55	0.41
47:DZ:4:THR:CG2	47:DZ:5:ILE:N	2.83	0.41
22:DA:1422:G:H4'	22:DA:1493:C:OP2	2.19	0.41
39:DR:42:ALA:HA	39:DR:46:GLU:HA	2.01	0.41
30:BI:28:LEU:HD21	30:BI:35:ILE:HG23	2.02	0.41
1:CA:584:G:H2'	1:CA:585:G:H8	1.85	0.41
1:AA:775:G:C2'	1:AA:776:G:H5'	2.50	0.41
6:CF:29:ILE:HG22	6:CF:34:GLY:O	2.19	0.41
15:CO:49:ASP:OD2	15:CO:52:SER:HB2	2.20	0.41
17:AQ:10:GLY:HA3	17:AQ:24:ALA:O	2.19	0.41
22:BA:176:A:C2'	22:BA:177:G:H5'	2.50	0.41
6:CF:8:PHE:CE1	6:CF:60:VAL:HB	2.56	0.41
14:AN:16:LEU:HD23	14:AN:19:LYS:HD2	2.02	0.41
1:AA:588:G:C2	1:AA:589:U:C2	3.08	0.41
22:BA:1786:A:C4	22:BA:1938:A:C6	3.08	0.41
23:BB:57:A:C4	27:BF:26:MET:HB3	2.55	0.41
22:BA:954:G:C5	22:BA:955:U:C5	3.08	0.41
1:AA:500:G:C6	1:AA:501:C:N4	2.88	0.41
22:BA:125:A:OP2	50:B2:19:ARG:NH2	2.53	0.41
22:DA:1921:G:N1	22:DA:1922:G:C5	2.87	0.41
9:AI:22:LYS:HB3	9:AI:22:LYS:HE3	1.91	0.41
22:BA:2211:A:OP2	22:BA:2211:A:H4'	2.19	0.41
31:BJ:114:LEU:HD12	31:BJ:114:LEU:O	2.20	0.41
4:AD:89:ASN:O	4:AD:93:LEU:HB2	2.20	0.41
1:CA:219:U:H2'	1:CA:220:G:C8	2.55	0.41
13:CM:4:ILE:HG22	13:CM:53:ILE:HG23	2.02	0.41
24:DC:57:GLY:O	24:DC:58:HIS:O	2.38	0.41
8:AH:22:LYS:O	8:AH:65:TYR:OH	2.34	0.41
20:CT:44:LYS:HD3	20:CT:87:ALA:HA	2.01	0.41
22:BA:1924:C:H2'	22:BA:1925:C:H5''	2.02	0.41
22:BA:1510:G:H2'	22:BA:1511:G:O4'	2.21	0.41
18:CR:22:ASP:OD2	18:CR:24:LYS:NZ	2.45	0.41
1:AA:923:A:C6	1:AA:924:C:C4	3.08	0.41
2:AB:164:ILE:HG23	2:AB:165:ASP:N	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:CU:18:ARG:O	21:CU:19:PHE:C	2.58	0.41
22:DA:1526:C:H2'	22:DA:1527:G:C8	2.55	0.41
22:DA:1545:A:H2'	22:DA:1546:G:H5'	2.03	0.41
1:CA:933:G:OP2	7:CG:3:ARG:HB3	2.20	0.41
22:DA:2420:C:C4	51:D3:31:HIS:HB3	2.55	0.41
20:AT:3:ASN:ND2	20:AT:3:ASN:C	2.74	0.41
22:DA:2521:C:C2	22:DA:2545:G:C2	3.08	0.41
8:AH:18:GLN:C	8:AH:20:ALA:N	2.72	0.41
1:AA:1464:U:OP2	37:BP:109:ARG:NH1	2.48	0.41
36:BO:80:GLU:O	36:BO:81:ARG:C	2.57	0.41
22:DA:483:A:H4'	42:DU:48:PRO:HD3	2.03	0.41
22:BA:2114:A:N3	22:BA:2114:A:C2'	2.83	0.41
13:AM:69:LEU:O	13:AM:73:ILE:HG13	2.20	0.41
22:DA:270:A:N1	22:DA:369:U:H1'	2.35	0.41
31:DJ:30:THR:HG22	31:DJ:31:GLU:N	2.35	0.41
22:DA:2321:U:H5'	22:DA:2322:A:OP2	2.20	0.41
26:BE:181:ILE:HG22	33:BL:2:ARG:HB3	2.02	0.41
1:CA:1227:A:OP2	13:CM:110:LYS:CE	2.68	0.41
39:DR:52:PRO:O	39:DR:53:PHE:HB2	2.20	0.41
1:AA:258:G:C4	1:AA:259:G:C8	3.08	0.41
2:AB:168:HIS:ND1	2:AB:168:HIS:O	2.53	0.41
4:AD:90:LEU:HD12	4:AD:90:LEU:O	2.20	0.41
1:AA:556:C:O2'	1:AA:557:G:H5'	2.20	0.41
7:AG:31:MET:HG3	7:AG:32:VAL:H	1.85	0.41
22:DA:973:A:OP2	39:DR:81:LYS:HD2	2.20	0.41
6:CF:6:ILE:HD13	6:CF:62:MET:HG2	2.01	0.41
15:AO:2:SER:O	15:AO:3:LEU:HB2	2.19	0.41
22:BA:495:G:C1'	40:BS:57:ASN:ND2	2.83	0.41
1:CA:741:G:H2'	1:CA:742:G:O4'	2.20	0.41
22:BA:391:A:C5	22:BA:392:U:C5	3.08	0.41
22:DA:693:A:C5	22:DA:694:U:C5	3.08	0.41
22:DA:851:C:H2'	22:DA:852:U:C6	2.55	0.41
1:CA:121:U:H3'	1:CA:122:G:C5'	2.49	0.41
46:DY:17:GLU:HB2	46:DY:53:VAL:HG11	2.02	0.41
27:BF:36:LEU:HD11	27:BF:154:ILE:HD13	2.02	0.41
38:DQ:112:LYS:HB2	39:DR:48:LYS:HD2	2.03	0.41
1:CA:147:G:H2'	1:CA:148:G:C8	2.55	0.41
1:CA:242:G:N2	1:CA:285:C:C2	2.88	0.41
19:AS:41:PHE:HB3	19:AS:43:ASN:OD1	2.20	0.41
22:BA:819:A:C4	22:BA:1189:A:C2	3.08	0.41
22:BA:819:A:H2'	22:BA:820:A:H5'	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:AJ:57:VAL:O	10:AJ:58:ASN:HB2	2.19	0.41
1:AA:753:A:H4'	1:AA:754:C:O5'	2.21	0.41
22:DA:1990:C:H2'	22:DA:1991:U:O4'	2.20	0.41
35:DN:73:ASN:O	35:DN:76:VAL:HG12	2.20	0.41
3:AC:47:LEU:O	3:AC:52:VAL:HG12	2.20	0.41
22:DA:1445:G:N2	22:DA:1547:C:C2	2.88	0.41
22:DA:2692:G:O4'	22:DA:2846:G:N2	2.53	0.41
22:BA:857:G:N3	44:BW:26:PHE:HE2	2.18	0.41
22:BA:950:G:C6	22:BA:951:C:C4	3.08	0.41
22:DA:1653:G:O6	35:DN:11:ASN:N	2.48	0.41
30:BI:17:MET:HA	30:BI:17:MET:HE2	2.02	0.41
1:AA:559:A:H2'	1:AA:559:A:N3	2.35	0.41
37:BP:80:VAL:O	37:BP:80:VAL:HG12	2.20	0.41
1:CA:406:G:C2	1:CA:407:U:C5	3.08	0.41
1:AA:1450:U:H2'	1:AA:1452:C:C5	2.55	0.41
23:DB:74:U:O2	43:DV:29:ILE:CD1	2.68	0.41
29:BH:90:LEU:HD13	29:BH:125:THR:HA	2.03	0.41
12:AL:24:LEU:O	12:AL:25:GLU:C	2.58	0.41
22:BA:1917:U:H2'	22:BA:1917:U:O2	2.20	0.41
28:BG:155:GLU:OE2	28:BG:157:TYR:C	2.56	0.41
22:DA:2230:G:H2'	22:DA:2231:U:C6	2.55	0.41
1:AA:413:G:N1	4:AD:32:CYS:O	2.48	0.41
22:DA:511:U:O4	22:DA:512:G:N1	2.54	0.41
22:DA:184:C:H2'	22:DA:185:G:C8	2.55	0.41
22:DA:1773:A:H2'	22:DA:1774:C:C5'	2.51	0.41
1:CA:995:C:O2	1:CA:1047:G:H5'	2.21	0.41
22:DA:1798:U:H5	24:DC:271:ARG:CZ	2.33	0.41
1:CA:1002:G:H2'	1:CA:1003:G:O4'	2.21	0.41
22:DA:783:A:O2'	22:DA:785:G:OP1	2.38	0.41
1:CA:734:G:N3	1:CA:735:C:C6	2.89	0.41
30:BI:50:GLU:C	30:BI:51:LYS:HD3	2.40	0.41
22:DA:2386:A:H2'	22:DA:2387:U:C6	2.55	0.41
22:BA:338:G:N2	22:BA:339:U:H1'	2.35	0.41
22:DA:299:A:N3	22:DA:319:G:O2'	2.38	0.41
1:CA:582:C:N3	1:CA:760:G:N1	2.68	0.41
22:DA:2291:U:H5''	22:DA:2380:C:O2'	2.19	0.41
22:DA:699:A:H2'	22:DA:700:G:O4'	2.21	0.41
22:BA:1185:G:C4'	22:BA:1186:G:OP1	2.66	0.41
3:CC:150:LYS:HE3	3:CC:201:TRP:CZ3	2.55	0.41
15:CO:37:ASN:O	15:CO:40:GLN:N	2.51	0.41
22:DA:2774:C:N4	22:DA:2775:G:C6	2.88	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:CP:43:ALA:O	16:CP:46:LYS:CG	2.68	0.41
1:AA:1350:A:C6	1:AA:1351:U:N3	2.88	0.41
15:AO:45:GLU:O	15:AO:46:HIS:HB2	2.20	0.41
13:CM:106:ALA:O	13:CM:110:LYS:HB3	2.21	0.41
30:BI:80:LEU:CD1	30:BI:132:THR:O	2.68	0.41
1:CA:76:G:N2	1:CA:95:C:C2	2.89	0.41
22:BA:2557:G:H2'	22:BA:2558:C:H6	1.85	0.41
22:BA:533:G:H5'	38:BQ:24:TYR:CE1	2.56	0.41
33:DL:82:LEU:O	33:DL:82:LEU:HG	2.20	0.41
7:AG:27:VAL:HG23	7:AG:28:ASN:N	2.34	0.41
15:CO:42:HIS:ND1	15:CO:46:HIS:CD2	2.88	0.41
22:DA:379:G:N2	22:DA:380:G:H1'	2.36	0.41
22:DA:2201:G:C4	22:DA:2202:U:C5	3.08	0.41
30:BI:97:LYS:HB3	30:BI:139:VAL:HG22	2.01	0.41
52:D4:4:ARG:O	52:D4:37:GLN:NE2	2.53	0.41
22:BA:854:C:O2	22:BA:924:G:C2	2.74	0.41
22:BA:2771:C:H2'	22:BA:2772:C:C6	2.55	0.41
20:AT:69:LYS:HB2	20:AT:69:LYS:HZ3	1.84	0.41
30:DI:54:PRO:HG2	30:DI:78:VAL:HG21	2.02	0.41
3:AC:42:TYR:OH	3:AC:90:VAL:HG21	2.20	0.41
12:CL:58:THR:CG2	12:CL:59:ASN:N	2.83	0.41
6:AF:46:GLN:HB2	6:AF:56:LYS:HE2	2.02	0.41
11:AK:110:ILE:HG22	11:AK:111:THR:N	2.35	0.41
4:CD:53:VAL:HG23	4:CD:54:GLN:N	2.35	0.41
22:DA:282:A:C2	22:DA:359:G:C2	3.08	0.41
36:DO:7:ARG:CZ	36:DO:97:PHE:CZ	3.03	0.41
1:AA:927:G:C2	1:AA:1391:U:O2	2.73	0.41
22:DA:836:G:C5	22:DA:837:C:C5	3.08	0.41
45:BX:68:LEU:HD23	45:BX:68:LEU:HA	1.84	0.41
22:BA:734:A:C5	22:BA:735:A:C8	3.09	0.41
46:DY:17:GLU:OE1	46:DY:50:VAL:HG13	2.20	0.41
9:AI:30:ILE:HA	9:AI:65:ILE:O	2.20	0.41
22:DA:2825:G:C3'	22:DA:2826:A:H5'	2.51	0.41
1:CA:78:A:C6	1:CA:79:G:C5	3.08	0.41
22:BA:2154:A:H2'	22:BA:2155:U:C6	2.56	0.41
10:AJ:57:VAL:HG22	10:AJ:58:ASN:H	1.86	0.41
22:DA:121:G:N3	22:DA:131:A:C2	2.88	0.41
46:DY:36:GLN:O	46:DY:37:LEU:C	2.59	0.41
22:DA:1471:G:N2	22:DA:1521:G:H1'	2.36	0.41
30:BI:103:ARG:HB3	30:BI:142:ASP:O	2.20	0.41
28:DG:2:SER:N	28:DG:5:ALA:HB3	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:CN:48:LEU:HD22	14:CN:48:LEU:O	2.20	0.41
30:DI:97:LYS:HE3	30:DI:97:LYS:HA	2.03	0.41
30:DI:140:VAL:O	30:DI:140:VAL:HG13	2.20	0.41
29:DH:135:HIS:CG	29:DH:136:SER:N	2.89	0.41
43:BV:63:ILE:CD1	43:BV:72:VAL:HG21	2.50	0.41
25:BD:122:VAL:HG12	25:BD:123:LYS:N	2.34	0.41
29:BH:33:GLN:O	29:BH:35:LYS:N	2.53	0.41
12:AL:59:ASN:OD1	12:AL:59:ASN:C	2.58	0.41
22:BA:271:G:C6	22:BA:367:G:C2	3.09	0.41
22:DA:2229:U:H2'	22:DA:2230:G:H8	1.85	0.41
22:DA:201:C:C5	22:DA:202:U:C4	3.08	0.41
1:AA:372:C:H4'	1:AA:373:A:OP1	2.20	0.41
5:CE:100:SER:O	5:CE:101:GLU:C	2.59	0.41
32:BK:76:VAL:HB	37:BP:73:VAL:HG13	2.01	0.41
21:CU:37:PHE:HB3	21:CU:41:PRO:CG	2.50	0.41
2:CB:134:ALA:O	2:CB:138:THR:HG23	2.20	0.41
1:CA:687:A:C5	1:CA:701:U:C5	3.08	0.41
1:AA:201:G:H2'	1:AA:202:G:O4'	2.20	0.41
1:CA:32:A:H3'	1:CA:33:A:H8	1.84	0.41
22:DA:487:C:N4	22:DA:488:G:C6	2.88	0.41
26:DE:24:ASN:ND2	26:DE:27:LEU:HB2	2.35	0.41
25:BD:2:ILE:HG13	25:BD:100:LEU:CD2	2.46	0.41
22:DA:2780:G:C6	31:DJ:102:GLU:OE2	2.73	0.41
1:CA:811:C:N4	1:CA:812:G:C6	2.89	0.41
33:BL:84:LYS:HG3	33:BL:84:LYS:O	2.19	0.41
21:AU:37:PHE:O	21:AU:38:TYR:CB	2.67	0.41
21:AU:38:TYR:O	21:AU:41:PRO:HD2	2.20	0.41
22:DA:1257:C:C4	22:DA:1258:U:C4	3.08	0.41
22:DA:329:G:P	42:DU:66:GLN:HG3	2.61	0.41
25:BD:149:ASN:CG	25:BD:150:GLN:H	2.23	0.41
29:DH:130:VAL:CG1	29:DH:131:SER:N	2.82	0.41
1:CA:1240:U:OP2	7:CG:116:MET:CB	2.68	0.41
30:BI:43:ASN:OD1	30:BI:46:THR:HB	2.21	0.41
22:DA:1313:U:C2'	22:DA:1313:U:O2	2.69	0.41
22:DA:2344:U:H4'	22:DA:2345:G:OP1	2.20	0.41
22:DA:634:C:H2'	22:DA:635:C:C6	2.56	0.41
22:DA:71:A:C2	22:DA:73:A:C2	3.08	0.41
13:AM:66:GLU:HA	13:AM:66:GLU:OE1	2.20	0.41
22:DA:844:A:C2	22:DA:845:A:N7	2.88	0.41
22:DA:833:A:OP2	33:DL:39:LYS:HE3	2.20	0.41
1:CA:682:G:O2'	1:CA:683:G:H5'	2.19	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AB:101:LEU:HD11	2:AB:158:PRO:HG2	2.03	0.41
30:BI:80:LEU:HD13	30:BI:136:MET:SD	2.60	0.41
22:BA:613:A:HO2'	22:BA:614:A:P	2.41	0.41
22:DA:223:A:H2'	22:DA:408:G:N3	2.35	0.41
22:DA:19:A:OP1	38:DQ:22:LYS:NZ	2.53	0.41
22:DA:972:A:N1	22:DA:973:A:N6	2.69	0.41
1:CA:1364:U:O2	1:CA:1364:U:O2'	2.32	0.41
12:CL:4:VAL:HG13	12:CL:5:ASN:H	1.85	0.41
2:AB:61:ALA:HA	2:AB:65:GLY:CA	2.51	0.41
50:D2:39:ARG:HB2	50:D2:42:LEU:HD22	2.03	0.41
22:BA:1577:C:H2'	22:BA:1578:U:C1'	2.51	0.41
1:AA:862:C:N4	1:AA:863:U:O4	2.54	0.41
11:AK:35:THR:OG1	11:AK:40:ASN:C	2.58	0.41
22:DA:2282:G:C2	22:DA:2425:A:C6	3.08	0.41
1:CA:573:A:H2'	1:CA:574:A:C8	2.55	0.41
43:BV:40:ILE:HA	43:BV:40:ILE:HD13	1.80	0.41
1:AA:620:C:H1'	4:AD:132:ILE:CD1	2.50	0.41
42:DU:84:GLY:O	42:DU:94:ARG:HA	2.20	0.41
1:CA:1225:A:C2'	1:CA:1225:A:N3	2.83	0.41
1:AA:19:A:C5	1:AA:20:U:C5	3.09	0.41
41:BT:57:VAL:CG2	41:BT:58:VAL:N	2.83	0.41
42:DU:40:ASN:HB3	42:DU:63:ALA:HB3	2.02	0.41
1:AA:601:G:H2'	1:AA:602:A:C8	2.56	0.41
40:BS:37:THR:OG1	40:BS:48:LYS:NZ	2.54	0.41
41:BT:27:SER:O	41:BT:28:ASN:C	2.59	0.41
36:BO:49:VAL:HG21	36:BO:82:ALA:HA	2.02	0.41
31:BJ:89:PHE:CE1	31:BJ:100:VAL:HG11	2.56	0.41
6:AF:22:ILE:O	6:AF:23:GLU:C	2.56	0.41
3:AC:128:VAL:O	3:AC:129:MET:C	2.58	0.41
8:CH:101:ILE:C	8:CH:101:ILE:HD12	2.40	0.41
30:DI:110:ALA:HB2	30:DI:129:ILE:HD12	2.03	0.41
34:DM:33:LEU:HB2	34:DM:117:PHE:CD2	2.56	0.41
37:BP:32:VAL:HG23	37:BP:34:GLU:HG3	2.01	0.41
6:CF:39:LEU:O	6:CF:40:GLU:HG3	2.20	0.41
22:BA:2819:G:H2'	22:BA:2821:A:N7	2.35	0.41
53:B5:43:GLU:HA	53:B5:178:LYS:HA	2.03	0.41
29:BH:95:GLY:HA2	29:BH:117:LEU:CD2	2.51	0.41
22:BA:2031:A:C6	22:BA:2498:C:H1'	2.56	0.41
22:BA:1076:C:H2'	22:BA:1077:A:N9	2.36	0.41
22:DA:2840:C:H2'	22:DA:2841:C:C6	2.56	0.41
18:CR:20:GLU:C	18:CR:22:ASP:H	2.24	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AP:6:LEU:HB3	16:AP:17:TYR:HB3	2.03	0.41
22:BA:2636:C:H2'	22:BA:2637:U:C6	2.55	0.41
22:DA:580:U:C2'	22:DA:581:C:H5'	2.50	0.41
29:DH:53:GLU:C	29:DH:55:GLU:N	2.72	0.41
2:CB:131:LYS:HE2	2:CB:131:LYS:HA	2.02	0.41
22:DA:1526:C:N4	22:DA:1527:G:C6	2.88	0.41
1:AA:179:A:C6	1:AA:180:U:N3	2.88	0.41
22:DA:301:G:N2	22:DA:302:C:O2	2.53	0.41
24:DC:101:ARG:O	24:DC:102:ARG:CG	2.68	0.41
22:DA:773:U:H5''	22:DA:774:G:OP2	2.20	0.41
24:DC:43:ARG:HB3	24:DC:48:ARG:O	2.21	0.41
24:DC:153:GLN:O	24:DC:156:ARG:HD2	2.20	0.41
5:CE:144:LEU:O	5:CE:147:MET:HB3	2.21	0.41
1:CA:805:C:O2'	1:CA:806:C:H5'	2.20	0.41
2:CB:207:ILE:C	2:CB:210:VAL:HG22	2.41	0.41
1:AA:651:C:N4	1:AA:652:U:O4	2.53	0.41
1:CA:1220:G:H2'	1:CA:1221:G:H8	1.85	0.41
38:BQ:47:TYR:CZ	38:BQ:51:ARG:NH1	2.88	0.41
1:CA:1385:G:C2'	1:CA:1386:G:H5'	2.51	0.41
2:AB:91:PHE:CE1	2:AB:150:GLY:CA	3.04	0.41
1:AA:945:G:H2'	1:AA:945:G:N3	2.36	0.41
1:AA:1074:G:H2'	1:AA:1075:U:O4'	2.21	0.41
22:DA:1857:G:O2'	22:DA:1884:G:N2	2.54	0.41
22:BA:1832:C:C4	22:BA:1833:C:C5	3.08	0.41
1:CA:1317:C:N3	1:CA:1318:A:H1'	2.36	0.41
22:BA:1867:G:C2'	22:BA:1868:C:H5'	2.51	0.41
16:CP:19:VAL:HG13	16:CP:36:VAL:HG12	2.01	0.41
16:CP:36:VAL:O	16:CP:36:VAL:HG13	2.21	0.41
22:DA:847:U:O2	22:DA:934:U:H1'	2.21	0.41
48:B0:48:TYR:CE2	48:B0:53:LYS:HB2	2.55	0.41
35:BN:33:ILE:HD11	48:B0:55:ILE:CD1	2.51	0.41
22:DA:648:G:N2	22:DA:649:G:C4	2.89	0.41
22:BA:1061:U:O2	30:BI:10:LYS:HD2	2.20	0.41
22:BA:2520:C:C5	22:BA:2567:G:C4	3.09	0.41
38:DQ:87:SER:O	39:DR:52:PRO:HD3	2.20	0.41
43:BV:21:ARG:HA	43:BV:25:LYS:O	2.20	0.41
13:CM:23:TYR:O	13:CM:66:GLU:N	2.53	0.41
31:BJ:23:LYS:HE2	31:BJ:142:ILE:OXT	2.21	0.41
27:DF:108:VAL:N	27:DF:109:PRO:HD2	2.36	0.41
9:CI:7:TYR:CD1	9:CI:19:VAL:O	2.74	0.41
22:DA:644:A:N1	22:DA:2369:A:H1'	2.34	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:1869:G:H2'	22:DA:1870:C:H5'	2.02	0.41
24:BC:33:LEU:HA	24:BC:33:LEU:HD23	1.84	0.41
22:BA:851:C:H2'	22:BA:852:U:H6	1.86	0.41
1:AA:270:A:C5	1:AA:271:C:C4	3.08	0.41
1:CA:731:G:O2'	1:CA:732:C:H5'	2.20	0.41
1:AA:942:G:C6	1:AA:943:U:C4	3.09	0.41
22:BA:714:U:O2'	22:BA:716:A:N7	2.50	0.41
6:CF:50:PRO:CD	18:CR:74:HIS:HB3	2.51	0.41
26:DE:184:ASP:O	26:DE:185:LYS:HG2	2.21	0.41
22:DA:2077:A:N3	22:DA:2078:C:C6	2.88	0.41
22:BA:1840:G:C2	22:BA:1841:U:C2	3.08	0.41
22:BA:2341:G:H2'	22:BA:2342:C:C6	2.55	0.41
25:BD:125:TRP:HB2	25:BD:127:PHE:CD1	2.56	0.41
9:AI:19:VAL:HA	9:AI:65:ILE:HG22	2.02	0.41
22:DA:1520:U:O4	22:DA:1521:G:C6	2.74	0.41
22:BA:102:U:C2	46:BY:2:LYS:HE3	2.54	0.41
4:AD:157:ALA:O	4:AD:160:GLU:HB3	2.20	0.41
8:AH:124:GLU:O	8:AH:126:ILE:HD12	2.21	0.41
37:DP:43:PHE:CE2	37:DP:72:ARG:HD3	2.54	0.41
3:AC:191:THR:N	3:AC:194:GLY:O	2.54	0.41
22:BA:744:U:H2'	22:BA:745:G:O4'	2.21	0.41
1:CA:186:C:O4'	20:CT:76:LYS:HD2	2.20	0.41
1:AA:121:U:H3'	1:AA:122:G:H5'	2.02	0.41
1:AA:751:U:H4'	15:AO:24:SER:HA	2.02	0.41
7:CG:84:THR:HG22	7:CG:86:GLN:OE1	2.20	0.41
47:DZ:52:SER:HA	47:DZ:55:VAL:HG22	2.03	0.41
41:DT:20:ALA:CB	41:DT:31:VAL:HG21	2.50	0.41
3:CC:189:ALA:HB3	3:CC:196:ILE:HB	2.02	0.41
1:CA:1111:A:C5	1:CA:1112:C:C5	3.08	0.41
23:BB:75:G:H2'	23:BB:76:G:O4'	2.20	0.41
22:BA:343:C:O2	22:BA:343:C:H2'	2.20	0.41
22:BA:2880:C:O2	22:BA:2880:C:H2'	2.21	0.41
15:AO:32:LEU:HA	15:AO:32:LEU:HD23	1.86	0.41
22:BA:1960:A:H5''	22:BA:1961:C:OP2	2.21	0.41
25:BD:77:ARG:NH2	25:BD:200:ASP:OD1	2.48	0.41
44:BW:36:ILE:HG23	44:BW:58:THR:HG23	2.02	0.41
1:CA:876:C:H1'	8:CH:12:THR:HG21	2.01	0.41
8:CH:96:MET:HB2	8:CH:99:LEU:O	2.20	0.41
22:DA:1695:G:H1'	24:DC:8:PRO:O	2.20	0.41
22:DA:1688:U:C4	22:DA:1698:A:C2	3.09	0.41
39:DR:97:LYS:O	39:DR:99:THR:HG23	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:2431:U:N3	22:DA:2434:A:OP2	2.46	0.41
1:CA:1394:A:H2'	1:CA:1501:C:O2'	2.21	0.41
25:DD:150:GLN:HG3	25:DD:151:THR:N	2.32	0.41
25:DD:148:GLN:OE1	25:DD:152:PRO:HG2	2.21	0.41
1:AA:554:A:H5''	12:AL:26:ALA:HB1	2.01	0.41
2:CB:106:THR:HA	2:CB:109:GLN:OE1	2.21	0.41
45:DX:12:PRO:HB3	45:DX:28:ARG:NH2	2.36	0.41
5:AE:25:VAL:C	5:AE:27:GLY:N	2.74	0.41
5:CE:101:GLU:C	5:CE:103:THR:N	2.74	0.41
22:DA:2823:A:C6	22:DA:2824:C:C5	3.09	0.41
1:AA:69:G:H2'	1:AA:70:U:C6	2.56	0.41
22:DA:2209:G:C5	22:DA:2210:U:C4	3.09	0.41
22:DA:1544:A:N1	22:DA:1545:A:C2	2.88	0.41
22:DA:600:G:H5'	26:DE:27:LEU:HD22	2.03	0.41
22:BA:674:G:O2'	26:BE:69:ARG:CB	2.69	0.41
22:DA:771:G:N1	22:DA:772:C:C5	2.89	0.41
38:BQ:82:GLY:O	38:BQ:83:LEU:C	2.57	0.41
22:DA:319:G:C6	22:DA:333:G:C2	3.08	0.41
5:AE:89:HIS:CD2	5:AE:138:ARG:HD3	2.56	0.41
10:AJ:15:HIS:C	10:AJ:17:LEU:H	2.23	0.41
22:BA:1722:A:C4	22:BA:1739:A:C2	3.09	0.41
22:DA:1860:G:N2	22:DA:1883:U:H1'	2.35	0.41
30:DI:21:SER:HB3	30:DI:22:PRO:HD3	2.03	0.41
15:AO:46:HIS:O	15:AO:47:LYS:HB2	2.20	0.41
22:DA:2867:G:C6	37:DP:21:ARG:NH2	2.88	0.41
38:BQ:9:ILE:CG1	38:BQ:10:ALA:N	2.83	0.41
49:D1:8:LYS:HG3	49:D1:24:THR:HG22	2.03	0.41
1:AA:257:G:C2	1:AA:258:G:C8	3.08	0.41
1:AA:258:G:O2'	1:AA:259:G:H5'	2.20	0.41
5:AE:149:SER:C	5:AE:151:GLU:H	2.24	0.41
2:AB:16:PHE:HD1	2:AB:17:GLY:N	2.18	0.41
27:BF:46:ASP:HB3	27:BF:49:LEU:HB2	2.03	0.41
4:AD:98:LEU:HD23	4:AD:118:VAL:HG13	2.01	0.41
1:AA:464:U:N3	1:AA:466:A:H5''	2.35	0.41
22:DA:1087:G:N9	22:DA:1089:A:H1'	2.36	0.41
2:CB:96:TRP:CE2	2:CB:172:ALA:HB2	2.56	0.41
22:BA:801:G:C8	26:BE:50:ALA:HB2	2.55	0.41
1:CA:572:A:H5'	1:CA:573:A:P	2.61	0.41
22:DA:817:C:O2'	22:DA:839:U:H5''	2.21	0.41
14:AN:45:VAL:HG23	14:AN:46:LEU:H	1.86	0.41
1:CA:1118:U:C1'	1:CA:1179:A:C4	3.04	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:1229:C:H2'	22:DA:1230:A:C8	2.54	0.41
11:AK:102:ALA:C	11:AK:104:GLY:N	2.73	0.41
22:BA:645:C:O2'	22:BA:646:U:H5''	2.21	0.41
17:CQ:39:LYS:O	17:CQ:40:ARG:HD2	2.21	0.41
22:DA:1783:A:C5'	22:DA:2608:G:H4'	2.50	0.41
1:CA:490:C:H2'	1:CA:491:G:H8	1.84	0.41
22:BA:1356:G:C2	22:BA:1357:C:C2	3.09	0.41
22:BA:47:C:H2'	22:BA:48:G:H5'	2.03	0.41
3:AC:54:ARG:O	3:AC:69:HIS:HB2	2.20	0.41
3:AC:77:ILE:O	3:AC:77:ILE:CG2	2.69	0.41
22:DA:416:U:H2'	22:DA:417:C:O4'	2.21	0.41
24:DC:71:LYS:HB2	24:DC:96:TYR:CE2	2.56	0.41
6:CF:85:ILE:HB	6:CF:86:ARG:H	1.73	0.41
22:DA:2310:C:C4	27:DF:77:PHE:CZ	3.08	0.41
32:BK:1:MET:HE2	32:BK:32:TYR:CE1	2.56	0.41
22:BA:1946:U:H2'	22:BA:1947:C:C6	2.56	0.41
23:BB:61:G:H2'	23:BB:62:C:H6	1.84	0.41
22:BA:120:U:H5''	22:BA:122:G:OP2	2.19	0.41
1:CA:1057:G:C5	1:CA:1204:A:C2	3.08	0.41
36:DO:58:ILE:O	36:DO:58:ILE:HG22	2.21	0.41
22:BA:523:C:O2'	22:BA:524:G:H5'	2.20	0.41
29:BH:100:ALA:HB2	29:BH:115:VAL:CG2	2.50	0.41
29:BH:82:SER:HB3	29:BH:146:VAL:HG12	2.03	0.41
22:BA:1171:G:C5	22:BA:1172:C:C4	3.08	0.41
22:BA:1179:G:N7	22:BA:1180:U:O4'	2.53	0.41
22:DA:1361:G:C5	22:DA:1362:C:C5	3.08	0.41
22:DA:2147:A:N7	22:DA:2148:G:C5	2.89	0.41
45:DX:3:ARG:HB3	45:DX:31:PRO:CG	2.50	0.41
1:CA:792:A:O2'	1:CA:794:A:N7	2.38	0.41
38:BQ:58:ARG:HH11	38:BQ:62:ILE:HD11	1.85	0.41
1:CA:31:G:C5	1:CA:306:A:H1'	2.55	0.41
22:DA:83:A:H2'	22:DA:84:A:N7	2.35	0.41
1:AA:406:G:C4	1:AA:495:A:C5	3.08	0.41
1:CA:32:A:C2	1:CA:33:A:N7	2.89	0.41
22:BA:142:A:C8	22:BA:143:C:C5	3.08	0.41
1:CA:1413:A:C2	1:CA:1488:G:C2	3.09	0.41
22:DA:1268:A:C6	22:DA:2013:A:C8	3.09	0.41
1:AA:111:G:O6	1:AA:330:C:N3	2.54	0.41
22:BA:1416:G:O2'	22:BA:1417:C:H6	2.03	0.41
33:BL:111:ILE:CD1	33:BL:111:ILE:N	2.83	0.41
11:CK:88:GLY:N	11:CK:114:THR:HG22	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:1734:G:C5	22:BA:1735:A:N7	2.89	0.41
1:CA:1461:G:H2'	1:CA:1462:C:H6	1.84	0.41
8:AH:11:LEU:HB3	8:AH:75:ILE:HG13	2.02	0.41
36:DO:34:HIS:N	36:DO:65:THR:O	2.52	0.41
1:AA:474:G:N2	1:AA:475:C:H1'	2.35	0.41
30:DI:103:ARG:HB3	30:DI:142:ASP:OD1	2.20	0.41
22:BA:137:U:P	22:BA:140:C:C5	3.14	0.41
22:BA:1812:U:C2'	22:BA:1812:U:O2	2.66	0.41
22:BA:1115:G:N3	22:BA:1116:G:C8	2.89	0.41
1:AA:232:G:C5	1:AA:233:C:C5	3.08	0.41
22:DA:2409:G:C6	22:DA:2410:G:C5	3.09	0.41
29:DH:2:GLN:O	29:DH:3:VAL:O	2.38	0.41
22:DA:2079:U:O4	22:DA:2241:A:N1	2.54	0.41
7:CG:101:MET:HA	7:CG:104:ILE:HD12	2.02	0.41
3:CC:179:ARG:O	3:CC:206:GLU:O	2.38	0.41
28:DG:107:LEU:HB2	28:DG:109:PHE:CZ	2.56	0.41
7:CG:83:SER:O	7:CG:85:TYR:N	2.53	0.41
1:CA:1494:G:C2	1:CA:1495:U:C6	3.09	0.41
47:BZ:24:LEU:HD11	47:BZ:54:MET:HE1	2.02	0.41
4:AD:85:ASN:OD1	4:AD:88:GLU:HG3	2.21	0.41
1:AA:819:A:N7	1:AA:1529:G:C2	2.89	0.41
22:BA:2493:U:H2'	22:BA:2494:G:O5'	2.21	0.41
1:CA:524:G:H2'	1:CA:525:C:H6	1.86	0.41
1:CA:182:A:C5	1:CA:184:G:C8	3.09	0.41
22:DA:566:U:C5	22:DA:567:U:C5	3.09	0.41
1:AA:1421:G:H2'	1:AA:1422:G:O5'	2.21	0.41
1:AA:830:G:H2'	1:AA:831:A:H8	1.85	0.41
24:BC:162:VAL:CG1	24:BC:163:GLN:N	2.83	0.41
36:DO:7:ARG:O	36:DO:10:ARG:N	2.54	0.41
22:BA:2214:C:C5	22:BA:2215:C:C4	3.09	0.41
26:BE:149:ILE:HA	26:BE:170:ARG:O	2.20	0.41
22:BA:2527:C:H2'	22:BA:2528:U:H5'	2.02	0.41
28:DG:138:LYS:HA	28:DG:141:ILE:HG12	2.01	0.41
22:BA:1654:A:C1'	22:BA:2823:A:H5'	2.50	0.41
1:AA:815:A:N3	1:AA:1527:U:O2'	2.47	0.41
1:AA:338:A:N1	1:AA:351:G:O6	2.53	0.41
22:DA:1604:C:O2'	22:DA:1610:A:N1	2.43	0.41
22:BA:1930:G:O2'	22:BA:1931:U:P	2.79	0.41
22:BA:739:A:C2	57:BA:3305:HOH:O	2.74	0.41
35:BN:72:ASP:OD2	35:BN:75:ILE:HD13	2.21	0.41
1:AA:1384:C:H2'	1:AA:1385:G:C8	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:823:C:C4	22:BA:824:U:C4	3.09	0.41
22:BA:2788:C:O2'	22:BA:2809:A:N3	2.47	0.41
22:DA:2599:G:C8	24:DC:236:GLU:HB2	2.55	0.41
43:BV:15:GLY:O	43:BV:19:ARG:HG3	2.20	0.41
22:BA:1290:C:H2'	22:BA:1291:C:H6	1.85	0.41
22:BA:1274:A:N3	22:BA:1297:C:H1'	2.36	0.41
1:AA:780:A:C8	1:AA:800:G:O6	2.74	0.41
22:BA:2335:A:N6	22:BA:2337:G:H1'	2.35	0.41
13:CM:50:GLU:HA	13:CM:50:GLU:OE2	2.21	0.41
26:BE:124:PHE:CD1	26:BE:124:PHE:C	2.93	0.41
4:AD:159:LEU:HD23	4:AD:159:LEU:HA	1.96	0.41
24:DC:267:ILE:O	24:DC:267:ILE:HG22	2.20	0.41
22:BA:2812:G:H2'	22:BA:2813:A:O4'	2.21	0.41
22:BA:2540:C:C2'	22:BA:2541:A:H5'	2.51	0.41
1:CA:827:U:H2'	1:CA:870:U:O4	2.21	0.41
51:D3:34:THR:HG23	51:D3:35:LYS:N	2.36	0.41
11:AK:16:VAL:HG12	11:AK:77:TYR:HB3	2.02	0.41
29:BH:121:VAL:HA	29:BH:128:HIS:CB	2.51	0.41
1:CA:1394:A:N1	1:CA:1500:A:O2'	2.42	0.41
25:DD:148:GLN:CD	25:DD:148:GLN:N	2.74	0.41
22:DA:1153:C:P	57:DA:3361:HOH:O	2.79	0.41
22:DA:58:G:C2	22:DA:70:G:C2	3.08	0.41
28:BG:155:GLU:HG2	28:BG:156:PRO:HD2	2.03	0.41
22:BA:1482:G:N3	22:BA:1483:G:C8	2.89	0.41
22:DA:2842:G:H2'	22:DA:2843:G:O4'	2.21	0.41
22:BA:572:A:C5'	22:BA:573:U:OP2	2.63	0.41
22:DA:1365:A:OP2	45:DX:3:ARG:N	2.51	0.41
22:DA:396:G:H5'	45:DX:13:VAL:HG22	2.02	0.41
37:DP:53:ARG:CB	37:DP:56:HIS:HB2	2.51	0.41
22:DA:54:G:N2	22:DA:117:G:H1'	2.35	0.41
1:AA:374:A:H5''	1:AA:452:A:H2	1.86	0.41
42:BU:18:ASP:O	42:BU:19:LYS:C	2.59	0.41
11:CK:123:PRO:HB2	11:CK:124:PRO:HD2	2.02	0.41
5:CE:157:ARG:C	5:CE:159:LYS:N	2.74	0.41
29:DH:1:MET:HB3	29:DH:21:VAL:O	2.20	0.41
1:CA:62:U:H1'	1:CA:379:C:H1'	2.02	0.41
1:AA:92:U:H2'	1:AA:93:U:C6	2.56	0.41
13:AM:10:PRO:O	13:AM:11:ASP:HB2	2.20	0.41
22:DA:2210:U:H4'	22:DA:2211:A:H5'	2.03	0.41
22:DA:593:U:C2	22:DA:594:U:C4	3.09	0.41
38:BQ:76:TYR:CD2	38:BQ:76:TYR:C	2.94	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:2061:G:C2	22:DA:2063:C:C4	3.09	0.41
22:DA:1814:G:C6	22:DA:1815:A:C6	3.08	0.41
1:AA:469:C:H2'	1:AA:470:C:O4'	2.21	0.41
22:DA:563:A:N1	22:DA:564:C:C2	2.89	0.41
1:CA:247:G:C5	1:CA:278:G:C2	3.08	0.41
22:DA:2467:C:H41	22:DA:2468:A:N6	2.19	0.41
22:BA:2271:G:H2'	22:BA:2272:U:C6	2.56	0.41
1:CA:676:A:N1	1:CA:677:U:C4	2.89	0.41
1:CA:1377:A:C5	7:CG:7:ILE:HD11	2.55	0.41
33:BL:95:LEU:HD22	33:BL:100:ILE:HD11	2.02	0.41
33:BL:82:LEU:HG	33:BL:90:VAL:HG21	2.02	0.41
21:AU:38:TYR:C	21:AU:41:PRO:HD2	2.41	0.41
22:DA:2258:C:H4'	22:DA:2259:U:OP2	2.21	0.41
24:DC:72:ASP:O	24:DC:74:ILE:HD12	2.21	0.41
22:DA:310:A:OP1	42:DU:15:THR:HG22	2.21	0.41
22:DA:2131:U:H4'	22:DA:2133:G:C1'	2.51	0.41
17:CQ:17:MET:O	17:CQ:18:GLU:CB	2.68	0.41
1:CA:784:A:N3	1:CA:785:G:C8	2.88	0.41
1:CA:1220:G:H1'	19:CS:52:HIS:HD2	1.86	0.41
41:BT:61:LEU:CD1	41:BT:61:LEU:C	2.86	0.41
22:BA:1800:C:H3'	24:BC:146:MET:HE1	2.02	0.41
22:DA:1202:G:C5	22:DA:1203:U:C5	3.09	0.41
1:CA:919:A:C2	1:CA:920:U:C5	3.08	0.41
11:CK:112:ASP:OD1	11:CK:114:THR:HG23	2.21	0.41
35:DN:92:GLY:HA2	35:DN:94:TYR:CZ	2.55	0.41
22:DA:1905:C:C4	22:DA:1930:G:C2	3.09	0.41
1:AA:193:C:H2'	1:AA:194:C:H6	1.86	0.41
22:DA:2812:G:N2	22:DA:2889:C:C2	2.89	0.41
12:AL:43:LYS:O	12:AL:44:LYS:C	2.60	0.41
22:BA:2615:U:H1'	48:B0:4:GLN:HB3	2.02	0.41
22:DA:30:G:H2'	22:DA:31:C:C6	2.55	0.41
29:BH:30:LEU:C	29:BH:32:PRO:HD2	2.41	0.41
30:DI:22:PRO:CB	30:DI:23:PRO:HD3	2.50	0.41
1:AA:1118:U:H1'	1:AA:1179:A:C4	2.56	0.41
22:DA:67:U:C2	22:DA:68:G:C8	3.08	0.41
22:DA:68:G:H2'	22:DA:69:C:O4'	2.21	0.41
20:CT:67:ILE:CG1	20:CT:71:LYS:HD3	2.49	0.41
1:AA:972:C:H2'	1:AA:972:C:O2	2.21	0.41
25:DD:125:TRP:CE3	25:DD:160:LYS:HD3	2.55	0.41
10:CJ:87:LEU:HD13	10:CJ:88:MET:N	2.36	0.41
10:CJ:78:GLU:OE2	10:CJ:80:THR:OG1	2.39	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:BF:114:PHE:CE1	27:BF:116:GLY:HA2	2.56	0.41
27:BF:107:ALA:C	27:BF:109:PRO:HD2	2.41	0.41
22:DA:13:A:N7	22:DA:525:U:C4	2.89	0.41
1:CA:706:A:H1'	11:CK:31:ILE:HD11	2.03	0.41
22:DA:523:C:H2'	22:DA:524:G:H8	1.85	0.41
22:DA:1033:U:O2'	22:DA:2750:A:N6	2.54	0.41
22:DA:972:A:C6	22:DA:973:A:C6	3.09	0.41
22:DA:1190:G:OP1	33:DL:32:GLY:CA	2.69	0.41
2:AB:64:LYS:HD3	2:AB:65:GLY:N	2.36	0.41
22:DA:2307:G:H4'	22:DA:2308:G:O5'	2.21	0.41
42:BU:54:GLN:N	42:BU:55:PRO:HD2	2.36	0.41
18:AR:40:VAL:HG13	18:AR:41:PRO:HD2	2.03	0.41
22:BA:207:A:H2'	22:BA:208:C:O4'	2.20	0.41
22:BA:214:G:N2	22:BA:216:A:N3	2.69	0.41
1:CA:209:U:C4'	1:CA:210:C:OP2	2.69	0.41
34:BM:136:MET:CE	43:BV:57:TYR:CE1	3.03	0.41
1:CA:525:C:N3	1:CA:526:C:C4	2.89	0.41
22:DA:713:G:N2	22:DA:719:C:C4	2.89	0.41
22:DA:2446:G:C3'	22:DA:2447:G:H5''	2.51	0.41
1:AA:1421:G:C2'	1:AA:1422:G:O5'	2.69	0.41
46:DY:28:LEU:HD12	46:DY:46:VAL:HG21	2.02	0.41
9:CI:35:LEU:HD11	9:CI:48:VAL:HG21	2.03	0.41
19:AS:58:VAL:O	19:AS:58:VAL:HG22	2.21	0.41
6:AF:54:LEU:HD22	6:AF:55:HIS:N	2.36	0.41
1:AA:1392:G:C6	1:AA:1393:U:C4	3.09	0.41
1:AA:927:G:N2	1:AA:1391:U:H1'	2.35	0.41
22:DA:835:C:C4	22:DA:836:G:N7	2.89	0.41
1:CA:87:C:H2'	1:CA:88:U:N1	2.35	0.41
22:DA:694:U:C3'	22:DA:695:G:H5''	2.50	0.41
26:BE:170:ARG:NH2	26:BE:176:ASP:OD2	2.50	0.41
4:CD:58:LYS:HG3	4:CD:59:GLN:N	2.33	0.41
48:D0:13:ARG:O	48:D0:17:ARG:HD2	2.21	0.41
24:DC:135:ILE:O	24:DC:167:ARG:NH2	2.53	0.41
3:AC:14:ILE:C	3:AC:16:LYS:H	2.25	0.41
6:AF:10:VAL:CG1	6:AF:11:HIS:N	2.84	0.41
13:AM:22:ILE:HG21	13:AM:65:VAL:HG21	2.02	0.41
1:CA:560:A:H5'	1:CA:566:G:N2	2.36	0.41
22:BA:875:G:C2	22:BA:903:C:C2	3.09	0.41
22:DA:1198:U:O2	38:DQ:4:VAL:HG11	2.20	0.41
11:CK:89:PRO:HD3	21:CU:29:LEU:CD1	2.51	0.41
1:AA:376:G:N3	1:AA:389:A:C2	2.89	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:DD:22:ILE:HG22	25:DD:24:VAL:HG13	2.03	0.41
22:DA:2619:C:H4'	25:DD:156:PHE:O	2.21	0.41
1:AA:700:G:O4'	1:AA:704:A:H1'	2.21	0.41
22:BA:1352:U:C2'	22:BA:1353:A:H5'	2.51	0.41
22:BA:819:A:C2'	22:BA:820:A:H5'	2.51	0.41
1:AA:277:C:C2'	1:AA:278:G:H5'	2.50	0.41
15:AO:14:GLU:HB3	15:AO:84:ARG:HH22	1.85	0.41
22:DA:155:A:C2	22:DA:172:A:C2	3.09	0.41
22:BA:1289:C:O2'	22:BA:1330:C:H4'	2.20	0.41
1:AA:1489:G:H2'	1:AA:1490:U:O4'	2.21	0.41
22:BA:1474:U:C2'	22:BA:1475:G:H5'	2.50	0.41
22:BA:2531:A:C6	22:BA:2532:G:C5	3.09	0.41
28:DG:50:LEU:HD13	28:DG:72:LEU:HD23	2.03	0.41
30:DI:5:VAL:HA	30:DI:8:TYR:CE1	2.55	0.41
1:CA:766:A:H2'	1:CA:767:A:O4'	2.21	0.41
40:DS:61:ASN:O	40:DS:62:ASP:HB3	2.21	0.41
25:DD:186:LEU:HD21	37:DP:4:ILE:HG21	2.03	0.41
36:BO:43:ASN:OD1	36:BO:45:SER:HB2	2.19	0.41
28:DG:167:GLU:HG2	28:DG:169:VAL:CG2	2.51	0.41
22:DA:871:U:H4'	34:DM:68:PHE:CD2	2.56	0.41
1:CA:425:G:H2'	1:CA:426:U:O4'	2.20	0.41
23:BB:49:C:O3'	36:BO:68:LYS:HE2	2.21	0.41
1:AA:459:A:H2'	1:AA:460:A:C8	2.56	0.41
22:DA:1503:A:H3'	22:DA:1504:A:H5''	2.03	0.41
22:DA:153:U:H2'	22:DA:154:U:C6	2.56	0.41
1:AA:661:G:O2'	1:AA:662:U:H5'	2.21	0.41
25:DD:3:GLY:O	25:DD:82:PHE:CE2	2.74	0.41
53:B5:87:ALA:HB2	53:B5:153:ILE:CB	2.51	0.41
10:AJ:73:LEU:O	10:AJ:74:VAL:HB	2.21	0.41
1:CA:554:A:H2'	1:CA:555:U:H6	1.86	0.41
47:DZ:47:MET:O	47:DZ:51:VAL:HG22	2.21	0.41
23:BB:78:A:C2	23:BB:99:A:C4	3.09	0.41
9:CI:97:GLU:CD	9:CI:97:GLU:N	2.74	0.41
42:BU:95:PHE:C	42:BU:95:PHE:CD1	2.94	0.41
1:CA:240:G:OP1	1:CA:240:G:H4'	2.20	0.41
32:BK:17:ARG:HD3	32:BK:17:ARG:HA	1.86	0.41
7:AG:139:GLU:HA	7:AG:139:GLU:OE1	2.21	0.41
7:CG:67:GLU:O	7:CG:67:GLU:HG2	2.20	0.41
2:CB:22:TYR:N	2:CB:22:TYR:CD1	2.89	0.41
12:CL:33:VAL:O	12:CL:33:VAL:HG12	2.21	0.41
29:BH:88:GLY:C	29:BH:125:THR:OG1	2.59	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:BH:129:GLU:C	29:BH:130:VAL:CG2	2.90	0.41
29:BH:129:GLU:C	29:BH:130:VAL:HG23	2.42	0.41
29:BH:90:LEU:HG	29:BH:92:GLY:C	2.42	0.41
1:CA:55:A:C6	1:CA:56:U:C2	3.09	0.41
25:BD:129:THR:CG2	25:BD:140:HIS:O	2.68	0.41
22:DA:1791:A:H2'	22:DA:1792:G:O4'	2.21	0.41
22:DA:396:G:H1'	45:DX:29:PHE:CD2	2.56	0.41
22:DA:2683:C:H4'	25:DD:13:ARG:HH12	1.86	0.41
1:AA:1345:U:C2	1:AA:1377:A:C6	3.08	0.41
1:AA:1377:A:C5	7:AG:7:ILE:HD11	2.55	0.41
11:AK:127:ARG:HB2	21:AU:34:ARG:HD2	2.03	0.41
25:BD:103:ASP:C	25:BD:104:VAL:HG22	2.42	0.41
22:DA:2823:A:C2	22:DA:2824:C:N1	2.89	0.41
22:DA:842:U:C2	22:DA:843:G:N7	2.89	0.41
2:AB:203:ASN:OD1	2:AB:204:ASP:N	2.54	0.41
22:DA:594:U:H2'	22:DA:595:C:C6	2.57	0.41
24:DC:246:THR:C	24:DC:248:TRP:H	2.24	0.41
2:AB:72:THR:O	2:AB:73:LYS:CG	2.68	0.41
22:DA:740:C:H5'	22:DA:1784:A:C2'	2.51	0.41
21:AU:40:LYS:N	21:AU:41:PRO:CD	2.84	0.41
22:DA:2546:U:O4'	22:DA:2565:A:C2	2.74	0.41
22:DA:1818:U:H2'	24:DC:156:ARG:HD3	2.02	0.41
1:CA:822:U:C2	1:CA:823:C:C5	3.09	0.41
1:AA:1238:A:H5''	1:AA:1239:A:OP2	2.21	0.41
33:DL:90:VAL:HB	33:DL:122:VAL:HA	2.03	0.41
31:DJ:4:PHE:CG	38:DQ:100:VAL:HG11	2.56	0.41
22:BA:2554:U:C6	22:BA:2555:U:C5	3.10	0.41
6:AF:5:GLU:HB3	6:AF:90:MET:HB2	2.03	0.41
22:DA:1429:G:C2	22:DA:1430:G:C5	3.09	0.41
1:CA:783:C:H2'	1:CA:784:A:H8	1.86	0.41
1:CA:1386:G:O2'	1:CA:1387:G:C5'	2.67	0.41
22:DA:2372:U:H2'	22:DA:2373:G:C8	2.55	0.41
29:BH:1:MET:HE3	29:BH:26:ALA:HB3	2.02	0.41
16:CP:4:ILE:HD13	16:CP:57:ILE:HG12	2.02	0.41
1:AA:96:U:O2'	1:AA:97:G:O5'	2.35	0.41
40:BS:29:VAL:HG11	40:BS:55:ILE:CD1	2.51	0.41
1:CA:1315:U:C5	1:CA:1316:G:C6	3.09	0.41
20:AT:67:ILE:HD11	20:AT:71:LYS:CD	2.50	0.41
1:AA:1374:A:C2	1:AA:1375:A:N9	2.89	0.41
30:BI:80:LEU:HD11	30:BI:132:THR:O	2.21	0.41
27:DF:117:LEU:O	27:DF:177:PHE:HA	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:1113:U:H2'	22:BA:1114:C:H6	1.86	0.41
22:DA:1000:A:C6	22:DA:1001:A:C6	3.09	0.41
16:CP:38:PHE:CZ	16:CP:51:ARG:HB3	2.56	0.41
22:DA:1136:G:H1'	22:DA:2038:G:H4'	2.03	0.41
1:AA:697:U:C6	1:AA:698:G:C8	3.09	0.41
6:CF:2:ARG:O	6:CF:4:TYR:CE2	2.74	0.41
52:B4:36:ARG:CG	52:B4:37:GLN:N	2.84	0.41
53:B5:42:VAL:O	53:B5:42:VAL:HG23	2.21	0.41
30:BI:96:ASP:CG	30:BI:97:LYS:N	2.74	0.41
34:BM:41:LEU:HD22	34:BM:125:PRO:HD2	2.02	0.41
1:CA:1392:G:H2'	1:CA:1393:U:H5'	2.02	0.41
22:DA:1384:A:H1'	22:DA:1405:U:H1'	2.03	0.41
22:BA:754:U:H2'	22:BA:755:U:H6	1.85	0.41
22:DA:567:U:C4	22:DA:568:U:C5	3.09	0.41
22:DA:2538:C:H2'	22:DA:2539:C:H6	1.85	0.41
35:BN:32:GLU:HA	35:BN:115:LEU:HD12	2.02	0.41
2:CB:186:ILE:HA	2:CB:200:ILE:O	2.21	0.41
22:DA:1435:G:O2'	22:DA:1436:G:H5'	2.21	0.41
7:AG:57:SER:OG	7:AG:58:GLU:N	2.53	0.41
1:CA:631:C:H3'	1:CA:632:U:H5'	2.03	0.41
12:CL:94:ARG:HG2	12:CL:94:ARG:H	1.78	0.41
22:BA:780:G:H21	22:BA:783:A:H62	1.69	0.41
22:DA:836:G:C6	22:DA:837:C:C4	3.09	0.41
22:BA:1857:G:N3	22:BA:1884:G:C2	2.89	0.41
12:CL:61:PHE:N	12:CL:61:PHE:HD1	2.19	0.41
26:DE:136:GLN:O	26:DE:137:LYS:C	2.59	0.41
1:AA:186:C:H2'	1:AA:187:G:O4'	2.20	0.41
1:AA:251:G:H4'	1:AA:252:U:O5'	2.17	0.41
53:B5:167:ASP:CB	53:B5:176:VAL:O	2.69	0.41
22:BA:1428:C:C5	22:BA:1569:A:H5''	2.55	0.41
24:BC:119:GLY:C	24:BC:121:ASP:H	2.25	0.41
22:DA:121:G:N2	22:DA:131:A:C4	2.89	0.41
22:BA:2474:U:H5''	22:BA:2475:C:OP2	2.21	0.41
31:DJ:96:ARG:O	31:DJ:98:GLU:N	2.54	0.41
26:BE:198:GLU:O	26:BE:199:MET:C	2.60	0.41
22:DA:1885:A:H2'	22:DA:1886:U:O4'	2.21	0.41
1:CA:1258:G:H2'	1:CA:1259:C:C6	2.56	0.41
22:BA:2673:G:C2	22:BA:2674:G:C8	3.09	0.41
38:BQ:66:ASN:O	38:BQ:70:ARG:HB2	2.20	0.41
22:DA:244:A:C2	22:DA:255:A:C4	3.08	0.41
23:BB:43:C:O2	27:BF:92:ARG:HD2	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:DO:106:LEU:O	36:DO:109:ALA:HB3	2.21	0.41
25:DD:5:VAL:HG11	25:DD:80:TRP:CE3	2.57	0.41
22:BA:655:A:H4'	22:BA:656:G:OP1	2.21	0.41
22:DA:1453:A:N1	35:DN:74:GLU:HG3	2.36	0.41
28:BG:143:GLN:O	28:BG:144:VAL:C	2.59	0.41
42:DU:10:GLU:HG3	42:DU:23:GLY:O	2.21	0.41
40:DS:41:LYS:O	40:DS:42:LYS:C	2.58	0.41
28:BG:164:TYR:HB2	28:BG:167:GLU:HB2	2.04	0.41
28:DG:139:GLN:OE1	28:DG:140:VAL:HG22	2.21	0.41
24:BC:78:VAL:HG21	24:BC:110:LEU:HG	2.03	0.41
26:DE:179:SER:O	26:DE:183:PHE:HD1	2.04	0.41
22:BA:894:U:H2'	22:BA:895:U:C6	2.56	0.41
27:DF:17:MET:SD	27:DF:22:TYR:HB2	2.60	0.41
1:CA:1360:A:C8	14:CN:58:SER:HB3	2.55	0.41
22:DA:2252:G:H2'	22:DA:2253:G:O4'	2.20	0.41
29:BH:132:PHE:CE2	29:BH:142:VAL:CG2	3.04	0.40
22:DA:1355:G:N2	22:DA:1356:G:H1'	2.36	0.40
22:BA:2243:U:O2	22:BA:2434:A:C2	2.74	0.40
29:BH:83:LYS:CE	1:CA:56:U:H5'	2.51	0.40
1:CA:57:G:C5	1:CA:58:C:C4	3.10	0.40
22:BA:1170:C:H2'	22:BA:1171:G:C8	2.56	0.40
22:DA:1607:C:H4'	22:DA:1608:A:O5'	2.21	0.40
22:DA:1373:A:C4	22:DA:1374:G:H1'	2.57	0.40
22:BA:1062:G:OP1	22:BA:1070:A:H4'	2.20	0.40
1:AA:1223:C:OP2	1:AA:1224:U:H2'	2.21	0.40
22:DA:249:C:P	22:DA:2394:C:O2'	2.79	0.40
4:CD:188:ARG:HA	4:CD:188:ARG:HD2	1.81	0.40
22:DA:1020:A:C2	22:DA:1141:U:O2	2.74	0.40
5:CE:137:VAL:O	5:CE:138:ARG:HB2	2.19	0.40
1:AA:772:U:C2'	1:AA:773:G:O5'	2.68	0.40
22:DA:2199:A:O4'	29:DH:28:ASN:ND2	2.54	0.40
22:DA:945:A:H8	57:DA:3258:HOH:O	2.04	0.40
44:BW:19:LYS:O	44:BW:20:ARG:C	2.58	0.40
1:CA:1492:A:N6	1:CA:1493:A:C2	2.89	0.40
1:AA:1299:A:C5	1:AA:1301:U:O2	2.74	0.40
22:DA:2747:G:O2'	28:DG:67:THR:HG22	2.22	0.40
22:DA:236:C:H4'	22:DA:431:U:O2'	2.21	0.40
11:CK:23:ILE:HD11	11:CK:86:VAL:HG13	2.03	0.40
22:DA:1430:G:C5	22:DA:1431:A:N7	2.89	0.40
22:DA:1431:A:C2	22:DA:1432:G:N9	2.89	0.40
1:CA:770:C:C2'	1:CA:771:G:H5'	2.51	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:771:G:O2'	1:CA:772:U:H5'	2.20	0.40
1:AA:653:U:O2'	1:AA:654:G:H8	2.04	0.40
13:AM:64:VAL:O	13:AM:69:LEU:HB2	2.20	0.40
35:DN:12:ARG:HE	35:DN:16:HIS:CE1	2.39	0.40
22:DA:732:C:N4	22:DA:733:G:C6	2.89	0.40
22:DA:583:G:C5	22:DA:584:C:C5	3.10	0.40
1:AA:730:G:N2	1:AA:766:A:OP1	2.47	0.40
20:CT:70:ASN:O	20:CT:71:LYS:C	2.59	0.40
2:CB:203:ASN:ND2	2:CB:206:ALA:HB2	2.36	0.40
13:CM:66:GLU:HB3	13:CM:67:GLY:H	1.76	0.40
32:DK:2:ILE:HG21	32:DK:8:LEU:HD21	2.03	0.40
1:CA:1014:A:N7	1:CA:1015:G:C6	2.89	0.40
27:BF:107:ALA:O	27:BF:108:VAL:C	2.58	0.40
20:AT:83:ILE:O	20:AT:87:ALA:CB	2.69	0.40
5:AE:151:GLU:C	5:AE:153:VAL:H	2.24	0.40
22:DA:7:G:H4'	31:DJ:15:TRP:CH2	2.56	0.40
22:DA:818:G:H2'	22:DA:819:A:H5''	2.04	0.40
1:AA:903:G:H2'	1:AA:904:U:H6	1.86	0.40
15:CO:45:GLU:O	15:CO:46:HIS:HB2	2.21	0.40
1:AA:1067:A:H3'	1:AA:1094:G:OP1	2.21	0.40
23:DB:41:G:H8	27:DF:66:LEU:HD11	1.86	0.40
1:CA:66:A:C6	1:CA:67:C:C4	3.09	0.40
1:CA:66:A:C6	1:CA:67:C:C5	3.10	0.40
22:DA:2389:G:C5'	22:DA:2390:U:H5'	2.51	0.40
1:AA:1048:G:C2	1:AA:1050:G:N7	2.89	0.40
31:BJ:122:LEU:HG	31:BJ:124:VAL:HG12	2.03	0.40
1:CA:1343:G:C6	1:CA:1344:C:C4	3.09	0.40
4:CD:206:LYS:OXT	4:CD:206:LYS:CG	2.69	0.40
22:BA:780:G:H2'	22:BA:782:A:N7	2.36	0.40
22:DA:2077:A:C2	22:DA:2078:C:C4	3.09	0.40
22:DA:1255:U:H3'	22:DA:1256:G:H5''	2.03	0.40
1:CA:438:U:HO2'	1:CA:439:U:P	2.44	0.40
21:CU:15:ALA:O	21:CU:16:LEU:C	2.59	0.40
23:DB:46:A:C5	23:DB:47:C:C4	3.08	0.40
1:AA:746:A:H2'	1:AA:747:A:C8	2.56	0.40
1:CA:392:C:H2'	1:CA:393:A:O4'	2.21	0.40
22:BA:2627:G:O2'	22:BA:2781:A:N1	2.51	0.40
3:AC:47:LEU:HD22	3:AC:76:VAL:HG22	2.02	0.40
3:CC:79:LYS:O	3:CC:81:GLY:N	2.54	0.40
1:AA:886:G:C2	1:AA:912:C:O2	2.74	0.40
22:DA:2533:U:H2'	22:DA:2534:A:O4'	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:43:C:H2'	1:AA:44:A:O4'	2.20	0.40
3:CC:140:ASN:HA	3:CC:143:ARG:HB3	2.02	0.40
1:AA:935:A:O2'	1:AA:1383:C:N3	2.43	0.40
1:CA:1456:A:H2'	1:CA:1457:G:O4'	2.21	0.40
18:CR:46:GLY:O	18:CR:47:THR:O	2.39	0.40
22:BA:936:A:H2'	22:BA:937:C:C6	2.56	0.40
25:DD:121:THR:HG21	25:DD:143:PRO:HB3	2.03	0.40
22:DA:2313:C:H5''	27:DF:88:LYS:HD3	2.03	0.40
1:AA:140:U:H2'	1:AA:141:G:O4'	2.20	0.40
22:DA:1562:U:H2'	22:DA:1563:U:O4'	2.21	0.40
1:CA:1208:C:C4	1:CA:1209:C:C4	3.08	0.40
22:DA:2880:C:H2'	22:DA:2880:C:O2	2.21	0.40
24:BC:212:ARG:HA	24:BC:212:ARG:HD2	1.71	0.40
22:BA:2112:G:H2'	22:BA:2112:G:N3	2.36	0.40
38:BQ:91:ASP:OD1	38:BQ:91:ASP:C	2.59	0.40
4:AD:55:LEU:HD23	4:AD:55:LEU:C	2.41	0.40
33:DL:46:VAL:HB	33:DL:50:PHE:CD2	2.56	0.40
22:DA:698:C:C4	22:DA:762:U:O4	2.75	0.40
8:CH:94:LYS:HD3	8:CH:98:GLY:N	2.36	0.40
25:DD:114:LYS:HE2	25:DD:196:ALA:CB	2.51	0.40
1:AA:811:C:H4'	1:AA:900:A:N6	2.37	0.40
38:DQ:62:ILE:CG1	38:DQ:92:ARG:HD3	2.51	0.40
39:BR:51:VAL:HG23	39:BR:52:PRO:HD2	2.02	0.40
24:DC:219:THR:HG22	24:DC:220:VAL:H	1.86	0.40
22:BA:1910:G:N2	22:BA:1921:G:C1'	2.85	0.40
22:BA:1076:C:H2'	22:BA:1077:A:C8	2.55	0.40
22:BA:511:U:C5	22:BA:512:G:C5	3.09	0.40
1:AA:266:G:H3'	17:AQ:69:LYS:HB2	2.04	0.40
22:BA:1603:A:H5''	22:BA:1604:C:OP2	2.21	0.40
22:DA:35:G:H1'	22:DA:454:A:C4	2.56	0.40
5:CE:104:GLY:O	5:CE:105:ILE:HG22	2.22	0.40
5:CE:80:THR:CB	5:CE:122:ASN:OD1	2.69	0.40
35:BN:67:PHE:CE1	35:BN:73:ASN:ND2	2.90	0.40
22:BA:2318:G:C6	22:BA:2319:G:C6	3.09	0.40
33:DL:55:MET:SD	33:DL:59:ARG:HB3	2.61	0.40
1:CA:834:U:H2'	1:CA:835:U:C6	2.55	0.40
46:BY:18:LEU:HD21	46:BY:22:LEU:HD22	2.04	0.40
1:CA:1211:U:HO2'	1:CA:1212:U:P	2.43	0.40
5:CE:133:PRO:O	5:CE:136:VAL:N	2.53	0.40
22:DA:195:A:C5	22:DA:198:C:C5	3.08	0.40
50:D2:11:LYS:HA	50:D2:14:ARG:HB2	2.03	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:945:A:N1	57:DA:3686:HOH:O	2.51	0.40
22:BA:1386:C:H5''	22:BA:1396:U:O2	2.22	0.40
23:BB:29:A:OP2	36:BO:32:PRO:HD2	2.21	0.40
1:CA:913:A:H4'	1:CA:914:A:H4'	2.02	0.40
8:AH:18:GLN:NE2	8:AH:70:ALA:HB1	2.36	0.40
6:AF:90:MET:O	6:AF:91:ARG:O	2.38	0.40
22:DA:235:U:C2	22:DA:236:C:C6	3.09	0.40
2:CB:166:ALA:HB2	2:CB:187:VAL:HG12	2.02	0.40
22:BA:2799:A:HO2'	22:BA:2800:A:P	2.43	0.40
1:CA:582:C:C2	1:CA:760:G:N1	2.90	0.40
1:CA:582:C:C2	1:CA:760:G:C2	3.08	0.40
22:DA:2221:G:H2'	22:DA:2222:C:H5'	2.03	0.40
40:BS:84:ARG:O	40:BS:96:ILE:N	2.45	0.40
4:AD:150:LYS:NZ	4:AD:178:MET:HB2	2.35	0.40
1:AA:1241:G:N2	1:AA:1242:G:C4	2.90	0.40
15:CO:40:GLN:OE1	22:DA:716:A:H1'	2.22	0.40
22:DA:734:A:O2'	22:DA:1635:A:C5'	2.70	0.40
23:DB:76:G:H2'	23:DB:77:U:O4'	2.22	0.40
22:DA:226:A:H5'	22:DA:257:C:O3'	2.21	0.40
22:BA:2080:A:C5'	45:BX:19:SER:HB2	2.52	0.40
30:BI:77:ALA:HB2	30:BI:132:THR:HG22	2.03	0.40
17:CQ:62:ARG:HD3	17:CQ:76:VAL:HG11	2.02	0.40
31:BJ:142:ILE:HA	31:BJ:142:ILE:HD12	1.94	0.40
19:AS:32:ARG:HG2	19:AS:57:HIS:CD2	2.57	0.40
1:CA:987:G:H2'	1:CA:988:G:O4'	2.22	0.40
17:AQ:80:GLU:C	17:AQ:81:LYS:HD3	2.41	0.40
1:AA:397:A:C5	1:AA:548:G:N7	2.89	0.40
51:D3:6:THR:O	51:D3:8:ARG:N	2.54	0.40
10:CJ:34:ALA:O	10:CJ:35:GLN:CB	2.68	0.40
19:CS:36:ARG:HG2	19:CS:51:VAL:CG1	2.51	0.40
43:BV:10:LYS:HE2	43:BV:10:LYS:H	1.85	0.40
22:DA:1483:G:C4	22:DA:1484:U:C5	3.09	0.40
22:BA:497:A:H2'	22:BA:498:G:O4'	2.21	0.40
9:CI:84:THR:HG22	9:CI:98:LEU:HD11	2.04	0.40
22:DA:2598:A:OP1	24:DC:234:GLY:O	2.40	0.40
22:BA:1183:U:H2'	22:BA:1184:U:C6	2.56	0.40
1:CA:709:U:H2'	1:CA:710:G:C8	2.56	0.40
1:AA:1048:G:OP1	14:AN:4:GLN:N	2.51	0.40
22:DA:2636:C:H2'	22:DA:2637:U:C6	2.55	0.40
42:BU:6:ARG:O	42:BU:7:ARG:C	2.59	0.40
24:BC:10:SER:O	24:BC:13:ARG:HB3	2.20	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:950:G:C6	22:DA:951:C:N3	2.89	0.40
23:BB:109:A:C5	23:BB:110:C:C5	3.09	0.40
1:AA:1391:U:H2'	1:AA:1392:G:C8	2.56	0.40
32:BK:103:VAL:O	32:BK:122:VAL:HB	2.22	0.40
36:BO:100:HIS:O	36:BO:104:GLN:HB3	2.21	0.40
1:AA:380:G:N2	1:AA:384:G:C5	2.90	0.40
22:DA:2106:U:H2'	22:DA:2107:G:C8	2.57	0.40
22:DA:279:A:H61	22:DA:361:G:H1'	1.86	0.40
28:DG:85:LYS:HG2	28:DG:141:ILE:HD12	2.02	0.40
10:AJ:9:ARG:NH2	10:AJ:71:LEU:HD11	2.35	0.40
1:AA:628:G:H2'	1:AA:629:A:O4'	2.21	0.40
22:DA:1539:U:H2'	22:DA:1540:G:C8	2.56	0.40
28:DG:44:LYS:HE3	28:DG:51:THR:OG1	2.22	0.40
22:BA:2585:U:O2'	22:BA:2586:U:C5'	2.69	0.40
1:AA:791:G:N2	1:AA:1497:G:O3'	2.48	0.40
1:AA:277:C:H2'	1:AA:278:G:H5'	2.03	0.40
1:CA:608:A:C8	57:CA:1798:HOH:O	2.71	0.40
1:AA:1126:U:C6	1:AA:1281:C:N3	2.90	0.40
26:BE:148:ILE:HG21	26:BE:157:LEU:HD21	2.04	0.40
3:CC:9:GLY:HA3	14:CN:89:MET:SD	2.61	0.40
22:BA:386:G:H4'	22:BA:387:U:OP2	2.21	0.40
28:DG:41:VAL:HG22	28:DG:64:GLN:O	2.22	0.40
20:CT:48:GLN:O	20:CT:52:ASN:ND2	2.54	0.40
22:DA:1996:C:H5	32:DK:32:TYR:OH	2.04	0.40
35:BN:57:THR:HG22	35:BN:57:THR:O	2.21	0.40
42:DU:53:ASN:N	42:DU:53:ASN:OD1	2.53	0.40
22:DA:2615:U:H2'	22:DA:2615:U:O2	2.20	0.40
53:B5:78:ILE:O	53:B5:78:ILE:HG12	2.22	0.40
25:DD:33:ARG:HB3	25:DD:95:SER:OG	2.22	0.40
1:AA:577:G:C8	1:AA:816:A:C6	3.10	0.40
29:BH:93:SER:C	29:BH:122:LEU:HG	2.35	0.40
22:BA:1921:G:N3	22:BA:1922:G:C8	2.89	0.40
22:BA:1922:G:C6	22:BA:1923:U:C5	3.09	0.40
17:AQ:14:SER:OG	17:AQ:17:MET:HE2	2.21	0.40
22:DA:2235:G:H2'	22:DA:2236:U:H6	1.86	0.40
22:DA:247:G:N7	22:DA:249:C:C2	2.90	0.40
1:CA:1195:C:H2'	1:CA:1197:A:O4'	2.22	0.40
1:CA:1302:C:N4	13:CM:17:ILE:HD11	2.37	0.40
21:CU:34:ARG:O	21:CU:35:ARG:O	2.40	0.40
22:DA:24:G:C5	22:DA:25:U:C5	3.09	0.40
22:DA:514:A:C2	22:DA:515:A:C2	3.10	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:2318:G:O2'	22:BA:2319:G:H5'	2.21	0.40
22:DA:183:C:C4	22:DA:184:C:C4	3.09	0.40
1:CA:577:G:N3	1:CA:578:C:C5	2.89	0.40
1:CA:247:G:C6	1:CA:278:G:N1	2.89	0.40
24:DC:74:ILE:HG22	24:DC:75:PRO:O	2.22	0.40
22:BA:2430:A:H5'	22:BA:2431:U:OP2	2.21	0.40
36:DO:31:THR:O	36:DO:32:PRO:C	2.60	0.40
30:BI:39:CYS:SG	30:BI:43:ASN:ND2	2.94	0.40
1:AA:1102:A:O2'	2:AB:98:GLY:O	2.34	0.40
16:CP:20:VAL:CG2	16:CP:32:PHE:CD1	3.04	0.40
22:BA:1985:C:N3	22:BA:1986:C:C5	2.89	0.40
10:AJ:65:TYR:OH	14:AN:85:ARG:HG3	2.22	0.40
22:DA:1657:U:OP2	25:DD:141:ARG:HG3	2.21	0.40
22:BA:2520:C:C2'	22:BA:2521:C:O5'	2.70	0.40
8:AH:78:VAL:HG11	8:AH:125:ILE:HD11	2.02	0.40
41:BT:18:GLU:O	41:BT:22:THR:HG23	2.22	0.40
1:CA:462:G:N7	1:CA:463:U:C5	2.89	0.40
22:BA:1095:A:H2'	22:BA:1096:A:C8	2.56	0.40
22:DA:480:A:N3	22:DA:480:A:H2'	2.36	0.40
22:BA:1991:U:C2'	22:BA:1992:G:H5'	2.51	0.40
28:DG:107:LEU:HB2	28:DG:109:PHE:CE2	2.57	0.40
26:BE:48:THR:C	26:BE:50:ALA:H	2.25	0.40
36:BO:24:THR:HG23	36:BO:42:PRO:HD3	2.04	0.40
1:AA:965:U:P	57:AA:1833:HOH:O	2.78	0.40
4:AD:26:ARG:HD3	4:AD:31:LYS:HE3	2.03	0.40
1:AA:161:A:H2'	1:AA:162:A:C8	2.56	0.40
22:BA:753:A:C2'	22:BA:754:U:O5'	2.70	0.40
22:DA:1659:G:C5	22:DA:1660:G:C8	3.10	0.40
22:BA:1187:G:P	57:BA:3373:HOH:O	2.79	0.40
26:BE:149:ILE:HD12	26:BE:150:THR:N	2.37	0.40
22:BA:624:C:O2'	22:BA:657:U:H5''	2.20	0.40
44:DW:56:ASP:O	44:DW:57:HIS:HB2	2.21	0.40
22:DA:2474:U:H5''	22:DA:2475:C:OP2	2.21	0.40
22:DA:1783:A:C2	22:DA:2587:A:C4	3.09	0.40
51:B3:63:PRO:HG2	51:B3:64:TYR:CD2	2.56	0.40
1:CA:554:A:H2'	1:CA:555:U:C6	2.56	0.40
3:AC:20:SER:HB3	14:AN:94:PRO:HG3	2.02	0.40
26:BE:46:GLN:O	26:BE:88:ARG:NH1	2.54	0.40
26:DE:47:LYS:HB3	26:DE:51:GLU:HB2	2.03	0.40
4:CD:80:ALA:HA	4:CD:86:THR:OG1	2.20	0.40
1:AA:291:U:O2'	1:AA:292:G:H5'	2.22	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:442:G:C6	1:CA:443:C:C4	3.10	0.40
22:DA:1454:C:H1'	35:DN:60:VAL:HG13	2.02	0.40
22:DA:1549:A:C6	22:DA:1550:C:N3	2.90	0.40
1:CA:104:G:C2	1:CA:105:G:C8	3.10	0.40
11:CK:82:LEU:HD22	11:CK:105:PHE:CD1	2.57	0.40
22:BA:1593:A:H2'	22:BA:1594:U:C6	2.57	0.40
51:B3:45:ARG:HH11	51:B3:45:ARG:HG3	1.86	0.40
22:DA:1021:A:H2'	22:DA:1021:A:N3	2.35	0.40
19:AS:17:LYS:O	19:AS:21:LYS:HB2	2.20	0.40
29:DH:96:THR:O	29:DH:98:ASP:N	2.54	0.40
27:BF:53:ALA:O	27:BF:56:ASP:HB2	2.21	0.40
6:AF:70:VAL:HA	6:AF:73:GLU:HG2	2.03	0.40
1:AA:892:A:C2	1:AA:907:A:C4	3.10	0.40
1:AA:909:A:H2'	1:AA:910:C:O4'	2.21	0.40
22:BA:1916:A:C4	22:BA:1917:U:H1'	2.55	0.40
1:CA:1070:U:H2'	1:CA:1071:C:C6	2.54	0.40
1:AA:536:C:H2'	1:AA:537:G:C8	2.57	0.40
22:DA:2840:C:H2'	22:DA:2841:C:H6	1.87	0.40
22:DA:1394:U:C4	22:DA:1395:A:C5	3.10	0.40
22:DA:1365:A:O5'	45:DX:12:PRO:HG2	2.21	0.40
22:DA:53:A:C8	22:DA:54:G:C8	3.10	0.40
22:DA:2124:G:O6	22:DA:2125:G:N3	2.54	0.40
22:DA:581:C:H2'	22:DA:582:A:C8	2.56	0.40
1:AA:404:G:H2'	1:AA:405:U:O4'	2.22	0.40
24:DC:258:ARG:NH1	24:DC:264:ASP:OD1	2.53	0.40
1:CA:109:A:N1	1:CA:327:A:C6	2.90	0.40
22:DA:589:U:C4	22:DA:590:A:N7	2.89	0.40
1:CA:577:G:C5	1:CA:816:A:C2	3.08	0.40
22:DA:825:A:H4'	22:DA:2428:G:C5	2.56	0.40
5:AE:80:THR:HB	5:AE:122:ASN:ND2	2.36	0.40
1:CA:901:A:N7	1:CA:902:G:H1'	2.36	0.40
1:AA:864:A:H4'	5:AE:90:THR:HG23	2.04	0.40
22:DA:2073:C:O2'	22:DA:2074:U:H5'	2.22	0.40
1:AA:1157:A:H5'	1:AA:1158:C:C5	2.56	0.40
22:DA:106:C:C2'	22:DA:294:A:O2'	2.70	0.40
2:AB:47:VAL:C	2:AB:49:MET:N	2.75	0.40
4:CD:177:LYS:O	4:CD:178:MET:HB2	2.22	0.40
22:DA:1568:G:O6	24:DC:28:LYS:NZ	2.53	0.40
22:BA:2799:A:C6	22:BA:2801:G:C5	3.10	0.40
12:AL:117:TYR:O	12:AL:119:VAL:HG23	2.21	0.40
5:AE:133:PRO:O	5:AE:135:ASN:N	2.53	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:DA:1515:A:O2'	22:DA:1556:C:O2'	2.16	0.40
42:DU:47:LYS:CG	42:DU:48:PRO:HD2	2.52	0.40
22:DA:527:C:N3	22:DA:2779:U:H2'	2.37	0.40
22:BA:1869:G:C3'	22:BA:1870:C:H5'	2.51	0.40
40:BS:29:VAL:CG1	40:BS:55:ILE:CD1	2.99	0.40
6:CF:14:GLN:C	6:CF:16:GLU:H	2.24	0.40
6:AF:29:ILE:HD13	6:AF:64:VAL:CG1	2.51	0.40
1:CA:570:G:C6	1:CA:873:A:N1	2.89	0.40
22:BA:2478:A:H5'	52:B4:32:LYS:HD3	2.02	0.40
32:DK:107:LEU:CD2	32:DK:115:ILE:HG21	2.52	0.40
27:DF:106:ILE:C	27:DF:109:PRO:HD2	2.42	0.40
22:BA:1973:G:H2'	22:BA:1974:C:C6	2.57	0.40
4:AD:188:ARG:CZ	4:AD:197:GLU:OE1	2.69	0.40
42:DU:65:ILE:O	42:DU:65:ILE:HG23	2.22	0.40
22:DA:404:A:H1'	22:DA:405:U:OP2	2.22	0.40
22:DA:1483:G:C5	22:DA:1484:U:C5	3.10	0.40
9:AI:97:GLU:H	9:AI:97:GLU:CD	2.25	0.40
22:DA:663:G:C6	22:DA:664:G:C5	3.09	0.40
30:BI:97:LYS:HB3	30:BI:139:VAL:CG2	2.51	0.40
22:DA:167:A:C4	22:DA:168:G:C8	3.09	0.40
1:CA:1508:A:H2'	1:CA:1509:C:C6	2.56	0.40
1:AA:614:C:C2	1:AA:615:G:C8	3.09	0.40
34:BM:95:LEU:HA	34:BM:95:LEU:HD23	1.91	0.40
1:AA:1251:A:H2'	1:AA:1252:A:O4'	2.22	0.40
22:DA:2478:A:C8	22:DA:2529:G:C6	3.09	0.40
3:CC:65:ARG:O	3:CC:101:ILE:HA	2.21	0.40
22:DA:875:G:N2	22:DA:903:C:C2	2.89	0.40
9:AI:19:VAL:HG22	9:AI:65:ILE:CG2	2.52	0.40
22:DA:2150:C:C4	22:DA:2151:U:C4	3.10	0.40
1:AA:1432:G:H3'	37:BP:106:LYS:HE2	2.03	0.40
1:AA:1070:U:C2	1:AA:1071:C:C5	3.10	0.40
22:BA:1403:A:H2'	22:BA:1404:C:C6	2.56	0.40
22:DA:2618:G:C6	22:DA:2619:C:C4	3.10	0.40
6:CF:49:TYR:CE2	18:CR:66:SER:HA	2.57	0.40
1:AA:642:A:C5	8:AH:107:SER:HA	2.57	0.40
22:DA:2742:G:OP1	52:D4:36:ARG:HD2	2.21	0.40
22:BA:1444:G:H2'	22:BA:1445:G:O4'	2.22	0.40
13:CM:71:ARG:HA	13:CM:74:SER:HB2	2.02	0.40
35:BN:92:GLY:HA2	35:BN:94:TYR:CZ	2.57	0.40
22:DA:1893:C:C5	22:DA:1894:C:C5	3.09	0.40
8:AH:105:SER:O	8:AH:123:GLY:HA3	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:CK:16:VAL:HG12	11:CK:79:ILE:HG12	2.02	0.40
39:BR:4:VAL:HA	39:BR:12:HIS:O	2.22	0.40
6:CF:70:VAL:O	6:CF:73:GLU:HG2	2.21	0.40
37:DP:46:VAL:HG12	37:DP:47:VAL:N	2.37	0.40
30:BI:64:ASP:O	30:BI:65:ARG:HB3	2.21	0.40
31:DJ:28:LEU:HD23	31:DJ:101:ILE:HD12	2.02	0.40
7:CG:123:GLU:OE1	7:CG:126:ASP:HB2	2.21	0.40
4:CD:160:GLU:O	4:CD:163:GLU:HB3	2.21	0.40
32:BK:2:ILE:HD12	32:BK:6:THR:HG21	2.03	0.40
17:CQ:74:THR:HG22	17:CQ:75:LEU:N	2.36	0.40
27:BF:119:ALA:HB1	27:BF:167:ARG:HD2	2.02	0.40
27:DF:131:GLY:HA2	27:DF:153:ASP:HA	2.03	0.40
22:DA:651:G:C6	22:DA:652:U:C4	3.09	0.40
22:BA:1790:C:H6	22:BA:1790:C:O5'	2.05	0.40
22:BA:880:G:N2	22:BA:898:C:C2	2.90	0.40
35:DN:86:ARG:HG2	35:DN:86:ARG:O	2.20	0.40
24:BC:44:ASN:C	24:BC:44:ASN:OD1	2.59	0.40
12:CL:43:LYS:HB3	12:CL:43:LYS:HE2	1.91	0.40
46:BY:28:LEU:O	46:BY:31:GLN:HB3	2.22	0.40
22:BA:1243:C:H2'	22:BA:1244:A:O4'	2.21	0.40
1:AA:1263:C:H2'	1:AA:1264:U:O4'	2.22	0.40
45:BX:53:ALA:O	45:BX:54:LYS:C	2.59	0.40
22:BA:611:C:C2'	22:BA:612:G:H5'	2.51	0.40
1:AA:519:C:H2'	1:AA:520:A:O4'	2.22	0.40
35:DN:35:LYS:HE3	35:DN:110:MET:HB3	2.03	0.40
28:DG:14:GLY:O	28:DG:28:GLY:HA2	2.22	0.40
25:BD:85:ALA:O	25:BD:86:GLU:C	2.60	0.40
25:DD:151:THR:HB	25:DD:152:PRO:HD2	2.03	0.40
22:DA:1009:A:C6	22:DA:1010:A:N1	2.90	0.40
22:BA:1180:U:H2'	22:BA:1181:U:O4'	2.21	0.40
1:AA:921:U:H2'	1:AA:922:G:O4'	2.21	0.40
1:CA:1523:G:H2'	1:CA:1524:C:C6	2.56	0.40
22:DA:580:U:H4'	38:DQ:31:VAL:HG11	2.04	0.40
22:DA:581:C:OP1	38:DQ:33:ARG:HG3	2.22	0.40
22:DA:374:A:N6	22:DA:401:A:C8	2.89	0.40
22:DA:570:G:C4	22:DA:2030:A:C8	3.09	0.40
22:DA:571:U:H1'	22:DA:573:U:C6	2.56	0.40
1:AA:144:G:H2'	1:AA:145:G:O4'	2.21	0.40
1:AA:178:C:H2'	1:AA:179:A:O4'	2.22	0.40
22:BA:784:G:O2'	22:BA:785:G:H5''	2.21	0.40
22:BA:2151:U:H2'	22:BA:2152:G:N7	2.36	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:716:A:N3	11:CK:120:GLY:HA2	2.37	0.40
1:CA:587:G:H4'	8:CH:4:GLN:HB3	2.03	0.40
22:BA:45:G:H5''	22:BA:46:G:H5'	2.03	0.40
22:BA:2515:C:O2'	22:BA:2516:A:H5'	2.22	0.40
11:AK:87:LYS:HA	11:AK:114:THR:HG22	2.03	0.40
1:CA:1438:G:OP1	20:CT:29:ARG:HD3	2.21	0.40
22:BA:1415:U:O2'	22:BA:1416:G:H4'	2.21	0.40
22:BA:1590:A:C2	22:BA:1591:A:C5	3.09	0.40
1:CA:582:C:N3	1:CA:760:G:C2	2.90	0.40
1:AA:1154:G:H2'	1:AA:1155:A:C8	2.57	0.40
48:B0:34:SER:O	48:B0:35:GLY:C	2.60	0.40
22:DA:1027:A:N7	22:DA:1126:A:C2	2.88	0.40
22:BA:2114:A:C4	22:BA:2167:U:H5'	2.57	0.40
23:DB:38:C:O4'	36:DO:100:HIS:CE1	2.74	0.40
1:AA:1118:U:H2'	1:AA:1119:C:O4'	2.21	0.40
7:AG:120:LEU:HD22	7:AG:120:LEU:O	2.22	0.40
1:AA:1349:A:C6	1:AA:1374:A:C8	3.09	0.40
22:DA:547:A:H3'	22:DA:548:G:H5'	2.02	0.40
21:AU:42:THR:O	21:AU:46:LYS:HB2	2.22	0.40
41:DT:44:LYS:HG3	41:DT:55:VAL:HB	2.03	0.40
14:AN:28:LYS:N	14:AN:31:ILE:HB	2.37	0.40
22:DA:972:A:C2	22:DA:973:A:N6	2.89	0.40
22:DA:982:C:H4'	22:DA:983:A:OP1	2.22	0.40
12:AL:38:TYR:HB2	12:AL:52:VAL:HG23	2.03	0.40
9:CI:84:THR:HG21	9:CI:103:PHE:HB3	2.03	0.40
1:AA:1347:G:H2'	9:AI:110:GLN:O	2.21	0.40
24:DC:185:GLU:O	24:DC:188:CYS:SG	2.64	0.40
57:AA:1783:HOH:O	14:AN:3:LYS:HA	2.20	0.40
1:AA:942:G:C4	1:AA:943:U:C5	3.10	0.40
1:CA:209:U:H5''	1:CA:210:C:OP2	2.21	0.40
22:DA:2084:C:H2'	22:DA:2085:U:H6	1.87	0.40
22:BA:2582:G:H3'	22:BA:2582:G:OP2	2.22	0.40
22:BA:988:A:P	47:BZ:12:SER:HB2	2.61	0.40
4:AD:165:ARG:O	4:AD:167:LYS:N	2.54	0.40
22:DA:1422:G:C6	22:DA:1423:G:C5	3.09	0.40
31:DJ:75:TYR:CE2	31:DJ:86:GLN:OE1	2.75	0.40
41:DT:57:VAL:HG12	41:DT:86:THR:OG1	2.21	0.40
10:AJ:7:ARG:CB	10:AJ:75:ASP:OD1	2.69	0.40
8:CH:101:ILE:HD12	8:CH:101:ILE:O	2.22	0.40
34:DM:67:VAL:HG11	34:DM:96:ILE:HD12	2.04	0.40
22:BA:359:G:C5	22:BA:360:U:C5	3.09	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:BA:185:G:H2'	22:BA:186:G:O4'	2.21	0.40
1:CA:619:U:N3	4:CD:132:ILE:CD1	2.84	0.40
22:BA:2131:U:H5'	22:BA:2132:U:H5''	2.03	0.40
22:BA:108:G:C2'	22:BA:109:C:H5'	2.51	0.40
1:AA:414:A:H2'	1:AA:415:A:H8	1.87	0.40
22:BA:599:A:O2'	22:BA:600:G:H5'	2.21	0.40
22:BA:1408:G:C6	22:BA:1409:U:C4	3.09	0.40
43:BV:92:VAL:O	43:BV:92:VAL:HG12	2.21	0.40
18:CR:25:ASP:OD1	18:CR:25:ASP:N	2.54	0.40
1:AA:1047:G:N3	1:AA:1047:G:H2'	2.35	0.40
22:BA:419:U:H2'	22:BA:420:C:C6	2.56	0.40

All (5) symmetry-related close contacts are listed below. The label for Atom-2 includes the symmetry operator and encoded unit-cell translations to be applied.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:368:U:OP2	29:DH:123:ARG:NH2[4_455]	2.03	0.17
1:AA:368:U:OP2	29:DH:123:ARG:NE[4_455]	2.07	0.13
1:AA:368:U:OP1	29:DH:93:SER:OG[4_455]	2.08	0.12
1:AA:368:U:OP2	29:DH:123:ARG:CZ[4_455]	2.18	0.02
1:AA:359:G:OP1	29:DH:89:LYS:NZ[4_455]	2.19	0.01

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 X-ray entries followed by that with respect to entries of similar resolution.

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	
2	AB	216/218 (99%)	140 (65%)	41 (19%)	35 (16%)	0	0
2	CB	216/218 (99%)	144 (67%)	44 (20%)	28 (13%)	0	1
3	AC	204/206 (99%)	161 (79%)	32 (16%)	11 (5%)	2	14
3	CC	204/206 (99%)	165 (81%)	29 (14%)	10 (5%)	3	16

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
4	AD	203/205 (99%)	149 (73%)	29 (14%)	25 (12%)	0	2
4	CD	203/205 (99%)	155 (76%)	34 (17%)	14 (7%)	1	8
5	AE	148/150 (99%)	109 (74%)	24 (16%)	15 (10%)	1	4
5	CE	148/150 (99%)	104 (70%)	29 (20%)	15 (10%)	1	4
6	AF	98/100 (98%)	75 (76%)	17 (17%)	6 (6%)	2	11
6	CF	98/100 (98%)	72 (74%)	17 (17%)	9 (9%)	1	4
7	AG	149/151 (99%)	111 (74%)	31 (21%)	7 (5%)	3	17
7	CG	149/151 (99%)	127 (85%)	18 (12%)	4 (3%)	6	31
8	AH	127/129 (98%)	97 (76%)	20 (16%)	10 (8%)	1	6
8	CH	127/129 (98%)	100 (79%)	24 (19%)	3 (2%)	7	33
9	AI	125/127 (98%)	93 (74%)	28 (22%)	4 (3%)	5	26
9	CI	125/127 (98%)	95 (76%)	23 (18%)	7 (6%)	2	13
10	AJ	96/98 (98%)	67 (70%)	13 (14%)	16 (17%)	0	0
10	CJ	96/98 (98%)	69 (72%)	15 (16%)	12 (12%)	0	1
11	AK	115/117 (98%)	91 (79%)	15 (13%)	9 (8%)	1	6
11	CK	115/117 (98%)	87 (76%)	23 (20%)	5 (4%)	3	19
12	AL	121/123 (98%)	84 (69%)	31 (26%)	6 (5%)	3	16
12	CL	121/123 (98%)	90 (74%)	16 (13%)	15 (12%)	0	1
13	AM	112/114 (98%)	84 (75%)	20 (18%)	8 (7%)	1	8
13	CM	112/114 (98%)	83 (74%)	16 (14%)	13 (12%)	0	2
14	AN	92/100 (92%)	65 (71%)	17 (18%)	10 (11%)	0	3
14	CN	92/100 (92%)	71 (77%)	13 (14%)	8 (9%)	1	5
15	AO	86/88 (98%)	72 (84%)	10 (12%)	4 (5%)	3	17
15	CO	86/88 (98%)	69 (80%)	14 (16%)	3 (4%)	4	24
16	AP	80/82 (98%)	49 (61%)	18 (22%)	13 (16%)	0	0
16	CP	80/82 (98%)	61 (76%)	14 (18%)	5 (6%)	2	10
17	AQ	78/80 (98%)	53 (68%)	20 (26%)	5 (6%)	2	10
17	CQ	78/80 (98%)	59 (76%)	12 (15%)	7 (9%)	1	5
18	AR	53/55 (96%)	44 (83%)	8 (15%)	1 (2%)	10	40
18	CR	53/55 (96%)	41 (77%)	8 (15%)	4 (8%)	1	7
19	AS	77/79 (98%)	61 (79%)	13 (17%)	3 (4%)	4	22

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
19	CS	77/79 (98%)	66 (86%)	6 (8%)	5 (6%)	1	9
20	AT	83/85 (98%)	68 (82%)	8 (10%)	7 (8%)	1	6
20	CT	83/85 (98%)	67 (81%)	10 (12%)	6 (7%)	1	7
21	AU	49/51 (96%)	29 (59%)	11 (22%)	9 (18%)	0	0
21	CU	49/51 (96%)	24 (49%)	11 (22%)	14 (29%)	0	0
24	BC	269/271 (99%)	219 (81%)	38 (14%)	12 (4%)	3	17
24	DC	269/271 (99%)	200 (74%)	43 (16%)	26 (10%)	1	4
25	BD	207/209 (99%)	186 (90%)	18 (9%)	3 (1%)	14	48
25	DD	207/209 (99%)	165 (80%)	32 (16%)	10 (5%)	3	17
26	BE	199/201 (99%)	174 (87%)	22 (11%)	3 (2%)	13	46
26	DE	199/201 (99%)	167 (84%)	24 (12%)	8 (4%)	4	21
27	BF	175/177 (99%)	144 (82%)	19 (11%)	12 (7%)	1	8
27	DF	175/177 (99%)	143 (82%)	24 (14%)	8 (5%)	3	17
28	BG	174/176 (99%)	150 (86%)	18 (10%)	6 (3%)	5	25
28	DG	174/176 (99%)	145 (83%)	21 (12%)	8 (5%)	3	17
29	BH	147/149 (99%)	91 (62%)	35 (24%)	21 (14%)	0	1
29	DH	147/149 (99%)	100 (68%)	32 (22%)	15 (10%)	1	4
30	BI	139/141 (99%)	85 (61%)	37 (27%)	17 (12%)	0	2
30	DI	139/141 (99%)	89 (64%)	39 (28%)	11 (8%)	1	6
31	BJ	140/142 (99%)	127 (91%)	9 (6%)	4 (3%)	6	29
31	DJ	140/142 (99%)	116 (83%)	22 (16%)	2 (1%)	14	48
32	BK	120/122 (98%)	98 (82%)	16 (13%)	6 (5%)	3	16
32	DK	120/122 (98%)	100 (83%)	14 (12%)	6 (5%)	3	16
33	BL	141/143 (99%)	114 (81%)	21 (15%)	6 (4%)	3	19
33	DL	141/143 (99%)	106 (75%)	28 (20%)	7 (5%)	3	16
34	BM	134/136 (98%)	118 (88%)	13 (10%)	3 (2%)	8	36
34	DM	134/136 (98%)	111 (83%)	18 (13%)	5 (4%)	4	23
35	BN	118/120 (98%)	98 (83%)	18 (15%)	2 (2%)	11	43
35	DN	118/120 (98%)	93 (79%)	18 (15%)	7 (6%)	2	12
36	BO	114/116 (98%)	94 (82%)	18 (16%)	2 (2%)	11	42
36	DO	114/116 (98%)	94 (82%)	16 (14%)	4 (4%)	4	24

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
37	BP	112/114 (98%)	104 (93%)	5 (4%)	3 (3%)	6	31
37	DP	112/114 (98%)	89 (80%)	18 (16%)	5 (4%)	3	17
38	BQ	115/117 (98%)	107 (93%)	5 (4%)	3 (3%)	7	32
38	DQ	115/117 (98%)	97 (84%)	15 (13%)	3 (3%)	7	32
39	BR	101/103 (98%)	90 (89%)	5 (5%)	6 (6%)	2	12
39	DR	101/103 (98%)	79 (78%)	17 (17%)	5 (5%)	3	16
40	BS	108/110 (98%)	99 (92%)	7 (6%)	2 (2%)	10	40
40	DS	108/110 (98%)	91 (84%)	13 (12%)	4 (4%)	4	23
41	BT	91/93 (98%)	73 (80%)	9 (10%)	9 (10%)	1	4
41	DT	91/93 (98%)	65 (71%)	18 (20%)	8 (9%)	1	5
42	BU	100/102 (98%)	79 (79%)	15 (15%)	6 (6%)	2	11
42	DU	100/102 (98%)	74 (74%)	13 (13%)	13 (13%)	0	1
43	BV	92/94 (98%)	86 (94%)	5 (5%)	1 (1%)	17	55
43	DV	92/94 (98%)	83 (90%)	7 (8%)	2 (2%)	8	36
44	BW	74/76 (97%)	66 (89%)	7 (10%)	1 (1%)	14	48
44	DW	73/76 (96%)	61 (84%)	11 (15%)	1 (1%)	14	48
45	BX	75/77 (97%)	68 (91%)	6 (8%)	1 (1%)	15	50
45	DX	75/77 (97%)	54 (72%)	17 (23%)	4 (5%)	2	14
46	BY	61/63 (97%)	47 (77%)	5 (8%)	9 (15%)	0	1
46	DY	61/63 (97%)	50 (82%)	8 (13%)	3 (5%)	3	16
47	BZ	56/58 (97%)	52 (93%)	4 (7%)	0	100	100
47	DZ	56/58 (97%)	49 (88%)	5 (9%)	2 (4%)	4	24
48	B0	54/56 (96%)	44 (82%)	8 (15%)	2 (4%)	4	23
48	D0	54/56 (96%)	37 (68%)	14 (26%)	3 (6%)	2	13
49	B1	48/50 (96%)	39 (81%)	7 (15%)	2 (4%)	3	19
49	D1	48/50 (96%)	35 (73%)	9 (19%)	4 (8%)	1	6
50	B2	44/46 (96%)	39 (89%)	3 (7%)	2 (4%)	3	17
50	D2	44/46 (96%)	38 (86%)	3 (7%)	3 (7%)	1	8
51	B3	62/64 (97%)	57 (92%)	4 (6%)	1 (2%)	12	44
51	D3	62/64 (97%)	54 (87%)	6 (10%)	2 (3%)	5	26
52	B4	36/38 (95%)	34 (94%)	2 (6%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
52	D4	36/38 (95%)	31 (86%)	3 (8%)	2 (6%)	2	13
53	B5	183/207 (88%)	100 (55%)	58 (32%)	25 (14%)	0	1
All	All	11418/11651 (98%)	8949 (78%)	1727 (15%)	742 (6%)	1	9

All (742) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
2	AB	16	PHE
2	AB	34	ALA
2	AB	73	LYS
2	AB	74	ARG
2	AB	120	GLN
2	AB	134	ALA
2	AB	148	LEU
2	AB	152	LYS
2	AB	194	ASP
2	AB	220	THR
3	AC	26	THR
3	AC	61	ALA
3	AC	141	ALA
4	AD	23	SER
4	AD	24	GLY
4	AD	33	LYS
4	AD	35	GLU
4	AD	160	GLU
4	AD	168	PRO
4	AD	191	LEU
4	AD	192	SER
5	AE	26	LYS
5	AE	76	LEU
5	AE	105	ILE
5	AE	122	ASN
6	AF	91	ARG
6	AF	92	THR
8	AH	66	PHE
8	AH	67	GLN
10	AJ	33	GLY
10	AJ	57	VAL
10	AJ	61	ALA
10	AJ	101	SER
11	AK	52	PHE

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Mol	Chain	Res	Type
11	AK	73	ALA
11	AK	127	ARG
12	AL	24	LEU
12	AL	25	GLU
12	AL	44	LYS
13	AM	4	ILE
13	AM	112	PRO
14	AN	28	LYS
14	AN	52	PRO
14	AN	62	ASN
14	AN	92	GLU
15	AO	3	LEU
16	AP	11	ALA
16	AP	48	GLU
17	AQ	68	SER
19	AS	6	LYS
20	AT	4	ILE
20	AT	6	SER
21	AU	35	ARG
21	AU	38	TYR
21	AU	40	LYS
24	BC	71	LYS
24	BC	233	GLY
24	BC	244	PRO
25	BD	104	VAL
25	BD	152	PRO
27	BF	172	ALA
28	BG	119	ALA
29	BH	10	ALA
29	BH	34	GLY
29	BH	53	GLU
29	BH	87	GLU
29	BH	90	LEU
29	BH	119	ASN
29	BH	121	VAL
29	BH	140	ALA
30	BI	45	LYS
30	BI	63	ALA
30	BI	90	SER
30	BI	113	LYS
32	BK	35	VAL
32	BK	91	SER

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Mol	Chain	Res	Type
32	BK	108	ARG
33	BL	69	ARG
33	BL	88	GLY
33	BL	115	GLU
34	BM	69	PRO
35	BN	118	ARG
36	BO	87	ILE
36	BO	88	LYS
37	BP	94	LYS
39	BR	49	ILE
40	BS	64	ALA
41	BT	17	SER
41	BT	71	GLY
41	BT	72	GLN
41	BT	88	LYS
42	BU	17	LYS
42	BU	39	ILE
42	BU	100	SER
45	BX	3	ARG
46	BY	22	LEU
46	BY	23	ARG
46	BY	24	GLU
46	BY	36	GLN
50	B2	44	VAL
53	B5	53	ARG
53	B5	134	PRO
53	B5	141	PRO
53	B5	174	ALA
53	B5	175	PRO
53	B5	181	PHE
53	B5	221	PRO
2	CB	16	PHE
2	CB	51	ASN
2	CB	74	ARG
2	CB	87	CYS
2	CB	88	ASP
2	CB	103	ASN
2	CB	120	GLN
2	CB	193	PRO
2	CB	194	ASP
4	CD	25	VAL
4	CD	26	ARG

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Mol	Chain	Res	Type
4	CD	33	LYS
4	CD	36	GLN
5	CE	45	ARG
5	CE	51	GLY
5	CE	100	SER
5	CE	101	GLU
5	CE	103	THR
5	CE	123	VAL
5	CE	138	ARG
5	CE	158	GLY
6	CF	14	GLN
6	CF	56	LYS
6	CF	91	ARG
6	CF	92	THR
6	CF	98	GLU
9	CI	129	LYS
10	CJ	36	VAL
10	CJ	57	VAL
10	CJ	90	LEU
10	CJ	92	LEU
11	CK	52	PHE
11	CK	127	ARG
12	CL	17	ALA
12	CL	24	LEU
12	CL	34	CYS
12	CL	43	LYS
12	CL	76	GLU
12	CL	77	HIS
12	CL	89	ASP
13	CM	7	ILE
13	CM	11	ASP
13	CM	41	GLU
13	CM	49	SER
14	CN	92	GLU
15	CO	17	ARG
17	CQ	70	THR
18	CR	21	ILE
18	CR	47	THR
19	CS	5	LEU
20	CT	4	ILE
20	CT	6	SER
20	CT	68	HIS

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Mol	Chain	Res	Type
21	CU	10	GLU
21	CU	13	ASP
21	CU	35	ARG
21	CU	36	GLU
21	CU	39	GLU
21	CU	40	LYS
24	DC	29	PRO
24	DC	35	GLU
24	DC	58	HIS
24	DC	71	LYS
25	DD	104	VAL
25	DD	151	THR
25	DD	152	PRO
26	DE	83	VAL
26	DE	86	ALA
27	DF	9	LYS
27	DF	123	ASP
28	DG	119	ALA
29	DH	3	VAL
29	DH	10	ALA
29	DH	33	GLN
29	DH	35	LYS
29	DH	41	LYS
29	DH	53	GLU
29	DH	54	LEU
29	DH	83	LYS
29	DH	109	GLU
30	DI	93	PRO
31	DJ	42	ALA
31	DJ	81	ILE
32	DK	92	GLU
32	DK	108	ARG
34	DM	69	PRO
34	DM	77	PRO
35	DN	88	ALA
35	DN	104	ALA
37	DP	24	ASP
37	DP	66	ASN
39	DR	31	GLU
39	DR	82	HIS
40	DS	29	VAL
40	DS	62	ASP

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Mol	Chain	Res	Type
41	DT	37	ASP
42	DU	21	LYS
42	DU	55	PRO
42	DU	89	ASP
45	DX	3	ARG
45	DX	62	LYS
50	D2	44	VAL
50	D2	45	SER
2	AB	13	GLY
2	AB	83	ALA
2	AB	117	LEU
2	AB	126	PHE
2	AB	150	GLY
2	AB	155	GLY
2	AB	170	HIS
2	AB	193	PRO
2	AB	212	LEU
3	AC	15	VAL
3	AC	139	GLN
3	AC	140	ASN
4	AD	26	ARG
4	AD	29	ASP
4	AD	126	ASN
4	AD	153	SER
4	AD	175	ALA
5	AE	12	GLN
5	AE	43	ASN
5	AE	138	ARG
7	AG	56	LYS
7	AG	130	ASN
8	AH	57	PRO
8	AH	69	LYS
9	AI	13	LYS
9	AI	58	VAL
11	AK	27	PHE
11	AK	89	PRO
12	AL	89	ASP
13	AM	12	HIS
13	AM	105	ASN
13	AM	111	GLY
13	AM	114	LYS
14	AN	27	LEU

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Mol	Chain	Res	Type
14	AN	47	LYS
14	AN	53	ARG
16	AP	46	LYS
16	AP	68	SER
17	AQ	13	VAL
18	AR	48	ARG
20	AT	5	LYS
20	AT	68	HIS
21	AU	10	GLU
21	AU	12	PHE
24	BC	124	ILE
24	BC	169	GLY
24	BC	196	GLY
24	BC	253	LYS
25	BD	86	GLU
26	BE	11	ALA
27	BF	41	GLY
27	BF	42	GLU
27	BF	62	GLY
27	BF	176	PRO
28	BG	39	ASP
28	BG	79	VAL
28	BG	175	LYS
29	BH	3	VAL
29	BH	11	ASN
29	BH	14	SER
29	BH	15	LEU
29	BH	66	ASN
29	BH	123	ARG
30	BI	7	ALA
30	BI	58	VAL
31	BJ	95	ARG
32	BK	5	GLN
33	BL	68	SER
37	BP	35	GLY
38	BQ	7	GLY
38	BQ	102	ASP
39	BR	53	PHE
39	BR	57	GLY
40	BS	63	GLY
42	BU	50	PRO
44	BW	20	ARG

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Mol	Chain	Res	Type
46	BY	46	VAL
46	BY	57	LEU
46	BY	62	GLY
48	B0	55	ILE
50	B2	42	LEU
53	B5	62	THR
53	B5	86	GLU
53	B5	126	SER
53	B5	144	GLY
53	B5	205	ALA
53	B5	215	VAL
53	B5	217	THR
2	CB	36	ASN
2	CB	124	GLY
2	CB	166	ALA
2	CB	170	HIS
2	CB	220	THR
3	CC	64	ILE
3	CC	141	ALA
3	CC	146	ALA
4	CD	24	GLY
4	CD	27	ALA
4	CD	35	GLU
4	CD	175	ALA
5	CE	126	LYS
5	CE	143	GLY
6	CF	13	ASP
7	CG	130	ASN
8	CH	114	ARG
9	CI	41	ARG
9	CI	58	VAL
9	CI	91	ASP
10	CJ	17	LEU
10	CJ	38	GLY
11	CK	92	GLY
12	CL	23	ALA
12	CL	93	VAL
12	CL	105	SER
13	CM	10	PRO
13	CM	44	LYS
14	CN	50	THR
14	CN	52	PRO

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Mol	Chain	Res	Type
16	CP	42	ILE
17	CQ	82	ALA
19	CS	6	LYS
20	CT	5	LYS
20	CT	67	ILE
21	CU	12	PHE
21	CU	24	GLU
21	CU	42	THR
21	CU	52	ALA
21	CU	53	VAL
24	DC	36	LYS
24	DC	48	ARG
24	DC	159	GLY
24	DC	196	GLY
24	DC	239	ASN
24	DC	240	PHE
24	DC	255	LYS
25	DD	36	GLN
25	DD	194	PRO
26	DE	6	LYS
26	DE	18	THR
26	DE	48	THR
27	DF	176	PRO
28	DG	61	GLY
29	DH	31	VAL
29	DH	77	THR
29	DH	118	PRO
30	DI	7	ALA
30	DI	84	ALA
30	DI	90	SER
32	DK	35	VAL
33	DL	4	ASN
33	DL	53	GLY
33	DL	111	ILE
34	DM	57	VAL
34	DM	58	LYS
35	DN	2	ARG
35	DN	3	HIS
35	DN	70	THR
37	DP	114	LEU
38	DQ	102	ASP
40	DS	63	GLY

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Mol	Chain	Res	Type
41	DT	77	ARG
42	DU	9	ASP
42	DU	57	GLY
42	DU	78	GLY
45	DX	32	ASN
46	DY	37	LEU
47	DZ	4	THR
47	DZ	30	ARG
49	D1	11	LEU
50	D2	38	GLY
51	D3	7	VAL
2	AB	53	ALA
2	AB	68	LEU
2	AB	75	ALA
2	AB	79	ALA
2	AB	87	CYS
2	AB	97	LEU
2	AB	210	VAL
3	AC	54	ARG
4	AD	30	THR
4	AD	34	ILE
4	AD	149	ALA
4	AD	151	LYS
5	AE	24	THR
5	AE	98	PRO
7	AG	33	ASP
7	AG	79	ARG
7	AG	80	VAL
8	AH	19	ALA
9	AI	44	ALA
10	AJ	45	ARG
10	AJ	75	ASP
11	AK	103	ALA
11	AK	128	ARG
14	AN	81	ARG
15	AO	46	HIS
16	AP	36	VAL
16	AP	44	SER
16	AP	49	GLY
16	AP	80	LYS
17	AQ	17	MET
17	AQ	51	ASN

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Mol	Chain	Res	Type
17	AQ	70	THR
19	AS	76	PRO
21	AU	33	ARG
21	AU	37	PHE
21	AU	52	ALA
24	BC	205	LEU
24	BC	231	PRO
27	BF	146	VAL
27	BF	177	PHE
28	BG	158	LYS
29	BH	9	VAL
29	BH	30	LEU
29	BH	85	GLY
29	BH	93	SER
29	BH	105	ALA
30	BI	75	PRO
30	BI	84	ALA
30	BI	98	VAL
30	BI	115	ALA
30	BI	117	MET
31	BJ	81	ILE
32	BK	93	GLN
37	BP	114	LEU
39	BR	55	ASP
41	BT	89	GLU
48	B0	26	THR
49	B1	28	ARG
51	B3	28	ASN
53	B5	36	ALA
53	B5	176	VAL
53	B5	180	SER
2	CB	22	TYR
2	CB	64	LYS
2	CB	73	LYS
2	CB	86	SER
2	CB	126	PHE
2	CB	141	LEU
3	CC	14	ILE
3	CC	80	LYS
5	CE	12	GLN
5	CE	98	PRO
5	CE	122	ASN

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Mol	Chain	Res	Type
6	CF	85	ILE
7	CG	84	THR
9	CI	42	GLU
10	CJ	7	ARG
10	CJ	35	GLN
12	CL	25	GLU
12	CL	78	SER
12	CL	118	GLY
13	CM	114	LYS
14	CN	22	ALA
14	CN	42	TRP
15	CO	46	HIS
18	CR	25	ASP
19	CS	32	ARG
21	CU	11	PRO
24	DC	46	ASN
24	DC	85	PRO
24	DC	105	LEU
24	DC	205	LEU
24	DC	218	PRO
24	DC	238	ARG
24	DC	260	ASN
24	DC	271	ARG
25	DD	57	ALA
25	DD	118	PHE
26	DE	42	GLY
26	DE	69	ARG
26	DE	129	PRO
27	DF	175	PHE
28	DG	20	ASN
28	DG	46	ALA
29	DH	16	GLY
29	DH	40	THR
30	DI	140	VAL
33	DL	29	LYS
33	DL	30	THR
35	DN	106	ASP
36	DO	34	HIS
37	DP	111	LYS
38	DQ	39	VAL
39	DR	70	GLU
39	DR	102	SER

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Mol	Chain	Res	Type
40	DS	65	ASP
41	DT	22	THR
42	DU	7	ARG
42	DU	41	LEU
42	DU	98	SER
42	DU	99	ASN
46	DY	12	GLU
49	D1	51	GLU
2	AB	133	GLU
2	AB	147	SER
3	AC	66	VAL
3	AC	157	LEU
4	AD	166	GLU
4	AD	167	LYS
5	AE	100	SER
5	AE	152	MET
6	AF	5	GLU
6	AF	54	LEU
7	AG	15	ASP
8	AH	15	ARG
8	AH	18	GLN
8	AH	21	ASN
8	AH	98	GLY
10	AJ	36	VAL
10	AJ	38	GLY
10	AJ	43	PRO
10	AJ	47	GLU
10	AJ	79	PRO
11	AK	108	THR
12	AL	26	ALA
14	AN	3	LYS
15	AO	20	ASN
20	AT	7	ALA
21	AU	11	PRO
24	BC	29	PRO
27	BF	134	GLU
27	BF	147	ASP
27	BF	175	PHE
28	BG	152	ARG
29	BH	83	LYS
30	BI	4	LYS
30	BI	60	THR

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Mol	Chain	Res	Type
32	BK	119	ALA
41	BT	3	ARG
41	BT	28	ASN
41	BT	52	GLU
42	BU	8	ASP
42	BU	52	LEU
49	B1	5	ILE
53	B5	133	GLY
2	CB	34	ALA
2	CB	82	ASP
2	CB	149	GLY
2	CB	203	ASN
3	CC	101	ILE
4	CD	32	CYS
4	CD	52	GLY
4	CD	192	SER
5	CE	24	THR
6	CF	54	LEU
8	CH	97	ALA
9	CI	120	LYS
11	CK	15	GLN
11	CK	93	ARG
13	CM	5	ALA
13	CM	13	LYS
15	CO	21	ASP
16	CP	76	LYS
17	CQ	18	GLU
19	CS	28	LYS
19	CS	30	PRO
24	DC	122	ALA
24	DC	143	ASN
24	DC	189	ARG
24	DC	190	ALA
27	DF	3	LYS
27	DF	21	ASN
27	DF	174	ASP
29	DH	9	VAL
30	DI	101	ILE
32	DK	93	GLN
32	DK	110	GLU
32	DK	120	PRO
33	DL	9	ALA

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Mol	Chain	Res	Type
36	DO	101	GLY
38	DQ	46	ALA
39	DR	53	PHE
44	DW	28	GLY
46	DY	57	LEU
49	D1	27	LYS
49	D1	52	ALA
52	D4	20	ASP
2	AB	64	LYS
2	AB	201	PRO
3	AC	17	PRO
4	AD	17	THR
5	AE	157	ARG
6	AF	6	ILE
6	AF	36	ILE
7	AG	113	ASP
10	AJ	35	GLN
10	AJ	42	LEU
12	AL	15	LYS
13	AM	11	ASP
14	AN	69	ARG
16	AP	10	GLY
16	AP	77	GLU
19	AS	29	LYS
20	AT	69	LYS
24	BC	240	PHE
26	BE	199	MET
31	BJ	39	LYS
33	BL	114	GLY
38	BQ	116	ALA
41	BT	2	ILE
46	BY	20	ASN
53	B5	50	ILE
53	B5	65	LEU
53	B5	90	ALA
53	B5	185	LYS
2	CB	21	ARG
2	CB	33	GLY
2	CB	104	TRP
3	CC	17	PRO
3	CC	166	GLU
4	CD	4	TYR

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Mol	Chain	Res	Type
4	CD	167	LYS
4	CD	182	PHE
7	CG	82	GLY
7	CG	146	GLU
9	CI	38	TYR
10	CJ	42	LEU
10	CJ	58	ASN
12	CL	40	THR
13	CM	25	VAL
16	CP	80	LYS
17	CQ	80	GLU
18	CR	34	THR
20	CT	7	ALA
21	CU	37	PHE
24	DC	261	LYS
24	DC	262	ARG
25	DD	205	PRO
27	DF	159	THR
28	DG	12	PRO
28	DG	118	PRO
33	DL	42	SER
41	DT	21	SER
41	DT	38	ALA
41	DT	72	GLN
41	DT	73	ARG
41	DT	88	LYS
42	DU	53	ASN
48	D0	4	GLN
2	AB	183	VAL
4	AD	125	VAL
10	AJ	32	THR
10	AJ	41	PRO
15	AO	73	LYS
16	AP	15	PRO
16	AP	16	PHE
27	BF	84	PRO
27	BF	121	SER
30	BI	21	SER
30	BI	39	CYS
31	BJ	98	GLU
43	BV	67	GLY
46	BY	37	LEU

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Mol	Chain	Res	Type
53	B5	146	VAL
53	B5	162	ILE
2	CB	25	PRO
3	CC	66	VAL
10	CJ	79	PRO
13	CM	110	LYS
14	CN	3	LYS
14	CN	62	ASN
16	CP	49	GLY
17	CQ	5	ILE
30	DI	9	VAL
30	DI	22	PRO
35	DN	82	GLU
37	DP	32	VAL
43	DV	84	PRO
51	D3	20	GLY
2	AB	182	PRO
4	AD	7	PRO
5	AE	51	GLY
10	AJ	74	VAL
11	AK	90	GLY
20	AT	58	VAL
34	BM	87	GLY
10	CJ	6	ILE
13	CM	24	GLY
14	CN	34	VAL
17	CQ	13	VAL
24	DC	19	VAL
25	DD	195	GLY
42	DU	58	ILE
3	AC	101	ILE
13	AM	64	VAL
24	BC	137	VAL
26	BE	151	GLY
30	BI	122	ILE
33	BL	130	GLY
35	BN	109	PRO
12	CL	4	VAL
28	DG	154	PRO
30	DI	74	PRO
36	DO	66	GLY
45	DX	31	PRO

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Mol	Chain	Res	Type
48	D0	55	ILE
52	D4	23	ILE
4	AD	64	ILE
4	AD	101	VAL
5	AE	106	ILE
16	AP	78	VAL
39	BR	51	VAL
53	B5	182	PRO
5	CE	133	PRO
6	CF	7	VAL
13	CM	94	GLY
17	CQ	76	VAL
30	DI	13	VAL
42	DU	25	VAL
43	DV	81	PRO
4	AD	25	VAL
9	AI	51	PRO
30	BI	59	ILE
34	BM	26	VAL
3	CC	174	PRO
8	CH	92	LEU
16	CP	33	ILE
21	CU	41	PRO
25	DD	120	GLY
34	DM	3	GLN
48	D0	8	PRO
2	AB	41	ILE
2	AB	124	GLY
2	AB	224	GLY
5	AE	134	ILE
8	AH	14	ILE
29	BH	118	PRO
39	BR	64	VAL
28	DG	17	VAL
30	DI	89	GLY
36	DO	32	PRO

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 X-ray entries followed by that with respect to entries of similar resolution.

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	
2	AB	180/180 (100%)	142 (79%)	38 (21%)	1	6
2	CB	180/180 (100%)	141 (78%)	39 (22%)	1	5
3	AC	170/170 (100%)	138 (81%)	32 (19%)	2	8
3	CC	170/170 (100%)	150 (88%)	20 (12%)	6	25
4	AD	172/172 (100%)	148 (86%)	24 (14%)	4	18
4	CD	172/172 (100%)	149 (87%)	23 (13%)	5	20
5	AE	113/113 (100%)	89 (79%)	24 (21%)	1	6
5	CE	113/113 (100%)	94 (83%)	19 (17%)	2	11
6	AF	87/87 (100%)	71 (82%)	16 (18%)	2	9
6	CF	87/87 (100%)	67 (77%)	20 (23%)	1	4
7	AG	124/124 (100%)	103 (83%)	21 (17%)	2	11
7	CG	124/124 (100%)	102 (82%)	22 (18%)	2	10
8	AH	104/104 (100%)	92 (88%)	12 (12%)	7	27
8	CH	104/104 (100%)	88 (85%)	16 (15%)	3	14
9	AI	105/105 (100%)	82 (78%)	23 (22%)	1	5
9	CI	105/105 (100%)	84 (80%)	21 (20%)	1	7
10	AJ	86/86 (100%)	71 (83%)	15 (17%)	2	11
10	CJ	86/86 (100%)	75 (87%)	11 (13%)	5	21
11	AK	90/90 (100%)	73 (81%)	17 (19%)	2	8
11	CK	90/90 (100%)	75 (83%)	15 (17%)	3	11
12	AL	103/103 (100%)	90 (87%)	13 (13%)	5	22
12	CL	103/103 (100%)	81 (79%)	22 (21%)	1	5
13	AM	92/92 (100%)	82 (89%)	10 (11%)	8	30
13	CM	92/92 (100%)	82 (89%)	10 (11%)	8	30
14	AN	79/83 (95%)	71 (90%)	8 (10%)	9	33
14	CN	79/83 (95%)	74 (94%)	5 (6%)	22	58
15	AO	76/76 (100%)	66 (87%)	10 (13%)	5	20
15	CO	76/76 (100%)	61 (80%)	15 (20%)	1	7
16	AP	65/65 (100%)	54 (83%)	11 (17%)	2	11
16	CP	65/65 (100%)	55 (85%)	10 (15%)	3	14

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
17	AQ	74/74 (100%)	58 (78%)	16 (22%)	1	5
17	CQ	74/74 (100%)	59 (80%)	15 (20%)	1	6
18	AR	48/48 (100%)	44 (92%)	4 (8%)	14	46
18	CR	48/48 (100%)	43 (90%)	5 (10%)	9	32
19	AS	70/70 (100%)	63 (90%)	7 (10%)	9	34
19	CS	70/70 (100%)	63 (90%)	7 (10%)	9	34
20	AT	65/65 (100%)	55 (85%)	10 (15%)	3	14
20	CT	65/65 (100%)	55 (85%)	10 (15%)	3	14
21	AU	44/44 (100%)	29 (66%)	15 (34%)	0	0
21	CU	44/44 (100%)	31 (70%)	13 (30%)	0	1
24	BC	216/216 (100%)	195 (90%)	21 (10%)	10	36
24	DC	216/216 (100%)	191 (88%)	25 (12%)	7	27
25	BD	164/164 (100%)	154 (94%)	10 (6%)	23	59
25	DD	164/164 (100%)	147 (90%)	17 (10%)	9	32
26	BE	165/165 (100%)	152 (92%)	13 (8%)	15	49
26	DE	165/165 (100%)	152 (92%)	13 (8%)	15	49
27	BF	148/148 (100%)	127 (86%)	21 (14%)	4	18
27	DF	148/148 (100%)	131 (88%)	17 (12%)	7	27
28	BG	137/137 (100%)	122 (89%)	15 (11%)	8	30
28	DG	137/137 (100%)	124 (90%)	13 (10%)	11	38
29	BH	114/114 (100%)	89 (78%)	25 (22%)	1	5
29	DH	114/114 (100%)	89 (78%)	25 (22%)	1	5
30	BI	109/109 (100%)	86 (79%)	23 (21%)	1	6
30	DI	109/109 (100%)	91 (84%)	18 (16%)	3	12
31	BJ	116/116 (100%)	106 (91%)	10 (9%)	13	45
31	DJ	116/116 (100%)	110 (95%)	6 (5%)	29	65
32	BK	103/103 (100%)	93 (90%)	10 (10%)	10	36
32	DK	103/103 (100%)	96 (93%)	7 (7%)	20	55
33	BL	102/102 (100%)	88 (86%)	14 (14%)	4	19
33	DL	102/102 (100%)	94 (92%)	8 (8%)	16	49
34	BM	109/109 (100%)	98 (90%)	11 (10%)	9	33

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles
34	DM	109/109 (100%)	104 (95%)	5 (5%)	33 70
35	BN	100/100 (100%)	87 (87%)	13 (13%)	5 21
35	DN	100/100 (100%)	89 (89%)	11 (11%)	8 30
36	BO	86/86 (100%)	72 (84%)	14 (16%)	3 12
36	DO	86/86 (100%)	78 (91%)	8 (9%)	11 39
37	BP	99/99 (100%)	89 (90%)	10 (10%)	9 33
37	DP	99/99 (100%)	89 (90%)	10 (10%)	9 33
38	BQ	89/89 (100%)	81 (91%)	8 (9%)	12 41
38	DQ	89/89 (100%)	84 (94%)	5 (6%)	26 62
39	BR	84/84 (100%)	71 (84%)	13 (16%)	3 14
39	DR	84/84 (100%)	79 (94%)	5 (6%)	24 60
40	BS	93/93 (100%)	80 (86%)	13 (14%)	4 18
40	DS	93/93 (100%)	86 (92%)	7 (8%)	17 51
41	BT	80/80 (100%)	68 (85%)	12 (15%)	3 15
41	DT	80/80 (100%)	68 (85%)	12 (15%)	3 15
42	BU	83/83 (100%)	74 (89%)	9 (11%)	8 30
42	DU	83/83 (100%)	73 (88%)	10 (12%)	6 24
43	BV	78/78 (100%)	68 (87%)	10 (13%)	5 21
43	DV	78/78 (100%)	71 (91%)	7 (9%)	12 41
44	BW	57/58 (98%)	55 (96%)	2 (4%)	43 78
44	DW	56/58 (97%)	52 (93%)	4 (7%)	18 54
45	BX	67/67 (100%)	64 (96%)	3 (4%)	34 70
45	DX	67/67 (100%)	61 (91%)	6 (9%)	12 41
46	BY	55/55 (100%)	47 (86%)	8 (14%)	4 16
46	DY	55/55 (100%)	46 (84%)	9 (16%)	3 12
47	BZ	48/48 (100%)	41 (85%)	7 (15%)	4 16
47	DZ	48/48 (100%)	40 (83%)	8 (17%)	3 11
48	B0	47/47 (100%)	40 (85%)	7 (15%)	4 15
48	D0	47/47 (100%)	42 (89%)	5 (11%)	8 31
49	B1	45/45 (100%)	39 (87%)	6 (13%)	5 20
49	D1	45/45 (100%)	43 (96%)	2 (4%)	35 71

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
50	B2	38/38 (100%)	34 (90%)	4 (10%)	8	31
50	D2	38/38 (100%)	32 (84%)	6 (16%)	3	13
51	B3	51/51 (100%)	47 (92%)	4 (8%)	16	49
51	D3	51/51 (100%)	48 (94%)	3 (6%)	24	60
52	B4	34/34 (100%)	29 (85%)	5 (15%)	4	16
52	D4	34/34 (100%)	31 (91%)	3 (9%)	12	43
53	B5	61/161 (38%)	50 (82%)	11 (18%)	2	10
All	All	9388/9499 (99%)	8137 (87%)	1251 (13%)	5	20

All (1251) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
2	AB	14	VAL
2	AB	15	HIS
2	AB	21	ARG
2	AB	23	TRP
2	AB	27	MET
2	AB	32	PHE
2	AB	38	VAL
2	AB	39	HIS
2	AB	41	ILE
2	AB	43	LEU
2	AB	44	GLU
2	AB	46	THR
2	AB	49	MET
2	AB	50	PHE
2	AB	56	GLU
2	AB	64	LYS
2	AB	66	LYS
2	AB	88	ASP
2	AB	89	GLN
2	AB	100	MET
2	AB	107	VAL
2	AB	117	LEU
2	AB	123	ASP
2	AB	126	PHE
2	AB	129	LEU
2	AB	130	THR
2	AB	139	ARG
2	AB	146	ASN

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Mol	Chain	Res	Type
2	AB	153	ASP
2	AB	164	ILE
2	AB	174	LYS
2	AB	186	ILE
2	AB	199	VAL
2	AB	207	ILE
2	AB	208	ARG
2	AB	210	VAL
2	AB	213	TYR
2	AB	225	ARG
3	AC	3	GLN
3	AC	14	ILE
3	AC	15	VAL
3	AC	18	TRP
3	AC	20	SER
3	AC	21	THR
3	AC	26	THR
3	AC	27	LYS
3	AC	28	GLU
3	AC	33	LEU
3	AC	37	PHE
3	AC	55	ILE
3	AC	58	GLU
3	AC	59	ARG
3	AC	69	HIS
3	AC	86	LYS
3	AC	88	ARG
3	AC	107	ARG
3	AC	121	THR
3	AC	131	ARG
3	AC	140	ASN
3	AC	142	MET
3	AC	143	ARG
3	AC	151	VAL
3	AC	162	ILE
3	AC	165	THR
3	AC	167	TRP
3	AC	168	TYR
3	AC	185	ASN
3	AC	191	THR
3	AC	193	TYR
3	AC	200	VAL

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Mol	Chain	Res	Type
4	AD	9	LEU
4	AD	13	ARG
4	AD	17	THR
4	AD	23	SER
4	AD	31	LYS
4	AD	35	GLU
4	AD	48	LEU
4	AD	58	LYS
4	AD	60	LYS
4	AD	63	ARG
4	AD	83	LYS
4	AD	93	LEU
4	AD	104	ARG
4	AD	116	GLN
4	AD	123	ILE
4	AD	128	ARG
4	AD	143	VAL
4	AD	152	GLN
4	AD	161	LEU
4	AD	163	GLU
4	AD	190	ASP
4	AD	198	HIS
4	AD	199	LEU
4	AD	206	LYS
5	AE	10	GLU
5	AE	14	LYS
5	AE	15	LEU
5	AE	18	VAL
5	AE	19	ASN
5	AE	30	ILE
5	AE	32	SER
5	AE	34	THR
5	AE	38	VAL
5	AE	46	VAL
5	AE	69	ARG
5	AE	77	ASN
5	AE	81	LEU
5	AE	83	HIS
5	AE	85	VAL
5	AE	88	VAL
5	AE	97	GLN
5	AE	114	VAL

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Mol	Chain	Res	Type
5	AE	115	LEU
5	AE	122	ASN
5	AE	124	LEU
5	AE	134	ILE
5	AE	136	VAL
5	AE	149	SER
6	AF	5	GLU
6	AF	7	VAL
6	AF	14	GLN
6	AF	15	SER
6	AF	17	GLN
6	AF	24	ARG
6	AF	39	LEU
6	AF	45	ARG
6	AF	51	ILE
6	AF	54	LEU
6	AF	55	HIS
6	AF	68	GLN
6	AF	77	THR
6	AF	84	VAL
6	AF	86	ARG
6	AF	96	VAL
7	AG	4	ARG
7	AG	9	GLN
7	AG	32	VAL
7	AG	36	LYS
7	AG	37	SER
7	AG	38	THR
7	AG	52	GLN
7	AG	59	LEU
7	AG	62	PHE
7	AG	70	ARG
7	AG	73	VAL
7	AG	78	ARG
7	AG	79	ARG
7	AG	80	VAL
7	AG	95	ARG
7	AG	120	LEU
7	AG	133	THR
7	AG	135	VAL
7	AG	136	LYS
7	AG	142	HIS

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Mol	Chain	Res	Type
7	AG	144	MET
8	AH	13	ARG
8	AH	22	LYS
8	AH	30	SER
8	AH	42	GLU
8	AH	49	PHE
8	AH	58	GLU
8	AH	83	LEU
8	AH	87	LYS
8	AH	90	ASP
8	AH	99	LEU
8	AH	104	VAL
8	AH	121	LEU
9	AI	12	ARG
9	AI	22	LYS
9	AI	30	ILE
9	AI	36	GLU
9	AI	42	GLU
9	AI	43	THR
9	AI	46	MET
9	AI	48	VAL
9	AI	49	ARG
9	AI	55	VAL
9	AI	57	MET
9	AI	63	LEU
9	AI	65	ILE
9	AI	68	LYS
9	AI	88	MET
9	AI	89	GLU
9	AI	90	TYR
9	AI	94	LEU
9	AI	99	ARG
9	AI	100	LYS
9	AI	111	VAL
9	AI	114	LYS
9	AI	130	ARG
10	AJ	6	ILE
10	AJ	8	ILE
10	AJ	17	LEU
10	AJ	32	THR
10	AJ	42	LEU
10	AJ	52	LEU

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Mol	Chain	Res	Type
10	AJ	57	VAL
10	AJ	59	LYS
10	AJ	63	ASP
10	AJ	73	LEU
10	AJ	75	ASP
10	AJ	83	THR
10	AJ	84	VAL
10	AJ	89	ARG
10	AJ	91	ASP
11	AK	17	SER
11	AK	18	ASP
11	AK	31	ILE
11	AK	38	GLN
11	AK	52	PHE
11	AK	58	SER
11	AK	65	VAL
11	AK	81	ASN
11	AK	95	SER
11	AK	97	ILE
11	AK	100	LEU
11	AK	101	ASN
11	AK	107	ILE
11	AK	108	THR
11	AK	111	THR
11	AK	126	LYS
11	AK	128	ARG
12	AL	10	LYS
12	AL	21	VAL
12	AL	25	GLU
12	AL	29	GLN
12	AL	44	LYS
12	AL	54	ARG
12	AL	58	THR
12	AL	62	GLU
12	AL	64	THR
12	AL	89	ASP
12	AL	102	LEU
12	AL	105	SER
12	AL	121	ARG
13	AM	3	ARG
13	AM	4	ILE
13	AM	7	ILE

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Mol	Chain	Res	Type
13	AM	13	LYS
13	AM	27	LYS
13	AM	29	ARG
13	AM	59	GLU
13	AM	72	GLU
13	AM	85	CYS
13	AM	107	ARG
14	AN	7	LYS
14	AN	26	GLU
14	AN	28	LYS
14	AN	48	LEU
14	AN	62	ASN
14	AN	63	ARG
14	AN	81	ARG
14	AN	98	LYS
15	AO	6	GLU
15	AO	17	ARG
15	AO	31	LEU
15	AO	39	LEU
15	AO	40	GLN
15	AO	48	LYS
15	AO	57	LEU
15	AO	75	VAL
15	AO	83	GLU
15	AO	87	LEU
16	AP	1	MET
16	AP	2	VAL
16	AP	5	ARG
16	AP	6	LEU
16	AP	8	ARG
16	AP	19	VAL
16	AP	20	VAL
16	AP	29	ASN
16	AP	46	LYS
16	AP	71	VAL
16	AP	75	ILE
17	AQ	4	LYS
17	AQ	13	VAL
17	AQ	16	LYS
17	AQ	17	MET
17	AQ	29	VAL
17	AQ	40	ARG

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Mol	Chain	Res	Type
17	AQ	51	ASN
17	AQ	52	GLU
17	AQ	55	ILE
17	AQ	63	GLU
17	AQ	68	SER
17	AQ	70	THR
17	AQ	74	THR
17	AQ	75	LEU
17	AQ	76	VAL
17	AQ	81	LYS
18	AR	29	LEU
18	AR	30	LYS
18	AR	43	ARG
18	AR	55	LEU
19	AS	6	LYS
19	AS	15	LEU
19	AS	21	LYS
19	AS	41	PHE
19	AS	55	ARG
19	AS	63	THR
19	AS	65	GLU
20	AT	3	ASN
20	AT	5	LYS
20	AT	12	ILE
20	AT	27	MET
20	AT	34	LYS
20	AT	36	TYR
20	AT	54	MET
20	AT	70	ASN
20	AT	74	ARG
20	AT	76	LYS
21	AU	5	LYS
21	AU	9	ASN
21	AU	10	GLU
21	AU	12	PHE
21	AU	16	LEU
21	AU	18	ARG
21	AU	19	PHE
21	AU	28	VAL
21	AU	34	ARG
21	AU	36	GLU
21	AU	37	PHE

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Mol	Chain	Res	Type
21	AU	38	TYR
21	AU	43	THR
21	AU	47	ARG
21	AU	54	LYS
24	BC	3	VAL
24	BC	14	ARG
24	BC	18	LYS
24	BC	24	LEU
24	BC	97	LYS
24	BC	105	LEU
24	BC	118	SER
24	BC	141	VAL
24	BC	156	ARG
24	BC	167	ARG
24	BC	174	LEU
24	BC	187	ASP
24	BC	195	VAL
24	BC	197	ASN
24	BC	213	TRP
24	BC	221	ARG
24	BC	223	THR
24	BC	244	PRO
24	BC	245	VAL
24	BC	258	ARG
24	BC	265	LYS
25	BD	12	THR
25	BD	32	ASN
25	BD	73	VAL
25	BD	95	SER
25	BD	97	SER
25	BD	121	THR
25	BD	129	THR
25	BD	141	ARG
25	BD	157	LYS
25	BD	177	VAL
26	BE	4	VAL
26	BE	40	ARG
26	BE	44	ARG
26	BE	95	LYS
26	BE	107	SER
26	BE	108	ILE
26	BE	123	LYS

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Mol	Chain	Res	Type
26	BE	126	VAL
26	BE	136	GLN
26	BE	149	ILE
26	BE	159	LEU
26	BE	198	GLU
26	BE	200	LEU
27	BF	3	LYS
27	BF	14	LYS
27	BF	17	MET
27	BF	21	ASN
27	BF	31	VAL
27	BF	35	THR
27	BF	36	LEU
27	BF	42	GLU
27	BF	44	ILE
27	BF	48	LYS
27	BF	51	ASP
27	BF	61	SER
27	BF	66	LEU
27	BF	67	ILE
27	BF	81	GLN
27	BF	83	TYR
27	BF	92	ARG
27	BF	95	ARG
27	BF	133	ARG
27	BF	147	ASP
27	BF	154	ILE
28	BG	9	VAL
28	BG	39	ASP
28	BG	67	THR
28	BG	69	ARG
28	BG	77	ILE
28	BG	80	THR
28	BG	89	LEU
28	BG	116	GLN
28	BG	124	GLU
28	BG	127	THR
28	BG	139	GLN
28	BG	149	ARG
28	BG	155	GLU
28	BG	166	ASP
28	BG	168	VAL

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Mol	Chain	Res	Type
29	BH	1	MET
29	BH	3	VAL
29	BH	6	LEU
29	BH	12	LEU
29	BH	15	LEU
29	BH	27	ARG
29	BH	50	ARG
29	BH	60	GLU
29	BH	62	LEU
29	BH	66	ASN
29	BH	75	LEU
29	BH	77	THR
29	BH	79	THR
29	BH	86	ASP
29	BH	91	PHE
29	BH	112	LYS
29	BH	119	ASN
29	BH	122	LEU
29	BH	125	THR
29	BH	129	GLU
29	BH	131	SER
29	BH	137	GLU
29	BH	142	VAL
29	BH	145	ASN
29	BH	146	VAL
30	BI	9	VAL
30	BI	11	LEU
30	BI	12	GLN
30	BI	24	VAL
30	BI	31	GLN
30	BI	34	ASN
30	BI	38	PHE
30	BI	45	LYS
30	BI	50	GLU
30	BI	58	VAL
30	BI	62	TYR
30	BI	67	PHE
30	BI	69	PHE
30	BI	72	LYS
30	BI	82	LYS
30	BI	86	ILE
30	BI	87	LYS

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Mol	Chain	Res	Type
30	BI	96	ASP
30	BI	101	ILE
30	BI	103	ARG
30	BI	108	GLU
30	BI	111	GLN
30	BI	136	MET
31	BJ	4	PHE
31	BJ	5	THR
31	BJ	11	VAL
31	BJ	30	THR
31	BJ	40	HIS
31	BJ	61	LYS
31	BJ	64	VAL
31	BJ	76	HIS
31	BJ	81	ILE
31	BJ	135	GLN
32	BK	3	GLN
32	BK	35	VAL
32	BK	49	ARG
32	BK	58	LEU
32	BK	66	LYS
32	BK	80	ASP
32	BK	91	SER
32	BK	92	GLU
32	BK	117	SER
32	BK	121	GLU
33	BL	2	ARG
33	BL	12	SER
33	BL	17	LYS
33	BL	19	LEU
33	BL	21	ARG
33	BL	40	SER
33	BL	42	SER
33	BL	82	LEU
33	BL	86	GLU
33	BL	89	VAL
33	BL	91	ASP
33	BL	100	ILE
33	BL	115	GLU
33	BL	144	GLU
34	BM	1	MET
34	BM	2	LEU

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Mol	Chain	Res	Type
34	BM	6	ARG
34	BM	10	ARG
34	BM	18	ARG
34	BM	55	ARG
34	BM	69	PRO
34	BM	70	ASP
34	BM	78	LEU
34	BM	106	ASP
34	BM	115	GLU
35	BN	2	ARG
35	BN	6	SER
35	BN	8	ARG
35	BN	15	SER
35	BN	38	LEU
35	BN	69	ARG
35	BN	71	ARG
35	BN	95	THR
35	BN	96	ARG
35	BN	116	VAL
35	BN	117	ASP
35	BN	118	ARG
35	BN	120	GLU
36	BO	2	ASP
36	BO	3	LYS
36	BO	4	LYS
36	BO	9	ARG
36	BO	17	LYS
36	BO	18	LEU
36	BO	24	THR
36	BO	27	VAL
36	BO	45	SER
36	BO	47	VAL
36	BO	65	THR
36	BO	78	VAL
36	BO	88	LYS
36	BO	102	ARG
37	BP	3	ASN
37	BP	19	SER
37	BP	26	VAL
37	BP	68	GLU
37	BP	73	VAL
37	BP	85	SER

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Mol	Chain	Res	Type
37	BP	93	ARG
37	BP	109	ARG
37	BP	110	ILE
37	BP	114	LEU
38	BQ	3	ARG
38	BQ	8	VAL
38	BQ	18	LEU
38	BQ	41	LYS
38	BQ	51	ARG
38	BQ	78	LYS
38	BQ	87	SER
38	BQ	92	ARG
39	BR	1	MET
39	BR	10	LYS
39	BR	14	VAL
39	BR	15	SER
39	BR	16	GLU
39	BR	38	VAL
39	BR	46	GLU
39	BR	48	LYS
39	BR	64	VAL
39	BR	81	LYS
39	BR	85	LYS
39	BR	89	HIS
39	BR	102	SER
40	BS	4	ILE
40	BS	6	LYS
40	BS	7	HIS
40	BS	19	LEU
40	BS	23	LEU
40	BS	30	SER
40	BS	47	VAL
40	BS	53	SER
40	BS	59	GLU
40	BS	69	LEU
40	BS	81	SER
40	BS	96	ILE
40	BS	97	LEU
41	BT	3	ARG
41	BT	5	GLU
41	BT	18	GLU
41	BT	30	ILE

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Mol	Chain	Res	Type
41	BT	36	LYS
41	BT	49	LYS
41	BT	50	LEU
41	BT	53	VAL
41	BT	68	LYS
41	BT	70	HIS
41	BT	76	ARG
41	BT	86	THR
42	BU	9	ASP
42	BU	21	LYS
42	BU	52	LEU
42	BU	61	LYS
42	BU	62	GLU
42	BU	68	SER
42	BU	72	ILE
42	BU	99	ASN
42	BU	100	SER
43	BV	8	VAL
43	BV	10	LYS
43	BV	20	LEU
43	BV	29	ILE
43	BV	41	GLU
43	BV	53	LYS
43	BV	61	LEU
43	BV	65	VAL
43	BV	70	ILE
43	BV	92	VAL
44	BW	20	ARG
44	BW	81	SER
45	BX	48	THR
45	BX	76	GLU
45	BX	77	LYS
46	BY	6	LEU
46	BY	13	GLU
46	BY	16	THR
46	BY	17	GLU
46	BY	22	LEU
46	BY	25	GLN
46	BY	39	GLN
46	BY	59	GLU
47	BZ	3	LYS
47	BZ	10	THR

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Mol	Chain	Res	Type
47	BZ	25	LEU
47	BZ	36	VAL
47	BZ	45	ARG
47	BZ	52	SER
47	BZ	57	VAL
48	B0	10	ARG
48	B0	18	SER
48	B0	23	THR
48	B0	27	SER
48	B0	40	ARG
48	B0	46	ASP
48	B0	53	LYS
49	B1	22	THR
49	B1	28	ARG
49	B1	43	VAL
49	B1	46	HIS
49	B1	47	VAL
49	B1	51	GLU
50	B2	8	SER
50	B2	25	LYS
50	B2	43	THR
50	B2	45	SER
51	B3	13	ARG
51	B3	15	LYS
51	B3	30	ARG
51	B3	31	HIS
52	B4	4	ARG
52	B4	6	SER
52	B4	18	LYS
52	B4	20	ASP
52	B4	37	GLN
53	B5	21	TYR
53	B5	38	PHE
53	B5	39	ASP
53	B5	47	LYS
53	B5	48	LEU
53	B5	50	ILE
53	B5	59	VAL
53	B5	65	LEU
53	B5	73	VAL
53	B5	78	ILE
53	B5	100	ILE

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Mol	Chain	Res	Type
2	CB	14	VAL
2	CB	16	PHE
2	CB	19	GLN
2	CB	20	THR
2	CB	21	ARG
2	CB	23	TRP
2	CB	27	MET
2	CB	43	LEU
2	CB	49	MET
2	CB	50	PHE
2	CB	66	LYS
2	CB	68	LEU
2	CB	72	THR
2	CB	81	LYS
2	CB	88	ASP
2	CB	91	PHE
2	CB	94	HIS
2	CB	95	ARG
2	CB	96	TRP
2	CB	102	THR
2	CB	103	ASN
2	CB	106	THR
2	CB	117	LEU
2	CB	121	SER
2	CB	126	PHE
2	CB	127	ASP
2	CB	130	THR
2	CB	136	MET
2	CB	137	ARG
2	CB	141	LEU
2	CB	144	LEU
2	CB	163	VAL
2	CB	169	GLU
2	CB	179	LEU
2	CB	188	ASP
2	CB	205	ASP
2	CB	207	ILE
2	CB	220	THR
2	CB	222	ARG
3	CC	3	GLN
3	CC	11	ARG
3	CC	18	TRP

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Mol	Chain	Res	Type
3	CC	27	LYS
3	CC	29	PHE
3	CC	33	LEU
3	CC	36	ASP
3	CC	37	PHE
3	CC	43	LEU
3	CC	46	GLU
3	CC	80	LYS
3	CC	103	ILE
3	CC	107	ARG
3	CC	121	THR
3	CC	129	MET
3	CC	131	ARG
3	CC	168	TYR
3	CC	179	ARG
3	CC	190	HIS
3	CC	193	TYR
4	CD	9	LEU
4	CD	10	LYS
4	CD	32	CYS
4	CD	33	LYS
4	CD	54	GLN
4	CD	55	LEU
4	CD	56	ARG
4	CD	58	LYS
4	CD	59	GLN
4	CD	60	LYS
4	CD	70	ARG
4	CD	83	LYS
4	CD	125	VAL
4	CD	138	SER
4	CD	143	VAL
4	CD	152	GLN
4	CD	155	VAL
4	CD	161	LEU
4	CD	163	GLU
4	CD	191	LEU
4	CD	192	SER
4	CD	200	ILE
4	CD	206	LYS
5	CE	10	GLU
5	CE	15	LEU

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Mol	Chain	Res	Type
5	CE	18	VAL
5	CE	19	ASN
5	CE	26	LYS
5	CE	34	THR
5	CE	46	VAL
5	CE	65	GLU
5	CE	69	ARG
5	CE	77	ASN
5	CE	85	VAL
5	CE	114	VAL
5	CE	115	LEU
5	CE	116	GLU
5	CE	120	VAL
5	CE	124	LEU
5	CE	126	LYS
5	CE	131	THR
5	CE	149	SER
6	CF	7	VAL
6	CF	9	MET
6	CF	15	SER
6	CF	17	GLN
6	CF	23	GLU
6	CF	35	LYS
6	CF	36	ILE
6	CF	38	ARG
6	CF	51	ILE
6	CF	53	LYS
6	CF	54	LEU
6	CF	55	HIS
6	CF	63	ASN
6	CF	68	GLN
6	CF	69	GLU
6	CF	75	GLU
6	CF	80	PHE
6	CF	85	ILE
6	CF	93	LYS
6	CF	97	THR
7	CG	4	ARG
7	CG	6	VAL
7	CG	22	LEU
7	CG	23	LEU
7	CG	26	PHE

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Mol	Chain	Res	Type
7	CG	30	LEU
7	CG	47	LEU
7	CG	53	ARG
7	CG	62	PHE
7	CG	66	LEU
7	CG	70	ARG
7	CG	75	VAL
7	CG	78	ARG
7	CG	84	THR
7	CG	92	ARG
7	CG	97	ASN
7	CG	120	LEU
7	CG	123	GLU
7	CG	129	GLU
7	CG	135	VAL
7	CG	140	ASP
7	CG	146	GLU
8	CH	3	MET
8	CH	13	ARG
8	CH	22	LYS
8	CH	25	VAL
8	CH	31	LYS
8	CH	47	GLU
8	CH	49	PHE
8	CH	55	THR
8	CH	67	GLN
8	CH	75	ILE
8	CH	77	ARG
8	CH	87	LYS
8	CH	94	LYS
8	CH	111	MET
8	CH	121	LEU
8	CH	125	ILE
9	CI	11	ARG
9	CI	18	ARG
9	CI	36	GLU
9	CI	38	TYR
9	CI	43	THR
9	CI	45	ARG
9	CI	46	MET
9	CI	55	VAL
9	CI	56	ASP

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Mol	Chain	Res	Type
9	CI	57	MET
9	CI	61	LEU
9	CI	63	LEU
9	CI	68	LYS
9	CI	85	ARG
9	CI	88	MET
9	CI	90	TYR
9	CI	94	LEU
9	CI	97	GLU
9	CI	99	ARG
9	CI	106	ARG
9	CI	129	LYS
10	CJ	22	THR
10	CJ	25	ILE
10	CJ	27	GLU
10	CJ	32	THR
10	CJ	57	VAL
10	CJ	59	LYS
10	CJ	80	THR
10	CJ	84	VAL
10	CJ	87	LEU
10	CJ	89	ARG
10	CJ	92	LEU
11	CK	14	LYS
11	CK	15	GLN
11	CK	31	ILE
11	CK	33	THR
11	CK	52	PHE
11	CK	65	VAL
11	CK	81	ASN
11	CK	96	THR
11	CK	105	PHE
11	CK	106	ARG
11	CK	107	ILE
11	CK	125	LYS
11	CK	126	LYS
11	CK	127	ARG
11	CK	128	ARG
12	CL	4	VAL
12	CL	5	ASN
12	CL	10	LYS
12	CL	12	ARG

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Mol	Chain	Res	Type
12	CL	18	LYS
12	CL	29	GLN
12	CL	44	LYS
12	CL	47	SER
12	CL	58	THR
12	CL	59	ASN
12	CL	61	PHE
12	CL	63	VAL
12	CL	78	SER
12	CL	82	ILE
12	CL	83	ARG
12	CL	89	ASP
12	CL	94	ARG
12	CL	97	THR
12	CL	103	ASP
12	CL	110	ARG
12	CL	114	ARG
12	CL	121	ARG
13	CM	25	VAL
13	CM	29	ARG
13	CM	41	GLU
13	CM	48	LEU
13	CM	59	GLU
13	CM	63	PHE
13	CM	68	ASP
13	CM	80	LEU
13	CM	91	HIS
13	CM	101	ARG
14	CN	4	GLN
14	CN	16	LEU
14	CN	23	LYS
14	CN	26	GLU
14	CN	48	LEU
15	CO	2	SER
15	CO	4	SER
15	CO	17	ARG
15	CO	18	ASP
15	CO	24	SER
15	CO	31	LEU
15	CO	35	GLN
15	CO	39	LEU
15	CO	62	GLN

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Mol	Chain	Res	Type
15	CO	64	ARG
15	CO	66	LEU
15	CO	70	LEU
15	CO	73	LYS
15	CO	85	LEU
15	CO	87	LEU
16	CP	2	VAL
16	CP	18	GLN
16	CP	31	ARG
16	CP	36	VAL
16	CP	46	LYS
16	CP	51	ARG
16	CP	63	GLN
16	CP	66	THR
16	CP	77	GLU
16	CP	80	LYS
17	CQ	5	ILE
17	CQ	14	SER
17	CQ	17	MET
17	CQ	18	GLU
17	CQ	20	SER
17	CQ	29	VAL
17	CQ	40	ARG
17	CQ	51	ASN
17	CQ	52	GLU
17	CQ	55	ILE
17	CQ	65	ARG
17	CQ	75	LEU
17	CQ	78	VAL
17	CQ	79	VAL
17	CQ	81	LYS
18	CR	25	ASP
18	CR	33	ILE
18	CR	36	SER
18	CR	47	THR
18	CR	68	LEU
19	CS	6	LYS
19	CS	11	ILE
19	CS	14	HIS
19	CS	20	GLU
19	CS	28	LYS
19	CS	49	ILE

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Mol	Chain	Res	Type
19	CS	63	THR
20	CT	6	SER
20	CT	14	SER
20	CT	27	MET
20	CT	36	TYR
20	CT	49	LYS
20	CT	58	VAL
20	CT	64	LYS
20	CT	69	LYS
20	CT	76	LYS
20	CT	84	ASN
21	CU	5	LYS
21	CU	10	GLU
21	CU	12	PHE
21	CU	16	LEU
21	CU	19	PHE
21	CU	28	VAL
21	CU	34	ARG
21	CU	36	GLU
21	CU	37	PHE
21	CU	38	TYR
21	CU	42	THR
21	CU	47	ARG
21	CU	53	VAL
24	DC	14	ARG
24	DC	20	VAL
24	DC	58	HIS
24	DC	80	ARG
24	DC	103	TYR
24	DC	110	LEU
24	DC	111	LYS
24	DC	114	ASP
24	DC	130	LEU
24	DC	139	SER
24	DC	156	ARG
24	DC	160	THR
24	DC	167	ARG
24	DC	182	ARG
24	DC	194	GLU
24	DC	195	VAL
24	DC	202	LEU
24	DC	213	TRP

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Mol	Chain	Res	Type
24	DC	221	ARG
24	DC	229	ASP
24	DC	236	GLU
24	DC	246	THR
24	DC	256	LYS
24	DC	262	ARG
24	DC	267	ILE
25	DD	1	MET
25	DD	4	LEU
25	DD	12	THR
25	DD	28	GLU
25	DD	33	ARG
25	DD	67	HIS
25	DD	77	ARG
25	DD	86	GLU
25	DD	95	SER
25	DD	98	VAL
25	DD	118	PHE
25	DD	139	SER
25	DD	141	ARG
25	DD	150	GLN
25	DD	170	VAL
25	DD	181	ASP
25	DD	189	VAL
26	DE	25	GLU
26	DE	44	ARG
26	DE	69	ARG
26	DE	72	SER
26	DE	77	ILE
26	DE	78	TRP
26	DE	83	VAL
26	DE	84	THR
26	DE	91	ASP
26	DE	108	ILE
26	DE	149	ILE
26	DE	170	ARG
26	DE	171	ASP
27	DF	10	ASP
27	DF	14	LYS
27	DF	26	MET
27	DF	28	VAL
27	DF	35	THR

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Mol	Chain	Res	Type
27	DF	46	ASP
27	DF	64	LYS
27	DF	67	ILE
27	DF	74	VAL
27	DF	83	TYR
27	DF	106	ILE
27	DF	117	LEU
27	DF	125	ARG
27	DF	147	ASP
27	DF	152	LEU
27	DF	162	SER
27	DF	178	ARG
28	DG	11	VAL
28	DG	29	LYS
28	DG	30	ASN
28	DG	39	ASP
28	DG	44	LYS
28	DG	45	HIS
28	DG	95	ARG
28	DG	124	GLU
28	DG	127	THR
28	DG	133	LEU
28	DG	155	GLU
28	DG	166	ASP
28	DG	168	VAL
29	DH	7	ASP
29	DH	12	LEU
29	DH	41	LYS
29	DH	42	LYS
29	DH	48	GLU
29	DH	50	ARG
29	DH	53	GLU
29	DH	54	LEU
29	DH	57	LYS
29	DH	62	LEU
29	DH	77	THR
29	DH	78	VAL
29	DH	87	GLU
29	DH	89	LYS
29	DH	94	ILE
29	DH	109	GLU
29	DH	114	GLU

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Mol	Chain	Res	Type
29	DH	116	ARG
29	DH	117	LEU
29	DH	119	ASN
29	DH	121	VAL
29	DH	124	THR
29	DH	125	THR
29	DH	129	GLU
29	DH	142	VAL
30	DI	8	TYR
30	DI	11	LEU
30	DI	17	MET
30	DI	24	VAL
30	DI	31	GLN
30	DI	38	PHE
30	DI	40	LYS
30	DI	68	THR
30	DI	69	PHE
30	DI	72	LYS
30	DI	86	ILE
30	DI	87	LYS
30	DI	96	ASP
30	DI	97	LYS
30	DI	117	MET
30	DI	125	MET
30	DI	127	ARG
30	DI	134	ARG
31	DJ	17	VAL
31	DJ	30	THR
31	DJ	40	HIS
31	DJ	70	THR
31	DJ	80	HIS
31	DJ	88	THR
32	DK	49	ARG
32	DK	90	ASN
32	DK	91	SER
32	DK	92	GLU
32	DK	95	ILE
32	DK	121	GLU
32	DK	122	VAL
33	DL	7	SER
33	DL	47	ARG
33	DL	59	ARG

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Mol	Chain	Res	Type
33	DL	82	LEU
33	DL	100	ILE
33	DL	117	THR
33	DL	118	THR
33	DL	126	ARG
34	DM	31	PHE
34	DM	59	ARG
34	DM	70	ASP
34	DM	78	LEU
34	DM	108	VAL
35	DN	14	SER
35	DN	15	SER
35	DN	33	ILE
35	DN	53	THR
35	DN	63	ARG
35	DN	69	ARG
35	DN	70	THR
35	DN	71	ARG
35	DN	76	VAL
35	DN	106	ASP
35	DN	116	VAL
36	DO	18	LEU
36	DO	28	VAL
36	DO	31	THR
36	DO	45	SER
36	DO	47	VAL
36	DO	74	VAL
36	DO	100	HIS
36	DO	103	VAL
37	DP	4	ILE
37	DP	7	GLN
37	DP	8	LEU
37	DP	26	VAL
37	DP	39	ARG
37	DP	66	ASN
37	DP	75	GLN
37	DP	80	VAL
37	DP	81	VAL
37	DP	110	ILE
38	DQ	9	ILE
38	DQ	29	SER
38	DQ	41	LYS

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Mol	Chain	Res	Type
38	DQ	47	TYR
38	DQ	51	ARG
39	DR	29	THR
39	DR	43	ASN
39	DR	46	GLU
39	DR	48	LYS
39	DR	58	VAL
40	DS	3	THR
40	DS	19	LEU
40	DS	22	ASP
40	DS	23	LEU
40	DS	78	GLU
40	DS	96	ILE
40	DS	109	ASP
41	DT	3	ARG
41	DT	7	LEU
41	DT	16	VAL
41	DT	24	MET
41	DT	27	SER
41	DT	30	ILE
41	DT	31	VAL
41	DT	32	LEU
41	DT	49	LYS
41	DT	70	HIS
41	DT	77	ARG
41	DT	91	GLN
42	DU	7	ARG
42	DU	15	THR
42	DU	18	ASP
42	DU	40	ASN
42	DU	45	HIS
42	DU	53	ASN
42	DU	54	GLN
42	DU	72	ILE
42	DU	81	ASP
42	DU	99	ASN
43	DV	41	GLU
43	DV	42	LEU
43	DV	50	MET
43	DV	53	LYS
43	DV	61	LEU
43	DV	62	THR

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Mol	Chain	Res	Type
43	DV	65	VAL
44	DW	16	SER
44	DW	20	ARG
44	DW	38	VAL
44	DW	39	ARG
45	DX	23	ASN
45	DX	33	LEU
45	DX	40	VAL
45	DX	46	PHE
45	DX	64	ILE
45	DX	71	LEU
46	DY	2	LYS
46	DY	6	LEU
46	DY	13	GLU
46	DY	16	THR
46	DY	23	ARG
46	DY	39	GLN
46	DY	40	SER
46	DY	49	ASP
46	DY	58	ASN
47	DZ	3	LYS
47	DZ	6	LYS
47	DZ	25	LEU
47	DZ	31	ARG
47	DZ	36	VAL
47	DZ	39	GLU
47	DZ	41	THR
47	DZ	45	ARG
48	D0	3	VAL
48	D0	27	SER
48	D0	28	LEU
48	D0	46	ASP
48	D0	52	ARG
49	D1	28	ARG
49	D1	51	GLU
50	D2	1	MET
50	D2	4	THR
50	D2	19	ARG
50	D2	24	THR
50	D2	41	ARG
50	D2	42	LEU
51	D3	8	ARG

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Mol	Chain	Res	Type
51	D3	13	ARG
51	D3	30	ARG
52	D4	3	VAL
52	D4	26	ILE
52	D4	37	GLN

Some sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (32) such sidechains are listed below:

Mol	Chain	Res	Type
2	AB	89	GLN
3	AC	6	HIS
4	AD	54	GLN
5	AE	82	GLN
5	AE	89	HIS
5	AE	122	ASN
7	AG	148	ASN
8	AH	18	GLN
15	AO	46	HIS
16	AP	59	HIS
24	BC	142	HIS
24	BC	243	HIS
25	BD	136	ASN
26	BE	41	GLN
27	BF	21	ASN
29	BH	135	HIS
36	BO	29	HIS
6	CF	37	HIS
15	CO	46	HIS
17	CQ	31	HIS
20	CT	84	ASN
26	DE	24	ASN
29	DH	28	ASN
29	DH	128	HIS
32	DK	29	HIS
36	DO	29	HIS
36	DO	100	HIS
37	DP	66	ASN
39	DR	66	HIS
40	DS	7	HIS
45	DX	34	HIS
48	D0	38	HIS

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	AA	1537/1539 (99%)	304 (19%)	13 (0%)
1	CA	1538/1539 (99%)	303 (19%)	7 (0%)
22	BA	2895/2903 (99%)	518 (17%)	34 (1%)
22	DA	2895/2903 (99%)	536 (18%)	22 (0%)
23	BB	118/119 (99%)	16 (13%)	1 (0%)
23	DB	117/119 (98%)	18 (15%)	0
All	All	9100/9122 (99%)	1695 (18%)	77 (0%)

All (1695) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	AA	4	U
1	AA	5	U
1	AA	7	A
1	AA	9	G
1	AA	13	U
1	AA	22	G
1	AA	32	A
1	AA	39	G
1	AA	47	C
1	AA	48	C
1	AA	51	A
1	AA	70	U
1	AA	71	A
1	AA	72	A
1	AA	75	G
1	AA	76	G
1	AA	77	A
1	AA	81	A
1	AA	82	G
1	AA	83	C
1	AA	85	U
1	AA	86	G
1	AA	89	U
1	AA	90	C
1	AA	91	U
1	AA	95	C
1	AA	97	G
1	AA	108	G
1	AA	109	A
1	AA	111	G

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Mol	Chain	Res	Type
1	AA	116	A
1	AA	117	G
1	AA	120	A
1	AA	121	U
1	AA	130	A
1	AA	131	A
1	AA	137	U
1	AA	138	G
1	AA	143	A
1	AA	144	G
1	AA	159	G
1	AA	162	A
1	AA	163	C
1	AA	168	G
1	AA	182	A
1	AA	183	C
1	AA	188	C
1	AA	195	A
1	AA	205	A
1	AA	210	C
1	AA	212	G
1	AA	226	G
1	AA	240	G
1	AA	245	U
1	AA	247	G
1	AA	251	G
1	AA	263	A
1	AA	266	G
1	AA	267	C
1	AA	289	G
1	AA	298	A
1	AA	321	A
1	AA	328	C
1	AA	329	A
1	AA	330	C
1	AA	332	G
1	AA	341	C
1	AA	346	G
1	AA	347	G
1	AA	352	C
1	AA	353	A
1	AA	354	G

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Mol	Chain	Res	Type
1	AA	356	A
1	AA	365	U
1	AA	367	U
1	AA	372	C
1	AA	384	G
1	AA	398	U
1	AA	406	G
1	AA	411	A
1	AA	412	A
1	AA	413	G
1	AA	414	A
1	AA	421	U
1	AA	422	C
1	AA	423	G
1	AA	429	U
1	AA	430	A
1	AA	435	A
1	AA	453	G
1	AA	454	G
1	AA	456	A
1	AA	457	G
1	AA	458	U
1	AA	462	G
1	AA	463	U
1	AA	465	A
1	AA	466	A
1	AA	467	U
1	AA	468	A
1	AA	474	G
1	AA	479	U
1	AA	481	G
1	AA	482	A
1	AA	485	U
1	AA	486	U
1	AA	495	A
1	AA	499	A
1	AA	505	G
1	AA	511	C
1	AA	518	C
1	AA	524	G
1	AA	527	G
1	AA	530	G

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Mol	Chain	Res	Type
1	AA	532	A
1	AA	533	A
1	AA	541	G
1	AA	547	A
1	AA	559	A
1	AA	562	U
1	AA	573	A
1	AA	576	C
1	AA	577	G
1	AA	579	A
1	AA	615	G
1	AA	650	G
1	AA	653	U
1	AA	657	U
1	AA	665	A
1	AA	671	G
1	AA	703	G
1	AA	717	U
1	AA	721	G
1	AA	723	U
1	AA	731	G
1	AA	733	G
1	AA	734	G
1	AA	747	A
1	AA	755	G
1	AA	772	U
1	AA	773	G
1	AA	774	G
1	AA	777	A
1	AA	792	A
1	AA	793	U
1	AA	794	A
1	AA	802	A
1	AA	814	A
1	AA	815	A
1	AA	817	C
1	AA	821	G
1	AA	828	U
1	AA	841	C
1	AA	842	U
1	AA	843	U
1	AA	845	A

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Mol	Chain	Res	Type
1	AA	846	G
1	AA	849	G
1	AA	853	C
1	AA	859	G
1	AA	860	A
1	AA	868	C
1	AA	902	G
1	AA	906	A
1	AA	914	A
1	AA	925	G
1	AA	926	G
1	AA	927	G
1	AA	932	C
1	AA	934	C
1	AA	935	A
1	AA	960	U
1	AA	964	A
1	AA	965	U
1	AA	966	G
1	AA	968	A
1	AA	969	A
1	AA	971	G
1	AA	972	C
1	AA	975	A
1	AA	976	G
1	AA	977	A
1	AA	983	A
1	AA	992	U
1	AA	993	G
1	AA	1004	A
1	AA	1008	U
1	AA	1009	U
1	AA	1024	G
1	AA	1025	U
1	AA	1026	G
1	AA	1027	C
1	AA	1028	C
1	AA	1030	U
1	AA	1031	C
1	AA	1032	G
1	AA	1033	G
1	AA	1034	G

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Mol	Chain	Res	Type
1	AA	1036	A
1	AA	1037	C
1	AA	1043	G
1	AA	1044	A
1	AA	1047	G
1	AA	1050	G
1	AA	1054	C
1	AA	1055	A
1	AA	1056	U
1	AA	1061	G
1	AA	1065	U
1	AA	1066	C
1	AA	1086	U
1	AA	1089	G
1	AA	1091	U
1	AA	1094	G
1	AA	1095	U
1	AA	1098	C
1	AA	1101	A
1	AA	1108	G
1	AA	1113	C
1	AA	1124	G
1	AA	1125	U
1	AA	1133	G
1	AA	1136	C
1	AA	1137	C
1	AA	1139	G
1	AA	1140	C
1	AA	1141	C
1	AA	1142	G
1	AA	1145	A
1	AA	1146	A
1	AA	1149	C
1	AA	1151	A
1	AA	1152	A
1	AA	1159	U
1	AA	1160	G
1	AA	1161	C
1	AA	1168	U
1	AA	1181	G
1	AA	1183	U
1	AA	1184	G

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Mol	Chain	Res	Type
1	AA	1193	G
1	AA	1196	A
1	AA	1197	A
1	AA	1202	U
1	AA	1212	U
1	AA	1213	A
1	AA	1214	C
1	AA	1215	G
1	AA	1221	G
1	AA	1227	A
1	AA	1228	C
1	AA	1238	A
1	AA	1239	A
1	AA	1253	G
1	AA	1256	A
1	AA	1261	A
1	AA	1280	A
1	AA	1286	U
1	AA	1287	A
1	AA	1299	A
1	AA	1300	G
1	AA	1302	C
1	AA	1304	G
1	AA	1305	G
1	AA	1317	C
1	AA	1320	C
1	AA	1321	U
1	AA	1322	C
1	AA	1323	G
1	AA	1329	A
1	AA	1332	A
1	AA	1335	U
1	AA	1336	C
1	AA	1337	G
1	AA	1338	G
1	AA	1346	A
1	AA	1353	G
1	AA	1363	A
1	AA	1364	U
1	AA	1368	A
1	AA	1370	G
1	AA	1397	C

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Mol	Chain	Res	Type
1	AA	1398	A
1	AA	1401	G
1	AA	1441	A
1	AA	1442	G
1	AA	1446	A
1	AA	1452	C
1	AA	1453	G
1	AA	1454	G
1	AA	1472	U
1	AA	1493	A
1	AA	1497	G
1	AA	1499	A
1	AA	1505	G
1	AA	1506	U
1	AA	1517	G
1	AA	1526	G
1	AA	1529	G
1	AA	1530	G
1	AA	1533	C
1	AA	1534	A
1	AA	1535	C
1	AA	1538	C
22	BA	10	A
22	BA	12	U
22	BA	13	A
22	BA	27	G
22	BA	34	U
22	BA	46	G
22	BA	57	C
22	BA	58	G
22	BA	63	A
22	BA	71	A
22	BA	74	A
22	BA	75	G
22	BA	101	A
22	BA	102	U
22	BA	118	A
22	BA	119	A
22	BA	120	U
22	BA	122	G
22	BA	138	U
22	BA	139	U

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Mol	Chain	Res	Type
22	BA	140	C
22	BA	141	G
22	BA	142	A
22	BA	181	A
22	BA	196	A
22	BA	208	C
22	BA	215	G
22	BA	216	A
22	BA	221	A
22	BA	222	A
22	BA	248	G
22	BA	255	A
22	BA	265	A
22	BA	266	G
22	BA	272	A
22	BA	273	G
22	BA	276	U
22	BA	277	G
22	BA	279	A
22	BA	302	C
22	BA	310	A
22	BA	311	A
22	BA	325	G
22	BA	329	G
22	BA	330	A
22	BA	343	C
22	BA	353	C
22	BA	361	G
22	BA	362	A
22	BA	371	A
22	BA	372	G
22	BA	380	G
22	BA	386	G
22	BA	396	G
22	BA	404	A
22	BA	405	U
22	BA	411	G
22	BA	412	A
22	BA	420	C
22	BA	424	G
22	BA	442	G
22	BA	451	U

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Mol	Chain	Res	Type
22	BA	455	C
22	BA	457	A
22	BA	467	G
22	BA	480	A
22	BA	481	G
22	BA	491	G
22	BA	504	A
22	BA	505	A
22	BA	509	C
22	BA	510	C
22	BA	528	A
22	BA	531	C
22	BA	532	A
22	BA	533	G
22	BA	543	G
22	BA	544	C
22	BA	546	U
22	BA	547	A
22	BA	548	G
22	BA	549	G
22	BA	563	A
22	BA	573	U
22	BA	575	A
22	BA	583	G
22	BA	586	A
22	BA	603	A
22	BA	613	A
22	BA	614	A
22	BA	615	U
22	BA	622	G
22	BA	627	A
22	BA	631	A
22	BA	637	A
22	BA	645	C
22	BA	646	U
22	BA	647	G
22	BA	654	A
22	BA	655	A
22	BA	686	U
22	BA	694	U
22	BA	716	A
22	BA	722	A

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Mol	Chain	Res	Type
22	BA	730	A
22	BA	738	G
22	BA	747	U
22	BA	754	U
22	BA	759	G
22	BA	764	A
22	BA	765	C
22	BA	770	G
22	BA	775	G
22	BA	776	G
22	BA	782	A
22	BA	784	G
22	BA	785	G
22	BA	791	C
22	BA	792	A
22	BA	802	A
22	BA	805	G
22	BA	810	U
22	BA	812	C
22	BA	819	A
22	BA	827	U
22	BA	828	U
22	BA	830	G
22	BA	843	G
22	BA	845	A
22	BA	846	U
22	BA	847	U
22	BA	855	G
22	BA	858	G
22	BA	859	G
22	BA	860	U
22	BA	861	A
22	BA	866	A
22	BA	869	G
22	BA	871	U
22	BA	878	A
22	BA	879	G
22	BA	885	C
22	BA	896	A
22	BA	910	A
22	BA	914	G
22	BA	915	C

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Mol	Chain	Res	Type
22	BA	932	U
22	BA	934	U
22	BA	941	A
22	BA	946	C
22	BA	959	A
22	BA	961	C
22	BA	974	G
22	BA	983	A
22	BA	984	A
22	BA	985	C
22	BA	995	C
22	BA	996	A
22	BA	1005	C
22	BA	1012	U
22	BA	1013	C
22	BA	1022	G
22	BA	1026	G
22	BA	1033	U
22	BA	1040	A
22	BA	1046	A
22	BA	1047	G
22	BA	1057	A
22	BA	1061	U
22	BA	1062	G
22	BA	1063	G
22	BA	1066	U
22	BA	1067	A
22	BA	1068	G
22	BA	1070	A
22	BA	1071	G
22	BA	1072	C
22	BA	1073	A
22	BA	1074	G
22	BA	1075	C
22	BA	1077	A
22	BA	1081	U
22	BA	1087	G
22	BA	1088	A
22	BA	1089	A
22	BA	1091	G
22	BA	1092	C
22	BA	1097	U

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Mol	Chain	Res	Type
22	BA	1098	A
22	BA	1100	C
22	BA	1104	C
22	BA	1112	G
22	BA	1132	U
22	BA	1133	A
22	BA	1135	C
22	BA	1136	G
22	BA	1138	G
22	BA	1142	A
22	BA	1144	A
22	BA	1156	A
22	BA	1168	G
22	BA	1171	G
22	BA	1172	C
22	BA	1173	U
22	BA	1174	U
22	BA	1175	A
22	BA	1176	U
22	BA	1178	C
22	BA	1179	G
22	BA	1180	U
22	BA	1186	G
22	BA	1218	G
22	BA	1221	C
22	BA	1238	G
22	BA	1244	A
22	BA	1253	A
22	BA	1256	G
22	BA	1266	G
22	BA	1271	G
22	BA	1272	A
22	BA	1273	U
22	BA	1281	G
22	BA	1294	U
22	BA	1300	G
22	BA	1301	A
22	BA	1303	G
22	BA	1327	A
22	BA	1332	G
22	BA	1334	G
22	BA	1342	A

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Mol	Chain	Res	Type
22	BA	1345	C
22	BA	1352	U
22	BA	1353	A
22	BA	1359	A
22	BA	1365	A
22	BA	1368	G
22	BA	1374	G
22	BA	1378	A
22	BA	1379	U
22	BA	1383	A
22	BA	1386	C
22	BA	1397	U
22	BA	1405	U
22	BA	1407	G
22	BA	1416	G
22	BA	1419	A
22	BA	1420	A
22	BA	1427	A
22	BA	1428	C
22	BA	1435	G
22	BA	1441	G
22	BA	1451	C
22	BA	1452	G
22	BA	1453	A
22	BA	1467	U
22	BA	1478	G
22	BA	1482	G
22	BA	1483	G
22	BA	1493	C
22	BA	1504	A
22	BA	1508	A
22	BA	1510	G
22	BA	1515	A
22	BA	1523	U
22	BA	1527	G
22	BA	1532	A
22	BA	1533	C
22	BA	1534	U
22	BA	1535	A
22	BA	1536	C
22	BA	1554	U
22	BA	1558	C

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Mol	Chain	Res	Type
22	BA	1569	A
22	BA	1578	U
22	BA	1583	A
22	BA	1584	U
22	BA	1585	C
22	BA	1606	C
22	BA	1607	C
22	BA	1608	A
22	BA	1610	A
22	BA	1619	G
22	BA	1647	U
22	BA	1648	U
22	BA	1649	G
22	BA	1669	A
22	BA	1674	G
22	BA	1682	G
22	BA	1715	G
22	BA	1729	U
22	BA	1730	C
22	BA	1738	G
22	BA	1739	A
22	BA	1744	A
22	BA	1758	U
22	BA	1764	C
22	BA	1773	A
22	BA	1776	G
22	BA	1791	A
22	BA	1800	C
22	BA	1801	A
22	BA	1802	A
22	BA	1808	A
22	BA	1816	C
22	BA	1829	A
22	BA	1847	A
22	BA	1870	C
22	BA	1872	A
22	BA	1873	G
22	BA	1882	U
22	BA	1884	G
22	BA	1885	A
22	BA	1890	A
22	BA	1906	G

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Mol	Chain	Res	Type
22	BA	1909	C
22	BA	1912	A
22	BA	1914	C
22	BA	1915	U
22	BA	1916	A
22	BA	1917	U
22	BA	1919	A
22	BA	1921	G
22	BA	1922	G
22	BA	1923	U
22	BA	1925	C
22	BA	1926	U
22	BA	1927	A
22	BA	1929	G
22	BA	1930	G
22	BA	1931	U
22	BA	1932	A
22	BA	1936	A
22	BA	1937	A
22	BA	1938	A
22	BA	1955	U
22	BA	1960	A
22	BA	1963	U
22	BA	1964	G
22	BA	1965	C
22	BA	1967	C
22	BA	1970	A
22	BA	1972	G
22	BA	1976	U
22	BA	1977	A
22	BA	1991	U
22	BA	1992	G
22	BA	1993	U
22	BA	1997	C
22	BA	2017	U
22	BA	2022	U
22	BA	2023	C
22	BA	2031	A
22	BA	2033	A
22	BA	2042	A
22	BA	2043	C
22	BA	2054	A

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Mol	Chain	Res	Type
22	BA	2055	C
22	BA	2056	G
22	BA	2060	A
22	BA	2061	G
22	BA	2062	A
22	BA	2063	C
22	BA	2069	G
22	BA	2093	G
22	BA	2096	C
22	BA	2102	G
22	BA	2110	G
22	BA	2111	U
22	BA	2112	G
22	BA	2113	U
22	BA	2115	G
22	BA	2116	G
22	BA	2117	A
22	BA	2118	U
22	BA	2119	A
22	BA	2122	U
22	BA	2123	G
22	BA	2126	A
22	BA	2127	G
22	BA	2128	G
22	BA	2132	U
22	BA	2133	G
22	BA	2136	G
22	BA	2147	A
22	BA	2148	G
22	BA	2157	G
22	BA	2162	G
22	BA	2164	C
22	BA	2165	C
22	BA	2167	U
22	BA	2169	A
22	BA	2170	A
22	BA	2171	A
22	BA	2172	U
22	BA	2173	A
22	BA	2178	C
22	BA	2181	U
22	BA	2187	U

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Mol	Chain	Res	Type
22	BA	2190	G
22	BA	2198	A
22	BA	2203	U
22	BA	2204	G
22	BA	2211	A
22	BA	2212	A
22	BA	2220	U
22	BA	2225	A
22	BA	2233	U
22	BA	2235	G
22	BA	2238	G
22	BA	2239	G
22	BA	2245	U
22	BA	2268	A
22	BA	2278	A
22	BA	2280	G
22	BA	2283	C
22	BA	2287	A
22	BA	2295	C
22	BA	2296	U
22	BA	2297	A
22	BA	2305	U
22	BA	2308	G
22	BA	2311	A
22	BA	2325	G
22	BA	2326	C
22	BA	2327	A
22	BA	2333	A
22	BA	2335	A
22	BA	2345	G
22	BA	2347	C
22	BA	2350	C
22	BA	2357	G
22	BA	2361	G
22	BA	2383	G
22	BA	2385	C
22	BA	2402	U
22	BA	2406	A
22	BA	2420	C
22	BA	2421	G
22	BA	2424	C
22	BA	2425	A

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Mol	Chain	Res	Type
22	BA	2426	A
22	BA	2429	G
22	BA	2430	A
22	BA	2434	A
22	BA	2435	A
22	BA	2441	U
22	BA	2448	A
22	BA	2453	A
22	BA	2465	C
22	BA	2476	A
22	BA	2478	A
22	BA	2484	G
22	BA	2486	C
22	BA	2490	G
22	BA	2491	U
22	BA	2501	C
22	BA	2502	G
22	BA	2504	U
22	BA	2505	G
22	BA	2518	A
22	BA	2520	C
22	BA	2521	C
22	BA	2522	U
22	BA	2525	G
22	BA	2529	G
22	BA	2554	U
22	BA	2556	C
22	BA	2566	A
22	BA	2567	G
22	BA	2573	C
22	BA	2576	G
22	BA	2579	C
22	BA	2582	G
22	BA	2585	U
22	BA	2586	U
22	BA	2602	A
22	BA	2603	G
22	BA	2604	U
22	BA	2609	U
22	BA	2613	U
22	BA	2629	U
22	BA	2638	G

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Mol	Chain	Res	Type
22	BA	2661	G
22	BA	2663	G
22	BA	2689	U
22	BA	2690	U
22	BA	2707	U
22	BA	2714	G
22	BA	2716	C
22	BA	2718	G
22	BA	2721	A
22	BA	2726	A
22	BA	2729	G
22	BA	2733	A
22	BA	2744	G
22	BA	2748	A
22	BA	2757	A
22	BA	2765	A
22	BA	2769	U
22	BA	2778	A
22	BA	2791	G
22	BA	2798	U
22	BA	2799	A
22	BA	2800	A
22	BA	2811	G
22	BA	2820	A
22	BA	2821	A
22	BA	2834	G
22	BA	2850	A
22	BA	2867	G
22	BA	2873	A
22	BA	2874	C
22	BA	2880	C
22	BA	2883	A
22	BA	2884	U
22	BA	2885	G
22	BA	2886	A
22	BA	2891	U
23	BB	13	G
23	BB	15	A
23	BB	16	G
23	BB	25	U
23	BB	28	C
23	BB	35	C

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Mol	Chain	Res	Type
23	BB	42	C
23	BB	45	A
23	BB	56	G
23	BB	65	U
23	BB	67	G
23	BB	89	U
23	BB	90	C
23	BB	98	G
23	BB	99	A
23	BB	109	A
1	CA	5	U
1	CA	9	G
1	CA	22	G
1	CA	32	A
1	CA	39	G
1	CA	45	G
1	CA	47	C
1	CA	48	C
1	CA	50	A
1	CA	51	A
1	CA	70	U
1	CA	71	A
1	CA	74	A
1	CA	80	A
1	CA	83	C
1	CA	84	U
1	CA	85	U
1	CA	87	C
1	CA	88	U
1	CA	91	U
1	CA	94	G
1	CA	95	C
1	CA	108	G
1	CA	115	G
1	CA	116	A
1	CA	117	G
1	CA	119	A
1	CA	120	A
1	CA	121	U
1	CA	122	G
1	CA	130	A
1	CA	131	A

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Mol	Chain	Res	Type
1	CA	137	U
1	CA	143	A
1	CA	144	G
1	CA	149	A
1	CA	155	A
1	CA	156	C
1	CA	159	G
1	CA	182	A
1	CA	183	C
1	CA	189	A
1	CA	195	A
1	CA	197	A
1	CA	201	G
1	CA	204	G
1	CA	207	C
1	CA	208	U
1	CA	209	U
1	CA	210	C
1	CA	211	G
1	CA	212	G
1	CA	240	G
1	CA	241	G
1	CA	245	U
1	CA	247	G
1	CA	251	G
1	CA	266	G
1	CA	267	C
1	CA	289	G
1	CA	298	A
1	CA	321	A
1	CA	328	C
1	CA	329	A
1	CA	330	C
1	CA	332	G
1	CA	339	C
1	CA	352	C
1	CA	354	G
1	CA	367	U
1	CA	370	C
1	CA	372	C
1	CA	378	G
1	CA	382	A

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Mol	Chain	Res	Type
1	CA	384	G
1	CA	389	A
1	CA	390	U
1	CA	398	U
1	CA	399	G
1	CA	404	G
1	CA	406	G
1	CA	411	A
1	CA	412	A
1	CA	413	G
1	CA	421	U
1	CA	422	C
1	CA	424	G
1	CA	429	U
1	CA	430	A
1	CA	438	U
1	CA	441	A
1	CA	457	G
1	CA	458	U
1	CA	459	A
1	CA	466	A
1	CA	467	U
1	CA	468	A
1	CA	469	C
1	CA	478	A
1	CA	479	U
1	CA	481	G
1	CA	484	G
1	CA	485	U
1	CA	486	U
1	CA	494	G
1	CA	495	A
1	CA	498	A
1	CA	499	A
1	CA	500	G
1	CA	509	A
1	CA	511	C
1	CA	518	C
1	CA	519	C
1	CA	527	G
1	CA	532	A
1	CA	533	A

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Mol	Chain	Res	Type
1	CA	547	A
1	CA	550	G
1	CA	558	G
1	CA	559	A
1	CA	564	C
1	CA	568	G
1	CA	572	A
1	CA	573	A
1	CA	576	C
1	CA	577	G
1	CA	580	C
1	CA	581	G
1	CA	615	G
1	CA	621	A
1	CA	622	A
1	CA	650	G
1	CA	653	U
1	CA	656	G
1	CA	665	A
1	CA	666	G
1	CA	687	A
1	CA	695	A
1	CA	702	A
1	CA	703	G
1	CA	717	U
1	CA	719	C
1	CA	721	G
1	CA	723	U
1	CA	724	G
1	CA	731	G
1	CA	747	A
1	CA	752	G
1	CA	755	G
1	CA	758	C
1	CA	778	G
1	CA	792	A
1	CA	793	U
1	CA	794	A
1	CA	799	G
1	CA	802	A
1	CA	804	U
1	CA	810	C

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Mol	Chain	Res	Type
1	CA	812	G
1	CA	815	A
1	CA	817	C
1	CA	819	A
1	CA	827	U
1	CA	828	U
1	CA	841	C
1	CA	843	U
1	CA	844	G
1	CA	845	A
1	CA	846	G
1	CA	849	G
1	CA	853	C
1	CA	855	U
1	CA	859	G
1	CA	872	A
1	CA	873	A
1	CA	876	C
1	CA	885	G
1	CA	899	C
1	CA	902	G
1	CA	909	A
1	CA	914	A
1	CA	916	U
1	CA	919	A
1	CA	922	G
1	CA	926	G
1	CA	931	C
1	CA	934	C
1	CA	935	A
1	CA	960	U
1	CA	966	G
1	CA	969	A
1	CA	971	G
1	CA	975	A
1	CA	976	G
1	CA	977	A
1	CA	981	U
1	CA	983	A
1	CA	987	G
1	CA	993	G
1	CA	1004	A

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Mol	Chain	Res	Type
1	CA	1008	U
1	CA	1009	U
1	CA	1017	U
1	CA	1018	G
1	CA	1022	A
1	CA	1025	U
1	CA	1026	G
1	CA	1027	C
1	CA	1028	C
1	CA	1030	U
1	CA	1031	C
1	CA	1032	G
1	CA	1033	G
1	CA	1034	G
1	CA	1037	C
1	CA	1043	G
1	CA	1044	A
1	CA	1050	G
1	CA	1053	G
1	CA	1054	C
1	CA	1065	U
1	CA	1069	C
1	CA	1072	G
1	CA	1073	U
1	CA	1080	A
1	CA	1086	U
1	CA	1094	G
1	CA	1095	U
1	CA	1101	A
1	CA	1113	C
1	CA	1124	G
1	CA	1125	U
1	CA	1126	U
1	CA	1133	G
1	CA	1134	G
1	CA	1136	C
1	CA	1137	C
1	CA	1139	G
1	CA	1140	C
1	CA	1141	C
1	CA	1142	G
1	CA	1145	A

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Mol	Chain	Res	Type
1	CA	1154	G
1	CA	1159	U
1	CA	1160	G
1	CA	1161	C
1	CA	1184	G
1	CA	1196	A
1	CA	1202	U
1	CA	1212	U
1	CA	1213	A
1	CA	1238	A
1	CA	1240	U
1	CA	1241	G
1	CA	1256	A
1	CA	1257	A
1	CA	1260	G
1	CA	1275	A
1	CA	1280	A
1	CA	1286	U
1	CA	1287	A
1	CA	1293	C
1	CA	1299	A
1	CA	1300	G
1	CA	1302	C
1	CA	1305	G
1	CA	1317	C
1	CA	1318	A
1	CA	1322	C
1	CA	1337	G
1	CA	1338	G
1	CA	1346	A
1	CA	1362	A
1	CA	1363	A
1	CA	1365	G
1	CA	1377	A
1	CA	1379	G
1	CA	1398	A
1	CA	1419	G
1	CA	1429	A
1	CA	1441	A
1	CA	1446	A
1	CA	1452	C
1	CA	1454	G

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Mol	Chain	Res	Type
1	CA	1475	G
1	CA	1480	A
1	CA	1487	G
1	CA	1491	G
1	CA	1493	A
1	CA	1494	G
1	CA	1497	G
1	CA	1503	A
1	CA	1505	G
1	CA	1506	U
1	CA	1507	A
1	CA	1517	G
1	CA	1518	A
1	CA	1520	C
1	CA	1529	G
1	CA	1530	G
1	CA	1531	A
1	CA	1533	C
1	CA	1535	C
22	DA	10	A
22	DA	11	C
22	DA	12	U
22	DA	34	U
22	DA	39	G
22	DA	42	A
22	DA	46	G
22	DA	55	G
22	DA	58	G
22	DA	61	C
22	DA	66	C
22	DA	71	A
22	DA	73	A
22	DA	74	A
22	DA	75	G
22	DA	80	G
22	DA	82	U
22	DA	84	A
22	DA	96	C
22	DA	98	G
22	DA	101	A
22	DA	103	A
22	DA	118	A

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Mol	Chain	Res	Type
22	DA	119	A
22	DA	120	U
22	DA	125	A
22	DA	138	U
22	DA	139	U
22	DA	140	C
22	DA	141	G
22	DA	142	A
22	DA	162	U
22	DA	166	U
22	DA	178	G
22	DA	181	A
22	DA	196	A
22	DA	199	A
22	DA	206	U
22	DA	215	G
22	DA	216	A
22	DA	222	A
22	DA	223	A
22	DA	224	U
22	DA	225	C
22	DA	229	C
22	DA	233	A
22	DA	248	G
22	DA	249	C
22	DA	255	A
22	DA	256	A
22	DA	264	C
22	DA	266	G
22	DA	271	G
22	DA	272	A
22	DA	276	U
22	DA	279	A
22	DA	284	U
22	DA	285	G
22	DA	294	A
22	DA	311	A
22	DA	317	G
22	DA	329	G
22	DA	330	A
22	DA	361	G
22	DA	362	A

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Mol	Chain	Res	Type
22	DA	367	G
22	DA	371	A
22	DA	372	G
22	DA	380	G
22	DA	383	C
22	DA	385	C
22	DA	386	G
22	DA	387	U
22	DA	396	G
22	DA	399	U
22	DA	405	U
22	DA	411	G
22	DA	424	G
22	DA	430	A
22	DA	435	C
22	DA	447	A
22	DA	451	U
22	DA	455	C
22	DA	480	A
22	DA	481	G
22	DA	490	C
22	DA	491	G
22	DA	504	A
22	DA	505	A
22	DA	508	A
22	DA	510	C
22	DA	528	A
22	DA	532	A
22	DA	543	G
22	DA	544	C
22	DA	546	U
22	DA	547	A
22	DA	548	G
22	DA	549	G
22	DA	550	C
22	DA	563	A
22	DA	569	U
22	DA	573	U
22	DA	575	A
22	DA	586	A
22	DA	603	A
22	DA	615	U

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Mol	Chain	Res	Type
22	DA	627	A
22	DA	628	G
22	DA	630	G
22	DA	631	A
22	DA	637	A
22	DA	645	C
22	DA	646	U
22	DA	647	G
22	DA	648	G
22	DA	654	A
22	DA	655	A
22	DA	685	A
22	DA	686	U
22	DA	695	G
22	DA	717	C
22	DA	726	G
22	DA	730	A
22	DA	740	C
22	DA	747	U
22	DA	748	G
22	DA	751	A
22	DA	752	A
22	DA	757	G
22	DA	762	U
22	DA	764	A
22	DA	771	G
22	DA	775	G
22	DA	776	G
22	DA	777	G
22	DA	782	A
22	DA	784	G
22	DA	785	G
22	DA	789	A
22	DA	792	A
22	DA	798	G
22	DA	802	A
22	DA	805	G
22	DA	810	U
22	DA	812	C
22	DA	819	A
22	DA	826	U
22	DA	827	U

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Mol	Chain	Res	Type
22	DA	828	U
22	DA	829	A
22	DA	830	G
22	DA	844	A
22	DA	845	A
22	DA	846	U
22	DA	858	G
22	DA	859	G
22	DA	878	A
22	DA	882	G
22	DA	896	A
22	DA	897	C
22	DA	902	C
22	DA	910	A
22	DA	914	G
22	DA	922	C
22	DA	932	U
22	DA	934	U
22	DA	941	A
22	DA	946	C
22	DA	961	C
22	DA	974	G
22	DA	982	C
22	DA	983	A
22	DA	990	A
22	DA	995	C
22	DA	996	A
22	DA	1012	U
22	DA	1013	C
22	DA	1022	G
22	DA	1025	G
22	DA	1026	G
22	DA	1033	U
22	DA	1046	A
22	DA	1047	G
22	DA	1051	G
22	DA	1053	C
22	DA	1058	U
22	DA	1060	U
22	DA	1061	U
22	DA	1062	G
22	DA	1065	U

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Mol	Chain	Res	Type
22	DA	1066	U
22	DA	1068	G
22	DA	1070	A
22	DA	1072	C
22	DA	1075	C
22	DA	1079	C
22	DA	1088	A
22	DA	1089	A
22	DA	1090	A
22	DA	1092	C
22	DA	1094	U
22	DA	1097	U
22	DA	1100	C
22	DA	1104	C
22	DA	1110	G
22	DA	1111	A
22	DA	1112	G
22	DA	1122	G
22	DA	1128	G
22	DA	1132	U
22	DA	1133	A
22	DA	1134	A
22	DA	1135	C
22	DA	1136	G
22	DA	1139	G
22	DA	1153	C
22	DA	1155	A
22	DA	1171	G
22	DA	1172	C
22	DA	1175	A
22	DA	1176	U
22	DA	1180	U
22	DA	1186	G
22	DA	1204	A
22	DA	1212	G
22	DA	1236	G
22	DA	1238	G
22	DA	1241	A
22	DA	1250	G
22	DA	1253	A
22	DA	1256	G
22	DA	1266	G

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Mol	Chain	Res	Type
22	DA	1268	A
22	DA	1271	G
22	DA	1272	A
22	DA	1275	A
22	DA	1276	A
22	DA	1300	G
22	DA	1301	A
22	DA	1305	C
22	DA	1321	A
22	DA	1343	G
22	DA	1345	C
22	DA	1352	U
22	DA	1355	G
22	DA	1359	A
22	DA	1365	A
22	DA	1372	U
22	DA	1374	G
22	DA	1376	C
22	DA	1378	A
22	DA	1379	U
22	DA	1380	G
22	DA	1383	A
22	DA	1386	C
22	DA	1387	A
22	DA	1395	A
22	DA	1403	A
22	DA	1411	U
22	DA	1414	C
22	DA	1416	G
22	DA	1427	A
22	DA	1428	C
22	DA	1434	A
22	DA	1451	C
22	DA	1452	G
22	DA	1455	G
22	DA	1458	U
22	DA	1468	U
22	DA	1471	G
22	DA	1478	G
22	DA	1482	G
22	DA	1493	C
22	DA	1504	A

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Mol	Chain	Res	Type
22	DA	1509	A
22	DA	1510	G
22	DA	1515	A
22	DA	1523	U
22	DA	1533	C
22	DA	1534	U
22	DA	1535	A
22	DA	1536	C
22	DA	1537	G
22	DA	1560	G
22	DA	1565	C
22	DA	1569	A
22	DA	1578	U
22	DA	1581	G
22	DA	1583	A
22	DA	1584	U
22	DA	1585	C
22	DA	1602	U
22	DA	1603	A
22	DA	1606	C
22	DA	1607	C
22	DA	1608	A
22	DA	1610	A
22	DA	1613	G
22	DA	1616	A
22	DA	1646	C
22	DA	1647	U
22	DA	1648	U
22	DA	1649	G
22	DA	1651	G
22	DA	1661	G
22	DA	1664	A
22	DA	1674	G
22	DA	1694	C
22	DA	1695	G
22	DA	1705	A
22	DA	1715	G
22	DA	1729	U
22	DA	1730	C
22	DA	1732	C
22	DA	1738	G
22	DA	1744	A

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Mol	Chain	Res	Type
22	DA	1750	G
22	DA	1758	U
22	DA	1764	C
22	DA	1773	A
22	DA	1782	U
22	DA	1794	A
22	DA	1798	U
22	DA	1800	C
22	DA	1801	A
22	DA	1802	A
22	DA	1808	A
22	DA	1816	C
22	DA	1823	G
22	DA	1827	U
22	DA	1828	G
22	DA	1829	A
22	DA	1847	A
22	DA	1848	A
22	DA	1858	A
22	DA	1869	G
22	DA	1870	C
22	DA	1871	A
22	DA	1873	G
22	DA	1874	C
22	DA	1889	A
22	DA	1903	G
22	DA	1906	G
22	DA	1914	C
22	DA	1927	A
22	DA	1929	G
22	DA	1930	G
22	DA	1934	C
22	DA	1937	A
22	DA	1955	U
22	DA	1966	A
22	DA	1967	C
22	DA	1970	A
22	DA	1972	G
22	DA	1991	U
22	DA	1993	U
22	DA	1997	C
22	DA	2018	G

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Mol	Chain	Res	Type
22	DA	2020	A
22	DA	2022	U
22	DA	2023	C
22	DA	2030	A
22	DA	2031	A
22	DA	2033	A
22	DA	2043	C
22	DA	2049	G
22	DA	2055	C
22	DA	2056	G
22	DA	2057	G
22	DA	2060	A
22	DA	2061	G
22	DA	2062	A
22	DA	2069	G
22	DA	2073	C
22	DA	2083	G
22	DA	2092	U
22	DA	2093	G
22	DA	2108	A
22	DA	2110	G
22	DA	2111	U
22	DA	2112	G
22	DA	2113	U
22	DA	2115	G
22	DA	2116	G
22	DA	2117	A
22	DA	2118	U
22	DA	2119	A
22	DA	2125	G
22	DA	2126	A
22	DA	2127	G
22	DA	2128	G
22	DA	2131	U
22	DA	2132	U
22	DA	2133	G
22	DA	2135	A
22	DA	2137	U
22	DA	2146	C
22	DA	2147	A
22	DA	2149	U
22	DA	2158	A

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Mol	Chain	Res	Type
22	DA	2162	G
22	DA	2163	A
22	DA	2164	C
22	DA	2165	C
22	DA	2169	A
22	DA	2170	A
22	DA	2171	A
22	DA	2172	U
22	DA	2173	A
22	DA	2178	C
22	DA	2184	A
22	DA	2189	U
22	DA	2190	G
22	DA	2198	A
22	DA	2204	G
22	DA	2211	A
22	DA	2212	A
22	DA	2213	U
22	DA	2225	A
22	DA	2226	C
22	DA	2238	G
22	DA	2239	G
22	DA	2242	G
22	DA	2243	U
22	DA	2250	G
22	DA	2267	A
22	DA	2268	A
22	DA	2273	A
22	DA	2280	G
22	DA	2283	C
22	DA	2287	A
22	DA	2288	A
22	DA	2297	A
22	DA	2305	U
22	DA	2307	G
22	DA	2309	A
22	DA	2310	C
22	DA	2311	A
22	DA	2312	U
22	DA	2321	U
22	DA	2322	A
22	DA	2325	G

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Mol	Chain	Res	Type
22	DA	2327	A
22	DA	2333	A
22	DA	2345	G
22	DA	2347	C
22	DA	2350	C
22	DA	2354	C
22	DA	2357	G
22	DA	2361	G
22	DA	2383	G
22	DA	2385	C
22	DA	2402	U
22	DA	2403	C
22	DA	2406	A
22	DA	2407	A
22	DA	2417	C
22	DA	2422	C
22	DA	2423	U
22	DA	2424	C
22	DA	2425	A
22	DA	2426	A
22	DA	2428	G
22	DA	2429	G
22	DA	2430	A
22	DA	2431	U
22	DA	2434	A
22	DA	2435	A
22	DA	2436	G
22	DA	2441	U
22	DA	2442	C
22	DA	2447	G
22	DA	2448	A
22	DA	2449	U
22	DA	2476	A
22	DA	2482	A
22	DA	2484	G
22	DA	2491	U
22	DA	2502	G
22	DA	2503	A
22	DA	2504	U
22	DA	2505	G
22	DA	2518	A
22	DA	2525	G

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Mol	Chain	Res	Type
22	DA	2529	G
22	DA	2547	A
22	DA	2554	U
22	DA	2556	C
22	DA	2564	A
22	DA	2566	A
22	DA	2567	G
22	DA	2572	A
22	DA	2573	C
22	DA	2575	C
22	DA	2578	G
22	DA	2582	G
22	DA	2585	U
22	DA	2586	U
22	DA	2596	U
22	DA	2600	A
22	DA	2602	A
22	DA	2603	G
22	DA	2609	U
22	DA	2613	U
22	DA	2614	A
22	DA	2629	U
22	DA	2630	G
22	DA	2642	G
22	DA	2646	C
22	DA	2689	U
22	DA	2690	U
22	DA	2714	G
22	DA	2716	C
22	DA	2726	A
22	DA	2727	A
22	DA	2733	A
22	DA	2748	A
22	DA	2757	A
22	DA	2778	A
22	DA	2791	G
22	DA	2798	U
22	DA	2799	A
22	DA	2812	G
22	DA	2818	U
22	DA	2820	A
22	DA	2826	A

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Mol	Chain	Res	Type
22	DA	2835	A
22	DA	2861	U
22	DA	2867	G
22	DA	2872	A
22	DA	2873	A
22	DA	2879	A
22	DA	2880	C
22	DA	2891	U
22	DA	2903	U
23	DB	13	G
23	DB	15	A
23	DB	16	G
23	DB	24	G
23	DB	35	C
23	DB	36	C
23	DB	41	G
23	DB	44	G
23	DB	51	G
23	DB	56	G
23	DB	58	A
23	DB	66	A
23	DB	73	A
23	DB	88	C
23	DB	89	U
23	DB	90	C
23	DB	99	A
23	DB	109	A

All (77) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
1	AA	115	G
1	AA	209	U
1	AA	351	G
1	AA	353	A
1	AA	429	U
1	AA	481	G
1	AA	484	G
1	AA	733	G
1	AA	1031	C
1	AA	1145	A
1	AA	1201	A

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Mol	Chain	Res	Type
1	AA	1211	U
1	AA	1533	C
22	BA	70	G
22	BA	199	A
22	BA	271	G
22	BA	310	A
22	BA	404	A
22	BA	479	A
22	BA	503	A
22	BA	613	A
22	BA	764	A
22	BA	776	G
22	BA	784	G
22	BA	846	U
22	BA	858	G
22	BA	960	A
22	BA	984	A
22	BA	995	C
22	BA	1185	G
22	BA	1301	A
22	BA	1344	U
22	BA	1378	A
22	BA	1606	C
22	BA	1610	A
22	BA	1738	G
22	BA	1757	A
22	BA	2127	G
22	BA	2282	G
22	BA	2286	G
22	BA	2326	C
22	BA	2353	G
22	BA	2406	A
22	BA	2425	A
22	BA	2756	U
22	BA	2800	A
22	BA	2873	A
23	BB	15	A
1	CA	115	G
1	CA	209	U
1	CA	429	U
1	CA	438	U
1	CA	1049	U

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Mol	Chain	Res	Type
1	CA	1201	A
1	CA	1211	U
22	DA	196	A
22	DA	271	G
22	DA	404	A
22	DA	479	A
22	DA	982	C
22	DA	1240	U
22	DA	1275	A
22	DA	1344	U
22	DA	1378	A
22	DA	1606	C
22	DA	1900	A
22	DA	2109	U
22	DA	2127	G
22	DA	2146	C
22	DA	2162	G
22	DA	2211	A
22	DA	2225	A
22	DA	2286	G
22	DA	2308	G
22	DA	2326	C
22	DA	2425	A
22	DA	2756	U

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 carbohydrates in this entry.

5.6 Ligand geometry [i](#)

Of 502 ligands modelled in this entry, 501 are monoatomic - leaving 1 for Mogul analysis.

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	$\# Z > 2$
56	NEG	CA	1657	54	11,16,16	0.94	1 (9%)	12,20,20	0.71	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
56	NEG	CA	1657	54	-	0/15/18/18	0/0/0/0

All (1) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
56	CA	1657	NEG	C6-N2	2.47	1.37	1.34

There are no bond angle outliers.

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

1 monomer is involved in 1 short contact:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
56	CA	1657	NEG	1	0

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 Fit of model and data ⓘ

6.1 Protein, DNA and RNA chains ⓘ

In the following table, the column labelled ‘#RSRZ> 2’ contains the number (and percentage) of RSRZ outliers, followed by percent RSRZ outliers for the chain as percentile scores relative to all X-ray entries and entries of similar resolution. The OWAB column contains the minimum, median, 95th percentile and maximum values of the occupancy-weighted average B-factor per residue. The column labelled ‘Q< 0.9’ lists the number of (and percentage) of residues with an average occupancy less than 0.9.

Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
1	AA	1538/1539 (99%)	-0.05	57 (3%) 45 22	12, 56, 146, 187	0
1	CA	1539/1539 (100%)	0.41	146 (9%) 10 4	24, 78, 151, 182	0
2	AB	218/218 (100%)	0.29	14 (6%) 23 9	38, 79, 106, 130	0
2	CB	218/218 (100%)	0.53	25 (11%) 6 2	62, 97, 114, 127	0
3	AC	206/206 (100%)	0.40	16 (7%) 16 5	57, 83, 97, 110	0
3	CC	206/206 (100%)	1.45	62 (30%) 1 0	83, 101, 112, 120	0
4	AD	205/205 (100%)	-0.07	3 (1%) 76 58	39, 64, 88, 106	0
4	CD	205/205 (100%)	-0.19	4 (1%) 68 46	21, 41, 70, 92	0
5	AE	150/150 (100%)	-0.06	1 (0%) 89 78	31, 50, 83, 105	0
5	CE	150/150 (100%)	-0.03	0 100 100	28, 58, 89, 109	0
6	AF	100/100 (100%)	0.20	6 (6%) 25 10	36, 60, 80, 92	0
6	CF	100/100 (100%)	0.15	2 (2%) 68 46	48, 81, 103, 108	0
7	AG	151/151 (100%)	0.79	28 (18%) 2 1	68, 93, 106, 116	0
7	CG	151/151 (100%)	2.77	92 (60%) 0 0	95, 117, 127, 130	0
8	AH	129/129 (100%)	-0.16	1 (0%) 87 75	31, 52, 72, 90	0
8	CH	129/129 (100%)	0.12	7 (5%) 29 12	55, 75, 93, 105	0
9	AI	127/127 (100%)	1.51	42 (33%) 0 0	70, 95, 110, 122	0
9	CI	127/127 (100%)	2.10	50 (39%) 0 0	100, 112, 123, 131	0
10	AJ	98/98 (100%)	0.94	20 (20%) 1 0	76, 90, 107, 127	0
10	CJ	98/98 (100%)	3.28	56 (57%) 0 0	97, 114, 122, 133	0
11	AK	117/117 (100%)	0.11	3 (2%) 59 35	26, 70, 98, 117	0
11	CK	117/117 (100%)	0.09	3 (2%) 59 35	36, 75, 94, 98	0
12	AL	123/123 (100%)	-0.10	3 (2%) 62 39	23, 39, 70, 100	0
12	CL	123/123 (100%)	0.22	5 (4%) 41 19	34, 56, 84, 108	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
13	AM	114/114 (100%)	0.97	26 (22%) 1 0	70, 87, 104, 111	0
13	CM	114/114 (100%)	3.40	83 (72%) 0 0	105, 122, 130, 134	0
14	AN	96/100 (96%)	1.09	24 (25%) 1 0	79, 91, 108, 117	0
14	CN	96/100 (96%)	2.64	62 (64%) 0 0	98, 112, 125, 129	0
15	AO	88/88 (100%)	-0.13	0 100 100	30, 51, 72, 92	0
15	CO	88/88 (100%)	0.28	5 (5%) 27 11	45, 71, 89, 106	0
16	AP	82/82 (100%)	0.13	2 (2%) 62 39	34, 51, 92, 108	0
16	CP	82/82 (100%)	0.78	13 (15%) 3 1	52, 73, 105, 117	0
17	AQ	80/80 (100%)	0.29	2 (2%) 61 37	30, 53, 77, 120	0
17	CQ	80/80 (100%)	0.93	14 (17%) 2 1	48, 88, 102, 109	0
18	AR	55/55 (100%)	0.23	3 (5%) 29 12	44, 57, 87, 108	0
18	CR	55/55 (100%)	0.25	4 (7%) 18 6	41, 58, 85, 109	0
19	AS	79/79 (100%)	1.46	22 (27%) 1 0	78, 95, 107, 113	0
19	CS	79/79 (100%)	2.51	45 (56%) 0 0	105, 123, 130, 133	0
20	AT	85/85 (100%)	0.24	3 (3%) 48 23	35, 52, 79, 98	0
20	CT	85/85 (100%)	1.47	22 (25%) 1 0	61, 86, 102, 106	0
21	AU	51/51 (100%)	0.32	4 (7%) 16 5	39, 73, 105, 115	0
21	CU	51/51 (100%)	0.30	3 (5%) 26 11	46, 73, 101, 107	0
22	BA	2897/2903 (99%)	0.11	131 (4%) 37 17	1, 14, 135, 195	0
22	DA	2897/2903 (99%)	0.57	230 (7%) 15 5	38, 91, 152, 183	0
23	BB	119/119 (100%)	-0.37	0 100 100	2, 25, 59, 89	0
23	DB	118/119 (99%)	0.47	5 (4%) 40 19	80, 119, 136, 145	0
24	BC	271/271 (100%)	-0.13	12 (4%) 38 17	2, 21, 52, 65	0
24	DC	271/271 (100%)	0.51	27 (9%) 9 3	45, 70, 90, 96	0
25	BD	209/209 (100%)	-0.33	0 100 100	1, 12, 38, 77	0
25	DD	209/209 (100%)	0.82	36 (17%) 2 1	56, 82, 97, 107	0
26	BE	201/201 (100%)	-0.32	1 (0%) 91 83	1, 27, 64, 92	0
26	DE	201/201 (100%)	1.18	47 (23%) 1 0	63, 98, 114, 122	0
27	BF	177/177 (100%)	-0.10	6 (3%) 49 24	20, 51, 84, 100	0
27	DF	177/177 (100%)	2.16	75 (42%) 0 0	100, 119, 131, 137	0
28	BG	176/176 (100%)	-0.28	2 (1%) 82 66	12, 38, 66, 79	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
28	DG	176/176 (100%)	1.72	63 (35%) 0 0	87, 105, 116, 126	0
29	BH	149/149 (100%)	2.53	65 (43%) 0 0	25, 102, 121, 129	0
29	DH	149/149 (100%)	0.97	22 (14%) 3 1	25, 92, 107, 115	0
30	BI	141/141 (100%)	3.02	73 (51%) 0 0	103, 122, 132, 139	0
30	DI	141/141 (100%)	4.37	126 (89%) 0 0	109, 130, 140, 149	0
31	BJ	142/142 (100%)	-0.31	0 100 100	1, 8, 28, 41	0
31	DJ	142/142 (100%)	0.49	11 (7%) 16 5	57, 79, 92, 103	0
32	BK	122/122 (100%)	-0.35	0 100 100	4, 13, 35, 66	0
32	DK	122/122 (100%)	0.60	21 (17%) 2 1	56, 75, 94, 106	0
33	BL	143/143 (100%)	-0.31	0 100 100	1, 22, 52, 87	0
33	DL	143/143 (100%)	1.31	34 (23%) 1 0	58, 92, 106, 127	0
34	BM	136/136 (100%)	-0.40	0 100 100	1, 11, 34, 82	0
34	DM	136/136 (100%)	0.39	11 (8%) 15 5	50, 79, 94, 113	0
35	BN	120/120 (100%)	-0.38	0 100 100	2, 10, 21, 70	0
35	DN	120/120 (100%)	0.88	19 (15%) 3 1	65, 88, 100, 123	0
36	BO	116/116 (100%)	-0.30	0 100 100	12, 30, 47, 57	0
36	DO	116/116 (100%)	1.83	50 (43%) 0 0	91, 106, 114, 121	0
37	BP	114/114 (100%)	-0.33	0 100 100	5, 18, 46, 67	0
37	DP	114/114 (100%)	0.87	22 (19%) 2 1	65, 81, 94, 103	0
38	BQ	117/117 (100%)	-0.26	0 100 100	1, 4, 15, 52	0
38	DQ	117/117 (100%)	0.90	17 (14%) 3 1	62, 79, 89, 94	0
39	BR	103/103 (100%)	-0.41	0 100 100	1, 13, 35, 67	0
39	DR	103/103 (100%)	1.35	26 (25%) 1 0	67, 91, 101, 103	0
40	BS	110/110 (100%)	-0.38	0 100 100	1, 5, 27, 79	0
40	DS	110/110 (100%)	1.32	28 (25%) 1 0	68, 88, 105, 112	0
41	BT	93/93 (100%)	-0.02	3 (3%) 51 27	8, 27, 77, 106	0
41	DT	93/93 (100%)	1.71	31 (33%) 0 0	80, 98, 117, 125	0
42	BU	102/102 (100%)	0.12	9 (8%) 12 4	13, 32, 67, 96	0
42	DU	102/102 (100%)	2.36	44 (43%) 0 0	88, 106, 122, 129	0
43	BV	94/94 (100%)	-0.34	0 100 100	7, 23, 51, 63	0
43	DV	94/94 (100%)	0.53	8 (8%) 13 4	74, 95, 107, 115	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
44	BW	76/76 (100%)	-0.25	1 (1%) 79 62	3, 13, 32, 61	0
44	DW	75/76 (98%)	1.64	21 (28%) 1 0	60, 90, 99, 114	0
45	BX	77/77 (100%)	-0.32	0 100 100	5, 26, 54, 77	0
45	DX	77/77 (100%)	0.78	9 (11%) 6 2	57, 80, 94, 101	0
46	BY	63/63 (100%)	0.06	2 (3%) 51 27	17, 40, 74, 92	0
46	DY	63/63 (100%)	1.85	29 (46%) 0 0	89, 105, 112, 115	0
47	BZ	58/58 (100%)	-0.27	0 100 100	2, 8, 35, 48	0
47	DZ	58/58 (100%)	0.67	6 (10%) 9 3	64, 81, 96, 102	0
48	B0	56/56 (100%)	-0.45	0 100 100	1, 11, 44, 65	0
48	D0	56/56 (100%)	1.18	16 (28%) 1 0	58, 89, 102, 109	0
49	B1	50/50 (100%)	0.36	3 (6%) 25 10	25, 38, 63, 84	0
49	D1	50/50 (100%)	2.37	25 (50%) 0 0	79, 99, 108, 112	0
50	B2	46/46 (100%)	-0.19	1 (2%) 65 42	3, 8, 17, 80	0
50	D2	46/46 (100%)	0.98	7 (15%) 3 1	58, 80, 89, 105	0
51	B3	64/64 (100%)	-0.27	0 100 100	5, 11, 20, 33	0
51	D3	64/64 (100%)	0.97	12 (18%) 2 1	66, 82, 93, 98	0
52	B4	38/38 (100%)	-0.15	0 100 100	4, 13, 33, 51	0
52	D4	38/38 (100%)	1.54	10 (26%) 1 0	65, 84, 99, 109	0
53	B5	191/207 (92%)	5.07	168 (87%) 0 0	108, 127, 138, 143	0
All	All	20734/20773 (99%)	0.55	2548 (12%) 5 2	1, 73, 130, 195	0

All (2548) RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
22	BA	2184	A	20.2
30	BI	3	LYS	17.2
22	BA	2101	A	16.7
30	BI	2	ALA	15.7
30	BI	4	LYS	15.3
53	B5	147	GLY	14.9
53	B5	111	PHE	14.7
22	BA	2100	G	14.5
10	AJ	102	LEU	13.5
30	BI	12	GLN	13.2
22	BA	2144	G	13.1

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Mol	Chain	Res	Type	RSRZ
30	DI	69	PHE	13.0
22	BA	2117	A	12.9
53	B5	218	THR	12.8
22	BA	2135	A	12.8
22	BA	2174	C	12.7
22	BA	2185	U	12.6
29	BH	96	THR	12.4
53	B5	122	GLY	12.3
9	CI	129	LYS	12.3
9	CI	128	SER	12.3
22	BA	2104	C	12.2
30	DI	77	ALA	12.0
30	BI	53	LEU	11.9
22	BA	2183	A	11.8
10	CJ	75	ASP	11.8
1	CA	1539	C	11.7
22	BA	2136	G	11.7
13	CM	45	ILE	11.6
53	B5	66	PRO	11.6
22	BA	2158	A	11.4
29	BH	97	ARG	11.4
1	CA	1536	C	11.2
53	B5	107	GLY	11.2
33	DL	144	GLU	11.2
22	BA	2182	U	11.1
30	DI	68	THR	11.1
53	B5	85	LYS	11.1
22	BA	2143	C	11.0
9	CI	125	PRO	11.0
42	DU	20	GLY	10.9
1	AA	1030	U	10.9
22	BA	2107	G	10.8
27	DF	23	ASN	10.7
53	B5	157	ILE	10.5
53	B5	84	ILE	10.4
24	DC	241	GLY	10.4
30	BI	5	VAL	10.3
22	BA	2169	A	10.3
53	B5	149	ASN	10.3
30	DI	2	ALA	10.3
22	BA	2103	C	10.2
1	CA	1537	U	10.1

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Mol	Chain	Res	Type	RSRZ
53	B5	95	VAL	10.0
53	B5	110	ASP	10.0
22	BA	2102	G	10.0
29	BH	86	ASP	9.8
53	B5	225	ILE	9.8
22	BA	2159	G	9.7
30	BI	22	PRO	9.7
22	BA	2148	G	9.7
29	BH	130	VAL	9.7
53	B5	106	ASP	9.6
29	BH	146	VAL	9.6
29	BH	148	ALA	9.6
30	DI	10	LYS	9.6
53	B5	150	ILE	9.5
20	CT	4	ILE	9.5
29	BH	120	GLY	9.4
22	BA	2106	U	9.4
7	CG	133	THR	9.4
1	CA	1535	C	9.3
53	B5	77	ALA	9.3
49	D1	52	ALA	9.3
30	DI	47	ASP	9.3
3	CC	155	GLY	9.2
42	DU	60	GLU	9.2
9	CI	124	ARG	9.1
1	CA	1538	C	9.1
30	DI	3	LYS	9.1
38	DQ	29	SER	9.0
10	CJ	77	VAL	9.0
22	BA	2114	A	9.0
22	DA	1537	G	9.0
22	BA	2145	C	9.0
3	CC	192	THR	8.9
53	B5	212	SER	8.9
9	CI	127	PHE	8.9
22	BA	2139	U	8.9
53	B5	55	SER	8.9
19	CS	74	PHE	8.9
10	CJ	76	ILE	8.9
19	CS	66	MET	8.9
30	BI	23	PRO	8.8
30	DI	57	VAL	8.8

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Mol	Chain	Res	Type	RSRZ
22	BA	2115	G	8.8
27	DF	128	TYR	8.8
30	DI	58	VAL	8.7
53	B5	173	HIS	8.7
1	CA	1302	C	8.7
53	B5	43	GLU	8.7
10	CJ	8	ILE	8.7
22	BA	2175	C	8.7
53	B5	220	GLY	8.6
29	BH	102	ALA	8.6
7	CG	2	PRO	8.6
53	B5	140	ASN	8.6
29	BH	113	SER	8.5
22	BA	2177	C	8.5
9	CI	126	GLN	8.5
13	CM	85	CYS	8.5
10	CJ	74	VAL	8.4
2	AB	156	GLY	8.4
42	DU	79	LYS	8.4
10	CJ	60	ASP	8.4
10	CJ	41	PRO	8.4
53	B5	182	PRO	8.3
22	BA	2178	C	8.3
24	DC	239	ASN	8.3
53	B5	203	GLU	8.3
7	CG	49	THR	8.3
22	BA	2140	G	8.3
9	CI	117	GLY	8.2
22	BA	2113	U	8.2
30	DI	53	LEU	8.2
1	CA	1032	G	8.2
53	B5	109	MET	8.2
10	CJ	72	ARG	8.2
53	B5	184	GLU	8.2
10	CJ	39	PRO	8.1
22	DA	1067	A	8.1
30	DI	12	GLN	8.1
10	CJ	7	ARG	8.0
22	BA	2123	G	8.0
29	BH	144	VAL	8.0
30	BI	67	PHE	7.9
30	DI	67	PHE	7.9

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Mol	Chain	Res	Type	RSRZ
42	DU	36	VAL	7.8
9	AI	123	ARG	7.8
53	B5	39	ASP	7.8
22	BA	2130	U	7.8
53	B5	121	MET	7.8
22	BA	2156	G	7.8
33	DL	78	ARG	7.8
22	DA	1536	C	7.8
30	DI	96	ASP	7.8
22	BA	2157	G	7.8
30	BI	87	LYS	7.7
1	AA	1535	C	7.7
22	BA	2176	A	7.7
30	BI	14	ALA	7.7
53	B5	52	PRO	7.7
22	BA	2121	G	7.7
53	B5	86	GLU	7.6
14	CN	100	SER	7.6
13	CM	95	LEU	7.6
13	CM	10	PRO	7.6
42	DU	50	PRO	7.6
30	BI	54	PRO	7.5
22	BA	2127	G	7.5
53	B5	183	PRO	7.5
22	BA	2181	U	7.5
30	DI	95	LYS	7.5
53	B5	195	ARG	7.5
9	AI	128	SER	7.5
33	DL	92	LEU	7.5
22	BA	2112	G	7.4
30	DI	7	ALA	7.4
30	DI	130	GLU	7.4
22	BA	2138	G	7.4
7	CG	16	PRO	7.4
22	BA	2099	U	7.4
22	BA	2155	U	7.4
29	BH	85	GLY	7.4
44	DW	54	GLY	7.4
30	DI	126	THR	7.3
53	B5	156	GLU	7.3
29	BH	91	PHE	7.3
53	B5	217	THR	7.3

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Mol	Chain	Res	Type	RSRZ
22	BA	2150	C	7.3
53	B5	81	GLY	7.3
1	AA	1538	C	7.3
30	BI	81	LYS	7.3
20	CT	3	ASN	7.3
53	B5	76	LEU	7.3
29	BH	101	ASP	7.3
22	BA	2105	U	7.3
22	BA	2147	A	7.3
10	CJ	99	GLN	7.2
22	BA	2124	G	7.2
22	BA	2165	C	7.2
22	DA	1085	A	7.2
27	DF	176	PRO	7.2
30	DI	6	GLN	7.2
53	B5	94	TYR	7.2
22	BA	2131	U	7.2
22	BA	2172	U	7.1
12	CL	25	GLU	7.1
2	AB	157	LEU	7.1
53	B5	72	GLN	7.1
1	CA	1031	C	7.1
30	BI	11	LEU	7.1
7	CG	39	ALA	7.1
30	DI	85	GLY	7.1
14	CN	20	TYR	7.0
13	CM	11	ASP	7.0
53	B5	20	VAL	7.0
22	DA	345	A	7.0
53	B5	152	GLU	7.0
13	CM	111	GLY	7.0
18	AR	20	GLU	7.0
7	CG	62	PHE	7.0
2	AB	155	GLY	7.0
53	B5	97	GLY	7.0
1	CA	1030	U	7.0
22	BA	2166	U	7.0
53	B5	105	LEU	7.0
22	DA	2126	A	7.0
10	CJ	10	LEU	7.0
22	BA	2125	G	7.0
9	AI	130	ARG	6.9

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Mol	Chain	Res	Type	RSRZ
53	B5	79	ALA	6.9
30	DI	4	LYS	6.9
19	CS	29	LYS	6.9
24	DC	233	GLY	6.9
12	CL	124	ALA	6.9
22	BA	2162	G	6.9
53	B5	38	PHE	6.9
30	DI	20	PRO	6.9
40	DS	84	ARG	6.8
27	DF	155	THR	6.8
29	BH	58	LEU	6.8
22	BA	2179	C	6.8
10	CJ	87	LEU	6.8
30	DI	59	ILE	6.8
7	CG	134	ALA	6.8
30	DI	70	VAL	6.8
27	DF	95	ARG	6.8
36	DO	117	PHE	6.8
53	B5	181	PHE	6.7
9	CI	43	THR	6.7
29	BH	136	SER	6.7
30	BI	13	VAL	6.7
13	CM	94	GLY	6.7
22	BA	2116	G	6.7
22	DA	1073	A	6.7
7	CG	108	ALA	6.7
22	BA	2126	A	6.7
29	BH	69	ALA	6.6
1	CA	209	U	6.6
13	CM	46	SER	6.6
30	DI	15	ALA	6.6
46	DY	33	ALA	6.6
22	DA	1535	A	6.6
13	CM	24	GLY	6.6
19	CS	30	PRO	6.6
22	BA	2120	G	6.5
9	CI	108	ALA	6.5
22	BA	2118	U	6.5
27	DF	154	ILE	6.5
1	AA	1536	C	6.5
30	DI	34	ASN	6.5
13	CM	112	PRO	6.5

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Mol	Chain	Res	Type	RSRZ
2	CB	88	ASP	6.5
30	DI	76	ALA	6.5
53	B5	92	ALA	6.5
24	DC	232	HIS	6.4
53	B5	148	PHE	6.4
9	CI	123	ARG	6.4
22	BA	885	C	6.4
53	B5	187	ALA	6.4
10	CJ	101	SER	6.4
53	B5	213	VAL	6.4
19	AS	56	GLN	6.4
14	CN	69	ARG	6.4
41	DT	60	THR	6.4
42	DU	51	ALA	6.4
42	DU	77	THR	6.4
24	DC	237	GLY	6.3
7	CG	111	ARG	6.3
13	CM	108	THR	6.3
53	B5	133	GLY	6.3
1	CA	1209	C	6.3
53	B5	204	GLY	6.3
1	CA	1540	U	6.3
29	BH	115	VAL	6.3
7	CG	52	GLN	6.3
22	DA	1175	A	6.3
27	DF	86	GLY	6.3
46	DY	32	ALA	6.2
52	D4	10	LEU	6.2
1	CA	1020	G	6.2
53	B5	28	ARG	6.2
49	D1	53	LYS	6.2
49	D1	36	LEU	6.2
3	CC	127	ARG	6.2
30	DI	48	SER	6.2
22	BA	1065	U	6.2
1	AA	1534	A	6.2
7	CG	137	LYS	6.2
22	DA	331	C	6.2
22	DA	1068	G	6.2
29	BH	98	ASP	6.2
28	DG	80	THR	6.2
22	DA	1066	U	6.2

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Mol	Chain	Res	Type	RSRZ
30	DI	118	THR	6.1
22	DA	878	A	6.1
42	DU	43	LYS	6.1
13	CM	12	HIS	6.1
30	DI	99	GLY	6.1
53	B5	40	GLU	6.1
27	DF	157	THR	6.1
7	CG	82	GLY	6.1
29	BH	68	ARG	6.1
9	CI	118	LEU	6.1
22	BA	2163	A	6.1
33	DL	114	GLY	6.1
28	DG	62	TRP	6.1
53	B5	179	ALA	6.1
27	DF	24	SER	6.1
10	CJ	59	LYS	6.1
22	DA	1094	U	6.1
9	AI	127	PHE	6.1
22	DA	2174	C	6.0
14	CN	16	LEU	6.0
14	CN	58	SER	6.0
1	CA	1018	G	6.0
29	BH	119	ASN	6.0
53	B5	87	ALA	6.0
13	CM	33	ILE	6.0
53	B5	132	LEU	6.0
29	BH	54	LEU	6.0
53	B5	67	HIS	6.0
53	B5	104	ILE	6.0
39	DR	96	VAL	6.0
29	BH	72	ILE	6.0
22	BA	2180	U	5.9
10	CJ	102	LEU	5.9
22	BA	2122	U	5.9
9	CI	67	VAL	5.9
1	CA	983	A	5.9
1	CA	1534	A	5.9
42	DU	33	LYS	5.9
30	DI	129	ILE	5.9
14	CN	10	GLU	5.9
7	CG	53	ARG	5.9
14	CN	36	ALA	5.9

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Mol	Chain	Res	Type	RSRZ
10	CJ	89	ARG	5.9
36	DO	88	LYS	5.9
30	BI	68	THR	5.8
10	CJ	61	ALA	5.8
28	DG	33	LEU	5.8
22	BA	2171	A	5.8
7	CG	37	SER	5.8
53	B5	151	GLY	5.8
22	DA	1172	C	5.8
1	CA	1021	A	5.8
7	CG	48	GLU	5.8
22	BA	2153	C	5.8
9	CI	66	THR	5.8
1	AA	1539	C	5.8
14	AN	43	ASN	5.8
53	B5	90	ALA	5.8
14	CN	101	TRP	5.8
53	B5	89	GLU	5.8
22	BA	2161	C	5.8
39	DR	27	ILE	5.8
53	B5	70	GLY	5.8
13	CM	31	LYS	5.8
28	DG	52	PHE	5.7
1	AA	1003	G	5.7
30	DI	46	THR	5.7
29	BH	110	VAL	5.7
22	BA	2154	A	5.7
29	BH	124	THR	5.7
30	DI	89	GLY	5.7
53	B5	93	ASP	5.7
13	CM	30	SER	5.7
30	BI	66	SER	5.7
14	CN	47	LYS	5.7
53	B5	194	ILE	5.7
10	CJ	71	LEU	5.7
2	AB	135	LEU	5.7
13	CM	32	ALA	5.7
22	BA	2134	A	5.7
22	BA	2168	G	5.7
53	B5	185	LYS	5.7
44	DW	52	GLY	5.7
24	DC	242	LYS	5.7

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Mol	Chain	Res	Type	RSRZ
42	DU	40	ASN	5.7
27	DF	25	VAL	5.7
53	B5	62	THR	5.7
53	B5	141	PRO	5.7
22	BA	2137	U	5.7
29	BH	95	GLY	5.7
9	CI	122	ARG	5.6
1	CA	1024	G	5.6
53	B5	73	VAL	5.6
13	CM	96	PRO	5.6
40	DS	3	THR	5.6
1	CA	1017	U	5.6
53	B5	197	LEU	5.6
42	DU	39	ILE	5.6
30	DI	60	THR	5.6
49	D1	24	THR	5.6
44	DW	25	ARG	5.6
22	DA	1171	G	5.6
42	DU	31	SER	5.6
22	BA	2108	A	5.6
22	DA	2124	G	5.6
1	AA	1031	C	5.6
5	AE	159	LYS	5.6
13	CM	63	PHE	5.6
30	BI	21	SER	5.6
48	D0	27	SER	5.6
49	D1	34	LEU	5.6
53	B5	108	TRP	5.6
7	CG	17	LYS	5.5
33	DL	81	ASP	5.5
46	DY	40	SER	5.5
53	B5	155	ARG	5.5
53	B5	96	GLY	5.5
19	CS	15	LEU	5.5
44	DW	38	VAL	5.5
1	CA	999	C	5.5
2	CB	9	MET	5.5
30	BI	41	ALA	5.5
1	CA	1296	C	5.5
9	CI	112	GLU	5.5
30	DI	66	SER	5.5
7	CG	59	LEU	5.5

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Mol	Chain	Res	Type	RSRZ
49	D1	37	LYS	5.5
53	B5	219	MET	5.5
13	CM	77	ILE	5.5
9	CI	58	VAL	5.5
53	B5	83	LYS	5.5
53	B5	200	HIS	5.5
1	AA	78	A	5.5
28	DG	32	GLU	5.5
30	BI	115	ALA	5.5
1	CA	1028	C	5.5
42	DU	78	GLY	5.5
1	CA	1025	U	5.5
1	CA	82	G	5.5
53	B5	46	ALA	5.4
22	DA	1103	A	5.4
22	DA	1065	U	5.4
24	DC	235	GLY	5.4
29	BH	149	GLU	5.4
10	CJ	6	ILE	5.4
3	CC	193	TYR	5.4
7	AG	69	VAL	5.4
1	CA	1022	A	5.4
22	BA	138	U	5.4
42	BU	53	ASN	5.4
7	CG	41	SER	5.4
30	BI	114	ALA	5.4
22	DA	2402	U	5.4
30	DI	11	LEU	5.4
53	B5	131	ILE	5.4
13	CM	109	ARG	5.4
29	BH	83	LYS	5.4
22	DA	1084	A	5.4
29	BH	123	ARG	5.4
14	CN	33	ASP	5.4
30	DI	78	VAL	5.4
22	DA	1087	G	5.4
30	DI	14	ALA	5.3
53	B5	222	SER	5.3
13	CM	39	ILE	5.3
14	AN	21	PHE	5.3
1	AA	87	C	5.3
46	DY	13	GLU	5.3

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Mol	Chain	Res	Type	RSRZ
29	BH	55	GLU	5.3
34	DM	136	MET	5.3
49	D1	23	THR	5.3
30	DI	45	LYS	5.3
19	AS	49	ILE	5.3
22	DA	1107	G	5.3
22	DA	1870	C	5.3
30	DI	139	VAL	5.3
9	AI	20	PHE	5.2
22	DA	1093	G	5.2
13	CM	80	LEU	5.2
22	DA	1083	U	5.2
19	CS	60	VAL	5.2
53	B5	82	GLU	5.2
53	B5	98	GLU	5.2
22	BA	2146	C	5.2
28	DG	2	SER	5.2
1	CA	1027	C	5.2
19	CS	49	ILE	5.2
7	CG	5	ARG	5.2
22	BA	884	U	5.2
10	CJ	26	VAL	5.2
27	DF	35	THR	5.2
46	DY	59	GLU	5.2
7	CG	148	ASN	5.2
14	CN	99	ALA	5.2
22	BA	2142	A	5.2
30	DI	120	ALA	5.2
53	B5	223	VAL	5.2
10	CJ	58	ASN	5.2
48	D0	57	LYS	5.2
22	BA	2111	U	5.2
13	CM	60	VAL	5.2
53	B5	88	GLU	5.2
53	B5	65	LEU	5.2
36	DO	65	THR	5.2
19	AS	74	PHE	5.2
53	B5	161	ARG	5.2
14	CN	27	LEU	5.2
42	DU	32	GLY	5.2
42	DU	80	ALA	5.2
7	CG	130	ASN	5.1

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Mol	Chain	Res	Type	RSRZ
14	CN	31	ILE	5.1
36	DO	51	ALA	5.1
42	DU	38	GLY	5.1
30	BI	135	SER	5.1
30	BI	69	PHE	5.1
53	B5	78	ILE	5.1
10	CJ	97	ASP	5.1
22	DA	2172	U	5.1
24	BC	234	GLY	5.1
30	DI	131	GLY	5.1
7	CG	103	TRP	5.1
24	DC	49	ILE	5.1
7	CG	15	ASP	5.1
53	B5	221	PRO	5.1
9	AI	43	THR	5.1
14	CN	44	ALA	5.1
36	DO	64	TYR	5.1
13	CM	55	THR	5.1
53	B5	145	THR	5.1
14	CN	30	ILE	5.1
7	CG	4	ARG	5.1
30	DI	31	GLN	5.1
22	BA	2132	U	5.0
42	DU	52	LEU	5.0
1	CA	942	G	5.0
7	CG	18	PHE	5.0
9	CI	38	TYR	5.0
27	DF	67	ILE	5.0
22	DA	1104	C	5.0
31	DJ	142	ILE	5.0
43	DV	94	ALA	5.0
22	DA	2585	U	5.0
19	AS	9	PRO	5.0
10	CJ	81	GLU	5.0
26	DE	103	GLY	5.0
27	DF	62	GLY	5.0
22	BA	2167	U	5.0
29	DH	142	VAL	5.0
30	DI	13	VAL	5.0
46	DY	45	GLN	5.0
24	DC	240	PHE	5.0
22	BA	2189	U	5.0

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Mol	Chain	Res	Type	RSRZ
22	DA	1086	A	5.0
22	BA	2160	C	5.0
53	B5	180	SER	5.0
27	DF	129	SER	5.0
10	CJ	62	ARG	5.0
7	CG	85	TYR	5.0
30	DI	112	THR	5.0
24	BC	236	GLU	4.9
1	CA	984	C	4.9
7	CG	87	VAL	4.9
27	DF	106	ILE	4.9
22	DA	2123	G	4.9
30	DI	133	ALA	4.9
9	AI	120	LYS	4.9
10	CJ	27	GLU	4.9
40	DS	85	ILE	4.9
30	BI	79	LEU	4.9
30	BI	8	TYR	4.9
30	DI	8	TYR	4.9
30	DI	62	TYR	4.9
1	CA	1235	U	4.9
10	CJ	90	LEU	4.9
29	BH	67	ALA	4.9
53	B5	60	ARG	4.9
14	CN	49	GLN	4.9
19	CS	51	VAL	4.9
28	DG	10	VAL	4.9
13	CM	29	ARG	4.9
14	CN	68	GLY	4.9
22	DA	1077	A	4.9
22	DA	2173	A	4.9
42	DU	89	ASP	4.9
7	CG	116	MET	4.9
46	DY	10	SER	4.9
13	CM	83	LEU	4.9
27	DF	132	VAL	4.9
53	B5	42	VAL	4.9
28	DG	157	TYR	4.9
3	CC	109	PRO	4.9
3	CC	172	ARG	4.9
24	BC	242	LYS	4.9
30	BI	100	LYS	4.9

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Mol	Chain	Res	Type	RSRZ
30	DI	97	LYS	4.9
53	B5	158	LYS	4.9
30	DI	17	MET	4.9
27	DF	87	CYS	4.9
9	CI	120	LYS	4.9
53	B5	130	ARG	4.9
24	BC	240	PHE	4.8
22	DA	1075	C	4.8
30	BI	20	PRO	4.8
10	AJ	34	ALA	4.8
33	DL	101	ILE	4.8
7	AG	5	ARG	4.8
30	BI	96	ASP	4.8
19	CS	23	VAL	4.8
42	BU	49	VAL	4.8
53	B5	160	GLY	4.8
14	CN	17	ALA	4.8
9	AI	119	ARG	4.8
30	DI	80	LEU	4.8
30	DI	9	VAL	4.8
30	DI	56	PRO	4.8
22	DA	138	U	4.8
30	DI	54	PRO	4.8
36	DO	66	GLY	4.8
40	DS	68	ASP	4.8
7	CG	20	SER	4.8
19	CS	22	ALA	4.8
27	DF	26	MET	4.8
1	CA	1006	G	4.8
24	DC	248	TRP	4.8
22	BA	2149	U	4.8
7	CG	107	ALA	4.8
9	CI	39	PHE	4.8
13	CM	58	ASP	4.8
28	DG	45	HIS	4.8
14	AN	31	ILE	4.7
29	DH	79	THR	4.7
39	DR	35	PHE	4.7
30	DI	121	ASP	4.7
30	DI	22	PRO	4.7
30	DI	5	VAL	4.7
53	B5	64	SER	4.7

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Mol	Chain	Res	Type	RSRZ
22	BA	2133	G	4.7
19	CS	24	GLU	4.7
42	DU	49	VAL	4.7
46	DY	56	LEU	4.7
19	CS	59	PRO	4.7
42	DU	58	ILE	4.7
49	D1	47	VAL	4.7
2	CB	83	ALA	4.7
30	BI	17	MET	4.7
24	DC	245	VAL	4.7
27	DF	31	VAL	4.7
30	DI	32	GLY	4.7
17	AQ	20	SER	4.7
44	DW	32	LEU	4.7
44	DW	26	PHE	4.7
22	DA	101	A	4.7
22	DA	2797	U	4.7
29	BH	87	GLU	4.7
29	BH	142	VAL	4.7
28	DG	105	LEU	4.6
30	BI	133	ALA	4.6
53	B5	207	GLY	4.6
10	CJ	22	THR	4.6
1	CA	1242	G	4.6
7	CG	132	GLY	4.6
27	DF	122	PHE	4.6
46	DY	28	LEU	4.6
10	CJ	40	ILE	4.6
39	DR	32	THR	4.6
41	DT	34	VAL	4.6
41	DT	58	VAL	4.6
42	BU	52	LEU	4.6
4	CD	36	GLN	4.6
13	AM	99	GLY	4.6
13	CM	75	MET	4.6
3	CC	156	ARG	4.6
37	DP	9	GLU	4.6
27	DF	164	GLU	4.6
28	DG	6	LYS	4.6
53	B5	146	VAL	4.6
41	DT	1	MET	4.6
22	DA	2168	G	4.6

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Mol	Chain	Res	Type	RSRZ
36	DO	62	LEU	4.5
53	B5	172	ILE	4.5
30	BI	38	PHE	4.5
1	CA	1236	A	4.5
14	CN	29	ALA	4.5
53	B5	123	ALA	4.5
53	B5	216	THR	4.5
22	DA	613	A	4.5
17	AQ	83	VAL	4.5
30	BI	59	ILE	4.5
30	BI	99	GLY	4.5
49	D1	35	GLU	4.5
1	AA	1002	G	4.5
11	CK	55	SER	4.5
27	DF	85	ILE	4.5
37	DP	111	LYS	4.5
10	CJ	80	THR	4.5
30	DI	98	VAL	4.5
44	DW	53	CYS	4.5
19	AS	40	ILE	4.5
53	B5	80	LYS	4.5
22	DA	2120	G	4.5
13	AM	115	PRO	4.5
7	CG	43	VAL	4.5
9	AI	41	ARG	4.5
10	AJ	35	GLN	4.5
19	CS	43	ASN	4.5
3	CC	53	SER	4.5
13	CM	23	TYR	4.5
29	BH	106	ALA	4.5
41	DT	35	ALA	4.5
53	B5	196	ALA	4.5
7	AG	62	PHE	4.5
1	CA	1243	C	4.4
10	AJ	75	ASP	4.4
10	CJ	91	ASP	4.4
14	CN	2	ALA	4.4
36	DO	85	LYS	4.4
12	AL	25	GLU	4.4
53	B5	102	GLN	4.4
53	B5	134	PRO	4.4
28	DG	9	VAL	4.4

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Mol	Chain	Res	Type	RSRZ
53	B5	202	PRO	4.4
1	CA	985	C	4.4
22	DA	318	C	4.4
27	DF	130	MET	4.4
13	CM	44	LYS	4.4
30	DI	25	GLY	4.4
14	CN	4	GLN	4.4
22	DA	1105	U	4.4
7	CG	66	LEU	4.4
22	DA	866	A	4.4
7	CG	119	ARG	4.4
53	B5	210	LEU	4.4
3	CC	197	GLY	4.4
13	CM	105	ASN	4.4
13	AM	10	PRO	4.4
20	CT	39	ILE	4.4
10	CJ	16	ARG	4.4
45	DX	20	HIS	4.4
53	B5	54	ARG	4.4
22	BA	2110	G	4.4
53	B5	136	GLY	4.4
27	DF	32	GLU	4.4
28	DG	174	ALA	4.4
42	DU	41	LEU	4.4
14	AN	20	TYR	4.4
1	CA	1313	U	4.4
37	DP	110	ILE	4.4
13	CM	64	VAL	4.4
9	AI	126	GLN	4.4
22	DA	1078	U	4.4
30	DI	90	SER	4.4
7	CG	118	LEU	4.4
13	CM	48	LEU	4.4
19	CS	63	THR	4.3
30	BI	78	VAL	4.3
9	CI	113	ARG	4.3
29	BH	63	ALA	4.3
53	B5	174	ALA	4.3
22	BA	1926	U	4.3
48	D0	34	SER	4.3
53	B5	199	ALA	4.3
22	DA	1100	C	4.3

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Mol	Chain	Res	Type	RSRZ
22	DA	12	U	4.3
53	B5	188	ASP	4.3
1	CA	1026	G	4.3
30	DI	21	SER	4.3
3	CC	79	LYS	4.3
30	BI	92	LYS	4.3
10	AJ	74	VAL	4.3
53	B5	61	GLY	4.3
7	AG	88	PRO	4.3
46	DY	37	LEU	4.3
8	CH	2	SER	4.3
19	CS	48	THR	4.3
2	CB	67	ILE	4.3
12	AL	124	ALA	4.3
2	CB	32	PHE	4.3
29	DH	149	GLU	4.3
30	BI	80	LEU	4.3
7	CG	75	VAL	4.3
53	B5	91	GLY	4.3
20	CT	72	ALA	4.3
19	AS	39	THR	4.3
22	DA	1174	U	4.3
27	DF	65	PRO	4.3
36	DO	102	ARG	4.3
28	DG	59	ALA	4.3
28	DG	104	ASN	4.3
36	DO	40	ILE	4.3
9	AI	122	ARG	4.3
40	DS	92	ARG	4.3
30	BI	138	LEU	4.3
3	CC	129	MET	4.2
30	BI	101	ILE	4.2
22	DA	2110	G	4.2
30	DI	88	SER	4.2
30	DI	119	GLY	4.2
27	DF	34	ILE	4.2
39	DR	20	VAL	4.2
19	AS	33	THR	4.2
30	DI	52	GLY	4.2
52	D4	38	GLY	4.2
53	B5	68	GLY	4.2
29	BH	59	ALA	4.2

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Mol	Chain	Res	Type	RSRZ
1	CA	988	G	4.2
22	DA	2125	G	4.2
24	DC	246	THR	4.2
1	AA	990	C	4.2
14	CN	24	ARG	4.2
27	DF	94	GLU	4.2
3	CC	144	LEU	4.2
30	DI	44	ALA	4.2
15	CO	17	ARG	4.2
53	B5	37	LYS	4.2
30	DI	18	ALA	4.2
1	AA	86	G	4.2
36	DO	104	GLN	4.2
14	CN	21	PHE	4.2
1	CA	1019	A	4.2
14	AN	19	LYS	4.2
1	CA	1029	U	4.2
22	DA	1217	U	4.2
24	DC	244	PRO	4.2
41	DT	71	GLY	4.2
53	B5	24	ASP	4.2
3	CC	195	VAL	4.2
13	CM	36	ALA	4.2
41	DT	2	ILE	4.2
14	AN	32	SER	4.2
30	DI	128	SER	4.2
27	DF	37	ASN	4.1
40	DS	40	ASN	4.1
10	CJ	98	VAL	4.1
36	DO	105	ALA	4.1
42	DU	62	GLU	4.1
1	CA	211	G	4.1
25	DD	75	ALA	4.1
7	CG	63	GLU	4.1
22	DA	1106	G	4.1
25	DD	6	GLY	4.1
22	BA	2164	C	4.1
34	DM	60	GLN	4.1
10	CJ	63	ASP	4.1
24	BC	243	HIS	4.1
29	BH	109	GLU	4.1
10	CJ	19	ASP	4.1

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Mol	Chain	Res	Type	RSRZ
28	DG	111	HIS	4.1
30	BI	34	ASN	4.1
53	B5	99	GLU	4.1
14	CN	72	GLY	4.1
25	DD	166	GLY	4.1
7	CG	45	SER	4.1
53	B5	142	LYS	4.1
53	B5	224	ARG	4.1
13	AM	43	VAL	4.1
22	DA	280	U	4.1
36	DO	60	GLU	4.1
1	CA	204	G	4.1
46	DY	36	GLN	4.1
29	BH	80	ILE	4.1
1	CA	1007	U	4.1
7	AG	79	ARG	4.1
50	D2	33	ARG	4.1
14	CN	34	VAL	4.1
22	DA	1538	G	4.1
28	DG	56	ASP	4.1
41	BT	1	MET	4.1
51	D3	64	TYR	4.1
13	CM	76	SER	4.1
1	CA	1050	G	4.1
7	CG	55	GLY	4.1
29	DH	123	ARG	4.1
22	DA	2630	G	4.1
25	DD	199	SER	4.0
13	CM	84	GLY	4.0
3	CC	76	VAL	4.0
10	AJ	89	ARG	4.0
24	BC	235	GLY	4.0
26	DE	172	ALA	4.0
27	DF	153	ASP	4.0
19	CS	20	GLU	4.0
10	CJ	100	ILE	4.0
18	CR	74	HIS	4.0
30	DI	140	VAL	4.0
49	D1	45	GLN	4.0
13	CM	99	GLY	4.0
22	DA	1057	A	4.0
53	B5	166	ASN	4.0

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Mol	Chain	Res	Type	RSRZ
22	DA	2127	G	4.0
53	B5	192	ALA	4.0
29	BH	112	LYS	4.0
1	AA	994	A	4.0
3	CC	46	GLU	4.0
7	CG	80	VAL	4.0
22	DA	846	U	4.0
28	DG	102	VAL	4.0
35	DN	119	SER	4.0
22	DA	2627	G	4.0
30	BI	9	VAL	4.0
1	CA	1314	C	4.0
22	BA	2173	A	4.0
22	DA	2150	C	4.0
14	CN	19	LYS	4.0
53	B5	27	ALA	4.0
35	DN	120	GLU	4.0
1	AA	842	U	4.0
22	BA	1094	U	4.0
1	CA	959	A	4.0
42	DU	71	ALA	4.0
13	CM	68	ASP	4.0
53	B5	23	ILE	4.0
1	AA	1020	G	4.0
1	CA	4	U	4.0
7	CG	88	PRO	4.0
34	DM	80	VAL	4.0
2	AB	9	MET	4.0
48	D0	46	ASP	4.0
35	DN	63	ARG	3.9
22	DA	2891	U	3.9
22	DA	880	G	3.9
22	BA	2402	U	3.9
37	DP	104	THR	3.9
7	CG	145	ALA	3.9
30	BI	142	ASP	3.9
13	CM	71	ARG	3.9
4	AD	36	GLN	3.9
7	CG	91	VAL	3.9
21	AU	4	ILE	3.9
30	DI	35	ILE	3.9
27	DF	156	ILE	3.9

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Mol	Chain	Res	Type	RSRZ
53	B5	26	ALA	3.9
53	B5	215	VAL	3.9
22	DA	2128	G	3.9
7	CG	123	GLU	3.9
33	DL	70	LYS	3.9
9	CI	119	ARG	3.9
41	DT	72	GLN	3.9
45	DX	49	LEU	3.9
53	B5	100	ILE	3.9
53	B5	101	ILE	3.9
1	AA	88	U	3.9
7	CG	81	GLY	3.9
22	BA	2141	G	3.9
22	DA	549	G	3.9
20	CT	38	ALA	3.9
24	DC	251	GLN	3.9
53	B5	211	ARG	3.9
26	DE	119	ILE	3.9
8	CH	90	ASP	3.9
22	DA	1095	A	3.9
18	CR	20	GLU	3.9
29	BH	129	GLU	3.9
1	AA	995	C	3.9
1	CA	1210	C	3.9
53	B5	165	ARG	3.9
7	CG	141	VAL	3.9
13	AM	5	ALA	3.9
27	DF	133	ARG	3.9
28	DG	78	GLY	3.9
10	CJ	73	LEU	3.8
13	CM	19	LEU	3.8
42	DU	48	PRO	3.8
13	CM	9	ILE	3.8
53	B5	126	SER	3.8
2	AB	131	LYS	3.8
14	CN	51	LEU	3.8
28	DG	103	ILE	3.8
1	AA	1017	U	3.8
1	AA	1032	G	3.8
1	CA	1305	G	3.8
22	DA	361	G	3.8
41	DT	55	VAL	3.8

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Mol	Chain	Res	Type	RSRZ
33	DL	79	LEU	3.8
2	AB	136	MET	3.8
30	BI	36	MET	3.8
22	DA	546	U	3.8
1	AA	1001	C	3.8
7	CG	144	MET	3.8
22	DA	2112	G	3.8
30	DI	36	MET	3.8
9	AI	32	GLN	3.8
29	BH	89	LYS	3.8
13	CM	54	ASP	3.8
30	BI	47	ASP	3.8
41	DT	15	HIS	3.8
17	CQ	8	LEU	3.8
29	DH	130	VAL	3.8
9	AI	17	ALA	3.8
14	AN	47	LYS	3.8
45	DX	22	LEU	3.8
20	CT	8	LYS	3.8
19	CS	11	ILE	3.8
22	DA	1056	G	3.8
30	DI	63	ALA	3.8
53	B5	154	ILE	3.8
1	CA	1271	A	3.8
13	CM	113	ARG	3.8
28	DG	166	ASP	3.8
29	BH	121	VAL	3.8
29	DH	12	LEU	3.8
30	BI	82	LYS	3.8
13	CM	38	GLY	3.8
1	AA	1537	U	3.8
10	AJ	59	LYS	3.8
3	CC	159	GLY	3.8
19	CS	12	ASP	3.8
26	DE	48	THR	3.8
7	CG	19	GLY	3.8
30	DI	110	ALA	3.8
44	DW	62	LYS	3.8
22	DA	1170	C	3.8
3	CC	120	ILE	3.8
26	DE	164	LEU	3.7
30	DI	106	LEU	3.7

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Mol	Chain	Res	Type	RSRZ
53	B5	69	LEU	3.7
1	AA	991	U	3.7
22	BA	139	U	3.7
41	DT	42	GLU	3.7
53	B5	53	ARG	3.7
9	AI	125	PRO	3.7
51	D3	2	PRO	3.7
39	DR	50	GLY	3.7
44	DW	63	ALA	3.7
19	CS	13	LEU	3.7
22	BA	2192	U	3.7
27	DF	117	LEU	3.7
26	DE	144	GLU	3.7
41	DT	59	ASN	3.7
51	D3	52	LYS	3.7
36	DO	58	ILE	3.7
13	CM	40	ALA	3.7
22	DA	2163	A	3.7
24	BC	237	GLY	3.7
35	DN	111	ALA	3.7
41	DT	83	ALA	3.7
13	AM	114	LYS	3.7
22	DA	34	U	3.7
30	DI	26	PRO	3.7
14	CN	11	VAL	3.7
47	DZ	8	THR	3.7
25	DD	76	GLY	3.7
3	CC	45	LYS	3.7
26	DE	118	LEU	3.7
36	DO	108	ASP	3.7
39	DR	19	THR	3.7
8	CH	122	GLY	3.7
13	CM	70	ARG	3.7
14	AN	78	GLY	3.7
19	AS	57	HIS	3.7
53	B5	198	GLU	3.7
30	BI	6	GLN	3.7
22	DA	896	A	3.7
1	AA	1037	C	3.7
30	DI	94	ASN	3.7
51	D3	65	ALA	3.7
1	CA	1351	U	3.7

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Mol	Chain	Res	Type	RSRZ
27	DF	93	GLY	3.7
30	DI	91	GLY	3.7
53	B5	144	GLY	3.7
25	DD	60	VAL	3.7
53	B5	48	LEU	3.7
30	DI	19	ASN	3.7
18	AR	74	HIS	3.7
6	CF	79	ARG	3.7
9	CI	47	VAL	3.7
22	BA	1925	C	3.7
9	CI	109	ARG	3.7
25	DD	56	LYS	3.7
26	DE	21	ARG	3.7
30	BI	71	THR	3.7
53	B5	45	HIS	3.7
22	DA	1530	G	3.7
26	DE	30	GLN	3.6
9	AI	121	ALA	3.6
1	CA	1217	C	3.6
9	AI	92	GLU	3.6
30	BI	30	GLN	3.6
36	DO	116	GLN	3.6
27	DF	121	SER	3.6
45	DX	78	TYR	3.6
50	D2	1	MET	3.6
40	DS	4	ILE	3.6
7	AG	4	ARG	3.6
9	AI	129	LYS	3.6
16	CP	39	PHE	3.6
7	CG	57	SER	3.6
13	CM	74	SER	3.6
26	DE	55	SER	3.6
19	CS	42	PRO	3.6
30	BI	83	ALA	3.6
9	AI	90	TYR	3.6
9	CI	130	ARG	3.6
38	DQ	74	ILE	3.6
53	B5	214	TYR	3.6
30	BI	91	GLY	3.6
40	DS	97	LEU	3.6
36	DO	38	GLN	3.6
36	DO	52	SER	3.6

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Mol	Chain	Res	Type	RSRZ
7	CG	38	THR	3.6
7	CG	86	GLN	3.6
30	DI	41	ALA	3.6
52	D4	9	LYS	3.6
1	AA	1492	A	3.6
22	DA	1026	G	3.6
22	DA	2799	A	3.6
22	DA	288	U	3.6
13	CM	81	MET	3.6
14	AN	33	ASP	3.6
20	CT	65	GLY	3.6
24	BC	233	GLY	3.6
29	DH	13	GLY	3.6
7	CG	54	SER	3.6
26	DE	98	LYS	3.6
1	AA	1036	A	3.6
33	DL	71	ALA	3.6
47	DZ	9	GLN	3.6
28	DG	57	GLY	3.6
36	DO	63	LYS	3.6
29	BH	93	SER	3.6
1	CA	1132	C	3.6
3	CC	71	ALA	3.6
22	DA	2313	C	3.6
1	CA	989	U	3.6
36	DO	87	ILE	3.6
7	CG	79	ARG	3.6
19	CS	58	VAL	3.6
28	DG	43	VAL	3.6
7	CG	131	LYS	3.6
46	DY	24	GLU	3.6
29	BH	11	ASN	3.6
19	AS	75	ALA	3.6
10	AJ	8	ILE	3.5
53	B5	19	LYS	3.5
30	BI	77	ALA	3.5
41	DT	43	ILE	3.5
42	DU	12	ILE	3.5
27	DF	151	GLY	3.5
9	CI	48	VAL	3.5
24	BC	238	ARG	3.5
1	AA	1021	A	3.5

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Mol	Chain	Res	Type	RSRZ
22	BA	882	G	3.5
22	BA	1068	G	3.5
13	AM	92	ARG	3.5
31	DJ	47	HIS	3.5
10	CJ	82	LYS	3.5
27	DF	12	VAL	3.5
9	CI	90	TYR	3.5
9	CI	106	ARG	3.5
1	CA	843	U	3.5
30	BI	24	VAL	3.5
18	CR	51	TYR	3.5
13	CM	35	ALA	3.5
13	CM	69	LEU	3.5
7	CG	8	GLY	3.5
36	DO	103	VAL	3.5
9	CI	103	PHE	3.5
20	CT	64	LYS	3.5
28	DG	177	LYS	3.5
1	CA	1275	A	3.5
2	CB	108	ARG	3.5
13	CM	8	ASN	3.5
27	DF	40	VAL	3.5
29	BH	147	VAL	3.5
53	B5	41	THR	3.5
28	DG	106	SER	3.5
46	DY	14	LEU	3.5
13	CM	98	ARG	3.5
8	AH	2	SER	3.5
22	DA	289	G	3.5
22	DA	356	G	3.5
22	DA	2801	G	3.5
32	DK	37	ASP	3.5
26	DE	143	LEU	3.5
38	DQ	73	GLY	3.5
42	BU	56	GLY	3.5
42	DU	19	LYS	3.5
9	AI	118	LEU	3.5
22	DA	885	C	3.5
52	D4	33	HIS	3.5
13	CM	37	ALA	3.5
30	DI	24	VAL	3.5
42	DU	13	VAL	3.5

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Mol	Chain	Res	Type	RSRZ
39	DR	29	THR	3.4
22	BA	879	G	3.4
22	BA	2128	G	3.4
53	B5	57	GLN	3.4
11	AK	19	GLY	3.4
13	CM	43	VAL	3.4
24	DC	249	GLY	3.4
46	DY	7	ARG	3.4
3	CC	108	LYS	3.4
30	DI	72	LYS	3.4
1	CA	1441	A	3.4
23	DB	18	G	3.4
36	DO	2	ASP	3.4
40	DS	9	HIS	3.4
27	DF	91	LEU	3.4
53	B5	178	LYS	3.4
1	CA	1214	C	3.4
22	DA	653	U	3.4
26	DE	131	THR	3.4
42	DU	27	ASN	3.4
14	CN	9	ARG	3.4
20	CT	75	HIS	3.4
22	DA	1045	C	3.4
27	DF	13	VAL	3.4
30	DI	73	THR	3.4
30	DI	28	LEU	3.4
3	AC	158	GLY	3.4
13	CM	67	GLY	3.4
22	DA	7	G	3.4
19	CS	76	PRO	3.4
24	DC	243	HIS	3.4
1	AA	1027	C	3.4
22	DA	1053	C	3.4
37	DP	102	GLU	3.4
51	D3	61	CYS	3.4
9	CI	57	MET	3.4
53	B5	71	LYS	3.4
1	CA	1244	G	3.4
7	CG	77	SER	3.4
24	DC	247	PRO	3.4
52	D4	1	MET	3.4
3	CC	87	LEU	3.4

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Mol	Chain	Res	Type	RSRZ
22	DA	1321	A	3.4
26	DE	41	GLN	3.4
7	AG	68	ASN	3.4
30	DI	43	ASN	3.4
20	CT	5	LYS	3.4
26	DE	22	ASP	3.4
28	DG	82	GLY	3.4
40	DS	94	ASP	3.4
22	BA	1847	A	3.4
48	D0	37	LYS	3.4
1	CA	1226	C	3.3
1	CA	1312	G	3.3
22	DA	88	G	3.3
29	DH	144	VAL	3.3
30	DI	92	LYS	3.3
13	AM	8	ASN	3.3
19	AS	13	LEU	3.3
29	BH	105	ALA	3.3
4	AD	28	ILE	3.3
22	DA	1046	A	3.3
41	DT	76	ARG	3.3
30	DI	23	PRO	3.3
13	CM	47	GLU	3.3
26	DE	47	LYS	3.3
30	DI	105	GLN	3.3
17	CQ	50	ASN	3.3
44	DW	55	ARG	3.3
1	CA	1023	U	3.3
1	CA	1295	U	3.3
14	AN	23	LYS	3.3
20	CT	34	LYS	3.3
27	BF	116	GLY	3.3
22	BA	1175	A	3.3
13	CM	2	ALA	3.3
22	DA	267	C	3.3
27	DF	64	LYS	3.3
40	DS	2	GLU	3.3
53	B5	143	ALA	3.3
17	CQ	5	ILE	3.3
1	CA	1240	U	3.3
9	CI	115	LYS	3.3
26	DE	42	GLY	3.3

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Mol	Chain	Res	Type	RSRZ
22	DA	2667	C	3.3
30	DI	30	GLN	3.3
44	DW	57	HIS	3.3
1	CA	1247	U	3.3
22	DA	2118	U	3.3
1	AA	1033	G	3.3
1	CA	971	G	3.3
53	B5	75	VAL	3.3
19	CS	41	PHE	3.3
26	DE	102	ARG	3.3
17	CQ	53	CYS	3.3
1	AA	1029	U	3.3
19	CS	67	VAL	3.3
29	BH	82	SER	3.3
33	DL	89	VAL	3.3
19	CS	44	MET	3.3
29	BH	132	PHE	3.3
3	AC	168	TYR	3.3
7	CG	68	ASN	3.3
22	BA	2186	G	3.3
1	CA	1016	A	3.3
20	CT	87	ALA	3.3
22	DA	1048	A	3.3
22	DA	2109	U	3.3
19	CS	31	LEU	3.3
28	DG	58	TYR	3.3
10	CJ	23	ALA	3.3
32	DK	99	ILE	3.3
49	D1	44	ARG	3.3
9	AI	117	GLY	3.3
41	DT	89	GLU	3.3
19	CS	25	SER	3.3
19	CS	64	ASP	3.3
30	BI	116	ASP	3.3
44	DW	56	ASP	3.3
25	DD	125	TRP	3.3
3	CC	207	ILE	3.3
46	BY	2	LYS	3.3
30	DI	116	ASP	3.3
27	DF	8	TYR	3.2
13	CM	51	GLY	3.2
53	B5	22	THR	3.2

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Mol	Chain	Res	Type	RSRZ
53	B5	193	PHE	3.2
47	DZ	56	LYS	3.2
1	AA	1004	A	3.2
22	DA	931	U	3.2
30	DI	37	GLU	3.2
1	CA	953	G	3.2
1	CA	1270	G	3.2
22	DA	117	G	3.2
10	AJ	73	LEU	3.2
10	CJ	57	VAL	3.2
14	CN	60	GLN	3.2
30	BI	95	LYS	3.2
44	DW	59	LEU	3.2
35	DN	20	MET	3.2
9	AI	89	GLU	3.2
22	BA	892	A	3.2
3	CC	203	PHE	3.2
13	CM	56	LEU	3.2
36	DO	20	GLU	3.2
9	AI	115	LYS	3.2
13	AM	32	ALA	3.2
13	CM	25	VAL	3.2
29	DH	61	VAL	3.2
30	DI	27	ALA	3.2
45	DX	18	ARG	3.2
22	BA	880	G	3.2
16	CP	47	GLU	3.2
33	DL	141	LYS	3.2
22	BA	2151	U	3.2
35	DN	28	LEU	3.2
27	DF	113	ASP	3.2
22	DA	2300	C	3.2
7	AG	109	ARG	3.2
13	AM	84	GLY	3.2
14	AN	48	LEU	3.2
17	CQ	44	LEU	3.2
2	CB	159	ASP	3.2
9	AI	33	ARG	3.2
13	CM	59	GLU	3.2
22	DA	228	C	3.2
22	DA	2000	C	3.2
53	B5	170	GLY	3.2

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Mol	Chain	Res	Type	RSRZ
20	CT	9	LYS	3.2
7	AG	74	GLU	3.2
7	CG	71	PRO	3.2
40	DS	82	MET	3.2
1	CA	90	C	3.2
16	AP	22	ALA	3.2
41	DT	87	LEU	3.2
22	BA	1067	A	3.2
22	DA	884	U	3.2
30	DI	71	THR	3.2
46	DY	55	THR	3.2
53	B5	208	THR	3.2
27	DF	51	ASP	3.2
27	DF	175	PHE	3.2
41	DT	16	VAL	3.2
17	CQ	4	LYS	3.2
22	DA	879	G	3.2
7	CG	143	ARG	3.2
26	DE	11	ALA	3.2
27	DF	2	ALA	3.2
22	DA	344	A	3.1
3	CC	205	GLY	3.1
13	AM	33	ILE	3.1
24	BC	241	GLY	3.1
35	DN	83	LEU	3.1
7	CG	122	ASN	3.1
9	AI	21	ILE	3.1
27	DF	105	THR	3.1
30	DI	33	VAL	3.1
22	DA	1089	A	3.1
22	DA	2800	A	3.1
3	CC	206	GLU	3.1
27	DF	120	LYS	3.1
17	CQ	61	ILE	3.1
26	DE	114	ARG	3.1
29	BH	122	LEU	3.1
1	AA	1019	A	3.1
1	CA	1287	A	3.1
22	BA	2109	U	3.1
10	CJ	78	GLU	3.1
13	CM	72	GLU	3.1
22	DA	2169	A	3.1

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Mol	Chain	Res	Type	RSRZ
14	CN	3	LYS	3.1
36	DO	55	GLU	3.1
39	DR	33	VAL	3.1
3	CC	181	ASP	3.1
14	CN	22	ALA	3.1
39	DR	28	ALA	3.1
22	BA	881	G	3.1
22	DA	1047	G	3.1
52	D4	12	ARG	3.1
7	CG	27	VAL	3.1
24	BC	239	ASN	3.1
2	CB	132	LYS	3.1
14	CN	12	LYS	3.1
24	DC	238	ARG	3.1
30	DI	123	GLU	3.1
33	DL	121	THR	3.1
15	CO	89	ARG	3.1
30	DI	38	PHE	3.1
49	B1	4	GLY	3.1
22	DA	1074	G	3.1
19	CS	65	GLU	3.1
25	DD	209	ALA	3.1
30	DI	55	ILE	3.1
35	DN	26	GLY	3.1
3	CC	126	ARG	3.1
22	DA	1092	C	3.1
53	B5	74	ARG	3.1
22	BA	1087	G	3.1
22	DA	1088	A	3.1
28	DG	8	PRO	3.1
36	DO	24	THR	3.1
2	CB	129	LEU	3.1
25	DD	126	ASN	3.1
7	AG	6	VAL	3.1
27	DF	10	ASP	3.1
30	DI	61	VAL	3.1
2	AB	139	ARG	3.1
10	AJ	33	GLY	3.1
22	DA	1210	G	3.1
42	DU	88	GLU	3.1
49	D1	46	HIS	3.1
1	CA	1218	C	3.1

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Mol	Chain	Res	Type	RSRZ
1	CA	1342	C	3.1
14	AN	50	THR	3.1
6	AF	61	LEU	3.1
26	DE	165	HIS	3.1
32	DK	68	GLY	3.1
37	DP	3	ASN	3.1
52	D4	37	GLN	3.1
3	CC	119	SER	3.0
49	D1	21	TYR	3.0
1	AA	1008	U	3.0
22	BA	1171	G	3.0
30	BI	94	ASN	3.0
7	AG	151	PHE	3.0
53	B5	125	GLY	3.0
17	CQ	63	GLU	3.0
22	DA	1622	G	3.0
26	DE	104	ALA	3.0
28	DG	50	LEU	3.0
32	DK	112	PHE	3.0
17	CQ	78	VAL	3.0
30	BI	55	ILE	3.0
30	DI	82	LYS	3.0
1	CA	950	U	3.0
22	DA	1090	A	3.0
19	AS	71	LEU	3.0
32	DK	83	ALA	3.0
25	DD	10	GLY	3.0
22	DA	1044	C	3.0
14	CN	23	LYS	3.0
30	DI	83	ALA	3.0
49	B1	53	LYS	3.0
14	CN	95	GLY	3.0
27	DF	131	GLY	3.0
49	D1	29	THR	3.0
1	CA	94	G	3.0
6	AF	35	LYS	3.0
22	DA	1238	G	3.0
35	DN	62	ASN	3.0
22	BA	2188	U	3.0
22	DA	1847	A	3.0
35	DN	24	MET	3.0
36	DO	92	PHE	3.0

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Mol	Chain	Res	Type	RSRZ
42	DU	87	PHE	3.0
30	DI	74	PRO	3.0
19	AS	3	ARG	3.0
1	CA	998	C	3.0
39	DR	101	ILE	3.0
22	DA	1071	G	3.0
1	CA	1321	U	3.0
22	DA	1173	U	3.0
1	CA	994	A	3.0
14	CN	55	SER	3.0
33	DL	8	PRO	3.0
12	CL	44	LYS	3.0
3	CC	3	GLN	3.0
19	CS	39	THR	3.0
27	DF	114	PHE	3.0
28	DG	116	GLN	3.0
1	CA	1276	G	3.0
10	CJ	45	ARG	3.0
13	AM	95	LEU	3.0
14	CN	65	ARG	3.0
22	BA	1179	G	3.0
37	DP	115	ASN	3.0
27	DF	115	ARG	3.0
38	DQ	59	GLN	3.0
46	DY	16	THR	3.0
42	BU	51	ALA	3.0
1	CA	1343	G	3.0
13	CM	86	TYR	3.0
31	DJ	95	ARG	3.0
3	AC	91	VAL	3.0
19	CS	72	GLY	3.0
33	DL	106	GLU	3.0
23	DB	118	C	3.0
49	D1	48	ILE	3.0
53	B5	129	GLY	3.0
2	AB	129	LEU	2.9
1	CA	1043	G	2.9
3	CC	102	ASN	2.9
1	CA	101	A	2.9
14	CN	26	GLU	2.9
9	CI	99	ARG	2.9
38	DQ	25	TYR	2.9

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Mol	Chain	Res	Type	RSRZ
1	CA	1320	C	2.9
28	DG	54	PRO	2.9
38	DQ	101	PHE	2.9
49	D1	14	SER	2.9
9	CI	92	GLU	2.9
25	DD	200	ASP	2.9
33	DL	38	GLN	2.9
39	DR	18	GLN	2.9
42	DU	28	VAL	2.9
29	BH	84	ALA	2.9
42	DU	35	ILE	2.9
48	D0	35	GLY	2.9
10	CJ	30	LYS	2.9
20	CT	76	LYS	2.9
1	CA	121	U	2.9
3	CC	91	VAL	2.9
23	DB	119	A	2.9
25	DD	145	SER	2.9
7	AG	48	GLU	2.9
3	AC	81	GLY	2.9
29	DH	81	ALA	2.9
22	DA	2666	C	2.9
25	DD	186	LEU	2.9
28	DG	25	THR	2.9
42	DU	14	LEU	2.9
25	DD	74	GLU	2.9
37	DP	33	VAL	2.9
37	DP	103	ARG	2.9
19	AS	50	ALA	2.9
34	DM	79	ALA	2.9
44	DW	33	ALA	2.9
30	DI	42	PHE	2.9
34	DM	88	ASN	2.9
8	CH	55	THR	2.9
28	DG	49	THR	2.9
30	BI	140	VAL	2.9
1	AA	1000	A	2.9
21	AU	7	ARG	2.9
13	CM	13	LYS	2.9
20	CT	71	LYS	2.9
3	CC	179	ARG	2.9
10	AJ	60	ASP	2.9

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Mol	Chain	Res	Type	RSRZ
3	CC	125	GLU	2.9
10	CJ	15	HIS	2.9
20	AT	68	HIS	2.9
25	DD	88	GLU	2.9
26	DE	24	ASN	2.9
27	DF	14	LYS	2.9
30	BI	40	LYS	2.9
40	DS	106	VAL	2.9
42	DU	42	VAL	2.9
46	DY	41	HIS	2.9
22	DA	1064	C	2.9
22	DA	1076	C	2.9
39	DR	1	MET	2.9
44	DW	61	ALA	2.9
28	DG	151	TYR	2.9
30	BI	98	VAL	2.9
30	DI	16	GLY	2.9
13	CM	57	ARG	2.9
14	CN	6	MET	2.9
29	DH	84	ALA	2.9
14	CN	7	LYS	2.9
50	D2	37	LYS	2.9
1	CA	83	C	2.9
1	CA	1208	C	2.9
26	DE	127	GLU	2.9
1	CA	954	G	2.9
22	BA	883	G	2.9
22	DA	881	G	2.9
51	D3	58	VAL	2.9
22	BA	2098	U	2.9
30	BI	97	LYS	2.9
46	DY	9	LYS	2.9
7	CG	84	THR	2.9
7	CG	73	VAL	2.9
14	CN	45	VAL	2.9
14	CN	54	ASP	2.9
20	AT	36	TYR	2.9
11	AK	14	LYS	2.8
40	DS	90	LYS	2.8
2	CB	104	TRP	2.8
13	CM	5	ALA	2.8
14	CN	8	ALA	2.8

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Mol	Chain	Res	Type	RSRZ
22	DA	914	G	2.8
13	CM	42	ASP	2.8
26	DE	88	ARG	2.8
30	BI	48	SER	2.8
32	DK	111	LYS	2.8
1	CA	1492	A	2.8
38	DQ	37	GLN	2.8
22	BA	1729	U	2.8
14	CN	63	ARG	2.8
2	CB	40	ILE	2.8
3	CC	78	GLY	2.8
29	DH	15	LEU	2.8
36	DO	107	ALA	2.8
22	DA	1211	C	2.8
22	DA	2158	A	2.8
24	DC	47	GLY	2.8
3	AC	93	ASP	2.8
22	DA	45	G	2.8
29	DH	117	LEU	2.8
30	DI	64	ASP	2.8
1	CA	949	A	2.8
22	DA	1111	A	2.8
13	CM	22	ILE	2.8
22	BA	1066	U	2.8
22	DA	2181	U	2.8
26	DE	17	THR	2.8
30	DI	49	ILE	2.8
26	DE	23	PHE	2.8
28	DG	84	THR	2.8
16	CP	48	GLU	2.8
48	D0	2	ALA	2.8
7	AG	75	VAL	2.8
7	AG	80	VAL	2.8
42	DU	25	VAL	2.8
3	AC	192	THR	2.8
22	DA	1439	A	2.8
22	DA	2170	A	2.8
28	DG	51	THR	2.8
28	DG	86	LYS	2.8
30	BI	106	LEU	2.8
7	CG	51	ALA	2.8
9	CI	37	GLN	2.8

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Mol	Chain	Res	Type	RSRZ
41	DT	67	VAL	2.8
9	CI	4	ASN	2.8
42	BU	57	GLY	2.8
9	AI	124	ARG	2.8
48	D0	36	GLU	2.8
49	B1	52	ALA	2.8
1	CA	207	C	2.8
35	DN	72	ASP	2.8
19	AS	5	LEU	2.8
21	CU	38	TYR	2.8
29	DH	83	LYS	2.8
1	AA	1018	G	2.8
1	CA	1033	G	2.8
22	DA	882	G	2.8
12	AL	14	ARG	2.8
27	DF	78	LYS	2.8
13	CM	97	VAL	2.8
26	DE	199	MET	2.8
27	DF	149	VAL	2.8
3	CC	190	HIS	2.8
20	CT	24	ARG	2.8
39	DR	87	GLN	2.8
46	DY	44	LYS	2.8
22	DA	2903	U	2.8
7	CG	47	LEU	2.8
10	CJ	34	ALA	2.8
26	DE	201	ALA	2.8
25	DD	8	LYS	2.8
30	DI	127	ARG	2.8
53	B5	124	VAL	2.8
28	DG	31	GLY	2.8
31	DJ	89	PHE	2.8
14	CN	57	PRO	2.8
3	CC	39	VAL	2.8
9	AI	19	VAL	2.8
10	CJ	9	ARG	2.8
25	DD	55	LYS	2.8
37	DP	92	VAL	2.8
42	DU	21	LYS	2.8
32	DK	50	GLY	2.7
41	DT	74	ILE	2.7
7	CG	139	GLU	2.7

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Mol	Chain	Res	Type	RSRZ
34	DM	103	TYR	2.7
9	AI	23	PRO	2.7
13	CM	61	ALA	2.7
13	AM	24	GLY	2.7
22	DA	1052	C	2.7
42	DU	29	LEU	2.7
1	AA	81	A	2.7
1	AA	1022	A	2.7
1	CA	1339	A	2.7
9	CI	89	GLU	2.7
12	CL	123	LYS	2.7
22	DA	2877	G	2.7
25	DD	154	LYS	2.7
39	DR	34	GLU	2.7
14	CN	35	ASN	2.7
25	DD	77	ARG	2.7
7	CG	72	THR	2.7
41	DT	39	THR	2.7
1	CA	1331	G	2.7
22	DA	317	G	2.7
22	DA	329	G	2.7
3	CC	202	ILE	2.7
14	CN	28	LYS	2.7
22	DA	2122	U	2.7
30	BI	137	GLY	2.7
52	D4	8	LYS	2.7
28	DG	110	SER	2.7
28	DG	176	LYS	2.7
1	CA	975	A	2.7
30	DI	117	MET	2.7
13	CM	34	LEU	2.7
28	DG	26	ILE	2.7
30	DI	101	ILE	2.7
1	AA	993	G	2.7
14	CN	43	ASN	2.7
16	AP	80	LYS	2.7
28	DG	44	LYS	2.7
28	DG	12	PRO	2.7
14	CN	32	SER	2.7
9	AI	5	GLN	2.7
13	CM	52	GLN	2.7
1	AA	1039	G	2.7

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Mol	Chain	Res	Type	RSRZ
1	CA	1310	G	2.7
22	BA	2152	G	2.7
22	DA	776	G	2.7
43	DV	6	ALA	2.7
34	DM	124	LEU	2.7
30	DI	102	SER	2.7
29	BH	73	ASN	2.7
39	DR	63	VAL	2.7
30	BI	7	ALA	2.7
30	BI	120	ALA	2.7
14	AN	72	GLY	2.7
26	DE	12	LEU	2.7
28	DG	167	GLU	2.7
3	CC	165	THR	2.7
7	AG	133	THR	2.7
40	DS	6	LYS	2.7
38	DQ	6	ARG	2.7
1	CA	1245	C	2.7
36	DO	26	LEU	2.7
3	CC	118	ASP	2.7
25	DD	105	LYS	2.7
25	DD	121	THR	2.7
46	DY	31	GLN	2.7
36	DO	39	VAL	2.7
48	D0	3	VAL	2.7
13	AM	111	GLY	2.7
1	CA	1237	C	2.7
1	CA	1286	U	2.7
22	DA	32	C	2.7
22	DA	1049	C	2.7
1	CA	1248	A	2.7
36	DO	54	VAL	2.7
49	D1	43	VAL	2.7
16	CP	17	TYR	2.7
13	AM	83	LEU	2.7
15	CO	15	PHE	2.7
26	DE	72	SER	2.7
53	B5	135	ARG	2.7
14	AN	54	ASP	2.7
19	CS	27	ASP	2.7
27	DF	28	VAL	2.7
38	DQ	44	GLN	2.7

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Mol	Chain	Res	Type	RSRZ
9	AI	39	PHE	2.6
10	CJ	11	LYS	2.6
22	DA	2171	A	2.6
30	DI	84	ALA	2.6
30	DI	104	ALA	2.6
20	CT	79	LEU	2.6
22	DA	275	C	2.6
24	DC	48	ARG	2.6
28	DG	40	ALA	2.6
38	DQ	2	ALA	2.6
51	D3	22	PHE	2.6
43	DV	33	GLY	2.6
44	BW	10	THR	2.6
53	B5	50	ILE	2.6
1	CA	1241	G	2.6
22	BA	2193	G	2.6
30	BI	139	VAL	2.6
35	DN	116	VAL	2.6
38	DQ	39	VAL	2.6
43	DV	34	LYS	2.6
53	B5	44	VAL	2.6
35	DN	73	ASN	2.6
44	DW	50	ASN	2.6
32	DK	49	ARG	2.6
33	DL	107	PHE	2.6
39	DR	37	GLU	2.6
19	CS	75	ALA	2.6
26	DE	128	ALA	2.6
33	DL	49	GLY	2.6
37	DP	95	ALA	2.6
3	AC	62	LYS	2.6
16	CP	80	LYS	2.6
41	DT	33	LYS	2.6
14	CN	13	ARG	2.6
28	DG	22	GLN	2.6
17	CQ	73	TRP	2.6
19	CS	68	GLY	2.6
23	DB	20	G	2.6
3	CC	196	ILE	2.6
22	DA	877	A	2.6
26	DE	50	ALA	2.6
40	DS	5	ALA	2.6

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Mol	Chain	Res	Type	RSRZ
22	DA	1523	U	2.6
28	BG	111	HIS	2.6
29	BH	128	HIS	2.6
29	DH	82	SER	2.6
37	DP	109	ARG	2.6
29	BH	139	PHE	2.6
37	DP	12	GLN	2.6
32	DK	89	ASN	2.6
7	AG	7	ILE	2.6
30	DI	86	ILE	2.6
30	DI	114	ALA	2.6
9	AI	104	VAL	2.6
22	DA	2802	G	2.6
30	DI	65	ARG	2.6
3	CC	70	THR	2.6
10	CJ	50	THR	2.6
19	CS	61	PHE	2.6
3	CC	145	GLY	2.6
29	BH	92	GLY	2.6
38	DQ	7	GLY	2.6
1	CA	210	C	2.6
1	CA	962	C	2.6
3	CC	93	ASP	2.6
7	CG	23	LEU	2.6
35	DN	21	PHE	2.6
14	AN	60	GLN	2.6
22	BA	2191	A	2.6
22	DA	2121	G	2.6
27	DF	118	SER	2.6
10	AJ	7	ARG	2.6
27	DF	112	ARG	2.6
29	DH	143	ILE	2.6
31	DJ	93	ILE	2.6
26	DE	126	VAL	2.6
36	DO	56	LYS	2.6
40	DS	36	LEU	2.6
1	AA	997	U	2.6
2	AB	74	ARG	2.6
7	CG	78	ARG	2.6
32	DK	101	GLY	2.6
16	CP	57	ILE	2.6
37	DP	84	ILE	2.6

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Mol	Chain	Res	Type	RSRZ
20	CT	63	ALA	2.6
36	DO	109	ALA	2.6
37	DP	85	SER	2.6
1	CA	968	A	2.6
22	BA	1093	G	2.6
22	DA	2116	G	2.6
22	DA	2895	G	2.6
26	DE	129	PRO	2.6
16	CP	52	LEU	2.6
27	DF	177	PHE	2.6
51	D3	14	PHE	2.6
1	CA	948	C	2.6
19	AS	55	ARG	2.6
22	BA	1172	C	2.6
22	DA	902	C	2.6
7	CG	9	GLN	2.6
28	DG	27	LYS	2.6
30	DI	87	LYS	2.6
17	CQ	23	VAL	2.6
1	CA	1004	A	2.6
7	CG	40	GLU	2.6
9	AI	88	MET	2.6
30	DI	141	GLU	2.6
36	DO	106	LEU	2.6
3	CC	131	ARG	2.6
22	DA	75	G	2.6
22	DA	1452	G	2.6
22	DA	2803	G	2.6
22	DA	268	C	2.6
26	DE	65	THR	2.6
30	BI	76	ALA	2.6
44	DW	31	VAL	2.6
22	DA	1176	U	2.6
13	CM	110	LYS	2.5
20	CT	82	GLN	2.5
22	BA	1089	A	2.5
33	DL	73	ILE	2.5
52	D4	32	LYS	2.5
53	B5	153	ILE	2.5
1	CA	202	G	2.5
40	DS	95	ARG	2.5
27	DF	96	MET	2.5

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Mol	Chain	Res	Type	RSRZ
22	DA	1534	U	2.5
27	DF	97	TRP	2.5
45	DX	17	ASN	2.5
49	D1	40	ASP	2.5
24	BC	247	PRO	2.5
1	CA	1452	C	2.5
14	CN	42	TRP	2.5
22	DA	897	C	2.5
26	DE	191	ASP	2.5
27	DF	107	ALA	2.5
52	D4	35	GLN	2.5
3	CC	106	VAL	2.5
3	CC	173	VAL	2.5
9	AI	6	TYR	2.5
50	D2	46	LYS	2.5
1	CA	81	A	2.5
48	D0	55	ILE	2.5
3	CC	88	ARG	2.5
7	AG	141	VAL	2.5
13	AM	106	ALA	2.5
19	AS	69	HIS	2.5
28	DG	20	ASN	2.5
29	BH	137	GLU	2.5
29	DH	140	ALA	2.5
11	CK	126	LYS	2.5
29	BH	90	LEU	2.5
7	CG	14	PRO	2.5
25	DD	133	THR	2.5
26	DE	13	THR	2.5
27	DF	159	THR	2.5
33	DL	68	SER	2.5
49	D1	13	SER	2.5
7	CG	67	GLU	2.5
13	CM	78	LYS	2.5
14	CN	15	ALA	2.5
39	DR	66	HIS	2.5
48	D0	31	ASP	2.5
14	AN	64	CYS	2.5
22	DA	11	C	2.5
22	DA	2833	U	2.5
34	DM	99	GLY	2.5
46	DY	1	MET	2.5

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Mol	Chain	Res	Type	RSRZ
7	CG	12	ILE	2.5
14	AN	24	ARG	2.5
51	D3	21	GLY	2.5
19	CS	4	SER	2.5
9	AI	64	TYR	2.5
10	CJ	20	GLN	2.5
13	CM	65	VAL	2.5
27	DF	7	TYR	2.5
29	DH	6	LEU	2.5
41	DT	37	ASP	2.5
22	DA	1214	A	2.5
32	DK	90	ASN	2.5
7	AG	78	ARG	2.5
13	CM	87	ARG	2.5
38	DQ	30	ARG	2.5
13	CM	62	LYS	2.5
2	CB	213	TYR	2.5
6	CF	39	LEU	2.5
7	CG	83	SER	2.5
14	AN	2	ALA	2.5
14	AN	55	SER	2.5
19	CS	38	SER	2.5
29	BH	14	SER	2.5
36	DO	61	GLN	2.5
36	DO	113	ALA	2.5
26	DE	91	ASP	2.5
22	DA	332	A	2.5
22	DA	2766	A	2.5
8	CH	60	GLU	2.5
21	CU	8	GLU	2.5
33	DL	50	PHE	2.5
36	DO	74	VAL	2.5
27	DF	54	ALA	2.5
29	DH	133	GLN	2.5
33	DL	132	ARG	2.5
14	AN	12	LYS	2.5
22	BA	1063	G	2.5
13	AM	4	ILE	2.5
13	CM	50	GLU	2.5
7	AG	71	PRO	2.5
13	AM	19	LEU	2.5
22	BA	613	A	2.5

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Mol	Chain	Res	Type	RSRZ
42	BU	50	PRO	2.5
7	AG	110	LYS	2.5
30	DI	81	LYS	2.5
27	DF	142	ASP	2.5
29	BH	17	ASP	2.5
10	CJ	25	ILE	2.5
22	BA	2190	G	2.5
22	DA	139	U	2.5
3	CC	42	TYR	2.5
22	DA	2119	A	2.5
10	CJ	38	GLY	2.5
9	CI	116	VAL	2.5
1	AA	4	U	2.4
1	AA	1026	G	2.4
1	CA	1297	G	2.4
22	DA	1450	G	2.4
22	DA	1524	G	2.4
1	AA	1042	A	2.4
3	AC	161	GLU	2.4
9	CI	41	ARG	2.4
13	AM	82	ASP	2.4
22	BA	896	A	2.4
25	DD	59	ARG	2.4
2	AB	90	PHE	2.4
13	AM	89	LEU	2.4
22	DA	867	C	2.4
27	DF	21	ASN	2.4
29	BH	20	ASN	2.4
33	DL	127	VAL	2.4
41	DT	88	LYS	2.4
2	CB	136	MET	2.4
2	CB	69	PHE	2.4
30	BI	118	THR	2.4
42	DU	59	VAL	2.4
1	AA	1041	G	2.4
1	AA	412	A	2.4
22	DA	1278	C	2.4
37	DP	91	ALA	2.4
10	AJ	78	GLU	2.4
27	DF	79	ILE	2.4
47	DZ	48	ILE	2.4
9	AI	48	VAL	2.4

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Mol	Chain	Res	Type	RSRZ
3	CC	83	ASP	2.4
6	AF	66	ALA	2.4
20	AT	3	ASN	2.4
29	DH	74	ALA	2.4
30	BI	19	ASN	2.4
1	CA	1368	A	2.4
21	AU	38	TYR	2.4
22	DA	245	G	2.4
22	DA	2860	A	2.4
43	DV	57	TYR	2.4
10	AJ	10	LEU	2.4
14	CN	48	LEU	2.4
29	DH	58	LEU	2.4
7	AG	140	ASP	2.4
19	AS	38	SER	2.4
19	AS	48	THR	2.4
26	DE	173	THR	2.4
4	AD	44	ARG	2.4
12	CL	14	ARG	2.4
29	BH	116	ARG	2.4
42	DU	63	ALA	2.4
30	DI	137	GLY	2.4
10	CJ	24	GLU	2.4
1	CA	1000	A	2.4
22	DA	666	A	2.4
25	DD	165	MET	2.4
39	DR	92	TRP	2.4
25	DD	25	THR	2.4
39	DR	103	ALA	2.4
30	DI	29	GLY	2.4
24	DC	231	PRO	2.4
10	AJ	90	LEU	2.4
16	CP	54	LEU	2.4
50	D2	42	LEU	2.4
53	B5	186	LEU	2.4
10	AJ	62	ARG	2.4
14	CN	53	ARG	2.4
1	CA	1002	G	2.4
3	CC	122	SER	2.4
30	DI	51	LYS	2.4
37	DP	93	ARG	2.4
46	DY	47	ARG	2.4

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Mol	Chain	Res	Type	RSRZ
30	BI	39	CYS	2.4
13	AM	41	GLU	2.4
1	AA	1035	A	2.4
4	CD	28	ILE	2.4
13	AM	7	ILE	2.4
22	DA	357	C	2.4
13	CM	3	ARG	2.4
22	DA	501	A	2.4
41	DT	8	LEU	2.4
1	AA	82	G	2.4
34	DM	17	ASN	2.4
49	D1	31	PRO	2.4
36	DO	80	GLU	2.4
53	B5	159	ALA	2.4
6	AF	36	ILE	2.4
7	CG	50	LEU	2.4
9	AI	63	LEU	2.4
10	AJ	36	VAL	2.4
28	DG	83	PHE	2.4
22	DA	33	C	2.4
22	DA	476	G	2.4
36	DO	59	ALA	2.4
36	DO	82	ALA	2.4
49	D1	25	LYS	2.4
31	DJ	119	PHE	2.4
41	DT	50	LEU	2.4
53	B5	164	PHE	2.4
30	DI	142	ASP	2.4
28	DG	74	SER	2.4
1	AA	998	C	2.4
22	BA	1072	C	2.4
22	BA	2129	C	2.4
13	AM	3	ARG	2.4
19	CS	28	LYS	2.4
22	DA	2712	C	2.4
22	DA	2767	C	2.4
25	DD	38	LYS	2.4
30	DI	79	LEU	2.4
37	DP	97	LEU	2.4
1	CA	68	G	2.3
22	DA	1179	G	2.3
22	DA	2890	G	2.3

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Mol	Chain	Res	Type	RSRZ
33	DL	80	SER	2.3
7	AG	53	ARG	2.3
9	CI	68	LYS	2.3
9	CI	53	GLU	2.3
2	CB	148	LEU	2.3
20	CT	68	HIS	2.3
29	BH	99	ILE	2.3
22	DA	513	A	2.3
44	DW	51	VAL	2.3
3	CC	164	ARG	2.3
32	DK	14	SER	2.3
1	CA	102	G	2.3
2	CB	84	ALA	2.3
29	BH	145	ASN	2.3
1	CA	1264	U	2.3
3	AC	39	VAL	2.3
22	DA	1539	U	2.3
42	DU	47	LYS	2.3
13	CM	41	GLU	2.3
16	CP	53	ASP	2.3
25	DD	89	GLU	2.3
26	BE	7	ASP	2.3
38	DQ	21	ALA	2.3
40	DS	48	LYS	2.3
1	AA	79	G	2.3
22	DA	70	G	2.3
22	DA	1112	G	2.3
22	DA	41	C	2.3
22	DA	1518	C	2.3
36	DO	114	GLY	2.3
44	DW	64	ASP	2.3
53	B5	167	ASP	2.3
2	CB	164	ILE	2.3
7	CG	151	PHE	2.3
32	DK	104	THR	2.3
49	D1	39	PHE	2.3
19	CS	81	ARG	2.3
36	DO	30	ARG	2.3
22	DA	1108	U	2.3
22	DA	2111	U	2.3
1	AA	1138	G	2.3
14	CN	18	ASP	2.3

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Mol	Chain	Res	Type	RSRZ
1	CA	1137	C	2.3
7	CG	65	ALA	2.3
9	CI	44	ALA	2.3
22	DA	1407	G	2.3
30	BI	42	PHE	2.3
32	DK	53	LYS	2.3
36	DO	89	ASP	2.3
22	DA	2165	C	2.3
25	DD	31	ALA	2.3
25	DD	132	ALA	2.3
28	DG	7	ALA	2.3
10	CJ	96	VAL	2.3
46	DY	48	ARG	2.3
1	CA	1035	A	2.3
39	DR	59	ILE	2.3
28	DG	126	PRO	2.3
22	BA	1071	G	2.3
23	DB	19	C	2.3
9	AI	112	GLU	2.3
1	CA	205	A	2.3
22	DA	505	A	2.3
39	DR	88	GLY	2.3
40	DS	110	ARG	2.3
46	DY	29	ARG	2.3
8	CH	130	ALA	2.3
22	BA	546	U	2.3
40	DS	109	ASP	2.3
33	DL	143	GLU	2.3
35	DN	56	LYS	2.3
33	DL	102	GLY	2.3
36	DO	19	GLN	2.3
1	CA	1013	G	2.3
14	AN	27	LEU	2.3
47	DZ	53	PHE	2.3
30	BI	86	ILE	2.3
46	DY	4	LYS	2.3
9	CI	110	GLN	2.3
43	DV	32	GLY	2.3
51	D3	49	MET	2.3
22	DA	1764	C	2.3
1	CA	1246	A	2.3
3	AC	193	TYR	2.3

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Mol	Chain	Res	Type	RSRZ
15	CO	22	THR	2.3
42	DU	17	LYS	2.3
1	CA	1136	C	2.3
2	CB	18	HIS	2.3
7	AG	66	LEU	2.3
19	CS	26	GLY	2.3
22	BA	846	U	2.3
27	DF	66	LEU	2.3
41	DT	75	GLY	2.3
2	CB	107	VAL	2.3
9	CI	65	ILE	2.3
22	DA	654	A	2.3
22	DA	1245	G	2.3
51	D3	57	LEU	2.3
14	CN	67	THR	2.3
37	DP	38	LYS	2.3
41	BT	69	ARG	2.2
50	D2	36	ALA	2.2
53	B5	191	ARG	2.2
20	CT	12	ILE	2.2
30	DI	109	ILE	2.2
48	D0	30	VAL	2.2
14	AN	63	ARG	2.2
14	CN	41	ARG	2.2
35	DN	46	ARG	2.2
3	AC	157	LEU	2.2
9	CI	64	TYR	2.2
9	CI	91	ASP	2.2
27	BF	83	TYR	2.2
31	DJ	38	GLY	2.2
19	CS	3	ARG	2.2
27	DF	92	ARG	2.2
44	DW	23	VAL	2.2
7	CG	60	GLU	2.2
7	CG	117	ALA	2.2
16	CP	45	GLU	2.2
32	DK	106	GLU	2.2
19	CS	77	THR	2.2
40	DS	108	SER	2.2
53	B5	169	THR	2.2
1	CA	969	A	2.2
7	CG	110	LYS	2.2

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Mol	Chain	Res	Type	RSRZ
1	AA	963	G	2.2
22	DA	883	G	2.2
27	BF	130	MET	2.2
28	DG	132	VAL	2.2
40	DS	20	VAL	2.2
46	DY	38	GLN	2.2
53	B5	162	ILE	2.2
3	CC	92	ALA	2.2
35	DN	82	GLU	2.2
1	CA	1038	C	2.2
20	CT	6	SER	2.2
22	DA	90	U	2.2
22	DA	316	C	2.2
28	BG	177	LYS	2.2
10	CJ	32	THR	2.2
2	CB	135	LEU	2.2
2	CB	161	LEU	2.2
26	DE	200	LEU	2.2
33	DL	140	GLY	2.2
22	DA	631	A	2.2
36	DO	90	VAL	2.2
25	DD	168	GLU	2.2
49	D1	7	GLU	2.2
1	CA	1309	G	2.2
1	AA	1025	U	2.2
22	BA	893	C	2.2
22	DA	281	C	2.2
30	BI	61	VAL	2.2
42	DU	99	ASN	2.2
3	AC	46	GLU	2.2
4	CD	177	LYS	2.2
7	CG	36	LYS	2.2
10	AJ	30	LYS	2.2
30	DI	100	LYS	2.2
34	DM	84	LYS	2.2
29	BH	64	ALA	2.2
53	B5	51	ASP	2.2
22	DA	6	A	2.2
22	DA	2602	A	2.2
3	CC	37	PHE	2.2
41	DT	69	ARG	2.2
7	CG	56	LYS	2.2

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Mol	Chain	Res	Type	RSRZ
11	AK	126	LYS	2.2
22	DA	2129	C	2.2
22	DA	2177	C	2.2
41	DT	25	GLU	2.2
9	AI	44	ALA	2.2
25	DD	144	GLY	2.2
19	CS	40	ILE	2.2
31	DJ	55	ILE	2.2
31	DJ	88	THR	2.2
9	CI	32	GLN	2.2
1	CA	87	C	2.2
1	CA	203	G	2.2
13	CM	90	ARG	2.2
19	AS	32	ARG	2.2
22	DA	2106	U	2.2
46	DY	20	ASN	2.2
50	D2	34	ARG	2.2
42	DU	81	ASP	2.2
27	DF	77	PHE	2.2
33	DL	124	GLY	2.2
53	B5	103	LYS	2.2
2	CB	151	ILE	2.2
25	DD	96	ILE	2.2
28	DG	169	VAL	2.2
33	DL	77	ILE	2.2
38	DQ	98	ILE	2.2
40	DS	105	VAL	2.2
7	AG	49	THR	2.2
22	BA	2119	A	2.2
36	DO	13	ARG	2.2
36	DO	57	ALA	2.2
1	AA	121	U	2.2
39	DR	26	ASP	2.2
8	CH	110	VAL	2.2
22	DA	277	G	2.2
22	DA	333	G	2.2
35	DN	52	ILE	2.2
40	DS	8	ARG	2.2
25	DD	97	SER	2.2
9	CI	114	LYS	2.2
18	AR	68	LEU	2.2
3	CC	85	GLU	2.2

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Mol	Chain	Res	Type	RSRZ
7	CG	3	ARG	2.2
29	BH	76	GLU	2.2
22	DA	1531	C	2.2
1	CA	1048	G	2.2
9	AI	22	LYS	2.2
26	DE	9	GLN	2.2
27	BF	72	LYS	2.2
33	DL	72	ALA	2.2
14	CN	78	GLY	2.2
29	BH	118	PRO	2.2
2	AB	127	ASP	2.2
2	CB	111	ILE	2.2
22	DA	2131	U	2.1
22	DA	2796	U	2.1
29	DH	94	ILE	2.2
42	BU	58	ILE	2.2
45	DX	47	VAL	2.2
16	CP	1	MET	2.1
3	CC	123	GLN	2.1
22	DA	183	C	2.1
22	DA	314	C	2.1
37	DP	2	SER	2.1
53	B5	137	LEU	2.1
7	CG	112	GLY	2.1
14	CN	61	ARG	2.1
17	CQ	70	THR	2.1
7	CG	90	GLU	2.1
9	AI	107	ASP	2.1
22	DA	859	G	2.1
32	DK	56	ASP	2.1
40	DS	52	GLU	2.1
45	DX	4	VAL	2.1
53	B5	120	VAL	2.1
28	DG	99	LYS	2.1
43	DV	43	ASP	2.1
1	CA	216	U	2.1
10	AJ	71	LEU	2.1
18	CR	48	ARG	2.1
21	CU	7	ARG	2.1
27	BF	117	LEU	2.1
34	DM	6	ARG	2.1
36	DO	25	ARG	2.1

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Mol	Chain	Res	Type	RSRZ
46	DY	21	LEU	2.1
9	AI	34	SER	2.1
49	D1	18	GLY	2.1
2	AB	67	ILE	2.1
37	DP	42	ALA	2.1
11	CK	43	GLY	2.1
22	DA	89	A	2.1
24	DC	38	SER	2.1
28	DG	130	GLU	2.1
31	DJ	73	VAL	2.1
48	D0	8	PRO	2.1
3	AC	43	LEU	2.1
3	CC	142	MET	2.1
13	CM	93	ARG	2.1
27	BF	113	ASP	2.1
36	DO	93	ASP	2.1
1	CA	961	U	2.1
10	CJ	64	GLN	2.1
30	DI	111	GLN	2.1
33	DL	113	ALA	2.1
22	DA	1099	G	2.1
7	AG	2	PRO	2.1
17	CQ	6	ARG	2.1
17	CQ	65	ARG	2.1
19	CS	14	HIS	2.1
22	BA	2170	A	2.1
22	DA	213	A	2.1
22	DA	1213	A	2.1
38	DQ	33	ARG	2.1
3	CC	124	LEU	2.1
33	DL	19	LEU	2.1
1	CA	470	C	2.1
1	CA	1265	C	2.1
9	CI	121	ALA	2.1
30	DI	39	CYS	2.1
39	DR	24	LYS	2.1
39	DR	62	GLU	2.1
3	CC	77	ILE	2.1
14	AN	52	PRO	2.1
1	CA	1133	G	2.1
7	CG	35	LYS	2.1
19	AS	31	LEU	2.1

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Mol	Chain	Res	Type	RSRZ
22	DA	1984	G	2.1
22	DA	2107	G	2.1
48	D0	38	HIS	2.1
1	CA	1340	A	2.1
22	DA	1134	A	2.1
24	DC	225	MET	2.1
1	AA	1038	C	2.1
4	CD	25	VAL	2.1
14	CN	66	GLN	2.1
36	DO	78	VAL	2.1
7	CG	10	ARG	2.1
26	DE	138	LEU	2.1
26	DE	158	PHE	2.1
36	DO	76	LYS	2.1
1	CA	79	G	2.1
3	CC	84	VAL	2.1
1	CA	1042	A	2.1
1	CA	1274	A	2.1
3	AC	94	ILE	2.1
7	CG	92	ARG	2.1
9	CI	83	ILE	2.1
9	CI	95	ARG	2.1
14	AN	101	TRP	2.1
26	DE	162	ARG	2.1
32	DK	98	ARG	2.1
35	DN	118	ARG	2.1
39	DR	47	VAL	2.1
48	D0	5	GLN	2.1
38	DQ	117	LEU	2.1
26	DE	75	SER	2.1
37	DP	65	SER	2.1
31	DJ	98	GLU	2.1
46	DY	3	ALA	2.1
9	AI	67	VAL	2.1
9	AI	116	VAL	2.1
14	CN	14	VAL	2.1
1	CA	86	G	2.1
1	CA	1044	A	2.1
16	CP	38	PHE	2.1
22	DA	544	C	2.1
22	DA	901	C	2.1
2	CB	206	ALA	2.1

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Mol	Chain	Res	Type	RSRZ
24	DC	172	VAL	2.1
33	DL	15	ALA	2.1
42	DU	76	ALA	2.1
47	DZ	2	ALA	2.1
15	CO	25	THR	2.1
40	DS	49	LYS	2.1
44	DW	43	THR	2.1
27	DF	152	LEU	2.1
27	DF	29	PRO	2.1
27	DF	143	TYR	2.1
2	AB	27	MET	2.1
6	AF	95	ALA	2.1
13	AM	64	VAL	2.1
19	CS	18	LYS	2.1
22	DA	1069	A	2.1
32	DK	60	ALA	2.1
49	D1	30	LYS	2.1
1	AA	999	C	2.1
1	CA	967	C	2.1
1	CA	1454	G	2.1
22	DA	539	G	2.1
22	DA	1072	C	2.1
22	DA	1468	U	2.1
3	CC	140	ASN	2.1
32	DK	82	ASN	2.1
7	CG	76	LYS	2.1
28	DG	42	GLU	2.1
30	BI	52	GLY	2.1
9	CI	17	ALA	2.1
24	DC	250	VAL	2.1
28	DG	11	VAL	2.1
28	DG	97	ALA	2.1
32	DK	52	VAL	2.1
1	AA	992	U	2.0
1	CA	632	U	2.0
7	AG	52	GLN	2.0
13	AM	85	CYS	2.0
32	DK	91	SER	2.0
22	DA	353	C	2.0
33	DL	57	LEU	2.0
19	AS	17	LYS	2.0
22	DA	1091	G	2.0

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Mol	Chain	Res	Type	RSRZ
22	DA	1983	G	2.0
41	BT	68	LYS	2.0
28	DG	94	TYR	2.0
30	DI	108	GLU	2.0
2	CB	55	ALA	2.0
3	CC	128	VAL	2.0
10	AJ	98	VAL	2.0
27	DF	89	VAL	2.0
3	CC	149	ILE	2.0
26	DE	175	ILE	2.0
7	CG	109	ARG	2.0
41	DT	91	GLN	2.0
14	CN	46	LEU	2.0
22	DA	354	A	2.0
33	DL	10	GLU	2.0
6	AF	96	VAL	2.0
7	AG	73	VAL	2.0
25	DD	180	VAL	2.0
51	D3	37	ALA	2.0
53	B5	63	VAL	2.0
1	CA	1034	G	2.0
13	CM	4	ILE	2.0
22	BA	277	G	2.0
43	DV	44	HIS	2.0
50	B2	46	LYS	2.0
21	AU	24	GLU	2.0
24	DC	234	GLY	2.0
28	DG	108	GLY	2.0
1	CA	461	A	2.0
1	CA	979	C	2.0
22	DA	876	C	2.0
22	DA	1571	A	2.0
25	DD	162	ALA	2.0
29	BH	81	ALA	2.0
30	DI	134	ARG	2.0
45	DX	50	ARG	2.0
46	BY	63	ALA	2.0
1	CA	1139	G	2.0
3	AC	155	GLY	2.0
10	CJ	51	VAL	2.0
1	AA	84	U	2.0
1	CA	1049	U	2.0

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Mol	Chain	Res	Type	RSRZ
7	CG	150	ALA	2.0
19	AS	21	LYS	2.0
22	DA	646	U	2.0
42	BU	55	PRO	2.0
1	CA	1273	C	2.0
22	DA	1043	C	2.0
7	AG	8	GLY	2.0
3	AC	22	TRP	2.0
13	AM	30	SER	2.0
28	DG	69	ARG	2.0
41	DT	68	LYS	2.0
48	D0	29	SER	2.0
16	CP	81	ALA	2.0
24	DC	44	ASN	2.0
28	DG	34	THR	2.0
40	DS	1	MET	2.0
27	DF	173	PHE	2.0

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

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

6.3 Carbohydrates [i](#)

There are no carbohydrates in this entry.

6.4 Ligands [i](#)

In the following table, the Atoms column lists the number of modelled atoms in the group and the number defined in the chemical component dictionary. LLDF column lists the quality of electron density of the group with respect to its neighbouring residues in protein, DNA or RNA chains. The B-factors column lists the minimum, median, 95th percentile and maximum values of B factors of atoms in the group. The column labelled 'Q< 0.9' lists the number of atoms with occupancy less than 0.9.

Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
54	MG	BA	3174	1/1	0.87	0.82	33.52	25,25,25,25	0
54	MG	DA	3135	1/1	0.78	0.62	29.36	112,112,112,112	0
54	MG	DA	3155	1/1	0.71	0.74	27.39	51,51,51,51	0
54	MG	BA	3057	1/1	0.83	0.48	27.27	40,40,40,40	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
54	MG	AM	201	1/1	0.53	1.29	24.26	63,63,63,63	0
54	MG	BA	3146	1/1	0.93	0.36	22.91	32,32,32,32	0
54	MG	DA	3057	1/1	0.70	0.85	22.11	99,99,99,99	0
54	MG	AA	1670	1/1	0.75	0.43	20.90	49,49,49,49	0
54	MG	BA	3040	1/1	0.80	0.47	20.37	1,1,1,1	0
54	MG	BA	3167	1/1	0.73	0.41	18.14	25,25,25,25	0
54	MG	BA	3083	1/1	0.91	0.33	17.10	44,44,44,44	0
54	MG	AA	1662	1/1	0.65	0.75	13.87	76,76,76,76	0
54	MG	AA	1622	1/1	0.96	0.26	13.70	21,21,21,21	0
54	MG	BA	3160	1/1	0.80	0.35	13.26	7,7,7,7	0
54	MG	BA	3152	1/1	0.96	0.45	13.09	1,1,1,1	0
54	MG	BA	3162	1/1	0.94	0.39	12.66	30,30,30,30	0
54	MG	DA	3027	1/1	0.84	0.56	11.10	106,106,106,106	0
54	MG	BA	3104	1/1	0.99	0.29	9.51	1,1,1,1	0
54	MG	BA	3185	1/1	0.86	0.28	8.76	17,17,17,17	0
54	MG	AA	1630	1/1	0.90	0.31	8.21	78,78,78,78	0
54	MG	BA	3150	1/1	0.83	0.34	7.63	23,23,23,23	0
54	MG	BA	3177	1/1	0.93	0.22	5.57	24,24,24,24	0
54	MG	AA	1617	1/1	0.27	0.85	5.24	124,124,124,124	0
54	MG	DA	3147	1/1	0.81	0.41	5.18	49,49,49,49	0
54	MG	AA	1635	1/1	0.86	0.26	4.95	73,73,73,73	0
54	MG	BA	3131	1/1	0.70	0.29	4.38	49,49,49,49	0
54	MG	DA	3071	1/1	0.52	0.33	4.33	99,99,99,99	0
54	MG	BA	3158	1/1	0.95	0.26	4.27	15,15,15,15	0
54	MG	DA	3104	1/1	0.86	0.46	4.12	97,97,97,97	0
54	MG	BA	3130	1/1	0.94	0.28	3.99	7,7,7,7	0
54	MG	CA	1614	1/1	0.80	0.27	3.66	60,60,60,60	0
54	MG	BA	3136	1/1	0.93	0.26	3.16	50,50,50,50	0
54	MG	DA	3152	1/1	0.76	0.28	3.08	59,59,59,59	0
54	MG	BA	3124	1/1	0.96	0.33	2.71	44,44,44,44	0
54	MG	DA	3048	1/1	0.82	0.36	2.65	109,109,109,109	0
54	MG	DA	3115	1/1	0.88	0.29	2.54	79,79,79,79	0
54	MG	DA	3109	1/1	0.96	0.36	2.41	64,64,64,64	0
54	MG	DA	3127	1/1	0.93	0.38	2.34	91,91,91,91	0
54	MG	BA	3169	1/1	0.88	0.18	1.89	33,33,33,33	0
54	MG	BA	3013	1/1	0.97	0.22	1.77	1,1,1,1	0
54	MG	BA	3154	1/1	0.90	0.25	1.67	18,18,18,18	0
54	MG	BA	3053	1/1	0.99	0.23	1.46	1,1,1,1	0
54	MG	CA	1641	1/1	0.87	0.56	1.34	49,49,49,49	0
54	MG	BA	3105	1/1	0.87	0.22	1.23	10,10,10,10	0
54	MG	BA	3151	1/1	0.92	0.24	1.00	7,7,7,7	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
54	MG	BA	3012	1/1	0.95	0.24	0.96	1,1,1,1	0
54	MG	DA	3008	1/1	0.76	0.27	0.89	96,96,96,96	0
54	MG	BA	3108	1/1	0.93	0.21	0.88	1,1,1,1	0
54	MG	DA	3108	1/1	0.96	0.25	0.55	22,22,22,22	0
54	MG	BA	3109	1/1	0.96	0.21	0.37	1,1,1,1	0
54	MG	DA	3114	1/1	0.42	0.42	0.37	126,126,126,126	0
54	MG	DA	3005	1/1	0.79	0.20	0.33	113,113,113,113	0
54	MG	DL	202	1/1	0.85	0.39	0.24	116,116,116,116	0
54	MG	CA	1611	1/1	0.97	0.12	0.06	42,42,42,42	0
54	MG	BA	3132	1/1	0.96	0.37	0.01	45,45,45,45	0
54	MG	DA	3024	1/1	0.45	0.25	-0.01	77,77,77,77	0
54	MG	CA	1638	1/1	0.96	0.17	-0.21	28,28,28,28	0
54	MG	DA	3131	1/1	0.90	0.30	-0.30	79,79,79,79	0
54	MG	BA	3017	1/1	0.98	0.20	-0.31	1,1,1,1	0
54	MG	CA	1621	1/1	0.90	0.17	-0.32	75,75,75,75	0
54	MG	BA	3049	1/1	0.98	0.20	-0.34	3,3,3,3	0
54	MG	CA	1633	1/1	0.61	0.29	-0.36	134,134,134,134	0
54	MG	DA	3070	1/1	0.68	0.20	-0.36	105,105,105,105	0
54	MG	DA	3101	1/1	0.94	0.20	-0.41	85,85,85,85	0
54	MG	DA	3013	1/1	0.65	0.22	-0.47	72,72,72,72	0
54	MG	BB	201	1/1	0.95	0.15	-0.52	32,32,32,32	0
54	MG	DA	3012	1/1	0.95	0.22	-0.55	46,46,46,46	0
54	MG	BA	3079	1/1	0.90	0.18	-0.59	29,29,29,29	0
54	MG	DA	3023	1/1	0.92	0.21	-0.63	57,57,57,57	0
55	ZN	B4	101	1/1	1.00	0.17	-0.77	28,28,28,28	0
54	MG	AA	1629	1/1	0.93	0.20	-0.95	73,73,73,73	0
54	MG	DA	3077	1/1	0.84	0.14	-1.01	114,114,114,114	0
54	MG	DA	3107	1/1	0.88	0.18	-1.04	70,70,70,70	0
54	MG	DA	3018	1/1	0.78	0.11	-1.05	88,88,88,88	0
54	MG	BA	3116	1/1	0.96	0.19	-1.05	1,1,1,1	0
54	MG	BA	3097	1/1	0.99	0.17	-1.13	1,1,1,1	0
54	MG	AA	1607	1/1	0.94	0.06	-1.18	48,48,48,48	0
56	NEG	CA	1657	17/17	0.87	0.13	-1.27	62,88,103,104	0
54	MG	BA	3008	1/1	0.95	0.16	-1.30	3,3,3,3	0
54	MG	DA	3017	1/1	0.94	0.23	-1.35	71,71,71,71	0
54	MG	BA	3023	1/1	0.98	0.18	-1.37	1,1,1,1	0
55	ZN	D4	101	1/1	0.97	0.09	-1.38	95,95,95,95	0
54	MG	BA	3022	1/1	0.96	0.14	-1.55	1,1,1,1	0
54	MG	DA	3129	1/1	0.93	0.17	-1.55	93,93,93,93	0
54	MG	DB	202	1/1	0.90	0.08	-1.58	89,89,89,89	0
54	MG	CA	1628	1/1	0.40	0.14	-1.64	137,137,137,137	0
54	MG	BA	3107	1/1	0.99	0.16	-1.73	3,3,3,3	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
54	MG	BA	3039	1/1	0.97	0.19	-1.73	1,1,1,1	0
54	MG	AA	1641	1/1	0.99	0.13	-1.76	12,12,12,12	0
54	MG	DA	3046	1/1	0.90	0.13	-1.79	80,80,80,80	0
54	MG	AA	1616	1/1	0.95	0.08	-1.92	94,94,94,94	0
54	MG	AA	1618	1/1	0.87	0.10	-1.93	89,89,89,89	0
54	MG	BA	3164	1/1	0.92	0.15	-1.98	13,13,13,13	0
54	MG	BA	3113	1/1	0.99	0.15	-2.04	4,4,4,4	0
54	MG	BA	3032	1/1	0.95	0.18	-2.08	4,4,4,4	0
54	MG	DA	3003	1/1	0.95	0.10	-2.20	83,83,83,83	0
54	MG	DA	3042	1/1	0.94	0.11	-2.32	67,67,67,67	0
54	MG	CA	1656	1/1	0.90	0.08	-2.33	88,88,88,88	0
54	MG	DA	3105	1/1	0.75	0.18	-2.33	77,77,77,77	0
54	MG	DA	3097	1/1	0.90	0.09	-2.42	67,67,67,67	0
54	MG	BA	3134	1/1	0.95	0.18	-2.48	3,3,3,3	0
54	MG	DA	3078	1/1	0.80	0.09	-2.50	103,103,103,103	0
54	MG	BA	3050	1/1	0.99	0.12	-2.51	7,7,7,7	0
54	MG	DA	3053	1/1	0.85	0.19	-2.52	49,49,49,49	0
54	MG	BA	3009	1/1	0.95	0.14	-2.57	1,1,1,1	0
54	MG	AA	1642	1/1	0.99	0.11	-2.63	23,23,23,23	0
54	MG	BA	3062	1/1	0.98	0.12	-2.68	4,4,4,4	0
54	MG	BA	3018	1/1	0.98	0.10	-2.69	14,14,14,14	0
54	MG	CA	1630	1/1	0.80	0.06	-2.73	79,79,79,79	0
54	MG	DA	3021	1/1	0.94	0.09	-2.75	69,69,69,69	0
54	MG	DA	3073	1/1	0.93	0.15	-2.79	69,69,69,69	0
54	MG	DA	3095	1/1	0.85	0.08	-2.81	77,77,77,77	0
54	MG	BA	3114	1/1	0.86	0.14	-2.90	14,14,14,14	0
54	MG	BB	202	1/1	0.98	0.08	-3.08	10,10,10,10	0
54	MG	DA	3022	1/1	0.95	0.09	-3.17	74,74,74,74	0
54	MG	CA	1616	1/1	0.87	0.11	-3.19	43,43,43,43	0
54	MG	DA	3050	1/1	0.93	0.09	-3.20	41,41,41,41	0
54	MG	CA	1606	1/1	0.97	0.11	-3.23	61,61,61,61	0
54	MG	BA	3101	1/1	0.88	0.12	-3.26	14,14,14,14	0
54	MG	DA	3058	1/1	0.95	0.10	-3.46	64,64,64,64	0
54	MG	BA	3129	1/1	0.97	0.17	-3.59	1,1,1,1	0
54	MG	BA	3028	1/1	0.97	0.16	-3.65	1,1,1,1	0
54	MG	DA	3065	1/1	0.93	0.16	-3.65	44,44,44,44	0
54	MG	DA	3068	1/1	0.89	0.08	-3.69	81,81,81,81	0
54	MG	CA	1609	1/1	0.93	0.10	-3.78	56,56,56,56	0
54	MG	AA	1604	1/1	0.84	0.09	-3.87	81,81,81,81	0
54	MG	DA	3133	1/1	0.88	0.14	-4.03	78,78,78,78	0
54	MG	DA	3026	1/1	0.33	0.14	-4.07	93,93,93,93	0
54	MG	DA	3062	1/1	0.94	0.13	-4.14	67,67,67,67	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
54	MG	CA	1603	1/1	0.93	0.06	-4.22	40,40,40,40	0
54	MG	DA	3028	1/1	0.74	0.14	-4.23	83,83,83,83	0
54	MG	BA	3027	1/1	0.95	0.14	-4.36	10,10,10,10	0
54	MG	CA	1601	1/1	0.94	0.12	-4.47	45,45,45,45	0
54	MG	AA	1633	1/1	0.95	0.10	-4.49	25,25,25,25	0
54	MG	BA	3096	1/1	0.98	0.09	-4.57	2,2,2,2	0
54	MG	BA	3066	1/1	0.96	0.12	-4.69	3,3,3,3	0
54	MG	BA	3112	1/1	0.91	0.10	-4.70	25,25,25,25	0
54	MG	BA	3002	1/1	0.95	0.09	-4.82	11,11,11,11	0
54	MG	CA	1618	1/1	0.98	0.10	-4.94	35,35,35,35	0
54	MG	DA	3038	1/1	0.97	0.15	-4.94	68,68,68,68	0
54	MG	DA	3063	1/1	0.97	0.14	-5.02	64,64,64,64	0
54	MG	AA	1613	1/1	0.95	0.08	-5.03	13,13,13,13	0
54	MG	BA	3073	1/1	0.82	0.12	-5.08	16,16,16,16	0
54	MG	BA	3065	1/1	0.97	0.13	-5.34	1,1,1,1	0
54	MG	AA	1606	1/1	0.97	0.08	-5.44	42,42,42,42	0
54	MG	DB	201	1/1	0.73	0.09	-5.54	118,118,118,118	0
54	MG	BA	3024	1/1	0.98	0.10	-5.55	13,13,13,13	0
54	MG	DA	3079	1/1	0.95	0.11	-5.55	91,91,91,91	0
54	MG	BA	3021	1/1	0.94	0.13	-5.60	1,1,1,1	0
54	MG	BA	3093	1/1	0.98	0.12	-5.74	11,11,11,11	0
54	MG	BA	3058	1/1	0.98	0.09	-6.12	14,14,14,14	0
54	MG	AA	1625	1/1	0.83	0.08	-6.75	47,47,47,47	0
54	MG	DA	3049	1/1	0.88	0.10	-6.98	61,61,61,61	0
54	MG	BA	3071	1/1	0.98	0.08	-7.08	13,13,13,13	0
54	MG	BA	3005	1/1	0.97	0.06	-7.20	44,44,44,44	0
54	MG	CA	1625	1/1	0.96	0.04	-9.66	44,44,44,44	0
54	MG	AA	1609	1/1	0.97	0.05	-10.06	23,23,23,23	0
54	MG	BA	3070	1/1	0.96	0.11	-14.80	6,6,6,6	0
54	MG	BA	3118	1/1	0.94	0.07	-33.28	14,14,14,14	0
54	MG	DA	3092	1/1	-0.22	0.48	-	123,123,123,123	0
54	MG	DA	3002	1/1	0.83	0.48	-	94,94,94,94	0
54	MG	DA	3146	1/1	0.23	0.19	-	68,68,68,68	0
54	MG	DA	3145	1/1	0.70	1.24	-	60,60,60,60	0
54	MG	CA	1626	1/1	0.54	0.94	-	125,125,125,125	0
54	MG	BA	3085	1/1	0.92	0.13	-	9,9,9,9	0
54	MG	BA	3100	1/1	0.97	0.17	-	4,4,4,4	0
54	MG	BA	3043	1/1	0.99	0.08	-	11,11,11,11	0
54	MG	DA	3031	1/1	0.85	0.52	-	95,95,95,95	0
54	MG	BA	3120	1/1	0.94	0.14	-	10,10,10,10	0
54	MG	DA	3084	1/1	0.76	0.42	-	104,104,104,104	0
54	MG	BA	3123	1/1	0.97	0.14	-	6,6,6,6	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
54	MG	DA	3076	1/1	0.68	0.52	-	114,114,114,114	0
54	MG	BA	3077	1/1	0.94	0.12	-	16,16,16,16	0
54	MG	DA	3099	1/1	0.78	0.13	-	69,69,69,69	0
54	MG	CA	1645	1/1	0.86	0.38	-	52,52,52,52	0
54	MG	BA	3182	1/1	0.94	0.33	-	25,25,25,25	0
54	MG	BA	3175	1/1	0.93	0.23	-	38,38,38,38	0
54	MG	DA	3032	1/1	0.97	0.06	-	67,67,67,67	0
54	MG	BA	3142	1/1	0.95	0.61	-	17,17,17,17	0
54	MG	DA	3067	1/1	0.84	0.17	-	70,70,70,70	0
54	MG	AA	1624	1/1	0.74	0.12	-	36,36,36,36	0
54	MG	AA	1651	1/1	0.90	0.80	-	52,52,52,52	0
54	MG	DA	3139	1/1	0.90	0.58	-	49,49,49,49	0
54	MG	BA	3048	1/1	0.97	0.15	-	10,10,10,10	0
54	MG	DA	3037	1/1	0.86	0.25	-	66,66,66,66	0
54	MG	BA	3155	1/1	0.89	0.26	-	21,21,21,21	0
54	MG	BA	3060	1/1	0.63	0.71	-	72,72,72,72	0
54	MG	BA	3051	1/1	0.98	0.15	-	2,2,2,2	0
54	MG	DA	3086	1/1	0.91	0.20	-	81,81,81,81	0
54	MG	CA	1605	1/1	0.72	0.20	-	110,110,110,110	0
54	MG	DA	3132	1/1	0.27	1.08	-	123,123,123,123	0
54	MG	DA	3125	1/1	0.94	0.27	-	83,83,83,83	0
54	MG	CT	101	1/1	0.94	0.18	-	91,91,91,91	0
54	MG	DA	3051	1/1	0.94	0.19	-	51,51,51,51	0
54	MG	BA	3166	1/1	0.88	0.45	-	36,36,36,36	0
54	MG	BA	3194	1/1	0.91	0.16	-	42,42,42,42	0
54	MG	DA	3025	1/1	0.67	1.85	-	120,120,120,120	0
54	MG	DA	3141	1/1	0.97	0.56	-	45,45,45,45	0
54	MG	DA	3035	1/1	0.95	0.27	-	54,54,54,54	0
54	MG	BA	3138	1/1	0.96	0.54	-	1,1,1,1	0
54	MG	BQ	201	1/1	0.88	0.33	-	17,17,17,17	0
54	MG	BA	3106	1/1	0.98	0.18	-	1,1,1,1	0
54	MG	BA	3135	1/1	0.95	0.15	-	23,23,23,23	0
54	MG	DA	3102	1/1	0.88	0.15	-	77,77,77,77	0
54	MG	DA	3015	1/1	0.88	0.33	-	90,90,90,90	0
54	MG	DA	3019	1/1	0.96	0.32	-	38,38,38,38	0
54	MG	BA	3064	1/1	0.96	0.13	-	5,5,5,5	0
54	MG	AA	1660	1/1	0.93	0.37	-	72,72,72,72	0
54	MG	CA	1639	1/1	0.88	0.15	-	59,59,59,59	0
54	MG	CA	1653	1/1	0.70	0.39	-	54,54,54,54	0
54	MG	CA	1607	1/1	0.85	0.20	-	96,96,96,96	0
54	MG	AA	1611	1/1	0.93	0.10	-	32,32,32,32	0
54	MG	DA	3007	1/1	0.83	0.29	-	114,114,114,114	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
54	MG	DA	3137	1/1	0.88	0.49	-	38,38,38,38	0
54	MG	BA	3192	1/1	0.96	0.17	-	30,30,30,30	0
54	MG	DA	3163	1/1	0.74	0.25	-	45,45,45,45	0
54	MG	DA	3004	1/1	0.48	0.53	-	112,112,112,112	0
54	MG	BA	3178	1/1	0.94	0.77	-	41,41,41,41	0
54	MG	DA	3153	1/1	0.80	0.42	-	58,58,58,58	0
54	MG	BA	3190	1/1	0.89	0.27	-	46,46,46,46	0
54	MG	BA	3171	1/1	0.95	0.23	-	29,29,29,29	0
54	MG	DA	3009	1/1	0.84	0.07	-	91,91,91,91	0
54	MG	BA	3006	1/1	0.98	0.14	-	29,29,29,29	0
54	MG	BA	3054	1/1	0.97	0.14	-	4,4,4,4	0
54	MG	BA	3176	1/1	0.70	0.45	-	37,37,37,37	0
54	MG	AA	1627	1/1	0.93	0.47	-	68,68,68,68	0
54	MG	DA	3128	1/1	0.89	0.14	-	60,60,60,60	0
54	MG	BA	3168	1/1	0.69	0.74	-	43,43,43,43	0
54	MG	BA	3059	1/1	0.91	0.41	-	27,27,27,27	0
54	MG	AA	1652	1/1	0.79	0.21	-	63,63,63,63	0
54	MG	DA	3120	1/1	0.72	0.27	-	82,82,82,82	0
54	MG	DA	3119	1/1	0.90	0.24	-	83,83,83,83	0
54	MG	DA	3093	1/1	0.77	0.23	-	119,119,119,119	0
54	MG	BA	3010	1/1	0.97	0.13	-	1,1,1,1	0
54	MG	CA	1634	1/1	0.65	0.59	-	143,143,143,143	0
54	MG	BA	3041	1/1	0.99	0.16	-	3,3,3,3	0
54	MG	AA	1638	1/1	0.82	0.15	-	88,88,88,88	0
54	MG	CA	1624	1/1	0.99	0.28	-	4,4,4,4	0
54	MG	BA	3125	1/1	0.97	0.20	-	4,4,4,4	0
54	MG	DA	3110	1/1	0.93	0.24	-	89,89,89,89	0
54	MG	CA	1651	1/1	0.16	0.46	-	70,70,70,70	0
54	MG	BA	3094	1/1	0.97	0.07	-	19,19,19,19	0
54	MG	DA	3144	1/1	0.08	0.33	-	96,96,96,96	0
54	MG	CA	1627	1/1	0.69	0.07	-	116,116,116,116	0
54	MG	DA	3055	1/1	0.43	0.60	-	109,109,109,109	0
54	MG	CA	1647	1/1	0.95	0.56	-	72,72,72,72	0
54	MG	BA	3188	1/1	0.88	0.20	-	5,5,5,5	0
54	MG	BA	3075	1/1	0.99	0.17	-	3,3,3,3	0
54	MG	DA	3162	1/1	0.67	0.49	-	66,66,66,66	0
54	MG	DA	3061	1/1	0.76	1.24	-	91,91,91,91	0
54	MG	DA	3117	1/1	0.93	0.21	-	79,79,79,79	0
54	MG	BA	3143	1/1	0.95	0.48	-	18,18,18,18	0
54	MG	DA	3052	1/1	0.88	0.09	-	50,50,50,50	0
54	MG	AA	1665	1/1	0.89	0.57	-	61,61,61,61	0
54	MG	BA	3163	1/1	0.86	0.99	-	36,36,36,36	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
54	MG	AA	1655	1/1	0.96	0.41	-	55,55,55,55	0
54	MG	DA	3098	1/1	0.70	0.21	-	89,89,89,89	0
54	MG	CA	1610	1/1	0.96	0.04	-	57,57,57,57	0
54	MG	BA	3137	1/1	0.97	0.59	-	11,11,11,11	0
54	MG	DA	3143	1/1	0.85	0.45	-	63,63,63,63	0
54	MG	DA	3157	1/1	0.86	0.37	-	61,61,61,61	0
54	MG	BA	3126	1/1	0.98	0.15	-	3,3,3,3	0
54	MG	BA	3180	1/1	0.92	0.28	-	45,45,45,45	0
54	MG	DA	3054	1/1	0.89	0.14	-	64,64,64,64	0
54	MG	AA	1634	1/1	0.85	0.12	-	58,58,58,58	0
54	MG	CA	1632	1/1	0.81	0.16	-	85,85,85,85	0
54	MG	BA	3026	1/1	0.99	0.09	-	2,2,2,2	0
54	MG	DA	3151	1/1	0.94	0.27	-	42,42,42,42	0
54	MG	DA	3122	1/1	0.95	0.10	-	51,51,51,51	0
54	MG	DA	3124	1/1	0.87	0.26	-	55,55,55,55	0
54	MG	DA	3087	1/1	0.76	0.10	-	84,84,84,84	0
54	MG	BA	3148	1/1	0.85	0.62	-	44,44,44,44	0
54	MG	AA	1639	1/1	0.97	0.12	-	60,60,60,60	0
54	MG	BA	3153	1/1	0.89	0.57	-	33,33,33,33	0
54	MG	BA	3052	1/1	0.97	0.08	-	1,1,1,1	0
54	MG	AA	1614	1/1	0.73	0.65	-	97,97,97,97	0
54	MG	DA	3118	1/1	0.68	0.61	-	113,113,113,113	0
54	MG	AA	1632	1/1	0.97	0.09	-	60,60,60,60	0
54	MG	BA	3007	1/1	0.93	0.06	-	29,29,29,29	0
54	MG	BA	3016	1/1	0.96	0.12	-	13,13,13,13	0
54	MG	BA	3128	1/1	0.99	0.41	-	1,1,1,1	0
54	MG	BA	3172	1/1	0.90	0.19	-	29,29,29,29	0
54	MG	BA	3031	1/1	0.98	0.04	-	11,11,11,11	0
54	MG	CA	1623	1/1	0.97	0.06	-	48,48,48,48	0
54	MG	CA	1642	1/1	0.83	0.28	-	53,53,53,53	0
54	MG	DA	3123	1/1	0.91	0.23	-	88,88,88,88	0
54	MG	DA	3047	1/1	0.52	0.16	-	132,132,132,132	0
54	MG	DA	3020	1/1	0.90	0.74	-	76,76,76,76	0
54	MG	DA	3116	1/1	0.78	0.19	-	106,106,106,106	0
54	MG	DA	3161	1/1	0.85	0.36	-	65,65,65,65	0
54	MG	BB	203	1/1	0.98	0.08	-	12,12,12,12	0
54	MG	DA	3148	1/1	0.90	0.25	-	53,53,53,53	0
54	MG	BA	3067	1/1	0.91	0.16	-	2,2,2,2	0
54	MG	CA	1635	1/1	0.89	0.76	-	99,99,99,99	0
54	MG	DA	3033	1/1	0.89	0.20	-	77,77,77,77	0
54	MG	BA	3089	1/1	0.97	0.07	-	8,8,8,8	0
54	MG	DA	3016	1/1	0.76	0.18	-	103,103,103,103	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
54	MG	DA	3130	1/1	0.65	1.52	-	108,108,108,108	0
54	MG	BA	3183	1/1	0.98	0.42	-	35,35,35,35	0
54	MG	DA	3111	1/1	0.89	0.12	-	71,71,71,71	0
54	MG	AA	1666	1/1	0.69	0.37	-	55,55,55,55	0
54	MG	DA	3075	1/1	0.97	0.28	-	78,78,78,78	0
54	MG	CA	1637	1/1	0.97	0.12	-	35,35,35,35	0
54	MG	AA	1663	1/1	0.84	0.66	-	36,36,36,36	0
54	MG	DA	3001	1/1	0.80	0.12	-	93,93,93,93	0
54	MG	CA	1602	1/1	0.89	0.11	-	93,93,93,93	0
54	MG	BA	3173	1/1	0.89	0.24	-	40,40,40,40	0
54	MG	DA	3030	1/1	0.88	0.18	-	71,71,71,71	0
54	MG	AA	1644	1/1	0.62	0.60	-	60,60,60,60	0
54	MG	BA	3191	1/1	0.82	0.30	-	17,17,17,17	0
54	MG	AA	1620	1/1	0.79	0.13	-	100,100,100,100	0
54	MG	AA	1658	1/1	0.52	0.51	-	71,71,71,71	0
54	MG	CA	1620	1/1	0.79	0.10	-	81,81,81,81	0
54	MG	BA	3044	1/1	0.99	0.34	-	1,1,1,1	0
54	MG	CA	1608	1/1	0.81	0.17	-	86,86,86,86	0
54	MG	CA	1612	1/1	0.93	0.19	-	20,20,20,20	0
54	MG	BA	3086	1/1	0.97	0.17	-	2,2,2,2	0
54	MG	AA	1669	1/1	0.76	0.72	-	83,83,83,83	0
54	MG	DA	3080	1/1	0.94	0.17	-	74,74,74,74	0
54	MG	CA	1646	1/1	0.91	0.80	-	21,21,21,21	0
54	MG	AA	1628	1/1	0.69	0.11	-	84,84,84,84	0
54	MG	AA	1657	1/1	0.73	0.44	-	42,42,42,42	0
54	MG	AA	1636	1/1	0.96	0.16	-	46,46,46,46	0
54	MG	BA	3119	1/1	0.95	0.29	-	40,40,40,40	0
54	MG	AA	1612	1/1	0.96	0.10	-	42,42,42,42	0
54	MG	DA	3041	1/1	0.74	0.15	-	96,96,96,96	0
54	MG	CA	1654	1/1	0.69	0.67	-	70,70,70,70	0
54	MG	BA	3063	1/1	0.95	0.11	-	6,6,6,6	0
54	MG	AA	1649	1/1	0.70	0.66	-	52,52,52,52	0
54	MG	AA	1631	1/1	0.95	0.08	-	32,32,32,32	0
54	MG	DA	3064	1/1	0.90	0.12	-	61,61,61,61	0
54	MG	CA	1631	1/1	0.90	0.18	-	85,85,85,85	0
54	MG	DA	3150	1/1	0.98	0.16	-	86,86,86,86	0
54	MG	BA	3038	1/1	0.96	0.16	-	1,1,1,1	0
54	MG	BA	3082	1/1	0.91	0.07	-	8,8,8,8	0
54	MG	DA	3156	1/1	0.90	0.16	-	72,72,72,72	0
54	MG	CA	1619	1/1	0.64	0.10	-	85,85,85,85	0
54	MG	DA	3066	1/1	0.94	0.15	-	58,58,58,58	0
54	MG	CA	1650	1/1	0.76	0.41	-	85,85,85,85	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
54	MG	CA	1649	1/1	0.88	0.63	-	50,50,50,50	0
54	MG	BA	3029	1/1	0.95	0.08	-	20,20,20,20	0
54	MG	BA	3020	1/1	0.97	0.05	-	1,1,1,1	0
54	MG	DA	3072	1/1	0.77	0.27	-	78,78,78,78	0
54	MG	BA	3187	1/1	0.95	0.22	-	41,41,41,41	0
54	MG	BA	3078	1/1	0.96	0.07	-	48,48,48,48	0
54	MG	AA	1615	1/1	0.94	0.23	-	52,52,52,52	0
54	MG	BA	3004	1/1	0.94	0.20	-	55,55,55,55	0
54	MG	BA	3133	1/1	0.77	0.55	-	63,63,63,63	0
54	MG	DA	3164	1/1	0.53	0.56	-	60,60,60,60	0
54	MG	AA	1640	1/1	0.92	0.06	-	39,39,39,39	0
54	MG	AA	1602	1/1	0.96	0.10	-	37,37,37,37	0
54	MG	AA	1647	1/1	0.96	0.19	-	51,51,51,51	0
54	MG	BA	3042	1/1	0.93	0.14	-	1,1,1,1	0
54	MG	CA	1615	1/1	0.96	0.17	-	39,39,39,39	0
54	MG	BA	3145	1/1	0.94	0.56	-	35,35,35,35	0
54	MG	BA	3088	1/1	0.95	0.12	-	25,25,25,25	0
54	MG	DA	3074	1/1	0.72	0.18	-	96,96,96,96	0
54	MG	DA	3056	1/1	0.58	0.63	-	103,103,103,103	0
54	MG	AA	1661	1/1	0.90	0.55	-	40,40,40,40	0
54	MG	DA	3059	1/1	0.56	0.34	-	87,87,87,87	0
54	MG	DA	3121	1/1	0.94	0.19	-	62,62,62,62	0
54	MG	DA	3089	1/1	0.61	0.15	-	106,106,106,106	0
54	MG	BA	3055	1/1	0.85	0.64	-	89,89,89,89	0
54	MG	DA	3126	1/1	0.96	0.13	-	58,58,58,58	0
54	MG	DA	3043	1/1	0.64	0.31	-	104,104,104,104	0
54	MG	CA	1617	1/1	0.94	0.15	-	38,38,38,38	0
54	MG	BA	3011	1/1	0.89	0.07	-	16,16,16,16	0
54	MG	AA	1608	1/1	0.97	0.17	-	26,26,26,26	0
54	MG	CA	1604	1/1	0.87	0.07	-	102,102,102,102	0
54	MG	DA	3160	1/1	0.98	0.30	-	86,86,86,86	0
54	MG	BA	3080	1/1	0.96	0.15	-	24,24,24,24	0
54	MG	DA	3036	1/1	0.66	0.14	-	114,114,114,114	0
54	MG	CA	1622	1/1	0.94	0.60	-	84,84,84,84	0
54	MG	BA	3141	1/1	0.98	0.65	-	2,2,2,2	0
54	MG	BA	3159	1/1	0.96	0.32	-	18,18,18,18	0
54	MG	AA	1648	1/1	0.88	0.32	-	34,34,34,34	0
54	MG	CA	1644	1/1	-0.03	1.58	-	86,86,86,86	0
54	MG	BA	3147	1/1	0.96	0.39	-	13,13,13,13	0
54	MG	DA	3158	1/1	0.77	0.33	-	52,52,52,52	0
54	MG	BA	3102	1/1	0.29	0.37	-	68,68,68,68	0
54	MG	BA	3061	1/1	0.78	0.33	-	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
54	MG	DA	3140	1/1	0.78	0.48	-	51,51,51,51	0
54	MG	AA	1626	1/1	0.98	0.25	-	1,1,1,1	0
54	MG	BA	3090	1/1	0.99	0.18	-	13,13,13,13	0
54	MG	AA	1654	1/1	0.95	0.41	-	46,46,46,46	0
54	MG	BA	3193	1/1	0.90	0.29	-	46,46,46,46	0
54	MG	D2	101	1/1	0.81	0.18	-	111,111,111,111	0
54	MG	BA	3184	1/1	0.89	0.19	-	24,24,24,24	0
54	MG	DA	3159	1/1	0.57	0.45	-	71,71,71,71	0
54	MG	CA	1643	1/1	0.92	0.22	-	43,43,43,43	0
54	MG	DA	3044	1/1	0.97	0.30	-	51,51,51,51	0
54	MG	DA	3090	1/1	0.87	0.09	-	85,85,85,85	0
54	MG	DA	3094	1/1	0.76	0.38	-	99,99,99,99	0
54	MG	DA	3106	1/1	0.87	0.10	-	60,60,60,60	0
54	MG	DA	3138	1/1	0.92	0.42	-	41,41,41,41	0
54	MG	AA	1667	1/1	0.88	0.49	-	54,54,54,54	0
54	MG	BA	3186	1/1	0.88	0.35	-	29,29,29,29	0
54	MG	BA	3115	1/1	0.72	0.16	-	58,58,58,58	0
54	MG	AA	1601	1/1	0.85	0.08	-	59,59,59,59	0
54	MG	AA	1621	1/1	0.92	0.08	-	32,32,32,32	0
54	MG	CA	1636	1/1	0.77	0.18	-	93,93,93,93	0
54	MG	BA	3068	1/1	0.95	0.11	-	1,1,1,1	0
54	MG	DA	3014	1/1	0.90	0.30	-	86,86,86,86	0
54	MG	DA	3088	1/1	0.71	0.22	-	81,81,81,81	0
54	MG	DL	201	1/1	0.38	0.48	-	95,95,95,95	0
54	MG	BA	3170	1/1	0.86	0.25	-	44,44,44,44	0
54	MG	BA	3081	1/1	0.97	0.15	-	1,1,1,1	0
54	MG	AA	1619	1/1	0.88	0.24	-	57,57,57,57	0
54	MG	AA	1610	1/1	0.91	0.09	-	64,64,64,64	0
54	MG	AA	1653	1/1	0.93	0.38	-	27,27,27,27	0
54	MG	BA	3074	1/1	0.97	0.08	-	24,24,24,24	0
54	MG	BA	3033	1/1	0.97	0.23	-	1,1,1,1	0
54	MG	BA	3140	1/1	0.93	0.29	-	34,34,34,34	0
54	MG	DA	3100	1/1	0.76	0.17	-	74,74,74,74	0
54	MG	BA	3084	1/1	0.93	0.26	-	33,33,33,33	0
54	MG	BA	3015	1/1	0.65	0.31	-	62,62,62,62	0
54	MG	DA	3136	1/1	0.72	0.54	-	46,46,46,46	0
54	MG	AA	1645	1/1	0.95	0.89	-	52,52,52,52	0
54	MG	CA	1655	1/1	0.89	0.08	-	85,85,85,85	0
54	MG	AA	1659	1/1	0.73	0.45	-	68,68,68,68	0
54	MG	BA	3092	1/1	0.95	0.14	-	33,33,33,33	0
54	MG	BA	3014	1/1	0.94	0.12	-	8,8,8,8	0
54	MG	BA	3003	1/1	0.85	0.09	-	29,29,29,29	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
54	MG	BA	3046	1/1	0.96	0.20	-	5,5,5,5	0
54	MG	BA	3045	1/1	0.93	0.25	-	5,5,5,5	0
54	MG	BA	3122	1/1	0.99	0.24	-	1,1,1,1	0
54	MG	BA	3091	1/1	0.97	0.27	-	53,53,53,53	0
54	MG	DA	3039	1/1	0.89	0.18	-	96,96,96,96	0
54	MG	AA	1605	1/1	0.94	0.18	-	35,35,35,35	0
54	MG	AA	1637	1/1	0.98	0.10	-	13,13,13,13	0
54	MG	BA	3149	1/1	0.80	0.23	-	48,48,48,48	0
54	MG	CA	1648	1/1	0.76	0.43	-	47,47,47,47	0
54	MG	BA	3030	1/1	0.99	0.17	-	4,4,4,4	0
54	MG	BA	3025	1/1	0.92	0.34	-	38,38,38,38	0
54	MG	DA	3060	1/1	0.60	0.67	-	92,92,92,92	0
54	MG	DA	3040	1/1	0.07	0.64	-	94,94,94,94	0
54	MG	BA	3056	1/1	0.90	0.36	-	27,27,27,27	0
54	MG	AA	1643	1/1	0.96	0.71	-	7,7,7,7	0
54	MG	DB	203	1/1	0.93	0.07	-	106,106,106,106	0
54	MG	AA	1671	1/1	0.60	0.54	-	52,52,52,52	0
54	MG	BA	3098	1/1	0.96	0.36	-	41,41,41,41	0
54	MG	BA	3087	1/1	0.97	0.07	-	18,18,18,18	0
54	MG	DA	3045	1/1	0.95	0.11	-	74,74,74,74	0
54	MG	BA	3117	1/1	0.98	0.21	-	1,1,1,1	0
54	MG	BA	3144	1/1	0.98	0.46	-	34,34,34,34	0
54	MG	BA	3019	1/1	0.95	0.41	-	1,1,1,1	0
54	MG	BA	3157	1/1	0.95	0.23	-	26,26,26,26	0
54	MG	BA	3110	1/1	0.88	0.23	-	61,61,61,61	0
54	MG	AA	1656	1/1	0.99	0.15	-	46,46,46,46	0
54	MG	AA	1668	1/1	0.89	0.24	-	39,39,39,39	0
54	MG	AA	1603	1/1	0.98	0.04	-	30,30,30,30	0
54	MG	BA	3103	1/1	0.99	0.13	-	2,2,2,2	0
54	MG	BB	204	1/1	0.97	0.69	-	6,6,6,6	0
54	MG	BA	3037	1/1	0.99	0.25	-	1,1,1,1	0
54	MG	BA	3072	1/1	0.97	0.13	-	1,1,1,1	0
54	MG	DA	3081	1/1	0.69	0.12	-	75,75,75,75	0
54	MG	BA	3035	1/1	0.97	0.21	-	1,1,1,1	0
54	MG	BA	3181	1/1	0.96	0.49	-	33,33,33,33	0
54	MG	DA	3082	1/1	0.96	0.09	-	80,80,80,80	0
54	MG	BA	3069	1/1	0.92	0.10	-	72,72,72,72	0
54	MG	DA	3112	1/1	0.92	0.28	-	78,78,78,78	0
54	MG	AA	1623	1/1	0.93	0.21	-	60,60,60,60	0
54	MG	AA	1664	1/1	0.98	0.29	-	58,58,58,58	0
54	MG	AA	1646	1/1	0.71	0.61	-	49,49,49,49	0
54	MG	DA	3085	1/1	0.92	0.14	-	71,71,71,71	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
54	MG	CA	1652	1/1	0.88	0.45	-	73,73,73,73	0
54	MG	BA	3189	1/1	0.84	0.16	-	43,43,43,43	0
54	MG	DA	3113	1/1	0.90	0.13	-	54,54,54,54	0
54	MG	DA	3010	1/1	0.92	0.16	-	60,60,60,60	0
54	MG	DA	3029	1/1	0.80	0.18	-	83,83,83,83	0
54	MG	DA	3096	1/1	0.90	0.26	-	81,81,81,81	0
54	MG	DA	3154	1/1	0.83	0.36	-	63,63,63,63	0
54	MG	BA	3034	1/1	0.88	0.50	-	39,39,39,39	0
54	MG	DA	3083	1/1	0.57	0.28	-	116,116,116,116	0
54	MG	DA	3006	1/1	0.43	0.13	-	127,127,127,127	0
54	MG	DA	3034	1/1	0.93	0.12	-	90,90,90,90	0
54	MG	BA	3179	1/1	0.32	1.42	-	29,29,29,29	0
54	MG	BA	3036	1/1	0.87	0.52	-	20,20,20,20	0
54	MG	BA	3156	1/1	0.96	0.88	-	40,40,40,40	0
54	MG	BA	3139	1/1	0.91	0.50	-	1,1,1,1	0
54	MG	DA	3091	1/1	0.51	0.71	-	128,128,128,128	0
54	MG	DA	3103	1/1	0.81	0.10	-	85,85,85,85	0
54	MG	BA	3121	1/1	0.85	0.14	-	50,50,50,50	0
54	MG	DA	3069	1/1	0.77	0.06	-	121,121,121,121	0
54	MG	CA	1613	1/1	0.85	0.10	-	51,51,51,51	0
54	MG	BA	3047	1/1	0.90	0.13	-	43,43,43,43	0
54	MG	BA	3099	1/1	0.94	0.18	-	1,1,1,1	0
54	MG	DA	3134	1/1	0.71	0.30	-	112,112,112,112	0
54	MG	DA	3011	1/1	0.95	0.19	-	98,98,98,98	0
54	MG	BA	3095	1/1	0.98	0.07	-	6,6,6,6	0
54	MG	BA	3076	1/1	0.92	0.25	-	60,60,60,60	0
54	MG	DA	3142	1/1	0.95	0.44	-	53,53,53,53	0
54	MG	DQ	201	1/1	0.84	0.77	-	51,51,51,51	0
54	MG	BA	3111	1/1	0.96	0.10	-	40,40,40,40	0
54	MG	BA	3127	1/1	0.95	0.13	-	3,3,3,3	0
54	MG	DA	3149	1/1	0.83	0.38	-	77,77,77,77	0
54	MG	BA	3161	1/1	0.78	0.41	-	30,30,30,30	0
54	MG	AA	1650	1/1	0.81	0.79	-	49,49,49,49	0
54	MG	BA	3001	1/1	0.90	0.14	-	39,39,39,39	0
54	MG	BA	3165	1/1	0.89	0.30	-	29,29,29,29	0
54	MG	CA	1629	1/1	0.70	0.52	-	123,123,123,123	0
54	MG	CA	1640	1/1	0.88	0.35	-	29,29,29,29	0

6.5 Other polymers ⓘ

There are no such residues in this entry.