



Full wwPDB X-ray Structure Validation Report ⓘ

Oct 6, 2020 – 03:07 AM EDT

PDB ID : 6NUO
Title : Modified tRNA(Pro) bound to Thermus thermophilus 70S (cognate)
Authors : Hoffer, E.D.; Subaramanian, S.; Hong, S.; Maehigashi, T.; Dunham, C.M.
Deposited on : 2019-02-01
Resolution : 3.20 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

<https://www.wwpdb.org/validation/2017/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 : 1.8.5 (274361), CSD as541be (2020)
Xtriage (Phenix) : 1.13
EDS : **FAILED**
Percentile statistics : 20191225.v01 (using entries in the PDB archive December 25th 2019)
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.14.6

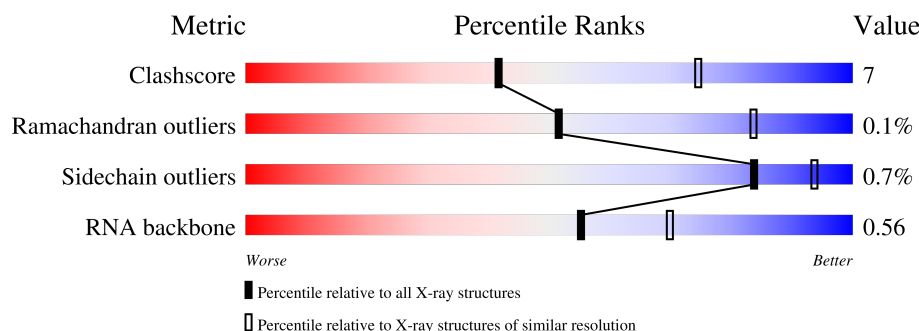
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 3.20 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))
Clashscore	141614	1253 (3.20-3.20)
Ramachandran outliers	138981	1234 (3.20-3.20)
Sidechain outliers	138945	1233 (3.20-3.20)
RNA backbone	3102	1010 (3.50-2.90)


























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 respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions $\leq 5\%$

Note EDS failed to run properly.

Mol	Chain	Length	Quality of chain
1	QA	1521	
1	XA	1521	
2	QB	256	
2	XB	256	
3	QC	239	
3	XC	239	












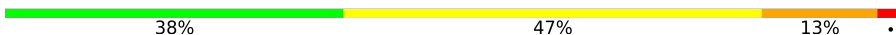
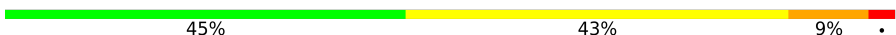
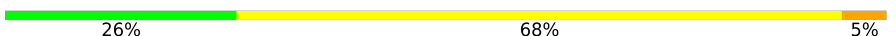











Continued on next page...

Continued from previous page...

Mol	Chain	Length	Quality of chain
4	QD	209	 77% 22% .
4	XD	209	 66% 30% .
5	QE	162	 67% 25% . 7%
5	XE	162	 76% 17% . 7%
6	QF	101	 74% 25% .
6	XF	101	 80% 18% .
7	QG	156	 79% 19% ..
7	XG	156	 72% 25% ..
8	QH	138	 78% 22% .
8	XH	138	 76% 22% ..
9	QI	128	 68% 27% . .
9	XI	128	 60% 32% 6% .
10	QJ	105	 56% 34% . 6%
10	XJ	105	 68% 21% . 9%
11	QK	129	 64% 25% . 8%
11	XK	129	 71% 18% . 10%
12	QL	132	 78% 15% . 5%
12	XL	132	 67% 20% 5% 8%
13	QM	126	 63% 28% . 5%
13	XM	126	 53% 37% . 6%
14	QN	61	 51% 43% 5% .
14	XN	61	 75% 18% 5% .
15	QO	89	 88% 10% ..
15	XO	89	 82% 16% .
16	QP	88	 68% 26% . 5%












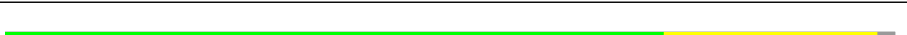

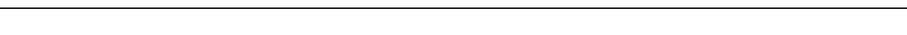
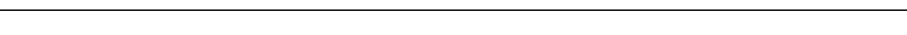
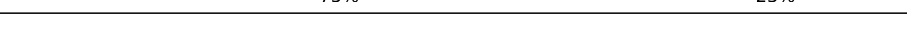

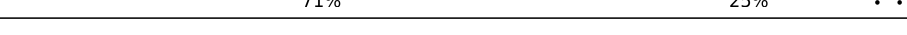







Continued on next page...

Continued from previous page...

Mol	Chain	Length	Quality of chain
16	XP	88	
17	QQ	105	
17	XQ	105	
18	QR	88	
18	XR	88	
19	QS	93	
19	XS	93	
20	QT	106	
20	XT	106	
21	QU	27	
21	XU	27	
22	QV	77	
22	XV	77	
23	QX	19	
23	XX	19	
24	RA	2915	
24	YA	2915	
25	RB	122	
25	YB	122	
26	RD	276	
26	YD	276	
27	RE	206	
27	YE	206	
28	RF	210	
28	YF	210	


























Continued on next page...

Continued from previous page...

Mol	Chain	Length	Quality of chain
29	RG	182	
29	YG	182	
30	RH	180	
30	YH	180	
31	RI	148	
31	YI	148	
32	RN	140	
32	YN	140	
33	RO	122	
33	YO	122	
34	RP	150	
34	YP	150	
35	RQ	141	
35	YQ	141	
36	RR	118	
36	YR	118	
37	RS	112	
37	YS	112	
38	RT	146	
38	YT	146	
39	RU	118	
39	YU	118	
40	RV	101	
40	YV	101	
41	RW	113	



Continued on next page...

Continued from previous page...

Mol	Chain	Length	Quality of chain
41	YW	113	 82% 17% .
42	RX	96	 82% 14% .
42	YX	96	 86% 11% .
43	RY	110	 78% 18% . .
43	YY	110	 81% 15% . .
44	RZ	206	 54% 32% . 11%
44	YZ	206	 67% 21% 11%
45	R0	85	 71% 24% . 5%
45	Y0	85	 75% 13% 12%
46	R1	98	 79% 16% . .
46	Y1	98	 72% 20% . 5%
47	R2	72	 72% 24% .
47	Y2	72	 69% 22% 8%
48	R3	60	 72% 27% .
48	Y3	60	 78% 20% .
49	R4	71	 58% 37% . .
49	Y4	71	 56% 35% 6% .
50	R5	60	 83% 15% .
50	Y5	60	 83% 15% .
51	R6	54	 81% 15% . .
51	Y6	54	 78% 19% . .
52	R7	49	 80% 16% .
52	Y7	49	 90% 8% .
53	R8	65	 60% 38% .
53	Y8	65	 62% 34% . . .

Continued on next page...

Continued from previous page...

Mol	Chain	Length	Quality of chain
54	R9	37	 78%22%
54	Y9	37	 70%30%

2 Entry composition

There are 57 unique types of molecules in this entry. The entry contains 291964 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	QA	1500	Total	C	N	O	P	0	0	0
			32247	14353	5981	10414	1499			
1	XA	1500	Total	C	N	O	P	0	0	0
			32249	14354	5984	10412	1499			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
2	QB	235	Total	C	N	O	S	0	0	0
			1907	1217	342	343	5			
2	XB	236	Total	C	N	O	S	0	0	0
			1915	1223	343	344	5			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
3	QC	205	Total	C	N	O	S	0	0	0
			1605	1011	313	280	1			
3	XC	205	Total	C	N	O	S	0	0	0
			1605	1011	313	280	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
4	QD	208	Total	C	N	O	S	0	0	0
			1703	1066	339	291	7			
4	XD	208	Total	C	N	O	S	0	0	0
			1703	1066	339	291	7			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
5	QE	151	Total	C	N	O	S	0	0	0
			1155	729	218	204	4			
5	XE	151	Total	C	N	O	S	0	0	0
			1155	729	218	204	4			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
6	QF	101	Total	C	N	O	S	0	0	0
			843	531	155	154	3			
6	XF	101	Total	C	N	O	S	0	0	0
			843	531	155	154	3			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
7	QG	155	Total	C	N	O	S	0	0	0
			1257	781	252	218	6			
7	XG	155	Total	C	N	O	S	0	0	0
			1257	781	252	218	6			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
8	QH	137	Total	C	N	O	S	0	0	0
			1108	700	214	192	2			
8	XH	137	Total	C	N	O	S	0	0	0
			1108	700	214	192	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
9	QI	127	Total	C	N	O		0	0	0
			1010	639	197	174				
9	XI	126	Total	C	N	O		0	0	0
			998	633	193	172				

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	QJ	99	Total	C	N	O	S	0	0	0
			801	504	157	139	1			

Continued on next page...

Continued from previous page...

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	XJ	96	Total	C	N	O	S	0	0	0
			777	487	153	136	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
11	QK	119	Total	C	N	O	S	0	0	0
			885	549	168	165	3			
11	XK	116	Total	C	N	O	S	0	0	0
			864	537	164	160	3			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
12	QL	125	Total	C	N	O	S	0	0	0
			975	614	196	164	1			
12	XL	122	Total	C	N	O	S	0	0	0
			956	603	193	159	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
13	QM	120	Total	C	N	O	S	0	0	0
			955	591	197	165	2			
13	XM	119	Total	C	N	O	S	0	0	0
			946	585	195	164	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
14	QN	60	Total	C	N	O	S	0	0	0
			492	312	104	72	4			
14	XN	60	Total	C	N	O	S	0	0	0
			492	312	104	72	4			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
15	QO	88	Total	C	N	O	S	0	0	0
			734	459	147	126	2			
15	XO	87	Total	C	N	O	S	0	0	0
			729	457	146	124	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
16	QP	84	Total	C	N	O	S	0	0	0
			705	446	140	118	1			
16	XP	84	Total	C	N	O	S	0	0	0
			705	446	140	118	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
17	QQ	100	Total	C	N	O	S	0	0	0
			834	534	155	143	2			
17	XQ	100	Total	C	N	O	S	0	0	0
			834	534	155	143	2			

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

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
18	QR	70	Total	C	N	O	0	0	0
			574	367	112	95			
18	XR	70	Total	C	N	O	0	0	0
			574	367	112	95			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
19	QS	83	Total	C	N	O	S	0	0	0
			665	424	124	115	2			
19	XS	84	Total	C	N	O	S	0	0	0
			674	430	126	116	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
20	QT	99	Total	C	N	O	S	0	0	0
			763	470	162	129	2			
20	XT	99	Total	C	N	O	S	0	0	0
			763	470	162	129	2			

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

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
21	QU	25	Total	C	N	O	0	0	0
			217	134	52	31			
21	XU	25	Total	C	N	O	0	0	0
			217	134	52	31			

- Molecule 22 is a RNA chain called P-site tRNA-Pro.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
22	QV	77	Total	C	N	O	P	0	0	0
			1648	734	295	542	77			
22	XV	77	Total	C	N	O	P	0	0	0
			1648	734	295	542	77			

- Molecule 23 is a RNA chain called mRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
23	QX	19	Total	C	N	O	P	0	0	0
			418	186	86	127	19			
23	XX	19	Total	C	N	O	P	0	0	0
			418	186	86	127	19			

- Molecule 24 is a RNA chain called 23S rRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
24	RA	2881	Total	C	N	O	P	0	0	0
			62051	27618	11609	19944	2880			
24	YA	2883	Total	C	N	O	P	0	0	0
			62091	27636	11613	19960	2882			

- Molecule 25 is a RNA chain called 5S rRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
25	RB	120	Total	C	N	O	P	0	0	0
			2573	1146	476	832	119			
25	YB	120	Total	C	N	O	P	0	0	0
			2573	1146	476	832	119			

- Molecule 26 is a protein called 50S ribosomal protein L2.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	RD	274	Total	C	N	O	S	0	0	0
			2135	1347	426	359	3			

Continued on next page...

Continued from previous page...

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	YD	274	Total	C	N	O	S	0	0	0
			2135	1347	426	359	3			

- Molecule 27 is a protein called 50S ribosomal protein L3.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
27	RE	205	Total	C	N	O	S	0	0	0
			1568	991	300	271	6			
27	YE	204	Total	C	N	O	S	0	0	0
			1563	988	299	270	6			

- Molecule 28 is a protein called 50S ribosomal protein L4.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
28	RF	202	Total	C	N	O	S	0	0	0
			1585	1011	297	275	2			
28	YF	202	Total	C	N	O	S	0	0	0
			1585	1011	297	275	2			

- Molecule 29 is a protein called 50S ribosomal protein L5.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
29	RG	181	Total	C	N	O	S	0	0	0
			1474	942	268	260	4			
29	YG	181	Total	C	N	O	S	0	0	0
			1474	942	268	260	4			

- Molecule 30 is a protein called 50S ribosomal protein L6.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
30	RH	174	Total	C	N	O	S	0	0	0
			1336	848	251	236	1			
30	YH	173	Total	C	N	O	S	0	0	0
			1330	845	250	234	1			

- Molecule 31 is a protein called 50S ribosomal protein L9.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
31	RI	146	Total	C	N	O	S	0	0	0
			1136	726	201	208	1			
31	YI	146	Total	C	N	O	S	0	0	0
			1136	726	201	208	1			

- Molecule 32 is a protein called 50S ribosomal protein L13.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
32	RN	138	Total	C	N	O	S	0	0	0
			1104	712	206	182	4			
32	YN	140	Total	C	N	O	S	0	0	0
			1121	722	208	187	4			

- Molecule 33 is a protein called 50S ribosomal protein L14.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
33	RO	122	Total	C	N	O	S	0	0	0
			933	588	171	170	4			
33	YO	122	Total	C	N	O	S	0	0	0
			933	588	171	170	4			

- Molecule 34 is a protein called 50S ribosomal protein L15.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
34	RP	149	Total	C	N	O	S	0	0	0
			1139	709	231	196	3			
34	YP	147	Total	C	N	O	S	0	0	0
			1122	698	229	192	3			

- Molecule 35 is a protein called 50S ribosomal protein L16.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
35	RQ	141	Total	C	N	O	S	0	0	0
			1122	715	212	188	7			
35	YQ	141	Total	C	N	O	S	0	0	0
			1122	715	212	188	7			

- Molecule 36 is a protein called 50S ribosomal protein L17.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
36	RR	117	Total	C	N	O	0	0	0
			960	599	202	159			
36	YR	117	Total	C	N	O	0	0	0
			960	599	202	159			

- Molecule 37 is a protein called 50S ribosomal protein L18.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
37	RS	111	Total	C	N	O	0	0	0
			882	556	176	150			
37	YS	110	Total	C	N	O	0	0	0
			877	553	175	149			

- Molecule 38 is a protein called 50S ribosomal protein L19.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
38	RT	137	Total	C	N	O	S	0	0	0
			1141	710	234	196	1			
38	YT	137	Total	C	N	O	S	0	0	0
			1141	710	234	196	1			

- Molecule 39 is a protein called 50S ribosomal protein L20.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
39	RU	117	Total	C	N	O	S	0	0	0
			964	610	202	151	1			
39	YU	117	Total	C	N	O	S	0	0	0
			964	610	202	151	1			

- Molecule 40 is a protein called 50S ribosomal protein L21.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
40	RV	101	Total	C	N	O	S	0	0	0
			779	501	142	135	1			
40	YV	101	Total	C	N	O	S	0	0	0
			779	501	142	135	1			

- Molecule 41 is a protein called 50S ribosomal protein L22.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
41	RW	113	Total	C	N	O	S	0	0	0
			900	566	177	155	2			
41	YW	113	Total	C	N	O	S	0	0	0
			900	566	177	155	2			

- Molecule 42 is a protein called 50S ribosomal protein L23.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
42	RX	92	Total	C	N	O	0	0	0
			725	471	131	123			

Continued on next page...

Continued from previous page...

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
42	YX	94	Total	C	N	O	S	0	0	0
			742	482	134	125	1			

- Molecule 43 is a protein called 50S ribosomal protein L24.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
43	RY	107	Total	C	N	O	S	0	0	0
			818	525	155	132	6			
43	YY	107	Total	C	N	O	S	0	0	0
			818	525	155	132	6			

- Molecule 44 is a protein called 50S ribosomal protein L25.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
44	RZ	183	Total	C	N	O	S	0	0	0
			1461	933	260	265	3			
44	YZ	183	Total	C	N	O	S	0	0	0
			1461	933	260	265	3			

- Molecule 45 is a protein called 50S ribosomal protein L27.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
45	R0	81	Total	C	N	O	S	0	0	0
			643	398	137	107	1			
45	Y0	75	Total	C	N	O	S	0	0	0
			599	370	127	101	1			

- Molecule 46 is a protein called 50S ribosomal protein L28.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
46	R1	97	Total	C	N	O	S	0	0	0
			763	481	150	131	1			
46	Y1	93	Total	C	N	O	S	0	0	0
			729	457	145	126	1			

- Molecule 47 is a protein called 50S ribosomal protein L29.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
47	R2	69	Total	C	N	O	S	0	0	0
			581	358	118	104	1			
47	Y2	66	Total	C	N	O	S	0	0	0
			558	346	113	98	1			

- Molecule 48 is a protein called 50S ribosomal protein L30.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
48	R3	59	Total	C	N	O	0	0	0
			469	298	90	81			
48	Y3	59	Total	C	N	O	0	0	0
			469	298	90	81			

- Molecule 49 is a protein called 50S ribosomal protein L31.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
49	R4	69	Total	C	N	O	S	0	0	0
			565	356	103	101	5			
49	Y4	69	Total	C	N	O	S	0	0	0
			565	356	103	101	5			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
50	R5	59	Total	C	N	O	S	0	0	0
			459	288	90	76	5			
50	Y5	59	Total	C	N	O	S	0	0	0
			459	288	90	76	5			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
51	R6	53	Total	C	N	O	S	0	0	0
			453	281	91	77	4			
51	Y6	53	Total	C	N	O	S	0	0	0
			453	281	91	77	4			

- Molecule 52 is a protein called 50S ribosomal protein L34.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
52	R7	47	Total	C	N	O	S	0	0	0
			409	251	102	54	2			
52	Y7	48	Total	C	N	O	S	0	0	0
			418	257	104	55	2			

- Molecule 53 is a protein called 50S ribosomal protein L35.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
53	R8	64	Total	C	N	O	S	0	0	0
			517	331	102	82	2			
53	Y8	64	Total	C	N	O	S	0	0	0
			517	331	102	82	2			

- Molecule 54 is a protein called 50S ribosomal protein L36.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
54	R9	37	Total	C	N	O	S	0	0	0
			307	188	68	47	4			
54	Y9	37	Total	C	N	O	S	0	0	0
			307	188	68	47	4			

- Molecule 55 is MAGNESIUM ION (three-letter code: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
55	QA	64	Total	Mg	0	0
			64	64		
55	YA	510	Total	Mg	0	0
			510	510		
55	Y5	1	Total	Mg	0	0
			1	1		
55	YR	1	Total	Mg	0	0
			1	1		
55	RT	1	Total	Mg	0	0
			1	1		
55	RN	1	Total	Mg	0	0
			1	1		
55	XE	1	Total	Mg	0	0
			1	1		
55	Y1	2	Total	Mg	0	0
			2	2		
55	YD	2	Total	Mg	0	0
			2	2		
55	QV	1	Total	Mg	0	0
			1	1		
55	RX	1	Total	Mg	0	0
			1	1		
55	Y8	1	Total	Mg	0	0
			1	1		
55	XA	78	Total	Mg	0	0
			78	78		
55	RQ	1	Total	Mg	0	0
			1	1		

Continued on next page...

Continued from previous page...

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
55	R0	2	Total 2	Mg 2	0	0
55	Y0	2	Total 2	Mg 2	0	0
55	YQ	3	Total 3	Mg 3	0	0
55	R8	2	Total 2	Mg 2	0	0
55	RR	1	Total 1	Mg 1	0	0
55	Y7	1	Total 1	Mg 1	0	0
55	RA	444	Total 444	Mg 444	0	0
55	Y3	1	Total 1	Mg 1	0	0
55	YP	3	Total 3	Mg 3	0	0
55	RE	5	Total 5	Mg 5	0	0
55	YB	7	Total 7	Mg 7	0	0
55	XV	1	Total 1	Mg 1	0	0
55	RB	7	Total 7	Mg 7	0	0
55	RF	1	Total 1	Mg 1	0	0
55	XM	2	Total 2	Mg 2	0	0
55	YE	5	Total 5	Mg 5	0	0

- Molecule 56 is IRON/SULFUR CLUSTER (three-letter code: SF4) (formula: Fe₄S₄).



Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
56	QD	1	Total	Fe	S	0	0
			8	4	4		
56	XD	1	Total	Fe	S	0	0
			8	4	4		

- Molecule 57 is ZINC ION (three-letter code: ZN) (formula: Zn).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	Y9	1	Total	Zn	0	0
			1	1		
57	YY	1	Total	Zn	0	0
			1	1		
57	Y6	1	Total	Zn	0	0
			1	1		
57	QN	1	Total	Zn	0	0
			1	1		
57	XN	1	Total	Zn	0	0
			1	1		
57	RY	1	Total	Zn	0	0
			1	1		
57	Y4	1	Total	Zn	0	0
			1	1		
57	R6	1	Total	Zn	0	0
			1	1		
57	Y5	1	Total	Zn	0	0
			1	1		
57	R5	1	Total	Zn	0	0
			1	1		

Continued on next page...

Continued from previous page...

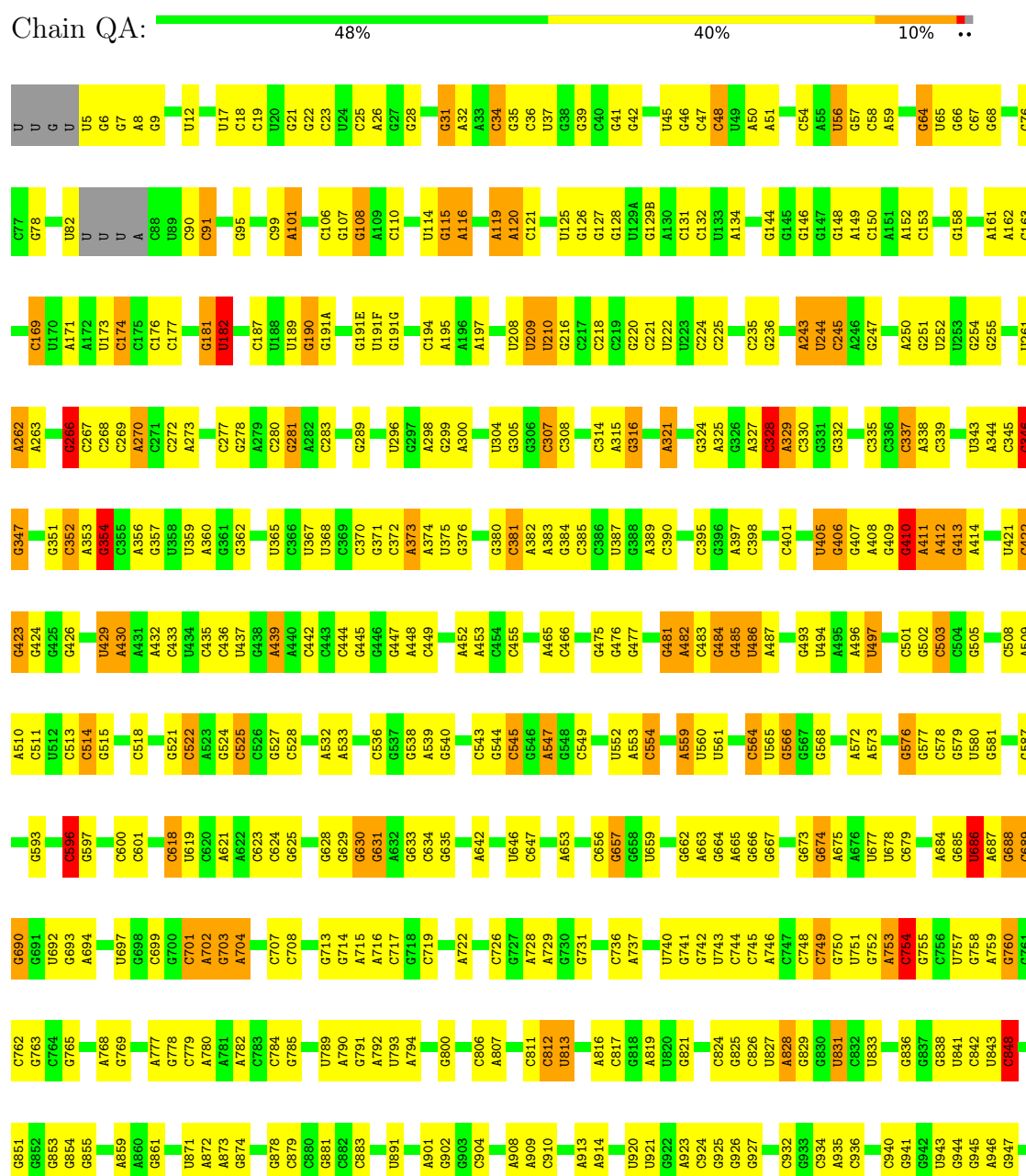
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	R4	1	Total 1	Zn 1	0	0
57	R9	1	Total 1	Zn 1	0	0

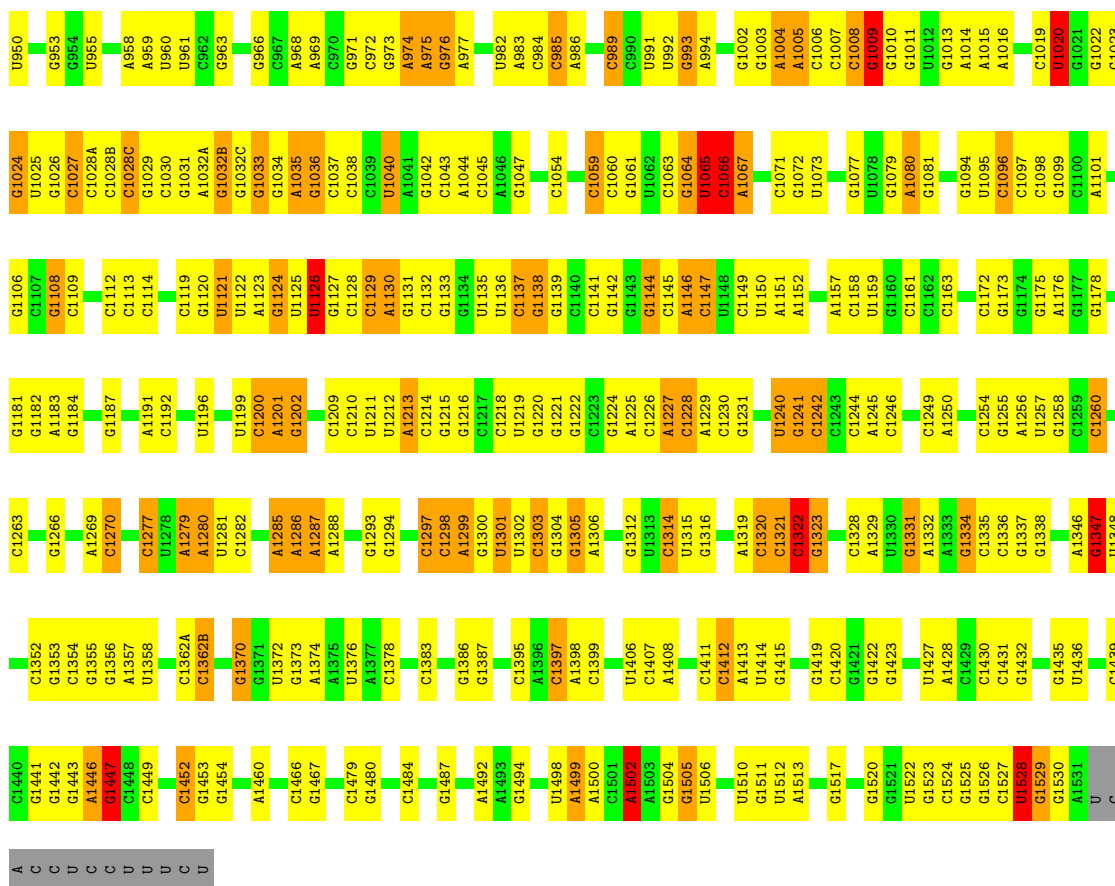
3 Residue-property plots

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

Note EDS failed to run properly.

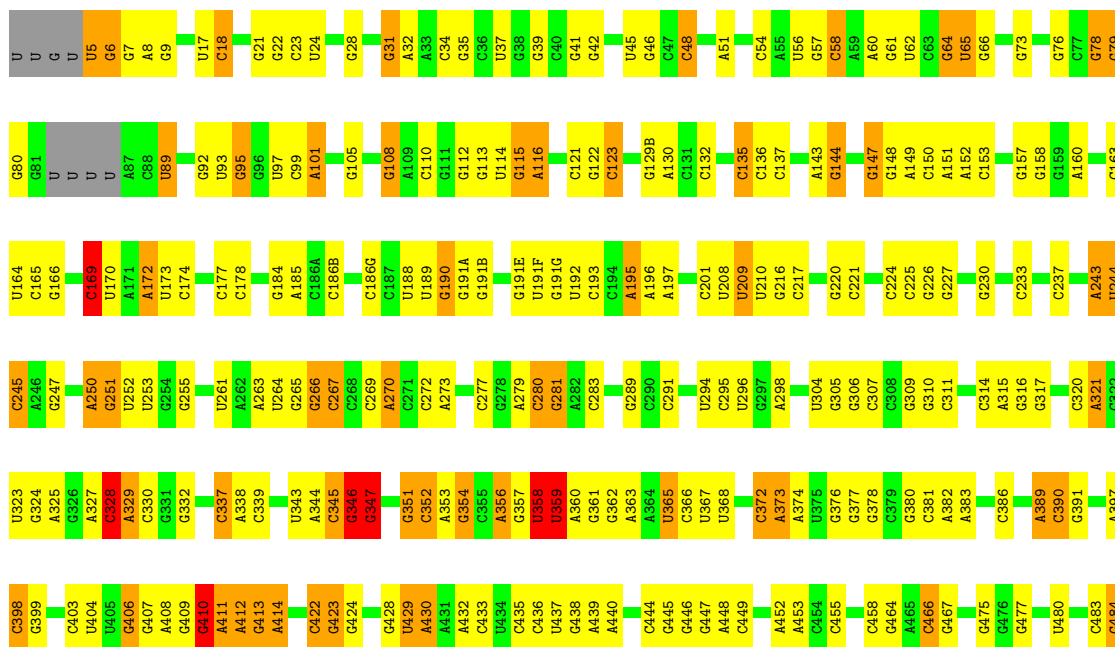
• Molecule 1: 16S rRNA

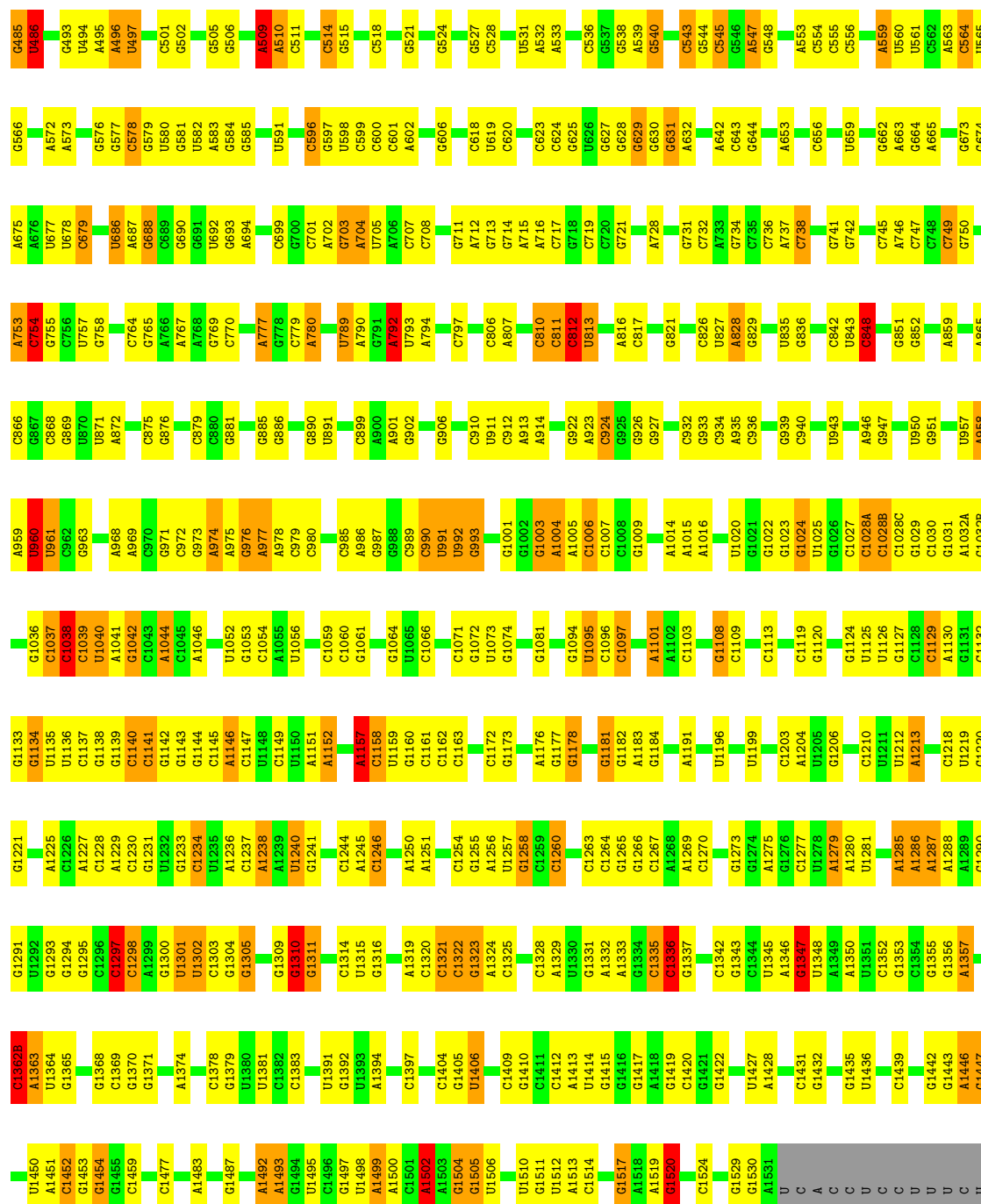




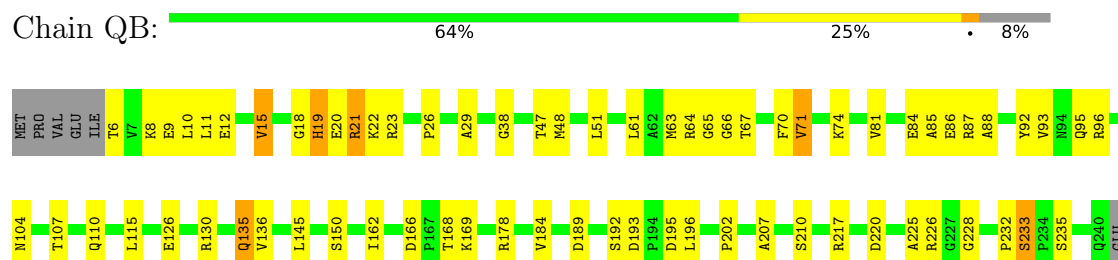
- Molecule 1: 16S rRNA

Chain XA: 49% 38% 10%





- Molecule 2: 30S ribosomal protein S2



ALA
GLU
ALA
THR
GLU
THR
PRO
GLY
GLU
SER
GLU
VAL
GLU
ALA

• Molecule 2: 30S ribosomal protein S2

Chain XB:  65% 25% 8%

MET PRO VAL VAL I5 T6 L11 V15 H16 F17 G18 H19 E20 F27 F28 A29 A30 Y31 I32 I40 T47 R64 T67 I68 L69 F70 K75 Q76 A77 Q78 R82 M83 E84 A85 E86 R87 A88 G89 M90 P91 Y92 Q95 R96 N104 F105 K106 L118

P131 Q135 V136 R137 L138 K139 H140 E141 L142 K147 L154 L155 D159 P160 F163 T168 I172 R178 I185 A186 L187 S192 D193 P194 D195 D198 Y199 I200 I201 P202 D206 A207 I208 I211 S216 R217 A225 R226 G227 P232 S233 P234 S235

V239 Q240 GLU ALA ALA THR GLU PRO GLY GLU SER VAL VAL ALA

• Molecule 3: 30S ribosomal protein S3

Chain QC:  63% 21% 14%

MET G2 N3 I8 G9 F10 F11 L12 L14 I15 T16 R16 D17 W18 E19 S20 A24 Y29 L32 L40 L43 E44 L47 A80 G51 L52 R59 D82 N63 V70 A71 G74 G78 R88 L91 K97 N98 L101 V106 R126 R127

F128 R131 R132 A133 I134 K135 Q136 V151 S154 G155 R164 W18 T165 E166 W167 A168 R172 T177 I182 D183 Y184 A189 L196 A200 Y201 I202 E206 VAL ILE GLY GLN LYS LYS ALA ARG PRO GLU LEU PRO LYS N98 GLU ARG GLU ARG PRO ARG ARG

ARG PRO ALA VAL ARG VAL LYS LYS GLU


• Molecule 3: 30S ribosomal protein S3

Chain XC:  67% 18% 14%

MET G2 N3 G9 F10 F11 L12 D17 W18 E19 G25 Y29 E35 R40 G41 L42 K45 R54 V55 D56 R59 D62 T67 V70 A71 K72 G78 A92 K93 L94 S112 A113 P114 A117 Q118 Q136 A137 V138 Q139 R140 K147 G148

A149 K150 V151 I152 E166 W167 A168 P174 T177 Y184 A187 R25 L188 A189 L196 K199 A200 Y201 E206 VAL ILE GLY GLN LYS PRO LYS ALA ARG PRO GLU LEU PRO LYS ALA ALA ARG ARG ARG ARG PRO ALA VAL ARG VAL VAL LYS LYS GLU

• Molecule 4: 30S ribosomal protein S4

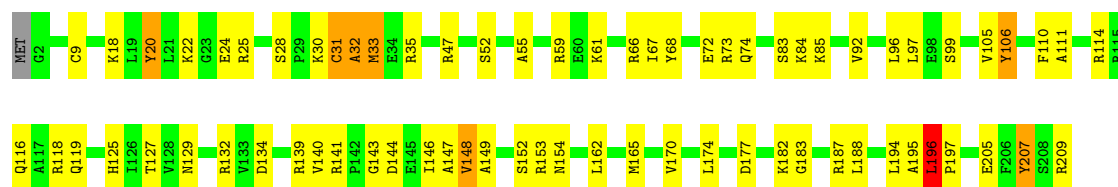
Chain QD:  77% 22%

MET G2 R3 Y4 R10 R13 R14 E15 K18 G23 E24 R25 C26 A32 M33 E34 R35 R36 P37 P39 P40 R47 A48 A49 R59 L64 Y68 G69 R73 S83 K84 V88 E98 L101 L108 R115 Q119 N129 D134

Y138 E145 R153 N154 L155 G167 G180 M181 K182 G183 N199 L202 E205 S208 R209

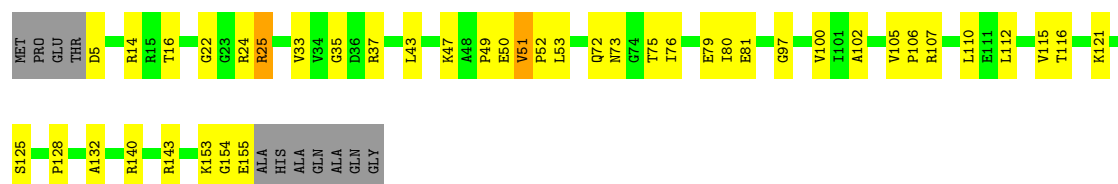
• Molecule 4: 30S ribosomal protein S4

Chain XD:  66% 30% .




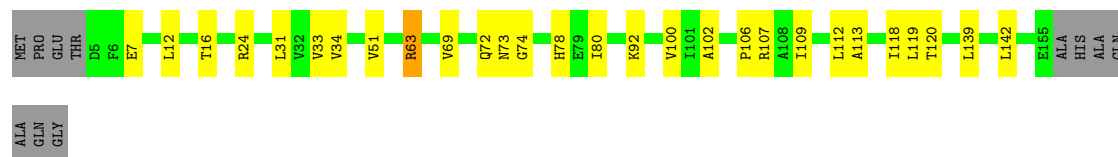
- Molecule 5: 30S ribosomal protein S5

Chain QE:  67% 25% . 7%




- Molecule 5: 30S ribosomal protein S5

Chain XE:  76% 17% . 7%




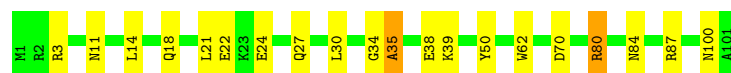
- Molecule 6: 30S ribosomal protein S6

Chain QF:  74% 25% .




- Molecule 6: 30S ribosomal protein S6

Chain XF:  80% 18% .

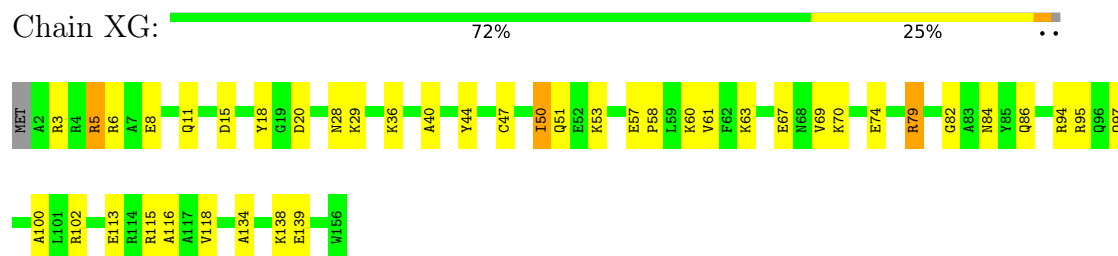


- Molecule 7: 30S ribosomal protein S7

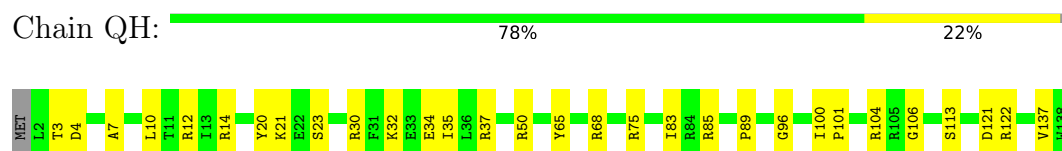
Chain QG:  79% 19% ..



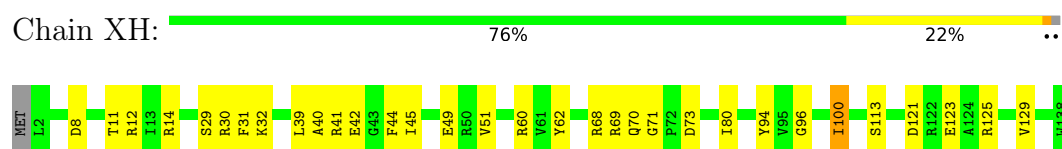
- Molecule 7: 30S ribosomal protein S7



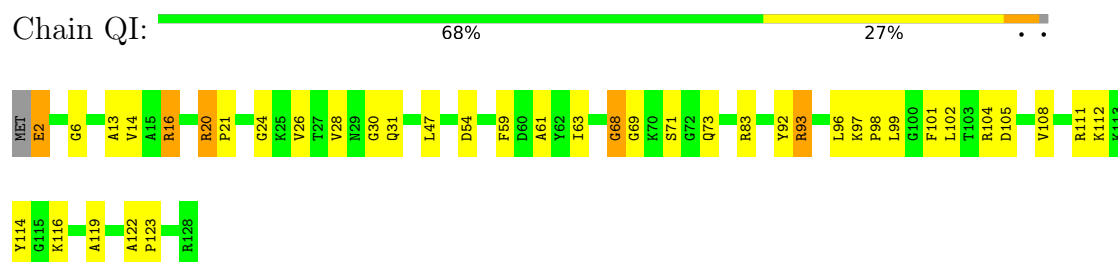
- Molecule 8: 30S ribosomal protein S8



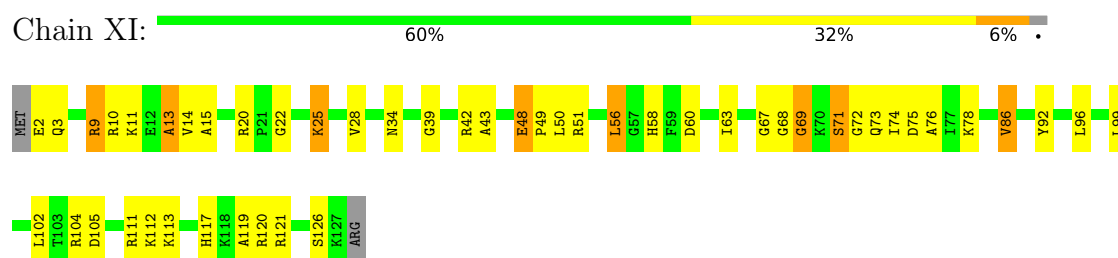
- Molecule 8: 30S ribosomal protein S8



- Molecule 9: 30S ribosomal protein S9

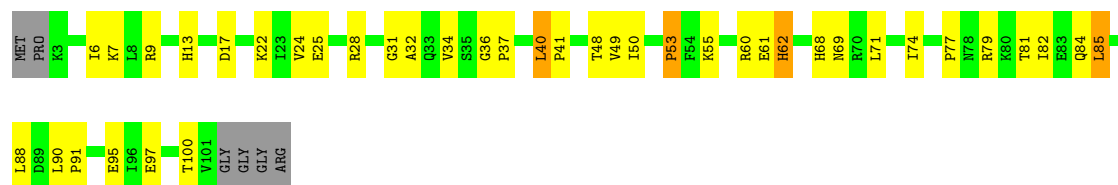


- Molecule 9: 30S ribosomal protein S9



- Molecule 10: 30S ribosomal protein S10





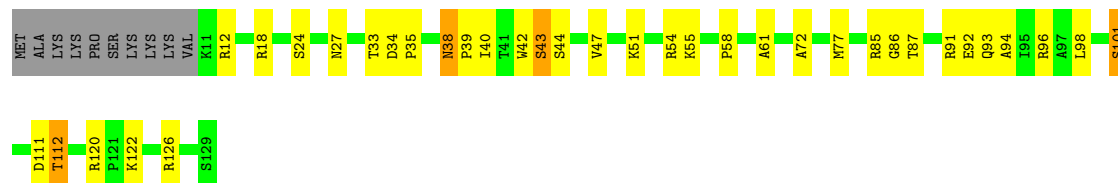
- Molecule 10: 30S ribosomal protein S10

Chain XJ: 68% 21% 9%



- Molecule 11: 30S ribosomal protein S11

Chain QK: 64% 25% 8%



- Molecule 11: 30S ribosomal protein S11

Chain XK: 71% 18% 10%



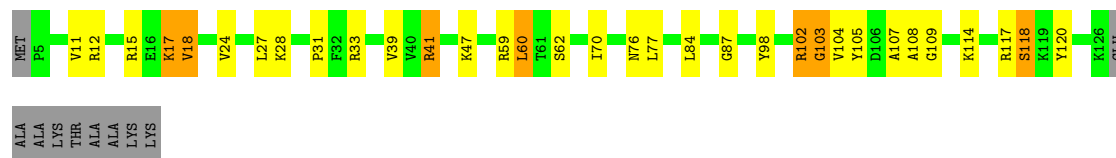
- Molecule 12: 30S ribosomal protein S12

Chain QL: 78% 15% 5%



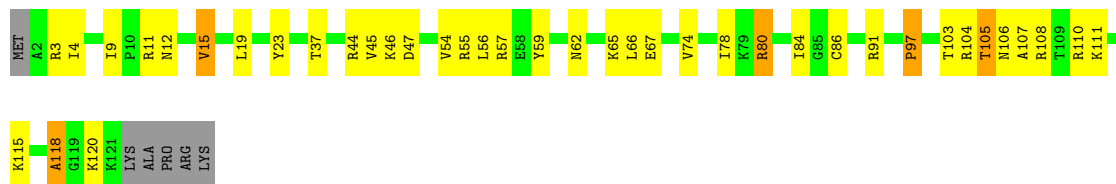
- Molecule 12: 30S ribosomal protein S12

Chain XL: 67% 20% 5% 8%



- Molecule 13: 30S ribosomal protein S13

Chain QM:  63% 28% 5%



- Molecule 13: 30S ribosomal protein S13

Chain XM:  53% 37% 6%




- Molecule 14: 30S ribosomal protein S14 type Z

Chain QN:  51% 43% 5%




- Molecule 14: 30S ribosomal protein S14 type Z

Chain XN:  75% 18% 5%




- Molecule 15: 30S ribosomal protein S15

Chain QO:  88% 10% 2%



- Molecule 15: 30S ribosomal protein S15

Chain XO:  82% 16% 2%



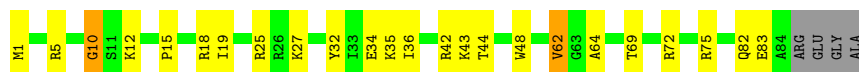
- Molecule 16: 30S ribosomal protein S16

Chain QP:  68% 26% 5%



- Molecule 16: 30S ribosomal protein S16

Chain XP: 68% 25% • 5%



- Molecule 17: 30S ribosomal protein S17

Chain QQ: 61% 32% • 5%



- Molecule 17: 30S ribosomal protein S17

Chain XQ: 79% 15% • 5%



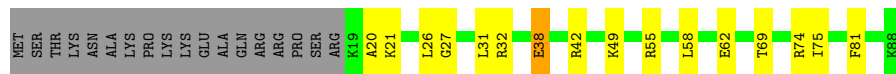
- Molecule 18: 30S ribosomal protein S18

Chain QR: 61% 18% 20%



- Molecule 18: 30S ribosomal protein S18

Chain XR: 61% 17% 20%

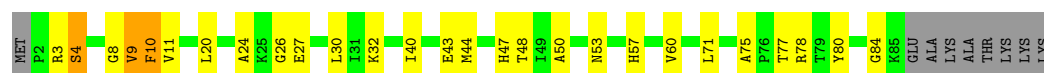


- Molecule 19: 30S ribosomal protein S19

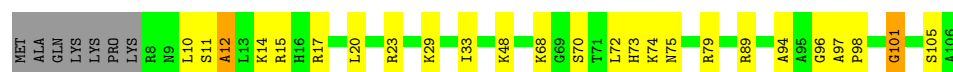
Chain QS: 66% 22% • 11%



- Molecule 19: 30S ribosomal protein S19



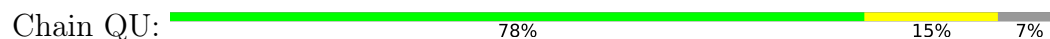
- Molecule 20: 30S ribosomal protein S20



- Molecule 20: 30S ribosomal protein S20



- Molecule 21: 30S ribosomal protein Thx



- Molecule 21: 30S ribosomal protein Thx



- Molecule 22: P-site tRNA-Pro

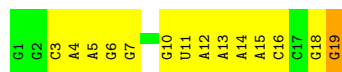


- Molecule 22: P-site tRNA-Pro



- Molecule 23: mRNA

Chain QX: 26% 68% 5%



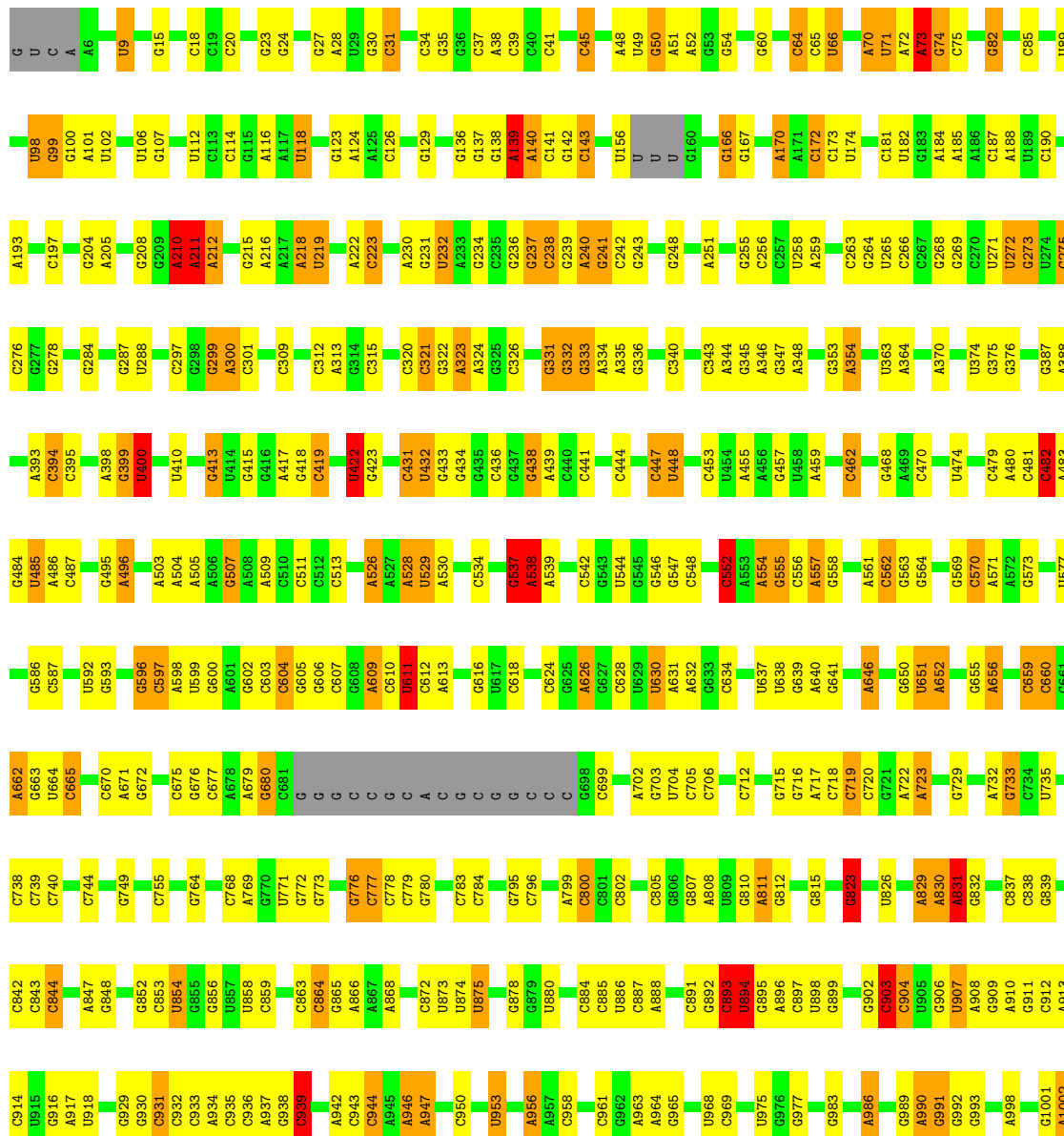
• Molecule 23: mRNA

Chain XX: 26% 58% 11% 5%



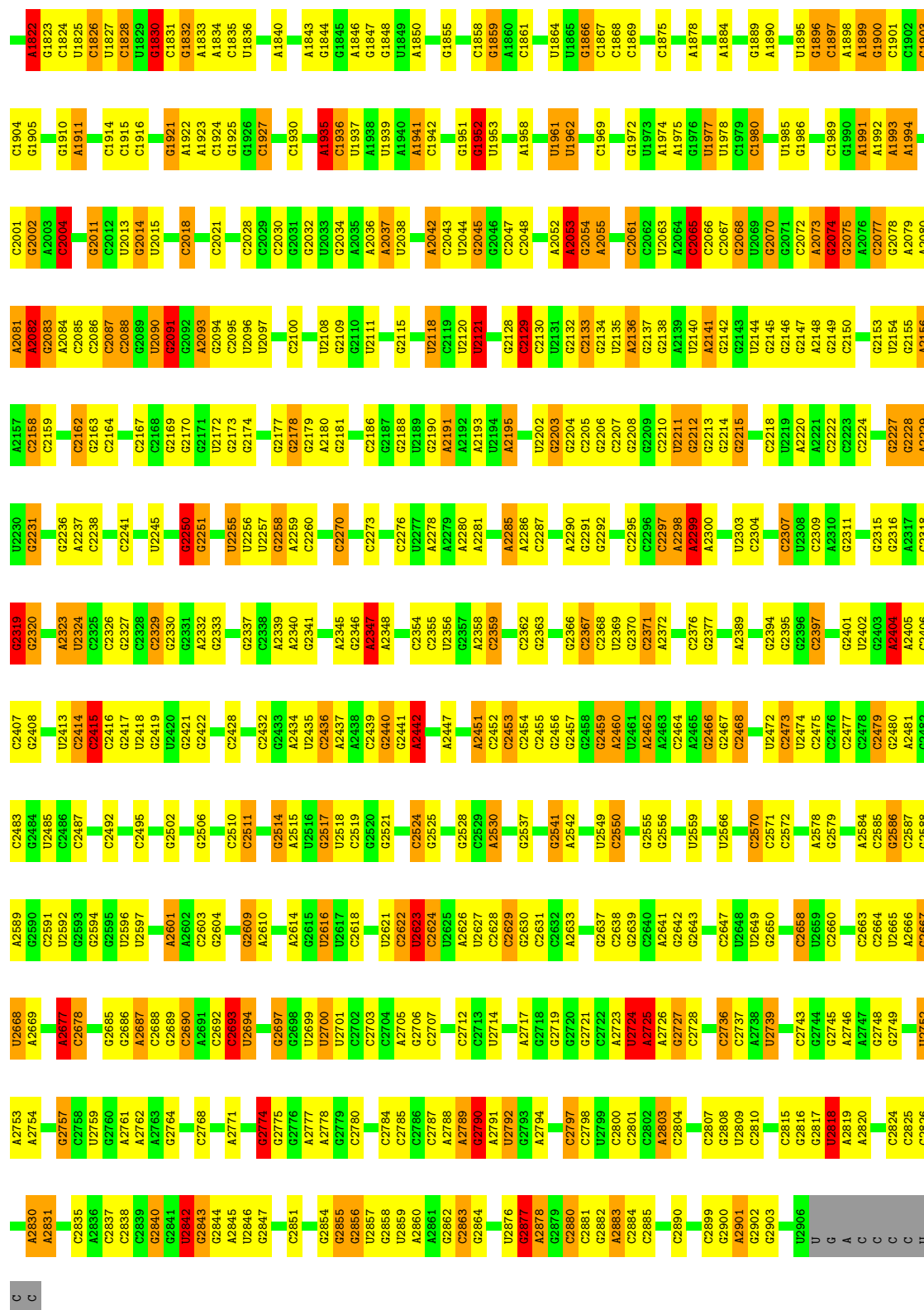
• Molecule 24: 23S rRNA

Chain RA: 51% 35% 11% ..



A2157	A2081	A1993	C1904	G1823	U1726	G1639	A1491	C1316	G1230	G1152	C1087	U1003
C2158	A2082	A1994	G1910	C1824	U1727	G1640	A1491	G1317	G1231	G1153	C1088	A1004
C2159	A2083	A1994	G1910	C1824	G1728	G1641	C1492	A1318	G1232	U1154	C1089	A1005
C2160	A2084	C2001	A1911	U1827	G1729	A1642	C1493	G1415	U1233	G1155	G1090	C1006
	C2085	C2002	A1912	C1828	C1730	A1643	G1415	A1320	U1234	G1156	A1091	
C2164	C2086	A2093		U1829	C1731	C1844	A1496	A1321	G1235	A1157	A1092	C1010
C2165	C2087	C2004	C1916	G1830	C1732	C1645	G1497			G1157	G1093	
U2166	C2088		C1831	C1831	C1733	C1645	G1497			G1158	A1094	U1014
C2167	U2089	U2013	G1921	G1832	C1734	U1648	U1501	U1328	G1240	U1159	A1094	C1015
C2168	U2090	G2014	A1922	G1833	A1736	A1649	G1502	G1160	G1248	G1161		A1018
C2169	C2091	U2015	A1923	A1834	C1650	C1574	A1574	C1331	A1249	G1331	C1098	G1019
C2170	C2092	C2016	A1924	C1835	A1745	C1653	G1576	A1332	U1250	G1163	C1099	C1020
		U2017	C1925		G1746	A1654	U1577	A1333	C1252	G1164	A1100	
C2171	G2094	C2018	G1928	A1843		A1655	G1578	C1336	G1254	G1168	A1103	G1021
C2172	C2095	C2018	G1929	G1844	G1763	A1656	C1579	C1336	C1255	G1169	G1104	C1022
C2173	U2096	G1930	U1846	G1845	G1766	C1657	G1580		U1256	C1170	G1105	
G2174	U2097	C1930	G1847	G1846	A1767	C1657	U1581	U1340	G1257	G1171	U1106	A1026
	U2098	C1931		G1847			U1581	C1431	U1258			
	A2099	C1931		G1847			U1581	C1432	G1259		U	A1029
C2178		G2024				A1662	C1514		A1173	A1173		A1030
C2179	U2108	C2028	A1935	U1848	G1769	C1663	C1515	C1341	A1174	A1174	G1108	C1031
A2180	C2109	C2029	C1936	G1849	G1769	C1663	C1515	C1342	A1175	A1175	G1109	
C2183	G2110	C2030	U1937	U1851	G1771			U1440	U1176	U1176	C1110	
C2184	U2111	G2031	A1938	A1852		G1668	G1585	A1346	G1263	G1177	U1111	G1035
C2185		C2032	U1939	G1853	C1776		C1521	A1347	G1264		U1112	A1036
C2186	G2115	U2033	C1942	G1854	G1776	U1675	G1522	C1350	A1265		A1113	C1037
	C2187	G2034		G1855		G1676	A1589	C1351		C1180	G1114	G1038
C2188	U2118	A2035	C1942	A1856	G1781	C1683	A1591	C1352	G1269	C1181	A1115	G1039
C2189	U2189	A2036	A1949	G1857	C1782	G1692	G1526	C1448	G1270	G1182	A1116	C1040
C2190	G2120	C2037	C1950	C1858	C1783	A1684	A1592	C1449	G1271	G1183	G1117	C1041
A2191	U2124	A2037	G1951				G1527	C1450	A1272	G1184	C1118	A1042
A2192	C2125	A2041	U1952	C1861	G1787	C1687	U1528	U1368	G1273	U1185	A1119	G1043
	G2126	A2042	U1953	G1862			C1595	U1369		U1186	G1120	C1044
C2199	C2129	C2043	C1956	C1863	A1793	C1691	G1530	C1361	C1276	A1187	C1121	U1045
C2200	U2130	G2045	G1957	C1864	G1794	G1692	A1532	U1362	G1277	A1188	C1122	A1046
C2201	U2131	C2046	A1958	G1865	G1795	C1693	A1532	U1362	G1278	A1188	A1123	G1047
	G2132	A2047	A1959	G1866	C1796	G1894	G1531	C1361	C1279	G1195	A1124	G1048
C2202	C2133	C2048	U1960	C1868	U1797	C1695	G1606	C1364	U1280	C1196	C1125	G1049
	G2134	G2049	U1961				G1607		G1281	G1197	C1126	C1051
G2204	U2135	G2050	C1962	C1875	U1799	G1698	G1612	G1473	G1296	G1214	A1130	A1055
C2205	A2136	G2051	C1963		G1801	A1699	A1541	A1473	G1297	G1214	A1130	
G2206	G2137	A2052		A1878	C1802	A1701	A1613	A1473	G1297	G1214	A1130	
C2207	G2138	C2053	C1969	A1879	G1803	A1702	G1615	G1471	G1289	U1211	A1131	U1068
C2208	A2139	G2054		G1880	A1804	C1703	A1545	G1374	G1290	C1212	A1132	C1059
C2209	U2140	A2055	G1972	G1881	C1805	U1806	G1617	U1375	G1291	U1213	G1133	
C2210	A2141	U2056			U1807	C1707	C1547	G1214			A1134	U1065
C2211	G2142		U1977	A1884	G1807	G1708	C1548	A1473	G1296	G1214	A1134	
C2212	G2143	G2059	C1978	A1885		C1709	U1625	A1473	G1296	G1214	A1134	U1066
C2213	U2144	G2060	C1979		U1810		A1626	A1473	G1297	G1214	A1134	A1067
G2214	G2145		C1980	G1889	A1811	A1712	A1626	G1475	G1298	G1218	G1137	G1068
G2215	C2065	C2065	G1981	C1812	C1812	C1713	C1551	U1476	A1299	A1219	C1138	U1069
	C2066	C2066	C1982	C1813	C1813	G1714	C1552	U1477	A1300	U1220	G1139	G1070
	G2147		C1983	A1814	A1814	C1681	A1553	G1478	G1394	U1221	U1140	G1071
A2220	G2148	C1896	C1984	C1897	A1815	A1716	A1554	U1479	G1302	A1222	A1141	U1072
G2221	G2149		U1985	A1898	A1816	A1717	C1555	A1480	C1303	C1223	A1142	A1073
C2222	G2150	G2075	C2076	A1899	A1817	C1717	A1556	A1481		C1224	U1143	
	G2153	G2076	C2077	A1900	C1634	C1721	A1557	G1482	G1310	C1225	C1148	G1077
C2226	U2154	C2077	G1990	A1818	C1635	G1588	C1558	C1483	A1311	C1226	A1149	U1078
C2228	G2155	A1991	C1902	G1901	U1636	C1722	G1589	C1483	G1312	A1227	A1149	U1079
	G2156	C1903	C1903	C1903	G1637	U1560	C1560	G1489	U1313	G1228	C1150	C1086

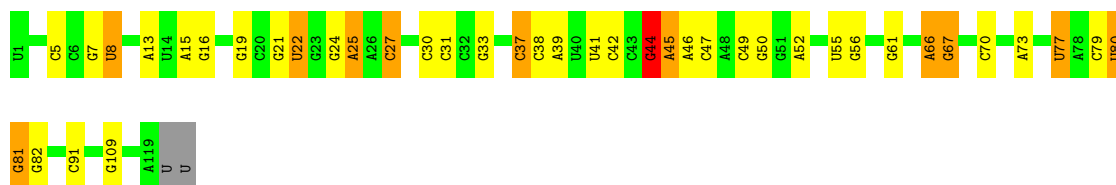
G1725	C1730	C1645	U1560	U1479	A1405	C1303	A1219	U1128	G1057	G983	G902	A830	C726	G659	G584
G1731	C1646	G1481	C1561	A1480	A1405	C1304	U1220	U1129	U1058	G984	G903	A831	C730	C660	U585
G1732	A1649	G1482	U1562	G1481	C1408	G1310	G1221	A1130	C1069	G985	C904	G832	G731	G661	G586
C1733	C1650	C1490	G1563	G1481	G1410	A1312	A1222	A1131	G1061	A986	U905	U834	A732	A662	C588
	G1651	A1491	G1568	A1490	G1411	G1311	C1223	A1132	C1064	G989	G906	U835	A733	G663	U589
	G1652	C1482	U1569	C1482	A1411	U1313	C1224	A1134	U1065	A990	U907	A836	C734	U664	A590
	C1653	C1493	A1574	C1493	G1414	G1317	C1225	A1135	U1066	A991	A908	A837	U735	C665	U591
	A1654	A1496	A1574	A1496	G1415	U1318	G1229	U1136	A1066	G991	A909	C838	U736	C666	U592
	A1655	G1497	C1577	G1497	G1416	A1319	G1230	G1139	A1067	G992	A909	C839	A737		
	A1656	C1498	C1578	C1498	G1417	U1320	G1231	U1140	U1068	G993	A910	A840	C738		
	C1657	A1498	C1579	A1498	G1418	A1321	G1232	U1141	U1069	C994	A913	G841	C739		
	C1658	A1500	G1580	A1500	U1419	U1322	U1233	A1142	G1071	A998	C914	G842			
		U501	U1581	U501	C1421	A1332	U1234	U1143	U1072	G999	U924	C843	C745		
		A1502	A1581	A1502	C1422	A1333	G1235	U1143	A1073	C1000		C844	A746		
		G1503	C1583	G1503	G1423	U1334	A1236	A1149	A1074	G1001			G749		
		A1504	G1584	A1504	A1424	C1335	C1245	A1150	A1075	U1002			G764		
		C1505	G1585	C1505	A1425	C1336	G1246	U1159	A1076	U1003					
		G1506	G1586	G1506	G1426	C1336	C1247	G1166	G1088	A1004					
		A1507	U1587	A1507		C1339	G1248	G1167	C1089	A1006					
		G1508	G1588	G1508	A1430	U1340	A1249	G1168		A1008					
		C1509	A1589	C1509	G1431	C1341	U1250	C1169	A1092	U1009					
		C1510	C1590	C1510	C1432	G1342	G1251	U1175	G1093	C939					
		C1511	A1591	C1511	C1433	C1343	C1252	A1177	A1096	C940					
			A1592					A1178	C1097	U941					
			A1592					A1179	G1098	A942					
			C1595	C1514	U1436	U1346	A1255	G1178	C1098	A943					
			C1596	C1515	U1437	A1347	U1256	A1179	A1099	A944					
			C1596		A1438	C1352	G1260	G1171	C1099	A945					
			C1603	A1518	A1439	A1352	G1261	C1171	C944	A946					
			C1604	C1521	U1440	A1354	C1263	A1172	C944	A947					
			A1605	C1526	A1441	G1355	G1264	A1173	C944	A947					
			G1606	G1527	C1444	G1355	A1265	A1174	C944	A947					
				G1528		U1359	C1266	A1175	C944	A947					
			A1613	U1528	C1449	C1360	G1269	G1183	A1097	C950					
				G1529	C1450	U1361	G1270	G1184	C1028	C800					
			A1616	G1590	U1451	U1362	C1271	U1179	A1029	C801					
			A1617	G1531	U1452	A1363	G1271	C1180	A1030	C802					
			A1618	C1453	C1453	C1364	A1272	G1181	C1032	C805					
				C1454	C1454		C1276	G1182	G1033	C805					
			C1624	A1536	C1455	U1372	C1276	G1183	A1034	C805					
			U1625	C1539	G1456	U1375	G1277	G1184	A1035	C805					
			A1627	A1542	C1457	U1378	G1278	C1185	A1036	C805					
			G1628	U1543	G1462	G1378	C1279	U1186	C1037	C805					
				C1544	C1463	U1388	C1280	U1187	C1038	C805					
			C1631	C1545	G1464	G1384	G1281	A1187	C1039	C805					
			A1632	G1546	A1465	U1385	G1282	A1112	G961	C805					
			A1633		U1466	U1386	A1283	A1113	C962	C805					
			C1634		G1467	U1387	G1284	G1114	A963	C805					
				C1550	G1467	U1387	G1285	A1115	A964	C805					
				C1551	C1468	A1388	U1286	A1116	G965	C805					
				A1552	G1472	G1389	U1286	G1117	C1044	C805					
			C1637	C1553	G1472	U1394	G1289	U1187	C1044	C805					
			C1638	A1554	A1473	A1395	G1298	A1188	C1044	C805					
			G1639	C1555	C1474	A1395	A1299	G1206	C1044	C805					
			A1640	C1556	G1475	A1398	A1299	G1207	U1045	C805					
			G1641	A1557	G1476	U1398	A1299	C1208	A1046	C805					
			A1642	G1558	U1477	A1399	A1299	G1209	U1046	C805					
			A1643		C1478	A1399	A1299	G1210	A1046	C805					
			C1644	C1559	C1478	A1400	G1302	C1212	U1124	C805					



- Molecule 25: 5S rRNA

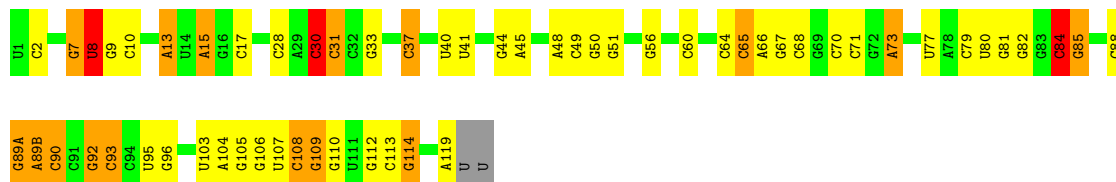
Chain RB:





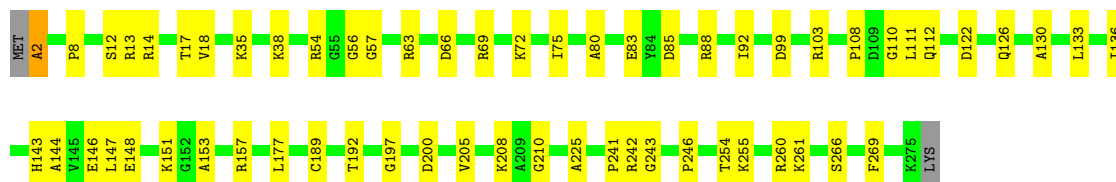
• Molecule 25: 5S rRNA

Chain YB: 51% 32% 13% ..



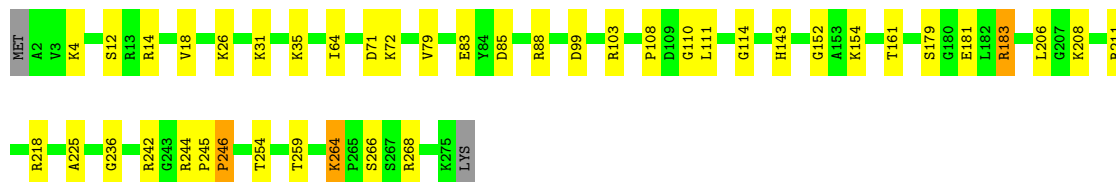
• Molecule 26: 50S ribosomal protein L2

Chain RD: 78% 21% .



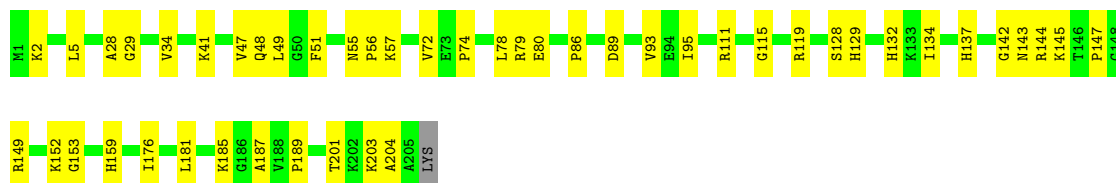
• Molecule 26: 50S ribosomal protein L2

Chain YD: 84% 14% ..



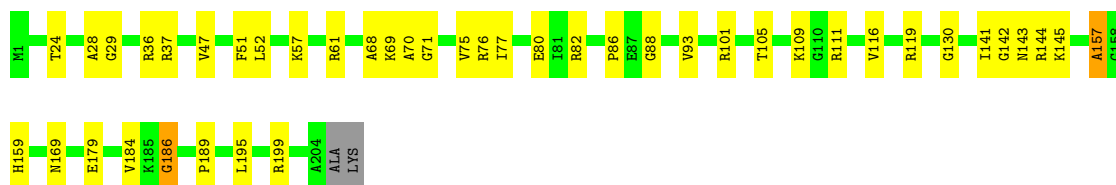
• Molecule 27: 50S ribosomal protein L3

Chain RE: 77% 23%



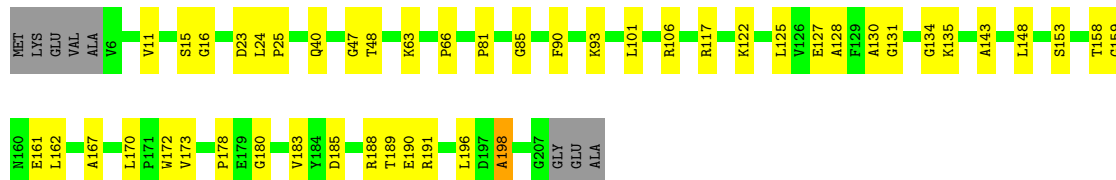
• Molecule 27: 50S ribosomal protein L3

Chain YE: 78% 20% ..



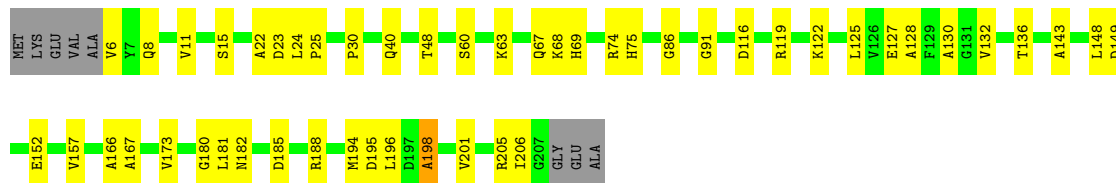
- Molecule 28: 50S ribosomal protein L4

Chain RF: 74% 22% .



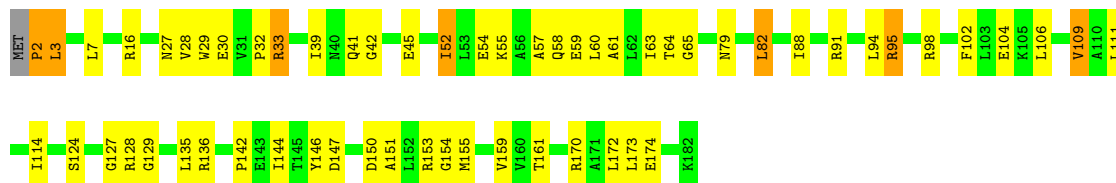
- Molecule 28: 50S ribosomal protein L4

Chain YF: 73% 23% .



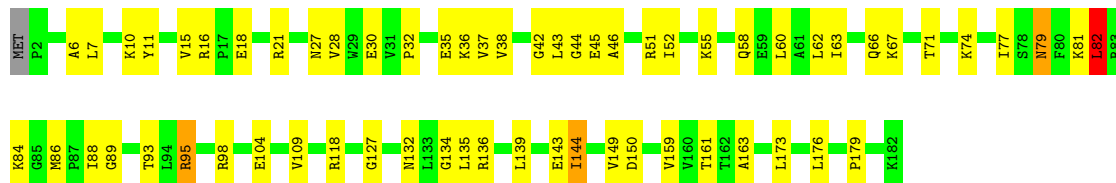
- Molecule 29: 50S ribosomal protein L5

Chain RG: 67% 29% . .



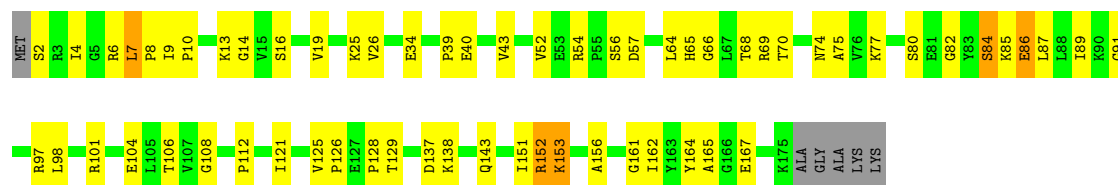
- Molecule 29: 50S ribosomal protein L5

Chain YG: 65% 32% . . .



- Molecule 30: 50S ribosomal protein L6

Chain RH: 62% 32% . . .



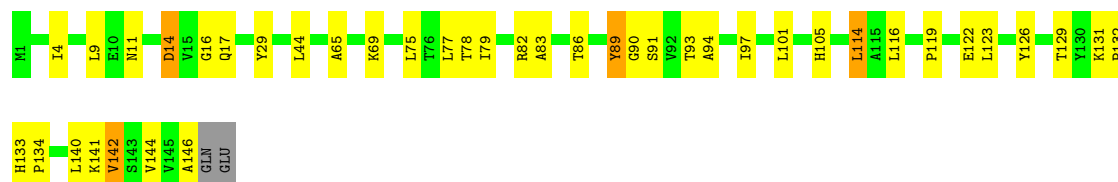
- Molecule 30: 50S ribosomal protein L6

Chain YH: ..



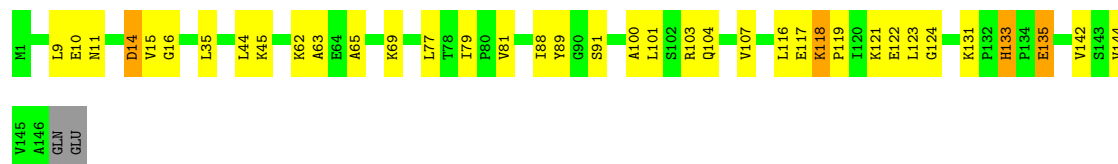
- Molecule 31: 50S ribosomal protein L9

Chain RI: ..



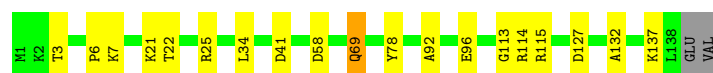
- Molecule 31: 50S ribosomal protein L9

Chain YI: ..



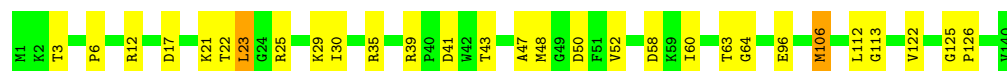
- Molecule 32: 50S ribosomal protein L13

Chain RN: ..




- Molecule 32: 50S ribosomal protein L13

Chain YN: ..




- Molecule 33: 50S ribosomal protein L14

Chain RO:  75% 23%




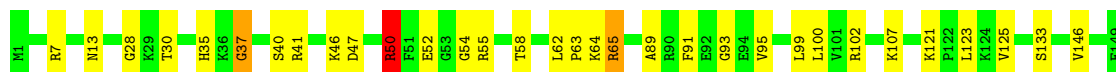
- Molecule 33: 50S ribosomal protein L14

Chain YO:  80% 19%




- Molecule 34: 50S ribosomal protein L15

Chain RP:  78% 19%



ALA

- Molecule 34: 50S ribosomal protein L15

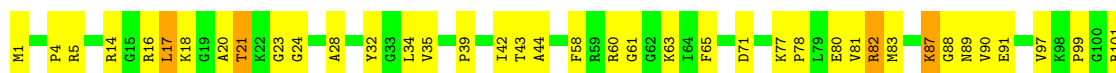
Chain YP:  74% 24%



S133, A141, L147, LEU, GLU, ALA


- Molecule 35: 50S ribosomal protein L16

Chain RQ:  62% 35%




V102, M103, F104, E105, G108, V109, T110, E111, E112, Q113, E116, A117, L118, R119, K130, R134, D135, A136, Y137, Q141

- Molecule 35: 50S ribosomal protein L16

Chain YQ:  86% 14%




- Molecule 36: 50S ribosomal protein L17

Chain RR:  75% 23% ..



- Molecule 36: 50S ribosomal protein L17

Chain YR:  82% 15% ..




- Molecule 37: 50S ribosomal protein L18

Chain RS:  71% 25% . .



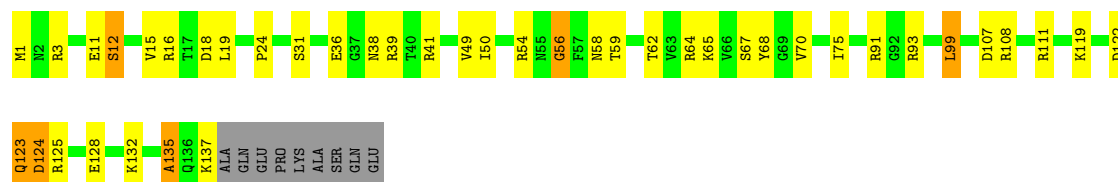
- Molecule 37: 50S ribosomal protein L18

Chain YS:  79% 20% .



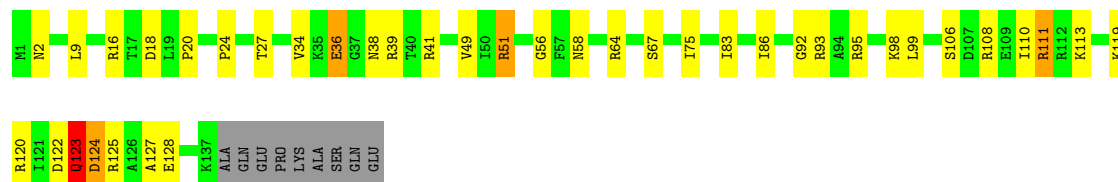
- Molecule 38: 50S ribosomal protein L19

Chain RT:  65% 25% . 6%




- Molecule 38: 50S ribosomal protein L19

Chain YT:  67% 23% . . 6%



- Molecule 39: 50S ribosomal protein L20

Chain RU:  78% 19% . .



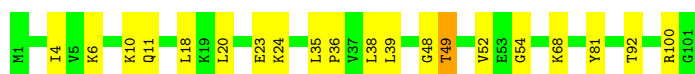
- Molecule 39: 50S ribosomal protein L20

Chain YU: 82% 14% . .



- Molecule 40: 50S ribosomal protein L21

Chain RV: 80% 19% .



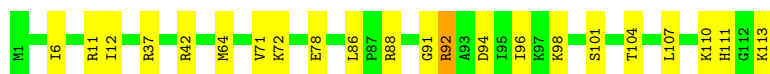
- Molecule 40: 50S ribosomal protein L21

Chain YV: 82% 13% 5%



- Molecule 41: 50S ribosomal protein L22

Chain RW: 81% 19% .



- Molecule 41: 50S ribosomal protein L22

Chain YW: 82% 17% .



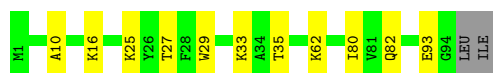
- Molecule 42: 50S ribosomal protein L23

Chain RX: 82% 14% .



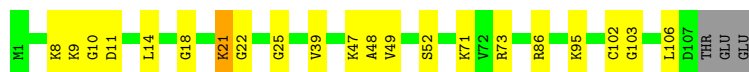
- Molecule 42: 50S ribosomal protein L23

Chain YX: 86% 11% .



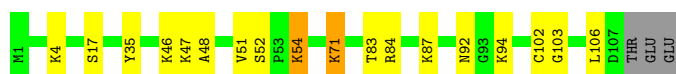
- Molecule 43: 50S ribosomal protein L24

Chain RY: 78% 18% ..



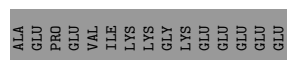
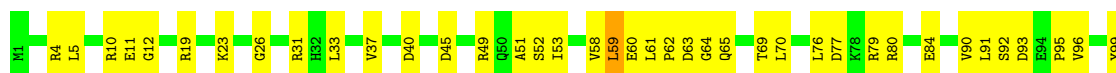
- Molecule 43: 50S ribosomal protein L24

Chain YY: 81% 15% ..



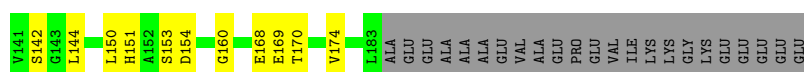
- Molecule 44: 50S ribosomal protein L25

Chain RZ: 54% 32% 11%



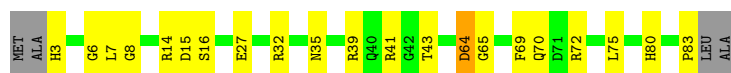
- Molecule 44: 50S ribosomal protein L25

Chain YZ: 67% 21% 11%



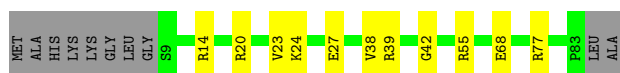
- Molecule 45: 50S ribosomal protein L27

Chain R0: 71% 24% 5%



- Molecule 45: 50S ribosomal protein L27

Chain Y0: 75% 13% 12%



- Molecule 46: 50S ribosomal protein L28

Chain R1: 79% 16%



- Molecule 46: 50S ribosomal protein L28

Chain Y1: 72% 20% 5%



- Molecule 47: 50S ribosomal protein L29

Chain R2: 72% 24%



- Molecule 47: 50S ribosomal protein L29

Chain Y2: 69% 22% 8%



- Molecule 48: 50S ribosomal protein L30

Chain R3: 72% 27%



- Molecule 48: 50S ribosomal protein L30

Chain Y3: 78% 20%

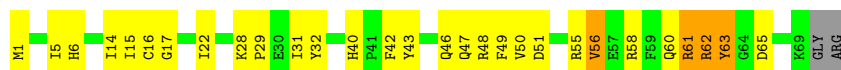


- Molecule 49: 50S ribosomal protein L31

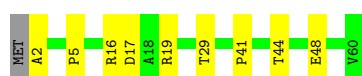
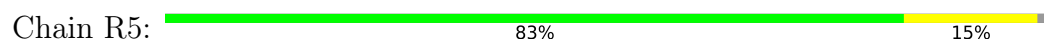
Chain R4: 58% 37%



- Molecule 49: 50S ribosomal protein L31



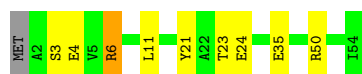
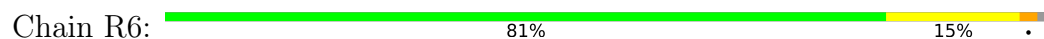
- Molecule 50: 50S ribosomal protein L32



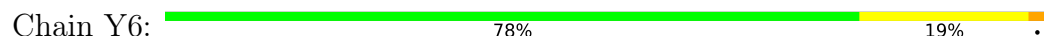
- Molecule 50: 50S ribosomal protein L32



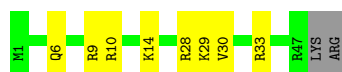
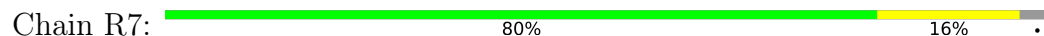
- Molecule 51: 50S ribosomal protein L33



- Molecule 51: 50S ribosomal protein L33



- Molecule 52: 50S ribosomal protein L34



- Molecule 52: 50S ribosomal protein L34





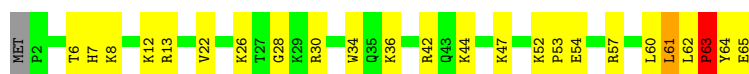
- Molecule 53: 50S ribosomal protein L35

Chain R8: 60% 38%



- Molecule 53: 50S ribosomal protein L35

Chain Y8: 62% 34%



- Molecule 54: 50S ribosomal protein L36

Chain R9: 78% 22%



- Molecule 54: 50S ribosomal protein L36

Chain Y9: 70% 30%



4 Data and refinement statistics

EDS failed to run properly - this section is therefore incomplete.

Property	Value	Source
Space group	P 21 21 21	Depositor
Cell constants a, b, c, α , β , γ	210.17Å 451.47Å 620.21Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	152.46 – 3.20	Depositor
% Data completeness (in resolution range)	99.1 (152.46-3.20)	Depositor
R_{merge}	0.26	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.69 (at 3.19Å)	Xtriage
Refinement program	PHENIX 1.14_3260	Depositor
R, R_{free}	0.224 , 0.256	Depositor
Wilson B-factor (Å ²)	71.2	Xtriage
Anisotropy	0.167	Xtriage
L-test for twinning ²	$\langle L \rangle = 0.42$, $\langle L^2 \rangle = 0.25$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
Total number of atoms	291964	wwPDB-VP
Average B, all atoms (Å ²)	103.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.60% 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, 0.333 respectively for untwinned datasets, and 0.375, 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: MG, ZN, 1MG, SF4

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	QA	0.96	1/36098 (0.0%)	1.22	232/56341 (0.4%)
1	XA	0.95	3/36101 (0.0%)	1.23	252/56346 (0.4%)
2	QB	0.42	0/1942	0.63	0/2619
2	XB	0.42	0/1950	0.57	0/2630
3	QC	0.38	0/1629	0.62	1/2195 (0.0%)
3	XC	0.37	0/1629	0.58	0/2195
4	QD	0.52	0/1733	0.58	0/2318
4	XD	0.54	1/1733 (0.1%)	0.64	0/2318
5	QE	0.45	0/1171	0.62	0/1576
5	XE	0.49	0/1171	0.57	0/1576
6	QF	0.45	0/856	0.59	0/1154
6	XF	0.55	0/856	0.54	0/1154
7	QG	0.37	0/1276	0.55	0/1709
7	XG	0.37	0/1276	0.54	0/1709
8	QH	0.51	0/1128	0.61	0/1517
8	XH	0.54	0/1128	0.62	0/1517
9	QI	0.34	0/1029	0.64	1/1379 (0.1%)
9	XI	0.39	0/1017	0.64	0/1365
10	QJ	0.36	0/814	0.62	0/1095
10	XJ	0.34	0/790	0.60	0/1063
11	QK	0.47	0/900	0.61	0/1213
11	XK	0.49	0/879	0.54	0/1187
12	QL	0.62	0/991	0.61	0/1327
12	XL	0.57	0/972	0.64	1/1301 (0.1%)
13	QM	0.37	0/965	0.64	0/1292
13	XM	0.33	0/956	0.65	0/1281
14	QN	0.43	0/501	0.70	1/664 (0.2%)
14	XN	0.44	0/501	0.63	1/664 (0.2%)
15	QO	0.46	0/745	0.59	0/992
15	XO	0.49	0/740	0.52	0/987
16	QP	0.55	0/721	0.60	0/970
16	XP	0.46	0/721	0.60	0/970

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
17	QQ	0.54	0/847	0.60	0/1131
17	XQ	0.55	0/847	0.58	0/1131
18	QR	0.46	0/579	0.64	0/768
18	XR	0.49	0/579	0.60	0/768
19	QS	0.36	0/680	0.66	0/915
19	XS	0.35	0/689	0.65	0/926
20	QT	0.43	0/765	0.57	0/1007
20	XT	0.36	0/765	0.58	0/1007
21	QU	0.39	0/221	0.88	1/288 (0.3%)
21	XU	0.34	0/221	0.58	0/288
22	QV	0.62	1/1814 (0.1%)	1.10	6/2825 (0.2%)
22	XV	0.57	1/1814 (0.1%)	1.21	8/2825 (0.3%)
23	QX	0.46	0/470	1.00	1/733 (0.1%)
23	XX	0.52	0/470	1.22	6/733 (0.8%)
24	RA	1.32	53/69498 (0.1%)	1.31	773/108491 (0.7%)
24	YA	1.59	262/69543 (0.4%)	1.39	981/108563 (0.9%)
25	RB	0.92	0/2878	1.24	26/4490 (0.6%)
25	YB	1.86	31/2878 (1.1%)	6.09	81/4490 (1.8%)
26	RD	0.75	1/2185 (0.0%)	0.62	0/2944
26	YD	0.87	1/2185 (0.0%)	0.67	0/2944
27	RE	0.71	0/1601	0.61	0/2160
27	YE	0.83	1/1596 (0.1%)	0.66	0/2153
28	RF	0.68	0/1620	0.61	0/2194
28	YF	0.85	0/1620	0.63	0/2194
29	RG	0.41	0/1499	0.64	0/2016
29	YG	0.51	0/1499	0.63	1/2016 (0.0%)
30	RH	0.48	0/1362	0.65	1/1841 (0.1%)
30	YH	0.72	0/1356	0.59	0/1833
31	RI	0.44	0/1151	0.69	1/1558 (0.1%)
31	YI	0.50	0/1151	0.70	1/1558 (0.1%)
32	RN	0.60	0/1131	0.59	0/1525
32	YN	0.82	1/1148 (0.1%)	0.67	0/1547
33	RO	0.73	0/943	0.61	0/1269
33	YO	0.82	1/943 (0.1%)	0.66	1/1269 (0.1%)
34	RP	0.61	0/1156	0.62	0/1537
34	YP	0.74	0/1139	0.64	0/1514
35	RQ	0.60	0/1143	0.63	0/1527
35	YQ	0.82	0/1143	0.59	0/1527
36	RR	0.67	0/974	0.64	1/1302 (0.1%)
36	YR	0.77	0/974	0.72	0/1302
37	RS	0.49	0/892	0.73	1/1187 (0.1%)
37	YS	0.68	0/887	0.58	0/1180
38	RT	0.62	0/1155	0.66	1/1542 (0.1%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
38	YT	0.70	0/1155	0.67	1/1542 (0.1%)
39	RU	0.68	0/982	0.61	0/1306
39	YU	0.93	0/982	0.62	0/1306
40	RV	0.62	0/790	0.61	0/1057
40	YV	0.86	0/790	0.67	0/1057
41	RW	0.69	0/911	0.61	0/1220
41	YW	0.89	0/911	0.64	0/1220
42	RX	0.67	0/739	0.61	0/993
42	YX	0.83	0/756	0.57	0/1014
43	RY	0.60	0/831	0.58	0/1108
43	YY	0.74	0/831	0.60	0/1108
44	RZ	0.48	0/1493	0.71	2/2026 (0.1%)
44	YZ	0.63	0/1493	0.57	0/2026
45	R0	0.60	0/652	0.60	0/867
45	Y0	0.84	0/607	0.63	0/809
46	R1	0.67	0/770	0.65	0/1022
46	Y1	0.78	0/736	0.65	0/978
47	R2	0.47	0/583	0.52	0/771
47	Y2	0.65	0/560	0.61	0/741
48	R3	0.62	1/474 (0.2%)	0.61	0/635
48	Y3	0.73	0/474	0.67	0/635
49	R4	0.39	0/578	0.64	0/776
49	Y4	0.41	0/578	0.68	0/776
50	R5	0.71	0/473	0.58	0/639
50	Y5	0.91	1/473 (0.2%)	0.65	1/639 (0.2%)
51	R6	0.62	0/460	0.65	1/613 (0.2%)
51	Y6	0.75	0/460	0.65	1/613 (0.2%)
52	R7	0.74	0/417	0.57	0/550
52	Y7	0.91	0/426	0.61	0/561
53	R8	0.68	0/525	0.71	0/691
53	Y8	0.77	0/525	0.76	1/691 (0.1%)
54	R9	0.62	0/310	0.53	0/407
54	Y9	0.84	0/310	0.55	0/407
All	All	1.14	360/315985 (0.1%)	1.30	2387/472446 (0.5%)

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
1	XA	0	1

Continued on next page...

Continued from previous page...

Mol	Chain	#Chirality outliers	#Planarity outliers
2	QB	0	22
2	XB	0	15
3	QC	0	15
3	XC	0	9
4	QD	0	12
4	XD	0	15
5	QE	0	7
5	XE	0	6
6	QF	0	5
6	XF	0	3
7	QG	0	9
7	XG	0	10
8	QH	0	3
8	XH	0	5
9	QI	0	15
9	XI	0	16
10	QJ	0	12
10	XJ	0	8
11	QK	0	5
11	XK	0	4
12	QL	0	8
12	XL	0	11
13	QM	0	13
13	XM	0	18
14	QN	0	6
14	XN	0	2
15	QO	0	1
15	XO	0	2
16	QP	0	5
16	XP	0	10
17	QQ	0	3
17	XQ	0	1
18	QR	0	5
18	XR	0	6
19	QS	0	13
19	XS	0	14
20	QT	0	10
20	XT	0	12
21	XU	0	4
24	YA	0	2
26	RD	0	13
26	YD	0	3

Continued on next page...

Continued from previous page...

Mol	Chain	#Chirality outliers	#Planarity outliers
27	RE	0	7
27	YE	0	10
28	RF	0	12
28	YF	0	15
29	RG	0	17
29	YG	0	20
30	RH	0	23
30	YH	0	1
31	RI	0	17
31	YI	0	21
32	RN	0	10
32	YN	0	7
33	RO	0	2
33	YO	0	1
34	RP	0	8
34	YP	0	5
35	RQ	0	18
35	YQ	0	6
36	RR	0	4
36	YR	0	5
37	RS	0	11
37	YS	0	1
38	RT	0	15
38	YT	0	11
39	RU	0	5
39	YU	0	5
40	RV	0	6
40	YV	0	7
41	RW	0	5
41	YW	0	5
42	RX	0	2
42	YX	0	1
43	RY	0	7
43	YY	0	5
44	RZ	0	38
44	YZ	0	10
45	R0	0	2
45	Y0	0	1
46	R1	0	10
46	Y1	0	7
47	R2	0	5
47	Y2	0	6

Continued on next page...

Continued from previous page...

Mol	Chain	#Chirality outliers	#Planarity outliers
48	R3	0	1
48	Y3	0	2
49	R4	0	15
49	Y4	0	11
50	R5	0	1
52	R7	0	1
53	R8	0	11
53	Y8	0	6
All	All	0	781

All (360) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
25	YB	112	G	N9-C4	30.70	1.62	1.38
25	YB	114	G	N9-C4	28.24	1.60	1.38
25	YB	112	G	N9-C8	-27.57	1.18	1.37
25	YB	114	G	N9-C8	-26.21	1.19	1.37
25	YB	112	G	C8-N7	14.77	1.39	1.30
25	YB	112	G	N3-C4	14.68	1.45	1.35
25	YB	114	G	C8-N7	14.13	1.39	1.30
24	RA	1649	A	C5'-C4'	13.69	1.67	1.51
25	YB	114	G	N3-C4	13.38	1.44	1.35
25	YB	114	G	N7-C5	-10.72	1.32	1.39
22	XV	1	C	OP3-P	-10.59	1.48	1.61
22	QV	1	C	OP3-P	-10.44	1.48	1.61
25	YB	96	G	N3-C4	-9.85	1.28	1.35
25	YB	112	G	N7-C5	-9.33	1.33	1.39
25	YB	92	G	N3-C4	-9.17	1.29	1.35
26	YD	264	LYS	C-N	9.02	1.51	1.34
4	XD	196	LEU	C-N	8.92	1.51	1.34
24	YA	1067	A	N9-C4	-8.68	1.32	1.37
1	QA	1528	U	C5'-C4'	8.54	1.61	1.51
25	YB	92	G	C2-N3	-8.47	1.25	1.32
25	YB	104	A	N9-C4	-8.37	1.32	1.37
24	YA	1642	A	N9-C4	-8.06	1.33	1.37
24	YA	2725	A	N9-C8	-8.05	1.31	1.37
24	YA	723	A	N9-C4	-7.96	1.33	1.37
24	RA	1649	A	C1'-N9	7.83	1.60	1.48
24	YA	830	A	N9-C4	-7.78	1.33	1.37
24	YA	1188	A	N9-C4	-7.72	1.33	1.37
25	YB	112	G	C5-C4	-7.69	1.32	1.38
24	YA	2083	G	N7-C5	-7.67	1.34	1.39

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
25	YB	114	G	C5-C4	-7.57	1.33	1.38
24	RA	1649	A	O4'-C1'	7.50	1.51	1.41
24	YA	733	G	C6-N1	-7.39	1.34	1.39
24	YA	553	A	N9-C4	-7.32	1.33	1.37
25	YB	96	G	C2-N3	-7.28	1.26	1.32
24	YA	1814	A	N9-C4	-7.27	1.33	1.37
24	YA	138	G	N3-C4	-7.23	1.30	1.35
24	YA	2083	G	N9-C8	-7.20	1.32	1.37
24	RA	1817	A	N7-C5	-7.16	1.34	1.39
1	XA	358	U	C4'-C3'	7.14	1.61	1.53
25	YB	96	G	C2-N2	-7.09	1.27	1.34
24	YA	354	A	N9-C4	-7.06	1.33	1.37
24	YA	70	A	N9-C4	-7.02	1.33	1.37
24	RA	1188	A	N9-C4	-6.98	1.33	1.37
25	YB	73	A	N7-C5	-6.98	1.35	1.39
24	YA	1817	A	N7-C5	-6.84	1.35	1.39
24	RA	1921	G	N9-C4	-6.80	1.32	1.38
24	YA	2073	A	N9-C4	-6.65	1.33	1.37
24	YA	140	A	N7-C5	-6.63	1.35	1.39
24	RA	1067	A	N9-C4	-6.63	1.33	1.37
25	YB	81	G	C5-C6	-6.61	1.35	1.42
24	YA	723	A	N3-C4	-6.61	1.30	1.34
24	YA	2037	A	N7-C5	-6.58	1.35	1.39
24	YA	125	A	N9-C4	-6.58	1.33	1.37
24	YA	1177	G	N7-C5	-6.54	1.35	1.39
24	YA	1188	A	N3-C4	-6.53	1.30	1.34
24	YA	718	C	N1-C6	-6.52	1.33	1.37
24	YA	590	A	N9-C4	-6.47	1.33	1.37
24	YA	646	A	N9-C4	-6.45	1.33	1.37
25	YB	96	G	N9-C4	-6.43	1.32	1.38
24	YA	776	G	C2-N3	-6.43	1.27	1.32
24	RA	1649	A	C4'-O4'	6.42	1.53	1.45
24	YA	48	A	N7-C5	-6.40	1.35	1.39
24	YA	1067	A	N3-C4	-6.38	1.31	1.34
24	YA	1617	A	N9-C4	-6.37	1.34	1.37
25	YB	112	G	N1-C2	-6.37	1.32	1.37
24	YA	1665	G	N7-C5	-6.36	1.35	1.39
25	YB	85	G	C2-N3	-6.33	1.27	1.32
24	YA	1378	G	C6-N1	-6.26	1.35	1.39
24	YA	738	C	N1-C6	-6.25	1.33	1.37
24	YA	2462	A	N7-C5	-6.25	1.35	1.39
24	YA	829	A	N9-C4	-6.24	1.34	1.37

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	YA	553	A	N7-C5	-6.23	1.35	1.39
24	YA	1027	A	N9-C4	-6.23	1.34	1.37
24	YA	800	C	C4-C5	-6.19	1.38	1.43
24	YA	1463	C	N1-C6	-6.18	1.33	1.37
24	YA	610	C	N1-C6	-6.16	1.33	1.37
24	YA	606	G	N7-C5	-6.16	1.35	1.39
24	RA	2726	A	N9-C4	-6.16	1.34	1.37
24	RA	1921	G	N3-C4	-6.15	1.31	1.35
24	YA	2464	C	N1-C6	-6.10	1.33	1.37
24	YA	73	A	N7-C5	-6.09	1.35	1.39
24	YA	1378	G	N7-C5	-6.07	1.35	1.39
24	YA	1643	A	N9-C4	-6.07	1.34	1.37
25	YB	114	G	C6-N1	-6.06	1.35	1.39
24	RA	1310	G	N7-C5	-6.05	1.35	1.39
24	YA	1067	A	N7-C5	-6.03	1.35	1.39
24	YA	2530	A	N9-C4	-6.02	1.34	1.37
24	YA	829	A	N7-C5	-6.00	1.35	1.39
24	RA	1378	G	N7-C5	-6.00	1.35	1.39
24	YA	2299	A	N9-C4	-5.99	1.34	1.37
24	YA	579	G	N7-C5	-5.98	1.35	1.39
24	YA	2618	C	N1-C6	-5.97	1.33	1.37
24	YA	745	C	N1-C6	-5.96	1.33	1.37
24	YA	593	G	N7-C5	-5.95	1.35	1.39
25	YB	92	G	N9-C4	-5.95	1.33	1.38
24	YA	553	A	N3-C4	-5.94	1.31	1.34
24	YA	733	G	N1-C2	-5.93	1.33	1.37
24	YA	723	A	N7-C5	-5.89	1.35	1.39
24	YA	112	U	C4-C5	-5.89	1.38	1.43
24	YA	849	A	N7-C5	-5.87	1.35	1.39
24	YA	2018	C	N1-C6	-5.86	1.33	1.37
24	YA	538	A	N7-C5	-5.86	1.35	1.39
24	YA	821	A	N9-C4	-5.86	1.34	1.37
24	YA	830	A	N7-C5	-5.83	1.35	1.39
24	YA	2415	C	C4-C5	-5.83	1.38	1.43
24	YA	2633	A	C5-C4	-5.82	1.34	1.38
24	YA	2093	A	N9-C4	-5.82	1.34	1.37
24	YA	2068	G	N7-C5	-5.81	1.35	1.39
24	YA	2468	C	N1-C6	-5.80	1.33	1.37
24	RA	2554	A	N9-C4	-5.80	1.34	1.37
24	YA	2100	C	N1-C6	-5.80	1.33	1.37
24	YA	73	A	N3-C4	-5.79	1.31	1.34
25	YB	114	G	N1-C2	-5.78	1.33	1.37

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
50	Y5	6	VAL	CB-CG1	-5.77	1.40	1.52
24	YA	187	C	C4-C5	-5.76	1.38	1.43
24	YA	479	C	N1-C6	-5.74	1.33	1.37
24	YA	2072	C	N1-C6	-5.73	1.33	1.37
24	YA	1173	A	N9-C4	-5.71	1.34	1.37
24	YA	18	C	N1-C6	-5.71	1.33	1.37
24	YA	835	A	N9-C4	-5.71	1.34	1.37
24	YA	607	C	C4-C5	-5.70	1.38	1.43
24	RA	776	G	C2-N3	-5.70	1.28	1.32
24	RA	2530	A	N9-C4	-5.70	1.34	1.37
24	RA	1617	A	N9-C4	-5.68	1.34	1.37
24	YA	2455	C	N1-C6	-5.68	1.33	1.37
24	RA	831	A	N9-C4	-5.67	1.34	1.37
24	YA	1422	C	N1-C6	-5.66	1.33	1.37
24	YA	2074	G	N1-C2	-5.66	1.33	1.37
24	YA	2004	C	C4-C5	-5.66	1.38	1.43
24	YA	733	G	C5-C4	-5.65	1.34	1.38
24	YA	1618	A	N9-C4	-5.65	1.34	1.37
27	YE	184	VAL	CB-CG1	-5.65	1.41	1.52
24	YA	1835	C	C4-C5	-5.64	1.38	1.43
24	YA	596	G	C2-N3	-5.63	1.28	1.32
24	YA	2725	A	N7-C5	-5.63	1.35	1.39
24	YA	853	C	C4-C5	-5.62	1.38	1.43
24	YA	603	C	C4-C5	-5.62	1.38	1.43
24	YA	1027	A	N7-C5	-5.61	1.35	1.39
24	YA	2601	A	N9-C4	-5.60	1.34	1.37
24	YA	2550	C	N1-C6	-5.60	1.33	1.37
24	YA	2462	A	C6-N1	-5.59	1.31	1.35
1	XA	1502	A	N7-C5	-5.59	1.35	1.39
24	YA	32	C	N1-C6	-5.59	1.33	1.37
1	XA	358	U	O4'-C1'	-5.58	1.34	1.41
24	YA	1975	A	N9-C4	-5.58	1.34	1.37
24	YA	855	G	N7-C5	-5.56	1.35	1.39
24	RA	1921	G	C2-N3	-5.56	1.28	1.32
24	YA	1922	A	N7-C5	-5.56	1.35	1.39
24	YA	73	A	N9-C4	-5.55	1.34	1.37
24	YA	1822	A	N9-C4	-5.55	1.34	1.37
24	YA	827	G	N9-C8	-5.55	1.33	1.37
24	YA	2629	C	N1-C6	-5.55	1.33	1.37
24	YA	2468	C	C4-C5	-5.54	1.38	1.43
24	YA	841	G	N7-C5	-5.54	1.35	1.39
25	YB	85	G	N3-C4	-5.54	1.31	1.35

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	YA	1070	G	N7-C5	-5.52	1.35	1.39
24	YA	2298	A	N7-C5	-5.51	1.35	1.39
24	YA	2830	A	N7-C5	-5.51	1.35	1.39
24	YA	1725	G	N7-C5	-5.50	1.35	1.39
24	YA	1454	C	N1-C6	-5.50	1.33	1.37
24	YA	2700	U	N3-C4	-5.49	1.33	1.38
24	YA	828	A	N9-C4	-5.49	1.34	1.37
24	RA	2083	G	N9-C8	-5.49	1.34	1.37
24	RA	2004	C	C4-C5	-5.49	1.38	1.43
24	YA	1814	A	N3-C4	-5.49	1.31	1.34
24	YA	1020	C	N3-C4	-5.48	1.30	1.33
24	RA	783	C	C4-C5	-5.48	1.38	1.43
24	YA	2609	G	N7-C5	-5.48	1.35	1.39
24	YA	2618	C	C4-C5	-5.47	1.38	1.43
24	YA	1719	C	N1-C6	-5.46	1.33	1.37
24	YA	2066	C	C4-C5	-5.46	1.38	1.43
24	YA	196	A	C5-C4	-5.46	1.34	1.38
24	YA	847	A	N9-C4	-5.46	1.34	1.37
24	YA	1819	C	C4-C5	-5.46	1.38	1.43
24	YA	2442	A	N7-C5	-5.46	1.35	1.39
24	YA	825	G	N7-C5	-5.46	1.35	1.39
24	RA	1715	A	N3-C4	-5.46	1.31	1.34
24	YA	1804	A	C5-C4	-5.45	1.34	1.38
24	YA	2028	C	N1-C6	-5.45	1.33	1.37
24	YA	2737	C	N3-C4	-5.44	1.30	1.33
24	YA	1015	C	N1-C6	-5.44	1.33	1.37
24	YA	2087	C	N1-C6	-5.44	1.33	1.37
24	YA	736	A	N7-C5	-5.42	1.35	1.39
24	YA	2466	G	N7-C5	-5.42	1.35	1.39
24	RA	187	C	C4-C5	-5.42	1.38	1.43
24	RA	1855	G	N7-C5	-5.42	1.36	1.39
24	RA	1067	A	N3-C4	-5.41	1.31	1.34
24	YA	730	C	C4-C5	-5.41	1.38	1.43
24	YA	2737	C	N1-C6	-5.41	1.33	1.37
24	YA	596	G	C8-N7	-5.41	1.27	1.30
24	YA	1002	A	C5-C6	-5.40	1.36	1.41
24	YA	597	C	N1-C6	-5.40	1.33	1.37
24	YA	1828	C	N1-C6	-5.40	1.33	1.37
24	RA	723	A	N7-C5	-5.39	1.36	1.39
24	YA	2452	C	N1-C6	-5.38	1.33	1.37
24	RA	70	A	N9-C4	-5.38	1.34	1.37
24	RA	2285	A	N9-C4	-5.37	1.34	1.37

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	YA	138	G	N7-C5	-5.36	1.36	1.39
24	YA	2082	A	N7-C5	-5.36	1.36	1.39
24	YA	2419	G	N7-C5	-5.36	1.36	1.39
24	YA	596	G	N7-C5	-5.35	1.36	1.39
24	YA	2345	A	C5-C4	-5.35	1.35	1.38
24	RA	1835	C	C4-C5	-5.35	1.38	1.43
24	YA	354	A	N3-C4	-5.35	1.31	1.34
24	YA	1475	G	N7-C5	-5.34	1.36	1.39
24	RA	609	A	N9-C4	-5.34	1.34	1.37
24	YA	140	A	N9-C4	-5.32	1.34	1.37
24	YA	730	C	N3-C4	-5.32	1.30	1.33
25	YB	81	G	N1-C2	-5.32	1.33	1.37
25	YB	81	G	C8-N7	-5.32	1.27	1.30
24	RA	1804	A	N7-C5	-5.32	1.36	1.39
24	YA	1817	A	N3-C4	-5.31	1.31	1.34
24	YA	1814	A	N7-C5	-5.31	1.36	1.39
25	YB	96	G	N7-C5	-5.30	1.36	1.39
32	YN	106	MET	C-N	-5.30	1.21	1.34
24	YA	989	G	N7-C5	-5.30	1.36	1.39
24	YA	2586	G	N7-C5	-5.30	1.36	1.39
24	RA	2037	A	N7-C5	-5.29	1.36	1.39
24	YA	733	G	N7-C5	-5.29	1.36	1.39
24	YA	552	C	N1-C6	-5.29	1.33	1.37
24	YA	535	C	C4-C5	-5.29	1.38	1.43
24	YA	1710	C	N1-C6	-5.28	1.33	1.37
24	YA	2697	G	N9-C4	-5.28	1.33	1.38
24	YA	733	G	C6-O6	-5.28	1.19	1.24
24	YA	190	C	C4-C5	-5.28	1.38	1.43
24	YA	1654	A	N7-C5	-5.28	1.36	1.39
24	YA	139	A	N7-C5	-5.27	1.36	1.39
24	YA	1646	C	N1-C6	-5.27	1.33	1.37
24	YA	2831	A	N7-C5	-5.27	1.36	1.39
24	YA	495	G	C6-N1	-5.27	1.35	1.39
24	YA	2042	A	N7-C5	-5.27	1.36	1.39
24	YA	832	G	C5-C4	-5.27	1.34	1.38
24	YA	117	A	N9-C4	-5.27	1.34	1.37
24	RA	2052	A	N9-C4	-5.26	1.34	1.37
24	YA	596	G	C5-C4	-5.26	1.34	1.38
24	RA	552	C	N1-C6	-5.26	1.33	1.37
24	YA	2002	G	C5-C4	-5.26	1.34	1.38
24	YA	211	A	C5'-C4'	5.26	1.57	1.51
24	YA	2459	G	N9-C8	-5.26	1.34	1.37

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	YA	2359	C	N1-C6	-5.25	1.33	1.37
24	RA	1188	A	N3-C4	-5.25	1.31	1.34
24	YA	2394	G	N7-C5	-5.25	1.36	1.39
24	YA	594	A	C5-C4	-5.25	1.35	1.38
24	YA	2790	G	N7-C5	-5.25	1.36	1.39
24	RA	2032	G	N7-C5	-5.25	1.36	1.39
24	YA	1310	G	N7-C5	-5.25	1.36	1.39
33	YO	70	LYS	C-N	-5.25	1.22	1.34
24	YA	2521	G	N7-C5	-5.24	1.36	1.39
24	YA	2054	G	N9-C4	-5.24	1.33	1.38
24	YA	829	A	C5-C4	-5.24	1.35	1.38
24	YA	2065	C	C4-C5	-5.24	1.38	1.43
24	YA	2083	G	C8-N7	-5.24	1.27	1.30
24	RA	73	A	N7-C5	-5.23	1.36	1.39
24	RA	1828	C	N1-C6	-5.23	1.34	1.37
24	YA	140	A	C5-C6	-5.23	1.36	1.41
24	YA	197	C	C4-C5	-5.23	1.38	1.43
24	YA	732	A	C5-C4	-5.23	1.35	1.38
24	YA	2633	A	N7-C5	-5.23	1.36	1.39
24	YA	2594	G	N7-C5	-5.22	1.36	1.39
24	YA	198	C	C4-C5	-5.22	1.38	1.43
24	YA	2371	C	N3-C4	-5.22	1.30	1.33
24	YA	795	G	N7-C5	-5.22	1.36	1.39
24	YA	2460	A	N7-C5	-5.22	1.36	1.39
24	YA	2028	C	C4-C5	-5.21	1.38	1.43
25	YB	7	G	N7-C5	-5.21	1.36	1.39
26	RD	255	LYS	CA-CB	-5.21	1.42	1.53
24	YA	1255	A	N7-C5	-5.21	1.36	1.39
24	YA	1272	A	N9-C4	-5.21	1.34	1.37
24	YA	1016	C	C4-C5	-5.21	1.38	1.43
24	YA	2075	G	N9-C8	-5.21	1.34	1.37
24	YA	2421	G	N7-C5	-5.20	1.36	1.39
24	YA	20	C	C4-C5	-5.20	1.38	1.43
24	YA	113	C	N1-C6	-5.20	1.34	1.37
24	YA	1231	G	N7-C5	-5.20	1.36	1.39
24	YA	2633	A	N9-C4	-5.19	1.34	1.37
24	RA	830	A	N7-C5	-5.19	1.36	1.39
24	YA	2028	C	N3-C4	-5.18	1.30	1.33
24	YA	2079	A	C5-C4	-5.18	1.35	1.38
24	YA	68	C	N3-C4	-5.18	1.30	1.33
24	YA	596	G	N9-C8	-5.18	1.34	1.37
24	YA	444	C	C4-C5	-5.17	1.38	1.43

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	YA	2285	A	N7-C5	-5.17	1.36	1.39
24	YA	1255	A	C5-C6	-5.17	1.36	1.41
24	RA	2083	G	N7-C5	-5.17	1.36	1.39
24	RA	830	A	N9-C4	-5.17	1.34	1.37
24	YA	579	G	N9-C8	-5.17	1.34	1.37
24	RA	2036	A	N9-C4	-5.16	1.34	1.37
24	YA	1332	A	N9-C4	-5.16	1.34	1.37
24	YA	178	G	C6-N1	-5.16	1.35	1.39
24	YA	825	G	N9-C8	-5.16	1.34	1.37
24	YA	2347	A	C5-C4	-5.16	1.35	1.38
24	YA	2070	G	N7-C5	-5.15	1.36	1.39
24	YA	801	C	C4-C5	-5.15	1.38	1.43
24	YA	2631	C	N1-C6	-5.15	1.34	1.37
24	YA	480	A	N7-C5	-5.14	1.36	1.39
24	YA	732	A	N3-C4	-5.14	1.31	1.34
24	YA	1574	A	N7-C5	-5.14	1.36	1.39
24	YA	2511	C	C4-C5	-5.13	1.38	1.43
24	YA	1313	U	C4-C5	-5.13	1.39	1.43
24	YA	2697	G	N3-C4	-5.12	1.31	1.35
24	YA	226	C	C4-C5	-5.12	1.38	1.43
24	YA	640	A	N9-C4	-5.12	1.34	1.37
24	YA	119	G	N7-C5	-5.12	1.36	1.39
24	YA	237	G	N7-C5	-5.12	1.36	1.39
24	YA	1666	G	N9-C8	-5.12	1.34	1.37
24	YA	2462	A	C5-C6	-5.12	1.36	1.41
24	YA	2055	A	N9-C4	-5.12	1.34	1.37
24	YA	46	C	C4-C5	-5.11	1.38	1.43
24	YA	126	C	C4-C5	-5.10	1.38	1.43
24	YA	811	A	C5-C4	-5.10	1.35	1.38
24	YA	1234	A	N7-C5	-5.10	1.36	1.39
24	YA	1170	C	N1-C6	-5.10	1.34	1.37
24	YA	2297	C	N1-C6	-5.10	1.34	1.37
24	RA	479	C	N1-C6	-5.09	1.34	1.37
24	YA	2070	G	N9-C8	-5.09	1.34	1.37
24	RA	28	A	N7-C5	-5.09	1.36	1.39
24	YA	1706	U	C2-N3	-5.09	1.34	1.37
24	RA	1336	C	N1-C6	-5.08	1.34	1.37
24	RA	2460	A	N9-C4	-5.07	1.34	1.37
48	R3	39	ASP	C-N	-5.07	1.22	1.34
24	YA	1644	C	C4-C5	-5.07	1.38	1.43
24	RA	237	G	N7-C5	-5.07	1.36	1.39
24	YA	1645	C	C4-C5	-5.07	1.38	1.43

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	RA	2593	G	C6-N1	-5.06	1.36	1.39
24	YA	604	C	C4-C5	-5.06	1.39	1.43
24	RA	2042	A	N7-C5	-5.06	1.36	1.39
24	YA	797	A	N7-C5	-5.06	1.36	1.39
24	YA	1354	A	N7-C5	-5.06	1.36	1.39
24	YA	842	C	N3-C4	-5.05	1.30	1.33
24	YA	2077	C	C4-C5	-5.05	1.39	1.43
24	RA	31	C	N1-C6	-5.05	1.34	1.37
24	YA	184	A	N9-C4	-5.05	1.34	1.37
24	YA	1657	C	N1-C6	-5.05	1.34	1.37
24	YA	2067	C	C4-C5	-5.05	1.39	1.43
24	YA	2080	A	N7-C5	-5.05	1.36	1.39
24	YA	2389	A	N9-C4	-5.05	1.34	1.37
24	YA	2530	A	N7-C5	-5.05	1.36	1.39
24	YA	2794	A	N9-C4	-5.05	1.34	1.37
24	RA	2509	A	N9-C4	-5.04	1.34	1.37
24	YA	1141	A	N9-C4	5.04	1.40	1.37
24	YA	2297	C	C4-C5	-5.04	1.39	1.43
24	YA	475	A	C5-C4	-5.04	1.35	1.38
24	YA	709	G	N9-C8	-5.04	1.34	1.37
24	RA	609	A	N3-C4	-5.04	1.31	1.34
24	YA	1320	A	N7-C5	-5.04	1.36	1.39
24	YA	2728	C	N1-C6	-5.03	1.34	1.37
24	YA	1388	A	C5-C4	-5.03	1.35	1.38
24	YA	2083	G	C5-C4	-5.03	1.34	1.38
24	YA	2091	G	C6-N1	-5.03	1.36	1.39
24	YA	1423	G	N7-C5	-5.03	1.36	1.39
24	YA	1661	C	N1-C6	-5.03	1.34	1.37
24	YA	545	G	C5-C4	-5.03	1.34	1.38
24	YA	2757	G	N7-C5	-5.02	1.36	1.39
24	YA	2270	C	N1-C6	-5.02	1.34	1.37
24	RA	596	G	N7-C5	-5.02	1.36	1.39
24	YA	1247	C	N1-C6	-5.02	1.34	1.37
24	YA	1658	C	C4-C5	-5.01	1.39	1.43
24	YA	556	C	N1-C6	-5.01	1.34	1.37
24	YA	1300	A	N9-C4	-5.00	1.34	1.37
24	YA	2440	G	C5-C4	-5.00	1.34	1.38

All (2387) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
25	YB	114	G	C4-C5-N7	-176.50	40.20	110.80

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
25	YB	112	G	C4-C5-N7	-173.34	41.46	110.80
25	YB	114	G	N7-C8-N9	-148.48	38.86	113.10
25	YB	112	G	N7-C8-N9	-146.23	39.98	113.10
25	YB	112	G	C8-N9-C4	-129.76	54.49	106.40
25	YB	114	G	C8-N9-C4	-123.21	57.12	106.40
25	YB	114	G	C5-N7-C8	70.62	139.61	104.30
25	YB	112	G	C5-N7-C8	62.30	135.45	104.30
25	YB	112	G	C6-C5-N7	47.29	158.77	130.40
25	YB	114	G	C6-C5-N7	45.05	157.43	130.40
25	YB	114	G	N3-C4-N9	32.61	145.57	126.00
25	YB	112	G	N3-C4-N9	31.24	144.75	126.00
25	YB	114	G	N9-C4-C5	-28.68	93.93	105.40
25	YB	114	G	C4-N9-C1'	26.52	160.98	126.50
25	YB	112	G	C4-N9-C1'	23.20	156.66	126.50
25	YB	112	G	N9-C4-C5	-22.50	96.40	105.40
1	QA	1528	U	C2-N1-C1'	22.40	144.58	117.70
25	YB	112	G	N3-C4-C5	-19.61	118.79	128.60
1	QA	1528	U	C6-N1-C1'	-19.51	93.89	121.20
25	YB	92	G	N3-C2-N2	-19.36	106.35	119.90
25	YB	112	G	C8-N9-C1'	-19.05	102.23	127.00
1	XA	359	U	O5'-P-OP1	-18.75	88.19	110.70
25	YB	114	G	C8-N9-C1'	-17.79	103.88	127.00
25	YB	114	G	N3-C4-C5	-16.20	120.50	128.60
24	RA	1649	A	O4'-C1'-N9	15.95	120.96	108.20
1	XA	359	U	C2-N1-C1'	-15.44	99.17	117.70
1	XA	359	U	C6-N1-C1'	14.99	142.19	121.20
25	YB	96	G	N3-C2-N2	-14.75	109.57	119.90
24	YA	903	C	C6-N1-C2	-14.53	114.49	120.30
25	YB	112	G	C5-C6-N1	14.46	118.73	111.50
25	YB	112	G	C2-N3-C4	13.99	118.90	111.90
24	YA	2700	U	N3-C2-O2	-13.88	112.48	122.20
24	RA	903	C	C6-N1-C2	-13.33	114.97	120.30
24	RA	1649	A	C5'-C4'-O4'	13.32	125.09	109.10
25	YB	96	G	C8-N9-C4	-13.06	101.18	106.40
24	YA	2774	G	C8-N9-C1'	-12.99	110.11	127.00
1	XA	359	U	OP1-P-OP2	-12.78	100.43	119.60
24	YA	1378	G	N7-C8-N9	12.68	119.44	113.10
24	YA	1581	U	N1-C2-O2	12.54	131.58	122.80
24	YA	1581	U	N3-C2-O2	-12.54	113.42	122.20
1	QA	1301	U	N1-C2-O2	12.44	131.51	122.80
24	YA	140	A	N7-C8-N9	12.28	119.94	113.80
24	YA	903	C	C5-C6-N1	11.95	126.97	121.00

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	2774	G	C4-N9-C1'	11.89	141.96	126.50
25	YB	96	G	N9-C4-C5	11.81	110.12	105.40
24	RA	1953	U	N3-C2-O2	-11.77	113.96	122.20
25	YB	85	G	N3-C2-N2	-11.76	111.67	119.90
1	XA	328	C	N1-C2-O2	11.68	125.91	118.90
24	RA	1649	A	C4-N9-C1'	11.66	147.29	126.30
1	QA	1301	U	N3-C2-O2	-11.53	114.13	122.20
1	XA	1158	C	N1-C2-O2	11.43	125.76	118.90
25	YB	96	G	N3-C4-N9	-11.34	119.20	126.00
1	QA	328	C	N1-C2-O2	11.28	125.67	118.90
24	YA	1627	A	C4-N9-C1'	11.25	146.56	126.30
25	YB	92	G	N1-C2-N2	11.21	126.29	116.20
25	YB	81	G	C4-C5-N7	11.11	115.25	110.80
24	YA	2087	C	C6-N1-C2	-11.09	115.87	120.30
24	RA	1649	A	C8-N9-C1'	-11.08	107.76	127.70
1	QA	1066	C	N1-C2-O2	11.02	125.51	118.90
24	RA	903	C	C5-C6-N1	10.86	126.43	121.00
24	RA	1817	A	N7-C8-N9	10.86	119.23	113.80
24	YA	800	C	C6-N1-C2	-10.84	115.97	120.30
24	YA	140	A	C5-N7-C8	-10.83	98.49	103.90
24	YA	907	U	N3-C2-O2	-10.76	114.67	122.20
24	YA	1003	U	C2-N1-C1'	-10.76	104.79	117.70
24	YA	118	U	N3-C2-O2	-10.74	114.68	122.20
24	YA	1627	A	C8-N9-C1'	-10.72	108.41	127.70
1	QA	1027	C	N1-C2-O2	10.71	125.33	118.90
1	XA	328	C	N3-C2-O2	-10.65	114.44	121.90
24	YA	1378	G	C8-N9-C4	-10.61	102.16	106.40
24	RA	1110	C	C6-N1-C2	-10.59	116.06	120.30
24	YA	1003	U	C6-N1-C1'	10.59	136.02	121.20
24	RA	1110	C	C5-C6-N1	10.54	126.27	121.00
24	RA	1138	C	N1-C2-O2	10.38	125.12	118.90
24	RA	875	U	N3-C2-O2	-10.36	114.95	122.20
25	YB	96	G	N1-C2-N3	10.28	130.07	123.90
1	QA	1066	C	C5-C6-N1	10.26	126.13	121.00
24	RA	1936	C	N1-C2-O2	10.26	125.06	118.90
24	RA	1921	G	N3-C4-N9	-10.26	119.84	126.00
24	YA	1725	G	N7-C8-N9	10.25	118.23	113.10
24	RA	1953	U	N1-C2-O2	10.14	129.90	122.80
25	RB	31	C	N1-C2-O2	10.06	124.94	118.90
1	XA	1158	C	N3-C2-O2	-10.06	114.86	121.90
24	RA	2818	U	N1-C2-O2	9.92	129.75	122.80
24	RA	875	U	N1-C2-O2	9.92	129.74	122.80

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	359	U	O5'-P-OP2	9.92	122.60	110.70
24	YA	1453	C	C5-C6-N1	9.89	125.94	121.00
24	RA	1282	G	C4-N9-C1'	9.88	139.34	126.50
1	QA	1158	C	N1-C2-O2	9.82	124.79	118.90
24	YA	2087	C	C5-C6-N1	9.79	125.89	121.00
24	RA	118	U	N3-C2-O2	-9.77	115.36	122.20
24	RA	1359	U	N3-C2-O2	-9.77	115.36	122.20
24	YA	2700	U	C5-C4-O4	9.71	131.72	125.90
24	RA	2700	U	N3-C2-O2	-9.68	115.43	122.20
25	YB	92	G	N9-C4-C5	9.67	109.27	105.40
24	YA	1476	C	C5-C6-N1	9.66	125.83	121.00
1	QA	328	C	N3-C2-O2	-9.62	115.17	121.90
24	RA	723	A	N7-C8-N9	9.61	118.60	113.80
24	YA	2825	C	C6-N1-C2	-9.60	116.46	120.30
24	YA	2065	C	C5-C6-N1	9.58	125.79	121.00
24	YA	675	C	C5-C6-N1	9.56	125.78	121.00
24	YA	1817	A	N7-C8-N9	9.51	118.55	113.80
24	YA	1579	C	C5-C6-N1	9.49	125.75	121.00
24	YA	553	A	C8-N9-C4	-9.49	102.00	105.80
24	RA	1310	G	C8-N9-C4	-9.48	102.61	106.40
1	XA	1141	C	N3-C2-O2	-9.47	115.27	121.90
24	YA	1378	G	C5-N7-C8	-9.42	99.59	104.30
24	YA	637	U	N3-C2-O2	-9.42	115.61	122.20
24	RA	1453	C	C6-N1-C2	-9.40	116.54	120.30
24	YA	2792	U	N3-C2-O2	-9.39	115.62	122.20
25	YB	85	G	N1-C2-N2	9.39	124.65	116.20
24	YA	800	C	C5-C6-N1	9.31	125.65	121.00
24	RA	2678	C	N1-C2-O2	9.30	124.48	118.90
24	YA	1453	C	C6-N1-C2	-9.30	116.58	120.30
24	RA	1817	A	C5-N7-C8	-9.28	99.26	103.90
24	YA	603	C	C5-C6-N1	9.27	125.63	121.00
1	QA	525	C	C5-C6-N1	9.24	125.62	121.00
24	RA	118	U	N1-C2-O2	9.23	129.26	122.80
24	YA	1835	C	C5-C6-N1	9.21	125.60	121.00
24	YA	1936	C	N1-C2-O2	9.20	124.42	118.90
24	YA	12	U	N3-C2-O2	-9.19	115.77	122.20
24	YA	2725	A	N7-C8-N9	9.17	118.39	113.80
24	RA	1281	G	C4-N9-C1'	9.17	138.42	126.50
24	YA	1935	A	C4-N9-C1'	9.16	142.80	126.30
1	XA	358	U	P-O3'-C3'	9.16	130.69	119.70
24	RA	1835	C	C5-C6-N1	9.14	125.57	121.00
25	YB	92	G	N3-C4-N9	-9.11	120.53	126.00

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	358	U	O4'-C1'-N1	-9.10	100.92	108.20
24	RA	2739	U	N3-C2-O2	-9.09	115.84	122.20
25	YB	81	G	N9-C4-C5	-9.09	101.77	105.40
24	RA	1128	U	N1-C2-O2	9.08	129.16	122.80
24	YA	939	C	N1-C2-O2	9.04	124.32	118.90
24	YA	1452	U	C5-C6-N1	9.02	127.21	122.70
1	QA	307	C	N1-C2-O2	9.02	124.31	118.90
24	YA	197	C	C5-C6-N1	9.01	125.50	121.00
24	YA	660	C	C6-N1-C2	-9.00	116.70	120.30
24	YA	675	C	C6-N1-C2	-9.00	116.70	120.30
24	YA	2739	U	N3-C2-O2	-8.98	115.91	122.20
24	YA	2065	C	C6-N1-C2	-8.97	116.71	120.30
24	YA	1903	C	C6-N1-C2	-8.96	116.72	120.30
24	RA	2230	U	N1-C2-O2	8.96	129.07	122.80
1	QA	1027	C	N3-C2-O2	-8.90	115.67	121.90
24	RA	907	U	N3-C2-O2	-8.90	115.97	122.20
24	RA	1281	G	C8-N9-C1'	-8.90	115.43	127.00
24	RA	1282	G	C8-N9-C1'	-8.88	115.45	127.00
1	XA	358	U	OP2-P-O3'	8.88	124.74	105.20
24	RA	1835	C	C6-N1-C2	-8.88	116.75	120.30
25	YB	30	C	C6-N1-C2	-8.88	116.75	120.30
24	YA	1020	C	N1-C2-O2	8.83	124.20	118.90
24	YA	1935	A	C8-N9-C1'	-8.83	111.81	127.70
24	RA	400	U	N3-C2-O2	-8.82	116.02	122.20
24	YA	1453	C	N1-C2-O2	8.81	124.19	118.90
1	QA	1066	C	C6-N1-C2	-8.78	116.79	120.30
1	XA	307	C	N1-C2-O2	8.77	124.16	118.90
24	YA	2596	U	N3-C2-O2	-8.76	116.07	122.20
24	RA	1921	G	N3-C4-C5	8.74	132.97	128.60
24	YA	875	U	N1-C2-O2	8.73	128.91	122.80
24	YA	140	A	C8-N9-C4	-8.73	102.31	105.80
24	YA	1110	C	C6-N1-C2	-8.73	116.81	120.30
25	YB	114	G	C5-C6-N1	8.72	115.86	111.50
24	YA	118	U	N1-C2-O2	8.71	128.90	122.80
24	RA	796	C	N1-C2-O2	8.70	124.12	118.90
24	RA	1359	U	N1-C2-O2	8.70	128.89	122.80
25	RB	31	C	C2-N1-C1'	8.68	128.35	118.80
24	YA	796	C	N1-C2-O2	8.65	124.09	118.90
24	YA	2792	U	N1-C2-O2	8.65	128.86	122.80
24	YA	1942	C	C5-C6-N1	8.61	125.31	121.00
24	RA	2818	U	N3-C2-O2	-8.60	116.18	122.20
1	XA	1502	A	N7-C8-N9	8.58	118.09	113.80

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	1453	C	C5-C6-N1	8.57	125.29	121.00
24	YA	258	U	N1-C2-O2	8.56	128.79	122.80
1	QA	1528	U	C5'-C4'-O4'	8.55	119.36	109.10
1	QA	1323	G	N3-C4-N9	8.54	131.12	126.00
24	RA	1095	C	C6-N1-C2	-8.54	116.88	120.30
24	RA	1805	C	N1-C2-O2	8.54	124.03	118.90
24	YA	1476	C	C6-N1-C2	-8.54	116.89	120.30
24	YA	2519	C	C5-C6-N1	8.54	125.27	121.00
24	YA	2468	C	C5-C6-N1	8.53	125.27	121.00
24	RA	1921	G	C2-N3-C4	-8.52	107.64	111.90
24	RA	187	C	C5-C6-N1	8.52	125.26	121.00
24	YA	1903	C	C5-C6-N1	8.51	125.26	121.00
24	YA	444	C	C6-N1-C2	-8.51	116.90	120.30
24	RA	1936	C	N3-C2-O2	-8.50	115.95	121.90
1	QA	328	C	C2-N1-C1'	8.50	128.15	118.80
25	YB	92	G	C8-N9-C4	-8.49	103.00	106.40
24	YA	197	C	C6-N1-C2	-8.48	116.91	120.30
24	RA	1550	C	C5-C6-N1	8.47	125.24	121.00
24	YA	1904	C	C5-C6-N1	8.47	125.24	121.00
1	XA	1158	C	C6-N1-C2	-8.47	116.91	120.30
24	RA	1138	C	C5-C6-N1	8.46	125.23	121.00
24	YA	1067	A	C8-N9-C4	-8.45	102.42	105.80
24	YA	555	G	N1-C6-O6	-8.45	114.83	119.90
24	YA	884	C	C6-N1-C2	-8.44	116.92	120.30
1	QA	1066	C	C2-N1-C1'	8.44	128.08	118.80
24	YA	1980	C	C6-N1-C2	-8.42	116.93	120.30
24	YA	554	A	C4-N9-C1'	8.40	141.43	126.30
24	YA	2700	U	N1-C2-O2	8.40	128.68	122.80
24	YA	1725	G	C8-N9-C4	-8.39	103.04	106.40
1	QA	1263	C	N1-C2-O2	8.38	123.93	118.90
24	YA	462	C	C6-N1-C2	-8.38	116.95	120.30
24	RA	2485	U	N3-C2-O2	-8.37	116.34	122.20
1	XA	328	C	C2-N1-C1'	8.37	128.01	118.80
24	YA	2319	G	C4-N9-C1'	8.37	137.38	126.50
24	YA	187	C	C5-C6-N1	8.34	125.17	121.00
24	YA	659	C	C6-N1-C2	-8.34	116.97	120.30
24	RA	2736	C	C6-N1-C2	-8.33	116.97	120.30
25	YB	68	C	C6-N1-C2	-8.33	116.97	120.30
24	RA	675	C	C6-N1-C2	-8.31	116.97	120.30
24	YA	1463	C	C5-C6-N1	8.30	125.15	121.00
25	YB	96	G	N7-C8-N9	8.30	117.25	113.10
1	QA	328	C	C6-N1-C2	-8.29	116.98	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	2230	U	N3-C2-O2	-8.29	116.40	122.20
24	RA	1378	G	C4-N9-C1'	8.29	137.27	126.50
1	QA	1158	C	N3-C2-O2	-8.28	116.10	121.90
24	RA	723	A	C5-N7-C8	-8.29	99.76	103.90
24	RA	400	U	C5-C6-N1	8.28	126.84	122.70
24	YA	1453	C	C2-N1-C1'	8.28	127.91	118.80
24	YA	1942	C	C6-N1-C2	-8.28	116.99	120.30
24	YA	1937	U	N1-C2-O2	8.27	128.59	122.80
24	YA	2428	C	C6-N1-C2	-8.27	116.99	120.30
24	RA	2004	C	C5-C6-N1	8.23	125.11	121.00
1	XA	328	C	C6-N1-C2	-8.22	117.01	120.30
24	YA	1952	G	C8-N9-C1'	8.20	137.66	127.00
24	YA	863	C	C6-N1-C2	-8.20	117.02	120.30
24	YA	1835	C	C6-N1-C2	-8.20	117.02	120.30
24	RA	400	U	N1-C2-O2	8.19	128.54	122.80
25	YB	96	G	C2-N3-C4	-8.17	107.81	111.90
24	YA	12	U	N1-C2-O2	8.17	128.52	122.80
1	XA	110	C	N1-C2-O2	8.17	123.80	118.90
24	RA	1453	C	C2-N1-C1'	8.15	127.77	118.80
24	YA	2468	C	C6-N1-C2	-8.15	117.04	120.30
22	XV	65	U	N1-C2-O2	8.14	128.50	122.80
24	RA	1378	G	N7-C8-N9	8.13	117.17	113.10
24	RA	618	C	C5-C6-N1	8.13	125.07	121.00
24	YA	112	U	C5-C6-N1	8.13	126.77	122.70
24	YA	1936	C	N3-C2-O2	-8.13	116.21	121.90
24	YA	1725	G	C5-N7-C8	-8.12	100.24	104.30
24	RA	1972	G	C4-N9-C1'	8.11	137.05	126.50
24	YA	887	C	C5-C6-N1	8.11	125.05	121.00
24	YA	1590	C	N1-C2-O2	8.11	123.76	118.90
24	YA	2678	C	N1-C2-O2	8.10	123.76	118.90
22	XV	65	U	N3-C2-O2	-8.09	116.54	122.20
24	RA	802	C	C5-C6-N1	8.09	125.05	121.00
1	QA	58	C	C6-N1-C2	-8.08	117.07	120.30
24	RA	1111	U	O4'-C1'-N1	8.07	114.66	108.20
24	RA	2768	C	C5-C6-N1	8.06	125.03	121.00
24	YA	1067	A	C2-N3-C4	-8.06	106.57	110.60
24	YA	1952	G	C4-N9-C1'	-8.05	116.03	126.50
24	RA	197	C	C5-C6-N1	8.03	125.01	121.00
24	YA	2690	C	C5-C6-N1	8.00	125.00	121.00
24	YA	2837	C	C5-C6-N1	8.00	125.00	121.00
24	RA	562	C	C5-C6-N1	8.00	125.00	121.00
24	RA	1817	A	C8-N9-C4	-8.00	102.60	105.80

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	1904	C	C6-N1-C2	-7.99	117.10	120.30
24	RA	2085	C	C6-N1-C2	-7.97	117.11	120.30
24	YA	562	C	C6-N1-C2	-7.97	117.11	120.30
24	YA	535	C	C5-C6-N1	7.97	124.99	121.00
24	RA	1805	C	C5-C6-N1	7.96	124.98	121.00
24	RA	1904	C	C6-N1-C2	-7.96	117.12	120.30
24	RA	1378	G	C6-C5-N7	-7.95	125.63	130.40
24	YA	875	U	N3-C2-O2	-7.95	116.63	122.20
24	RA	2571	C	C5-C6-N1	7.93	124.96	121.00
24	YA	103	C	C6-N1-C2	-7.92	117.13	120.30
24	YA	1310	G	C8-N9-C4	-7.92	103.23	106.40
24	YA	2571	C	C5-C6-N1	7.92	124.96	121.00
24	RA	1125	C	C6-N1-C2	-7.92	117.13	120.30
24	RA	1098	C	C6-N1-C2	-7.92	117.13	120.30
24	RA	1138	C	C6-N1-C2	-7.92	117.13	120.30
24	RA	1110	C	N1-C2-O2	7.91	123.65	118.90
24	YA	535	C	N1-C2-O2	7.90	123.64	118.90
1	QA	1303	C	N1-C2-O2	7.90	123.64	118.90
24	YA	2877	G	P-O3'-C3'	7.89	129.17	119.70
24	YA	1980	C	C5-C6-N1	7.88	124.94	121.00
25	YB	93	C	N1-C2-O2	7.88	123.63	118.90
38	YT	99	LEU	CA-CB-CG	7.88	133.43	115.30
24	YA	1110	C	C5-C6-N1	7.88	124.94	121.00
24	YA	1020	C	C2-N1-C1'	7.87	127.46	118.80
1	QA	1066	C	N3-C2-O2	-7.87	116.39	121.90
24	YA	2480	G	C4-N9-C1'	7.87	136.72	126.50
1	QA	1322	C	N1-C2-O2	7.86	123.62	118.90
1	QA	1347	G	C8-N9-C1'	-7.86	116.78	127.00
24	RA	2001	C	C6-N1-C2	-7.86	117.16	120.30
24	YA	884	C	N3-C2-O2	-7.85	116.40	121.90
24	RA	1969	C	C5-C6-N1	7.85	124.93	121.00
24	YA	887	C	C6-N1-C2	-7.83	117.17	120.30
24	YA	1579	C	C6-N1-C2	-7.83	117.17	120.30
1	QA	789	U	N3-C2-O2	-7.82	116.73	122.20
24	YA	2724	U	N3-C2-O2	-7.81	116.73	122.20
24	YA	2571	C	C6-N1-C2	-7.80	117.18	120.30
1	XA	789	U	N3-C2-O2	-7.80	116.74	122.20
24	YA	232	U	C5-C6-N1	7.80	126.60	122.70
24	RA	637	U	N1-C2-O2	7.79	128.26	122.80
24	YA	603	C	C6-N1-C2	-7.79	117.18	120.30
24	RA	2678	C	N3-C2-O2	-7.79	116.45	121.90
24	RA	570	C	C5-C6-N1	7.79	124.89	121.00

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	1978	U	N1-C2-O2	7.78	128.24	122.80
1	XA	1381	U	N3-C2-O2	-7.77	116.76	122.20
24	YA	1817	A	C8-N9-C4	-7.76	102.69	105.80
24	YA	1824	C	C6-N1-C2	-7.76	117.20	120.30
1	QA	283	C	N1-C2-O2	7.76	123.56	118.90
1	QA	307	C	N3-C2-O2	-7.76	116.47	121.90
24	YA	2480	G	O4'-C1'-N9	7.72	114.38	108.20
24	RA	2196	C	C6-N1-C2	-7.72	117.21	120.30
24	YA	1625	U	N3-C2-O2	-7.71	116.80	122.20
24	RA	783	C	C5-C6-N1	7.71	124.85	121.00
24	YA	2678	C	N3-C2-O2	-7.70	116.51	121.90
24	YA	1812	C	N1-C2-O2	7.70	123.52	118.90
24	YA	553	A	C2-N3-C4	-7.70	106.75	110.60
24	RA	1358	U	C5-C6-N1	7.69	126.55	122.70
24	YA	1691	C	C5-C6-N1	7.69	124.84	121.00
24	RA	755	C	C5-C6-N1	7.67	124.84	121.00
24	RA	1903	C	C5-C6-N1	7.66	124.83	121.00
24	YA	139	A	N7-C8-N9	7.66	117.63	113.80
24	YA	554	A	N7-C8-N9	7.65	117.63	113.80
24	YA	723	A	C5-N7-C8	-7.65	100.07	103.90
1	QA	699	C	C6-N1-C2	-7.65	117.24	120.30
24	RA	1937	U	N3-C2-O2	-7.65	116.84	122.20
24	YA	220	C	C6-N1-C2	-7.65	117.24	120.30
24	YA	1099	C	C5-C6-N1	7.65	124.83	121.00
24	YA	907	U	N1-C2-O2	7.64	128.15	122.80
24	YA	2415	C	C6-N1-C2	-7.64	117.24	120.30
24	RA	1560	U	N3-C2-O2	-7.64	116.86	122.20
24	RA	2878	A	N7-C8-N9	7.64	117.62	113.80
1	QA	1347	G	C4-N9-C1'	7.63	136.43	126.50
1	XA	1260	C	N3-C2-O2	-7.63	116.56	121.90
24	RA	1579	C	N1-C2-O2	7.63	123.48	118.90
1	QA	1301	U	C2-N1-C1'	7.63	126.86	117.70
24	RA	1937	U	N1-C2-O2	7.63	128.14	122.80
24	RA	844	C	C5-C6-N1	7.62	124.81	121.00
1	XA	330	C	N1-C2-O2	7.62	123.47	118.90
24	RA	2485	U	N1-C2-O2	7.62	128.13	122.80
24	YA	562	C	C5-C6-N1	7.61	124.81	121.00
24	YA	1914	C	C6-N1-C2	-7.61	117.26	120.30
24	RA	802	C	C6-N1-C2	-7.61	117.26	120.30
24	RA	2255	U	N3-C2-O2	-7.61	116.88	122.20
24	RA	1154	U	N3-C2-O2	-7.60	116.88	122.20
24	YA	2260	C	C6-N1-C2	-7.59	117.26	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	1478	C	C5-C6-N1	7.59	124.80	121.00
24	YA	1552	C	C5-C6-N1	7.59	124.80	121.00
24	RA	1138	C	C2-N1-C1'	7.59	127.14	118.80
24	RA	630	U	N3-C2-O2	-7.58	116.89	122.20
24	RA	903	C	N1-C2-O2	7.58	123.45	118.90
1	XA	1502	A	C5-N7-C8	-7.58	100.11	103.90
1	XA	497	U	N3-C2-O2	-7.58	116.90	122.20
24	YA	2028	C	C6-N1-C2	-7.56	117.27	120.30
1	XA	359	U	O4'-C1'-N1	7.55	114.24	108.20
24	YA	579	G	N7-C8-N9	7.55	116.88	113.10
24	YA	1457	C	C5-C6-N1	7.55	124.78	121.00
24	YA	223	C	N1-C2-O2	7.55	123.43	118.90
1	QA	1027	C	C6-N1-C2	-7.54	117.29	120.30
24	RA	2065	C	C6-N1-C2	-7.52	117.29	120.30
24	RA	482	C	C5-C6-N1	7.52	124.76	121.00
24	RA	2571	C	N1-C2-O2	7.52	123.41	118.90
24	YA	884	C	N1-C2-O2	7.52	123.41	118.90
24	YA	1378	G	C6-C5-N7	-7.51	125.89	130.40
24	YA	2851	C	C6-N1-C2	-7.51	117.30	120.30
24	YA	1817	A	C5-N7-C8	-7.49	100.15	103.90
1	XA	1381	U	N1-C2-O2	7.49	128.04	122.80
24	RA	1352	C	C6-N1-C2	-7.48	117.31	120.30
24	YA	1310	G	N7-C8-N9	7.48	116.84	113.10
24	RA	618	C	C6-N1-C2	-7.48	117.31	120.30
24	RA	1279	C	C6-N1-C2	-7.47	117.31	120.30
24	YA	2428	C	C5-C6-N1	7.47	124.74	121.00
1	XA	797	C	C6-N1-C2	-7.47	117.31	120.30
24	RA	637	U	N3-C2-O2	-7.46	116.98	122.20
24	YA	554	A	C8-N9-C4	-7.46	102.82	105.80
24	RA	1805	C	C6-N1-C2	-7.46	117.32	120.30
24	RA	1138	C	N3-C2-O2	-7.45	116.69	121.90
24	RA	1379	C	N1-C2-O2	7.45	123.37	118.90
24	YA	1020	C	N3-C2-O2	-7.45	116.69	121.90
24	RA	1478	C	C5-C6-N1	7.44	124.72	121.00
1	XA	749	C	C6-N1-C2	-7.44	117.32	120.30
1	XA	827	U	C6-N1-C2	-7.44	116.54	121.00
24	RA	1360	C	C5-C6-N1	7.42	124.71	121.00
1	QA	1020	U	N3-C2-O2	-7.42	117.01	122.20
1	QA	1323	G	N3-C4-C5	-7.42	124.89	128.60
24	YA	1725	G	C6-C5-N7	-7.42	125.95	130.40
24	RA	2004	C	C6-N1-C2	-7.41	117.33	120.30
1	XA	135	C	N1-C2-O2	7.41	123.35	118.90

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	258	U	N3-C2-O2	-7.41	117.01	122.20
24	YA	2596	U	N1-C2-O2	7.41	127.99	122.80
1	QA	1040	U	N3-C2-O2	-7.41	117.01	122.20
1	QA	1147	C	N3-C2-O2	-7.40	116.72	121.90
24	RA	555	G	O4'-C1'-N9	7.39	114.12	108.20
24	YA	1730	C	C5-C6-N1	7.39	124.70	121.00
1	QA	110	C	N1-C2-O2	7.39	123.33	118.90
1	XA	135	C	C6-N1-C2	-7.39	117.34	120.30
38	RT	99	LEU	CA-CB-CG	7.39	132.29	115.30
24	YA	1826	C	C5-C6-N1	7.39	124.69	121.00
24	RA	1521	C	N1-C2-O2	7.38	123.33	118.90
24	YA	2404	A	N7-C8-N9	7.38	117.49	113.80
24	RA	447	C	C5-C6-N1	7.38	124.69	121.00
24	RA	1360	C	C6-N1-C2	-7.38	117.35	120.30
24	YA	1819	C	C5-C6-N1	7.38	124.69	121.00
1	XA	754	C	C2-N1-C1'	7.37	126.91	118.80
1	XA	110	C	N3-C2-O2	-7.37	116.74	121.90
24	YA	139	A	C5-N7-C8	-7.37	100.22	103.90
24	RA	1310	G	N7-C8-N9	7.36	116.78	113.10
24	YA	660	C	C5-C6-N1	7.35	124.68	121.00
24	YA	1574	A	N7-C8-N9	7.35	117.47	113.80
24	YA	2798	C	C6-N1-C2	-7.34	117.36	120.30
24	YA	723	A	N7-C8-N9	7.34	117.47	113.80
24	RA	482	C	N1-C2-O2	7.34	123.31	118.90
24	YA	1972	G	C4-N9-C1'	7.34	136.04	126.50
1	XA	1024	G	C2-N3-C4	7.34	115.57	111.90
24	RA	1722	C	C6-N1-C2	-7.34	117.36	120.30
1	XA	58	C	N1-C2-O2	7.34	123.30	118.90
24	YA	1738	C	C6-N1-C2	-7.33	117.37	120.30
24	RA	181	C	C6-N1-C2	-7.33	117.37	120.30
1	QA	58	C	C5-C6-N1	7.32	124.66	121.00
24	YA	340	C	C6-N1-C2	-7.31	117.38	120.30
24	RA	1128	U	N3-C2-O2	-7.31	117.08	122.20
24	YA	875	U	C2-N1-C1'	7.30	126.46	117.70
24	YA	1705	C	C5-C6-N1	7.30	124.65	121.00
24	YA	444	C	C5-C6-N1	7.30	124.65	121.00
24	RA	394	C	C6-N1-C2	-7.29	117.38	120.30
24	RA	755	C	N1-C2-O2	7.29	123.27	118.90
24	RA	660	C	C6-N1-C2	-7.28	117.39	120.30
24	YA	2884	C	C6-N1-C2	-7.28	117.39	120.30
24	YA	1596	C	N1-C2-O2	7.28	123.27	118.90
24	RA	1709	C	C6-N1-C2	-7.27	117.39	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	1904	C	C5-C6-N1	7.27	124.64	121.00
24	YA	2877	G	C8-N9-C1'	-7.27	117.55	127.00
24	RA	1551	C	N1-C2-O2	7.26	123.26	118.90
24	YA	2324	U	C5-C6-N1	7.26	126.33	122.70
24	YA	553	A	N7-C8-N9	7.26	117.43	113.80
24	RA	447	C	C6-N1-C2	-7.26	117.40	120.30
1	QA	1163	C	C6-N1-C2	-7.26	117.40	120.30
24	RA	2486	C	N1-C2-O2	7.25	123.25	118.90
24	YA	566	C	C6-N1-C2	-7.25	117.40	120.30
24	YA	1457	C	C6-N1-C2	-7.23	117.41	120.30
1	QA	754	C	C2-N1-C1'	7.22	126.75	118.80
24	RA	884	C	C6-N1-C2	-7.22	117.41	120.30
24	RA	958	C	C6-N1-C2	-7.22	117.41	120.30
22	QV	67	U	N3-C2-O2	-7.21	117.15	122.20
1	XA	1147	C	N1-C2-O2	7.21	123.23	118.90
24	YA	1059	C	C6-N1-C2	-7.20	117.42	120.30
24	RA	1359	U	C2-N1-C1'	7.20	126.33	117.70
24	RA	2028	C	C6-N1-C2	-7.19	117.42	120.30
24	RA	1866	G	N3-C4-C5	-7.19	125.00	128.60
24	YA	819	C	C5-C6-N1	7.19	124.60	121.00
24	RA	1560	U	N1-C2-O2	7.19	127.83	122.80
1	QA	1040	U	N1-C2-O2	7.19	127.83	122.80
1	QA	1260	C	C6-N1-C2	-7.19	117.42	120.30
24	RA	141	C	C6-N1-C2	-7.19	117.42	120.30
24	RA	1350	C	C5-C6-N1	7.18	124.59	121.00
24	RA	1154	U	N1-C2-O2	7.18	127.82	122.80
24	RA	1550	C	C6-N1-C2	-7.17	117.43	120.30
24	YA	2255	U	N3-C2-O2	-7.17	117.18	122.20
24	YA	1937	U	N3-C2-O2	-7.16	117.19	122.20
24	RA	2065	C	C5-C6-N1	7.16	124.58	121.00
25	YB	114	G	C2-N3-C4	7.15	115.48	111.90
24	RA	944	C	N1-C2-O2	7.14	123.19	118.90
24	YA	1067	A	N3-C4-N9	-7.14	121.69	127.40
1	QA	1132	C	N1-C2-O2	7.14	123.18	118.90
1	XA	979	C	N3-C2-O2	-7.14	116.90	121.90
24	YA	2668	U	C5-C6-N1	7.13	126.27	122.70
24	RA	1580	G	N3-C4-C5	-7.13	125.03	128.60
24	RA	1978	U	N3-C2-O2	-7.13	117.21	122.20
24	YA	778	C	C5-C6-N1	7.13	124.56	121.00
1	XA	738	C	C6-N1-C2	-7.13	117.45	120.30
24	RA	140	A	C5-N7-C8	-7.12	100.34	103.90
24	RA	1421	C	C6-N1-C2	-7.12	117.45	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	1457	C	N1-C2-O2	7.12	123.17	118.90
24	RA	395	C	C6-N1-C2	-7.12	117.45	120.30
24	YA	1580	G	N3-C4-C5	-7.12	125.04	128.60
24	YA	1581	U	C2-N1-C1'	7.12	126.24	117.70
24	RA	1523	C	C5-C6-N1	7.12	124.56	121.00
24	RA	907	U	N1-C2-O2	7.11	127.78	122.80
24	YA	1542	A	N7-C8-N9	7.11	117.36	113.80
1	XA	1158	C	C2-N1-C1'	7.11	126.62	118.80
24	YA	2818	U	N3-C2-O2	-7.10	117.23	122.20
24	RA	482	C	C6-N1-C2	-7.10	117.46	120.30
24	RA	1645	C	C6-N1-C2	-7.10	117.46	120.30
24	YA	1463	C	C6-N1-C2	-7.10	117.46	120.30
24	YA	1691	C	N1-C2-O2	7.10	123.16	118.90
24	RA	1956	C	C5-C6-N1	7.10	124.55	121.00
24	YA	1444	C	N1-C2-O2	7.10	123.16	118.90
24	RA	630	U	N1-C2-O2	7.09	127.77	122.80
24	YA	2880	C	C6-N1-C2	-7.09	117.46	120.30
24	YA	2011	G	N3-C2-N2	-7.09	114.94	119.90
24	RA	2028	C	N1-C2-O2	7.09	123.16	118.90
1	XA	979	C	N1-C2-O2	7.09	123.16	118.90
24	RA	2857	U	N1-C2-O2	7.09	127.76	122.80
1	QA	449	C	N1-C2-O2	7.08	123.15	118.90
24	RA	1969	C	C6-N1-C2	-7.08	117.47	120.30
24	YA	2528	G	N3-C4-C5	-7.07	125.07	128.60
24	RA	400	U	C6-N1-C2	-7.07	116.76	121.00
24	RA	1121	C	N1-C2-O2	7.07	123.14	118.90
24	YA	2768	C	C5-C6-N1	7.06	124.53	121.00
25	RB	31	C	N3-C2-O2	-7.05	116.96	121.90
24	RA	31	C	C5-C6-N1	7.05	124.53	121.00
24	RA	2185	C	C5-C6-N1	7.05	124.53	121.00
1	QA	346	G	N3-C4-N9	7.05	130.23	126.00
24	YA	778	C	C6-N1-C2	-7.04	117.48	120.30
24	YA	1378	G	C4-N9-C1'	7.04	135.65	126.50
24	RA	1901	C	C6-N1-C2	-7.04	117.48	120.30
24	YA	2502	G	C4-C5-N7	7.04	113.61	110.80
24	YA	2818	U	N1-C2-O2	7.04	127.73	122.80
24	YA	2877	G	C4-N9-C1'	7.04	135.65	126.50
24	RA	2201	C	N1-C2-O2	7.03	123.12	118.90
24	YA	1067	A	C5-N7-C8	-7.03	100.38	103.90
24	YA	1709	C	C6-N1-C2	-7.02	117.49	120.30
1	QA	1378	C	C6-N1-C2	-7.02	117.49	120.30
24	RA	1819	C	C5-C6-N1	7.02	124.51	121.00

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	497	U	N1-C2-O2	7.02	127.71	122.80
24	YA	554	A	C8-N9-C1'	-7.02	115.07	127.70
24	YA	41	C	C6-N1-C2	-7.01	117.49	120.30
1	XA	528	C	C6-N1-C2	-7.01	117.50	120.30
24	YA	2222	C	C6-N1-C2	-7.01	117.50	120.30
24	RA	112	U	C5-C6-N1	7.00	126.20	122.70
24	RA	258	U	N3-C2-O2	-7.00	117.30	122.20
1	XA	135	C	N3-C2-O2	-7.00	117.00	121.90
24	YA	1478	C	C6-N1-C2	-7.00	117.50	120.30
24	YA	2479	C	C2-N3-C4	-7.00	116.40	119.90
1	QA	346	G	N3-C4-C5	-6.99	125.10	128.60
1	XA	307	C	N3-C2-O2	-6.98	117.01	121.90
24	YA	1067	A	N7-C8-N9	6.98	117.29	113.80
24	RA	1864	U	N3-C2-O2	-6.98	117.31	122.20
1	XA	283	C	N1-C2-O2	6.98	123.09	118.90
24	YA	1022	C	C6-N1-C2	-6.97	117.51	120.30
24	RA	914	C	N1-C2-O2	6.97	123.08	118.90
24	YA	2519	C	C6-N1-C2	-6.97	117.51	120.30
25	RB	31	C	C6-N1-C2	-6.97	117.51	120.30
25	YB	81	G	C5-N7-C8	-6.97	100.82	104.30
24	RA	555	G	N1-C6-O6	-6.96	115.72	119.90
24	RA	939	C	N1-C2-O2	6.96	123.08	118.90
1	XA	679	C	C5-C6-N1	6.96	124.48	121.00
24	YA	637	U	N1-C2-O2	6.96	127.67	122.80
24	RA	2150	C	N1-C2-O2	6.96	123.08	118.90
24	YA	994	C	C5-C6-N1	6.96	124.48	121.00
1	QA	503	C	C6-N1-C2	-6.96	117.52	120.30
24	YA	2063	U	C5-C6-N1	6.96	126.18	122.70
1	XA	449	C	C2-N1-C1'	6.95	126.45	118.80
1	QA	1279	A	N7-C8-N9	6.95	117.28	113.80
24	RA	1956	C	C6-N1-C2	-6.95	117.52	120.30
24	YA	2222	C	C5-C6-N1	6.95	124.47	121.00
24	RA	1457	C	N1-C2-O2	6.95	123.07	118.90
24	YA	830	A	C5-N7-C8	-6.94	100.43	103.90
24	YA	2319	G	C8-N9-C1'	-6.94	117.97	127.00
22	QV	67	U	N1-C2-O2	6.94	127.66	122.80
24	YA	66	U	C5-C6-N1	6.93	126.17	122.70
24	YA	187	C	C6-N1-C2	-6.93	117.53	120.30
24	RA	2736	C	C5-C6-N1	6.93	124.46	121.00
24	YA	2083	G	N3-C4-N9	6.93	130.16	126.00
24	YA	1978	U	N3-C2-O2	-6.92	117.35	122.20
24	YA	719	C	C6-N1-C2	-6.92	117.53	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	554	C	C6-N1-C2	-6.92	117.53	120.30
24	YA	2066	C	C6-N1-C2	-6.92	117.53	120.30
24	YA	139	A	C8-N9-C4	-6.92	103.03	105.80
24	RA	604	C	C6-N1-C2	-6.91	117.53	120.30
1	XA	410	G	C8-N9-C4	-6.90	103.64	106.40
1	QA	1439	C	C6-N1-C2	-6.90	117.54	120.30
24	RA	2880	C	C6-N1-C2	-6.90	117.54	120.30
24	RA	562	C	C6-N1-C2	-6.89	117.55	120.30
1	XA	1452	C	N1-C2-O2	6.88	123.03	118.90
25	YB	112	G	C6-N1-C2	-6.88	120.97	125.10
24	YA	1703	C	C6-N1-C2	-6.88	117.55	120.30
24	RA	755	C	C6-N1-C2	-6.88	117.55	120.30
1	XA	37	U	N3-C2-O2	-6.88	117.39	122.20
1	QA	266	G	C8-N9-C1'	-6.87	118.06	127.00
24	RA	2857	U	N3-C2-O2	-6.87	117.39	122.20
24	YA	2884	C	C5-C6-N1	6.87	124.44	121.00
24	YA	1707	C	C6-N1-C2	-6.87	117.55	120.30
24	YA	1099	C	N1-C2-O2	6.87	123.02	118.90
24	RA	1904	C	C2-N1-C1'	6.86	126.35	118.80
24	RA	872	C	C6-N1-C2	-6.86	117.56	120.30
24	YA	864	C	C6-N1-C2	-6.86	117.56	120.30
1	XA	738	C	C5-C6-N1	6.86	124.43	121.00
24	YA	1009	C	C6-N1-C2	-6.85	117.56	120.30
24	YA	511	C	C6-N1-C2	-6.85	117.56	120.30
1	QA	525	C	C6-N1-C2	-6.84	117.56	120.30
1	QA	780	A	C8-N9-C4	-6.84	103.06	105.80
24	YA	12	U	C2-N1-C1'	6.84	125.91	117.70
24	YA	1099	C	C6-N1-C2	-6.84	117.56	120.30
24	RA	2739	U	N1-C2-O2	6.84	127.58	122.80
1	QA	1020	U	N1-C2-O2	6.83	127.58	122.80
24	YA	2807	C	N1-C2-O2	6.83	123.00	118.90
24	RA	2596	U	N3-C2-O2	-6.83	117.42	122.20
24	YA	1550	C	C6-N1-C2	-6.83	117.57	120.30
24	RA	140	A	N7-C8-N9	6.83	117.21	113.80
24	YA	2001	C	C6-N1-C2	-6.83	117.57	120.30
24	YA	659	C	C5-C6-N1	6.82	124.41	121.00
24	YA	1204	C	C6-N1-C2	-6.82	117.57	120.30
24	YA	2899	C	C5-C6-N1	6.82	124.41	121.00
24	RA	1360	C	N1-C2-O2	6.82	122.99	118.90
24	RA	1942	C	C5-C6-N1	6.82	124.41	121.00
24	RA	1550	C	N1-C2-O2	6.81	122.99	118.90
1	QA	1362(A)	C	C6-N1-C2	-6.81	117.58	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	1580	G	N3-C4-N9	6.81	130.09	126.00
24	YA	2596	U	C2-N1-C1'	6.81	125.87	117.70
24	YA	1812	C	N3-C2-O2	-6.80	117.14	121.90
24	RA	1378	G	C8-N9-C4	-6.80	103.68	106.40
1	QA	554	C	C5-C6-N1	6.80	124.40	121.00
1	QA	1147	C	N1-C2-O2	6.80	122.98	118.90
24	RA	675	C	C5-C6-N1	6.80	124.40	121.00
24	YA	2238	C	N1-C2-O2	6.80	122.98	118.90
1	QA	1163	C	C2-N1-C1'	6.79	126.28	118.80
1	XA	346	G	N3-C4-N9	6.79	130.08	126.00
24	RA	672	G	C6-C5-N7	-6.79	126.33	130.40
24	YA	553	A	N3-C4-N9	-6.79	121.97	127.40
24	YA	103	C	C5-C6-N1	6.79	124.39	121.00
24	YA	739	C	C6-N1-C2	-6.79	117.58	120.30
24	YA	2083	G	N3-C4-C5	-6.79	125.21	128.60
24	RA	1657	C	C6-N1-C2	-6.78	117.59	120.30
24	YA	2004	C	C5-C6-N1	6.78	124.39	121.00
1	QA	266	G	N3-C4-N9	-6.78	121.93	126.00
1	QA	1158	C	C2-N1-C1'	6.78	126.26	118.80
24	RA	1730	C	C5-C6-N1	6.78	124.39	121.00
1	XA	58	C	C6-N1-C2	-6.78	117.59	120.30
24	YA	2629	C	N1-C2-O2	6.78	122.97	118.90
24	YA	1521	C	C6-N1-C2	-6.77	117.59	120.30
1	XA	186(G)	C	N3-C2-O2	-6.77	117.16	121.90
1	QA	1502	A	C4-C5-N7	6.76	114.08	110.70
24	RA	187	C	C6-N1-C2	-6.76	117.59	120.30
24	RA	297	C	C5-C6-N1	6.76	124.38	121.00
24	RA	1155	C	N1-C2-O2	6.76	122.96	118.90
24	RA	1717	C	N1-C2-O2	6.76	122.96	118.90
24	YA	1031	C	C5-C6-N1	6.76	124.38	121.00
24	YA	1453	C	N3-C2-O2	-6.76	117.17	121.90
24	RA	659	C	C6-N1-C2	-6.75	117.60	120.30
24	RA	2319	G	C4-N9-C1'	6.75	135.28	126.50
25	YB	114	G	N3-C2-N2	6.75	124.63	119.90
24	RA	2030	C	C6-N1-C2	-6.75	117.60	120.30
24	RA	1457	C	C6-N1-C2	-6.75	117.60	120.30
24	YA	853	C	C5-C6-N1	6.75	124.37	121.00
1	QA	1452	C	N1-C2-O2	6.75	122.95	118.90
24	YA	2592	U	N3-C2-O2	-6.74	117.48	122.20
24	YA	1542	A	C8-N9-C4	-6.74	103.11	105.80
1	XA	1260	C	C6-N1-C2	-6.74	117.61	120.30
24	RA	2165	C	C5-C6-N1	6.73	124.37	121.00

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	181	C	N3-C2-O2	-6.73	117.19	121.90
24	YA	2899	C	C6-N1-C2	-6.73	117.61	120.30
24	YA	1730	C	C6-N1-C2	-6.72	117.61	120.30
1	QA	1038	C	N1-C2-O2	6.72	122.93	118.90
24	RA	2835	C	N3-C2-O2	-6.72	117.19	121.90
24	YA	2066	C	C5-C6-N1	6.72	124.36	121.00
1	XA	1113	C	C6-N1-C2	-6.72	117.61	120.30
24	YA	1864	U	N3-C2-O2	-6.72	117.50	122.20
24	RA	1350	C	C6-N1-C2	-6.71	117.61	120.30
24	RA	800	C	C5-C6-N1	6.71	124.36	121.00
24	YA	1836	U	C6-N1-C2	-6.71	116.97	121.00
1	XA	1028(C)	C	C6-N1-C2	-6.71	117.62	120.30
24	YA	1521	C	N1-C2-O2	6.71	122.92	118.90
25	RB	31	C	C5-C6-N1	6.70	124.35	121.00
24	RA	844	C	C6-N1-C2	-6.70	117.62	120.30
1	XA	58	C	C5-C6-N1	6.70	124.35	121.00
24	RA	723	A	C8-N9-C4	-6.70	103.12	105.80
1	XA	1439	C	C6-N1-C2	-6.70	117.62	120.30
24	YA	1552	C	C6-N1-C2	-6.70	117.62	120.30
1	QA	1113	C	C6-N1-C2	-6.70	117.62	120.30
24	YA	2472	U	N1-C2-O2	6.70	127.49	122.80
24	YA	1378	G	C2-N3-C4	-6.70	108.55	111.90
24	YA	1930	C	C6-N1-C2	-6.69	117.62	120.30
24	YA	1725	G	C4-C5-N7	6.69	113.48	110.80
24	RA	2571	C	C6-N1-C2	-6.69	117.62	120.30
24	RA	297	C	C6-N1-C2	-6.69	117.62	120.30
24	RA	783	C	C6-N1-C2	-6.68	117.63	120.30
1	XA	1362(B)	C	C6-N1-C2	-6.68	117.63	120.30
24	YA	1867	C	C6-N1-C2	-6.68	117.63	120.30
24	YA	1819	C	C6-N1-C2	-6.68	117.63	120.30
24	YA	1271	G	C8-N9-C4	-6.67	103.73	106.40
24	RA	258	U	N1-C2-O2	6.67	127.47	122.80
1	XA	620	C	N1-C2-O2	6.67	122.90	118.90
24	YA	1580	G	C2-N3-C4	6.67	115.24	111.90
24	YA	2061	C	C5-C6-N1	6.67	124.34	121.00
24	RA	1279	C	C5-C6-N1	6.67	124.33	121.00
24	YA	1263	C	C6-N1-C2	-6.67	117.63	120.30
24	RA	1352	C	C5-C6-N1	6.67	124.33	121.00
24	RA	2130	C	C5-C6-N1	6.66	124.33	121.00
1	XA	1037	C	C6-N1-C2	-6.66	117.64	120.30
24	YA	2260	C	N3-C2-O2	-6.66	117.24	121.90
24	RA	1810	U	C2-N1-C1'	6.66	125.69	117.70

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	826	C	N3-C2-O2	-6.66	117.24	121.90
24	RA	1579	C	C5-C6-N1	6.66	124.33	121.00
24	YA	2528	G	C2-N3-C4	6.65	115.23	111.90
24	RA	2333	G	N3-C4-C5	-6.65	125.27	128.60
24	RA	1483	C	C6-N1-C2	-6.65	117.64	120.30
24	YA	1972	G	C6-C5-N7	-6.65	126.41	130.40
24	RA	340	C	C6-N1-C2	-6.65	117.64	120.30
24	RA	570	C	N1-C2-O2	6.65	122.89	118.90
24	YA	1263	C	C5-C6-N1	6.64	124.32	121.00
24	YA	1581	U	C6-N1-C2	-6.64	117.01	121.00
1	QA	1314	C	N1-C2-O2	6.64	122.89	118.90
24	YA	2480	G	C8-N9-C1'	-6.64	118.37	127.00
24	YA	730	C	C6-N1-C2	-6.64	117.64	120.30
24	YA	1020	C	C6-N1-C1'	-6.64	112.83	120.80
24	RA	75	C	C6-N1-C2	-6.64	117.64	120.30
24	YA	1590	C	N3-C2-O2	-6.63	117.26	121.90
1	QA	266	G	C4-N9-C1'	6.63	135.12	126.50
24	YA	958	C	C6-N1-C2	-6.63	117.65	120.30
24	YA	1450	C	C6-N1-C2	-6.62	117.65	120.30
24	YA	646	A	C5-N7-C8	-6.62	100.59	103.90
24	RA	1703	C	C6-N1-C2	-6.62	117.65	120.30
24	RA	2780	C	C6-N1-C2	-6.62	117.65	120.30
24	RA	1548	C	C6-N1-C2	-6.61	117.65	120.30
24	RA	1931	C	C6-N1-C2	-6.61	117.66	120.30
24	YA	1819	C	N1-C2-O2	6.61	122.86	118.90
24	YA	512	C	C5-C6-N1	6.61	124.30	121.00
1	QA	110	C	N3-C2-O2	-6.60	117.28	121.90
1	QA	522	C	N1-C2-O2	6.60	122.86	118.90
24	YA	223	C	N3-C2-O2	-6.60	117.28	121.90
1	QA	623	C	C6-N1-C2	-6.60	117.66	120.30
24	RA	885	C	C5-C6-N1	6.60	124.30	121.00
24	RA	603	C	C6-N1-C2	-6.59	117.66	120.30
23	XX	17	C	N1-C2-O2	6.59	122.86	118.90
1	QA	993	G	C5-C6-O6	-6.59	124.64	128.60
24	RA	830	A	N7-C8-N9	6.59	117.10	113.80
24	RA	931	C	C6-N1-C2	-6.59	117.66	120.30
1	QA	1439	C	C5-C6-N1	6.58	124.29	121.00
1	QA	1449	C	C6-N1-C2	-6.58	117.67	120.30
24	YA	1824	C	C5-C6-N1	6.58	124.29	121.00
1	QA	1502	A	C5-N7-C8	-6.58	100.61	103.90
24	RA	570	C	C2-N1-C1'	6.58	126.03	118.80
1	XA	826	C	C6-N1-C2	-6.58	117.67	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	2228	G	C4-N9-C1'	6.57	135.04	126.50
24	YA	2518	U	N3-C2-O2	-6.57	117.60	122.20
24	YA	73	A	N7-C8-N9	6.57	117.08	113.80
24	RA	1453	C	N1-C2-O2	6.57	122.84	118.90
1	XA	1260	C	N1-C2-O2	6.57	122.84	118.90
24	YA	1521	C	C5-C6-N1	6.57	124.28	121.00
24	YA	1738	C	C5-C6-N1	6.57	124.28	121.00
24	YA	2700	U	N1-C2-N3	6.57	118.84	114.90
24	YA	2700	U	C2-N1-C1'	6.57	125.58	117.70
1	QA	1163	C	N1-C2-O2	6.56	122.83	118.90
24	RA	1903	C	C6-N1-C2	-6.56	117.68	120.30
24	RA	2715	C	N1-C2-O2	6.55	122.83	118.90
1	XA	705	U	N3-C2-O2	-6.55	117.61	122.20
24	RA	2130	C	C6-N1-C2	-6.54	117.68	120.30
24	RA	2238	C	N1-C2-O2	6.54	122.83	118.90
24	RA	2638	C	C6-N1-C2	-6.54	117.68	120.30
24	RA	1523	C	C6-N1-C2	-6.54	117.69	120.30
1	XA	797	C	C5-C6-N1	6.54	124.27	121.00
24	YA	2724	U	N1-C2-O2	6.53	127.37	122.80
24	YA	735	U	C5-C6-N1	6.53	125.96	122.70
24	RA	723	A	C4-N9-C1'	6.53	138.05	126.30
24	RA	939	C	N3-C2-O2	-6.53	117.33	121.90
24	RA	1010	C	C6-N1-C2	-6.52	117.69	120.30
24	RA	2598	C	N1-C2-O2	6.52	122.81	118.90
24	YA	447	C	N1-C2-O2	6.52	122.81	118.90
24	RA	738	C	C5-C6-N1	6.51	124.25	121.00
25	YB	112	G	N1-C6-O6	-6.51	115.99	119.90
1	XA	891	U	N3-C2-O2	-6.51	117.64	122.20
24	RA	321	C	C6-N1-C2	-6.50	117.70	120.30
24	RA	1450	C	C6-N1-C2	-6.50	117.70	120.30
1	QA	972	C	C6-N1-C2	-6.50	117.70	120.30
24	YA	1916	C	C6-N1-C2	-6.50	117.70	120.30
24	YA	1141	A	C2-N3-C4	6.50	113.85	110.60
1	QA	1430	C	C6-N1-C2	-6.50	117.70	120.30
24	RA	2087	C	C6-N1-C2	-6.49	117.70	120.30
24	YA	1098	C	C6-N1-C2	-6.49	117.70	120.30
24	YA	2890	C	C6-N1-C2	-6.49	117.71	120.30
24	RA	1722	C	N3-C2-O2	-6.48	117.36	121.90
24	YA	68	C	N1-C2-O2	6.48	122.79	118.90
24	YA	41	C	C5-C6-N1	6.48	124.24	121.00
24	RA	2454	C	C6-N1-C2	-6.48	117.71	120.30
24	YA	530	A	N1-C6-N6	-6.47	114.72	118.60

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
25	RB	27	C	N1-C2-O2	6.47	122.78	118.90
24	YA	1421	C	C5-C6-N1	6.47	124.24	121.00
24	YA	1936	C	C2-N1-C1'	6.47	125.92	118.80
24	YA	2228	G	N3-C4-N9	6.47	129.88	126.00
1	QA	1263	C	C5-C6-N1	6.47	124.23	121.00
24	RA	796	C	C5-C6-N1	6.47	124.23	121.00
24	RA	665	C	C5-C6-N1	6.47	124.23	121.00
1	QA	36	C	C6-N1-C2	-6.47	117.71	120.30
1	XA	827	U	N1-C2-N3	6.46	118.78	114.90
24	YA	1436	U	N1-C2-O2	6.46	127.32	122.80
24	YA	118	U	C4-C5-C6	6.46	123.58	119.70
24	RA	1691	C	C5-C6-N1	6.46	124.23	121.00
24	RA	1313	U	N3-C2-O2	-6.45	117.68	122.20
24	YA	535	C	C6-N1-C2	-6.45	117.72	120.30
24	YA	2347	A	O4'-C1'-N9	6.45	113.36	108.20
24	YA	1705	C	C6-N1-C2	-6.45	117.72	120.30
24	YA	2309	C	C6-N1-C2	-6.45	117.72	120.30
24	RA	1092	A	C8-N9-C4	-6.44	103.22	105.80
1	QA	1027	C	C2-N1-C1'	6.44	125.89	118.80
1	XA	679	C	C6-N1-C2	-6.44	117.72	120.30
24	YA	511	C	C5-C6-N1	6.44	124.22	121.00
24	YA	862	C	C6-N1-C2	-6.44	117.72	120.30
24	YA	18	C	C6-N1-C2	-6.44	117.72	120.30
24	YA	2700	U	C6-N1-C2	-6.44	117.14	121.00
24	YA	2524	C	C5-C6-N1	6.44	124.22	121.00
24	RA	2222	C	C6-N1-C2	-6.43	117.73	120.30
24	YA	853	C	C6-N1-C2	-6.43	117.73	120.30
24	YA	2260	C	N1-C2-O2	6.43	122.76	118.90
24	YA	2404	A	C5-N7-C8	-6.43	100.68	103.90
24	YA	2792	U	C2-N1-C1'	6.43	125.42	117.70
24	YA	893	C	C6-N1-C2	-6.43	117.73	120.30
24	RA	2519	C	C5-C6-N1	6.43	124.22	121.00
1	XA	1140	C	N1-C2-O2	6.43	122.76	118.90
25	YB	79	C	C6-N1-C2	-6.43	117.73	120.30
24	YA	2623	U	C5-C6-N1	6.43	125.91	122.70
24	RA	1817	A	C4-C5-N7	6.43	113.91	110.70
24	RA	1866	G	N3-C4-N9	6.43	129.86	126.00
1	XA	578	C	C6-N1-C2	-6.42	117.73	120.30
24	RA	1868	C	C6-N1-C2	-6.42	117.73	120.30
24	YA	1123	A	C2-N3-C4	6.42	113.81	110.60
1	QA	1149	C	N1-C2-O2	6.42	122.75	118.90
24	YA	939	C	N3-C2-O2	-6.42	117.41	121.90

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
25	RB	37	C	N1-C2-O2	6.41	122.75	118.90
24	YA	802	C	C6-N1-C2	-6.41	117.73	120.30
24	RA	2480	G	C4-N9-C1'	6.41	134.84	126.50
24	YA	841	G	N7-C8-N9	6.41	116.31	113.10
25	YB	93	C	N3-C2-O2	-6.41	117.41	121.90
24	RA	2598	C	C5-C6-N1	6.41	124.20	121.00
24	YA	553	A	C5-N7-C8	-6.41	100.70	103.90
24	YA	138	G	N9-C4-C5	6.40	107.96	105.40
24	RA	875	U	C2-N1-C1'	6.40	125.38	117.70
24	YA	1580	G	N3-C4-N9	6.40	129.84	126.00
1	QA	328	C	C5-C6-N1	6.40	124.20	121.00
1	XA	1502	A	C8-N9-C4	-6.40	103.24	105.80
23	XX	17	C	C6-N1-C2	-6.40	117.74	120.30
24	YA	796	C	C5-C6-N1	6.40	124.20	121.00
24	YA	1364	C	C5-C6-N1	6.40	124.20	121.00
1	XA	346	G	N3-C4-C5	-6.40	125.40	128.60
24	YA	2088	C	C5-C6-N1	6.40	124.20	121.00
24	RA	126	C	C5-C6-N1	6.40	124.20	121.00
1	QA	932	C	N1-C2-O2	6.40	122.74	118.90
24	YA	885	C	C6-N1-C2	-6.39	117.74	120.30
1	QA	56	U	C5-C6-N1	6.39	125.90	122.70
24	RA	242	C	C6-N1-C2	-6.39	117.74	120.30
24	YA	1188	A	N3-C4-C5	6.39	131.27	126.80
24	YA	2453	C	C6-N1-C2	-6.39	117.74	120.30
1	QA	252	U	N3-C2-O2	-6.38	117.73	122.20
25	YB	7	G	C4-C5-N7	6.38	113.35	110.80
25	YB	85	G	N9-C4-C5	6.38	107.95	105.40
24	YA	140	A	C4-C5-N7	6.38	113.89	110.70
24	RA	1805	C	N3-C2-O2	-6.38	117.43	121.90
24	YA	2088	C	N1-C2-O2	6.38	122.73	118.90
24	RA	961	C	C6-N1-C2	-6.38	117.75	120.30
24	RA	1125	C	C5-C6-N1	6.38	124.19	121.00
24	RA	2700	U	N1-C2-O2	6.38	127.26	122.80
24	YA	2479	C	N1-C2-N3	6.38	123.66	119.20
24	YA	394	C	N1-C2-O2	6.38	122.72	118.90
24	RA	2596	U	C2-N1-C1'	6.37	125.35	117.70
23	XX	3	C	C6-N1-C2	-6.37	117.75	120.30
24	YA	2177	G	N3-C4-C5	-6.37	125.42	128.60
51	Y6	6	ARG	NE-CZ-NH1	-6.37	117.11	120.30
1	XA	1420	C	C6-N1-C2	-6.37	117.75	120.30
24	YA	1691	C	C6-N1-C2	-6.37	117.75	120.30
25	RB	30	C	C6-N1-C2	-6.36	117.75	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	2785	C	C5-C6-N1	6.36	124.18	121.00
25	YB	60	C	C5-C6-N1	6.36	124.18	121.00
24	YA	854	U	N1-C2-O2	6.36	127.25	122.80
24	RA	2087	C	C5-C6-N1	6.36	124.18	121.00
1	XA	960	U	N1-C2-O2	6.36	127.25	122.80
1	XA	486	U	N3-C2-O2	-6.36	117.75	122.20
24	RA	2210	C	C6-N1-C2	-6.35	117.76	120.30
1	QA	449	C	C2-N1-C1'	6.35	125.78	118.80
24	RA	723	A	O4'-C1'-N9	6.35	113.28	108.20
24	YA	950	C	N1-C2-O2	6.35	122.71	118.90
24	YA	1733	C	C5-C6-N1	6.35	124.17	121.00
24	RA	2587	C	C6-N1-C2	-6.35	117.76	120.30
24	YA	2727	G	C4-C5-N7	6.35	113.34	110.80
25	YB	68	C	C5-C6-N1	6.35	124.17	121.00
24	RA	830	A	C5-N7-C8	-6.34	100.73	103.90
24	RA	1861	C	C6-N1-C2	-6.34	117.76	120.30
23	QX	14	A	C8-N9-C4	-6.33	103.27	105.80
24	YA	2518	U	N1-C2-O2	6.33	127.23	122.80
24	RA	2088	C	N1-C2-O2	6.33	122.70	118.90
24	YA	914	C	N1-C2-O2	6.33	122.70	118.90
24	RA	1972	G	C8-N9-C1'	-6.33	118.77	127.00
24	RA	2257	U	C5-C6-N1	6.33	125.86	122.70
24	YA	634	C	C6-N1-C2	-6.33	117.77	120.30
24	YA	1953	U	N3-C2-O2	-6.33	117.77	122.20
24	RA	1478	C	C6-N1-C2	-6.33	117.77	120.30
24	RA	1095	C	N1-C2-O2	6.32	122.69	118.90
1	XA	330	C	N3-C2-O2	-6.32	117.47	121.90
25	YB	30	C	C5-C6-N1	6.32	124.16	121.00
37	RS	110	LEU	CA-CB-CG	6.32	129.84	115.30
1	XA	381	C	N1-C2-O2	6.32	122.69	118.90
24	RA	45	C	C5-C6-N1	6.32	124.16	121.00
24	YA	1015	C	C5-C6-N1	6.32	124.16	121.00
1	QA	932	C	C6-N1-C2	-6.31	117.77	120.30
24	YA	157	U	N1-C2-O2	6.31	127.22	122.80
24	YA	1969	C	C6-N1-C2	-6.31	117.78	120.30
24	RA	1010	C	C5-C6-N1	6.30	124.15	121.00
24	YA	670	C	N1-C2-O2	6.30	122.68	118.90
24	YA	2628	C	C5-C6-N1	6.30	124.15	121.00
1	QA	346	G	C2-N3-C4	6.30	115.05	111.90
24	RA	1232	G	N7-C8-N9	6.30	116.25	113.10
24	RA	2333	G	C4-N9-C1'	6.30	134.69	126.50
1	XA	221	C	C6-N1-C2	-6.30	117.78	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	1969	C	N1-C2-O2	6.30	122.68	118.90
1	XA	1097	C	C6-N1-C2	-6.30	117.78	120.30
24	RA	1806	U	C5-C4-O4	-6.29	122.13	125.90
24	RA	903	C	N3-C2-O2	-6.29	117.50	121.90
24	YA	419	C	C5-C6-N1	6.29	124.14	121.00
24	YA	1032	C	C6-N1-C2	-6.28	117.79	120.30
24	YA	1016	C	C6-N1-C2	-6.28	117.79	120.30
1	QA	514	C	C6-N1-C2	-6.28	117.79	120.30
24	YA	2473	C	N1-C2-O2	6.28	122.67	118.90
24	YA	2739	U	N1-C2-O2	6.28	127.19	122.80
24	RA	2780	C	C5-C6-N1	6.27	124.14	121.00
24	RA	1110	C	C2-N1-C1'	6.26	125.69	118.80
23	XX	17	C	N3-C2-O2	-6.26	117.52	121.90
24	YA	2752	U	N3-C2-O2	-6.26	117.82	122.20
1	XA	1028(C)	C	N3-C2-O2	-6.26	117.52	121.90
24	YA	1378	G	N1-C2-N3	6.26	127.66	123.90
24	RA	2587	C	C5-C6-N1	6.26	124.13	121.00
51	R6	6	ARG	NE-CZ-NH1	-6.26	117.17	120.30
24	YA	2592	U	N1-C2-O2	6.26	127.18	122.80
24	YA	884	C	C5-C6-N1	6.25	124.13	121.00
24	YA	1550	C	C5-C6-N1	6.25	124.13	121.00
24	RA	1921	G	N3-C2-N2	-6.25	115.52	119.90
25	RB	27	C	C6-N1-C2	-6.25	117.80	120.30
24	RA	395	C	C5-C6-N1	6.25	124.12	121.00
1	QA	910	C	C5-C6-N1	6.25	124.12	121.00
24	RA	141	C	C5-C6-N1	6.25	124.12	121.00
24	RA	1775	C	C6-N1-C2	-6.25	117.80	120.30
24	YA	864	C	C5-C6-N1	6.25	124.12	121.00
24	RA	570	C	C6-N1-C2	-6.25	117.80	120.30
1	XA	812	C	P-O3'-C3'	6.25	127.19	119.70
24	YA	975	U	N3-C2-O2	-6.24	117.83	122.20
24	RA	1691	C	C6-N1-C2	-6.24	117.80	120.30
24	RA	197	C	C6-N1-C2	-6.24	117.80	120.30
24	RA	1977	U	C2-N1-C1'	6.24	125.18	117.70
24	YA	2630	G	N3-C4-C5	-6.24	125.48	128.60
24	RA	2095	C	C6-N1-C2	-6.23	117.81	120.30
24	RA	2183	C	N1-C2-O2	6.23	122.64	118.90
24	YA	1022	C	C5-C6-N1	6.23	124.12	121.00
24	RA	1703	C	C5-C6-N1	6.23	124.12	121.00
1	XA	54	C	C6-N1-C2	-6.23	117.81	120.30
24	RA	1340	U	N3-C2-O2	-6.23	117.84	122.20
24	RA	2029	C	C5-C6-N1	6.23	124.11	121.00

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	1303	C	N3-C2-O2	-6.22	117.55	121.90
24	YA	1492	C	C6-N1-C2	-6.22	117.81	120.30
24	RA	2596	U	N1-C2-O2	6.22	127.15	122.80
24	YA	2227	G	C4-N9-C1'	6.22	134.58	126.50
1	XA	1157	A	C4-N9-C1'	6.21	137.48	126.30
24	YA	2724	U	C2-N1-C1'	6.21	125.16	117.70
1	QA	1163	C	C5-C6-N1	6.21	124.11	121.00
24	YA	665	C	C5-C6-N1	6.21	124.11	121.00
1	XA	455	C	N1-C2-O2	6.21	122.63	118.90
24	RA	487	C	C6-N1-C2	-6.21	117.82	120.30
22	XV	65	U	C5-C6-N1	6.21	125.80	122.70
24	YA	2001	C	C5-C6-N1	6.21	124.10	121.00
24	YA	2725	A	C5-N7-C8	-6.21	100.80	103.90
1	QA	1397	C	N1-C2-O2	6.21	122.62	118.90
24	RA	2201	C	C6-N1-C2	-6.21	117.82	120.30
24	YA	359	C	C6-N1-C2	-6.20	117.82	120.30
24	RA	1180	C	N1-C2-O2	6.20	122.62	118.90
25	RB	44	G	C4-N9-C1'	-6.20	118.44	126.50
24	YA	256	C	C6-N1-C2	-6.20	117.82	120.30
24	RA	66	U	C5-C6-N1	6.20	125.80	122.70
24	YA	1978	U	N1-C2-O2	6.20	127.14	122.80
24	RA	864	C	C6-N1-C2	-6.20	117.82	120.30
24	YA	392	U	N1-C2-O2	6.19	127.14	122.80
24	RA	223	C	N3-C2-O2	-6.19	117.57	121.90
24	RA	2695	C	C6-N1-C2	-6.19	117.82	120.30
1	XA	18	C	C5-C6-N1	6.19	124.09	121.00
24	YA	1914	C	C5-C6-N1	6.18	124.09	121.00
24	YA	2571	C	N1-C2-O2	6.18	122.61	118.90
1	XA	1109	C	N3-C2-O2	-6.18	117.58	121.90
24	RA	1121	C	C5-C6-N1	6.18	124.09	121.00
24	YA	2837	C	C6-N1-C2	-6.17	117.83	120.30
24	RA	853	C	C6-N1-C2	-6.17	117.83	120.30
24	RA	2695	C	N1-C2-O2	6.17	122.60	118.90
24	YA	1644	C	C5-C6-N1	6.17	124.08	121.00
24	RA	1448	C	C6-N1-C2	-6.17	117.83	120.30
24	RA	1483	C	C5-C6-N1	6.17	124.08	121.00
24	YA	802	C	C5-C6-N1	6.17	124.08	121.00
24	YA	1304	C	C5-C6-N1	6.17	124.08	121.00
25	YB	85	G	C8-N9-C4	-6.17	103.93	106.40
1	QA	1383	C	N1-C2-O2	6.16	122.60	118.90
24	RA	1378	G	C8-N9-C1'	-6.16	118.99	127.00
24	RA	2679	C	C6-N1-C2	-6.16	117.83	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	181	C	N1-C2-O2	6.16	122.60	118.90
24	YA	1805	C	C6-N1-C2	-6.16	117.83	120.30
24	YA	1861	C	N1-C2-O2	6.16	122.60	118.90
24	YA	2690	C	C6-N1-C2	-6.16	117.83	120.30
1	QA	1439	C	N1-C2-O2	6.16	122.60	118.90
1	QA	1505	G	N3-C4-N9	-6.16	122.30	126.00
24	RA	719	C	C6-N1-C2	-6.16	117.84	120.30
1	XA	137	C	C5-C6-N1	6.16	124.08	121.00
24	YA	2483	C	C6-N1-C2	-6.16	117.84	120.30
24	YA	340	C	C5-C6-N1	6.16	124.08	121.00
24	YA	1028	C	C6-N1-C2	-6.16	117.84	120.30
1	XA	555	C	C6-N1-C2	-6.16	117.84	120.30
24	YA	1703	C	C5-C6-N1	6.16	124.08	121.00
24	YA	1733	C	C6-N1-C2	-6.16	117.84	120.30
24	RA	562	C	N1-C2-O2	6.15	122.59	118.90
24	RA	1551	C	C6-N1-C2	-6.15	117.84	120.30
1	XA	1147	C	N3-C2-O2	-6.15	117.59	121.90
1	XA	540	G	N7-C8-N9	6.15	116.18	113.10
1	XA	1109	C	C6-N1-C2	-6.15	117.84	120.30
1	QA	1406	U	N3-C2-O2	-6.15	117.89	122.20
24	RA	1930	C	C6-N1-C2	-6.15	117.84	120.30
24	YA	394	C	C6-N1-C2	-6.15	117.84	120.30
24	YA	914	C	C6-N1-C2	-6.15	117.84	120.30
24	RA	2768	C	C6-N1-C2	-6.15	117.84	120.30
1	XA	365	U	C2-N1-C1'	6.15	125.08	117.70
24	YA	2502	G	C4-N9-C1'	6.15	134.49	126.50
1	XA	433	C	C6-N1-C2	-6.14	117.84	120.30
24	YA	637	U	C2-N1-C1'	6.14	125.07	117.70
24	YA	2077	C	C5-C6-N1	6.14	124.07	121.00
24	RA	2196	C	C5-C6-N1	6.14	124.07	121.00
1	XA	314	C	C6-N1-C2	-6.14	117.84	120.30
24	YA	903	C	N1-C2-O2	6.14	122.58	118.90
24	RA	830	A	C8-N9-C4	-6.14	103.34	105.80
24	RA	2627	U	C5-C6-N1	6.14	125.77	122.70
24	RA	1450	C	N1-C2-O2	6.14	122.58	118.90
24	RA	1725	G	C8-N9-C4	-6.14	103.94	106.40
30	RH	7	LEU	CA-CB-CG	6.14	129.42	115.30
24	RA	1379	C	N3-C2-O2	-6.14	117.61	121.90
24	RA	1858	C	N3-C2-O2	-6.13	117.61	121.90
24	YA	2475	C	C6-N1-C2	-6.13	117.85	120.30
24	YA	1574	A	C8-N9-C4	-6.13	103.35	105.80
24	RA	2527	C	C6-N1-C2	-6.13	117.85	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	960	U	C2-N1-C1'	6.13	125.06	117.70
1	QA	1395	C	C6-N1-C2	-6.13	117.85	120.30
24	RA	447	C	N1-C2-O2	6.13	122.58	118.90
24	RA	1087	C	N1-C2-O2	6.13	122.58	118.90
24	YA	562	C	C2-N1-C1'	6.12	125.54	118.80
24	YA	614	C	C6-N1-C2	-6.12	117.85	120.30
24	RA	2483	C	C6-N1-C2	-6.12	117.85	120.30
24	RA	2629	C	N1-C2-O2	6.12	122.57	118.90
24	RA	2472	U	N1-C2-O2	6.12	127.08	122.80
24	RA	1225	C	N1-C2-O2	6.12	122.57	118.90
24	YA	73	A	C8-N9-C4	-6.12	103.35	105.80
1	XA	1439	C	N1-C2-O2	6.12	122.57	118.90
24	YA	482	C	C6-N1-C2	6.12	122.75	120.30
24	YA	1255	A	C5-N7-C8	-6.12	100.84	103.90
24	RA	31	C	C6-N1-C2	-6.11	117.86	120.30
25	YB	10	C	C6-N1-C2	-6.11	117.86	120.30
24	YA	1252	C	C6-N1-C2	-6.11	117.86	120.30
24	RA	1253	C	C6-N1-C2	-6.11	117.86	120.30
24	RA	415	G	N3-C4-N9	-6.11	122.34	126.00
24	RA	1953	U	C5-C4-O4	6.11	129.56	125.90
24	RA	2884	C	C5-C6-N1	6.10	124.05	121.00
24	YA	263	C	N1-C2-O2	6.10	122.56	118.90
25	YB	112	G	C5-C6-O6	-6.10	124.94	128.60
24	RA	1817	A	C4-N9-C1'	6.10	137.28	126.30
24	YA	308	U	N3-C2-O2	-6.10	117.93	122.20
24	YA	1574	A	C5-N7-C8	-6.10	100.85	103.90
24	RA	2631	C	C6-N1-C2	-6.09	117.86	120.30
25	RB	77	U	C5-C6-N1	6.09	125.75	122.70
24	YA	826	U	N3-C2-O2	-6.09	117.93	122.20
24	YA	2397	C	C5-C6-N1	6.09	124.05	121.00
24	RA	914	C	C6-N1-C2	-6.09	117.86	120.30
25	RB	70	C	C6-N1-C2	-6.09	117.86	120.30
24	YA	46	C	C5-C6-N1	6.09	124.05	121.00
1	QA	1528	U	P-O3'-C3'	6.09	127.00	119.70
24	YA	257	C	C5-C6-N1	6.08	124.04	121.00
24	RA	210	A	C4-N9-C1'	6.08	137.24	126.30
24	RA	634	C	C6-N1-C2	-6.08	117.87	120.30
24	YA	2803	A	C2-N3-C4	6.08	113.64	110.60
1	QA	826	C	N1-C2-O2	6.07	122.54	118.90
24	RA	1098	C	C5-C6-N1	6.07	124.04	121.00
24	RA	2185	C	C6-N1-C2	-6.07	117.87	120.30
24	RA	237	G	C4-C5-N7	6.07	113.23	110.80

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	1931	C	N1-C2-O2	6.07	122.54	118.90
24	YA	1339	C	C6-N1-C2	-6.07	117.87	120.30
24	RA	1313	U	N1-C2-O2	6.07	127.05	122.80
24	RA	1722	C	N1-C2-O2	6.07	122.54	118.90
24	YA	1249	A	O4'-C1'-N9	6.07	113.06	108.20
24	RA	1942	C	C6-N1-C2	-6.07	117.87	120.30
24	YA	854	U	N3-C2-O2	-6.07	117.95	122.20
24	YA	2053	A	O4'-C1'-N9	6.07	113.05	108.20
1	QA	1113	C	C5-C6-N1	6.06	124.03	121.00
24	RA	1095	C	C5-C6-N1	6.06	124.03	121.00
24	RA	1095	C	N3-C2-O2	-6.06	117.66	121.90
24	RA	1232	G	C8-N9-C4	-6.06	103.97	106.40
24	RA	1341	C	C6-N1-C2	-6.06	117.88	120.30
24	RA	1819	C	C6-N1-C2	-6.06	117.88	120.30
1	XA	1024	G	N3-C4-C5	-6.06	125.57	128.60
24	RA	1463	C	C5-C6-N1	6.05	124.02	121.00
1	QA	528	C	N1-C2-O2	6.05	122.53	118.90
24	RA	1812	C	N1-C2-O2	6.04	122.53	118.90
24	RA	1817	A	C6-C5-N7	-6.04	128.07	132.30
1	XA	54	C	N3-C2-O2	-6.04	117.67	121.90
24	YA	485	U	C5-C6-N1	6.04	125.72	122.70
24	YA	1225	C	C5-C6-N1	6.04	124.02	121.00
24	YA	2228	G	C8-N9-C1'	-6.04	119.14	127.00
24	YA	2415	C	C5-C6-N1	6.04	124.02	121.00
24	RA	777	C	C6-N1-C2	-6.04	117.88	120.30
24	YA	436	C	C5-C6-N1	6.04	124.02	121.00
24	RA	2711	C	C5-C6-N1	6.03	124.02	121.00
1	QA	1242	C	C5-C6-N1	6.03	124.02	121.00
1	XA	54	C	N1-C2-O2	6.03	122.52	118.90
1	QA	91	C	C5-C6-N1	6.03	124.01	121.00
24	RA	939	C	C6-N1-C2	-6.03	117.89	120.30
24	RA	1574	A	N7-C8-N9	6.03	116.81	113.80
1	XA	1347	G	C4-N9-C1'	-6.03	118.67	126.50
24	YA	462	C	N3-C2-O2	-6.03	117.68	121.90
24	RA	2028	C	C5-C6-N1	6.02	124.01	121.00
24	YA	1805	C	N3-C2-O2	-6.02	117.68	121.90
24	YA	2085	C	C6-N1-C2	-6.02	117.89	120.30
24	RA	2570	C	C6-N1-C2	-6.02	117.89	120.30
1	QA	674	G	N7-C8-N9	6.02	116.11	113.10
1	QA	1270	C	C6-N1-C2	-6.02	117.89	120.30
25	YB	104	A	C2-N3-C4	-6.02	107.59	110.60
24	RA	1225	C	C5-C6-N1	6.01	124.01	121.00

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	690	G	C4-N9-C1'	6.01	134.32	126.50
24	RA	223	C	N1-C2-O2	6.01	122.51	118.90
24	RA	1120	G	C6-C5-N7	-6.01	126.79	130.40
24	YA	989	G	C4-N9-C1'	6.01	134.31	126.50
24	YA	1099	C	C2-N1-C1'	6.01	125.41	118.80
24	YA	1303	C	C5-C6-N1	6.01	124.00	121.00
24	RA	1589	A	O4'-C1'-N9	6.01	113.00	108.20
24	YA	591	U	C5-C6-N1	6.01	125.70	122.70
24	YA	1188	A	N3-C4-N9	-6.00	122.60	127.40
24	YA	1972	G	C8-N9-C1'	-6.00	119.19	127.00
24	YA	359	C	C5-C6-N1	6.00	124.00	121.00
1	XA	749	C	C5-C6-N1	6.00	124.00	121.00
24	YA	1343	C	C6-N1-C2	-6.00	117.90	120.30
24	YA	1644	C	C6-N1-C2	-6.00	117.90	120.30
24	RA	210	A	C8-N9-C1'	-6.00	116.90	127.70
1	XA	433	C	N3-C2-O2	-6.00	117.70	121.90
24	YA	583	C	C6-N1-C2	-6.00	117.90	120.30
24	YA	1861	C	C5-C6-N1	6.00	124.00	121.00
24	YA	624	C	C6-N1-C2	-6.00	117.90	120.30
24	YA	2800	C	C6-N1-C2	-6.00	117.90	120.30
24	RA	139	A	N7-C8-N9	5.99	116.80	113.80
24	YA	2736	C	C6-N1-C2	-5.99	117.90	120.30
24	RA	1559	C	N1-C2-O2	5.99	122.49	118.90
53	Y8	61	LEU	CA-CB-CG	-5.99	101.53	115.30
1	XA	1234	C	C5-C6-N1	5.99	123.99	121.00
1	QA	514	C	C5-C6-N1	5.99	123.99	121.00
24	RA	400	U	C2-N1-C1'	5.99	124.88	117.70
24	YA	2462	A	N1-C2-N3	-5.99	126.31	129.30
24	YA	2854	G	C8-N9-C4	-5.99	104.01	106.40
24	RA	1521	C	C6-N1-C2	-5.98	117.91	120.30
1	QA	449	C	N3-C2-O2	-5.98	117.71	121.90
24	RA	738	C	C6-N1-C2	-5.98	117.91	120.30
24	RA	1916	C	C6-N1-C2	-5.98	117.91	120.30
24	RA	1457	C	N3-C2-O2	-5.98	117.72	121.90
1	XA	556	C	C6-N1-C2	-5.98	117.91	120.30
24	YA	857	U	C5-C6-N1	5.98	125.69	122.70
24	YA	462	C	N1-C2-O2	5.97	122.48	118.90
24	YA	2354	C	C5-C6-N1	5.97	123.99	121.00
24	RA	990	A	C4-C5-N7	5.97	113.69	110.70
24	RA	2406	C	N1-C2-O2	5.97	122.48	118.90
1	XA	1158	C	C5-C6-N1	5.97	123.99	121.00
24	YA	1398	U	N3-C2-O2	-5.97	118.02	122.20

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
25	YB	71	C	N1-C2-O2	5.97	122.48	118.90
1	QA	1109	C	N1-C2-O2	5.97	122.48	118.90
24	YA	853	C	N1-C2-O2	5.97	122.48	118.90
24	RA	864	C	C5-C6-N1	5.97	123.98	121.00
24	YA	1866	G	N3-C4-C5	-5.97	125.62	128.60
1	XA	347	G	O4'-C1'-N9	5.96	112.97	108.20
24	RA	2196	C	N3-C2-O2	-5.96	117.73	121.90
24	RA	485	U	C5-C6-N1	5.96	125.68	122.70
24	YA	257	C	C6-N1-C2	-5.96	117.92	120.30
24	YA	2121	U	C5-C6-N1	5.96	125.68	122.70
1	QA	266	G	N3-C4-C5	5.96	131.58	128.60
24	YA	1625	U	N1-C2-O2	5.96	126.97	122.80
24	YA	2177	G	N3-C4-N9	5.96	129.57	126.00
25	YB	8	U	C5-C6-N1	5.96	125.68	122.70
1	QA	1263	C	C6-N1-C2	-5.95	117.92	120.30
24	RA	958	C	C5-C6-N1	5.95	123.97	121.00
24	RA	1983	C	N1-C2-O2	5.95	122.47	118.90
1	XA	891	U	N1-C2-O2	5.95	126.96	122.80
24	YA	2028	C	C5-C6-N1	5.95	123.97	121.00
24	RA	1596	C	N1-C2-O2	5.94	122.47	118.90
1	QA	943	U	N3-C2-O2	-5.94	118.04	122.20
24	YA	555	G	C5-C6-O6	5.94	132.16	128.60
24	YA	227	C	C6-N1-C2	-5.94	117.92	120.30
24	YA	453	C	N1-C2-O2	5.94	122.46	118.90
24	YA	2129	C	N1-C2-O2	5.94	122.46	118.90
24	RA	1137	G	C8-N9-C4	-5.94	104.03	106.40
24	RA	1796	C	C6-N1-C2	-5.94	117.93	120.30
1	QA	410	G	C8-N9-C4	-5.93	104.03	106.40
24	RA	143	C	C6-N1-C2	-5.93	117.93	120.30
24	YA	220	C	C5-C6-N1	5.93	123.97	121.00
1	XA	932	C	C6-N1-C2	-5.93	117.93	120.30
24	YA	1641	G	N1-C6-O6	-5.93	116.34	119.90
24	YA	1817	A	C4-N9-C1'	5.93	136.98	126.30
24	YA	826	U	N1-C2-O2	5.93	126.95	122.80
24	YA	1068	G	N9-C4-C5	5.93	107.77	105.40
24	YA	1436	U	N3-C2-O2	-5.93	118.05	122.20
24	YA	2074	G	C4-C5-N7	5.93	113.17	110.80
24	YA	565	C	C6-N1-C2	-5.93	117.93	120.30
24	YA	801	C	C5-C6-N1	5.93	123.97	121.00
1	QA	932	C	C2-N1-C1'	5.93	125.32	118.80
1	XA	1141	C	N1-C2-N3	5.93	123.35	119.20
24	YA	730	C	C5-C6-N1	5.93	123.96	121.00

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	955	U	C5-C6-N1	5.92	125.66	122.70
24	YA	893	C	N1-C2-O2	5.92	122.45	118.90
24	YA	1577	C	C6-N1-C2	-5.92	117.93	120.30
24	YA	1444	C	C5-C6-N1	5.92	123.96	121.00
24	YA	2319	G	C6-C5-N7	-5.92	126.85	130.40
24	YA	2784	C	C6-N1-C2	-5.92	117.93	120.30
1	QA	381	C	N1-C2-O2	5.91	122.45	118.90
1	XA	536	C	N1-C2-O2	5.91	122.45	118.90
24	YA	1045	U	N3-C2-O2	-5.91	118.06	122.20
24	YA	1510	C	C6-N1-C2	-5.91	117.94	120.30
24	RA	1824	C	C6-N1-C2	-5.91	117.94	120.30
24	RA	2354	C	C5-C6-N1	5.91	123.95	121.00
1	XA	1439	C	C5-C6-N1	5.90	123.95	121.00
24	YA	447	C	C5-C6-N1	5.90	123.95	121.00
24	RA	1121	C	C6-N1-C2	-5.90	117.94	120.30
1	XA	554	C	C5-C6-N1	5.90	123.95	121.00
1	XA	1310	G	N3-C4-N9	5.90	129.54	126.00
24	YA	646	A	N7-C8-N9	5.90	116.75	113.80
24	YA	12	U	C6-N1-C2	-5.90	117.46	121.00
24	RA	2272	C	C6-N1-C2	-5.90	117.94	120.30
1	XA	486	U	N1-C2-O2	5.90	126.93	122.80
1	QA	337	C	C5-C6-N1	5.89	123.94	121.00
24	RA	2714	U	C2-N1-C1'	-5.89	110.63	117.70
24	YA	454	U	N3-C2-O2	-5.89	118.08	122.20
24	RA	1463	C	C6-N1-C2	-5.88	117.95	120.30
24	RA	1556	A	C2-N3-C4	5.88	113.54	110.60
1	XA	435	C	C5-C6-N1	5.88	123.94	121.00
25	YB	28	C	C6-N1-C2	-5.88	117.95	120.30
24	RA	914	C	N3-C2-O2	-5.88	117.78	121.90
1	QA	1225	A	N7-C8-N9	5.88	116.74	113.80
25	YB	28	C	C5-C6-N1	5.88	123.94	121.00
24	RA	340	C	C5-C6-N1	5.88	123.94	121.00
24	RA	1972	G	O4'-C1'-N9	5.88	112.90	108.20
24	RA	2260	C	C6-N1-C2	-5.88	117.95	120.30
24	RA	2772	G	N3-C4-N9	5.88	129.53	126.00
1	XA	233	C	C6-N1-C2	-5.88	117.95	120.30
24	RA	2835	C	C6-N1-C2	-5.87	117.95	120.30
24	YA	2877	G	OP2-P-O3'	5.87	118.12	105.20
1	XA	827	U	N3-C2-O2	-5.87	118.09	122.20
24	YA	1271	G	N7-C8-N9	5.87	116.04	113.10
1	QA	1279	A	C8-N9-C4	-5.87	103.45	105.80
25	RB	47	C	N1-C2-O2	5.87	122.42	118.90

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	725	C	C5-C6-N1	5.87	123.93	121.00
1	XA	528	C	C5-C6-N1	5.87	123.93	121.00
24	YA	2668	U	N3-C4-O4	5.87	123.51	119.40
24	YA	903	C	N3-C2-O2	-5.86	117.80	121.90
24	RA	2196	C	N1-C2-O2	5.86	122.42	118.90
24	RA	744	C	C6-N1-C2	-5.86	117.96	120.30
24	YA	607	C	C5-C6-N1	5.86	123.93	121.00
25	YB	31	C	N1-C2-O2	5.86	122.42	118.90
24	RA	872	C	C5-C6-N1	5.86	123.93	121.00
24	RA	1731	C	C6-N1-C2	-5.86	117.96	120.30
24	YA	267	C	C6-N1-C2	-5.86	117.96	120.30
24	YA	537	G	N1-C6-O6	-5.86	116.39	119.90
24	RA	1120	G	N9-C4-C5	-5.86	103.06	105.40
24	RA	2406	C	N3-C2-O2	-5.86	117.80	121.90
1	XA	623	C	C6-N1-C2	-5.85	117.96	120.30
1	QA	91	C	N1-C2-O2	5.85	122.41	118.90
24	YA	1544	C	N1-C2-O2	5.85	122.41	118.90
24	RA	1448	C	C5-C6-N1	5.85	123.92	121.00
24	YA	2454	C	C6-N1-C2	-5.85	117.96	120.30
1	QA	433	C	C6-N1-C2	-5.85	117.96	120.30
24	YA	1398	U	C6-N1-C2	-5.85	117.49	121.00
24	RA	1551	C	C5-C6-N1	5.84	123.92	121.00
25	RB	37	C	N3-C2-O2	-5.84	117.81	121.90
24	YA	258	U	C5-C6-N1	5.84	125.62	122.70
24	RA	2030	C	C5-C6-N1	5.84	123.92	121.00
1	QA	686	U	N3-C2-O2	-5.84	118.11	122.20
22	QV	4	U	O4'-C1'-N1	5.84	112.87	108.20
24	RA	2472	U	N3-C2-O2	-5.84	118.11	122.20
24	RA	1716	A	C2-N3-C4	5.84	113.52	110.60
24	YA	2095	C	C6-N1-C2	-5.84	117.96	120.30
24	RA	1120	G	C4-C5-N7	5.83	113.13	110.80
1	XA	980	C	N1-C2-O2	5.83	122.40	118.90
24	YA	1687	C	C6-N1-C2	-5.83	117.97	120.30
1	QA	1452	C	N3-C2-O2	-5.83	117.82	121.90
1	XA	137	C	C6-N1-C2	-5.83	117.97	120.30
1	XA	169	C	C6-N1-C2	-5.83	117.97	120.30
24	YA	796	C	N3-C2-O2	-5.83	117.82	121.90
24	YA	1436	U	C5-C6-N1	5.83	125.61	122.70
24	YA	2697	G	N3-C4-N9	-5.83	122.50	126.00
1	XA	528	C	N1-C2-O2	5.83	122.40	118.90
24	YA	31	C	C5-C6-N1	5.83	123.91	121.00
24	YA	1136	U	N1-C2-O2	5.83	126.88	122.80

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	1186	U	C2-N1-C1'	-5.82	110.71	117.70
25	YB	84	C	C6-N1-C2	-5.82	117.97	120.30
24	YA	178	G	N1-C6-O6	-5.82	116.41	119.90
1	QA	1161	C	N1-C2-O2	5.82	122.39	118.90
1	QA	1502	A	N9-C4-C5	-5.82	103.47	105.80
24	YA	203	G	N1-C6-O6	-5.82	116.41	119.90
24	RA	854	U	N3-C2-O2	-5.82	118.13	122.20
1	XA	792	A	O4'-C1'-N9	5.82	112.85	108.20
1	XA	1310	G	C6-C5-N7	-5.82	126.91	130.40
24	YA	634	C	C5-C6-N1	5.82	123.91	121.00
24	YA	708	C	C5-C6-N1	5.82	123.91	121.00
24	YA	1915	C	C6-N1-C2	-5.82	117.97	120.30
25	RB	27	C	N3-C2-O2	-5.81	117.83	121.90
24	YA	630	U	C2-N3-C4	5.81	130.49	127.00
1	QA	91	C	C6-N1-C2	-5.81	117.97	120.30
24	YA	482	C	N3-C4-C5	5.81	124.22	121.90
1	QA	1132	C	N3-C2-O2	-5.81	117.83	121.90
24	RA	2570	C	C5-C6-N1	5.81	123.90	121.00
24	YA	2780	C	C6-N1-C2	-5.81	117.98	120.30
24	YA	1378	G	C4-C5-N7	5.81	113.12	110.80
24	YA	2404	A	C8-N9-C4	-5.80	103.48	105.80
1	QA	455	C	N1-C2-O2	5.80	122.38	118.90
24	RA	1515	C	C6-N1-C2	-5.80	117.98	120.30
24	RA	1921	G	C8-N9-C1'	5.80	134.54	127.00
24	RA	2878	A	C8-N9-C4	-5.80	103.48	105.80
24	YA	1457	C	C2-N1-C1'	5.80	125.18	118.80
1	XA	1362(B)	C	N3-C2-O2	-5.80	117.84	121.90
24	YA	2111	U	C5-C6-N1	5.80	125.60	122.70
24	RA	2241	C	C6-N1-C2	-5.80	117.98	120.30
24	YA	607	C	N1-C2-O2	5.80	122.38	118.90
1	QA	1260	C	N3-C2-O2	-5.80	117.84	121.90
24	YA	457	G	C8-N9-C4	-5.80	104.08	106.40
24	YA	579	G	C6-C5-N7	-5.80	126.92	130.40
24	RA	1726	U	N3-C2-O2	-5.79	118.14	122.20
24	YA	485	U	C6-N1-C2	-5.79	117.52	121.00
1	QA	1270	C	C5-C6-N1	5.79	123.90	121.00
24	RA	1559	C	C6-N1-C2	-5.79	117.98	120.30
24	YA	1423	G	C8-N9-C4	-5.79	104.08	106.40
24	YA	2256	U	N3-C4-O4	5.79	123.45	119.40
24	YA	2004	C	C6-N1-C2	-5.79	117.98	120.30
24	YA	2678	C	C6-N1-C2	-5.79	117.98	120.30
24	YA	646	A	C2-N3-C4	-5.79	107.70	110.60

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	1141	A	N3-C4-C5	-5.79	122.75	126.80
24	RA	1691	C	N1-C2-O2	5.79	122.37	118.90
1	XA	1279	A	N7-C8-N9	5.79	116.69	113.80
24	YA	2329	C	N1-C2-O2	5.79	122.37	118.90
1	QA	697	U	N3-C2-O2	-5.78	118.15	122.20
24	YA	666	C	C6-N1-C2	-5.78	117.99	120.30
24	YA	1595	C	C6-N1-C2	-5.78	117.99	120.30
24	RA	1806	U	N3-C4-O4	5.78	123.45	119.40
24	YA	1927	C	C5-C6-N1	5.78	123.89	121.00
24	RA	1425	A	C4-C5-N7	5.78	113.59	110.70
25	RB	44	G	C8-N9-C1'	5.78	134.51	127.00
1	QA	106	C	C6-N1-C2	-5.78	117.99	120.30
24	YA	564	G	C4-N9-C1'	5.78	134.01	126.50
24	YA	975	U	N1-C2-O2	5.78	126.84	122.80
24	YA	1038	C	C6-N1-C2	-5.78	117.99	120.30
24	RA	2638	C	C5-C6-N1	5.78	123.89	121.00
24	YA	419	C	C6-N1-C2	-5.78	117.99	120.30
24	RA	256	C	C6-N1-C2	-5.77	117.99	120.30
24	YA	1031	C	N1-C2-O2	5.77	122.36	118.90
24	YA	528	A	P-O3'-C3'	5.77	126.63	119.70
24	YA	2083	G	C4-C5-C6	5.77	122.26	118.80
24	RA	2603	C	C6-N1-C2	-5.77	117.99	120.30
24	RA	863	C	C6-N1-C2	-5.77	117.99	120.30
24	YA	1683	C	C6-N1-C2	-5.77	117.99	120.30
1	QA	943	U	N1-C2-O2	5.76	126.83	122.80
24	YA	1645	C	C6-N1-C2	-5.76	117.99	120.30
25	YB	92	G	N1-C2-N3	5.76	127.36	123.90
24	RA	1360	C	C2-N1-C1'	5.76	125.14	118.80
24	RA	2354	C	C6-N1-C2	-5.76	118.00	120.30
24	YA	872	C	C5-C6-N1	5.76	123.88	121.00
24	YA	1509	C	N1-C2-O2	5.76	122.36	118.90
24	YA	2273	C	C6-N1-C2	-5.76	118.00	120.30
24	YA	1867	C	C5-C6-N1	5.76	123.88	121.00
24	RA	1936	C	C6-N1-C2	-5.75	118.00	120.30
1	QA	1411	C	C6-N1-C2	-5.75	118.00	120.30
24	RA	1031	C	C5-C6-N1	5.75	123.88	121.00
24	RA	1916	C	N1-C2-O2	5.75	122.35	118.90
1	XA	330	C	C6-N1-C2	-5.75	118.00	120.30
1	XA	1520	G	C4-C5-N7	5.75	113.10	110.80
24	YA	1089	C	C6-N1-C2	-5.75	118.00	120.30
24	YA	2054	G	N3-C4-N9	-5.75	122.55	126.00
24	YA	844	C	C6-N1-C2	-5.75	118.00	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	75	C	C6-N1-C2	-5.75	118.00	120.30
24	YA	217	A	C8-N9-C4	-5.75	103.50	105.80
24	RA	1921	G	C4-N9-C1'	-5.74	119.03	126.50
24	YA	2854	G	N3-C4-C5	-5.74	125.73	128.60
1	QA	1323	G	C6-C5-N7	-5.74	126.95	130.40
24	RA	1022	C	C5-C6-N1	5.74	123.87	121.00
24	RA	2598	C	C6-N1-C2	-5.74	118.00	120.30
24	RA	275	C	N1-C2-O2	5.74	122.34	118.90
24	YA	863	C	C5-C6-N1	5.74	123.87	121.00
24	YA	1002	A	C5-C6-N6	-5.74	119.11	123.70
24	RA	1687	C	N1-C2-O2	5.74	122.34	118.90
24	RA	2222	C	C5-C6-N1	5.74	123.87	121.00
24	YA	2014	G	C2'-C3'-O3'	5.74	122.88	113.70
1	XA	1028(B)	C	N3-C2-O2	-5.74	117.88	121.90
1	QA	904	C	C6-N1-C2	-5.74	118.01	120.30
24	YA	842	C	C6-N1-C2	-5.74	118.00	120.30
24	YA	1514	C	C6-N1-C2	-5.74	118.01	120.30
24	YA	1645	C	C5-C6-N1	5.74	123.87	121.00
24	YA	2631	C	C6-N1-C2	-5.74	118.01	120.30
24	RA	2407	C	C6-N1-C2	-5.73	118.01	120.30
24	YA	2083	G	C6-C5-N7	-5.73	126.96	130.40
24	YA	2324	U	C6-N1-C2	-5.73	117.56	121.00
24	RA	665	C	C6-N1-C2	-5.73	118.01	120.30
24	YA	1972	G	C4-C5-N7	5.73	113.09	110.80
1	XA	1028(A)	C	N3-C2-O2	-5.73	117.89	121.90
24	RA	1493	C	C6-N1-C2	-5.73	118.01	120.30
1	QA	252	U	N1-C2-O2	5.73	126.81	122.80
1	QA	780	A	N7-C8-N9	5.73	116.66	113.80
24	RA	719	C	C5-C6-N1	5.73	123.86	121.00
24	RA	1924	C	N3-C4-N4	-5.72	113.99	118.00
1	XA	656	C	C6-N1-C2	-5.72	118.01	120.30
24	YA	166	G	C4-N9-C1'	5.72	133.94	126.50
24	YA	284	G	C8-N9-C4	-5.72	104.11	106.40
24	RA	256	C	C5-C6-N1	5.72	123.86	121.00
24	RA	805	C	C6-N1-C2	-5.72	118.01	120.30
24	RA	2504	U	N3-C2-O2	-5.72	118.20	122.20
44	RZ	59	LEU	CA-CB-CG	5.72	128.45	115.30
24	YA	1531	G	C4-C5-N7	5.72	113.09	110.80
24	YA	2774	G	N9-C1'-C2'	-5.72	105.71	112.00
1	QA	405	U	N1-C2-O2	5.72	126.80	122.80
24	RA	1972	G	N7-C8-N9	5.72	115.96	113.10
1	QA	1430	C	C5-C6-N1	5.71	123.86	121.00

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	232	U	C5-C6-N1	5.71	125.56	122.70
24	RA	2618	C	N1-C2-O2	5.71	122.33	118.90
24	YA	1092	A	C2-N3-C4	5.71	113.46	110.60
1	XA	1452	C	N3-C2-O2	-5.71	117.90	121.90
24	RA	2807	C	N1-C2-O2	5.71	122.33	118.90
24	YA	2329	C	N3-C2-O2	-5.71	117.90	121.90
1	QA	618	C	N1-C2-O2	5.71	122.32	118.90
1	QA	754	C	C6-N1-C1'	-5.70	113.96	120.80
24	RA	897	C	C5-C6-N1	5.70	123.85	121.00
24	RA	1421	C	C5-C6-N1	5.70	123.85	121.00
24	RA	2857	U	C5-C6-N1	5.70	125.55	122.70
24	YA	579	G	C8-N9-C4	-5.70	104.12	106.40
24	YA	830	A	C4-C5-N7	5.70	113.55	110.70
1	XA	1246	C	C6-N1-C2	-5.70	118.02	120.30
24	RA	181	C	C5-C6-N1	5.70	123.85	121.00
24	YA	2091	G	N1-C6-O6	-5.70	116.48	119.90
1	XA	280	C	C6-N1-C2	5.69	122.58	120.30
24	YA	1372	U	N1-C2-O2	5.69	126.78	122.80
1	QA	1109	C	N3-C2-O2	-5.69	117.92	121.90
1	XA	1161	C	N3-C2-O2	-5.69	117.92	121.90
24	RA	1110	C	N3-C4-N4	5.69	121.98	118.00
24	YA	1205	U	N3-C2-O2	-5.69	118.22	122.20
24	RA	729	G	N1-C6-O6	-5.69	116.49	119.90
24	YA	280	C	N1-C2-O2	5.69	122.31	118.90
24	YA	2397	C	C6-N1-C2	-5.69	118.03	120.30
24	RA	2201	C	N3-C2-O2	-5.69	117.92	121.90
24	YA	907	U	C2-N1-C1'	5.69	124.52	117.70
1	QA	37	U	N3-C2-O2	-5.68	118.22	122.20
1	QA	1301	U	C5-C6-N1	5.68	125.54	122.70
24	YA	512	C	C6-N1-C2	-5.68	118.03	120.30
24	RA	778	C	C5-C6-N1	5.68	123.84	121.00
24	RA	2890	C	C6-N1-C2	-5.68	118.03	120.30
25	RB	22	U	N3-C2-O2	-5.68	118.22	122.20
1	XA	543	C	C6-N1-C2	-5.68	118.03	120.30
24	RA	842	C	C6-N1-C2	-5.68	118.03	120.30
24	RA	1579	C	C2-N1-C1'	5.68	125.05	118.80
24	YA	286	C	N1-C2-O2	5.68	122.31	118.90
24	YA	579	G	N3-C4-N9	5.68	129.41	126.00
24	YA	1225	C	C2-N1-C1'	5.68	125.05	118.80
1	QA	497	U	N3-C2-O2	-5.67	118.23	122.20
1	XA	749	C	N1-C2-O2	5.67	122.30	118.90
24	YA	1952	G	O4'-C1'-N9	5.67	112.74	108.20

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	1002	A	N1-C6-N6	5.67	122.00	118.60
25	YB	107	U	N3-C2-O2	-5.67	118.23	122.20
24	RA	2519	C	C6-N1-C2	-5.67	118.03	120.30
24	YA	1432	C	C6-N1-C2	-5.67	118.03	120.30
24	RA	1931	C	C5-C6-N1	5.67	123.83	121.00
1	XA	754	C	C6-N1-C1'	-5.67	114.00	120.80
1	QA	1383	C	N3-C2-O2	-5.67	117.93	121.90
24	RA	611	U	C5-C6-N1	5.67	125.53	122.70
24	RA	975	U	N3-C2-O2	-5.67	118.23	122.20
24	RA	1552	C	N1-C2-O2	5.67	122.30	118.90
24	YA	1339	C	C5-C6-N1	5.67	123.83	121.00
24	RA	875	U	C6-N1-C2	-5.67	117.60	121.00
1	XA	358	U	C5'-C4'-C3'	5.67	125.06	116.00
1	XA	1397	C	C6-N1-C2	-5.67	118.03	120.30
1	QA	335	C	C6-N1-C2	-5.66	118.03	120.30
24	RA	1583	C	C6-N1-C2	-5.66	118.03	120.30
24	YA	2228	G	C6-C5-N7	-5.66	127.00	130.40
24	YA	2835	C	N3-C2-O2	-5.66	117.94	121.90
33	YO	8	LEU	CA-CB-CG	5.66	128.32	115.30
1	QA	1420	C	C6-N1-C2	-5.66	118.04	120.30
24	RA	1730	C	C6-N1-C2	-5.66	118.04	120.30
24	YA	1002	A	C4-C5-N7	5.66	113.53	110.70
24	RA	1572	G	C6-C5-N7	-5.66	127.00	130.40
24	YA	73	A	C5-N7-C8	-5.66	101.07	103.90
24	RA	1230	C	C6-N1-C2	-5.66	118.04	120.30
24	YA	420	C	C6-N1-C2	-5.65	118.04	120.30
24	YA	1921	G	N1-C2-N2	-5.65	111.11	116.20
1	XA	435	C	N1-C2-O2	5.65	122.29	118.90
24	RA	1521	C	C5-C6-N1	5.65	123.82	121.00
24	RA	2700	U	C2-N1-C1'	5.65	124.48	117.70
25	RB	31	C	C6-N1-C1'	-5.65	114.02	120.80
14	XN	44	LEU	CA-CB-CG	5.65	128.28	115.30
1	QA	1322	C	N3-C2-O2	-5.64	117.95	121.90
24	YA	1176	U	P-O3'-C3'	5.64	126.47	119.70
24	RA	1263	C	C6-N1-C2	-5.64	118.04	120.30
1	XA	620	C	N3-C2-O2	-5.64	117.95	121.90
24	RA	1548	C	N1-C2-O2	5.64	122.28	118.90
24	YA	46	C	N1-C2-O2	5.64	122.28	118.90
24	YA	893	C	N3-C2-O2	-5.64	117.95	121.90
24	YA	1196	C	C6-N1-C2	-5.64	118.05	120.30
24	YA	141	C	C5-C6-N1	5.64	123.82	121.00
24	YA	189	U	N3-C2-O2	-5.63	118.25	122.20

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	2029	C	C6-N1-C2	-5.63	118.05	120.30
24	YA	939	C	C5-C6-N1	5.63	123.82	121.00
24	YA	2367	C	N1-C2-O2	5.63	122.28	118.90
24	RA	1155	C	N3-C2-O2	-5.63	117.96	121.90
24	RA	1154	U	C6-N1-C2	-5.62	117.62	121.00
1	XA	137	C	N1-C2-O2	5.62	122.28	118.90
1	QA	1027	C	C5-C6-N1	5.62	123.81	121.00
24	RA	555	G	C5-C6-O6	5.62	131.97	128.60
1	XA	186(B)	C	C6-N1-C2	-5.62	118.05	120.30
1	XA	307	C	C6-N1-C2	-5.62	118.05	120.30
24	YA	1341	C	C6-N1-C2	-5.62	118.05	120.30
1	QA	1121	U	C5-C6-N1	5.62	125.51	122.70
24	RA	1924	C	N1-C2-O2	5.62	122.27	118.90
24	YA	132	C	C6-N1-C2	-5.62	118.05	120.30
24	YA	1000	C	C6-N1-C2	-5.62	118.05	120.30
24	YA	354	A	C2-N3-C4	-5.62	107.79	110.60
1	XA	449	C	N3-C2-O2	-5.62	117.97	121.90
24	YA	182	U	N3-C2-O2	-5.62	118.27	122.20
24	RA	1956	C	N1-C2-O2	5.61	122.27	118.90
24	RA	1972	G	C8-N9-C4	-5.61	104.15	106.40
24	RA	2884	C	N1-C2-O2	5.61	122.27	118.90
24	YA	2798	C	C5-C6-N1	5.61	123.81	121.00
1	QA	717	C	C5-C6-N1	5.61	123.81	121.00
1	QA	1378	C	N3-C2-O2	-5.61	117.97	121.90
24	RA	1560	U	C2-N1-C1'	5.61	124.43	117.70
24	YA	2550	C	C6-N1-C2	-5.61	118.06	120.30
1	QA	989	C	N1-C2-O2	5.61	122.27	118.90
24	RA	2859	U	C2-N1-C1'	-5.61	110.97	117.70
25	RB	77	U	N3-C2-O2	-5.61	118.28	122.20
24	RA	1067	A	N3-C4-N9	-5.60	122.92	127.40
24	RA	2453	C	C6-N1-C2	-5.60	118.06	120.30
24	YA	354	A	C5-N7-C8	-5.60	101.10	103.90
24	YA	2851	C	C5-C6-N1	5.60	123.80	121.00
1	XA	1097	C	C5-C6-N1	5.60	123.80	121.00
24	YA	1805	C	N1-C2-O2	5.60	122.26	118.90
24	YA	2587	C	C6-N1-C2	-5.60	118.06	120.30
24	YA	1121	C	N1-C2-O2	5.60	122.26	118.90
24	YA	2095	C	C5-C6-N1	5.60	123.80	121.00
24	YA	2588	G	C4-N9-C1'	5.60	133.78	126.50
1	XA	390	C	C6-N1-C2	-5.60	118.06	120.30
24	RA	1548	C	C2-N1-C1'	5.60	124.96	118.80
24	YA	1279	C	C5-C6-N1	5.60	123.80	121.00

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	1528	U	C5-C6-N1	5.60	125.50	122.70
24	YA	2570	C	C5-C6-N1	5.60	123.80	121.00
24	YA	454	U	N1-C2-O2	5.60	126.72	122.80
1	XA	848	C	N1-C2-O2	5.59	122.26	118.90
24	YA	885	C	C5-C6-N1	5.59	123.80	121.00
24	YA	1188	A	C2-N3-C4	-5.59	107.80	110.60
24	YA	113	C	C6-N1-C2	-5.59	118.06	120.30
24	YA	2250	G	N3-C4-C5	-5.59	125.80	128.60
24	YA	1836	U	C5-C6-N1	5.59	125.50	122.70
25	YB	104	A	C5-N7-C8	-5.59	101.10	103.90
24	RA	1002	A	C4-C5-N7	5.59	113.49	110.70
24	YA	436	C	N1-C2-O2	5.59	122.25	118.90
24	YA	1141	A	N3-C4-N9	5.59	131.87	127.40
22	XV	65	U	C6-N1-C2	-5.59	117.65	121.00
1	QA	596	C	C6-N1-C2	-5.59	118.07	120.30
24	RA	1154	U	C5-C6-N1	5.59	125.49	122.70
1	QA	697	U	N1-C2-O2	5.58	126.71	122.80
24	RA	706	C	C5-C6-N1	5.58	123.79	121.00
24	RA	187	C	N1-C2-O2	5.58	122.25	118.90
24	YA	1302	G	C4-C5-N7	5.58	113.03	110.80
24	YA	2570	C	C6-N1-C2	-5.58	118.07	120.30
1	XA	1383	C	N1-C2-O2	5.58	122.25	118.90
24	YA	157	U	N3-C2-O2	-5.58	118.30	122.20
24	RA	1452	U	C5-C6-N1	5.58	125.49	122.70
24	RA	2743	C	C6-N1-C2	-5.58	118.07	120.30
24	YA	2752	U	N1-C2-O2	5.58	126.70	122.80
24	RA	1983	C	N3-C2-O2	-5.57	118.00	121.90
24	YA	844	C	C5-C6-N1	5.57	123.79	121.00
24	YA	1861	C	C6-N1-C2	-5.57	118.07	120.30
24	YA	2354	C	C6-N1-C2	-5.57	118.07	120.30
24	YA	1336	C	C6-N1-C2	-5.57	118.07	120.30
24	YA	604	C	C5-C6-N1	5.57	123.78	121.00
24	YA	190	C	C6-N1-C2	-5.57	118.07	120.30
24	RA	2884	C	C6-N1-C2	-5.57	118.07	120.30
1	QA	1412	C	C6-N1-C2	-5.57	118.07	120.30
24	RA	482	C	C2-N1-C1'	5.57	124.92	118.80
24	YA	73	A	C2-N3-C4	-5.57	107.82	110.60
24	YA	2307	C	C6-N1-C2	-5.57	118.07	120.30
24	YA	2807	C	C6-N1-C2	-5.57	118.07	120.30
24	RA	2118	U	N1-C2-O2	5.56	126.69	122.80
24	YA	463	C	C6-N1-C2	-5.56	118.08	120.30
1	XA	191(B)	G	N1-C6-O6	-5.56	116.56	119.90

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	2028	C	N1-C2-O2	5.56	122.24	118.90
24	YA	1927	C	C6-N1-C2	-5.56	118.08	120.30
25	YB	114	G	N1-C2-N2	-5.56	111.20	116.20
1	XA	1113	C	C5-C6-N1	5.56	123.78	121.00
24	YA	830	A	C2-N3-C4	-5.56	107.82	110.60
24	RA	826	U	C5-C6-N1	5.55	125.48	122.70
24	RA	1162	C	N3-C2-O2	-5.55	118.01	121.90
1	XA	1157	A	C8-N9-C1'	-5.55	117.71	127.70
24	YA	961	C	N1-C2-O2	5.55	122.23	118.90
24	YA	2456	G	N1-C6-O6	-5.55	116.57	119.90
24	RA	190	C	C6-N1-C2	-5.55	118.08	120.30
24	RA	511	C	C6-N1-C2	-5.55	118.08	120.30
24	RA	1707	C	C6-N1-C2	-5.55	118.08	120.30
24	YA	1901	C	C6-N1-C2	-5.55	118.08	120.30
24	RA	1037	C	C6-N1-C2	-5.55	118.08	120.30
24	RA	2904	U	N1-C2-O2	5.55	126.68	122.80
1	XA	1038	C	C5-C6-N1	5.55	123.77	121.00
24	YA	1279	C	C6-N1-C2	-5.54	118.08	120.30
24	YA	2218	C	C5-C6-N1	5.54	123.77	121.00
24	RA	1810	U	C6-N1-C1'	-5.54	113.44	121.20
1	QA	699	C	C5-C6-N1	5.54	123.77	121.00
1	QA	936	C	N1-C2-O2	5.54	122.22	118.90
24	RA	944	C	C5-C6-N1	5.54	123.77	121.00
24	RA	363	U	C5-C6-N1	5.54	125.47	122.70
24	RA	853	C	C5-C6-N1	5.54	123.77	121.00
24	YA	1760	U	N3-C2-O2	-5.54	118.32	122.20
24	RA	777	C	C5-C6-N1	5.54	123.77	121.00
24	YA	2528	G	N3-C4-N9	5.54	129.32	126.00
24	RA	1707	C	N1-C2-O2	5.53	122.22	118.90
24	YA	104	C	C6-N1-C2	-5.53	118.09	120.30
24	RA	1031	C	C6-N1-C2	-5.53	118.09	120.30
1	XA	979	C	C6-N1-C2	-5.53	118.09	120.30
24	RA	1921	G	N1-C2-N3	5.53	127.22	123.90
25	RB	77	U	N1-C2-O2	5.53	126.67	122.80
1	XA	789	U	C2-N1-C1'	5.53	124.33	117.70
1	XA	810	C	C6-N1-C2	-5.52	118.09	120.30
24	YA	65	C	C5-C6-N1	5.52	123.76	121.00
1	QA	985	C	C6-N1-C2	-5.52	118.09	120.30
24	RA	1154	U	C2-N1-C1'	5.52	124.33	117.70
44	RZ	157	LEU	CA-CB-CG	5.52	128.00	115.30
24	YA	1408	C	C6-N1-C2	-5.52	118.09	120.30
24	RA	1722	C	C5-C6-N1	5.52	123.76	121.00

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	126	C	C6-N1-C2	-5.52	118.09	120.30
24	RA	1635	C	C6-N1-C2	-5.52	118.09	120.30
24	YA	1067	A	N3-C4-C5	5.52	130.66	126.80
24	YA	1225	C	N1-C2-O2	5.52	122.21	118.90
24	YA	1408	C	C5-C6-N1	5.52	123.76	121.00
24	YA	2790	G	C8-N9-C4	-5.52	104.19	106.40
24	YA	1032	C	C5-C6-N1	5.51	123.76	121.00
24	RA	1867	C	C6-N1-C2	-5.51	118.09	120.30
24	RA	20	C	C6-N1-C2	-5.51	118.09	120.30
24	RA	487	C	C5-C6-N1	5.51	123.76	121.00
24	RA	2678	C	C6-N1-C2	-5.51	118.09	120.30
24	YA	1423	G	N7-C8-N9	5.51	115.86	113.10
31	YI	14	ASP	CB-CG-OD1	5.51	123.26	118.30
24	RA	893	C	N1-C2-O2	5.51	122.21	118.90
24	RA	1725	G	N7-C8-N9	5.51	115.86	113.10
24	YA	1136	U	N3-C2-O2	-5.51	118.34	122.20
24	YA	2736	C	C5-C6-N1	5.51	123.75	121.00
24	RA	2772	G	N9-C4-C5	-5.51	103.20	105.40
24	YA	211	A	C5'-C4'-O4'	5.51	115.71	109.10
24	YA	862	C	C5-C6-N1	5.51	123.75	121.00
24	YA	1904	C	C2-N1-C1'	5.51	124.86	118.80
24	YA	2479	C	N3-C2-O2	-5.51	118.04	121.90
24	RA	172	C	C6-N1-C2	-5.51	118.10	120.30
24	RA	2213	G	N3-C4-N9	5.51	129.30	126.00
1	QA	932	C	C5-C6-N1	5.50	123.75	121.00
24	RA	958	C	N1-C2-O2	5.50	122.20	118.90
24	RA	2695	C	N3-C2-O2	-5.50	118.05	121.90
25	YB	93	C	N3-C4-C5	5.50	124.10	121.90
1	QA	337	C	C6-N1-C2	-5.50	118.10	120.30
1	XA	186(G)	C	N1-C2-O2	5.50	122.20	118.90
24	RA	2539	C	C6-N1-C2	-5.50	118.10	120.30
24	YA	1581	U	C5-C6-N1	5.50	125.45	122.70
24	RA	2513	C	C6-N1-C1'	5.50	127.39	120.80
1	XA	18	C	C6-N1-C2	-5.50	118.10	120.30
22	XV	59	A	C2-N3-C4	5.49	113.35	110.60
24	YA	226	C	C6-N1-C2	-5.49	118.10	120.30
24	RA	139	A	C5-N7-C8	-5.49	101.15	103.90
24	RA	712	C	N3-C2-O2	-5.49	118.06	121.90
24	YA	1450	C	C5-C6-N1	5.49	123.75	121.00
24	YA	2660	C	C5-C6-N1	5.49	123.75	121.00
24	RA	2480	G	O4'-C1'-N9	5.49	112.59	108.20
1	XA	169	C	N1-C2-O2	5.49	122.19	118.90

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	723	A	O4'-C1'-N9	5.49	112.59	108.20
24	YA	830	A	N7-C8-N9	5.49	116.54	113.80
24	YA	1304	C	C6-N1-C2	-5.49	118.11	120.30
25	YB	81	G	C6-C5-N7	-5.49	127.11	130.40
24	RA	1232	G	C6-C5-N7	-5.49	127.11	130.40
24	YA	552	C	N1-C2-O2	5.49	122.19	118.90
24	YA	2628	C	C6-N1-C2	-5.49	118.11	120.30
1	QA	689	C	C6-N1-C2	-5.49	118.11	120.30
1	QA	1263	C	C2-N1-C1'	5.49	124.83	118.80
24	RA	2118	U	N3-C2-O2	-5.49	118.36	122.20
24	YA	232	U	C6-N1-C2	-5.48	117.71	121.00
1	XA	1056	U	N3-C2-O2	-5.48	118.36	122.20
24	YA	665	C	C6-N1-C2	-5.48	118.11	120.30
24	YA	1682	G	C6-C5-N7	-5.48	127.11	130.40
24	RA	2752	U	N3-C2-O2	-5.48	118.36	122.20
24	RA	276	C	C6-N1-C2	-5.48	118.11	120.30
1	XA	455	C	C5-C6-N1	5.48	123.74	121.00
24	YA	2502	G	C5-N7-C8	-5.48	101.56	104.30
24	YA	2347	A	C4-N9-C1'	-5.47	116.45	126.30
1	QA	1009	G	N3-C4-N9	5.47	129.28	126.00
22	XV	65	U	C2-N1-C1'	5.47	124.27	117.70
24	YA	2724	U	C6-N1-C2	-5.47	117.72	121.00
1	QA	1020	U	C2-N1-C1'	5.47	124.27	117.70
24	YA	66	U	C2-N3-C4	5.47	130.28	127.00
24	RA	672	G	N3-C4-N9	5.47	129.28	126.00
24	RA	1812	C	N3-C2-O2	-5.47	118.07	121.90
1	QA	91	C	C2-N1-C1'	5.46	124.81	118.80
24	RA	1548	C	C5-C6-N1	5.46	123.73	121.00
24	YA	976	G	C8-N9-C4	-5.46	104.21	106.40
24	YA	2030	C	C6-N1-C2	-5.46	118.11	120.30
24	RA	1551	C	C2-N1-C1'	5.46	124.81	118.80
24	YA	267	C	N1-C2-O2	5.46	122.18	118.90
1	QA	1395	C	C5-C6-N1	5.46	123.73	121.00
24	RA	1916	C	N3-C2-O2	-5.46	118.08	121.90
1	XA	172	A	C8-N9-C4	-5.46	103.62	105.80
24	YA	723	A	C4-N9-C1'	5.46	136.13	126.30
23	XX	7	G	N7-C8-N9	5.46	115.83	113.10
24	YA	1098	C	C5-C6-N1	5.46	123.73	121.00
24	RA	907	U	C4-C5-C6	5.46	122.97	119.70
24	RA	1067	A	C2-N3-C4	-5.46	107.87	110.60
24	YA	31	C	C6-N1-C2	-5.46	118.12	120.30
24	YA	64	C	C6-N1-C2	-5.46	118.12	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	1374	G	C4-C5-N7	5.45	112.98	110.80
24	YA	1019	G	O4'-C1'-N9	5.45	112.56	108.20
24	YA	1303	C	C6-N1-C2	-5.45	118.12	120.30
1	XA	433	C	N1-C2-O2	5.45	122.17	118.90
24	YA	723	A	C8-N9-C4	-5.45	103.62	105.80
24	YA	1569	U	C5-C6-N1	5.45	125.43	122.70
1	QA	497	U	N1-C2-O2	5.45	126.61	122.80
24	RA	990	A	C5-N7-C8	-5.45	101.18	103.90
1	QA	328	C	P-O3'-C3'	5.45	126.23	119.70
24	RA	735	U	C5-C6-N1	5.45	125.42	122.70
24	RA	2404	A	N7-C8-N9	5.45	116.52	113.80
1	XA	328	C	P-O3'-C3'	5.45	126.23	119.70
24	YA	1665	G	C4-C5-N7	5.44	112.98	110.80
24	YA	2480	G	C4-C5-N7	5.44	112.98	110.80
1	XA	705	U	C6-N1-C2	-5.44	117.74	121.00
1	XA	1149	C	N1-C2-O2	5.44	122.17	118.90
1	XA	337	C	C5-C6-N1	5.44	123.72	121.00
24	YA	1725	G	C2-N3-C4	-5.44	109.18	111.90
24	RA	2703	C	C6-N1-C2	-5.44	118.12	120.30
24	YA	1642	A	C4-C5-C6	-5.44	114.28	117.00
1	XA	150	C	C6-N1-C2	-5.44	118.13	120.30
24	YA	1452	U	C2-N1-C1'	5.44	124.22	117.70
24	YA	2824	C	C6-N1-C2	-5.44	118.13	120.30
24	RA	309	C	C5-C6-N1	5.43	123.72	121.00
1	QA	883	C	C6-N1-C2	-5.43	118.13	120.30
24	YA	2472	U	C5-C6-N1	5.43	125.42	122.70
22	QV	30	C	N1-C2-O2	5.43	122.16	118.90
24	RA	1453	C	N3-C2-O2	-5.43	118.10	121.90
24	RA	1861	C	N1-C2-O2	5.43	122.16	118.90
24	YA	513	C	C6-N1-C2	-5.43	118.13	120.30
24	YA	1805	C	C5-C6-N1	5.43	123.71	121.00
24	RA	1134	A	C2-N3-C4	5.43	113.31	110.60
24	YA	583	C	C5-C6-N1	5.43	123.71	121.00
24	YA	805	C	C6-N1-C2	-5.43	118.13	120.30
24	YA	1927	C	N1-C2-O2	5.43	122.16	118.90
24	RA	712	C	C6-N1-C2	-5.42	118.13	120.30
24	RA	975	U	N1-C2-O2	5.42	126.60	122.80
24	RA	1559	C	C5-C6-N1	5.42	123.71	121.00
36	RR	75	LEU	CA-CB-CG	5.42	127.78	115.30
24	YA	471	C	C6-N1-C2	-5.42	118.13	120.30
24	YA	2255	U	N1-C2-N3	5.42	118.15	114.90
1	QA	824	C	C6-N1-C2	-5.42	118.13	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	1550	C	C2-N1-C1'	5.42	124.76	118.80
24	YA	914	C	C5-C6-N1	5.42	123.71	121.00
1	XA	455	C	C6-N1-C2	-5.42	118.13	120.30
24	YA	1977	U	C2-N1-C1'	5.42	124.20	117.70
24	YA	2857	U	N3-C2-O2	-5.42	118.41	122.20
1	QA	690	G	C4-N9-C1'	5.42	133.54	126.50
24	RA	1378	G	C5-N7-C8	-5.42	101.59	104.30
24	RA	2083	G	N3-C4-N9	5.42	129.25	126.00
24	YA	2004	C	N1-C2-O2	5.42	122.15	118.90
24	RA	989	G	C4-N9-C1'	5.42	133.54	126.50
24	YA	45	C	C6-N1-C2	-5.42	118.13	120.30
24	YA	1252	C	C5-C6-N1	5.42	123.71	121.00
1	XA	980	C	N3-C2-O2	-5.41	118.11	121.90
24	YA	2241	C	C6-N1-C2	-5.41	118.13	120.30
24	YA	607	C	N3-C4-N4	5.41	121.79	118.00
24	YA	541	C	C5-C6-N1	5.41	123.70	121.00
24	RA	2426	G	C4-C5-N7	5.41	112.96	110.80
1	XA	358	U	C2-N1-C1'	-5.41	111.21	117.70
24	YA	2838	C	C6-N1-C2	-5.41	118.14	120.30
24	RA	1450	C	C5-C6-N1	5.41	123.70	121.00
1	QA	1449	C	C5-C6-N1	5.41	123.70	121.00
24	YA	1233	U	C5-C6-N1	5.41	125.40	122.70
24	YA	2054	G	N3-C4-C5	5.41	131.30	128.60
24	YA	2495	C	C5-C6-N1	5.41	123.70	121.00
1	QA	827	U	C6-N1-C2	-5.40	117.76	121.00
24	RA	1450	C	C2-N1-C1'	5.40	124.74	118.80
24	RA	1936	C	C2-N1-C1'	5.40	124.74	118.80
1	XA	747	C	N1-C2-O2	5.40	122.14	118.90
24	YA	1044	C	C5-C6-N1	5.40	123.70	121.00
24	RA	784	C	C5-C6-N1	5.40	123.70	121.00
24	RA	1463	C	N1-C2-O2	5.40	122.14	118.90
24	RA	1542	A	N7-C8-N9	5.40	116.50	113.80
24	YA	1364	C	C4-C5-C6	-5.40	114.70	117.40
24	RA	2330	G	N7-C8-N9	5.40	115.80	113.10
1	QA	1065	U	P-O3'-C3'	5.40	126.18	119.70
24	RA	182	U	N3-C2-O2	-5.40	118.42	122.20
24	RA	2332	A	C2-N3-C4	5.40	113.30	110.60
24	RA	2690	C	C5-C6-N1	5.40	123.70	121.00
24	RA	2878	A	C5-N7-C8	-5.40	101.20	103.90
24	YA	1693	C	N1-C2-O2	5.40	122.14	118.90
24	YA	2061	C	C6-N1-C2	-5.40	118.14	120.30
1	QA	789	U	C2-N1-C1'	5.40	124.18	117.70

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	QV	70	C	C6-N1-C2	-5.39	118.14	120.30
1	XA	217	C	N1-C2-O2	5.39	122.14	118.90
1	XA	381	C	N3-C2-O2	-5.39	118.12	121.90
1	XA	717	C	N1-C2-O2	5.39	122.14	118.90
24	YA	470	C	C5-C6-N1	5.39	123.70	121.00
24	YA	739	C	C5-C6-N1	5.39	123.70	121.00
25	YB	28	C	N1-C2-O2	5.39	122.14	118.90
24	RA	2483	C	C5-C6-N1	5.39	123.70	121.00
24	RA	1022	C	C6-N1-C2	-5.39	118.14	120.30
24	RA	2330	G	C8-N9-C4	-5.39	104.25	106.40
1	XA	1028(C)	C	N1-C2-O2	5.39	122.13	118.90
24	YA	400	U	N3-C2-O2	-5.39	118.43	122.20
24	YA	1352	C	C6-N1-C2	-5.39	118.14	120.30
1	QA	308	C	N1-C2-O2	5.39	122.13	118.90
24	YA	542	C	C6-N1-C2	-5.39	118.15	120.30
24	YA	2693	C	P-O3'-C3'	5.39	126.17	119.70
24	RA	1824	C	C5-C6-N1	5.38	123.69	121.00
1	XA	1237	C	N1-C2-O2	5.38	122.13	118.90
24	YA	1069	U	N3-C2-O2	-5.38	118.43	122.20
24	YA	1398	U	C5-C6-N1	5.38	125.39	122.70
24	YA	2333	G	N3-C4-C5	-5.38	125.91	128.60
24	YA	735	U	C5-C4-O4	-5.38	122.67	125.90
24	YA	73	A	N1-C2-N3	5.38	131.99	129.30
24	YA	1245	C	C6-N1-C2	-5.38	118.15	120.30
24	YA	2419	G	C6-C5-N7	-5.38	127.17	130.40
1	XA	910	C	C5-C6-N1	5.38	123.69	121.00
24	YA	1416	C	C6-N1-C2	-5.38	118.15	120.30
24	YA	2472	U	N3-C2-O2	-5.38	118.44	122.20
24	RA	1707	C	N3-C2-O2	-5.38	118.14	121.90
1	XA	792	A	C4-N9-C1'	5.38	135.98	126.30
24	YA	211	A	N1-C6-N6	-5.38	115.37	118.60
24	YA	2739	U	C2-N1-C1'	5.38	124.15	117.70
1	QA	221	C	N1-C2-O2	5.38	122.12	118.90
1	QA	307	C	C6-N1-C2	-5.37	118.15	120.30
1	QA	528	C	C2-N1-C1'	5.37	124.71	118.80
24	RA	2480	G	C8-N9-C1'	-5.37	120.02	127.00
1	XA	1109	C	N1-C2-O2	5.37	122.12	118.90
24	YA	969	C	C5-C6-N1	5.37	123.69	121.00
1	QA	1019	C	C6-N1-C2	-5.37	118.15	120.30
24	RA	309	C	N1-C2-O2	5.37	122.12	118.90
24	RA	562	C	C2-N1-C1'	5.37	124.71	118.80
1	XA	1059	C	C6-N1-C2	-5.37	118.15	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	2227	G	N3-C4-N9	5.37	129.22	126.00
24	YA	2415	C	C2-N1-C1'	5.37	124.71	118.80
24	RA	1479	U	N1-C2-O2	5.37	126.56	122.80
24	RA	2266	C	N3-C2-O2	-5.37	118.14	121.90
1	QA	989	C	N3-C2-O2	-5.37	118.14	121.90
1	QA	623	C	C5-C6-N1	5.37	123.68	121.00
24	YA	462	C	C5-C6-N1	5.37	123.68	121.00
24	RA	2266	C	N1-C2-O2	5.36	122.12	118.90
24	RA	2603	C	C5-C6-N1	5.36	123.68	121.00
24	YA	994	C	C6-N1-C2	-5.36	118.16	120.30
1	QA	405	U	N3-C2-O2	-5.36	118.45	122.20
24	RA	1726	U	N1-C2-O2	5.36	126.55	122.80
24	YA	1313	U	C5-C6-N1	5.35	125.38	122.70
24	YA	1731	C	C6-N1-C2	-5.35	118.16	120.30
1	QA	218	C	C6-N1-C2	-5.35	118.16	120.30
21	QU	15	ARG	NE-CZ-NH1	5.35	122.98	120.30
24	RA	2849	G	N3-C4-C5	-5.35	125.92	128.60
24	YA	45	C	C5-C6-N1	5.35	123.68	121.00
24	YA	66	U	N1-C2-O2	5.35	126.55	122.80
24	YA	1577	C	C5-C6-N1	5.35	123.67	121.00
24	RA	1775	C	C5-C6-N1	5.35	123.67	121.00
1	XA	1237	C	C6-N1-C2	-5.35	118.16	120.30
24	YA	1180	C	N1-C2-O2	5.35	122.11	118.90
1	XA	943	U	N3-C2-O2	-5.35	118.46	122.20
24	YA	2224	C	C6-N1-C2	-5.35	118.16	120.30
24	RA	660	C	C5-C6-N1	5.34	123.67	121.00
24	RA	2513	C	C2-N1-C1'	-5.34	112.92	118.80
24	YA	2121	U	N1-C2-O2	5.34	126.54	122.80
24	YA	2407	C	C6-N1-C2	-5.34	118.16	120.30
24	YA	2588	G	C8-N9-C1'	-5.34	120.05	127.00
24	YA	2858	G	O4'-C1'-N9	5.34	112.47	108.20
24	RA	482	C	N3-C2-O2	-5.34	118.16	121.90
24	YA	903	C	C2-N3-C4	5.34	122.57	119.90
24	RA	784	C	C6-N1-C2	-5.34	118.16	120.30
1	XA	509	A	C8-N9-C4	-5.34	103.66	105.80
24	YA	280	C	N3-C2-O2	-5.34	118.16	121.90
24	YA	1550	C	N1-C2-O2	5.34	122.11	118.90
24	YA	2712	C	N1-C2-O2	5.34	122.11	118.90
24	RA	2085	C	C5-C6-N1	5.34	123.67	121.00
14	QN	44	LEU	CA-CB-CG	5.34	127.57	115.30
24	RA	1951	G	C4-N9-C1'	5.34	133.44	126.50
24	YA	1651	C	C6-N1-C2	-5.34	118.17	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	1797	U	N3-C2-O2	-5.34	118.47	122.20
24	YA	2591	C	C5-C6-N1	5.34	123.67	121.00
24	RA	1422	C	C5-C6-N1	5.33	123.67	121.00
24	YA	712	C	N1-C2-O2	5.33	122.10	118.90
24	YA	894	U	C4-C5-C6	5.33	122.90	119.70
24	YA	2485	U	N3-C2-O2	-5.33	118.47	122.20
24	RA	950	C	C2-N1-C1'	5.33	124.67	118.80
24	RA	2091	G	C8-N9-C4	-5.33	104.27	106.40
1	XA	390	C	C5-C6-N1	5.33	123.67	121.00
24	YA	1352	C	C5-C6-N1	5.33	123.67	121.00
1	XA	1347	G	C8-N9-C1'	5.33	133.93	127.00
24	YA	796	C	C6-N1-C2	-5.33	118.17	120.30
1	XA	123	C	C6-N1-C2	-5.33	118.17	120.30
1	QA	1263	C	N3-C2-O2	-5.33	118.17	121.90
24	YA	447	C	C6-N1-C2	-5.33	118.17	120.30
24	YA	1579	C	C2-N1-C1'	5.33	124.66	118.80
24	RA	1037	C	C5-C6-N1	5.33	123.66	121.00
24	RA	1521	C	N3-C2-O2	-5.33	118.17	121.90
24	RA	2183	C	N3-C2-O2	-5.33	118.17	121.90
1	XA	540	G	C8-N9-C4	-5.33	104.27	106.40
24	YA	1936	C	C6-N1-C2	-5.33	118.17	120.30
24	YA	1972	G	O4'-C1'-N9	5.33	112.46	108.20
1	XA	449	C	C6-N1-C2	-5.32	118.17	120.30
24	YA	908	A	C8-N9-C4	-5.32	103.67	105.80
24	YA	2807	C	C5-C6-N1	5.32	123.66	121.00
24	YA	924	U	C5-C6-N1	5.32	125.36	122.70
1	QA	690	G	C8-N9-C4	-5.32	104.27	106.40
24	RA	2319	G	C8-N9-C1'	-5.32	120.08	127.00
24	RA	2800	C	N1-C2-O2	5.32	122.09	118.90
24	YA	588	C	N1-C2-O2	5.32	122.09	118.90
24	YA	872	C	C6-N1-C2	-5.32	118.17	120.30
24	YA	2492	C	N1-C2-O2	5.32	122.09	118.90
24	YA	2752	U	C6-N1-C2	-5.32	117.81	121.00
24	RA	2066	C	C6-N1-C2	-5.32	118.17	120.30
24	RA	2413	U	N3-C2-O2	-5.32	118.48	122.20
1	XA	960	U	N3-C2-O2	-5.32	118.48	122.20
24	RA	1365	G	N3-C4-N9	5.32	129.19	126.00
24	RA	2502	G	C4-N9-C1'	5.32	133.41	126.50
24	YA	1207	C	C6-N1-C2	-5.32	118.17	120.30
1	XA	404	U	N3-C2-O2	-5.31	118.48	122.20
24	YA	2694	U	C5-C6-N1	5.31	125.36	122.70
25	RB	91	C	C6-N1-C2	-5.31	118.17	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	705	C	C6-N1-C2	-5.31	118.17	120.30
1	QA	182	U	C5-C6-N1	5.31	125.36	122.70
24	YA	939	C	C2-N1-C1'	5.31	124.64	118.80
24	RA	462	C	N1-C2-O2	5.31	122.08	118.90
24	RA	1904	C	N1-C2-O2	5.31	122.08	118.90
24	YA	226	C	C5-C6-N1	5.31	123.66	121.00
24	YA	1559	C	C5-C6-N1	5.31	123.66	121.00
24	YA	1875	C	C6-N1-C2	-5.31	118.18	120.30
24	RA	2016	C	C6-N1-C2	-5.31	118.18	120.30
24	RA	2899	C	N1-C2-O2	5.31	122.08	118.90
25	RB	31	C	C2-N3-C4	5.31	122.55	119.90
24	RA	2456	G	N1-C6-O6	-5.30	116.72	119.90
24	RA	1214	G	C6-C5-N7	-5.30	127.22	130.40
24	YA	2111	U	C6-N1-C2	-5.30	117.82	121.00
24	RA	893	C	P-O3'-C3'	5.30	126.06	119.70
1	XA	699	C	C6-N1-C2	-5.30	118.18	120.30
24	YA	112	U	N3-C4-O4	5.30	123.11	119.40
24	YA	720	C	N1-C2-O2	5.30	122.08	118.90
1	QA	314	C	C6-N1-C2	-5.30	118.18	120.30
24	YA	2587	C	C5-C6-N1	5.30	123.65	121.00
24	RA	2846	U	N3-C2-O2	-5.30	118.49	122.20
24	YA	2502	G	C8-N9-C1'	-5.30	120.11	127.00
24	RA	894	U	C2-N1-C1'	-5.30	111.34	117.70
1	XA	1397	C	N3-C2-O2	-5.30	118.19	121.90
24	YA	726	C	C5-C6-N1	5.30	123.65	121.00
24	RA	65	C	C5-C6-N1	5.29	123.65	121.00
24	RA	242	C	C5-C6-N1	5.29	123.65	121.00
24	YA	1939	U	N3-C2-O2	-5.29	118.50	122.20
24	YA	2835	C	C6-N1-C2	-5.29	118.18	120.30
24	YA	2863	C	C5-C6-N1	5.29	123.65	121.00
24	RA	587	C	C6-N1-C2	-5.29	118.18	120.30
24	RA	1425	A	C5-N7-C8	-5.29	101.25	103.90
1	XA	186(B)	C	N3-C2-O2	-5.29	118.20	121.90
24	YA	132	C	N3-C2-O2	-5.29	118.20	121.90
24	RA	2428	C	C5-C6-N1	5.29	123.64	121.00
1	XA	435	C	C6-N1-C2	-5.29	118.19	120.30
23	XX	7	G	C8-N9-C4	-5.29	104.28	106.40
24	RA	795	G	O4'-C1'-N9	5.29	112.43	108.20
24	RA	944	C	C6-N1-C2	-5.29	118.19	120.30
24	RA	2230	U	C2-N1-C1'	5.29	124.04	117.70
24	YA	1426	G	N3-C4-N9	5.29	129.17	126.00
24	RA	2298	A	C5-N7-C8	-5.28	101.26	103.90

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	394	C	N3-C2-O2	-5.28	118.20	121.90
24	YA	2067	C	C5-C6-N1	5.28	123.64	121.00
1	QA	1502	A	N7-C8-N9	5.28	116.44	113.80
24	YA	819	C	C6-N1-C2	-5.28	118.19	120.30
24	RA	2164	C	C6-N1-C2	-5.28	118.19	120.30
1	XA	1502	A	C4-C5-N7	5.28	113.34	110.70
24	RA	1650	C	C6-N1-C2	-5.28	118.19	120.30
24	RA	823	G	C8-N9-C4	-5.28	104.29	106.40
24	RA	1015	C	C5-C6-N1	5.28	123.64	121.00
24	YA	1068	G	C8-N9-C4	-5.28	104.29	106.40
25	YB	37	C	N1-C2-O2	5.28	122.06	118.90
24	RA	634	C	C5-C6-N1	5.27	123.64	121.00
24	YA	2759	U	C5-C6-N1	5.27	125.34	122.70
24	RA	172	C	N1-C2-O2	5.27	122.06	118.90
24	RA	1067	A	C8-N9-C4	-5.27	103.69	105.80
1	XA	186(B)	C	N1-C2-O2	5.27	122.06	118.90
24	YA	182	U	N1-C2-O2	5.27	126.49	122.80
24	YA	2063	U	C6-N1-C2	-5.27	117.84	121.00
1	QA	1362(A)	C	C5-C6-N1	5.27	123.63	121.00
24	YA	550	U	C6-N1-C2	-5.27	117.84	121.00
24	RA	2846	U	C6-N1-C2	-5.27	117.84	121.00
1	XA	252	U	N3-C2-O2	-5.27	118.51	122.20
24	YA	2319	G	O4'-C1'-N9	5.27	112.41	108.20
24	RA	796	C	N3-C2-O2	-5.27	118.21	121.90
24	YA	1479	U	N3-C2-O2	-5.27	118.51	122.20
24	RA	805	C	N3-C2-O2	-5.26	118.22	121.90
1	XA	1495	U	C5-C6-N1	5.26	125.33	122.70
24	YA	2616	U	N3-C2-O2	-5.26	118.52	122.20
31	RI	114	LEU	CA-CB-CG	5.26	127.41	115.30
24	RA	768	C	C5-C6-N1	5.26	123.63	121.00
24	RA	2407	C	C5-C6-N1	5.26	123.63	121.00
25	YB	92	G	C6-N1-C2	-5.26	121.94	125.10
25	YB	96	G	C6-N1-C2	-5.26	121.94	125.10
24	YA	881	C	C6-N1-C2	-5.26	118.20	120.30
24	YA	1231	G	N7-C8-N9	5.26	115.73	113.10
24	RA	1336	C	N1-C2-O2	5.26	122.06	118.90
24	RA	1551	C	N3-C2-O2	-5.26	118.22	121.90
1	XA	209	U	C5-C6-N1	5.26	125.33	122.70
24	YA	733	G	C5-C6-N1	5.26	114.13	111.50
1	XA	1336	C	C6-N1-C2	-5.26	118.20	120.30
24	YA	749	G	N3-C4-N9	5.26	129.15	126.00
24	YA	1904	C	N1-C2-O2	5.26	122.05	118.90

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	2677	A	N7-C8-N9	5.26	116.43	113.80
1	QA	1277	C	N1-C2-O2	5.25	122.05	118.90
1	QA	1322	C	C5-C6-N1	5.25	123.63	121.00
24	YA	665	C	N1-C2-O2	5.25	122.05	118.90
24	YA	909	G	N3-C4-C5	-5.25	125.97	128.60
24	YA	1038	C	C5-C6-N1	5.25	123.63	121.00
24	RA	884	C	C5-C6-N1	5.25	123.62	121.00
24	RA	1903	C	C2-N1-C1'	5.25	124.57	118.80
24	RA	2160	C	C6-N1-C2	-5.25	118.20	120.30
1	XA	989	C	C6-N1-C2	-5.25	118.20	120.30
1	XA	1135	U	C5-C6-N1	5.25	125.32	122.70
1	QA	34	C	N1-C2-O2	5.25	122.05	118.90
24	YA	1798	C	C6-N1-C2	-5.25	118.20	120.30
24	YA	2884	C	N1-C2-O2	5.25	122.05	118.90
24	RA	422	U	N3-C2-O2	-5.25	118.53	122.20
24	RA	1270	C	C6-N1-C2	-5.25	118.20	120.30
1	XA	936	C	N1-C2-O2	5.25	122.05	118.90
24	YA	141	C	C6-N1-C2	-5.25	118.20	120.30
24	RA	1978	U	C5-C6-N1	5.24	125.32	122.70
24	YA	1475	G	C6-C5-N7	-5.24	127.25	130.40
24	YA	2066	C	N1-C2-O2	5.24	122.05	118.90
24	YA	2842	U	P-O3'-C3'	5.24	125.99	119.70
24	RA	1798	C	C6-N1-C2	-5.24	118.20	120.30
24	RA	2486	C	N3-C2-O2	-5.24	118.23	121.90
1	XA	346	G	C4-N9-C1'	5.24	133.31	126.50
1	XA	764	C	C6-N1-C2	-5.24	118.20	120.30
1	QA	1502	A	C6-C5-N7	-5.24	128.63	132.30
1	XA	556	C	C5-C6-N1	5.24	123.62	121.00
1	XA	477	G	N3-C4-N9	5.24	129.14	126.00
1	QA	554	C	C6-N1-C2	-5.24	118.21	120.30
24	RA	1303	C	C6-N1-C2	-5.23	118.21	120.30
1	XA	449	C	N1-C2-O2	5.23	122.04	118.90
12	XL	60	LEU	CA-CB-CG	5.23	127.34	115.30
24	YA	1002	A	N9-C4-C5	-5.23	103.71	105.80
24	YA	1868	C	C6-N1-C2	-5.23	118.21	120.30
1	XA	1103	C	C6-N1-C2	-5.23	118.21	120.30
1	XA	1161	C	N1-C2-O2	5.23	122.04	118.90
1	QA	749	C	N3-C2-O2	-5.23	118.24	121.90
24	RA	744	C	C5-C6-N1	5.23	123.61	121.00
24	RA	1110	C	N3-C2-O2	-5.23	118.24	121.90
24	YA	528	A	OP2-P-O3'	5.23	116.71	105.20
24	YA	1255	A	C4-C5-N7	5.23	113.31	110.70

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	2826	C	C5-C6-N1	5.23	123.62	121.00
1	QA	1260	C	N1-C2-O2	5.23	122.04	118.90
24	RA	1358	U	C6-N1-C2	-5.23	117.86	121.00
1	XA	328	C	C5-C6-N1	5.23	123.61	121.00
24	YA	65	C	C6-N1-C2	-5.23	118.21	120.30
24	YA	1335	C	N1-C2-O2	5.23	122.04	118.90
24	RA	2737	C	C6-N1-C2	-5.23	118.21	120.30
1	XA	1199	U	C6-N1-C2	-5.23	117.86	121.00
24	YA	1510	C	C5-C6-N1	5.23	123.61	121.00
1	QA	222	U	N3-C2-O2	-5.22	118.54	122.20
24	RA	2442	A	C8-N9-C4	-5.22	103.71	105.80
24	YA	471	C	N1-C2-O2	5.22	122.03	118.90
24	RA	2432	C	C5-C6-N1	5.22	123.61	121.00
1	XA	217	C	C5-C6-N1	5.22	123.61	121.00
24	YA	102	U	N3-C2-O2	-5.22	118.54	122.20
24	YA	478	G	C8-N9-C4	-5.22	104.31	106.40
24	YA	1233	U	N1-C2-O2	5.22	126.45	122.80
24	YA	2270	C	C6-N1-C2	-5.22	118.21	120.30
24	RA	1581	U	C2-N1-C1'	5.22	123.96	117.70
1	QA	395	C	N1-C2-O2	5.22	122.03	118.90
24	RA	944	C	N3-C2-O2	-5.22	118.25	121.90
24	RA	1252	C	C5-C6-N1	5.22	123.61	121.00
24	RA	1578	C	C6-N1-C2	-5.22	118.21	120.30
24	RA	854	U	N1-C2-O2	5.21	126.45	122.80
24	RA	2772	G	C4-C5-N7	5.21	112.89	110.80
24	YA	267	C	C5-C6-N1	5.21	123.61	121.00
25	YB	104	A	N3-C4-C5	5.21	130.45	126.80
24	RA	659	C	C5-C6-N1	5.21	123.61	121.00
24	YA	800	C	C2-N3-C4	5.21	122.51	119.90
24	RA	1590	C	N1-C2-O2	5.21	122.03	118.90
24	YA	2707	C	C5-C6-N1	5.21	123.61	121.00
24	RA	45	C	C6-N1-C2	-5.21	118.22	120.30
1	QA	536	C	C6-N1-C2	-5.21	118.22	120.30
24	RA	64	C	N1-C2-O2	5.21	122.03	118.90
24	RA	2097	U	N3-C4-O4	5.21	123.05	119.40
24	RA	2586	G	C6-C5-N7	-5.21	127.28	130.40
24	YA	830	A	N1-C6-N6	5.21	121.72	118.60
1	QA	545	C	C6-N1-C2	-5.21	118.22	120.30
24	YA	98	U	OP2-P-O3'	5.21	116.66	105.20
29	YG	82	LEU	CB-CG-CD1	-5.21	102.15	111.00
24	YA	603	C	C2-N3-C4	5.21	122.50	119.90
24	RA	554	A	N7-C8-N9	5.20	116.40	113.80

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	446	C	C5-C6-N1	5.20	123.60	121.00
24	YA	1260	G	C8-N9-C4	-5.20	104.32	106.40
24	RA	795	G	C8-N9-C1'	5.20	133.76	127.00
1	XA	924	C	C6-N1-C2	-5.20	118.22	120.30
24	YA	975	U	C2-N1-C1'	5.20	123.94	117.70
24	YA	1247	C	C6-N1-C2	-5.20	118.22	120.30
1	QA	283	C	N3-C2-O2	-5.20	118.26	121.90
25	RB	80	U	N3-C2-O2	-5.20	118.56	122.20
1	XA	990	C	C6-N1-C2	-5.20	118.22	120.30
24	YA	411	U	N1-C2-O2	5.20	126.44	122.80
24	YA	2227	G	C8-N9-C1'	-5.20	120.24	127.00
1	QA	54	C	C6-N1-C2	-5.20	118.22	120.30
1	QA	1045	C	N1-C2-O2	5.20	122.02	118.90
1	QA	1096	C	N3-C2-O2	-5.20	118.26	121.90
24	YA	2550	C	N1-C2-O2	5.20	122.02	118.90
24	YA	1634	C	C5-C6-N1	5.19	123.60	121.00
1	QA	827	U	N3-C2-O2	-5.19	118.56	122.20
1	QA	1114	C	N1-C2-O2	5.19	122.02	118.90
1	XA	1477	C	C6-N1-C2	-5.19	118.22	120.30
24	YA	654	G	C8-N9-C4	-5.19	104.32	106.40
24	YA	2030	C	C5-C6-N1	5.19	123.60	121.00
24	RA	275	C	N3-C2-O2	-5.19	118.27	121.90
24	YA	343	C	N1-C2-O2	5.19	122.01	118.90
24	RA	1044	C	C5-C6-N1	5.19	123.59	121.00
22	XV	3	G	N3-C4-C5	-5.19	126.01	128.60
24	YA	1711	A	C8-N9-C4	-5.19	103.72	105.80
24	RA	538	A	N7-C8-N9	5.19	116.39	113.80
1	XA	1237	C	N3-C2-O2	-5.19	118.27	121.90
24	YA	187	C	N1-C2-O2	5.19	122.01	118.90
24	YA	1595	C	C5-C6-N1	5.19	123.59	121.00
24	YA	2480	G	N7-C8-N9	5.19	115.69	113.10
24	YA	2511	C	C5-C6-N1	5.19	123.59	121.00
24	RA	1875	C	C6-N1-C2	-5.19	118.22	120.30
25	YB	96	G	C5-N7-C8	-5.18	101.71	104.30
1	QA	1447	G	C6-C5-N7	-5.18	127.29	130.40
24	RA	1644	C	N1-C2-O2	5.18	122.01	118.90
24	RA	2724	U	C2-N1-C1'	5.18	123.92	117.70
9	QI	47	LEU	CA-CB-CG	5.18	127.22	115.30
1	QA	354	G	C4-N9-C1'	5.18	133.23	126.50
24	RA	2028	C	N3-C2-O2	-5.18	118.28	121.90
24	RA	2085	C	N3-C2-O2	-5.18	118.27	121.90
24	YA	604	C	C6-N1-C2	-5.18	118.23	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	848	C	C5-C6-N1	5.18	123.59	121.00
1	XA	690	G	O4'-C1'-N9	5.18	112.34	108.20
24	RA	1214	G	C4-N9-C1'	5.18	133.23	126.50
24	RA	1521	C	C2-N1-C1'	5.18	124.50	118.80
1	XA	936	C	N3-C2-O2	-5.18	118.28	121.90
24	YA	470	C	C6-N1-C2	-5.18	118.23	120.30
24	YA	670	C	C5-C6-N1	5.18	123.59	121.00
24	YA	2250	G	N3-C4-N9	5.18	129.11	126.00
24	RA	706	C	N1-C2-O2	5.17	122.00	118.90
1	XA	1357	A	N7-C8-N9	5.17	116.39	113.80
24	YA	326	C	C6-N1-C2	-5.17	118.23	120.30
24	RA	1133	G	C8-N9-C1'	5.17	133.72	127.00
1	XA	732	C	C6-N1-C2	-5.17	118.23	120.30
1	XA	1163	C	N1-C2-O2	5.17	122.00	118.90
24	YA	711	C	C6-N1-C2	-5.17	118.23	120.30
24	YA	399	G	C8-N9-C1'	5.17	133.72	127.00
24	YA	562	C	N1-C2-O2	5.17	122.00	118.90
24	YA	2571	C	C2-N1-C1'	5.17	124.48	118.80
24	RA	603	C	C5-C6-N1	5.17	123.58	121.00
24	YA	768	C	C5-C6-N1	5.17	123.58	121.00
24	RA	419	C	C5-C6-N1	5.17	123.58	121.00
24	YA	1923	A	C2-N3-C4	5.17	113.18	110.60
24	RA	548	C	C6-N1-C2	-5.16	118.23	120.30
24	YA	907	U	C6-N1-C2	-5.16	117.90	121.00
24	YA	1453	C	C2-N3-C4	5.16	122.48	119.90
24	RA	2739	U	C2-N1-C1'	5.16	123.89	117.70
24	RA	1963	C	N1-C2-O2	5.16	122.00	118.90
24	YA	1858	C	C6-N1-C2	-5.16	118.24	120.30
24	YA	2355	C	N3-C2-O2	-5.16	118.29	121.90
24	RA	554	A	C8-N9-C4	-5.16	103.74	105.80
24	RA	893	C	N3-C2-O2	-5.16	118.29	121.90
24	YA	2837	C	C4-C5-C6	-5.16	114.82	117.40
1	QA	1246	C	C6-N1-C2	-5.16	118.24	120.30
24	RA	1359	U	C6-N1-C2	-5.16	117.91	121.00
24	YA	139	A	C4-C5-N7	5.16	113.28	110.70
25	YB	103	U	N3-C2-O2	-5.15	118.59	122.20
24	YA	725	C	C6-N1-C2	-5.15	118.24	120.30
24	YA	2442	A	N1-C2-N3	5.15	131.88	129.30
24	YA	2502	G	C6-C5-N7	-5.15	127.31	130.40
24	RA	863	C	C5-C6-N1	5.15	123.58	121.00
24	RA	2088	C	N3-C2-O2	-5.15	118.29	121.90
24	RA	950	C	N1-C2-O2	5.15	121.99	118.90

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	566	C	C5-C6-N1	5.15	123.58	121.00
24	YA	1806	U	C6-N1-C2	-5.15	117.91	121.00
24	RA	1138	C	C2-N3-C4	5.15	122.47	119.90
24	YA	1122	C	N1-C2-O2	5.15	121.99	118.90
24	RA	2014	G	P-O3'-C3'	5.15	125.88	119.70
24	RA	2743	C	C5-C6-N1	5.15	123.57	121.00
24	YA	1389	G	C8-N9-C4	-5.15	104.34	106.40
24	YA	1072	U	P-O3'-C3'	5.14	125.87	119.70
24	RA	2111	U	C5-C6-N1	5.14	125.27	122.70
24	YA	2088	C	C6-N1-C2	-5.14	118.24	120.30
1	XA	1406	U	N3-C2-O2	-5.14	118.60	122.20
24	YA	1070	G	N3-C4-N9	5.14	129.09	126.00
1	QA	1126	U	C5-C6-N1	5.14	125.27	122.70
24	RA	826	U	N1-C2-O2	5.14	126.40	122.80
24	YA	1813	C	C5-C6-N1	5.14	123.57	121.00
1	XA	169	C	C5-C6-N1	5.14	123.57	121.00
24	YA	1637	G	C4-C5-N7	5.14	112.86	110.80
50	Y5	6	VAL	CG1-CB-CG2	-5.14	102.68	110.90
24	RA	1128	U	C2-N1-C1'	5.14	123.86	117.70
24	YA	961	C	C6-N1-C2	-5.14	118.25	120.30
24	RA	2837	C	N1-C2-O2	5.13	121.98	118.90
24	YA	2840	G	C8-N9-C4	-5.13	104.35	106.40
1	XA	358	U	C1'-O4'-C4'	-5.13	105.80	109.90
1	XA	812	C	OP2-P-O3'	5.13	116.49	105.20
24	RA	1733	C	N1-C2-O2	5.13	121.98	118.90
24	YA	637	U	C6-N1-C2	-5.13	117.92	121.00
24	YA	950	C	C6-N1-C2	-5.13	118.25	120.30
24	RA	885	C	C6-N1-C2	-5.13	118.25	120.30
1	XA	23	C	C6-N1-C2	-5.13	118.25	120.30
24	YA	1767	A	O4'-C1'-N9	5.13	112.30	108.20
24	YA	2485	U	N1-C2-O2	5.13	126.39	122.80
1	QA	891	U	N1-C2-O2	5.12	126.39	122.80
24	YA	1903	C	C2-N1-C1'	5.12	124.44	118.80
1	QA	1282	C	C6-N1-C2	-5.12	118.25	120.30
24	RA	1155	C	C2-N1-C1'	5.12	124.44	118.80
24	YA	1421	C	N1-C2-O2	5.12	121.97	118.90
24	YA	2688	C	C6-N1-C2	-5.12	118.25	120.30
1	QA	1406	U	N1-C2-O2	5.12	126.39	122.80
24	RA	950	C	C6-N1-C2	-5.12	118.25	120.30
24	YA	2550	C	C5-C6-N1	5.12	123.56	121.00
1	QA	717	C	N1-C2-O2	5.12	121.97	118.90
24	YA	140	A	C6-C5-N7	-5.12	128.72	132.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	YA	989	G	C8-N9-C1'	-5.12	120.35	127.00
24	YA	2111	U	N3-C2-O2	-5.12	118.62	122.20
24	YA	985	G	C8-N9-C4	-5.12	104.35	106.40
25	YB	10	C	C5-C6-N1	5.12	123.56	121.00
24	RA	444	C	N1-C2-O2	5.11	121.97	118.90
24	RA	1249	A	O4'-C1'-N9	5.11	112.29	108.20
1	XA	528	C	C2-N1-C1'	5.11	124.43	118.80
24	YA	18	C	C5-C6-N1	5.11	123.56	121.00
24	YA	808	A	C5-N7-C8	-5.11	101.34	103.90
24	YA	2369	U	C5-C6-N1	5.11	125.26	122.70
24	RA	118	U	C2-N1-C1'	5.11	123.83	117.70
24	RA	2082	A	P-O3'-C3'	5.11	125.83	119.70
24	YA	1806	U	N3-C4-O4	5.11	122.98	119.40
24	YA	118	U	C2-N1-C1'	5.11	123.83	117.70
1	QA	1059	C	C6-N1-C2	-5.11	118.26	120.30
24	RA	893	C	C6-N1-C2	-5.11	118.26	120.30
24	YA	2088	C	N3-C2-O2	-5.11	118.33	121.90
24	YA	2238	C	N3-C2-O2	-5.11	118.33	121.90
24	YA	2884	C	C2-N1-C1'	5.10	124.42	118.80
24	RA	1120	G	N3-C4-N9	5.10	129.06	126.00
24	RA	2712	C	N1-C2-O2	5.10	121.96	118.90
24	RA	2859	U	O4'-C1'-N1	5.10	112.28	108.20
24	YA	1086	C	N1-C2-O2	5.10	121.96	118.90
24	YA	1452	U	C2-N3-C4	5.10	130.06	127.00
24	YA	1531	G	C6-C5-N7	-5.10	127.34	130.40
1	XA	1246	C	C5-C6-N1	5.10	123.55	121.00
1	XA	1267	C	C6-N1-C2	-5.10	118.26	120.30
1	XA	780	A	C8-N9-C4	-5.10	103.76	105.80
24	YA	672	G	N3-C4-C5	-5.10	126.05	128.60
24	YA	950	C	N3-C2-O2	-5.10	118.33	121.90
24	RA	712	C	N1-C2-O2	5.10	121.96	118.90
1	XA	1505	G	N3-C4-N9	-5.10	122.94	126.00
24	YA	2368	C	C6-N1-C2	-5.10	118.26	120.30
24	RA	2021	C	C5-C6-N1	5.09	123.55	121.00
24	YA	98	U	P-O3'-C3'	5.09	125.81	119.70
24	YA	140	A	C6-N1-C2	5.09	121.66	118.60
24	YA	399	G	C4-N9-C1'	-5.09	119.88	126.50
24	YA	1452	U	C6-N1-C2	-5.09	117.94	121.00
24	YA	2407	C	C5-C6-N1	5.09	123.55	121.00
24	RA	544	U	C5-C6-N1	5.09	125.25	122.70
24	YA	287	G	P-O3'-C3'	5.09	125.81	119.70
24	RA	1977	U	C6-N1-C1'	-5.09	114.07	121.20

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	637	U	C2-N1-C1'	5.09	123.81	117.70
24	RA	1070	G	N3-C4-N9	5.09	129.05	126.00
24	RA	2591	C	C5-C6-N1	5.09	123.55	121.00
3	QC	101	LEU	CA-CB-CG	5.09	127.00	115.30
1	XA	563	A	C4-N9-C1'	5.09	135.45	126.30
24	YA	112	U	C6-N1-C2	-5.09	117.95	121.00
24	RA	624	C	C6-N1-C2	-5.08	118.27	120.30
24	RA	1766	G	N3-C4-N9	5.08	129.05	126.00
24	YA	607	C	C6-N1-C2	-5.08	118.27	120.30
24	YA	670	C	C6-N1-C2	-5.08	118.27	120.30
24	YA	1433	C	C6-N1-C2	-5.08	118.27	120.30
24	YA	2800	C	N1-C2-O2	5.08	121.95	118.90
24	RA	723	A	C4-C5-N7	5.08	113.24	110.70
24	YA	887	C	C2-N3-C4	5.08	122.44	119.90
24	YA	2118	U	N1-C2-O2	5.08	126.36	122.80
24	RA	1225	C	C6-N1-C2	-5.08	118.27	120.30
24	RA	2150	C	N3-C2-O2	-5.08	118.34	121.90
24	RA	2648	U	N3-C2-O2	-5.08	118.64	122.20
24	YA	2797	C	C5-C6-N1	5.08	123.54	121.00
24	RA	1186	U	C6-N1-C1'	5.08	128.31	121.20
24	YA	1830	G	C2'-C3'-O3'	5.08	121.82	113.70
24	RA	513	C	C5-C6-N1	5.08	123.54	121.00
24	YA	1556	A	N3-C4-N9	5.08	131.46	127.40
24	RA	1444	C	C6-N1-C2	-5.08	118.27	120.30
24	RA	1596	C	C6-N1-C2	-5.08	118.27	120.30
24	YA	630	U	C6-N1-C2	-5.08	117.95	121.00
24	YA	1810	U	N1-C2-O2	5.08	126.35	122.80
24	YA	1972	G	N7-C8-N9	5.08	115.64	113.10
24	RA	211	A	P-O3'-C3'	5.07	125.79	119.70
24	RA	1133	G	C4-N9-C1'	-5.07	119.91	126.50
24	RA	2483	C	C2-N1-C1'	5.07	124.38	118.80
24	RA	1164	C	C6-N1-C2	-5.07	118.27	120.30
24	RA	1939	U	C5-C6-N1	5.07	125.24	122.70
24	RA	2421	G	N3-C4-N9	5.07	129.04	126.00
24	YA	342	C	C6-N1-C2	-5.07	118.27	120.30
24	YA	1051	C	N1-C2-O2	5.07	121.94	118.90
1	XA	153	C	N1-C2-O2	5.07	121.94	118.90
24	YA	138	G	C4-C5-N7	-5.07	108.77	110.80
24	YA	940	C	N1-C2-O2	5.07	121.94	118.90
24	RA	2004	C	N1-C2-O2	5.07	121.94	118.90
24	YA	542	C	N1-C2-O2	5.07	121.94	118.90
24	RA	309	C	C6-N1-C2	-5.07	118.27	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	755	C	C2-N1-C1'	5.07	124.38	118.80
1	XA	961	U	N3-C2-O2	-5.07	118.65	122.20
1	XA	1381	U	C6-N1-C2	-5.07	117.96	121.00
24	RA	41	C	N1-C2-O2	5.07	121.94	118.90
24	YA	2492	C	C6-N1-C2	-5.07	118.27	120.30
25	YB	84	C	C5-C6-N1	5.07	123.53	121.00
1	XA	910	C	C6-N1-C2	-5.06	118.27	120.30
24	RA	537	G	O4'-C1'-N9	5.06	112.25	108.20
24	RA	2703	C	C5-C6-N1	5.06	123.53	121.00
24	YA	2752	U	C5-C6-N1	5.06	125.23	122.70
22	QV	54	U	C5-C6-N1	5.06	125.23	122.70
24	RA	237	G	C6-C5-N7	-5.06	127.36	130.40
24	YA	392	U	N3-C2-O2	-5.06	118.66	122.20
24	YA	2090	U	C2-N3-C4	5.06	130.04	127.00
1	XA	1037	C	C5-C6-N1	5.06	123.53	121.00
24	YA	1953	U	N1-C2-O2	5.06	126.34	122.80
1	QA	174	C	C6-N1-C2	-5.06	118.28	120.30
24	RA	2554	A	C4-C5-C6	-5.06	114.47	117.00
24	YA	1455	C	C5-C6-N1	5.06	123.53	121.00
24	YA	2768	C	C6-N1-C2	-5.06	118.28	120.30
24	RA	410	U	C2-N1-C1'	-5.06	111.63	117.70
24	YA	1515	C	C6-N1-C2	-5.06	118.28	120.30
24	RA	1379	C	C6-N1-C2	-5.05	118.28	120.30
24	RA	2159	C	C2-N1-C1'	5.05	124.36	118.80
1	XA	291	C	C6-N1-C2	-5.05	118.28	120.30
24	YA	664	U	C5-C6-N1	5.05	125.23	122.70
24	YA	1212	C	N1-C2-O2	5.05	121.93	118.90
22	XV	68	C	C6-N1-C2	-5.05	118.28	120.30
24	YA	2456	G	C5-C6-O6	5.05	131.63	128.60
24	RA	2473	C	N1-C2-O2	5.05	121.93	118.90
24	YA	2863	C	C6-N1-C2	-5.05	118.28	120.30
1	QA	56	U	N3-C4-O4	5.05	122.94	119.40
24	RA	1942	C	N1-C2-O2	5.05	121.93	118.90
24	YA	68	C	C6-N1-C2	-5.05	118.28	120.30
1	QA	1378	C	N1-C2-O2	5.05	121.93	118.90
24	YA	897	C	C5-C6-N1	5.05	123.52	121.00
24	YA	1866	G	N3-C4-N9	5.05	129.03	126.00
24	YA	864	C	N1-C2-O2	5.04	121.93	118.90
24	YA	2061	C	N1-C2-O2	5.04	121.93	118.90
24	YA	2785	C	C6-N1-C2	-5.04	118.28	120.30
1	QA	717	C	C6-N1-C2	-5.04	118.28	120.30
24	RA	315	C	C6-N1-C2	-5.04	118.28	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	108	G	C4-N9-C1'	5.04	133.05	126.50
24	RA	729	G	C5-C6-O6	5.04	131.62	128.60
24	RA	2333	G	C8-N9-C4	-5.04	104.38	106.40
25	RB	47	C	N3-C2-O2	-5.04	118.37	121.90
1	XA	1024	G	N3-C4-N9	5.04	129.03	126.00
24	RA	297	C	N1-C2-O2	5.04	121.92	118.90
1	QA	1301	U	C6-N1-C2	-5.04	117.98	121.00
24	RA	2700	U	C5-C4-O4	5.04	128.92	125.90
24	YA	961	C	N3-C2-O2	-5.04	118.37	121.90
24	RA	1895	U	N3-C2-O2	-5.04	118.67	122.20
25	RB	27	C	C5-C6-N1	5.04	123.52	121.00
24	YA	2311	G	C6-C5-N7	-5.04	127.38	130.40
24	RA	555	G	C8-N9-C1'	5.04	133.54	127.00
24	RA	2714	U	N1-C2-N3	5.04	117.92	114.90
24	RA	2014	G	OP2-P-O3'	5.03	116.27	105.20
24	RA	2516	U	N1-C2-O2	5.03	126.32	122.80
24	YA	1921	G	C6-C5-N7	-5.03	127.38	130.40
24	YA	2250	G	C2-N3-C4	5.03	114.42	111.90
24	RA	2125	C	C4-C5-C6	5.03	119.92	117.40
24	YA	1059	C	C5-C6-N1	5.03	123.52	121.00
1	QA	545	C	N1-C2-O2	5.03	121.92	118.90
24	RA	394	C	C5-C6-N1	5.03	123.52	121.00
24	YA	1378	G	N1-C2-N2	-5.03	111.67	116.20
24	YA	2227	G	N3-C4-C5	-5.03	126.08	128.60
24	YA	2883	A	O4'-C1'-N9	5.03	112.22	108.20
25	YB	7	G	C6-C5-N7	-5.03	127.38	130.40
1	QA	827	U	N1-C2-N3	5.03	117.92	114.90
24	RA	2714	U	C6-N1-C1'	5.03	128.24	121.20
24	YA	471	C	N3-C2-O2	-5.03	118.38	121.90
24	YA	1255	A	N7-C8-N9	5.03	116.31	113.80
24	YA	1416	C	N1-C2-O2	5.03	121.92	118.90
24	RA	768	C	N1-C2-O2	5.03	121.92	118.90
24	RA	2213	G	N3-C4-C5	-5.03	126.09	128.60
1	XA	514	C	C6-N1-C2	-5.03	118.29	120.30
24	YA	1037	C	C6-N1-C2	-5.03	118.29	120.30
1	XA	1362(B)	C	N1-C2-O2	5.02	121.91	118.90
1	QA	381	C	N3-C2-O2	-5.02	118.38	121.90
24	RA	453	C	N1-C2-O2	5.02	121.91	118.90
1	XA	169	C	C2-N3-C4	5.02	122.41	119.90
1	XA	811	C	N3-C2-O2	-5.02	118.39	121.90
24	YA	1450	C	N1-C2-O2	5.02	121.91	118.90
24	YA	1656	A	N7-C8-N9	5.02	116.31	113.80

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	RA	2432	C	C6-N1-C2	-5.02	118.29	120.30
1	XA	1297	C	P-O3'-C3'	5.02	125.72	119.70
24	YA	1605	A	P-O3'-C3'	5.02	125.72	119.70
24	YA	1658	C	C5-C6-N1	5.02	123.51	121.00
24	YA	98	U	N1-C2-O2	5.02	126.31	122.80
24	YA	2256	U	C5-C4-O4	-5.02	122.89	125.90
24	YA	2743	C	C6-N1-C2	-5.02	118.29	120.30
25	RB	80	U	C6-N1-C2	-5.01	117.99	121.00
24	YA	704	U	N3-C2-O2	-5.01	118.69	122.20
24	YA	795	G	C8-N9-C1'	5.01	133.52	127.00
24	YA	1030	A	C8-N9-C4	-5.01	103.80	105.80
24	RA	2442	A	N7-C8-N9	5.01	116.31	113.80
24	YA	444	C	N1-C2-O2	5.01	121.91	118.90
25	YB	70	C	C6-N1-C2	-5.01	118.30	120.30
24	YA	720	C	N3-C2-O2	-5.01	118.39	121.90
24	RA	1775	C	N1-C2-O2	5.01	121.91	118.90
24	RA	2257	U	C6-N1-C2	-5.01	118.00	121.00
24	RA	2327	G	N3-C4-N9	5.01	129.00	126.00
24	YA	861	C	C5-C6-N1	5.01	123.50	121.00
24	YA	2065	C	N1-C2-O2	5.01	121.91	118.90
24	RA	240	A	C8-N9-C4	-5.01	103.80	105.80
24	RA	739	C	C6-N1-C2	-5.01	118.30	120.30
24	RA	1225	C	N3-C2-O2	-5.01	118.39	121.90
24	YA	805	C	N3-C2-O2	-5.01	118.39	121.90
24	YA	1921	G	N3-C2-N2	5.01	123.41	119.90
1	QA	328	C	C6-N1-C1'	-5.00	114.79	120.80
1	QA	831	U	C6-N1-C2	-5.00	118.00	121.00
24	RA	2807	C	C5-C6-N1	5.00	123.50	121.00
24	YA	646	A	C4-N9-C1'	5.00	135.31	126.30
24	YA	1037	C	C5-C6-N1	5.00	123.50	121.00
24	YA	2011	G	N9-C4-C5	5.00	107.40	105.40
24	YA	2260	C	C5-C6-N1	5.00	123.50	121.00
24	YA	2663	C	C6-N1-C2	-5.00	118.30	120.30
24	RA	1068	G	P-O3'-C3'	5.00	125.70	119.70

There are no chirality outliers.

All (781) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
2	QB	126	GLU	Peptide
2	QB	135	GLN	Peptide
2	QB	136	VAL	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
2	QB	15	VAL	Peptide
2	QB	18	GLY	Peptide
2	QB	19	HIS	Peptide
2	QB	207	ALA	Peptide
2	QB	21	ARG	Peptide
2	QB	228	GLY	Peptide
2	QB	232	PRO	Peptide
2	QB	233	SER	Peptide
2	QB	235	SER	Peptide
2	QB	38	GLY	Peptide
2	QB	64	ARG	Peptide
2	QB	66	GLY	Peptide
2	QB	67	THR	Peptide
2	QB	71	VAL	Peptide
2	QB	74	LYS	Peptide
2	QB	81	VAL	Peptide
2	QB	86	GLU	Peptide
2	QB	95	GLN	Peptide
2	QB	96	ARG	Peptide
3	QC	11	ARG	Peptide
3	QC	126	ARG	Peptide
3	QC	132	ARG	Peptide
3	QC	14	ILE	Peptide
3	QC	164	ARG	Peptide
3	QC	166	GLU	Peptide
3	QC	168	ALA	Peptide
3	QC	189	ALA	Peptide
3	QC	51	GLY	Peptide
3	QC	59	ARG	Peptide
3	QC	62	ASP	Peptide
3	QC	74	GLY	Peptide
3	QC	78	GLY	Peptide
3	QC	9	GLY	Peptide
3	QC	97	LYS	Peptide
4	QD	153	ARG	Peptide
4	QD	154	ASN	Peptide
4	QD	167	GLY	Peptide
4	QD	180	GLY	Peptide
4	QD	205	GLU	Peptide
4	QD	208	SER	Peptide
4	QD	32	ALA	Peptide
4	QD	33	MET	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
4	QD	69	GLY	Peptide
4	QD	83	SER	Peptide
4	QD	84	LYS	Peptide
4	QD	88	VAL	Peptide
5	QE	116	THR	Peptide
5	QE	154	GLY	Peptide
5	QE	22	GLY	Peptide
5	QE	25	ARG	Peptide
5	QE	72	GLN	Peptide
5	QE	73	ASN	Peptide
5	QE	97	GLY	Peptide
6	QF	41	GLU	Peptide
6	QF	42	GLU	Peptide
6	QF	49	ALA	Peptide
6	QF	75	LEU	Peptide
6	QF	99	ALA	Peptide
7	QG	11	GLN	Peptide
7	QG	113	GLU	Peptide
7	QG	132	GLY	Peptide
7	QG	19	GLY	Peptide
7	QG	30	ILE	Peptide
7	QG	53	LYS	Peptide
7	QG	6	ARG	Peptide
7	QG	7	ALA	Peptide
7	QG	8	GLU	Peptide
8	QH	137	VAL	Peptide
8	QH	23	SER	Peptide
8	QH	96	GLY	Peptide
9	QI	114	TYR	Peptide
9	QI	123	PRO	Peptide
9	QI	14	VAL	Peptide
9	QI	16	ARG	Peptide
9	QI	2	GLU	Peptide
9	QI	20	ARG	Peptide
9	QI	24	GLY	Peptide
9	QI	30	GLY	Peptide
9	QI	54	ASP	Peptide
9	QI	6	GLY	Peptide
9	QI	68	GLY	Peptide
9	QI	69	GLY	Peptide
9	QI	92	TYR	Peptide
9	QI	93	ARG	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
9	QI	99	LEU	Peptide
10	QJ	100	THR	Peptide
10	QJ	25	GLU	Peptide
10	QJ	31	GLY	Peptide
10	QJ	32	ALA	Peptide
10	QJ	40	LEU	Peptide
10	QJ	53	PRO	Peptide
10	QJ	68	HIS	Peptide
10	QJ	69	ASN	Peptide
10	QJ	85	LEU	Peptide
10	QJ	88	LEU	Peptide
10	QJ	90	LEU	Peptide
10	QJ	91	PRO	Peptide
11	QK	101	SER	Peptide
11	QK	112	THR	Peptide
11	QK	126	ARG	Peptide
11	QK	38	ASN	Peptide
11	QK	43	SER	Peptide
12	QL	104	VAL	Peptide
12	QL	118	SER	Peptide
12	QL	120	TYR	Peptide
12	QL	126	LYS	Peptide
12	QL	17	LYS	Peptide
12	QL	24	VAL	Peptide
12	QL	29	GLY	Peptide
12	QL	47	LYS	Peptide
13	QM	105	THR	Peptide
13	QM	11	ARG	Peptide
13	QM	118	ALA	Peptide
13	QM	12	ASN	Peptide
13	QM	120	LYS	Peptide
13	QM	3	ARG	Peptide
13	QM	4	ILE	Peptide
13	QM	59	TYR	Peptide
13	QM	62	ASN	Peptide
13	QM	65	LYS	Peptide
13	QM	66	LEU	Peptide
13	QM	9	ILE	Peptide
13	QM	97	PRO	Peptide
14	QN	14	PRO	Peptide
14	QN	15	LYS	Peptide
14	QN	2	ALA	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
14	QN	42	ILE	Peptide
14	QN	55	GLY	Peptide
14	QN	59	ALA	Peptide
15	QO	88	ARG	Peptide
16	QP	15	PRO	Peptide
16	QP	19	ILE	Peptide
16	QP	29	ASP	Peptide
16	QP	30	GLY	Peptide
16	QP	83	GLU	Peptide
17	QQ	100	LYS	Peptide
17	QQ	43	LEU	Peptide
17	QQ	80	GLY	Peptide
18	QR	19	LYS	Peptide
18	QR	31	LEU	Peptide
18	QR	54	ARG	Peptide
18	QR	56	THR	Peptide
18	QR	87	ARG	Peptide
19	QS	10	PHE	Peptide
19	QS	11	VAL	Peptide
19	QS	2	PRO	Peptide
19	QS	26	GLY	Peptide
19	QS	27	GLU	Peptide
19	QS	29	ARG	Peptide
19	QS	30	LEU	Peptide
19	QS	42	PRO	Peptide
19	QS	46	GLY	Peptide
19	QS	63	THR	Peptide
19	QS	71	LEU	Peptide
19	QS	72	GLY	Peptide
19	QS	9	VAL	Peptide
20	QT	101	GLY	Peptide
20	QT	11	SER	Peptide
20	QT	12	ALA	Peptide
20	QT	48	LYS	Peptide
20	QT	72	LEU	Peptide
20	QT	73	HIS	Peptide
20	QT	94	ALA	Peptide
20	QT	96	GLY	Peptide
20	QT	97	ALA	Peptide
20	QT	98	PRO	Peptide
45	R0	43	THR	Peptide
45	R0	8	GLY	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
46	R1	53	VAL	Peptide
46	R1	54	ALA	Peptide
46	R1	55	GLY	Peptide
46	R1	71	TYR	Peptide
46	R1	73	LEU	Peptide
46	R1	74	VAL	Peptide
46	R1	80	LEU	Peptide
46	R1	85	LEU	Peptide
46	R1	87	PRO	Peptide
46	R1	97	LEU	Peptide
47	R2	13	ALA	Peptide
47	R2	43	GLN	Peptide
47	R2	46	GLN	Peptide
47	R2	70	GLN	Peptide
47	R2	71	ASN	Peptide
48	R3	2	PRO	Peptide
49	R4	11	PRO	Peptide
49	R4	19	GLY	Peptide
49	R4	23	GLU	Peptide
49	R4	34	GLU	Peptide
49	R4	37	SER	Peptide
49	R4	42	PHE	Peptide
49	R4	46	GLN	Peptide
49	R4	47	GLN	Peptide
49	R4	48	ARG	Peptide
49	R4	52	THR	Peptide
49	R4	55	ARG	Peptide
49	R4	61	ARG	Peptide
49	R4	62	ARG	Peptide
49	R4	63	TYR	Peptide
49	R4	65	ASP	Peptide
50	R5	5	PRO	Peptide
52	R7	6	GLN	Peptide
53	R8	27	THR	Peptide
53	R8	28	GLY	Peptide
53	R8	29	LYS	Peptide
53	R8	32	LEU	Peptide
53	R8	35	GLN	Peptide
53	R8	44	LYS	Peptide
53	R8	51	ALA	Peptide
53	R8	52	LYS	Peptide
53	R8	60	LEU	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
53	R8	62	LEU	Peptide
53	R8	64	TYR	Peptide
26	RD	110	GLY	Peptide
26	RD	112	GLN	Peptide
26	RD	12	SER	Peptide
26	RD	122	ASP	Peptide
26	RD	126	GLN	Peptide
26	RD	197	GLY	Peptide
26	RD	2	ALA	Peptide
26	RD	241	PRO	Peptide
26	RD	243	GLY	Peptide
26	RD	246	PRO	Peptide
26	RD	56	GLY	Peptide
26	RD	57	GLY	Peptide
26	RD	80	ALA	Peptide
27	RE	115	GLY	Peptide
27	RE	132	HIS	Peptide
27	RE	142	GLY	Peptide
27	RE	153	GLY	Peptide
27	RE	187	ALA	Peptide
27	RE	204	ALA	Peptide
27	RE	74	PRO	Peptide
28	RF	127	GLU	Peptide
28	RF	128	ALA	Peptide
28	RF	130	ALA	Peptide
28	RF	131	GLY	Peptide
28	RF	135	LYS	Peptide
28	RF	196	LEU	Peptide
28	RF	198	ALA	Peptide
28	RF	24	LEU	Peptide
28	RF	25	PRO	Peptide
28	RF	47	GLY	Peptide
28	RF	66	PRO	Peptide
28	RF	85	GLY	Peptide
29	RG	109	VAL	Peptide
29	RG	127	GLY	Peptide
29	RG	128	ARG	Peptide
29	RG	135	LEU	Peptide
29	RG	136	ARG	Peptide
29	RG	147	ASP	Peptide
29	RG	151	ALA	Peptide
29	RG	159	VAL	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
29	RG	2	PRO	Peptide
29	RG	3	LEU	Peptide
29	RG	39	ILE	Peptide
29	RG	45	GLU	Peptide
29	RG	52	ILE	Peptide
29	RG	54	GLU	Peptide
29	RG	79	ASN	Peptide
29	RG	82	LEU	Peptide
29	RG	95	ARG	Peptide
30	RH	10	PRO	Peptide
30	RH	108	GLY	Peptide
30	RH	126	PRO	Peptide
30	RH	128	PRO	Peptide
30	RH	129	THR	Peptide
30	RH	13	LYS	Peptide
30	RH	14	GLY	Peptide
30	RH	151	ILE	Peptide
30	RH	152	ARG	Peptide
30	RH	153	LYS	Peptide
30	RH	156	ALA	Peptide
30	RH	16	SER	Peptide
30	RH	161	GLY	Peptide
30	RH	19	VAL	Peptide
30	RH	39	PRO	Peptide
30	RH	40	GLU	Peptide
30	RH	66	GLY	Peptide
30	RH	7	LEU	Peptide
30	RH	8	PRO	Peptide
30	RH	82	GLY	Peptide
30	RH	84	SER	Peptide
30	RH	87	LEU	Peptide
30	RH	91	GLY	Peptide
31	RI	11	ASN	Peptide
31	RI	116	LEU	Peptide
31	RI	119	PRO	Peptide
31	RI	123	LEU	Peptide
31	RI	131	LYS	Peptide
31	RI	132	PRO	Peptide
31	RI	133	HIS	Peptide
31	RI	134	PRO	Peptide
31	RI	14	ASP	Peptide
31	RI	142	VAL	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
31	RI	144	VAL	Peptide
31	RI	16	GLY	Peptide
31	RI	29	TYR	Peptide
31	RI	86	THR	Peptide
31	RI	89	TYR	Peptide
31	RI	9	LEU	Peptide
31	RI	90	GLY	Peptide
32	RN	113	GLY	Peptide
32	RN	114	ARG	Peptide
32	RN	127	ASP	Peptide
32	RN	132	ALA	Peptide
32	RN	137	LYS	Peptide
32	RN	21	LYS	Peptide
32	RN	22	THR	Peptide
32	RN	34	LEU	Peptide
32	RN	69	GLN	Peptide
32	RN	92	ALA	Peptide
33	RO	48	PRO	Peptide
33	RO	96	THR	Peptide
34	RP	100	LEU	Peptide
34	RP	146	VAL	Peptide
34	RP	28	GLY	Peptide
34	RP	35	HIS	Peptide
34	RP	37	GLY	Peptide
34	RP	50	ARG	Peptide
34	RP	54	GLY	Peptide
34	RP	65	ARG	Peptide
35	RQ	1	MET	Peptide
35	RQ	103	MET	Peptide
35	RQ	104	PHE	Peptide
35	RQ	108	GLY	Peptide
35	RQ	17	LEU	Peptide
35	RQ	20	ALA	Peptide
35	RQ	21	THR	Peptide
35	RQ	23	GLY	Peptide
35	RQ	24	GLY	Peptide
35	RQ	4	PRO	Peptide
35	RQ	60	ARG	Peptide
35	RQ	61	GLY	Peptide
35	RQ	82	ARG	Peptide
35	RQ	87	LYS	Peptide
35	RQ	88	GLY	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
35	RQ	89	ASN	Peptide
35	RQ	90	VAL	Peptide
35	RQ	91	GLU	Peptide
36	RR	117	VAL	Peptide
36	RR	24	GLN	Peptide
36	RR	75	LEU	Peptide
36	RR	92	GLY	Peptide
37	RS	105	ALA	Peptide
37	RS	109	GLY	Peptide
37	RS	110	LEU	Peptide
37	RS	111	GLU	Peptide
37	RS	22	GLY	Peptide
37	RS	42	ASP	Peptide
37	RS	56	LEU	Peptide
37	RS	58	LEU	Peptide
37	RS	61	ASN	Peptide
37	RS	62	LYS	Peptide
37	RS	72	ALA	Peptide
38	RT	1	MET	Peptide
38	RT	11	GLU	Peptide
38	RT	111	ARG	Peptide
38	RT	12	SER	Peptide
38	RT	122	ASP	Peptide
38	RT	123	GLN	Peptide
38	RT	135	ALA	Peptide
38	RT	3	ARG	Peptide
38	RT	31	SER	Peptide
38	RT	36	GLU	Peptide
38	RT	38	ASN	Peptide
38	RT	56	GLY	Peptide
38	RT	58	ASN	Peptide
38	RT	91	ARG	Peptide
38	RT	93	ARG	Peptide
39	RU	116	ALA	Peptide
39	RU	91	ASP	Peptide
39	RU	92	ARG	Peptide
39	RU	95	LEU	Peptide
39	RU	96	ALA	Peptide
40	RV	100	ARG	Peptide
40	RV	18	LEU	Peptide
40	RV	48	GLY	Peptide
40	RV	49	THR	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
40	RV	52	VAL	Peptide
40	RV	54	GLY	Peptide
41	RW	110	LYS	Peptide
41	RW	64	MET	Peptide
41	RW	72	LYS	Peptide
41	RW	91	GLY	Peptide
41	RW	92	ARG	Peptide
42	RX	27	THR	Peptide
42	RX	93	GLU	Peptide
43	RY	10	GLY	Peptide
43	RY	106	LEU	Peptide
43	RY	25	GLY	Peptide
43	RY	39	VAL	Peptide
43	RY	52	SER	Peptide
43	RY	71	LYS	Peptide
43	RY	9	LYS	Peptide
44	RZ	108	PRO	Peptide
44	RZ	11	GLU	Peptide
44	RZ	110	GLY	Peptide
44	RZ	113	ALA	Peptide
44	RZ	114	GLY	Peptide
44	RZ	12	GLY	Peptide
44	RZ	127	LYS	Peptide
44	RZ	130	PRO	Peptide
44	RZ	148	ASP	Peptide
44	RZ	152	ALA	Peptide
44	RZ	153	SER	Peptide
44	RZ	154	ASP	Peptide
44	RZ	156	LYS	Peptide
44	RZ	158	PRO	Peptide
44	RZ	160	GLY	Peptide
44	RZ	162	GLU	Peptide
44	RZ	164	ALA	Peptide
44	RZ	165	VAL	Peptide
44	RZ	167	PRO	Peptide
44	RZ	168	GLU	Peptide
44	RZ	178	GLU	Peptide
44	RZ	180	VAL	Peptide
44	RZ	4	ARG	Peptide
44	RZ	5	LEU	Peptide
44	RZ	51	ALA	Peptide
44	RZ	52	SER	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
44	RZ	53	ILE	Peptide
44	RZ	58	VAL	Peptide
44	RZ	59	LEU	Peptide
44	RZ	60	GLU	Peptide
44	RZ	62	PRO	Peptide
44	RZ	63	ASP	Peptide
44	RZ	65	GLN	Peptide
44	RZ	70	LEU	Peptide
44	RZ	76	LEU	Peptide
44	RZ	92	SER	Peptide
44	RZ	93	ASP	Peptide
44	RZ	95	PRO	Peptide
1	XA	358	U	Sidechain
2	XB	15	VAL	Peptide
2	XB	160	ASP	Peptide
2	XB	168	THR	Peptide
2	XB	18	GLY	Peptide
2	XB	19	HIS	Peptide
2	XB	225	ALA	Peptide
2	XB	227	GLY	Peptide
2	XB	232	PRO	Peptide
2	XB	233	SER	Peptide
2	XB	235	SER	Peptide
2	XB	239	VAL	Peptide
2	XB	6	THR	Peptide
2	XB	64	ARG	Peptide
2	XB	95	GLN	Peptide
2	XB	96	ARG	Peptide
3	XC	11	ARG	Peptide
3	XC	112	SER	Peptide
3	XC	147	LYS	Peptide
3	XC	166	GLU	Peptide
3	XC	168	ALA	Peptide
3	XC	189	ALA	Peptide
3	XC	62	ASP	Peptide
3	XC	78	GLY	Peptide
3	XC	92	ALA	Peptide
4	XD	141	ARG	Peptide
4	XD	143	GLY	Peptide
4	XD	146	ILE	Peptide
4	XD	148	VAL	Peptide
4	XD	153	ARG	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
4	XD	154	ASN	Peptide
4	XD	170	VAL	Peptide
4	XD	174	LEU	Peptide
4	XD	177	ASP	Peptide
4	XD	183	GLY	Peptide
4	XD	207	TYR	Peptide
4	XD	30	LYS	Mainchain
4	XD	47	ARG	Peptide
4	XD	83	SER	Peptide
4	XD	84	LYS	Peptide
5	XE	112	LEU	Peptide
5	XE	16	THR	Peptide
5	XE	7	GLU	Peptide
5	XE	72	GLN	Peptide
5	XE	73	ASN	Peptide
5	XE	74	GLY	Peptide
6	XF	34	GLY	Peptide
6	XF	35	ALA	Peptide
6	XF	80	ARG	Peptide
7	XG	11	GLN	Peptide
7	XG	113	GLU	Peptide
7	XG	139	GLU	Peptide
7	XG	5	ARG	Peptide
7	XG	57	GLU	Peptide
7	XG	79	ARG	Peptide
7	XG	8	GLU	Peptide
7	XG	82	GLY	Peptide
7	XG	86	GLN	Peptide
7	XG	97	GLN	Peptide
8	XH	100	ILE	Peptide
8	XH	42	GLU	Peptide
8	XH	70	GLN	Peptide
8	XH	71	GLY	Peptide
8	XH	96	GLY	Peptide
9	XI	113	LYS	Peptide
9	XI	126	SER	Peptide
9	XI	13	ALA	Peptide
9	XI	15	ALA	Peptide
9	XI	25	LYS	Peptide
9	XI	34	ASN	Peptide
9	XI	42	ARG	Peptide
9	XI	48	GLU	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
9	XI	49	PRO	Peptide
9	XI	56	LEU	Peptide
9	XI	58	HIS	Peptide
9	XI	69	GLY	Peptide
9	XI	71	SER	Peptide
9	XI	86	VAL	Peptide
9	XI	9	ARG	Peptide
9	XI	99	LEU	Peptide
10	XJ	30	SER	Peptide
10	XJ	32	ALA	Peptide
10	XJ	36	GLY	Peptide
10	XJ	52	GLY	Peptide
10	XJ	54	PHE	Peptide
10	XJ	85	LEU	Peptide
10	XJ	91	PRO	Peptide
10	XJ	99	LYS	Peptide
11	XK	100	ALA	Peptide
11	XK	102	GLY	Peptide
11	XK	91	ARG	Peptide
11	XK	94	ALA	Peptide
12	XL	102	ARG	Peptide
12	XL	103	GLY	Peptide
12	XL	104	VAL	Peptide
12	XL	118	SER	Peptide
12	XL	120	TYR	Peptide
12	XL	17	LYS	Peptide
12	XL	18	VAL	Peptide
12	XL	28	LYS	Peptide
12	XL	41	ARG	Peptide
12	XL	47	LYS	Peptide
12	XL	59	ARG	Peptide
13	XM	10	PRO	Peptide
13	XM	105	THR	Peptide
13	XM	106	ASN	Peptide
13	XM	107	ALA	Peptide
13	XM	11	ARG	Peptide
13	XM	113	PRO	Peptide
13	XM	115	LYS	Peptide
13	XM	116	THR	Peptide
13	XM	30	ALA	Peptide
13	XM	4	ILE	Peptide
13	XM	46	LYS	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
13	XM	5	ALA	Peptide
13	XM	62	ASN	Peptide
13	XM	65	LYS	Peptide
13	XM	66	LEU	Peptide
13	XM	70	LEU	Peptide
13	XM	9	ILE	Peptide
13	XM	98	VAL	Peptide
14	XN	42	ILE	Peptide
14	XN	44	LEU	Peptide
15	XO	19	PRO	Peptide
15	XO	77	ARG	Peptide
16	XP	10	GLY	Peptide
16	XP	12	LYS	Peptide
16	XP	19	ILE	Peptide
16	XP	34	GLU	Peptide
16	XP	36	ILE	Peptide
16	XP	44	THR	Peptide
16	XP	62	VAL	Peptide
16	XP	64	ALA	Peptide
16	XP	82	GLN	Peptide
16	XP	83	GLU	Peptide
17	XQ	66	SER	Peptide
18	XR	20	ALA	Peptide
18	XR	21	LYS	Peptide
18	XR	26	LEU	Peptide
18	XR	31	LEU	Peptide
18	XR	38	GLU	Peptide
18	XR	55	ARG	Peptide
19	XS	10	PHE	Peptide
19	XS	11	VAL	Peptide
19	XS	20	LEU	Peptide
19	XS	24	ALA	Peptide
19	XS	26	GLY	Peptide
19	XS	27	GLU	Peptide
19	XS	30	LEU	Peptide
19	XS	4	SER	Peptide
19	XS	44	MET	Peptide
19	XS	47	HIS	Peptide
19	XS	48	THR	Peptide
19	XS	60	VAL	Peptide
19	XS	84	GLY	Peptide
19	XS	9	VAL	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
20	XT	11	SER	Peptide
20	XT	12	ALA	Peptide
20	XT	33	ILE	Peptide
20	XT	48	LYS	Peptide
20	XT	49	ALA	Peptide
20	XT	71	THR	Peptide
20	XT	72	LEU	Peptide
20	XT	73	HIS	Peptide
20	XT	8	ARG	Peptide
20	XT	94	ALA	Peptide
20	XT	95	ALA	Peptide
20	XT	96	GLY	Peptide
21	XU	14	TRP	Peptide
21	XU	16	GLY	Peptide
21	XU	19	GLY	Peptide
21	XU	21	TYR	Peptide
45	Y0	42	GLY	Peptide
46	Y1	30	VAL	Peptide
46	Y1	53	VAL	Peptide
46	Y1	54	ALA	Peptide
46	Y1	74	VAL	Peptide
46	Y1	80	LEU	Peptide
46	Y1	85	LEU	Peptide
46	Y1	87	PRO	Peptide
47	Y2	14	ARG	Peptide
47	Y2	15	LYS	Peptide
47	Y2	4	SER	Peptide
47	Y2	43	GLN	Peptide
47	Y2	46	GLN	Peptide
47	Y2	47	ASN	Peptide
48	Y3	2	PRO	Peptide
48	Y3	33	GLN	Peptide
49	Y4	46	GLN	Peptide
49	Y4	48	ARG	Peptide
49	Y4	50	VAL	Peptide
49	Y4	51	ASP	Peptide
49	Y4	55	ARG	Peptide
49	Y4	56	VAL	Peptide
49	Y4	60	GLN	Peptide
49	Y4	61	ARG	Peptide
49	Y4	62	ARG	Peptide
49	Y4	63	TYR	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
49	Y4	65	ASP	Peptide
53	Y8	44	LYS	Peptide
53	Y8	52	LYS	Peptide
53	Y8	60	LEU	Peptide
53	Y8	62	LEU	Peptide
53	Y8	63	PRO	Peptide
53	Y8	64	TYR	Peptide
24	YA	1003	U	Sidechain
24	YA	2774	G	Sidechain
26	YD	110	GLY	Peptide
26	YD	236	GLY	Peptide
26	YD	246	PRO	Peptide
27	YE	130	GLY	Peptide
27	YE	142	GLY	Peptide
27	YE	157	ALA	Peptide
27	YE	186	GLY	Peptide
27	YE	29	GLY	Peptide
27	YE	51	PHE	Peptide
27	YE	68	ALA	Peptide
27	YE	69	LYS	Peptide
27	YE	70	ALA	Peptide
27	YE	71	GLY	Peptide
28	YF	127	GLU	Peptide
28	YF	128	ALA	Peptide
28	YF	130	ALA	Peptide
28	YF	132	VAL	Peptide
28	YF	15	SER	Peptide
28	YF	196	LEU	Peptide
28	YF	198	ALA	Peptide
28	YF	206	ILE	Peptide
28	YF	22	ALA	Peptide
28	YF	23	ASP	Peptide
28	YF	25	PRO	Peptide
28	YF	6	VAL	Peptide
28	YF	67	GLN	Peptide
28	YF	86	GLY	Peptide
28	YF	91	GLY	Peptide
29	YG	109	VAL	Peptide
29	YG	118	ARG	Peptide
29	YG	127	GLY	Peptide
29	YG	134	GLY	Peptide
29	YG	135	LEU	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
29	YG	136	ARG	Peptide
29	YG	144	ILE	Peptide
29	YG	149	VAL	Peptide
29	YG	150	ASP	Peptide
29	YG	35	GLU	Peptide
29	YG	43	LEU	Peptide
29	YG	44	GLY	Peptide
29	YG	45	GLU	Peptide
29	YG	46	ALA	Peptide
29	YG	51	ARG	Peptide
29	YG	52	ILE	Peptide
29	YG	58	GLN	Peptide
29	YG	79	ASN	Peptide
29	YG	82	LEU	Peptide
29	YG	95	ARG	Peptide
30	YH	174	GLY	Peptide
31	YI	10	GLU	Peptide
31	YI	103	ARG	Peptide
31	YI	11	ASN	Peptide
31	YI	116	LEU	Peptide
31	YI	119	PRO	Peptide
31	YI	121	LYS	Peptide
31	YI	122	GLU	Peptide
31	YI	123	LEU	Peptide
31	YI	124	GLY	Peptide
31	YI	131	LYS	Peptide
31	YI	133	HIS	Peptide
31	YI	135	GLU	Peptide
31	YI	14	ASP	Peptide
31	YI	142	VAL	Peptide
31	YI	144	VAL	Peptide
31	YI	15	VAL	Peptide
31	YI	16	GLY	Peptide
31	YI	35	LEU	Peptide
31	YI	63	ALA	Peptide
31	YI	79	ILE	Peptide
31	YI	9	LEU	Peptide
32	YN	113	GLY	Peptide
32	YN	125	GLY	Peptide
32	YN	126	PRO	Peptide
32	YN	17	ASP	Peptide
32	YN	21	LYS	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
32	YN	22	THR	Peptide
32	YN	23	LEU	Peptide
33	YO	96	THR	Peptide
34	YP	115	LEU	Peptide
34	YP	141	ALA	Peptide
34	YP	28	GLY	Peptide
34	YP	35	HIS	Peptide
34	YP	43	GLY	Peptide
35	YQ	108	GLY	Peptide
35	YQ	15	GLY	Peptide
35	YQ	22	LYS	Peptide
35	YQ	40	ALA	Peptide
35	YQ	59	ARG	Peptide
35	YQ	62	GLY	Peptide
36	YR	117	VAL	Peptide
36	YR	3	HIS	Peptide
36	YR	73	VAL	Peptide
36	YR	75	LEU	Peptide
36	YR	8	ARG	Peptide
37	YS	109	GLY	Peptide
38	YT	110	ILE	Peptide
38	YT	111	ARG	Peptide
38	YT	122	ASP	Peptide
38	YT	123	GLN	Peptide
38	YT	127	ALA	Peptide
38	YT	27	THR	Peptide
38	YT	36	GLU	Peptide
38	YT	38	ASN	Peptide
38	YT	56	GLY	Peptide
38	YT	58	ASN	Peptide
38	YT	93	ARG	Peptide
39	YU	117	GLN	Peptide
39	YU	90	VAL	Peptide
39	YU	91	ASP	Peptide
39	YU	95	LEU	Peptide
39	YU	96	ALA	Peptide
40	YV	30	GLY	Peptide
40	YV	31	ALA	Peptide
40	YV	44	LYS	Peptide
40	YV	49	THR	Peptide
40	YV	54	GLY	Peptide
40	YV	55	ALA	Peptide

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
40	YV	65	GLY	Peptide
41	YW	110	LYS	Peptide
41	YW	64	MET	Peptide
41	YW	90	ARG	Peptide
41	YW	91	GLY	Peptide
41	YW	92	ARG	Peptide
42	YX	93	GLU	Peptide
43	YY	106	LEU	Peptide
43	YY	51	VAL	Peptide
43	YY	52	SER	Peptide
43	YY	54	LYS	Peptide
43	YY	71	LYS	Peptide
44	YZ	109	ALA	Peptide
44	YZ	111	VAL	Peptide
44	YZ	114	GLY	Peptide
44	YZ	136	PHE	Peptide
44	YZ	160	GLY	Peptide
44	YZ	168	GLU	Peptide
44	YZ	29	TYR	Peptide
44	YZ	37	VAL	Peptide
44	YZ	50	GLN	Peptide
44	YZ	51	ALA	Peptide

5.2 Too-close contacts [i](#)

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	QA	32247	0	16277	448	0
1	XA	32249	0	16278	456	3
2	QB	1907	0	1958	34	0
2	XB	1915	0	1969	35	0
3	QC	1605	0	1668	29	0
3	XC	1605	0	1668	21	0
4	QD	1703	0	1766	27	0
4	XD	1703	0	1764	37	0
5	QE	1155	0	1213	27	0
5	XE	1155	0	1213	12	0
6	QF	843	0	857	14	0

Continued on next page...

Continued from previous page...

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
6	XF	843	0	857	11	0
7	QG	1257	0	1296	15	0
7	XG	1257	0	1296	22	0
8	QH	1108	0	1165	19	0
8	XH	1108	0	1165	23	0
9	QI	1010	0	1037	19	0
9	XI	998	0	1024	28	0
10	QJ	801	0	849	23	0
10	XJ	777	0	816	15	0
11	QK	885	0	904	28	0
11	XK	864	0	881	13	0
12	QL	975	0	1062	13	0
12	XL	956	0	1046	22	0
13	QM	955	0	1021	21	0
13	XM	946	0	1008	24	0
14	QN	492	0	529	19	0
14	XN	492	0	529	11	0
15	QO	734	0	771	9	0
15	XO	729	0	768	8	0
16	QP	705	0	725	15	0
16	XP	705	0	725	12	0
17	QQ	834	0	904	24	0
17	XQ	834	0	904	15	0
18	QR	574	0	644	7	0
18	XR	574	0	644	7	0
19	QS	665	0	686	8	0
19	XS	674	0	699	10	0
20	QT	763	0	861	13	0
20	XT	763	0	861	22	0
21	QU	217	0	234	4	0
21	XU	217	0	234	4	0
22	QV	1648	0	834	28	0
22	XV	1648	0	834	23	0
23	QX	418	0	209	7	0
23	XX	418	0	209	7	0
24	RA	62051	0	31279	607	3
24	YA	62091	0	31292	530	0
25	RB	2573	0	1306	22	0
25	YB	2573	0	1304	41	0
26	RD	2135	0	2221	35	0
26	YD	2135	0	2221	32	0
27	RE	1568	0	1634	29	0

Continued on next page...

Continued from previous page...

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
27	YE	1563	0	1628	27	0
28	RF	1585	0	1632	23	0
28	YF	1585	0	1632	24	0
29	RG	1474	0	1535	35	0
29	YG	1474	0	1535	30	0
30	RH	1336	0	1418	27	0
30	YH	1330	0	1413	18	0
31	RI	1136	0	1223	13	2
31	YI	1136	0	1223	12	0
32	RN	1104	0	1180	8	0
32	YN	1121	0	1194	15	0
33	RO	933	0	996	25	0
33	YO	933	0	995	17	0
34	RP	1139	0	1223	26	0
34	YP	1122	0	1206	26	0
35	RQ	1122	0	1179	27	0
35	YQ	1122	0	1179	10	0
36	RR	960	0	1021	18	0
36	YR	960	0	1021	11	0
37	RS	882	0	943	17	0
37	YS	877	0	938	15	0
38	RT	1141	0	1202	23	0
38	YT	1141	0	1202	26	0
39	RU	964	0	1022	25	0
39	YU	964	0	1021	19	0
40	RV	779	0	852	14	0
40	YV	779	0	852	12	0
41	RW	900	0	964	11	0
41	YW	900	0	964	11	0
42	RX	725	0	778	7	0
42	YX	742	0	803	8	0
43	RY	818	0	909	11	0
43	YY	818	0	909	13	0
44	RZ	1461	0	1493	26	0
44	YZ	1461	0	1493	22	0
45	R0	643	0	667	12	0
45	Y0	599	0	617	7	0
46	R1	763	0	848	9	0
46	Y1	729	0	801	10	0
47	R2	581	0	629	7	0
47	Y2	558	0	610	5	2
48	R3	469	0	517	7	0

Continued on next page...

Continued from previous page...

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
48	Y3	469	0	518	8	0
49	R4	565	0	558	9	0
49	Y4	565	0	557	13	0
50	R5	459	0	476	7	0
50	Y5	459	0	476	9	0
51	R6	453	0	473	7	0
51	Y6	453	0	473	10	0
52	R7	409	0	454	6	0
52	Y7	418	0	467	3	0
53	R8	517	0	582	12	0
53	Y8	517	0	582	13	0
54	R9	307	0	335	6	0
54	Y9	307	0	335	7	0
55	QA	64	0	0	0	0
55	QV	1	0	0	0	0
55	R0	2	0	0	0	0
55	R8	2	0	0	0	0
55	RA	444	0	0	0	0
55	RB	7	0	0	0	0
55	RE	5	0	0	0	0
55	RF	1	0	0	0	0
55	RN	1	0	0	0	0
55	RQ	1	0	0	0	0
55	RR	1	0	0	0	0
55	RT	1	0	0	0	0
55	RX	1	0	0	0	0
55	XA	78	0	0	0	0
55	XE	1	0	0	0	0
55	XM	2	0	0	0	0
55	XV	1	0	0	0	0
55	Y0	2	0	0	0	0
55	Y1	2	0	0	0	0
55	Y3	1	0	0	0	0
55	Y5	1	0	0	0	0
55	Y7	1	0	0	0	0
55	Y8	1	0	0	0	0
55	YA	510	0	0	0	0
55	YB	7	0	0	0	0
55	YD	2	0	0	0	0
55	YE	5	0	0	0	0
55	YP	3	0	0	0	0
55	YQ	3	0	0	0	0

Continued on next page...

Continued from previous page...

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
55	YR	1	0	0	0	0
56	QD	8	0	0	0	0
56	XD	8	0	0	0	0
57	QN	1	0	0	0	0
57	R4	1	0	0	0	0
57	R5	1	0	0	0	0
57	R6	1	0	0	0	0
57	R9	1	0	0	0	0
57	RY	1	0	0	0	0
57	XN	1	0	0	0	0
57	Y4	1	0	0	0	0
57	Y5	1	0	0	0	0
57	Y6	1	0	0	0	0
57	Y9	1	0	0	0	0
57	YY	1	0	0	0	0
All	All	291964	0	197872	3283	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 7.

All (3283) 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)
24:RA:1712:A:C2'	24:RA:1713:G:H5'	1.74	1.18
43:YY:92:ASN:ND2	43:YY:94:LYS:HG2	1.60	1.15
24:YA:1405:A:N6	24:YA:1418:U:H3	1.48	1.10
24:RA:2818:U:C2	24:RA:2901:A:N6	2.27	1.02
24:RA:2226:C:H1'	24:RA:2232:G:N2	1.74	1.01
43:YY:92:ASN:HD22	43:YY:94:LYS:HG2	1.25	1.01
24:RA:1712:A:H2'	24:RA:1713:G:H5'	1.02	1.00
24:RA:2157:A:H62	24:RA:2178:G:N2	1.59	0.99
1:XA:358:U:H3'	1:XA:359:U:C6	1.97	0.99
24:RA:331:G:H8	24:RA:331:G:H5''	1.27	0.98
24:RA:2157:A:N6	24:RA:2178:G:H21	1.60	0.98
25:YB:109:G:H8	25:YB:109:G:H5''	1.26	0.98
1:QA:1009:G:H1	1:QA:1020:U:H3	1.02	0.98
1:XA:438:G:H21	1:XA:496:A:H62	1.03	0.97
24:RA:1712:A:H2'	24:RA:1713:G:C5'	1.95	0.96
24:RA:2157:A:H62	24:RA:2178:G:H21	0.96	0.96
1:XA:835:U:H3	1:XA:851:G:H1	1.12	0.96
25:RB:22:U:H3	25:RB:61:G:H1	0.98	0.95

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:157:G:H1	1:XA:164:U:H3	1.14	0.94
24:YA:2120:U:H3	24:YA:2213:G:H1	1.07	0.94
1:XA:1238:A:H62	1:XA:1301:U:H3	1.15	0.92
1:XA:1052:U:H3	1:XA:1206:G:H1	1.18	0.91
1:XA:992:U:H3	1:XA:1044:A:H62	1.15	0.90
1:XA:452:A:H62	1:XA:480:U:H3	1.08	0.90
24:RA:1712:A:C2'	24:RA:1713:G:C5'	2.51	0.89
43:YY:92:ASN:HD22	43:YY:94:LYS:CG	1.85	0.89
24:RA:2257:U:H5'	24:RA:2258:G:H5'	1.56	0.88
1:XA:152:A:H62	1:XA:169:C:N4	1.71	0.87
43:YY:92:ASN:ND2	43:YY:94:LYS:CG	2.38	0.87
24:YA:2118:U:H3	24:YA:2215:G:H1	0.88	0.87
1:QA:666:G:H1	1:QA:740:U:H3	1.18	0.87
24:RA:334:A:OP1	43:RY:18:GLY:N	2.06	0.86
24:RA:1713:G:N2	24:RA:2016:C:O2	2.08	0.86
1:XA:438:G:N2	1:XA:496:A:H62	1.71	0.86
1:XA:452:A:N6	1:XA:480:U:H3	1.72	0.86
19:QS:50:ALA:HA	19:QS:58:VAL:O	1.75	0.85
24:RA:2118:U:H3	24:RA:2215:G:H1	1.23	0.85
1:XA:358:U:H3'	1:XA:359:U:H6	1.40	0.85
24:YA:2121:U:H3	24:YA:2212:G:H1	1.24	0.85
24:RA:2226:C:H1'	24:RA:2232:G:H22	1.41	0.84
1:QA:565:U:OP2	1:QA:566:G:O2'	1.96	0.84
1:XA:438:G:H21	1:XA:496:A:N6	1.75	0.83
24:RA:354:A:H2	24:RA:1255:A:H2'	1.44	0.83
24:YA:1058:U:O2'	24:YA:1059:C:OP2	1.97	0.82
30:YH:56:SER:HG	30:YH:61:HIS:HD1	1.26	0.82
24:RA:1405:A:H62	24:RA:1418:U:H3	1.28	0.82
24:YA:1310:G:OP1	50:Y5:19:ARG:NH2	2.09	0.81
24:YA:2257:U:H5'	24:YA:2258:G:H5'	1.61	0.81
24:RA:331:G:C8	24:RA:331:G:H5''	2.15	0.80
4:XD:32:ALA:O	4:XD:35:ARG:N	2.15	0.80
25:YB:8:U:H3	25:YB:113:C:H42	1.28	0.80
1:XA:152:A:N6	1:XA:169:C:H42	1.80	0.79
24:RA:2713:C:H3'	24:RA:2714:U:H5''	1.62	0.79
1:XA:1238:A:N6	1:XA:1301:U:H3	1.80	0.79
24:YA:2877:G:O2'	24:YA:2878:A:OP2	1.99	0.79
1:XA:1266:G:N2	1:XA:1269:A:OP2	2.16	0.79
1:XA:152:A:H62	1:XA:169:C:H42	1.28	0.79
24:RA:2074:G:H4'	27:RE:143:ASN:H	1.48	0.79
24:YA:1040:C:OP1	39:YU:53:ARG:NH2	2.16	0.78

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:RA:1653:C:N4	24:RA:1668:G:OP2	2.15	0.78
24:YA:1653:C:N4	24:YA:1668:G:OP2	2.17	0.78
7:XG:51:GLN:HE22	7:XG:58:PRO:HG3	1.49	0.77
24:YA:542:C:OP1	50:Y5:16:ARG:NH2	2.16	0.77
1:QA:1528:U:O2'	1:QA:1529:G:OP2	2.02	0.77
24:YA:1039:G:OP1	39:YU:50:ARG:NH2	2.17	0.77
24:YA:2032:G:H5''	41:YW:42:ARG:HB2	1.65	0.77
24:YA:2877:G:OP2	38:YT:119:LYS:NZ	2.13	0.77
24:RA:1714:G:N2	24:RA:2015:U:O4	2.17	0.76
24:RA:2801:C:O2'	24:RA:2819:A:N3	2.19	0.76
1:QA:1316:G:N2	1:QA:1319:A:OP2	2.19	0.76
1:XA:950:U:H3	1:XA:1231:G:H1	1.30	0.76
1:XA:372:C:H42	1:XA:389:A:H62	1.32	0.76
1:XA:686:U:H1'	11:XK:42:TRP:HE1	1.50	0.76
24:RA:2760:G:H21	24:RA:2770:A:H62	1.33	0.75
25:YB:82:G:H1	25:YB:95:U:H3	1.34	0.75
24:RA:2226:C:O2	24:RA:2232:G:C2	2.40	0.75
1:QA:686:U:H1'	11:QK:42:TRP:HE1	1.51	0.75
24:RA:552:C:N4	24:RA:2792:U:OP2	2.20	0.75
24:RA:331:G:H8	24:RA:331:G:C5'	2.00	0.74
1:QA:1441:G:H21	1:QA:1460:A:H62	1.32	0.74
24:RA:823:G:N1	24:RA:2094:G:OP1	2.15	0.74
1:XA:73:G:H1	1:XA:97:U:H3	1.33	0.74
1:QA:593:G:H1	1:QA:646:U:H3	1.33	0.74
1:XA:410:G:H21	1:XA:432:A:H62	1.33	0.74
24:RA:1040:C:OP1	39:RU:53:ARG:NH2	2.21	0.74
24:YA:723:A:H8	24:YA:2091:G:H21	1.33	0.74
1:QA:1216:G:OP1	14:QN:2:ALA:N	2.21	0.74
1:XA:1356:G:H2'	1:XA:1357:A:H8	1.52	0.73
25:RB:80:U:H2'	25:RB:81:G:H21	1.53	0.73
1:XA:151:A:H62	1:XA:170:U:H3	1.36	0.73
32:YN:12:ARG:NH2	32:YN:50:ASP:OD2	2.20	0.73
24:RA:555:G:O2'	24:RA:557:A:N7	2.22	0.73
24:RA:2818:U:N3	24:RA:2901:A:N6	2.37	0.73
24:YA:1627:A:H5''	24:YA:1628:G:N7	2.02	0.73
25:YB:109:G:C8	25:YB:109:G:H5''	2.17	0.73
1:XA:152:A:N6	1:XA:169:C:N4	2.37	0.73
1:XA:358:U:C6	1:XA:358:U:C5'	2.72	0.73
24:RA:680:G:H1	24:RA:699:C:H42	1.37	0.72
24:RA:2849:G:H5'	36:RR:46:GLY:HA2	1.71	0.72
24:YA:1766:G:N1	24:YA:1768:U:OP2	2.22	0.72

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:YA:1405:A:N7	24:YA:1418:U:O4	2.22	0.72
24:RA:1810:U:OP2	24:RA:1815:A:N6	2.18	0.72
14:QN:29:ARG:HG3	14:QN:31:ARG:H	1.53	0.71
24:YA:1555:C:H3'	24:YA:1556:A:H5''	1.72	0.71
24:YA:2074:G:H4'	27:YE:143:ASN:H	1.55	0.71
24:RA:2892:A:OP1	36:RR:96:ARG:NH1	2.22	0.71
1:XA:553:A:H5''	12:XL:24:VAL:HG21	1.72	0.71
24:YA:2090:U:H3	24:YA:2442:A:H2	1.37	0.71
1:XA:992:U:H3	1:XA:1044:A:N6	1.88	0.71
1:XA:544:G:OP1	4:XD:59:ARG:NH2	2.23	0.71
1:XA:410:G:N2	1:XA:432:A:H62	1.88	0.70
1:QA:31:G:O2'	1:QA:48:C:N4	2.24	0.70
14:QN:27:CYS:SG	14:QN:28:GLY:N	2.64	0.70
24:RA:1848:G:OP1	26:RD:88:ARG:NH2	2.24	0.70
1:QA:674:G:H2'	1:QA:675:A:H8	1.56	0.70
24:RA:2818:U:N3	24:RA:2901:A:C6	2.60	0.70
1:XA:1210:C:O2'	1:XA:1213:A:O2'	2.08	0.70
6:QF:75:LEU:O	6:QF:78:GLU:HB3	1.90	0.70
24:RA:27:G:N2	24:RA:538:A:OP2	2.25	0.70
24:YA:248:G:O2'	24:YA:646:A:O2'	2.09	0.70
24:YA:2692:C:OP2	27:YE:111:ARG:NH2	2.25	0.70
24:YA:553:A:HO2'	24:YA:554:A:C5'	2.04	0.70
1:QA:1133:G:H1	1:QA:1141:C:H42	1.40	0.69
24:YA:301:C:H6	24:YA:301:C:H5''	1.57	0.69
24:YA:1716:A:N3	24:YA:1716:A:H2'	2.06	0.69
1:QA:410:G:H21	1:QA:432:A:H62	1.40	0.69
24:YA:1499:C:O3'	24:YA:1504:A:N6	2.26	0.69
1:QA:1227:A:H5'	13:QM:111:LYS:HD3	1.75	0.69
24:RA:815:G:O2'	24:RA:1425:A:N6	2.25	0.69
24:YA:28:A:N6	24:YA:537:G:O2'	2.25	0.69
1:XA:359:U:H2'	1:XA:360:A:C8	2.28	0.69
22:QV:30:C:H2'	22:QV:31:G:H8	1.57	0.69
1:XA:359:U:H2'	1:XA:360:A:H8	1.57	0.69
23:XX:3:C:H2'	23:XX:4:A:H8	1.57	0.69
24:YA:1132:A:O2'	24:YA:1149:A:N6	2.22	0.69
37:YS:61:ASN:HD22	37:YS:64:GLU:HG3	1.57	0.69
1:QA:1106:G:H5''	3:QC:172:ARG:HG2	1.74	0.69
1:XA:439:A:OP2	1:XA:493:G:N1	2.21	0.69
22:XV:58:A:O2'	22:XV:60:U:OP2	2.10	0.69
1:QA:833:U:H3	1:QA:853:G:H1	1.41	0.69
49:Y4:16:CYS:SG	49:Y4:17:GLY:N	2.66	0.69

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:664:G:H22	1:QA:741:G:H1	1.39	0.68
24:YA:1042:A:OP2	39:YU:92:ARG:NH2	2.26	0.68
24:YA:301:C:H5'	24:YA:302:A:H5''	1.74	0.68
9:XI:112:LYS:HA	9:XI:119:ALA:HB2	1.75	0.68
24:YA:1104:G:H2'	24:YA:1105:G:H8	1.58	0.68
1:QA:376:G:H5''	16:QP:5:ARG:HB2	1.76	0.68
1:XA:358:U:C6	1:XA:358:U:H5''	2.28	0.68
24:RA:238:C:O2'	34:RP:64:LYS:NZ	2.27	0.68
43:RY:18:GLY:O	43:RY:21:LYS:HD3	1.94	0.68
1:QA:1422:G:H5''	33:RO:48:PRO:HB3	1.76	0.68
1:XA:674:G:H2'	1:XA:675:A:H8	1.59	0.68
34:YP:63:PRO:HB2	53:Y8:30:ARG:HH21	1.59	0.68
24:RA:723:A:H8	24:RA:2091:G:H21	1.40	0.68
34:RP:93:GLY:H	34:RP:123:LEU:HD22	1.58	0.68
23:XX:5:A:H2'	23:XX:6:G:H8	1.58	0.68
24:YA:241:G:OP2	34:YP:50:ARG:NH2	2.26	0.68
24:RA:2764:G:N7	30:RH:2:SER:OG	2.26	0.67
26:RD:8:PRO:HB3	26:RD:14:ARG:HB3	1.76	0.67
28:RF:143:ALA:HB1	28:RF:148:LEU:HB2	1.75	0.67
24:YA:935:C:H3'	24:YA:936:C:H4'	1.75	0.67
7:QG:116:ALA:HA	7:QG:119:ARG:HE	1.60	0.67
1:XA:76:G:H1	1:XA:93:U:H3	1.42	0.67
1:QA:1224:G:O2'	1:QA:1322:C:OP2	2.13	0.67
13:QM:23:TYR:HB3	13:QM:67:GLU:HG2	1.76	0.67
1:XA:362:G:N2	1:XA:365:U:OP2	2.27	0.67
1:XA:452:A:N6	1:XA:480:U:N3	2.36	0.67
1:QA:316:G:OP2	1:QA:351:G:O2'	2.11	0.67
1:QA:838:G:H1	1:QA:848:C:H42	1.42	0.67
1:QA:1080:A:H5'	5:QE:16:THR:HG21	1.77	0.67
22:QV:30:C:C2	22:QV:31:G:C8	2.83	0.67
1:XA:323:U:OP1	20:XT:26:ASN:ND2	2.26	0.67
1:QA:152:A:H62	1:QA:169:C:N4	1.92	0.67
24:RA:539:A:N3	24:RA:604:C:O2'	2.25	0.67
1:XA:452:A:N7	1:XA:480:U:O4	2.28	0.67
1:QA:262:A:H5'	20:QT:74:LYS:HG3	1.77	0.67
24:RA:1555:C:H3'	24:RA:1556:A:H5''	1.77	0.67
1:XA:359:U:C6	1:XA:359:U:OP2	2.48	0.67
24:RA:2032:G:H5''	41:RW:42:ARG:HB2	1.77	0.66
1:XA:272:C:H2'	1:XA:273:A:H8	1.59	0.66
24:YA:2658:C:OP2	24:YA:2745:G:O2'	2.12	0.66
25:YB:84:C:OP1	48:Y3:15:TYR:OH	2.14	0.66

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:YA:1058:U:C2'	24:YA:1059:C:OP2	2.42	0.66
39:YU:50:ARG:O	39:YU:54:LYS:NZ	2.27	0.66
1:QA:1077:G:N2	1:QA:1080:A:OP2	2.26	0.66
24:YA:553:A:O2'	24:YA:554:A:O5'	2.13	0.66
1:QA:673:G:H2'	1:QA:674:G:C8	2.31	0.66
1:QA:1321:C:H5''	1:QA:1322:C:H5''	1.76	0.66
24:RA:656:A:OP1	34:RP:65:ARG:NH1	2.28	0.66
24:YA:2842:U:H4'	24:YA:2843:G:H5''	1.78	0.66
24:RA:2002:G:O2'	24:RA:2004:C:OP2	2.14	0.66
24:RA:23:G:OP1	24:RA:529:U:N3	2.25	0.66
1:QA:1306:A:N6	1:QA:1331:G:O2'	2.29	0.66
1:XA:380:G:N2	1:XA:383:A:OP2	2.27	0.66
24:RA:1713:G:H1'	33:RO:3:GLN:HE21	1.59	0.66
24:RA:1531:G:H1	24:RA:1550:C:H42	1.44	0.65
1:XA:618:C:H5'	1:XA:619:U:H5''	1.77	0.65
24:YA:2002:G:O2'	24:YA:2004:C:OP2	2.12	0.65
24:YA:2136:A:H62	24:YA:2191:A:H61	1.44	0.65
1:XA:1305:G:N2	1:XA:1332:A:OP2	2.29	0.65
24:RA:2226:C:C2	24:RA:2232:G:N1	2.64	0.65
24:RA:333:G:OP2	24:RA:333:G:H8	1.78	0.65
24:YA:248:G:H21	24:YA:646:A:H8	1.44	0.65
1:XA:1432:G:OP1	38:YT:108:ARG:N	2.29	0.65
1:QA:1128:C:H1'	1:QA:1146:A:H61	1.62	0.65
24:RA:2404:A:OP2	24:RA:2434:A:N6	2.30	0.65
1:QA:1182:G:H5''	1:QA:1183:A:H5'	1.79	0.65
1:QA:1226:C:OP2	13:QM:103:THR:OG1	2.11	0.65
1:QA:825:G:O2'	8:QH:12:ARG:NH2	2.29	0.65
33:RO:64:ARG:HH12	38:RT:70:VAL:HG21	1.62	0.65
1:XA:1348:U:H4'	9:XI:120:ARG:HD2	1.79	0.65
11:QK:51:LYS:H	11:QK:54:ARG:HE	1.43	0.65
1:XA:372:C:N4	1:XA:389:A:H62	1.94	0.65
1:XA:811:C:O2'	1:XA:901:A:N1	2.30	0.65
14:YN:27:CYS:SG	14:YN:28:GLY:N	2.69	0.65
24:YA:2316:G:H22	24:YA:2324:U:H3	1.45	0.65
24:YA:248:G:HO2'	24:YA:646:A:HO2'	1.44	0.65
12:QL:113:ARG:HH21	12:QL:116:SER:HB2	1.61	0.64
44:YZ:52:SER:OG	44:YZ:53:ILE:N	2.28	0.64
1:QA:1446:A:O2'	1:QA:1447:G:O5'	2.16	0.64
3:QC:182:ILE:HA	3:QC:202:ILE:O	1.97	0.64
24:YA:1890:A:N6	24:YA:1905:G:O2'	2.30	0.64
24:YA:2559:U:O2	33:YO:23:ARG:NH2	2.30	0.64

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:RA:139:A:H8	24:RA:1454:C:HO2'	1.45	0.64
24:RA:1648:U:O2'	24:RA:1649:A:O5'	2.12	0.64
24:YA:24:G:O2'	41:YW:78:GLU:O	2.16	0.64
24:YA:2820:A:H61	24:YA:2900:G:H2'	1.61	0.64
5:QE:105:VAL:HG21	5:QE:128:PRO:HB3	1.78	0.64
1:QA:1329:A:N7	21:QU:7:ARG:NH2	2.46	0.64
48:R3:15:TYR:O	48:R3:20:LYS:NZ	2.31	0.64
24:RA:1140:U:OP1	24:RA:1142:A:N6	2.30	0.64
44:RZ:10:ARG:NH2	44:RZ:37:VAL:O	2.29	0.64
1:XA:1141:C:H2'	1:XA:1142:G:H8	1.62	0.64
1:XA:321:A:N6	1:XA:329:A:OP2	2.29	0.64
1:QA:1314:C:N4	19:QS:2:PRO:O	2.28	0.64
1:XA:579:G:H5'	1:XA:728:A:H1'	1.77	0.64
24:YA:1556:A:O2'	24:YA:1558:G:N7	2.30	0.64
24:YA:1440:U:H4'	24:YA:1649:A:H4'	1.79	0.64
34:YP:52:GLU:OE1	34:YP:55:ARG:NH1	2.31	0.64
24:RA:542:C:OP1	50:R5:16:ARG:NH2	2.30	0.64
1:XA:339:C:OP2	33:YO:97:ARG:NH1	2.29	0.64
24:YA:2297:C:OP2	51:Y6:6:ARG:NH2	2.30	0.64
1:QA:1129:C:H5''	1:QA:1130:A:H8	1.63	0.64
24:RA:1316:C:H5''	24:RA:1317:G:H5'	1.80	0.64
1:XA:992:U:O4	1:XA:1044:A:N7	2.31	0.64
7:QG:143:ARG:HH12	7:QG:147:ALA:HB2	1.62	0.64
24:RA:2090:U:H3	24:RA:2442:A:H2	1.45	0.64
3:XC:11:ARG:NH2	3:XC:177:THR:O	2.31	0.64
24:YA:2315:G:N3	29:YG:132:ASN:ND2	2.46	0.64
24:YA:1076:G:OP2	35:YQ:128:LYS:NZ	2.27	0.64
19:QS:50:ALA:HB1	19:QS:57:HIS:HB3	1.79	0.64
22:QV:4:U:O2'	22:QV:5:G:O5'	2.14	0.64
34:RP:50:ARG:HD3	53:R8:59:LYS:HE2	1.80	0.64
1:XA:933:G:O6	7:XG:3:ARG:NH2	2.31	0.64
24:RA:1109:G:N2	24:RA:1122:C:O2'	2.32	0.63
37:RS:34:HIS:ND1	37:RS:53:SER:OG	2.31	0.63
1:XA:266:G:O2'	1:XA:267:C:OP2	2.15	0.63
24:YA:2797:C:H1'	27:YE:37:ARG:HH12	1.63	0.63
1:XA:946:A:O2'	1:XA:1333:A:N3	2.28	0.63
1:XA:31:G:O2'	1:XA:48:C:N4	2.31	0.63
1:QA:1266:G:N2	1:QA:1269:A:OP2	2.30	0.63
24:RA:2137:G:N1	24:RA:2186:C:OP2	2.31	0.63
24:RA:2692:C:OP2	27:RE:111:ARG:NH2	2.29	0.63
25:YB:2:C:H42	25:YB:119:A:H61	1.45	0.63

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:YS:3:ARG:NH1	37:YS:4:LEU:O	2.32	0.63
1:XA:358:U:H2'	1:XA:358:U:O2	1.98	0.63
53:Y8:54:GLU:OE2	53:Y8:57:ARG:NH2	2.32	0.63
24:YA:880:U:O2	34:YP:55:ARG:NH2	2.29	0.63
1:QA:1399:C:C2	1:QA:1502:A:N6	2.66	0.63
35:RQ:17:LEU:HD13	35:RQ:39:PRO:HB2	1.81	0.63
25:YB:109:G:H8	25:YB:109:G:C5'	2.05	0.63
26:YD:85:ASP:OD2	26:YD:88:ARG:NH1	2.32	0.63
28:YF:167:ALA:HB1	28:YF:173:VAL:HG11	1.81	0.63
17:QQ:66:SER:O	17:QQ:70:ARG:NH1	2.31	0.63
1:QA:1338:G:N3	22:QV:41:A:O2'	2.31	0.63
24:RA:2877:G:OP2	38:RT:119:LYS:NZ	2.20	0.63
1:XA:264:U:O2'	17:XQ:64:PRO:O	2.15	0.63
54:R9:27:CYS:SG	54:R9:28:GLU:N	2.72	0.63
24:RA:2514:G:H5''	24:RA:2515:A:H5''	1.81	0.63
4:XD:134:ASP:OD1	4:XD:134:ASP:N	2.31	0.63
13:XM:88:ARG:HG3	13:XM:98:VAL:HG11	1.81	0.63
36:RR:3:HIS:O	36:RR:5:LYS:N	2.32	0.62
1:XA:714:G:H2'	1:XA:715:A:C8	2.34	0.62
1:XA:578:C:O2'	1:XA:728:A:N3	2.29	0.62
24:YA:2298:A:OP1	51:Y6:29:ASN:ND2	2.32	0.62
24:YA:998:A:OP2	35:YQ:16:ARG:NH1	2.31	0.62
1:QA:579:G:H5'	1:QA:728:A:H1'	1.80	0.62
1:QA:953:G:N7	13:QM:104:ARG:NH2	2.46	0.62
13:QM:118:ALA:HB1	22:QV:28:U:H4'	1.80	0.62
24:RA:354:A:H2	24:RA:1255:A:C2'	2.12	0.62
1:XA:978:A:H61	1:XA:1316:G:H1'	1.63	0.62
1:QA:745:C:OP1	1:QA:851:G:O2'	2.17	0.62
24:RA:2226:C:C2	24:RA:2232:G:C2	2.87	0.62
24:YA:2347:A:O2'	24:YA:2348:A:O5'	2.17	0.62
4:QD:13:ARG:HB3	4:QD:39:PRO:HA	1.81	0.62
24:YA:2667:G:N2	24:YA:2677:A:OP2	2.32	0.62
24:RA:2226:C:C1'	24:RA:2232:G:H22	2.12	0.62
1:XA:1133:G:H2'	1:XA:1134:G:C8	2.34	0.62
1:XA:1342:C:H2'	1:XA:1343:G:H8	1.65	0.62
1:XA:539:A:H2'	1:XA:540:G:H8	1.63	0.62
1:XA:745:C:H2'	1:XA:746:A:H8	1.63	0.62
1:QA:1023:G:H3'	1:QA:1024:G:H5''	1.80	0.62
1:QA:624:C:H2'	1:QA:625:G:H8	1.64	0.62
13:XM:81:LEU:HD21	13:XM:88:ARG:HG2	1.81	0.62
2:QB:107:THR:HA	2:QB:110:GLN:HG3	1.81	0.62

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:673:G:H2'	1:XA:674:G:C8	2.35	0.62
1:QA:581:G:OP1	15:QO:65:ARG:NH1	2.28	0.62
17:XQ:66:SER:O	17:XQ:70:ARG:NH1	2.32	0.62
24:YA:2097:U:OP2	24:YA:2250:G:O2'	2.14	0.62
39:YU:92:ARG:NH1	40:YV:11:GLN:O	2.32	0.62
1:QA:266:G:H5'	1:QA:268:C:H41	1.65	0.62
1:QA:659:U:OP1	15:QO:9:GLN:NE2	2.32	0.62
24:RA:903:C:O2'	24:RA:904:C:OP1	2.16	0.62
27:RE:201:THR:HG22	27:RE:203:LYS:H	1.65	0.62
1:XA:261:U:OP2	20:XT:79:ARG:NH2	2.33	0.62
1:XA:1352:C:OP1	21:XU:3:LYS:NZ	2.33	0.62
24:YA:2367:C:H1'	45:Y0:39:ARG:HH21	1.63	0.62
24:YA:1993:A:OP2	26:YD:242:ARG:NH2	2.32	0.62
31:YI:117:GLU:HG2	31:YI:118:LYS:HG2	1.82	0.62
22:XV:1:C:H2'	22:XV:2:G:H8	1.65	0.62
24:YA:2404:A:H2	24:YA:2436:C:H42	1.48	0.62
30:YH:46:GLU:HB2	30:YH:49:VAL:HG12	1.82	0.62
16:QP:1:MET:SD	16:QP:3:LYS:NZ	2.66	0.61
24:RA:2123:G:H2'	24:RA:2124:U:C2	2.34	0.61
24:RA:651:U:H5''	24:RA:652:A:H5'	1.82	0.61
35:RQ:83:MET:HB2	45:R0:7:LEU:HD23	1.81	0.61
47:R2:4:SER:OG	47:R2:5:GLU:N	2.33	0.61
1:XA:1238:A:N6	1:XA:1301:U:N3	2.38	0.61
1:XA:978:A:OP2	1:XA:1362(B):C:N4	2.33	0.61
1:XA:1073:U:O2'	2:XB:104:ASN:OD1	2.15	0.61
48:Y3:18:ASP:OD1	48:Y3:18:ASP:N	2.33	0.61
24:YA:2297:C:H5	51:Y6:6:ARG:HH12	1.48	0.61
26:YD:71:ASP:OD2	26:YD:103:ARG:NH2	2.33	0.61
11:QK:43:SER:OG	11:QK:44:SER:N	2.34	0.61
17:QQ:41:LYS:NZ	17:QQ:88:TYR:OH	2.33	0.61
30:RH:70:THR:O	30:RH:74:ASN:ND2	2.33	0.61
24:RA:1042:A:OP2	39:RU:92:ARG:NH2	2.33	0.61
9:XI:2:GLU:HG3	9:XI:3:GLN:HG2	1.80	0.61
12:XL:27:LEU:O	12:XL:33:ARG:NH2	2.33	0.61
24:RA:1072:U:H1'	24:RA:1073:A:H5''	1.81	0.61
24:YA:140:A:H8	24:YA:1641:G:H21	1.46	0.61
24:YA:2801:C:OP1	27:YE:61:ARG:NH2	2.34	0.61
33:YO:80:ASP:OD2	38:YT:64:ARG:NH2	2.33	0.61
6:QF:2:ARG:NH1	6:QF:3:ARG:O	2.34	0.61
9:XI:111:ARG:NH1	9:XI:112:LYS:O	2.33	0.61
37:YS:93:LYS:HG2	37:YS:95:HIS:HB2	1.81	0.61

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:RA:1115:A:H2'	24:RA:1119:A:H62	1.65	0.61
24:RA:1310:G:OP1	50:R5:19:ARG:NH2	2.33	0.61
5:XE:100:VAL:O	5:XE:107:ARG:NH2	2.34	0.61
46:Y1:87:PRO:HA	46:Y1:90:ILE:HG22	1.83	0.61
24:YA:1627:A:H5''	24:YA:1628:G:C8	2.36	0.61
24:YA:312:C:H2'	24:YA:313:A:H8	1.65	0.61
1:QA:578:C:O2'	1:QA:728:A:N3	2.28	0.61
19:QS:36:ARG:NH2	19:QS:75:ALA:O	2.34	0.61
22:QV:9:G:O2'	22:QV:10:G:N7	2.34	0.61
24:RA:2240:G:OP1	26:RD:261:LYS:NZ	2.30	0.61
6:XF:30:LEU:HB3	6:XF:35:ALA:HB3	1.82	0.61
12:XL:60:LEU:HD12	12:XL:62:SER:H	1.65	0.61
27:YE:116:VAL:HG22	27:YE:157:ALA:HB2	1.82	0.61
1:QA:1297:C:O2	7:QG:114:ARG:NH1	2.34	0.61
1:XA:1446:A:O2'	1:XA:1447:G:O5'	2.15	0.61
1:QA:1124:G:H3'	1:QA:1145:C:H42	1.66	0.61
1:QA:666:G:N2	1:QA:740:U:O2	2.32	0.61
8:QH:14:ARG:HE	8:QH:83:ILE:HG23	1.65	0.61
24:RA:1269:G:N2	24:RA:1272:A:OP2	2.30	0.61
24:RA:2318:C:N4	29:RG:42:GLY:O	2.33	0.61
1:XA:243:A:H4'	1:XA:244:U:O5'	2.01	0.61
1:QA:831:U:H3	1:QA:855:G:H1	1.49	0.61
3:QC:9:GLY:HA3	14:QN:49:HIS:HA	1.82	0.61
1:QA:544:G:OP1	4:QD:59:ARG:NH2	2.33	0.61
22:QV:36:G:H1	23:QX:16:C:H42	1.49	0.61
24:RA:597:C:O2	27:RE:145:LYS:NZ	2.34	0.60
24:RA:829:A:O2'	26:RD:225:ALA:O	2.19	0.60
1:XA:1305:G:O2'	1:XA:1332:A:N6	2.34	0.60
24:RA:892:G:O2'	24:RA:894:U:O4	2.19	0.60
53:Y8:6:THR:OG1	53:Y8:8:LYS:NZ	2.34	0.60
24:YA:1102:G:HO2'	24:YA:1132:A:H8	1.49	0.60
24:RA:441:C:O2	24:RA:1895:U:O2'	2.19	0.60
38:RT:135:ALA:HA	38:RT:137:LYS:HG2	1.83	0.60
1:XA:924:C:O2'	1:XA:1502:A:N6	2.34	0.60
2:XB:82:ARG:NH1	2:XB:92:TYR:OH	2.34	0.60
1:XA:619:U:N3	4:XD:134:ASP:OD1	2.31	0.60
13:XM:81:LEU:HD11	13:XM:88:ARG:HB3	1.83	0.60
24:YA:139:A:H8	24:YA:1454:C:HO2'	1.48	0.60
24:YA:2860:A:N7	24:YA:2878:A:O2'	2.27	0.60
7:QG:77:SER:HA	7:QG:85:TYR:O	2.02	0.60
16:QP:3:LYS:O	16:QP:21:VAL:HA	2.02	0.60

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:XM:3:ARG:HG3	13:XM:9:ILE:HD13	1.84	0.60
24:YA:1378:G:N2	24:YA:1655:A:O2'	2.34	0.60
4:QD:4:TYR:OH	4:QD:10:ARG:NH2	2.35	0.60
54:R9:25:VAL:HB	54:R9:34:GLN:HB2	1.82	0.60
24:RA:324:A:OP1	43:RY:86:ARG:NH2	2.34	0.60
24:RA:332:G:C8	24:RA:526:A:H1'	2.36	0.60
24:RA:1846:A:OP2	26:RD:54:ARG:NH2	2.34	0.60
28:RF:134:GLY:H	28:RF:162:LEU:HD22	1.67	0.60
30:RH:152:ARG:HG2	30:RH:153:LYS:HD3	1.82	0.60
1:XA:359:U:P	1:XA:359:U:H3'	2.41	0.60
1:XA:659:U:OP1	15:XO:8:LYS:NZ	2.34	0.60
1:QA:1399:C:N3	1:QA:1502:A:N6	2.49	0.60
3:QC:134:ILE:HG23	3:QC:151:VAL:HB	1.83	0.60
8:XH:51:VAL:HG21	8:XH:60:ARG:HG3	1.83	0.60
24:YA:2517:G:N7	50:Y5:3:LYS:NZ	2.48	0.60
24:YA:1736:A:H62	24:YA:1745:A:H2	1.50	0.60
1:QA:278:G:N7	17:QQ:92:ARG:NH1	2.50	0.60
46:R1:83:GLU:HG2	46:R1:85:LEU:H	1.66	0.60
24:RA:354:A:C2	24:RA:1255:A:H2'	2.32	0.60
2:XB:76:GLN:NE2	2:XB:206:ASP:OD1	2.34	0.60
24:YA:1991:A:O2'	24:YA:1994:A:N3	2.29	0.60
24:YA:238:C:O2	53:Y8:12:LYS:NZ	2.28	0.60
32:YN:58:ASP:N	32:YN:58:ASP:OD1	2.35	0.60
37:YS:11:LYS:HG3	37:YS:91:PRO:HD3	1.82	0.60
1:QA:1522:U:H2'	1:QA:1523:G:H8	1.67	0.60
1:QA:1528:U:HO2'	1:QA:1529:G:P	2.23	0.60
31:RI:82:ARG:HD3	31:RI:146:ALA:HB3	1.84	0.60
1:XA:539:A:H2'	1:XA:540:G:C8	2.37	0.60
21:XU:6:ARG:HD3	21:XU:15:ARG:HH12	1.66	0.60
54:Y9:27:CYS:SG	54:Y9:28:GLU:N	2.74	0.60
1:QA:1286:A:H2'	1:QA:1287:A:H4'	1.84	0.60
24:RA:1035:G:OP2	48:R3:11:SER:OG	2.16	0.60
29:RG:32:PRO:HB2	29:RG:172:LEU:HD13	1.83	0.60
33:RO:80:ASP:OD2	38:RT:64:ARG:NH2	2.35	0.60
34:RP:52:GLU:OE1	34:RP:55:ARG:NH1	2.34	0.60
1:XA:1238:A:N7	1:XA:1301:U:O4	2.35	0.60
1:QA:689:C:OP2	11:QK:55:LYS:NZ	2.34	0.60
1:QA:878:G:H5'	8:QH:89:PRO:HG2	1.84	0.60
2:QB:87:ARG:NH1	2:QB:220:ASP:OD1	2.34	0.60
3:QC:63:ASN:HA	3:QC:98:ASN:HB2	1.84	0.60
52:R7:10:ARG:HE	52:R7:14:LYS:HD2	1.67	0.60

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:RA:1072:U:H4'	24:RA:1073:A:OP1	2.02	0.60
25:RB:37:C:O2	37:RS:95:HIS:NE2	2.35	0.60
1:XA:279:A:OP1	1:XA:280:C:O2'	2.15	0.60
7:XG:60:LYS:HD3	7:XG:63:LYS:HB3	1.83	0.60
15:XO:82:ILE:HB	15:XO:87:ILE:HB	1.83	0.60
1:QA:244:U:OP2	17:QQ:100:LYS:NZ	2.34	0.59
34:YP:87:ASP:O	34:YP:90:ARG:NH1	2.35	0.59
24:YA:1209:G:OP1	40:YV:24:LYS:NZ	2.30	0.59
24:YA:1269:G:N2	24:YA:1272:A:OP2	2.29	0.59
1:QA:1414:U:H2'	1:QA:1415:G:H8	1.67	0.59
11:QK:51:LYS:HA	11:QK:55:LYS:HE2	1.84	0.59
11:QK:86:GLY:O	11:QK:91:ARG:NH1	2.35	0.59
22:QV:20:G:N3	22:QV:57:G:N2	2.51	0.59
24:RA:345:G:O2'	24:RA:364:A:N3	2.35	0.59
9:XI:3:GLN:OE1	9:XI:20:ARG:NH2	2.34	0.59
1:XA:1454:G:OP1	20:XT:39:LYS:NZ	2.35	0.59
1:QA:1047:G:OP1	14:QN:4:LYS:NZ	2.29	0.59
1:QA:1200:C:O2'	1:QA:1201:A:OP2	2.18	0.59
5:QE:100:VAL:O	5:QE:107:ARG:NH2	2.34	0.59
5:QE:79:GLU:O	8:QH:104:ARG:NH1	2.34	0.59
24:RA:865:G:N1	24:RA:1233:U:OP2	2.28	0.59
24:RA:140:A:H8	24:RA:1641:G:H21	1.50	0.59
24:RA:211:A:H3'	24:RA:448:U:H5'	1.84	0.59
2:XB:208:ILE:HA	2:XB:211:ILE:HD12	1.84	0.59
11:XK:18:ARG:NH2	11:XK:35:PRO:O	2.36	0.59
24:YA:2065:C:OP1	24:YA:2790:G:O2'	2.17	0.59
1:QA:1356:G:H2'	1:QA:1357:A:C8	2.38	0.59
1:QA:437:U:O2	4:QD:119:GLN:NE2	2.36	0.59
18:QR:33:ASP:OD2	18:QR:36:ASN:ND2	2.35	0.59
24:RA:232:U:OP2	53:R8:8:LYS:NZ	2.30	0.59
24:RA:2518:U:OP2	24:RA:2588:G:N1	2.31	0.59
24:RA:1058:U:OP1	39:RU:75:ASN:ND2	2.35	0.59
16:XP:18:ARG:NH1	16:XP:32:TYR:OH	2.35	0.59
51:Y6:3:SER:OG	51:Y6:4:GLU:N	2.35	0.59
24:YA:1001:G:OP2	35:YQ:14:ARG:NH2	2.35	0.59
44:YZ:45:ASP:OD1	44:YZ:49:ARG:NH1	2.36	0.59
22:QV:58:A:O2'	22:QV:60:U:OP2	2.14	0.59
24:RA:2340:A:H2'	24:RA:2341:G:C8	2.37	0.59
26:RD:2:ALA:N	26:RD:200:ASP:OD2	2.36	0.59
1:XA:1459:C:OP1	20:XT:27:LYS:NZ	2.35	0.59
24:YA:1405:A:OP2	24:YA:1417:G:N1	2.30	0.59

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:YA:388:A:H2'	24:YA:389:G:H8	1.67	0.59
1:QA:642:A:N3	8:QH:113:SER:OG	2.34	0.59
24:RA:2297:C:OP2	51:R6:6:ARG:NH2	2.31	0.59
1:XA:1157:A:N7	1:XA:1178:G:N2	2.50	0.59
51:Y6:3:SER:HB3	51:Y6:6:ARG:HB3	1.85	0.59
24:YA:1961:U:OP1	24:YA:2616:U:O2'	2.20	0.59
48:R3:6:VAL:HG12	48:R3:56:VAL:HG12	1.83	0.59
49:R4:16:CYS:SG	49:R4:17:GLY:N	2.76	0.59
24:RA:2297:C:H5	51:R6:6:ARG:HH12	1.50	0.59
31:RI:79:ILE:HB	31:RI:142:VAL:HG23	1.84	0.59
24:YA:616:G:H4'	53:Y8:61:LEU:HD13	1.84	0.59
28:YF:185:ASP:OD1	28:YF:188:ARG:NH1	2.35	0.59
1:QA:346:G:H1'	1:QA:347:G:H5'	1.85	0.59
1:QA:689:C:H3'	1:QA:690:G:H21	1.68	0.59
2:QB:84:GLU:OE1	2:QB:87:ARG:NH2	2.36	0.59
46:R1:65:SER:HG	46:R1:66:HIS:HD1	1.48	0.59
50:R5:16:ARG:NH1	50:R5:17:ASP:OD1	2.35	0.59
24:RA:592:U:O2'	24:RA:1029:A:N1	2.32	0.59
24:RA:238:C:OP2	24:RA:2406:C:O2'	2.20	0.59
2:XB:30:ARG:NH2	2:XB:195:ASP:OD1	2.36	0.59
2:XB:185:ILE:HG22	2:XB:199:TYR:HB2	1.85	0.59
24:YA:2801:C:O2'	24:YA:2819:A:N3	2.35	0.59
1:QA:677:U:O2	1:QA:777:A:O2'	2.21	0.59
24:RA:1289:G:O2'	34:RP:7:ARG:NH2	2.34	0.59
26:RD:17:THR:HB	26:RD:205:VAL:H	1.68	0.59
24:RA:24:G:O2'	41:RW:78:GLU:O	2.20	0.59
1:XA:701:C:OP1	1:XA:702:A:O2'	2.15	0.59
1:XA:662:G:O2'	1:XA:836:G:OP1	2.20	0.59
11:XK:33:THR:HA	11:XK:39:PRO:HA	1.85	0.59
12:XL:114:LYS:O	12:XL:117:ARG:NH1	2.36	0.59
25:YB:93:C:OP1	44:YZ:80:ARG:NH2	2.36	0.59
17:QQ:90:ILE:O	17:QQ:94:ASN:ND2	2.36	0.58
20:XT:100:ILE:HG23	20:XT:102:GLY:H	1.68	0.58
37:YS:81:GLY:O	37:YS:83:LYS:NZ	2.35	0.58
1:QA:811:C:O2'	1:QA:901:A:N1	2.37	0.58
38:YT:20:PRO:HD2	38:YT:86:ILE:HG23	1.83	0.58
24:RA:66:U:H3	24:RA:73:A:H2	1.49	0.58
30:RH:101:ARG:NH2	30:RH:121:ILE:O	2.36	0.58
3:XC:93:LYS:HE2	3:XC:94:LEU:HD23	1.86	0.58
36:YR:75:LEU:HA	36:YR:78:LYS:HB3	1.83	0.58
1:QA:1441:G:N2	1:QA:1460:A:H62	2.01	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:XD:105:VAL:HG13	4:XD:110:PHE:HB2	1.84	0.58
1:XA:191(G):G:O2'	20:XT:101:GLY:O	2.20	0.58
24:YA:2622:C:H4'	24:YA:2623:U:OP2	2.03	0.58
33:YO:107:ARG:NH1	38:YT:36:GLU:OE1	2.37	0.58
1:QA:684:A:O2'	11:QK:39:PRO:O	2.20	0.58
24:RA:1847:G:O6	26:RD:35:LYS:NZ	2.32	0.58
24:RA:2858:G:O2'	24:RA:2877:G:N2	2.37	0.58
30:RH:84:SER:OG	30:RH:85:LYS:N	2.37	0.58
1:XA:28:G:O2'	1:XA:296:U:OP1	2.19	0.58
24:YA:82:G:H1	24:YA:100:G:HO2'	1.49	0.58
38:YT:124:ASP:N	38:YT:124:ASP:OD1	2.36	0.58
38:YT:16:ARG:NH2	38:YT:83:ILE:O	2.36	0.58
1:QA:410:G:N2	1:QA:432:A:H62	2.00	0.58
1:QA:1280:A:H5''	10:QJ:40:LEU:HD23	1.84	0.58
24:RA:1065:U:H3	24:RA:1188:A:H62	1.48	0.58
7:XG:28:ASN:OD1	7:XG:36:LYS:NZ	2.34	0.58
24:YA:2414:C:H1'	24:YA:2415:C:H5	1.67	0.58
24:YA:2639:G:N2	24:YA:2790:G:OP1	2.35	0.58
28:YF:198:ALA:HA	28:YF:201:VAL:HG12	1.84	0.58
1:QA:1376:U:OP1	7:QG:94:ARG:NH1	2.37	0.58
1:QA:559:A:H4'	1:QA:560:U:H3'	1.86	0.58
24:RA:1109:G:OP1	24:RA:1111:U:O2'	2.18	0.58
50:Y5:16:ARG:NH1	50:Y5:17:ASP:OD1	2.35	0.58
24:YA:1155:C:O2'	24:YA:1156:G:OP1	2.19	0.58
24:YA:2132:G:O2'	24:YA:2133:C:O4'	2.20	0.58
25:YB:37:C:O2	37:YS:95:HIS:NE2	2.35	0.58
36:RR:56:LYS:NZ	36:RR:90:ARG:O	2.37	0.58
1:XA:62:U:H3	1:XA:105:G:H1	1.51	0.58
24:YA:115:G:OP2	24:YA:117:A:O2'	2.20	0.58
18:QR:70:ILE:O	18:QR:74:ARG:HB2	2.04	0.58
24:RA:2367:C:H1'	45:R0:39:ARG:HH21	1.69	0.58
24:RA:2453:C:OP2	24:RA:2598:C:O2'	2.21	0.58
1:XA:1450:U:O2'	1:XA:1451:A:N7	2.36	0.58
1:XA:664:G:H22	1:XA:741:G:H1	1.51	0.58
35:YQ:122:GLY:HA2	35:YQ:125:LEU:HD12	1.86	0.58
1:QA:946:A:H2'	1:QA:947:G:C8	2.39	0.58
24:RA:1481:G:N2	24:RA:1524:A:O2'	2.36	0.58
1:XA:1347:G:N2	1:XA:1374:A:OP2	2.36	0.58
4:XD:209:ARG:NH1	4:XD:209:ARG:O	2.37	0.58
24:YA:2211:U:H2'	24:YA:2212:G:C8	2.39	0.58
1:QA:1095:U:OP1	1:QA:1108:G:N2	2.27	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:765:G:N2	1:QA:813:U:OP2	2.30	0.57
13:XM:49:THR:HG22	13:XM:51:ALA:H	1.69	0.57
24:YA:662:A:OP1	34:YP:133:SER:OG	2.18	0.57
35:YQ:31:ASP:OD1	35:YQ:134:ARG:NH1	2.33	0.57
38:YT:124:ASP:O	38:YT:128:GLU:N	2.36	0.57
1:QA:973:G:O6	1:QA:974:A:N6	2.37	0.57
3:QC:184:TYR:HA	3:QC:200:ALA:O	2.03	0.57
19:QS:36:ARG:NH2	19:QS:72:GLY:O	2.38	0.57
24:RA:208:G:N3	24:RA:223:C:O2'	2.36	0.57
24:RA:241:G:OP2	34:RP:50:ARG:NH2	2.29	0.57
24:RA:2533:C:O2'	24:RA:2576:A:N3	2.34	0.57
44:YZ:119:GLU:OE2	44:YZ:122:ARG:NH1	2.37	0.57
24:RA:2148:A:H4'	24:RA:2149:G:O5'	2.05	0.57
5:XE:102:ALA:HB1	5:XE:106:PRO:HG2	1.85	0.57
24:YA:1176:U:O2'	24:YA:1177:G:O5'	2.19	0.57
24:YA:2318:C:N4	29:YG:42:GLY:O	2.37	0.57
1:QA:742:G:OP2	15:QO:35:ARG:NH2	2.35	0.57
1:QA:128:G:O2'	17:QQ:3:LYS:NZ	2.37	0.57
24:RA:2339:A:H2'	24:RA:2340:A:C8	2.39	0.57
24:RA:2404:A:H2	24:RA:2436:C:H42	1.50	0.57
1:XA:152:A:N6	1:XA:169:C:C4	2.73	0.57
29:YG:74:LYS:O	29:YG:84:LYS:NZ	2.37	0.57
32:YN:25:ARG:O	32:YN:29:LYS:NZ	2.37	0.57
5:QE:153:LYS:HE2	5:QE:155:GLU:HB3	1.86	0.57
24:RA:1132:A:O2'	24:RA:1149:A:N1	2.36	0.57
24:RA:2860:A:N7	24:RA:2878:A:O2'	2.33	0.57
30:RH:106:THR:HG22	30:RH:112:PRO:HB3	1.86	0.57
1:XA:624:C:H4'	16:XP:10:GLY:HA2	1.87	0.57
24:YA:2859:U:OP2	38:YT:95:ARG:NH1	2.38	0.57
44:YZ:19:ARG:NH1	44:YZ:84:GLU:O	2.37	0.57
24:RA:2054:G:H1'	27:RE:145:LYS:HD2	1.86	0.57
24:RA:2798:C:OP1	27:RE:41:LYS:NZ	2.34	0.57
39:RU:90:VAL:HG22	40:RV:38:LEU:HB3	1.86	0.57
1:XA:316:G:OP2	1:XA:351:G:O2'	2.15	0.57
8:XH:11:THR:HG22	8:XH:14:ARG:HH12	1.69	0.57
1:QA:411:A:H62	1:QA:413:G:H21	1.51	0.57
5:QE:153:LYS:HG3	5:QE:155:GLU:H	1.68	0.57
4:XD:114:ARG:O	4:XD:118:ARG:N	2.36	0.57
24:YA:1052:C:O2'	32:YN:106:MET:O	2.23	0.57
24:RA:2270:C:O2'	24:RA:2439:C:OP2	2.22	0.57
24:RA:907:U:H2'	24:RA:908:A:H8	1.70	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:99:C:H2'	1:XA:101:A:C8	2.40	0.57
34:YP:62:LEU:O	53:Y8:13:ARG:NH1	2.38	0.57
33:YO:76:ALA:HB3	38:YT:75:ILE:HD12	1.87	0.57
2:QB:92:TYR:HH	2:QB:150:SER:HG	1.50	0.57
8:QH:7:ALA:HB2	8:QH:85:ARG:HD3	1.86	0.57
24:RA:331:G:H21	24:RA:354:A:H62	1.52	0.57
24:YA:1627:A:H3'	24:YA:1628:G:H8	1.70	0.57
24:YA:419:C:H5''	24:YA:436:C:H5''	1.86	0.57
24:YA:484:G:N2	24:YA:496:A:OP2	2.25	0.57
28:YF:116:ASP:OD1	28:YF:119:ARG:NH2	2.38	0.57
1:QA:1422:G:O3'	33:RO:49:ARG:NH1	2.37	0.57
8:QH:106:GLY:O	8:QH:122:ARG:NH2	2.37	0.57
27:RE:176:ILE:HB	27:RE:181:LEU:HB2	1.87	0.57
7:XG:69:VAL:HG13	7:XG:100:ALA:HB1	1.87	0.57
45:Y0:27:GLU:HG3	45:Y0:68:GLU:HA	1.85	0.57
24:YA:2120:U:O4	24:YA:2213:G:O6	2.23	0.57
1:QA:1028(C):C:N4	1:QA:1032(B):G:O6	2.38	0.56
24:RA:210:A:H4'	24:RA:211:A:O5'	2.04	0.56
27:RE:128:SER:OG	27:RE:129:HIS:N	2.37	0.56
9:XI:48:GLU:O	9:XI:51:ARG:N	2.37	0.56
29:YG:67:LYS:HD3	49:Y4:5:ILE:HD12	1.85	0.56
25:YB:48:A:H4'	37:YS:95:HIS:HD2	1.70	0.56
1:QA:132:C:O3'	20:QT:74:LYS:NZ	2.38	0.56
6:QF:100:ASN:ND2	18:QR:23:LYS:O	2.33	0.56
24:RA:1588:G:O6	24:RA:1589:A:N6	2.38	0.56
24:RA:484:G:N2	24:RA:496:A:OP2	2.36	0.56
30:RH:86:GLU:HB3	30:RH:165:ALA:H	1.70	0.56
24:YA:1501:U:O2'	24:YA:1502:G:N7	2.38	0.56
24:YA:2318:C:H3'	24:YA:2319:G:H5''	1.87	0.56
24:YA:895:G:H2'	24:YA:896:A:C8	2.39	0.56
24:YA:964:A:N3	25:YB:80:U:O2'	2.36	0.56
1:QA:1287:A:H2	1:QA:1353:G:H1'	1.69	0.56
13:QM:47:ASP:N	13:QM:47:ASP:OD1	2.38	0.56
24:RA:2826:C:O2	24:RA:2893:A:O2'	2.23	0.56
1:XA:1356:G:H2'	1:XA:1357:A:C8	2.36	0.56
1:XA:24:U:HO2'	1:XA:524:G:HO2'	1.53	0.56
23:XX:6:G:H2'	23:XX:7:G:C8	2.40	0.56
24:YA:238:C:OP2	24:YA:2406:C:O2'	2.20	0.56
24:YA:2479:C:OP1	54:Y9:6:SER:OG	2.20	0.56
2:QB:61:LEU:O	2:QB:65:GLY:N	2.36	0.56
12:QL:49:ASN:ND2	12:QL:92:ASP:OD2	2.38	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:QM:15:VAL:HG12	13:QM:45:VAL:HB	1.87	0.56
24:RA:1014:U:H2'	24:RA:1015:C:C6	2.41	0.56
24:RA:1291:G:OP1	34:RP:13:ASN:ND2	2.33	0.56
24:RA:2901:A:H2'	24:RA:2902:G:O4'	2.06	0.56
27:RE:47:VAL:HG21	27:RE:86:PRO:HD2	1.86	0.56
44:RZ:108:PRO:HG3	44:RZ:141:VAL:HG22	1.88	0.56
1:XA:1236:A:H4'	1:XA:1304:G:H4'	1.87	0.56
24:YA:1108:G:H2'	24:YA:1109:G:H8	1.69	0.56
24:YA:331:G:H21	24:YA:354:A:H62	1.53	0.56
1:QA:1386:G:H2'	1:QA:1387:G:H8	1.71	0.56
12:QL:60:LEU:HD12	12:QL:62:SER:H	1.71	0.56
48:R3:40:THR:HB	48:R3:43:ILE:HG12	1.86	0.56
1:XA:243:A:H4'	1:XA:244:U:H3'	1.87	0.56
19:XS:3:ARG:NH1	19:XS:8:GLY:O	2.37	0.56
24:YA:484:G:O2'	24:YA:495:G:O6	2.20	0.56
1:QA:1079:G:O2'	5:QE:14:ARG:NH2	2.38	0.56
1:QA:1147:C:O2	9:QI:16:ARG:NH2	2.39	0.56
17:QQ:7:THR:HG22	17:QQ:58:GLU:HG2	1.87	0.56
1:XA:606:G:H22	1:XA:631:G:H5''	1.71	0.56
2:XB:135:GLN:HB3	2:XB:139:LYS:HE3	1.87	0.56
2:XB:95:GLN:HA	2:XB:96:ARG:HH21	1.70	0.56
3:XC:150:LYS:HD3	3:XC:152:ILE:HD11	1.87	0.56
24:YA:1449:C:H5''	24:YA:1518:A:H1'	1.87	0.56
24:YA:1896:G:H5'	24:YA:1897:C:OP2	2.05	0.56
29:YG:161:THR:HG22	29:YG:163:ALA:H	1.70	0.56
35:YQ:32:TYR:OH	35:YQ:111:GLU:OE1	2.24	0.56
1:QA:714:G:H2'	1:QA:715:A:C8	2.39	0.56
17:QQ:83:ASP:OD1	17:QQ:83:ASP:N	2.38	0.56
35:RQ:134:ARG:HH22	44:RZ:122:ARG:HH21	1.54	0.56
7:XG:5:ARG:HH21	7:XG:6:ARG:HB2	1.69	0.56
24:YA:1033:G:O2'	24:YA:1046:A:N3	2.36	0.56
24:YA:2052:A:H4'	24:YA:2053:A:H8	1.70	0.56
24:YA:441:C:O2	24:YA:1895:U:O2'	2.22	0.56
1:QA:243:A:H4'	1:QA:244:U:O5'	2.04	0.56
1:QA:409:G:OP1	4:QD:24:GLU:N	2.35	0.56
24:RA:616:G:H4'	53:R8:61:LEU:HD13	1.87	0.56
24:RA:2559:U:O2	33:RO:23:ARG:NH2	2.38	0.56
24:RA:137:G:N2	42:RX:44:GLU:OE1	2.35	0.56
1:XA:881:G:OP2	12:XL:12:ARG:NH2	2.39	0.56
24:YA:1604:C:OP2	24:YA:1605:A:O2'	2.19	0.56
24:YA:2203:G:H2'	24:YA:2204:G:C8	2.40	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:410:G:H3'	4:QD:25:ARG:HH12	1.70	0.56
1:QA:547:A:OP1	4:QD:73:ARG:NH2	2.39	0.56
1:QA:56:U:H2'	1:QA:57:G:H8	1.70	0.56
3:QC:19:GLU:O	3:QC:40:ARG:NH2	2.39	0.56
4:QD:47:ARG:HB3	4:QD:49:ARG:HH21	1.71	0.56
9:QI:13:ALA:HB2	9:QI:68:GLY:HA3	1.87	0.56
24:RA:1087:C:H42	24:RA:1160:G:H1	1.54	0.56
24:RA:2073:A:H5'	24:RA:2590:G:O4'	2.05	0.56
24:RA:2877:G:HO2'	24:RA:2878:A:H8	1.53	0.56
33:RO:71:ARG:HB3	33:RO:73:ASP:OD1	2.05	0.56
13:XM:60:VAL:HG22	13:XM:66:LEU:HD11	1.86	0.56
28:YF:122:LYS:NZ	28:YF:152:GLU:OE2	2.39	0.56
1:QA:359:U:H2'	1:QA:360:A:H8	1.70	0.56
3:QC:88:ARG:HA	3:QC:91:LEU:HD12	1.88	0.56
9:QI:21:PRO:HA	9:QI:59:PHE:HA	1.87	0.56
24:RA:1714:G:H8	24:RA:1714:G:OP2	1.88	0.56
24:RA:1736:A:H62	24:RA:1745:A:H2	1.54	0.56
24:RA:2760:G:OP1	30:RH:138:LYS:NZ	2.33	0.56
26:RD:153:ALA:O	26:RD:157:ARG:NH1	2.39	0.56
26:RD:146:GLU:HB2	26:RD:189:CYS:HB3	1.86	0.56
24:RA:98:U:O4	43:RY:8:LYS:NZ	2.39	0.56
24:YA:1927:C:H3'	24:YA:1952:G:C8	2.40	0.56
24:YA:2669:A:O2'	30:YH:160:LYS:NZ	2.39	0.56
24:YA:904:C:OP2	45:Y0:77:ARG:NH2	2.39	0.56
10:QJ:28:ARG:NH1	10:QJ:34:VAL:O	2.36	0.56
45:R0:70:GLN:OE1	45:R0:80:HIS:NE2	2.38	0.56
24:RA:1001:G:OP2	35:RQ:14:ARG:NH2	2.39	0.56
24:RA:1068:G:N2	24:RA:1069:U:O4	2.38	0.56
25:RB:27:C:H4'	37:RS:54:LEU:HD11	1.87	0.56
33:RO:68:GLU:HG3	33:RO:78:ARG:HD3	1.88	0.56
38:RT:16:ARG:HH21	38:RT:19:LEU:HD21	1.71	0.56
44:RZ:45:ASP:OD1	44:RZ:49:ARG:NE	2.39	0.56
1:XA:1291:G:H4'	9:XI:39:GLY:HA3	1.89	0.56
1:XA:359:U:H3'	1:XA:359:U:OP2	2.06	0.56
2:XB:19:HIS:HB3	2:XB:20:GLU:HG2	1.87	0.56
1:XA:875:C:O2'	8:XH:14:ARG:NH1	2.39	0.56
9:XI:9:ARG:HB3	9:XI:104:ARG:HH21	1.70	0.56
24:YA:537:G:OP1	24:YA:1280:U:O2'	2.13	0.56
26:YD:108:PRO:HG2	26:YD:111:LEU:HG	1.88	0.56
26:YD:72:LYS:HG3	26:YD:103:ARG:HH21	1.71	0.56
1:QA:1123:A:H4'	10:QJ:36:GLY:HA3	1.88	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:RA:2136:A:C2	24:RA:2139:A:H5''	2.40	0.55
1:XA:1316:G:N2	1:XA:1319:A:OP2	2.38	0.55
1:XA:148:G:H2'	1:XA:149:A:C8	2.41	0.55
1:XA:547:A:OP1	4:XD:73:ARG:NH2	2.39	0.55
1:QA:1373:G:H5'	7:QG:36:LYS:HG2	1.88	0.55
24:RA:1117:G:N2	24:RA:1135:G:O6	2.38	0.55
24:RA:1449:C:H5''	24:RA:1518:A:H1'	1.86	0.55
24:RA:2818:U:C2	24:RA:2901:A:C6	2.92	0.55
2:XB:136:VAL:HA	2:XB:139:LYS:HD2	1.88	0.55
1:XA:642:A:N3	8:XH:113:SER:OG	2.39	0.55
45:Y0:20:ARG:O	45:Y0:24:LYS:NZ	2.37	0.55
3:XC:35:GLU:OE1	3:XC:59:ARG:NH2	2.37	0.55
24:YA:1686:U:H2'	24:YA:1687:C:H5''	1.89	0.55
24:YA:1830:G:H8	26:YD:181:GLU:OE1	1.88	0.55
24:RA:2377:G:N7	53:R8:39:LYS:NZ	2.55	0.55
43:RY:47:LYS:NZ	43:RY:48:ALA:O	2.35	0.55
9:XI:13:ALA:HB2	9:XI:68:GLY:HA3	1.89	0.55
24:YA:1289:G:O2'	34:YP:7:ARG:NH2	2.40	0.55
24:YA:360:C:HO2'	43:YY:35:TYR:HH	1.52	0.55
1:QA:1240:U:O4	7:QG:109:ASN:ND2	2.39	0.55
20:QT:89:ARG:NH2	20:QT:105:SER:O	2.38	0.55
38:RT:16:ARG:NH2	38:RT:18:ASP:OD2	2.40	0.55
1:XA:624:C:H2'	1:XA:625:G:H8	1.71	0.55
24:YA:85:C:H4'	24:YA:102:U:H1'	1.87	0.55
24:YA:2034:G:OP1	41:YW:11:ARG:NH2	2.38	0.55
1:QA:346:G:OP1	38:RT:41:ARG:NH1	2.33	0.55
5:QE:81:GLU:OE1	8:QH:104:ARG:NH2	2.38	0.55
1:XA:1251:A:N3	1:XA:1369:C:O2'	2.32	0.55
1:XA:973:G:H3'	1:XA:974:A:H5''	1.88	0.55
2:XB:11:LEU:O	2:XB:16:HIS:ND1	2.39	0.55
1:XA:132:C:O3'	20:XT:74:LYS:NZ	2.39	0.55
24:YA:596:G:N1	24:YA:2053:A:OP2	2.25	0.55
1:QA:1009:G:O6	1:QA:1020:U:O4	2.25	0.55
1:QA:439:A:OP2	1:QA:493:G:N1	2.24	0.55
2:QB:63:MET:HG3	2:QB:225:ALA:HB1	1.88	0.55
47:R2:20:GLU:HA	47:R2:23:LYS:HG3	1.88	0.55
24:RA:1132:A:OP1	24:RA:1150:C:O2'	2.23	0.55
24:RA:184:A:OP1	34:RP:46:LYS:NZ	2.31	0.55
33:RO:13:ASN:HD21	33:RO:96:THR:HG1	1.55	0.55
44:RZ:132:ASN:ND2	44:RZ:159:PRO:O	2.39	0.55
1:XA:1071:C:H2'	1:XA:1072:G:H8	1.72	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:XB:32:ILE:HD11	2:XB:40:HIS:HB3	1.89	0.55
24:YA:278:G:H1'	46:Y1:78:LYS:HD3	1.89	0.55
1:QA:150:C:H42	1:QA:171:A:H62	1.54	0.55
1:QA:321:A:N6	1:QA:329:A:OP2	2.38	0.55
1:QA:1316:G:H4'	14:QN:18:VAL:HG11	1.88	0.55
10:QJ:61:GLU:OE2	14:QN:45:ARG:NE	2.40	0.55
24:RA:1003:U:OP2	35:RQ:14:ARG:NH1	2.40	0.55
9:XI:22:GLY:HA3	9:XI:60:ASP:HB2	1.87	0.55
1:QA:782:A:H62	1:QA:800:G:H21	1.54	0.55
24:RA:1039:G:OP1	39:RU:50:ARG:NH2	2.38	0.55
24:YA:2299:A:H62	24:YA:2356:U:H3	1.54	0.55
24:YA:2416:C:O3'	34:YP:77:ARG:NH2	2.40	0.55
42:YX:27:THR:HG23	42:YX:80:ILE:HG13	1.88	0.55
1:QA:1305:G:N2	1:QA:1332:A:OP2	2.38	0.55
1:QA:359:U:H2'	1:QA:360:A:C8	2.41	0.55
2:QB:9:GLU:HA	2:QB:12:GLU:HB2	1.88	0.55
12:QL:114:LYS:O	12:QL:117:ARG:NH1	2.39	0.55
22:QV:34:C:H2'	22:QV:35:G:H8	1.72	0.55
24:RA:1047:A:H62	24:RA:1200:G:H21	1.53	0.55
24:RA:2013:U:H2'	24:RA:2014:G:H5''	1.89	0.55
24:RA:251:A:N3	24:RA:457:G:O2'	2.31	0.55
1:XA:422:C:HO2'	1:XA:423:G:N2	2.05	0.55
1:QA:1328:C:OP1	21:QU:21:TYR:OH	2.22	0.54
1:XA:352:C:O2'	1:XA:354:G:OP1	2.15	0.54
3:XC:17:ASP:O	3:XC:54:ARG:NH2	2.40	0.54
24:YA:2885:C:O2'	38:YT:2:ASN:OD1	2.24	0.54
30:YH:51:ARG:HB3	30:YH:51:ARG:HH11	1.72	0.54
6:QF:3:ARG:HA	6:QF:65:VAL:O	2.08	0.54
8:QH:121:ASP:OD1	8:QH:121:ASP:N	2.37	0.54
20:QT:14:LYS:HA	20:QT:17:ARG:HE	1.72	0.54
30:RH:56:SER:OG	30:RH:57:ASP:N	2.38	0.54
31:RI:78:THR:HG22	31:RI:141:LYS:HE3	1.88	0.54
4:XD:187:ARG:NH1	4:XD:188:LEU:O	2.40	0.54
22:XV:2:G:H2'	22:XV:3:G:H8	1.72	0.54
53:Y8:22:VAL:HB	53:Y8:53:PRO:HB3	1.90	0.54
1:QA:1172:C:H2'	1:QA:1173:G:H8	1.72	0.54
2:QB:6:THR:O	2:QB:217:ARG:NH1	2.38	0.54
3:QC:8:ILE:HG23	3:QC:16:ARG:HG2	1.89	0.54
10:QJ:48:THR:HG22	10:QJ:62:HIS:HB3	1.89	0.54
14:QN:24:CYS:SG	14:QN:25:VAL:N	2.79	0.54
1:QA:1304:G:OP1	21:QU:2:GLY:N	2.39	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:RQ:28:ALA:N	35:RQ:105:GLU:OE2	2.39	0.54
1:XA:1129:C:N4	1:XA:1134:G:O6	2.40	0.54
1:XA:1406:U:O2	1:XA:1517:G:N2	2.36	0.54
1:XA:129(B):G:N2	1:XA:188:U:O2'	2.41	0.54
2:QB:20:GLU:OE1	2:QB:23:ARG:NH1	2.40	0.54
29:RG:170:ARG:NH2	29:RG:174:GLU:OE2	2.38	0.54
33:RO:2:ILE:HB	33:RO:33:ALA:HB3	1.89	0.54
1:QA:1510:U:H2'	1:QA:1511:G:C8	2.42	0.54
3:QC:128:PHE:HB3	3:QC:132:ARG:HH21	1.72	0.54
24:RA:1854:G:OP1	26:RD:54:ARG:NH1	2.40	0.54
24:RA:606:G:OP2	39:RU:10:ARG:NH1	2.40	0.54
24:RA:655:G:OP1	53:R8:46:ARG:NH1	2.40	0.54
1:XA:1336:C:O2	1:XA:1337:G:N1	2.41	0.54
8:XH:49:GLU:OE2	8:XH:62:TYR:OH	2.24	0.54
1:XA:790:A:OP1	22:XV:38:A:O2'	2.24	0.54
24:YA:600:G:O2'	24:YA:1300:A:OP1	2.26	0.54
24:YA:485:U:H2'	24:YA:486:A:H8	1.71	0.54
27:YE:105:THR:OG1	27:YE:199:ARG:NH2	2.40	0.54
1:QA:701:C:H1'	1:QA:703:G:C6	2.43	0.54
1:QA:701:C:OP1	1:QA:702:A:O2'	2.20	0.54
24:RA:953:U:HO2'	35:RQ:101:ARG:HH22	1.53	0.54
1:XA:76:G:O6	1:XA:93:U:O4	2.25	0.54
13:XM:93:ARG:HG3	13:XM:94:ARG:HE	1.73	0.54
24:YA:1361:C:O2'	24:YA:1438:A:N3	2.39	0.54
24:YA:831:A:OP2	24:YA:2601:A:OP1	2.26	0.54
24:YA:721:G:H1'	28:YF:74:ARG:HD3	1.90	0.54
1:QA:362:G:N2	1:QA:365:U:OP2	2.40	0.54
17:QQ:66:SER:OG	17:QQ:67:LYS:O	2.25	0.54
50:R5:41:PRO:O	50:R5:44:THR:OG1	2.24	0.54
51:R6:3:SER:OG	51:R6:4:GLU:N	2.41	0.54
24:RA:1884:A:N3	24:RA:2245:U:O2'	2.33	0.54
25:RB:7:G:N3	37:RS:38:GLN:NE2	2.48	0.54
29:RG:60:LEU:HA	29:RG:63:ILE:HG12	1.90	0.54
1:XA:677:U:H3	1:XA:713:G:H22	1.55	0.54
4:XD:99:SER:OG	4:XD:140:VAL:O	2.26	0.54
11:XK:34:ASP:OD1	11:XK:38:ASN:N	2.40	0.54
24:YA:209:G:O2'	24:YA:222:A:N3	2.36	0.54
24:YA:2347:A:O2'	24:YA:2348:A:H2'	2.07	0.54
24:YA:489:G:N2	24:YA:492:A:OP2	2.36	0.54
38:YT:64:ARG:NH1	38:YT:106:SER:OG	2.41	0.54
1:QA:1150:U:H4'	10:QJ:41:PRO:HB3	1.89	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:RA:1105:G:OP2	24:RA:1106:U:H3'	2.08	0.54
24:RA:2099:A:OP1	24:RA:2250:G:N2	2.33	0.54
1:XA:129(B):G:H1'	1:XA:190:G:H5''	1.90	0.54
1:XA:324:G:N2	1:XA:327:A:OP2	2.41	0.54
1:XA:957:U:N3	1:XA:960:U:OP2	2.41	0.54
24:YA:626:A:H5''	24:YA:702:A:H61	1.72	0.54
1:XA:143:A:H5''	1:XA:144:G:H5'	1.90	0.54
1:XA:8:A:N6	4:XD:205:GLU:O	2.41	0.54
29:YG:11:TYR:OH	29:YG:32:PRO:O	2.22	0.54
30:YH:23:ARG:NH1	30:YH:34:GLU:OE1	2.36	0.54
24:RA:2575:U:N3	24:RA:2578:A:OP2	2.28	0.54
24:RA:2710:U:H2'	24:RA:2711:C:C6	2.42	0.54
24:RA:334:A:C4	24:RA:336:G:C8	2.96	0.54
24:RA:73:A:H4'	24:RA:74:G:O5'	2.07	0.54
28:RF:185:ASP:HA	28:RF:188:ARG:HD3	1.89	0.54
4:XD:20:TYR:CD1	4:XD:20:TYR:N	2.73	0.54
24:YA:1312:G:O2'	24:YA:2034:G:O6	2.19	0.54
24:YA:597:C:N3	27:YE:145:LYS:NZ	2.50	0.54
1:QA:1026:G:O6	1:QA:1036:G:N2	2.41	0.53
1:QA:1224:G:H1	1:QA:1362(B):C:H42	1.56	0.53
24:RA:2159:C:H2'	24:RA:2160:C:C6	2.43	0.53
24:RA:873:U:OP1	24:RA:2441:G:OP1	2.27	0.53
37:RS:25:ARG:NH1	37:RS:42:ASP:OD2	2.41	0.53
1:QA:345:C:OP2	38:RT:39:ARG:NH2	2.42	0.53
1:XA:148:G:H2'	1:XA:149:A:H8	1.73	0.53
17:XQ:66:SER:OG	17:XQ:67:LYS:O	2.23	0.53
54:Y9:25:VAL:HB	54:Y9:34:GLN:HB2	1.91	0.53
24:YA:1088:G:H1	24:YA:1159:U:H3	1.55	0.53
24:YA:815:G:N2	24:YA:1425:A:O2'	2.41	0.53
25:YB:109:G:C8	25:YB:109:G:C5'	2.85	0.53
1:QA:1002:G:H2'	1:QA:1003:G:H8	1.71	0.53
1:QA:1228:C:OP1	13:QM:115:LYS:N	2.38	0.53
24:RA:1110:C:H3'	24:RA:1111:U:C5'	2.38	0.53
24:RA:2308:U:OP2	37:RS:9:ARG:NH1	2.42	0.53
32:RN:58:ASP:N	32:RN:58:ASP:OD1	2.33	0.53
24:RA:1039:G:N2	40:RV:23:GLU:OE2	2.41	0.53
1:XA:1502:A:H2	1:XA:1505:G:H1	1.56	0.53
1:XA:409:G:OP1	4:XD:24:GLU:N	2.41	0.53
1:XA:514:C:H2'	1:XA:515:G:H8	1.72	0.53
11:XK:17:GLY:HA2	11:XK:35:PRO:HD3	1.89	0.53
13:XM:47:ASP:N	13:XM:47:ASP:OD1	2.38	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:YA:236:G:OP2	24:YA:238:C:N4	2.41	0.53
29:YG:38:VAL:HG22	29:YG:93:THR:HG23	1.89	0.53
5:QE:140:ARG:O	5:QE:143:ARG:NH1	2.40	0.53
5:QE:5:ASP:OD1	5:QE:5:ASP:N	2.40	0.53
9:QI:97:LYS:HG3	9:QI:98:PRO:HD3	1.90	0.53
1:QA:689:C:OP1	11:QK:44:SER:OG	2.27	0.53
24:RA:82:G:O2'	24:RA:100:G:N2	2.41	0.53
24:RA:1911:A:N3	24:RA:2108:U:O2'	2.40	0.53
24:RA:612:C:H2'	24:RA:613:A:C8	2.44	0.53
24:RA:831:A:OP2	24:RA:2601:A:OP1	2.27	0.53
31:RI:4:ILE:HD11	31:RI:44:LEU:HD12	1.90	0.53
1:XA:1095:U:OP2	1:XA:1108:G:N1	2.37	0.53
2:XB:85:ALA:O	2:XB:89:GLY:N	2.41	0.53
13:XM:15:VAL:HG12	13:XM:45:VAL:HB	1.89	0.53
46:Y1:85:LEU:HB3	46:Y1:88:LYS:HD2	1.90	0.53
47:Y2:23:LYS:NZ	47:Y2:27:GLU:OE2	2.41	0.53
24:YA:1822:A:N6	24:YA:1859:G:O2'	2.40	0.53
24:YA:1501:U:H5'	36:YR:63:ARG:NH2	2.23	0.53
1:QA:745:C:H5''	1:QA:851:G:H1'	1.90	0.53
7:QG:60:LYS:HA	7:QG:63:LYS:HG2	1.91	0.53
9:QI:2:GLU:HB3	9:QI:20:ARG:HG3	1.89	0.53
24:RA:326:C:OP2	43:RY:73:ARG:NH1	2.34	0.53
39:RU:66:ASN:HD21	39:RU:70:ARG:HH21	1.55	0.53
44:RZ:10:ARG:NH1	44:RZ:26:GLY:O	2.42	0.53
2:XB:69:LEU:O	2:XB:163:PHE:N	2.39	0.53
3:XC:56:ASP:HB2	3:XC:67:THR:HB	1.89	0.53
20:XT:10:LEU:HD23	20:XT:12:ALA:H	1.72	0.53
36:YR:19:ALA:O	36:YR:23:ASN:ND2	2.42	0.53
1:QA:630:G:H2'	1:QA:631:G:H4'	1.89	0.53
20:QT:10:LEU:HG	20:QT:12:ALA:H	1.72	0.53
49:R4:59:PHE:O	49:R4:62:ARG:NE	2.41	0.53
24:RA:173:C:H2'	24:RA:174:U:H6	1.72	0.53
24:RA:811:A:H5'	26:RD:210:GLY:HA3	1.91	0.53
26:RD:148:GLU:HB2	26:RD:151:LYS:HD2	1.90	0.53
1:XA:1510:U:H2'	1:XA:1511:G:C8	2.43	0.53
3:XC:19:GLU:O	3:XC:40:ARG:NH2	2.41	0.53
5:XE:33:VAL:HG11	5:XE:109:ILE:HA	1.90	0.53
17:XQ:88:TYR:OH	17:XQ:92:ARG:NH1	2.41	0.53
18:XR:38:GLU:O	18:XR:42:ARG:NH1	2.41	0.53
20:XT:53:LEU:O	20:XT:57:ARG:NH1	2.42	0.53
20:XT:89:ARG:O	20:XT:93:GLU:N	2.38	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:Y3:39:ASP:OD2	48:Y3:44:ARG:NH2	2.41	0.53
24:YA:1014:U:H2'	24:YA:1015:C:C6	2.44	0.53
24:YA:2073:A:H4'	27:YE:141:ILE:HG12	1.91	0.53
1:QA:127:G:O2'	17:QQ:2:PRO:O	2.24	0.53
22:QV:16:C:H5'	22:QV:18:U:H5	1.73	0.53
24:RA:2452:C:H5''	24:RA:2599:A:H4'	1.91	0.53
27:RE:147:PRO:HB2	27:RE:149:ARG:HG2	1.90	0.53
36:RR:54:LEU:HB3	36:RR:62:ALA:HB1	1.91	0.53
39:RU:95:LEU:HD13	40:RV:4:ILE:HD12	1.90	0.53
7:XG:50:ILE:HG21	7:XG:61:VAL:HG11	1.90	0.53
10:XJ:61:GLU:OE1	14:XN:58:LYS:NZ	2.37	0.53
24:YA:2147:G:O2'	24:YA:2195:A:N6	2.38	0.53
24:YA:2156:A:O2'	24:YA:2181:G:N2	2.33	0.53
24:YA:390:G:H2'	24:YA:391:G:H8	1.73	0.53
34:YP:47:ASP:OD2	34:YP:50:ARG:NH1	2.36	0.53
1:QA:12:U:H3	1:QA:22:G:H1	1.56	0.53
24:RA:1569:U:H2'	24:RA:1570:G:H8	1.73	0.53
24:RA:1692:G:H5''	24:RA:1693:C:H5'	1.91	0.53
24:RA:2692:C:H5'	27:RE:189:PRO:HA	1.89	0.53
26:RD:66:ASP:OD2	26:RD:103:ARG:NH1	2.40	0.53
37:RS:18:ILE:HG21	37:RS:88:ASP:HA	1.90	0.53
4:XD:196:LEU:HD22	4:XD:197:PRO:HD2	1.89	0.53
24:YA:2320:G:H22	24:YA:2323:A:H2	1.56	0.53
24:YA:907:U:H1'	24:YA:2280:A:H5'	1.89	0.53
1:QA:1137:C:O2'	1:QA:1138:G:H5''	2.08	0.53
1:QA:28:G:O2'	1:QA:296:U:OP1	2.21	0.53
2:QB:70:PHE:O	2:QB:93:VAL:N	2.41	0.53
3:QC:11:ARG:NH2	3:QC:177:THR:O	2.42	0.53
16:QP:21:VAL:HG23	16:QP:34:GLU:H	1.73	0.53
24:RA:1440:U:H4'	24:RA:1649:A:H4'	1.91	0.53
1:XA:250:A:H4'	1:XA:251:G:O5'	2.09	0.53
1:XA:358:U:C2'	1:XA:358:U:O2	2.55	0.53
24:YA:2697:G:P	38:YT:51:ARG:HH22	2.31	0.53
1:QA:401:C:O2'	1:QA:621:A:N3	2.33	0.53
4:QD:15:GLU:OE2	4:QD:59:ARG:NH1	2.42	0.53
8:QH:30:ARG:HH21	8:QH:50:ARG:HH21	1.57	0.53
54:R9:6:SER:OG	54:R9:6:SER:O	2.22	0.53
24:RA:600:G:O2'	24:RA:1300:A:OP1	2.26	0.53
24:RA:2529:C:N3	24:RA:2554:A:N6	2.56	0.53
24:RA:2713:C:H3'	24:RA:2714:U:C5'	2.36	0.53
7:XG:115:ARG:HB2	7:XG:118:VAL:HG22	1.89	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:376:G:H5''	16:XP:5:ARG:HB2	1.91	0.53
22:XV:35:G:H2'	22:XV:36:G:C8	2.44	0.53
29:YG:98:ARG:NH1	49:Y4:1:MET:SD	2.81	0.53
53:Y8:26:LYS:HD2	53:Y8:47:LYS:HD3	1.91	0.53
24:YA:854:U:OP2	34:YP:41:ARG:NH2	2.42	0.53
24:YA:907:U:OP2	24:YA:962:G:N1	2.38	0.53
24:RA:2450:U:O3'	24:RA:2451:A:H3'	2.09	0.53
27:RE:2:LYS:HB3	27:RE:95:ILE:HD12	1.91	0.53
1:XA:1446:A:HO2'	1:XA:1447:G:P	2.31	0.53
1:XA:757:U:H2'	1:XA:758:G:O4'	2.09	0.53
1:XA:745:C:OP1	1:XA:851:G:O2'	2.26	0.53
2:XB:83:MET:O	2:XB:87:ARG:N	2.37	0.53
14:XN:24:CYS:HB3	14:XN:29:ARG:H	1.74	0.53
24:YA:1692:G:H5''	24:YA:1693:C:H5'	1.90	0.53
24:YA:1810:U:OP2	24:YA:1815:A:N6	2.30	0.53
24:YA:823:G:N7	24:YA:840:A:O2'	2.42	0.53
24:YA:1613:A:OP1	26:YD:211:ARG:NH1	2.42	0.53
28:RF:167:ALA:HB1	28:RF:173:VAL:HG11	1.92	0.52
41:RW:12:ILE:O	41:RW:101:SER:OG	2.25	0.52
1:XA:135:C:H2'	1:XA:136:C:H5'	1.90	0.52
1:XA:338:A:OP2	33:YO:97:ARG:NH2	2.40	0.52
45:Y0:23:VAL:HG13	45:Y0:38:VAL:HG22	1.90	0.52
47:Y2:28:LYS:HD2	47:Y2:53:LEU:HD11	1.91	0.52
1:QA:946:A:H2'	1:QA:947:G:H8	1.74	0.52
5:QE:102:ALA:HB1	5:QE:106:PRO:HG2	1.91	0.52
15:QO:26:GLU:OE2	15:QO:77:ARG:NH1	2.43	0.52
24:RA:1511:C:HO2'	24:RA:1574:A:H8	1.56	0.52
24:RA:1857:G:H4'	26:RD:242:ARG:HE	1.74	0.52
24:RA:215:G:H2'	24:RA:216:A:C8	2.45	0.52
24:RA:2455:C:H2'	24:RA:2456:G:H8	1.74	0.52
24:RA:18:C:O2'	24:RA:577:U:OP1	2.25	0.52
10:XJ:33:GLN:NE2	10:XJ:34:VAL:O	2.42	0.52
10:XJ:24:VAL:HG23	10:XJ:34:VAL:HG21	1.91	0.52
46:Y1:17:SER:HB2	46:Y1:40:ARG:HD2	1.90	0.52
26:YD:12:SER:HB3	26:YD:208:LYS:HB3	1.91	0.52
1:QA:405:U:O4	4:QD:2:GLY:N	2.43	0.52
8:QH:21:LYS:O	8:QH:65:TYR:OH	2.21	0.52
45:R0:15:ASP:OD1	45:R0:16:SER:N	2.43	0.52
46:R1:52:ARG:NH2	46:R1:55:GLY:O	2.43	0.52
24:RA:1526:G:OP2	24:RA:1556:A:N6	2.41	0.52
44:RZ:91:LEU:HD22	44:RZ:96:VAL:HG21	1.91	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1145:C:H4'	1:XA:1146:A:H8	1.73	0.52
1:XA:17:U:H2'	1:XA:18:C:C6	2.44	0.52
1:XA:414:A:OP2	1:XA:428:G:N2	2.36	0.52
1:XA:643:C:H2'	1:XA:644:G:H8	1.73	0.52
7:XG:134:ALA:O	7:XG:138:LYS:N	2.42	0.52
9:XI:9:ARG:HG2	9:XI:14:VAL:HG12	1.91	0.52
10:XJ:26:ALA:O	10:XJ:30:SER:OG	2.28	0.52
22:XV:76:A:OP1	24:YA:2451:A:N6	2.43	0.52
24:YA:2307:C:OP1	37:YS:10:ARG:NH1	2.42	0.52
24:YA:2693:C:OP2	27:YE:109:LYS:NZ	2.26	0.52
26:YD:206:LEU:HA	26:YD:211:ARG:HE	1.74	0.52
32:YN:43:THR:HG22	39:YU:64:ARG:HH11	1.73	0.52
1:QA:736:C:H2'	1:QA:737:A:C8	2.44	0.52
9:QI:26:VAL:HG12	9:QI:61:ALA:HB3	1.91	0.52
22:QV:47:U:O4	22:QV:49:G:O2'	2.26	0.52
24:RA:218:A:H4'	24:RA:219:U:O5'	2.10	0.52
24:RA:722:A:OP1	28:RF:63:LYS:NZ	2.34	0.52
35:RQ:16:ARG:HB2	35:RQ:18:LYS:HE3	1.91	0.52
1:XA:753:A:H4'	1:XA:754:C:O5'	2.09	0.52
8:XH:41:ARG:NH2	8:XH:123:GLU:OE2	2.42	0.52
24:YA:2404:A:OP2	24:YA:2434:A:N6	2.36	0.52
24:YA:2717:A:O2'	24:YA:2862:G:OP1	2.19	0.52
25:YB:40:U:H1'	25:YB:45:A:H61	1.74	0.52
24:YA:650:G:O6	34:YP:107:LYS:NZ	2.42	0.52
1:QA:1064:G:O2'	1:QA:1065:U:O5'	2.27	0.52
1:QA:1254:C:H2'	1:QA:1255:G:H8	1.73	0.52
1:QA:447:G:H21	1:QA:487:A:H62	1.56	0.52
7:QG:20:ASP:HB3	7:QG:23:VAL:HG23	1.91	0.52
1:QA:656:C:O2'	15:QO:28:GLN:OE1	2.21	0.52
24:RA:2028:C:O2'	24:RA:2833:A:N3	2.42	0.52
24:RA:332:G:N3	24:RA:332:G:H2'	2.24	0.52
1:XA:1310:G:O2'	1:XA:1311:G:OP1	2.26	0.52
1:XA:422:C:O2'	1:XA:423:G:N2	2.43	0.52
1:XA:719:C:H1'	18:XR:49:LYS:HB3	1.92	0.52
22:XV:19:G:O2'	22:XV:57:G:N2	2.26	0.52
24:YA:1311:A:OP1	24:YA:1311:A:H8	1.93	0.52
24:YA:827:G:H21	24:YA:830:A:H62	1.56	0.52
32:YN:47:ALA:HB2	32:YN:112:LEU:HD11	1.92	0.52
33:YO:8:LEU:HB2	33:YO:19:ILE:HG13	1.92	0.52
1:QA:1172:C:H2'	1:QA:1173:G:C8	2.45	0.52
1:QA:1412:C:H2'	1:QA:1413:A:C8	2.44	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:152:A:N6	1:QA:169:C:H42	2.08	0.52
11:QK:18:ARG:NH2	11:QK:35:PRO:O	2.42	0.52
17:QQ:10:VAL:HA	17:QQ:20:THR:O	2.09	0.52
22:QV:31:G:H1	22:QV:39:C:H42	1.56	0.52
45:R0:64:ASP:N	45:R0:64:ASP:OD1	2.41	0.52
24:RA:1463:C:H2'	24:RA:1464:G:O4'	2.09	0.52
24:RA:2172:U:H2'	24:RA:2173:G:H8	1.75	0.52
29:RG:16:ARG:NH2	29:RG:28:VAL:O	2.43	0.52
1:XA:201:C:H42	1:XA:216:G:H1	1.57	0.52
1:XA:769:G:H4'	1:XA:1513:A:H4'	1.91	0.52
2:XB:87:ARG:NH2	2:XB:216:SER:O	2.43	0.52
24:YA:2086:C:H2'	24:YA:2087:C:C6	2.45	0.52
24:YA:2044:U:O2'	24:YA:2629:C:H5'	2.09	0.52
25:YB:85:G:H1	25:YB:92:G:H22	1.57	0.52
1:QA:1022:G:H2'	1:QA:1023:G:C8	2.45	0.52
1:QA:674:G:H2'	1:QA:675:A:C8	2.41	0.52
24:RA:1843:A:H2'	24:RA:1844:G:H8	1.74	0.52
1:XA:963:G:H21	10:XJ:55:LYS:HG2	1.75	0.52
24:YA:1921:G:H21	24:YA:1924:C:H41	1.57	0.52
1:QA:1036:G:N7	1:QA:1037:C:N4	2.58	0.52
1:QA:501:C:H1'	1:QA:549:C:H1'	1.92	0.52
1:QA:565:U:OP2	1:QA:566:G:C2'	2.58	0.52
16:QP:37:GLY:HA3	16:QP:50:LYS:O	2.10	0.52
24:RA:2855:G:H2'	24:RA:2856:G:H8	1.75	0.52
24:RA:417:A:H4'	24:RA:418:G:H5'	1.91	0.52
24:RA:438:G:OP2	24:RA:2418:U:O2'	2.24	0.52
31:RI:65:ALA:O	31:RI:69:LYS:N	2.43	0.52
6:XF:70:ASP:OD1	6:XF:70:ASP:N	2.36	0.52
49:Y4:47:GLN:HG3	49:Y4:49:PHE:HB3	1.92	0.52
24:YA:1848:G:OP1	26:YD:88:ARG:NH2	2.41	0.52
24:YA:1974:A:N3	24:YA:2572:C:O2'	2.38	0.52
24:YA:2414:C:O2'	24:YA:2415:C:O5'	2.27	0.52
25:YB:13:A:O2'	25:YB:15:A:OP2	2.25	0.52
1:QA:543:C:OP2	4:QD:10:ARG:NH1	2.43	0.52
1:QA:662:G:H2'	1:QA:663:A:C8	2.45	0.52
24:RA:1079:U:OP1	54:R9:9:ARG:NH2	2.43	0.52
24:RA:1639:G:H2'	24:RA:1640:G:C8	2.45	0.52
1:XA:484:G:H4'	1:XA:485:G:O5'	2.10	0.52
5:XE:78:HIS:HE1	5:XE:80:ILE:HD13	1.75	0.52
8:XH:100:ILE:O	8:XH:125:ARG:NH2	2.42	0.52
46:Y1:12:PRO:HB3	46:Y1:43:TYR:HD1	1.75	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:YA:1581:U:H2'	24:YA:1582:A:C8	2.45	0.52
24:YA:2326:C:H2'	24:YA:2327:G:H8	1.74	0.52
24:YA:64:C:O2'	24:YA:482:C:N3	2.35	0.52
28:YF:63:LYS:NZ	28:YF:75:HIS:O	2.40	0.52
1:QA:134:A:H61	16:QP:25:ARG:HH12	1.57	0.52
1:QA:152:A:N6	1:QA:169:C:N4	2.57	0.52
1:QA:618:C:H5'	1:QA:619:U:H5''	1.92	0.52
2:QB:19:HIS:HB3	2:QB:20:GLU:HG2	1.92	0.52
36:RR:33:ARG:HG3	36:RR:115:GLU:HB3	1.92	0.52
42:RX:5:TYR:O	47:R2:36:ARG:NH2	2.43	0.52
6:XF:3:ARG:NH1	6:XF:38:GLU:OE2	2.43	0.52
17:XQ:81:ARG:NH2	17:XQ:83:ASP:OD2	2.42	0.52
48:Y3:15:TYR:CE2	48:Y3:53:LEU:HD21	2.45	0.52
1:QA:64:G:OP1	1:QA:382:A:N6	2.40	0.51
25:RB:50:G:OP1	37:RS:62:LYS:N	2.44	0.51
27:RE:29:GLY:HA3	27:RE:51:PHE:HE1	1.74	0.51
29:RG:52:ILE:HG23	29:RG:55:LYS:HB3	1.92	0.51
1:XA:437:U:O3'	4:XD:125:HIS:NE2	2.39	0.51
24:YA:641:G:OP1	28:YF:40:GLN:NE2	2.44	0.51
28:YF:181:LEU:O	28:YF:205:ARG:NH2	2.42	0.51
28:YF:157:VAL:HB	28:YF:194:MET:HB3	1.92	0.51
29:YG:27:ASN:HB3	29:YG:30:GLU:HG3	1.92	0.51
31:YI:65:ALA:O	31:YI:69:LYS:N	2.44	0.51
33:YO:97:ARG:H	33:YO:117:LEU:HD22	1.76	0.51
45:R0:72:ARG:HB2	45:R0:75:LEU:HB2	1.92	0.51
24:RA:1041:C:O2	32:RN:3:THR:OG1	2.23	0.51
24:RA:2034:G:OP1	41:RW:11:ARG:NH2	2.43	0.51
24:RA:2804:C:H4'	24:RA:2805:G:H5'	1.92	0.51
24:RA:331:G:C8	24:RA:331:G:C5'	2.85	0.51
29:RG:27:ASN:HB3	29:RG:30:GLU:HG3	1.93	0.51
1:XA:1435:G:H2'	1:XA:1436:U:C6	2.46	0.51
1:XA:407:G:H2'	1:XA:408:A:H8	1.74	0.51
1:XA:713:G:H2'	1:XA:714:G:C8	2.45	0.51
1:XA:742:G:OP2	15:XO:35:ARG:NH2	2.42	0.51
19:XS:50:ALA:HB1	19:XS:57:HIS:HB3	1.91	0.51
24:YA:1526:G:N7	24:YA:1556:A:N6	2.58	0.51
24:YA:2162:C:H2'	24:YA:2163:G:H8	1.75	0.51
24:YA:673:G:O2'	24:YA:2363:G:OP1	2.21	0.51
1:QA:448:A:H62	1:QA:486:U:H3	1.59	0.51
2:QB:192:SER:OG	2:QB:193:ASP:N	2.38	0.51
17:QQ:77:VAL:HG12	17:QQ:78:GLU:HG2	1.92	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:RA:2877:G:O2'	24:RA:2878:A:H8	1.93	0.51
24:YA:1003:U:OP2	35:YQ:14:ARG:NH1	2.42	0.51
24:YA:1911:A:N3	24:YA:2108:U:O2'	2.44	0.51
24:YA:2118:U:O4	24:YA:2215:G:O6	2.28	0.51
24:YA:2258:G:H2'	24:YA:2259:A:C8	2.45	0.51
24:YA:2376:C:OP1	45:Y0:55:ARG:NH1	2.43	0.51
41:YW:18:ARG:HD2	41:YW:76:VAL:HB	1.91	0.51
1:QA:1151:A:H2'	1:QA:1152:A:C8	2.45	0.51
1:QA:581:G:N2	1:QA:759:A:OP2	2.37	0.51
24:RA:1120:G:C2	24:RA:1121:C:H1'	2.45	0.51
24:RA:2317:A:N6	29:RG:154:GLY:O	2.44	0.51
1:XA:1037:C:H2'	1:XA:1038:C:C6	2.46	0.51
1:XA:678:U:H2'	1:XA:679:C:C6	2.46	0.51
1:XA:977:A:OP1	14:XN:31:ARG:NH2	2.44	0.51
24:YA:828:A:OP1	26:YD:218:ARG:NH2	2.44	0.51
28:YF:185:ASP:HA	28:YF:188:ARG:HD3	1.93	0.51
28:YF:195:ASP:OD1	28:YF:195:ASP:N	2.39	0.51
1:QA:1209:C:O2'	1:QA:1214:C:N4	2.43	0.51
10:QJ:34:VAL:HG13	10:QJ:74:ILE:HG12	1.93	0.51
10:QJ:77:PRO:O	10:QJ:79:ARG:NH1	2.43	0.51
22:QV:30:C:N3	22:QV:31:G:N7	2.59	0.51
22:QV:30:C:C2	22:QV:31:G:N7	2.79	0.51
47:R2:32:LEU:O	47:R2:36:ARG:N	2.43	0.51
24:RA:2118:U:O4	24:RA:2215:G:O6	2.28	0.51
35:RQ:135:ASP:HB3	35:RQ:137:TYR:HD2	1.76	0.51
39:RU:87:GLY:HA3	40:RV:49:THR:HG23	1.93	0.51
3:XC:19:GLU:HB3	3:XC:40:ARG:HH21	1.76	0.51
12:XL:87:GLY:HA2	12:XL:98:TYR:HD1	1.76	0.51
24:YA:2297:C:OP2	51:Y6:26:ASN:ND2	2.43	0.51
25:YB:88:C:H2'	25:YB:89(A):G:C8	2.46	0.51
5:QE:75:THR:OG1	5:QE:76:ILE:N	2.44	0.51
24:RA:98:U:H4'	24:RA:99:G:O5'	2.10	0.51
27:RE:5:LEU:HD12	27:RE:51:PHE:HB2	1.92	0.51
34:RP:52:GLU:OE2	34:RP:58:THR:OG1	2.25	0.51
1:XA:1001:G:N2	1:XA:1040:U:O2	2.43	0.51
1:XA:1323:G:H2'	1:XA:1324:A:C8	2.46	0.51
3:XC:150:LYS:HB3	3:XC:201:TYR:HB2	1.93	0.51
8:XH:69:ARG:NH2	8:XH:73:ASP:O	2.44	0.51
23:XX:5:A:H2'	23:XX:6:G:C8	2.43	0.51
2:QB:130:ARG:O	2:QB:135:GLN:NE2	2.43	0.51
6:QF:8:ILE:HD13	6:QF:26:ILE:HD13	1.91	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1347:G:H3'	9:QI:108:VAL:O	2.11	0.51
10:QJ:24:VAL:HG11	10:QJ:37:PRO:HD3	1.93	0.51
22:QV:3:G:O2'	22:QV:4:U:H5'	2.10	0.51
24:RA:1387:U:OP1	24:RA:1443:U:N3	2.31	0.51
29:RG:63:ILE:HG13	29:RG:64:THR:HG23	1.93	0.51
38:RT:128:GLU:O	38:RT:132:LYS:N	2.41	0.51
44:RZ:148:ASP:N	44:RZ:148:ASP:OD1	2.43	0.51
1:XA:1244:C:H2'	1:XA:1245:A:C8	2.46	0.51
1:XA:1348:U:H3	1:XA:1374:A:H2	1.56	0.51
14:XN:40:CYS:SG	14:XN:42:ILE:N	2.79	0.51
24:YA:1176:U:HO2'	24:YA:1177:G:P	2.34	0.51
24:YA:830:A:H8	24:YA:831:A:H4'	1.76	0.51
26:YD:108:PRO:HB3	26:YD:143:HIS:HE1	1.75	0.51
6:QF:71:ARG:O	6:QF:75:LEU:N	2.43	0.51
24:RA:1615:G:P	26:RD:63:ARG:HH12	2.34	0.51
24:RA:2207:C:H2'	24:RA:2208:G:H8	1.75	0.51
1:XA:1342:C:H2'	1:XA:1343:G:C8	2.44	0.51
1:XA:627:G:H2'	1:XA:628:G:H8	1.76	0.51
24:YA:1069:U:OP2	24:YA:1070:G:N7	2.44	0.51
24:YA:1463:C:H2'	24:YA:1464:G:O4'	2.11	0.51
32:YN:6:PRO:HG3	32:YN:41:ASP:HB2	1.93	0.51
1:QA:707:C:H2'	1:QA:708:C:H6	1.76	0.51
24:RA:2140:U:H3	24:RA:2170:G:H4'	1.76	0.51
24:RA:332:G:H3'	24:RA:333:G:C8	2.46	0.51
24:RA:238:C:HO2'	34:RP:64:LYS:NZ	2.08	0.51
1:XA:1060:C:H2'	1:XA:1061:G:H8	1.76	0.51
1:XA:692:U:O2'	1:XA:694:A:N7	2.35	0.51
24:YA:1334:U:O2'	24:YA:1694:G:N2	2.44	0.51
24:YA:2068:G:H5'	50:Y5:19:ARG:HG3	1.92	0.51
24:YA:2318:C:H2'	24:YA:2319:G:H21	1.76	0.51
33:YO:2:ILE:HB	33:YO:33:ALA:HB3	1.92	0.51
1:QA:1210:C:O2'	1:QA:1213:A:O2'	2.24	0.51
1:QA:1244:C:H2'	1:QA:1245:A:C8	2.46	0.51
1:QA:985:C:H2'	1:QA:986:A:C8	2.46	0.51
4:QD:199:ASN:HB3	4:QD:202:LEU:HD23	1.93	0.51
24:RA:2399:U:O2'	45:R0:41:ARG:NH2	2.44	0.51
24:RA:895:G:H2'	24:RA:896:A:C8	2.45	0.51
25:RB:38:C:H2'	25:RB:39:A:H8	1.75	0.51
25:RB:39:A:O2'	25:RB:46:A:N1	2.40	0.51
31:RI:94:ALA:HB1	31:RI:114:LEU:HD23	1.93	0.51
41:RW:88:ARG:NH1	41:RW:94:ASP:OD2	2.44	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1052:U:O4	1:XA:1206:G:O6	2.29	0.51
1:XA:1325:C:H4'	21:XU:17:THR:HG21	1.93	0.51
1:XA:165:C:H2'	1:XA:166:G:H8	1.76	0.51
2:XB:67:THR:HG23	2:XB:159:PRO:HA	1.93	0.51
4:XD:111:ALA:HB1	4:XD:116:GLN:HB3	1.93	0.51
9:XI:10:ARG:HG2	9:XI:11:LYS:HB2	1.93	0.51
1:XA:973:G:H4'	14:YN:41:ARG:HH22	1.75	0.51
20:XT:57:ARG:NH1	20:XT:102:GLY:O	2.39	0.51
54:Y9:2:LYS:NZ	54:Y9:31:LYS:O	2.35	0.51
24:YA:2541:G:H5''	24:YA:2542:A:H5''	1.93	0.51
1:QA:503:C:OP2	12:QL:116:SER:OG	2.25	0.50
8:QH:10:LEU:HD22	8:QH:83:ILE:HD11	1.93	0.50
24:RA:1217:G:N7	24:RA:1219:A:N6	2.59	0.50
24:RA:2136:A:C8	24:RA:2190:G:N2	2.79	0.50
24:RA:2136:A:N7	24:RA:2190:G:N2	2.60	0.50
24:RA:2863:C:H2'	24:RA:2864:G:H8	1.76	0.50
39:RU:97:ASP:OD1	39:RU:101:ARG:NH1	2.45	0.50
39:RU:17:ILE:HG13	39:RU:32:PHE:HE1	1.75	0.50
1:XA:701:C:O2	1:XA:703:G:N1	2.44	0.50
24:YA:231:G:H1'	24:YA:232:U:OP2	2.11	0.50
44:YZ:93:ASP:HB2	44:YZ:131:ARG:HH12	1.76	0.50
44:YZ:61:LEU:HB2	44:YZ:65:GLN:HB2	1.93	0.50
1:QA:272:C:H2'	1:QA:273:A:H8	1.75	0.50
1:QA:656:C:O2	15:QO:28:GLN:NE2	2.42	0.50
24:RA:1103:A:O2'	24:RA:1104:G:OP1	2.25	0.50
24:RA:1186:U:H1'	24:RA:1188:A:C6	2.46	0.50
24:RA:2149:G:OP2	24:RA:2149:G:H8	1.94	0.50
24:RA:722:A:N3	24:RA:2455:C:O2'	2.38	0.50
29:RG:16:ARG:HH21	29:RG:28:VAL:HB	1.75	0.50
34:RP:91:PHE:O	34:RP:121:LYS:NZ	2.35	0.50
1:XA:662:G:H2'	1:XA:663:A:C8	2.46	0.50
1:XA:715:A:H2'	1:XA:716:A:C8	2.46	0.50
1:XA:806:C:H2'	1:XA:807:A:H8	1.76	0.50
6:XF:22:GLU:OE2	6:XF:84:ASN:ND2	2.36	0.50
8:XH:8:ASP:OD2	8:XH:12:ARG:NH2	2.44	0.50
13:XM:49:THR:HB	13:XM:52:GLU:HG3	1.92	0.50
22:XV:68:C:H2'	22:XV:69:A:C8	2.46	0.50
25:YB:77:U:OP1	44:YZ:19:ARG:NH2	2.43	0.50
24:YA:2678:C:H42	30:YH:109:PHE:HA	1.75	0.50
44:YZ:140:ASP:OD1	44:YZ:142:SER:OG	2.29	0.50
1:QA:254:G:O2'	17:QQ:16:GLN:O	2.29	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:339:C:OP2	33:RO:97:ARG:NH1	2.44	0.50
1:QA:191(G):G:O2'	20:QT:101:GLY:O	2.29	0.50
24:RA:419:C:H5''	24:RA:436:C:H5''	1.93	0.50
24:RA:70:A:H5''	24:RA:71:U:H3'	1.91	0.50
1:XA:565:U:OP2	1:XA:566:G:O2'	2.27	0.50
11:XK:22:HIS:HB3	11:XK:29:ILE:HG23	1.93	0.50
12:XL:76:ASN:OD1	12:XL:76:ASN:N	2.45	0.50
22:XV:5:G:H2'	22:XV:6:A:C8	2.46	0.50
49:Y4:58:ARG:HA	49:Y4:61:ARG:HE	1.75	0.50
25:YB:65:C:H2'	25:YB:66:A:H5'	1.93	0.50
29:YG:82:LEU:HD21	29:YG:88:ILE:HG21	1.92	0.50
30:YH:164:TYR:HB2	30:YH:167:GLU:HB2	1.92	0.50
30:YH:25:LYS:NZ	30:YH:34:GLU:OE2	2.43	0.50
1:QA:152:A:H62	1:QA:169:C:H42	1.60	0.50
1:QA:524:G:H2'	1:QA:525:C:C6	2.46	0.50
3:QC:71:ALA:HB2	3:QC:106:VAL:HB	1.92	0.50
24:RA:1356:G:OP2	52:R7:9:ARG:NE	2.40	0.50
24:RA:2024:G:OP2	36:RR:9:LYS:NZ	2.41	0.50
24:RA:172:C:O2'	24:RA:459:A:N3	2.41	0.50
26:RD:260:ARG:NH2	26:RD:266:SER:OG	2.39	0.50
35:RQ:34:LEU:HB2	35:RQ:118:LEU:HD22	1.93	0.50
40:RV:24:LYS:HA	40:RV:92:THR:HG23	1.94	0.50
44:RZ:77:ASP:OD2	44:RZ:80:ARG:NE	2.45	0.50
1:XA:1412:C:H2'	1:XA:1413:A:C8	2.47	0.50
1:XA:505:G:H2'	1:XA:506:G:H8	1.76	0.50
1:XA:842:C:O2'	1:XA:848:C:N4	2.45	0.50
16:XP:1:MET:SD	16:XP:1:MET:N	2.78	0.50
1:XA:227:G:N2	16:XP:62:VAL:O	2.43	0.50
24:YA:1725:G:N2	24:YA:2011:G:H22	2.10	0.50
25:YB:45:A:O4'	29:YG:95:ARG:NH1	2.45	0.50
26:YD:259:THR:OG1	26:YD:259:THR:O	2.28	0.50
24:YA:2647:C:O2'	27:YE:80:GLU:OE1	2.24	0.50
40:YV:8:GLY:O	40:YV:10:LYS:NZ	2.43	0.50
44:YZ:153:SER:OG	44:YZ:154:ASP:OD1	2.27	0.50
24:RA:1018:A:H5'	24:RA:1233:U:H1'	1.93	0.50
24:RA:1554:A:O2'	24:RA:1555:C:O4'	2.29	0.50
24:RA:1855:G:N3	26:RD:254:THR:OG1	2.44	0.50
24:RA:346:A:H5'	24:RA:364:A:H1'	1.92	0.50
24:RA:732:A:OP1	24:RA:733:G:N2	2.44	0.50
27:RE:119:ARG:NH1	27:RE:159:HIS:O	2.45	0.50
1:XA:1143:G:H2'	1:XA:1144:G:C8	2.47	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:358:U:H6	1:XA:358:U:O5'	1.93	0.50
10:XJ:38:ILE:HD11	10:XJ:71:LEU:HD23	1.94	0.50
14:XN:6:LEU:HB3	14:XN:23:ARG:HH22	1.76	0.50
24:YA:2054:G:OP2	24:YA:2466:G:O2'	2.28	0.50
24:YA:468:G:H1'	28:YF:48:THR:HG21	1.94	0.50
1:QA:119:A:H4'	1:QA:120:A:O5'	2.11	0.50
1:QA:501:C:H2'	1:QA:502:G:C8	2.45	0.50
24:RA:1255:A:H4'	24:RA:1256:U:O5'	2.12	0.50
24:RA:1473:A:H4'	24:RA:1474:C:O5'	2.10	0.50
24:RA:2631:C:OP1	27:RE:152:LYS:HE2	2.12	0.50
24:RA:170:A:H1'	24:RA:462:C:H5'	1.94	0.50
1:XA:41:G:H2'	1:XA:42:G:H8	1.77	0.50
1:XA:545:C:OP1	4:XD:61:LYS:NZ	2.44	0.50
1:XA:559:A:H4'	1:XA:560:U:H3'	1.93	0.50
24:YA:1235:G:H5'	34:YP:32:THR:HA	1.93	0.50
24:YA:2555:G:H2'	24:YA:2556:G:C8	2.47	0.50
24:YA:2789:A:H4'	24:YA:2790:G:O5'	2.10	0.50
24:YA:2819:A:H2'	24:YA:2820:A:C8	2.47	0.50
30:YH:51:ARG:HB3	30:YH:51:ARG:NH1	2.26	0.50
5:QE:33:VAL:HG13	5:QE:112:LEU:HD22	1.92	0.50
13:QM:105:THR:OG1	13:QM:106:ASN:N	2.44	0.50
22:QV:43:G:H2'	22:QV:44:G:C8	2.47	0.50
49:R4:58:ARG:HA	49:R4:61:ARG:HH21	1.77	0.50
24:RA:1571:G:H2'	24:RA:1572:G:C8	2.47	0.50
24:RA:321:C:H2'	24:RA:322:G:O4'	2.12	0.50
33:RO:120:GLU:OE1	38:RT:67:SER:OG	2.27	0.50
33:RO:34:THR:OG1	33:RO:35:VAL:N	2.45	0.50
36:RR:30:THR:O	36:RR:78:LYS:NZ	2.42	0.50
1:XA:812:C:H1'	1:XA:813:U:OP2	2.12	0.50
3:XC:9:GLY:HA2	3:XC:12:LEU:HG	1.93	0.50
4:XD:96:LEU:HD22	4:XD:139:ARG:HH21	1.76	0.50
22:XV:43:G:H2'	22:XV:44:G:C8	2.47	0.50
24:YA:1108:G:H22	24:YA:1124:U:H1'	1.77	0.50
32:YN:63:THR:OG1	32:YN:64:GLY:N	2.44	0.50
1:QA:1355:G:H2'	1:QA:1356:G:C8	2.47	0.50
2:QB:166:ASP:HB3	2:QB:169:LYS:HB3	1.92	0.50
7:QG:115:ARG:HG3	7:QG:117:ALA:H	1.75	0.50
46:R1:87:PRO:HA	46:R1:90:ILE:HG22	1.93	0.50
24:RA:1571:G:H2'	24:RA:1572:G:H8	1.77	0.50
24:RA:2148:A:N6	24:RA:2185:C:O2'	2.42	0.50
24:RA:299:G:H3'	24:RA:300:A:H5''	1.94	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:RO:13:ASN:ND2	33:RO:96:THR:OG1	2.35	0.50
24:YA:2158:C:H42	24:YA:2178:G:H21	1.59	0.50
29:YG:66:GLN:OE1	29:YG:98:ARG:NH1	2.44	0.50
41:YW:12:ILE:O	41:YW:101:SER:OG	2.29	0.50
1:QA:514:C:H2'	1:QA:515:G:H8	1.77	0.50
1:QA:634:C:H2'	1:QA:635:G:H8	1.76	0.50
49:R4:24:THR:OG1	49:R4:25:TYR:N	2.42	0.50
24:RA:1046:A:H2'	24:RA:1047:A:C8	2.47	0.50
38:RT:39:ARG:HH21	38:RT:41:ARG:HD3	1.77	0.50
4:XD:162:LEU:HA	4:XD:165:MET:HB2	1.93	0.50
11:XK:15:ALA:HA	11:XK:77:MET:HA	1.93	0.50
47:Y2:29:LYS:HE3	47:Y2:57:ILE:HG21	1.94	0.50
24:YA:2846:U:H2'	24:YA:2847:G:C8	2.47	0.50
24:YA:589:U:H5''	34:YP:29:LYS:HE3	1.93	0.50
6:QF:9:VAL:HA	6:QF:59:TYR:O	2.12	0.49
14:QN:27:CYS:SG	14:QN:29:ARG:N	2.83	0.49
24:RA:1361:C:O2'	24:RA:1438:A:N3	2.37	0.49
24:RA:2155:G:O2'	24:RA:2180:A:N1	2.39	0.49
24:RA:236:G:H4'	24:RA:413:G:C5	2.47	0.49
30:RH:153:LYS:HB2	30:RH:162:ILE:H	1.77	0.49
1:XA:410:G:H21	1:XA:432:A:N6	2.05	0.49
1:XA:736:C:H2'	1:XA:737:A:C8	2.47	0.49
22:XV:2:G:H2'	22:XV:3:G:C8	2.47	0.49
24:YA:1475:G:H2'	24:YA:1476:C:C6	2.47	0.49
24:YA:815:G:O2'	24:YA:1425:A:N6	2.42	0.49
25:YB:89(B):A:H2'	25:YB:90:C:H1'	1.94	0.49
33:YO:104:ARG:HH11	33:YO:121:VAL:HG12	1.75	0.49
43:YY:102:CYS:SG	43:YY:103:GLY:N	2.85	0.49
1:QA:576:G:N2	1:QA:760:G:OP2	2.44	0.49
1:QA:8:A:N6	4:QD:205:GLU:O	2.46	0.49
14:QN:47:LEU:HA	14:QN:50:LYS:HB2	1.94	0.49
53:R8:49:VAL:HG23	53:R8:53:PRO:HD3	1.94	0.49
24:RA:1112:U:O2'	24:RA:1114:G:OP2	2.28	0.49
24:RA:1714:G:O2'	24:RA:2013:U:O4	2.29	0.49
24:RA:650:G:O6	34:RP:107:LYS:NZ	2.41	0.49
44:RZ:33:LEU:HD11	44:RZ:90:VAL:HG21	1.94	0.49
1:XA:1129:C:O2	1:XA:1132:C:N4	2.45	0.49
1:XA:1227:A:OP1	19:XS:80:TYR:OH	2.22	0.49
2:XB:47:THR:HG23	2:XB:202:PRO:HG2	1.93	0.49
1:QA:501:C:H2'	1:QA:502:G:H8	1.76	0.49
49:R4:28:LYS:HB3	49:R4:31:ILE:HD11	1.94	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:RA:1715:A:H62	24:RA:2013:U:H3	1.59	0.49
24:RA:2141:A:N6	24:RA:2192:A:N7	2.58	0.49
24:RA:887:C:H2'	24:RA:888:A:C8	2.47	0.49
24:RA:992:G:H2'	24:RA:993:G:C8	2.47	0.49
39:RU:49:HIS:HA	39:RU:52:ARG:HB3	1.95	0.49
1:XA:493:G:N2	1:XA:494:U:O4	2.45	0.49
1:XA:757:U:O2'	1:XA:879:C:O2	2.29	0.49
24:YA:1070:G:OP2	24:YA:1071:G:C2'	2.60	0.49
1:QA:1407:C:H2'	1:QA:1408:A:H8	1.76	0.49
23:QX:19:G:H2'	23:QX:19:G:N3	2.27	0.49
24:RA:1604:C:OP2	24:RA:1605:A:O2'	2.24	0.49
24:RA:2340:A:H2'	24:RA:2341:G:H8	1.76	0.49
24:RA:210:A:C4	24:RA:255:G:N7	2.81	0.49
28:RF:117:ARG:NH2	28:RF:189:THR:O	2.46	0.49
30:RH:25:LYS:NZ	30:RH:34:GLU:OE1	2.46	0.49
36:YR:59:ASP:OD1	36:YR:59:ASP:N	2.37	0.49
39:YU:92:ARG:CZ	40:YV:11:GLN:H	2.25	0.49
1:QA:1298:C:H4'	1:QA:1299:A:C4	2.48	0.49
9:QI:28:VAL:O	9:QI:31:GLN:N	2.46	0.49
24:RA:1617:A:H2'	24:RA:1618:A:C8	2.48	0.49
9:XI:50:LEU:HD23	9:XI:56:LEU:HA	1.94	0.49
24:YA:1068:G:N2	24:YA:1069:U:O4	2.45	0.49
24:YA:1386:U:OP1	42:YX:16:LYS:NZ	2.41	0.49
24:YA:312:C:H2'	24:YA:313:A:C8	2.45	0.49
25:YB:89(B):A:H2'	25:YB:90:C:C1'	2.43	0.49
29:YG:37:VAL:HG13	29:YG:159:VAL:HG12	1.94	0.49
29:YG:16:ARG:NH2	29:YG:28:VAL:O	2.46	0.49
43:YY:17:SER:OG	43:YY:71:LYS:NZ	2.31	0.49
1:QA:1096:C:H2'	1:QA:1097:C:H6	1.77	0.49
1:QA:1254:C:H2'	1:QA:1255:G:C8	2.47	0.49
1:QA:1435:G:H2'	1:QA:1436:U:C6	2.47	0.49
1:QA:483:C:OP2	1:QA:484:G:O2'	2.18	0.49
2:QB:193:ASP:N	2:QB:193:ASP:OD1	2.44	0.49
16:QP:23:ASP:OD2	16:QP:25:ARG:NH2	2.46	0.49
24:RA:1130:A:N7	24:RA:1131:A:O2'	2.42	0.49
24:RA:2303:U:H2'	24:RA:2304:C:C6	2.47	0.49
24:RA:2376:C:H2'	24:RA:2377:G:O4'	2.13	0.49
2:XB:118:LEU:HD23	2:XB:142:LEU:HD23	1.94	0.49
24:YA:2173:G:H2'	24:YA:2174:G:C8	2.48	0.49
24:YA:2251:G:OP1	26:YD:244:ARG:NH1	2.35	0.49
27:YE:24:THR:OG1	27:YE:186:GLY:O	2.25	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
43:YY:47:LYS:NZ	43:YY:48:ALA:O	2.46	0.49
1:QA:1297:C:O2'	1:QA:1298:C:OP2	2.28	0.49
1:QA:41:G:H2'	1:QA:42:G:H8	1.77	0.49
1:QA:7:G:H21	5:QE:121:LYS:HG2	1.78	0.49
24:RA:1991:A:O2'	24:RA:1994:A:N3	2.42	0.49
24:RA:602:G:O2'	24:RA:2041:A:OP1	2.26	0.49
34:RP:47:ASP:OD2	34:RP:50:ARG:NE	2.44	0.49
35:RQ:81:VAL:O	35:RQ:82:ARG:NE	2.33	0.49
1:XA:123:C:OP1	1:XA:311:C:O2'	2.30	0.49
1:XA:1502:A:H2'	1:XA:1504:G:N7	2.28	0.49
2:XB:27:LYS:O	2:XB:30:ARG:NH1	2.45	0.49
3:XC:174:PRO:O	3:XC:177:THR:OG1	2.31	0.49
4:XD:85:LYS:HD2	4:XD:92:VAL:HG11	1.94	0.49
24:YA:1248:G:H3'	24:YA:1249:A:H5''	1.95	0.49
1:QA:1249:C:O2'	9:QL:73:GLN:OE1	2.27	0.49
11:QK:98:LEU:O	11:QK:101:SER:OG	2.26	0.49
1:XA:195:A:H2'	1:XA:196:A:C4	2.48	0.49
2:XB:28:PHE:HD1	2:XB:194:PRO:HD3	1.77	0.49
24:YA:2584:A:OP1	24:YA:2586:G:O2'	2.28	0.49
1:QA:384:G:H2'	1:QA:385:C:C6	2.48	0.49
24:RA:1525:G:H2'	24:RA:1526:G:H8	1.77	0.49
24:RA:2131:U:O4	24:RA:2201:C:N4	2.41	0.49
24:RA:2420:U:H2'	24:RA:2421:G:H8	1.77	0.49
24:RA:946:A:H3'	24:RA:947:A:H8	1.78	0.49
1:XA:407:G:H2'	1:XA:408:A:C8	2.48	0.49
4:XD:106:TYR:C	4:XD:106:TYR:CD2	2.86	0.49
8:XH:121:ASP:N	8:XH:121:ASP:OD1	2.43	0.49
24:YA:1041:C:O2	32:YN:3:THR:OG1	2.30	0.49
24:YA:1855:G:N3	26:YD:254:THR:OG1	2.46	0.49
24:YA:869:U:H2'	24:YA:870:G:H8	1.78	0.49
30:YH:3:ARG:HH21	30:YH:5:GLY:H	1.61	0.49
31:YI:81:VAL:HG21	31:YI:88:ILE:HD12	1.94	0.49
38:YT:24:PRO:HA	38:YT:49:VAL:HG13	1.95	0.49
1:QA:1013:G:N2	1:QA:1016:A:OP2	2.38	0.49
1:QA:1192:C:O2	5:QE:25:ARG:NH2	2.45	0.49
1:QA:1352:C:H2'	1:QA:1353:G:C8	2.47	0.49
1:QA:224:C:H2'	1:QA:225:C:H6	1.78	0.49
1:QA:280:C:N3	17:QQ:39:SER:OG	2.36	0.49
1:QA:976:G:OP2	1:QA:1358:U:O2'	2.30	0.49
3:QC:131:ARG:NH1	3:QC:166:GLU:OE2	2.45	0.49
3:QC:9:GLY:HA2	3:QC:12:LEU:HG	1.95	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:719:C:O2'	18:QR:50:ILE:O	2.26	0.49
24:RA:2258:G:H2'	24:RA:2259:A:C8	2.48	0.49
24:RA:626:A:N1	24:RA:650:G:O2'	2.39	0.49
1:XA:1371:G:H4'	9:XI:69:GLY:HA3	1.94	0.49
2:XB:131:PRO:O	2:XB:135:GLN:N	2.45	0.49
8:XH:39:LEU:O	8:XH:44:PHE:N	2.37	0.49
24:YA:1511:C:HO2'	24:YA:1574:A:H8	1.56	0.49
24:YA:2285:A:H2'	24:YA:2286:A:C8	2.48	0.49
24:YA:2340:A:H2'	24:YA:2341:G:C8	2.48	0.49
25:YB:108:C:H5''	25:YB:108:C:H6	1.77	0.49
1:QA:148:G:H2'	1:QA:149:A:C8	2.48	0.48
1:QA:619:U:N3	4:QD:134:ASP:OD1	2.42	0.48
24:RA:1183:G:H2'	24:RA:1184:G:O4'	2.13	0.48
24:RA:1829:U:O2'	24:RA:1833:A:N3	2.40	0.48
24:RA:2258:G:H2'	24:RA:2259:A:H8	1.78	0.48
24:RA:2455:C:H2'	24:RA:2456:G:C8	2.48	0.48
24:RA:902:G:H1	24:RA:968:U:H3	1.61	0.48
26:RD:108:PRO:HD2	26:RD:111:LEU:HD13	1.95	0.48
29:RG:82:LEU:HD21	29:RG:88:ILE:HD13	1.95	0.48
1:XA:591:U:OP1	8:XH:30:ARG:NH1	2.46	0.48
16:XP:5:ARG:NH2	16:XP:27:LYS:O	2.46	0.48
47:Y2:17:SER:HB3	47:Y2:20:GLU:HG2	1.94	0.48
24:YA:1249:A:O2'	24:YA:1250:U:O5'	2.30	0.48
24:YA:208:G:N3	24:YA:223:C:O2'	2.42	0.48
1:QA:1355:G:H2'	1:QA:1356:G:H8	1.77	0.48
1:QA:422:C:O2'	1:QA:423:G:N2	2.46	0.48
23:QX:3:C:H2'	23:QX:4:A:H8	1.78	0.48
24:RA:1148:C:H2'	24:RA:1149:A:C8	2.48	0.48
27:RE:89:ASP:N	27:RE:89:ASP:OD1	2.34	0.48
38:RT:65:LYS:HE3	38:RT:67:SER:HB2	1.95	0.48
1:XA:1391:U:H2'	1:XA:1392:G:C8	2.49	0.48
1:XA:596:C:H2'	1:XA:597:G:H8	1.78	0.48
4:XD:129:ASN:HD21	4:XD:144:ASP:HB3	1.79	0.48
4:XD:25:ARG:O	4:XD:28:SER:HB3	2.12	0.48
9:XI:28:VAL:HA	9:XI:63:ILE:HB	1.94	0.48
19:XS:32:LYS:HA	19:XS:50:ALA:HB3	1.95	0.48
22:XV:32:U:H3	22:XV:38:A:H61	1.61	0.48
24:YA:1716:A:N3	24:YA:1716:A:C2'	2.75	0.48
24:YA:2270:C:O2'	24:YA:2439:C:OP2	2.26	0.48
1:QA:190:G:H8	1:QA:190:G:OP1	1.96	0.48
1:QA:565:U:H5''	1:QA:566:G:H2'	1.96	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
45:R0:27:GLU:HB2	45:R0:69:PHE:HD1	1.77	0.48
24:RA:1576:G:H1	24:RA:1587:U:H3	1.61	0.48
24:RA:2155:G:H1'	24:RA:2180:A:H61	1.78	0.48
28:RF:15:SER:OG	28:RF:16:GLY:N	2.46	0.48
29:RG:124:SER:OG	29:RG:124:SER:O	2.26	0.48
1:XA:1287:A:H2	1:XA:1353:G:H1'	1.76	0.48
1:XA:505:G:H2'	1:XA:506:G:C8	2.48	0.48
4:XD:68:TYR:HD2	4:XD:97:LEU:HD22	1.78	0.48
13:XM:107:ALA:H	13:XM:108:ARG:HG2	1.78	0.48
24:YA:236:G:H4'	24:YA:413:G:C5	2.48	0.48
1:QA:1073:U:O2	2:QB:104:ASN:ND2	2.41	0.48
8:QH:20:TYR:HE2	8:QH:75:ARG:HD2	1.78	0.48
13:QM:19:LEU:HD21	13:QM:56:LEU:HD21	1.95	0.48
19:QS:49:ILE:HG23	19:QS:60:VAL:HB	1.95	0.48
24:RA:139:A:H8	24:RA:1454:C:O2'	1.95	0.48
24:RA:628:C:O2	24:RA:704:U:O2'	2.32	0.48
24:RA:887:C:H2'	24:RA:888:A:H8	1.77	0.48
24:RA:1713:G:N3	33:RO:3:GLN:NE2	2.61	0.48
37:RS:10:ARG:NH1	37:RS:91:PRO:O	2.42	0.48
44:RZ:181:GLU:HG3	44:RZ:183:LEU:HG	1.95	0.48
1:XA:674:G:H2'	1:XA:675:A:C8	2.45	0.48
1:XA:950:U:H2'	1:XA:951:G:H8	1.79	0.48
9:XI:112:LYS:NZ	9:XI:117:HIS:O	2.37	0.48
18:XR:58:LEU:HB3	18:XR:62:GLU:HB2	1.95	0.48
22:XV:8:U:H5'	22:XV:49:G:H5'	1.94	0.48
24:YA:1787:G:H4'	24:YA:1789:G:O4'	2.14	0.48
27:YE:47:VAL:HG11	27:YE:86:PRO:HD2	1.95	0.48
1:QA:1130:A:H62	1:QA:1144:G:H21	1.60	0.48
1:QA:59:A:H5''	1:QA:387:U:H5''	1.94	0.48
3:QC:3:ASN:N	3:QC:3:ASN:OD1	2.46	0.48
5:QE:80:ILE:HD12	8:QH:104:ARG:HH22	1.77	0.48
13:QM:84:ILE:HG13	13:QM:86:CYS:H	1.78	0.48
23:QX:4:A:H2'	23:QX:5:A:C8	2.49	0.48
27:RE:48:GLN:HA	27:RE:80:GLU:HA	1.94	0.48
36:RR:33:ARG:HD3	36:RR:113:LEU:HD11	1.95	0.48
37:RS:68:GLN:O	37:RS:72:ALA:N	2.38	0.48
43:RY:11:ASP:OD1	43:RY:11:ASP:N	2.45	0.48
1:XA:1512:U:H2'	1:XA:1513:A:C8	2.49	0.48
24:YA:1042:A:H4'	39:YU:92:ARG:NE	2.28	0.48
24:YA:1825:U:H2'	24:YA:1826:C:H6	1.78	0.48
24:YA:491:G:H2'	24:YA:492:A:C8	2.49	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:YA:937:A:H2'	24:YA:938:G:H8	1.77	0.48
25:YB:50:G:OP1	37:YS:63:THR:OG1	2.24	0.48
31:YI:101:LEU:HG	31:YI:107:VAL:HG13	1.95	0.48
34:YP:1:MET:HG3	34:YP:5:ASP:HB2	1.95	0.48
41:YW:33:ARG:NE	41:YW:52:GLU:OE2	2.44	0.48
1:QA:1226:C:O2'	13:QM:111:LYS:NZ	2.42	0.48
1:QA:380:G:N2	1:QA:383:A:OP2	2.37	0.48
1:XA:1431:C:H2'	1:XA:1432:G:O4'	2.13	0.48
7:XG:47:CYS:O	7:XG:51:GLN:NE2	2.47	0.48
24:YA:1171:G:OP2	24:YA:1172:A:O2'	2.25	0.48
24:YA:1451:U:H2'	24:YA:1452:U:H6	1.79	0.48
24:YA:1825:U:H2'	24:YA:1826:C:C6	2.49	0.48
24:YA:1850:A:H5''	26:YD:161:THR:HG21	1.95	0.48
24:YA:2236:G:OP1	26:YD:268:ARG:NH2	2.45	0.48
16:QP:4:ILE:HA	16:QP:20:VAL:O	2.13	0.48
24:RA:1911:A:H2'	24:RA:1912:A:C8	2.49	0.48
24:RA:2480:G:O2'	24:RA:2493:G:N2	2.44	0.48
24:RA:2804:C:OP1	24:RA:2902:G:N2	2.46	0.48
24:RA:30:G:H2'	24:RA:31:C:C6	2.48	0.48
24:RA:631:A:H2'	24:RA:632:A:C8	2.49	0.48
24:RA:248:G:H21	24:RA:646:A:H8	1.62	0.48
30:RH:64:LEU:O	30:RH:68:THR:OG1	2.24	0.48
1:XA:1512:U:H2'	1:XA:1513:A:H8	1.79	0.48
1:XA:377:G:H2'	1:XA:378:G:H8	1.78	0.48
1:XA:565:U:H5''	1:XA:566:G:H2'	1.96	0.48
24:YA:2405:A:H5'	34:YP:63:PRO:HB3	1.96	0.48
24:YA:142:G:H4'	42:YX:35:THR:HG21	1.95	0.48
1:QA:1032(C):G:N2	1:QA:1033:G:O6	2.47	0.48
1:QA:235:C:H2'	1:QA:236:G:H8	1.78	0.48
1:QA:666:G:H5'	1:QA:726:C:H1'	1.95	0.48
18:QR:22:VAL:HA	18:QR:25:THR:HG23	1.96	0.48
24:RA:2708:U:H2'	24:RA:2709:G:C8	2.49	0.48
24:RA:2701:U:H5'	24:RA:2726:A:C2	2.49	0.48
24:RA:2846:U:H2'	24:RA:2847:G:C8	2.48	0.48
24:RA:956:A:N3	24:RA:2276:C:O2'	2.45	0.48
28:RF:161:GLU:HG2	28:RF:162:LEU:HG	1.96	0.48
1:XA:1277:C:H2'	1:XA:1279:A:H8	1.79	0.48
1:XA:177:C:H2'	1:XA:178:C:H6	1.78	0.48
10:XJ:49:VAL:HG23	14:YN:41:ARG:HB2	1.96	0.48
14:YN:40:CYS:SG	14:YN:41:ARG:N	2.87	0.48
24:YA:1072:U:O2	24:YA:1073:A:H3'	2.14	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:YA:354:A:H2	24:YA:1255:A:H2'	1.78	0.48
24:YA:840:A:OP2	24:YA:2093:A:O2'	2.31	0.48
24:YA:2413:U:OP1	51:Y6:18:ARG:NH2	2.43	0.48
24:YA:628:C:O2	24:YA:704:U:O2'	2.30	0.48
25:YB:17:C:H42	25:YB:109:G:H21	1.60	0.48
27:YE:144:ARG:HB3	27:YE:145:LYS:H	1.48	0.48
1:QA:1222:G:H5'	19:QS:77:THR:HG21	1.95	0.48
1:QA:713:G:H2'	1:QA:714:G:C8	2.49	0.48
1:QA:715:A:H2'	1:QA:716:A:C8	2.49	0.48
1:QA:806:C:H2'	1:QA:807:A:H8	1.78	0.48
10:QJ:13:HIS:O	10:QJ:17:ASP:HB2	2.14	0.48
24:RA:1332:A:O2'	24:RA:1334:U:OP2	2.19	0.48
24:RA:1590:C:HO2'	24:RA:1591:A:P	2.36	0.48
29:RG:144:ILE:HG22	29:RG:146:TYR:H	1.78	0.48
29:RG:29:TRP:O	29:RG:33:ARG:NH1	2.42	0.48
34:RP:62:LEU:O	53:R8:13:ARG:NH1	2.46	0.48
1:XA:373:A:H2'	1:XA:374:A:H8	1.78	0.48
1:XA:985:C:H2'	1:XA:986:A:H8	1.79	0.48
6:XF:50:TYR:OH	18:XR:75:ILE:O	2.32	0.48
24:YA:1581:U:H3	24:YA:1583:C:H1'	1.79	0.48
24:YA:1884:A:N3	24:YA:2245:U:O2'	2.41	0.48
30:YH:80:SER:OG	30:YH:81:GLU:N	2.47	0.48
2:QB:11:LEU:HD23	2:QB:217:ARG:HH22	1.79	0.48
5:QE:51:VAL:HG23	5:QE:52:PRO:HD3	1.95	0.48
3:QC:29:TYR:OH	14:QN:54:PRO:O	2.30	0.48
24:RA:1457:C:H42	24:RA:1637:G:H1	1.61	0.48
43:RY:102:CYS:SG	43:RY:103:GLY:N	2.87	0.48
13:XM:13:LYS:HA	13:XM:44:ARG:HD2	1.94	0.48
53:Y8:28:GLY:O	53:Y8:36:LYS:NZ	2.41	0.48
24:YA:1409:C:O2'	24:YA:1840:A:N3	2.41	0.48
24:YA:2207:C:H2'	24:YA:2208:G:H8	1.79	0.48
24:YA:2840:G:OP1	27:YE:76:ARG:NH2	2.47	0.48
28:YF:143:ALA:HB1	28:YF:148:LEU:HB2	1.96	0.48
41:YW:86:LEU:HD22	41:YW:96:ILE:HD11	1.94	0.48
1:QA:1527:C:O2'	1:QA:1528:U:H5'	2.13	0.47
1:QA:539:A:H2'	1:QA:540:G:C8	2.49	0.47
1:QA:985:C:H2'	1:QA:986:A:H8	1.79	0.47
4:QD:14:ARG:HD2	4:QD:40:PRO:HD2	1.96	0.47
28:RF:185:ASP:OD1	28:RF:188:ARG:NH1	2.45	0.47
31:RI:122:GLU:O	31:RI:126:TYR:OH	2.29	0.47
24:RA:2262:G:C4	35:RQ:82:ARG:HG3	2.49	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:RT:62:THR:HG22	38:RT:75:ILE:HG23	1.96	0.47
44:RZ:163:LEU:HD13	44:RZ:167:PRO:HD3	1.95	0.47
1:XA:309:G:H2'	1:XA:310:G:H8	1.79	0.47
1:XA:398:C:H2'	1:XA:399:G:C8	2.49	0.47
24:YA:1462:G:H2'	24:YA:1463:C:C6	2.49	0.47
24:YA:2855:G:H2'	24:YA:2856:G:C8	2.49	0.47
24:YA:2327:G:OP1	29:YG:36:LYS:NZ	2.46	0.47
38:YT:16:ARG:NH1	38:YT:18:ASP:OD2	2.47	0.47
40:YV:49:THR:O	40:YV:49:THR:OG1	2.31	0.47
1:QA:281:G:OP2	1:QA:281:G:H8	1.97	0.47
1:QA:34:C:H2'	1:QA:35:G:H8	1.79	0.47
1:QA:580:U:H2'	1:QA:581:G:O4'	2.14	0.47
3:QC:47:LEU:HB3	3:QC:50:ALA:HB3	1.95	0.47
17:QQ:3:LYS:HB3	17:QQ:61:GLU:HB3	1.96	0.47
24:RA:1411:A:O2'	46:R1:11:ARG:NH2	2.45	0.47
24:RA:1119:A:O2'	24:RA:1120:G:O5'	2.30	0.47
24:RA:1414:G:OP1	52:R7:28:ARG:NH2	2.45	0.47
24:RA:2065:C:OP1	24:RA:2790:G:O2'	2.22	0.47
24:RA:2172:U:H2'	24:RA:2173:G:C8	2.49	0.47
26:RD:147:LEU:HD23	26:RD:148:GLU:HG3	1.96	0.47
28:RF:153:SER:OG	28:RF:190:GLU:N	2.46	0.47
30:RH:98:LEU:HD23	30:RH:125:VAL:HG23	1.96	0.47
40:RV:6:LYS:HB2	40:RV:38:LEU:HD11	1.96	0.47
1:XA:1422:G:H5''	33:YO:48:PRO:HB3	1.96	0.47
9:XI:72:GLY:O	9:XI:76:ALA:N	2.42	0.47
11:XK:91:ARG:NH2	11:XK:110:ASP:OD1	2.34	0.47
24:YA:2173:G:H2'	24:YA:2174:G:H8	1.79	0.47
1:QA:304:U:H2'	1:QA:305:G:C8	2.50	0.47
22:QV:53:G:O2'	22:QV:54:U:O5'	2.29	0.47
24:RA:2667:G:N2	24:RA:2677:A:OP2	2.47	0.47
30:RH:70:THR:HG22	30:RH:74:ASN:HD21	1.78	0.47
34:RP:37:GLY:O	34:RP:40:SER:OG	2.29	0.47
44:RZ:10:ARG:HH22	44:RZ:26:GLY:H	1.61	0.47
1:XA:1255:G:O2'	1:XA:1258:G:N3	2.36	0.47
1:XA:22:G:H4'	1:XA:885:G:C8	2.49	0.47
1:XA:483:C:OP2	1:XA:484:G:O2'	2.20	0.47
3:XC:118:GLN:HG2	3:XC:187:ALA:HB2	1.96	0.47
18:XR:32:ARG:HA	18:XR:69:THR:HG21	1.95	0.47
24:YA:1581:U:H2'	24:YA:1582:A:N7	2.29	0.47
24:YA:1843:A:H2'	24:YA:1844:G:H8	1.79	0.47
24:YA:268:G:H2'	24:YA:269:G:C8	2.48	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:YA:604:C:H2'	24:YA:605:G:C8	2.49	0.47
24:YA:612:C:H2'	24:YA:613:A:C8	2.49	0.47
24:YA:863:C:O2'	24:YA:977:G:O6	2.30	0.47
30:YH:159:GLU:HG2	30:YH:169:VAL:HG11	1.96	0.47
33:YO:120:GLU:OE1	38:YT:67:SER:OG	2.26	0.47
37:YS:34:HIS:ND1	37:YS:53:SER:OG	2.45	0.47
1:QA:1175:G:H2'	1:QA:1176:A:H8	1.78	0.47
5:QE:35:GLY:H	5:QE:112:LEU:HD23	1.79	0.47
9:QL:28:VAL:HG12	9:QL:63:ILE:HB	1.94	0.47
14:QN:37:PHE:HD2	14:QN:44:LEU:HD22	1.78	0.47
52:R7:30:VAL:HG22	52:R7:33:ARG:HH21	1.79	0.47
24:RA:1514:C:H5	24:RA:1593:C:H2'	1.80	0.47
24:RA:231:G:O2'	24:RA:243:G:O6	2.27	0.47
24:RA:2863:C:H2'	24:RA:2864:G:C8	2.49	0.47
24:RA:484:G:O2'	24:RA:495:G:O6	2.26	0.47
24:RA:907:U:H2'	24:RA:908:A:C8	2.49	0.47
24:RA:776:G:OP2	26:RD:13:ARG:NH1	2.47	0.47
38:RT:124:ASP:O	38:RT:128:GLU:N	2.47	0.47
38:RT:50:ILE:HG13	38:RT:99:LEU:HD12	1.96	0.47
24:RA:1644:C:O3'	42:RX:35:THR:OG1	2.32	0.47
1:XA:359:U:H6	1:XA:359:U:OP2	1.96	0.47
2:XB:67:THR:HA	2:XB:90:MET:HE1	1.95	0.47
3:XC:25:GLY:O	3:XC:29:TYR:N	2.48	0.47
24:YA:1587:U:H2'	24:YA:1588:G:O4'	2.15	0.47
24:YA:1921:G:H21	24:YA:1924:C:N4	2.13	0.47
24:YA:2687:A:H5'	33:YO:31:LYS:HE3	1.96	0.47
1:QA:1010:G:H2'	1:QA:1011:G:H8	1.79	0.47
1:QA:1315:U:H2'	1:QA:1316:G:O4'	2.15	0.47
1:QA:975:A:N1	10:QJ:48:THR:OG1	2.46	0.47
3:QC:18:TRP:HB3	3:QC:20:SER:H	1.79	0.47
24:RA:2624:C:OP2	50:R5:2:ALA:N	2.48	0.47
24:RA:2705:A:H2'	24:RA:2706:G:H8	1.79	0.47
24:RA:320:C:O3'	43:RY:95:LYS:NZ	2.47	0.47
37:RS:67:ARG:O	37:RS:71:ARG:N	2.44	0.47
1:XA:1178:G:N1	1:XA:1181:G:N7	2.62	0.47
1:XA:129(B):G:N2	1:XA:188:U:HO2'	2.12	0.47
1:XA:422:C:H4'	1:XA:423:G:C4	2.49	0.47
16:XP:69:THR:OG1	16:XP:69:THR:O	2.31	0.47
24:YA:240:A:C5	24:YA:241:G:H1'	2.49	0.47
27:YE:36:ARG:HG2	27:YE:47:VAL:HG12	1.95	0.47
28:YF:149:ASP:OD1	28:YF:149:ASP:N	2.45	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:56:U:H2'	1:QA:57:G:C8	2.49	0.47
13:QM:44:ARG:HG3	13:QM:46:LYS:HZ3	1.78	0.47
53:R8:22:VAL:HB	53:R8:53:PRO:HB3	1.96	0.47
24:RA:1174:A:N7	24:RA:2501:G:O2'	2.47	0.47
24:RA:166:G:H3'	24:RA:167:G:H8	1.80	0.47
24:RA:1311:A:H61	24:RA:2035:A:H5''	1.78	0.47
24:RA:2044:U:O2'	24:RA:2629:C:H5'	2.15	0.47
24:RA:968:U:H2'	24:RA:969:C:C6	2.49	0.47
29:RG:150:ASP:N	29:RG:150:ASP:OD1	2.39	0.47
36:RR:75:LEU:HA	36:RR:78:LYS:HB3	1.95	0.47
38:RT:24:PRO:HA	38:RT:49:VAL:HG13	1.96	0.47
1:XA:412:A:H4'	1:XA:413:G:O5'	2.15	0.47
1:XA:991:U:O2'	1:XA:992:U:O5'	2.30	0.47
22:XV:17:C:H5'	22:XV:61:C:OP1	2.15	0.47
24:YA:2129:C:H2'	24:YA:2130:C:O4'	2.15	0.47
25:YB:85:G:H22	25:YB:92:G:H22	1.61	0.47
25:YB:85:G:H22	25:YB:92:G:N2	2.12	0.47
29:YG:7:LEU:N	29:YG:104:GLU:OE2	2.46	0.47
1:QA:299:G:H2'	1:QA:300:A:C8	2.50	0.47
1:QA:375:U:O2	16:QP:28:ARG:NH1	2.48	0.47
53:R8:37:SER:OG	53:R8:38:GLY:N	2.48	0.47
24:RA:1462:G:H2'	24:RA:1463:C:C6	2.49	0.47
1:QA:1484:C:HO2'	24:RA:1982:A:HO2'	1.63	0.47
24:RA:2555:G:H21	24:RA:2658:C:H5''	1.79	0.47
26:RD:108:PRO:HB3	26:RD:143:HIS:CE1	2.50	0.47
29:RG:109:VAL:HG11	29:RG:142:PRO:HB3	1.96	0.47
29:RG:59:GLU:OE1	29:RG:153:ARG:NH2	2.47	0.47
1:XA:269:C:H2'	1:XA:270:A:H8	1.79	0.47
24:YA:2467:G:H2'	24:YA:2468:C:C6	2.49	0.47
24:YA:390:G:H2'	24:YA:391:G:C8	2.49	0.47
24:YA:528:A:H1'	24:YA:529:U:OP2	2.14	0.47
24:YA:602:G:H2'	24:YA:603:C:C6	2.49	0.47
24:YA:965:G:N2	24:YA:2281:A:OP2	2.43	0.47
38:YT:123:GLN:O	38:YT:125:ARG:N	2.48	0.47
1:QA:1133:G:N2	1:QA:1141:C:N3	2.54	0.47
2:QB:115:LEU:HD22	2:QB:145:LEU:HB3	1.97	0.47
2:QB:192:SER:OG	2:QB:193:ASP:OD1	2.30	0.47
24:RA:1103:A:HO2'	24:RA:1104:G:P	2.35	0.47
24:RA:2156:A:N7	24:RA:2179:G:O2'	2.47	0.47
33:RO:64:ARG:HE	33:RO:83:ALA:HB3	1.80	0.47
1:XA:1003:G:H21	1:XA:1005:A:H5'	1.80	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1264:C:H2'	1:XA:1265:G:C8	2.50	0.47
1:XA:600:C:H5'	8:XH:129:VAL:O	2.15	0.47
1:XA:971:G:N1	1:XA:1364:U:O2'	2.48	0.47
24:YA:1399:A:H2'	24:YA:1400:A:C8	2.50	0.47
24:YA:992:G:H2'	24:YA:993:G:C8	2.49	0.47
1:QA:412:A:H4'	1:QA:413:G:O5'	2.15	0.47
3:QC:24:ALA:HB2	3:QC:32:LEU:HD12	1.97	0.47
24:RA:1321:A:N1	24:RA:1341:C:O2'	2.41	0.47
24:RA:1487:G:H2'	24:RA:1488:G:H8	1.79	0.47
24:RA:1514:C:C5	24:RA:1593:C:H2'	2.50	0.47
24:RA:2124:U:C2	24:RA:2125:C:H5	2.32	0.47
24:RA:776:G:O2'	24:RA:810:G:H4'	2.15	0.47
26:RD:85:ASP:OD2	26:RD:88:ARG:NH1	2.45	0.47
42:RX:55:ASN:HB2	42:RX:80:ILE:HG23	1.96	0.47
1:XA:1096:C:H2'	1:XA:1097:C:H6	1.79	0.47
1:XA:115:G:H4'	1:XA:116:A:O5'	2.15	0.47
1:XA:320:C:H2'	1:XA:321:A:C8	2.50	0.47
1:XA:372:C:H42	1:XA:389:A:N6	2.08	0.47
7:XG:15:ASP:H	7:XG:20:ASP:H	1.62	0.47
10:XJ:78:ASN:O	10:XJ:81:THR:OG1	2.24	0.47
13:XM:23:TYR:HB3	13:XM:67:GLU:HG3	1.97	0.47
17:XQ:83:ASP:N	17:XQ:83:ASP:OD1	2.41	0.47
24:YA:1355:G:HO2'	24:YA:1657:C:HO2'	1.60	0.47
26:YD:4:LYS:HB3	26:YD:18:VAL:HG23	1.97	0.47
1:QA:958:A:H2'	1:QA:959:A:C8	2.50	0.47
6:QF:72:VAL:O	6:QF:75:LEU:HB3	2.15	0.47
9:QI:83:ARG:HG2	9:QI:102:LEU:HG	1.97	0.47
12:QL:77:LEU:HD23	12:QL:77:LEU:HA	1.81	0.47
24:RA:2096:U:H2'	24:RA:2097:U:C6	2.50	0.47
24:RA:640:A:C4	28:RF:180:GLY:HA3	2.50	0.47
1:XA:1006:C:H2'	1:XA:1007:C:C6	2.50	0.47
1:XA:464:G:C6	1:XA:466:C:H5'	2.50	0.47
1:XA:627:G:H2'	1:XA:628:G:C8	2.49	0.47
12:XL:103:GLY:HA2	12:XL:108:ALA:HA	1.97	0.47
46:Y1:83:GLU:HG2	46:Y1:85:LEU:H	1.80	0.47
33:YO:88:ASN:HD21	33:YO:92:GLU:HB2	1.79	0.47
1:QA:1221:G:OP1	1:QA:1321:C:N4	2.41	0.47
1:QA:1432:G:OP1	38:RT:108:ARG:N	2.37	0.47
24:RA:1405:A:H2'	24:RA:1406:A:H5'	1.96	0.47
24:RA:2248:C:H2'	24:RA:2249:G:O4'	2.15	0.47
28:RF:178:PRO:HB3	28:RF:198:ALA:HB1	1.96	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:184:G:H2'	1:XA:185:A:H8	1.80	0.47
1:XA:583:A:O2'	17:XQ:91:ARG:NE	2.39	0.47
1:XA:628:G:H2'	1:XA:629:G:C8	2.50	0.47
20:XT:75:ASN:OD1	20:XT:75:ASN:N	2.40	0.47
1:XA:1230:C:H5'	22:XV:30:C:H5''	1.97	0.47
48:Y3:10:LYS:NZ	48:Y3:15:TYR:OH	2.48	0.47
24:YA:746:A:N3	24:YA:1680:G:O2'	2.42	0.47
34:YP:46:LYS:HB3	34:YP:46:LYS:HE3	1.71	0.47
1:QA:768:A:H4'	1:QA:1523:G:N2	2.30	0.46
2:QB:178:ARG:NH2	2:QB:196:LEU:O	2.48	0.46
17:QQ:79:SER:OG	17:QQ:80:GLY:N	2.48	0.46
46:R1:40:ARG:HH12	46:R1:42:GLN:HE21	1.62	0.46
24:RA:2855:G:H2'	24:RA:2856:G:C8	2.50	0.46
24:RA:399:G:O2'	24:RA:400:U:O5'	2.33	0.46
24:RA:2687:A:H5'	33:RO:31:LYS:HE2	1.96	0.46
1:XA:1238:A:N6	1:XA:1301:U:C2	2.82	0.46
1:XA:1499:A:H1'	1:XA:1520:G:H5'	1.97	0.46
2:XB:172:ILE:HG13	2:XB:172:ILE:H	1.53	0.46
24:YA:1362:U:H2'	24:YA:1363:A:C8	2.50	0.46
24:YA:2108:U:H2'	24:YA:2109:G:C8	2.50	0.46
24:YA:829:A:O2'	26:YD:225:ALA:O	2.30	0.46
38:YT:51:ARG:HB2	38:YT:98:LYS:HG3	1.97	0.46
1:QA:673:G:H2'	1:QA:674:G:H8	1.78	0.46
1:QA:963:G:H21	10:QJ:55:LYS:HG2	1.81	0.46
24:RA:1451:U:H2'	24:RA:1452:U:H6	1.79	0.46
24:RA:911:G:H21	24:RA:913:A:N6	2.12	0.46
28:RF:170:LEU:HD22	28:RF:172:TRP:HE1	1.80	0.46
25:RB:55:U:O3'	29:RG:27:ASN:ND2	2.48	0.46
24:RA:998:A:OP2	35:RQ:16:ARG:HD3	2.16	0.46
44:RZ:104:PHE:HA	44:RZ:139:VAL:HG13	1.97	0.46
1:XA:1229:A:O2'	22:XV:30:C:OP1	2.34	0.46
1:XA:21:G:H2'	1:XA:22:G:C8	2.49	0.46
1:XA:328:C:H4'	1:XA:329:A:O5'	2.16	0.46
1:XA:360:A:H2'	1:XA:361:G:C8	2.50	0.46
1:XA:45:U:H2'	1:XA:46:G:C8	2.51	0.46
6:XF:24:GLU:HA	6:XF:27:GLN:HG2	1.96	0.46
24:YA:1177:G:HO2'	24:YA:1178:A:H8	1.61	0.46
24:YA:2136:A:N1	24:YA:2141:A:N6	2.63	0.46
24:YA:2863:C:H2'	24:YA:2864:G:H8	1.80	0.46
24:YA:389:G:H2'	24:YA:390:G:H8	1.79	0.46
24:YA:968:U:H2'	24:YA:969:C:C6	2.50	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1141:C:H2'	1:QA:1142:G:C8	2.51	0.46
1:QA:1066:C:N4	1:QA:1191:A:N6	2.63	0.46
1:QA:769:G:H4'	1:QA:1513:A:H4'	1.98	0.46
1:QA:436:C:H2'	1:QA:437:U:C6	2.51	0.46
46:R1:73:LEU:HB3	46:R1:90:ILE:HG12	1.96	0.46
48:R3:3:ARG:HD3	48:R3:60:GLU:O	2.14	0.46
24:RA:48:A:H61	24:RA:166:G:H2'	1.80	0.46
24:RA:2294:G:H4'	24:RA:2401:G:O2'	2.14	0.46
24:RA:528:A:H4'	24:RA:529:U:H5'	1.97	0.46
28:RF:11:VAL:HG22	28:RF:125:LEU:HD12	1.97	0.46
33:RO:1:MET:HE3	33:RO:67:LYS:HG2	1.98	0.46
24:RA:561:A:OP1	39:RU:53:ARG:NH1	2.49	0.46
1:XA:1038:C:H2'	1:XA:1039:C:C6	2.50	0.46
9:XI:71:SER:HA	9:XI:74:ILE:HG13	1.98	0.46
24:YA:1108:G:H1	24:YA:1122:C:H42	1.63	0.46
24:YA:2156:A:HO2'	24:YA:2181:G:H21	1.58	0.46
26:YD:26:LYS:NZ	26:YD:83:GLU:OE2	2.36	0.46
1:QA:1043:C:H2'	1:QA:1044:A:H8	1.80	0.46
1:QA:1077:G:O6	5:QE:47:LYS:NZ	2.48	0.46
1:QA:277:C:H5''	17:QQ:68:ARG:HH21	1.81	0.46
1:QA:426:G:OP1	4:QD:36:ARG:NH1	2.44	0.46
1:QA:41:G:H2'	1:QA:42:G:C8	2.49	0.46
1:QA:600:C:H2'	1:QA:601:C:H6	1.81	0.46
1:QA:662:G:O2'	1:QA:836:G:OP1	2.32	0.46
12:QL:24:VAL:HG13	12:QL:98:TYR:HE1	1.80	0.46
14:QN:47:LEU:O	14:QN:51:GLY:N	2.49	0.46
16:QP:43:LYS:HG2	16:QP:48:TRP:CD2	2.50	0.46
24:RA:173:C:H2'	24:RA:174:U:C6	2.49	0.46
24:RA:2401:G:H5''	24:RA:2402:U:O4'	2.16	0.46
25:RB:22:U:O4	25:RB:61:G:O6	2.33	0.46
28:RF:122:LYS:HB3	28:RF:191:ARG:HA	1.98	0.46
1:XA:328:C:H1'	1:XA:329:A:OP2	2.15	0.46
1:XA:509:A:N3	1:XA:543:C:O2'	2.42	0.46
1:XA:78:G:O2'	1:XA:79:G:OP1	2.27	0.46
2:XB:192:SER:OG	2:XB:193:ASP:N	2.48	0.46
3:XC:136:GLN:O	3:XC:140:ARG:N	2.35	0.46
13:XM:52:GLU:HG2	13:XM:55:ARG:HH21	1.80	0.46
13:XM:69:GLU:O	13:XM:73:GLU:N	2.41	0.46
20:XT:74:LYS:HD3	20:XT:74:LYS:HA	1.76	0.46
24:YA:199:C:OP2	52:Y7:29:LYS:NZ	2.49	0.46
24:YA:1046:A:H2'	24:YA:1047:A:C8	2.50	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:YA:2096:U:H2'	24:YA:2097:U:C6	2.51	0.46
31:YI:45:LYS:HA	31:YI:45:LYS:HD2	1.78	0.46
1:QA:1187:G:N3	14:QN:60:SER:OG	2.49	0.46
1:QA:1285:A:H1'	1:QA:1286:A:OP2	2.16	0.46
1:QA:1347:G:N2	1:QA:1374:A:OP2	2.35	0.46
3:QC:88:ARG:HH12	3:QC:101:LEU:N	2.13	0.46
9:QI:104:ARG:NH1	9:QI:105:ASP:O	2.48	0.46
1:QA:522:C:H41	12:QL:53:ARG:HH22	1.64	0.46
13:QM:54:VAL:HG23	13:QM:57:ARG:HH21	1.80	0.46
24:RA:1585:G:H2'	24:RA:1586:G:H8	1.81	0.46
24:RA:1880:G:H2'	24:RA:1881:G:H8	1.79	0.46
24:RA:2137:G:OP1	24:RA:2188:G:O2'	2.34	0.46
29:RG:2:PRO:HB2	29:RG:3:LEU:HB2	1.97	0.46
1:XA:1264:C:H2'	1:XA:1265:G:H8	1.80	0.46
1:XA:767:A:O2'	1:XA:1524:C:O2	2.31	0.46
1:XA:325:A:OP2	20:XT:70:SER:OG	2.19	0.46
1:XA:789:U:H1'	1:XA:792:A:H2	1.80	0.46
3:XC:42:LEU:HA	3:XC:45:LYS:HB3	1.96	0.46
4:XD:148:VAL:HG12	4:XD:152:SER:HG	1.79	0.46
9:XI:25:LYS:HE3	9:XI:25:LYS:HB2	1.69	0.46
24:YA:831:A:N6	24:YA:2094:G:O2'	2.48	0.46
24:YA:570:C:H3'	24:YA:571:A:C8	2.50	0.46
29:YG:11:TYR:HA	29:YG:15:VAL:HB	1.97	0.46
1:QA:1218:C:H2'	1:QA:1219:U:C6	2.50	0.46
1:QA:1314:C:H2'	1:QA:1315:U:C6	2.51	0.46
1:QA:587:G:N2	1:QA:754:C:OP2	2.48	0.46
12:QL:117:ARG:HB2	12:QL:122:THR:HB	1.96	0.46
24:RA:656:A:OP2	53:R8:46:ARG:NH2	2.48	0.46
24:RA:1258:A:N3	24:RA:1284:G:O2'	2.46	0.46
24:RA:1585:G:H2'	24:RA:1586:G:C8	2.51	0.46
24:RA:2326:C:H2'	24:RA:2327:G:H8	1.81	0.46
24:RA:2555:G:H2'	24:RA:2556:G:C8	2.50	0.46
24:RA:312:C:H2'	24:RA:313:A:H8	1.80	0.46
34:RP:89:ALA:O	34:RP:121:LYS:NZ	2.47	0.46
37:RS:100:ALA:HA	37:RS:103:GLU:HB2	1.98	0.46
1:XA:437:U:H3	1:XA:495:A:H62	1.62	0.46
1:XA:737:A:H2'	1:XA:738:C:C6	2.51	0.46
4:XD:127:THR:HA	4:XD:132:ARG:HA	1.97	0.46
8:XH:49:GLU:HG2	8:XH:62:TYR:HE2	1.79	0.46
10:XJ:54:PHE:HB3	10:XJ:55:LYS:HD2	1.98	0.46
24:YA:902:G:O2'	45:Y0:27:GLU:OE2	2.26	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:YA:1112:U:H5'	24:YA:1113:A:OP2	2.15	0.46
24:YA:2045:G:H5'	24:YA:2629:C:H4'	1.98	0.46
24:YA:321:C:H2'	24:YA:322:G:O4'	2.16	0.46
24:YA:510:C:H2'	24:YA:511:C:H6	1.80	0.46
26:YD:254:THR:O	26:YD:254:THR:OG1	2.34	0.46
40:YV:68:LYS:HD3	40:YV:68:LYS:HA	1.73	0.46
41:YW:10:VAL:HG12	41:YW:12:ILE:HG22	1.97	0.46
41:YW:35:ILE:O	41:YW:39:THR:OG1	2.29	0.46
42:YX:25:LYS:NZ	42:YX:82:GLN:OE1	2.39	0.46
1:QA:1199:U:O2'	1:QA:1202:G:OP1	2.33	0.46
1:QA:35:G:N3	12:QL:118:SER:OG	2.45	0.46
1:QA:560:U:H4'	1:QA:561:U:H5''	1.98	0.46
1:QA:743:U:H2'	1:QA:744:C:C6	2.51	0.46
24:RA:1311:A:H8	24:RA:1311:A:OP1	1.98	0.46
24:RA:1362:U:H2'	24:RA:1363:A:C8	2.51	0.46
24:RA:1530:G:H2'	24:RA:1531:G:H8	1.81	0.46
24:RA:1728:G:HO2'	24:RA:1793:A:HO2'	1.62	0.46
24:RA:2209:G:H2'	24:RA:2210:C:C6	2.51	0.46
28:RF:93:LYS:HA	28:RF:93:LYS:HD3	1.78	0.46
30:RH:4:ILE:HG22	30:RH:6:ARG:HG2	1.96	0.46
32:RN:6:PRO:HG3	32:RN:41:ASP:HB2	1.96	0.46
35:RQ:63:LYS:HD2	44:RZ:175:VAL:HG21	1.97	0.46
1:XA:564:C:OP2	12:XL:15:ARG:NH2	2.27	0.46
1:XA:688:G:O2'	1:XA:704:A:N1	2.37	0.46
1:XA:950:U:H2'	1:XA:951:G:C8	2.51	0.46
1:XA:985:C:H2'	1:XA:986:A:C8	2.51	0.46
10:XJ:84:GLN:HB2	10:XJ:85:LEU:HD12	1.97	0.46
24:YA:1211:U:H3	24:YA:1229:G:H1	1.63	0.46
24:YA:2549:U:H2'	24:YA:2550:C:C6	2.50	0.46
24:YA:998:A:OP1	35:YQ:18:LYS:NZ	2.38	0.46
28:YF:30:PRO:HB3	34:YP:1:MET:HE1	1.98	0.46
31:YI:62:LYS:HE3	31:YI:133:HIS:CE1	2.50	0.46
33:YO:106:LEU:O	33:YO:110:GLY:N	2.47	0.46
44:YZ:123:ASP:N	44:YZ:123:ASP:OD1	2.46	0.46
44:YZ:151:HIS:HA	44:YZ:170:THR:HA	1.96	0.46
1:QA:1479:C:H2'	1:QA:1480:G:H8	1.81	0.46
24:RA:1148:C:H2'	24:RA:1149:A:H8	1.81	0.46
24:RA:2140:U:O2	24:RA:2170:G:O2'	2.31	0.46
24:RA:2165:C:N4	24:RA:2171:G:O6	2.49	0.46
31:RI:129:THR:OG1	31:RI:129:THR:O	2.31	0.46
24:RA:9:U:OP1	32:RN:115:ARG:NH2	2.49	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:RV:68:LYS:HA	40:RV:68:LYS:HD2	1.67	0.46
20:XT:105:SER:O	20:XT:105:SER:OG	2.34	0.46
23:XX:2:G:H2'	23:XX:3:C:C2	2.51	0.46
46:Y1:86:SER:O	46:Y1:89:GLU:N	2.49	0.46
48:Y3:12:PRO:HA	48:Y3:15:TYR:HD2	1.81	0.46
24:YA:47:G:N1	24:YA:166:G:OP2	2.47	0.46
24:YA:1890:A:N6	24:YA:1905:G:HO2'	2.14	0.46
28:YF:40:GLN:HE22	28:YF:182:ASN:HD22	1.63	0.46
32:YN:30:ILE:HG23	32:YN:52:VAL:HG11	1.97	0.46
1:QA:328:C:H4'	1:QA:329:A:H5'	1.98	0.46
1:QA:757:U:O2'	1:QA:879:C:O2	2.34	0.46
6:QF:30:LEU:HB3	6:QF:35:ALA:HB3	1.98	0.46
10:QJ:84:GLN:HG3	10:QJ:85:LEU:HD12	1.98	0.46
10:QJ:7:LYS:HB3	10:QJ:97:GLU:HB2	1.96	0.46
11:QK:85:ARG:HE	11:QK:111:ASP:HB3	1.80	0.46
47:R2:37:PHE:O	47:R2:40:SER:OG	2.32	0.46
42:RX:53:LYS:HB3	42:RX:82:GLN:HB3	1.97	0.46
1:XA:1321:C:H5''	1:XA:1322:C:H2'	1.98	0.46
1:XA:448:A:OP2	1:XA:485:G:N2	2.34	0.46
2:XB:138:LEU:HA	2:XB:141:GLU:HB2	1.98	0.46
9:XI:75:ASP:OD2	9:XI:78:LYS:NZ	2.45	0.46
12:XL:102:ARG:HB3	12:XL:109:GLY:HA2	1.98	0.46
22:XV:23:C:H2'	22:XV:24:G:H8	1.80	0.46
46:Y1:18:ILE:HG12	46:Y1:37:ILE:HG12	1.97	0.46
24:YA:2207:C:H2'	24:YA:2208:G:C8	2.51	0.46
24:YA:2371:C:H2'	24:YA:2372:A:O4'	2.15	0.46
25:YB:40:U:N3	25:YB:44:G:OP2	2.29	0.46
39:YU:92:ARG:HD2	40:YV:11:GLN:HB2	1.97	0.46
1:QA:107:G:N7	20:QT:15:ARG:NH2	2.64	0.46
1:QA:243:A:H4'	1:QA:244:U:H3'	1.98	0.46
1:QA:370:C:H2'	1:QA:371:G:H8	1.80	0.46
1:QA:444:C:H2'	1:QA:445:G:H8	1.81	0.46
2:QB:71:VAL:HA	2:QB:93:VAL:HB	1.98	0.46
1:QA:407:G:H5''	4:QD:115:ARG:HB3	1.98	0.46
7:QG:42:ILE:HA	7:QG:115:ARG:HH21	1.81	0.46
11:QK:33:THR:OG1	11:QK:34:ASP:OD1	2.33	0.46
1:QA:1305:G:OP1	21:QU:2:GLY:N	2.48	0.46
24:RA:2407:C:O2'	46:R1:30:VAL:O	2.33	0.46
24:RA:2760:G:N2	24:RA:2770:A:H62	2.06	0.46
24:RA:344:A:H4'	24:RA:346:A:N7	2.31	0.46
24:RA:911:G:H2'	24:RA:912:C:C6	2.51	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:RT:12:SER:HB3	38:RT:15:VAL:HG13	1.97	0.46
1:XA:1314:C:H2'	1:XA:1315:U:C6	2.51	0.46
1:XA:444:C:H2'	1:XA:445:G:H8	1.80	0.46
1:XA:452:A:H2'	1:XA:453:A:H8	1.80	0.46
1:XA:972:C:H5'	10:XJ:57:LYS:HZ2	1.81	0.46
24:YA:1072:U:H1'	24:YA:1073:A:O5'	2.16	0.46
24:YA:1472:G:N7	26:YD:31:LYS:NZ	2.62	0.46
26:YD:152:GLY:O	26:YD:154:LYS:HG2	2.16	0.46
1:QA:1127:G:H1	1:QA:1144:G:H1	1.64	0.45
1:QA:1422:G:H2'	1:QA:1423:G:H8	1.81	0.45
1:QA:493:G:N2	1:QA:494:U:O4	2.49	0.45
1:QA:564:C:OP2	12:QL:15:ARG:NH2	2.36	0.45
10:QJ:22:LYS:HB3	10:QJ:22:LYS:HE2	1.74	0.45
24:RA:1068:G:P	32:RN:69:GLN:HE22	2.39	0.45
24:RA:123:G:OP1	52:R7:14:LYS:NZ	2.36	0.45
29:RG:57:ALA:O	29:RG:61:ALA:N	2.41	0.45
35:RQ:116:GLU:OE2	35:RQ:119:ARG:NH2	2.47	0.45
1:XA:1203:C:H2'	1:XA:1204:A:H8	1.81	0.45
1:XA:939:G:H2'	1:XA:940:C:C6	2.51	0.45
1:XA:992:U:C2	1:XA:1044:A:N6	2.84	0.45
24:YA:1574:A:OP2	24:YA:1588:G:N1	2.49	0.45
1:QA:1354:C:H2'	1:QA:1355:G:H8	1.81	0.45
1:QA:384:G:H2'	1:QA:385:C:H6	1.81	0.45
1:QA:736:C:H2'	1:QA:737:A:H8	1.81	0.45
2:QB:168:THR:HB	2:QB:192:SER:HB2	1.98	0.45
4:QD:155:LEU:HD23	4:QD:155:LEU:HA	1.78	0.45
1:QA:1071:C:H5''	5:QE:49:PRO:HG2	1.97	0.45
7:QG:123:GLU:OE1	7:QG:133:GLY:N	2.44	0.45
14:QN:53:LEU:HA	14:QN:53:LEU:HD12	1.86	0.45
24:RA:106:U:H2'	24:RA:107:G:H8	1.81	0.45
24:RA:1896:G:H5'	24:RA:1897:C:OP2	2.17	0.45
24:RA:854:U:O2'	24:RA:2082:A:N1	2.41	0.45
24:RA:2326:C:H2'	24:RA:2327:G:C8	2.51	0.45
24:RA:2299:A:N6	24:RA:2356:U:H3	2.13	0.45
30:RH:26:VAL:HG21	30:RH:75:ALA:HB1	1.97	0.45
31:RI:77:LEU:HG	31:RI:140:LEU:HB2	1.97	0.45
31:RI:83:ALA:O	31:RI:89:TYR:CE2	2.69	0.45
44:RZ:79:ARG:HB2	44:RZ:80:ARG:HG3	1.98	0.45
1:XA:410:G:H2'	1:XA:429:U:C5	2.51	0.45
2:XB:69:LEU:HD12	2:XB:70:PHE:H	1.80	0.45
4:XD:147:ALA:HA	4:XD:182:LYS:HA	1.98	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:750:G:O2'	15:XO:21:ASP:OD1	2.34	0.45
20:XT:16:HIS:O	20:XT:19:SER:OG	2.31	0.45
24:YA:2432:C:OP1	53:Y8:34:TRP:HB3	2.14	0.45
24:YA:2589:A:O4'	50:Y5:3:LYS:HB2	2.16	0.45
24:YA:36:G:N3	24:YA:476:G:O2'	2.47	0.45
24:YA:599:U:H2'	24:YA:600:G:C8	2.51	0.45
24:YA:776:G:O2'	24:YA:810:G:H4'	2.17	0.45
25:YB:85:G:N2	25:YB:92:G:H22	2.14	0.45
29:YG:139:LEU:HD12	29:YG:144:ILE:HG23	1.98	0.45
37:YS:27:SER:HA	37:YS:88:ASP:HB3	1.99	0.45
1:QA:1007:C:H2'	1:QA:1008:C:C6	2.51	0.45
1:QA:853:G:H2'	1:QA:854:G:H8	1.81	0.45
1:QA:779:C:H5''	11:QK:122:LYS:HD3	1.97	0.45
13:QM:80:ARG:NH2	19:QS:65:ASN:O	2.50	0.45
23:QX:5:A:H2'	23:QX:6:G:H8	1.80	0.45
24:RA:1110:C:H3'	24:RA:1111:U:H5'	1.97	0.45
24:RA:1122:C:H4'	24:RA:1123:A:OP1	2.16	0.45
24:RA:230:A:OP1	24:RA:232:U:H1'	2.16	0.45
24:RA:2699:U:H2'	24:RA:2700:U:O4'	2.16	0.45
24:RA:2701:U:H4'	24:RA:2702:C:O5'	2.16	0.45
27:RE:134:ILE:HA	27:RE:137:HIS:CD2	2.51	0.45
33:RO:10:VAL:HG11	33:RO:16:ALA:HB3	1.99	0.45
38:RT:107:ASP:OD1	38:RT:107:ASP:N	2.48	0.45
1:XA:1074:G:O2'	1:XA:1101:A:N1	2.46	0.45
1:XA:1221:G:OP1	1:XA:1321:C:N4	2.33	0.45
24:YA:1116:A:N6	24:YA:1143:U:O2'	2.49	0.45
24:YA:1359:U:H3'	24:YA:1359:U:O2	2.16	0.45
24:YA:1491:A:H4'	24:YA:1507:A:H2'	1.97	0.45
24:YA:1468:G:H4'	24:YA:1539:C:OP1	2.15	0.45
24:YA:1833:A:H2'	24:YA:1834:A:C8	2.51	0.45
24:YA:18:C:O2'	24:YA:577:U:OP1	2.30	0.45
1:QA:382:A:H2'	1:QA:383:A:C8	2.52	0.45
10:QJ:81:THR:OG1	10:QJ:82:ILE:N	2.50	0.45
24:RA:138:G:H22	24:RA:1642:A:H4'	1.82	0.45
24:RA:263:C:H2'	24:RA:264:G:C8	2.52	0.45
24:RA:880:U:O2	34:RP:55:ARG:NH2	2.38	0.45
26:RD:144:ALA:HB3	26:RD:192:THR:HB	1.98	0.45
25:RB:33:G:H5'	29:RG:2:PRO:HG3	1.98	0.45
39:RU:50:ARG:O	39:RU:54:LYS:NZ	2.34	0.45
1:XA:1014:A:H2	1:XA:1219:U:H1'	1.82	0.45
2:XB:216:SER:OG	2:XB:217:ARG:N	2.49	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:XE:100:VAL:HA	5:XE:118:ILE:HG22	1.97	0.45
12:XL:11:VAL:HG11	17:XQ:36:ILE:HG21	1.99	0.45
49:Y4:15:ILE:HB	49:Y4:32:TYR:HD1	1.80	0.45
24:YA:1832:G:N3	24:YA:1832:G:H2'	2.32	0.45
24:YA:662:A:H4'	24:YA:663:G:O5'	2.16	0.45
32:YN:96:GLU:HB2	32:YN:122:VAL:HG12	1.99	0.45
1:QA:1063:C:H2'	1:QA:1064:G:C8	2.51	0.45
1:QA:1201:A:HO2'	1:QA:1202:G:P	2.40	0.45
1:QA:1320:C:H2'	1:QA:1321:C:C6	2.51	0.45
1:QA:484:G:H4'	1:QA:485:G:O5'	2.16	0.45
1:QA:749:C:H2'	1:QA:750:G:H8	1.82	0.45
3:QC:155:GLY:HA3	3:QC:196:LEU:HD13	1.99	0.45
11:QK:85:ARG:HA	11:QK:112:THR:HG22	1.98	0.45
9:QI:111:ARG:NH1	14:QN:61:TRP:OXT	2.39	0.45
24:RA:30:G:O2'	24:RA:1259:A:N3	2.41	0.45
24:RA:1712:A:C3'	24:RA:1713:G:H5'	2.42	0.45
30:RH:43:VAL:HG23	30:RH:52:VAL:HG12	1.98	0.45
41:RW:6:ILE:HG12	41:RW:104:THR:HG23	1.98	0.45
1:XA:263:A:OP1	20:XT:79:ARG:HD3	2.17	0.45
49:Y4:62:ARG:HG3	49:Y4:63:TYR:H	1.81	0.45
24:YA:1683:C:H2'	24:YA:1684:A:H8	1.82	0.45
36:YR:26:LYS:O	36:YR:30:THR:OG1	2.26	0.45
1:QA:1414:U:H2'	1:QA:1415:G:C8	2.48	0.45
1:QA:1512:U:H2'	1:QA:1513:A:C8	2.52	0.45
1:QA:181:G:HO2'	1:QA:182:U:P	2.39	0.45
1:QA:269:C:H2'	1:QA:270:A:H8	1.81	0.45
1:QA:356:A:N3	1:QA:368:U:O2'	2.36	0.45
1:QA:45:U:OP1	1:QA:307:C:O2'	2.28	0.45
1:QA:703:G:H4'	1:QA:704:A:O5'	2.15	0.45
2:QB:195:ASP:O	8:QH:68:ARG:NH2	2.40	0.45
24:RA:1342:G:OP1	24:RA:2721:G:O2'	2.14	0.45
24:RA:740:C:O2'	24:RA:1399:A:N3	2.44	0.45
24:RA:1921:G:H22	24:RA:1924:C:H41	1.65	0.45
24:RA:468:G:H1'	28:RF:48:THR:HG21	1.98	0.45
1:XA:1095:U:P	1:XA:1108:G:H1	2.37	0.45
24:YA:1501:U:H5'	36:YR:63:ARG:HH21	1.81	0.45
24:YA:1701:A:N1	24:YA:2070:G:O2'	2.50	0.45
24:YA:1717:C:O5'	24:YA:1717:C:H6	1.99	0.45
24:YA:2603:C:H2'	24:YA:2604:G:C8	2.51	0.45
36:YR:28:LEU:HD23	36:YR:48:VAL:HG21	1.97	0.45
1:QA:1446:A:HO2'	1:QA:1447:G:P	2.39	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:21:G:H2'	1:QA:22:G:C8	2.52	0.45
1:QA:254:G:H2'	1:QA:255:G:H8	1.82	0.45
22:QV:43:G:H2'	22:QV:44:G:H8	1.80	0.45
24:RA:1331:G:N2	24:RA:1374:G:H5''	2.32	0.45
24:RA:166:G:H5'	24:RA:167:G:C8	2.52	0.45
24:RA:2204:G:H2'	24:RA:2205:C:C6	2.52	0.45
24:RA:2409:G:O6	24:RA:2431:U:O2	2.35	0.45
24:YA:1079:U:OP1	54:Y9:9:ARG:NH2	2.41	0.45
24:YA:1961:U:H3'	24:YA:1962:U:H5'	1.98	0.45
24:YA:956:A:N3	24:YA:2276:C:O2'	2.39	0.45
27:YE:52:LEU:O	27:YE:76:ARG:N	2.48	0.45
33:YO:34:THR:OG1	33:YO:35:VAL:N	2.49	0.45
34:YP:100:LEU:HD12	34:YP:112:LEU:HD11	1.98	0.45
24:YA:506:A:O2'	43:YY:46:LYS:O	2.34	0.45
44:YZ:144:LEU:HD11	44:YZ:150:LEU:HD23	1.98	0.45
1:QA:1287:A:H2'	1:QA:1288:A:C8	2.51	0.45
1:QA:692:U:O2'	1:QA:694:A:N7	2.42	0.45
6:QF:50:TYR:OH	18:QR:75:ILE:O	2.35	0.45
24:RA:1921:G:O2'	24:RA:1922:A:H5''	2.16	0.45
24:RA:2075:G:OP1	27:RE:144:ARG:HG2	2.17	0.45
24:RA:239:G:H2'	24:RA:240:A:C8	2.51	0.45
24:RA:503:A:H2'	24:RA:504:A:C8	2.52	0.45
39:RU:15:LYS:HB3	39:RU:15:LYS:HE3	1.77	0.45
1:XA:1119:C:H2'	1:XA:1120:G:H8	1.82	0.45
1:XA:1297:C:H1'	1:XA:1298:C:OP2	2.16	0.45
1:XA:1346:A:N1	1:XA:1374:A:H5''	2.32	0.45
1:XA:323:U:H2'	1:XA:324:G:O4'	2.16	0.45
2:XB:178:ARG:HH22	8:XH:68:ARG:HH22	1.65	0.45
3:XC:138:VAL:HG13	3:XC:149:ALA:HB3	1.99	0.45
4:XD:52:SER:OG	4:XD:55:ALA:N	2.44	0.45
15:XO:7:GLU:OE2	15:XO:38:ARG:NH2	2.40	0.45
6:XF:100:ASN:ND2	18:XR:27:GLY:O	2.43	0.45
21:XU:12:LYS:O	21:XU:16:GLY:N	2.45	0.45
49:Y4:28:LYS:HA	49:Y4:29:PRO:HD3	1.79	0.45
24:YA:2075:G:OP1	27:YE:144:ARG:HG2	2.17	0.45
1:QA:1499:A:OP1	1:QA:1500:A:OP2	2.35	0.45
1:QA:67:C:H2'	1:QA:68:G:C8	2.52	0.45
11:QK:87:THR:HA	11:QK:91:ARG:HD2	1.98	0.45
24:RA:1588:G:H5''	24:RA:1589:A:OP2	2.17	0.45
24:RA:1636:U:H2'	24:RA:1637:G:C8	2.51	0.45
24:RA:2589:A:H2'	24:RA:2626:A:N6	2.32	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:RB:44:G:H5''	25:RB:45:A:OP1	2.17	0.45
31:RI:97:ILE:O	31:RI:101:LEU:N	2.42	0.45
35:RQ:32:TYR:CZ	35:RQ:111:GLU:HB3	2.52	0.45
1:XA:377:G:H2'	1:XA:378:G:C8	2.52	0.45
5:XE:69:VAL:HG11	5:XE:113:ALA:HB1	1.99	0.45
24:YA:48:A:N7	24:YA:118:U:H5	2.15	0.45
24:YA:1639:G:H2'	24:YA:1640:G:C8	2.52	0.45
27:YE:57:LYS:HD2	27:YE:57:LYS:HA	1.73	0.45
1:QA:244:U:H4'	1:QA:245:C:O5'	2.17	0.45
18:QR:53:ARG:HH21	18:QR:59:SER:HA	1.82	0.45
24:RA:1580:G:O3'	24:RA:1581:U:H4'	2.17	0.45
24:RA:387:G:H2'	24:RA:388:A:H8	1.81	0.45
1:XA:1151:A:H2'	1:XA:1152:A:H8	1.81	0.45
1:XA:7:G:H5'	1:XA:298:A:O4'	2.17	0.45
1:XA:851:G:H2'	1:XA:852:G:H8	1.82	0.45
1:XA:881:G:P	12:XL:12:ARG:HH22	2.39	0.45
24:YA:1546:G:N2	26:YD:99:ASP:O	2.44	0.45
39:YU:92:ARG:HD3	39:YU:94:ASN:HB3	1.99	0.45
24:YA:1385:G:H5''	42:YX:16:LYS:HD2	1.99	0.45
1:QA:325:A:OP2	20:QT:70:SER:OG	2.23	0.44
3:QC:132:ARG:NH1	3:QC:136:GLN:HG3	2.32	0.44
11:QK:24:SER:HB3	11:QK:27:ASN:H	1.83	0.44
16:QP:1:MET:N	16:QP:1:MET:SD	2.85	0.44
24:RA:1553:A:H5'	24:RA:1554:A:OP2	2.16	0.44
24:RA:986:A:O2'	24:RA:1235:G:O3'	2.35	0.44
24:RA:2428:C:H5''	34:RP:65:ARG:HE	1.82	0.44
36:RR:88:ARG:NH2	36:RR:89:ASP:OD1	2.49	0.44
39:RU:92:ARG:HH12	40:RV:10:LYS:HA	1.82	0.44
42:RX:57:LEU:HG	42:RX:78:LYS:HB2	1.98	0.44
44:RZ:138:GLU:HB3	44:RZ:156:LYS:HE2	1.99	0.44
1:XA:1286:A:H2'	1:XA:1287:A:H4'	1.98	0.44
1:XA:538:G:H2'	1:XA:539:A:H8	1.82	0.44
5:XE:34:VAL:HG11	5:XE:63:ARG:HD2	1.98	0.44
12:XL:17:LYS:HD2	12:XL:18:VAL:H	1.82	0.44
1:XA:1309:G:P	13:XM:92:HIS:HE2	2.40	0.44
24:YA:1231:G:H2'	24:YA:1232:G:O4'	2.18	0.44
24:YA:2514:G:H5''	24:YA:2515:A:H5''	1.99	0.44
24:YA:2649:U:H2'	24:YA:2650:G:O4'	2.17	0.44
24:YA:539:A:N3	24:YA:604:C:O2'	2.39	0.44
25:YB:105:G:OP1	44:YZ:72:ARG:NH2	2.50	0.44
25:YB:92:G:H5''	44:YZ:79:ARG:NH2	2.32	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:19:C:OP1	5:QE:125:SER:OG	2.27	0.44
1:QA:261:U:OP2	20:QT:79:ARG:NH2	2.48	0.44
1:QA:411:A:C5	1:QA:413:G:H1'	2.52	0.44
48:R3:45:GLY:HA2	48:R3:48:GLU:HG2	1.99	0.44
24:RA:332:G:OP1	24:RA:332:G:C8	2.69	0.44
25:RB:24:G:H5''	25:RB:25:A:OP1	2.17	0.44
30:RH:54:ARG:HD3	30:RH:65:HIS:HD1	1.82	0.44
1:XA:1310:G:HO2'	1:XA:1311:G:P	2.40	0.44
1:XA:1427:U:H2'	1:XA:1428:A:C8	2.51	0.44
1:XA:444:C:H2'	1:XA:445:G:C8	2.52	0.44
1:XA:56:U:H2'	1:XA:57:G:H8	1.83	0.44
1:XA:828:A:H2'	1:XA:829:G:O4'	2.17	0.44
1:XA:851:G:H2'	1:XA:852:G:C8	2.52	0.44
1:XA:1345:U:H5''	9:XI:120:ARG:NH1	2.32	0.44
13:XM:97:PRO:HD3	13:XM:110:ARG:HB3	1.98	0.44
53:Y8:63:PRO:HB2	53:Y8:65:GLU:H	1.82	0.44
24:YA:1102:G:H5''	24:YA:1103:A:H5'	1.99	0.44
24:YA:795:G:OP1	24:YA:2624:C:N4	2.50	0.44
24:YA:2880:C:H2'	24:YA:2881:C:O4'	2.17	0.44
24:YA:302:A:H2'	24:YA:303:C:C6	2.52	0.44
26:YD:35:LYS:HD3	26:YD:64:ILE:HD11	1.99	0.44
36:YR:67:LEU:HD13	36:YR:76:VAL:HG21	1.97	0.44
38:YT:36:GLU:HG3	38:YT:41:ARG:HH21	1.82	0.44
1:QA:1127:G:H22	1:QA:1144:G:H22	1.64	0.44
1:QA:1277:C:O2'	1:QA:1279:A:H8	2.00	0.44
1:QA:25:C:H2'	1:QA:26:A:C8	2.52	0.44
1:QA:448:A:N7	1:QA:486:U:O4	2.50	0.44
9:QI:112:LYS:HA	9:QI:119:ALA:HB2	1.99	0.44
11:QK:72:ALA:HB1	11:QK:77:MET:HG2	1.99	0.44
20:QT:29:LYS:O	20:QT:33:ILE:HG12	2.18	0.44
24:RA:1103:A:H2'	24:RA:1104:G:H8	1.83	0.44
24:RA:1130:A:H3'	24:RA:1131:A:H4'	1.99	0.44
24:RA:1235:G:OP1	34:RP:30:THR:OG1	2.29	0.44
24:RA:2164:C:H2'	24:RA:2165:C:C6	2.52	0.44
24:RA:323:A:N3	24:RA:343:C:O2'	2.46	0.44
24:RA:51:A:H2'	24:RA:52:A:C8	2.51	0.44
25:RB:8:U:O2'	37:RS:25:ARG:NH2	2.50	0.44
1:XA:779:C:H2'	1:XA:780:A:O4'	2.17	0.44
1:XA:991:U:HO2'	1:XA:992:U:P	2.41	0.44
6:XF:11:ASN:HB3	6:XF:14:LEU:HG	1.99	0.44
7:XG:74:GLU:OE1	7:XG:95:ARG:NH2	2.49	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:XI:96:LEU:HB3	9:XI:102:LEU:HD23	1.98	0.44
10:XJ:24:VAL:HG13	10:XJ:25:GLU:HG2	1.99	0.44
16:XP:18:ARG:HD3	16:XP:35:LYS:HG3	1.99	0.44
20:XT:74:LYS:HB3	20:XT:76:ALA:H	1.82	0.44
24:YA:1064:C:O3'	24:YA:1166:G:N2	2.51	0.44
24:YA:1332:A:H1'	24:YA:1334:U:OP2	2.17	0.44
24:YA:1496:A:OP1	24:YA:1589:A:OP1	2.36	0.44
24:YA:1830:G:N7	26:YD:179:SER:OG	2.49	0.44
24:YA:2303:U:H2'	24:YA:2304:C:C6	2.52	0.44
24:YA:2689:G:H2'	24:YA:2690:C:C6	2.52	0.44
24:YA:606:G:OP2	39:YU:10:ARG:NH1	2.50	0.44
28:YF:136:THR:HG22	28:YF:166:ALA:HA	1.98	0.44
38:YT:111:ARG:H	38:YT:111:ARG:HG2	1.65	0.44
1:QA:1098:C:H2'	1:QA:1099:G:C8	2.51	0.44
1:QA:944:G:N1	1:QA:1338:G:OP2	2.50	0.44
1:QA:1250:A:H2	1:QA:1370:G:H1'	1.82	0.44
48:R3:51:ALA:HA	48:R3:54:VAL:HG12	1.97	0.44
24:RA:1378:G:H2'	24:RA:1378:G:H8	1.65	0.44
24:RA:1675:U:H2'	24:RA:1676:G:C8	2.53	0.44
24:RA:1714:G:C8	24:RA:1714:G:OP2	2.70	0.44
24:RA:718:C:H2'	24:RA:719:C:H6	1.82	0.44
24:RA:909:G:H2'	24:RA:910:A:O4'	2.17	0.44
1:XA:64:G:H5''	1:XA:65:U:OP1	2.17	0.44
18:XR:74:ARG:HD3	18:XR:81:PHE:HA	1.99	0.44
24:YA:1065:U:H3	24:YA:1188:A:H62	1.64	0.44
24:YA:1113:A:H8	24:YA:1113:A:OP1	2.01	0.44
24:YA:2609:G:H2'	24:YA:2610:A:C8	2.52	0.44
24:YA:2753:A:H2'	24:YA:2754:A:C8	2.52	0.44
24:YA:2797:C:H1'	27:YE:37:ARG:NH1	2.31	0.44
24:YA:388:A:H2'	24:YA:389:G:C8	2.52	0.44
1:QA:596:C:OP2	1:QA:597:G:OP2	2.35	0.44
5:QE:37:ARG:HG2	5:QE:112:LEU:HA	2.00	0.44
24:RA:1501:U:H5'	36:RR:63:ARG:HH22	1.83	0.44
24:RA:2184:G:H4'	24:RA:2195:A:OP2	2.17	0.44
24:RA:718:C:H2'	24:RA:719:C:C6	2.53	0.44
25:RB:77:U:OP1	44:RZ:19:ARG:NH2	2.50	0.44
1:XA:1003:G:N2	1:XA:1005:A:H5'	2.32	0.44
1:XA:1323:G:H2'	1:XA:1324:A:H8	1.81	0.44
1:XA:304:U:H2'	1:XA:305:G:C8	2.53	0.44
1:XA:343:U:O2	1:XA:346:G:N1	2.51	0.44
11:XK:21:ILE:HG23	11:XK:30:VAL:HG12	1.99	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:Y4:28:LYS:HD3	49:Y4:31:ILE:HD11	2.00	0.44
24:YA:1675:U:H2'	24:YA:1676:G:C8	2.53	0.44
24:YA:2145:G:H2'	24:YA:2146:G:H8	1.81	0.44
24:YA:716:G:N3	24:YA:716:G:H2'	2.32	0.44
1:XA:345:C:OP2	38:YT:39:ARG:NH2	2.50	0.44
1:QA:989:C:H1'	1:QA:1016:A:H2	1.83	0.44
1:QA:923:A:O2'	1:QA:1399:C:OP2	2.29	0.44
1:QA:553:A:H2'	1:QA:554:C:C6	2.52	0.44
1:QA:99:C:H2'	1:QA:101:A:C8	2.52	0.44
5:QE:110:LEU:HD23	5:QE:115:VAL:HG21	2.00	0.44
11:QK:34:ASP:OD1	11:QK:38:ASN:N	2.51	0.44
51:R6:35:GLU:OE2	51:R6:50:ARG:NH2	2.50	0.44
24:RA:593:G:H2'	24:RA:2052:A:C5	2.53	0.44
24:RA:2803:A:H2'	24:RA:2804:C:H5''	1.99	0.44
24:RA:212:A:O2'	24:RA:447:C:O2	2.32	0.44
34:RP:95:VAL:HA	34:RP:99:LEU:HD12	1.98	0.44
1:XA:1363:A:H4'	1:XA:1364:U:H2'	1.99	0.44
1:XA:34:C:H2'	1:XA:35:G:H8	1.83	0.44
1:XA:363:A:C6	12:XL:31:PRO:HD2	2.52	0.44
1:XA:485:G:O2'	1:XA:486:U:O5'	2.32	0.44
3:XC:189:ALA:HB3	3:XC:196:LEU:HB2	1.99	0.44
7:XG:15:ASP:OD2	7:XG:18:TYR:N	2.47	0.44
12:XL:77:LEU:HD21	12:XL:107:ALA:HB2	1.99	0.44
24:YA:485:U:H2'	24:YA:486:A:C8	2.52	0.44
35:YQ:110:THR:OG1	35:YQ:113:GLN:N	2.42	0.44
1:QA:1064:G:HO2'	1:QA:1065:U:P	2.40	0.44
1:QA:422:C:HO2'	1:QA:423:G:N2	2.14	0.44
1:QA:552:U:H2'	1:QA:553:A:H8	1.83	0.44
1:QA:629:G:H2'	1:QA:630:G:C8	2.52	0.44
1:QA:828:A:H2'	1:QA:829:G:O4'	2.18	0.44
24:RA:1077:G:O2'	54:R9:7:VAL:O	2.33	0.44
24:RA:259:A:OP2	24:RA:284:G:N1	2.47	0.44
5:XE:139:LEU:HA	5:XE:142:LEU:HD12	1.99	0.44
7:XG:29:LYS:HZ1	7:XG:102:ARG:NH2	2.15	0.44
7:XG:53:LYS:HD3	7:XG:53:LYS:HA	1.81	0.44
9:XI:10:ARG:HE	9:XI:105:ASP:HB2	1.82	0.44
24:YA:2038:U:O2'	50:Y5:7:PRO:O	2.29	0.44
24:YA:1683:C:H2'	24:YA:1684:A:C8	2.52	0.44
24:YA:1822:A:H4'	26:YD:206:LEU:HB2	1.99	0.44
24:YA:2086:C:H2'	24:YA:2087:C:H6	1.82	0.44
24:YA:37:C:H2'	24:YA:38:A:C8	2.53	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:YA:645:G:H4'	24:YA:646:A:H5''	1.99	0.44
25:YB:89(A):G:H3'	25:YB:89(B):A:C5'	2.47	0.44
37:YS:46:VAL:HG12	37:YS:48:LEU:HD22	2.00	0.44
38:YT:113:LYS:HA	38:YT:113:LYS:HD3	1.76	0.44
1:QA:1079:G:H2'	1:QA:1080:A:C8	2.53	0.44
1:QA:1479:C:H2'	1:QA:1480:G:C8	2.53	0.44
1:QA:1505:G:H2'	23:QX:15:A:OP2	2.17	0.44
1:QA:704:A:H8	1:QA:704:A:OP2	1.99	0.44
16:QP:45:THR:HG23	16:QP:47:ASP:H	1.83	0.44
24:RA:114:C:O2'	24:RA:124:A:N3	2.36	0.44
24:RA:1399:A:H2'	24:RA:1400:A:C8	2.53	0.44
24:RA:2082:A:O2'	24:RA:2083:G:OP1	2.32	0.44
24:RA:2173:G:H2'	24:RA:2174:G:C8	2.53	0.44
24:RA:659:C:H2'	24:RA:660:C:C6	2.53	0.44
24:RA:2591:C:H1'	27:RE:134:ILE:HD13	2.00	0.44
34:RP:99:LEU:HD23	34:RP:102:ARG:HH21	1.82	0.44
35:RQ:43:THR:OG1	35:RQ:44:ALA:N	2.51	0.44
1:XA:46:G:H2'	1:XA:366:C:H5	1.82	0.44
1:XA:410:G:H4'	1:XA:411:A:OP1	2.18	0.44
1:XA:31:G:HO2'	1:XA:48:C:N4	2.15	0.44
3:XC:114:PRO:HA	3:XC:117:ALA:HB3	1.99	0.44
1:XA:7:G:O2'	5:XE:120:THR:O	2.35	0.44
12:XL:84:LEU:HB2	12:XL:105:TYR:CE2	2.52	0.44
13:XM:114:ARG:O	13:XM:116:THR:OG1	2.30	0.44
23:XX:8:A:H2'	23:XX:9:G:C8	2.53	0.44
24:YA:2329:C:H2'	24:YA:2330:G:O4'	2.17	0.44
24:YA:2299:A:N6	24:YA:2356:U:H3	2.15	0.44
24:YA:851:A:H2'	24:YA:853:C:C4	2.53	0.44
29:YG:77:ILE:HG22	29:YG:79:ASN:H	1.83	0.44
34:YP:94:GLU:HG3	34:YP:124:LYS:HD3	1.99	0.44
24:YA:950:C:O2'	44:YZ:169:GLU:OE2	2.36	0.44
1:QA:368:U:OP1	31:YI:91:SER:OG	2.27	0.44
3:QC:44:GLU:HG3	3:QC:52:LEU:HD21	1.99	0.44
24:RA:1199:C:H2'	24:RA:1200:G:O4'	2.18	0.44
24:RA:1712:A:C3'	24:RA:1713:G:C5'	2.94	0.44
24:RA:2818:U:O2	24:RA:2901:A:C6	2.71	0.44
24:RA:715:G:H5'	24:RA:716:G:OP2	2.16	0.44
24:RA:930:G:H1	24:RA:939:C:H42	1.66	0.44
27:RE:78:LEU:HD23	27:RE:78:LEU:HA	1.87	0.44
28:RF:23:ASP:OD1	28:RF:23:ASP:N	2.39	0.44
33:RO:12:ASP:N	33:RO:12:ASP:OD1	2.51	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:RR:33:ARG:NE	36:RR:115:GLU:OE1	2.45	0.44
36:RR:67:LEU:HD13	36:RR:76:VAL:HG21	1.99	0.44
1:XA:1022:G:H2'	1:XA:1023:G:C8	2.53	0.44
1:XA:1036:G:N7	1:XA:1037:C:N4	2.66	0.44
1:XA:1404:C:H2'	1:XA:1405:G:C8	2.53	0.44
1:XA:281:G:OP2	1:XA:281:G:H8	2.01	0.44
1:XA:514:C:H2'	1:XA:515:G:C8	2.52	0.44
19:XS:9:VAL:HG12	19:XS:10:PHE:HB2	2.00	0.44
29:YG:67:LYS:H	49:Y4:6:HIS:CE1	2.36	0.44
24:YA:2859:U:H4'	24:YA:2878:A:C2	2.53	0.44
24:YA:2877:G:HO2'	24:YA:2878:A:P	2.34	0.44
24:YA:604:C:H2'	24:YA:605:G:H8	1.83	0.44
25:YB:89(B):A:C8	25:YB:89(B):A:H3'	2.52	0.44
24:YA:2649:U:H5''	27:YE:82:ARG:NH1	2.33	0.44
29:YG:60:LEU:HA	29:YG:60:LEU:HD12	1.85	0.44
36:YR:54:LEU:HB3	36:YR:62:ALA:HB1	1.99	0.44
1:QA:1298:C:H4'	1:QA:1299:A:C8	2.53	0.43
17:QQ:81:ARG:HH21	17:QQ:84:LEU:HD11	1.83	0.43
51:R6:11:LEU:HB2	51:R6:21:TYR:HB2	2.00	0.43
24:RA:1729:G:OP2	24:RA:1746:G:N2	2.50	0.43
24:RA:268:G:H2'	24:RA:269:G:C8	2.53	0.43
24:RA:2878:A:H2'	24:RA:2879:G:C8	2.53	0.43
24:RA:334:A:N3	24:RA:336:G:C8	2.85	0.43
24:RA:1612:C:H5''	26:RD:18:VAL:HG11	1.99	0.43
27:RE:28:ALA:HB3	27:RE:93:VAL:HG12	1.99	0.43
30:RH:89:ILE:HG22	30:RH:162:ILE:HG23	1.99	0.43
1:XA:1427:U:H2'	1:XA:1428:A:H8	1.83	0.43
2:XB:154:LEU:HB3	2:XB:155:LEU:HD23	2.00	0.43
13:XM:68:GLY:O	13:XM:72:ALA:N	2.48	0.43
24:YA:1394:G:H2'	24:YA:1395:A:H5''	2.00	0.43
24:YA:1480:A:H2'	24:YA:1481:G:C8	2.53	0.43
24:YA:2172:U:H2'	24:YA:2173:G:C8	2.53	0.43
24:YA:2637:G:H2'	24:YA:2638:C:O4'	2.18	0.43
24:YA:64:C:H2'	24:YA:65:C:H6	1.83	0.43
36:YR:3:HIS:O	36:YR:5:LYS:N	2.51	0.43
1:QA:600:C:H2'	1:QA:601:C:C6	2.53	0.43
6:QF:79:LEU:HB3	6:QF:88:VAL:HG21	1.99	0.43
28:RF:101:LEU:O	28:RF:106:ARG:NH2	2.46	0.43
41:RW:111:HIS:CD2	41:RW:113:LYS:H	2.36	0.43
1:XA:1233:G:H2'	1:XA:1234:C:C6	2.53	0.43
1:XA:189:U:H3	17:XQ:63:ARG:HB2	1.82	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:398:C:H2'	1:XA:399:G:H8	1.82	0.43
1:XA:560:U:H4'	1:XA:561:U:H5''	1.99	0.43
24:YA:1935:A:H2'	24:YA:1935:A:H8	1.37	0.43
24:YA:2163:G:H2'	24:YA:2164:C:C6	2.54	0.43
24:YA:299:G:H22	24:YA:301:C:H41	1.67	0.43
38:YT:34:VAL:HG12	38:YT:36:GLU:HG2	2.00	0.43
1:QA:153:C:N3	1:QA:169:C:N4	2.65	0.43
1:QA:762:C:H2'	1:QA:763:G:C8	2.53	0.43
10:QJ:50:ILE:HA	10:QJ:60:ARG:HG2	2.00	0.43
1:QA:778:G:O2'	11:QK:120:ARG:O	2.26	0.43
16:QP:34:GLU:OE1	16:QP:59:TRP:NE1	2.45	0.43
51:R6:23:THR:OG1	51:R6:24:GLU:N	2.51	0.43
24:RA:106:U:H2'	24:RA:107:G:C8	2.53	0.43
24:RA:1170:C:H2'	24:RA:1171:G:O4'	2.17	0.43
24:RA:2124:U:H2'	24:RA:2125:C:H6	1.83	0.43
24:RA:2842:U:HO2'	24:RA:2843:G:C5'	2.31	0.43
1:XA:1355:G:H2'	1:XA:1356:G:C8	2.54	0.43
1:XA:191(E):G:H2'	1:XA:191(F):U:C6	2.54	0.43
1:XA:315:A:H5''	1:XA:317:G:OP2	2.18	0.43
7:XG:79:ARG:NE	7:XG:84:ASN:OD1	2.51	0.43
13:XM:69:GLU:HA	13:XM:72:ALA:HB3	2.00	0.43
24:YA:1704:C:H2'	24:YA:1705:C:H6	1.83	0.43
24:YA:1830:G:O6	26:YD:179:SER:OG	2.35	0.43
24:YA:2081:A:H5'	24:YA:2082:A:OP2	2.18	0.43
24:YA:2642:G:H2'	24:YA:2643:G:H8	1.82	0.43
24:YA:2699:U:H2'	24:YA:2700:U:O4'	2.18	0.43
1:QA:1095:U:H2'	1:QA:1096:C:O4'	2.18	0.43
1:QA:1210:C:HO2'	1:QA:1213:A:HO2'	1.63	0.43
1:QA:7:G:H5'	1:QA:298:A:O4'	2.18	0.43
1:QA:657:G:N2	15:QO:22:THR:OG1	2.49	0.43
24:RA:1405:A:OP2	24:RA:1417:G:N1	2.47	0.43
24:RA:37:C:H2'	24:RA:38:A:C8	2.53	0.43
24:RA:485:U:H2'	24:RA:486:A:H8	1.83	0.43
24:RA:662:A:OP1	34:RP:133:SER:OG	2.30	0.43
26:RD:69:ARG:HD3	26:RD:130:ALA:HB2	2.01	0.43
29:RG:91:ARG:HB3	29:RG:91:ARG:HE	1.58	0.43
1:XA:151:A:N6	1:XA:170:U:H3	2.12	0.43
1:XA:413:G:OP2	1:XA:413:G:H8	2.01	0.43
1:XA:41:G:H2'	1:XA:42:G:C8	2.53	0.43
2:XB:198:ASP:OD1	8:XH:68:ARG:NH1	2.49	0.43
4:XD:59:ARG:HH12	4:XD:66:ARG:HH12	1.67	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:Y6:35:GLU:OE2	51:Y6:50:ARG:NH1	2.43	0.43
24:YA:196:A:H2'	24:YA:197:C:O4'	2.17	0.43
24:YA:2162:C:H2'	24:YA:2163:G:C8	2.52	0.43
24:YA:2589:A:H2'	24:YA:2626:A:N6	2.34	0.43
24:YA:659:C:H2'	24:YA:660:C:C6	2.53	0.43
29:YG:6:ALA:O	29:YG:10:LYS:N	2.49	0.43
34:YP:97:PRO:HA	34:YP:112:LEU:HD12	2.00	0.43
1:QA:1219:U:H2'	1:QA:1220:G:C8	2.54	0.43
1:QA:1250:A:N3	1:QA:1370:G:O2'	2.43	0.43
1:QA:475:G:H2'	1:QA:476:G:H8	1.83	0.43
1:QA:940:C:H2'	1:QA:941:G:C8	2.52	0.43
3:QC:154:SER:HB2	3:QC:165:THR:HG22	2.00	0.43
24:RA:1047:A:H62	24:RA:1200:G:N2	2.16	0.43
24:RA:49:U:H3'	24:RA:50:G:H5'	1.99	0.43
24:RA:660:C:O2'	24:RA:664:U:OP1	2.37	0.43
27:RE:47:VAL:HG12	27:RE:49:LEU:HD12	1.99	0.43
24:RA:1501:U:H5'	36:RR:63:ARG:NH2	2.33	0.43
1:XA:253:U:OP1	17:XQ:67:LYS:NZ	2.30	0.43
1:XA:582:U:OP1	15:XO:68:ARG:NH2	2.47	0.43
3:XC:70:VAL:HG12	3:XC:72:LYS:H	1.83	0.43
48:Y3:15:TYR:O	48:Y3:20:LYS:NZ	2.39	0.43
49:Y4:40:HIS:HB3	49:Y4:43:TYR:HB2	2.00	0.43
24:YA:1473:A:H4'	24:YA:1474:C:O5'	2.18	0.43
24:YA:1899:A:H5'	24:YA:1900:G:OP2	2.19	0.43
24:YA:2172:U:H2'	24:YA:2173:G:H8	1.83	0.43
24:YA:2229:A:H1'	24:YA:2231:G:C4	2.53	0.43
24:YA:389:G:H2'	24:YA:390:G:C8	2.53	0.43
24:YA:236:G:H4'	24:YA:413:G:C6	2.53	0.43
24:YA:664:U:H2'	24:YA:665:C:C6	2.53	0.43
38:YT:18:ASP:OD1	38:YT:18:ASP:N	2.33	0.43
1:QA:1035:A:C5	1:QA:1036:G:H1'	2.53	0.43
1:QA:673:G:O3'	6:QF:87:ARG:NH2	2.51	0.43
1:QA:768:A:N3	1:QA:1512:U:O2'	2.51	0.43
2:QB:48:MET:HA	2:QB:51:LEU:HD12	2.00	0.43
3:QC:10:PHE:HD2	3:QC:11:ARG:HD2	1.83	0.43
24:RA:505:A:O2'	24:RA:507:G:H5'	2.19	0.43
29:RG:55:LYS:HA	29:RG:58:GLN:HG2	2.01	0.43
38:RT:54:ARG:HA	38:RT:59:THR:HG23	2.00	0.43
39:RU:90:VAL:HG11	40:RV:39:LEU:HB2	1.99	0.43
1:XA:1314:C:OP2	19:XS:4:SER:OG	2.30	0.43
1:XA:1492:A:H1'	1:XA:1493:A:H8	1.83	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:195:A:OP1	20:XT:65:LYS:NZ	2.45	0.43
1:XA:390:C:H2'	1:XA:391:G:H8	1.84	0.43
1:XA:411:A:C4	1:XA:413:G:H1'	2.54	0.43
10:XJ:38:ILE:HG13	10:XJ:71:LEU:HB3	2.01	0.43
12:XL:39:VAL:HG12	12:XL:41:ARG:HG3	2.00	0.43
24:YA:1065:U:O2'	24:YA:1067:A:H2	2.02	0.43
24:YA:1183:G:H2'	24:YA:1184:G:O4'	2.19	0.43
24:YA:1233:U:H4'	40:YV:79:VAL:HG22	1.99	0.43
24:YA:554:A:H8	24:YA:555:G:C6	2.37	0.43
25:YB:13:A:O2'	25:YB:15:A:H5''	2.19	0.43
1:QA:1241:G:H2'	1:QA:1242:C:C6	2.53	0.43
1:QA:1279:A:OP1	10:QJ:7:LYS:NZ	2.47	0.43
1:QA:1298:C:H4'	1:QA:1299:A:C5	2.53	0.43
1:QA:429:U:H1'	1:QA:430:A:H5''	2.00	0.43
24:RA:1211:U:H2'	24:RA:1212:C:C6	2.54	0.43
24:RA:1593:C:H5'	24:RA:1594:C:H5'	2.01	0.43
24:RA:1639:G:H2'	24:RA:1640:G:H8	1.84	0.43
24:RA:331:G:N2	24:RA:333:G:H3'	2.33	0.43
24:RA:1853:G:H5''	26:RD:54:ARG:HH22	1.84	0.43
27:RE:152:LYS:HB3	32:RN:78:TYR:HA	2.01	0.43
24:RA:2757:G:N2	30:RH:143:GLN:OE1	2.46	0.43
1:XA:564:C:C4	17:XQ:31:LEU:HD11	2.53	0.43
1:XA:890:G:O2'	1:XA:906:G:O6	2.32	0.43
24:YA:574:G:O2'	24:YA:1265:A:N3	2.43	0.43
24:YA:2754:A:OP1	54:Y9:22:ARG:NH1	2.51	0.43
30:YH:28:GLY:HA3	30:YH:79:VAL:HB	2.00	0.43
39:YU:54:LYS:HG3	39:YU:54:LYS:H	1.66	0.43
39:YU:98:LEU:O	39:YU:102:GLU:N	2.51	0.43
40:YV:30:GLY:H	40:YV:61:VAL:HG23	1.83	0.43
2:QB:162:ILE:HD11	2:QB:184:VAL:HG22	1.99	0.43
11:QK:86:GLY:H	11:QK:112:THR:HG22	1.84	0.43
17:QQ:59:ILE:HG22	17:QQ:73:VAL:HA	2.01	0.43
20:QT:75:ASN:N	20:QT:75:ASN:OD1	2.44	0.43
24:RA:2125:C:H42	24:RA:2209:G:H1	1.66	0.43
24:RA:2694:U:H5'	24:RA:2694:U:C6	2.53	0.43
24:RA:891:C:H2'	24:RA:892:G:O4'	2.17	0.43
28:RF:40:GLN:HE22	28:RF:183:VAL:H	1.65	0.43
29:RG:102:PHE:O	29:RG:106:LEU:N	2.46	0.43
42:RX:29:TRP:HZ3	42:RX:59:VAL:HG11	1.84	0.43
1:XA:1151:A:H2'	1:XA:1152:A:C8	2.54	0.43
1:XA:1172:C:H2'	1:XA:1173:G:H8	1.84	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:201:C:N4	1:XA:216:G:H1	2.17	0.43
1:XA:922:G:H2'	1:XA:923:A:C8	2.54	0.43
1:XA:429:U:H5'	4:XD:9:CYS:SG	2.59	0.43
6:XF:18:GLN:HA	6:XF:21:LEU:HD12	1.99	0.43
13:XM:32:GLU:O	13:XM:36:LYS:N	2.38	0.43
36:YR:33:ARG:NH2	50:Y5:57:VAL:O	2.39	0.43
51:Y6:11:LEU:HB2	51:Y6:21:TYR:HB2	2.01	0.43
24:YA:1542:A:H8	24:YA:1624:C:O2'	2.01	0.43
24:YA:2205:C:H2'	24:YA:2206:G:C8	2.54	0.43
24:YA:2376:C:H2'	24:YA:2377:G:O4'	2.19	0.43
24:YA:2748:G:H2'	24:YA:2749:G:H8	1.83	0.43
1:QA:1071:C:H2'	1:QA:1072:G:H8	1.84	0.43
1:QA:1293:G:H2'	1:QA:1294:G:H8	1.83	0.43
5:QE:121:LYS:HD2	5:QE:121:LYS:HA	1.71	0.43
9:QI:116:LYS:HD3	9:QI:122:ALA:HB2	2.01	0.43
11:QK:92:GLU:O	11:QK:96:ARG:HG3	2.18	0.43
22:QV:1:C:O2'	45:R0:6:GLY:O	2.36	0.43
22:QV:54:U:O4	22:QV:58:A:N7	2.52	0.43
46:R1:81:LYS:HB2	46:R1:81:LYS:HE3	1.82	0.43
24:RA:138:G:N2	24:RA:1642:A:H4'	2.34	0.43
24:RA:1566:U:H2'	24:RA:1567:G:O4'	2.18	0.43
24:RA:604:C:H2'	24:RA:605:G:C8	2.54	0.43
24:RA:612:C:H2'	24:RA:613:A:H8	1.83	0.43
31:RI:75:LEU:HD12	31:RI:105:HIS:CE1	2.54	0.43
39:RU:66:ASN:ND2	39:RU:70:ARG:HH21	2.16	0.43
1:XA:1240:U:OP2	7:XG:116:ALA:N	2.51	0.43
1:XA:17:U:H2'	1:XA:18:C:H6	1.83	0.43
1:XA:346:G:H1'	1:XA:347:G:H5'	2.00	0.43
1:XA:80:G:H1	1:XA:89:U:H3	1.65	0.43
24:YA:169:G:OP1	52:Y7:32:LYS:HG3	2.18	0.43
24:YA:1897:C:H2'	24:YA:1898:A:O4'	2.18	0.43
24:YA:909:G:H2'	24:YA:910:A:O4'	2.18	0.43
1:QA:357:G:O2'	31:YI:89:TYR:O	2.30	0.43
34:YP:91:PHE:O	34:YP:121:LYS:NZ	2.40	0.43
1:QA:1119:C:H2'	1:QA:1120:G:H8	1.84	0.43
1:QA:115:G:H4'	1:QA:116:A:O5'	2.18	0.43
1:QA:476:G:H2'	1:QA:477:G:H8	1.84	0.43
1:QA:757:U:H2'	1:QA:758:G:O4'	2.19	0.43
10:QJ:49:VAL:HG23	14:QN:41:ARG:HB2	2.00	0.43
24:RA:1108:G:O2'	24:RA:1109:G:H5'	2.19	0.43
24:RA:1712:A:O2'	24:RA:1713:G:H5''	2.19	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:RA:2082:A:H3'	24:RA:2082:A:OP1	2.18	0.43
24:RA:2168:C:H4'	24:RA:2169:G:C5	2.54	0.43
24:RA:605:G:H2'	24:RA:606:G:H8	1.84	0.43
33:RO:73:ASP:N	33:RO:73:ASP:OD1	2.36	0.43
1:XA:703:G:H4'	1:XA:704:A:O5'	2.19	0.43
1:XA:911:U:H2'	1:XA:912:C:C6	2.53	0.43
1:XA:501:C:P	12:XL:117:ARG:HH21	2.42	0.43
22:XV:3:G:H2'	22:XV:4:U:H5'	1.99	0.43
24:YA:1869:C:N4	24:YA:1921:G:OP2	2.27	0.43
24:YA:230:A:H4'	24:YA:231:G:OP1	2.19	0.43
24:YA:2818:U:C2	24:YA:2901:A:N6	2.87	0.43
24:YA:553:A:HO2'	24:YA:554:A:H5'	1.81	0.43
25:YB:30:C:H2'	25:YB:31:C:H5'	2.01	0.43
27:YE:119:ARG:NH1	27:YE:159:HIS:O	2.51	0.43
35:YQ:34:LEU:HB2	35:YQ:118:LEU:HD22	2.00	0.43
39:YU:15:LYS:HB3	39:YU:15:LYS:HE3	1.77	0.43
24:YA:1043:G:OP1	39:YU:93:LYS:HD3	2.19	0.43
1:QA:352:C:O2'	1:QA:354:G:OP1	2.22	0.42
1:QA:751:U:H2'	1:QA:752:G:O4'	2.19	0.42
4:QD:18:LYS:NZ	4:QD:34:GLU:OE2	2.45	0.42
9:QI:93:ARG:HB2	9:QI:97:LYS:HD3	2.01	0.42
49:R4:56:VAL:HG23	49:R4:58:ARG:HG2	2.01	0.42
24:RA:1958:A:OP2	24:RA:1984:C:N4	2.52	0.42
24:RA:2185:C:N4	24:RA:2186:C:H41	2.17	0.42
24:RA:2227:G:H3'	24:RA:2228:G:C8	2.54	0.42
24:RA:2473:C:H2'	24:RA:2474:U:C6	2.54	0.42
24:RA:702:A:H2'	24:RA:703:G:H5'	2.01	0.42
24:RA:843:C:H2'	24:RA:844:C:C6	2.54	0.42
25:RB:49:C:H2'	25:RB:50:G:C8	2.54	0.42
25:RB:66:A:HO2'	25:RB:67:G:P	2.42	0.42
25:RB:79:C:H2'	25:RB:80:U:O4'	2.19	0.42
39:RU:92:ARG:HD3	39:RU:95:LEU:HD12	2.01	0.42
1:XA:1218:C:H2'	1:XA:1219:U:C6	2.54	0.42
1:XA:269:C:H2'	1:XA:270:A:C8	2.54	0.42
24:YA:1265:A:H5'	24:YA:1266:C:OP2	2.18	0.42
24:YA:2206:G:H2'	24:YA:2207:C:C6	2.54	0.42
24:YA:2787:C:H2'	24:YA:2788:A:O4'	2.19	0.42
24:YA:331:G:N1	24:YA:334:A:OP2	2.52	0.42
24:YA:571:A:H2'	24:YA:572:A:C8	2.54	0.42
33:YO:24:VAL:HG13	33:YO:33:ALA:HB2	2.01	0.42
1:QA:1014:A:H2'	1:QA:1015:A:C8	2.54	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1151:A:H2'	1:QA:1152:A:H8	1.84	0.42
1:QA:1524:C:H2'	1:QA:1525:G:C8	2.53	0.42
1:QA:131:C:O2'	1:QA:262:A:N3	2.40	0.42
1:QA:753:A:H1'	1:QA:754:C:OP2	2.18	0.42
1:QA:861:G:HO2'	1:QA:874:G:HO2'	1.64	0.42
4:QD:101:LEU:HB2	4:QD:138:TYR:HB3	2.01	0.42
1:QA:1376:U:OP2	7:QG:94:ARG:NH2	2.52	0.42
24:RA:1921:G:N2	24:RA:1924:C:H41	2.17	0.42
24:RA:2129:C:H2'	24:RA:2130:C:C6	2.54	0.42
24:RA:2143:G:H2'	24:RA:2144:U:C6	2.54	0.42
29:RG:7:LEU:N	29:RG:104:GLU:OE2	2.36	0.42
34:RP:95:VAL:HG22	34:RP:125:VAL:HA	2.00	0.42
35:RQ:65:PHE:HB2	35:RQ:105:GLU:HB3	2.00	0.42
35:RQ:42:ILE:HD12	35:RQ:97:VAL:HG21	2.01	0.42
1:XA:1517:G:N3	24:YA:1941:A:O2'	2.40	0.42
1:XA:93:U:H2'	1:XA:95:G:O4'	2.20	0.42
1:XA:946:A:H2'	1:XA:947:G:C8	2.54	0.42
29:YG:179:PRO:HB2	49:Y4:42:PHE:HE2	1.84	0.42
24:YA:414:U:P	46:Y1:20:ARG:HH12	2.41	0.42
24:YA:640:A:C4	28:YF:180:GLY:HA3	2.54	0.42
1:QA:1095:U:P	1:QA:1108:G:H1	2.41	0.42
1:QA:114:U:H2'	1:QA:115:G:C8	2.55	0.42
1:QA:1372:U:H5''	9:QI:71:SER:HB3	2.01	0.42
1:QA:513:C:H2'	1:QA:514:C:C6	2.55	0.42
1:QA:628:G:H2'	1:QA:629:G:C8	2.54	0.42
8:QH:34:GLU:OE1	8:QH:37:ARG:NH2	2.53	0.42
13:QM:91:ARG:HA	13:QM:91:ARG:HD3	1.77	0.42
20:QT:20:LEU:HA	20:QT:23:ARG:HG2	2.00	0.42
23:QX:5:A:H2'	23:QX:6:G:C8	2.53	0.42
41:RW:37:ARG:NH2	50:R5:48:GLU:OE2	2.52	0.42
24:RA:2538:G:H1	24:RA:2549:U:H3	1.67	0.42
25:RB:5:C:OP1	25:RB:61:G:O2'	2.34	0.42
43:RY:14:LEU:HD11	43:RY:22:GLY:HA2	2.00	0.42
1:XA:1219:U:H2'	1:XA:1220:G:C8	2.55	0.42
1:XA:1285:A:H1'	1:XA:1286:A:OP2	2.19	0.42
1:XA:986:A:H2'	1:XA:987:G:C8	2.54	0.42
1:XA:1191:A:OP2	3:XC:3:ASN:ND2	2.52	0.42
7:XG:58:PRO:HA	7:XG:61:VAL:HG12	2.02	0.42
13:XM:54:VAL:O	13:XM:58:GLU:N	2.41	0.42
19:XS:43:GLU:OE2	19:XS:43:GLU:N	2.53	0.42
24:YA:1362:U:H2'	24:YA:1363:A:H8	1.84	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:YA:1822:A:H3'	24:YA:1823:G:H8	1.84	0.42
32:YN:23:LEU:HD13	32:YN:60:ILE:HD13	2.00	0.42
40:YV:24:LYS:HB3	40:YV:24:LYS:HE2	1.88	0.42
1:QA:1067:A:N1	1:QA:1108:G:O2'	2.43	0.42
1:QA:150:C:N4	1:QA:171:A:H62	2.17	0.42
1:QA:262:A:H2'	1:QA:263:A:C8	2.53	0.42
1:QA:381:C:H2'	1:QA:382:A:O4'	2.20	0.42
1:QA:538:G:H2'	1:QA:539:A:H8	1.84	0.42
1:QA:816:A:OP1	1:QA:1526:G:O2'	2.31	0.42
45:R0:32:ARG:H	45:R0:35:ASN:ND2	2.18	0.42
49:R4:8:LYS:HA	49:R4:8:LYS:HD2	1.89	0.42
51:R6:4:GLU:HG2	51:R6:4:GLU:H	1.73	0.42
24:RA:85:C:H4'	24:RA:102:U:H1'	2.01	0.42
24:RA:1698:G:OP1	36:RR:40:LYS:NZ	2.41	0.42
24:RA:1884:A:H2'	24:RA:1885:A:C8	2.55	0.42
24:RA:431:C:H1'	24:RA:432:U:OP2	2.20	0.42
24:RA:546:G:H2'	24:RA:547:G:H8	1.85	0.42
24:RA:664:U:H2'	24:RA:665:C:C6	2.53	0.42
24:RA:895:G:H2'	24:RA:896:A:H8	1.82	0.42
29:RG:129:GLY:O	29:RG:161:THR:OG1	2.37	0.42
30:RH:164:TYR:HB2	30:RH:167:GLU:HB2	2.01	0.42
24:RA:1188:A:H4'	32:RN:25:ARG:NH2	2.34	0.42
35:RQ:110:THR:OG1	35:RQ:113:GLN:OE1	2.26	0.42
37:RS:15:ARG:NH1	37:RS:88:ASP:OD2	2.52	0.42
1:XA:192:U:H2'	1:XA:193:C:C6	2.54	0.42
1:XA:265:G:H2'	1:XA:266:G:H5''	2.00	0.42
1:XA:458:C:N4	1:XA:475:G:O6	2.53	0.42
1:XA:501:C:H2'	1:XA:502:G:H8	1.83	0.42
2:XB:106:LYS:HD3	2:XB:106:LYS:HA	1.86	0.42
7:XG:40:ALA:HB1	7:XG:44:TYR:HE2	1.84	0.42
24:YA:1405:A:N6	24:YA:1418:U:N3	2.29	0.42
24:YA:1480:A:H61	24:YA:1605:A:H62	1.67	0.42
24:YA:184:A:OP1	34:YP:46:LYS:NZ	2.37	0.42
24:YA:259:A:OP2	24:YA:284:G:N1	2.47	0.42
24:YA:985:G:H2'	24:YA:986:A:O4'	2.19	0.42
24:YA:2692:C:H5'	27:YE:189:PRO:HA	2.02	0.42
29:YG:62:LEU:HA	29:YG:62:LEU:HD12	1.89	0.42
43:YY:83:THR:OG1	43:YY:84:ARG:O	2.30	0.42
1:QA:1466:C:H2'	1:QA:1467:G:O4'	2.19	0.42
1:QA:176:C:H2'	1:QA:177:C:H6	1.85	0.42
1:QA:685:G:H2'	1:QA:686:U:C6	2.55	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:QE:43:LEU:HD11	5:QE:132:ALA:HB1	2.02	0.42
49:R4:15:ILE:HG13	49:R4:32:TYR:HD1	1.84	0.42
24:RA:1055:A:N3	24:RA:1199:C:O2'	2.44	0.42
24:RA:847:A:O4'	24:RA:848:G:H3'	2.19	0.42
1:XA:1238:A:OP1	1:XA:1335:C:O2'	2.29	0.42
1:XA:1417:G:O2'	1:XA:1483:A:N6	2.53	0.42
1:XA:277:C:H5''	17:XQ:68:ARG:HH21	1.84	0.42
1:XA:382:A:H2'	1:XA:383:A:H8	1.85	0.42
1:XA:707:C:H2'	1:XA:708:C:H6	1.85	0.42
1:XA:35:G:N2	12:XL:118:SER:OG	2.46	0.42
12:XL:33:ARG:HA	12:XL:33:ARG:HD3	1.87	0.42
24:YA:139:A:H8	24:YA:1454:C:O2'	2.00	0.42
24:YA:1405:A:N6	24:YA:1418:U:C2	2.86	0.42
24:YA:1563:G:H4'	24:YA:1603:C:O2'	2.19	0.42
24:YA:240:A:OP1	53:Y8:7:HIS:NE2	2.52	0.42
24:YA:2815:C:H2'	24:YA:2816:G:C8	2.55	0.42
24:YA:374:U:H2'	24:YA:375:G:O4'	2.19	0.42
25:YB:7:G:H1	25:YB:114:G:H21	1.67	0.42
1:QA:373:A:H2'	1:QA:374:A:H8	1.84	0.42
10:QJ:40:LEU:HD13	10:QJ:71:LEU:HD13	2.01	0.42
24:RA:1249:A:HO2'	24:RA:1250:U:P	2.42	0.42
24:RA:1531:G:H1	24:RA:1550:C:N4	2.14	0.42
24:RA:2059:G:H2'	24:RA:2060:G:C8	2.54	0.42
24:RA:2081:A:H5'	24:RA:2082:A:OP2	2.19	0.42
24:RA:2524:C:H2'	24:RA:2525:G:O4'	2.19	0.42
24:RA:2859:U:H4'	24:RA:2878:A:C2	2.54	0.42
25:RB:8:U:O5'	37:RS:15:ARG:NH2	2.52	0.42
30:RH:97:ARG:NH2	30:RH:104:GLU:OE2	2.53	0.42
35:RQ:17:LEU:HA	35:RQ:17:LEU:HD23	1.79	0.42
35:RQ:71:ASP:N	35:RQ:71:ASP:OD1	2.44	0.42
1:XA:358:U:C4'	1:XA:358:U:C6	3.02	0.42
1:XA:452:A:N6	1:XA:480:U:C2	2.86	0.42
1:XA:885:G:H2'	1:XA:886:G:H8	1.84	0.42
4:XD:149:ALA:N	4:XD:152:SER:OG	2.52	0.42
9:XI:86:VAL:HG12	9:XI:92:TYR:HB2	2.02	0.42
54:Y9:18:ARG:HG3	54:Y9:23:VAL:HG22	2.01	0.42
24:YA:1627:A:H3'	24:YA:1628:G:C8	2.51	0.42
24:YA:2326:C:H2'	24:YA:2327:G:C8	2.53	0.42
24:YA:967:G:H4'	24:YA:2281:A:C6	2.55	0.42
27:YE:179:GLU:HG3	38:YT:9:LEU:HD21	2.01	0.42
24:YA:1298:G:C2	39:YU:33:ARG:HB3	2.55	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:YZ:35:ARG:HD2	44:YZ:35:ARG:HA	1.86	0.42
1:QA:1121:U:H2'	1:QA:1122:U:C6	2.55	0.42
1:QA:17:U:H2'	1:QA:18:C:H6	1.84	0.42
1:QA:269:C:H2'	1:QA:270:A:C8	2.55	0.42
1:QA:745:C:H2'	1:QA:746:A:C8	2.55	0.42
1:QA:812:C:H1'	1:QA:813:U:OP2	2.19	0.42
1:QA:920:U:H2'	1:QA:921:U:C6	2.55	0.42
11:QK:38:ASN:HA	11:QK:39:PRO:HD3	1.85	0.42
14:QN:9:LYS:HE3	14:QN:9:LYS:HB2	1.84	0.42
24:RA:1066:A:N1	24:RA:1186:U:H2'	2.34	0.42
24:RA:1394:G:H2'	24:RA:1395:A:H5''	2.01	0.42
24:RA:1583:C:H2'	24:RA:1584:G:O4'	2.20	0.42
24:RA:1683:C:H2'	24:RA:1684:A:C8	2.54	0.42
24:RA:1698:G:H2'	24:RA:1699:A:O4'	2.19	0.42
24:RA:2047:C:H2'	24:RA:2048:C:C6	2.54	0.42
24:RA:211:A:O2'	24:RA:212:A:O5'	2.38	0.42
24:RA:2173:G:H2'	24:RA:2174:G:H8	1.83	0.42
24:RA:965:G:N2	24:RA:2281:A:OP2	2.51	0.42
24:RA:265:U:H2'	24:RA:266:C:C6	2.55	0.42
24:RA:2869:G:H2'	24:RA:2870:A:C8	2.54	0.42
24:RA:779:C:H2'	24:RA:780:G:O4'	2.20	0.42
24:RA:898:U:H2'	24:RA:899:G:H8	1.84	0.42
26:RD:72:LYS:NZ	26:RD:99:ASP:OD2	2.36	0.42
39:RU:46:ALA:O	39:RU:50:ARG:HB2	2.20	0.42
44:RZ:61:LEU:O	44:RZ:64:GLY:N	2.52	0.42
1:XA:1228:C:H2'	1:XA:1229:A:H8	1.85	0.42
1:XA:1245:A:H2'	1:XA:1246:C:H6	1.84	0.42
1:XA:501:C:H2'	1:XA:502:G:C8	2.54	0.42
1:XA:5:U:O2'	1:XA:6:G:O5'	2.37	0.42
8:XH:40:ALA:HB2	8:XH:45:ILE:HG13	2.01	0.42
9:XI:43:ALA:HA	9:XI:74:ILE:HD13	2.01	0.42
20:XT:11:SER:HA	20:XT:13:LEU:HD23	2.02	0.42
22:XV:1:C:H2'	22:XV:2:G:C8	2.51	0.42
22:XV:61:C:H2'	22:XV:62:C:C6	2.55	0.42
50:Y5:41:PRO:O	50:Y5:44:THR:OG1	2.37	0.42
24:YA:160:G:H2'	24:YA:161:C:C6	2.55	0.42
24:YA:184:A:H5''	24:YA:185:A:O5'	2.20	0.42
24:YA:793:A:HO2'	24:YA:2623:U:HO2'	1.65	0.42
26:YD:183:ARG:NH2	26:YD:266:SER:HB2	2.35	0.42
24:YA:2762:A:H5''	30:YH:3:ARG:HH12	1.85	0.42
1:QA:1126:U:H3'	1:QA:1127:G:H8	1.85	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1226:C:H2'	13:QM:103:THR:HB	2.01	0.42
1:QA:328:C:H1'	1:QA:329:A:OP2	2.20	0.42
1:QA:908:A:H2'	1:QA:909:A:H8	1.84	0.42
2:QB:15:VAL:HG13	2:QB:210:SER:HB2	2.01	0.42
2:QB:26:PRO:O	2:QB:29:ALA:CB	2.67	0.42
4:QD:180:GLY:HA3	4:QD:182:LYS:HG2	2.01	0.42
13:QM:97:PRO:HD3	13:QM:110:ARG:HB3	2.01	0.42
17:QQ:98:LEU:HD23	17:QQ:98:LEU:HA	1.85	0.42
24:RA:1800:G:O2'	24:RA:1980:C:OP1	2.35	0.42
24:RA:656:A:N3	24:RA:2427:G:O2'	2.42	0.42
29:RG:41:GLN:NE2	29:RG:153:ARG:O	2.52	0.42
25:RB:45:A:O4'	29:RG:95:ARG:NH1	2.52	0.42
1:XA:1328:C:H2'	1:XA:1329:A:C8	2.55	0.42
1:XA:224:C:H2'	1:XA:225:C:C6	2.55	0.42
13:XM:91:ARG:HA	13:XM:91:ARG:HD2	1.90	0.42
22:XV:30:C:H2'	22:XV:31:G:H8	1.84	0.42
24:YA:1057:G:OP1	39:YU:77:SER:N	2.50	0.42
24:YA:2053:A:N3	24:YA:2467:G:O2'	2.41	0.42
24:YA:321:C:OP1	43:YY:87:LYS:NZ	2.50	0.42
31:YI:69:LYS:NZ	31:YI:135:GLU:OE1	2.53	0.42
24:YA:1053:C:H5''	32:YN:35:ARG:NH1	2.34	0.42
44:YZ:16:SER:OG	44:YZ:17:ALA:N	2.52	0.42
1:QA:1112:C:H42	3:QC:177:THR:HA	1.84	0.42
1:QA:17:U:H2'	1:QA:18:C:C6	2.55	0.42
1:QA:522:C:H41	12:QL:53:ARG:NH2	2.18	0.42
1:QA:881:G:OP2	12:QL:12:ARG:NH2	2.52	0.42
1:QA:909:A:N3	1:QA:1413:A:O2'	2.38	0.42
1:QA:924:C:H2'	1:QA:925:G:H8	1.85	0.42
1:QA:984:C:H2'	1:QA:985:C:C6	2.55	0.42
3:QC:43:LEU:O	3:QC:47:LEU:N	2.52	0.42
13:QM:37:THR:HA	13:QM:55:ARG:HH21	1.85	0.42
13:QM:74:VAL:O	13:QM:78:ILE:HG12	2.18	0.42
15:QO:7:GLU:H	15:QO:7:GLU:HG2	1.65	0.42
24:RA:1177:G:OP2	24:RA:2527:C:H4'	2.20	0.42
24:RA:2867:G:N2	24:RA:2870:A:OP2	2.35	0.42
24:RA:917:A:H2'	24:RA:918:U:O4'	2.20	0.42
26:RD:133:LEU:HD23	26:RD:136:ILE:HD12	2.02	0.42
30:RH:64:LEU:HD23	30:RH:64:LEU:HA	1.89	0.42
39:RU:92:ARG:CZ	40:RV:11:GLN:H	2.32	0.42
24:RA:509:A:O2'	43:RY:49:VAL:O	2.28	0.42
44:RZ:123:ASP:N	44:RZ:123:ASP:OD1	2.53	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1004:A:H1'	1:XA:1036:G:H22	1.85	0.42
1:XA:1346:A:H1'	1:XA:1348:U:C2	2.55	0.42
1:XA:390:C:H2'	1:XA:391:G:C8	2.55	0.42
8:XH:44:PHE:HB3	8:XH:80:ILE:HD11	2.02	0.42
1:XA:598:U:H4'	8:XH:94:TYR:CG	2.55	0.42
10:XJ:40:LEU:HA	10:XJ:40:LEU:HD23	1.93	0.42
16:XP:72:ARG:HA	16:XP:75:ARG:HB2	2.00	0.42
24:YA:1133:G:C4	24:YA:1135:G:H1'	2.54	0.42
24:YA:1766:G:H3'	24:YA:1767:A:C5'	2.50	0.42
24:YA:553:A:C2'	24:YA:554:A:O5'	2.68	0.42
1:QA:666:G:H2'	1:QA:667:G:H8	1.84	0.42
1:QA:688:G:H2'	1:QA:689:C:H6	1.85	0.42
24:RA:1636:U:H2'	24:RA:1637:G:H8	1.84	0.42
24:RA:2227:G:H3'	24:RA:2228:G:H8	1.85	0.42
26:RD:72:LYS:HD3	26:RD:75:ILE:HD12	2.02	0.42
29:RG:27:ASN:N	29:RG:30:GLU:OE1	2.53	0.42
24:RA:2735:G:O2'	36:RR:3:HIS:HB2	2.20	0.42
33:RO:120:GLU:HB2	38:RT:68:TYR:HE2	1.85	0.42
44:RZ:31:ARG:HG3	44:RZ:31:ARG:H	1.66	0.42
1:XA:1254:C:H2'	1:XA:1255:G:C8	2.55	0.42
1:XA:192:U:H2'	1:XA:193:C:H6	1.85	0.42
1:XA:237:C:OP2	17:XQ:40:LYS:NZ	2.50	0.42
7:XG:67:GLU:HA	7:XG:70:LYS:HB2	2.02	0.42
9:XI:67:GLY:O	9:XI:73:GLN:NE2	2.45	0.42
52:Y7:6:GLN:HA	52:Y7:7:PRO:HD3	1.89	0.42
24:YA:1476:C:H2'	24:YA:1477:U:C6	2.55	0.42
24:YA:1650:C:H2'	24:YA:1651:C:H6	1.85	0.42
24:YA:1765:U:H2'	24:YA:1766:G:O4'	2.20	0.42
24:YA:2549:U:H2'	24:YA:2550:C:H6	1.84	0.42
29:YG:63:ILE:HG22	29:YG:143:GLU:HB2	2.02	0.42
29:YG:82:LEU:HD23	29:YG:86:MET:HE2	2.02	0.42
30:YH:3:ARG:HG2	30:YH:6:ARG:HD2	2.02	0.42
30:YH:51:ARG:CB	30:YH:51:ARG:NH1	2.83	0.42
32:YN:39:ARG:NH1	32:YN:41:ASP:OD1	2.53	0.42
44:YZ:110:GLY:HA3	44:YZ:174:VAL:HG11	2.02	0.42
44:YZ:63:ASP:OD1	44:YZ:63:ASP:N	2.45	0.42
1:QA:1347:G:H2'	1:QA:1347:G:H8	1.58	0.41
1:QA:745:C:H2'	1:QA:746:A:H8	1.84	0.41
3:QC:189:ALA:HB3	3:QC:196:LEU:HB2	2.02	0.41
1:QA:1059:C:O2'	10:QJ:53:PRO:HD3	2.19	0.41
11:QK:33:THR:OG1	11:QK:34:ASP:N	2.53	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:RA:136:G:H2'	24:RA:138:G:N7	2.35	0.41
24:RA:1634:C:H2'	24:RA:1635:C:H6	1.85	0.41
24:RA:272:U:OP1	24:RA:273:G:N2	2.52	0.41
24:RA:2880:C:H2'	24:RA:2881:C:O4'	2.20	0.41
24:RA:611:U:H1'	28:RF:90:PHE:HB3	2.02	0.41
1:XA:1071:C:H2'	1:XA:1072:G:C8	2.55	0.41
1:XA:1293:G:H2'	1:XA:1294:G:C8	2.55	0.41
1:XA:1374:A:O2'	7:XG:28:ASN:HB3	2.20	0.41
1:XA:294:U:H2'	1:XA:295:C:C6	2.55	0.41
1:XA:770:C:O2'	1:XA:899:C:N3	2.43	0.41
4:XD:196:LEU:HA	4:XD:196:LEU:HD23	1.74	0.41
11:XK:19:ALA:HA	11:XK:32:ILE:HA	2.02	0.41
11:XK:81:ASP:OD1	11:XK:107:SER:OG	2.32	0.41
24:YA:1035:G:OP2	48:Y3:11:SER:OG	2.24	0.41
24:YA:1067:A:H62	24:YA:1186:U:H3	1.68	0.41
24:YA:1493:C:HO2'	24:YA:1592:A:HO2'	1.67	0.41
24:YA:1827:U:H2'	24:YA:1828:C:H6	1.85	0.41
24:YA:2457:G:P	28:YF:74:ARG:HH22	2.43	0.41
34:YP:57:THR:O	34:YP:61:ARG:HG3	2.20	0.41
42:YX:33:LYS:HA	42:YX:33:LYS:HD3	1.93	0.41
1:QA:1130:A:N6	1:QA:1144:G:H21	2.18	0.41
1:QA:191(E):G:H2'	1:QA:191(F):U:C6	2.56	0.41
1:QA:254:G:OP1	17:QQ:68:ARG:HB3	2.21	0.41
2:QB:21:ARG:HG2	2:QB:22:LYS:HB3	2.03	0.41
11:QK:61:ALA:HB1	11:QK:94:ALA:HB2	2.02	0.41
1:QA:376:G:P	16:QP:67:THR:HG21	2.60	0.41
24:RA:1549:U:H2'	24:RA:1550:C:H6	1.85	0.41
24:RA:1464:G:N1	24:RA:1626:A:OP2	2.43	0.41
24:RA:1727:U:H2'	24:RA:1728:G:O4'	2.20	0.41
24:RA:2842:U:O2'	24:RA:2843:G:O5'	2.37	0.41
24:RA:937:A:H2'	24:RA:938:G:H8	1.84	0.41
24:RA:2323:A:H1'	29:RG:82:LEU:HD11	2.02	0.41
1:XA:1250:A:H2'	1:XA:1251:A:C8	2.55	0.41
1:XA:345:C:H1'	1:XA:346:G:OP2	2.19	0.41
4:XD:194:LEU:HD12	4:XD:195:ALA:H	1.84	0.41
4:XD:72:GLU:OE2	4:XD:207:TYR:OH	2.24	0.41
1:XA:230:G:O2'	16:XP:25:ARG:NH2	2.53	0.41
24:YA:1122:C:H2'	24:YA:1123:A:H5''	2.02	0.41
24:YA:1342:G:OP1	24:YA:2721:G:O2'	2.20	0.41
24:YA:1766:G:H3'	24:YA:1767:A:H5''	2.02	0.41
28:YF:11:VAL:HG22	28:YF:125:LEU:HB2	2.02	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:YG:173:LEU:HA	29:YG:176:LEU:HB2	2.02	0.41
31:YI:77:LEU:HD13	31:YI:101:LEU:HD13	2.01	0.41
40:YV:65:GLY:HA3	40:YV:91:TYR:CZ	2.55	0.41
1:QA:1032(A):A:H3'	1:QA:1032(B):G:H4'	2.02	0.41
1:QA:1443:G:H5'	1:QA:1446:A:OP2	2.20	0.41
1:QA:337:C:H2'	1:QA:338:A:H8	1.86	0.41
1:QA:410:G:H2'	1:QA:429:U:C5	2.54	0.41
1:QA:45:U:H2'	1:QA:46:G:C8	2.56	0.41
1:QA:646:U:H2'	1:QA:647:C:C6	2.54	0.41
1:QA:662:G:H2'	1:QA:663:A:H8	1.85	0.41
10:QJ:6:ILE:HA	10:QJ:97:GLU:O	2.20	0.41
24:RA:1674:G:H2'	24:RA:1675:U:C6	2.55	0.41
24:RA:1862:G:H2'	24:RA:1863:C:C6	2.54	0.41
24:RA:2708:U:H2'	24:RA:2709:G:H8	1.85	0.41
24:RA:607:C:OP2	39:RU:10:ARG:NH2	2.53	0.41
24:RA:864:C:O2'	24:RA:886:U:H5''	2.19	0.41
30:RH:77:LYS:HD2	30:RH:77:LYS:HA	1.86	0.41
35:RQ:35:VAL:HG13	35:RQ:130:LYS:HB3	2.02	0.41
1:XA:1245:A:H2'	1:XA:1246:C:C6	2.56	0.41
1:XA:1368:G:OP1	9:XI:111:ARG:NH1	2.52	0.41
1:XA:337:C:H2'	1:XA:338:A:C8	2.55	0.41
1:XA:356:A:N3	1:XA:368:U:O2'	2.32	0.41
24:YA:240:A:P	53:Y8:7:HIS:HE2	2.43	0.41
24:YA:1104:G:O2'	24:YA:1105:G:H5'	2.20	0.41
24:YA:1276:C:H2'	24:YA:1277:G:C8	2.56	0.41
24:YA:2524:C:H2'	24:YA:2525:G:O4'	2.20	0.41
24:YA:287:G:H1'	24:YA:288:U:OP2	2.20	0.41
29:YG:18:GLU:OE2	29:YG:21:ARG:NH2	2.39	0.41
31:YI:100:ALA:O	31:YI:104:GLN:N	2.53	0.41
34:YP:106:LEU:HD23	34:YP:106:LEU:HA	1.94	0.41
1:QA:1080:A:O5'	5:QE:14:ARG:NH2	2.52	0.41
1:QA:337:C:H2'	1:QA:338:A:C8	2.55	0.41
2:QB:8:LYS:HD3	2:QB:9:GLU:HG3	2.01	0.41
8:QH:100:ILE:HA	8:QH:101:PRO:HD3	1.89	0.41
22:QV:15:G:N2	22:QV:21(B):A:N3	2.69	0.41
24:RA:2824:C:O2'	50:R5:29:THR:OG1	2.18	0.41
29:RG:65:GLY:HA3	49:R4:9:LEU:HD21	2.02	0.41
24:RA:854:U:OP2	34:RP:41:ARG:NH2	2.53	0.41
1:XA:1295:G:N2	1:XA:1302:U:O4	2.49	0.41
1:XA:1357:A:H61	1:XA:1365:G:H1	1.69	0.41
1:XA:34:C:H2'	1:XA:35:G:C8	2.55	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:580:U:H2'	1:XA:581:G:O4'	2.21	0.41
1:XA:677:U:O2	1:XA:777:A:O2'	2.38	0.41
1:XA:745:C:H2'	1:XA:746:A:C8	2.49	0.41
1:XA:990:C:H2'	1:XA:991:U:C2	2.55	0.41
8:XH:29:SER:HB3	8:XH:32:LYS:HG3	2.03	0.41
11:XK:38:ASN:HA	11:XK:39:PRO:HD3	1.83	0.41
24:YA:2401:G:H5''	24:YA:2402:U:O4'	2.20	0.41
24:YA:2623:U:OP2	24:YA:2623:U:H6	2.03	0.41
38:YT:92:GLY:O	38:YT:120:ARG:NH2	2.53	0.41
1:QA:1427:U:H2'	1:QA:1428:A:C8	2.55	0.41
2:QB:8:LYS:HG3	2:QB:10:LEU:HB3	2.02	0.41
2:QB:47:THR:HA	2:QB:202:PRO:HG2	2.02	0.41
2:QB:85:ALA:O	2:QB:88:ALA:N	2.49	0.41
4:QD:38:TYR:HA	4:QD:39:PRO:HD3	1.85	0.41
10:QJ:9:ARG:HE	10:QJ:95:GLU:HG2	1.85	0.41
11:QK:33:THR:HA	11:QK:40:ILE:HG12	2.02	0.41
24:RA:2382:G:H2'	24:RA:2383:G:O4'	2.19	0.41
24:RA:2549:U:H2'	24:RA:2550:C:C6	2.54	0.41
24:RA:2842:U:H4'	24:RA:2843:G:H5''	2.02	0.41
24:RA:38:A:H2'	24:RA:39:C:C6	2.55	0.41
27:RE:55:ASN:HA	27:RE:56:PRO:HD3	1.94	0.41
33:RO:104:ARG:NH1	33:RO:121:VAL:O	2.53	0.41
34:RP:63:PRO:HB2	53:R8:30:ARG:HH21	1.84	0.41
44:RZ:164:ALA:H	44:RZ:165:VAL:HG23	1.86	0.41
1:XA:1499:A:OP1	1:XA:1500:A:OP2	2.38	0.41
1:XA:157:G:H2'	1:XA:158:G:H8	1.85	0.41
1:XA:255:G:OP1	17:XQ:69:LYS:NZ	2.53	0.41
1:XA:509:A:H2'	1:XA:510:A:C8	2.55	0.41
8:XH:11:THR:HG22	8:XH:14:ARG:NH1	2.34	0.41
12:XL:70:ILE:HG12	12:XL:77:LEU:HD12	2.03	0.41
15:XO:67:LEU:HB3	15:XO:78:TYR:HE1	1.86	0.41
17:XQ:5:VAL:HG22	17:XQ:60:ILE:HD13	2.03	0.41
24:YA:278:G:H2'	24:YA:279:G:H8	1.86	0.41
25:YB:48:A:H2'	25:YB:49:C:C6	2.56	0.41
27:YE:36:ARG:NH2	27:YE:88:GLY:O	2.53	0.41
1:QA:481:G:O2'	1:QA:482:A:O5'	2.39	0.41
6:QF:12:PRO:HB3	6:QF:58:GLY:HA2	2.03	0.41
8:QH:3:THR:OG1	8:QH:4:ASP:N	2.54	0.41
17:QQ:60:ILE:HA	17:QQ:60:ILE:HD12	1.91	0.41
54:R9:14:CYS:HA	54:R9:27:CYS:HB2	2.02	0.41
24:RA:1328:U:H2'	24:RA:1329:G:O4'	2.21	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:RA:2816:G:H2'	24:RA:2817:G:C8	2.55	0.41
24:RA:776:G:C5	26:RD:208:LYS:HB2	2.56	0.41
24:RA:964:A:N3	25:RB:80:U:O2'	2.44	0.41
27:RE:78:LEU:O	27:RE:79:ARG:NH1	2.45	0.41
38:RT:56:GLY:O	38:RT:59:THR:OG1	2.34	0.41
39:RU:34:LYS:HA	39:RU:34:LYS:HD3	1.80	0.41
1:XA:1277:C:O2'	1:XA:1279:A:H1'	2.20	0.41
1:XA:1409:C:H2'	1:XA:1410:G:H8	1.86	0.41
1:XA:378:G:O6	1:XA:386:C:N4	2.53	0.41
1:XA:673:G:O3'	6:XF:87:ARG:NH2	2.54	0.41
1:XA:865:A:H2'	1:XA:866:C:C6	2.56	0.41
1:XA:406:G:N2	4:XD:119:GLN:HE22	2.17	0.41
8:XH:29:SER:OG	8:XH:31:PHE:N	2.50	0.41
19:XS:40:ILE:HD11	19:XS:71:LEU:HA	2.03	0.41
47:Y2:18:PRO:HA	47:Y2:21:LEU:HB2	2.03	0.41
49:Y4:14:ILE:HB	49:Y4:22:ILE:HB	2.03	0.41
24:YA:1109:G:N2	24:YA:1122:C:H1'	2.35	0.41
24:YA:594:A:OP2	24:YA:2052:A:N6	2.53	0.41
24:YA:2603:C:H2'	24:YA:2604:G:H8	1.85	0.41
27:YE:77:ILE:HD13	27:YE:195:LEU:HD13	2.02	0.41
29:YG:71:THR:N	29:YG:89:GLY:O	2.53	0.41
1:QA:22:G:H2'	1:QA:23:C:H6	1.86	0.41
1:QA:407:G:H2'	1:QA:408:A:C8	2.55	0.41
22:QV:28:U:H2'	22:QV:29:U:C6	2.56	0.41
47:R2:63:VAL:HA	47:R2:66:GLU:HB2	2.02	0.41
24:RA:1125:C:H2'	24:RA:1126:C:O4'	2.21	0.41
24:RA:1195:G:H2'	24:RA:1196:C:C6	2.56	0.41
24:RA:1802:C:H2'	24:RA:1803:G:C8	2.55	0.41
24:RA:2312:G:N1	24:RA:2329:C:N3	2.69	0.41
24:RA:2662:U:H2'	24:RA:2663:C:H6	1.86	0.41
24:RA:2810:C:H2'	24:RA:2811:A:C8	2.55	0.41
24:RA:64:C:H1'	24:RA:482:C:H42	1.85	0.41
26:RD:177:LEU:HD23	26:RD:177:LEU:HA	1.87	0.41
30:RH:137:ASP:OD1	30:RH:138:LYS:N	2.53	0.41
33:RO:63:VAL:HB	33:RO:102:VAL:HG12	2.01	0.41
1:XA:765:G:N2	1:XA:813:U:OP2	2.54	0.41
1:XA:958:A:H2'	1:XA:959:A:C8	2.56	0.41
5:XE:12:LEU:HB3	5:XE:31:LEU:HB3	2.02	0.41
6:XF:39:LYS:HG3	6:XF:62:TRP:HZ3	1.85	0.41
24:YA:1493:C:O2'	24:YA:1592:A:O2'	2.37	0.41
24:YA:2036:A:H2'	24:YA:2037:A:C8	2.55	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:YA:2686:G:H2'	24:YA:2687:A:C8	2.56	0.41
24:YA:325:G:OP2	43:YY:84:ARG:NH2	2.53	0.41
26:YD:14:ARG:HH11	26:YD:14:ARG:HD2	1.74	0.41
25:YB:48:A:P	37:YS:30:ARG:HH12	2.43	0.41
42:YX:10:ALA:HB3	42:YX:29:TRP:HB2	2.02	0.41
44:YZ:10:ARG:HH21	44:YZ:26:GLY:H	1.68	0.41
4:QD:23:GLY:N	4:QD:26:CYS:SG	2.91	0.41
24:RA:1827:U:H2'	24:RA:1828:C:H6	1.86	0.41
24:RA:2050:U:H2'	24:RA:2051:G:C8	2.56	0.41
24:RA:2318:C:H3'	24:RA:2319:G:H5''	2.01	0.41
24:RA:2318:C:OP2	24:RA:2319:G:H2'	2.21	0.41
24:RA:2389:A:H2'	24:RA:2390:A:C8	2.55	0.41
24:RA:609:A:N1	24:RA:856:G:O2'	2.41	0.41
24:RA:807:G:H2'	24:RA:808:A:O4'	2.19	0.41
27:RE:185:LYS:HA	27:RE:185:LYS:HD2	1.87	0.41
31:RI:14:ASP:H	31:RI:17:GLN:HB3	1.86	0.41
35:RQ:77:LYS:NZ	35:RQ:80:GLU:OE1	2.41	0.41
1:XA:1022:G:H2'	1:XA:1023:G:H8	1.86	0.41
1:XA:1412:C:H2'	1:XA:1413:A:H8	1.86	0.41
1:XA:244:U:H4'	1:XA:245:C:O5'	2.20	0.41
1:XA:294:U:H2'	1:XA:295:C:H6	1.85	0.41
1:XA:406:G:H2'	1:XA:407:G:H8	1.85	0.41
1:XA:711:G:H2'	1:XA:712:A:C8	2.56	0.41
1:XA:868:C:H2'	1:XA:869:G:O4'	2.21	0.41
5:XE:78:HIS:CE1	5:XE:80:ILE:HD13	2.55	0.41
7:XG:79:ARG:HD3	7:XG:79:ARG:HA	1.86	0.41
16:XP:15:PRO:HD2	16:XP:42:ARG:HD2	2.02	0.41
23:XX:19:G:H8	23:XX:19:G:O5'	2.03	0.41
24:YA:1195:G:H2'	24:YA:1196:C:C6	2.55	0.41
24:YA:2417:G:H5'	34:YP:75:ILE:HD13	2.03	0.41
24:YA:2473:C:H2'	24:YA:2474:U:C6	2.55	0.41
25:YB:89(B):A:H2'	25:YB:90:C:O4'	2.21	0.41
28:YF:68:LYS:HG2	28:YF:69:HIS:CD2	2.56	0.41
30:YH:126:PRO:HG2	30:YH:130:ARG:HH21	1.85	0.41
1:QA:1066:C:H42	1:QA:1191:A:N6	2.19	0.41
1:QA:1446:A:H4'	38:RT:125:ARG:HH12	1.86	0.41
1:QA:209:U:O2'	1:QA:210:U:H4'	2.21	0.41
1:QA:452:A:H2'	1:QA:453:A:H8	1.85	0.41
2:QB:189:ASP:OD1	2:QB:189:ASP:N	2.53	0.41
8:QH:32:LYS:HA	8:QH:35:ILE:HD12	2.03	0.41
9:QI:96:LEU:O	9:QI:101:PHE:N	2.47	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:RA:142:G:H2'	24:RA:143:C:C6	2.55	0.41
24:RA:2619:G:H2'	24:RA:2620:G:O4'	2.21	0.41
24:RA:266:C:H42	24:RA:278:G:H1	1.69	0.41
24:RA:771:U:H2'	24:RA:772:G:O4'	2.21	0.41
27:RE:34:VAL:HG12	27:RE:72:VAL:HG11	2.03	0.41
29:RG:111:LEU:HA	29:RG:114:ILE:HG12	2.02	0.41
29:RG:55:LYS:HD2	29:RG:58:GLN:HG2	2.02	0.41
24:RA:2686:G:H5''	33:RO:26:LYS:HE3	2.03	0.41
37:RS:39:ILE:O	37:RS:48:LEU:N	2.47	0.41
1:XA:112:G:H4'	1:XA:389:A:H4'	2.03	0.41
1:XA:1443:G:H5'	1:XA:1446:A:OP2	2.21	0.41
1:XA:584:G:H2'	1:XA:585:G:C8	2.55	0.41
2:XB:95:GLN:HG3	2:XB:147:LYS:HG3	2.03	0.41
5:XE:92:LYS:N	5:XE:119:LEU:O	2.41	0.41
10:XJ:42:THR:HG21	10:XJ:66:ARG:HD2	2.03	0.41
22:XV:32:U:H3	22:XV:38:A:N6	2.18	0.41
46:Y1:5:CYS:SG	46:Y1:8:SER:OG	2.71	0.41
24:YA:1101:G:H1	24:YA:1150:C:H42	1.69	0.41
24:YA:2021:C:H5''	24:YA:2736:C:O2'	2.21	0.41
24:YA:2047:C:H2'	24:YA:2048:C:C6	2.55	0.41
24:YA:2816:G:H2'	24:YA:2817:G:H8	1.86	0.41
24:YA:844:C:OP1	28:YF:60:SER:OG	2.33	0.41
26:YD:79:VAL:HG23	26:YD:114:GLY:H	1.85	0.41
29:YG:55:LYS:HA	29:YG:55:LYS:HD2	1.87	0.41
41:YW:113:LYS:HD2	41:YW:113:LYS:HA	1.93	0.41
44:YZ:23:LYS:NZ	44:YZ:40:ASP:OD1	2.37	0.41
1:QA:1064:G:H1'	1:QA:1066:C:C6	2.56	0.41
1:QA:1358:U:OP1	14:QN:35:ARG:HB2	2.21	0.41
15:QO:87:ILE:HG22	15:QO:88:ARG:H	1.85	0.41
16:QP:55:ARG:HD2	16:QP:55:ARG:HA	1.89	0.41
24:RA:1119:A:H4'	24:RA:1120:G:OP1	2.21	0.41
24:RA:1248:G:H3'	24:RA:1249:A:H5''	2.02	0.41
24:RA:1276:C:H2'	24:RA:1277:G:C8	2.55	0.41
24:RA:2658:C:OP2	24:RA:2745:G:O2'	2.30	0.41
24:RA:2694:U:H6	24:RA:2694:U:C5'	2.33	0.41
24:RA:323:A:OP2	24:RA:323:A:H8	2.04	0.41
24:RA:537:G:H4'	24:RA:538:A:O5'	2.20	0.41
24:RA:910:A:H2'	24:RA:911:G:H8	1.85	0.41
28:RF:158:THR:OG1	28:RF:159:GLY:N	2.54	0.41
29:RG:172:LEU:HD23	29:RG:173:LEU:HD23	2.03	0.41
35:RQ:5:ARG:HA	35:RQ:5:ARG:HD3	1.80	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:358:U:C6	1:XA:358:U:O5'	2.73	0.41
1:XA:599:C:O2'	8:XH:129:VAL:O	2.28	0.41
12:XL:77:LEU:HD23	12:XL:77:LEU:HA	1.78	0.41
1:XA:976:G:P	14:XN:32:SER:H	2.44	0.41
16:XP:43:LYS:HG2	16:XP:48:TRP:CE3	2.56	0.41
24:YA:1480:A:H61	24:YA:1605:A:N6	2.19	0.41
24:YA:30:G:H2'	24:YA:31:C:C6	2.56	0.41
24:YA:598:A:OP2	24:YA:2511:C:O2'	2.31	0.41
25:YB:89(A):G:C8	25:YB:89(A):G:H3'	2.56	0.41
40:YV:72:VAL:HG13	40:YV:85:LYS:HB3	2.03	0.41
1:QA:1060:C:H2'	1:QA:1061:G:H8	1.86	0.41
1:QA:1150:U:H2'	1:QA:1151:A:C8	2.56	0.41
1:QA:125:U:H2'	1:QA:126:G:C8	2.56	0.41
1:QA:382:A:H2'	1:QA:383:A:H8	1.86	0.41
1:QA:406:G:H21	4:QD:119:GLN:HE22	1.68	0.41
1:QA:790:A:H2'	1:QA:791:G:C8	2.56	0.41
3:QC:50:ALA:HB1	3:QC:70:VAL:HG11	2.03	0.41
4:QD:108:LEU:HD21	4:QD:183:GLY:HA3	2.03	0.41
1:QA:690:G:H22	11:QK:55:LYS:HZ1	1.68	0.41
22:QV:35:G:H2'	22:QV:36:G:C8	2.56	0.41
22:QV:12:G:H5''	24:RA:1930:C:O2'	2.21	0.41
24:RA:2424:A:H2'	24:RA:2425:G:O4'	2.21	0.41
24:RA:2705:A:H2'	24:RA:2706:G:C8	2.55	0.41
24:RA:374:U:H2'	24:RA:375:G:O4'	2.20	0.41
24:RA:720:C:H5''	28:RF:81:PRO:HD2	2.03	0.41
24:RA:893:C:O2'	24:RA:894:U:OP2	2.31	0.41
35:RQ:39:PRO:HB3	35:RQ:99:PRO:HD3	2.03	0.41
37:RS:26:LEU:HB3	37:RS:87:PHE:HA	2.02	0.41
44:RZ:69:THR:HG22	44:RZ:90:VAL:HA	2.03	0.41
1:XA:113:G:H2'	1:XA:114:U:C6	2.56	0.41
1:XA:1288:A:N1	1:XA:1371:G:H1'	2.36	0.41
1:XA:1513:A:H2'	1:XA:1514:C:C6	2.56	0.41
1:XA:406:G:H21	4:XD:119:GLN:HE22	1.67	0.41
2:XB:75:LYS:HA	2:XB:78:GLN:HG3	2.03	0.41
19:XS:53:ASN:HB3	19:XS:75:ALA:HB1	2.03	0.41
24:YA:1490:G:H2'	24:YA:1492:C:C5	2.56	0.41
24:YA:1580:G:N3	24:YA:1580:G:H2'	2.36	0.41
24:YA:2258:G:H2'	24:YA:2259:A:H8	1.83	0.41
24:YA:2705:A:H2'	24:YA:2706:G:H8	1.86	0.41
24:YA:362:G:OP1	43:YY:4:LYS:NZ	2.44	0.41
27:YE:101:ARG:NH1	27:YE:169:ASN:O	2.54	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1002:G:H2'	1:QA:1003:G:C8	2.52	0.40
1:QA:1004:A:H2'	1:QA:1005:A:O4'	2.21	0.40
1:QA:1431:C:H2'	1:QA:1432:G:O4'	2.21	0.40
1:QA:410:G:H21	1:QA:432:A:N6	2.15	0.40
6:QF:15:ASP:OD2	6:QF:17:SER:OG	2.30	0.40
7:QG:13:GLN:HA	7:QG:14:PRO:HD3	1.94	0.40
7:QG:15:ASP:HB2	7:QG:23:VAL:HB	2.04	0.40
17:QQ:57:VAL:HG12	17:QQ:76:LEU:HA	2.02	0.40
22:QV:37:1MG:HN21	22:QV:37:1MG:HM11	1.71	0.40
24:RA:1545:C:H2'	24:RA:1546:G:H8	1.86	0.40
24:RA:1549:U:H2'	24:RA:1550:C:C6	2.56	0.40
24:RA:2167:C:H2'	24:RA:2169:G:H22	1.86	0.40
24:RA:2540:U:H2'	24:RA:2542:A:O5'	2.21	0.40
24:RA:2552:C:H2'	24:RA:2553:A:O4'	2.20	0.40
24:RA:422:U:H2'	24:RA:423:G:C8	2.56	0.40
29:RG:94:LEU:HD22	29:RG:98:ARG:HB3	2.03	0.40
32:RN:7:LYS:HE3	32:RN:7:LYS:HB3	1.94	0.40
41:RW:86:LEU:HD22	41:RW:96:ILE:HD11	2.02	0.40
1:XA:359:U:P	1:XA:359:U:C3'	3.08	0.40
15:XO:4:THR:OG1	15:XO:5:LYS:N	2.54	0.40
19:XS:77:THR:H	19:XS:78:ARG:NH1	2.19	0.40
20:XT:79:ARG:O	20:XT:83:ARG:HG3	2.21	0.40
24:YA:2087:C:H2'	24:YA:2088:C:H6	1.86	0.40
24:YA:2764:G:H4'	30:YH:4:ILE:HD11	2.03	0.40
24:YA:469:A:H5''	24:YA:470:C:OP1	2.21	0.40
24:YA:821:A:H2	24:YA:834:U:O2'	2.03	0.40
26:YD:245:PRO:HA	26:YD:246:PRO:HD3	1.85	0.40
32:YN:39:ARG:HD3	32:YN:48:MET:HE2	2.02	0.40
1:QA:1010:G:H2'	1:QA:1011:G:C8	2.54	0.40
1:QA:324:G:N2	1:QA:327:A:OP2	2.53	0.40
2:QB:233:SER:OG	2:QB:233:SER:O	2.31	0.40
4:QD:129:ASN:HA	4:QD:145:GLU:HB3	2.02	0.40
9:QI:20:ARG:HA	9:QI:21:PRO:HD3	1.94	0.40
13:QM:107:ALA:H	13:QM:108:ARG:HH11	1.68	0.40
17:QQ:6:LEU:HD22	17:QQ:23:VAL:HG11	2.03	0.40
24:RA:1067:A:H8	24:RA:1068:G:H5''	1.86	0.40
24:RA:1151:U:H2'	24:RA:1152:G:C8	2.56	0.40
24:RA:1232:G:H5''	40:RV:81:TYR:CE1	2.56	0.40
24:RA:1273:G:OP2	39:RU:16:LYS:NZ	2.38	0.40
1:QA:784:C:H4'	24:RA:1868:C:OP1	2.21	0.40
24:RA:2235:G:H5''	26:RD:269:PHE:HZ	1.86	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:RD:83:GLU:HB2	26:RD:92:ILE:HG13	2.03	0.40
24:RA:1298:G:C2	39:RU:33:ARG:HB3	2.56	0.40
40:RV:20:LEU:HA	40:RV:20:LEU:HD12	1.89	0.40
1:XA:1287:A:H2'	1:XA:1288:A:C8	2.56	0.40
1:XA:1355:G:H2'	1:XA:1356:G:H8	1.85	0.40
1:XA:403:C:OP2	4:XD:74:GLN:NE2	2.52	0.40
11:XK:21:ILE:HB	11:XK:84:VAL:HG12	2.02	0.40
22:XV:57:G:O2'	22:XV:58:A:H5'	2.20	0.40
24:YA:1068:G:H1'	24:YA:1069:U:OP2	2.21	0.40
24:YA:1255:A:H4'	24:YA:1256:U:O5'	2.21	0.40
24:YA:1298:G:N3	39:YU:33:ARG:HD2	2.36	0.40
24:YA:1378:G:H21	24:YA:1656:A:H8	1.69	0.40
24:YA:1741:C:H1'	24:YA:1742:G:OP2	2.21	0.40
24:YA:2414:C:O2'	24:YA:2415:C:P	2.79	0.40
24:YA:1684:A:H4'	24:YA:2723:A:O2'	2.21	0.40
24:YA:422:U:H2'	24:YA:423:G:N7	2.37	0.40
24:YA:471:C:H2'	24:YA:472:G:O4'	2.20	0.40
24:YA:82:G:O2'	24:YA:100:G:N2	2.53	0.40
27:YE:28:ALA:HB3	27:YE:93:VAL:HG12	2.02	0.40
28:YF:24:LEU:HD23	28:YF:24:LEU:HA	1.83	0.40
5:QE:50:GLU:HB2	5:QE:53:LEU:HD13	2.03	0.40
11:QK:44:SER:HB3	11:QK:47:VAL:HG23	2.03	0.40
22:QV:40:G:H2'	22:QV:41:A:H8	1.86	0.40
53:R8:7:HIS:HB3	53:R8:59:LYS:HB3	2.04	0.40
24:RA:1369:U:OP1	41:RW:98:LYS:NZ	2.47	0.40
24:RA:2108:U:H2'	24:RA:2109:G:C8	2.56	0.40
24:RA:2303:U:H2'	24:RA:2304:C:H6	1.86	0.40
24:RA:664:U:H2'	24:RA:665:C:H6	1.87	0.40
30:RH:80:SER:OG	30:RH:80:SER:O	2.34	0.40
40:RV:35:LEU:HA	40:RV:36:PRO:HD3	1.85	0.40
40:RV:6:LYS:HD2	40:RV:11:GLN:HG2	2.04	0.40
1:XA:1350:A:OP1	9:XI:121:ARG:NH1	2.54	0.40
1:XA:147:G:H21	1:XA:1451:A:H61	1.70	0.40
1:XA:436:C:H2'	1:XA:437:U:C6	2.56	0.40
1:XA:596:C:H2'	1:XA:597:G:C8	2.57	0.40
1:XA:601:C:H2'	1:XA:602:A:C8	2.57	0.40
13:XM:87:TYR:HA	13:XM:90:LEU:HG	2.03	0.40
14:XN:27:CYS:HB3	14:XN:43:CYS:SG	2.62	0.40
1:XA:108:G:C6	20:XT:15:ARG:HD3	2.56	0.40
51:Y6:18:ARG:HH11	51:Y6:18:ARG:HD2	1.74	0.40
24:YA:1107:U:H3'	24:YA:1108:G:C5'	2.51	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:YA:1249:A:H1'	24:YA:1251:G:C5	2.56	0.40
24:YA:1910:G:N2	24:YA:1910:G:OP2	2.52	0.40
24:YA:2339:A:H2'	24:YA:2340:A:C8	2.56	0.40
24:YA:2664:C:H2'	24:YA:2665:U:O4'	2.21	0.40
24:YA:2724:U:HO2'	24:YA:2725:A:H8	1.63	0.40
24:YA:441:C:H2'	24:YA:442:A:C8	2.56	0.40
24:YA:585:U:H6	24:YA:585:U:H2'	1.74	0.40
25:YB:89(B):A:C3'	25:YB:89(B):A:C8	3.05	0.40
37:YS:10:ARG:HG2	37:YS:91:PRO:HA	2.03	0.40
37:YS:67:ARG:HG2	37:YS:71:ARG:HE	1.86	0.40
1:QA:1003:G:N2	1:QA:1004:A:O2'	2.55	0.40
1:QA:444:C:H2'	1:QA:445:G:C8	2.56	0.40
1:QA:728:A:H2'	1:QA:729:A:H8	1.86	0.40
1:QA:194:C:O3'	20:QT:68:LYS:HD2	2.20	0.40
1:QA:1230:C:H5'	22:QV:30:C:H5''	2.04	0.40
45:R0:65:GLY:HA3	45:R0:83:PRO:HA	2.04	0.40
47:R2:29:LYS:HE3	47:R2:57:ILE:HG21	2.04	0.40
52:R7:29:LYS:HE2	52:R7:29:LYS:HB3	1.86	0.40
24:RA:1227:A:H2'	24:RA:1228:G:C8	2.56	0.40
24:RA:1362:U:H2'	24:RA:1363:A:H8	1.85	0.40
24:RA:1471:G:H2'	24:RA:1472:G:C8	2.57	0.40
24:RA:1712:A:O2'	24:RA:1713:G:C5'	2.69	0.40
24:RA:1782:C:H2'	24:RA:1783:C:C6	2.57	0.40
24:RA:868:A:H2'	24:RA:991:G:H5''	2.03	0.40
30:RH:9:ILE:N	30:RH:69:ARG:HH21	2.19	0.40
36:RR:104:ARG:HG3	36:RR:107:ASP:HB3	2.03	0.40
41:RW:71:VAL:HA	41:RW:107:LEU:HD23	2.03	0.40
44:RZ:23:LYS:NZ	44:RZ:40:ASP:OD2	2.54	0.40
44:RZ:19:ARG:NH1	44:RZ:84:GLU:O	2.54	0.40
1:XA:1015:A:H2'	1:XA:1016:A:C8	2.57	0.40
1:XA:1414:U:H2'	1:XA:1415:G:H8	1.86	0.40
1:XA:446:G:H2'	1:XA:447:G:O4'	2.21	0.40
1:XA:992:U:O2'	1:XA:993:G:OP2	2.34	0.40
2:XB:187:LEU:HA	2:XB:201:ILE:HB	2.04	0.40
3:XC:184:TYR:HE1	3:XC:199:LYS:HB3	1.87	0.40
24:YA:1074:A:N6	24:YA:1171:G:H2'	2.36	0.40
24:YA:2087:C:H2'	24:YA:2088:C:C6	2.56	0.40
24:YA:2748:G:H2'	24:YA:2749:G:C8	2.57	0.40
24:YA:869:U:H2'	24:YA:870:G:C8	2.57	0.40
24:YA:903:C:HO2'	24:YA:904:C:P	2.44	0.40
25:YB:85:G:H1	25:YB:92:G:N2	2.17	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:YI:44:LEU:HA	31:YI:44:LEU:HD23	1.92	0.40
1:QA:1228:C:H2'	1:QA:1229:A:H8	1.86	0.40
1:QA:1322:C:OP1	1:QA:1322:C:H6	2.03	0.40
1:QA:1526:G:O2'	1:QA:1527:C:H5'	2.22	0.40
1:QA:678:U:H2'	1:QA:679:C:C6	2.56	0.40
1:QA:782:A:H62	1:QA:800:G:N2	2.19	0.40
1:QA:945:G:N2	1:QA:1334:G:O2'	2.55	0.40
1:QA:950:U:H3	1:QA:1231:G:H1	1.69	0.40
4:QD:68:TYR:OH	4:QD:98:GLU:OE1	2.37	0.40
11:QK:58:PRO:HB2	11:QK:93:GLN:HG3	2.02	0.40
24:RA:1400:A:OP1	26:RD:38:LYS:NZ	2.45	0.40
24:RA:1475:G:H2'	24:RA:1476:C:C6	2.56	0.40
24:RA:2506:G:OP1	45:R0:3:HIS:N	2.54	0.40
24:RA:552:C:N3	24:RA:2792:U:H5''	2.37	0.40
24:RA:599:U:H2'	24:RA:600:G:C8	2.56	0.40
35:RQ:141:GLN:OXT	44:RZ:99:TYR:OH	2.32	0.40
1:XA:225:C:H2'	1:XA:226:G:H8	1.87	0.40
1:XA:429:U:H1'	1:XA:430:A:H5''	2.04	0.40
1:XA:601:C:H2'	1:XA:602:A:H8	1.85	0.40
24:YA:1285:G:H2'	24:YA:1286:U:O4'	2.21	0.40
24:YA:1384:G:N7	42:YX:62:LYS:NZ	2.56	0.40
24:YA:1507:A:HO2'	24:YA:1508:G:P	2.45	0.40
24:YA:1800:G:O2'	24:YA:1980:C:OP1	2.31	0.40
24:YA:229:G:OP2	24:YA:230:A:O2'	2.25	0.40
39:YU:8:VAL:HG23	39:YU:11:ARG:HH21	1.85	0.40
24:YA:798:A:H5'	41:YW:90:ARG:HG2	2.04	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)
31:RI:91:SER:OG	1:XA:368:U:OP1[4_555]	2.00	0.20
24:RA:331:G:OP2	47:Y2:17:SER:CA[3_555]	2.01	0.19
31:RI:89:TYR:O	1:XA:357:G:O2'[4_555]	2.10	0.10
24:RA:331:G:OP2	47:Y2:17:SER:CB[3_555]	2.10	0.10
24:RA:2158:C:O2'	1:XA:1042:G:O2'[4_555]	2.13	0.07

5.3 Torsion angles

5.3.1 Protein backbone

In the following table, the Percentiles column shows the percent Ramachandran outliers of the chain as a percentile score with respect to all 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	QB	233/256 (91%)	205 (88%)	28 (12%)	0	100	100
2	XB	234/256 (91%)	207 (88%)	27 (12%)	0	100	100
3	QC	203/239 (85%)	186 (92%)	17 (8%)	0	100	100
3	XC	203/239 (85%)	182 (90%)	21 (10%)	0	100	100
4	QD	206/209 (99%)	198 (96%)	8 (4%)	0	100	100
4	XD	206/209 (99%)	191 (93%)	12 (6%)	3 (2%)	10	44
5	QE	149/162 (92%)	140 (94%)	9 (6%)	0	100	100
5	XE	149/162 (92%)	140 (94%)	9 (6%)	0	100	100
6	QF	99/101 (98%)	97 (98%)	2 (2%)	0	100	100
6	XF	99/101 (98%)	97 (98%)	2 (2%)	0	100	100
7	QG	153/156 (98%)	150 (98%)	3 (2%)	0	100	100
7	XG	153/156 (98%)	143 (94%)	10 (6%)	0	100	100
8	QH	135/138 (98%)	131 (97%)	4 (3%)	0	100	100
8	XH	135/138 (98%)	129 (96%)	6 (4%)	0	100	100
9	QI	125/128 (98%)	115 (92%)	10 (8%)	0	100	100
9	XI	124/128 (97%)	107 (86%)	17 (14%)	0	100	100
10	QJ	97/105 (92%)	88 (91%)	9 (9%)	0	100	100
10	XJ	94/105 (90%)	88 (94%)	6 (6%)	0	100	100
11	QK	117/129 (91%)	101 (86%)	16 (14%)	0	100	100
11	XK	114/129 (88%)	102 (90%)	12 (10%)	0	100	100
12	QL	123/132 (93%)	110 (89%)	13 (11%)	0	100	100
12	XL	120/132 (91%)	100 (83%)	20 (17%)	0	100	100
13	QM	118/126 (94%)	102 (86%)	16 (14%)	0	100	100
13	XM	117/126 (93%)	99 (85%)	17 (14%)	1 (1%)	17	56
14	QN	58/61 (95%)	51 (88%)	7 (12%)	0	100	100

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
14	XN	58/61 (95%)	50 (86%)	8 (14%)	0	100	100
15	QO	86/89 (97%)	83 (96%)	3 (4%)	0	100	100
15	XO	85/89 (96%)	84 (99%)	1 (1%)	0	100	100
16	QP	82/88 (93%)	79 (96%)	3 (4%)	0	100	100
16	XP	82/88 (93%)	74 (90%)	8 (10%)	0	100	100
17	QQ	98/105 (93%)	91 (93%)	7 (7%)	0	100	100
17	XQ	98/105 (93%)	93 (95%)	5 (5%)	0	100	100
18	QR	68/88 (77%)	63 (93%)	5 (7%)	0	100	100
18	XR	68/88 (77%)	65 (96%)	3 (4%)	0	100	100
19	QS	81/93 (87%)	71 (88%)	10 (12%)	0	100	100
19	XS	82/93 (88%)	71 (87%)	11 (13%)	0	100	100
20	QT	97/106 (92%)	90 (93%)	7 (7%)	0	100	100
20	XT	97/106 (92%)	91 (94%)	6 (6%)	0	100	100
21	QU	23/27 (85%)	22 (96%)	1 (4%)	0	100	100
21	XU	23/27 (85%)	23 (100%)	0	0	100	100
26	RD	272/276 (99%)	259 (95%)	13 (5%)	0	100	100
26	YD	272/276 (99%)	253 (93%)	19 (7%)	0	100	100
27	RE	203/206 (98%)	187 (92%)	16 (8%)	0	100	100
27	YE	202/206 (98%)	189 (94%)	13 (6%)	0	100	100
28	RF	200/210 (95%)	186 (93%)	14 (7%)	0	100	100
28	YF	200/210 (95%)	185 (92%)	15 (8%)	0	100	100
29	RG	179/182 (98%)	155 (87%)	24 (13%)	0	100	100
29	YG	179/182 (98%)	151 (84%)	27 (15%)	1 (1%)	25	64
30	RH	172/180 (96%)	142 (83%)	29 (17%)	1 (1%)	25	64
30	YH	171/180 (95%)	163 (95%)	8 (5%)	0	100	100
31	RI	144/148 (97%)	124 (86%)	20 (14%)	0	100	100
31	YI	144/148 (97%)	120 (83%)	24 (17%)	0	100	100
32	RN	136/140 (97%)	119 (88%)	17 (12%)	0	100	100
32	YN	138/140 (99%)	126 (91%)	12 (9%)	0	100	100
33	RO	120/122 (98%)	114 (95%)	6 (5%)	0	100	100
33	YO	120/122 (98%)	114 (95%)	6 (5%)	0	100	100

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
34	RP	147/150 (98%)	142 (97%)	5 (3%)	0	100	100
34	YP	145/150 (97%)	138 (95%)	6 (4%)	1 (1%)	22	61
35	RQ	139/141 (99%)	116 (84%)	22 (16%)	1 (1%)	22	61
35	YQ	139/141 (99%)	136 (98%)	3 (2%)	0	100	100
36	RR	115/118 (98%)	106 (92%)	8 (7%)	1 (1%)	17	56
36	YR	115/118 (98%)	108 (94%)	7 (6%)	0	100	100
37	RS	109/112 (97%)	89 (82%)	20 (18%)	0	100	100
37	YS	108/112 (96%)	105 (97%)	3 (3%)	0	100	100
38	RT	135/146 (92%)	119 (88%)	14 (10%)	2 (2%)	10	44
38	YT	135/146 (92%)	120 (89%)	13 (10%)	2 (2%)	10	44
39	RU	115/118 (98%)	103 (90%)	12 (10%)	0	100	100
39	YU	115/118 (98%)	107 (93%)	7 (6%)	1 (1%)	17	56
40	RV	99/101 (98%)	94 (95%)	5 (5%)	0	100	100
40	YV	99/101 (98%)	94 (95%)	5 (5%)	0	100	100
41	RW	111/113 (98%)	106 (96%)	5 (4%)	0	100	100
41	YW	111/113 (98%)	104 (94%)	7 (6%)	0	100	100
42	RX	90/96 (94%)	86 (96%)	4 (4%)	0	100	100
42	YX	92/96 (96%)	90 (98%)	2 (2%)	0	100	100
43	RY	105/110 (96%)	100 (95%)	5 (5%)	0	100	100
43	YY	105/110 (96%)	97 (92%)	8 (8%)	0	100	100
44	RZ	181/206 (88%)	143 (79%)	38 (21%)	0	100	100
44	YZ	181/206 (88%)	176 (97%)	4 (2%)	1 (1%)	25	64
45	R0	79/85 (93%)	76 (96%)	3 (4%)	0	100	100
45	Y0	73/85 (86%)	70 (96%)	3 (4%)	0	100	100
46	R1	95/98 (97%)	84 (88%)	11 (12%)	0	100	100
46	Y1	91/98 (93%)	84 (92%)	7 (8%)	0	100	100
47	R2	67/72 (93%)	65 (97%)	2 (3%)	0	100	100
47	Y2	64/72 (89%)	61 (95%)	3 (5%)	0	100	100
48	R3	57/60 (95%)	54 (95%)	3 (5%)	0	100	100
48	Y3	57/60 (95%)	55 (96%)	2 (4%)	0	100	100
49	R4	67/71 (94%)	54 (81%)	13 (19%)	0	100	100

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
49	Y4	67/71 (94%)	55 (82%)	12 (18%)	0	100	100
50	R5	57/60 (95%)	54 (95%)	3 (5%)	0	100	100
50	Y5	57/60 (95%)	54 (95%)	3 (5%)	0	100	100
51	R6	51/54 (94%)	51 (100%)	0	0	100	100
51	Y6	51/54 (94%)	50 (98%)	1 (2%)	0	100	100
52	R7	45/49 (92%)	44 (98%)	1 (2%)	0	100	100
52	Y7	46/49 (94%)	45 (98%)	1 (2%)	0	100	100
53	R8	62/65 (95%)	49 (79%)	13 (21%)	0	100	100
53	Y8	62/65 (95%)	53 (86%)	8 (13%)	1 (2%)	9	43
54	R9	35/37 (95%)	35 (100%)	0	0	100	100
54	Y9	35/37 (95%)	34 (97%)	1 (3%)	0	100	100
All	All	11453/12128 (94%)	10502 (92%)	935 (8%)	16 (0%)	51	83

All (16) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
36	RR	4	LEU
4	XD	31	CYS
44	YZ	52	SER
38	RT	123	GLN
38	RT	124	ASP
4	XD	32	ALA
38	YT	123	GLN
38	YT	124	ASP
29	YG	81	LYS
34	YP	36	LYS
13	XM	14	ARG
39	YU	93	LYS
35	RQ	78	PRO
4	XD	33	MET
30	RH	86	GLU
53	Y8	63	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	QB	203/220 (92%)	202 (100%)	1 (0%)	88	95
2	XB	204/220 (93%)	203 (100%)	1 (0%)	88	95
3	QC	159/188 (85%)	159 (100%)	0	100	100
3	XC	159/188 (85%)	159 (100%)	0	100	100
4	QD	180/181 (99%)	179 (99%)	1 (1%)	86	94
4	XD	180/181 (99%)	172 (96%)	8 (4%)	28	64
5	QE	116/123 (94%)	114 (98%)	2 (2%)	60	83
5	XE	116/123 (94%)	113 (97%)	3 (3%)	46	76
6	QF	90/90 (100%)	90 (100%)	0	100	100
6	XF	90/90 (100%)	89 (99%)	1 (1%)	73	88
7	QG	126/127 (99%)	123 (98%)	3 (2%)	49	77
7	XG	126/127 (99%)	124 (98%)	2 (2%)	62	84
8	QH	118/119 (99%)	118 (100%)	0	100	100
8	XH	118/119 (99%)	118 (100%)	0	100	100
9	QI	98/99 (99%)	98 (100%)	0	100	100
9	XI	97/99 (98%)	97 (100%)	0	100	100
10	QJ	89/92 (97%)	88 (99%)	1 (1%)	73	88
10	XJ	86/92 (94%)	86 (100%)	0	100	100
11	QK	90/99 (91%)	89 (99%)	1 (1%)	73	88
11	XK	88/99 (89%)	88 (100%)	0	100	100
12	QL	104/109 (95%)	104 (100%)	0	100	100
12	XL	103/109 (94%)	103 (100%)	0	100	100
13	QM	96/101 (95%)	94 (98%)	2 (2%)	53	79
13	XM	95/101 (94%)	95 (100%)	0	100	100
14	QN	49/50 (98%)	47 (96%)	2 (4%)	30	66
14	XN	49/50 (98%)	48 (98%)	1 (2%)	55	80
15	QO	79/80 (99%)	78 (99%)	1 (1%)	69	87
15	XO	79/80 (99%)	79 (100%)	0	100	100
16	QP	72/74 (97%)	71 (99%)	1 (1%)	67	86
16	XP	72/74 (97%)	72 (100%)	0	100	100

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
17	QQ	95/97 (98%)	94 (99%)	1 (1%)	73	88
17	XQ	95/97 (98%)	95 (100%)	0	100	100
18	QR	61/77 (79%)	61 (100%)	0	100	100
18	XR	61/77 (79%)	61 (100%)	0	100	100
19	QS	72/80 (90%)	72 (100%)	0	100	100
19	XS	73/80 (91%)	73 (100%)	0	100	100
20	QT	76/82 (93%)	76 (100%)	0	100	100
20	XT	76/82 (93%)	76 (100%)	0	100	100
21	QU	20/22 (91%)	20 (100%)	0	100	100
21	XU	20/22 (91%)	19 (95%)	1 (5%)	24	60
26	RD	216/218 (99%)	216 (100%)	0	100	100
26	YD	216/218 (99%)	214 (99%)	2 (1%)	78	91
27	RE	165/166 (99%)	164 (99%)	1 (1%)	86	94
27	YE	165/166 (99%)	164 (99%)	1 (1%)	86	94
28	RF	161/166 (97%)	161 (100%)	0	100	100
28	YF	161/166 (97%)	160 (99%)	1 (1%)	86	94
29	RG	155/156 (99%)	153 (99%)	2 (1%)	69	87
29	YG	155/156 (99%)	155 (100%)	0	100	100
30	RH	145/148 (98%)	145 (100%)	0	100	100
30	YH	144/148 (97%)	142 (99%)	2 (1%)	67	86
31	RI	122/124 (98%)	121 (99%)	1 (1%)	81	93
31	YI	122/124 (98%)	121 (99%)	1 (1%)	81	93
32	RN	117/119 (98%)	116 (99%)	1 (1%)	78	91
32	YN	119/119 (100%)	119 (100%)	0	100	100
33	RO	100/100 (100%)	98 (98%)	2 (2%)	55	80
33	YO	100/100 (100%)	100 (100%)	0	100	100
34	RP	116/116 (100%)	115 (99%)	1 (1%)	78	91
34	YP	114/116 (98%)	113 (99%)	1 (1%)	78	91
35	RQ	111/111 (100%)	108 (97%)	3 (3%)	44	75
35	YQ	111/111 (100%)	111 (100%)	0	100	100
36	RR	100/101 (99%)	99 (99%)	1 (1%)	76	90

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
36	YR	100/101 (99%)	100 (100%)	0	100	100
37	RS	87/88 (99%)	87 (100%)	0	100	100
37	YS	87/88 (99%)	87 (100%)	0	100	100
38	RT	120/127 (94%)	120 (100%)	0	100	100
38	YT	120/127 (94%)	119 (99%)	1 (1%)	81	93
39	RU	93/94 (99%)	92 (99%)	1 (1%)	73	88
39	YU	93/94 (99%)	91 (98%)	2 (2%)	52	79
40	RV	82/82 (100%)	82 (100%)	0	100	100
40	YV	82/82 (100%)	79 (96%)	3 (4%)	34	68
41	RW	92/92 (100%)	91 (99%)	1 (1%)	73	88
41	YW	92/92 (100%)	92 (100%)	0	100	100
42	RX	74/78 (95%)	74 (100%)	0	100	100
42	YX	76/78 (97%)	76 (100%)	0	100	100
43	RY	88/91 (97%)	87 (99%)	1 (1%)	73	88
43	YY	88/91 (97%)	87 (99%)	1 (1%)	73	88
44	RZ	162/179 (90%)	162 (100%)	0	100	100
44	YZ	162/179 (90%)	162 (100%)	0	100	100
45	R0	65/67 (97%)	63 (97%)	2 (3%)	40	72
45	Y0	61/67 (91%)	60 (98%)	1 (2%)	62	84
46	R1	82/83 (99%)	82 (100%)	0	100	100
46	Y1	78/83 (94%)	78 (100%)	0	100	100
47	R2	64/67 (96%)	64 (100%)	0	100	100
47	Y2	62/67 (92%)	62 (100%)	0	100	100
48	R3	51/52 (98%)	50 (98%)	1 (2%)	55	80
48	Y3	51/52 (98%)	50 (98%)	1 (2%)	55	80
49	R4	62/63 (98%)	62 (100%)	0	100	100
49	Y4	62/63 (98%)	61 (98%)	1 (2%)	62	84
50	R5	51/52 (98%)	51 (100%)	0	100	100
50	Y5	51/52 (98%)	51 (100%)	0	100	100
51	R6	51/52 (98%)	51 (100%)	0	100	100
51	Y6	51/52 (98%)	50 (98%)	1 (2%)	55	80

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
52	R7	40/42 (95%)	40 (100%)	0	100	100
52	Y7	41/42 (98%)	41 (100%)	0	100	100
53	R8	54/55 (98%)	53 (98%)	1 (2%)	57	81
53	Y8	54/55 (98%)	53 (98%)	1 (2%)	57	81
54	R9	34/34 (100%)	34 (100%)	0	100	100
54	Y9	34/34 (100%)	34 (100%)	0	100	100
All	All	9692/10066 (96%)	9620 (99%)	72 (1%)	84	94

All (72) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
2	QB	226	ARG
4	QD	64	LEU
5	QE	24	ARG
5	QE	51	VAL
7	QG	36	LYS
7	QG	94	ARG
7	QG	155	ARG
10	QJ	62	HIS
11	QK	12	ARG
13	QM	15	VAL
13	QM	80	ARG
14	QN	27	CYS
14	QN	33	VAL
15	QO	88	ARG
16	QP	21	VAL
17	QQ	101	ARG
27	RE	57	LYS
29	RG	33	ARG
29	RG	155	MET
31	RI	93	THR
32	RN	96	GLU
33	RO	24	VAL
33	RO	64	ARG
34	RP	50	ARG
35	RQ	21	THR
35	RQ	58	PHE
35	RQ	87	LYS
36	RR	2	ARG
39	RU	91	ASP

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
41	RW	92	ARG
43	RY	21	LYS
45	R0	14	ARG
45	R0	64	ASP
48	R3	30	ARG
53	R8	31	HIS
2	XB	178	ARG
4	XD	18	LYS
4	XD	20	TYR
4	XD	22	LYS
4	XD	31	CYS
4	XD	33	MET
4	XD	67	ILE
4	XD	106	TYR
4	XD	196	LEU
5	XE	24	ARG
5	XE	51	VAL
5	XE	63	ARG
6	XF	80	ARG
7	XG	50	ILE
7	XG	94	ARG
14	XN	27	CYS
21	XU	8	THR
26	YD	183	ARG
26	YD	264	LYS
27	YE	75	VAL
28	YF	8	GLN
30	YH	47	GLU
30	YH	49	VAL
31	YI	118	LYS
34	YP	15	ARG
38	YT	51	ARG
39	YU	91	ASP
39	YU	117	GLN
40	YV	72	VAL
40	YV	79	VAL
40	YV	80	GLN
43	YY	54	LYS
45	Y0	14	ARG
48	Y3	30	ARG
49	Y4	56	VAL
51	Y6	9	LEU

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
53	Y8	42	ARG

Some sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (17) such sidechains are listed below:

Mol	Chain	Res	Type
4	QD	119	GLN
4	QD	123	HIS
18	QR	36	ASN
19	QS	47	HIS
26	RD	253	GLN
30	RH	74	ASN
33	RO	3	GLN
38	RT	58	ASN
45	R0	35	ASN
53	R8	31	HIS
7	XG	51	GLN
10	XJ	33	GLN
26	YD	201	HIS
27	YE	129	HIS
28	YF	40	GLN
29	YG	132	ASN
43	YY	92	ASN

5.3.3 RNA [i](#)

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	QA	1499/1521 (98%)	289 (19%)	39 (2%)
1	XA	1499/1521 (98%)	271 (18%)	33 (2%)
22	QV	76/77 (98%)	22 (28%)	1 (1%)
22	XV	76/77 (98%)	18 (23%)	1 (1%)
23	QX	18/19 (94%)	7 (38%)	1 (5%)
23	XX	18/19 (94%)	6 (33%)	0
24	RA	2877/2915 (98%)	591 (20%)	37 (1%)
24	YA	2880/2915 (98%)	551 (19%)	43 (1%)
25	RB	119/122 (97%)	18 (15%)	1 (0%)
25	YB	119/122 (97%)	21 (17%)	1 (0%)
All	All	9181/9308 (98%)	1794 (19%)	157 (1%)

All (1794) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	QA	6	G
1	QA	9	G
1	QA	32	A
1	QA	39	G
1	QA	47	C
1	QA	48	C
1	QA	50	A
1	QA	51	A
1	QA	64	G
1	QA	65	U
1	QA	66	G
1	QA	76	G
1	QA	78	G
1	QA	82	U
1	QA	90	C
1	QA	91	C
1	QA	95	G
1	QA	101	A
1	QA	108	G
1	QA	116	A
1	QA	120	A
1	QA	121	C
1	QA	129(B)	G
1	QA	144	G
1	QA	146	G
1	QA	158	G
1	QA	161	A
1	QA	162	A
1	QA	163	C
1	QA	169	C
1	QA	173	U
1	QA	174	C
1	QA	182	U
1	QA	187	C
1	QA	189	U
1	QA	190	G
1	QA	191(A)	G
1	QA	195	A
1	QA	197	A
1	QA	208	U
1	QA	209	U
1	QA	210	U
1	QA	216	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	QA	220	G
1	QA	244	U
1	QA	245	C
1	QA	247	G
1	QA	250	A
1	QA	251	G
1	QA	262	A
1	QA	267	C
1	QA	270	A
1	QA	281	G
1	QA	289	G
1	QA	315	A
1	QA	316	G
1	QA	321	A
1	QA	328	C
1	QA	329	A
1	QA	330	C
1	QA	332	G
1	QA	343	U
1	QA	344	A
1	QA	346	G
1	QA	347	G
1	QA	352	C
1	QA	353	A
1	QA	354	G
1	QA	367	U
1	QA	372	C
1	QA	373	A
1	QA	389	A
1	QA	390	C
1	QA	397	A
1	QA	398	C
1	QA	406	G
1	QA	411	A
1	QA	412	A
1	QA	413	G
1	QA	414	A
1	QA	421	U
1	QA	422	C
1	QA	423	G
1	QA	424	G
1	QA	429	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	QA	430	A
1	QA	435	C
1	QA	439	A
1	QA	442	C
1	QA	465	A
1	QA	466	C
1	QA	482	A
1	QA	485	G
1	QA	486	U
1	QA	496	A
1	QA	497	U
1	QA	505	G
1	QA	508	C
1	QA	509	A
1	QA	510	A
1	QA	511	C
1	QA	518	C
1	QA	521	G
1	QA	527	G
1	QA	532	A
1	QA	533	A
1	QA	545	C
1	QA	547	A
1	QA	559	A
1	QA	564	C
1	QA	566	G
1	QA	568	G
1	QA	572	A
1	QA	573	A
1	QA	576	G
1	QA	577	G
1	QA	596	C
1	QA	630	G
1	QA	631	G
1	QA	633	G
1	QA	653	A
1	QA	657	G
1	QA	665	A
1	QA	686	U
1	QA	688	G
1	QA	693	G
1	QA	701	C

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	QA	702	A
1	QA	704	A
1	QA	722	A
1	QA	731	G
1	QA	748	C
1	QA	754	C
1	QA	755	G
1	QA	760	G
1	QA	785	G
1	QA	792	A
1	QA	793	U
1	QA	794	A
1	QA	813	U
1	QA	817	C
1	QA	819	A
1	QA	821	G
1	QA	828	A
1	QA	841	U
1	QA	842	C
1	QA	843	U
1	QA	848	C
1	QA	859	A
1	QA	871	U
1	QA	872	A
1	QA	873	A
1	QA	902	G
1	QA	914	A
1	QA	926	G
1	QA	927	G
1	QA	934	C
1	QA	935	A
1	QA	960	U
1	QA	961	U
1	QA	966	G
1	QA	968	A
1	QA	969	A
1	QA	971	G
1	QA	974	A
1	QA	975	A
1	QA	976	G
1	QA	977	A
1	QA	982	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	QA	983	A
1	QA	991	U
1	QA	992	U
1	QA	993	G
1	QA	994	A
1	QA	1004	A
1	QA	1005	A
1	QA	1006	C
1	QA	1008	C
1	QA	1009	G
1	QA	1020	U
1	QA	1024	G
1	QA	1025	U
1	QA	1027	C
1	QA	1028(A)	C
1	QA	1028(B)	C
1	QA	1028(C)	C
1	QA	1029	G
1	QA	1030	C
1	QA	1031	G
1	QA	1032(B)	G
1	QA	1033	G
1	QA	1034	G
1	QA	1035	A
1	QA	1036	G
1	QA	1040	U
1	QA	1042	G
1	QA	1054	C
1	QA	1064	G
1	QA	1065	U
1	QA	1066	C
1	QA	1067	A
1	QA	1080	A
1	QA	1081	G
1	QA	1094	G
1	QA	1101	A
1	QA	1108	G
1	QA	1124	G
1	QA	1125	U
1	QA	1126	U
1	QA	1129	C
1	QA	1130	A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	QA	1131	G
1	QA	1135	U
1	QA	1136	U
1	QA	1137	C
1	QA	1138	G
1	QA	1139	G
1	QA	1144	G
1	QA	1146	A
1	QA	1157	A
1	QA	1159	U
1	QA	1178	G
1	QA	1181	G
1	QA	1184	G
1	QA	1196	U
1	QA	1201	A
1	QA	1202	G
1	QA	1211	U
1	QA	1212	U
1	QA	1213	A
1	QA	1215	G
1	QA	1227	A
1	QA	1228	C
1	QA	1240	U
1	QA	1241	G
1	QA	1256	A
1	QA	1257	U
1	QA	1258	G
1	QA	1260	C
1	QA	1270	C
1	QA	1280	A
1	QA	1281	U
1	QA	1285	A
1	QA	1286	A
1	QA	1287	A
1	QA	1297	C
1	QA	1298	C
1	QA	1299	A
1	QA	1300	G
1	QA	1301	U
1	QA	1302	U
1	QA	1303	C
1	QA	1305	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	QA	1312	G
1	QA	1320	C
1	QA	1321	C
1	QA	1322	C
1	QA	1323	G
1	QA	1331	G
1	QA	1334	G
1	QA	1335	C
1	QA	1336	C
1	QA	1337	G
1	QA	1346	A
1	QA	1347	G
1	QA	1348	U
1	QA	1362(B)	C
1	QA	1370	G
1	QA	1397	C
1	QA	1398	A
1	QA	1419	G
1	QA	1442	G
1	QA	1446	A
1	QA	1447	G
1	QA	1452	C
1	QA	1453	G
1	QA	1454	G
1	QA	1487	G
1	QA	1492	A
1	QA	1494	G
1	QA	1499	A
1	QA	1502	A
1	QA	1504	G
1	QA	1506	U
1	QA	1517	G
1	QA	1520	G
1	QA	1528	U
1	QA	1529	G
1	QA	1530	G
22	QV	2	G
22	QV	4	U
22	QV	5	G
22	QV	8	U
22	QV	16	C
22	QV	17	C

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
22	QV	18	U
22	QV	19	G
22	QV	20	G
22	QV	21(A)	U
22	QV	21(B)	A
22	QV	38	A
22	QV	42	A
22	QV	46	G
22	QV	48	C
22	QV	50	G
22	QV	54	U
22	QV	59	A
22	QV	67	U
22	QV	69	A
22	QV	75	C
22	QV	76	A
23	QX	7	G
23	QX	10	G
23	QX	11	U
23	QX	12	A
23	QX	13	A
23	QX	18	G
23	QX	19	G
24	RA	9	U
24	RA	15	G
24	RA	34	C
24	RA	35	G
24	RA	45	C
24	RA	50	G
24	RA	54	G
24	RA	60	G
24	RA	71	U
24	RA	72	A
24	RA	73	A
24	RA	74	G
24	RA	82	G
24	RA	89	U
24	RA	99	G
24	RA	101	A
24	RA	116	A
24	RA	118	U
24	RA	129	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	139	A
24	RA	156	U
24	RA	166	G
24	RA	170	A
24	RA	185	A
24	RA	188	A
24	RA	193	A
24	RA	204	G
24	RA	205	A
24	RA	210	A
24	RA	211	A
24	RA	212	A
24	RA	218	A
24	RA	219	U
24	RA	222	A
24	RA	234	G
24	RA	237	G
24	RA	238	C
24	RA	241	G
24	RA	271	U
24	RA	272	U
24	RA	273	G
24	RA	275	C
24	RA	288	U
24	RA	299	G
24	RA	300	A
24	RA	301	C
24	RA	323	A
24	RA	331	G
24	RA	332	G
24	RA	333	G
24	RA	335	A
24	RA	347	G
24	RA	348	A
24	RA	353	G
24	RA	354	A
24	RA	370	A
24	RA	376	G
24	RA	393	A
24	RA	394	C
24	RA	398	A
24	RA	399	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	400	U
24	RA	413	G
24	RA	422	U
24	RA	432	U
24	RA	433	G
24	RA	434	G
24	RA	438	G
24	RA	439	A
24	RA	448	U
24	RA	455	A
24	RA	470	C
24	RA	474	U
24	RA	480	A
24	RA	481	C
24	RA	482	C
24	RA	483	A
24	RA	496	A
24	RA	507	G
24	RA	526	A
24	RA	529	U
24	RA	530	A
24	RA	534	C
24	RA	538	A
24	RA	552	C
24	RA	554	A
24	RA	556	C
24	RA	557	A
24	RA	558	G
24	RA	562	C
24	RA	563	G
24	RA	564	G
24	RA	569	G
24	RA	570	C
24	RA	571	A
24	RA	573	G
24	RA	586	G
24	RA	596	G
24	RA	597	C
24	RA	598	A
24	RA	611	U
24	RA	626	A
24	RA	630	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	638	U
24	RA	639	G
24	RA	641	G
24	RA	646	A
24	RA	651	U
24	RA	652	A
24	RA	656	A
24	RA	662	A
24	RA	663	G
24	RA	670	C
24	RA	671	A
24	RA	676	G
24	RA	677	C
24	RA	679	A
24	RA	680	G
24	RA	705	C
24	RA	717	A
24	RA	733	G
24	RA	749	G
24	RA	764	G
24	RA	769	A
24	RA	773	G
24	RA	777	C
24	RA	800	C
24	RA	811	A
24	RA	812	G
24	RA	823	G
24	RA	829	A
24	RA	830	A
24	RA	831	A
24	RA	832	G
24	RA	837	C
24	RA	838	C
24	RA	839	G
24	RA	852	G
24	RA	858	U
24	RA	859	C
24	RA	866	A
24	RA	874	U
24	RA	875	U
24	RA	878	G
24	RA	894	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	903	C
24	RA	904	C
24	RA	906	G
24	RA	916	G
24	RA	929	G
24	RA	931	C
24	RA	932	C
24	RA	933	C
24	RA	934	A
24	RA	935	C
24	RA	936	C
24	RA	939	C
24	RA	942	A
24	RA	943	C
24	RA	944	C
24	RA	946	A
24	RA	947	A
24	RA	953	U
24	RA	956	A
24	RA	963	A
24	RA	977	G
24	RA	983	G
24	RA	986	A
24	RA	990	A
24	RA	991	G
24	RA	1002	A
24	RA	1004	A
24	RA	1006	C
24	RA	1018	A
24	RA	1019	G
24	RA	1020	C
24	RA	1026	A
24	RA	1029	A
24	RA	1036	A
24	RA	1042	A
24	RA	1049	G
24	RA	1051	C
24	RA	1058	U
24	RA	1059	C
24	RA	1068	G
24	RA	1069	U
24	RA	1071	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	1072	U
24	RA	1073	A
24	RA	1079	U
24	RA	1086	C
24	RA	1088	G
24	RA	1090	G
24	RA	1091	A
24	RA	1092	A
24	RA	1093	G
24	RA	1099	C
24	RA	1100	A
24	RA	1103	A
24	RA	1104	G
24	RA	1105	G
24	RA	1106	U
24	RA	1109	G
24	RA	1110	C
24	RA	1111	U
24	RA	1112	U
24	RA	1113	A
24	RA	1114	G
24	RA	1116	A
24	RA	1117	G
24	RA	1119	A
24	RA	1120	G
24	RA	1121	C
24	RA	1122	C
24	RA	1124	U
24	RA	1125	C
24	RA	1126	C
24	RA	1128	U
24	RA	1129	U
24	RA	1130	A
24	RA	1131	A
24	RA	1132	A
24	RA	1134	A
24	RA	1136	U
24	RA	1137	G
24	RA	1138	C
24	RA	1140	U
24	RA	1142	A
24	RA	1143	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	1155	C
24	RA	1157	A
24	RA	1158	G
24	RA	1168	G
24	RA	1172	A
24	RA	1175	A
24	RA	1176	U
24	RA	1180	C
24	RA	1181	G
24	RA	1188	A
24	RA	1197	G
24	RA	1214	G
24	RA	1218	G
24	RA	1219	A
24	RA	1220	U
24	RA	1221	G
24	RA	1222	A
24	RA	1223	C
24	RA	1224	C
24	RA	1226	C
24	RA	1240	G
24	RA	1249	A
24	RA	1250	U
24	RA	1255	A
24	RA	1256	U
24	RA	1265	A
24	RA	1282	G
24	RA	1283	A
24	RA	1284	G
24	RA	1296	G
24	RA	1299	A
24	RA	1302	G
24	RA	1311	A
24	RA	1318	A
24	RA	1319	U
24	RA	1332	A
24	RA	1346	U
24	RA	1347	A
24	RA	1358	U
24	RA	1359	U
24	RA	1360	C
24	RA	1367	A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	1375	U
24	RA	1395	A
24	RA	1398	U
24	RA	1411	A
24	RA	1416	C
24	RA	1424	A
24	RA	1425	A
24	RA	1430	A
24	RA	1431	G
24	RA	1432	C
24	RA	1438	A
24	RA	1441	A
24	RA	1453	C
24	RA	1454	C
24	RA	1457	C
24	RA	1462	G
24	RA	1465	A
24	RA	1466	U
24	RA	1467	G
24	RA	1474	C
24	RA	1491	A
24	RA	1492	C
24	RA	1496	A
24	RA	1497	G
24	RA	1502	G
24	RA	1505	C
24	RA	1507	A
24	RA	1508	G
24	RA	1518	A
24	RA	1527	G
24	RA	1528	U
24	RA	1529	G
24	RA	1531	G
24	RA	1533	G
24	RA	1539	C
24	RA	1540	A
24	RA	1543	U
24	RA	1550	C
24	RA	1553	A
24	RA	1554	A
24	RA	1556	A
24	RA	1557	A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	1559	C
24	RA	1560	U
24	RA	1561	C
24	RA	1565	G
24	RA	1568	G
24	RA	1574	A
24	RA	1580	G
24	RA	1581	U
24	RA	1582	A
24	RA	1583	C
24	RA	1584	G
24	RA	1589	A
24	RA	1590	C
24	RA	1591	A
24	RA	1593	C
24	RA	1594	C
24	RA	1605	A
24	RA	1606	G
24	RA	1607	G
24	RA	1613	A
24	RA	1616	A
24	RA	1625	U
24	RA	1627	A
24	RA	1630	A
24	RA	1632	A
24	RA	1633	A
24	RA	1643	A
24	RA	1644	C
24	RA	1649	A
24	RA	1650	C
24	RA	1654	A
24	RA	1655	A
24	RA	1656	A
24	RA	1662	A
24	RA	1663	C
24	RA	1695	C
24	RA	1701	A
24	RA	1707	C
24	RA	1713	G
24	RA	1714	G
24	RA	1715	A
24	RA	1721	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	1763	G
24	RA	1766	G
24	RA	1767	A
24	RA	1768	U
24	RA	1769	G
24	RA	1771	G
24	RA	1775	C
24	RA	1776	G
24	RA	1781	G
24	RA	1787	G
24	RA	1793	A
24	RA	1794	G
24	RA	1795	G
24	RA	1804	A
24	RA	1807	G
24	RA	1811	A
24	RA	1812	C
24	RA	1813	C
24	RA	1822	A
24	RA	1830	G
24	RA	1831	C
24	RA	1832	G
24	RA	1847	G
24	RA	1851	U
24	RA	1866	G
24	RA	1878	A
24	RA	1879	A
24	RA	1889	G
24	RA	1896	G
24	RA	1897	C
24	RA	1899	A
24	RA	1900	G
24	RA	1904	C
24	RA	1910	G
24	RA	1911	A
24	RA	1921	G
24	RA	1925	G
24	RA	1928	G
24	RA	1935	A
24	RA	1949	A
24	RA	1951	G
24	RA	1952	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	1958	A
24	RA	1960	A
24	RA	1961	U
24	RA	1977	U
24	RA	1985	U
24	RA	1989	C
24	RA	1991	A
24	RA	1992	A
24	RA	1993	A
24	RA	1994	A
24	RA	2004	C
24	RA	2013	U
24	RA	2014	G
24	RA	2015	U
24	RA	2018	C
24	RA	2045	G
24	RA	2053	A
24	RA	2055	A
24	RA	2056	U
24	RA	2065	C
24	RA	2077	C
24	RA	2078	G
24	RA	2081	A
24	RA	2082	A
24	RA	2083	G
24	RA	2084	A
24	RA	2085	C
24	RA	2091	G
24	RA	2115	G
24	RA	2123	G
24	RA	2125	C
24	RA	2126	G
24	RA	2133	C
24	RA	2135	U
24	RA	2136	A
24	RA	2137	G
24	RA	2138	G
24	RA	2139	A
24	RA	2146	G
24	RA	2148	A
24	RA	2149	G
24	RA	2150	C

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	2153	G
24	RA	2154	U
24	RA	2155	G
24	RA	2157	A
24	RA	2158	C
24	RA	2159	C
24	RA	2160	C
24	RA	2167	C
24	RA	2168	C
24	RA	2169	G
24	RA	2170	G
24	RA	2179	G
24	RA	2180	A
24	RA	2183	C
24	RA	2188	G
24	RA	2191	A
24	RA	2192	A
24	RA	2195	A
24	RA	2196	C
24	RA	2199	C
24	RA	2200	C
24	RA	2211	U
24	RA	2212	G
24	RA	2213	G
24	RA	2220	A
24	RA	2227	G
24	RA	2228	G
24	RA	2229	A
24	RA	2230	U
24	RA	2233	G
24	RA	2234	G
24	RA	2237	A
24	RA	2250	G
24	RA	2251	G
24	RA	2255	U
24	RA	2278	A
24	RA	2287	C
24	RA	2290	A
24	RA	2292	G
24	RA	2295	C
24	RA	2299	A
24	RA	2300	A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	2315	G
24	RA	2317	A
24	RA	2319	G
24	RA	2320	G
24	RA	2321	A
24	RA	2323	A
24	RA	2324	U
24	RA	2331	G
24	RA	2332	A
24	RA	2337	G
24	RA	2346	G
24	RA	2348	A
24	RA	2354	C
24	RA	2357	G
24	RA	2358	A
24	RA	2359	C
24	RA	2362	C
24	RA	2366	G
24	RA	2395	G
24	RA	2397	C
24	RA	2415	C
24	RA	2418	U
24	RA	2437	A
24	RA	2441	G
24	RA	2442	A
24	RA	2447	A
24	RA	2451	A
24	RA	2452	C
24	RA	2453	C
24	RA	2460	A
24	RA	2481	A
24	RA	2482	G
24	RA	2486	C
24	RA	2487	C
24	RA	2492	C
24	RA	2506	G
24	RA	2514	G
24	RA	2517	G
24	RA	2530	A
24	RA	2531	U
24	RA	2537	G
24	RA	2541	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	2554	A
24	RA	2555	G
24	RA	2566	U
24	RA	2576	A
24	RA	2578	A
24	RA	2579	G
24	RA	2581	G
24	RA	2584	A
24	RA	2598	C
24	RA	2614	A
24	RA	2621	U
24	RA	2623	U
24	RA	2624	C
24	RA	2627	U
24	RA	2635	G
24	RA	2641	A
24	RA	2648	U
24	RA	2653	G
24	RA	2658	C
24	RA	2667	G
24	RA	2677	A
24	RA	2685	G
24	RA	2694	U
24	RA	2701	U
24	RA	2702	C
24	RA	2703	C
24	RA	2714	U
24	RA	2715	C
24	RA	2719	G
24	RA	2725	A
24	RA	2726	A
24	RA	2727	G
24	RA	2739	U
24	RA	2746	A
24	RA	2757	G
24	RA	2771	A
24	RA	2774	G
24	RA	2775	G
24	RA	2778	A
24	RA	2791	A
24	RA	2792	U
24	RA	2793	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	2800	C
24	RA	2802	C
24	RA	2803	A
24	RA	2804	C
24	RA	2805	G
24	RA	2809	U
24	RA	2810	C
24	RA	2811	A
24	RA	2818	U
24	RA	2828	G
24	RA	2830	A
24	RA	2831	A
24	RA	2843	G
24	RA	2844	G
24	RA	2857	U
24	RA	2859	U
24	RA	2877	G
24	RA	2878	A
24	RA	2882	G
24	RA	2890	C
24	RA	2900	G
24	RA	2901	A
24	RA	2903	G
24	RA	2904	U
24	RA	2905	C
24	RA	2906	U
25	RB	8	U
25	RB	13	A
25	RB	15	A
25	RB	16	G
25	RB	19	G
25	RB	21	G
25	RB	25	A
25	RB	41	U
25	RB	42	C
25	RB	44	G
25	RB	45	A
25	RB	52	A
25	RB	56	G
25	RB	67	G
25	RB	73	A
25	RB	81	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
25	RB	82	G
25	RB	109	G
1	XA	6	G
1	XA	9	G
1	XA	32	A
1	XA	39	G
1	XA	48	C
1	XA	51	A
1	XA	58	C
1	XA	61	G
1	XA	65	U
1	XA	66	G
1	XA	79	G
1	XA	89	U
1	XA	92	G
1	XA	95	G
1	XA	101	A
1	XA	108	G
1	XA	116	A
1	XA	121	C
1	XA	122	G
1	XA	130	A
1	XA	144	G
1	XA	147	G
1	XA	160	A
1	XA	163	C
1	XA	169	C
1	XA	172	A
1	XA	173	U
1	XA	174	C
1	XA	190	G
1	XA	191(A)	G
1	XA	195	A
1	XA	197	A
1	XA	208	U
1	XA	209	U
1	XA	210	U
1	XA	220	G
1	XA	244	U
1	XA	245	C
1	XA	247	G
1	XA	251	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	XA	267	C
1	XA	270	A
1	XA	281	G
1	XA	289	G
1	XA	306	G
1	XA	321	A
1	XA	328	C
1	XA	329	A
1	XA	332	G
1	XA	344	A
1	XA	346	G
1	XA	347	G
1	XA	351	G
1	XA	352	C
1	XA	353	A
1	XA	354	G
1	XA	356	A
1	XA	359	U
1	XA	367	U
1	XA	372	C
1	XA	373	A
1	XA	389	A
1	XA	397	A
1	XA	398	C
1	XA	406	G
1	XA	410	G
1	XA	411	A
1	XA	412	A
1	XA	413	G
1	XA	414	A
1	XA	422	C
1	XA	423	G
1	XA	424	G
1	XA	429	U
1	XA	430	A
1	XA	440	A
1	XA	466	C
1	XA	467	G
1	XA	485	G
1	XA	486	U
1	XA	496	A
1	XA	497	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	XA	509	A
1	XA	510	A
1	XA	511	C
1	XA	518	C
1	XA	521	G
1	XA	527	G
1	XA	531	U
1	XA	532	A
1	XA	533	A
1	XA	545	C
1	XA	547	A
1	XA	548	G
1	XA	559	A
1	XA	564	C
1	XA	572	A
1	XA	573	A
1	XA	576	G
1	XA	577	G
1	XA	596	C
1	XA	629	G
1	XA	630	G
1	XA	631	G
1	XA	632	A
1	XA	653	A
1	XA	665	A
1	XA	686	U
1	XA	688	G
1	XA	693	G
1	XA	703	G
1	XA	704	A
1	XA	721	G
1	XA	731	G
1	XA	734	G
1	XA	749	C
1	XA	753	A
1	XA	754	C
1	XA	755	G
1	XA	777	A
1	XA	792	A
1	XA	793	U
1	XA	794	A
1	XA	810	C

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	XA	813	U
1	XA	816	A
1	XA	817	C
1	XA	821	G
1	XA	828	A
1	XA	843	U
1	XA	848	C
1	XA	859	A
1	XA	871	U
1	XA	872	A
1	XA	876	G
1	XA	902	G
1	XA	914	A
1	XA	926	G
1	XA	927	G
1	XA	934	C
1	XA	935	A
1	XA	958	A
1	XA	960	U
1	XA	961	U
1	XA	968	A
1	XA	969	A
1	XA	974	A
1	XA	975	A
1	XA	976	G
1	XA	977	A
1	XA	991	U
1	XA	992	U
1	XA	993	G
1	XA	1003	G
1	XA	1004	A
1	XA	1006	C
1	XA	1009	G
1	XA	1020	U
1	XA	1024	G
1	XA	1025	U
1	XA	1028(A)	C
1	XA	1028(B)	C
1	XA	1029	G
1	XA	1030	C
1	XA	1031	G
1	XA	1032(A)	A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	XA	1032(B)	G
1	XA	1038	C
1	XA	1039	C
1	XA	1040	U
1	XA	1041	A
1	XA	1042	G
1	XA	1044	A
1	XA	1046	A
1	XA	1053	G
1	XA	1054	C
1	XA	1064	G
1	XA	1066	C
1	XA	1081	G
1	XA	1094	G
1	XA	1095	U
1	XA	1101	A
1	XA	1108	G
1	XA	1124	G
1	XA	1125	U
1	XA	1126	U
1	XA	1127	G
1	XA	1129	C
1	XA	1130	A
1	XA	1134	G
1	XA	1136	U
1	XA	1137	C
1	XA	1138	G
1	XA	1139	G
1	XA	1140	C
1	XA	1146	A
1	XA	1152	A
1	XA	1157	A
1	XA	1158	C
1	XA	1159	U
1	XA	1160	G
1	XA	1162	C
1	XA	1176	A
1	XA	1177	G
1	XA	1178	G
1	XA	1181	G
1	XA	1182	G
1	XA	1183	A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	XA	1184	G
1	XA	1196	U
1	XA	1212	U
1	XA	1213	A
1	XA	1225	A
1	XA	1238	A
1	XA	1240	U
1	XA	1241	G
1	XA	1256	A
1	XA	1257	U
1	XA	1258	G
1	XA	1260	C
1	XA	1263	C
1	XA	1270	C
1	XA	1273	G
1	XA	1275	A
1	XA	1280	A
1	XA	1281	U
1	XA	1286	A
1	XA	1287	A
1	XA	1290	G
1	XA	1298	C
1	XA	1300	G
1	XA	1301	U
1	XA	1302	U
1	XA	1303	C
1	XA	1305	G
1	XA	1310	G
1	XA	1311	G
1	XA	1320	C
1	XA	1321	C
1	XA	1322	C
1	XA	1323	G
1	XA	1331	G
1	XA	1335	C
1	XA	1336	C
1	XA	1347	G
1	XA	1362(B)	C
1	XA	1363	A
1	XA	1370	G
1	XA	1378	C
1	XA	1379	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	XA	1394	A
1	XA	1419	G
1	XA	1442	G
1	XA	1446	A
1	XA	1447	G
1	XA	1452	C
1	XA	1453	G
1	XA	1454	G
1	XA	1487	G
1	XA	1492	A
1	XA	1493	A
1	XA	1497	G
1	XA	1499	A
1	XA	1502	A
1	XA	1504	G
1	XA	1506	U
1	XA	1517	G
1	XA	1519	A
1	XA	1520	G
1	XA	1529	G
1	XA	1530	G
22	XV	3	G
22	XV	4	U
22	XV	5	G
22	XV	7	U
22	XV	8	U
22	XV	18	U
22	XV	19	G
22	XV	20	G
22	XV	21(B)	A
22	XV	47	U
22	XV	48	C
22	XV	49	G
22	XV	52	G
22	XV	54	U
22	XV	64	C
22	XV	68	C
22	XV	72	G
22	XV	76	A
23	XX	7	G
23	XX	10	G
23	XX	11	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
23	XX	12	A
23	XX	13	A
23	XX	19	G
24	YA	11	G
24	YA	15	G
24	YA	34	C
24	YA	45	C
24	YA	60	G
24	YA	62	U
24	YA	71	U
24	YA	73	A
24	YA	74	G
24	YA	90	A
24	YA	99	G
24	YA	100	G
24	YA	101	A
24	YA	116	A
24	YA	118	U
24	YA	123	G
24	YA	129	G
24	YA	139	A
24	YA	156	U
24	YA	157	U
24	YA	170	A
24	YA	185	A
24	YA	188	A
24	YA	205	A
24	YA	210	A
24	YA	211	A
24	YA	212	A
24	YA	217	A
24	YA	218	A
24	YA	219	U
24	YA	221	G
24	YA	231	G
24	YA	232	U
24	YA	237	G
24	YA	241	G
24	YA	258	U
24	YA	271	U
24	YA	272	U
24	YA	273	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	YA	275	C
24	YA	287	G
24	YA	288	U
24	YA	289	G
24	YA	298	G
24	YA	300	A
24	YA	302	A
24	YA	303	C
24	YA	323	A
24	YA	335	A
24	YA	347	G
24	YA	348	A
24	YA	353	G
24	YA	354	A
24	YA	356	A
24	YA	376	G
24	YA	387	G
24	YA	392	U
24	YA	394	C
24	YA	398	A
24	YA	399	G
24	YA	400	U
24	YA	413	G
24	YA	414	U
24	YA	422	U
24	YA	432	U
24	YA	433	G
24	YA	438	G
24	YA	439	A
24	YA	455	A
24	YA	469	A
24	YA	470	C
24	YA	474	U
24	YA	480	A
24	YA	483	A
24	YA	496	A
24	YA	507	G
24	YA	529	U
24	YA	530	A
24	YA	533	G
24	YA	534	C
24	YA	535	C

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	YA	537	G
24	YA	543	G
24	YA	554	A
24	YA	555	G
24	YA	556	C
24	YA	557	A
24	YA	562	C
24	YA	563	G
24	YA	564	G
24	YA	570	C
24	YA	571	A
24	YA	586	G
24	YA	594	A
24	YA	596	G
24	YA	598	A
24	YA	626	A
24	YA	630	U
24	YA	638	U
24	YA	641	G
24	YA	646	A
24	YA	647	G
24	YA	652	A
24	YA	656	A
24	YA	662	A
24	YA	663	G
24	YA	670	C
24	YA	671	A
24	YA	676	G
24	YA	679	A
24	YA	680	G
24	YA	681	C
24	YA	716	G
24	YA	717	A
24	YA	733	G
24	YA	764	G
24	YA	769	A
24	YA	773	G
24	YA	777	C
24	YA	800	C
24	YA	822	G
24	YA	823	G
24	YA	829	A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	YA	830	A
24	YA	831	A
24	YA	832	G
24	YA	837	C
24	YA	838	C
24	YA	839	G
24	YA	852	G
24	YA	858	U
24	YA	859	C
24	YA	866	A
24	YA	874	U
24	YA	875	U
24	YA	878	G
24	YA	894	U
24	YA	903	C
24	YA	904	C
24	YA	907	U
24	YA	913	A
24	YA	927	G
24	YA	928	G
24	YA	929	G
24	YA	931	C
24	YA	932	C
24	YA	933	C
24	YA	935	C
24	YA	936	C
24	YA	937	A
24	YA	942	A
24	YA	943	C
24	YA	944	C
24	YA	946	A
24	YA	947	A
24	YA	953	U
24	YA	956	A
24	YA	961	C
24	YA	963	A
24	YA	977	G
24	YA	983	G
24	YA	986	A
24	YA	990	A
24	YA	991	G
24	YA	998	A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	YA	1002	A
24	YA	1004	A
24	YA	1006	C
24	YA	1018	A
24	YA	1019	G
24	YA	1020	C
24	YA	1029	A
24	YA	1035	G
24	YA	1042	A
24	YA	1057	G
24	YA	1058	U
24	YA	1059	C
24	YA	1061	G
24	YA	1068	G
24	YA	1069	U
24	YA	1072	U
24	YA	1073	A
24	YA	1079	U
24	YA	1092	A
24	YA	1093	G
24	YA	1096	A
24	YA	1100	A
24	YA	1105	G
24	YA	1106	U
24	YA	1107	U
24	YA	1108	G
24	YA	1111	U
24	YA	1112	U
24	YA	1113	A
24	YA	1114	G
24	YA	1116	A
24	YA	1117	G
24	YA	1119	A
24	YA	1122	C
24	YA	1123	A
24	YA	1124	U
24	YA	1128	U
24	YA	1129	U
24	YA	1130	A
24	YA	1131	A
24	YA	1132	A
24	YA	1134	A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	YA	1135	G
24	YA	1139	G
24	YA	1141	A
24	YA	1142	A
24	YA	1143	U
24	YA	1149	A
24	YA	1156	G
24	YA	1168	G
24	YA	1175	A
24	YA	1177	G
24	YA	1180	C
24	YA	1181	G
24	YA	1184	G
24	YA	1187	U
24	YA	1188	A
24	YA	1219	A
24	YA	1220	U
24	YA	1221	G
24	YA	1222	A
24	YA	1224	C
24	YA	1249	A
24	YA	1250	U
24	YA	1256	U
24	YA	1265	A
24	YA	1282	G
24	YA	1284	G
24	YA	1299	A
24	YA	1302	G
24	YA	1311	A
24	YA	1317	G
24	YA	1318	A
24	YA	1319	U
24	YA	1321	A
24	YA	1346	U
24	YA	1347	A
24	YA	1375	U
24	YA	1395	A
24	YA	1398	U
24	YA	1411	A
24	YA	1414	G
24	YA	1416	C
24	YA	1424	A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	YA	1425	A
24	YA	1430	A
24	YA	1431	G
24	YA	1441	A
24	YA	1453	C
24	YA	1454	C
24	YA	1457	C
24	YA	1462	G
24	YA	1465	A
24	YA	1466	U
24	YA	1467	G
24	YA	1474	C
24	YA	1491	A
24	YA	1492	C
24	YA	1496	A
24	YA	1497	G
24	YA	1502	G
24	YA	1506	G
24	YA	1507	A
24	YA	1508	G
24	YA	1514	C
24	YA	1518	A
24	YA	1528	U
24	YA	1529	G
24	YA	1536	A
24	YA	1539	C
24	YA	1542	A
24	YA	1543	U
24	YA	1553	A
24	YA	1554	A
24	YA	1555	C
24	YA	1556	A
24	YA	1557	A
24	YA	1561	C
24	YA	1568	G
24	YA	1569	U
24	YA	1580	G
24	YA	1581	U
24	YA	1582	A
24	YA	1583	C
24	YA	1584	G
24	YA	1586	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	YA	1589	A
24	YA	1590	C
24	YA	1591	A
24	YA	1605	A
24	YA	1606	G
24	YA	1613	A
24	YA	1616	A
24	YA	1625	U
24	YA	1627	A
24	YA	1628	G
24	YA	1631	C
24	YA	1632	A
24	YA	1633	A
24	YA	1644	C
24	YA	1653	C
24	YA	1654	A
24	YA	1662	A
24	YA	1663	C
24	YA	1664	A
24	YA	1681	A
24	YA	1687	C
24	YA	1693	C
24	YA	1694	G
24	YA	1695	C
24	YA	1701	A
24	YA	1711	A
24	YA	1716	A
24	YA	1721	G
24	YA	1740	U
24	YA	1742	G
24	YA	1763	G
24	YA	1766	G
24	YA	1767	A
24	YA	1769	G
24	YA	1770	A
24	YA	1776	G
24	YA	1785	C
24	YA	1787	G
24	YA	1793	A
24	YA	1794	G
24	YA	1795	G
24	YA	1800	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	YA	1804	A
24	YA	1811	A
24	YA	1815	A
24	YA	1818	A
24	YA	1822	A
24	YA	1830	G
24	YA	1831	C
24	YA	1832	G
24	YA	1846	A
24	YA	1847	G
24	YA	1859	G
24	YA	1866	G
24	YA	1878	A
24	YA	1889	G
24	YA	1896	G
24	YA	1897	C
24	YA	1899	A
24	YA	1900	G
24	YA	1903	C
24	YA	1911	A
24	YA	1925	G
24	YA	1936	C
24	YA	1941	A
24	YA	1951	G
24	YA	1952	G
24	YA	1958	A
24	YA	1961	U
24	YA	1962	U
24	YA	1977	U
24	YA	1985	U
24	YA	1986	G
24	YA	1989	C
24	YA	1991	A
24	YA	1992	A
24	YA	1993	A
24	YA	1994	A
24	YA	2004	C
24	YA	2013	U
24	YA	2014	G
24	YA	2015	U
24	YA	2018	C
24	YA	2042	A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	YA	2043	C
24	YA	2045	G
24	YA	2053	A
24	YA	2055	A
24	YA	2061	C
24	YA	2065	C
24	YA	2074	G
24	YA	2077	C
24	YA	2078	G
24	YA	2081	A
24	YA	2082	A
24	YA	2083	G
24	YA	2084	A
24	YA	2091	G
24	YA	2115	G
24	YA	2121	U
24	YA	2128	G
24	YA	2129	C
24	YA	2133	C
24	YA	2134	G
24	YA	2135	U
24	YA	2136	A
24	YA	2137	G
24	YA	2138	G
24	YA	2140	U
24	YA	2141	A
24	YA	2142	G
24	YA	2144	U
24	YA	2148	A
24	YA	2149	G
24	YA	2150	C
24	YA	2153	G
24	YA	2154	U
24	YA	2155	G
24	YA	2156	A
24	YA	2158	C
24	YA	2159	C
24	YA	2162	C
24	YA	2167	C
24	YA	2169	G
24	YA	2170	G
24	YA	2178	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	YA	2179	G
24	YA	2180	A
24	YA	2186	C
24	YA	2188	G
24	YA	2190	G
24	YA	2191	A
24	YA	2193	A
24	YA	2195	A
24	YA	2202	U
24	YA	2203	G
24	YA	2210	C
24	YA	2211	U
24	YA	2212	G
24	YA	2214	G
24	YA	2215	G
24	YA	2220	A
24	YA	2227	G
24	YA	2228	G
24	YA	2229	A
24	YA	2231	G
24	YA	2237	A
24	YA	2250	G
24	YA	2251	G
24	YA	2255	U
24	YA	2258	G
24	YA	2278	A
24	YA	2287	C
24	YA	2290	A
24	YA	2291	G
24	YA	2292	G
24	YA	2295	C
24	YA	2299	A
24	YA	2300	A
24	YA	2319	G
24	YA	2320	G
24	YA	2323	A
24	YA	2332	A
24	YA	2337	G
24	YA	2346	G
24	YA	2347	A
24	YA	2358	A
24	YA	2359	C

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	YA	2362	C
24	YA	2366	G
24	YA	2370	G
24	YA	2395	G
24	YA	2397	C
24	YA	2404	A
24	YA	2408	G
24	YA	2415	C
24	YA	2418	U
24	YA	2422	G
24	YA	2435	U
24	YA	2436	C
24	YA	2437	A
24	YA	2440	G
24	YA	2441	G
24	YA	2442	A
24	YA	2447	A
24	YA	2451	A
24	YA	2453	C
24	YA	2459	G
24	YA	2460	A
24	YA	2462	A
24	YA	2477	C
24	YA	2481	A
24	YA	2487	C
24	YA	2506	G
24	YA	2510	C
24	YA	2514	G
24	YA	2517	G
24	YA	2530	A
24	YA	2537	G
24	YA	2541	G
24	YA	2566	U
24	YA	2570	C
24	YA	2579	G
24	YA	2585	C
24	YA	2597	U
24	YA	2614	A
24	YA	2621	U
24	YA	2623	U
24	YA	2624	C
24	YA	2627	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	YA	2641	A
24	YA	2658	C
24	YA	2666	A
24	YA	2667	G
24	YA	2668	U
24	YA	2677	A
24	YA	2685	G
24	YA	2687	A
24	YA	2694	U
24	YA	2701	U
24	YA	2703	C
24	YA	2714	U
24	YA	2719	G
24	YA	2724	U
24	YA	2725	A
24	YA	2726	A
24	YA	2727	G
24	YA	2739	U
24	YA	2746	A
24	YA	2752	U
24	YA	2757	G
24	YA	2761	A
24	YA	2771	A
24	YA	2774	G
24	YA	2775	G
24	YA	2777	A
24	YA	2778	A
24	YA	2790	G
24	YA	2791	A
24	YA	2792	U
24	YA	2803	A
24	YA	2804	C
24	YA	2808	G
24	YA	2809	U
24	YA	2810	C
24	YA	2818	U
24	YA	2830	A
24	YA	2831	A
24	YA	2843	G
24	YA	2844	G
24	YA	2845	A
24	YA	2855	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	YA	2856	G
24	YA	2876	U
24	YA	2878	A
24	YA	2882	G
24	YA	2883	A
24	YA	2901	A
24	YA	2902	G
24	YA	2903	G
25	YB	8	U
25	YB	9	G
25	YB	13	A
25	YB	15	A
25	YB	30	C
25	YB	33	G
25	YB	41	U
25	YB	51	G
25	YB	56	G
25	YB	64	C
25	YB	65	C
25	YB	67	G
25	YB	73	A
25	YB	84	C
25	YB	89(A)	G
25	YB	89(B)	A
25	YB	90	C
25	YB	106	G
25	YB	108	C
25	YB	109	G
25	YB	110	G

All (157) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
1	QA	5	U
1	QA	31	G
1	QA	64	G
1	QA	115	G
1	QA	119	A
1	QA	181	G
1	QA	243	A
1	QA	244	U
1	QA	250	A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	QA	266	G
1	QA	328	C
1	QA	410	G
1	QA	412	A
1	QA	429	U
1	QA	481	G
1	QA	484	G
1	QA	485	G
1	QA	509	A
1	QA	687	A
1	QA	703	G
1	QA	753	A
1	QA	792	A
1	QA	812	C
1	QA	913	A
1	QA	960	U
1	QA	992	U
1	QA	1064	G
1	QA	1065	U
1	QA	1200	C
1	QA	1201	A
1	QA	1285	A
1	QA	1297	C
1	QA	1336	C
1	QA	1346	A
1	QA	1347	G
1	QA	1446	A
1	QA	1453	G
1	QA	1498	U
1	QA	1528	U
22	QV	53	G
23	QX	12	A
24	RA	73	A
24	RA	98	U
24	RA	210	A
24	RA	211	A
24	RA	218	A
24	RA	287	G
24	RA	399	G
24	RA	431	C
24	RA	528	A
24	RA	537	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	RA	610	C
24	RA	662	A
24	RA	799	A
24	RA	893	C
24	RA	903	C
24	RA	1068	G
24	RA	1072	U
24	RA	1099	C
24	RA	1103	A
24	RA	1111	U
24	RA	1119	A
24	RA	1255	A
24	RA	1358	U
24	RA	1473	A
24	RA	1605	A
24	RA	1649	A
24	RA	1700	G
24	RA	1850	A
24	RA	2014	G
24	RA	2082	A
24	RA	2148	A
24	RA	2451	A
24	RA	2578	A
24	RA	2622	C
24	RA	2701	U
24	RA	2842	U
24	RA	2877	G
25	RB	66	A
1	XA	5	U
1	XA	31	G
1	XA	60	A
1	XA	64	G
1	XA	78	G
1	XA	115	G
1	XA	243	A
1	XA	244	U
1	XA	250	A
1	XA	266	G
1	XA	328	C
1	XA	345	C
1	XA	358	U
1	XA	410	G

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	XA	412	A
1	XA	429	U
1	XA	484	G
1	XA	485	G
1	XA	509	A
1	XA	687	A
1	XA	703	G
1	XA	753	A
1	XA	812	C
1	XA	913	A
1	XA	991	U
1	XA	992	U
1	XA	1027	C
1	XA	1126	U
1	XA	1285	A
1	XA	1297	C
1	XA	1310	G
1	XA	1446	A
1	XA	1498	U
22	XV	53	G
24	YA	98	U
24	YA	184	A
24	YA	210	A
24	YA	218	A
24	YA	231	G
24	YA	287	G
24	YA	302	A
24	YA	399	G
24	YA	431	C
24	YA	528	A
24	YA	533	G
24	YA	662	A
24	YA	799	A
24	YA	893	C
24	YA	903	C
24	YA	906	G
24	YA	1058	U
24	YA	1068	G
24	YA	1072	U
24	YA	1131	A
24	YA	1155	C
24	YA	1176	U

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
24	YA	1223	C
24	YA	1249	A
24	YA	1255	A
24	YA	1473	A
24	YA	1507	A
24	YA	1554	A
24	YA	1605	A
24	YA	1694	G
24	YA	1700	G
24	YA	1741	C
24	YA	1830	G
24	YA	1935	A
24	YA	2014	G
24	YA	2414	C
24	YA	2578	A
24	YA	2622	C
24	YA	2693	C
24	YA	2724	U
24	YA	2789	A
24	YA	2842	U
24	YA	2877	G
25	YB	109	G

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

2 non-standard protein/DNA/RNA residues are modelled in this entry.

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

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
22	1MG	XV	37	22	18,26,27	1.37	3 (16%)	19,39,42	1.70	3 (15%)
22	1MG	QV	37	22	18,26,27	1.34	2 (11%)	19,39,42	1.72	4 (21%)

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
22	1MG	XV	37	22	-	0/3/25/26	0/3/3/3
22	1MG	QV	37	22	-	0/3/25/26	0/3/3/3

All (5) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	XV	37	1MG	C6-C5	4.46	1.48	1.41
22	QV	37	1MG	C6-C5	4.44	1.48	1.41
22	QV	37	1MG	C5-C4	2.53	1.47	1.40
22	XV	37	1MG	C5-C4	2.48	1.47	1.40
22	XV	37	1MG	C6-N1	2.07	1.41	1.38

All (7) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	XV	37	1MG	C2-N3-C4	5.60	121.76	115.36
22	QV	37	1MG	C2-N3-C4	5.47	121.61	115.36
22	QV	37	1MG	C6-C5-C4	-2.94	118.07	119.96
22	XV	37	1MG	C6-C5-C4	-2.93	118.08	119.96
22	XV	37	1MG	C4-C5-N7	-2.91	106.36	109.40
22	QV	37	1MG	C4-C5-N7	-2.67	106.61	109.40
22	QV	37	1MG	C1'-N9-C4	2.01	130.18	126.64

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
22	QV	37	1MG	1	0

5.5 Carbohydrates [i](#)

There are no monosaccharides in this entry.

5.6 Ligand geometry [i](#)

Of 1166 ligands modelled in this entry, 1164 are monoatomic - leaving 2 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	SF4	QD	301	4	0,12,12	0.00	-	-		
56	SF4	XD	301	4	0,12,12	0.00	-	-		

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	SF4	QD	301	4	-	-	0/6/5/5
56	SF4	XD	301	4	-	-	0/6/5/5

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

No monomer is involved in short contacts.

5.7 Other polymers [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

EDS failed to run properly - this section is therefore empty.

6.2 Non-standard residues in protein, DNA, RNA chains

EDS failed to run properly - this section is therefore empty.

6.3 Carbohydrates

EDS failed to run properly - this section is therefore empty.

6.4 Ligands

EDS failed to run properly - this section is therefore empty.

6.5 Other polymers

EDS failed to run properly - this section is therefore empty.