



## Full wwPDB EM Validation Report ⓘ

Jan 24, 2023 – 08:08 PM EST

PDB ID : 4V7E  
EMDB ID : EMD-1780  
Title : Model of the small subunit RNA based on a 5.5 Å cryo-EM map of *Triticum aestivum* translating 80S ribosome  
Authors : Barrio-Garcia, C.; Armache, J.-P.; Jarasch, A.; Anger, A.M.; Villa, E.; Becker, T.; Bhushan, S.; Jossinet, F.; Habeck, M.; Dindar, G.; Franckenberg, S.; Marquez, V.; Mielke, T.; Thomm, M.; Berninghausen, O.; Beatrix, B.; Soeding, J.; Westhof, E.; Wilson, D.N.; Beckmann, R.  
Deposited on : 2013-11-22  
Resolution : 5.50 Å(reported)

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

We welcome your comments at [validation@mail.wwpdb.org](mailto:validation@mail.wwpdb.org)

A user guide is available at

<https://www.wwpdb.org/validation/2017/EMValidationReportHelp>

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

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

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

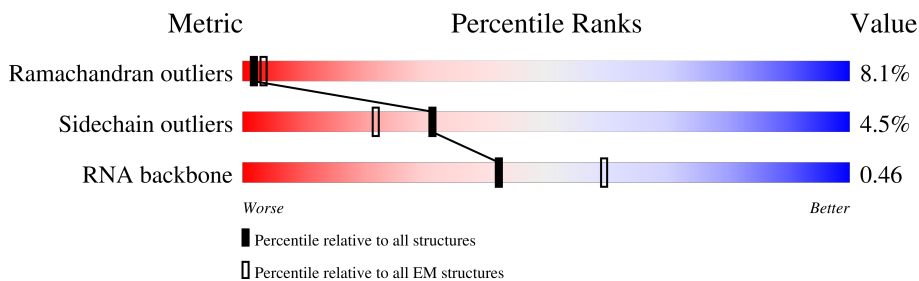
EMDB validation analysis : 0.0.1.dev43  
MolProbity : 4.02b-467  
Percentile statistics : 20191225.v01 (using entries in the PDB archive December 25th 2019)  
MapQ : 1.9.9  
Ideal geometry (proteins) : Engh & Huber (2001)  
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)  
Validation Pipeline (wwPDB-VP) : 2.31.2

# 1 Overall quality at a glance

The following experimental techniques were used to determine the structure:  
*ELECTRON MICROSCOPY*

The reported resolution of this entry is 5.50 Å.

Percentile scores (ranging between 0-100) for global validation metrics of the entry are shown in the following graphic. The table shows the number of entries on which the scores are based.



Metric	Whole archive (#Entries)	EM structures (#Entries)
Ramachandran outliers	154571	4023
Sidechain outliers	154315	3826
RNA backbone	4643	859

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

Mol	Chain	Length	Quality of chain
1	Ad	1810	
2	Ae	75	
3	Af	11	
4	BY	138	
5	BI	220	
6	BK	183	
7	BM	171	
8	Bf	155	

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Mol	Chain	Length	Quality of chain
9	BX	142	48% 95%
10	Bg	380	77% 92% 7%
11	BD	208	58% 75% 23%
12	BE	265	32% 70% 5% 25%
13	BF	191	38% 92% 7%
14	BQ	149	40% 68% 14% 15%
15	BU	128	59% 87% 10%
16	BO	151	41% 67% 11% 21%
17	BS	152	48% 78% 20%
18	BN	151	45% 66% 13% 20%
19	BL	160	31% 46% 7% 47%
20	BT	146	44% 89% 8%
21	BP	154	29% 51% 5% 41%
22	BZ	108	49% 77% 11% 5% 7%
23	Bc	65	60% 66% 18% 5% 11%
24	BW	130	56% 86% 12%
25	Bd	56	50% 62% 16% 7% 14%
26	Bb	86	49% 94% 6%
27	Be	62	58% 89% 8%
28	BA	260	42% 69% 7% 24%
29	BR	141	50% 73% 9% 18%
30	BB	262	38% 69% 11% 19%
31	BV	82	57% 84% 9% 7%
32	Ba	133	26% 59% 11% 30%
33	BJ	195	31% 90% 6%

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Mol	Chain	Length	Quality of chain
34	BC	263	44% 77% 19%
35	BG	245	43% 88% 5% • 6%
36	BH	189	52% 83% 12% • • •
37	CG	257	38% 84% 8% 8%
38	CT	164	47% 85% 12% •
39	CZ	136	33% 96% •
40	Cz	216	98% 92% 8%
41	CA	261	54% 86% 10% • •
42	CJ	180	34% 78% 12% • • 6%
43	CH	190	41% 93% 6% •
44	CV	140	74% 93% 7%
45	CN	200	35% 89% 8% •
46	Ca	144	50% 77% 22% • •
47	CQ	188	39% 73% 12% • • 13%
48	CD	304	37% 75% 18% 6% •
49	CR	209	44% 80% 8% • 10%
50	CP	171	42% 86% 12% •
51	CX	152	40% 73% 6% • 20%
52	CW	162	31% 43% • • 54%
53	CY	150	21% 79% 6% • 13%
54	Cr	147	18% 42% 6% • 50%
55	Cc	112	45% 92% 7% •
56	Cd	123	46% 84% 12% • • •
57	Ce	133	58% 89% 8% •
58	Cj	94	39% 80% 19% •

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Mol	Chain	Length	Quality of chain
59	Cl	51	47% 86% 12%
60	Co	105	48% 79% 18%
61	CM	134	35% 83% 14%
62	CS	178	37% 76% 14% 6%
63	CU	130	53% 62% 15% 5% 17%
64	Ci	112	31% 55% 11% 31%
65	CK	166	73% 65% 10% 23%
66	Cu	110	53% 51% 47%
66	Cv	110	53% 52% 47%
67	Cs	113	52% 51% 48%
67	Ct	113	52% 50% 48%
68	Ch	124	40% 89% 10%
69	CF	244	41% 92% 7%
70	Cq	319	76% 76% 5% 18%
71	CB	389	44% 83% 14%
72	CC	405	45% 81% 8% 8%
73	CO	206	42% 88% 8%
74	Cp	92	50% 95% 5%
75	CI	224	36% 74% 8% 18%
76	Cn	25	72% 92% 8%
77	Cm	53	30% 94% ..
78	CL	208	40% 84% 12%
79	CE	219	53% 80% 15% ..
80	Cf	111	45% 94% 5%
81	Ck	69	45% 90% 10%

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Mol	Chain	Length	Quality of chain
82	Cb	60	<p>57% 90% 10% 8%</p>
83	Cg	119	<p>41% 82% 10% 8%</p>
84	Aa	3391	<p>11% 71% 25%</p>
85	Ac	160	<p>5% 68% 28%</p>
86	Ab	120	<p>67% 32%</p>

## 2 Entry composition [i](#)

There are 86 unique types of molecules in this entry. The entry contains 212263 atoms, of which 0 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a RNA chain called 18S ribosomal RNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	P		
1	Ad	1762	37584	16788	6708	12327	1761	0	0

- Molecule 2 is a RNA chain called P-site tRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	P		
2	Ae	75	1595	712	280	529	74	0	0

- Molecule 3 is a RNA chain called 5'-R(\*AP\*AP\*AP\*AP\*GP\*AP\*CP\*UP\*UP\*CP\*A)-3'.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	P		
3	Af	11	232	106	45	71	10	0	0

- Molecule 4 is a protein called 40S ribosomal protein S24E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
4	BY	138	1108	703	212	189	4	0	0

- Molecule 5 is a protein called 40S ribosomal protein S8E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
5	BI	66	533	330	105	95	3	0	0

- Molecule 6 is a protein called 40S ribosomal protein S10E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
6	BK	96	818	535	137	143	3	0	0

- Molecule 7 is a protein called 40S ribosomal protein S12E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
7	BM	123	924	577	159	179	9	0	0

- Molecule 8 is a protein called 40S ribosomal protein S31e.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
8	Bf	71	577	367	107	98	5	0	0

- Molecule 9 is a protein called 40S ribosomal protein S12.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
9	BX	142	1103	698	214	187	4	0	0

- Molecule 10 is a protein called RACK1.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
10	Bg	380	2929	1813	530	567	19	0	0

- Molecule 11 is a protein called 40S ribosomal protein S3.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
11	BD	208	1629	1029	294	297	9	0	0

- Molecule 12 is a protein called 40S ribosomal protein S4E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
12	BE	200	1607	1030	290	283	4	0	0

- Molecule 13 is a protein called 40S ribosomal protein S7.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
13	BF	191	1489	928	281	273	7	0	0

- Molecule 14 is a protein called 40S ribosomal protein S9.



Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
14	BQ	126	1017	648	195	170	4	0	0

- Molecule 15 is a protein called 40S ribosomal protein S10.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
15	BU	128	982	613	176	187	6	0	0

- Molecule 16 is a protein called 40S ribosomal protein S11.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
16	BO	119	899	550	178	167	4	0	0

- Molecule 17 is a protein called 40S ribosomal protein S13.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
17	BS	152	1240	772	248	213	7	0	0

- Molecule 18 is a protein called 40S ribosomal protein S15.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
18	BN	121	977	627	180	167	3	0	0

- Molecule 19 is a protein called 40S ribosomal protein S17.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
19	BL	85	688	435	134	115	4	0	0

- Molecule 20 is a protein called 40S ribosomal protein S19E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
20	BT	146	1155	726	218	207	4	0	0

- Molecule 21 is a protein called 40S ribosomal protein S19.

Mol	Chain	Residues	Atoms					AltConf	Trace
21	BP	91	Total	C	N	O	S	0	0
			711	457	130	120	4		

- Molecule 22 is a protein called 40S ribosomal protein S25E.

Mol	Chain	Residues	Atoms					AltConf	Trace
22	BZ	100	Total	C	N	O	S	0	0
			779	489	146	144			

- Molecule 23 is a protein called 40S ribosomal protein S28E.

Mol	Chain	Residues	Atoms					AltConf	Trace
23	Bc	58	Total	C	N	O	S	0	0
			454	281	86	84	3		

- Molecule 24 is a protein called 40S ribosomal protein S8.

Mol	Chain	Residues	Atoms					AltConf	Trace
24	BW	130	Total	C	N	O	S	0	0
			1042	667	189	181	5		

- Molecule 25 is a protein called 40S ribosomal protein S14.

Mol	Chain	Residues	Atoms					AltConf	Trace
25	Bd	48	Total	C	N	O	S	0	0
			379	233	77	63	6		

- Molecule 26 is a protein called 40S ribosomal protein S27E.

Mol	Chain	Residues	Atoms					AltConf	Trace
26	Bb	86	Total	C	N	O	S	0	0
			663	414	119	122	8		

- Molecule 27 is a protein called 40S ribosomal protein S30E.

Mol	Chain	Residues	Atoms					AltConf	Trace
27	Be	60	Total	C	N	O	S	0	0
			469	289	104	75	1		

- Molecule 28 is a protein called 40S ribosomal protein S2.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
28	BA	197	1537	969	280	278	10	0	0

- Molecule 29 is a protein called 40S ribosomal protein S17E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
29	BR	116	945	589	178	171	7	0	0

- Molecule 30 is a protein called 40S ribosomal protein S1E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
30	BB	211	1707	1089	308	302	8	0	0

- Molecule 31 is a protein called 40S ribosomal protein S21E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
31	BV	76	601	371	112	115	3	0	0

- Molecule 32 is a protein called 40S ribosomal protein S26E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
32	Ba	93	753	461	163	122	7	0	0

- Molecule 33 is a protein called 40S ribosomal protein S4.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
33	BJ	187	1525	959	305	256	5	0	0

- Molecule 34 is a protein called 40S ribosomal protein S5.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
34	BC	214	1665	1074	297	287	7	0	0

- Molecule 35 is a protein called 40S ribosomal protein S6.

Mol	Chain	Residues	Atoms					AltConf	Trace
35	BG	231	Total	C	N	O	S	0	0
			1867	1164	367	328	8		

- Molecule 36 is a protein called 40S ribosomal protein S7E.

Mol	Chain	Residues	Atoms					AltConf	Trace
36	BH	184	Total	C	N	O	S	0	0
			1508	962	278	266	2		

- Molecule 37 is a protein called 60S ribosomal protein L8E.

Mol	Chain	Residues	Atoms					AltConf	Trace
37	CG	237	Total	C	N	O	S	0	0
			1906	1226	351	322	7		

- Molecule 38 is a protein called 60S ribosomal protein L21E.

Mol	Chain	Residues	Atoms					AltConf	Trace
38	CT	160	Total	C	N	O	S	0	0
			1288	814	251	219	4		

- Molecule 39 is a protein called 60S ribosomal protein L27E.

Mol	Chain	Residues	Atoms					AltConf	Trace
39	CZ	136	Total	C	N	O	S	0	0
			1090	704	205	176	5		

- Molecule 40 is a protein called 60S ribosomal protein L1.

Mol	Chain	Residues	Atoms					AltConf	Trace
40	Cz	216	Total	C	N	O	S	0	0
			1718	1092	309	304	13		

- Molecule 41 is a protein called 60S ribosomal protein L2.

Mol	Chain	Residues	Atoms					AltConf	Trace
41	CA	255	Total	C	N	O	S	0	0
			1946	1210	399	328	9		

- Molecule 42 is a protein called 60S ribosomal protein L5.

Mol	Chain	Residues	Atoms					AltConf	Trace
42	CJ	170	Total	C	N	O	S	0	0
			1380	869	256	246	9		

- Molecule 43 is a protein called 60S ribosomal protein L6.

Mol	Chain	Residues	Atoms					AltConf	Trace
43	CH	190	Total	C	N	O	S	0	0
			1500	947	270	277	6		

- Molecule 44 is a protein called 60S ribosomal protein L14.

Mol	Chain	Residues	Atoms					AltConf	Trace
44	CV	140	Total	C	N	O	S	0	0
			1048	658	199	181	10		

- Molecule 45 is a protein called 60S ribosomal protein L15E.

Mol	Chain	Residues	Atoms					AltConf	Trace
45	CN	194	Total	C	N	O	S	0	0
			1630	1027	342	257	4		

- Molecule 46 is a protein called 60S ribosomal protein L15.

Mol	Chain	Residues	Atoms					AltConf	Trace
46	Ca	144	Total	C	N	O	S	0	0
			1114	710	223	175	6		

- Molecule 47 is a protein called 60S ribosomal protein L18E.

Mol	Chain	Residues	Atoms					AltConf	Trace
47	CQ	163	Total	C	N	O	S	0	0
			1284	810	248	219	7		

- Molecule 48 is a protein called 60S ribosomal protein L18.

Mol	Chain	Residues	Atoms					AltConf	Trace
48	CD	304	Total	C	N	O	S	0	0
			2444	1531	440	466	7		

- Molecule 49 is a protein called 60S ribosomal protein L19E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
49	CR	189	1569	972	330	257	10	0	0

- Molecule 50 is a protein called 60S ribosomal protein L22.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
50	CP	171	1372	852	271	244	5	0	0

- Molecule 51 is a protein called 60S ribosomal protein L23.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
51	CX	122	987	634	178	173	2	0	0

- Molecule 52 is a protein called 60S ribosomal protein L24E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
52	CW	75	635	408	126	97	4	0	0

- Molecule 53 is a protein called 60S ribosomal protein L24.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
53	CY	130	1048	647	220	178	3	0	0

- Molecule 54 is a protein called 60S ribosomal protein L28E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
54	Cr	73	576	364	107	103	2	0	0

- Molecule 55 is a protein called 60S ribosomal protein L30E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
55	Cc	112	857	540	149	161	7	0	0

- Molecule 56 is a protein called 60S ribosomal protein L31E.

Mol	Chain	Residues	Atoms					AltConf	Trace
56	Cd	120	Total	C	N	O	S	0	0
			960	598	186	173	3		

- Molecule 57 is a protein called 60S ribosomal protein L32E.

Mol	Chain	Residues	Atoms					AltConf	Trace
57	Ce	133	Total	C	N	O	S	0	0
			1103	696	216	185	6		

- Molecule 58 is a protein called 60S ribosomal protein L37E.

Mol	Chain	Residues	Atoms					AltConf	Trace
58	Cj	94	Total	C	N	O	S	0	0
			755	459	166	123	7		

- Molecule 59 is a protein called 60S ribosomal protein L39E.

Mol	Chain	Residues	Atoms					AltConf	Trace
59	Cl	51	Total	C	N	O	S	0	0
			460	291	100	67	2		

- Molecule 60 is a protein called 60S ribosomal protein L44E.

Mol	Chain	Residues	Atoms					AltConf	Trace
60	Co	105	Total	C	N	O	S	0	0
			851	535	166	144	6		

- Molecule 61 is a protein called 60S ribosomal protein L14E.

Mol	Chain	Residues	Atoms					AltConf	Trace
61	CM	134	Total	C	N	O	S	0	0
			1081	690	201	185	5		

- Molecule 62 is a protein called 60S ribosomal protein L20.

Mol	Chain	Residues	Atoms					AltConf	Trace
62	CS	167	Total	C	N	O	S	0	0
			1419	916	263	233	7		

- Molecule 63 is a protein called 60S ribosomal protein L22E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
63	CU	108	864	551	155	156	2	0	0

- Molecule 64 is a protein called 60S ribosomal protein L36E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
64	Ci	77	613	383	128	100	2	0	0

- Molecule 65 is a protein called 60S ribosomal protein L11.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
65	CK	128	960	602	177	177	4	0	0

- Molecule 66 is a protein called 60S ribosomal protein P1.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
66	Cu	58	432	283	69	79	1	0	0
66	Cv	58	432	283	69	79	1	0	0

- Molecule 67 is a protein called Acidic ribosomal protein P2.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
67	Cs	59	441	278	69	90	4	0	0
67	Ct	59	441	278	69	90	4	0	0

- Molecule 68 is a protein called 60S ribosomal protein L29.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
68	Ch	124	1012	636	202	173	1	0	0

- Molecule 69 is a protein called 60S ribosomal protein L30.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
69	CF	244	1984	1271	368	339	6	0	0



- Molecule 70 is a protein called 60S acidic ribosomal protein P0.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
70	Cq	262	1993	1278	330	377	8	0	0

- Molecule 71 is a protein called 60S ribosomal protein L3.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
71	CB	389	3139	1997	584	540	18	0	0

- Molecule 72 is a protein called 60S ribosomal protein L4.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
72	CC	372	2898	1823	556	510	9	0	0

- Molecule 73 is a protein called 60S ribosomal protein L13.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
73	CO	206	1650	1045	320	274	11	0	0

- Molecule 74 is a protein called 60S ribosomal protein L43E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
74	Cp	92	715	447	137	124	7	0	0

- Molecule 75 is a protein called 60S ribosomal protein L16.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
75	CI	184	1490	941	290	247	12	0	0

- Molecule 76 is a protein called 60S ribosomal protein L41E.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
76	Cn	25	238	145	62	28	3	0	0

- Molecule 77 is a protein called 60S ribosomal protein L40E.

Mol	Chain	Residues	Atoms					AltConf	Trace
77	Cm	52	Total	C	N	O	S	0	0
			428	267	90	66	5		

- Molecule 78 is a protein called 60S ribosomal protein L13E.

Mol	Chain	Residues	Atoms					AltConf	Trace
78	CL	208	Total	C	N	O	S	0	0
			1691	1061	338	286	6		

- Molecule 79 is a protein called 60S ribosomal protein L6E.

Mol	Chain	Residues	Atoms					AltConf	Trace
79	CE	219	Total	C	N	O	S	0	0
			1731	1106	314	307	4		

- Molecule 80 is a protein called 60S ribosomal protein L33E.

Mol	Chain	Residues	Atoms					AltConf	Trace
80	Cf	111	Total	C	N	O	S	0	0
			891	561	170	156	4		

- Molecule 81 is a protein called 60S ribosomal protein L38E.

Mol	Chain	Residues	Atoms					AltConf	Trace
81	Ck	69	Total	C	N	O	S	0	0
			564	360	104	97	3		

- Molecule 82 is a protein called 60S ribosomal protein L29E.

Mol	Chain	Residues	Atoms					AltConf	Trace
82	Cb	58	Total	C	N	O	S	0	0
			477	288	103	85	1		

- Molecule 83 is a protein called 60S ribosomal protein L34E.

Mol	Chain	Residues	Atoms					AltConf	Trace
83	Cg	110	Total	C	N	O	S	0	0
			897	567	182	146	2		

- Molecule 84 is a RNA chain called 60S ribosomal RNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	P		
84	Aa	3391	72601	32373	13241	23598	3389	0	0

- Molecule 85 is a RNA chain called 5.8S ribosomal RNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	P		
85	Ac	160	3408	1522	614	1113	159	0	0

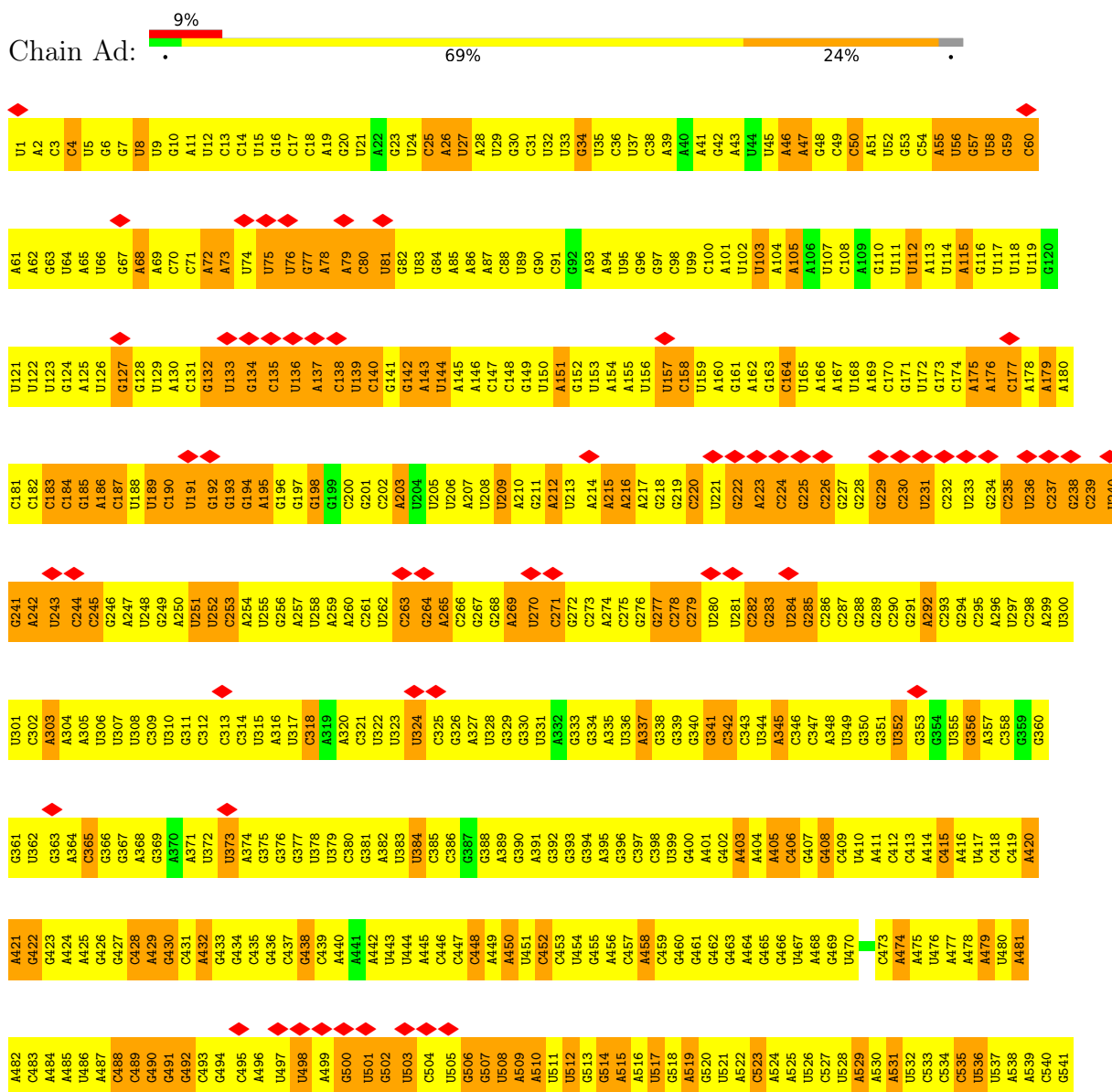
- Molecule 86 is a RNA chain called 5S ribosomal RNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	P		
86	Ab	120	2561	1144	461	837	119	0	0

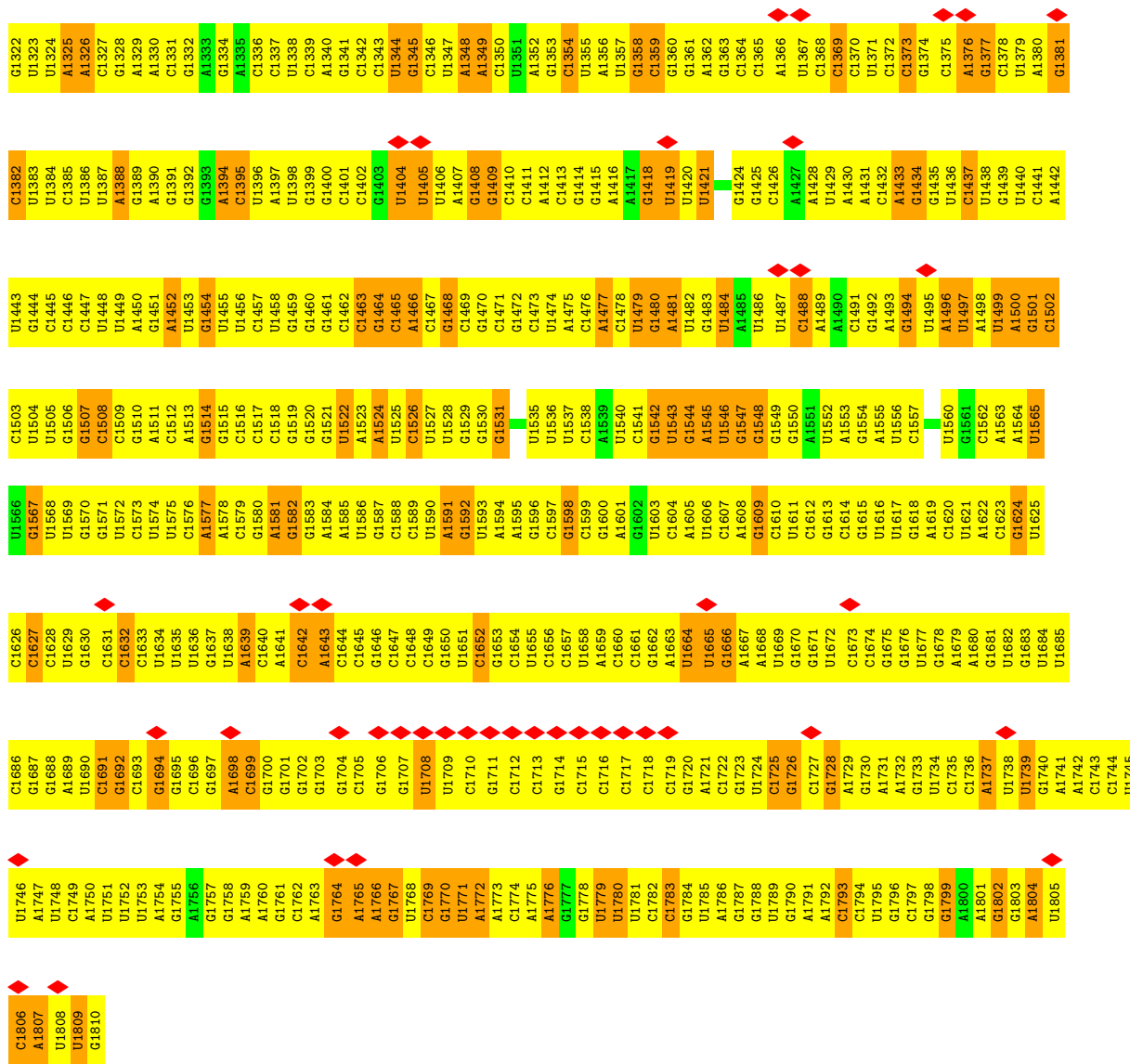
### 3 Residue-property plots [i](#)

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

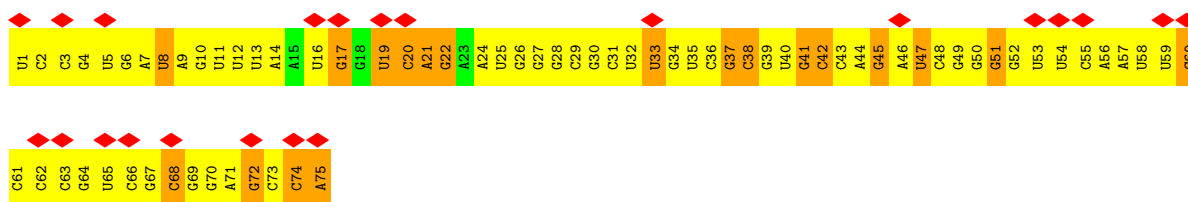
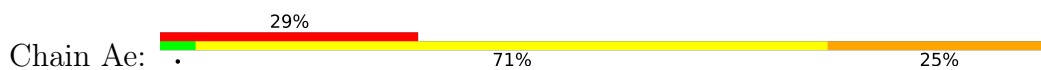
- Molecule 1: 18S ribosomal RNA



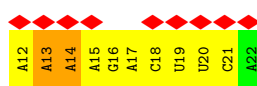
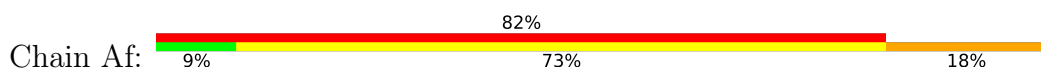
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A1142	A1143	A1144	G1145	G1146	A1147	A1148	U1149	U1150	A1151	A1152	C1153	G1154	G1155	A1156	A1157	G1158	G1159	G1160	C1161	A1162	C1163	C1164	A1165	C1166	C1167	A1168	G1169	C1170	C1171	G1172	U1173	G1174	G1175	A1176	G1177	C1178	C1179	C1239	A1240	G1241	A1242	C1243	U1244	G1245	A1246	G1247	A1248	U1249	C1250	C1251	C1252	U1253	U1254	U1255	C1256	U1257	U1258	C1199	A1200	C1280	U1261																																								
C1082	C1083	U1084	U1085	A1086	U1087	G1088	A1089	G1090	A1091	A1092	A1093	G1094	C1095	A1096	A1097	A1098	U1099	U1100	C1101	U1102	U1103	U1104	G1105	G1106	G1107	U1108	U1109	C1110	C1111	G1112	G1113	G1114	G1115	G1116	G1117	A1118	U1119	U1120	A1121	U1122	G1123	G1124	U1125	C1126	G1127	A1128	U1129	U1191	A1192	A1193	C1194	U1195	C1196	A1197	A1198	C1199	A1200	C1201	U1141																																										
U1022	C1023	A1024	A1025	U1026	C1027	A1028	U1029	A1030	A1031	A1032	A1033	G1034	A1035	U1036	G1037	C1038	C1039	G1040	A1041	C1042	C1043	A1044	G1045	G1046	G1047	A1048	U1049	G990	G991	G992	U993	U994	C995	G996	A997	A998	G999	U940	G941	C942	G943	U1004	C1005	A1006	G1007	A1008	U1009	A1010	C1011	G1012	G1013	U1014	C1015	C1016	U1017	A1018	G1019	U1020	C1021	A1081																																									
C902	A903	G904	A905	G906	G907	U908	G909	A910	G949	A911	A912	U913	U914	C915	U916	U917	G918	G919	A920	U921	U922	A923	U924	U925	G926	A927	G988	A929	G930	A931	C932	G933	A934	A935	C936	A937	A938	C939	U940	G941	C942	G943	A944	A945	A946	G947	C948	A949	U950	U951	U952	G953	C954	C955	A956	A957	G958	G959	A960	U961																																									
G82	C783	C784	A785	C845	U786	C787	G788	C848	G849	C850	C791	U792	G793	U794	A795	U796	A797	C798	A799	U800	U801	A802	G803	C804	A805	U806	G807	G808	G809	A810	U811	A812	A813	C814	A815	U816	C817	A818	U819	A820	G821	G822	U824	U825	C826	C827	G828	G829	U830	C831	C832	A833	U834	U835	U836	G837	U838	A839	U840	A781																																									
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U602	A603	U604	A605	U606	U607	U608	A609	A610	U611	U612	U613	G614	U615	U616	G617	C618	A619	G620	U621	A622	A623	A624	A625	A626	A627	G628	C629	U630	C631	G632	U633	A634	G635	U636	U637	G638	G639	A640	U641	C642	U643	U644	G645	U646	G647	C648	C649	G650	G651	U670	U671	U672	U673	U674	U675	U676	U677	U678	U679	U680	U681	U682	U683	U684	U685	U686	U687	U688	U689	U690	U691	U692	U693	U694	U695	U696	U697	U698	U699	U700	U701	U702	U703	C704	A705	U706	C707	G708	C709	G710	C711	U712	C713	G714	C715	U716	C717	C718	U719	U720	U721
A542	G543	G544	A545	U546	C547	C548	A549	U550	U551	G552	G553	A554	G555	U556	G557	C558	A559	A560	G561	U562	C563	U564	G565	U566	G567	G568	C569	C570	A571	G572	C573	A574	G575	U576	U577	G578	C579	G580	G581	U582	A583	A584	U585	U586	C587	C588	A589	G590	C591	U592	C593	C594	A595	U596	U597	A598	G599	C600	G601																																										



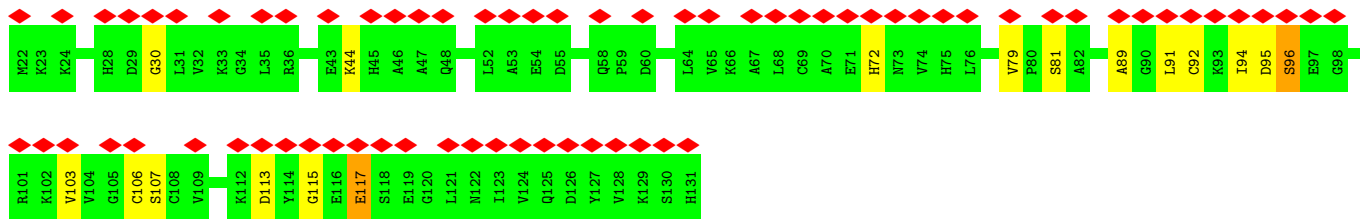
• Molecule 2: P-site tRNA



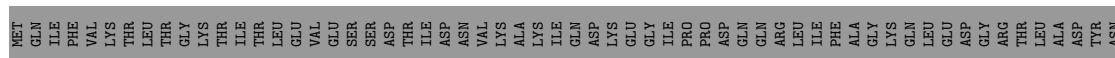
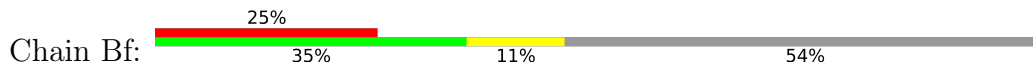
• Molecule 3: 5'-R(\*AP\*AP\*AP\*AP\*GP\*AP\*CP\*UP\*UP\*CP\*A)-3'



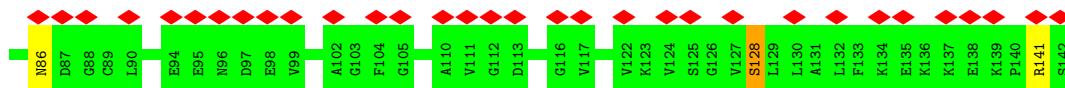
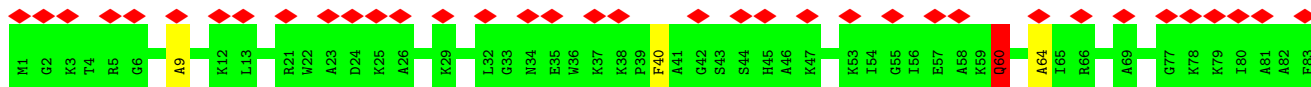




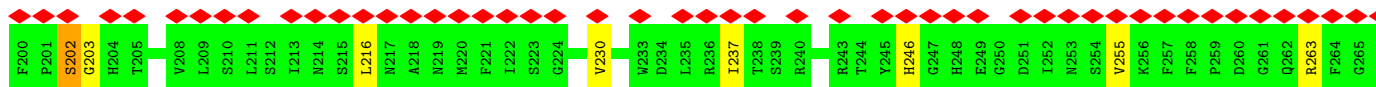
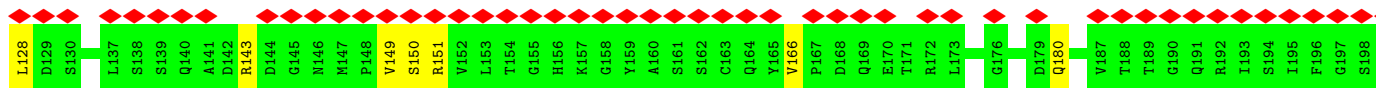
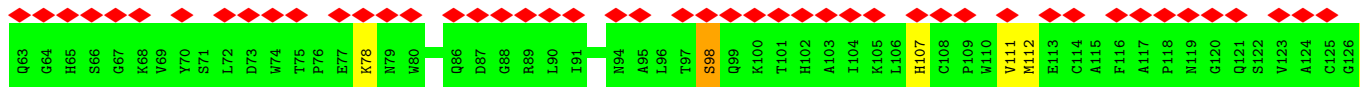
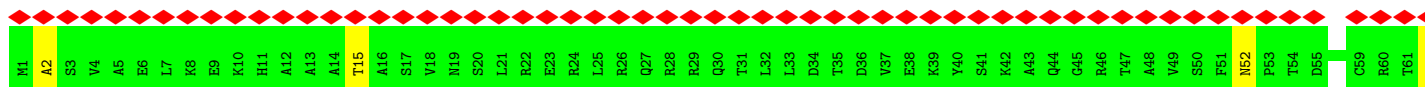
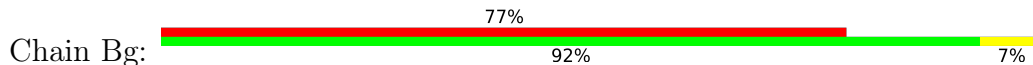
• Molecule 8: 40S ribosomal protein S31e



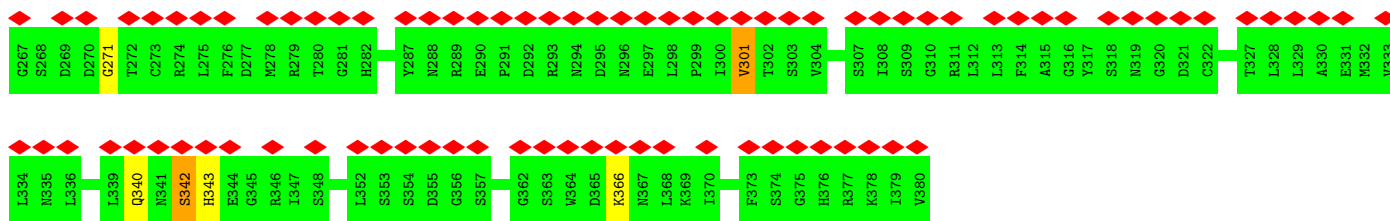
• Molecule 9: 40S ribosomal protein S12



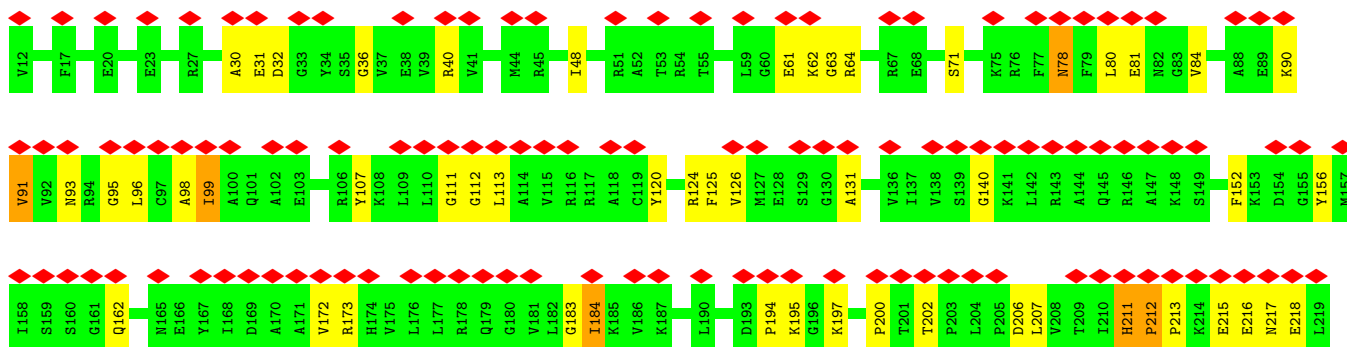
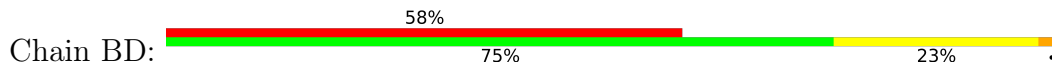
• Molecule 10: RACK1



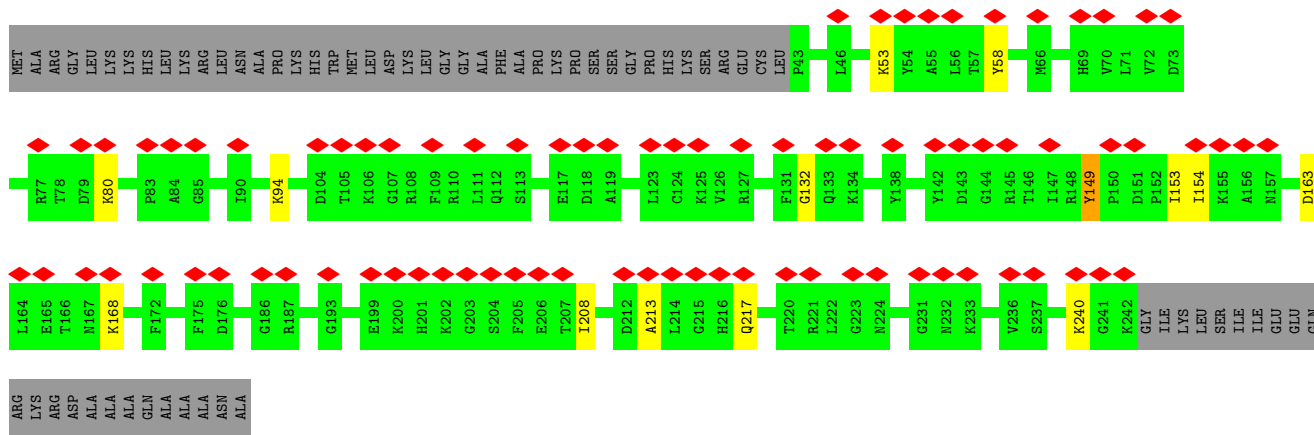




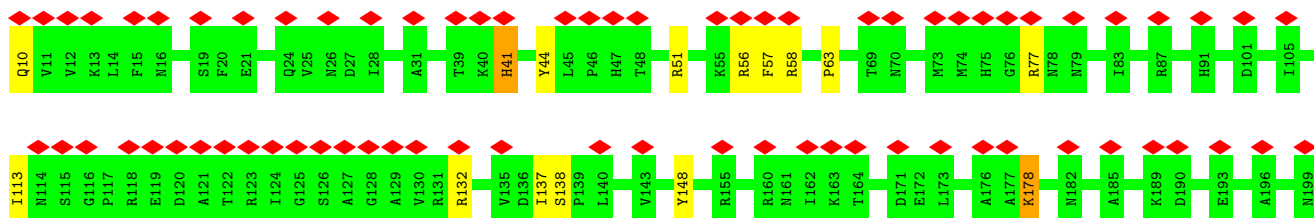
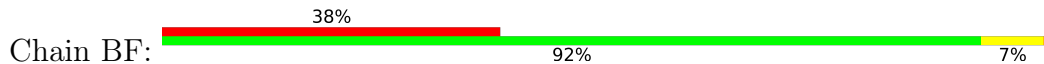
- Molecule 11: 40S ribosomal protein S3



- Molecule 12: 40S ribosomal protein S4E

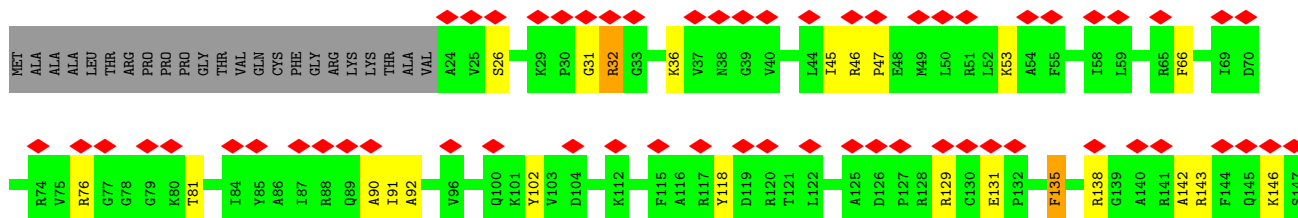
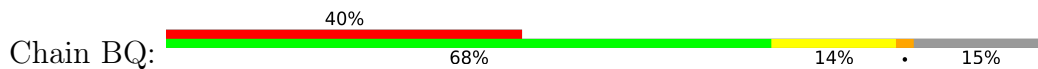


- Molecule 13: 40S ribosomal protein S7



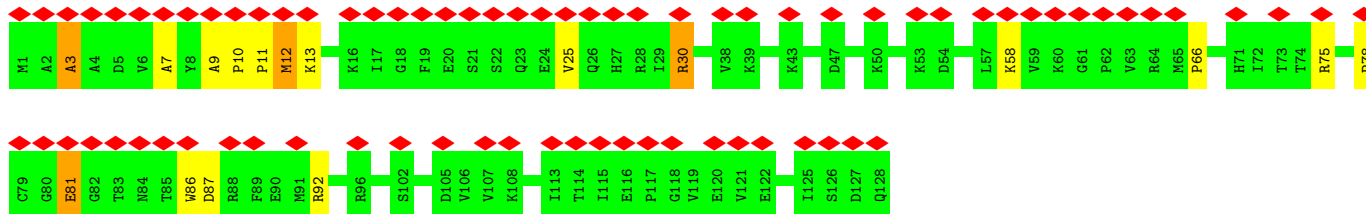
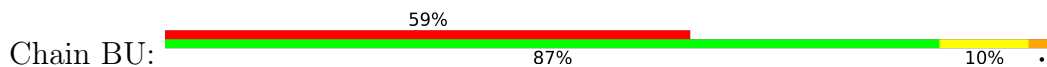
R200

- Molecule 14: 40S ribosomal protein S9

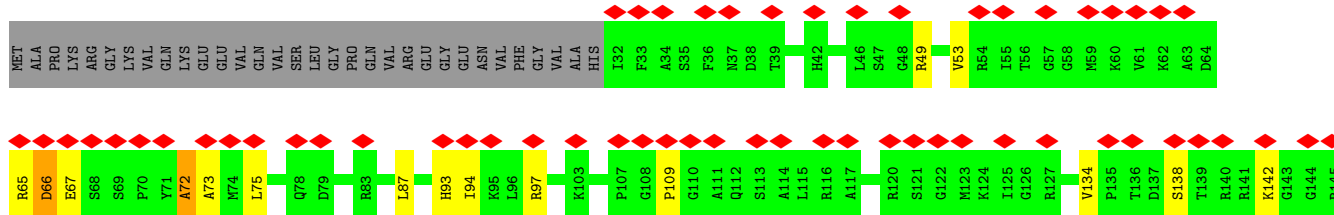
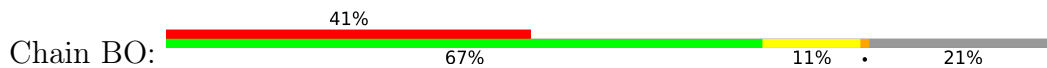


Y148 R149

- Molecule 15: 40S ribosomal protein S10

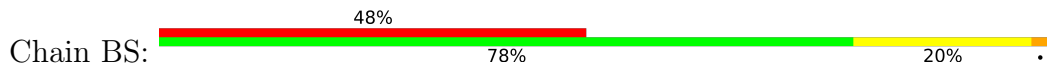


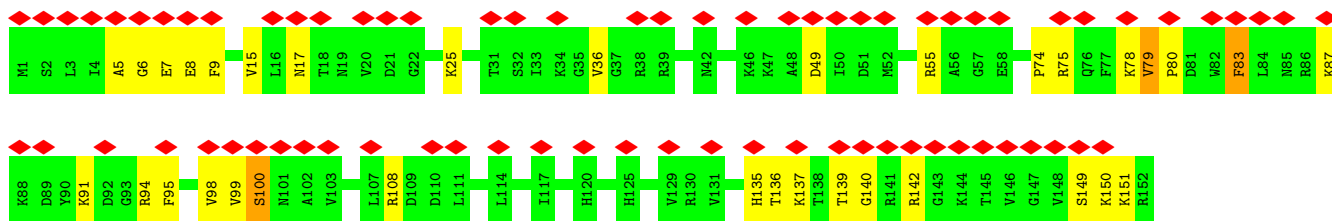
- Molecule 16: 40S ribosomal protein S11



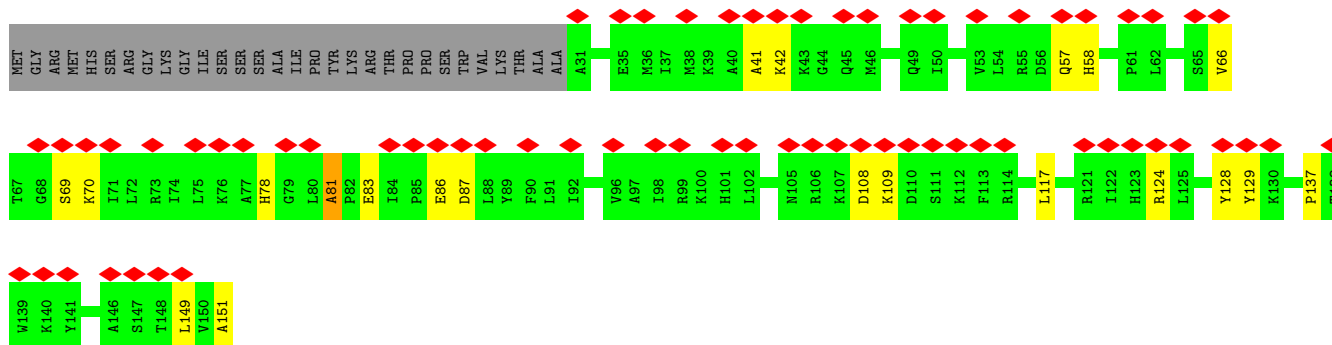
R146 G147 R148 R149 L150

- Molecule 17: 40S ribosomal protein S13

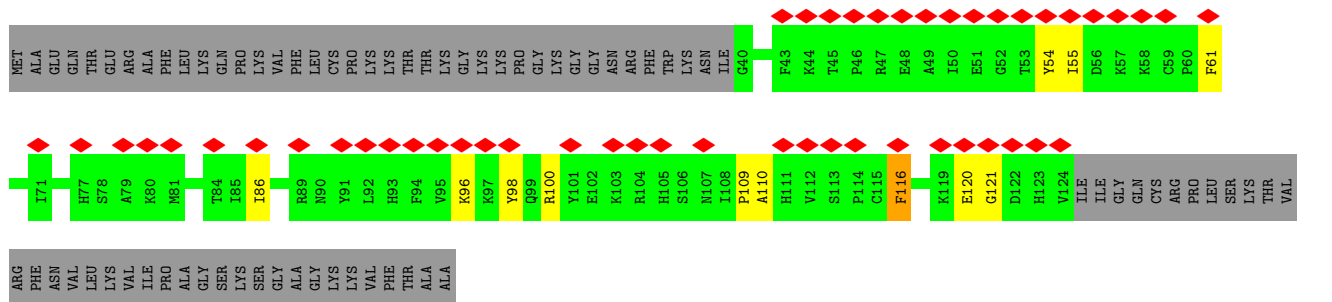




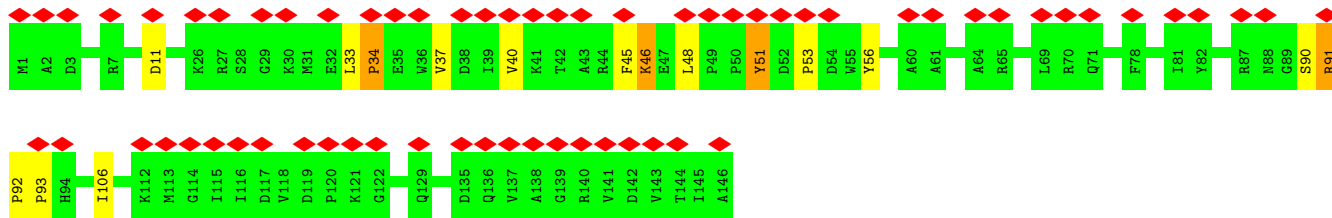
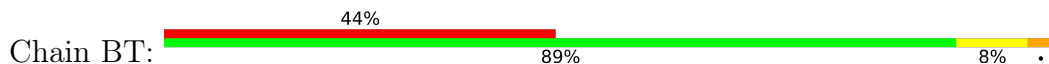
• Molecule 18: 40S ribosomal protein S15



• Molecule 19: 40S ribosomal protein S17



• Molecule 20: 40S ribosomal protein S19E

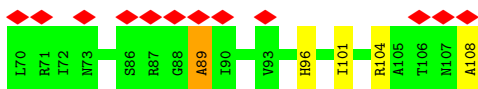
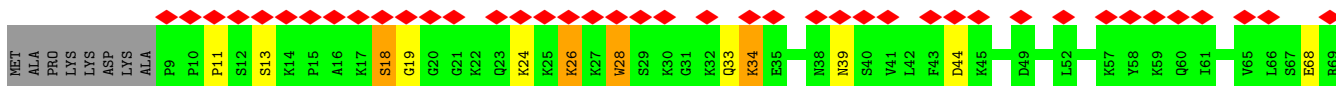
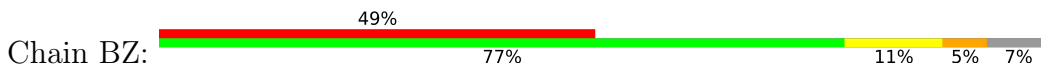


• Molecule 21: 40S ribosomal protein S19

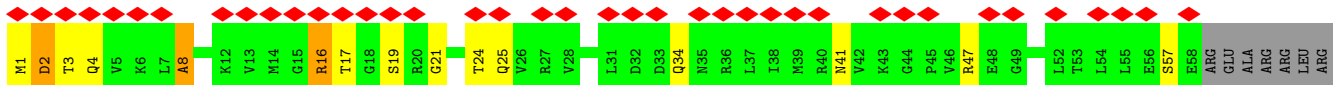




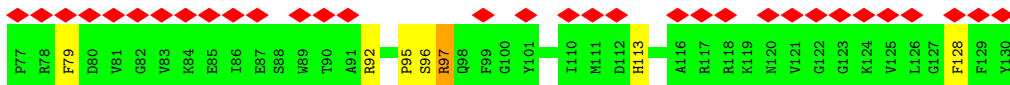
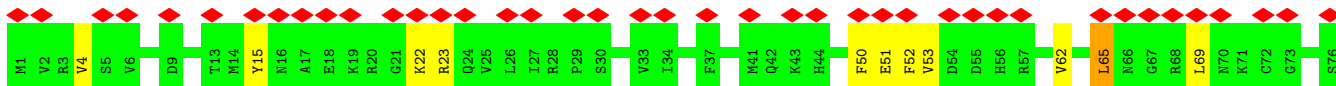
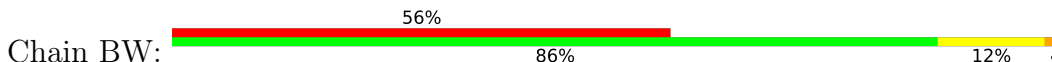
- Molecule 22: 40S ribosomal protein S25E



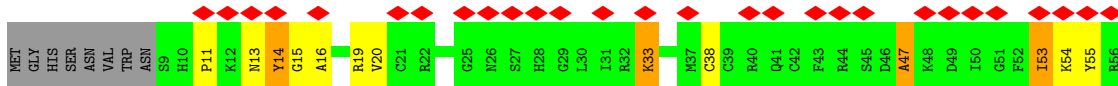
- Molecule 23: 40S ribosomal protein S28E



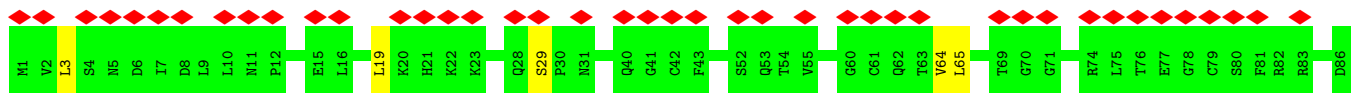
- Molecule 24: 40S ribosomal protein S8



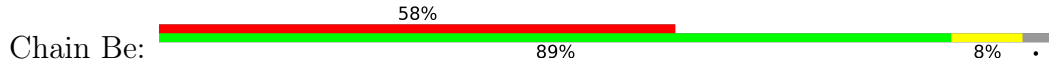
- Molecule 25: 40S ribosomal protein S14



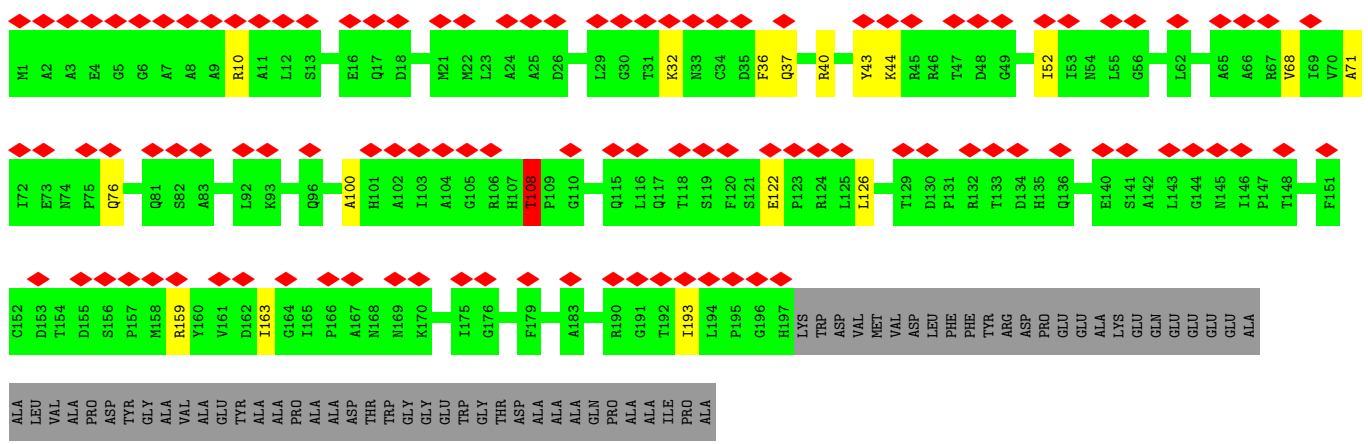
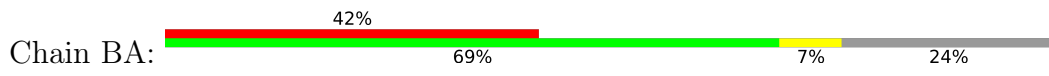
- Molecule 26: 40S ribosomal protein S27E



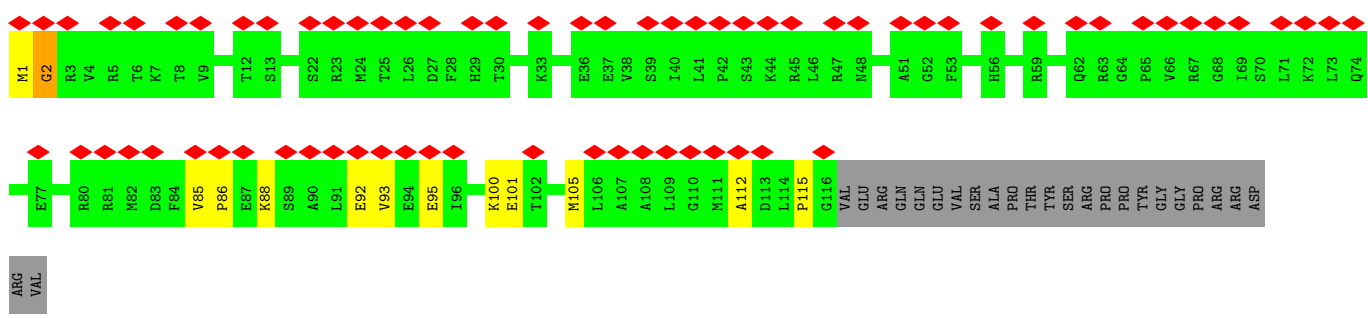
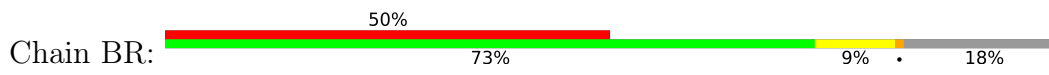
• Molecule 27: 40S ribosomal protein S30E



• Molecule 28: 40S ribosomal protein S2

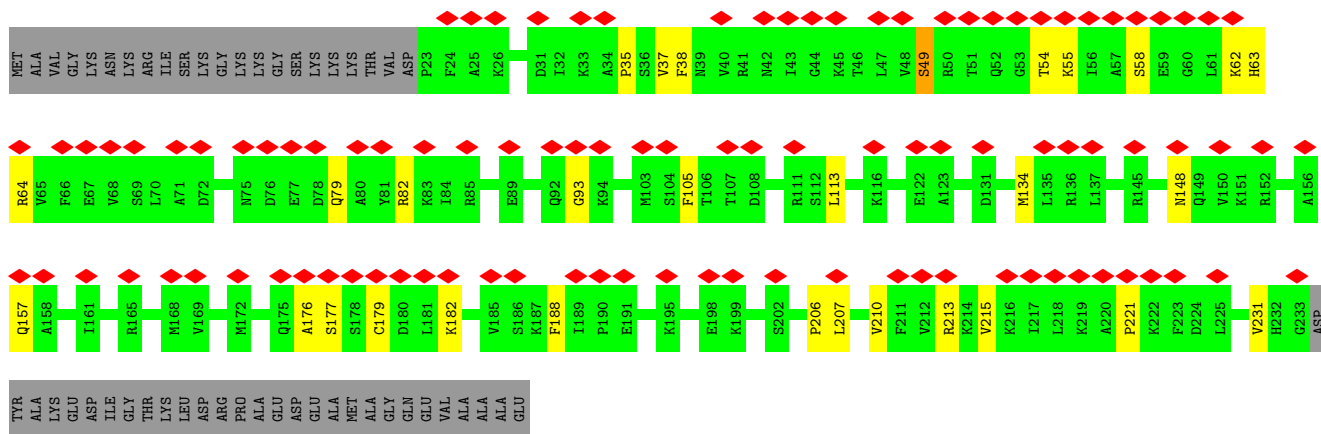


• Molecule 29: 40S ribosomal protein S17E

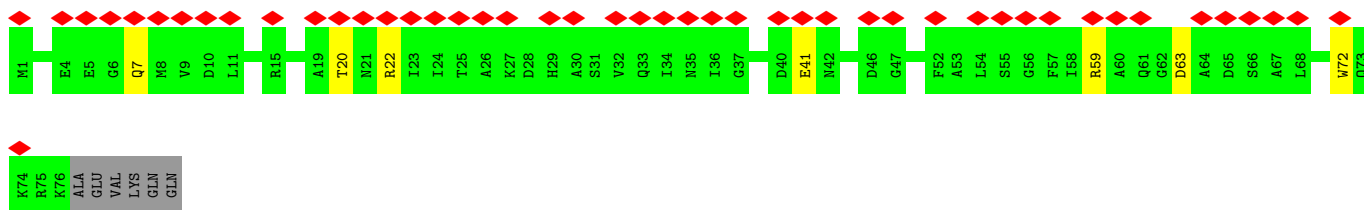
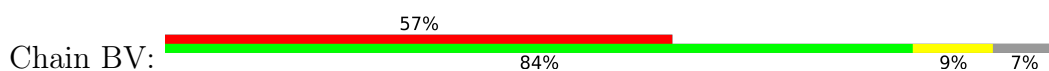


• Molecule 30: 40S ribosomal protein S1E

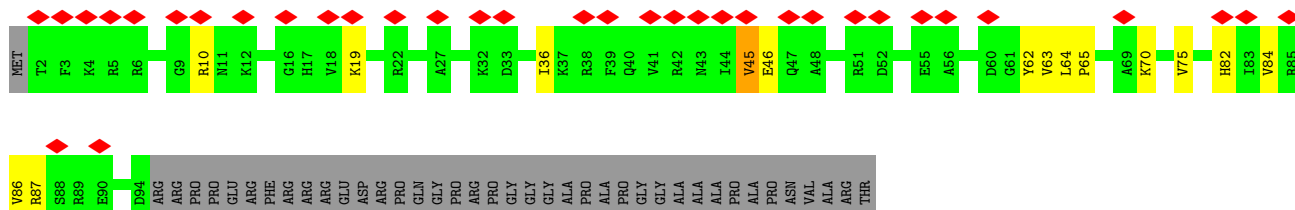




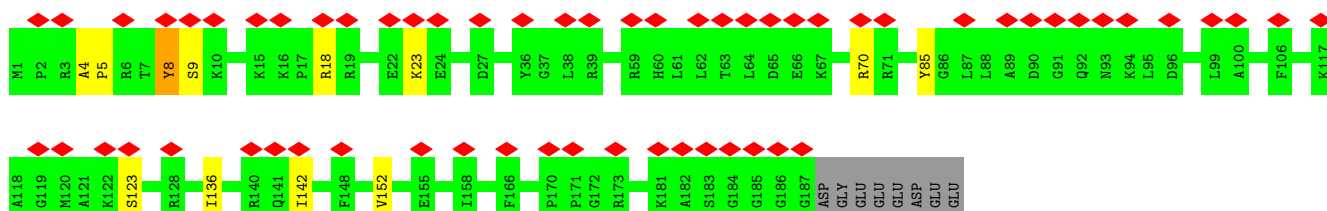
• Molecule 31: 40S ribosomal protein S21E



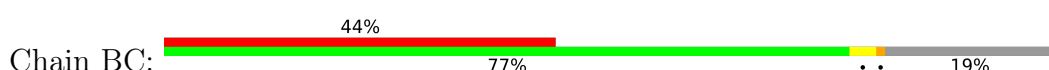
• Molecule 32: 40S ribosomal protein S26E

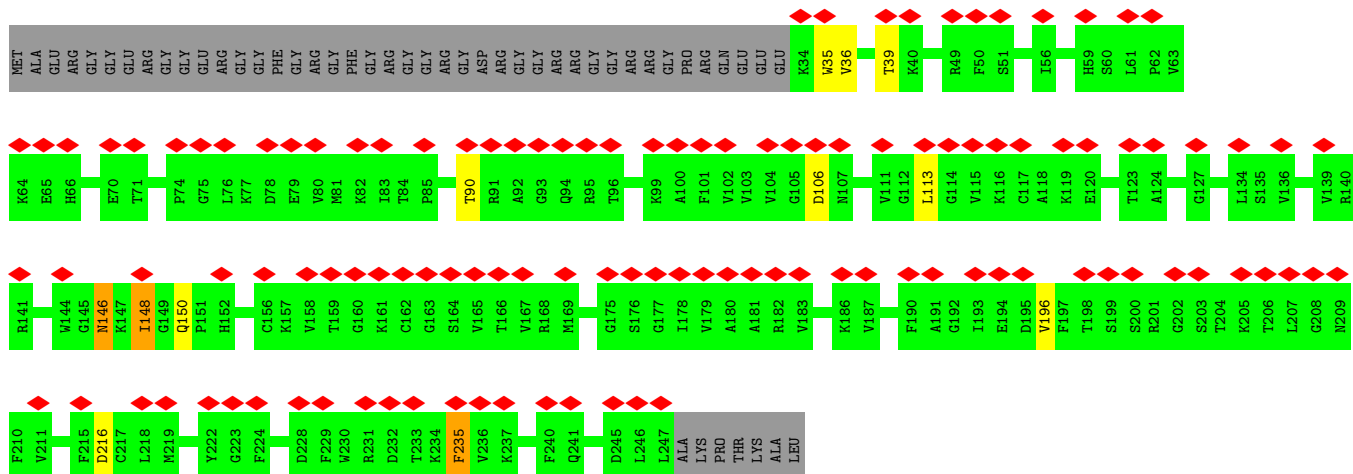


• Molecule 33: 40S ribosomal protein S4

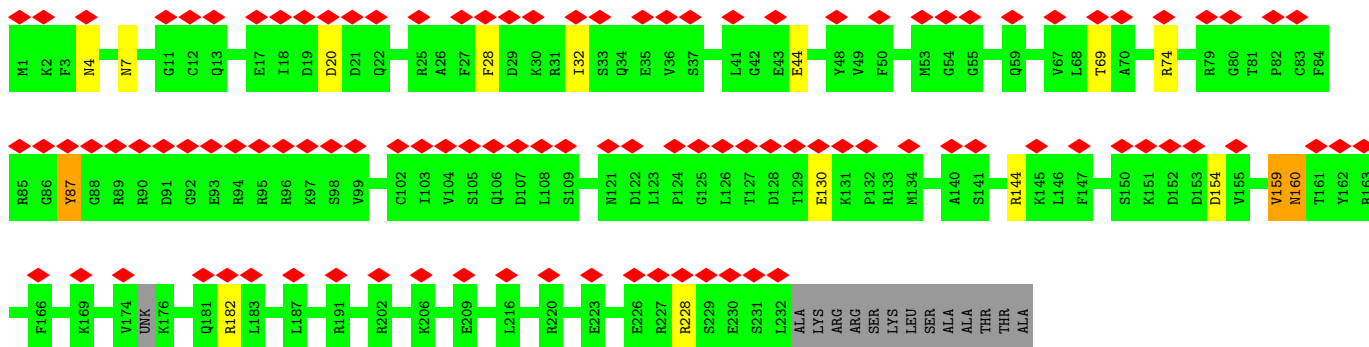
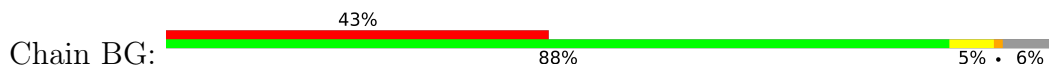


• Molecule 34: 40S ribosomal protein S5

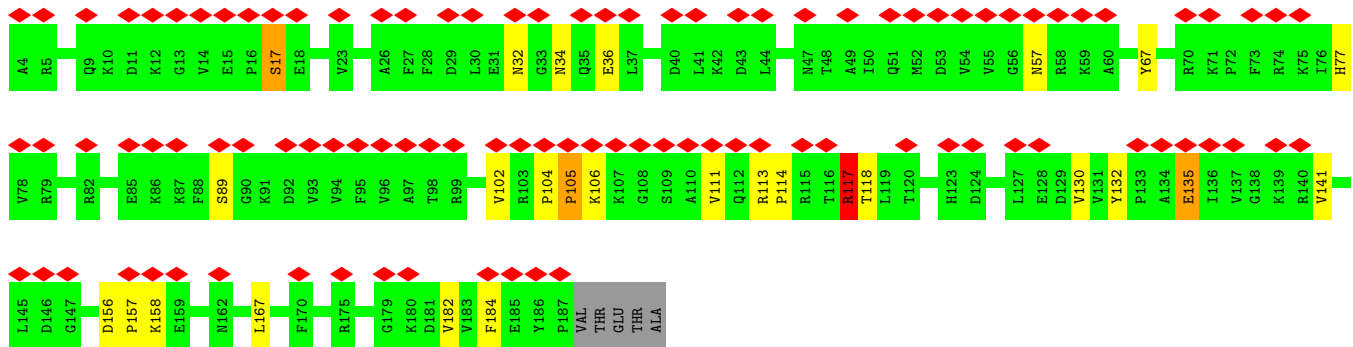
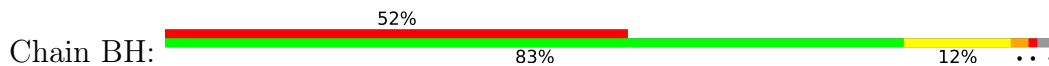




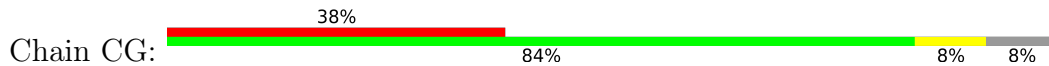
• Molecule 35: 40S ribosomal protein S6

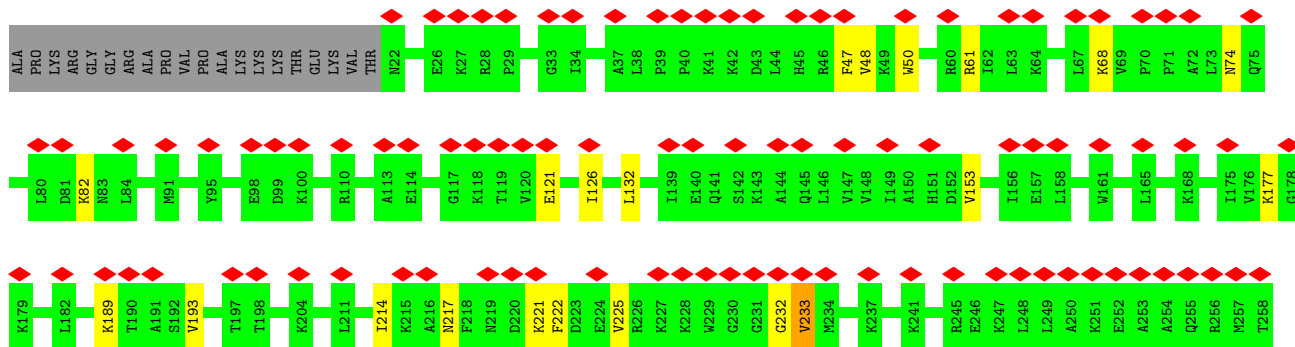


• Molecule 36: 40S ribosomal protein S7E

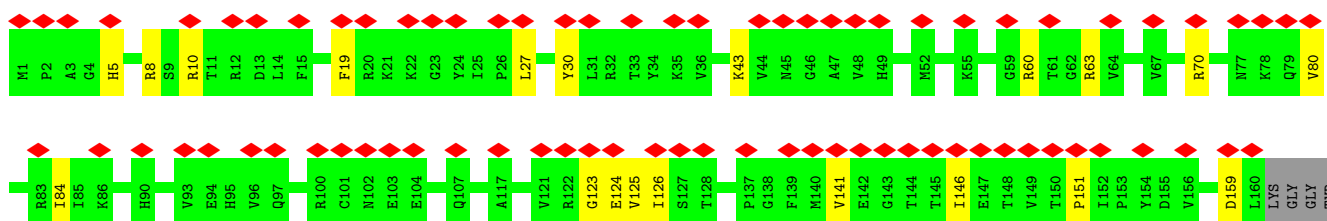
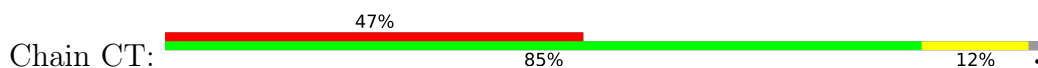


• Molecule 37: 60S ribosomal protein L8E

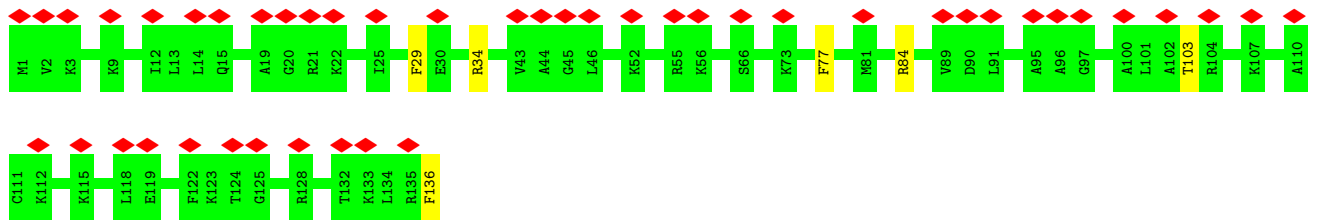




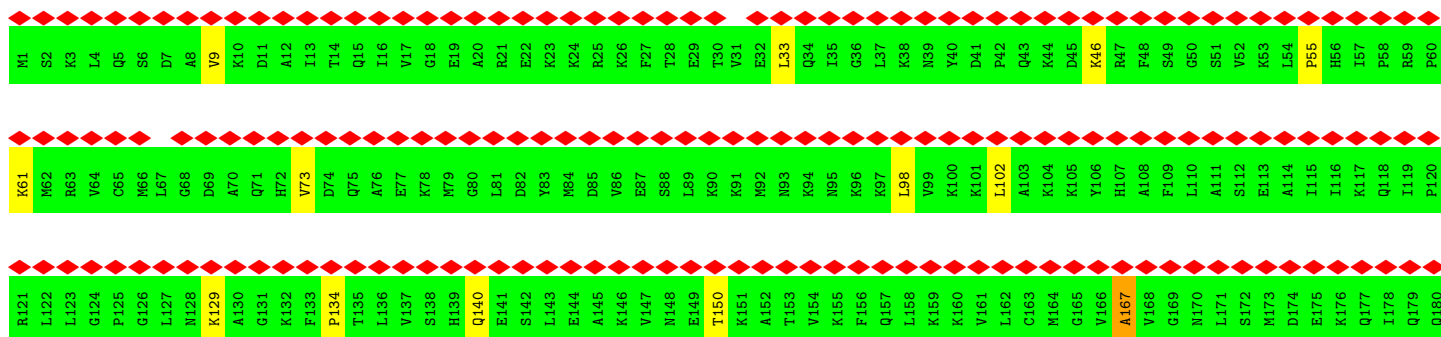
• Molecule 38: 60S ribosomal protein L21E



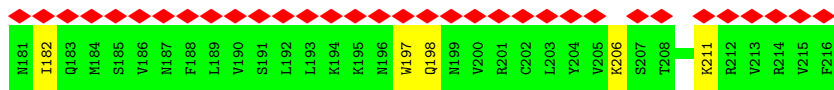
• Molecule 39: 60S ribosomal protein L27E



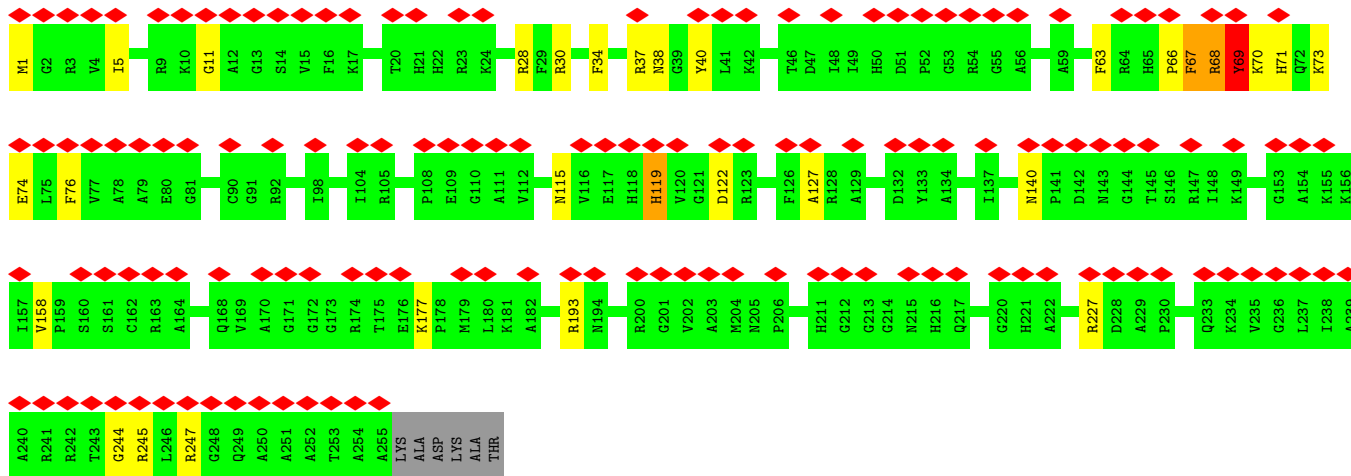
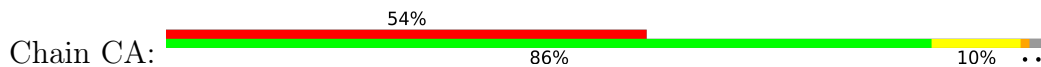
• Molecule 40: 60S ribosomal protein L1



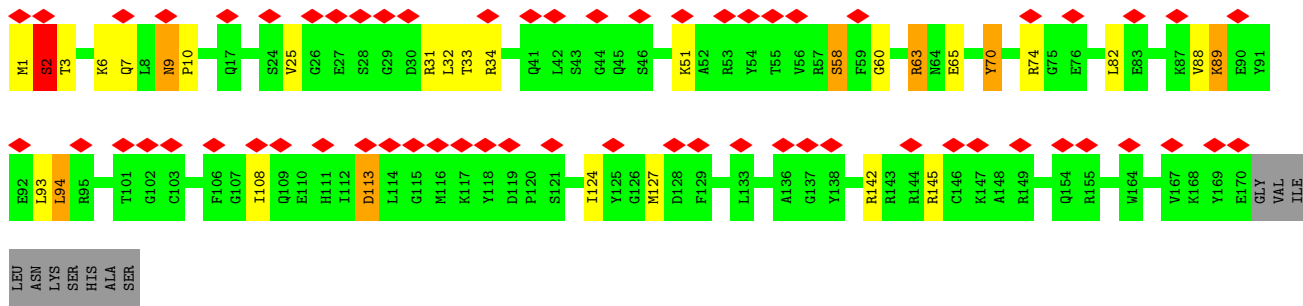
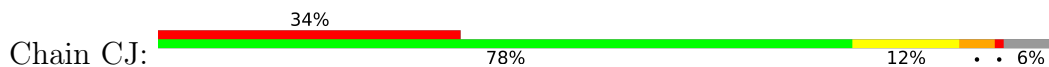




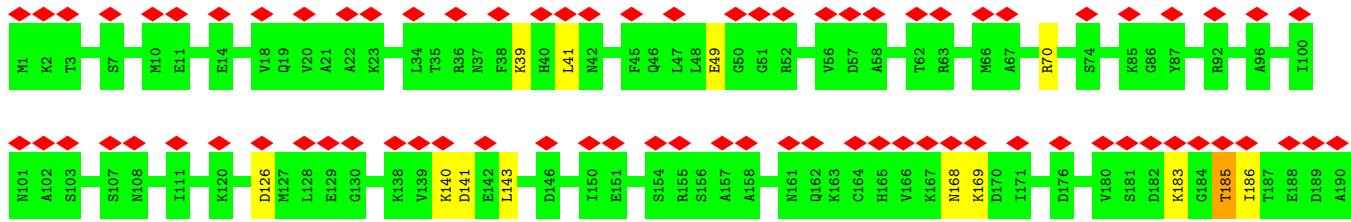
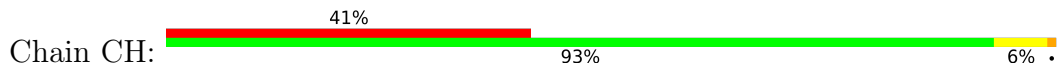
• Molecule 41: 60S ribosomal protein L2



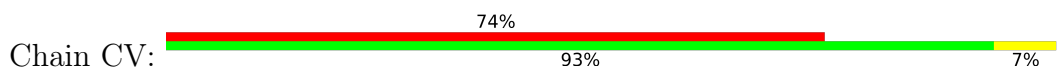
• Molecule 42: 60S ribosomal protein L5

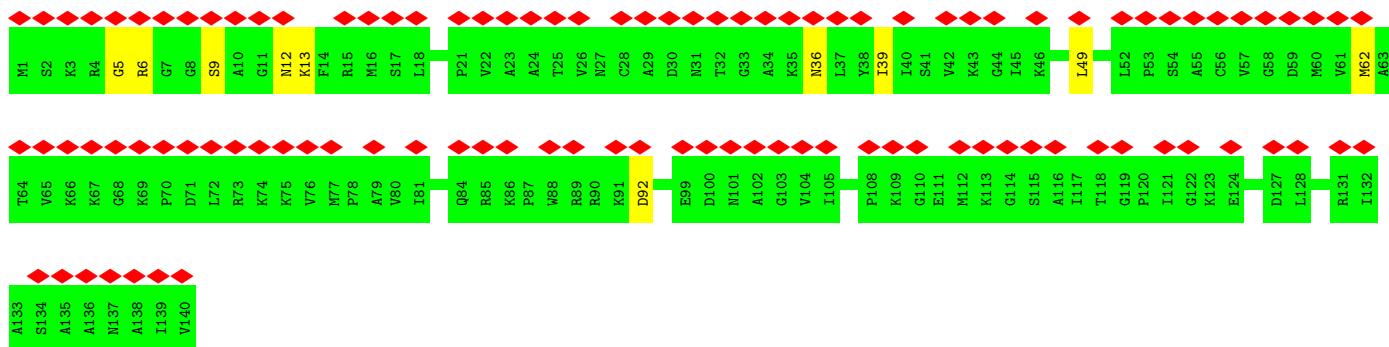


• Molecule 43: 60S ribosomal protein L6

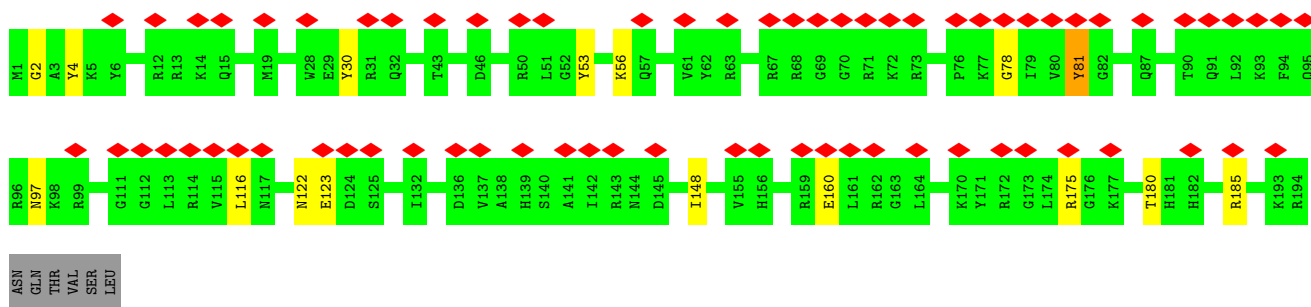
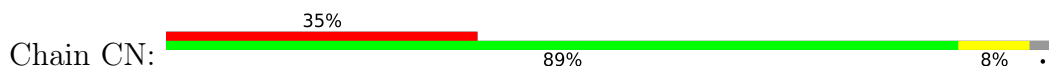


• Molecule 44: 60S ribosomal protein L14

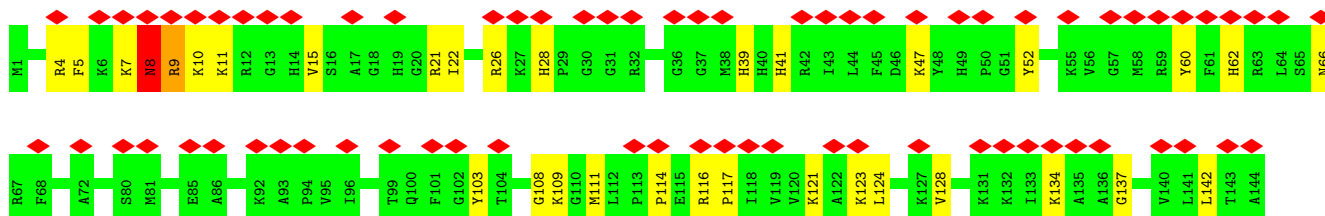
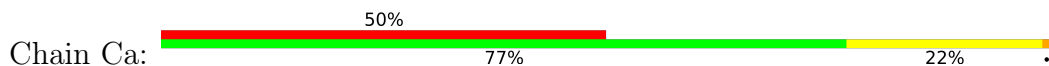




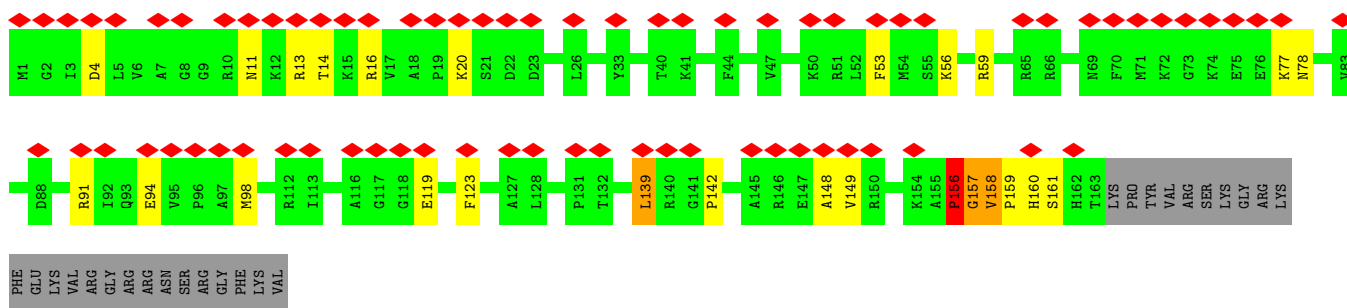
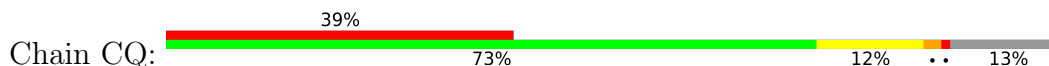
• Molecule 45: 60S ribosomal protein L15E



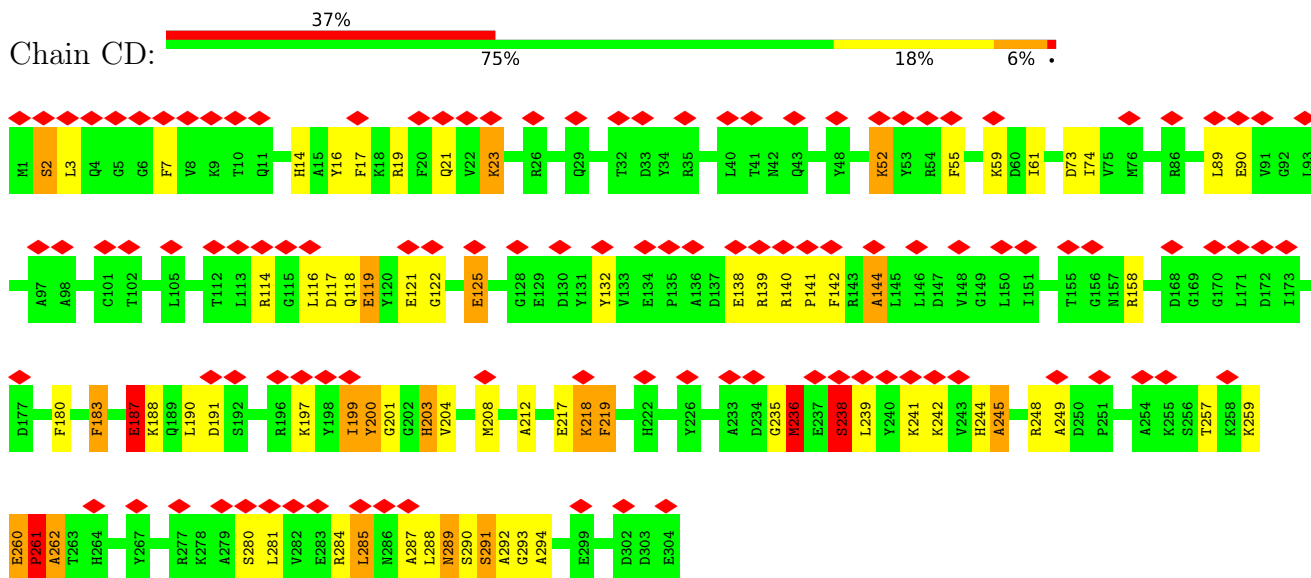
• Molecule 46: 60S ribosomal protein L15



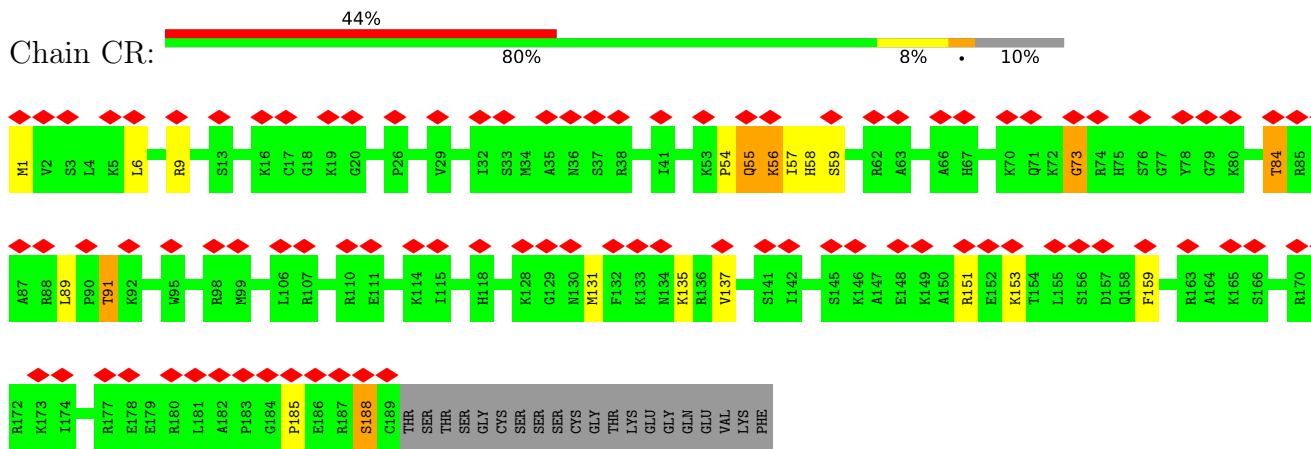
• Molecule 47: 60S ribosomal protein L18E



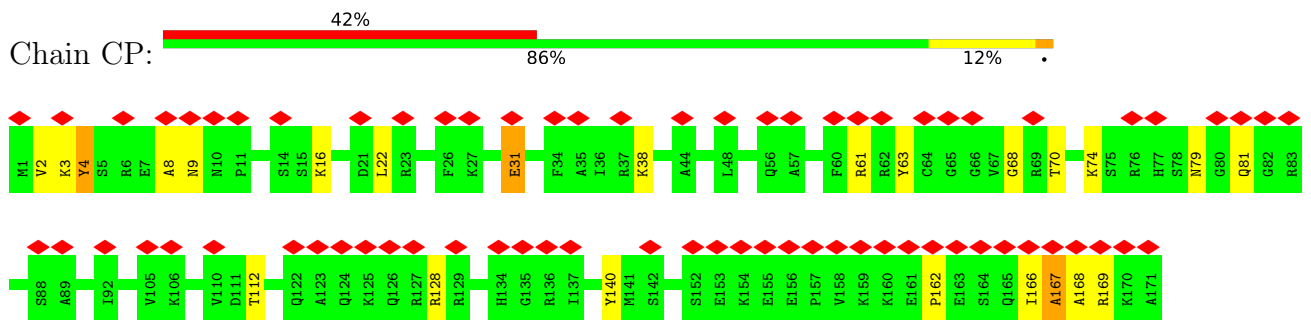
• Molecule 48: 60S ribosomal protein L18



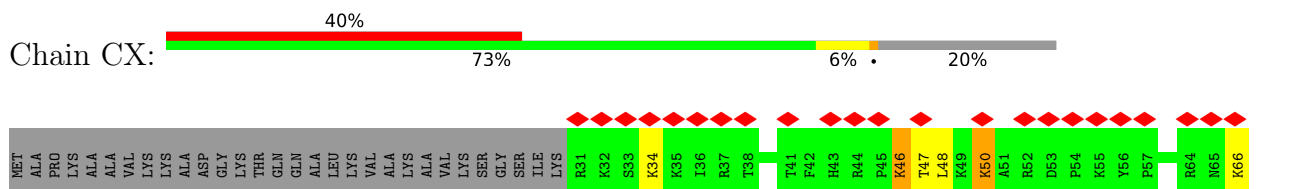
• Molecule 49: 60S ribosomal protein L19E

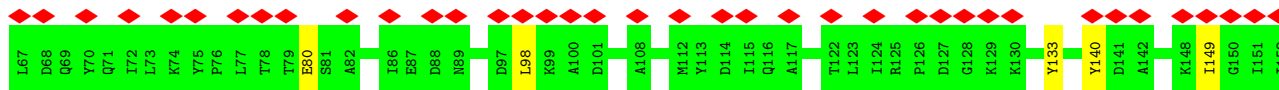


• Molecule 50: 60S ribosomal protein L22

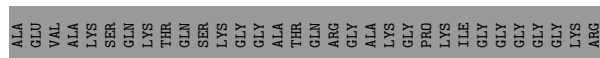
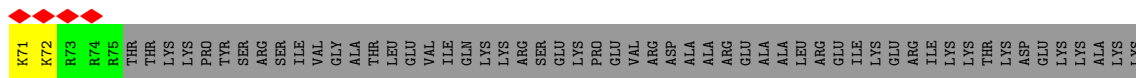
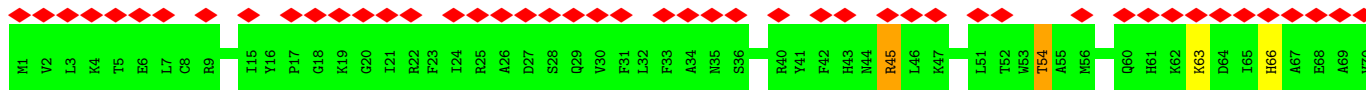


• Molecule 51: 60S ribosomal protein L23

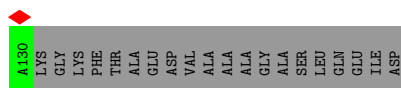
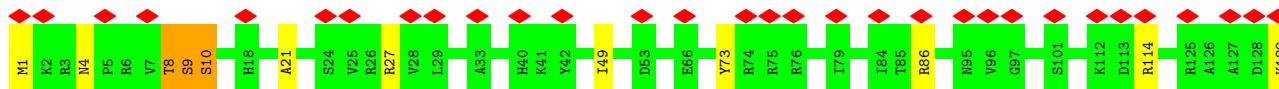
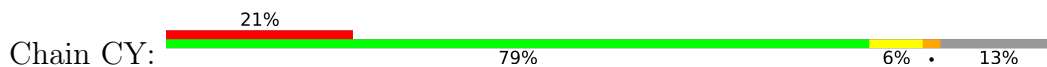




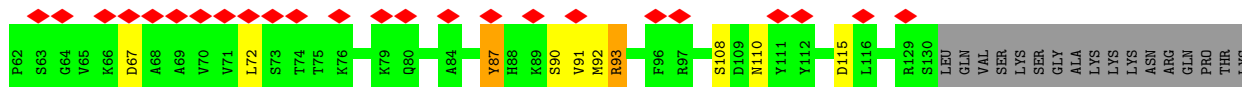
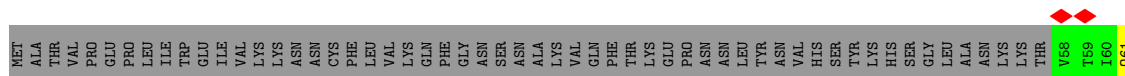
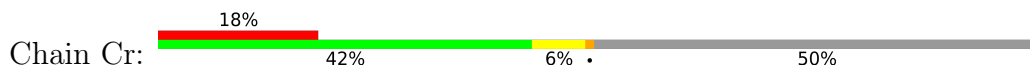
• Molecule 52: 60S ribosomal protein L24E



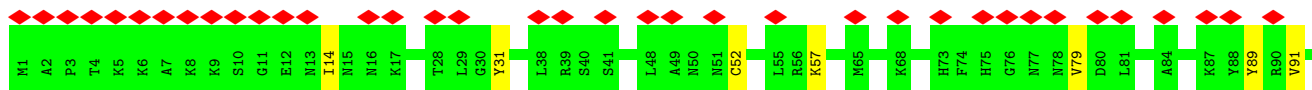
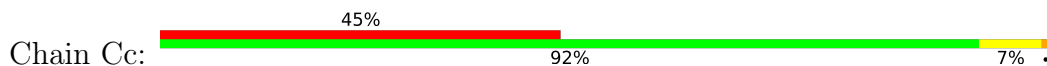
• Molecule 53: 60S ribosomal protein L24

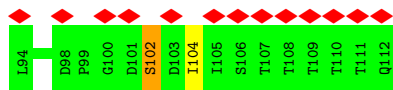


• Molecule 54: 60S ribosomal protein L28E

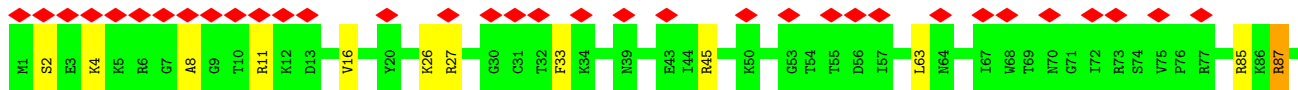
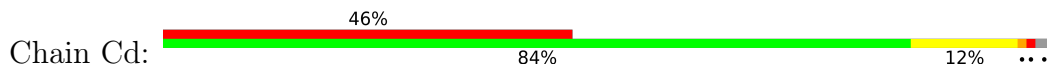


• Molecule 55: 60S ribosomal protein L30E

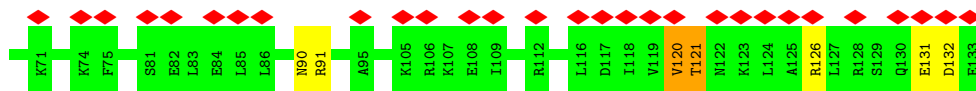
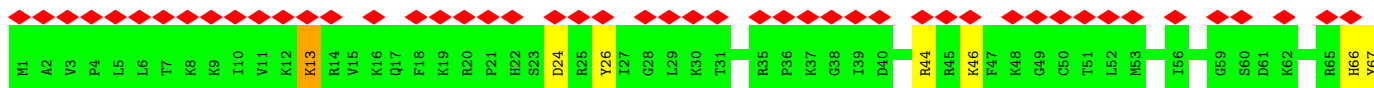
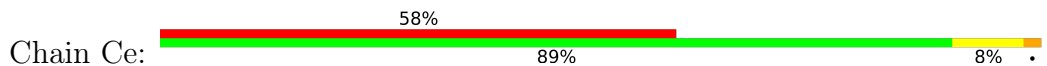




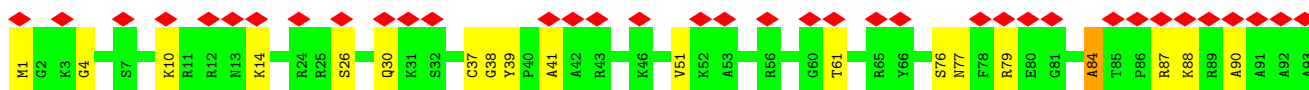
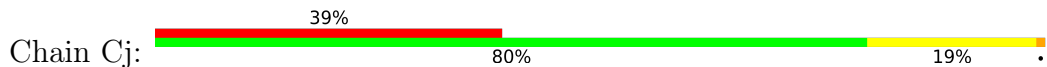
- Molecule 56: 60S ribosomal protein L31E



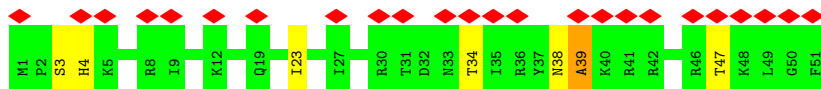
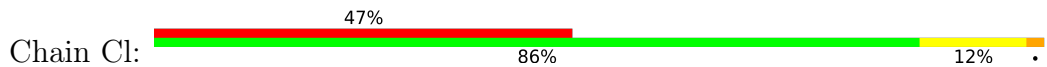
- Molecule 57: 60S ribosomal protein L32E



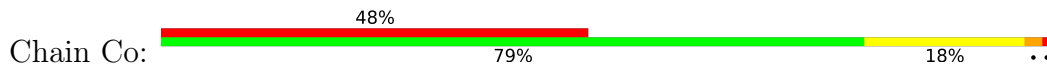
- Molecule 58: 60S ribosomal protein L37E

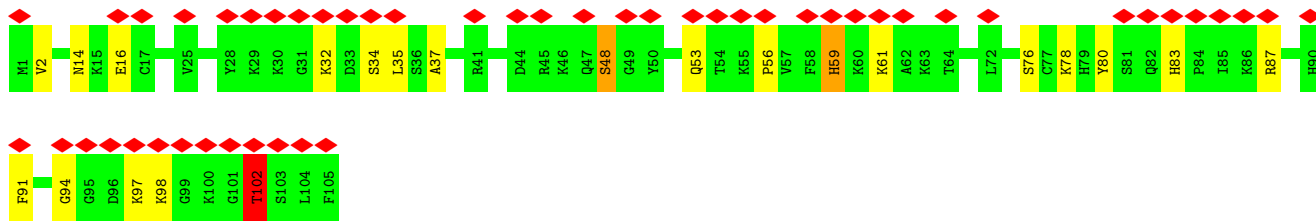


- Molecule 59: 60S ribosomal protein L39E

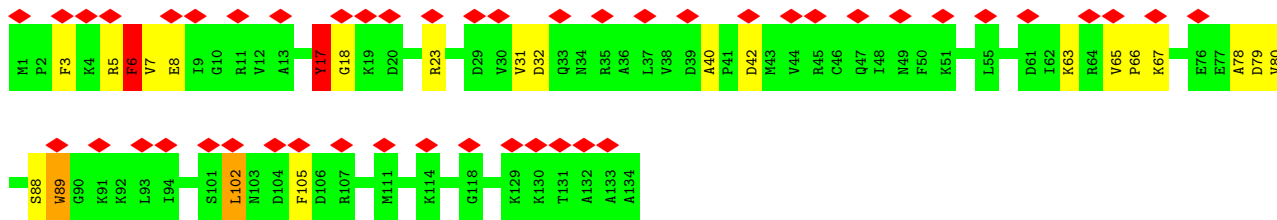
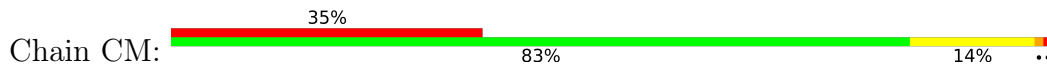


- Molecule 60: 60S ribosomal protein L44E

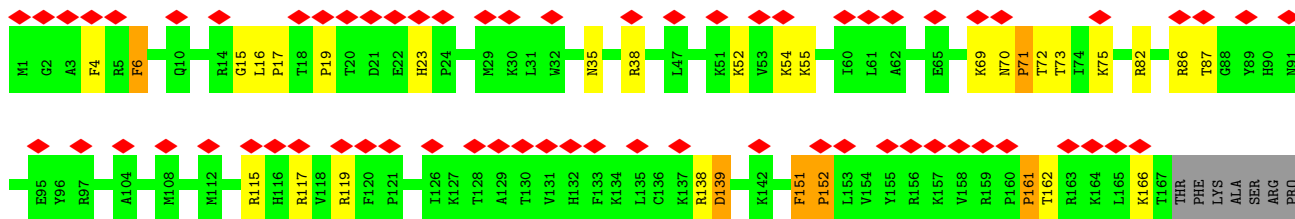
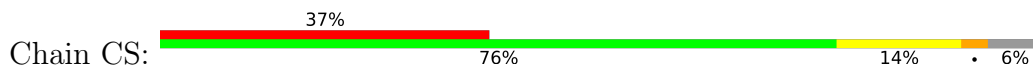




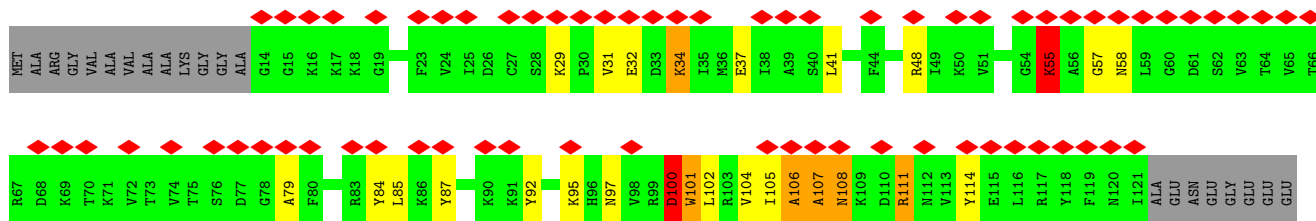
• Molecule 61: 60S ribosomal protein L14E



• Molecule 62: 60S ribosomal protein L20

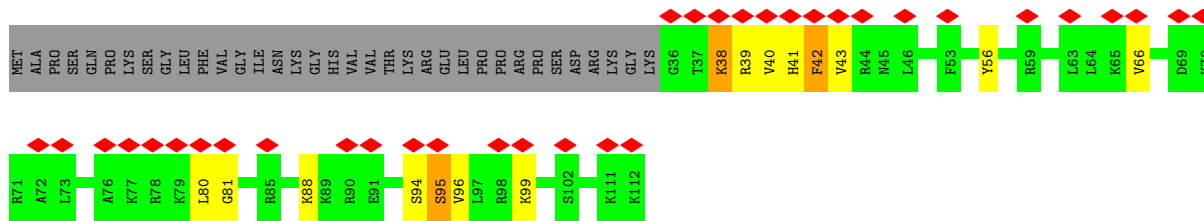


• Molecule 63: 60S ribosomal protein L22E

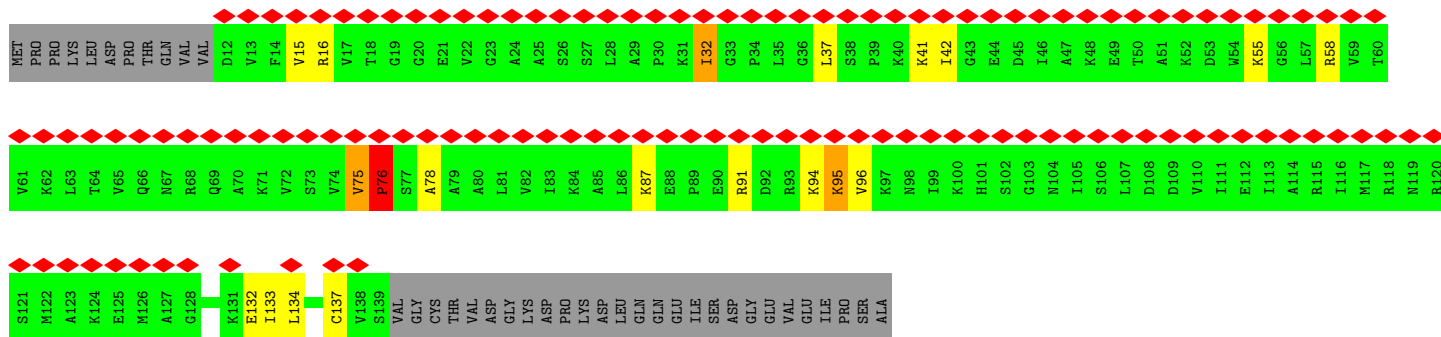
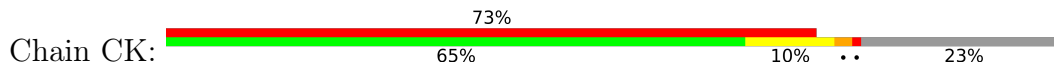


• Molecule 64: 60S ribosomal protein L36E

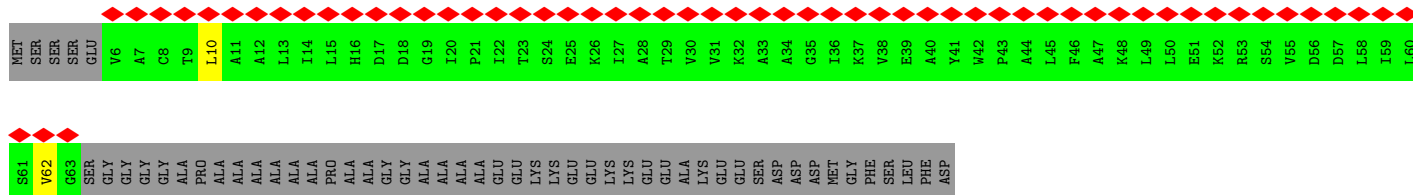




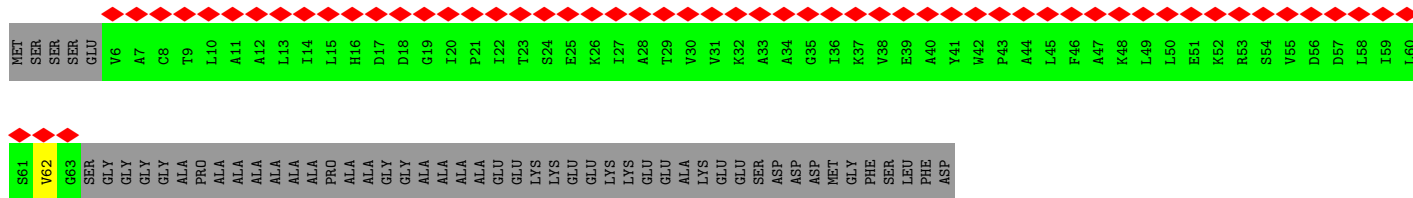
• Molecule 65: 60S ribosomal protein L11



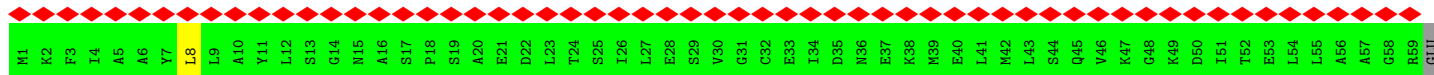
• Molecule 66: 60S ribosomal protein P1



• Molecule 66: 60S ribosomal protein P1

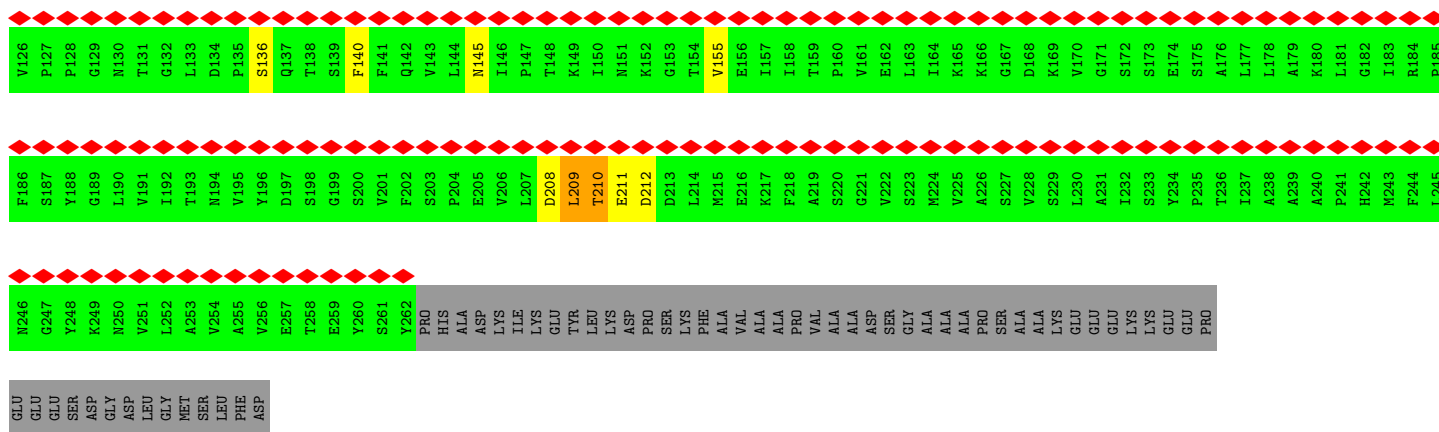


• Molecule 67: Acidic ribosomal protein P2

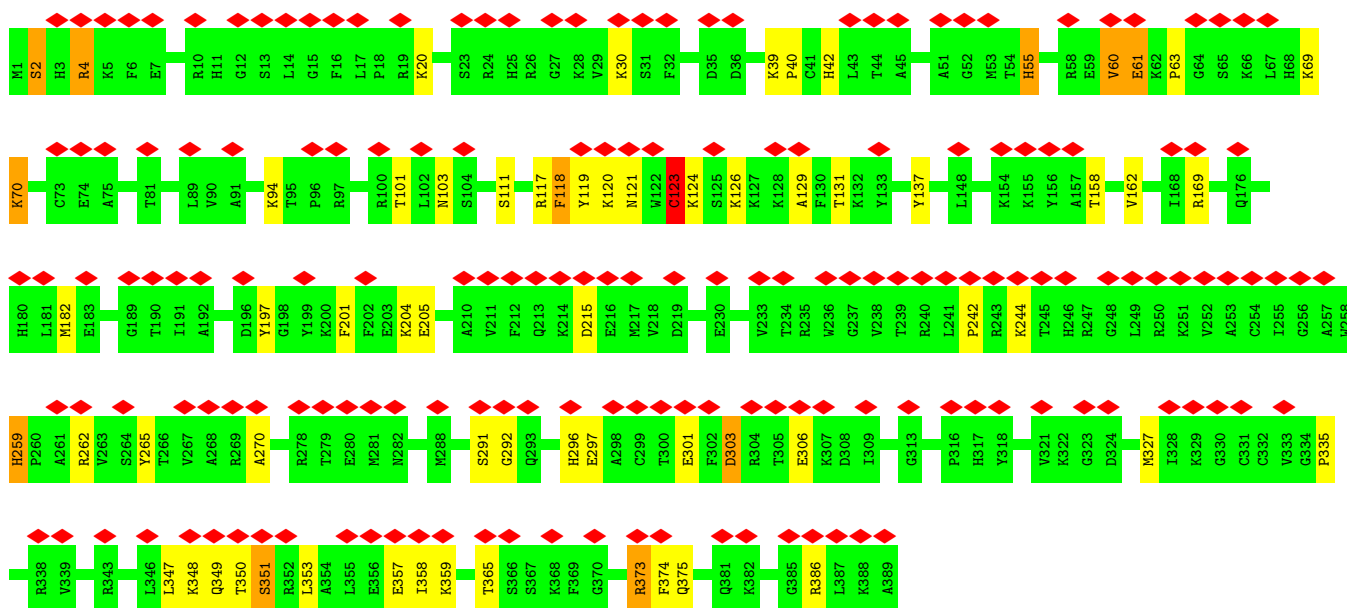
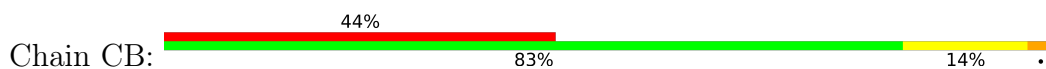




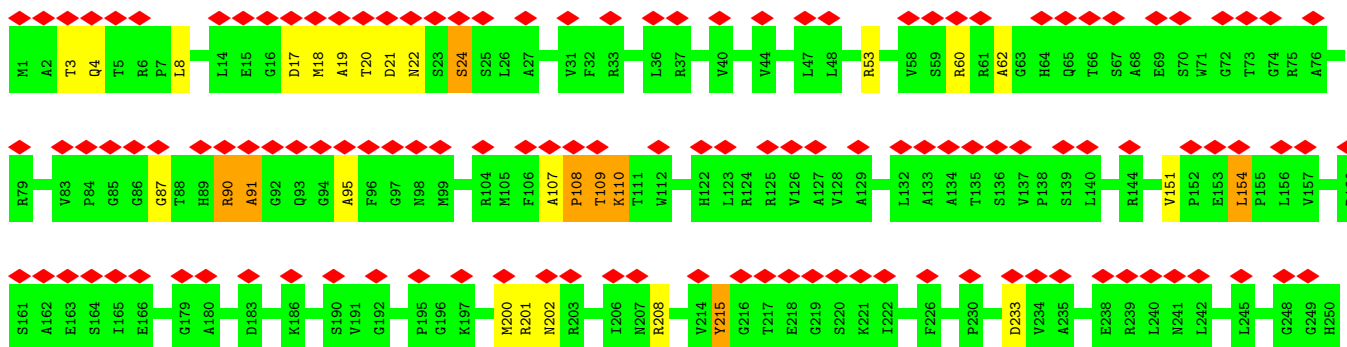
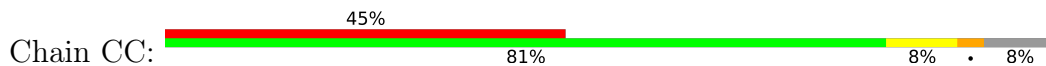


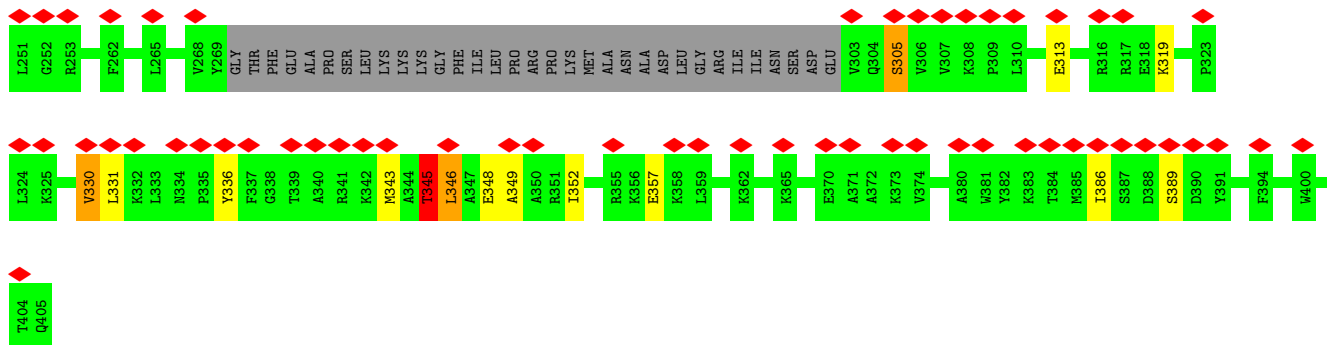


• Molecule 71: 60S ribosomal protein L3

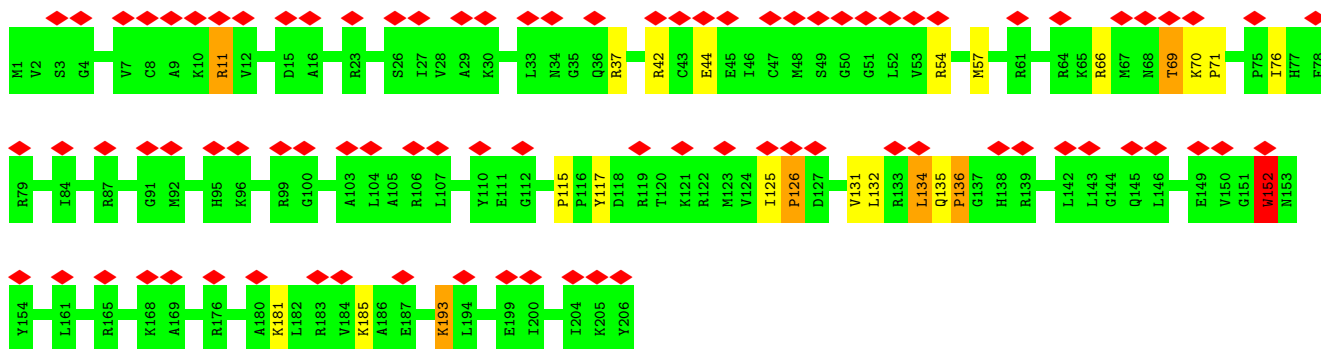


• Molecule 72: 60S ribosomal protein L4

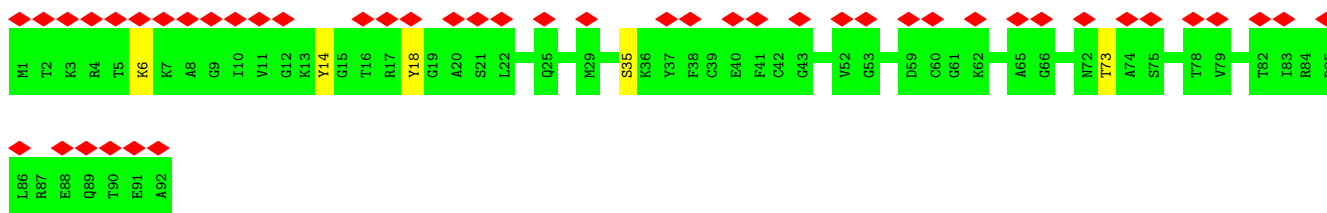




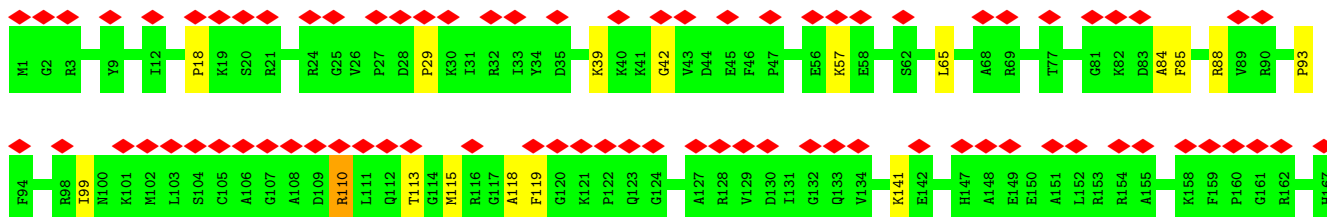
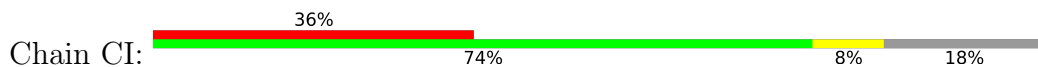
• Molecule 73: 60S ribosomal protein L13

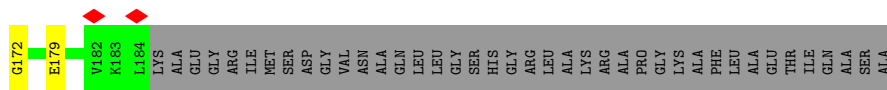


• Molecule 74: 60S ribosomal protein L43E

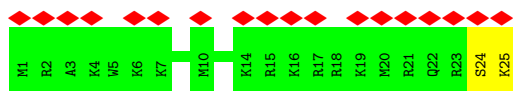
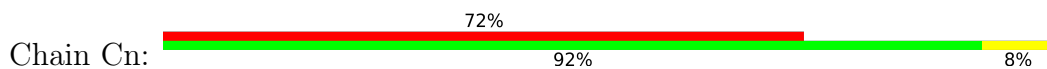


• Molecule 75: 60S ribosomal protein L16

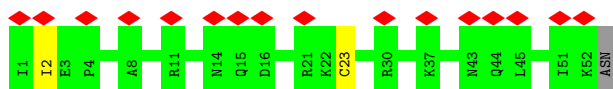




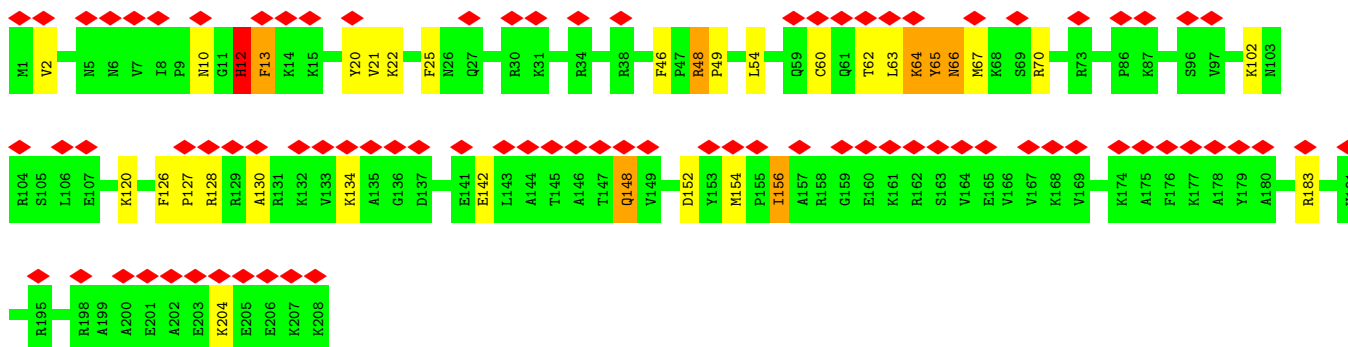
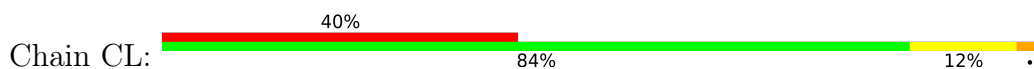
• Molecule 76: 60S ribosomal protein L41E



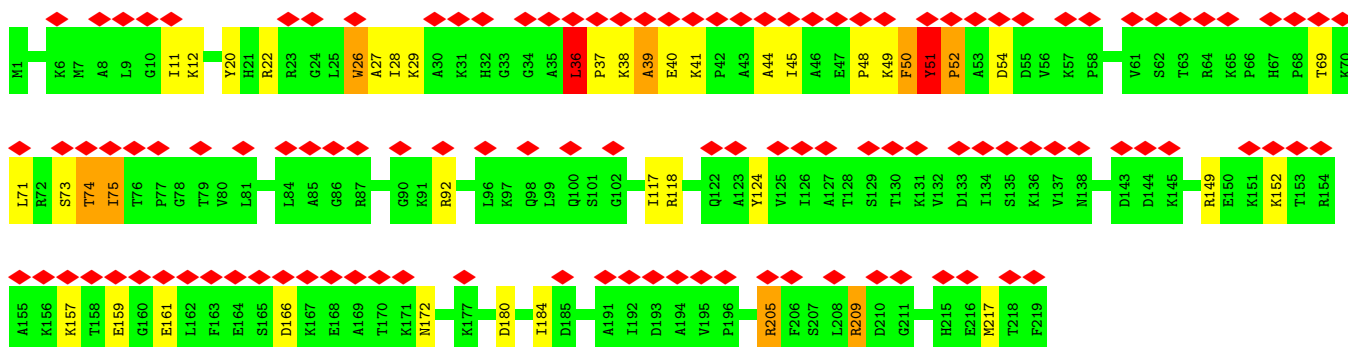
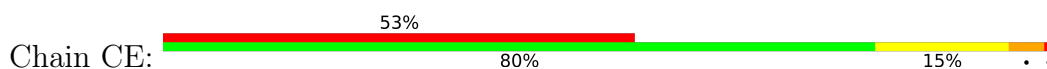
• Molecule 77: 60S ribosomal protein L40E



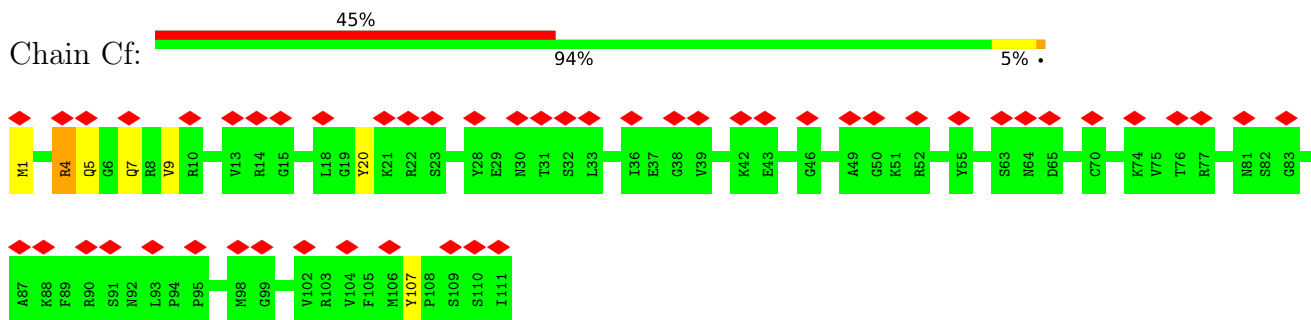
• Molecule 78: 60S ribosomal protein L13E



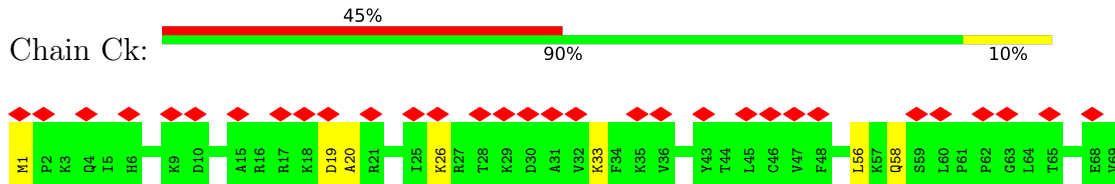
• Molecule 79: 60S ribosomal protein L6E



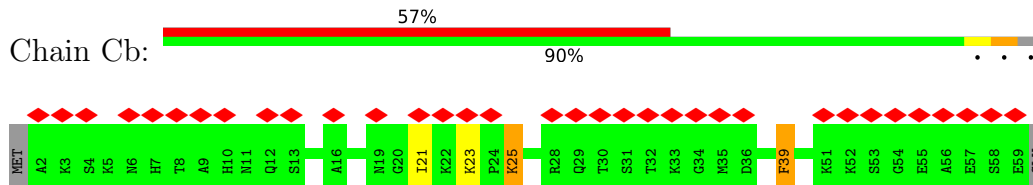
• Molecule 80: 60S ribosomal protein L33E



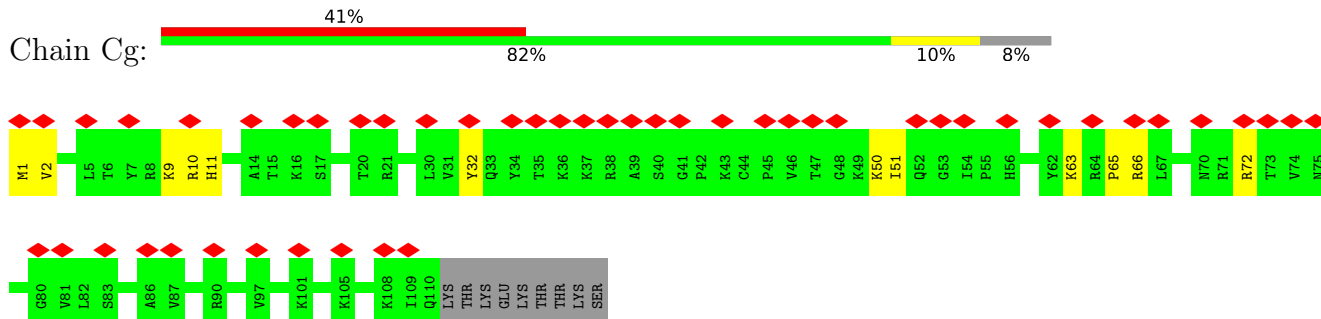
• Molecule 81: 60S ribosomal protein L38E



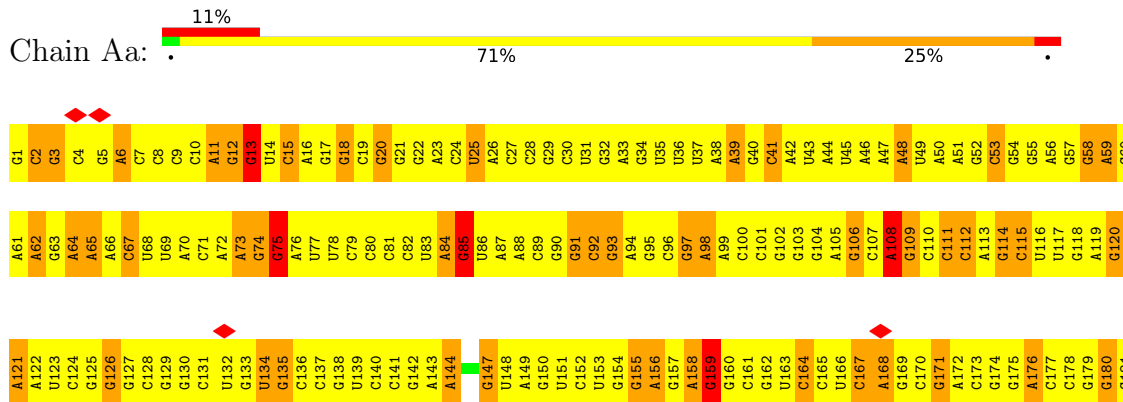
• Molecule 82: 60S ribosomal protein L29E



• Molecule 83: 60S ribosomal protein L34E



• Molecule 84: 60S ribosomal RNA



A985	A986	A987	A988	A989	A990	A991	A992	A993	A994	A995	A996	A997	A998	A999	A1000	A1001	A1002	A1003	A1004	A1005	A1006	A1007	A1008	A1009	A1010	A1011	A1012	A1013	A1014	A1015	A1016	A1017	A1018	A1019	A1020	A1021	A1022	A1023	A1024																					
G905	G906	G907	G908	G909	G910	G911	G912	G913	G914	G915	G916	G917	G918	G919	G920	G921	G922	G923	G924	G925	G926	G927	G928	G929	G930	G931	G932	G933	G934	G935	G936	G937	G938	G939	G940	G941	G942	G943	G944	G945	G946																			
G845	G846	G847	G848	G849	G850	G851	G852	G853	G854	G855	G856	G857	G858	G859	G860	G861	G862	G863	G864	G865	G866	G867	G868	G869	G870	G871	G872	G873	G874	G875	G876	G877	G878	G879	G880	G881	G882	G883	G884	G885	G886	G887	G888	G889	G890	G891	G892	G893	G894	G895	G896	G897	G898	G899	G900	G901	G902	G903	G904	
U785	U786	U787	U788	U789	U790	U791	U792	U793	U794	U795	U796	U797	U798	U799	U800	U801	U802	U803	U804	U805	U806	U807	U808	U809	U810	U811	U812	U813	U814	U815	U816	U817	U818	U819	U820	U821	U822	U823	U824	U825	U826	U827	U828	U829	U830	U831	U832	U833	U834	U835	U836	U837	U838	U839	U840	U841	U842	U843	U844	
G725	G726	G727	G728	G729	G730	G731	G732	G733	G734	G735	G736	G737	G738	G739	G740	G741	G742	G743	G744	G745	G746	G747	G748	G749	G750	G751	G752	G753	G754	G755	G756	G757	G758	G759	G760	G761	G762	G763	G764	G765	G766	G767	G768	G769	G770	G771	G772	G773	G774	G775	G776	G777	G778	G779	G780	G781	G782	G783	G784	
G665	G666	G667	G668	G669	G670	G671	G672	G673	G674	G675	G676	G677	G678	G679	G680	G681	G682	G683	G684	G685	G686	G687	G688	G689	G690	G691	G692	G693	G694	G695	G696	G697	G698	G699	G700	G701	G702	G703	G704	G705	G706	G707	G708	G709	G710	G711	G712	G713	G714	G715	G716	G717	G718	G719	G720	G721	G722	G723	G724	
C604	C605	C606	C607	C608	C609	C610	C611	C612	C613	C614	C615	C616	C617	C618	C619	C620	C621	C622	C623	C624	C625	C626	C627	C628	C629	C630	C631	C632	C633	C634	C635	C636	C637	C638	C639	C640	C641	C642	C643	C644	C645	C646	C647	C648	C649	C650	C651	C652	C653	C654	C655	C656	C657	C658	C659	C660	C661	C662	C663	C664
C544	C545	C546	C547	C548	C549	C550	C551	C552	C553	C554	C555	C556	C557	C558	C559	C560	C561	C562	C563	C564	C565	C566	C567	C568	C569	C570	C571	C572	C573	C574	C575	C576	C577	C578	C579	C580	C581	C582	C583	C584	C585	C586	C587	C588	C589	C590	C591	C592	C593	C594	C595	C596	C597	C598	C599	C600	C601	C602	C603	
G485	G486	G487	G488	G489	G490	G491	G492	G493	G494	G495	G496	G497	G498	G499	G500	G501	G502	G503	G504	G505	G506	G507	G508	G509	G510	G511	G512	G513	G514	G515	G516	G517	G518	G519	G520	G521	G522	G523	G524	G525	G526	G527	G528	G529	G530	G531	G532	G533	G534	G535	G536	G537	G538	G539	G540	G541	G542	G543		
G422	G423	G424	G425	G426	G427	G428	G429	G430	G431	G432	G433	G434	G435	G436	G437	G438	G439	G440	G441	G442	G443	G444	G445	G446	G447	G448	G449	G450	G451	G452	G453	G454	G455	G456	G457	G458	G459	G460	G461	G462	G463	G464	G465	G466	G467	G468	G469	G470	G471	G472	G473	G474	G475	G476	G477	G478	G479	G480	G481	G482
G362	G363	G364	G365	G366	G367	G368	G369	G370	G371	G372	G373	G374	G375	G376	G377	G378	G379	G380	G381	G382	G383	G384	G385	G386	G387	G388	G389	G390	G391	G392	G393	G394	G395	G396	G397	G398	G399	G400	G401	G402	G403	G404	G405	G406	G407	G408	G409	G410	G411	G412	G413	G414	G415	G416	G417	G418	G419	G420	G421	
U303	U304	U305	U306	U307	U308	U309	U310	U311	U312	U313	U314	U315	U316	U317	U318	U319	U320	U321	U322	U323	U324	U325	U326	U327	U328	U329	U330	U331	U332	U333	U334	U335	U336	U337	U338	U339	U340	U341	U342	U343	U344	U345	U346	U347	U348	U349	U350	U351	U352	U353	U354	U355	U356	U357	U358	U359	U360	U361		
G302	G303	G304	G305	G306	G307	G308	G309	G310	G311	G312	G313	G314	G315	G316	G317	G318	G319	G320	G321	G322	G323	G324	G325	G326	G327	G328	G329	G330	G331	G332	G333	G334	G335	G336	G337	G338	G339	G340	G341	G342	G343	G344	G345	G346	G347	G348	G349	G350	G351	G352	G353	G354	G355	G356	G357	G358	G359	G360	G361	
U242	U243	U244	U245	U246	U247	U248	U249	U250	U251	U252	U253	U254	U255	U256	U257	U258	U259	U260	U261	U262	U263	U264	U265	U266	U267	U268	U269	U270	U271	U272	U273	U274	U275	U276	U277	U278	U279	U280	U281	U282	U283	U284	U285	U286	U287	U288	U289	U290	U291	U292	U293	U294	U295	U296	U297	U298	U299	U300	U301	
C182	C183	C184	C185	C186	C187	C188	C189	C190	C191	C192	C193	C194	C195	C196	C197	C198	C199	C200	C201	C202	C203	C204	C205	C206	C207	C208	C209	C210	C211	C212	C213	C214	C215	C216	C217	C218	C219	C220	C221	C222	C223	C224	C225	C226	C227	C228	C229	C230	C231	C232	C233	C234	C235	C236	C237	C238	C239	C240	C241	

A1747	G1025	G1085	G1145	C1205	G1265	G1325	G1385	U1445	A1506	G1567	U1627	G1687	G1747
A1748	A1026	U1086	A1146	A1206	G1266	C1326	G1386	G1446	A1507	A1568	G1628	U1688	G1748
A1749	C1027	U1087	U1147	A1207	A1267	G1327	G1387	G1447	C1508	U1569	G1629	G1689	A1749
G1751	G1028	A1088	A1148	U1208	G1268	G1328	C1388	U1448	C1509	C1570	A1629	A1689	G1751
C1752	C1029	G1089	G1149	G1209	U1269	G1329	C1389	A1449	G1510	G1571	G1631	U1691	C1752
A1753	A1030	C1090	C1150	G1210	G1270	A1330	G1390	G1450	C1511	C1572	G1632	A1693	A1753
C1754	A1031	C1091	G1151	G1211	U1271	C1331	A1391	U1451	A1512	G1573	G1633	A1694	C1754
C1756	C1032	U1092	G1152	U1212	G1272	C1332	U1392	A1452	C1513	C1574	G1634	C1696	C1756
C1757	G1033	G1094	A1153	U1213	U1273	C1333	G1393	G1453	U1514	G1575	A1636	C1697	C1757
U1758	U1034	C1095	U1154	A1214	A1274	A1334	A1394	C1454	U1515	C1576	G1637	C1698	U1758
C1759	G1035	C1096	A1155	U1215	A1275	C1335	A1395	A1455	U1516	A1577	U1638	C1699	C1759
G1760	C1036	A1097	A1156	G1216	C1276	A1336	A1396	A1456	C1517	U1578	U1639	U1700	G1760
C1761	U1088	U1098	C1157	G1217	A1277	C1337	A1397	A1457	A1518	C1579	A1640	U1701	C1761
G1762	U1037	C1099	C1158	U1218	A1278	C1338	A1398	U1458	C1519	C1580	G1641	G1702	G1762
G1764	U1038	G1100	C1159	C1219	A1279	C1339	C1399	A1459	A1520	U1581	G1642	G1703	G1764
G1765	G1039	G1101	C1160	G1220	C1279	G1340	C1400	U1460	U1521	C1582	A1643	A1704	G1765
U1766	A1040	A1101	G1161	A1221	U1280	G1341	G1401	U1461	U1522	G1583	A1644	A1705	U1766
U1768	C1041	U1102	A1162	U1222	U1281	C1342	G1402	C1462	G1523	A1584	A1645	C1706	U1768
C1769	C1042	U1103	A1163	U1223	A1282	C1343	G1403	G1463	G1524	U1585	A1646	G1707	C1769
C1770	U1043	C1104	C1164	A1224	C1283	A1344	G1404	A1464	G1525	A1586	G1647	U1708	C1770
G1771	U1044	G1105	C1165	A1225	U1284	U1345	G1406	U1466	A1526	G1587	C1648	U1709	G1771
G1772	U1045	G1106	C1166	G1226	U1285	U1346	G1407	U1467	A1527	A1588	G1649	U1710	G1772
C1773	U1046	U1107	C1167	G1227	G1286	U1347	G1408	A1468	G1528	C1589	G1650	G1711	C1773
C1775	U1047	G1108	G1168	C1228	G1287	G1348	G1409	U1469	C1529	A1590	A1651	G1712	C1775
C1776	U1048	U1109	C1169	A1229	C1288	G1349	G1410	A1470	G1531	U1591	G1652	A1713	C1776
C1777	C1110	C1110	U1170	G1230	G1289	G1350	A1411	A1471	A1532	C1592	G1653	A1714	C1777
C1778	U1050	U1111	U1171	A1231	A1290	C1351	G1412	A1472	U1533	C1593	C1654	G1715	C1778
C1779	A1051	C1112	C1173	A1232	A1291	G1352	G1413	U1473	U1534	G1594	G1655	G1716	C1779
C1780	G1053	G1113	G1174	G1233	U1292	A1353	C1414	U1474	C1535	U1595	C1656	G1717	C1780
C1781	U1054	A1114	C1175	G1234	C1293	G1354	G1415	U1475	C1536	U1596	C1657	G1718	C1781
C1782	U1055	A1115	U1176	A1235	A1294	U1355	G1416	U1476	U1537	U1597	G1658	U1719	C1782
C1783	U1056	U1116	G1177	C1236	A1295	G1356	G1417	A1477	A1538	A1598	G1659	C1720	C1783
C1784	U1057	U1117	C1178	G1237	G1296	C1357	C1418	A1478	G1539	G1599	G1660	A1721	C1784
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C1786	A1059	U1119	C1180	U1239	A1298	C1359	G1420	G1480	G1541	G1601	G1662	G1723	C1786
C1787	U1060	C1120	A1181	G1240	G1299	U1360	A1421	G1481	A1542	U1602	G1663	G1724	C1787
C1788	A1061	C1121	C1182	G1241	C1300	G1361	G1422	G1482	G1543	U1603	G1664	G1725	C1788
U1791	G1062	A1123	C1183	U1242	C1301	C1362	A1423	G1483	G1544	U1604	C1665	G1726	U1791
C1792	G1063	U1124	G1184	C1243	C1302	C1363	G1424	A1484	G1545	U1605	C1666	G1727	C1792
A1793	U1064	U1125	C1185	A1244	C1303	C1364	G1425	A1485	G1546	C1606	C1667	G1728	A1793
A1795	A1065	U1126	C1186	U1245	G1304	C1365	G1426	G1486	G1547	C1607	U1668	G1729	A1795
U1797	G1066	U1127	C1187	U1246	A1305	G1366	G1427	A1487	U1548	U1608	U1669	U1730	U1797
C1798	G1067	U1128	G1188	G1247	A1306	G1367	G1428	G1488	A1549	G1609	G1670	G1731	C1798
G1800	A1068	U1129	C1189	G1248	A1307	U1368	U1429	G1489	A1550	C1611	G1671	G1732	G1800
A1801	U1069	G1130	C1190	A1249	U1308	G1369	C1430	A1490	C1551	G1612	U1672	A1733	A1801
A1802	G1070	U1131	U1191	G1250	G1310	A1370	G1431	A1491	C1552	C1613	A1673	G1734	A1802
G1804	A1071	A1132	A1192	U1251	G1311	U1371	G1432	A1492	C1553	G1614	A1674	U1735	G1804
A1805	C1072	A1133	A1193	U1252	G1312	U1372	G1433	A1493	C1554	G1615	A1675	C1736	A1805
C1082	G1073	A1134	C1194	G1253	U1313	A1373	G1434	A1494	C1555	G1616	G1677	C1737	C1082
C1083	U1074	C1135	C1195	A1254	G1314	G1374	G1435	A1495	G1556	U1617	U1678	U1738	C1083
C1075	G1075	U1136	U1196	A1255	G1315	G1375	A1436	A1496	G1557	U1618	U1679	G1739	C1075
C1076	G1076	A1137	U1197	A1256	C1316	A1376	G1437	C1497	U1558	U1619	U1680	G1740	C1076
U1078	C1077	U1138	G1198	U1257	G1317	G1377	A1438	C1498	A1559	G1620	U1681	U1741	U1078
G1079	U1078	A1139	U1199	U1258	C1318	U1378	U1439	A1501	U1561	U1621	G1682	G1742	G1079
C1080	G1079	C1140	A1200	C1259	U1319	G1379	C1440	U1502	U1562	G1622	U1683	G1743	C1080
U1081	C1080	U1141	C1201	G1260	G1320	C1380	U1441	G1503	G1563	C1623	U1684	C1744	U1081
C1082	U1081	G1142	C1202	G1261	A1321	G1381	U1442	U1504	G1564	G1624	U1685	G1745	C1082
C1083	U1082	C1143	C1203	C1261	A1322	G1382	G1443	U1505	C1565	G1625	U1686	G1746	C1083
G1084	C1083	C1144	A1204	U1262	C1323	G1383	G1444	G1505	C1566	U1626	U1686		G1084
				A1263		G1384							
				A1264									

G2527	U2528	A2467	U2407	A2347	U2287	G2167	A2107	A2047	C1987	A1927	U1867	C1807
U2528	C2529	G2468	G2408	U2348	C2288	C2168	C2108	C2048	G1988	A1928	C1868	G1808
C2529	C2469	C2469	U2409	C2349	U2289	U2169	G2109	C2049	G1989	A1929	U1869	A1809
G2530	G2470	C2470	U2410	C2350	A2290	G2170	G2110	G2050	U1990	G1930	G1870	G1810
G2531	C2471	C2471	G2411	G2351	A2291	A2171	A2111	G2051	A1991	A1931	C1871	U1811
A2532	U2472	C2472	G2412	G2352	U2292	C2172	C2112	G2052	U1992	U1932	C1872	A1812
G2533	C2473	C2473	G2413	C2353	U2293	G2173	G2113	A2053	U1993	U1933	C1873	C1813
G2534	A2474	C2474	C2414	G2354	U2294	C2174	G2114	A2054	G1993	G1934	A1874	G1814
C2535	A2475	C2475	U2415	G2355	G2295	U2175	G2115	A2055	U1994	G1935	U1875	G1815
G2536	G2476	A2476	G2416	A2356	U2296	A2176	G2116	A2056	C1994	G1936	U1876	U1816
G2537	G2477	C2477	U2417	U2357	U2297	U2177	G2117	U2056	U1995	U1937	G1877	A1817
G2538	G2478	C2478	A2418	U2358	A2298	G2178	G2118	C2056	C1996	U1938	G1878	C1818
G2539	U2479	C2479	G2419	C2359	U2299	U2179	A2119	G2057	U1997	U1940	A1879	A1819
C2540	G2479	C2479	U2420	A2360	G2300	G2180	G2120	C2058	A1998	U1941	A1880	C1820
A2541	U2542	G2480	C2421	C2361	C2301	U2181	U2121	C2059	A1982	A1942	C1881	G1821
U2542	G2481	G2481	U2422	G2362	G2302	G2182	C2122	C2060	G1999	G1943	A1882	C1822
G2543	A2482	C2482	G2423	C2363	C2303	A2183	C2123	C2061	C2000	G1944	A1883	C1823
C2544	A2483	C2483	U2424	G2364	G2304	U2184	G2124	C2062	G2001	A1945	A1884	C1824
C2545	A2484	C2484	C2425	C2365	U2305	U2185	A2125	U2062	U2002	C1946	G1885	G1825
C2546	G2484	A2484	A2426	A2366	G2306	U2186	C2126	U2063	G2003	U1947	U1886	G1826
C2547	U2484	G2484	G2427	A2367	A2307	C2187	U2127	U2064	C2004	G1948	G1887	U1827
U2548	U2485	G2485	G2428	G2368	U2308	U2188	U2128	U2065	U2005	G1949	G1888	G1828
C2549	G2486	G2486	C2429	U2369	U2309	U2189	U2129	C2066	U2006	G1950	G1889	G1829
C2550	A2487	C2487	U2430	G2370	U2310	C2190	U2130	G2067	C2005	G1951	C1890	U1830
U2551	A2488	C2488	A2431	A2371	A2251	G2191	U2131	G2068	A2006	C1952	A1891	A1831
U2552	A2489	C2489	U2432	A2372	C2252	U2192	A2132	U2072	G2007	U1952	A1892	C1832
U2553	A2490	C2490	U2433	C2373	U2253	C2193	A2133	G2073	C2008	G1953	G1893	U1833
U2554	A2491	C2491	G2434	G2374	A2254	U2194	U2134	G2074	G2009	G1954	G1894	C1834
U2555	A2492	C2492	U2435	G2375	U2255	G2195	U2135	G2075	C2010	G1955	G1895	A1835
U2556	C2493	C2493	G2436	G2376	G2256	G2196	A2136	C2076	G2011	G1956	A1897	U1836
U2557	A2494	C2494	A2437	C2377	A2257	U2197	U2137	U2077	G2012	G1957	A1898	A1837
U2558	A2495	C2495	U2438	U2378	U2258	U2198	A2138	U2078	C2013	G1958	U1899	A1838
U2559	A2496	C2496	G2439	U2379	U2259	C2199	A2139	U2079	G2014	G1959	C1900	G1840
U2560	A2497	C2497	U2440	G2380	C2260	G2200	C2140	C2079	A2015	U1959	G1901	G1841
A2561	U2497	C2497	G2441	C2381	U2261	G2201	A2141	C2081	G2016	C1960	C1902	C1842
U2562	U2498	C2498	U2442	G2382	U2262	A2202	A2142	C2082	G2017	C1961	C1903	A1843
G2563	U2500	C2498	C2443	C2383	U2263	A2203	G2143	C2083	A2018	C1962	A1904	U1844
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G2565	U2502	C2498	U2445	A2385	A2265	G2205	C2145	C2085	G2020	G1964	A1906	A1846
G2566	U2503	C2498	G2446	A2386	A2266	U2206	U2146	C2086	G2021	G1965	A1907	G1847
G2567	A2504	C2498	A2447	A2387	A2267	C2207	U2147	C2087	G2022	C1966	C1908	G1848
G2568	A2505	C2498	U2448	C2388	U2268	A2208	U2148	C2088	C2023	C1967	G1909	U1849
G2569	A2506	C2498	G2449	U2389	U2269	G2149	G2149	C2089	G2024	C1968	G1910	C1850
G2570	U2507	C2498	C2450	C2390	A2270	G2150	C2150	C2090	G2025	C1969	A1911	U1851
G2571	U2508	C2498	G2451	C2391	G2271	G2151	G2151	C2091	C2026	C1970	U1912	G1852
U2572	U2509	C2498	U2452	C2392	C2272	U2152	U2152	C2092	G2027	C1971	U1913	C1853
U2573	U2510	C2498	G2453	C2393	C2273	G2153	G2153	C2093	C2028	C1972	C1913	C1854
A2574	U2511	C2498	U2454	C2394	A2274	A2154	G2154	C2094	G2029	C1973	C1914	A1854
C2575	U2512	C2498	A2455	G2395	A2275	A2155	G2155	C2095	U2030	C1974	C1915	A1855
C2576	U2513	C2498	G2456	C2396	U2276	C2156	U2156	C2096	G2031	C1975	G1916	G1856
C2577	A2514	C2498	A2457	U2397	U2277	C2157	C2157	C2097	G2032	A1970	U1917	G1857
C2578	U2515	C2498	G2458	U2398	C2278	A2217	U2158	C2098	C2033	A1971	A1918	U1858
C2579	U2516	C2498	U2459	C2399	C2279	A2218	U2159	C2099	G2034	C1972	A1919	U1859
C2580	U2517	C2498	A2460	U2400	U2280	A2219	C2160	C2099	G2035	C1973	A1860	A1860
C2581	A2518	C2498	C2401	A2401	U2281	U2220	G2161	C2099	U2036	C1974	A1861	A1861
C2582	U2519	C2498	A2402	C2340	C2282	U2221	C2162	C2099	G2037	G1975	C1922	A1862
U2583	U2520	C2498	U2403	U2341	C2283	C2222	G2163	C2099	U2038	U1976	G1923	A1863
C2584	C2522	C2498	C2404	C2342	U2284	U2223	G2164	C2099	G2039	C1977	G1924	C1864
C2585	U2523	C2498	C2405	U2343	C2285	C2224	A2165	C2099	G2040	C1978	G1925	C1865
C2586	U2524	C2498	C2406	A2344	A2286	C2225	U2166	C2099	G2041	G1979	G1926	C1866
G2525	G2526	C2498	U2346	U2346	A2286	C2226		C2099	G2042	C1979	A1926	

C3367	C3368	C3369	U3370	C3371	C3372	C3373	C3374	C3375	C3376	C3377	C3378	C3379	C3380	C3381	C3382	C3383	C3384	C3385	C3386	U3387	C3388	C3389	C3390	U3391																																				
A3307	A3308	U3309	A3310	C3311	C3312	C3313	C3314	C3315	C3316	C3317	C3318	C3319	C3320	C3321	C3322	C3323	C3324	C3325	C3326	U3327	C3328	C3329	C3330	C3331	C3332	C3333	C3334	C3335	C3336	C3337	C3338	C3339	C3340	C3341	C3342	C3343	C3344	C3345	C3346	C3347	C3348	C3349	C3350	C3351	C3352	C3353	C3354	C3355	C3356	C3357	C3358	C3359	C3360	C3361	C3362	C3363	C3364	C3365	C3366	
C3247	C3248	C3249	C3250	C3251	C3252	C3253	C3254	C3255	C3256	C3257	C3258	C3259	C3260	C3261	C3262	C3263	C3264	C3265	C3266	U3267	C3268	C3269	C3270	C3271	C3272	C3273	C3274	C3275	C3276	C3277	C3278	C3279	C3280	C3281	C3282	C3283	C3284	C3285	C3286	C3287	C3288	C3289	C3290	C3291	C3292	C3293	C3294	C3295	C3296	C3297	C3298	C3299	C3300	C3301	C3302	C3303	C3304	C3305	A3306	
C3187	C3188	C3189	U3190	U3191	C3192	C3193	C3194	C3195	C3196	C3197	C3198	C3199	A3200	A3201	C3202	C3203	C3204	C3205	C3206	C3207	C3208	C3209	C3210	C3211	C3212	C3213	C3214	C3215	C3216	C3217	C3218	C3219	C3220	A3221	A3222	C3223	C3224	C3225	C3226	U3227	C3228	C3229	C3230	C3231	C3232	C3233	C3234	A3235	C3236	C3237	U3238	C3239	C3240	C3241	C3242	C3243	C3244	U3245		
C3127	A3128	C3129	A3130	U3131	C3132	C3133	C3134	A3135	C3136	C3137	C3138	C3139	A3140	C3141	C3142	C3143	U3144	C3145	C3146	C3147	C3148	C3149	C3150	C3151	C3152	U3153	C3154	C3155	C3156	C3157	C3158	C3159	C3160	C3161	C3162	C3163	C3164	C3165	C3166	C3167	C3168	C3169	C3170	C3171	C3172	A3173	C3174	C3175	C3176	A3177	C3178	C3179	U3180	U3181	A3182	C3183	C3184	C3185	C3186	
C3067	U3068	U3069	C3070	A3071	A3072	A3073	A3074	A3075	C3076	C3077	C3078	C3079	U3080	C3081	C3082	C3083	C3084	C3085	C3086	A3087	A3088	C3089	C3090	C3091	A3092	C3093	C3094	C3095	U3096	C3097	U3098	C3099	C3100	C3101	C3102	C3103	A3104	U3105	U3106	A3107	U3108	C3109	A3110	C3111	U3112	C3113	A3114	A3115	C3116	C3117	C3118	C3119	U3120	C3121	U3122	A3123	C3124	C3125	U3126	
G2947	A2948	G2949	C2950	U2951	C2952	C2953	G2954	U2955	U2956	U2957	A2958	C2959	A2960	C2961	C2962	C2963	U2964	C2965	C2966	U2967	C2968	C2969	C2970	A2971	C2972	A2973	C2974	C2975	U2976	U2977	A2978	C2979	U2980	U2981	U2982	U2983	A2984	C2985	C2986	C2987	C2988	C2989	U2990	U2991	C2992	C2993	C2994	U2995	U2996	C2997	A2998	C2999	U3000	G3001	C3002	U3003	C3004	C3005	U3006	C3006
G2827	U2828	U2829	G2830	U2831	C2832	C2833	C2834	A2835	U2836	C2837	C2838	A2839	A2840	G2841	C2842	C2843	U2844	U2845	C2846	A2847	U2848	A2849	C2850	C2851	C2852	A2853	C2854	C2855	C2856	U2857	C2858	C2859	U2860	U2861	U2862	U2863	U2864	C2865	A2866	U2867	C2868	C2869	U2870	U2871	C2872	C2873	A2874	U2875	U2876	U2877	C2878	C2879	G2880	C2881	C2882	C2883	U2884	U2885	C2886	
C2887	U2888	U2889	U2890	C2891	C2892	U2893	U2894	C2895	C2896	C2897	A2898	C2899	C2900	C2901	A3022	C2903	C2904	A2905	U2906	U2907	C2908	C2909	C2910	C2911	A2912	A2913	C2914	C2915	U2916	U2917	U2918	C2919	C2920	A2921	U2922	U2923	C2924	U2925	U2926	C2927	C2928	A2929	C2930	C2931	C2932	C2933	C2934	A2935	U2936	U2937	A2938	C2939	G2940	C2941	A2942	C2943	C2944	C2945	U2946	
G2767	C2768	U2769	C2770	U2771	A2772	C2773	A2774	C2775	U2776	U2777	C2778	G2779	C2780	A2781	C2782	U2783	U2784	U2785	U2786	A2787	C2788	C2789	U2791	A2792	A2793	C2794	C2795	C2796	U2797	C2798	U2799	C2800	A2801	C2802	A2803	A2804	A2805	A2806	C2807	U2808	U2809	A2810	C2811	C2812	A2813	A2814	U2815	A2816	C2817	C2818	A2819	U2820	U2821	C2822	C2823	U2824	C2825	G2826		
A2707	A2708	C2709	C2710	U2711	C2712	C2713	U2714	U2715	U2716	C2717	U2718	U2719	U2720	C2721	U2722	C2723	U2724	U2725	U2726	U2727	U2728	C2729	A2730	C2731	U2732	A2733	C2734	C2735	U2736	A2737	U2738	U2739	C2800	A2801	C2802	A2803	A2804	A2805	A2806	C2807	U2808	U2809	A2810	C2811	C2812	A2813	A2814	U2815	A2816	C2817	C2818	A2819	U2820	U2821	C2822	C2823	U2824	C2825	U2766	
C2647	C2648	C2649	A2650	C2650	C2651	C2652	U2653	U2654	U2655	U2656	C2657	U2658	A2659	U2660	C2661	A2662	U2663	U2664	A2665	C2666	U2667	U2668	C2669	A2670	C2671	C2672	C2673	C2674	C2675	U2676	C2677	C2678	C2679	A2680	C2681	A2682	A2683	U2684	C2685	C2686	C2687	C2688	U2689	A2690	A2691	C2692	C2693	U2694	U2695	U2696	A2697	C2698	A2699	U2700	C2701	C2702	C2703	U2704	A2705	A2706
G2587	C2588	C2589	C2590	C2591	C2592	A2593	A2594	C2595	A2596	C2597	U2598	A2599	U2600	C2601	U2602	U2603	A2604	C2605	C2606	U2607	U2608	C2609	C2610	C2611	A2612	C2613	U2614	U2615	U2616	C2617	C2618	C2619	C2620	C2621	C2622	C2623	C2624	C2625	C2626	C2627	C2628	C2629	A2630	C2631	C2632	C2633	U2634	C2635	U2636	U2637	A2638	A2639	U2640	C2641	C2642	C2643	U2644	A2645	A2646	





## 4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C1	Depositor
Number of particles used	2108230	Depositor
Resolution determination method	FSC 0.5 CUT-OFF	Depositor
CTF correction method	Wiener Filter on 3D volumes (SPIDER)	Depositor
Microscope	FEI TECNAI F30	Depositor
Voltage (kV)	300	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	25	Depositor
Minimum defocus (nm)	1000	Depositor
Maximum defocus (nm)	4500	Depositor
Magnification	38900	Depositor
Image detector	KODAK SO-163 FILM	Depositor
Maximum map value	0.454	Depositor
Minimum map value	-0.200	Depositor
Average map value	0.003	Depositor
Map value standard deviation	0.027	Depositor
Recommended contour level	0.11	Depositor
Map size ( $\text{\AA}$ )	455.4, 455.4, 455.4	wwPDB
Map dimensions	368, 368, 368	wwPDB
Map angles ( $^\circ$ )	90, 90, 90	wwPDB
Pixel spacing ( $\text{\AA}$ )	1.2375, 1.2375, 1.2375	Depositor

## 5 Model quality i

### 5.1 Standard geometry i

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	Ad	2.49	2030/42036 (4.8%)	2.11	2382/65520 (3.6%)
2	Ae	2.62	95/1781 (5.3%)	2.13	105/2775 (3.8%)
3	Af	2.45	12/260 (4.6%)	2.06	16/403 (4.0%)
4	BY	0.99	0/1123	1.10	1/1487 (0.1%)
5	BI	1.05	0/539	0.95	0/712
6	BK	0.93	0/840	1.21	6/1135 (0.5%)
7	BM	0.82	0/936	1.11	2/1260 (0.2%)
8	Bf	0.93	0/590	1.17	1/788 (0.1%)
9	BX	0.96	0/1122	1.05	4/1492 (0.3%)
10	Bg	0.97	0/2988	1.06	3/4049 (0.1%)
11	BD	1.01	0/1652	1.20	4/2222 (0.2%)
12	BE	0.99	0/1637	1.07	0/2202
13	BF	0.98	0/1509	1.00	3/2034 (0.1%)
14	BQ	1.11	0/1034	1.19	6/1379 (0.4%)
15	BU	0.93	0/995	1.14	3/1338 (0.2%)
16	BO	1.05	0/909	1.11	2/1217 (0.2%)
17	BS	1.04	0/1258	1.15	5/1674 (0.3%)
18	BN	0.96	0/994	1.13	5/1332 (0.4%)
19	BL	1.04	0/704	1.15	3/944 (0.3%)
20	BT	1.01	0/1179	1.08	3/1586 (0.2%)
21	BP	0.91	0/727	1.11	2/975 (0.2%)
22	BZ	0.94	0/791	1.18	7/1057 (0.7%)
23	Bc	1.04	0/455	1.26	2/609 (0.3%)
24	BW	1.02	0/1060	1.16	6/1419 (0.4%)
25	Bd	1.11	0/386	1.25	4/510 (0.8%)
26	Bb	0.92	0/674	1.04	0/905
27	Be	1.07	0/476	1.01	0/627
28	BA	0.96	0/1567	1.06	4/2121 (0.2%)
29	BR	1.03	0/955	1.03	1/1273 (0.1%)
30	BB	0.96	0/1736	1.12	4/2329 (0.2%)
31	BV	1.00	0/610	1.07	0/820
32	Ba	1.07	0/766	1.13	0/1023
33	BJ	1.09	0/1553	1.05	4/2079 (0.2%)
34	BC	0.93	0/1701	1.05	3/2298 (0.1%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
35	BG	1.06	0/1888	1.05	4/2507 (0.2%)
36	BH	3.40	1/1535 (0.1%)	1.14	4/2065 (0.2%)
37	CG	0.94	0/1939	1.01	5/2598 (0.2%)
38	CT	0.99	0/1316	1.11	2/1772 (0.1%)
39	CZ	1.00	0/1110	1.02	2/1480 (0.1%)
40	Cz	0.89	0/1741	1.00	1/2323 (0.0%)
41	CA	1.05	0/1992	1.15	10/2681 (0.4%)
42	CJ	1.06	0/1401	1.14	7/1869 (0.4%)
43	CH	0.96	0/1519	1.03	0/2042
44	CV	0.99	0/1064	1.07	0/1425
45	CN	1.12	0/1669	1.07	6/2235 (0.3%)
46	Ca	0.98	0/1143	1.17	4/1527 (0.3%)
47	CQ	1.04	0/1303	1.11	5/1748 (0.3%)
48	CD	1.00	0/2489	1.23	22/3342 (0.7%)
49	CR	1.09	0/1590	1.06	4/2100 (0.2%)
50	CP	1.03	0/1397	1.14	6/1871 (0.3%)
51	CX	0.90	0/1002	1.03	3/1340 (0.2%)
52	CW	1.04	0/649	1.07	1/861 (0.1%)
53	CY	1.10	0/1061	1.08	4/1418 (0.3%)
54	Cr	0.98	0/585	1.16	1/786 (0.1%)
55	Cc	0.86	0/869	0.98	1/1169 (0.1%)
56	Cd	1.01	0/970	1.10	4/1295 (0.3%)
57	Ce	1.01	0/1122	1.06	4/1497 (0.3%)
58	Cj	1.17	0/769	1.16	1/1019 (0.1%)
59	Cl	1.14	0/472	1.12	1/627 (0.2%)
60	Co	0.93	0/867	1.12	3/1144 (0.3%)
61	CM	0.99	0/1094	1.10	4/1461 (0.3%)
62	CS	1.01	0/1457	1.16	3/1957 (0.2%)
63	CU	0.98	0/876	1.27	12/1170 (1.0%)
64	Ci	1.07	0/618	1.16	5/809 (0.6%)
65	CK	0.92	0/968	1.11	1/1299 (0.1%)
66	Cu	0.78	0/438	0.91	0/596
66	Cv	0.79	0/438	0.90	0/596
67	Cs	0.83	0/444	0.82	0/596
67	Ct	0.84	0/444	0.81	0/596
68	Ch	1.02	0/1023	1.05	2/1359 (0.1%)
69	CF	0.98	0/2020	1.00	4/2708 (0.1%)
70	Cq	0.87	0/2023	0.96	5/2739 (0.2%)
71	CB	0.97	0/3207	1.14	16/4289 (0.4%)
72	CC	1.01	0/2951	1.11	8/3972 (0.2%)
73	CO	1.03	0/1678	1.07	6/2246 (0.3%)
74	Cp	1.00	0/724	1.02	1/958 (0.1%)
75	CI	1.01	0/1523	1.00	1/2036 (0.0%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
76	Cn	1.34	0/239	1.04	0/302
77	Cm	1.01	0/434	0.95	0/574
78	CL	1.03	0/1721	1.13	6/2299 (0.3%)
79	CE	0.95	0/1766	1.16	8/2374 (0.3%)
80	Cf	1.05	0/908	1.13	3/1215 (0.2%)
81	Ck	0.98	0/572	1.09	0/763
82	Cb	0.98	0/486	1.06	2/641 (0.3%)
83	Cg	1.07	0/913	1.02	0/1223
84	Aa	1.61	170/81235 (0.2%)	2.52	9121/126706 (7.2%)
85	Ac	1.61	7/3809 (0.2%)	2.48	426/5936 (7.2%)
86	Ab	2.31	125/2864 (4.4%)	2.91	380/4464 (8.5%)
All	All	1.65	2440/227878 (1.1%)	2.00	12710/334219 (3.8%)

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
4	BY	0	1
6	BK	0	4
7	BM	0	1
8	Bf	0	1
10	Bg	0	1
11	BD	0	3
12	BE	0	2
13	BF	0	2
14	BQ	0	1
15	BU	0	2
17	BS	0	1
19	BL	0	1
20	BT	0	4
23	Bc	0	1
24	BW	0	1
25	Bd	0	1
26	Bb	0	2
29	BR	0	1
30	BB	0	1
32	Ba	0	1
35	BG	0	1
36	BH	0	4
37	CG	0	2

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Mol	Chain	#Chirality outliers	#Planarity outliers
41	CA	0	4
42	CJ	0	2
43	CH	0	3
45	CN	0	1
46	Ca	0	7
47	CQ	0	6
48	CD	0	13
49	CR	0	3
50	CP	0	1
51	CX	0	1
55	Cc	0	1
57	Ce	0	1
58	Cj	0	1
59	Cl	0	1
60	Co	0	2
61	CM	0	4
62	CS	0	3
63	CU	0	2
65	CK	0	2
68	Ch	0	1
69	CF	0	3
70	Cq	0	2
71	CB	0	9
72	CC	0	4
73	CO	0	4
74	Cp	0	1
75	CI	0	4
78	CL	0	5
79	CE	0	7
80	Cf	0	2
83	Cg	0	1
84	Aa	0	309
85	Ac	0	18
86	Ab	0	19
All	All	0	486

All (2440) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BH	117	ARG	CZ-NH2	127.06	2.98	1.33
1	Ad	1203	G	C2'-C1'	23.55	1.79	1.53
2	Ae	28	G	C2'-C1'	-23.31	1.27	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	218	G	C2'-C1'	-23.01	1.28	1.53
1	Ad	999	G	C2'-C1'	-22.88	1.28	1.53
1	Ad	707	C	C2'-C1'	22.87	1.78	1.53
1	Ad	67	G	C2'-C1'	-22.47	1.28	1.53
1	Ad	1580	G	C2'-C1'	21.80	1.77	1.53
1	Ad	843	G	O4'-C1'	21.60	1.69	1.41
1	Ad	821	G	C2'-C1'	-21.45	1.29	1.53
1	Ad	1810	G	O4'-C1'	21.00	1.69	1.41
1	Ad	1796	G	C2'-C1'	-20.71	1.30	1.53
1	Ad	1080	C	O4'-C1'	20.56	1.68	1.41
1	Ad	1434	G	C2'-C1'	-20.34	1.30	1.53
1	Ad	260	A	C2'-C1'	-20.17	1.31	1.53
1	Ad	141	G	C2'-C1'	-20.08	1.31	1.53
1	Ad	457	C	C2'-C1'	-20.07	1.31	1.53
1	Ad	339	G	C2'-C1'	-19.92	1.31	1.53
1	Ad	1206	A	C2'-C1'	-19.53	1.31	1.53
1	Ad	1259	G	O4'-C1'	-19.39	1.16	1.41
2	Ae	19	U	O4'-C1'	19.14	1.66	1.41
1	Ad	179	A	O4'-C1'	18.97	1.66	1.41
1	Ad	836	U	O4'-C1'	18.92	1.66	1.41
1	Ad	67	G	O4'-C1'	18.85	1.66	1.41
1	Ad	617	G	C2'-C1'	-18.83	1.32	1.53
1	Ad	220	C	O4'-C1'	18.76	1.66	1.41
1	Ad	1206	A	O4'-C1'	18.70	1.66	1.41
1	Ad	1700	G	C2'-C1'	-18.54	1.32	1.53
1	Ad	782	G	C2'-C1'	-18.37	1.33	1.53
1	Ad	96	G	C2'-C1'	-18.25	1.33	1.53
1	Ad	648	C	O4'-C1'	18.05	1.65	1.41
1	Ad	1155	G	C2'-C1'	-17.96	1.33	1.53
1	Ad	176	A	C2'-C1'	17.93	1.73	1.53
1	Ad	87	A	C2'-C1'	17.86	1.73	1.53
2	Ae	69	G	C2'-C1'	-17.84	1.33	1.53
1	Ad	1260	A	C2'-C1'	-17.82	1.33	1.53
1	Ad	1466	A	C2'-C1'	-17.65	1.33	1.53
1	Ad	212	A	O4'-C1'	17.64	1.64	1.41
1	Ad	257	A	C2'-C1'	-17.60	1.33	1.53
1	Ad	1303	G	C2'-C1'	-17.57	1.34	1.53
1	Ad	238	G	O4'-C1'	17.52	1.64	1.41
1	Ad	238	G	C2'-C1'	-17.51	1.34	1.53
1	Ad	1002	G	C2'-C1'	-17.51	1.34	1.53
1	Ad	1464	G	C2'-C1'	-17.47	1.34	1.53
1	Ad	34	G	C2'-C1'	-17.43	1.34	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	617	G	O4'-C1'	17.41	1.64	1.41
2	Ae	30	G	C2'-C1'	-17.37	1.34	1.53
1	Ad	1444	G	C2'-C1'	-17.29	1.34	1.53
1	Ad	846	U	O4'-C1'	17.23	1.64	1.41
1	Ad	728	C	O4'-C1'	17.11	1.63	1.41
1	Ad	1580	G	O4'-C1'	-17.10	1.19	1.41
1	Ad	1445	C	O4'-C1'	17.10	1.63	1.41
1	Ad	140	C	O4'-C1'	17.02	1.63	1.41
1	Ad	524	A	C2'-C1'	-16.98	1.34	1.53
1	Ad	1311	U	C2'-C1'	16.95	1.72	1.53
1	Ad	1162	A	C2'-C1'	16.93	1.72	1.53
1	Ad	1375	C	O4'-C1'	16.91	1.63	1.41
1	Ad	174	C	C2'-C1'	16.86	1.71	1.53
1	Ad	613	U	O4'-C1'	16.82	1.63	1.41
1	Ad	254	A	C2'-C1'	-16.78	1.34	1.53
1	Ad	1161	C	O4'-C1'	16.71	1.63	1.41
1	Ad	1358	G	C2'-C1'	-16.67	1.35	1.53
1	Ad	1065	A	C2'-C1'	-16.66	1.35	1.53
1	Ad	1637	G	C2'-C1'	-16.66	1.35	1.53
1	Ad	1006	A	O4'-C1'	16.64	1.63	1.41
1	Ad	341	G	C2'-C1'	-16.53	1.35	1.53
1	Ad	570	C	C2'-C1'	-16.45	1.35	1.53
1	Ad	936	C	O4'-C1'	16.39	1.62	1.41
1	Ad	1447	C	C2'-C1'	-16.37	1.35	1.53
2	Ae	45	G	C2'-C1'	-16.31	1.35	1.53
1	Ad	385	C	O4'-C1'	16.30	1.62	1.41
1	Ad	998	A	C2'-C1'	-16.23	1.35	1.53
1	Ad	507	G	C2'-C1'	-16.18	1.35	1.53
1	Ad	14	C	C2'-C1'	-16.17	1.35	1.53
1	Ad	612	U	C2'-C1'	-16.14	1.35	1.53
1	Ad	861	A	C2'-C1'	16.03	1.71	1.53
1	Ad	1080	C	C2'-C1'	-15.99	1.35	1.53
1	Ad	280	U	O3'-P	-15.98	1.42	1.61
1	Ad	114	U	O4'-C1'	15.96	1.62	1.41
1	Ad	437	C	C2'-C1'	-15.93	1.35	1.53
1	Ad	1321	C	C2'-C1'	-15.88	1.35	1.53
1	Ad	1650	G	C2'-C1'	-15.80	1.35	1.53
1	Ad	281	U	O3'-P	-15.79	1.42	1.61
1	Ad	1687	G	C2'-C1'	-15.75	1.36	1.53
1	Ad	80	C	O4'-C1'	15.74	1.62	1.41
1	Ad	535	C	C2'-C1'	-15.71	1.36	1.53
1	Ad	504	C	O4'-C1'	15.69	1.62	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	839	G	C2'-C1'	-15.62	1.36	1.53
1	Ad	225	G	C2'-C1'	-15.62	1.36	1.53
1	Ad	1327	C	O4'-C1'	15.62	1.61	1.41
1	Ad	632	G	C2'-C1'	-15.61	1.36	1.53
1	Ad	1397	A	C2'-C1'	-15.57	1.36	1.53
1	Ad	135	C	O4'-C1'	15.56	1.61	1.41
1	Ad	1263	C	O4'-C1'	15.55	1.61	1.41
1	Ad	1391	G	C2'-C1'	-15.45	1.36	1.53
1	Ad	631	C	C2'-C1'	-15.44	1.36	1.53
1	Ad	282	C	C2'-C1'	15.42	1.70	1.53
1	Ad	1589	C	O4'-C1'	15.41	1.61	1.41
1	Ad	321	C	O4'-C1'	15.41	1.61	1.41
1	Ad	713	C	O4'-C1'	15.41	1.61	1.41
1	Ad	1354	C	C2'-C1'	-15.39	1.36	1.53
1	Ad	193	G	C2'-C1'	-15.36	1.36	1.53
1	Ad	1027	C	O4'-C1'	15.31	1.61	1.41
1	Ad	1184	C	O4'-C1'	15.30	1.61	1.41
1	Ad	861	A	O4'-C1'	15.28	1.61	1.41
1	Ad	1372	C	C2'-C1'	-15.15	1.36	1.53
1	Ad	287	C	O4'-C1'	15.14	1.61	1.41
1	Ad	1395	C	C2'-C1'	-15.14	1.36	1.53
1	Ad	158	C	C2'-C1'	-15.12	1.36	1.53
1	Ad	1730	G	O4'-C1'	15.05	1.61	1.41
1	Ad	25	C	O4'-C1'	15.04	1.61	1.41
1	Ad	457	C	O4'-C1'	15.04	1.61	1.41
1	Ad	1705	C	C2'-C1'	-15.03	1.36	1.53
1	Ad	1172	G	C2'-C1'	-15.01	1.36	1.53
1	Ad	223	A	C2'-C1'	-15.00	1.36	1.53
1	Ad	1154	G	C2'-C1'	-14.99	1.36	1.53
1	Ad	1023	C	O4'-C1'	14.98	1.61	1.41
1	Ad	753	C	C2'-C1'	-14.97	1.36	1.53
1	Ad	181	C	O4'-C1'	14.91	1.61	1.41
1	Ad	212	A	C2'-C1'	-14.86	1.37	1.53
1	Ad	1626	C	O4'-C1'	14.86	1.60	1.41
1	Ad	1778	G	C2'-C1'	-14.85	1.37	1.53
1	Ad	4	C	C2'-C1'	-14.77	1.37	1.53
1	Ad	381	G	C2'-C1'	-14.74	1.37	1.53
1	Ad	498	U	C2'-C1'	14.74	1.69	1.53
3	Af	17	A	C2'-C1'	-14.69	1.37	1.53
2	Ae	36	C	C2'-C1'	-14.68	1.37	1.53
1	Ad	1735	C	O4'-C1'	14.68	1.60	1.41
3	Af	17	A	O4'-C1'	14.64	1.60	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1358	G	O4'-C1'	14.63	1.60	1.41
1	Ad	290	C	O4'-C1'	14.62	1.60	1.41
1	Ad	480	U	C2'-C1'	-14.58	1.37	1.53
1	Ad	1783	C	C2'-C1'	-14.58	1.37	1.53
1	Ad	945	A	C2'-C1'	14.57	1.69	1.53
1	Ad	506	G	C2'-C1'	-14.56	1.37	1.53
1	Ad	509	A	O4'-C1'	14.55	1.60	1.41
1	Ad	1016	C	O4'-C1'	14.55	1.60	1.41
1	Ad	346	C	C2'-C1'	-14.53	1.37	1.53
1	Ad	183	C	O4'-C1'	14.49	1.60	1.41
1	Ad	1614	C	O4'-C1'	14.47	1.60	1.41
1	Ad	1346	C	O4'-C1'	14.44	1.60	1.41
1	Ad	1806	C	O4'-C1'	14.42	1.60	1.41
1	Ad	245	C	C2'-C1'	-14.39	1.37	1.53
1	Ad	774	C	O4'-C1'	14.36	1.60	1.41
1	Ad	1392	G	C2'-C1'	-14.35	1.37	1.53
1	Ad	1310	C	O4'-C1'	14.34	1.60	1.41
1	Ad	413	C	O4'-C1'	14.30	1.60	1.41
1	Ad	1263	C	C2'-C1'	-14.28	1.37	1.53
1	Ad	524	A	O4'-C1'	14.25	1.60	1.41
1	Ad	1461	G	C2'-C1'	-14.24	1.37	1.53
1	Ad	758	A	C2'-C1'	-14.22	1.37	1.53
1	Ad	257	A	O4'-C1'	14.21	1.60	1.41
1	Ad	1110	C	O4'-C1'	14.19	1.60	1.41
1	Ad	1444	G	O4'-C1'	14.14	1.60	1.41
1	Ad	938	A	C2'-C1'	-14.13	1.37	1.53
1	Ad	966	U	C2'-C1'	-14.12	1.37	1.53
1	Ad	1611	U	C2'-C1'	-14.10	1.37	1.53
1	Ad	201	G	C2'-C1'	-14.10	1.37	1.53
1	Ad	1620	C	C2'-C1'	-14.09	1.37	1.53
1	Ad	791	C	C2'-C1'	-14.07	1.37	1.53
1	Ad	1372	C	O4'-C1'	14.07	1.59	1.41
1	Ad	439	C	C2'-C1'	-14.06	1.37	1.53
1	Ad	187	C	O4'-C1'	14.04	1.59	1.41
1	Ad	1471	C	C2'-C1'	-14.02	1.38	1.53
1	Ad	1336	C	O4'-C1'	14.02	1.59	1.41
1	Ad	25	C	C2'-C1'	-13.99	1.38	1.53
1	Ad	1397	A	O4'-C1'	13.96	1.59	1.41
1	Ad	1416	A	C2'-C1'	-13.89	1.38	1.53
1	Ad	409	C	C2'-C1'	-13.89	1.38	1.53
1	Ad	1345	G	C2'-C1'	-13.88	1.38	1.53
1	Ad	1640	C	C2'-C1'	-13.86	1.38	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	205	U	C2'-C1'	13.84	1.68	1.53
1	Ad	914	U	C2'-C1'	-13.84	1.38	1.53
1	Ad	20	G	C2'-C1'	-13.83	1.38	1.53
1	Ad	719	C	O4'-C1'	13.78	1.59	1.41
1	Ad	1472	G	C2'-C1'	-13.77	1.38	1.53
1	Ad	849	G	C2'-C1'	-13.77	1.38	1.53
1	Ad	745	C	C2'-C1'	13.76	1.68	1.53
2	Ae	73	C	O4'-C1'	13.76	1.59	1.41
1	Ad	36	C	O4'-C1'	13.73	1.59	1.41
1	Ad	245	C	O4'-C1'	13.73	1.59	1.41
1	Ad	164	C	O4'-C1'	13.71	1.59	1.41
1	Ad	848	C	O4'-C1'	13.70	1.59	1.41
1	Ad	1069	G	C2'-C1'	-13.68	1.38	1.53
1	Ad	650	G	C2'-C1'	-13.64	1.38	1.53
1	Ad	839	G	O4'-C1'	13.63	1.59	1.41
1	Ad	1281	G	C2'-C1'	-13.63	1.38	1.53
1	Ad	1042	C	O4'-C1'	13.62	1.59	1.41
1	Ad	1705	C	O4'-C1'	13.59	1.59	1.41
1	Ad	533	C	C2'-C1'	-13.58	1.38	1.53
1	Ad	1296	G	C2'-C1'	-13.57	1.38	1.53
2	Ae	19	U	C2'-C1'	-13.57	1.38	1.53
1	Ad	644	U	C2'-C1'	-13.55	1.38	1.53
1	Ad	1329	A	O4'-C1'	13.54	1.59	1.41
1	Ad	237	C	C2'-C1'	-13.51	1.38	1.53
1	Ad	1654	C	O4'-C1'	13.51	1.59	1.41
1	Ad	220	C	C2'-C1'	-13.49	1.38	1.53
1	Ad	73	A	C2'-C1'	13.48	1.68	1.53
1	Ad	734	C	O4'-C1'	13.44	1.59	1.41
1	Ad	846	U	C2'-C1'	-13.43	1.38	1.53
1	Ad	1462	C	O4'-C1'	-13.42	1.24	1.41
1	Ad	1310	C	C2'-C1'	-13.41	1.38	1.53
1	Ad	793	G	C2'-C1'	-13.38	1.38	1.53
1	Ad	954	C	O4'-C1'	13.37	1.59	1.41
1	Ad	1038	C	O4'-C1'	13.34	1.58	1.41
1	Ad	877	G	C2'-C1'	-13.34	1.38	1.53
1	Ad	1033	C	O4'-C1'	13.34	1.58	1.41
1	Ad	1091	A	C2'-C1'	-13.31	1.38	1.53
1	Ad	286	C	O4'-C1'	13.31	1.58	1.41
1	Ad	944	A	O4'-C1'	13.30	1.58	1.41
1	Ad	94	A	C2'-C1'	-13.29	1.38	1.53
1	Ad	1231	A	C2'-C1'	-13.28	1.38	1.53
1	Ad	301	U	C2'-C1'	-13.27	1.38	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	256	G	O4'-C1'	13.26	1.58	1.41
2	Ae	55	C	O4'-C1'	13.26	1.58	1.41
1	Ad	1548	G	C2'-C1'	-13.25	1.38	1.53
1	Ad	1511	A	O4'-C1'	-13.25	1.24	1.41
1	Ad	704	C	O4'-C1'	13.23	1.58	1.41
1	Ad	1105	G	C2'-C1'	-13.21	1.38	1.53
1	Ad	954	C	C2'-C1'	-13.17	1.38	1.53
1	Ad	1302	C	O4'-C1'	13.16	1.58	1.41
1	Ad	1045	G	C2'-C1'	-13.14	1.38	1.53
2	Ae	60	C	C2'-C1'	-13.14	1.38	1.53
1	Ad	249	G	C2'-C1'	-13.14	1.38	1.53
1	Ad	1463	C	O4'-C1'	13.13	1.58	1.41
1	Ad	262	U	C2'-C1'	13.13	1.67	1.53
1	Ad	358	C	O4'-C1'	13.12	1.58	1.41
1	Ad	1801	A	C2'-C1'	13.12	1.67	1.53
1	Ad	202	C	O4'-C1'	13.11	1.58	1.41
1	Ad	1082	C	C2'-C1'	-13.11	1.39	1.53
1	Ad	1301	G	C2'-C1'	13.09	1.67	1.53
1	Ad	902	C	O4'-C1'	13.09	1.58	1.41
1	Ad	180	A	C2'-C1'	13.05	1.67	1.53
1	Ad	1704	G	C2'-C1'	-13.05	1.39	1.53
1	Ad	177	C	O4'-C1'	13.04	1.58	1.41
1	Ad	235	C	C2'-C1'	13.04	1.67	1.53
1	Ad	558	C	O4'-C1'	13.03	1.58	1.41
1	Ad	1727	C	O4'-C1'	13.02	1.58	1.41
1	Ad	536	U	C2'-C1'	-13.02	1.39	1.53
1	Ad	147	C	O4'-C1'	12.94	1.58	1.41
1	Ad	1599	C	O4'-C1'	12.94	1.58	1.41
1	Ad	1029	U	C2'-C1'	12.92	1.67	1.53
1	Ad	1409	G	C2'-C1'	-12.92	1.39	1.53
1	Ad	1184	C	C2'-C1'	-12.91	1.39	1.53
1	Ad	31	C	O4'-C1'	12.90	1.58	1.41
2	Ae	6	G	C2'-C1'	-12.90	1.39	1.53
1	Ad	757	G	C2'-C1'	-12.89	1.39	1.53
1	Ad	108	C	C2'-C1'	-12.88	1.39	1.53
2	Ae	45	G	O4'-C1'	12.84	1.58	1.41
1	Ad	1161	C	C2'-C1'	-12.84	1.39	1.53
1	Ad	1442	A	C2'-C1'	-12.84	1.39	1.53
1	Ad	1007	G	C2'-C1'	-12.84	1.39	1.53
1	Ad	1609	G	C2'-C1'	-12.82	1.39	1.53
1	Ad	1071	C	C2'-C1'	-12.78	1.39	1.53
1	Ad	971	A	O4'-C1'	12.77	1.58	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1615	G	C2'-C1'	-12.77	1.39	1.53
1	Ad	849	G	O4'-C1'	12.76	1.58	1.41
1	Ad	1572	U	O4'-C1'	12.76	1.58	1.41
1	Ad	1794	C	O4'-C1'	12.72	1.58	1.41
1	Ad	1758	G	C2'-C1'	-12.70	1.39	1.53
1	Ad	260	A	O4'-C1'	12.70	1.58	1.41
1	Ad	814	C	C2'-C1'	-12.70	1.39	1.53
1	Ad	1346	C	C2'-C1'	-12.68	1.39	1.53
1	Ad	296	A	C2'-C1'	-12.68	1.39	1.53
1	Ad	261	C	C2'-C1'	-12.66	1.39	1.53
1	Ad	1618	G	C2'-C1'	-12.62	1.39	1.53
1	Ad	881	G	C2'-C1'	-12.60	1.39	1.53
1	Ad	1715	C	O4'-C1'	12.60	1.58	1.41
1	Ad	1220	C	O4'-C1'	12.58	1.58	1.41
1	Ad	1082	C	O4'-C1'	12.57	1.57	1.41
1	Ad	1350	C	O4'-C1'	-12.57	1.25	1.41
1	Ad	1386	U	C2'-C1'	-12.57	1.39	1.53
1	Ad	253	C	O4'-C1'	12.57	1.57	1.41
1	Ad	1803	G	O4'-C1'	12.57	1.57	1.41
1	Ad	1020	U	C2'-C1'	-12.56	1.39	1.53
1	Ad	431	C	O4'-C1'	12.55	1.57	1.41
1	Ad	1593	U	C2'-C1'	-12.55	1.39	1.53
1	Ad	1232	G	C2'-C1'	-12.53	1.39	1.53
1	Ad	1631	C	O4'-C1'	12.53	1.57	1.41
1	Ad	388	G	C2'-C1'	-12.53	1.39	1.53
1	Ad	1047	G	C2'-C1'	-12.53	1.39	1.53
1	Ad	1656	C	O4'-C1'	12.52	1.57	1.41
1	Ad	570	C	O4'-C1'	12.52	1.57	1.41
1	Ad	845	C	C2'-C1'	12.52	1.67	1.53
1	Ad	834	A	C2'-C1'	12.52	1.67	1.53
1	Ad	1643	A	C2'-C1'	12.52	1.67	1.53
1	Ad	1773	A	O4'-C1'	12.52	1.57	1.41
1	Ad	1722	C	O4'-C1'	12.51	1.57	1.41
1	Ad	41	A	C2'-C1'	12.49	1.67	1.53
1	Ad	592	U	C2'-C1'	-12.48	1.39	1.53
1	Ad	342	C	O4'-C1'	12.48	1.57	1.41
1	Ad	1471	C	O4'-C1'	12.48	1.57	1.41
1	Ad	746	A	C2'-C1'	12.47	1.67	1.53
1	Ad	856	G	C2'-C1'	-12.47	1.39	1.53
1	Ad	259	A	O4'-C1'	12.47	1.57	1.41
1	Ad	762	A	O4'-C1'	12.46	1.57	1.41
1	Ad	12	U	C2'-C1'	-12.46	1.39	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	750	U	C2'-C1'	-12.46	1.39	1.53
1	Ad	1171	C	O4'-C1'	12.46	1.57	1.41
1	Ad	1336	C	C2'-C1'	-12.45	1.39	1.53
1	Ad	826	C	C2'-C1'	12.45	1.67	1.53
1	Ad	170	C	O4'-C1'	12.43	1.57	1.41
1	Ad	298	C	O4'-C1'	12.43	1.57	1.41
1	Ad	58	U	C2'-C1'	-12.43	1.39	1.53
1	Ad	369	G	C2'-C1'	12.43	1.67	1.53
1	Ad	1074	C	O4'-C1'	12.43	1.57	1.41
1	Ad	1500	A	C2'-C1'	12.42	1.67	1.53
1	Ad	297	U	O4'-C1'	12.40	1.57	1.41
1	Ad	1549	G	C2'-C1'	-12.39	1.39	1.53
1	Ad	547	C	O4'-C1'	-12.39	1.25	1.41
1	Ad	1582	G	O4'-C1'	12.36	1.57	1.41
1	Ad	1733	G	C2'-C1'	-12.36	1.39	1.53
1	Ad	1196	C	O4'-C1'	12.35	1.57	1.41
1	Ad	12	U	O4'-C1'	12.35	1.57	1.41
1	Ad	30	G	C2'-C1'	-12.35	1.39	1.53
1	Ad	753	C	O4'-C1'	12.34	1.57	1.41
1	Ad	1432	C	C2'-C1'	12.34	1.67	1.53
1	Ad	791	C	O4'-C1'	12.31	1.57	1.41
1	Ad	899	A	C2'-C1'	-12.29	1.39	1.53
1	Ad	1517	C	C2'-C1'	-12.25	1.39	1.53
1	Ad	351	G	C2'-C1'	-12.25	1.39	1.53
1	Ad	1213	C	O4'-C1'	12.23	1.57	1.41
1	Ad	1229	C	O4'-C1'	12.22	1.57	1.41
1	Ad	523	C	O4'-C1'	12.22	1.57	1.41
1	Ad	1644	C	C2'-C1'	-12.22	1.40	1.53
1	Ad	1240	A	O4'-C1'	12.19	1.57	1.41
1	Ad	356	G	O4'-C1'	12.19	1.57	1.41
1	Ad	1588	C	O4'-C1'	12.19	1.57	1.41
1	Ad	438	G	C2'-C1'	12.17	1.66	1.53
1	Ad	1156	A	C2'-C1'	-12.16	1.40	1.53
1	Ad	363	G	O4'-C1'	-12.16	1.25	1.41
1	Ad	439	C	O4'-C1'	12.15	1.57	1.41
1	Ad	1273	U	C2'-C1'	-12.13	1.40	1.53
1	Ad	1563	A	C2'-C1'	-12.13	1.40	1.53
1	Ad	336	U	C2'-C1'	12.12	1.66	1.53
1	Ad	237	C	O4'-C1'	12.12	1.57	1.41
1	Ad	1373	C	O4'-C1'	12.12	1.57	1.41
1	Ad	1253	U	C2'-C1'	-12.12	1.40	1.53
1	Ad	45	U	O4'-C1'	12.12	1.57	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	784	C	O4'-C1'	-12.10	1.25	1.41
1	Ad	1494	G	C2'-C1'	-12.09	1.40	1.53
1	Ad	361	G	C2'-C1'	-12.08	1.40	1.53
1	Ad	343	C	C2'-C1'	-12.08	1.40	1.53
1	Ad	394	G	C2'-C1'	12.05	1.66	1.53
1	Ad	922	U	C2'-C1'	-12.04	1.40	1.53
1	Ad	196	G	C2'-C1'	-12.04	1.40	1.53
3	Af	18	C	O4'-C1'	12.03	1.57	1.41
2	Ae	74	C	C2'-C1'	12.02	1.66	1.53
1	Ad	1021	C	O4'-C1'	12.01	1.57	1.41
1	Ad	1518	C	O4'-C1'	12.00	1.57	1.41
1	Ad	1252	C	C2'-C1'	-12.00	1.40	1.53
1	Ad	415	C	O4'-C1'	11.98	1.57	1.41
1	Ad	397	C	O4'-C1'	11.97	1.57	1.41
1	Ad	967	C	O4'-C1'	11.96	1.57	1.41
1	Ad	1652	C	O4'-C1'	11.96	1.57	1.41
1	Ad	1046	G	C2'-C1'	-11.96	1.40	1.53
1	Ad	1608	A	C2'-C1'	-11.94	1.40	1.53
1	Ad	334	G	C2'-C1'	-11.93	1.40	1.53
1	Ad	1488	C	O4'-C1'	11.92	1.57	1.41
1	Ad	1238	A	O4'-C1'	11.89	1.57	1.41
1	Ad	1342	C	C2'-C1'	-11.89	1.40	1.53
1	Ad	900	G	C2'-C1'	-11.86	1.40	1.53
1	Ad	588	C	O4'-C1'	11.85	1.57	1.41
1	Ad	1332	G	O4'-C1'	11.84	1.57	1.41
1	Ad	1610	C	O4'-C1'	11.83	1.57	1.41
1	Ad	535	C	O4'-C1'	11.83	1.57	1.41
1	Ad	644	U	O4'-C1'	11.81	1.57	1.41
1	Ad	259	A	C2'-C1'	-11.81	1.40	1.53
1	Ad	850	G	C2'-C1'	-11.79	1.40	1.53
1	Ad	870	A	C2'-C1'	11.79	1.66	1.53
1	Ad	700	C	O4'-C1'	11.75	1.56	1.41
1	Ad	1327	C	C2'-C1'	-11.74	1.40	1.53
1	Ad	117	U	C2'-C1'	-11.74	1.40	1.53
1	Ad	948	C	O4'-C1'	11.74	1.56	1.41
1	Ad	869	U	C2'-C1'	11.73	1.66	1.53
1	Ad	964	U	C2'-C1'	-11.73	1.40	1.53
2	Ae	29	C	O4'-C1'	11.71	1.56	1.41
1	Ad	915	C	O4'-C1'	11.70	1.56	1.41
1	Ad	955	C	O4'-C1'	11.70	1.56	1.41
1	Ad	1316	A	C2'-C1'	-11.70	1.40	1.53
1	Ad	1258	U	C2'-C1'	-11.69	1.40	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	202	C	C2'-C1'	-11.69	1.40	1.53
3	Af	20	U	C2'-C1'	-11.68	1.40	1.53
1	Ad	219	G	C2'-C1'	11.68	1.66	1.53
1	Ad	1557	C	O4'-C1'	11.65	1.56	1.41
1	Ad	832	C	C2'-C1'	-11.64	1.40	1.53
1	Ad	1686	C	C2'-C1'	-11.63	1.40	1.53
1	Ad	1589	C	C2'-C1'	-11.62	1.40	1.53
1	Ad	1710	C	O4'-C1'	11.62	1.56	1.41
2	Ae	43	C	C2'-C1'	-11.62	1.40	1.53
1	Ad	1648	C	O4'-C1'	11.61	1.56	1.41
1	Ad	1162	A	O4'-C1'	-11.61	1.26	1.41
1	Ad	346	C	O4'-C1'	11.59	1.56	1.41
1	Ad	1122	U	C2'-C1'	-11.59	1.40	1.53
1	Ad	1627	C	O4'-C1'	11.57	1.56	1.41
1	Ad	99	U	C2'-C1'	-11.57	1.40	1.53
1	Ad	1568	U	C2'-C1'	11.56	1.66	1.53
1	Ad	973	U	C2'-C1'	-11.55	1.40	1.53
1	Ad	488	C	O4'-C1'	11.55	1.56	1.41
1	Ad	1457	C	O4'-C1'	11.55	1.56	1.41
1	Ad	1110	C	C2'-C1'	-11.54	1.40	1.53
1	Ad	1407	A	C2'-C1'	11.54	1.66	1.53
1	Ad	1281	G	O4'-C1'	11.53	1.56	1.41
1	Ad	832	C	O4'-C1'	11.53	1.56	1.41
1	Ad	1731	A	O4'-C1'	11.52	1.56	1.41
1	Ad	146	A	C2'-C1'	11.50	1.66	1.53
1	Ad	801	U	C2'-C1'	11.50	1.66	1.53
1	Ad	1073	C	O4'-C1'	11.50	1.56	1.41
1	Ad	1032	A	C2'-C1'	11.48	1.66	1.53
1	Ad	968	A	C2'-C1'	11.48	1.66	1.53
1	Ad	1794	C	C2'-C1'	-11.47	1.40	1.53
1	Ad	1079	G	C2'-C1'	-11.47	1.40	1.53
1	Ad	1716	C	C2'-C1'	-11.46	1.40	1.53
1	Ad	1117	G	C2'-C1'	-11.46	1.40	1.53
1	Ad	1736	C	O4'-C1'	11.44	1.56	1.41
1	Ad	1451	G	O4'-C1'	11.42	1.56	1.41
2	Ae	4	G	C2'-C1'	-11.41	1.40	1.53
1	Ad	483	C	O4'-C1'	11.40	1.56	1.41
1	Ad	858	G	O4'-C1'	11.39	1.56	1.41
1	Ad	969	U	C2'-C1'	-11.38	1.40	1.53
1	Ad	1027	C	C2'-C1'	-11.36	1.40	1.53
1	Ad	288	G	C2'-C1'	-11.35	1.40	1.53
1	Ad	385	C	C2'-C1'	-11.35	1.40	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1058	G	C2'-C1'	-11.34	1.40	1.53
1	Ad	1447	C	O4'-C1'	11.34	1.56	1.41
1	Ad	1613	G	C2'-C1'	-11.34	1.40	1.53
1	Ad	1762	C	O4'-C1'	11.33	1.56	1.41
2	Ae	50	G	C2'-C1'	11.32	1.65	1.53
1	Ad	1426	C	O4'-C1'	11.30	1.56	1.41
1	Ad	239	C	O4'-C1'	11.30	1.56	1.41
2	Ae	22	G	C2'-C1'	-11.29	1.41	1.53
1	Ad	1715	C	C2'-C1'	-11.29	1.41	1.53
1	Ad	792	U	C2'-C1'	11.29	1.65	1.53
1	Ad	1064	U	C2'-C1'	11.28	1.65	1.53
1	Ad	133	U	O4'-C1'	11.28	1.56	1.41
1	Ad	711	C	C2'-C1'	-11.28	1.41	1.53
1	Ad	437	C	O4'-C1'	11.27	1.56	1.41
1	Ad	297	U	C2'-C1'	-11.26	1.41	1.53
1	Ad	1685	U	C2'-C1'	-11.26	1.41	1.53
1	Ad	1749	C	C2'-C1'	-11.26	1.41	1.53
2	Ae	60	C	O4'-C1'	11.25	1.56	1.41
1	Ad	949	A	O4'-C1'	11.22	1.56	1.41
1	Ad	1492	G	C2'-C1'	11.22	1.65	1.53
1	Ad	591	C	C2'-C1'	-11.21	1.41	1.53
1	Ad	885	C	O4'-C1'	11.21	1.56	1.41
1	Ad	651	G	C2'-C1'	-11.20	1.41	1.53
1	Ad	556	G	C2'-C1'	-11.20	1.41	1.53
1	Ad	1224	C	C2'-C1'	-11.18	1.41	1.53
1	Ad	81	U	O4'-C1'	11.18	1.56	1.41
1	Ad	995	C	C2'-C1'	-11.17	1.41	1.53
1	Ad	1545	A	C2'-C1'	11.17	1.65	1.53
1	Ad	1013	G	C2'-C1'	-11.17	1.41	1.53
1	Ad	1128	C	O4'-C1'	11.16	1.56	1.41
1	Ad	1239	C	O4'-C1'	11.16	1.56	1.41
1	Ad	804	C	O4'-C1'	11.15	1.56	1.41
1	Ad	182	C	O4'-C1'	11.15	1.56	1.41
1	Ad	1572	U	C2'-C1'	-11.15	1.41	1.53
1	Ad	175	A	O4'-C1'	11.15	1.56	1.41
1	Ad	45	U	C2'-C1'	-11.14	1.41	1.53
1	Ad	1414	G	C2'-C1'	-11.13	1.41	1.53
1	Ad	79	A	C2'-C1'	11.13	1.65	1.53
1	Ad	68	A	C2'-C1'	-11.12	1.41	1.53
1	Ad	1375	C	C2'-C1'	-11.12	1.41	1.53
1	Ad	1596	G	O4'-C1'	11.12	1.56	1.41
1	Ad	1476	C	C2'-C1'	-11.12	1.41	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	713	C	C2'-C1'	-11.11	1.41	1.53
1	Ad	879	C	C2'-C1'	-11.11	1.41	1.53
1	Ad	17	C	C2'-C1'	-11.11	1.41	1.53
1	Ad	169	A	O4'-C1'	11.11	1.56	1.41
1	Ad	587	C	C2'-C1'	-11.10	1.41	1.53
1	Ad	345	A	C2'-C1'	-11.10	1.41	1.53
1	Ad	1174	G	C2'-C1'	-11.10	1.41	1.53
1	Ad	1735	C	C2'-C1'	-11.08	1.41	1.53
1	Ad	1670	G	C2'-C1'	-11.07	1.41	1.53
1	Ad	428	C	C2'-C1'	-11.06	1.41	1.53
1	Ad	1037	G	C2'-C1'	-11.04	1.41	1.53
1	Ad	49	C	O4'-C1'	11.04	1.55	1.41
1	Ad	814	C	O4'-C1'	11.01	1.55	1.41
1	Ad	412	C	O4'-C1'	11.00	1.55	1.41
1	Ad	53	G	C2'-C1'	-10.98	1.41	1.53
1	Ad	121	U	O4'-C1'	10.98	1.55	1.41
1	Ad	1673	C	O4'-C1'	10.97	1.55	1.41
1	Ad	1516	C	C2'-C1'	-10.97	1.41	1.53
1	Ad	591	C	O4'-C1'	10.96	1.55	1.41
1	Ad	1507	G	C2'-C1'	-10.96	1.41	1.53
1	Ad	1319	U	C2'-C1'	-10.96	1.41	1.53
1	Ad	391	A	C2'-C1'	10.95	1.65	1.53
1	Ad	309	C	O4'-C1'	10.95	1.55	1.41
1	Ad	1443	U	C2'-C1'	-10.95	1.41	1.53
1	Ad	1502	C	O4'-C1'	10.94	1.55	1.41
1	Ad	384	U	O4'-C1'	-10.94	1.27	1.41
2	Ae	26	G	C2'-C1'	-10.92	1.41	1.53
2	Ae	56	A	O4'-C1'	10.91	1.55	1.41
1	Ad	1208	A	O4'-C1'	10.91	1.55	1.41
1	Ad	1765	A	C2'-C1'	10.91	1.65	1.53
2	Ae	58	U	C2'-C1'	10.91	1.65	1.53
1	Ad	1716	C	O4'-C1'	10.90	1.55	1.41
1	Ad	950	U	O4'-C1'	10.88	1.55	1.41
1	Ad	1515	G	C2'-C1'	-10.88	1.41	1.53
1	Ad	1699	C	O4'-C1'	10.87	1.55	1.41
1	Ad	649	C	C2'-C1'	-10.86	1.41	1.53
1	Ad	216	A	C2'-C1'	-10.85	1.41	1.53
1	Ad	1210	U	C2'-C1'	10.85	1.65	1.53
1	Ad	198	G	C2'-C1'	-10.83	1.41	1.53
1	Ad	158	C	O4'-C1'	10.83	1.55	1.41
1	Ad	1531	G	C2'-C1'	10.83	1.65	1.53
1	Ad	1644	C	O4'-C1'	10.82	1.55	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	135	C	C2'-C1'	-10.82	1.41	1.53
1	Ad	1517	C	O4'-C1'	10.81	1.55	1.41
1	Ad	1775	A	O4'-C1'	-10.80	1.27	1.41
1	Ad	1163	C	O4'-C1'	10.80	1.55	1.41
1	Ad	1119	G	C2'-C1'	10.80	1.65	1.53
1	Ad	1673	C	C2'-C1'	-10.80	1.41	1.53
1	Ad	198	G	O4'-C1'	10.79	1.55	1.41
1	Ad	448	C	O4'-C1'	10.79	1.55	1.41
1	Ad	1151	G	C2'-C1'	-10.78	1.41	1.53
1	Ad	249	G	O4'-C1'	10.77	1.55	1.41
1	Ad	1378	C	O4'-C1'	10.76	1.55	1.41
1	Ad	1174	G	O4'-C1'	10.75	1.55	1.41
1	Ad	484	A	O4'-C1'	10.74	1.55	1.41
1	Ad	593	C	O4'-C1'	10.74	1.55	1.41
1	Ad	179	A	C2'-C1'	-10.74	1.41	1.53
1	Ad	993	C	C2'-C1'	-10.72	1.41	1.53
1	Ad	121	U	C2'-C1'	-10.72	1.41	1.53
2	Ae	43	C	O4'-C1'	10.71	1.55	1.41
1	Ad	1567	G	O4'-C1'	-10.71	1.27	1.41
1	Ad	71	C	O4'-C1'	10.70	1.55	1.41
1	Ad	122	U	O4'-C1'	10.69	1.55	1.41
1	Ad	1603	U	O4'-C1'	-10.69	1.27	1.41
1	Ad	1585	A	O4'-C1'	10.67	1.55	1.41
1	Ad	796	U	C2'-C1'	-10.66	1.41	1.53
1	Ad	1219	C	O4'-C1'	10.66	1.55	1.41
1	Ad	380	C	O4'-C1'	10.64	1.55	1.41
1	Ad	539	A	C2'-C1'	-10.64	1.41	1.53
1	Ad	1760	A	C2'-C1'	-10.63	1.41	1.53
1	Ad	1077	C	O4'-C1'	10.63	1.55	1.41
1	Ad	18	C	O4'-C1'	10.61	1.55	1.41
1	Ad	36	C	C2'-C1'	-10.61	1.41	1.53
1	Ad	1193	A	C2'-C1'	10.61	1.65	1.53
1	Ad	17	C	O4'-C1'	10.60	1.55	1.41
1	Ad	1225	A	C2'-C1'	-10.60	1.41	1.53
1	Ad	1718	C	O4'-C1'	10.60	1.55	1.41
1	Ad	465	G	O4'-C1'	10.60	1.55	1.41
1	Ad	1410	C	O4'-C1'	10.60	1.55	1.41
1	Ad	1743	C	O4'-C1'	10.60	1.55	1.41
1	Ad	741	C	O4'-C1'	10.59	1.55	1.41
2	Ae	73	C	C2'-C1'	-10.58	1.41	1.53
1	Ad	296	A	O4'-C1'	10.57	1.55	1.41
1	Ad	1549	G	O4'-C1'	10.56	1.55	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1056	A	C2'-C1'	10.56	1.65	1.53
1	Ad	1096	A	C2'-C1'	-10.56	1.41	1.53
1	Ad	1702	G	C2'-C1'	-10.55	1.41	1.53
2	Ae	38	C	O4'-C1'	10.55	1.55	1.41
1	Ad	879	C	O4'-C1'	10.54	1.55	1.41
1	Ad	86	A	C2'-C1'	-10.53	1.41	1.53
1	Ad	999	G	O4'-C1'	10.53	1.55	1.41
1	Ad	342	C	C2'-C1'	-10.53	1.41	1.53
1	Ad	483	C	C2'-C1'	-10.52	1.41	1.53
1	Ad	1194	C	O4'-C1'	10.52	1.55	1.41
1	Ad	1012	C	O4'-C1'	10.52	1.55	1.41
1	Ad	136	U	C2'-C1'	-10.50	1.41	1.53
1	Ad	600	C	O4'-C1'	10.50	1.55	1.41
1	Ad	1232	G	O4'-C1'	10.50	1.55	1.41
1	Ad	321	C	C2'-C1'	-10.49	1.41	1.53
1	Ad	188	U	C2'-C1'	-10.49	1.41	1.53
1	Ad	183	C	C2'-C1'	-10.49	1.41	1.53
1	Ad	1001	C	O4'-C1'	10.49	1.55	1.41
1	Ad	473	C	O4'-C1'	10.48	1.55	1.41
1	Ad	889	C	C2'-C1'	-10.48	1.41	1.53
1	Ad	1122	U	O4'-C1'	10.48	1.55	1.41
1	Ad	141	G	O4'-C1'	10.48	1.55	1.41
1	Ad	504	C	C2'-C1'	-10.47	1.41	1.53
1	Ad	1751	U	C2'-C1'	-10.47	1.41	1.53
1	Ad	993	C	O4'-C1'	10.47	1.55	1.41
1	Ad	903	A	C2'-C1'	10.44	1.64	1.53
1	Ad	1574	U	C2'-C1'	10.43	1.64	1.53
1	Ad	1104	U	C2'-C1'	10.43	1.64	1.53
1	Ad	1610	C	C2'-C1'	-10.42	1.41	1.53
1	Ad	1304	A	C2'-C1'	10.42	1.64	1.53
1	Ad	148	C	C2'-C1'	-10.42	1.41	1.53
1	Ad	155	A	C2'-C1'	10.40	1.64	1.53
2	Ae	31	C	O4'-C1'	10.40	1.55	1.41
1	Ad	1051	G	C2'-C1'	-10.39	1.42	1.53
1	Ad	326	G	C2'-C1'	-10.39	1.42	1.53
1	Ad	316	A	C2'-C1'	10.37	1.64	1.53
1	Ad	1754	A	O4'-C1'	10.37	1.55	1.41
1	Ad	1708	U	O4'-C1'	10.36	1.55	1.41
1	Ad	533	C	O4'-C1'	10.36	1.55	1.41
1	Ad	1680	A	C2'-C1'	-10.36	1.42	1.53
1	Ad	1755	G	C2'-C1'	-10.36	1.42	1.53
1	Ad	642	C	C2'-C1'	-10.35	1.42	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
2	Ae	36	C	O4'-C1'	10.35	1.55	1.41
1	Ad	453	C	O4'-C1'	10.32	1.55	1.41
1	Ad	1388	A	C2'-C1'	10.32	1.64	1.53
1	Ad	1467	C	C2'-C1'	-10.32	1.42	1.53
1	Ad	301	U	O4'-C1'	10.31	1.55	1.41
1	Ad	415	C	C2'-C1'	-10.31	1.42	1.53
1	Ad	1382	C	O4'-C1'	10.31	1.55	1.41
1	Ad	889	C	O4'-C1'	10.31	1.55	1.41
1	Ad	1355	U	O4'-C1'	10.31	1.55	1.41
1	Ad	1503	C	O4'-C1'	10.30	1.55	1.41
1	Ad	1778	G	O4'-C1'	10.28	1.55	1.41
1	Ad	1007	G	O4'-C1'	10.27	1.55	1.41
1	Ad	1038	C	C2'-C1'	-10.26	1.42	1.53
1	Ad	1562	C	O4'-C1'	10.26	1.54	1.41
1	Ad	1124	G	C2'-C1'	-10.25	1.42	1.53
1	Ad	1678	G	C2'-C1'	-10.23	1.42	1.53
1	Ad	1571	G	C2'-C1'	-10.23	1.42	1.53
1	Ad	4	C	O4'-C1'	10.23	1.54	1.41
1	Ad	1612	C	O4'-C1'	10.22	1.54	1.41
1	Ad	1213	C	C2'-C1'	-10.22	1.42	1.53
1	Ad	798	C	O4'-C1'	10.21	1.54	1.41
1	Ad	936	C	C2'-C1'	-10.20	1.42	1.53
1	Ad	1187	A	C2'-C1'	-10.19	1.42	1.53
1	Ad	1255	U	C2'-C1'	10.19	1.64	1.53
1	Ad	1676	G	C2'-C1'	-10.19	1.42	1.53
1	Ad	376	G	C2'-C1'	-10.18	1.42	1.53
1	Ad	649	C	O4'-C1'	10.17	1.54	1.41
1	Ad	596	A	O4'-C1'	10.16	1.54	1.41
1	Ad	182	C	C2'-C1'	-10.16	1.42	1.53
1	Ad	1369	C	C2'-C1'	-10.16	1.42	1.53
1	Ad	1274	G	C2'-C1'	-10.14	1.42	1.53
1	Ad	735	G	C2'-C1'	-10.13	1.42	1.53
1	Ad	1725	C	C2'-C1'	10.14	1.64	1.53
1	Ad	1615	G	O4'-C1'	10.13	1.54	1.41
1	Ad	430	G	C2'-C1'	-10.12	1.42	1.53
1	Ad	1322	G	C2'-C1'	-10.11	1.42	1.53
1	Ad	714	C	O4'-C1'	10.11	1.54	1.41
1	Ad	810	A	O4'-C1'	10.10	1.54	1.41
1	Ad	1298	G	C2'-C1'	-10.08	1.42	1.53
1	Ad	604	U	C2'-C1'	-10.08	1.42	1.53
1	Ad	469	G	C2'-C1'	-10.07	1.42	1.53
1	Ad	393	G	C2'-C1'	-10.06	1.42	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	919	G	C2'-C1'	10.06	1.64	1.53
1	Ad	1385	C	O4'-C1'	10.05	1.54	1.41
1	Ad	1695	G	O4'-C1'	10.03	1.54	1.41
1	Ad	391	A	O4'-C1'	-10.03	1.28	1.41
1	Ad	404	A	O4'-C1'	-10.03	1.28	1.41
1	Ad	299	A	C2'-C1'	-10.02	1.42	1.53
1	Ad	408	G	C2'-C1'	-10.02	1.42	1.53
84	Aa	1747	A	O3'-P	-10.02	1.49	1.61
1	Ad	1083	C	O4'-C1'	10.02	1.54	1.41
1	Ad	1674	C	O4'-C1'	10.01	1.54	1.41
1	Ad	1282	G	C2'-C1'	-10.01	1.42	1.53
1	Ad	1625	U	C2'-C1'	-10.01	1.42	1.53
1	Ad	760	G	C2'-C1'	-10.00	1.42	1.53
1	Ad	1570	G	C2'-C1'	-10.00	1.42	1.53
1	Ad	313	C	C2'-C1'	-9.97	1.42	1.53
1	Ad	114	U	C2'-C1'	-9.95	1.42	1.53
1	Ad	538	A	O4'-C1'	9.94	1.54	1.41
1	Ad	1168	A	O4'-C1'	9.94	1.54	1.41
1	Ad	394	G	O4'-C1'	-9.94	1.28	1.41
1	Ad	1237	G	C2'-C1'	-9.94	1.42	1.53
1	Ad	435	C	O4'-C1'	9.93	1.54	1.41
1	Ad	526	U	C2'-C1'	-9.93	1.42	1.53
1	Ad	1139	C	O4'-C1'	9.92	1.54	1.41
1	Ad	527	C	O4'-C1'	9.91	1.54	1.41
1	Ad	93	A	C2'-C1'	9.91	1.64	1.53
2	Ae	21	A	O4'-C1'	9.91	1.54	1.41
1	Ad	462	G	C2'-C1'	-9.90	1.42	1.53
1	Ad	576	C	O4'-C1'	9.90	1.54	1.41
2	Ae	17	G	O4'-C1'	9.90	1.54	1.41
1	Ad	453	C	C2'-C1'	-9.89	1.42	1.53
1	Ad	1766	A	O4'-C1'	-9.89	1.28	1.41
1	Ad	914	U	O4'-C1'	9.88	1.54	1.41
1	Ad	630	U	C2'-C1'	-9.88	1.42	1.53
1	Ad	1599	C	C2'-C1'	-9.88	1.42	1.53
1	Ad	371	A	C2'-C1'	-9.88	1.42	1.53
1	Ad	1199	C	C2'-C1'	-9.88	1.42	1.53
1	Ad	1238	A	C2'-C1'	-9.86	1.42	1.53
1	Ad	588	C	C2'-C1'	-9.86	1.42	1.53
1	Ad	318	C	O4'-C1'	9.85	1.54	1.41
1	Ad	1766	A	C2'-C1'	9.85	1.64	1.53
1	Ad	1579	C	C2'-C1'	-9.83	1.42	1.53
86	Ab	46	C	N1-C6	9.83	1.43	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1349	A	C2'-C1'	-9.82	1.42	1.53
1	Ad	1687	G	O4'-C1'	9.81	1.54	1.41
1	Ad	980	C	O4'-C1'	9.81	1.54	1.41
1	Ad	1312	G	C2'-C1'	-9.81	1.42	1.53
1	Ad	1779	U	O4'-C1'	9.80	1.54	1.41
1	Ad	1765	A	O4'-C1'	-9.79	1.28	1.41
2	Ae	50	G	O4'-C1'	-9.79	1.28	1.41
1	Ad	289	G	C2'-C1'	-9.79	1.42	1.53
1	Ad	561	G	C2'-C1'	-9.79	1.42	1.53
2	Ae	75	A	C2'-C1'	9.78	1.64	1.53
1	Ad	229	G	C2'-C1'	-9.77	1.42	1.53
1	Ad	1462	C	C2'-C1'	9.77	1.64	1.53
1	Ad	1187	A	O4'-C1'	9.77	1.54	1.41
1	Ad	1199	C	O4'-C1'	9.77	1.54	1.41
1	Ad	1802	G	O4'-C1'	-9.77	1.28	1.41
1	Ad	550	U	O4'-C1'	9.76	1.54	1.41
1	Ad	1717	C	C2'-C1'	-9.76	1.42	1.53
1	Ad	1410	C	C2'-C1'	-9.76	1.42	1.53
2	Ae	9	A	O4'-C1'	9.75	1.54	1.41
1	Ad	966	U	O4'-C1'	9.75	1.54	1.41
1	Ad	1421	U	C2'-C1'	-9.75	1.42	1.53
1	Ad	165	U	O4'-C1'	9.74	1.54	1.41
1	Ad	187	C	C2'-C1'	-9.74	1.42	1.53
1	Ad	48	G	C2'-C1'	-9.74	1.42	1.53
1	Ad	548	C	C2'-C1'	-9.73	1.42	1.53
1	Ad	152	G	C2'-C1'	-9.73	1.42	1.53
2	Ae	61	C	O4'-C1'	9.73	1.54	1.41
1	Ad	881	G	O4'-C1'	9.73	1.54	1.41
1	Ad	569	C	O4'-C1'	9.72	1.54	1.41
1	Ad	641	C	O4'-C1'	9.72	1.54	1.41
1	Ad	1696	C	C2'-C1'	-9.71	1.42	1.53
1	Ad	1607	C	O4'-C1'	9.71	1.54	1.41
1	Ad	969	U	O4'-C1'	9.70	1.54	1.41
1	Ad	1123	G	C2'-C1'	-9.69	1.42	1.53
1	Ad	34	G	O4'-C1'	9.68	1.54	1.41
1	Ad	1538	C	O4'-C1'	9.68	1.54	1.41
1	Ad	1557	C	C2'-C1'	-9.67	1.42	1.53
1	Ad	1597	C	O4'-C1'	9.67	1.54	1.41
1	Ad	485	A	O4'-C1'	9.66	1.54	1.41
1	Ad	1043	C	O4'-C1'	9.66	1.54	1.41
1	Ad	632	G	O4'-C1'	9.65	1.54	1.41
1	Ad	820	A	C2'-C1'	-9.64	1.42	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1005	C	O4'-C1'	-9.64	1.29	1.41
1	Ad	1624	G	C2'-C1'	-9.64	1.42	1.53
1	Ad	458	A	C2'-C1'	9.63	1.64	1.53
1	Ad	942	C	C2'-C1'	-9.63	1.42	1.53
1	Ad	934	A	C2'-C1'	-9.61	1.42	1.53
1	Ad	537	U	O4'-C1'	9.60	1.54	1.41
1	Ad	1646	G	C2'-C1'	-9.60	1.42	1.53
1	Ad	406	C	C2'-C1'	-9.59	1.42	1.53
1	Ad	190	C	O4'-C1'	9.56	1.54	1.41
1	Ad	1645	C	O4'-C1'	9.55	1.54	1.41
1	Ad	1142	A	O4'-C1'	9.55	1.54	1.41
1	Ad	747	U	O4'-C1'	9.54	1.54	1.41
1	Ad	780	A	O4'-C1'	-9.54	1.29	1.41
1	Ad	235	C	O4'-C1'	9.54	1.54	1.41
1	Ad	1401	C	O4'-C1'	9.53	1.54	1.41
1	Ad	1095	C	O4'-C1'	9.53	1.54	1.41
1	Ad	1579	C	O4'-C1'	9.52	1.54	1.41
1	Ad	1109	U	C2'-C1'	-9.51	1.42	1.53
1	Ad	1354	C	O4'-C1'	9.51	1.54	1.41
1	Ad	97	G	O4'-C1'	9.49	1.53	1.41
1	Ad	1321	C	O4'-C1'	9.49	1.53	1.41
1	Ad	1686	C	O4'-C1'	9.48	1.53	1.41
1	Ad	1578	A	C2'-C1'	-9.48	1.43	1.53
1	Ad	540	C	C2'-C1'	-9.47	1.43	1.53
1	Ad	188	U	O4'-C1'	9.46	1.53	1.41
1	Ad	54	C	C2'-C1'	9.45	1.63	1.53
1	Ad	1305	U	O4'-C1'	9.45	1.53	1.41
86	Ab	24	G	N7-C5	-9.44	1.33	1.39
2	Ae	3	C	O4'-C1'	9.44	1.53	1.41
1	Ad	162	A	C2'-C1'	-9.43	1.43	1.53
1	Ad	317	U	O4'-C1'	9.43	1.53	1.41
1	Ad	1468	G	O4'-C1'	9.42	1.53	1.41
1	Ad	759	A	C2'-C1'	-9.42	1.43	1.53
1	Ad	896	C	O4'-C1'	9.42	1.53	1.41
1	Ad	772	C	C2'-C1'	-9.40	1.43	1.53
1	Ad	1649	C	O4'-C1'	9.40	1.53	1.41
1	Ad	98	C	O4'-C1'	9.40	1.53	1.41
1	Ad	286	C	C2'-C1'	-9.38	1.43	1.53
1	Ad	317	U	C2'-C1'	-9.38	1.43	1.53
1	Ad	1808	U	O4'-C1'	9.37	1.53	1.41
1	Ad	1379	U	O4'-C1'	9.37	1.53	1.41
1	Ad	1531	G	O4'-C1'	-9.36	1.29	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	804	C	C2'-C1'	-9.35	1.43	1.53
1	Ad	1451	G	C2'-C1'	-9.35	1.43	1.53
1	Ad	562	U	C2'-C1'	-9.34	1.43	1.53
1	Ad	821	G	O4'-C1'	9.34	1.53	1.41
1	Ad	277	G	C2'-C1'	-9.34	1.43	1.53
1	Ad	327	A	O4'-C1'	9.34	1.53	1.41
1	Ad	1015	C	O4'-C1'	9.34	1.53	1.41
1	Ad	1469	C	O4'-C1'	9.33	1.53	1.41
1	Ad	419	C	O4'-C1'	9.33	1.53	1.41
1	Ad	309	C	C2'-C1'	-9.33	1.43	1.53
1	Ad	953	G	C2'-C1'	-9.33	1.43	1.53
1	Ad	998	A	O4'-C1'	9.33	1.53	1.41
1	Ad	1520	G	C2'-C1'	-9.33	1.43	1.53
1	Ad	1252	C	O4'-C1'	9.32	1.53	1.41
1	Ad	711	C	O4'-C1'	9.32	1.53	1.41
1	Ad	31	C	C2'-C1'	-9.32	1.43	1.53
1	Ad	1337	C	O4'-C1'	9.32	1.53	1.41
1	Ad	116	G	C2'-C1'	-9.31	1.43	1.53
1	Ad	1418	G	O4'-C1'	9.31	1.53	1.41
1	Ad	1491	C	O4'-C1'	9.31	1.53	1.41
1	Ad	1754	A	C2'-C1'	-9.30	1.43	1.53
1	Ad	427	G	C2'-C1'	-9.30	1.43	1.53
1	Ad	573	C	O4'-C1'	9.29	1.53	1.41
1	Ad	860	A	C2'-C1'	9.29	1.63	1.53
1	Ad	1285	G	C2'-C1'	-9.29	1.43	1.53
1	Ad	116	G	O4'-C1'	9.29	1.53	1.41
1	Ad	278	C	C2'-C1'	-9.29	1.43	1.53
1	Ad	228	G	O4'-C1'	9.28	1.53	1.41
1	Ad	1783	C	O4'-C1'	9.27	1.53	1.41
2	Ae	72	G	O4'-C1'	-9.27	1.29	1.41
1	Ad	69	A	C2'-C1'	-9.27	1.43	1.53
1	Ad	421	A	O4'-C1'	9.26	1.53	1.41
1	Ad	795	A	O4'-C1'	9.26	1.53	1.41
1	Ad	460	G	C2'-C1'	-9.25	1.43	1.53
1	Ad	82	G	C2'-C1'	-9.23	1.43	1.53
86	Ab	40	A	N7-C5	-9.23	1.33	1.39
2	Ae	10	G	O4'-C1'	9.22	1.53	1.41
1	Ad	1059	U	C2'-C1'	-9.22	1.43	1.53
2	Ae	48	C	O4'-C1'	9.21	1.53	1.41
1	Ad	823	A	C2'-C1'	-9.20	1.43	1.53
1	Ad	1075	G	C2'-C1'	-9.20	1.43	1.53
1	Ad	416	A	C2'-C1'	9.18	1.63	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1269	G	C2'-C1'	-9.17	1.43	1.53
1	Ad	1680	A	O4'-C1'	9.17	1.53	1.41
1	Ad	918	G	C2'-C1'	-9.17	1.43	1.53
1	Ad	748	C	O4'-C1'	9.17	1.53	1.41
1	Ad	434	G	C2'-C1'	-9.16	1.43	1.53
1	Ad	254	A	O4'-C1'	9.16	1.53	1.41
1	Ad	278	C	O4'-C1'	9.14	1.53	1.41
1	Ad	1590	U	O4'-C1'	-9.14	1.29	1.41
2	Ae	71	A	C2'-C1'	-9.14	1.43	1.53
1	Ad	1125	U	C2'-C1'	9.13	1.63	1.53
1	Ad	1067	A	O4'-C1'	-9.13	1.29	1.41
1	Ad	1467	C	O4'-C1'	9.12	1.53	1.41
1	Ad	178	A	O4'-C1'	9.12	1.53	1.41
1	Ad	709	C	O4'-C1'	9.12	1.53	1.41
1	Ad	1179	C	O4'-C1'	9.12	1.53	1.41
1	Ad	426	G	C2'-C1'	-9.11	1.43	1.53
1	Ad	225	G	O4'-C1'	9.09	1.53	1.41
1	Ad	1491	C	C2'-C1'	-9.09	1.43	1.53
1	Ad	788	G	O4'-C1'	9.09	1.53	1.41
1	Ad	72	A	O4'-C1'	9.08	1.53	1.41
1	Ad	398	C	O4'-C1'	9.08	1.53	1.41
1	Ad	975	A	C2'-C1'	9.08	1.63	1.53
1	Ad	295	C	O4'-C1'	9.07	1.53	1.41
1	Ad	1562	C	C2'-C1'	-9.07	1.43	1.53
1	Ad	490	G	C2'-C1'	-9.05	1.43	1.53
1	Ad	261	C	O4'-C1'	9.05	1.53	1.41
1	Ad	702	G	C2'-C1'	-9.05	1.43	1.53
1	Ad	631	C	O4'-C1'	9.03	1.53	1.41
1	Ad	728	C	C2'-C1'	-9.03	1.43	1.53
1	Ad	150	U	C2'-C1'	9.03	1.63	1.53
1	Ad	1240	A	C2'-C1'	-9.03	1.43	1.53
1	Ad	406	C	O4'-C1'	9.02	1.53	1.41
1	Ad	1703	G	C2'-C1'	-9.02	1.43	1.53
1	Ad	1202	G	C2'-C1'	-9.00	1.43	1.53
1	Ad	149	G	C2'-C1'	-9.00	1.43	1.53
1	Ad	85	A	O4'-C1'	8.99	1.53	1.41
2	Ae	35	U	O4'-C1'	8.99	1.53	1.41
86	Ab	20	C	N3-C4	8.99	1.40	1.33
1	Ad	1712	C	O4'-C1'	8.98	1.53	1.41
1	Ad	1020	U	O4'-C1'	8.98	1.53	1.41
1	Ad	1614	C	C2'-C1'	-8.98	1.43	1.53
1	Ad	1398	U	O4'-C1'	8.97	1.53	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	965	U	O4'-C1'	8.97	1.53	1.41
2	Ae	72	G	C2'-C1'	8.97	1.63	1.53
1	Ad	474	A	O4'-C1'	8.96	1.53	1.41
1	Ad	727	G	O4'-C1'	8.96	1.53	1.41
1	Ad	558	C	C2'-C1'	-8.96	1.43	1.53
1	Ad	311	G	C2'-C1'	-8.95	1.43	1.53
1	Ad	506	G	O4'-C1'	8.94	1.53	1.41
1	Ad	1083	C	C2'-C1'	-8.94	1.43	1.53
1	Ad	126	U	O4'-C1'	8.92	1.53	1.41
1	Ad	43	A	C2'-C1'	8.91	1.63	1.53
1	Ad	903	A	O4'-C1'	-8.91	1.30	1.41
2	Ae	62	C	C2'-C1'	-8.91	1.43	1.53
1	Ad	1207	A	C2'-C1'	-8.89	1.43	1.53
1	Ad	1646	G	O4'-C1'	8.88	1.53	1.41
1	Ad	1355	U	C2'-C1'	-8.87	1.43	1.53
1	Ad	1166	C	C2'-C1'	-8.86	1.43	1.53
1	Ad	854	C	O4'-C1'	8.86	1.53	1.41
1	Ad	1060	U	C2'-C1'	-8.85	1.43	1.53
1	Ad	1552	U	O4'-C1'	8.85	1.53	1.41
1	Ad	932	C	O4'-C1'	8.84	1.53	1.41
1	Ad	376	G	O4'-C1'	8.84	1.53	1.41
1	Ad	979	A	O4'-C1'	8.84	1.53	1.41
1	Ad	24	U	O4'-C1'	8.84	1.53	1.41
1	Ad	860	A	O4'-C1'	-8.83	1.30	1.41
1	Ad	1442	A	O4'-C1'	8.83	1.53	1.41
1	Ad	1620	C	O4'-C1'	8.83	1.53	1.41
1	Ad	822	G	O4'-C1'	8.82	1.53	1.41
1	Ad	1475	A	O4'-C1'	8.82	1.53	1.41
1	Ad	345	A	O4'-C1'	8.81	1.53	1.41
1	Ad	1331	C	O4'-C1'	8.81	1.53	1.41
1	Ad	1523	A	O4'-C1'	8.81	1.53	1.41
2	Ae	7	A	C2'-C1'	-8.81	1.43	1.53
2	Ae	62	C	O4'-C1'	8.80	1.53	1.41
1	Ad	1113	G	C2'-C1'	-8.79	1.43	1.53
1	Ad	964	U	O4'-C1'	8.78	1.53	1.41
1	Ad	1137	A	O4'-C1'	8.78	1.53	1.41
1	Ad	623	A	C2'-C1'	-8.78	1.43	1.53
1	Ad	759	A	O4'-C1'	8.77	1.53	1.41
1	Ad	540	C	O4'-C1'	8.77	1.53	1.41
1	Ad	992	G	O4'-C1'	8.77	1.53	1.41
86	Ab	31	G	C6-N1	8.75	1.45	1.39
1	Ad	1399	G	C2'-C1'	-8.75	1.43	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1630	G	O4'-C1'	8.73	1.53	1.41
1	Ad	1685	U	O4'-C1'	8.71	1.52	1.41
1	Ad	1737	A	O4'-C1'	8.70	1.52	1.41
1	Ad	447	C	C2'-C1'	-8.70	1.43	1.53
1	Ad	1598	G	O4'-C1'	8.67	1.52	1.41
3	Af	16	G	O4'-C1'	8.66	1.52	1.41
1	Ad	1790	G	C2'-C1'	-8.66	1.43	1.53
1	Ad	344	U	C2'-C1'	8.64	1.62	1.53
1	Ad	1710	C	C2'-C1'	-8.64	1.43	1.53
1	Ad	847	U	C2'-C1'	-8.62	1.43	1.53
1	Ad	1395	C	O4'-C1'	8.61	1.52	1.41
1	Ad	1115	G	C2'-C1'	-8.61	1.43	1.53
1	Ad	23	G	C2'-C1'	-8.61	1.43	1.53
1	Ad	1026	C	O4'-C1'	8.60	1.52	1.41
1	Ad	1076	C	O4'-C1'	8.60	1.52	1.41
1	Ad	1749	C	O4'-C1'	8.60	1.52	1.41
1	Ad	932	C	C2'-C1'	-8.59	1.43	1.53
1	Ad	1033	C	C2'-C1'	-8.59	1.44	1.53
1	Ad	1700	G	O4'-C1'	8.59	1.52	1.41
1	Ad	1578	A	O4'-C1'	8.58	1.52	1.41
1	Ad	1600	G	C2'-C1'	-8.58	1.44	1.53
1	Ad	933	G	C2'-C1'	-8.58	1.44	1.53
1	Ad	1146	G	C2'-C1'	-8.57	1.44	1.53
1	Ad	1280	U	C2'-C1'	8.56	1.62	1.53
1	Ad	232	C	O4'-C1'	8.56	1.52	1.41
1	Ad	783	C	O4'-C1'	8.54	1.52	1.41
1	Ad	1744	C	O4'-C1'	8.54	1.52	1.41
1	Ad	195	A	O4'-C1'	8.53	1.52	1.41
1	Ad	589	A	C2'-C1'	-8.53	1.44	1.53
1	Ad	1633	C	O4'-C1'	8.53	1.52	1.41
1	Ad	1809	U	C2'-C1'	8.53	1.62	1.53
1	Ad	1111	C	O4'-C1'	8.53	1.52	1.41
2	Ae	27	G	C2'-C1'	-8.52	1.44	1.53
1	Ad	218	G	O4'-C1'	8.52	1.52	1.41
1	Ad	215	A	C2'-C1'	8.51	1.62	1.53
1	Ad	224	C	C2'-C1'	8.51	1.62	1.53
1	Ad	1696	C	O4'-C1'	8.51	1.52	1.41
1	Ad	1772	A	C2'-C1'	8.51	1.62	1.53
1	Ad	322	U	O4'-C1'	8.51	1.52	1.41
1	Ad	733	U	O4'-C1'	8.50	1.52	1.41
1	Ad	1797	C	O4'-C1'	8.50	1.52	1.41
1	Ad	1288	C	O4'-C1'	8.50	1.52	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1028	A	O4'-C1'	-8.50	1.30	1.41
2	Ae	35	U	C2'-C1'	-8.49	1.44	1.53
1	Ad	534	C	O4'-C1'	8.49	1.52	1.41
1	Ad	108	C	O4'-C1'	8.49	1.52	1.41
1	Ad	54	C	O4'-C1'	8.49	1.52	1.41
1	Ad	459	C	O4'-C1'	8.49	1.52	1.41
1	Ad	787	C	O4'-C1'	-8.48	1.30	1.41
1	Ad	561	G	O4'-C1'	8.48	1.52	1.41
1	Ad	973	U	O4'-C1'	8.48	1.52	1.41
1	Ad	1457	C	C2'-C1'	-8.48	1.44	1.53
1	Ad	1584	A	O4'-C1'	8.47	1.52	1.41
1	Ad	990	G	C2'-C1'	-8.47	1.44	1.53
1	Ad	1097	A	C2'-C1'	-8.47	1.44	1.53
1	Ad	110	G	C2'-C1'	-8.46	1.44	1.53
1	Ad	186	A	O4'-C1'	8.46	1.52	1.41
1	Ad	1587	G	C2'-C1'	-8.46	1.44	1.53
1	Ad	519	A	O4'-C1'	8.45	1.52	1.41
1	Ad	1672	U	C2'-C1'	-8.45	1.44	1.53
1	Ad	548	C	O4'-C1'	8.45	1.52	1.41
1	Ad	776	A	C2'-C1'	8.44	1.62	1.53
1	Ad	1205	G	C2'-C1'	8.43	1.62	1.53
1	Ad	350	G	O4'-C1'	8.43	1.52	1.41
1	Ad	1026	C	C2'-C1'	-8.42	1.44	1.53
1	Ad	1751	U	O4'-C1'	8.42	1.52	1.41
1	Ad	555	G	C2'-C1'	-8.41	1.44	1.53
1	Ad	68	A	O4'-C1'	8.41	1.52	1.41
1	Ad	57	G	C2'-C1'	-8.38	1.44	1.53
1	Ad	1059	U	O4'-C1'	8.38	1.52	1.41
1	Ad	1370	C	O4'-C1'	8.37	1.52	1.41
1	Ad	66	U	C2'-C1'	8.36	1.62	1.53
1	Ad	1506	G	C2'-C1'	-8.36	1.44	1.53
1	Ad	911	A	C2'-C1'	-8.36	1.44	1.53
1	Ad	646	G	C2'-C1'	-8.35	1.44	1.53
1	Ad	234	G	C2'-C1'	-8.35	1.44	1.53
1	Ad	1308	G	C2'-C1'	-8.34	1.44	1.53
1	Ad	1	U	O4'-C1'	8.34	1.52	1.41
1	Ad	285	G	C2'-C1'	-8.34	1.44	1.53
86	Ab	97	G	N7-C5	-8.33	1.34	1.39
1	Ad	835	U	O4'-C1'	-8.32	1.30	1.41
1	Ad	1543	U	O4'-C1'	8.31	1.52	1.41
1	Ad	1028	A	C2'-C1'	8.31	1.62	1.53
1	Ad	838	U	O4'-C1'	8.31	1.52	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	55	A	C2'-C1'	-8.30	1.44	1.53
1	Ad	1338	U	O4'-C1'	8.30	1.52	1.41
1	Ad	91	C	O4'-C1'	8.30	1.52	1.41
2	Ae	47	U	C2'-C1'	8.30	1.62	1.53
1	Ad	298	C	C2'-C1'	-8.28	1.44	1.53
1	Ad	727	G	C2'-C1'	-8.27	1.44	1.53
1	Ad	1223	A	O4'-C1'	8.27	1.52	1.41
86	Ab	81	G	N7-C5	-8.27	1.34	1.39
1	Ad	1628	C	O4'-C1'	8.26	1.52	1.41
86	Ab	99	G	C6-N1	8.26	1.45	1.39
1	Ad	66	U	O4'-C1'	-8.26	1.30	1.41
1	Ad	1378	C	C2'-C1'	-8.26	1.44	1.53
1	Ad	1094	U	O4'-C1'	8.25	1.52	1.41
1	Ad	1159	G	C2'-C1'	-8.25	1.44	1.53
1	Ad	1601	A	C2'-C1'	-8.25	1.44	1.53
1	Ad	1717	C	O4'-C1'	8.25	1.52	1.41
1	Ad	1727	C	C2'-C1'	-8.24	1.44	1.53
84	Aa	723	G	C2'-C1'	-8.24	1.44	1.53
1	Ad	1405	U	C2'-C1'	-8.24	1.44	1.53
1	Ad	628	G	C2'-C1'	-8.24	1.44	1.53
1	Ad	1164	C	O4'-C1'	8.24	1.52	1.41
2	Ae	1	U	O4'-C1'	8.23	1.52	1.41
1	Ad	1805	U	O4'-C1'	8.23	1.52	1.41
1	Ad	88	C	O4'-C1'	8.22	1.52	1.41
1	Ad	140	C	C2'-C1'	-8.22	1.44	1.53
1	Ad	1734	U	C2'-C1'	-8.22	1.44	1.53
1	Ad	1044	A	C2'-C1'	8.22	1.62	1.53
1	Ad	1524	A	O4'-C1'	-8.21	1.30	1.41
1	Ad	785	A	O4'-C1'	8.21	1.52	1.41
1	Ad	1262	U	C2'-C1'	8.20	1.62	1.53
1	Ad	810	A	C2'-C1'	-8.20	1.44	1.53
1	Ad	1630	G	C2'-C1'	-8.20	1.44	1.53
1	Ad	1401	C	C2'-C1'	-8.19	1.44	1.53
1	Ad	838	U	C2'-C1'	-8.18	1.44	1.53
1	Ad	161	G	O4'-C1'	-8.18	1.31	1.41
1	Ad	1439	G	C2'-C1'	-8.18	1.44	1.53
1	Ad	1594	A	C2'-C1'	8.17	1.62	1.53
1	Ad	343	C	O4'-C1'	8.17	1.52	1.41
1	Ad	530	A	O4'-C1'	8.16	1.52	1.41
1	Ad	1665	U	C2'-C1'	8.16	1.62	1.53
2	Ae	63	C	O4'-C1'	8.16	1.52	1.41
1	Ad	347	C	O4'-C1'	8.16	1.52	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1606	U	C2'-C1'	-8.16	1.44	1.53
2	Ae	51	G	C2'-C1'	-8.16	1.44	1.53
1	Ad	979	A	C2'-C1'	-8.16	1.44	1.53
1	Ad	790	U	C2'-C1'	-8.16	1.44	1.53
1	Ad	572	G	C2'-C1'	-8.15	1.44	1.53
1	Ad	466	G	C2'-C1'	-8.14	1.44	1.53
1	Ad	1368	C	O4'-C1'	-8.14	1.31	1.41
1	Ad	139	U	C2'-C1'	-8.14	1.44	1.53
1	Ad	1774	C	O4'-C1'	8.14	1.52	1.41
2	Ae	37	G	C2'-C1'	-8.14	1.44	1.53
1	Ad	96	G	O4'-C1'	8.13	1.52	1.41
1	Ad	248	U	O4'-C1'	8.13	1.52	1.41
1	Ad	537	U	C2'-C1'	-8.13	1.44	1.53
1	Ad	1062	C	C2'-C1'	8.13	1.62	1.53
1	Ad	1661	C	C2'-C1'	-8.12	1.44	1.53
1	Ad	1804	A	C2'-C1'	8.12	1.62	1.53
1	Ad	339	G	O4'-C1'	8.12	1.52	1.41
1	Ad	56	U	C4'-C3'	8.11	1.62	1.53
84	Aa	1827	U	P-O5'	-8.11	1.51	1.59
1	Ad	1203	G	O4'-C1'	-8.11	1.31	1.41
1	Ad	933	G	O3'-P	-8.11	1.51	1.61
1	Ad	312	C	O4'-C1'	8.11	1.52	1.41
1	Ad	511	U	C2'-C1'	8.10	1.62	1.53
1	Ad	779	C	O4'-C1'	8.10	1.52	1.41
1	Ad	619	A	C2'-C1'	8.10	1.62	1.53
1	Ad	1512	C	O4'-C1'	8.10	1.52	1.41
1	Ad	1720	G	C2'-C1'	-8.10	1.44	1.53
1	Ad	841	U	C2'-C1'	8.09	1.62	1.53
1	Ad	1430	A	C2'-C1'	-8.09	1.44	1.53
1	Ad	577	C	O4'-C1'	8.09	1.52	1.41
1	Ad	515	A	O4'-C1'	8.09	1.52	1.41
1	Ad	1102	U	C2'-C1'	-8.09	1.44	1.53
1	Ad	363	G	C2'-C1'	8.08	1.62	1.53
2	Ae	66	C	O4'-C1'	8.08	1.52	1.41
1	Ad	382	A	O4'-C1'	8.08	1.52	1.41
1	Ad	1402	C	O4'-C1'	8.08	1.52	1.41
1	Ad	1419	U	O4'-C1'	8.08	1.52	1.41
1	Ad	417	U	O4'-C1'	8.07	1.52	1.41
1	Ad	1071	C	O4'-C1'	8.07	1.52	1.41
1	Ad	629	C	C2'-C1'	-8.06	1.44	1.53
1	Ad	24	U	C2'-C1'	-8.06	1.44	1.53
1	Ad	1201	C	O4'-C1'	8.06	1.52	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	868	A	O4'-C1'	8.06	1.52	1.41
1	Ad	863	G	C2'-C1'	-8.05	1.44	1.53
1	Ad	1112	G	C2'-C1'	-8.06	1.44	1.53
1	Ad	1369	C	O4'-C1'	8.05	1.52	1.41
1	Ad	1750	A	C2'-C1'	-8.06	1.44	1.53
1	Ad	1669	U	C2'-C1'	-8.05	1.44	1.53
1	Ad	920	A	O4'-C1'	8.04	1.52	1.41
1	Ad	1433	A	C2'-C1'	8.04	1.62	1.53
1	Ad	1769	C	O4'-C1'	8.04	1.52	1.41
1	Ad	1092	A	C2'-C1'	-8.03	1.44	1.53
1	Ad	1170	G	C2'-C1'	-8.03	1.44	1.53
84	Aa	2354	G	N7-C5	-8.02	1.34	1.39
1	Ad	606	U	O4'-C1'	8.02	1.52	1.41
1	Ad	389	A	O4'-C1'	8.01	1.52	1.41
1	Ad	1050	C	C2'-C1'	-8.01	1.44	1.53
1	Ad	1595	A	O4'-C1'	8.01	1.52	1.41
86	Ab	18	C	N1-C6	8.01	1.42	1.37
1	Ad	147	C	C2'-C1'	-8.01	1.44	1.53
1	Ad	1649	C	C2'-C1'	-8.00	1.44	1.53
1	Ad	554	A	C2'-C1'	-8.00	1.44	1.53
1	Ad	1726	G	C5'-C4'	8.00	1.60	1.51
1	Ad	885	C	C2'-C1'	-8.00	1.44	1.53
1	Ad	1552	U	C2'-C1'	-7.99	1.44	1.53
1	Ad	1652	C	C2'-C1'	-7.99	1.44	1.53
1	Ad	341	G	O4'-C1'	7.99	1.52	1.41
1	Ad	304	A	C2'-C1'	-7.99	1.44	1.53
1	Ad	5	U	C2'-C1'	-7.98	1.44	1.53
1	Ad	990	G	O4'-C1'	7.98	1.52	1.41
1	Ad	1070	A	O4'-C1'	7.97	1.52	1.41
1	Ad	1512	C	C2'-C1'	-7.97	1.44	1.53
1	Ad	277	G	O4'-C1'	7.97	1.52	1.41
1	Ad	1165	A	C2'-C1'	-7.97	1.44	1.53
1	Ad	134	G	O4'-C1'	7.96	1.52	1.41
1	Ad	626	A	O4'-C1'	7.96	1.52	1.41
1	Ad	1413	C	C2'-C1'	-7.96	1.44	1.53
1	Ad	98	C	C2'-C1'	-7.96	1.44	1.53
1	Ad	1111	C	C2'-C1'	-7.96	1.44	1.53
1	Ad	285	G	O4'-C1'	7.96	1.51	1.41
1	Ad	447	C	O4'-C1'	7.96	1.51	1.41
1	Ad	807	G	C2'-C1'	-7.96	1.44	1.53
3	Af	21	C	O4'-C1'	7.95	1.51	1.41
1	Ad	566	G	C2'-C1'	-7.95	1.44	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
86	Ab	23	A	C6-N6	7.94	1.40	1.33
1	Ad	464	A	C2'-C1'	-7.94	1.44	1.53
1	Ad	967	C	C2'-C1'	-7.94	1.44	1.53
1	Ad	1126	C	O4'-C1'	7.93	1.51	1.41
1	Ad	714	C	C2'-C1'	-7.93	1.44	1.53
1	Ad	1171	C	C2'-C1'	-7.93	1.44	1.53
1	Ad	747	U	C2'-C1'	-7.93	1.44	1.53
1	Ad	717	G	C2'-C1'	-7.92	1.44	1.53
1	Ad	1214	C	C2'-C1'	-7.92	1.44	1.53
1	Ad	1664	U	C2'-C1'	7.92	1.62	1.53
1	Ad	1226	U	C2'-C1'	7.92	1.62	1.53
1	Ad	38	C	O4'-C1'	7.92	1.51	1.41
1	Ad	867	A	C2'-C1'	7.91	1.62	1.53
1	Ad	767	G	O4'-C1'	7.91	1.51	1.41
1	Ad	480	U	O4'-C1'	7.91	1.51	1.41
1	Ad	497	U	O4'-C1'	7.91	1.51	1.41
1	Ad	1636	U	C2'-C1'	-7.91	1.44	1.53
1	Ad	401	A	C2'-C1'	-7.90	1.44	1.53
86	Ab	27	A	N7-C5	-7.90	1.34	1.39
1	Ad	1269	G	O4'-C1'	7.89	1.51	1.41
1	Ad	81	U	C2'-C1'	-7.88	1.44	1.53
86	Ab	40	A	C6-N6	7.88	1.40	1.33
1	Ad	1191	U	O4'-C1'	7.88	1.51	1.41
1	Ad	1225	A	O4'-C1'	7.88	1.51	1.41
1	Ad	583	A	C2'-C1'	-7.87	1.44	1.53
1	Ad	1530	G	O4'-C1'	7.87	1.51	1.41
84	Aa	2084	G	O3'-P	-7.87	1.51	1.61
1	Ad	1294	U	O4'-C1'	7.87	1.51	1.41
1	Ad	1303	G	O4'-C1'	7.86	1.51	1.41
1	Ad	1747	A	C5'-C4'	7.84	1.60	1.51
1	Ad	1260	A	O4'-C1'	7.84	1.51	1.41
1	Ad	947	G	C5'-C4'	7.83	1.60	1.51
1	Ad	612	U	O4'-C1'	7.83	1.51	1.41
1	Ad	1207	A	O4'-C1'	7.83	1.51	1.41
1	Ad	937	A	C2'-C1'	7.83	1.61	1.53
1	Ad	1236	U	O4'-C1'	7.82	1.51	1.41
1	Ad	423	G	C2'-C1'	-7.80	1.44	1.53
1	Ad	915	C	C2'-C1'	-7.80	1.44	1.53
1	Ad	14	C	O4'-C1'	7.79	1.51	1.41
1	Ad	421	A	C2'-C1'	7.78	1.61	1.53
86	Ab	87	G	C2-N3	7.77	1.39	1.32
1	Ad	122	U	C2'-C1'	-7.77	1.44	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	629	C	O4'-C1'	7.77	1.51	1.41
1	Ad	450	A	O4'-C1'	7.76	1.51	1.41
1	Ad	1736	C	C2'-C1'	-7.76	1.44	1.53
1	Ad	1359	C	O4'-C1'	7.76	1.51	1.41
1	Ad	1554	G	C2'-C1'	-7.75	1.44	1.53
1	Ad	942	C	O4'-C1'	7.75	1.51	1.41
1	Ad	1492	G	O4'-C1'	-7.75	1.31	1.41
1	Ad	1728	G	C2'-C1'	-7.75	1.44	1.53
1	Ad	436	G	C2'-C1'	-7.75	1.44	1.53
1	Ad	189	U	C2'-C1'	7.75	1.61	1.53
1	Ad	1068	G	C2'-C1'	-7.75	1.44	1.53
1	Ad	1456	U	O4'-C1'	7.75	1.51	1.41
1	Ad	592	U	O4'-C1'	7.74	1.51	1.41
1	Ad	918	G	O4'-C1'	7.74	1.51	1.41
1	Ad	1542	G	O4'-C1'	-7.73	1.31	1.41
1	Ad	1000	A	C2'-C1'	-7.73	1.44	1.53
1	Ad	1789	U	O4'-C1'	7.72	1.51	1.41
1	Ad	1035	A	O3'-P	-7.72	1.51	1.61
1	Ad	1435	G	C2'-C1'	-7.72	1.44	1.53
1	Ad	374	A	C2'-C1'	-7.72	1.44	1.53
1	Ad	323	U	C2'-C1'	-7.70	1.44	1.53
1	Ad	1421	U	O4'-C1'	7.70	1.51	1.41
1	Ad	1640	C	O4'-C1'	7.69	1.51	1.41
1	Ad	1655	U	C2'-C1'	7.69	1.61	1.53
1	Ad	1086	A	O4'-C1'	7.69	1.51	1.41
1	Ad	875	C	O4'-C1'	7.69	1.51	1.41
1	Ad	1553	A	C2'-C1'	-7.68	1.44	1.53
2	Ae	29	C	C2'-C1'	-7.68	1.44	1.53
1	Ad	151	A	C2'-C1'	7.68	1.61	1.53
1	Ad	227	G	C2'-C1'	-7.67	1.45	1.53
1	Ad	292	A	O4'-C1'	7.67	1.51	1.41
1	Ad	923	U	O4'-C1'	7.67	1.51	1.41
1	Ad	1666	G	C2'-C1'	-7.67	1.45	1.53
1	Ad	148	C	O4'-C1'	7.67	1.51	1.41
1	Ad	771	G	C2'-C1'	7.65	1.61	1.53
86	Ab	37	G	C2-N3	7.65	1.38	1.32
1	Ad	1338	U	C2'-C1'	-7.65	1.45	1.53
1	Ad	1008	A	O4'-C1'	-7.64	1.31	1.41
1	Ad	1693	C	O4'-C1'	7.64	1.51	1.41
1	Ad	1339	C	O4'-C1'	7.64	1.51	1.41
1	Ad	210	A	C2'-C1'	-7.63	1.45	1.53
1	Ad	1642	C	O4'-C1'	7.63	1.51	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	514	G	C2'-C1'	-7.63	1.45	1.53
1	Ad	164	C	C2'-C1'	-7.62	1.45	1.53
1	Ad	1362	A	C2'-C1'	-7.62	1.45	1.53
86	Ab	5	G	C2-N3	7.61	1.38	1.32
1	Ad	137	A	O4'-C1'	-7.61	1.31	1.41
1	Ad	1496	A	C2'-C1'	7.61	1.61	1.53
1	Ad	456	A	O4'-C1'	-7.59	1.31	1.41
1	Ad	404	A	C2'-C1'	7.59	1.61	1.53
1	Ad	538	A	C2'-C1'	-7.59	1.45	1.53
1	Ad	1141	U	O4'-C1'	7.59	1.51	1.41
86	Ab	25	G	N1-C2	7.58	1.43	1.37
1	Ad	329	G	C2'-C1'	-7.57	1.45	1.53
1	Ad	167	A	C2'-C1'	7.57	1.61	1.53
1	Ad	1311	U	O4'-C1'	-7.57	1.31	1.41
1	Ad	412	C	C2'-C1'	-7.56	1.45	1.53
1	Ad	1278	C	O4'-C1'	-7.56	1.31	1.41
1	Ad	1398	U	C2'-C1'	-7.55	1.45	1.53
1	Ad	452	C	O4'-C1'	7.55	1.51	1.41
1	Ad	1738	U	C2'-C1'	-7.55	1.45	1.53
1	Ad	1682	U	O4'-C1'	7.54	1.51	1.41
1	Ad	495	C	O4'-C1'	7.54	1.51	1.41
86	Ab	99	G	C2-N3	7.54	1.38	1.32
1	Ad	1111	C	P-O5'	-7.53	1.52	1.59
1	Ad	1364	C	O4'-C1'	7.53	1.51	1.41
1	Ad	273	C	O4'-C1'	7.53	1.51	1.41
1	Ad	1647	C	O4'-C1'	7.53	1.51	1.41
1	Ad	797	A	O4'-C1'	7.52	1.51	1.41
1	Ad	3	C	C2'-C1'	7.52	1.61	1.53
1	Ad	1329	A	C2'-C1'	-7.52	1.45	1.53
2	Ae	52	G	C2'-C1'	-7.52	1.45	1.53
1	Ad	516	A	C2'-C1'	7.50	1.61	1.53
1	Ad	253	C	C2'-C1'	-7.49	1.45	1.53
1	Ad	1286	U	C2'-C1'	-7.49	1.45	1.53
1	Ad	1231	A	O4'-C1'	7.49	1.51	1.41
1	Ad	1592	G	O4'-C1'	-7.49	1.31	1.41
1	Ad	1594	A	O4'-C1'	7.49	1.51	1.41
1	Ad	1446	C	O4'-C1'	7.48	1.51	1.41
1	Ad	458	A	O4'-C1'	-7.47	1.31	1.41
1	Ad	817	C	C2'-C1'	7.47	1.61	1.53
1	Ad	1363	G	C2'-C1'	-7.47	1.45	1.53
3	Af	21	C	C2'-C1'	7.47	1.61	1.53
1	Ad	1230	A	O4'-C1'	7.46	1.51	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
84	Aa	290	C	P-O5'	-7.46	1.52	1.59
1	Ad	624	A	P-O5'	-7.46	1.52	1.59
1	Ad	712	U	C2'-C1'	7.46	1.61	1.53
1	Ad	1155	G	O4'-C1'	7.46	1.51	1.41
1	Ad	599	G	C2'-C1'	-7.46	1.45	1.53
1	Ad	1521	G	O4'-C1'	-7.45	1.31	1.41
1	Ad	386	C	O4'-C1'	7.45	1.51	1.41
1	Ad	414	A	C2'-C1'	-7.45	1.45	1.53
1	Ad	1136	A	O4'-C1'	7.45	1.51	1.41
1	Ad	33	U	C2'-C1'	7.44	1.61	1.53
1	Ad	1365	C	O4'-C1'	7.44	1.51	1.41
1	Ad	1469	C	C2'-C1'	-7.44	1.45	1.53
1	Ad	378	U	O4'-C1'	7.43	1.51	1.41
1	Ad	704	C	C2'-C1'	-7.43	1.45	1.53
1	Ad	1109	U	O4'-C1'	7.43	1.51	1.41
1	Ad	1555	A	O4'-C1'	7.43	1.51	1.41
1	Ad	1688	G	O4'-C1'	-7.43	1.31	1.41
2	Ae	55	C	C2'-C1'	-7.43	1.45	1.53
86	Ab	94	C	C4-C5	7.43	1.48	1.43
1	Ad	1099	G	C2'-C1'	-7.42	1.45	1.53
1	Ad	1623	C	O4'-C1'	7.42	1.51	1.41
84	Aa	2512	U	O3'-P	-7.42	1.52	1.61
2	Ae	68	C	O4'-C1'	7.42	1.51	1.41
1	Ad	858	G	C2'-C1'	-7.42	1.45	1.53
1	Ad	435	C	C2'-C1'	-7.41	1.45	1.53
1	Ad	380	C	C2'-C1'	-7.41	1.45	1.53
1	Ad	244	C	C2'-C1'	7.39	1.61	1.53
1	Ad	1041	A	C2'-C1'	7.39	1.61	1.53
1	Ad	1117	G	O4'-C1'	7.39	1.51	1.41
1	Ad	407	G	O4'-C1'	-7.39	1.32	1.41
1	Ad	1166	C	O4'-C1'	7.39	1.51	1.41
3	Af	20	U	O4'-C1'	7.39	1.51	1.41
1	Ad	960	A	O4'-C1'	7.38	1.51	1.41
1	Ad	117	U	O4'-C1'	7.38	1.51	1.41
1	Ad	596	A	C2'-C1'	-7.37	1.45	1.53
1	Ad	823	A	O3'-P	-7.37	1.52	1.61
86	Ab	10	C	N3-C4	7.37	1.39	1.33
1	Ad	1661	C	O4'-C1'	7.36	1.51	1.41
1	Ad	1362	A	O4'-C1'	7.36	1.51	1.41
1	Ad	308	U	O4'-C1'	7.34	1.51	1.41
1	Ad	1255	U	O4'-C1'	-7.33	1.32	1.41
1	Ad	651	G	O4'-C1'	7.33	1.51	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1244	U	O4'-C1'	7.33	1.51	1.41
1	Ad	1070	A	C2'-C1'	-7.33	1.45	1.53
1	Ad	217	A	C2'-C1'	-7.32	1.45	1.53
1	Ad	735	G	O4'-C1'	7.32	1.51	1.41
1	Ad	1779	U	C2'-C1'	-7.32	1.45	1.53
1	Ad	913	U	O4'-C1'	7.31	1.51	1.41
1	Ad	55	A	O4'-C1'	7.31	1.51	1.41
1	Ad	193	G	O4'-C1'	7.30	1.51	1.41
1	Ad	1364	C	C2'-C1'	-7.30	1.45	1.53
1	Ad	372	U	C2'-C1'	7.30	1.61	1.53
1	Ad	1316	A	O4'-C1'	7.30	1.51	1.41
1	Ad	1477	A	C2'-C1'	7.29	1.61	1.53
1	Ad	828	G	C2'-C1'	-7.29	1.45	1.53
1	Ad	388	G	O4'-C1'	7.29	1.51	1.41
1	Ad	1192	G	C2'-C1'	-7.29	1.45	1.53
1	Ad	368	A	C2'-C1'	7.29	1.61	1.53
1	Ad	1229	C	C2'-C1'	-7.28	1.45	1.53
1	Ad	1035	A	O4'-C1'	7.28	1.51	1.41
1	Ad	336	U	O4'-C1'	-7.27	1.32	1.41
1	Ad	755	U	O4'-C1'	7.27	1.51	1.41
1	Ad	1094	U	C2'-C1'	-7.27	1.45	1.53
1	Ad	1381	G	C2'-C1'	-7.27	1.45	1.53
1	Ad	772	C	O4'-C1'	7.27	1.51	1.41
1	Ad	49	C	C2'-C1'	-7.26	1.45	1.53
2	Ae	20	C	C5'-C4'	7.26	1.60	1.51
2	Ae	64	G	C2'-C1'	-7.26	1.45	1.53
1	Ad	269	A	O4'-C1'	-7.26	1.32	1.41
1	Ad	985	G	C2'-C1'	-7.25	1.45	1.53
1	Ad	1133	C	O4'-C1'	7.25	1.51	1.41
1	Ad	587	C	O4'-C1'	7.25	1.51	1.41
1	Ad	1499	U	C2'-C1'	-7.25	1.45	1.53
84	Aa	87	A	N7-C5	-7.25	1.34	1.39
1	Ad	590	G	O4'-C1'	7.24	1.51	1.41
1	Ad	1760	A	O4'-C1'	7.24	1.51	1.41
1	Ad	409	C	O4'-C1'	7.24	1.51	1.41
1	Ad	708	G	O4'-C1'	7.24	1.51	1.41
1	Ad	448	C	C5'-C4'	7.24	1.60	1.51
1	Ad	1526	C	O4'-C1'	7.24	1.51	1.41
2	Ae	17	G	C2'-C1'	-7.24	1.45	1.53
1	Ad	1453	U	O4'-C1'	7.23	1.51	1.41
2	Ae	59	U	O4'-C1'	7.23	1.51	1.41
1	Ad	706	U	O4'-C1'	7.22	1.51	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1152	A	C2'-C1'	-7.22	1.45	1.53
1	Ad	1660	C	O4'-C1'	7.22	1.51	1.41
1	Ad	517	U	O4'-C1'	7.22	1.51	1.41
86	Ab	107	C	N1-C6	7.22	1.41	1.37
3	Af	12	A	O4'-C1'	-7.22	1.32	1.41
1	Ad	20	G	O4'-C1'	7.22	1.51	1.41
1	Ad	1268	G	O4'-C1'	7.21	1.51	1.41
2	Ae	57	A	C2'-C1'	7.21	1.61	1.53
1	Ad	1040	G	C2'-C1'	-7.21	1.45	1.53
1	Ad	927	A	C2'-C1'	-7.21	1.45	1.53
1	Ad	1194	C	C2'-C1'	7.20	1.61	1.53
1	Ad	1576	C	C3'-C2'	7.20	1.60	1.52
1	Ad	1367	U	C2'-C1'	7.20	1.61	1.53
1	Ad	1605	A	C2'-C1'	-7.20	1.45	1.53
1	Ad	761	A	C2'-C1'	-7.20	1.45	1.53
1	Ad	1152	A	O4'-C1'	7.20	1.51	1.41
1	Ad	1054	G	O4'-C1'	7.19	1.51	1.41
1	Ad	1015	C	C2'-C1'	-7.18	1.45	1.53
2	Ae	66	C	C2'-C1'	-7.18	1.45	1.53
1	Ad	762	A	C2'-C1'	-7.18	1.45	1.53
1	Ad	1157	A	C2'-C1'	-7.18	1.45	1.53
1	Ad	840	U	O4'-C1'	7.16	1.50	1.41
1	Ad	835	U	C2'-C1'	7.16	1.61	1.53
1	Ad	1043	C	C2'-C1'	-7.16	1.45	1.53
1	Ad	333	G	C2'-C1'	-7.16	1.45	1.53
2	Ae	21	A	C2'-C1'	-7.16	1.45	1.53
1	Ad	607	U	C2'-C1'	-7.15	1.45	1.53
1	Ad	1307	U	C2'-C1'	-7.15	1.45	1.53
1	Ad	1524	A	C2'-C1'	7.15	1.61	1.53
1	Ad	239	C	C2'-C1'	-7.14	1.45	1.53
1	Ad	970	U	C2'-C1'	-7.14	1.45	1.53
1	Ad	1177	G	C2'-C1'	-7.14	1.45	1.53
1	Ad	313	C	O4'-C1'	7.14	1.50	1.41
1	Ad	1289	U	O4'-C1'	7.13	1.50	1.41
1	Ad	959	G	O4'-C1'	7.12	1.50	1.41
1	Ad	487	A	C2'-C1'	-7.12	1.45	1.53
1	Ad	1656	C	C2'-C1'	-7.12	1.45	1.53
3	Af	13	A	C2'-C1'	7.12	1.61	1.53
1	Ad	989	G	C2'-C1'	-7.11	1.45	1.53
85	Ac	59	A	N7-C5	-7.11	1.34	1.39
1	Ad	105	A	O4'-C1'	7.11	1.50	1.41
1	Ad	455	G	C2'-C1'	-7.11	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1045	G	C5'-C4'	7.11	1.59	1.51
1	Ad	1411	C	C2'-C1'	-7.11	1.45	1.53
1	Ad	997	A	C2'-C1'	-7.10	1.45	1.53
1	Ad	491	G	C2'-C1'	7.10	1.61	1.53
2	Ae	68	C	C2'-C1'	-7.10	1.45	1.53
1	Ad	223	A	O4'-C1'	7.09	1.50	1.41
1	Ad	1439	G	O4'-C1'	7.09	1.50	1.41
1	Ad	1296	G	O4'-C1'	7.08	1.50	1.41
1	Ad	1567	G	C2'-C1'	7.08	1.61	1.53
1	Ad	1678	G	O4'-C1'	7.08	1.50	1.41
1	Ad	347	C	C2'-C1'	-7.07	1.45	1.53
86	Ab	60	G	C2-N3	7.07	1.38	1.32
1	Ad	1004	U	O4'-C1'	7.07	1.50	1.41
86	Ab	90	A	C6-N1	7.07	1.40	1.35
1	Ad	1793	C	O4'-C1'	7.07	1.50	1.41
1	Ad	1476	C	O4'-C1'	7.06	1.50	1.41
1	Ad	1004	U	C2'-C1'	-7.06	1.45	1.53
1	Ad	465	G	C2'-C1'	-7.06	1.45	1.53
1	Ad	1139	C	C2'-C1'	-7.06	1.45	1.53
1	Ad	1413	C	O4'-C1'	7.05	1.50	1.41
1	Ad	1483	G	C2'-C1'	-7.05	1.45	1.53
1	Ad	1343	C	O4'-C1'	7.05	1.50	1.41
1	Ad	1436	U	O4'-C1'	7.05	1.50	1.41
84	Aa	2415	U	P-O5'	-7.04	1.52	1.59
2	Ae	3	C	C2'-C1'	-7.04	1.45	1.53
1	Ad	1224	C	O4'-C1'	7.04	1.50	1.41
2	Ae	7	A	O4'-C1'	7.03	1.50	1.41
1	Ad	1167	C	C2'-C1'	-7.01	1.45	1.53
1	Ad	978	A	O4'-C1'	7.00	1.50	1.41
1	Ad	207	A	C2'-C1'	-7.00	1.45	1.53
1	Ad	115	A	C2'-C1'	-7.00	1.45	1.53
1	Ad	948	C	C2'-C1'	-7.00	1.45	1.53
1	Ad	1158	G	C2'-C1'	-6.99	1.45	1.53
1	Ad	1720	G	O4'-C1'	6.99	1.50	1.41
1	Ad	230	C	O4'-C1'	6.99	1.50	1.41
1	Ad	305	A	O4'-C1'	6.98	1.50	1.41
1	Ad	1412	A	C2'-C1'	-6.98	1.45	1.53
1	Ad	958	G	C2'-C1'	-6.97	1.45	1.53
1	Ad	1654	C	C2'-C1'	-6.97	1.45	1.53
1	Ad	850	G	O4'-C1'	6.97	1.50	1.41
1	Ad	169	A	C2'-C1'	-6.96	1.45	1.53
1	Ad	327	A	C2'-C1'	-6.96	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	528	U	O4'-C1'	6.96	1.50	1.41
1	Ad	975	A	O4'-C1'	6.96	1.50	1.41
1	Ad	18	C	C2'-C1'	-6.96	1.45	1.53
1	Ad	525	A	C2'-C1'	-6.95	1.45	1.53
1	Ad	1236	U	C2'-C1'	-6.95	1.45	1.53
1	Ad	488	C	C2'-C1'	-6.94	1.45	1.53
84	Aa	2345	C	P-O5'	-6.94	1.52	1.59
1	Ad	546	U	O3'-P	-6.92	1.52	1.61
1	Ad	1677	U	O4'-C1'	6.92	1.50	1.41
1	Ad	1769	C	C5'-C4'	6.92	1.59	1.51
86	Ab	23	A	C6-N1	6.92	1.40	1.35
1	Ad	328	U	C2'-C1'	6.91	1.60	1.53
1	Ad	1767	G	C2'-C1'	-6.90	1.45	1.53
1	Ad	125	A	C2'-C1'	-6.90	1.45	1.53
1	Ad	888	U	C2'-C1'	-6.90	1.45	1.53
1	Ad	1146	G	O4'-C1'	6.89	1.50	1.41
1	Ad	1178	C	O4'-C1'	6.89	1.50	1.41
1	Ad	1741	A	C2'-C1'	-6.89	1.45	1.53
1	Ad	1436	U	C2'-C1'	-6.88	1.45	1.53
1	Ad	154	A	C2'-C1'	-6.88	1.45	1.53
1	Ad	47	A	C2'-C1'	-6.88	1.45	1.53
2	Ae	53	U	O4'-C1'	6.88	1.50	1.41
1	Ad	1384	U	C2'-C1'	6.87	1.60	1.53
1	Ad	1473	C	O4'-C1'	6.87	1.50	1.41
86	Ab	94	C	N3-C4	6.87	1.38	1.33
1	Ad	424	A	O4'-C1'	6.87	1.50	1.41
1	Ad	1556	U	C2'-C1'	6.86	1.60	1.53
1	Ad	127	G	O4'-C1'	6.85	1.50	1.41
1	Ad	1089	A	O4'-C1'	6.85	1.50	1.41
1	Ad	1684	U	O4'-C1'	6.84	1.50	1.41
1	Ad	1133	C	C2'-C1'	-6.84	1.45	1.53
1	Ad	1431	A	C5'-C4'	6.84	1.59	1.51
1	Ad	610	A	C2'-C1'	-6.83	1.45	1.53
1	Ad	1465	C	O4'-C1'	6.83	1.50	1.41
1	Ad	931	A	O4'-C1'	6.83	1.50	1.41
86	Ab	5	G	C2'-C1'	-6.82	1.45	1.53
1	Ad	1452	A	C5'-C4'	6.82	1.59	1.51
1	Ad	477	A	O4'-C1'	6.82	1.50	1.41
1	Ad	1563	A	O4'-C1'	6.81	1.50	1.41
1	Ad	282	C	C3'-C2'	6.81	1.60	1.52
1	Ad	293	C	O4'-C1'	6.81	1.50	1.41
1	Ad	200	C	O4'-C1'	6.81	1.50	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
86	Ab	66	G	N1-C2	6.81	1.43	1.37
1	Ad	1443	U	O4'-C1'	6.80	1.50	1.41
1	Ad	1002	G	O4'-C1'	6.80	1.50	1.41
1	Ad	1498	A	O4'-C1'	-6.79	1.32	1.41
1	Ad	976	A	C2'-C1'	-6.78	1.45	1.53
2	Ae	22	G	O4'-C1'	6.78	1.50	1.41
1	Ad	131	C	C2'-C1'	6.78	1.60	1.53
2	Ae	11	U	C2'-C1'	-6.78	1.45	1.53
86	Ab	58	G	C2'-C1'	-6.78	1.45	1.53
1	Ad	1085	U	C2'-C1'	6.77	1.60	1.53
86	Ab	83	A	C6-N6	6.77	1.39	1.33
1	Ad	479	A	O4'-C1'	6.77	1.50	1.41
1	Ad	165	U	C2'-C1'	-6.75	1.46	1.53
1	Ad	650	G	O4'-C1'	6.75	1.50	1.41
1	Ad	1639	A	O4'-C1'	6.75	1.50	1.41
86	Ab	6	C	N3-C4	6.75	1.38	1.33
1	Ad	266	C	O4'-C1'	6.75	1.50	1.41
86	Ab	85	G	C6-N1	6.74	1.44	1.39
84	Aa	1747	A	P-O5'	-6.74	1.53	1.59
1	Ad	418	C	C2'-C1'	-6.73	1.46	1.53
1	Ad	857	A	C2'-C1'	-6.71	1.46	1.53
1	Ad	1450	A	O4'-C1'	6.71	1.50	1.41
1	Ad	247	A	C2'-C1'	6.70	1.60	1.53
1	Ad	473	C	C2'-C1'	-6.70	1.46	1.53
1	Ad	1023	C	C2'-C1'	-6.70	1.46	1.53
1	Ad	166	A	O4'-C1'	6.70	1.50	1.41
1	Ad	984	A	C5'-C4'	6.70	1.59	1.51
1	Ad	531	A	O4'-C1'	6.70	1.50	1.41
2	Ae	48	C	C2'-C1'	-6.70	1.46	1.53
1	Ad	847	U	O4'-C1'	6.70	1.50	1.41
1	Ad	1350	C	C4'-C3'	6.69	1.60	1.53
1	Ad	226	C	O4'-C1'	6.69	1.50	1.41
1	Ad	470	U	C2'-C1'	-6.69	1.46	1.53
1	Ad	459	C	C2'-C1'	6.68	1.60	1.53
1	Ad	102	U	O4'-C1'	6.68	1.50	1.41
1	Ad	1441	C	O4'-C1'	6.68	1.50	1.41
1	Ad	1513	A	C2'-C1'	6.68	1.60	1.53
1	Ad	963	U	O4'-C1'	6.67	1.50	1.41
1	Ad	1386	U	O4'-C1'	6.67	1.50	1.41
1	Ad	83	U	C2'-C1'	6.66	1.60	1.53
1	Ad	639	G	C2'-C1'	-6.66	1.46	1.53
1	Ad	902	C	C2'-C1'	6.66	1.60	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1136	A	C2'-C1'	-6.66	1.46	1.53
1	Ad	1018	A	C5'-C4'	6.65	1.59	1.51
1	Ad	1338	U	C5'-C4'	6.65	1.59	1.51
1	Ad	1370	C	C2'-C1'	-6.65	1.46	1.53
1	Ad	365	C	O4'-C1'	6.64	1.50	1.41
1	Ad	625	A	P-O5'	-6.64	1.53	1.59
2	Ae	19	U	C5'-C4'	6.64	1.59	1.51
1	Ad	1424	G	C2'-C1'	-6.63	1.46	1.53
84	Aa	2092	C	C2'-C1'	-6.63	1.46	1.53
1	Ad	13	C	O4'-C1'	6.63	1.50	1.41
2	Ae	20	C	O4'-C1'	6.63	1.50	1.41
2	Ae	71	A	O4'-C1'	6.63	1.50	1.41
1	Ad	836	U	C5'-C4'	6.63	1.59	1.51
1	Ad	1741	A	O4'-C1'	6.63	1.50	1.41
1	Ad	1472	G	O4'-C1'	6.62	1.50	1.41
1	Ad	1049	U	O4'-C1'	6.61	1.50	1.41
86	Ab	29	C	C2-N3	6.61	1.41	1.35
1	Ad	307	U	O4'-C1'	6.61	1.50	1.41
1	Ad	959	G	C2'-C1'	-6.61	1.46	1.53
1	Ad	107	U	C2'-C1'	6.60	1.60	1.53
1	Ad	423	G	O4'-C1'	6.60	1.50	1.41
1	Ad	454	U	O4'-C1'	6.60	1.50	1.41
1	Ad	396	G	O4'-C1'	-6.60	1.33	1.41
86	Ab	101	A	C6-N6	6.60	1.39	1.33
1	Ad	1671	G	O3'-P	-6.60	1.53	1.61
1	Ad	732	G	O4'-C1'	-6.59	1.33	1.41
1	Ad	1535	U	O4'-C1'	6.59	1.50	1.41
1	Ad	748	C	C2'-C1'	-6.59	1.46	1.53
1	Ad	926	G	C2'-C1'	-6.58	1.46	1.53
1	Ad	1508	C	O4'-C1'	6.58	1.50	1.41
1	Ad	531	A	C5'-C4'	6.58	1.59	1.51
1	Ad	1077	C	C2'-C1'	-6.57	1.46	1.53
1	Ad	574	A	C2'-C1'	6.57	1.60	1.53
1	Ad	1185	U	O4'-C1'	6.57	1.50	1.41
1	Ad	761	A	C3'-C2'	6.57	1.60	1.52
86	Ab	71	A	C6-N1	6.56	1.40	1.35
1	Ad	163	G	C2'-C1'	-6.56	1.46	1.53
1	Ad	523	C	C2'-C1'	-6.56	1.46	1.53
1	Ad	614	G	C2'-C1'	-6.56	1.46	1.53
1	Ad	1672	U	O4'-C1'	6.56	1.50	1.41
1	Ad	337	A	O4'-C1'	6.55	1.50	1.41
1	Ad	927	A	O4'-C1'	6.55	1.50	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	768	A	O4'-C1'	6.55	1.50	1.41
1	Ad	793	G	C5'-C4'	6.55	1.59	1.51
1	Ad	899	A	O4'-C1'	6.55	1.50	1.41
1	Ad	468	A	O4'-C1'	-6.55	1.33	1.41
1	Ad	1537	U	C2'-C1'	6.54	1.60	1.53
1	Ad	1054	G	C2'-C1'	-6.54	1.46	1.53
86	Ab	76	U	N3-C4	6.53	1.44	1.38
1	Ad	1762	C	C2'-C1'	-6.53	1.46	1.53
1	Ad	1509	C	O4'-C1'	6.53	1.50	1.41
1	Ad	1271	G	O4'-C1'	-6.53	1.33	1.41
1	Ad	825	U	C2'-C1'	-6.52	1.46	1.53
2	Ae	11	U	O4'-C1'	6.52	1.50	1.41
1	Ad	1277	G	C2'-C1'	-6.52	1.46	1.53
1	Ad	741	C	C5'-C4'	6.52	1.59	1.51
1	Ad	1590	U	C2'-C1'	6.52	1.60	1.53
1	Ad	581	G	O3'-P	-6.51	1.53	1.61
1	Ad	113	A	C5'-C4'	6.51	1.59	1.51
1	Ad	133	U	C2'-C1'	-6.51	1.46	1.53
2	Ae	25	U	C2'-C1'	-6.51	1.46	1.53
1	Ad	366	G	O3'-P	-6.50	1.53	1.61
1	Ad	565	G	C2'-C1'	-6.50	1.46	1.53
84	Aa	1747	A	C2'-C1'	-6.50	1.46	1.53
1	Ad	801	U	O4'-C1'	-6.50	1.33	1.41
1	Ad	1167	C	O4'-C1'	6.50	1.50	1.41
1	Ad	586	U	C2'-C1'	-6.50	1.46	1.53
2	Ae	38	C	C2'-C1'	-6.50	1.46	1.53
86	Ab	74	A	C6-N6	6.50	1.39	1.33
1	Ad	1553	A	O4'-C1'	6.50	1.50	1.41
1	Ad	1062	C	O4'-C1'	6.49	1.50	1.41
1	Ad	937	A	P-O5'	-6.49	1.53	1.59
1	Ad	1420	U	C2'-C1'	-6.49	1.46	1.53
1	Ad	1663	A	C2'-C1'	6.49	1.60	1.53
1	Ad	1233	G	O4'-C1'	6.49	1.50	1.41
84	Aa	2374	G	N7-C5	-6.48	1.35	1.39
1	Ad	920	A	C2'-C1'	-6.47	1.46	1.53
1	Ad	1341	G	C2'-C1'	-6.47	1.46	1.53
1	Ad	974	C	O4'-C1'	6.47	1.50	1.41
1	Ad	1019	G	O4'-C1'	-6.47	1.33	1.41
1	Ad	1258	U	O4'-C1'	6.47	1.50	1.41
1	Ad	1358	G	O3'-P	-6.47	1.53	1.61
1	Ad	426	G	C5'-C4'	6.46	1.59	1.51
86	Ab	103	U	N3-C4	6.46	1.44	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
84	Aa	2527	G	N7-C5	-6.46	1.35	1.39
1	Ad	1253	U	O4'-C1'	6.46	1.50	1.41
1	Ad	1415	G	O4'-C1'	6.46	1.50	1.41
86	Ab	11	A	C6-N6	6.45	1.39	1.33
1	Ad	11	A	O4'-C1'	6.45	1.50	1.41
1	Ad	1118	A	C4'-C3'	6.45	1.60	1.53
1	Ad	27	U	O4'-C1'	6.45	1.50	1.41
1	Ad	622	U	C2'-C1'	-6.45	1.46	1.53
1	Ad	1448	U	C2'-C1'	-6.45	1.46	1.53
1	Ad	274	A	O4'-C1'	6.44	1.50	1.41
1	Ad	1703	G	O4'-C1'	6.44	1.50	1.41
86	Ab	74	A	C6-N1	6.44	1.40	1.35
1	Ad	529	A	O4'-C1'	6.44	1.50	1.41
84	Aa	1450	G	N7-C5	-6.44	1.35	1.39
1	Ad	411	A	O4'-C1'	6.44	1.50	1.41
1	Ad	15	U	O4'-C1'	6.43	1.50	1.41
1	Ad	1385	C	C2'-C1'	-6.43	1.46	1.53
1	Ad	357	A	O4'-C1'	6.42	1.50	1.41
1	Ad	402	G	O4'-C1'	6.42	1.50	1.41
84	Aa	1612	C	P-O5'	-6.42	1.53	1.59
1	Ad	1734	U	O4'-C1'	6.42	1.50	1.41
1	Ad	1651	U	C2'-C1'	-6.42	1.46	1.53
84	Aa	2093	G	C6-N1	6.42	1.44	1.39
1	Ad	263	C	C2'-C1'	6.42	1.60	1.53
1	Ad	584	A	C5'-C4'	6.41	1.59	1.51
1	Ad	598	A	O4'-C1'	6.41	1.50	1.41
1	Ad	723	A	C2'-C1'	6.41	1.60	1.53
1	Ad	1186	U	C2'-C1'	6.41	1.60	1.53
1	Ad	873	G	C2'-C1'	6.39	1.60	1.53
1	Ad	490	G	O4'-C1'	6.39	1.50	1.41
1	Ad	1658	U	O4'-C1'	6.39	1.50	1.41
1	Ad	331	U	C2'-C1'	6.39	1.60	1.53
1	Ad	1641	A	O3'-P	-6.39	1.53	1.61
84	Aa	423	C	N3-C4	6.38	1.38	1.33
1	Ad	501	U	C5'-C4'	6.38	1.59	1.51
1	Ad	852	A	O4'-C1'	6.38	1.50	1.41
1	Ad	1651	U	O4'-C1'	6.38	1.50	1.41
1	Ad	306	U	O4'-C1'	6.38	1.50	1.41
1	Ad	1550	G	O4'-C1'	-6.38	1.33	1.41
1	Ad	1723	G	C2'-C1'	-6.38	1.46	1.53
1	Ad	982	A	O4'-C1'	6.37	1.50	1.41
1	Ad	1613	G	O4'-C1'	6.37	1.50	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	252	U	C2'-C1'	6.36	1.60	1.53
84	Aa	3042	U	P-O5'	-6.36	1.53	1.59
1	Ad	1188	A	C2'-C1'	6.36	1.60	1.53
1	Ad	1128	C	C2'-C1'	-6.36	1.46	1.53
1	Ad	378	U	C5'-C4'	6.36	1.58	1.51
1	Ad	1054	G	P-O5'	-6.36	1.53	1.59
1	Ad	1246	A	O4'-C1'	6.35	1.50	1.41
1	Ad	1573	C	O4'-C1'	6.35	1.50	1.41
84	Aa	97	G	N7-C5	-6.35	1.35	1.39
1	Ad	482	A	O4'-C1'	6.35	1.50	1.41
1	Ad	1738	U	O4'-C1'	6.34	1.49	1.41
1	Ad	1713	C	O4'-C1'	6.34	1.49	1.41
1	Ad	54	C	P-O5'	-6.34	1.53	1.59
1	Ad	222	G	C2'-C1'	-6.33	1.46	1.53
1	Ad	1459	G	C2'-C1'	6.33	1.60	1.53
1	Ad	1278	C	C2'-C1'	-6.33	1.46	1.53
1	Ad	844	C	O4'-C1'	6.33	1.49	1.41
84	Aa	978	C	P-O5'	-6.32	1.53	1.59
86	Ab	42	A	C6-N6	6.32	1.39	1.33
1	Ad	802	A	O4'-C1'	6.32	1.49	1.41
1	Ad	1674	C	C2'-C1'	-6.32	1.46	1.53
1	Ad	200	C	C2'-C1'	-6.32	1.46	1.53
1	Ad	767	G	C2'-C1'	-6.32	1.46	1.53
86	Ab	75	G	N9-C8	6.32	1.42	1.37
1	Ad	1037	G	O4'-C1'	6.31	1.49	1.41
86	Ab	17	G	C6-N1	6.31	1.44	1.39
1	Ad	335	A	C5'-C4'	6.31	1.58	1.51
1	Ad	1259	G	C2'-C1'	6.31	1.60	1.53
1	Ad	1265	A	C2'-C1'	-6.31	1.46	1.53
86	Ab	8	A	C6-N6	6.31	1.39	1.33
1	Ad	1340	A	C2'-C1'	-6.30	1.46	1.53
1	Ad	1091	A	O4'-C1'	6.30	1.49	1.41
1	Ad	1106	G	C2'-C1'	-6.30	1.46	1.53
1	Ad	1419	U	C4'-C3'	6.30	1.60	1.53
1	Ad	1578	A	C5'-C4'	6.29	1.58	1.51
1	Ad	428	C	O4'-C1'	6.29	1.49	1.41
86	Ab	115	A	N7-C5	-6.29	1.35	1.39
1	Ad	1368	C	C2'-C1'	6.28	1.60	1.53
1	Ad	1540	U	P-O5'	-6.28	1.53	1.59
1	Ad	797	A	C2'-C1'	-6.28	1.46	1.53
1	Ad	6	G	C2'-C1'	-6.27	1.46	1.53
1	Ad	236	U	C2'-C1'	-6.27	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
84	Aa	2162	C	O3'-P	-6.26	1.53	1.61
1	Ad	1347	U	C2'-C1'	-6.26	1.46	1.53
1	Ad	1135	G	O4'-C1'	6.26	1.49	1.41
1	Ad	1334	G	C2'-C1'	6.25	1.60	1.53
86	Ab	109	U	N1-C6	6.25	1.43	1.38
3	Af	18	C	C2'-C1'	-6.25	1.46	1.53
1	Ad	269	A	C5'-C4'	6.25	1.58	1.51
1	Ad	1719	C	O4'-C1'	6.25	1.49	1.41
1	Ad	1004	U	C4'-C3'	6.24	1.60	1.53
1	Ad	1025	A	C2'-C1'	-6.24	1.46	1.53
1	Ad	1190	U	C2'-C1'	6.24	1.60	1.53
1	Ad	1430	A	C5'-C4'	6.24	1.58	1.51
1	Ad	1685	U	P-O5'	-6.24	1.53	1.59
1	Ad	1631	C	C2'-C1'	-6.23	1.46	1.53
86	Ab	60	G	O3'-P	-6.23	1.53	1.61
1	Ad	645	G	C2'-C1'	-6.22	1.46	1.53
1	Ad	844	C	C2'-C1'	6.22	1.60	1.53
1	Ad	1560	U	C2'-C1'	-6.22	1.46	1.53
1	Ad	1342	C	O4'-C1'	6.22	1.49	1.41
1	Ad	1660	C	C2'-C1'	-6.22	1.46	1.53
1	Ad	564	U	C2'-C1'	-6.21	1.46	1.53
1	Ad	1737	A	C5'-C4'	6.21	1.58	1.51
84	Aa	587	A	N7-C5	-6.21	1.35	1.39
84	Aa	1693	A	N7-C5	-6.21	1.35	1.39
86	Ab	56	G	N1-C2	6.20	1.42	1.37
1	Ad	563	C	C2'-C1'	-6.20	1.46	1.53
1	Ad	422	G	C5'-C4'	6.20	1.58	1.51
1	Ad	266	C	O3'-P	-6.19	1.53	1.61
1	Ad	90	G	C2'-C1'	-6.18	1.46	1.53
1	Ad	224	C	O4'-C1'	6.17	1.49	1.41
1	Ad	233	U	O4'-C1'	6.17	1.49	1.41
1	Ad	744	G	C2'-C1'	6.17	1.60	1.53
1	Ad	876	A	O4'-C1'	6.17	1.49	1.41
1	Ad	1131	G	C2'-C1'	6.17	1.60	1.53
1	Ad	72	A	C2'-C1'	-6.17	1.46	1.53
1	Ad	144	U	O4'-C1'	6.16	1.49	1.41
1	Ad	1785	U	C2'-C1'	-6.16	1.46	1.53
84	Aa	2177	U	O3'-P	-6.16	1.53	1.61
1	Ad	1796	G	O4'-C1'	6.16	1.49	1.41
86	Ab	28	U	C2'-C1'	-6.16	1.46	1.53
1	Ad	1221	A	C2'-C1'	6.16	1.60	1.53
1	Ad	1142	A	O3'-P	-6.15	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	834	A	P-O5'	-6.15	1.53	1.59
1	Ad	887	U	C2'-C1'	6.15	1.60	1.53
1	Ad	264	G	C5'-C4'	6.14	1.58	1.51
1	Ad	293	C	C2'-C1'	-6.14	1.46	1.53
1	Ad	1752	U	C2'-C1'	-6.14	1.46	1.53
86	Ab	50	A	C6-N1	6.14	1.39	1.35
1	Ad	1179	C	C2'-C1'	-6.13	1.46	1.53
86	Ab	109	U	P-O5'	-6.13	1.53	1.59
1	Ad	1571	G	O4'-C1'	6.13	1.49	1.41
86	Ab	93	U	O3'-P	-6.13	1.53	1.61
86	Ab	64	G	C2-N3	6.13	1.37	1.32
1	Ad	444	U	C2'-C1'	6.13	1.60	1.53
1	Ad	221	U	O4'-C1'	6.12	1.49	1.41
1	Ad	547	C	C2'-C1'	6.12	1.60	1.53
1	Ad	853	U	O4'-C1'	6.12	1.49	1.41
1	Ad	104	A	C2'-C1'	6.12	1.60	1.53
1	Ad	851	G	O4'-C1'	-6.12	1.33	1.41
1	Ad	1129	A	C2'-C1'	-6.12	1.46	1.53
1	Ad	1414	G	O4'-C1'	6.12	1.49	1.41
1	Ad	1698	A	C2'-C1'	-6.12	1.46	1.53
1	Ad	857	A	C5'-C4'	6.11	1.58	1.51
1	Ad	1438	U	C2'-C1'	-6.11	1.46	1.53
84	Aa	3234	G	N7-C5	-6.11	1.35	1.39
86	Ab	106	U	C2-N3	6.11	1.42	1.37
1	Ad	1250	C	O4'-C1'	-6.11	1.33	1.41
1	Ad	1536	U	O4'-C1'	6.10	1.49	1.41
1	Ad	1564	A	C2'-C1'	6.10	1.60	1.53
1	Ad	1044	A	O4'-C1'	-6.10	1.33	1.41
1	Ad	152	G	O3'-P	-6.09	1.53	1.61
84	Aa	279	G	C2-N3	6.09	1.37	1.32
1	Ad	730	G	O4'-C1'	-6.09	1.33	1.41
1	Ad	637	U	O4'-C1'	6.09	1.49	1.41
1	Ad	702	G	O4'-C1'	6.09	1.49	1.41
86	Ab	56	G	N7-C5	-6.08	1.35	1.39
1	Ad	1785	U	O4'-C1'	6.08	1.49	1.41
1	Ad	1170	G	C3'-C2'	-6.08	1.46	1.52
84	Aa	1079	G	C2-N3	6.08	1.37	1.32
1	Ad	555	G	O4'-C1'	6.07	1.49	1.41
1	Ad	326	G	O4'-C1'	6.07	1.49	1.41
1	Ad	625	A	C2'-C1'	6.07	1.60	1.53
1	Ad	1115	G	O4'-C1'	6.07	1.49	1.41
84	Aa	1526	A	N7-C5	-6.06	1.35	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1319	U	O4'-C1'	6.06	1.49	1.41
1	Ad	1071	C	C5'-C4'	6.06	1.58	1.51
1	Ad	1245	G	O4'-C1'	-6.06	1.33	1.41
1	Ad	783	C	C2'-C1'	-6.05	1.46	1.53
1	Ad	830	U	C2'-C1'	-6.05	1.46	1.53
1	Ad	957	A	O4'-C1'	6.04	1.49	1.41
1	Ad	843	G	C5'-C4'	6.04	1.58	1.51
1	Ad	494	G	C2'-C1'	6.03	1.59	1.53
1	Ad	1575	U	O3'-P	-6.03	1.53	1.61
84	Aa	3067	G	O3'-P	-6.03	1.53	1.61
1	Ad	1582	G	C2'-C1'	-6.03	1.46	1.53
86	Ab	71	A	C5-C4	6.02	1.43	1.38
86	Ab	26	C	N1-C6	6.02	1.40	1.37
1	Ad	113	A	O4'-C1'	6.01	1.49	1.41
1	Ad	1352	A	C2'-C1'	6.01	1.59	1.53
1	Ad	1408	G	O3'-P	-6.01	1.53	1.61
1	Ad	1768	U	C2'-C1'	-6.01	1.46	1.53
1	Ad	1249	G	C2'-C1'	-6.01	1.46	1.53
84	Aa	289	C	P-O5'	-6.01	1.53	1.59
1	Ad	1222	G	O4'-C1'	6.00	1.49	1.41
1	Ad	700	C	C2'-C1'	-6.00	1.46	1.53
1	Ad	1204	G	P-O5'	-6.00	1.53	1.59
86	Ab	44	C	C4-C5	6.00	1.47	1.43
86	Ab	69	A	N7-C5	-6.00	1.35	1.39
3	Af	15	A	C2'-C1'	-6.00	1.46	1.53
86	Ab	1	G	C6-N1	6.00	1.43	1.39
1	Ad	621	U	O4'-C1'	6.00	1.49	1.41
86	Ab	86	G	C5-C4	6.00	1.42	1.38
1	Ad	1203	G	C5'-C4'	6.00	1.58	1.51
1	Ad	46	A	C2'-C1'	-6.00	1.46	1.53
1	Ad	1679	A	O4'-C1'	5.99	1.49	1.41
84	Aa	1673	A	N7-C5	-5.99	1.35	1.39
1	Ad	1400	G	C2'-C1'	-5.99	1.46	1.53
86	Ab	93	U	C3'-C2'	-5.99	1.46	1.52
1	Ad	1718	C	C2'-C1'	-5.98	1.46	1.53
1	Ad	578	G	O4'-C1'	-5.98	1.33	1.41
1	Ad	1632	C	O3'-P	-5.98	1.53	1.61
86	Ab	57	C	C4-N4	5.98	1.39	1.33
86	Ab	69	A	N3-C4	-5.98	1.31	1.34
1	Ad	594	C	C2'-C1'	-5.98	1.46	1.53
1	Ad	189	U	O4'-C1'	-5.97	1.33	1.41
1	Ad	984	A	C2'-C1'	-5.97	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1569	U	C2'-C1'	5.97	1.59	1.53
86	Ab	58	G	C2-N3	5.97	1.37	1.32
1	Ad	1158	G	O4'-C1'	5.97	1.49	1.41
1	Ad	743	G	O4'-C1'	5.96	1.49	1.41
86	Ab	75	G	C5-C4	5.95	1.42	1.38
1	Ad	1194	C	C5'-C4'	5.95	1.58	1.51
1	Ad	243	U	O4'-C1'	-5.95	1.33	1.41
1	Ad	356	G	C5'-C4'	5.95	1.58	1.51
1	Ad	636	U	O4'-C1'	5.94	1.49	1.41
1	Ad	949	A	C2'-C1'	-5.94	1.46	1.53
1	Ad	1682	U	P-O5'	-5.94	1.53	1.59
2	Ae	65	U	C2'-C1'	-5.94	1.46	1.53
84	Aa	2132	A	N7-C5	-5.94	1.35	1.39
1	Ad	58	U	O4'-C1'	5.94	1.49	1.41
86	Ab	83	A	C6-N1	5.94	1.39	1.35
86	Ab	21	U	N3-C4	5.93	1.43	1.38
1	Ad	1320	C	P-O5'	-5.93	1.53	1.59
1	Ad	29	U	C2'-C1'	5.93	1.59	1.53
1	Ad	304	A	O4'-C1'	5.93	1.49	1.41
1	Ad	1577	A	C2'-C1'	-5.93	1.46	1.53
1	Ad	706	U	C2'-C1'	-5.92	1.46	1.53
84	Aa	860	G	N7-C5	-5.92	1.35	1.39
1	Ad	1791	A	O4'-C1'	5.92	1.49	1.41
1	Ad	1445	C	C2'-C1'	-5.91	1.46	1.53
1	Ad	112	U	C2'-C1'	-5.90	1.46	1.53
1	Ad	400	G	C2'-C1'	5.90	1.59	1.53
1	Ad	325	C	O4'-C1'	-5.90	1.33	1.41
86	Ab	119	C	C4-N4	5.90	1.39	1.33
1	Ad	112	U	O4'-C1'	5.89	1.49	1.41
1	Ad	374	A	O4'-C1'	5.89	1.49	1.41
1	Ad	563	C	O4'-C1'	5.89	1.49	1.41
1	Ad	1505	U	O4'-C1'	5.89	1.49	1.41
1	Ad	145	A	C2'-C1'	5.89	1.59	1.53
84	Aa	512	G	C2-N3	5.89	1.37	1.32
1	Ad	800	U	O3'-P	-5.88	1.54	1.61
86	Ab	71	A	N3-C4	5.88	1.38	1.34
1	Ad	90	G	O4'-C1'	5.88	1.49	1.41
1	Ad	382	A	C2'-C1'	-5.88	1.46	1.53
86	Ab	11	A	N9-C4	5.88	1.41	1.37
1	Ad	1086	A	O3'-P	-5.88	1.54	1.61
1	Ad	1156	A	O4'-C1'	5.87	1.49	1.41
1	Ad	1308	G	O4'-C1'	5.87	1.49	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	265	A	C2'-C1'	-5.87	1.46	1.53
1	Ad	1108	U	O3'-P	-5.87	1.54	1.61
1	Ad	981	G	C2'-C1'	-5.87	1.46	1.53
1	Ad	1638	U	C5'-C4'	5.87	1.58	1.51
2	Ae	69	G	O4'-C1'	5.87	1.49	1.41
1	Ad	1220	C	O3'-P	-5.86	1.54	1.61
1	Ad	1149	U	C2'-C1'	-5.86	1.47	1.53
1	Ad	103	U	O4'-C1'	5.86	1.49	1.41
1	Ad	604	U	O4'-C1'	5.86	1.49	1.41
84	Aa	494	C	C4'-C3'	5.86	1.59	1.53
1	Ad	1205	G	O4'-C1'	-5.86	1.34	1.41
84	Aa	97	G	C2-N3	5.85	1.37	1.32
86	Ab	83	A	C2'-C1'	-5.85	1.47	1.53
2	Ae	2	C	O4'-C1'	5.85	1.49	1.41
1	Ad	574	A	O4'-C1'	5.85	1.49	1.41
2	Ae	28	G	P-O5'	-5.84	1.53	1.59
1	Ad	621	U	C2'-C1'	-5.84	1.47	1.53
1	Ad	1276	U	O4'-C1'	5.84	1.49	1.41
2	Ae	33	U	O4'-C1'	5.84	1.49	1.41
1	Ad	1371	U	C2'-C1'	-5.83	1.47	1.53
1	Ad	1494	G	C5'-C4'	5.83	1.58	1.51
1	Ad	1709	U	O4'-C1'	5.83	1.49	1.41
1	Ad	1536	U	O3'-P	-5.83	1.54	1.61
84	Aa	1188	C	P-O5'	-5.83	1.53	1.59
1	Ad	618	C	O4'-C1'	5.83	1.49	1.41
1	Ad	928	A	C2'-C1'	-5.83	1.47	1.53
1	Ad	1588	C	C2'-C1'	-5.83	1.47	1.53
86	Ab	90	A	N7-C5	-5.83	1.35	1.39
1	Ad	137	A	C5'-C4'	5.82	1.58	1.51
84	Aa	2683	A	N7-C5	-5.82	1.35	1.39
1	Ad	1668	A	O4'-C1'	5.81	1.49	1.41
1	Ad	445	A	P-O5'	-5.80	1.53	1.59
1	Ad	1773	A	C2'-C1'	-5.80	1.47	1.53
1	Ad	1781	U	O4'-C1'	5.80	1.49	1.41
1	Ad	451	U	C2'-C1'	-5.80	1.47	1.53
84	Aa	2183	A	N7-C5	-5.80	1.35	1.39
1	Ad	227	G	O4'-C1'	5.80	1.49	1.41
1	Ad	1075	G	C5'-C4'	5.80	1.58	1.51
1	Ad	1175	G	C2'-C1'	-5.79	1.47	1.53
1	Ad	832	C	C5'-C4'	5.79	1.58	1.51
86	Ab	8	A	N7-C5	-5.79	1.35	1.39
1	Ad	816	U	O4'-C1'	5.79	1.49	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	890	G	O4'-C1'	5.79	1.49	1.41
86	Ab	50	A	N3-C4	-5.78	1.31	1.34
1	Ad	930	G	C2'-C1'	5.78	1.59	1.53
1	Ad	925	U	C2'-C1'	-5.78	1.47	1.53
1	Ad	1079	G	O4'-C1'	5.77	1.49	1.41
86	Ab	15	C	C5'-C4'	5.77	1.58	1.51
1	Ad	782	G	O4'-C1'	5.77	1.49	1.41
84	Aa	2569	G	P-O5'	-5.77	1.53	1.59
1	Ad	19	A	C2'-C1'	5.77	1.59	1.53
1	Ad	1643	A	O4'-C1'	-5.76	1.34	1.41
1	Ad	253	C	C4'-C3'	5.76	1.59	1.53
1	Ad	167	A	O4'-C1'	-5.76	1.34	1.41
1	Ad	870	A	O4'-C1'	-5.76	1.34	1.41
84	Aa	641	C	C4'-C3'	5.76	1.59	1.53
1	Ad	126	U	P-O5'	-5.76	1.53	1.59
1	Ad	1603	U	C2'-C1'	5.76	1.59	1.53
2	Ae	28	G	O4'-C1'	5.76	1.49	1.41
84	Aa	2527	G	C2-N3	5.75	1.37	1.32
1	Ad	98	C	O3'-P	-5.75	1.54	1.61
86	Ab	9	U	N3-C4	5.75	1.43	1.38
1	Ad	602	U	O4'-C1'	5.75	1.49	1.41
1	Ad	1714	G	C2'-C1'	-5.75	1.47	1.53
1	Ad	32	U	C2'-C1'	5.75	1.59	1.53
1	Ad	1009	U	C2'-C1'	5.74	1.59	1.53
1	Ad	401	A	O4'-C1'	5.74	1.49	1.41
84	Aa	721	A	P-O5'	-5.74	1.54	1.59
1	Ad	594	C	O4'-C1'	5.74	1.49	1.41
1	Ad	991	G	C2'-C1'	-5.74	1.47	1.53
1	Ad	6	G	C4'-C3'	5.73	1.59	1.53
1	Ad	104	A	O4'-C1'	-5.73	1.34	1.41
86	Ab	72	G	N7-C5	-5.73	1.35	1.39
84	Aa	2376	G	N1-C2	5.73	1.42	1.37
1	Ad	749	G	C2'-C1'	-5.72	1.47	1.53
2	Ae	72	G	C5'-C4'	5.72	1.58	1.51
84	Aa	1928	A	N7-C5	-5.72	1.35	1.39
1	Ad	377	G	O4'-C1'	5.72	1.49	1.41
1	Ad	1007	G	O3'-P	-5.72	1.54	1.61
1	Ad	1505	U	C2'-C1'	-5.72	1.47	1.53
1	Ad	1407	A	P-O5'	-5.72	1.54	1.59
84	Aa	2162	C	C2'-C1'	-5.72	1.47	1.53
84	Aa	2905	A	N7-C5	-5.71	1.35	1.39
1	Ad	911	A	O4'-C1'	5.71	1.49	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	271	C	O3'-P	-5.71	1.54	1.61
1	Ad	235	C	O3'-P	-5.71	1.54	1.61
1	Ad	745	C	O4'-C1'	5.71	1.49	1.41
86	Ab	99	G	C8-N7	-5.70	1.27	1.30
1	Ad	1484	U	O4'-C1'	5.70	1.49	1.41
1	Ad	1660	C	C5'-C4'	5.70	1.58	1.51
1	Ad	789	C	O4'-C1'	5.70	1.49	1.41
84	Aa	53	C	C2'-C1'	-5.70	1.47	1.53
1	Ad	914	U	P-O5'	-5.70	1.54	1.59
1	Ad	1456	U	C5'-C4'	5.70	1.58	1.51
1	Ad	335	A	O4'-C1'	5.69	1.49	1.41
1	Ad	837	G	C2'-C1'	-5.69	1.47	1.53
1	Ad	1489	A	O4'-C1'	5.69	1.49	1.41
1	Ad	1701	G	C2'-C1'	-5.69	1.47	1.53
1	Ad	1786	A	C3'-C2'	-5.69	1.46	1.52
86	Ab	36	C	N1-C6	5.69	1.40	1.37
1	Ad	83	U	O4'-C1'	5.68	1.49	1.41
1	Ad	566	G	O4'-C1'	5.68	1.49	1.41
1	Ad	780	A	C5'-C4'	5.68	1.58	1.51
85	Ac	5	U	P-O5'	-5.68	1.54	1.59
1	Ad	375	G	C2'-C1'	-5.68	1.47	1.53
1	Ad	1745	U	O4'-C1'	5.68	1.49	1.41
84	Aa	3335	G	C2-N3	5.68	1.37	1.32
1	Ad	750	U	O4'-C1'	5.68	1.49	1.41
1	Ad	928	A	O4'-C1'	5.68	1.49	1.41
1	Ad	1568	U	O4'-C1'	-5.68	1.34	1.41
1	Ad	1382	C	C2'-C1'	-5.67	1.47	1.53
1	Ad	51	A	C2'-C1'	-5.67	1.47	1.53
1	Ad	126	U	C5'-C4'	5.67	1.58	1.51
1	Ad	1000	A	O4'-C1'	5.67	1.49	1.41
1	Ad	1264	U	O4'-C1'	5.67	1.49	1.41
84	Aa	279	G	P-O5'	-5.67	1.54	1.59
1	Ad	101	A	O4'-C1'	5.66	1.49	1.41
1	Ad	1044	A	O3'-P	-5.66	1.54	1.61
1	Ad	987	U	C5'-C4'	5.65	1.58	1.51
1	Ad	1515	G	O4'-C1'	5.65	1.49	1.41
1	Ad	815	A	O4'-C1'	5.65	1.49	1.41
1	Ad	990	G	C4'-C3'	5.65	1.59	1.53
1	Ad	1544	G	O4'-C1'	5.65	1.49	1.41
84	Aa	716	A	N7-C5	-5.65	1.35	1.39
1	Ad	549	A	P-O5'	-5.65	1.54	1.59
1	Ad	1516	C	O4'-C1'	5.65	1.49	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	240	U	O4'-C1'	5.64	1.49	1.41
1	Ad	337	A	C5'-C4'	5.64	1.58	1.51
1	Ad	1604	C	O4'-C1'	-5.64	1.34	1.41
84	Aa	2095	C	O3'-P	-5.64	1.54	1.61
1	Ad	800	U	O4'-C1'	-5.63	1.34	1.41
1	Ad	1101	C	C2'-C1'	-5.63	1.47	1.53
84	Aa	694	U	P-O5'	-5.63	1.54	1.59
1	Ad	1302	C	C2'-C1'	-5.63	1.47	1.53
84	Aa	2196	G	C2-N3	5.63	1.37	1.32
1	Ad	116	G	C5'-C4'	5.63	1.58	1.51
1	Ad	1494	G	O3'-P	-5.63	1.54	1.61
1	Ad	59	G	C2'-C1'	-5.62	1.47	1.53
1	Ad	438	G	O3'-P	-5.62	1.54	1.61
1	Ad	981	G	O4'-C1'	5.61	1.49	1.41
1	Ad	1005	C	C2'-C1'	5.61	1.59	1.53
1	Ad	1209	C	O3'-P	-5.61	1.54	1.61
86	Ab	120	C	N1-C6	5.61	1.40	1.37
1	Ad	251	U	C2'-C1'	5.61	1.59	1.53
1	Ad	463	G	O4'-C1'	5.61	1.49	1.41
1	Ad	731	G	C2'-C1'	5.60	1.59	1.53
84	Aa	198	A	P-O5'	-5.60	1.54	1.59
86	Ab	99	G	C5'-C4'	5.60	1.58	1.51
1	Ad	1668	A	C2'-C1'	-5.60	1.47	1.53
86	Ab	100	A	N9-C4	-5.60	1.34	1.37
1	Ad	1264	U	C2'-C1'	-5.60	1.47	1.53
1	Ad	35	U	P-O5'	-5.60	1.54	1.59
1	Ad	76	U	O4'-C1'	5.60	1.49	1.41
1	Ad	377	G	C2'-C1'	-5.59	1.47	1.53
84	Aa	2354	G	C2-N3	5.59	1.37	1.32
1	Ad	719	C	C2'-C1'	-5.59	1.47	1.53
1	Ad	1153	C	C2'-C1'	-5.58	1.47	1.53
1	Ad	1634	U	O4'-C1'	5.58	1.49	1.41
1	Ad	828	G	O4'-C1'	-5.58	1.34	1.41
1	Ad	1016	C	C2'-C1'	-5.57	1.47	1.53
1	Ad	466	G	C5'-C4'	5.57	1.58	1.51
1	Ad	1705	C	C5'-C4'	5.57	1.58	1.51
1	Ad	184	C	C2'-C1'	-5.57	1.47	1.53
1	Ad	579	C	O4'-C1'	5.57	1.48	1.41
1	Ad	1575	U	C2'-C1'	5.57	1.59	1.53
1	Ad	446	C	O4'-C1'	5.57	1.48	1.41
2	Ae	14	A	C2'-C1'	-5.57	1.47	1.53
1	Ad	449	A	O4'-C1'	5.56	1.48	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1132	G	C4'-C3'	5.55	1.59	1.53
1	Ad	1632	C	O4'-C1'	5.55	1.48	1.41
1	Ad	1206	A	C4'-C3'	5.55	1.59	1.53
1	Ad	1498	A	C4'-O4'	-5.55	1.38	1.45
84	Aa	2266	A	N7-C5	-5.55	1.35	1.39
1	Ad	291	G	O4'-C1'	-5.54	1.34	1.41
86	Ab	107	C	C4-N4	5.54	1.39	1.33
1	Ad	852	A	C2'-C1'	-5.54	1.47	1.53
1	Ad	1056	A	O4'-C1'	-5.54	1.34	1.41
84	Aa	2163	G	C2-N3	5.54	1.37	1.32
1	Ad	1276	U	C2'-C1'	-5.54	1.47	1.53
1	Ad	1634	U	C2'-C1'	-5.54	1.47	1.53
1	Ad	508	U	O4'-C1'	5.53	1.48	1.41
1	Ad	708	G	C2'-C1'	-5.53	1.47	1.53
1	Ad	1724	U	O4'-C1'	5.53	1.48	1.41
85	Ac	70	G	O3'-P	-5.53	1.54	1.61
1	Ad	282	C	O5'-C5'	-5.53	1.33	1.42
84	Aa	1	G	N7-C5	-5.53	1.35	1.39
86	Ab	48	G	N7-C5	-5.53	1.35	1.39
84	Aa	773	G	P-O5'	-5.53	1.54	1.59
84	Aa	2513	U	C5'-C4'	-5.53	1.44	1.51
1	Ad	567	U	O4'-C1'	5.53	1.48	1.41
1	Ad	484	A	C2'-C1'	-5.52	1.47	1.53
1	Ad	1808	U	C5'-C4'	5.52	1.57	1.51
84	Aa	2634	U	P-O5'	-5.52	1.54	1.59
1	Ad	62	A	C5'-C4'	5.52	1.57	1.51
1	Ad	931	A	C5'-C4'	5.52	1.57	1.51
1	Ad	1286	U	O4'-C1'	5.52	1.48	1.41
1	Ad	1136	A	C3'-C2'	-5.52	1.46	1.52
1	Ad	1431	A	O4'-C1'	5.52	1.48	1.41
1	Ad	724	U	C2'-C1'	5.52	1.59	1.53
84	Aa	76	A	C6-N6	5.52	1.38	1.33
84	Aa	1356	G	C2-N3	5.51	1.37	1.32
1	Ad	1645	C	C2'-C1'	5.51	1.59	1.53
1	Ad	1383	U	C2'-C1'	5.51	1.59	1.53
84	Aa	935	U	C2-N3	5.51	1.41	1.37
86	Ab	90	A	C6-N6	5.51	1.38	1.33
1	Ad	200	C	C5'-C4'	5.50	1.57	1.51
1	Ad	905	A	O3'-P	-5.50	1.54	1.61
1	Ad	1235	U	O4'-C1'	5.50	1.48	1.41
84	Aa	2356	A	N7-C5	-5.50	1.35	1.39
84	Aa	2513	U	O5'-C5'	-5.50	1.34	1.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
86	Ab	72	G	O3'-P	-5.50	1.54	1.61
1	Ad	1018	A	O4'-C1'	5.49	1.48	1.41
1	Ad	1242	A	O4'-C1'	5.49	1.48	1.41
2	Ae	44	A	C2'-C1'	5.49	1.59	1.53
84	Aa	2167	G	C4'-O4'	-5.49	1.38	1.45
1	Ad	1479	U	O3'-P	-5.49	1.54	1.61
1	Ad	1570	G	O3'-P	-5.48	1.54	1.61
1	Ad	325	C	C5'-C4'	5.48	1.57	1.51
2	Ae	42	C	C4'-C3'	5.48	1.59	1.53
1	Ad	1121	A	C2'-C1'	-5.48	1.47	1.53
1	Ad	201	G	O4'-C1'	5.47	1.48	1.41
2	Ae	28	G	C5'-C4'	5.47	1.57	1.51
86	Ab	54	A	C2'-C1'	-5.47	1.47	1.53
1	Ad	283	G	P-O5'	-5.47	1.54	1.59
1	Ad	756	U	C2'-C1'	-5.47	1.47	1.53
1	Ad	53	G	O4'-C1'	5.47	1.48	1.41
1	Ad	868	A	C2'-C1'	5.47	1.59	1.53
84	Aa	1177	G	N3-C4	-5.47	1.31	1.35
1	Ad	758	A	O4'-C1'	5.47	1.48	1.41
1	Ad	1362	A	O3'-P	-5.47	1.54	1.61
1	Ad	550	U	C2'-C1'	-5.46	1.47	1.53
1	Ad	830	U	O4'-C1'	5.46	1.48	1.41
1	Ad	1766	A	O3'-P	-5.46	1.54	1.61
1	Ad	571	A	C3'-C2'	-5.46	1.46	1.52
1	Ad	1585	A	C2'-C1'	-5.46	1.47	1.53
1	Ad	1623	C	C2'-C1'	5.46	1.59	1.53
1	Ad	1479	U	O4'-C1'	-5.45	1.34	1.41
84	Aa	475	U	O3'-P	-5.45	1.54	1.61
1	Ad	303	A	C2'-C1'	5.45	1.59	1.53
1	Ad	802	A	C2'-C1'	-5.45	1.47	1.53
1	Ad	79	A	C5'-C4'	5.45	1.57	1.51
1	Ad	766	A	C2'-C1'	-5.45	1.47	1.53
1	Ad	1312	G	O4'-C1'	5.44	1.48	1.41
1	Ad	236	U	O3'-P	-5.44	1.54	1.61
84	Aa	1356	G	N7-C5	-5.44	1.35	1.39
84	Aa	836	G	N7-C5	-5.44	1.35	1.39
84	Aa	307	C	P-O5'	-5.44	1.54	1.59
1	Ad	1180	U	O4'-C1'	5.44	1.48	1.41
84	Aa	2806	A	N7-C5	-5.43	1.35	1.39
1	Ad	756	U	O4'-C1'	5.43	1.48	1.41
1	Ad	954	C	C3'-C2'	-5.43	1.46	1.52
1	Ad	1792	A	O4'-C1'	5.43	1.48	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
84	Aa	1319	U	P-O5'	-5.43	1.54	1.59
86	Ab	3	A	N9-C4	-5.43	1.34	1.37
1	Ad	429	A	O4'-C1'	5.42	1.48	1.41
1	Ad	970	U	O4'-C1'	5.42	1.48	1.41
1	Ad	1183	G	C4'-C3'	5.42	1.59	1.53
1	Ad	811	U	O4'-C1'	5.42	1.48	1.41
84	Aa	2566	C	O3'-P	-5.42	1.54	1.61
84	Aa	2756	G	C2-N3	5.42	1.37	1.32
86	Ab	42	A	N7-C5	-5.42	1.35	1.39
84	Aa	527	G	C2-N3	5.42	1.37	1.32
1	Ad	189	U	O3'-P	-5.41	1.54	1.61
1	Ad	485	A	C2'-C1'	-5.41	1.47	1.53
1	Ad	705	A	O3'-P	-5.41	1.54	1.61
1	Ad	1334	G	O4'-C1'	5.41	1.48	1.41
84	Aa	334	A	C6-N6	5.41	1.38	1.33
86	Ab	22	A	N9-C4	-5.41	1.34	1.37
1	Ad	1377	G	C5'-C4'	5.41	1.57	1.51
1	Ad	1247	G	O4'-C1'	-5.41	1.34	1.41
1	Ad	952	U	O4'-C1'	5.41	1.48	1.41
1	Ad	491	G	O4'-C1'	-5.40	1.34	1.41
86	Ab	3	A	C6-N6	5.40	1.38	1.33
1	Ad	60	C	O4'-C1'	5.40	1.48	1.41
84	Aa	2969	A	N7-C5	-5.40	1.36	1.39
85	Ac	15	G	P-O5'	-5.40	1.54	1.59
1	Ad	624	A	C2'-C1'	5.39	1.59	1.53
1	Ad	1629	U	O4'-C1'	5.39	1.48	1.41
1	Ad	1687	G	C4'-C3'	-5.39	1.47	1.52
1	Ad	405	A	O4'-C1'	5.39	1.48	1.41
1	Ad	739	U	O4'-C1'	5.39	1.48	1.41
1	Ad	1277	G	O4'-C1'	-5.39	1.34	1.41
86	Ab	56	G	C2-N3	5.39	1.37	1.32
84	Aa	2801	A	C6-N6	5.39	1.38	1.33
84	Aa	2502	U	O3'-P	-5.39	1.54	1.61
2	Ae	70	G	C5'-C4'	5.38	1.57	1.51
84	Aa	5	G	C2-N3	5.38	1.37	1.32
84	Aa	2513	U	O3'-P	-5.38	1.54	1.61
84	Aa	3177	A	C6-N6	5.38	1.38	1.33
1	Ad	1418	G	C5'-C4'	5.38	1.57	1.51
1	Ad	1176	A	O4'-C1'	5.38	1.48	1.41
1	Ad	100	C	O4'-C1'	5.38	1.48	1.41
1	Ad	206	U	O3'-P	-5.38	1.54	1.61
1	Ad	794	G	C2'-C1'	-5.37	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1669	U	O4'-C1'	5.37	1.48	1.41
1	Ad	519	A	C4'-C3'	5.37	1.59	1.53
84	Aa	1600	A	P-O5'	-5.37	1.54	1.59
1	Ad	543	G	O4'-C1'	5.37	1.48	1.41
1	Ad	545	A	C2'-C1'	-5.37	1.47	1.53
84	Aa	1317	G	C2'-C1'	-5.37	1.47	1.53
1	Ad	64	U	C2'-C1'	5.37	1.59	1.53
1	Ad	1807	A	C2'-C1'	5.37	1.59	1.53
84	Aa	2552	U	O3'-P	-5.37	1.54	1.61
1	Ad	323	U	O4'-C1'	5.37	1.48	1.41
1	Ad	778	G	O4'-C1'	-5.37	1.34	1.41
84	Aa	1309	U	O3'-P	-5.37	1.54	1.61
84	Aa	2610	G	O3'-P	-5.36	1.54	1.61
1	Ad	1143	A	O4'-C1'	5.36	1.48	1.41
1	Ad	99	U	O4'-C1'	5.36	1.48	1.41
1	Ad	901	U	C2'-C1'	-5.36	1.47	1.53
1	Ad	139	U	P-O5'	-5.36	1.54	1.59
1	Ad	405	A	C2'-C1'	-5.36	1.47	1.53
1	Ad	529	A	C2'-C1'	-5.35	1.47	1.53
1	Ad	1182	C	P-O5'	-5.35	1.54	1.59
1	Ad	1389	G	C2'-C1'	-5.34	1.47	1.53
1	Ad	1442	A	O3'-P	-5.34	1.54	1.61
1	Ad	1642	C	C2'-C1'	-5.34	1.47	1.53
84	Aa	2290	A	N7-C5	-5.34	1.36	1.39
1	Ad	1091	A	C4'-C3'	5.34	1.59	1.53
1	Ad	1309	U	O4'-C1'	-5.34	1.34	1.41
84	Aa	567	G	N7-C5	-5.34	1.36	1.39
84	Aa	87	A	C6-N6	5.33	1.38	1.33
1	Ad	1805	U	C5'-C4'	5.33	1.57	1.51
86	Ab	15	C	N3-C4	5.33	1.37	1.33
84	Aa	747	A	N7-C5	-5.33	1.36	1.39
1	Ad	418	C	O4'-C1'	5.32	1.48	1.41
1	Ad	479	A	C2'-C1'	-5.32	1.47	1.53
1	Ad	824	U	P-O5'	-5.32	1.54	1.59
1	Ad	1546	U	O4'-C1'	5.32	1.48	1.41
1	Ad	1132	G	P-O5'	-5.32	1.54	1.59
1	Ad	1525	U	C4'-C3'	5.32	1.59	1.53
84	Aa	73	A	N7-C5	-5.32	1.36	1.39
84	Aa	2216	G	C6-N1	5.32	1.43	1.39
1	Ad	630	U	O4'-C1'	5.32	1.48	1.41
1	Ad	790	U	O4'-C1'	5.32	1.48	1.41
84	Aa	144	A	N7-C5	-5.32	1.36	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	500	G	O3'-P	-5.31	1.54	1.61
2	Ae	4	G	P-O5'	-5.31	1.54	1.59
2	Ae	8	U	C2'-C1'	5.31	1.59	1.53
86	Ab	108	G	C2-N3	5.31	1.36	1.32
1	Ad	1805	U	C2'-C1'	-5.31	1.47	1.53
84	Aa	2178	G	O3'-P	-5.31	1.54	1.61
1	Ad	1576	C	C2'-C1'	5.31	1.59	1.53
86	Ab	27	A	N9-C8	5.31	1.42	1.37
1	Ad	338	G	O4'-C1'	-5.31	1.34	1.41
84	Aa	3135	A	N7-C5	-5.31	1.36	1.39
84	Aa	935	U	O3'-P	-5.30	1.54	1.61
84	Aa	2178	G	C3'-O3'	-5.30	1.34	1.42
1	Ad	318	C	O3'-P	-5.29	1.54	1.61
1	Ad	1478	C	O4'-C1'	5.29	1.48	1.41
1	Ad	493	C	O4'-C1'	5.29	1.48	1.41
84	Aa	267	G	N7-C5	-5.29	1.36	1.39
1	Ad	1738	U	P-O5'	-5.29	1.54	1.59
86	Ab	119	C	C3'-C2'	-5.29	1.47	1.52
84	Aa	1747	A	N7-C5	-5.29	1.36	1.39
86	Ab	93	U	N1-C2	5.29	1.43	1.38
1	Ad	1437	C	C4'-C3'	5.29	1.58	1.53
84	Aa	3177	A	N7-C5	-5.29	1.36	1.39
86	Ab	52	U	C2'-C1'	-5.28	1.47	1.53
1	Ad	752	A	O4'-C1'	5.28	1.48	1.41
1	Ad	527	C	C2'-C1'	-5.28	1.47	1.53
1	Ad	1295	G	C2'-C1'	-5.28	1.47	1.53
1	Ad	642	C	C3'-C2'	5.28	1.58	1.52
84	Aa	1310	G	N7-C5	-5.28	1.36	1.39
86	Ab	26	C	P-O5'	-5.28	1.54	1.59
1	Ad	1711	G	C2'-C1'	-5.27	1.47	1.53
84	Aa	2163	G	C2'-C1'	-5.27	1.47	1.53
1	Ad	160	A	C2'-C1'	-5.27	1.47	1.53
1	Ad	327	A	O3'-P	-5.27	1.54	1.61
86	Ab	111	U	C2-N3	5.27	1.41	1.37
1	Ad	1265	A	O4'-C1'	5.27	1.48	1.41
84	Aa	1250	G	C2'-C1'	-5.27	1.47	1.53
1	Ad	23	G	O3'-P	-5.27	1.54	1.61
1	Ad	168	U	C5'-C4'	5.26	1.57	1.51
1	Ad	775	A	O4'-C1'	5.26	1.48	1.41
86	Ab	12	U	C2-N3	5.26	1.41	1.37
1	Ad	1608	A	O4'-C1'	5.26	1.48	1.41
1	Ad	1786	A	O4'-C1'	5.26	1.48	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1251	U	O4'-C1'	5.26	1.48	1.41
1	Ad	1347	U	O4'-C1'	5.26	1.48	1.41
84	Aa	1241	G	O3'-P	-5.26	1.54	1.61
86	Ab	30	G	C6-N1	5.26	1.43	1.39
1	Ad	867	A	P-O5'	-5.25	1.54	1.59
1	Ad	240	U	C5'-C4'	5.25	1.57	1.51
1	Ad	266	C	C5'-C4'	5.25	1.57	1.51
1	Ad	827	C	O4'-C1'	5.25	1.48	1.41
1	Ad	9	U	O4'-C1'	5.25	1.48	1.41
1	Ad	1411	C	O4'-C1'	5.25	1.48	1.41
84	Aa	553	C	O3'-P	-5.25	1.54	1.61
1	Ad	926	G	O4'-C1'	5.25	1.48	1.41
1	Ad	1522	U	O4'-C1'	-5.25	1.34	1.41
84	Aa	294	A	N7-C5	-5.25	1.36	1.39
84	Aa	252	A	N7-C5	-5.24	1.36	1.39
1	Ad	8	U	C2'-C1'	5.24	1.59	1.53
1	Ad	951	U	O4'-C1'	5.24	1.48	1.41
1	Ad	539	A	O4'-C1'	5.24	1.48	1.41
1	Ad	545	A	O4'-C1'	-5.24	1.34	1.41
84	Aa	2239	A	N7-C5	-5.24	1.36	1.39
86	Ab	108	G	C5-C6	-5.24	1.37	1.42
1	Ad	1036	U	O4'-C1'	5.24	1.48	1.41
1	Ad	1799	G	C2'-C1'	-5.24	1.47	1.53
84	Aa	372	A	O3'-P	-5.23	1.54	1.61
84	Aa	2943	A	N7-C5	-5.23	1.36	1.39
1	Ad	1584	A	P-O5'	-5.23	1.54	1.59
1	Ad	1771	U	C2'-C1'	-5.23	1.47	1.53
86	Ab	64	G	C5-C4	5.23	1.42	1.38
1	Ad	131	C	O3'-P	-5.23	1.54	1.61
1	Ad	764	U	O4'-C1'	5.23	1.48	1.41
86	Ab	26	C	N3-C4	5.23	1.37	1.33
1	Ad	1114	G	O4'-C1'	5.23	1.48	1.41
1	Ad	1029	U	O4'-C1'	5.23	1.48	1.41
84	Aa	99	A	N7-C5	-5.23	1.36	1.39
84	Aa	2502	U	C2'-C1'	-5.23	1.47	1.53
1	Ad	215	A	O3'-P	-5.22	1.54	1.61
1	Ad	864	A	C5'-C4'	5.22	1.57	1.51
84	Aa	1062	G	P-O5'	-5.22	1.54	1.59
85	Ac	85	G	C2-N3	5.22	1.36	1.32
1	Ad	205	U	P-O5'	-5.22	1.54	1.59
84	Aa	1676	A	N7-C5	-5.22	1.36	1.39
1	Ad	1446	C	O3'-P	-5.22	1.54	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	1795	U	O4'-C1'	5.22	1.48	1.41
1	Ad	129	U	C2'-C1'	5.21	1.59	1.53
1	Ad	1078	G	C2'-C1'	-5.21	1.47	1.53
1	Ad	597	U	O4'-C1'	5.21	1.48	1.41
1	Ad	1217	G	O4'-C1'	-5.21	1.34	1.41
1	Ad	1729	A	C5'-C4'	5.21	1.57	1.51
84	Aa	819	A	O3'-P	-5.21	1.54	1.61
1	Ad	894	U	O3'-P	-5.21	1.54	1.61
2	Ae	2	C	C2'-C1'	-5.21	1.47	1.53
86	Ab	1	G	N7-C5	-5.21	1.36	1.39
1	Ad	1021	C	C2'-C1'	-5.21	1.47	1.53
1	Ad	1202	G	C5'-C4'	5.21	1.57	1.51
84	Aa	610	G	N7-C5	-5.21	1.36	1.39
1	Ad	597	U	C2'-C1'	-5.20	1.47	1.53
84	Aa	265	G	C2-N3	5.20	1.36	1.32
1	Ad	1314	U	O4'-C1'	5.20	1.48	1.41
1	Ad	811	U	C2'-C1'	-5.20	1.47	1.53
1	Ad	1337	C	C2'-C1'	-5.20	1.47	1.53
1	Ad	611	G	P-O5'	-5.19	1.54	1.59
84	Aa	1486	G	C2-N3	5.19	1.36	1.32
1	Ad	150	U	O3'-P	-5.19	1.54	1.61
1	Ad	793	G	O4'-C1'	5.19	1.48	1.41
84	Aa	1649	G	C2-N3	5.19	1.36	1.32
1	Ad	262	U	O3'-P	-5.19	1.54	1.61
1	Ad	909	G	C2'-C1'	-5.19	1.47	1.53
85	Ac	150	G	C2'-C1'	-5.18	1.47	1.53
1	Ad	1025	A	O4'-C1'	5.18	1.48	1.41
1	Ad	1296	G	O3'-P	-5.18	1.54	1.61
1	Ad	1098	A	P-O5'	5.18	1.65	1.59
1	Ad	1262	U	O4'-C1'	-5.18	1.34	1.41
1	Ad	845	C	O3'-P	-5.18	1.54	1.61
1	Ad	1360	G	C2'-C1'	5.18	1.59	1.53
1	Ad	1528	U	C2'-C1'	5.18	1.59	1.53
1	Ad	542	A	C4'-C3'	5.18	1.58	1.53
1	Ad	874	A	O4'-C1'	5.17	1.48	1.41
1	Ad	1116	G	O4'-C1'	-5.17	1.34	1.41
84	Aa	2968	G	O3'-P	-5.17	1.54	1.61
86	Ab	19	A	N3-C4	-5.17	1.31	1.34
1	Ad	211	G	O4'-C1'	5.17	1.48	1.41
1	Ad	517	U	C5'-C4'	5.17	1.57	1.51
1	Ad	945	A	C5'-C4'	5.17	1.57	1.51
1	Ad	1307	U	O4'-C1'	5.17	1.48	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	318	C	C2'-C1'	-5.17	1.47	1.53
1	Ad	1064	U	O3'-P	-5.17	1.54	1.61
1	Ad	898	U	O4'-C1'	5.16	1.48	1.41
1	Ad	1527	U	C2'-C1'	5.16	1.59	1.53
1	Ad	1658	U	C2'-C1'	-5.16	1.47	1.53
1	Ad	728	C	O3'-P	-5.16	1.54	1.61
1	Ad	1198	A	O4'-C1'	5.16	1.48	1.41
1	Ad	123	U	P-O5'	-5.16	1.54	1.59
1	Ad	446	C	C2'-C1'	-5.16	1.47	1.53
84	Aa	2093	G	C2-N3	5.16	1.36	1.32
86	Ab	51	G	C3'-C2'	5.16	1.58	1.52
1	Ad	838	U	C5'-C4'	5.16	1.57	1.51
1	Ad	1309	U	C2'-C1'	5.16	1.59	1.53
1	Ad	20	G	C5'-C4'	5.16	1.57	1.51
1	Ad	379	U	C2'-C1'	5.16	1.59	1.53
1	Ad	1155	G	C5'-C4'	5.16	1.57	1.51
1	Ad	1546	U	C2'-C1'	5.15	1.59	1.53
84	Aa	2494	A	N7-C5	-5.15	1.36	1.39
84	Aa	2523	G	O3'-P	-5.15	1.54	1.61
1	Ad	143	A	O3'-P	-5.15	1.54	1.61
1	Ad	732	G	C2'-C1'	5.15	1.59	1.53
84	Aa	3364	A	P-O5'	-5.15	1.54	1.59
1	Ad	231	U	O4'-C1'	5.15	1.48	1.41
84	Aa	1958	G	O3'-P	-5.15	1.54	1.61
86	Ab	114	C	N1-C6	5.15	1.40	1.37
1	Ad	726	G	C2'-C1'	-5.14	1.47	1.53
1	Ad	564	U	O4'-C1'	5.14	1.48	1.41
84	Aa	723	G	C4'-C3'	5.14	1.58	1.53
84	Aa	16	A	P-O5'	-5.14	1.54	1.59
84	Aa	641	C	C5'-C4'	5.14	1.57	1.51
1	Ad	133	U	O3'-P	-5.14	1.54	1.61
1	Ad	469	G	O4'-C1'	5.13	1.48	1.41
1	Ad	586	U	O4'-C1'	5.13	1.48	1.41
1	Ad	1320	C	O4'-C1'	5.13	1.48	1.41
1	Ad	161	G	C2'-C1'	5.13	1.58	1.53
84	Aa	1677	G	C2-N3	5.13	1.36	1.32
84	Aa	2374	G	N1-C2	5.13	1.41	1.37
1	Ad	536	U	O3'-P	-5.13	1.54	1.61
1	Ad	1440	U	C5'-C4'	5.13	1.57	1.51
84	Aa	579	G	C2-N3	5.12	1.36	1.32
86	Ab	71	A	N7-C5	-5.12	1.36	1.39
86	Ab	82	G	N9-C4	-5.12	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	522	A	C2'-C1'	-5.12	1.47	1.53
1	Ad	1011	C	O4'-C1'	5.12	1.48	1.41
1	Ad	1605	A	O4'-C1'	5.12	1.48	1.41
84	Aa	752	U	P-O5'	-5.12	1.54	1.59
1	Ad	440	A	C2'-C1'	5.12	1.58	1.53
86	Ab	47	C	N3-C4	5.12	1.37	1.33
1	Ad	1067	A	C2'-C1'	5.12	1.58	1.53
1	Ad	1356	A	C2'-C1'	-5.12	1.47	1.53
84	Aa	1945	A	O3'-P	-5.12	1.55	1.61
1	Ad	1493	A	C4'-C3'	5.11	1.58	1.53
1	Ad	1527	U	O4'-C1'	5.11	1.48	1.41
1	Ad	1544	G	C2'-C1'	-5.11	1.47	1.53
1	Ad	1770	G	C2'-C1'	-5.11	1.47	1.53
84	Aa	1274	A	N7-C5	-5.11	1.36	1.39
1	Ad	518	G	O3'-P	-5.11	1.55	1.61
84	Aa	2393	G	C2'-C1'	-5.11	1.47	1.53
1	Ad	1022	U	C2'-C1'	-5.11	1.47	1.53
86	Ab	38	U	C2'-C1'	-5.11	1.47	1.53
1	Ad	1132	G	C2'-C1'	5.11	1.58	1.53
1	Ad	235	C	C5'-C4'	5.10	1.57	1.51
1	Ad	1153	C	O4'-C1'	5.10	1.48	1.41
84	Aa	1674	A	C6-N6	5.10	1.38	1.33
1	Ad	1118	A	C2'-C1'	-5.10	1.47	1.53
1	Ad	279	C	C2'-C1'	5.10	1.58	1.53
1	Ad	532	U	O4'-C1'	5.10	1.48	1.41
1	Ad	833	U	C2'-C1'	-5.10	1.47	1.53
1	Ad	1481	A	O4'-C1'	5.10	1.48	1.41
1	Ad	1068	G	C5'-C4'	5.10	1.57	1.51
1	Ad	1228	G	O4'-C1'	5.10	1.48	1.41
1	Ad	1464	G	O4'-C1'	5.10	1.48	1.41
86	Ab	87	G	C6-N1	5.09	1.43	1.39
1	Ad	348	A	O4'-C1'	5.09	1.48	1.41
1	Ad	995	C	O3'-P	-5.09	1.55	1.61
1	Ad	1708	U	C2'-C1'	-5.09	1.47	1.53
84	Aa	1796	A	N7-C5	-5.09	1.36	1.39
1	Ad	925	U	O4'-C1'	5.09	1.48	1.41
1	Ad	298	C	O3'-P	-5.09	1.55	1.61
2	Ae	5	U	P-O5'	-5.09	1.54	1.59
84	Aa	2770	U	P-O5'	-5.09	1.54	1.59
86	Ab	6	C	O3'-P	-5.09	1.55	1.61
1	Ad	414	A	O4'-C1'	5.09	1.48	1.41
84	Aa	1676	A	C6-N6	5.09	1.38	1.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
84	Aa	2291	A	N7-C5	-5.09	1.36	1.39
84	Aa	2503	A	C6-N6	5.09	1.38	1.33
1	Ad	1017	U	O3'-P	-5.08	1.55	1.61
1	Ad	1350	C	P-O5'	5.08	1.64	1.59
84	Aa	1600	A	N7-C5	-5.08	1.36	1.39
84	Aa	2307	A	N7-C5	-5.08	1.36	1.39
86	Ab	86	G	C6-N1	5.08	1.43	1.39
1	Ad	414	A	C3'-C2'	-5.08	1.47	1.52
1	Ad	648	C	O3'-P	-5.08	1.55	1.61
2	Ae	13	U	O4'-C1'	5.08	1.48	1.41
1	Ad	532	U	C4'-C3'	5.08	1.58	1.53
84	Aa	936	A	N7-C5	-5.08	1.36	1.39
84	Aa	122	A	N7-C5	-5.08	1.36	1.39
84	Aa	2174	C	C2'-C1'	-5.08	1.47	1.53
84	Aa	2751	A	N7-C5	-5.08	1.36	1.39
1	Ad	10	G	O4'-C1'	5.08	1.48	1.41
2	Ae	57	A	C5'-C4'	5.08	1.57	1.51
84	Aa	293	A	N7-C5	-5.08	1.36	1.39
2	Ae	5	U	C2'-C1'	5.07	1.58	1.53
86	Ab	24	G	P-O5'	-5.07	1.54	1.59
1	Ad	1182	C	O4'-C1'	5.07	1.48	1.41
84	Aa	1486	G	N7-C5	-5.07	1.36	1.39
86	Ab	98	G	N1-C2	5.07	1.41	1.37
1	Ad	941	G	C2'-C1'	-5.07	1.47	1.53
1	Ad	1194	C	O3'-P	-5.07	1.55	1.61
84	Aa	1310	G	C2-N3	5.07	1.36	1.32
86	Ab	48	G	N1-C2	5.07	1.41	1.37
1	Ad	487	A	P-O5'	-5.07	1.54	1.59
1	Ad	1218	U	C2'-C1'	5.06	1.58	1.53
1	Ad	815	A	P-O5'	-5.06	1.54	1.59
1	Ad	489	C	O4'-C1'	5.06	1.48	1.41
1	Ad	89	U	O4'-C1'	5.06	1.48	1.41
1	Ad	897	A	C2'-C1'	5.06	1.58	1.53
1	Ad	917	U	O4'-C1'	5.06	1.48	1.41
86	Ab	26	C	C4-N4	5.06	1.38	1.33
1	Ad	618	C	C2'-C1'	-5.05	1.47	1.53
84	Aa	2515	C	O3'-P	-5.05	1.55	1.61
1	Ad	809	G	C5'-C4'	5.05	1.57	1.51
1	Ad	349	U	C3'-C2'	-5.05	1.47	1.52
86	Ab	32	A	C6-N1	5.05	1.39	1.35
1	Ad	358	C	C2'-C1'	-5.05	1.47	1.53
84	Aa	2622	G	C2-N3	5.05	1.36	1.32

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	Ad	247	A	O4'-C1'	-5.04	1.35	1.41
1	Ad	722	A	O3'-P	-5.04	1.55	1.61
86	Ab	91	C	N1-C6	5.04	1.40	1.37
1	Ad	729	C	O4'-C1'	5.04	1.48	1.41
1	Ad	1340	A	C4'-C3'	-5.04	1.47	1.52
86	Ab	79	A	N3-C4	-5.04	1.31	1.34
1	Ad	281	U	C3'-O3'	-5.04	1.35	1.42
1	Ad	1690	U	C4'-O4'	5.04	1.52	1.45
1	Ad	1305	U	C2'-C1'	5.04	1.58	1.53
84	Aa	478	G	C2-N3	5.04	1.36	1.32
84	Aa	3234	G	C2-N3	5.04	1.36	1.32
1	Ad	1526	C	C4'-C3'	5.04	1.58	1.53
84	Aa	75	G	C2-N3	5.03	1.36	1.32
86	Ab	1	G	C2'-C1'	-5.03	1.47	1.53
1	Ad	1439	G	O3'-P	-5.03	1.55	1.61
1	Ad	1791	A	C4'-C3'	5.03	1.58	1.53
84	Aa	423	C	O3'-P	-5.03	1.55	1.61
1	Ad	1651	U	C5'-C4'	5.03	1.57	1.51
1	Ad	256	G	C2'-C1'	-5.03	1.47	1.53
1	Ad	415	C	P-O5'	-5.03	1.54	1.59
1	Ad	581	G	C2'-C1'	-5.03	1.47	1.53
1	Ad	821	G	C5'-C4'	5.03	1.57	1.51
1	Ad	1229	C	C5'-C4'	5.03	1.57	1.51
1	Ad	371	A	O4'-C1'	5.02	1.48	1.41
84	Aa	2109	G	C2-N3	5.02	1.36	1.32
1	Ad	1806	C	C5'-C4'	5.02	1.57	1.51
85	Ac	88	A	N7-C5	-5.02	1.36	1.39
1	Ad	763	A	O3'-P	-5.02	1.55	1.61
1	Ad	866	U	C2'-C1'	-5.01	1.47	1.53
1	Ad	943	G	O3'-P	-5.01	1.55	1.61
1	Ad	1204	G	C4'-C3'	5.01	1.58	1.53
86	Ab	66	G	C6-N1	5.01	1.43	1.39
84	Aa	2604	A	N7-C5	-5.01	1.36	1.39
86	Ab	39	C	C2-O2	5.01	1.28	1.24
1	Ad	1711	G	O4'-C1'	5.01	1.48	1.41
2	Ae	57	A	C4'-C3'	5.01	1.58	1.53
84	Aa	3109	G	C2-N3	5.01	1.36	1.32
84	Aa	1644	A	N7-C5	-5.01	1.36	1.39
84	Aa	3070	G	C2-N3	5.00	1.36	1.32
84	Aa	543	C	P-O5'	-5.00	1.54	1.59
84	Aa	1892	A	N7-C5	-5.00	1.36	1.39
1	Ad	1619	A	C2'-C1'	-5.00	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
84	Aa	2460	A	N7-C5	-5.00	1.36	1.39

All (12710) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1005	C	O4'-C1'-N1	31.46	133.37	108.20
1	Ad	1462	C	O4'-C1'-N1	29.23	131.58	108.20
1	Ad	547	C	O4'-C1'-N1	28.96	131.37	108.20
1	Ad	1765	A	O4'-C1'-N9	28.77	131.22	108.20
84	Aa	2162	C	P-O3'-C3'	27.64	152.86	119.70
1	Ad	784	C	O4'-C1'-N1	27.46	130.17	108.20
1	Ad	511	U	O4'-C1'-N1	27.31	130.05	108.20
1	Ad	1580	G	O4'-C1'-N9	27.17	129.94	108.20
1	Ad	1604	C	O4'-C1'-N1	26.94	129.75	108.20
1	Ad	1064	U	O4'-C1'-N1	26.80	129.64	108.20
1	Ad	1368	C	O4'-C1'-N1	25.86	128.88	108.20
1	Ad	787	C	O4'-C1'-N1	25.30	128.44	108.20
1	Ad	67	G	N9-C1'-C2'	25.28	146.87	114.00
1	Ad	384	U	O4'-C1'-N1	24.94	128.15	108.20
1	Ad	1262	U	O4'-C1'-N1	24.71	127.97	108.20
1	Ad	1404	U	O4'-C1'-N1	24.63	127.90	108.20
1	Ad	835	U	O4'-C1'-N1	24.55	127.84	108.20
1	Ad	546	U	O4'-C1'-N1	24.37	127.69	108.20
1	Ad	1408	G	P-O3'-C3'	23.90	148.38	119.70
1	Ad	1622	A	O4'-C1'-N9	23.88	127.30	108.20
1	Ad	282	C	O4'-C1'-N1	23.75	127.20	108.20
1	Ad	1311	U	O4'-C1'-N1	23.61	127.09	108.20
1	Ad	1479	U	O4'-C1'-N1	23.55	127.04	108.20
1	Ad	1066	U	O4'-C1'-N1	23.14	126.71	108.20
2	Ae	73	C	P-O3'-C3'	22.96	147.25	119.70
1	Ad	1511	A	O4'-C1'-N9	22.82	126.45	108.20
1	Ad	869	U	O4'-C1'-N1	22.67	126.34	108.20
1	Ad	801	U	O4'-C1'-N1	22.63	126.31	108.20
1	Ad	1250	C	O4'-C1'-N1	22.52	126.21	108.20
1	Ad	1067	A	O4'-C1'-N9	22.18	125.94	108.20
1	Ad	823	A	P-O3'-C3'	21.59	145.61	119.70
1	Ad	1568	U	O4'-C1'-N1	21.57	125.46	108.20
1	Ad	1522	U	O4'-C1'-N1	21.30	125.24	108.20
1	Ad	325	C	O4'-C1'-N1	21.26	125.20	108.20
2	Ae	72	G	O4'-C1'-N9	21.26	125.20	108.20
1	Ad	1259	G	O4'-C1'-N9	21.24	125.19	108.20
1	Ad	1590	U	O4'-C1'-N1	21.23	125.19	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2502	U	P-O3'-C3'	21.14	145.07	119.70
1	Ad	189	U	O4'-C1'-N1	20.90	124.92	108.20
84	Aa	1957	G	C4'-C3'-O3'	20.90	154.79	113.00
1	Ad	1766	A	O4'-C1'-N9	20.80	124.84	108.20
84	Aa	1559	G	P-O3'-C3'	-20.79	94.76	119.70
84	Aa	2084	G	C4'-C3'-O3'	20.67	154.34	113.00
1	Ad	161	G	O4'-C1'-N9	20.64	124.71	108.20
1	Ad	1350	C	O4'-C1'-N1	20.53	124.63	108.20
84	Aa	641	C	P-O5'-C5'	20.30	153.39	120.90
1	Ad	1162	A	O4'-C1'-N9	20.18	124.34	108.20
1	Ad	1542	G	O4'-C1'-N9	20.15	124.32	108.20
1	Ad	1775	A	O4'-C1'-N9	20.10	124.28	108.20
1	Ad	363	G	O4'-C1'-N9	19.94	124.15	108.20
1	Ad	643	U	O4'-C1'-N1	19.78	124.03	108.20
1	Ad	773	U	O4'-C1'-N1	19.71	123.97	108.20
1	Ad	1531	G	O4'-C1'-N9	19.27	123.61	108.20
1	Ad	780	A	O4'-C1'-N9	19.18	123.55	108.20
1	Ad	456	A	O4'-C1'-N9	19.14	123.52	108.20
84	Aa	570	G	P-O3'-C3'	19.09	142.61	119.70
1	Ad	800	U	O4'-C1'-N1	19.07	123.46	108.20
1	Ad	457	C	N1-C1'-C2'	19.00	138.69	114.00
84	Aa	3182	A	P-O3'-C3'	18.92	142.41	119.70
1	Ad	861	A	O4'-C1'-C2'	-18.79	87.01	105.80
1	Ad	202	C	P-O3'-C3'	18.75	142.20	119.70
1	Ad	137	A	O4'-C1'-N9	18.72	123.17	108.20
84	Aa	1747	A	N1-C6-N6	18.69	129.82	118.60
1	Ad	76	U	P-O3'-C3'	18.55	141.96	119.70
1	Ad	1248	A	O4'-C1'-N9	18.43	122.94	108.20
1	Ad	1501	G	O4'-C1'-N9	18.23	122.78	108.20
1	Ad	1420	U	O4'-C1'-N1	18.12	122.69	108.20
1	Ad	1771	U	O4'-C1'-N1	18.08	122.67	108.20
1	Ad	585	U	O4'-C1'-N1	18.06	122.64	108.20
1	Ad	179	A	O4'-C1'-N9	-18.04	93.77	108.20
84	Aa	1576	C	P-O3'-C3'	17.78	141.03	119.70
1	Ad	1057	U	O4'-C1'-N1	17.68	122.34	108.20
84	Aa	2384	G	P-O3'-C3'	17.67	140.90	119.70
84	Aa	2216	G	N1-C6-O6	17.51	130.41	119.90
84	Aa	434	C	P-O3'-C3'	17.46	140.65	119.70
1	Ad	1802	G	O4'-C1'-N9	17.42	122.14	108.20
1	Ad	391	A	O4'-C1'-N9	17.35	122.08	108.20
1	Ad	939	C	O4'-C1'-N1	17.34	122.07	108.20
1	Ad	828	G	O4'-C1'-N9	17.19	121.95	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1567	G	O4'-C1'-N9	17.19	121.95	108.20
1	Ad	1496	A	O4'-C1'-N9	17.17	121.93	108.20
1	Ad	133	U	P-O3'-C3'	17.14	140.27	119.70
1	Ad	1309	U	O4'-C1'-N1	17.03	121.83	108.20
84	Aa	1950	G	P-O3'-C3'	16.98	140.07	119.70
1	Ad	821	G	N9-C1'-C2'	16.94	136.03	114.00
84	Aa	1263	A	N1-C6-N6	16.88	128.73	118.60
84	Aa	2086	A	O5'-P-OP1	16.79	130.85	110.70
84	Aa	506	U	P-O3'-C3'	16.76	139.82	119.70
84	Aa	2216	G	C5-C6-O6	-16.73	118.56	128.60
1	Ad	1086	A	P-O3'-C3'	16.73	139.77	119.70
1	Ad	1497	U	O4'-C1'-N1	16.69	121.56	108.20
84	Aa	423	C	P-O3'-C3'	16.66	139.69	119.70
1	Ad	394	G	O4'-C1'-N9	16.59	121.47	108.20
1	Ad	1203	G	O4'-C1'-N9	16.49	121.39	108.20
1	Ad	280	U	P-O3'-C3'	-16.48	99.92	119.70
84	Aa	158	A	P-O3'-C3'	16.41	139.39	119.70
1	Ad	396	G	O4'-C1'-N9	16.38	121.31	108.20
86	Ab	16	A	N1-C6-N6	16.37	128.42	118.60
1	Ad	1592	G	O4'-C1'-N9	16.36	121.28	108.20
1	Ad	1203	G	C1'-O4'-C4'	16.35	122.98	109.90
1	Ad	1353	G	P-O3'-C3'	16.33	139.30	119.70
1	Ad	906	G	O4'-C1'-N9	16.33	121.26	108.20
1	Ad	1619	A	O4'-C1'-N9	16.32	121.26	108.20
1	Ad	732	G	O4'-C1'-N9	16.26	121.21	108.20
1	Ad	1498	A	O4'-C1'-N9	16.20	121.16	108.20
1	Ad	1623	C	P-O3'-C3'	16.16	139.09	119.70
1	Ad	1405	U	O4'-C1'-N1	16.08	121.06	108.20
1	Ad	730	G	O4'-C1'-N9	16.00	121.00	108.20
86	Ab	115	A	N1-C6-N6	15.98	128.19	118.60
86	Ab	80	A	N1-C6-N6	15.96	128.18	118.60
84	Aa	144	A	N1-C6-N6	15.89	128.14	118.60
84	Aa	1019	A	N1-C6-N6	15.88	128.13	118.60
86	Ab	19	A	N1-C6-N6	15.84	128.11	118.60
86	Ab	31	G	N1-C6-O6	15.83	129.40	119.90
84	Aa	1726	G	P-O3'-C3'	15.73	138.58	119.70
1	Ad	83	U	O4'-C1'-N1	15.62	120.70	108.20
1	Ad	139	U	O4'-C1'-N1	15.62	120.69	108.20
1	Ad	1149	U	O4'-C1'-N1	15.59	120.67	108.20
1	Ad	1245	G	O4'-C1'-N9	15.52	120.62	108.20
1	Ad	800	U	P-O3'-C3'	15.47	138.27	119.70
1	Ad	458	A	O4'-C1'-N9	15.46	120.57	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1487	A	P-O3'-C3'	15.45	138.25	119.70
84	Aa	3153	U	P-O3'-C3'	15.41	138.19	119.70
1	Ad	444	U	O4'-C1'-N1	15.39	120.52	108.20
1	Ad	1205	G	O4'-C1'-N9	15.39	120.51	108.20
1	Ad	1220	C	P-O3'-C3'	15.35	138.12	119.70
1	Ad	792	U	O4'-C1'-N1	15.35	120.48	108.20
86	Ab	55	A	N1-C6-N6	15.35	127.81	118.60
84	Aa	2152	A	P-O3'-C3'	15.34	138.11	119.70
3	Af	12	A	O4'-C1'-N9	15.21	120.36	108.20
84	Aa	2167	G	C5-C6-O6	-15.19	119.48	128.60
84	Aa	571	G	P-O3'-C3'	15.18	137.91	119.70
84	Aa	2178	G	C4'-C3'-O3'	-15.15	77.58	109.40
84	Aa	3182	A	O4'-C1'-N9	15.14	120.31	108.20
1	Ad	707	C	O4'-C1'-N1	15.12	120.30	108.20
1	Ad	860	A	O4'-C1'-N9	15.12	120.30	108.20
1	Ad	344	U	O4'-C1'-N1	15.09	120.27	108.20
84	Aa	939	A	N1-C6-N6	15.05	127.63	118.60
84	Aa	2085	A	P-O5'-C5'	15.02	144.94	120.90
2	Ae	58	U	O4'-C1'-N1	15.01	120.21	108.20
1	Ad	1102	U	O4'-C1'-N1	15.00	120.20	108.20
86	Ab	79	A	C5-C6-N1	-14.95	110.22	117.70
1	Ad	845	C	P-O3'-C3'	14.92	137.61	119.70
84	Aa	1308	A	N1-C6-N6	14.92	127.55	118.60
84	Aa	1811	U	P-O3'-C3'	14.90	137.58	119.70
84	Aa	2086	A	O5'-P-OP2	-14.89	92.30	105.70
1	Ad	208	U	P-O3'-C3'	14.89	137.56	119.70
1	Ad	559	A	P-O3'-C3'	14.88	137.56	119.70
1	Ad	156	U	O4'-C1'-N1	14.87	120.10	108.20
84	Aa	1322	A	N1-C6-N6	14.87	127.52	118.60
86	Ab	43	A	N1-C6-N6	14.87	127.52	118.60
1	Ad	1344	U	O4'-C1'-N1	14.84	120.07	108.20
84	Aa	2100	A	P-O3'-C3'	14.78	137.43	119.70
1	Ad	1019	G	O4'-C1'-N9	14.77	120.02	108.20
1	Ad	80	C	P-O3'-C3'	14.77	137.42	119.70
2	Ae	12	U	O4'-C1'-N1	14.77	120.01	108.20
1	Ad	1226	U	O4'-C1'-N1	14.77	120.01	108.20
84	Aa	2086	A	C2'-C3'-O3'	14.70	141.84	109.50
1	Ad	1377	G	O4'-C1'-N9	14.69	119.95	108.20
1	Ad	1044	A	O4'-C1'-N9	14.66	119.93	108.20
84	Aa	2708	A	N1-C6-N6	14.65	127.39	118.60
1	Ad	261	C	N1-C1'-C2'	14.64	133.03	114.00
84	Aa	1057	A	N1-C6-N6	14.62	127.37	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1796	A	N1-C6-N6	14.60	127.36	118.60
85	Ac	71	A	N1-C6-N6	14.56	127.34	118.60
84	Aa	1526	A	N1-C6-N6	14.56	127.33	118.60
1	Ad	516	A	P-O3'-C3'	14.55	137.16	119.70
84	Aa	563	C	P-O3'-C3'	14.54	137.14	119.70
1	Ad	372	U	O4'-C1'-N1	14.53	119.83	108.20
84	Aa	3221	A	P-O3'-C3'	14.51	137.11	119.70
1	Ad	215	A	P-O3'-C3'	14.49	137.09	119.70
84	Aa	64	A	N1-C6-N6	14.48	127.29	118.60
84	Aa	51	A	N1-C6-N6	14.45	127.27	118.60
1	Ad	828	G	P-O3'-C3'	14.43	137.02	119.70
1	Ad	1675	G	O4'-C1'-N9	14.41	119.73	108.20
1	Ad	1492	G	O4'-C1'-N9	14.40	119.72	108.20
1	Ad	1169	G	O4'-C1'-N9	14.37	119.70	108.20
86	Ab	79	A	N1-C6-N6	14.37	127.22	118.60
84	Aa	2072	U	P-O3'-C3'	14.36	136.93	119.70
84	Aa	349	A	N1-C6-N6	14.35	127.21	118.60
1	Ad	1206	A	N9-C1'-C2'	14.34	132.64	114.00
1	Ad	843	G	C3'-C2'-C1'	14.32	112.96	101.50
1	Ad	131	C	P-O3'-C3'	14.30	136.86	119.70
84	Aa	3308	A	P-O3'-C3'	14.28	136.84	119.70
86	Ab	42	A	N1-C6-N6	14.27	127.16	118.60
1	Ad	707	C	C1'-O4'-C4'	14.27	121.31	109.90
84	Aa	2286	A	N1-C6-N6	14.27	127.16	118.60
1	Ad	1504	U	O4'-C1'-N1	14.26	119.61	108.20
1	Ad	1694	G	O4'-C1'-N9	14.25	119.60	108.20
84	Aa	3001	G	O4'-C1'-N9	14.24	119.59	108.20
84	Aa	2461	A	P-O3'-C3'	14.20	136.74	119.70
86	Ab	50	A	N1-C6-N6	14.19	127.11	118.60
1	Ad	851	G	O4'-C1'-N9	14.19	119.55	108.20
84	Aa	587	A	N1-C6-N6	14.19	127.11	118.60
84	Aa	2494	A	N1-C6-N6	14.18	127.11	118.60
84	Aa	2932	A	N1-C6-N6	14.14	127.08	118.60
84	Aa	1146	A	N1-C6-N6	14.10	127.06	118.60
1	Ad	1371	U	O4'-C1'-N1	14.09	119.47	108.20
84	Aa	716	A	N1-C6-N6	14.07	127.04	118.60
84	Aa	2053	A	P-O3'-C3'	14.06	136.57	119.70
84	Aa	849	A	N1-C6-N6	14.05	127.03	118.60
1	Ad	215	A	O4'-C1'-N9	14.05	119.44	108.20
1	Ad	150	U	O4'-C1'-N1	14.04	119.43	108.20
86	Ab	47	C	P-O3'-C3'	14.04	136.54	119.70
1	Ad	1421	U	O4'-C1'-N1	14.03	119.43	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	87	A	C1'-O4'-C4'	14.03	121.12	109.90
1	Ad	786	U	O4'-C1'-N1	14.03	119.42	108.20
84	Aa	3175	C	P-O3'-C3'	14.03	136.54	119.70
1	Ad	501	U	P-O3'-C3'	14.02	136.53	119.70
84	Aa	168	A	N1-C6-N6	14.02	127.01	118.60
1	Ad	349	U	O4'-C1'-N1	14.01	119.41	108.20
1	Ad	1621	U	O4'-C1'-N1	14.00	119.40	108.20
1	Ad	63	G	O4'-C1'-N9	14.00	119.40	108.20
1	Ad	1440	U	O4'-C1'-N1	14.00	119.40	108.20
1	Ad	502	G	P-O3'-C3'	13.98	136.47	119.70
84	Aa	2167	G	N1-C6-O6	13.97	128.28	119.90
84	Aa	3025	A	N1-C6-N6	13.93	126.96	118.60
84	Aa	1088	A	N1-C6-N6	13.93	126.96	118.60
84	Aa	488	U	P-O3'-C3'	13.92	136.40	119.70
1	Ad	1643	A	O4'-C1'-N9	13.91	119.33	108.20
84	Aa	2528	U	P-O3'-C3'	13.90	136.38	119.70
1	Ad	1464	G	N9-C1'-C2'	13.89	132.06	114.00
84	Aa	1812	A	P-O3'-C3'	13.88	136.36	119.70
1	Ad	884	G	O4'-C1'-N9	13.87	119.30	108.20
84	Aa	2143	A	N1-C6-N6	13.86	126.92	118.60
84	Aa	2311	A	N1-C6-N6	13.84	126.90	118.60
84	Aa	159	G	N1-C6-O6	13.80	128.18	119.90
1	Ad	1807	A	O4'-C1'-N9	13.80	119.24	108.20
84	Aa	1136	A	N1-C6-N6	13.79	126.88	118.60
1	Ad	1550	G	O4'-C1'-N9	13.78	119.22	108.20
84	Aa	2646	A	N1-C6-N6	13.78	126.87	118.60
84	Aa	686	A	N1-C6-N6	13.77	126.86	118.60
1	Ad	562	U	O4'-C1'-N1	13.77	119.21	108.20
1	Ad	744	G	O4'-C1'-N9	13.72	119.18	108.20
84	Aa	1694	A	N1-C6-N6	13.71	126.83	118.60
1	Ad	1101	C	O4'-C1'-N1	13.70	119.16	108.20
1	Ad	1387	U	O4'-C1'-N1	13.67	119.13	108.20
1	Ad	1810	G	C3'-C2'-C1'	13.67	112.43	101.50
1	Ad	79	A	O4'-C1'-N9	13.66	119.13	108.20
84	Aa	525	A	N1-C6-N6	13.66	126.79	118.60
84	Aa	1538	A	N1-C6-N6	13.66	126.79	118.60
86	Ab	11	A	N1-C6-N6	13.65	126.79	118.60
1	Ad	722	A	P-O3'-C3'	13.64	136.07	119.70
84	Aa	657	A	N1-C6-N6	13.64	126.78	118.60
1	Ad	1583	G	O4'-C1'-N9	13.64	119.11	108.20
1	Ad	1763	A	O4'-C1'-N9	13.61	119.09	108.20
86	Ab	3	A	N1-C6-N6	13.60	126.76	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2237	A	N1-C6-N6	13.59	126.75	118.60
84	Aa	2165	A	N1-C6-N6	13.58	126.75	118.60
84	Aa	304	A	N1-C6-N6	13.57	126.74	118.60
84	Aa	2142	A	N1-C6-N6	13.57	126.74	118.60
84	Aa	1463	A	N1-C6-N6	13.55	126.73	118.60
36	BH	117	ARG	NE-CZ-NH1	-13.55	113.53	120.30
1	Ad	139	U	P-O3'-C3'	13.55	135.96	119.70
84	Aa	2177	U	C4'-C3'-O3'	-13.52	81.01	109.40
85	Ac	85	G	P-O3'-C3'	13.52	135.92	119.70
1	Ad	1655	U	O4'-C1'-N1	13.51	119.00	108.20
84	Aa	119	A	N1-C6-N6	13.51	126.70	118.60
84	Aa	1644	A	N1-C6-N6	13.49	126.69	118.60
1	Ad	815	A	P-O3'-C3'	13.49	135.88	119.70
1	Ad	918	G	P-O3'-C3'	13.48	135.88	119.70
1	Ad	1343	C	P-O3'-C3'	13.48	135.88	119.70
1	Ad	29	U	O4'-C1'-N1	13.47	118.98	108.20
84	Aa	3057	A	N1-C6-N6	13.47	126.69	118.60
84	Aa	3211	C	P-O3'-C3'	13.46	135.86	119.70
84	Aa	3143	A	N1-C6-N6	13.46	126.68	118.60
1	Ad	185	G	P-O3'-C3'	13.46	135.85	119.70
1	Ad	336	U	O4'-C1'-N1	13.44	118.95	108.20
1	Ad	999	G	C1'-O4'-C4'	-13.43	99.15	109.90
84	Aa	3220	A	N1-C6-N6	13.43	126.66	118.60
84	Aa	1344	A	N1-C6-N6	13.42	126.65	118.60
1	Ad	461	G	O4'-C1'-N9	13.41	118.93	108.20
84	Aa	50	A	N1-C6-N6	13.41	126.65	118.60
84	Aa	540	G	N1-C6-O6	13.40	127.94	119.90
84	Aa	2751	A	N1-C6-N6	13.37	126.62	118.60
1	Ad	276	G	O4'-C1'-N9	13.37	118.89	108.20
84	Aa	920	A	N1-C6-N6	13.36	126.61	118.60
1	Ad	1162	A	P-O3'-C3'	13.35	135.72	119.70
86	Ab	68	G	N1-C6-O6	13.35	127.91	119.90
84	Aa	2313	U	P-O3'-C3'	13.34	135.71	119.70
84	Aa	2905	A	N1-C6-N6	13.33	126.60	118.60
84	Aa	1456	A	N1-C6-N6	13.31	126.58	118.60
84	Aa	1855	A	N1-C6-N6	13.30	126.58	118.60
84	Aa	73	A	N1-C6-N6	13.29	126.57	118.60
2	Ae	73	C	N1-C1'-C2'	13.28	131.27	114.00
1	Ad	145	A	O4'-C1'-N9	13.28	118.82	108.20
84	Aa	353	A	N1-C6-N6	13.28	126.57	118.60
1	Ad	1183	G	O4'-C1'-N9	13.27	118.82	108.20
84	Aa	747	A	N1-C6-N6	13.27	126.56	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2948	A	N1-C6-N6	13.27	126.56	118.60
84	Aa	3315	A	N1-C6-N6	13.26	126.55	118.60
84	Aa	1490	A	N1-C6-N6	13.25	126.55	118.60
84	Aa	454	A	N1-C6-N6	13.25	126.55	118.60
1	Ad	1367	U	O4'-C1'-N1	13.24	118.79	108.20
84	Aa	3351	A	N1-C6-N6	13.23	126.54	118.60
1	Ad	930	G	O4'-C1'-N9	13.22	118.78	108.20
84	Aa	2503	A	P-O3'-C3'	13.21	135.55	119.70
84	Aa	217	A	N1-C6-N6	13.21	126.53	118.60
84	Aa	2504	A	P-O3'-C3'	13.21	135.55	119.70
84	Aa	3012	A	N1-C6-N6	13.21	126.52	118.60
84	Aa	2483	A	N1-C6-N6	13.21	126.52	118.60
84	Aa	2125	A	N1-C6-N6	13.20	126.52	118.60
1	Ad	167	A	O4'-C1'-N9	13.19	118.75	108.20
84	Aa	2996	A	N1-C6-N6	13.19	126.51	118.60
1	Ad	1581	A	P-O3'-C3'	13.19	135.52	119.70
1	Ad	32	U	O4'-C1'-N1	13.18	118.75	108.20
84	Aa	1501	A	N1-C6-N6	13.18	126.51	118.60
84	Aa	2513	U	C5'-C4'-C3'	-13.18	94.92	116.00
84	Aa	2849	A	N1-C6-N6	13.17	126.50	118.60
1	Ad	723	A	O4'-C1'-N9	13.16	118.73	108.20
84	Aa	1704	A	N1-C6-N6	13.15	126.49	118.60
84	Aa	332	A	N1-C6-N6	13.15	126.49	118.60
1	Ad	634	A	O4'-C1'-N9	13.13	118.70	108.20
84	Aa	2898	A	N1-C6-N6	13.13	126.48	118.60
84	Aa	830	A	N1-C6-N6	13.12	126.47	118.60
1	Ad	2	A	O4'-C1'-N9	13.11	118.69	108.20
84	Aa	774	A	N1-C6-N6	13.10	126.46	118.60
85	Ac	52	A	N1-C6-N6	13.09	126.46	118.60
1	Ad	1134	U	O4'-C1'-N1	13.09	118.67	108.20
1	Ad	235	C	O4'-C1'-C2'	-13.09	92.72	105.80
84	Aa	1334	A	N1-C6-N6	13.07	126.44	118.60
84	Aa	1224	A	N1-C6-N6	13.03	126.42	118.60
84	Aa	3113	G	N1-C6-O6	13.03	127.72	119.90
85	Ac	12	A	N1-C6-N6	13.00	126.40	118.60
84	Aa	2989	A	N1-C6-N6	13.00	126.40	118.60
84	Aa	2331	A	N1-C6-N6	13.00	126.40	118.60
84	Aa	3136	A	N1-C6-N6	13.00	126.40	118.60
84	Aa	1838	A	N1-C6-N6	12.99	126.39	118.60
84	Aa	2351	A	N1-C6-N6	12.98	126.39	118.60
84	Aa	1123	A	N1-C6-N6	12.98	126.39	118.60
1	Ad	1254	U	P-O3'-C3'	12.96	135.25	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1195	C	P-O3'-C3'	12.96	135.25	119.70
84	Aa	2788	A	N1-C6-N6	12.96	126.37	118.60
84	Aa	1061	A	N1-C6-N6	12.95	126.37	118.60
84	Aa	2538	G	P-O3'-C3'	12.95	135.23	119.70
1	Ad	1376	A	O4'-C1'-N9	12.94	118.56	108.20
2	Ae	47	U	O4'-C1'-N1	12.94	118.55	108.20
84	Aa	723	G	O4'-C1'-N9	12.94	118.55	108.20
84	Aa	156	A	N1-C6-N6	12.93	126.36	118.60
84	Aa	2984	A	N1-C6-N6	12.93	126.36	118.60
84	Aa	2098	A	N1-C6-N6	12.92	126.35	118.60
1	Ad	486	U	O4'-C1'-N1	12.91	118.53	108.20
84	Aa	1367	A	N1-C6-N6	12.91	126.35	118.60
84	Aa	573	A	N1-C6-N6	12.91	126.35	118.60
84	Aa	1274	A	N1-C6-N6	12.91	126.34	118.60
84	Aa	3382	A	N1-C6-N6	12.90	126.34	118.60
84	Aa	2810	A	N1-C6-N6	12.90	126.34	118.60
84	Aa	2943	A	N1-C6-N6	12.90	126.34	118.60
84	Aa	697	A	N1-C6-N6	12.89	126.34	118.60
84	Aa	3336	A	N1-C6-N6	12.89	126.33	118.60
84	Aa	672	A	N1-C6-N6	12.87	126.32	118.60
2	Ae	55	C	N1-C1'-C2'	12.87	130.74	114.00
1	Ad	788	G	P-O3'-C3'	12.87	135.15	119.70
1	Ad	379	U	O4'-C1'-N1	12.87	118.49	108.20
84	Aa	2171	A	N1-C6-N6	12.86	126.31	118.60
84	Aa	294	A	N1-C6-N6	12.85	126.31	118.60
84	Aa	2679	A	N1-C6-N6	12.85	126.31	118.60
1	Ad	404	A	O4'-C1'-N9	12.83	118.46	108.20
84	Aa	2561	A	N1-C6-N6	12.81	126.29	118.60
84	Aa	2372	A	N1-C6-N6	12.81	126.28	118.60
1	Ad	1241	G	O4'-C1'-N9	12.80	118.44	108.20
84	Aa	1819	A	N1-C6-N6	12.80	126.28	118.60
84	Aa	2640	A	N1-C6-N6	12.80	126.28	118.60
1	Ad	1406	U	O4'-C1'-N1	12.80	118.44	108.20
1	Ad	557	G	O4'-C1'-N9	12.80	118.44	108.20
84	Aa	499	A	P-O3'-C3'	12.80	135.06	119.70
84	Aa	1727	A	P-O3'-C3'	12.79	135.05	119.70
84	Aa	1843	A	N1-C6-N6	12.79	126.28	118.60
84	Aa	1904	A	N1-C6-N6	12.79	126.27	118.60
84	Aa	2787	A	N1-C6-N6	12.79	126.27	118.60
1	Ad	1460	G	O4'-C1'-N9	12.78	118.42	108.20
1	Ad	1248	A	P-O3'-C3'	12.78	135.03	119.70
84	Aa	1225	A	N1-C6-N6	12.77	126.26	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2139	A	N1-C6-N6	12.77	126.26	118.60
84	Aa	3271	A	N1-C6-N6	12.77	126.26	118.60
84	Aa	56	A	N1-C6-N6	12.77	126.26	118.60
84	Aa	262	A	N1-C6-N6	12.77	126.26	118.60
84	Aa	1349	G	N1-C6-O6	12.76	127.56	119.90
84	Aa	385	A	N1-C6-N6	12.76	126.25	118.60
1	Ad	219	G	O4'-C1'-N9	12.75	118.40	108.20
84	Aa	121	A	N1-C6-N6	12.74	126.25	118.60
84	Aa	3041	A	N1-C6-N6	12.74	126.24	118.60
86	Ab	77	A	N1-C6-N6	12.74	126.24	118.60
1	Ad	1203	G	N9-C1'-C2'	-12.73	97.44	114.00
84	Aa	982	U	P-O3'-C3'	12.73	134.98	119.70
1	Ad	1689	A	O4'-C1'-N9	12.73	118.38	108.20
84	Aa	1114	A	N1-C6-N6	12.73	126.24	118.60
84	Aa	426	A	N1-C6-N6	12.73	126.24	118.60
1	Ad	1659	A	O4'-C1'-N9	12.72	118.38	108.20
84	Aa	39	A	N1-C6-N6	12.72	126.23	118.60
84	Aa	1918	A	N1-C6-N6	12.71	126.23	118.60
1	Ad	1688	G	O4'-C1'-N9	12.71	118.37	108.20
84	Aa	1221	A	N1-C6-N6	12.71	126.22	118.60
1	Ad	761	A	P-O3'-C3'	12.71	134.95	119.70
84	Aa	3045	A	N1-C6-N6	12.70	126.22	118.60
84	Aa	2396	A	N1-C6-N6	12.70	126.22	118.60
84	Aa	3358	A	N1-C6-N6	12.70	126.22	118.60
84	Aa	2562	A	N1-C6-N6	12.69	126.21	118.60
84	Aa	2385	A	N1-C6-N6	12.68	126.21	118.60
1	Ad	176	A	O4'-C1'-N9	12.68	118.34	108.20
85	Ac	157	A	N1-C6-N6	12.68	126.20	118.60
84	Aa	792	A	N1-C6-N6	12.67	126.20	118.60
84	Aa	886	A	N1-C6-N6	12.67	126.20	118.60
84	Aa	3113	G	C5-C6-O6	-12.67	121.00	128.60
85	Ac	126	A	N1-C6-N6	12.67	126.20	118.60
84	Aa	929	A	N1-C6-N6	12.66	126.20	118.60
84	Aa	2074	C	P-O3'-C3'	12.66	134.90	119.70
84	Aa	2512	U	C2'-C3'-O3'	12.66	137.35	109.50
2	Ae	28	G	C1'-O4'-C4'	-12.64	99.78	109.90
85	Ac	41	A	N1-C6-N6	12.64	126.19	118.60
84	Aa	3177	A	N1-C6-N6	12.63	126.18	118.60
1	Ad	878	U	O4'-C1'-N1	12.63	118.30	108.20
86	Ab	99	G	N1-C6-O6	12.62	127.47	119.90
84	Aa	395	A	N1-C6-N6	12.62	126.17	118.60
84	Aa	2491	A	N1-C6-N6	12.62	126.17	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2969	A	N1-C6-N6	12.61	126.17	118.60
84	Aa	3049	A	N1-C6-N6	12.61	126.17	118.60
1	Ad	1753	U	O4'-C1'-N1	12.61	118.29	108.20
84	Aa	1282	A	N1-C6-N6	12.61	126.16	118.60
84	Aa	322	A	N1-C6-N6	12.60	126.16	118.60
85	Ac	33	A	N1-C6-N6	12.60	126.16	118.60
84	Aa	2357	A	N1-C6-N6	12.59	126.16	118.60
1	Ad	635	G	O4'-C1'-N9	12.59	118.27	108.20
84	Aa	2439	A	N1-C6-N6	12.59	126.16	118.60
84	Aa	1024	G	N1-C6-O6	12.58	127.45	119.90
1	Ad	300	U	O4'-C1'-N1	12.58	118.26	108.20
84	Aa	2993	A	N1-C6-N6	12.58	126.14	118.60
84	Aa	2805	A	N1-C6-N6	12.57	126.14	118.60
84	Aa	2294	A	N1-C6-N6	12.57	126.14	118.60
84	Aa	3306	A	N1-C6-N6	12.57	126.14	118.60
84	Aa	481	G	P-O3'-C3'	12.56	134.78	119.70
84	Aa	1153	A	N1-C6-N6	12.56	126.14	118.60
84	Aa	1753	A	N1-C6-N6	12.56	126.14	118.60
84	Aa	1359	A	N1-C6-N6	12.56	126.14	118.60
84	Aa	586	A	N1-C6-N6	12.55	126.13	118.60
1	Ad	1041	A	O4'-C1'-N9	12.55	118.24	108.20
84	Aa	2645	A	N1-C6-N6	12.54	126.12	118.60
84	Aa	499	A	N1-C6-N6	12.53	126.11	118.60
84	Aa	1000	A	N1-C6-N6	12.53	126.12	118.60
84	Aa	1438	A	N1-C6-N6	12.53	126.12	118.60
84	Aa	1336	A	N1-C6-N6	12.52	126.11	118.60
84	Aa	2904	A	N1-C6-N6	12.52	126.11	118.60
84	Aa	1927	A	N1-C6-N6	12.52	126.11	118.60
84	Aa	1590	A	N1-C6-N6	12.51	126.11	118.60
84	Aa	65	A	N1-C6-N6	12.50	126.10	118.60
1	Ad	1014	U	O4'-C1'-N1	12.49	118.19	108.20
84	Aa	811	A	N1-C6-N6	12.49	126.09	118.60
84	Aa	1013	A	N1-C6-N6	12.49	126.09	118.60
84	Aa	2638	A	N1-C6-N6	12.49	126.09	118.60
84	Aa	3114	A	N1-C6-N6	12.49	126.09	118.60
84	Aa	99	A	N1-C6-N6	12.48	126.09	118.60
84	Aa	1640	A	N1-C6-N6	12.48	126.09	118.60
84	Aa	1398	A	N1-C6-N6	12.48	126.09	118.60
2	Ae	74	C	C1'-O4'-C4'	12.47	119.88	109.90
84	Aa	2356	A	N1-C6-N6	12.47	126.08	118.60
85	Ac	40	A	N1-C6-N6	12.47	126.08	118.60
84	Aa	887	A	N1-C6-N6	12.47	126.08	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	159	G	C5-C6-O6	-12.47	121.12	128.60
84	Aa	33	A	N1-C6-N6	12.46	126.08	118.60
84	Aa	721	A	N1-C6-N6	12.46	126.08	118.60
84	Aa	2630	A	N1-C6-N6	12.46	126.08	118.60
84	Aa	1002	A	N1-C6-N6	12.46	126.07	118.60
84	Aa	526	A	P-O5'-C5'	12.45	140.82	120.90
84	Aa	3092	A	N1-C6-N6	12.45	126.07	118.60
84	Aa	2958	A	N1-C6-N6	12.45	126.07	118.60
84	Aa	1929	A	N1-C6-N6	12.44	126.06	118.60
1	Ad	891	U	O4'-C1'-N1	12.44	118.15	108.20
1	Ad	45	U	C3'-C2'-C1'	12.44	111.45	101.50
1	Ad	845	C	C1'-O4'-C4'	12.44	119.85	109.90
84	Aa	2014	A	N1-C6-N6	12.44	126.06	118.60
84	Aa	2482	A	N1-C6-N6	12.43	126.06	118.60
84	Aa	746	C	O4'-C1'-N1	12.43	118.14	108.20
84	Aa	1831	A	N1-C6-N6	12.42	126.05	118.60
1	Ad	583	A	O4'-C1'-N9	12.42	118.14	108.20
84	Aa	3148	A	N1-C6-N6	12.42	126.05	118.60
84	Aa	2214	A	N1-C6-N6	12.41	126.05	118.60
84	Aa	3221	A	N1-C6-N6	12.41	126.05	118.60
1	Ad	190	C	P-O3'-C3'	12.41	134.59	119.70
84	Aa	879	A	N1-C6-N6	12.41	126.05	118.60
84	Aa	1932	A	N1-C6-N6	12.41	126.05	118.60
1	Ad	197	G	O4'-C1'-N9	12.40	118.12	108.20
84	Aa	985	C	P-O3'-C3'	12.40	134.58	119.70
86	Ab	112	U	O4'-C1'-N1	12.40	118.12	108.20
84	Aa	705	A	N1-C6-N6	12.40	126.04	118.60
1	Ad	986	U	O4'-C1'-N1	12.39	118.11	108.20
84	Aa	1040	A	N1-C6-N6	12.39	126.04	118.60
84	Aa	2912	A	N1-C6-N6	12.39	126.04	118.60
84	Aa	1157	A	N1-C6-N6	12.39	126.03	118.60
84	Aa	3201	A	N1-C6-N6	12.39	126.03	118.60
84	Aa	2695	A	N1-C6-N6	12.38	126.03	118.60
84	Aa	61	A	N1-C6-N6	12.38	126.03	118.60
84	Aa	3104	A	N1-C6-N6	12.38	126.03	118.60
1	Ad	1100	U	O4'-C1'-N1	12.38	118.10	108.20
84	Aa	1232	A	N1-C6-N6	12.37	126.02	118.60
84	Aa	1494	A	N1-C6-N6	12.37	126.02	118.60
84	Aa	2935	A	N1-C6-N6	12.36	126.02	118.60
1	Ad	1690	U	O4'-C1'-N1	12.36	118.09	108.20
84	Aa	2276	A	N1-C6-N6	12.36	126.02	118.60
84	Aa	487	C	P-O3'-C3'	12.35	134.53	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	Ab	49	A	N1-C6-N6	12.35	126.01	118.60
84	Aa	2344	A	N1-C6-N6	12.35	126.01	118.60
84	Aa	2397	A	N1-C6-N6	12.35	126.01	118.60
84	Aa	1731	A	N1-C6-N6	12.35	126.01	118.60
86	Ab	110	G	N1-C6-O6	12.35	127.31	119.90
85	Ac	54	A	N1-C6-N6	12.34	126.00	118.60
85	Ac	110	A	N1-C6-N6	12.34	126.00	118.60
84	Aa	2101	A	N1-C6-N6	12.33	126.00	118.60
84	Aa	639	A	N1-C6-N6	12.33	126.00	118.60
84	Aa	2889	A	N1-C6-N6	12.33	126.00	118.60
84	Aa	216	G	C5-C6-O6	-12.32	121.21	128.60
84	Aa	846	A	N1-C6-N6	12.32	125.99	118.60
2	Ae	37	G	P-O3'-C3'	12.32	134.48	119.70
1	Ad	315	U	O4'-C1'-N1	12.31	118.05	108.20
84	Aa	373	A	N1-C6-N6	12.31	125.99	118.60
84	Aa	2400	A	N1-C6-N6	12.31	125.99	118.60
84	Aa	393	A	N1-C6-N6	12.31	125.99	118.60
84	Aa	1512	A	N1-C6-N6	12.31	125.99	118.60
84	Aa	1097	A	N1-C6-N6	12.31	125.98	118.60
84	Aa	2670	A	N1-C6-N6	12.31	125.98	118.60
2	Ae	36	C	N1-C1'-C2'	12.30	130.00	114.00
84	Aa	2503	A	N1-C6-N6	12.30	125.98	118.60
1	Ad	383	U	O4'-C1'-N1	12.30	118.04	108.20
84	Aa	2389	A	N1-C6-N6	12.30	125.98	118.60
84	Aa	282	A	N1-C6-N6	12.30	125.98	118.60
84	Aa	2730	A	N1-C6-N6	12.29	125.98	118.60
2	Ae	50	G	O4'-C1'-N9	12.29	118.03	108.20
84	Aa	2223	A	N1-C6-N6	12.29	125.98	118.60
84	Aa	1651	A	N1-C6-N6	12.28	125.97	118.60
86	Ab	81	G	N1-C6-O6	12.28	127.27	119.90
84	Aa	2217	A	N1-C6-N6	12.27	125.96	118.60
1	Ad	331	U	O4'-C1'-N1	12.27	118.02	108.20
1	Ad	1537	U	O4'-C1'-N1	12.27	118.02	108.20
84	Aa	6	A	N1-C6-N6	12.27	125.96	118.60
84	Aa	661	A	N1-C6-N6	12.27	125.96	118.60
84	Aa	118	G	N1-C6-O6	12.27	127.26	119.90
84	Aa	1106	G	P-O3'-C3'	12.27	134.42	119.70
84	Aa	196	A	N1-C6-N6	12.26	125.96	118.60
1	Ad	842	G	O4'-C1'-N9	12.26	118.00	108.20
84	Aa	1370	A	N1-C6-N6	12.26	125.95	118.60
84	Aa	2210	A	N1-C6-N6	12.25	125.95	118.60
84	Aa	2724	A	N1-C6-N6	12.25	125.95	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	778	G	O4'-C1'-N9	12.24	118.00	108.20
84	Aa	2138	A	N1-C6-N6	12.24	125.95	118.60
84	Aa	1928	A	N1-C6-N6	12.23	125.94	118.60
84	Aa	2152	A	N1-C6-N6	12.23	125.94	118.60
84	Aa	2239	A	N1-C6-N6	12.23	125.94	118.60
84	Aa	3034	A	N1-C6-N6	12.23	125.94	118.60
84	Aa	64	A	P-O3'-C3'	12.23	134.38	119.70
1	Ad	1278	C	O4'-C1'-N1	12.23	117.98	108.20
84	Aa	3110	A	N1-C6-N6	12.22	125.93	118.60
1	Ad	606	U	O4'-C1'-N1	12.22	117.97	108.20
84	Aa	266	A	N1-C6-N6	12.21	125.93	118.60
84	Aa	2266	A	N1-C6-N6	12.21	125.93	118.60
84	Aa	3272	A	N1-C6-N6	12.21	125.93	118.60
84	Aa	3017	A	N1-C6-N6	12.21	125.93	118.60
84	Aa	2082	A	N1-C6-N6	12.20	125.92	118.60
84	Aa	2290	A	N1-C6-N6	12.20	125.92	118.60
1	Ad	507	G	P-O3'-C3'	12.20	134.34	119.70
84	Aa	2804	A	N1-C6-N6	12.19	125.91	118.60
84	Aa	917	A	N1-C6-N6	12.19	125.91	118.60
84	Aa	2089	A	N1-C6-N6	12.18	125.91	118.60
84	Aa	2487	A	N1-C6-N6	12.18	125.91	118.60
84	Aa	1882	A	N1-C6-N6	12.17	125.91	118.60
84	Aa	1373	A	N1-C6-N6	12.17	125.90	118.60
84	Aa	3087	A	N1-C6-N6	12.17	125.90	118.60
84	Aa	364	A	N1-C6-N6	12.16	125.90	118.60
84	Aa	3078	A	N1-C6-N6	12.16	125.90	118.60
1	Ad	1780	U	O4'-C1'-N1	12.16	117.93	108.20
84	Aa	383	A	N1-C6-N6	12.16	125.90	118.60
84	Aa	2100	A	N1-C6-N6	12.16	125.89	118.60
1	Ad	294	G	O4'-C1'-N9	12.15	117.92	108.20
84	Aa	955	A	N1-C6-N6	12.15	125.89	118.60
84	Aa	2227	A	N1-C6-N6	12.15	125.89	118.60
84	Aa	3018	A	N1-C6-N6	12.15	125.89	118.60
84	Aa	3107	A	N1-C6-N6	12.15	125.89	118.60
85	Ac	13	A	N1-C6-N6	12.15	125.89	118.60
1	Ad	1810	G	O4'-C1'-N9	-12.14	98.49	108.20
84	Aa	968	A	N1-C6-N6	12.14	125.89	118.60
84	Aa	1172	A	N1-C6-N6	12.14	125.89	118.60
84	Aa	1835	A	N1-C6-N6	12.13	125.88	118.60
84	Aa	3299	A	N1-C6-N6	12.13	125.88	118.60
84	Aa	2114	A	N1-C6-N6	12.13	125.88	118.60
84	Aa	2371	A	N1-C6-N6	12.13	125.88	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1506	A	N1-C6-N6	12.12	125.87	118.60
1	Ad	263	C	O4'-C1'-N1	12.12	117.90	108.20
84	Aa	2677	A	N1-C6-N6	12.10	125.86	118.60
84	Aa	158	A	N1-C6-N6	12.10	125.86	118.60
84	Aa	1249	A	N1-C6-N6	12.10	125.86	118.60
86	Ab	74	A	N1-C6-N6	12.10	125.86	118.60
84	Aa	440	U	P-O3'-C3'	12.10	134.22	119.70
84	Aa	1599	A	N1-C6-N6	12.10	125.86	118.60
84	Aa	2706	A	N1-C6-N6	12.10	125.86	118.60
84	Aa	2674	A	N1-C6-N6	12.09	125.86	118.60
84	Aa	3213	A	N1-C6-N6	12.09	125.86	118.60
84	Aa	2662	A	N1-C6-N6	12.09	125.86	118.60
84	Aa	154	G	N1-C6-O6	12.09	127.15	119.90
84	Aa	2545	C	P-O3'-C3'	12.09	134.21	119.70
84	Aa	3310	A	N1-C6-N6	12.09	125.85	118.60
1	Ad	492	G	O4'-C1'-N9	12.08	117.87	108.20
1	Ad	1580	G	C1'-O4'-C4'	12.08	119.56	109.90
84	Aa	143	A	N1-C6-N6	12.08	125.85	118.60
84	Aa	1755	A	N1-C6-N6	12.08	125.85	118.60
84	Aa	2971	A	N1-C6-N6	12.08	125.85	118.60
84	Aa	2228	A	N1-C6-N6	12.08	125.85	118.60
84	Aa	2853	A	N1-C6-N6	12.07	125.84	118.60
84	Aa	197	A	N1-C6-N6	12.07	125.84	118.60
2	Ae	13	U	O4'-C1'-N1	12.07	117.86	108.20
1	Ad	1574	U	O4'-C1'-N1	12.07	117.85	108.20
84	Aa	540	G	C5-C6-O6	-12.06	121.36	128.60
84	Aa	1291	A	N1-C6-N6	12.06	125.84	118.60
84	Aa	87	A	N1-C6-N6	12.06	125.84	118.60
84	Aa	46	A	N1-C6-N6	12.06	125.83	118.60
84	Aa	292	A	N1-C6-N6	12.06	125.83	118.60
84	Aa	738	A	N1-C6-N6	12.05	125.83	118.60
84	Aa	186	A	N1-C6-N6	12.05	125.83	118.60
84	Aa	2694	A	N1-C6-N6	12.05	125.83	118.60
84	Aa	924	A	N1-C6-N6	12.04	125.83	118.60
84	Aa	2386	A	N1-C6-N6	12.04	125.82	118.60
84	Aa	2458	A	N1-C6-N6	12.04	125.83	118.60
84	Aa	1492	A	N1-C6-N6	12.04	125.82	118.60
84	Aa	1410	A	N1-C6-N6	12.03	125.82	118.60
84	Aa	1653	A	N1-C6-N6	12.03	125.82	118.60
1	Ad	1804	A	O4'-C1'-N9	12.03	117.82	108.20
84	Aa	1305	A	N1-C6-N6	12.03	125.82	118.60
84	Aa	2978	A	N1-C6-N6	12.03	125.82	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1485	A	N1-C6-N6	12.03	125.81	118.60
84	Aa	94	A	N1-C6-N6	12.02	125.81	118.60
84	Aa	1307	A	N1-C6-N6	12.02	125.81	118.60
84	Aa	219	A	N1-C6-N6	12.02	125.81	118.60
84	Aa	2733	A	N1-C6-N6	12.02	125.81	118.60
84	Aa	2429	A	N1-C6-N6	12.02	125.81	118.60
86	Ab	27	A	N1-C6-N6	12.02	125.81	118.60
1	Ad	279	C	O4'-C1'-N1	12.01	117.81	108.20
84	Aa	820	A	N1-C6-N6	12.01	125.81	118.60
84	Aa	1200	A	N1-C6-N6	12.01	125.81	118.60
84	Aa	1610	A	N1-C6-N6	12.00	125.80	118.60
84	Aa	3073	A	N1-C6-N6	12.00	125.80	118.60
84	Aa	2088	C	O4'-C1'-N1	11.99	117.80	108.20
84	Aa	2107	A	N1-C6-N6	11.99	125.80	118.60
84	Aa	1353	A	N1-C6-N6	11.99	125.79	118.60
84	Aa	1139	A	N1-C6-N6	11.98	125.79	118.60
84	Aa	2016	A	N1-C6-N6	11.98	125.79	118.60
84	Aa	660	A	N1-C6-N6	11.97	125.78	118.60
84	Aa	1138	A	N1-C6-N6	11.97	125.78	118.60
84	Aa	1306	A	N1-C6-N6	11.97	125.78	118.60
84	Aa	420	A	N1-C6-N6	11.96	125.78	118.60
84	Aa	1192	A	N1-C6-N6	11.96	125.77	118.60
84	Aa	1256	A	N1-C6-N6	11.95	125.77	118.60
1	Ad	1028	A	O4'-C1'-N9	11.95	117.76	108.20
84	Aa	62	A	N1-C6-N6	11.95	125.77	118.60
84	Aa	873	A	N1-C6-N6	11.95	125.77	118.60
84	Aa	1311	G	P-O3'-C3'	11.95	134.04	119.70
84	Aa	1790	A	N1-C6-N6	11.95	125.77	118.60
84	Aa	3173	A	N1-C6-N6	11.94	125.77	118.60
84	Aa	766	C	P-O3'-C3'	11.94	134.03	119.70
84	Aa	1395	A	N1-C6-N6	11.94	125.76	118.60
84	Aa	2718	A	N1-C6-N6	11.94	125.76	118.60
84	Aa	839	A	N1-C6-N6	11.94	125.76	118.60
84	Aa	2449	A	N1-C6-N6	11.94	125.76	118.60
1	Ad	1664	U	O4'-C1'-N1	11.93	117.75	108.20
84	Aa	730	A	N1-C6-N6	11.93	125.75	118.60
84	Aa	2244	G	P-O3'-C3'	11.92	134.01	119.70
84	Aa	1206	A	N1-C6-N6	11.92	125.75	118.60
84	Aa	1520	A	N1-C6-N6	11.92	125.75	118.60
84	Aa	3074	A	N1-C6-N6	11.92	125.75	118.60
84	Aa	1254	A	N1-C6-N6	11.91	125.75	118.60
84	Aa	1543	A	N1-C6-N6	11.91	125.75	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1846	A	N1-C6-N6	11.91	125.74	118.60
84	Aa	2574	A	N1-C6-N6	11.91	125.74	118.60
84	Aa	3022	A	N1-C6-N6	11.91	125.74	118.60
84	Aa	113	A	N1-C6-N6	11.90	125.74	118.60
84	Aa	2247	A	N1-C6-N6	11.90	125.74	118.60
84	Aa	1568	A	N1-C6-N6	11.90	125.74	118.60
84	Aa	1676	A	N1-C6-N6	11.90	125.74	118.60
84	Aa	1906	A	N1-C6-N6	11.90	125.74	118.60
84	Aa	2533	A	N1-C6-N6	11.88	125.73	118.60
85	Ac	104	A	N1-C6-N6	11.88	125.73	118.60
85	Ac	44	A	N1-C6-N6	11.88	125.73	118.60
84	Aa	1802	A	N1-C6-N6	11.88	125.73	118.60
84	Aa	1030	A	N1-C6-N6	11.87	125.72	118.60
84	Aa	2086	A	P-O5'-C5'	11.88	139.90	120.90
84	Aa	2761	A	N1-C6-N6	11.87	125.72	118.60
84	Aa	3123	A	N1-C6-N6	11.87	125.72	118.60
1	Ad	271	C	P-O3'-C3'	11.87	133.94	119.70
85	Ac	92	A	N1-C6-N6	11.87	125.72	118.60
84	Aa	98	A	N1-C6-N6	11.87	125.72	118.60
84	Aa	2681	A	N1-C6-N6	11.86	125.72	118.60
84	Aa	813	A	N1-C6-N6	11.85	125.71	118.60
84	Aa	1295	A	N1-C6-N6	11.85	125.71	118.60
84	Aa	1917	A	N1-C6-N6	11.85	125.71	118.60
84	Aa	1227	A	N1-C6-N6	11.85	125.71	118.60
84	Aa	2665	A	N1-C6-N6	11.84	125.71	118.60
84	Aa	3071	A	N1-C6-N6	11.84	125.71	118.60
84	Aa	376	A	N1-C6-N6	11.84	125.70	118.60
84	Aa	3312	G	N1-C6-O6	11.84	127.00	119.90
84	Aa	382	A	N1-C6-N6	11.84	125.70	118.60
84	Aa	1875	A	N1-C6-N6	11.84	125.70	118.60
84	Aa	2794	A	N1-C6-N6	11.83	125.70	118.60
84	Aa	1905	A	N1-C6-N6	11.83	125.70	118.60
84	Aa	2815	A	N1-C6-N6	11.83	125.70	118.60
84	Aa	439	A	N1-C6-N6	11.82	125.69	118.60
84	Aa	2774	A	N1-C6-N6	11.82	125.69	118.60
84	Aa	2203	A	N1-C6-N6	11.81	125.69	118.60
1	Ad	1788	G	O4'-C1'-N9	11.81	117.65	108.20
84	Aa	1255	A	N1-C6-N6	11.81	125.69	118.60
84	Aa	1970	A	N1-C6-N6	11.81	125.69	118.60
84	Aa	2839	A	N1-C6-N6	11.81	125.69	118.60
84	Aa	3135	A	N1-C6-N6	11.81	125.69	118.60
85	Ac	109	A	N1-C6-N6	11.81	125.69	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2092	C	O4'-C1'-N1	11.81	117.65	108.20
84	Aa	3088	A	N1-C6-N6	11.81	125.69	118.60
1	Ad	749	G	O4'-C1'-N9	11.81	117.65	108.20
84	Aa	1713	A	N1-C6-N6	11.80	125.68	118.60
84	Aa	1793	A	N1-C6-N6	11.80	125.68	118.60
86	Ab	22	A	N1-C6-N6	11.80	125.68	118.60
84	Aa	1809	A	N1-C6-N6	11.80	125.68	118.60
1	Ad	333	G	O4'-C1'-N9	11.80	117.64	108.20
84	Aa	928	A	N1-C6-N6	11.80	125.68	118.60
84	Aa	1455	A	N1-C6-N6	11.80	125.68	118.60
84	Aa	394	A	N1-C6-N6	11.80	125.68	118.60
84	Aa	2518	A	N1-C6-N6	11.79	125.68	118.60
84	Aa	3342	C	P-O3'-C3'	11.79	133.85	119.70
85	Ac	105	A	N1-C6-N6	11.79	125.67	118.60
84	Aa	1207	A	N1-C6-N6	11.79	125.67	118.60
84	Aa	1727	A	N1-C6-N6	11.79	125.67	118.60
1	Ad	1162	A	C1'-O4'-C4'	11.78	119.33	109.90
84	Aa	1812	A	N1-C6-N6	11.78	125.67	118.60
1	Ad	252	U	O4'-C1'-N1	11.78	117.62	108.20
84	Aa	844	A	N1-C6-N6	11.78	125.67	118.60
84	Aa	3028	A	N1-C6-N6	11.78	125.67	118.60
84	Aa	1560	A	P-O5'-C5'	11.78	139.75	120.90
1	Ad	505	U	O4'-C1'-N1	11.78	117.62	108.20
84	Aa	23	A	N1-C6-N6	11.78	125.67	118.60
1	Ad	1293	U	O4'-C1'-N1	11.77	117.62	108.20
84	Aa	216	G	N1-C6-O6	11.77	126.96	119.90
84	Aa	1330	A	N1-C6-N6	11.77	125.66	118.60
84	Aa	1376	A	N1-C6-N6	11.77	125.66	118.60
84	Aa	363	A	N1-C6-N6	11.77	125.66	118.60
84	Aa	698	A	N1-C6-N6	11.77	125.66	118.60
1	Ad	1524	A	O4'-C1'-N9	11.76	117.61	108.20
84	Aa	783	A	N1-C6-N6	11.76	125.66	118.60
84	Aa	1863	A	N1-C6-N6	11.76	125.66	118.60
84	Aa	2781	A	N1-C6-N6	11.76	125.66	118.60
84	Aa	2113	A	N1-C6-N6	11.76	125.66	118.60
84	Aa	416	A	N1-C6-N6	11.76	125.66	118.60
1	Ad	1190	U	O4'-C1'-N1	11.76	117.61	108.20
84	Aa	711	A	N1-C6-N6	11.75	125.65	118.60
84	Aa	2596	A	P-O3'-C3'	11.75	133.81	119.70
84	Aa	327	A	N1-C6-N6	11.75	125.65	118.60
84	Aa	1162	A	N1-C6-N6	11.75	125.65	118.60
84	Aa	2193	A	N1-C6-N6	11.75	125.65	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	564	A	N1-C6-N6	11.75	125.65	118.60
84	Aa	1674	A	N1-C6-N6	11.75	125.65	118.60
84	Aa	365	A	N1-C6-N6	11.74	125.64	118.60
84	Aa	287	A	N1-C6-N6	11.74	125.64	118.60
1	Ad	1782	C	P-O3'-C3'	11.73	133.78	119.70
84	Aa	149	A	N1-C6-N6	11.73	125.64	118.60
84	Aa	3362	A	N1-C6-N6	11.72	125.64	118.60
1	Ad	284	U	P-O3'-C3'	11.72	133.77	119.70
84	Aa	2298	A	N1-C6-N6	11.72	125.63	118.60
85	Ac	121	A	N1-C6-N6	11.72	125.63	118.60
84	Aa	2141	A	N1-C6-N6	11.72	125.63	118.60
84	Aa	1468	A	N1-C6-N6	11.72	125.63	118.60
84	Aa	1643	A	N1-C6-N6	11.72	125.63	118.60
84	Aa	3115	A	N1-C6-N6	11.72	125.63	118.60
84	Aa	1602	A	N1-C6-N6	11.71	125.63	118.60
1	Ad	1617	U	O4'-C1'-N1	11.71	117.57	108.20
84	Aa	1193	A	N1-C6-N6	11.71	125.63	118.60
84	Aa	3327	A	N1-C6-N6	11.71	125.62	118.60
84	Aa	2749	A	N1-C6-N6	11.71	125.62	118.60
84	Aa	3235	A	N1-C6-N6	11.70	125.62	118.60
1	Ad	608	U	O4'-C1'-N1	11.70	117.56	108.20
84	Aa	1312	A	N1-C6-N6	11.70	125.62	118.60
84	Aa	1837	A	N1-C6-N6	11.70	125.62	118.60
84	Aa	1065	A	N1-C6-N6	11.70	125.62	118.60
84	Aa	1883	A	N1-C6-N6	11.70	125.62	118.60
1	Ad	95	U	O4'-C1'-N1	11.69	117.56	108.20
84	Aa	3072	A	N1-C6-N6	11.69	125.62	118.60
1	Ad	282	C	C1'-O4'-C4'	11.69	119.25	109.90
84	Aa	3386	A	N1-C6-N6	11.69	125.61	118.60
1	Ad	612	U	N1-C1'-C2'	11.68	129.18	114.00
84	Aa	850	A	N1-C6-N6	11.68	125.61	118.60
84	Aa	2111	A	N1-C6-N6	11.68	125.61	118.60
1	Ad	903	A	O4'-C1'-N9	11.67	117.54	108.20
84	Aa	1204	A	N1-C6-N6	11.67	125.60	118.60
84	Aa	2006	A	N1-C6-N6	11.67	125.60	118.60
86	Ab	66	G	N1-C6-O6	11.67	126.90	119.90
1	Ad	57	G	O4'-C1'-N9	11.66	117.53	108.20
84	Aa	572	U	P-O3'-C3'	11.66	133.69	119.70
84	Aa	2218	A	N1-C6-N6	11.66	125.59	118.60
84	Aa	1861	A	N1-C6-N6	11.65	125.59	118.60
84	Aa	325	A	N1-C6-N6	11.65	125.59	118.60
84	Aa	1591	A	N1-C6-N6	11.63	125.58	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1056	A	P-O3'-C3'	11.63	133.66	119.70
84	Aa	2251	A	N1-C6-N6	11.63	125.58	118.60
84	Aa	2960	A	N1-C6-N6	11.63	125.58	118.60
84	Aa	2765	A	N1-C6-N6	11.63	125.58	118.60
84	Aa	681	A	N1-C6-N6	11.63	125.58	118.60
1	Ad	282	C	P-O3'-C3'	11.62	133.65	119.70
84	Aa	1397	A	N1-C6-N6	11.62	125.57	118.60
84	Aa	1880	A	N1-C6-N6	11.62	125.57	118.60
84	Aa	2366	A	N1-C6-N6	11.61	125.57	118.60
84	Aa	249	A	N1-C6-N6	11.61	125.57	118.60
84	Aa	1926	A	N1-C6-N6	11.61	125.57	118.60
1	Ad	1267	G	O4'-C1'-N9	11.61	117.49	108.20
84	Aa	1738	A	N1-C6-N6	11.61	125.57	118.60
1	Ad	1582	G	C3'-C2'-C1'	11.61	110.78	101.50
84	Aa	981	A	N1-C6-N6	11.61	125.56	118.60
86	Ab	83	A	N1-C6-N6	11.61	125.56	118.60
84	Aa	1220	G	N1-C6-O6	11.60	126.86	119.90
85	Ac	89	A	N1-C6-N6	11.60	125.56	118.60
84	Aa	2736	A	N1-C6-N6	11.60	125.56	118.60
84	Aa	1278	A	N1-C6-N6	11.60	125.56	118.60
84	Aa	2840	A	N1-C6-N6	11.60	125.56	118.60
1	Ad	54	C	O4'-C1'-C2'	-11.59	94.21	105.80
84	Aa	2772	A	N1-C6-N6	11.59	125.55	118.60
84	Aa	2921	A	N1-C6-N6	11.59	125.55	118.60
1	Ad	1323	U	O4'-C1'-N1	11.58	117.47	108.20
84	Aa	1101	A	N1-C6-N6	11.58	125.55	118.60
2	Ae	65	U	O4'-C1'-N1	11.58	117.46	108.20
2	Ae	59	U	O4'-C1'-N1	11.57	117.46	108.20
1	Ad	817	C	P-O3'-C3'	11.57	133.59	119.70
84	Aa	2120	A	N1-C6-N6	11.57	125.54	118.60
84	Aa	1507	A	N1-C6-N6	11.57	125.54	118.60
1	Ad	1276	U	O4'-C1'-N1	11.56	117.45	108.20
1	Ad	743	G	O4'-C1'-N9	11.56	117.45	108.20
84	Aa	775	A	N1-C6-N6	11.56	125.54	118.60
84	Aa	2532	A	N1-C6-N6	11.56	125.54	118.60
84	Aa	53	C	O4'-C1'-N1	11.56	117.45	108.20
84	Aa	1571	A	N1-C6-N6	11.56	125.54	118.60
84	Aa	2835	A	N1-C6-N6	11.56	125.53	118.60
1	Ad	112	U	O4'-C1'-N1	11.55	117.44	108.20
84	Aa	932	A	N1-C6-N6	11.55	125.53	118.60
84	Aa	2474	A	N1-C6-N6	11.55	125.53	118.60
1	Ad	553	G	O4'-C1'-N9	11.55	117.44	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	355	U	O4'-C1'-N1	11.55	117.44	108.20
1	Ad	74	U	O4'-C1'-N1	11.54	117.44	108.20
2	Ae	42	C	O4'-C1'-N1	11.54	117.43	108.20
84	Aa	2054	A	N1-C6-N6	11.54	125.52	118.60
1	Ad	1576	C	P-O3'-C3'	11.54	133.54	119.70
84	Aa	2514	A	P-O3'-C3'	11.54	133.54	119.70
84	Aa	2412	A	N1-C6-N6	11.53	125.52	118.60
84	Aa	2202	A	N1-C6-N6	11.53	125.52	118.60
84	Aa	452	G	C4'-C3'-O3'	-11.53	85.20	109.40
84	Aa	3033	A	N1-C6-N6	11.53	125.52	118.60
84	Aa	1795	A	N1-C6-N6	11.52	125.51	118.60
86	Ab	20	C	N3-C4-C5	-11.52	117.29	121.90
84	Aa	2641	A	N1-C6-N6	11.51	125.50	118.60
85	Ac	53	A	N1-C6-N6	11.51	125.50	118.60
84	Aa	47	A	N1-C6-N6	11.50	125.50	118.60
1	Ad	1395	C	N1-C1'-C2'	11.50	128.95	114.00
1	Ad	509	A	C3'-C2'-C1'	11.50	110.70	101.50
84	Aa	1748	A	N1-C6-N6	11.50	125.50	118.60
1	Ad	156	U	P-O3'-C3'	11.50	133.50	119.70
84	Aa	122	A	N1-C6-N6	11.50	125.50	118.60
84	Aa	72	A	N1-C6-N6	11.50	125.50	118.60
84	Aa	1518	A	N1-C6-N6	11.50	125.50	118.60
84	Aa	885	A	N1-C6-N6	11.49	125.50	118.60
1	Ad	360	G	O4'-C1'-N9	11.49	117.39	108.20
84	Aa	550	C	P-O3'-C3'	11.49	133.49	119.70
84	Aa	1459	A	N1-C6-N6	11.49	125.49	118.60
84	Aa	2137	A	N1-C6-N6	11.49	125.49	118.60
84	Aa	670	A	N1-C6-N6	11.49	125.49	118.60
84	Aa	2813	A	N1-C6-N6	11.49	125.49	118.60
1	Ad	176	A	C1'-O4'-C4'	11.48	119.09	109.90
84	Aa	1163	A	N1-C6-N6	11.48	125.49	118.60
1	Ad	1556	U	O4'-C1'-N1	11.48	117.39	108.20
84	Aa	1860	A	N1-C6-N6	11.48	125.49	118.60
84	Aa	1107	G	N1-C6-O6	11.48	126.79	119.90
84	Aa	2250	A	N1-C6-N6	11.47	125.48	118.60
84	Aa	696	A	N1-C6-N6	11.46	125.48	118.60
84	Aa	2304	A	N1-C6-N6	11.46	125.48	118.60
1	Ad	744	G	N9-C1'-C2'	-11.46	99.10	114.00
85	Ac	135	A	N1-C6-N6	11.46	125.48	118.60
1	Ad	80	C	N1-C1'-C2'	11.45	128.89	114.00
1	Ad	262	U	O4'-C1'-N1	11.45	117.36	108.20
84	Aa	2527	G	N1-C6-O6	11.45	126.77	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1691	C	O4'-C1'-C2'	-11.45	94.35	105.80
84	Aa	210	G	N1-C6-O6	11.45	126.77	119.90
84	Aa	1391	A	N1-C6-N6	11.44	125.47	118.60
84	Aa	861	A	N1-C6-N6	11.44	125.46	118.60
84	Aa	2081	C	P-O3'-C3'	11.43	133.42	119.70
84	Aa	1550	A	N1-C6-N6	11.43	125.46	118.60
84	Aa	2215	A	N1-C6-N6	11.43	125.46	118.60
84	Aa	2750	A	N1-C6-N6	11.43	125.46	118.60
84	Aa	1464	A	N1-C6-N6	11.43	125.45	118.60
84	Aa	2275	A	N1-C6-N6	11.43	125.46	118.60
84	Aa	3309	U	P-O3'-C3'	11.42	133.41	119.70
85	Ac	48	A	N1-C6-N6	11.42	125.45	118.60
1	Ad	845	C	O4'-C1'-C2'	-11.42	94.38	105.80
1	Ad	1580	G	N9-C1'-C2'	-11.42	99.16	114.00
1	Ad	1388	A	O4'-C1'-N9	11.41	117.33	108.20
86	Ab	89	G	O4'-C1'-N9	11.41	117.33	108.20
1	Ad	1361	G	O4'-C1'-N9	11.41	117.33	108.20
84	Aa	66	A	N1-C6-N6	11.41	125.44	118.60
84	Aa	2162	C	O4'-C1'-N1	11.41	117.33	108.20
84	Aa	1805	A	N1-C6-N6	11.40	125.44	118.60
86	Ab	90	A	N1-C6-N6	11.40	125.44	118.60
84	Aa	875	A	N1-C6-N6	11.40	125.44	118.60
84	Aa	1584	A	N1-C6-N6	11.40	125.44	118.60
84	Aa	1629	A	N1-C6-N6	11.39	125.44	118.60
84	Aa	758	A	N1-C6-N6	11.39	125.44	118.60
84	Aa	2502	U	O4'-C1'-N1	11.39	117.31	108.20
1	Ad	846	U	N1-C1'-C2'	11.39	128.80	114.00
84	Aa	970	A	N1-C6-N6	11.39	125.43	118.60
84	Aa	2312	A	N1-C6-N6	11.39	125.43	118.60
84	Aa	976	A	N1-C6-N6	11.39	125.43	118.60
1	Ad	860	A	N9-C1'-C2'	-11.38	99.20	114.00
84	Aa	48	A	N1-C6-N6	11.38	125.43	118.60
1	Ad	1259	G	O4'-C1'-C2'	11.37	117.84	107.60
84	Aa	869	A	N1-C6-N6	11.37	125.42	118.60
86	Ab	72	G	P-O3'-C3'	11.37	133.34	119.70
86	Ab	97	G	C5-C6-O6	-11.37	121.78	128.60
84	Aa	2699	A	N1-C6-N6	11.37	125.42	118.60
84	Aa	1199	A	N1-C6-N6	11.37	125.42	118.60
84	Aa	2631	A	N1-C6-N6	11.37	125.42	118.60
84	Aa	405	A	N1-C6-N6	11.36	125.42	118.60
84	Aa	2367	A	N1-C6-N6	11.36	125.42	118.60
84	Aa	2819	A	N1-C6-N6	11.36	125.42	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	211	A	N1-C6-N6	11.35	125.41	118.60
84	Aa	585	A	N1-C6-N6	11.35	125.41	118.60
84	Aa	2596	A	N1-C6-N6	11.35	125.41	118.60
1	Ad	54	C	O4'-C1'-N1	11.34	117.27	108.20
84	Aa	2175	A	N1-C6-N6	11.34	125.41	118.60
84	Aa	1874	A	N1-C6-N6	11.34	125.40	118.60
84	Aa	2709	G	N1-C6-O6	11.34	126.70	119.90
84	Aa	1527	A	N1-C6-N6	11.34	125.40	118.60
84	Aa	2362	A	N1-C6-N6	11.33	125.40	118.60
84	Aa	2074	C	O4'-C1'-N1	11.33	117.26	108.20
1	Ad	155	A	O4'-C1'-N9	11.32	117.26	108.20
84	Aa	1024	G	C5-C6-O6	-11.32	121.81	128.60
84	Aa	2316	A	N1-C6-N6	11.32	125.39	118.60
84	Aa	2047	A	N1-C6-N6	11.32	125.39	118.60
84	Aa	2208	A	N1-C6-N6	11.32	125.39	118.60
84	Aa	918	A	N1-C6-N6	11.31	125.39	118.60
84	Aa	1229	A	N1-C6-N6	11.31	125.39	118.60
84	Aa	3345	G	P-O3'-C3'	11.31	133.28	119.70
1	Ad	1238	A	P-O3'-C3'	11.31	133.27	119.70
1	Ad	132	G	O4'-C1'-N9	11.31	117.25	108.20
84	Aa	3128	A	N1-C6-N6	11.31	125.39	118.60
86	Ab	51	G	P-O3'-C3'	11.31	133.27	119.70
84	Aa	916	A	N1-C6-N6	11.30	125.38	118.60
84	Aa	1600	A	N1-C6-N6	11.30	125.38	118.60
84	Aa	3236	A	N1-C6-N6	11.30	125.38	118.60
1	Ad	1810	G	O4'-C1'-C2'	-11.30	94.50	105.80
84	Aa	2053	A	N1-C6-N6	11.30	125.38	118.60
84	Aa	2542	U	P-O3'-C3'	11.30	133.26	119.70
1	Ad	964	U	N1-C1'-C2'	11.29	128.68	114.00
84	Aa	1990	A	N1-C6-N6	11.29	125.37	118.60
1	Ad	1009	U	O4'-C1'-N1	11.29	117.23	108.20
84	Aa	1264	A	N1-C6-N6	11.29	125.37	118.60
84	Aa	2973	A	N1-C6-N6	11.29	125.37	118.60
84	Aa	1457	A	N1-C6-N6	11.28	125.37	118.60
84	Aa	651	A	N1-C6-N6	11.27	125.36	118.60
84	Aa	1182	A	N1-C6-N6	11.27	125.36	118.60
1	Ad	1466	A	O4'-C1'-C2'	11.27	117.74	107.60
84	Aa	1971	A	N1-C6-N6	11.27	125.36	118.60
84	Aa	3009	A	N1-C6-N6	11.27	125.36	118.60
84	Aa	1747	A	C5-C6-N6	-11.26	114.69	123.70
1	Ad	491	G	O4'-C1'-N9	11.26	117.21	108.20
84	Aa	1044	A	N1-C6-N6	11.26	125.36	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2822	A	N1-C6-N6	11.26	125.35	118.60
84	Aa	704	G	N1-C6-O6	11.25	126.65	119.90
1	Ad	1589	C	C3'-C2'-C1'	11.25	110.50	101.50
84	Aa	789	A	N1-C6-N6	11.25	125.35	118.60
1	Ad	52	U	O4'-C1'-N1	11.25	117.20	108.20
84	Aa	567	G	N1-C6-O6	11.25	126.65	119.90
84	Aa	1937	C	P-O3'-C3'	11.25	133.19	119.70
84	Aa	1059	A	N1-C6-N6	11.24	125.35	118.60
84	Aa	1703	C	O4'-C1'-N1	11.24	117.19	108.20
1	Ad	1029	U	O4'-C1'-N1	11.24	117.19	108.20
1	Ad	71	C	P-O3'-C3'	11.23	133.18	119.70
84	Aa	1275	A	N1-C6-N6	11.23	125.34	118.60
84	Aa	2489	A	N1-C6-N6	11.23	125.34	118.60
1	Ad	907	G	O4'-C1'-N9	11.22	117.18	108.20
84	Aa	2612	A	N1-C6-N6	11.22	125.33	118.60
84	Aa	3007	A	N1-C6-N6	11.22	125.33	118.60
1	Ad	64	U	O4'-C1'-N1	11.22	117.17	108.20
84	Aa	70	A	N1-C6-N6	11.22	125.33	118.60
1	Ad	1681	G	O4'-C1'-N9	11.22	117.17	108.20
84	Aa	1689	G	N1-C6-O6	11.21	126.63	119.90
84	Aa	527	G	P-O5'-C5'	11.20	138.83	120.90
1	Ad	613	U	O4'-C1'-C2'	-11.20	94.60	105.80
84	Aa	731	G	N1-C6-O6	11.20	126.62	119.90
84	Aa	954	A	N1-C6-N6	11.20	125.32	118.60
84	Aa	3297	A	N1-C6-N6	11.20	125.32	118.60
1	Ad	1409	G	P-O5'-C5'	11.18	138.79	120.90
1	Ad	1389	G	O4'-C1'-N9	11.18	117.14	108.20
85	Ac	61	A	N1-C6-N6	11.18	125.31	118.60
1	Ad	578	G	O4'-C1'-N9	11.18	117.14	108.20
84	Aa	1026	A	N1-C6-N6	11.17	125.30	118.60
84	Aa	167	C	P-O3'-C3'	11.16	133.10	119.70
86	Ab	31	G	C5-C6-O6	-11.16	121.90	128.60
84	Aa	2149	G	P-O3'-C3'	11.16	133.09	119.70
84	Aa	616	A	N1-C6-N6	11.16	125.30	118.60
84	Aa	2509	A	N1-C6-N6	11.15	125.29	118.60
1	Ad	525	A	O4'-C1'-N9	11.15	117.12	108.20
84	Aa	767	U	P-O3'-C3'	11.15	133.08	119.70
84	Aa	1716	G	N1-C6-O6	11.15	126.59	119.90
84	Aa	868	A	N1-C6-N6	11.14	125.29	118.60
84	Aa	1052	A	N1-C6-N6	11.14	125.29	118.60
84	Aa	76	A	N1-C6-N6	11.14	125.28	118.60
84	Aa	2764	G	N1-C6-O6	11.14	126.58	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	Af	19	U	O4'-C1'-N1	11.14	117.11	108.20
84	Aa	2737	A	N1-C6-N6	11.13	125.28	118.60
84	Aa	1235	A	N1-C6-N6	11.13	125.28	118.60
1	Ad	888	U	O4'-C1'-N1	11.13	117.10	108.20
84	Aa	293	A	N1-C6-N6	11.13	125.28	118.60
1	Ad	284	U	O4'-C1'-N1	11.12	117.10	108.20
86	Ab	100	A	N1-C6-N6	11.12	125.28	118.60
84	Aa	3287	A	N1-C6-N6	11.12	125.27	118.60
84	Aa	3308	A	N1-C6-N6	11.11	125.27	118.60
84	Aa	1321	A	N1-C6-N6	11.11	125.27	118.60
84	Aa	2132	A	N1-C6-N6	11.10	125.26	118.60
84	Aa	3167	G	P-O3'-C3'	11.10	133.02	119.70
84	Aa	198	A	N1-C6-N6	11.10	125.26	118.60
84	Aa	2792	A	N1-C6-N6	11.09	125.26	118.60
84	Aa	3140	A	N1-C6-N6	11.09	125.26	118.60
1	Ad	67	G	P-O3'-C3'	11.09	133.00	119.70
84	Aa	387	A	N1-C6-N6	11.09	125.25	118.60
1	Ad	241	G	P-O3'-C3'	11.09	133.00	119.70
84	Aa	2705	A	N1-C6-N6	11.08	125.25	118.60
84	Aa	2928	A	N1-C6-N6	11.08	125.25	118.60
86	Ab	45	U	O4'-C1'-N1	11.08	117.06	108.20
1	Ad	1081	A	O4'-C1'-N9	11.08	117.06	108.20
84	Aa	2881	C	O4'-C1'-N1	11.08	117.06	108.20
1	Ad	1247	G	O4'-C1'-N9	11.07	117.06	108.20
84	Aa	653	A	N1-C6-N6	11.06	125.24	118.60
85	Ac	82	C	P-O3'-C3'	11.06	132.97	119.70
84	Aa	650	A	N1-C6-N6	11.06	125.23	118.60
1	Ad	316	A	O4'-C1'-N9	11.05	117.04	108.20
85	Ac	88	A	N1-C6-N6	11.05	125.23	118.60
84	Aa	2938	A	N1-C6-N6	11.04	125.22	118.60
1	Ad	1186	U	O4'-C1'-N1	11.04	117.03	108.20
1	Ad	1195	U	O4'-C1'-N1	11.04	117.03	108.20
84	Aa	2796	G	N1-C6-O6	11.04	126.52	119.90
84	Aa	1577	A	N1-C6-N6	11.03	125.22	118.60
84	Aa	2568	G	N1-C6-O6	11.03	126.52	119.90
1	Ad	985	G	O4'-C1'-N9	11.03	117.03	108.20
1	Ad	724	U	O4'-C1'-N1	11.03	117.02	108.20
1	Ad	1085	U	O4'-C1'-N1	11.02	117.02	108.20
86	Ab	110	G	C5-C6-O6	-11.02	121.99	128.60
1	Ad	1125	U	O4'-C1'-N1	11.02	117.01	108.20
84	Aa	2660	A	N1-C6-N6	11.01	125.21	118.60
1	Ad	1357	U	O4'-C1'-N1	11.01	117.01	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	350	A	N1-C6-N6	11.01	125.20	118.60
84	Aa	1115	A	N1-C6-N6	10.99	125.20	118.60
1	Ad	262	U	C1'-O4'-C4'	10.99	118.69	109.90
1	Ad	1658	U	O4'-C1'-N1	10.99	116.99	108.20
1	Ad	621	U	O4'-C1'-N1	10.98	116.99	108.20
86	Ab	79	A	C4-C5-C6	10.98	122.49	117.00
84	Aa	2325	A	N1-C6-N6	10.98	125.19	118.60
84	Aa	340	A	N1-C6-N6	10.98	125.19	118.60
84	Aa	984	A	N1-C6-N6	10.98	125.19	118.60
84	Aa	2079	A	N1-C6-N6	10.97	125.18	118.60
84	Aa	2094	A	C4'-C3'-O3'	-10.97	86.36	109.40
84	Aa	1635	A	N1-C6-N6	10.96	125.18	118.60
86	Ab	60	G	N1-C6-O6	10.96	126.47	119.90
1	Ad	940	U	O4'-C1'-N1	10.96	116.97	108.20
84	Aa	1051	A	N1-C6-N6	10.95	125.17	118.60
84	Aa	306	A	N1-C6-N6	10.95	125.17	118.60
84	Aa	288	G	N1-C6-O6	10.95	126.47	119.90
85	Ac	140	A	N1-C6-N6	10.95	125.17	118.60
84	Aa	615	A	N1-C6-N6	10.94	125.16	118.60
84	Aa	819	A	N1-C6-N6	10.93	125.16	118.60
85	Ac	79	A	N1-C6-N6	10.93	125.16	118.60
1	Ad	968	A	O4'-C1'-N9	10.92	116.94	108.20
85	Ac	77	A	N1-C6-N6	10.91	125.15	118.60
84	Aa	16	A	N1-C6-N6	10.91	125.15	118.60
84	Aa	3124	A	N1-C6-N6	10.91	125.14	118.60
84	Aa	1015	A	N1-C6-N6	10.91	125.14	118.60
85	Ac	129	C	P-O3'-C3'	10.90	132.78	119.70
1	Ad	107	U	O4'-C1'-N1	10.90	116.92	108.20
84	Aa	2661	G	N1-C6-O6	10.90	126.44	119.90
1	Ad	1625	U	O4'-C1'-N1	10.89	116.91	108.20
84	Aa	397	A	N1-C6-N6	10.89	125.14	118.60
84	Aa	1798	C	P-O3'-C3'	10.89	132.77	119.70
84	Aa	2936	A	N1-C6-N6	10.89	125.13	118.60
85	Ac	80	A	N1-C6-N6	10.89	125.13	118.60
1	Ad	212	A	C3'-C2'-C1'	10.89	110.21	101.50
84	Aa	2398	A	N1-C6-N6	10.88	125.13	118.60
84	Aa	2514	A	N1-C6-N6	10.88	125.13	118.60
84	Aa	2707	A	N1-C6-N6	10.88	125.13	118.60
1	Ad	870	A	O4'-C1'-N9	10.88	116.90	108.20
84	Aa	2254	A	N1-C6-N6	10.88	125.13	118.60
84	Aa	2307	A	N1-C6-N6	10.87	125.12	118.60
1	Ad	334	G	O4'-C1'-N9	10.86	116.89	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	72	A	P-O3'-C3'	10.86	132.73	119.70
86	Ab	65	G	N1-C6-O6	10.86	126.42	119.90
1	Ad	1635	U	O4'-C1'-N1	10.86	116.89	108.20
84	Aa	3368	A	N1-C6-N6	10.86	125.11	118.60
1	Ad	476	U	O4'-C1'-N1	10.85	116.88	108.20
86	Ab	103	U	O4'-C1'-N1	10.85	116.88	108.20
84	Aa	1537	A	N1-C6-N6	10.85	125.11	118.60
86	Ab	32	A	N1-C6-N6	10.85	125.11	118.60
84	Aa	1415	G	N1-C6-O6	10.84	126.40	119.90
2	Ae	40	U	O4'-C1'-N1	10.84	116.87	108.20
1	Ad	1808	U	P-O3'-C3'	10.83	132.70	119.70
1	Ad	77	G	O4'-C1'-N9	10.83	116.86	108.20
84	Aa	1712	A	N1-C6-N6	10.83	125.10	118.60
84	Aa	1470	A	N1-C6-N6	10.82	125.09	118.60
84	Aa	1714	A	N1-C6-N6	10.82	125.09	118.60
84	Aa	2246	G	N1-C6-O6	10.82	126.39	119.90
85	Ac	85	G	N1-C6-O6	10.82	126.39	119.90
84	Aa	1181	A	N1-C6-N6	10.82	125.09	118.60
84	Aa	1361	G	N1-C6-O6	10.82	126.39	119.90
1	Ad	1386	U	N1-C1'-C2'	10.81	128.06	114.00
84	Aa	3188	G	N1-C6-O6	10.81	126.39	119.90
84	Aa	1478	A	N1-C6-N6	10.80	125.08	118.60
85	Ac	151	G	N1-C6-O6	10.80	126.38	119.90
1	Ad	498	U	O4'-C1'-N1	10.80	116.84	108.20
84	Aa	1720	C	O4'-C1'-N1	10.80	116.84	108.20
84	Aa	2270	A	N1-C6-N6	10.80	125.08	118.60
1	Ad	824	U	P-O5'-C5'	10.80	138.18	120.90
1	Ad	1603	U	O4'-C1'-N1	10.80	116.84	108.20
84	Aa	2224	A	N1-C6-N6	10.80	125.08	118.60
1	Ad	241	G	O4'-C1'-N9	10.79	116.83	108.20
1	Ad	1535	U	O4'-C1'-N1	10.79	116.83	108.20
84	Aa	1705	A	N1-C6-N6	10.78	125.07	118.60
84	Aa	1197	A	N1-C6-N6	10.78	125.07	118.60
84	Aa	1569	U	P-O3'-C3'	10.78	132.63	119.70
84	Aa	118	G	C5-C6-O6	-10.77	122.14	128.60
85	Ac	17	A	N1-C6-N6	10.76	125.06	118.60
1	Ad	1745	U	O4'-C1'-N1	10.76	116.81	108.20
84	Aa	3282	G	N1-C6-O6	10.76	126.36	119.90
84	Aa	171	G	N1-C6-O6	10.76	126.36	119.90
1	Ad	39	A	O4'-C1'-N9	10.75	116.80	108.20
1	Ad	1637	G	O4'-C1'-C2'	10.75	117.28	107.60
84	Aa	1741	G	N1-C6-O6	10.75	126.35	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	336	A	N1-C6-N6	10.74	125.05	118.60
1	Ad	1065	A	O4'-C1'-N9	10.74	116.79	108.20
84	Aa	2588	G	N1-C6-O6	10.74	126.34	119.90
1	Ad	987	U	O4'-C1'-N1	10.73	116.79	108.20
1	Ad	421	A	P-O3'-C3'	10.73	132.57	119.70
84	Aa	88	A	N1-C6-N6	10.73	125.04	118.60
1	Ad	421	A	O4'-C1'-C2'	-10.72	95.08	105.80
84	Aa	1294	A	N1-C6-N6	10.72	125.03	118.60
85	Ac	122	G	N1-C6-O6	10.71	126.33	119.90
84	Aa	2119	A	N1-C6-N6	10.71	125.03	118.60
84	Aa	2423	A	N1-C6-N6	10.71	125.03	118.60
84	Aa	323	A	N1-C6-N6	10.70	125.02	118.60
84	Aa	899	A	N1-C6-N6	10.70	125.02	118.60
1	Ad	1290	U	O4'-C1'-N1	10.70	116.76	108.20
2	Ae	32	U	O4'-C1'-N1	10.70	116.76	108.20
84	Aa	2020	G	P-O3'-C3'	10.69	132.53	119.70
1	Ad	772	C	N1-C1'-C2'	10.68	127.89	114.00
84	Aa	372	A	N1-C6-N6	10.68	125.01	118.60
84	Aa	1477	A	N1-C6-N6	10.68	125.01	118.60
1	Ad	739	U	O4'-C1'-N1	10.68	116.74	108.20
84	Aa	1007	A	N1-C6-N6	10.68	125.01	118.60
84	Aa	2291	A	N1-C6-N6	10.68	125.01	118.60
84	Aa	2671	A	N1-C6-N6	10.68	125.01	118.60
86	Ab	37	G	N1-C6-O6	10.67	126.30	119.90
1	Ad	129	U	O4'-C1'-N1	10.66	116.73	108.20
1	Ad	633	U	P-O3'-C3'	10.66	132.50	119.70
84	Aa	384	A	N1-C6-N6	10.66	125.00	118.60
84	Aa	1593	C	O4'-C1'-N1	10.66	116.73	108.20
84	Aa	3131	A	N1-C6-N6	10.66	125.00	118.60
84	Aa	2892	A	N1-C6-N6	10.66	125.00	118.60
84	Aa	3200	A	N1-C6-N6	10.66	124.99	118.60
86	Ab	8	A	N1-C6-N6	10.64	124.99	118.60
84	Aa	1544	G	P-O3'-C3'	10.64	132.47	119.70
84	Aa	154	G	C5-C6-O6	-10.64	122.22	128.60
84	Aa	172	A	N1-C6-N6	10.64	124.98	118.60
86	Ab	92	C	N3-C4-C5	-10.63	117.65	121.90
84	Aa	967	G	N1-C6-O6	10.62	126.27	119.90
84	Aa	1320	G	N1-C6-O6	10.62	126.27	119.90
84	Aa	362	G	N1-C6-O6	10.62	126.27	119.90
84	Aa	718	C	O4'-C1'-N1	10.62	116.69	108.20
84	Aa	1911	A	N1-C6-N6	10.62	124.97	118.60
84	Aa	2623	G	N1-C6-O6	10.62	126.27	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1435	G	O4'-C1'-N9	10.61	116.69	108.20
84	Aa	105	A	N1-C6-N6	10.61	124.97	118.60
84	Aa	1887	A	N1-C6-N6	10.61	124.97	118.60
84	Aa	1237	G	N1-C6-O6	10.61	126.26	119.90
84	Aa	2593	A	N1-C6-N6	10.60	124.96	118.60
84	Aa	1248	A	N1-C6-N6	10.59	124.96	118.60
84	Aa	1943	G	C4'-C3'-O3'	10.59	134.18	113.00
84	Aa	1286	G	N1-C6-O6	10.58	126.25	119.90
84	Aa	1680	A	N1-C6-N6	10.58	124.95	118.60
84	Aa	2743	A	N1-C6-N6	10.58	124.95	118.60
1	Ad	1202	G	O4'-C1'-C2'	10.57	117.12	107.60
84	Aa	2354	G	N1-C6-O6	10.57	126.24	119.90
84	Aa	2360	A	N1-C6-N6	10.57	124.94	118.60
1	Ad	33	U	O4'-C1'-N1	10.57	116.65	108.20
84	Aa	1471	A	N1-C6-N6	10.57	124.94	118.60
84	Aa	3261	C	C4'-C3'-O3'	10.57	134.13	113.00
86	Ab	43	A	C5-C6-N6	-10.57	115.25	123.70
1	Ad	1255	U	O4'-C1'-N1	10.56	116.64	108.20
84	Aa	237	C	P-O3'-C3'	10.56	132.37	119.70
86	Ab	41	G	N1-C6-O6	10.55	126.23	119.90
84	Aa	1891	A	N1-C6-N6	10.55	124.93	118.60
1	Ad	1231	A	N9-C1'-C2'	10.54	127.70	114.00
84	Aa	138	G	N1-C6-O6	10.54	126.22	119.90
84	Aa	876	C	P-O3'-C3'	10.54	132.35	119.70
1	Ad	1336	C	N1-C1'-C2'	10.54	127.70	114.00
84	Aa	84	A	N1-C6-N6	10.54	124.92	118.60
1	Ad	715	U	O4'-C1'-N1	10.54	116.63	108.20
1	Ad	866	U	O4'-C1'-N1	10.53	116.63	108.20
84	Aa	2942	A	N1-C6-N6	10.54	124.92	118.60
86	Ab	19	A	C5-C6-N1	-10.54	112.43	117.70
84	Aa	1856	G	N1-C6-O6	10.53	126.22	119.90
84	Aa	1092	G	N1-C6-O6	10.53	126.22	119.90
84	Aa	1586	A	N1-C6-N6	10.53	124.92	118.60
1	Ad	764	U	O4'-C1'-N1	10.52	116.62	108.20
1	Ad	1105	G	N9-C1'-C2'	10.52	127.68	114.00
84	Aa	1585	A	N1-C6-N6	10.52	124.91	118.60
84	Aa	1047	C	O4'-C1'-N1	10.52	116.62	108.20
84	Aa	2605	G	N1-C6-O6	10.52	126.21	119.90
84	Aa	2801	A	N1-C6-N6	10.51	124.91	118.60
84	Aa	3047	A	N1-C6-N6	10.51	124.91	118.60
1	Ad	1321	C	C1'-O4'-C4'	-10.51	101.50	109.90
84	Aa	1322	A	C5-C6-N6	-10.50	115.30	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2265	A	N1-C6-N6	10.50	124.90	118.60
84	Aa	2919	G	N1-C6-O6	10.50	126.20	119.90
84	Aa	1217	G	N1-C6-O6	10.50	126.20	119.90
84	Aa	2222	C	P-O3'-C3'	10.49	132.29	119.70
84	Aa	2739	A	N1-C6-N6	10.49	124.89	118.60
1	Ad	831	C	O4'-C1'-N1	10.49	116.59	108.20
84	Aa	2319	A	N1-C6-N6	10.49	124.89	118.60
1	Ad	130	A	O4'-C1'-N9	10.48	116.59	108.20
84	Aa	346	A	N1-C6-N6	10.48	124.89	118.60
84	Aa	2136	A	N1-C6-N6	10.48	124.89	118.60
1	Ad	1162	A	N9-C1'-C2'	-10.48	100.38	114.00
84	Aa	1998	A	N1-C6-N6	10.48	124.89	118.60
84	Aa	723	G	C5'-C4'-O4'	-10.47	96.54	109.10
84	Aa	1019	A	C5-C6-N6	-10.47	115.32	123.70
1	Ad	872	G	O4'-C1'-N9	10.46	116.56	108.20
84	Aa	2697	A	N1-C6-N6	10.46	124.87	118.60
1	Ad	110	G	O4'-C1'-N9	10.45	116.56	108.20
1	Ad	1328	G	O4'-C1'-N9	10.45	116.56	108.20
84	Aa	359	A	N1-C6-N6	10.45	124.87	118.60
84	Aa	2374	G	N1-C6-O6	10.44	126.17	119.90
84	Aa	3225	G	N1-C6-O6	10.43	126.16	119.90
1	Ad	970	U	N1-C1'-C2'	10.43	127.56	114.00
1	Ad	1220	C	O4'-C1'-C2'	-10.43	95.37	105.80
84	Aa	2174	C	O4'-C1'-N1	10.43	116.54	108.20
1	Ad	1487	U	P-O3'-C3'	10.43	132.21	119.70
84	Aa	2513	U	C2'-C3'-O3'	10.43	132.44	109.50
86	Ab	86	G	N1-C6-O6	10.42	126.15	119.90
1	Ad	1161	C	N1-C1'-C2'	10.42	127.54	114.00
1	Ad	846	U	C1'-O4'-C4'	-10.41	101.57	109.90
1	Ad	731	G	O4'-C1'-N9	10.41	116.53	108.20
84	Aa	3179	G	N1-C6-O6	10.41	126.15	119.90
84	Aa	2257	A	N1-C6-N6	10.41	124.85	118.60
84	Aa	553	C	C4'-C3'-O3'	10.40	133.81	113.00
84	Aa	1523	G	N1-C6-O6	10.40	126.14	119.90
84	Aa	2150	C	O4'-C1'-N1	10.40	116.52	108.20
84	Aa	641	C	O4'-C1'-N1	10.40	116.52	108.20
84	Aa	436	G	N1-C6-O6	10.40	126.14	119.90
1	Ad	330	G	O4'-C1'-N9	10.40	116.52	108.20
84	Aa	1493	A	N1-C6-N6	10.39	124.84	118.60
84	Aa	26	A	N1-C6-N6	10.39	124.83	118.60
84	Aa	808	G	N1-C6-O6	10.39	126.13	119.90
1	Ad	1806	C	C3'-C2'-C1'	10.38	109.81	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2006	A	P-O3'-C3'	10.38	132.16	119.70
84	Aa	2132	A	O4'-C1'-N9	10.38	116.50	108.20
86	Ab	107	C	O4'-C1'-N1	10.38	116.50	108.20
1	Ad	707	C	N1-C1'-C2'	-10.38	100.51	114.00
1	Ad	1120	U	O4'-C1'-N1	10.37	116.50	108.20
84	Aa	1617	A	N1-C6-N6	10.37	124.82	118.60
84	Aa	2166	U	P-O3'-C3'	10.37	132.14	119.70
84	Aa	3334	A	N1-C6-N6	10.37	124.82	118.60
1	Ad	144	U	O4'-C1'-N1	10.37	116.49	108.20
1	Ad	712	U	O4'-C1'-N1	10.37	116.49	108.20
84	Aa	2306	G	N1-C6-O6	10.37	126.12	119.90
84	Aa	3307	A	N1-C6-N6	10.37	124.82	118.60
86	Ab	13	A	O4'-C1'-N9	10.36	116.49	108.20
1	Ad	1303	G	C1'-O4'-C4'	-10.36	101.61	109.90
84	Aa	843	C	O4'-C1'-N1	10.36	116.49	108.20
1	Ad	1297	U	O4'-C1'-N1	10.35	116.48	108.20
1	Ad	1358	G	N9-C1'-C2'	10.34	127.44	114.00
84	Aa	1879	A	N1-C6-N6	10.34	124.80	118.60
84	Aa	721	A	O4'-C1'-N9	10.34	116.47	108.20
84	Aa	1902	G	N1-C6-O6	10.34	126.10	119.90
1	Ad	1640	C	N1-C1'-C2'	10.33	127.43	114.00
85	Ac	59	A	N1-C6-N6	10.33	124.80	118.60
85	Ac	43	A	N1-C6-N6	10.33	124.80	118.60
84	Aa	316	A	N1-C6-N6	10.33	124.80	118.60
84	Aa	2850	G	N1-C6-O6	10.33	126.10	119.90
84	Aa	3013	A	N1-C6-N6	10.33	124.80	118.60
84	Aa	3333	C	O4'-C1'-N1	10.31	116.45	108.20
84	Aa	1349	G	C5-C6-O6	-10.30	122.42	128.60
1	Ad	172	U	O4'-C1'-N1	10.30	116.44	108.20
84	Aa	3248	G	N1-C6-O6	10.30	126.08	119.90
1	Ad	1520	G	O4'-C1'-N9	10.29	116.44	108.20
1	Ad	1757	G	O4'-C1'-N9	10.29	116.44	108.20
84	Aa	1316	C	O4'-C1'-N1	10.29	116.44	108.20
84	Aa	1870	G	N1-C6-O6	10.29	126.08	119.90
84	Aa	389	A	N1-C6-N6	10.29	124.77	118.60
84	Aa	1542	A	N1-C6-N6	10.29	124.77	118.60
84	Aa	1813	C	O4'-C1'-N1	10.29	116.43	108.20
84	Aa	2238	A	N1-C6-N6	10.29	124.77	118.60
84	Aa	3244	G	N1-C6-O6	10.29	126.07	119.90
84	Aa	2058	C	O4'-C1'-N1	10.29	116.43	108.20
84	Aa	2558	U	O4'-C1'-N1	10.28	116.43	108.20
86	Ab	37	G	C5-C6-O6	-10.28	122.43	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	754	U	O4'-C1'-N1	10.28	116.42	108.20
84	Aa	2583	A	N1-C6-N6	10.28	124.77	118.60
84	Aa	1031	A	N1-C6-N6	10.27	124.76	118.60
1	Ad	496	A	O4'-C1'-N9	10.27	116.41	108.20
84	Aa	1105	G	N1-C6-O6	10.27	126.06	119.90
1	Ad	870	A	N9-C1'-C2'	-10.26	100.66	114.00
1	Ad	1261	U	O4'-C1'-N1	10.26	116.41	108.20
84	Aa	1744	C	O4'-C1'-N1	10.26	116.41	108.20
84	Aa	339	G	N1-C6-O6	10.26	126.06	119.90
84	Aa	3030	A	N1-C6-N6	10.26	124.76	118.60
84	Aa	3175	C	O4'-C1'-N1	10.25	116.40	108.20
84	Aa	965	A	N1-C6-N6	10.25	124.75	118.60
86	Ab	19	A	C4-C5-C6	10.25	122.12	117.00
84	Aa	473	G	C4'-C3'-O3'	10.25	133.49	113.00
1	Ad	1004	U	O4'-C1'-N1	10.24	116.39	108.20
84	Aa	3288	A	N1-C6-N6	10.24	124.74	118.60
84	Aa	2909	A	N1-C6-N6	10.24	124.74	118.60
84	Aa	3253	C	O4'-C1'-N1	10.24	116.39	108.20
85	Ac	65	G	N1-C6-O6	10.24	126.04	119.90
1	Ad	281	U	P-O3'-C3'	10.23	131.98	119.70
1	Ad	73	A	N9-C1'-C2'	-10.23	100.70	114.00
84	Aa	1006	A	N1-C6-N6	10.23	124.74	118.60
84	Aa	2682	A	N1-C6-N6	10.22	124.73	118.60
86	Ab	8	A	C5-C6-N1	-10.22	112.59	117.70
84	Aa	1486	G	O4'-C1'-N9	10.22	116.37	108.20
84	Aa	2376	G	N1-C6-O6	10.22	126.03	119.90
1	Ad	7	G	O4'-C1'-N9	10.21	116.37	108.20
2	Ae	69	G	C1'-O4'-C4'	-10.21	101.73	109.90
84	Aa	2698	A	N1-C6-N6	10.21	124.73	118.60
1	Ad	1683	G	O4'-C1'-N9	10.20	116.36	108.20
84	Aa	634	A	N1-C6-N6	10.20	124.72	118.60
84	Aa	2442	A	N1-C6-N6	10.19	124.72	118.60
84	Aa	804	A	N1-C6-N6	10.19	124.72	118.60
84	Aa	1677	G	N1-C6-O6	10.19	126.01	119.90
1	Ad	457	C	C1'-O4'-C4'	-10.18	101.75	109.90
84	Aa	2515	C	P-O3'-C3'	10.18	131.92	119.70
84	Aa	1663	G	N1-C6-O6	10.18	126.01	119.90
84	Aa	275	G	N1-C6-O6	10.18	126.01	119.90
84	Aa	1017	G	N1-C6-O6	10.17	126.00	119.90
84	Aa	1832	C	O4'-C1'-N1	10.17	116.33	108.20
84	Aa	1389	C	O4'-C1'-N1	10.16	116.33	108.20
84	Aa	810	A	N1-C6-N6	10.16	124.70	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	925	U	O4'-C1'-N1	10.16	116.33	108.20
1	Ad	1582	G	O4'-C1'-N9	-10.15	100.08	108.20
1	Ad	707	C	O4'-C1'-C2'	-10.14	95.66	105.80
1	Ad	235	C	P-O3'-C3'	10.14	131.87	119.70
85	Ac	37	A	N1-C6-N6	10.14	124.68	118.60
84	Aa	2501	U	P-O3'-C3'	10.14	131.86	119.70
84	Aa	157	G	N1-C6-O6	10.13	125.98	119.90
84	Aa	297	G	N1-C6-O6	10.13	125.98	119.90
86	Ab	44	C	N3-C4-C5	-10.13	117.85	121.90
1	Ad	1414	G	C1'-O4'-C4'	-10.13	101.80	109.90
1	Ad	430	G	O4'-C1'-N9	10.12	116.30	108.20
84	Aa	2447	A	N1-C6-N6	10.12	124.67	118.60
84	Aa	3086	G	N1-C6-O6	10.12	125.97	119.90
84	Aa	3185	G	N1-C6-O6	10.12	125.97	119.90
84	Aa	315	A	N1-C6-N6	10.12	124.67	118.60
84	Aa	2998	A	N1-C6-N6	10.12	124.67	118.60
84	Aa	2133	A	N1-C6-N6	10.12	124.67	118.60
84	Aa	1897	A	N1-C6-N6	10.10	124.66	118.60
84	Aa	919	G	P-O3'-C3'	10.10	131.82	119.70
84	Aa	2516	U	P-O3'-C3'	10.10	131.82	119.70
84	Aa	59	A	N1-C6-N6	10.10	124.66	118.60
1	Ad	247	A	O4'-C1'-N9	10.10	116.28	108.20
1	Ad	1210	U	O4'-C1'-N1	10.10	116.28	108.20
1	Ad	1801	A	N9-C1'-C2'	-10.10	100.88	114.00
84	Aa	637	C	O4'-C1'-N1	10.10	116.28	108.20
1	Ad	616	U	O4'-C1'-N1	10.09	116.27	108.20
84	Aa	909	A	N1-C6-N6	10.09	124.65	118.60
84	Aa	2693	G	N1-C6-O6	10.09	125.95	119.90
84	Aa	494	C	C5'-C4'-C3'	10.08	132.13	116.00
1	Ad	1759	A	O4'-C1'-N9	10.08	116.27	108.20
84	Aa	2231	G	N1-C6-O6	10.08	125.95	119.90
1	Ad	1505	U	O4'-C1'-N1	10.07	116.26	108.20
3	Af	16	G	C3'-C2'-C1'	10.07	109.56	101.50
84	Aa	3130	A	N1-C6-N6	10.07	124.64	118.60
86	Ab	67	C	N3-C4-N4	10.07	125.05	118.00
1	Ad	1203	G	O4'-C1'-C2'	-10.07	95.73	105.80
84	Aa	1559	G	O3'-P-O5'	10.07	123.13	104.00
84	Aa	1717	G	N1-C6-O6	10.07	125.94	119.90
84	Aa	1721	A	O4'-C1'-N9	10.06	116.25	108.20
1	Ad	1315	U	O4'-C1'-N1	10.05	116.24	108.20
1	Ad	1565	U	O4'-C1'-N1	10.05	116.24	108.20
84	Aa	2659	A	N1-C6-N6	10.05	124.63	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3194	G	N1-C6-O6	10.05	125.93	119.90
1	Ad	504	C	C3'-C2'-C1'	10.04	109.53	101.50
84	Aa	108	A	N1-C6-N6	10.04	124.63	118.60
84	Aa	2177	U	P-O3'-C3'	10.04	131.75	119.70
84	Aa	3190	U	O4'-C1'-N1	10.04	116.23	108.20
1	Ad	1254	U	O4'-C1'-N1	10.03	116.23	108.20
84	Aa	1532	A	N1-C6-N6	10.03	124.62	118.60
1	Ad	235	C	C1'-O4'-C4'	10.03	117.92	109.90
84	Aa	1333	C	P-O3'-C3'	10.03	131.74	119.70
84	Aa	3318	G	N1-C6-O6	10.03	125.92	119.90
84	Aa	30	C	O4'-C1'-N1	10.03	116.22	108.20
84	Aa	2349	C	O4'-C1'-N1	10.03	116.22	108.20
84	Aa	2347	A	N1-C6-N6	10.02	124.61	118.60
84	Aa	3380	G	P-O3'-C3'	10.02	131.73	119.70
84	Aa	2242	G	N1-C6-O6	10.02	125.91	119.90
84	Aa	2504	A	N1-C6-N6	10.02	124.61	118.60
84	Aa	3385	G	O4'-C1'-N9	10.02	116.22	108.20
85	Ac	72	A	N1-C6-N6	10.02	124.61	118.60
84	Aa	2587	G	N1-C6-O6	10.02	125.91	119.90
84	Aa	1892	A	N1-C6-N6	10.01	124.61	118.60
1	Ad	66	U	O4'-C1'-N1	10.01	116.21	108.20
1	Ad	843	G	O4'-C1'-C2'	-10.01	95.79	105.80
84	Aa	840	A	N1-C6-N6	10.01	124.61	118.60
84	Aa	2970	G	N1-C6-O6	10.01	125.90	119.90
1	Ad	944	A	C3'-C2'-C1'	10.00	109.50	101.50
84	Aa	1412	C	O4'-C1'-N1	10.00	116.20	108.20
85	Ac	134	G	N1-C6-O6	10.00	125.90	119.90
84	Aa	878	G	N1-C6-O6	10.00	125.90	119.90
1	Ad	220	C	C3'-C2'-C1'	10.00	109.50	101.50
1	Ad	224	C	O4'-C1'-C2'	-9.99	95.81	105.80
84	Aa	1449	A	N1-C6-N6	9.99	124.59	118.60
84	Aa	2194	G	N1-C6-O6	9.99	125.89	119.90
1	Ad	1005	C	C3'-C2'-C1'	-9.98	93.51	101.50
84	Aa	2873	G	N1-C6-O6	9.98	125.89	119.90
84	Aa	2992	G	N1-C6-O6	9.98	125.89	119.90
86	Ab	41	G	C5-C6-O6	-9.98	122.61	128.60
1	Ad	1216	G	O4'-C1'-N9	9.97	116.18	108.20
84	Aa	3317	G	N1-C6-O6	9.97	125.89	119.90
1	Ad	1119	G	O4'-C1'-N9	9.97	116.18	108.20
84	Aa	431	G	N1-C6-O6	9.96	125.88	119.90
1	Ad	717	G	O4'-C1'-N9	-9.96	100.23	108.20
1	Ad	91	C	O4'-C1'-N1	9.95	116.16	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2639	A	N1-C6-N6	9.95	124.57	118.60
85	Ac	143	C	O4'-C1'-N1	9.95	116.16	108.20
1	Ad	1495	U	O4'-C1'-N1	9.94	116.15	108.20
84	Aa	11	A	N1-C6-N6	9.94	124.56	118.60
84	Aa	1421	A	N1-C6-N6	9.94	124.56	118.60
84	Aa	2084	G	C2'-C3'-O3'	-9.93	87.65	109.50
86	Ab	24	G	N1-C6-O6	9.93	125.86	119.90
84	Aa	1436	A	N1-C6-N6	9.92	124.56	118.60
84	Aa	2538	G	N1-C6-O6	9.92	125.85	119.90
85	Ac	131	G	N1-C6-O6	9.92	125.85	119.90
84	Aa	1383	G	N1-C6-O6	9.92	125.85	119.90
84	Aa	2523	G	N1-C6-O6	9.92	125.85	119.90
1	Ad	1575	U	O4'-C1'-N1	9.91	116.13	108.20
84	Aa	1396	A	N1-C6-N6	9.91	124.55	118.60
1	Ad	1797	C	O4'-C1'-N1	9.91	116.13	108.20
84	Aa	263	A	N1-C6-N6	9.91	124.55	118.60
84	Aa	1889	G	N1-C6-O6	9.91	125.85	119.90
84	Aa	3032	G	N1-C6-O6	9.91	125.85	119.90
84	Aa	993	A	N1-C6-N6	9.91	124.54	118.60
84	Aa	2650	A	N1-C6-N6	9.91	124.54	118.60
84	Aa	1211	G	N1-C6-O6	9.90	125.84	119.90
84	Aa	1298	A	N1-C6-N6	9.90	124.54	118.60
84	Aa	2619	C	O4'-C1'-N1	9.90	116.12	108.20
1	Ad	209	U	O4'-C1'-N1	9.90	116.12	108.20
1	Ad	468	A	O4'-C1'-N9	9.90	116.12	108.20
84	Aa	649	A	N1-C6-N6	9.89	124.54	118.60
1	Ad	243	U	O4'-C1'-N1	9.89	116.11	108.20
1	Ad	310	U	O4'-C1'-N1	9.89	116.11	108.20
84	Aa	425	G	N1-C6-O6	9.89	125.83	119.90
1	Ad	587	C	N1-C1'-C2'	9.89	126.85	114.00
84	Aa	974	G	N1-C6-O6	9.89	125.83	119.90
84	Aa	3375	G	N1-C6-O6	9.89	125.83	119.90
84	Aa	1723	C	P-O3'-C3'	9.88	131.55	119.70
84	Aa	398	G	N1-C6-O6	9.87	125.83	119.90
84	Aa	2443	C	O4'-C1'-N1	9.87	116.10	108.20
84	Aa	1487	A	N1-C6-N6	9.87	124.52	118.60
84	Aa	1901	G	N1-C6-O6	9.87	125.82	119.90
1	Ad	1498	A	P-O3'-C3'	9.87	131.54	119.70
84	Aa	253	G	O4'-C1'-N9	9.87	116.09	108.20
84	Aa	521	G	N1-C6-O6	9.87	125.82	119.90
84	Aa	1486	G	N1-C6-O6	9.87	125.82	119.90
1	Ad	1060	U	O4'-C1'-N1	9.86	116.09	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1455	U	O4'-C1'-N1	9.86	116.09	108.20
84	Aa	3067	G	N1-C6-O6	9.86	125.82	119.90
84	Aa	542	G	N1-C6-O6	9.86	125.82	119.90
1	Ad	536	U	N1-C1'-C2'	9.86	126.81	114.00
84	Aa	1452	A	N1-C6-N6	9.85	124.51	118.60
84	Aa	845	G	N1-C6-O6	9.85	125.81	119.90
84	Aa	911	G	N1-C6-O6	9.85	125.81	119.90
84	Aa	1958	G	O4'-C4'-C3'	-9.85	94.15	104.00
84	Aa	549	G	N1-C6-O6	9.85	125.81	119.90
84	Aa	579	G	N1-C6-O6	9.85	125.81	119.90
1	Ad	208	U	O4'-C1'-N1	9.84	116.07	108.20
84	Aa	283	A	N1-C6-N6	9.84	124.50	118.60
1	Ad	516	A	O4'-C1'-N9	9.84	116.07	108.20
84	Aa	1649	G	N1-C6-O6	9.84	125.80	119.90
84	Aa	1094	G	N1-C6-O6	9.83	125.80	119.90
84	Aa	1673	A	N1-C6-N6	9.83	124.50	118.60
1	Ad	219	G	N9-C1'-C2'	-9.82	101.19	112.00
84	Aa	2610	G	N1-C6-O6	9.81	125.78	119.90
1	Ad	820	A	O4'-C1'-N9	9.81	116.05	108.20
1	Ad	851	G	C3'-C2'-C1'	-9.80	93.66	101.50
1	Ad	924	A	O4'-C1'-N9	9.80	116.04	108.20
84	Aa	936	A	N1-C6-N6	9.80	124.48	118.60
84	Aa	2384	G	N1-C6-O6	9.81	125.78	119.90
84	Aa	2073	U	O4'-C1'-N1	9.80	116.04	108.20
84	Aa	2363	G	N1-C6-O6	9.80	125.78	119.90
84	Aa	1786	G	N1-C6-O6	9.79	125.78	119.90
1	Ad	916	U	O4'-C1'-N1	9.79	116.03	108.20
84	Aa	1746	G	N1-C6-O6	9.79	125.78	119.90
86	Ab	26	C	O4'-C1'-N1	9.79	116.03	108.20
84	Aa	1317	G	N1-C6-O6	9.79	125.78	119.90
86	Ab	60	G	O4'-C1'-N9	9.79	116.03	108.20
1	Ad	579	C	O4'-C1'-N1	9.79	116.03	108.20
1	Ad	1375	C	C3'-C2'-C1'	9.79	109.33	101.50
84	Aa	3099	G	N1-C6-O6	9.79	125.77	119.90
1	Ad	1462	C	C3'-C2'-C1'	-9.79	93.67	101.50
84	Aa	1359	A	P-O3'-C3'	9.79	131.44	119.70
84	Aa	2163	G	N1-C6-O6	9.78	125.77	119.90
84	Aa	1721	A	N1-C6-N6	9.78	124.47	118.60
1	Ad	1671	G	O4'-C1'-N9	9.78	116.02	108.20
84	Aa	388	G	N1-C6-O6	9.78	125.77	119.90
84	Aa	3345	G	N1-C6-O6	9.78	125.77	119.90
1	Ad	1771	U	P-O3'-C3'	9.77	131.43	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	526	A	N1-C6-N6	9.77	124.46	118.60
2	Ae	73	C	C1'-O4'-C4'	-9.77	102.08	109.90
84	Aa	3377	G	N1-C6-O6	9.77	125.76	119.90
84	Aa	3181	U	P-O3'-C3'	9.77	131.42	119.70
1	Ad	512	U	O4'-C1'-N1	9.77	116.01	108.20
1	Ad	1022	U	O4'-C1'-N1	9.77	116.01	108.20
84	Aa	2181	U	O4'-C1'-N1	9.77	116.01	108.20
84	Aa	2830	G	N1-C6-O6	9.77	125.76	119.90
86	Ab	24	G	C5-C6-O6	-9.76	122.74	128.60
84	Aa	618	G	N1-C6-O6	9.76	125.76	119.90
84	Aa	2172	C	O4'-C1'-N1	9.76	116.01	108.20
84	Aa	1352	G	N1-C6-O6	9.76	125.75	119.90
84	Aa	1446	G	N1-C6-O6	9.76	125.75	119.90
84	Aa	2234	G	N1-C6-O6	9.76	125.75	119.90
84	Aa	3278	G	N1-C6-O6	9.76	125.75	119.90
84	Aa	3354	A	N1-C6-N6	9.75	124.45	118.60
1	Ad	780	A	C3'-C2'-C1'	-9.75	93.70	101.50
1	Ad	1008	A	O4'-C1'-N9	9.75	116.00	108.20
84	Aa	2604	A	N1-C6-N6	9.75	124.45	118.60
86	Ab	108	G	N1-C6-O6	9.75	125.75	119.90
1	Ad	87	A	N9-C1'-C2'	-9.74	101.28	112.00
1	Ad	1322	G	O4'-C1'-N9	9.74	115.99	108.20
1	Ad	1200	A	O4'-C1'-N9	9.74	115.99	108.20
1	Ad	158	C	N1-C1'-C2'	9.74	126.66	114.00
84	Aa	2090	G	N1-C6-O6	9.73	125.74	119.90
84	Aa	334	A	N1-C6-N6	9.73	124.44	118.60
84	Aa	2302	G	N1-C6-O6	9.73	125.74	119.90
1	Ad	1657	C	O4'-C1'-N1	9.73	115.98	108.20
84	Aa	2692	G	N1-C6-O6	9.73	125.74	119.90
1	Ad	1382	C	O4'-C1'-N1	9.73	115.98	108.20
84	Aa	1923	G	N1-C6-O6	9.73	125.74	119.90
84	Aa	2613	G	N1-C6-O6	9.73	125.74	119.90
84	Aa	2069	G	N1-C6-O6	9.72	125.73	119.90
1	Ad	437	C	N1-C1'-C2'	9.72	126.64	114.00
84	Aa	3153	U	O4'-C1'-N1	9.72	115.98	108.20
85	Ac	130	G	N1-C6-O6	9.72	125.73	119.90
1	Ad	351	G	C1'-O4'-C4'	-9.72	102.13	109.90
84	Aa	2902	A	N1-C6-N6	9.72	124.43	118.60
84	Aa	1428	G	N1-C6-O6	9.71	125.73	119.90
1	Ad	503	U	O4'-C1'-N1	9.71	115.96	108.20
84	Aa	1756	C	O4'-C1'-N1	9.71	115.96	108.20
85	Ac	11	C	O4'-C1'-N1	9.71	115.96	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	251	U	O4'-C1'-N1	9.70	115.96	108.20
84	Aa	1801	G	N1-C6-O6	9.70	125.72	119.90
84	Aa	648	G	N1-C6-O6	9.69	125.72	119.90
84	Aa	907	A	N1-C6-N6	9.69	124.42	118.60
84	Aa	125	G	N1-C6-O6	9.69	125.72	119.90
84	Aa	423	C	N3-C4-N4	9.69	124.78	118.00
1	Ad	1684	U	O4'-C1'-N1	9.69	115.95	108.20
84	Aa	502	G	N1-C6-O6	9.69	125.71	119.90
84	Aa	2803	A	N1-C6-N6	9.69	124.41	118.60
84	Aa	133	G	N1-C6-O6	9.69	125.71	119.90
84	Aa	560	C	O4'-C1'-N1	9.68	115.95	108.20
84	Aa	2741	G	N1-C6-O6	9.68	125.71	119.90
1	Ad	841	U	O4'-C1'-N1	9.68	115.94	108.20
84	Aa	2945	G	N1-C6-O6	9.68	125.71	119.90
1	Ad	342	C	N1-C1'-C2'	9.67	126.58	114.00
1	Ad	505	U	P-O3'-C3'	9.67	131.30	119.70
84	Aa	1594	G	N1-C6-O6	9.67	125.70	119.90
84	Aa	474	G	O4'-C1'-N9	9.67	115.93	108.20
1	Ad	1806	C	O4'-C1'-C2'	-9.67	96.13	105.80
1	Ad	362	U	O4'-C1'-N1	9.66	115.93	108.20
86	Ab	70	G	N1-C6-O6	9.66	125.70	119.90
1	Ad	1257	U	O4'-C1'-N1	9.66	115.93	108.20
2	Ae	39	G	O4'-C1'-N9	9.66	115.93	108.20
84	Aa	753	G	N1-C6-O6	9.66	125.70	119.90
1	Ad	1500	A	O4'-C1'-N9	9.66	115.93	108.20
1	Ad	1084	U	O4'-C1'-N1	9.65	115.92	108.20
84	Aa	1650	G	N1-C6-O6	9.65	125.69	119.90
1	Ad	1176	A	O4'-C1'-N9	9.65	115.92	108.20
85	Ac	95	G	N1-C6-O6	9.64	125.69	119.90
1	Ad	854	C	O4'-C1'-C2'	-9.64	96.16	105.80
84	Aa	2146	A	N1-C6-N6	9.64	124.38	118.60
84	Aa	2174	C	P-O5'-C5'	9.64	136.32	120.90
84	Aa	764	A	N1-C6-N6	9.63	124.38	118.60
84	Aa	1772	G	N1-C6-O6	9.63	125.68	119.90
85	Ac	97	G	N1-C6-O6	9.63	125.68	119.90
1	Ad	887	U	O4'-C1'-N1	9.62	115.90	108.20
1	Ad	1560	U	O4'-C1'-N1	9.62	115.90	108.20
1	Ad	233	U	O4'-C1'-N1	9.62	115.90	108.20
84	Aa	1450	G	N1-C6-O6	9.62	125.67	119.90
84	Aa	1290	A	N1-C6-N6	9.62	124.37	118.60
1	Ad	1311	U	C1'-O4'-C4'	9.62	117.59	109.90
84	Aa	841	G	N1-C6-O6	9.61	125.67	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	Ae	74	C	O4'-C1'-N1	9.61	115.88	108.20
84	Aa	1309	U	P-O3'-C3'	9.61	131.23	119.70
84	Aa	3364	A	N1-C6-N6	9.61	124.36	118.60
84	Aa	3206	C	O4'-C1'-N1	9.61	115.88	108.20
84	Aa	2244	G	N1-C6-O6	9.60	125.66	119.90
84	Aa	3243	C	O4'-C1'-N1	9.60	115.88	108.20
84	Aa	1250	G	O4'-C1'-N9	9.60	115.88	108.20
1	Ad	1096	A	O4'-C1'-C2'	9.60	116.24	107.60
84	Aa	2361	C	O4'-C1'-N1	9.60	115.88	108.20
1	Ad	1031	A	O4'-C1'-N9	9.60	115.88	108.20
1	Ad	1448	U	O4'-C1'-N1	9.60	115.88	108.20
84	Aa	1092	G	C5-C6-O6	-9.60	122.84	128.60
84	Aa	851	A	N1-C6-N6	9.60	124.36	118.60
84	Aa	202	G	N1-C6-O6	9.59	125.65	119.90
84	Aa	1132	A	N1-C6-N6	9.59	124.35	118.60
2	Ae	49	G	O4'-C1'-N9	9.59	115.87	108.20
84	Aa	949	C	O4'-C1'-N1	9.59	115.87	108.20
84	Aa	1774	G	N1-C6-O6	9.59	125.65	119.90
86	Ab	42	A	C5-C6-N6	-9.59	116.03	123.70
86	Ab	116	U	O4'-C1'-N1	9.59	115.87	108.20
84	Aa	2087	A	N9-C1'-C2'	9.58	126.46	114.00
84	Aa	2525	G	P-O3'-C3'	9.58	131.20	119.70
1	Ad	1407	A	O4'-C1'-N9	9.58	115.86	108.20
84	Aa	1025	G	N1-C6-O6	9.58	125.65	119.90
84	Aa	1479	G	N1-C6-O6	9.58	125.65	119.90
84	Aa	2323	A	N1-C6-N6	9.58	124.35	118.60
1	Ad	35	U	O4'-C1'-N1	9.57	115.86	108.20
84	Aa	2473	C	O4'-C1'-N1	9.57	115.86	108.20
1	Ad	238	G	P-O3'-C3'	9.57	131.18	119.70
84	Aa	2093	G	N1-C6-O6	9.57	125.64	119.90
1	Ad	745	C	C1'-O4'-C4'	9.57	117.55	109.90
84	Aa	816	G	N1-C6-O6	9.57	125.64	119.90
84	Aa	953	G	N1-C6-O6	9.57	125.64	119.90
84	Aa	664	A	N1-C6-N6	9.56	124.34	118.60
84	Aa	1272	G	N1-C6-O6	9.56	125.64	119.90
84	Aa	2403	A	N1-C6-N6	9.56	124.34	118.60
84	Aa	3328	A	N1-C6-N6	9.56	124.34	118.60
86	Ab	76	U	O4'-C1'-N1	9.56	115.85	108.20
86	Ab	101	A	N1-C6-N6	9.56	124.34	118.60
84	Aa	662	G	N1-C6-O6	9.56	125.64	119.90
84	Aa	857	G	N1-C6-O6	9.56	125.64	119.90
84	Aa	423	C	C5-C4-N4	-9.55	113.51	120.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2011	G	P-O3'-C3'	9.55	131.16	119.70
84	Aa	740	G	N1-C6-O6	9.55	125.63	119.90
84	Aa	1384	G	N1-C6-O6	9.55	125.63	119.90
84	Aa	1480	G	N1-C6-O6	9.55	125.63	119.90
84	Aa	1562	A	P-O3'-C3'	9.55	131.16	119.70
1	Ad	582	U	O4'-C1'-N1	9.54	115.84	108.20
1	Ad	1067	A	C3'-C2'-C1'	-9.54	93.87	101.50
1	Ad	1765	A	N9-C1'-C2'	-9.54	101.51	112.00
85	Ac	107	G	N1-C6-O6	9.54	125.62	119.90
2	Ae	11	U	O4'-C1'-N1	9.53	115.83	108.20
84	Aa	2078	G	N1-C6-O6	9.53	125.62	119.90
1	Ad	238	G	N9-C1'-C2'	9.53	126.39	114.00
84	Aa	1244	A	N1-C6-N6	9.53	124.32	118.60
84	Aa	1946	C	O4'-C1'-N1	9.53	115.82	108.20
84	Aa	1244	A	P-O3'-C3'	9.52	131.13	119.70
1	Ad	87	A	O4'-C1'-C2'	-9.52	96.28	105.80
1	Ad	952	U	O4'-C1'-N1	9.52	115.81	108.20
84	Aa	3302	A	N1-C6-N6	9.52	124.31	118.60
86	Ab	54	A	N1-C6-N6	9.52	124.31	118.60
84	Aa	328	G	N1-C6-O6	9.52	125.61	119.90
84	Aa	1423	C	O4'-C1'-N1	9.52	115.81	108.20
1	Ad	868	A	P-O3'-C3'	9.51	131.12	119.70
84	Aa	1390	G	N1-C6-O6	9.50	125.60	119.90
84	Aa	2308	A	N1-C6-N6	9.50	124.30	118.60
84	Aa	430	G	N1-C6-O6	9.50	125.60	119.90
84	Aa	1431	G	N1-C6-O6	9.50	125.60	119.90
1	Ad	1637	G	C1'-O4'-C4'	-9.49	102.30	109.90
84	Aa	1750	A	N1-C6-N6	9.49	124.30	118.60
1	Ad	1691	C	O4'-C1'-N1	9.49	115.79	108.20
1	Ad	1765	A	C3'-C2'-C1'	-9.49	93.91	101.50
1	Ad	1475	A	C3'-C2'-C1'	9.49	109.09	101.50
84	Aa	235	G	N1-C6-O6	9.49	125.59	119.90
84	Aa	1955	G	N1-C6-O6	9.49	125.59	119.90
84	Aa	2462	G	O4'-C1'-N9	9.49	115.79	108.20
84	Aa	3254	C	O4'-C1'-N1	9.49	115.79	108.20
1	Ad	1730	G	O4'-C1'-C2'	-9.49	96.31	105.80
84	Aa	695	G	N1-C6-O6	9.49	125.59	119.90
84	Aa	456	G	N1-C6-O6	9.48	125.59	119.90
84	Aa	2690	G	N1-C6-O6	9.48	125.59	119.90
1	Ad	205	U	C1'-O4'-C4'	9.48	117.48	109.90
1	Ad	1656	C	C3'-C2'-C1'	9.48	109.08	101.50
84	Aa	590	C	C4'-C3'-O3'	9.48	131.96	113.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	862	G	N1-C6-O6	9.48	125.59	119.90
84	Aa	367	A	N1-C6-N6	9.47	124.28	118.60
84	Aa	627	G	N1-C6-O6	9.47	125.58	119.90
1	Ad	1310	C	C3'-C2'-C1'	9.47	109.08	101.50
84	Aa	514	G	N1-C6-O6	9.47	125.58	119.90
84	Aa	2764	G	C5-C6-O6	-9.47	122.92	128.60
1	Ad	1739	U	O4'-C1'-N1	9.47	115.78	108.20
2	Ae	67	G	O4'-C1'-N9	9.47	115.77	108.20
84	Aa	1641	G	N1-C6-O6	9.47	125.58	119.90
1	Ad	1608	A	O4'-C1'-N9	9.46	115.77	108.20
84	Aa	171	G	C5-C6-O6	-9.46	122.92	128.60
84	Aa	1247	G	N1-C6-O6	9.46	125.58	119.90
84	Aa	1652	G	N1-C6-O6	9.46	125.58	119.90
84	Aa	678	G	N1-C6-O6	9.46	125.58	119.90
1	Ad	1096	A	O4'-C1'-N9	9.46	115.77	108.20
84	Aa	449	G	N1-C6-O6	9.46	125.57	119.90
1	Ad	1200	A	P-O3'-C3'	9.45	131.04	119.70
84	Aa	1539	G	N1-C6-O6	9.46	125.57	119.90
84	Aa	1677	G	C5-C6-O6	-9.46	122.93	128.60
84	Aa	517	G	N1-C6-O6	9.45	125.57	119.90
84	Aa	1907	A	N1-C6-N6	9.45	124.27	118.60
84	Aa	1847	G	N1-C6-O6	9.45	125.57	119.90
1	Ad	1304	A	O4'-C1'-N9	9.45	115.76	108.20
84	Aa	1664	G	N1-C6-O6	9.45	125.57	119.90
1	Ad	921	U	O4'-C1'-N1	9.44	115.75	108.20
84	Aa	731	G	C5-C6-O6	-9.45	122.93	128.60
84	Aa	1130	G	N1-C6-O6	9.44	125.57	119.90
84	Aa	178	C	O4'-C1'-N1	9.44	115.75	108.20
84	Aa	3149	C	O4'-C1'-N1	9.44	115.75	108.20
86	Ab	51	G	N1-C6-O6	9.44	125.56	119.90
84	Aa	42	A	N1-C6-N6	9.44	124.26	118.60
84	Aa	1140	C	O4'-C1'-N1	9.44	115.75	108.20
84	Aa	1958	G	O4'-C1'-N9	9.44	115.75	108.20
84	Aa	318	G	N1-C6-O6	9.43	125.56	119.90
84	Aa	2295	G	N1-C6-O6	9.43	125.56	119.90
84	Aa	3371	C	O4'-C1'-N1	9.43	115.75	108.20
84	Aa	2661	G	C5-C6-O6	-9.43	122.94	128.60
84	Aa	3102	G	N1-C6-O6	9.43	125.56	119.90
86	Ab	105	C	O4'-C1'-N1	9.43	115.74	108.20
84	Aa	1089	G	N1-C6-O6	9.43	125.56	119.90
84	Aa	3157	C	O4'-C1'-N1	9.43	115.74	108.20
85	Ac	136	G	N1-C6-O6	9.43	125.56	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3390	G	N1-C6-O6	9.42	125.55	119.90
84	Aa	1243	C	O4'-C1'-N1	9.42	115.73	108.20
1	Ad	1397	A	N9-C1'-C2'	9.41	126.24	114.00
84	Aa	809	A	N1-C6-N6	9.41	124.25	118.60
1	Ad	588	C	N1-C1'-C2'	9.41	126.23	114.00
84	Aa	2183	A	N1-C6-N6	9.41	124.25	118.60
1	Ad	548	C	N1-C1'-C2'	9.41	126.23	114.00
1	Ad	947	G	O4'-C1'-N9	9.40	115.72	108.20
1	Ad	1039	C	O4'-C1'-N1	9.40	115.72	108.20
1	Ad	1271	G	O4'-C1'-N9	9.40	115.72	108.20
84	Aa	285	G	N1-C6-O6	9.40	125.54	119.90
84	Aa	38	A	N1-C6-N6	9.40	124.24	118.60
1	Ad	994	U	O4'-C1'-N1	9.39	115.72	108.20
84	Aa	21	G	N1-C6-O6	9.39	125.54	119.90
84	Aa	3312	G	C5-C6-O6	-9.39	122.96	128.60
1	Ad	104	A	O4'-C1'-N9	9.39	115.71	108.20
84	Aa	566	G	N1-C6-O6	9.39	125.53	119.90
84	Aa	432	G	N1-C6-O6	9.39	125.53	119.90
84	Aa	1066	G	N1-C6-O6	9.39	125.53	119.90
84	Aa	1145	G	N1-C6-O6	9.39	125.53	119.90
84	Aa	2087	A	O4'-C1'-N9	9.39	115.71	108.20
86	Ab	92	C	N3-C4-N4	9.39	124.57	118.00
1	Ad	785	A	C3'-C2'-C1'	9.39	109.01	101.50
84	Aa	1236	C	O4'-C1'-N1	9.38	115.71	108.20
84	Aa	2879	G	N1-C6-O6	9.38	125.53	119.90
84	Aa	3190	U	P-O3'-C3'	9.38	130.96	119.70
84	Aa	2392	G	N1-C6-O6	9.38	125.53	119.90
84	Aa	581	G	N1-C6-O6	9.38	125.53	119.90
84	Aa	1769	C	O4'-C1'-N1	9.38	115.70	108.20
1	Ad	1692	G	O4'-C1'-N9	9.38	115.70	108.20
1	Ad	1803	G	O4'-C1'-C2'	-9.38	96.42	105.80
84	Aa	836	G	N1-C6-O6	9.38	125.53	119.90
86	Ab	20	C	O4'-C1'-N1	9.38	115.70	108.20
84	Aa	3298	G	N1-C6-O6	9.38	125.53	119.90
84	Aa	213	G	N1-C6-O6	9.38	125.53	119.90
84	Aa	798	G	N1-C6-O6	9.38	125.53	119.90
84	Aa	1263	A	C5-C6-N6	-9.37	116.20	123.70
84	Aa	1675	G	N1-C6-O6	9.37	125.52	119.90
84	Aa	1780	C	O4'-C1'-N1	9.37	115.70	108.20
84	Aa	2525	G	N1-C6-O6	9.37	125.52	119.90
84	Aa	2278	G	N1-C6-O6	9.37	125.52	119.90
1	Ad	119	U	O4'-C1'-N1	9.37	115.69	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	937	A	O4'-C1'-N9	9.37	115.69	108.20
1	Ad	1570	G	O4'-C1'-N9	9.36	115.69	108.20
85	Ac	3	A	O4'-C1'-N9	9.36	115.69	108.20
84	Aa	548	G	N1-C6-O6	9.36	125.51	119.90
84	Aa	794	G	N1-C6-O6	9.35	125.51	119.90
84	Aa	2219	A	N1-C6-N6	9.35	124.21	118.60
84	Aa	1524	G	N1-C6-O6	9.35	125.51	119.90
3	Af	16	G	O4'-C1'-C2'	-9.35	96.45	105.80
1	Ad	459	C	O4'-C1'-C2'	-9.34	96.46	105.80
1	Ad	1334	G	O4'-C1'-C2'	-9.34	96.46	105.80
84	Aa	302	G	N1-C6-O6	9.34	125.50	119.90
86	Ab	87	G	N1-C6-O6	9.34	125.50	119.90
84	Aa	623	G	N1-C6-O6	9.33	125.50	119.90
84	Aa	2550	C	O4'-C1'-N1	9.33	115.67	108.20
84	Aa	1356	G	P-O3'-C3'	9.33	130.90	119.70
1	Ad	1301	G	O4'-C1'-N9	9.33	115.66	108.20
1	Ad	1129	A	O4'-C1'-N9	9.33	115.66	108.20
84	Aa	1575	G	N1-C6-O6	9.33	125.50	119.90
86	Ab	90	A	C8-N9-C4	-9.33	102.07	105.80
84	Aa	1220	G	C5-C6-O6	-9.32	123.01	128.60
84	Aa	2577	G	N1-C6-O6	9.32	125.49	119.90
1	Ad	1384	U	O4'-C1'-N1	9.32	115.66	108.20
84	Aa	1188	C	O4'-C1'-N1	9.32	115.66	108.20
84	Aa	509	G	N1-C6-O6	9.32	125.49	119.90
84	Aa	1106	G	N1-C6-O6	9.32	125.49	119.90
84	Aa	2701	G	N1-C6-O6	9.32	125.49	119.90
85	Ac	111	G	N1-C6-O6	9.31	125.49	119.90
84	Aa	2320	A	N1-C6-N6	9.31	124.19	118.60
1	Ad	50	C	O4'-C1'-N1	9.30	115.64	108.20
1	Ad	111	U	O4'-C1'-N1	9.30	115.64	108.20
84	Aa	2149	G	O4'-C1'-N9	9.30	115.64	108.20
2	Ae	5	U	O4'-C1'-N1	9.30	115.64	108.20
84	Aa	1208	A	N1-C6-N6	9.30	124.18	118.60
84	Aa	866	C	O4'-C1'-N1	9.29	115.63	108.20
84	Aa	2382	C	O4'-C1'-N1	9.29	115.63	108.20
86	Ab	97	G	N1-C6-O6	9.29	125.47	119.90
1	Ad	180	A	O4'-C1'-N9	9.29	115.63	108.20
1	Ad	894	U	O4'-C1'-N1	9.29	115.63	108.20
84	Aa	1107	G	C5-C6-O6	-9.29	123.03	128.60
86	Ab	108	G	C5-C6-O6	-9.29	123.03	128.60
1	Ad	192	G	C1'-O4'-C4'	-9.28	102.47	109.90
84	Aa	241	G	N1-C6-O6	9.28	125.47	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2453	G	N1-C6-O6	9.28	125.47	119.90
84	Aa	1375	G	N1-C6-O6	9.28	125.47	119.90
1	Ad	239	C	C3'-C2'-C1'	9.28	108.92	101.50
1	Ad	1509	C	O4'-C1'-N1	9.28	115.62	108.20
84	Aa	966	G	N1-C6-O6	9.28	125.47	119.90
1	Ad	1758	G	O4'-C1'-C2'	9.28	115.95	107.60
84	Aa	137	C	O4'-C1'-N1	9.28	115.62	108.20
84	Aa	682	G	N1-C6-O6	9.28	125.47	119.90
84	Aa	1277	A	N1-C6-N6	9.27	124.16	118.60
84	Aa	1814	C	O4'-C1'-N1	9.27	115.62	108.20
84	Aa	2151	G	N1-C6-O6	9.27	125.46	119.90
84	Aa	2652	G	N1-C6-O6	9.27	125.46	119.90
1	Ad	740	U	O4'-C1'-N1	9.27	115.62	108.20
84	Aa	2273	C	O4'-C1'-N1	9.27	115.62	108.20
86	Ab	109	U	O4'-C1'-N1	9.27	115.62	108.20
84	Aa	421	A	N1-C6-N6	9.27	124.16	118.60
84	Aa	2418	A	N1-C6-N6	9.27	124.16	118.60
84	Aa	3178	C	O4'-C1'-N1	9.27	115.61	108.20
84	Aa	724	A	N1-C6-N6	9.27	124.16	118.60
84	Aa	1374	G	N1-C6-O6	9.27	125.46	119.90
84	Aa	1595	G	N1-C6-O6	9.27	125.46	119.90
84	Aa	2796	G	C5-C6-O6	-9.26	123.04	128.60
1	Ad	745	C	O4'-C1'-C2'	-9.26	96.54	105.80
84	Aa	32	G	N1-C6-O6	9.26	125.46	119.90
84	Aa	823	A	N1-C6-N6	9.26	124.15	118.60
84	Aa	996	A	N1-C6-N6	9.26	124.15	118.60
85	Ac	70	G	N1-C6-O6	9.26	125.45	119.90
84	Aa	89	C	O4'-C1'-N1	9.25	115.60	108.20
84	Aa	1545	G	N1-C6-O6	9.25	125.45	119.90
84	Aa	2685	C	O4'-C1'-N1	9.25	115.60	108.20
1	Ad	121	U	N1-C1'-C2'	9.25	126.03	114.00
84	Aa	447	C	O4'-C1'-N1	9.25	115.60	108.20
84	Aa	1416	G	N1-C6-O6	9.25	125.45	119.90
1	Ad	1612	C	C3'-C2'-C1'	9.25	108.90	101.50
84	Aa	1545	G	P-O3'-C3'	9.25	130.79	119.70
84	Aa	2461	A	N1-C6-N6	9.25	124.15	118.60
1	Ad	647	G	O4'-C1'-N9	9.24	115.59	108.20
84	Aa	1253	G	N1-C6-O6	9.24	125.45	119.90
84	Aa	1286	G	C5-C6-O6	-9.24	123.05	128.60
1	Ad	1375	C	P-O3'-C3'	9.24	130.79	119.70
84	Aa	1469	G	N1-C6-O6	9.24	125.45	119.90
84	Aa	1993	G	N1-C6-O6	9.24	125.44	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3204	G	N1-C6-O6	9.24	125.44	119.90
1	Ad	1113	G	O4'-C1'-N9	9.23	115.59	108.20
84	Aa	2529	C	O4'-C1'-N1	9.23	115.59	108.20
85	Ac	150	G	N1-C6-O6	9.23	125.44	119.90
84	Aa	370	A	N1-C6-N6	9.23	124.14	118.60
84	Aa	593	G	N1-C6-O6	9.23	125.44	119.90
86	Ab	2	G	C5-C6-O6	-9.23	123.06	128.60
84	Aa	180	G	N1-C6-O6	9.23	125.44	119.90
84	Aa	288	G	C5-C6-O6	-9.23	123.06	128.60
84	Aa	281	G	N1-C6-O6	9.23	125.44	119.90
84	Aa	2345	C	O4'-C1'-N1	9.23	115.58	108.20
1	Ad	642	C	O4'-C1'-N1	9.23	115.58	108.20
84	Aa	3117	G	N1-C6-O6	9.23	125.44	119.90
1	Ad	893	U	O4'-C1'-N1	9.23	115.58	108.20
1	Ad	1029	U	C1'-O4'-C4'	9.23	117.28	109.90
1	Ad	102	U	O4'-C1'-N1	9.22	115.58	108.20
84	Aa	107	C	O4'-C1'-N1	9.22	115.58	108.20
84	Aa	530	C	O4'-C1'-N1	9.22	115.58	108.20
84	Aa	1260	G	N1-C6-O6	9.22	125.43	119.90
84	Aa	2353	C	O4'-C1'-N1	9.22	115.58	108.20
85	Ac	85	G	C5-C6-O6	-9.22	123.06	128.60
84	Aa	1116	G	N1-C6-O6	9.22	125.43	119.90
86	Ab	14	C	O4'-C1'-N1	9.22	115.58	108.20
84	Aa	2104	G	N1-C6-O6	9.22	125.43	119.90
84	Aa	3050	A	N1-C6-N6	9.22	124.13	118.60
86	Ab	28	U	O4'-C1'-N1	9.22	115.57	108.20
84	Aa	104	G	N1-C6-O6	9.22	125.43	119.90
84	Aa	3094	C	O4'-C1'-N1	9.22	115.57	108.20
84	Aa	979	C	O4'-C1'-N1	9.21	115.57	108.20
84	Aa	1169	G	N1-C6-O6	9.21	125.43	119.90
84	Aa	1386	G	N1-C6-O6	9.21	125.43	119.90
84	Aa	2036	C	O4'-C1'-N1	9.21	115.57	108.20
84	Aa	1377	G	N1-C6-O6	9.21	125.43	119.90
84	Aa	1622	G	N1-C6-O6	9.21	125.43	119.90
1	Ad	1056	A	N9-C1'-C2'	-9.21	101.87	112.00
84	Aa	1867	U	O4'-C1'-N1	9.21	115.57	108.20
84	Aa	512	G	N1-C6-O6	9.21	125.42	119.90
84	Aa	1799	C	O4'-C1'-N1	9.20	115.56	108.20
84	Aa	3127	C	O4'-C1'-N1	9.20	115.56	108.20
84	Aa	1662	G	N1-C6-O6	9.20	125.42	119.90
84	Aa	2950	C	O4'-C1'-N1	9.20	115.56	108.20
85	Ac	63	C	O4'-C1'-N1	9.20	115.56	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3010	G	N1-C6-O6	9.20	125.42	119.90
84	Aa	129	G	N1-C6-O6	9.19	125.42	119.90
84	Aa	1829	G	N1-C6-O6	9.19	125.42	119.90
84	Aa	601	G	N1-C6-O6	9.19	125.42	119.90
84	Aa	2334	G	N1-C6-O6	9.19	125.42	119.90
86	Ab	57	C	O4'-C1'-N1	9.19	115.55	108.20
84	Aa	375	G	N1-C6-O6	9.19	125.41	119.90
84	Aa	1658	G	N1-C6-O6	9.19	125.41	119.90
1	Ad	1456	U	O4'-C1'-N1	9.19	115.55	108.20
84	Aa	1770	C	O4'-C1'-N1	9.19	115.55	108.20
86	Ab	115	A	C5-C6-N6	-9.19	116.35	123.70
1	Ad	495	C	O4'-C1'-N1	9.18	115.55	108.20
84	Aa	1958	G	C4'-C3'-O3'	-9.18	90.11	109.40
1	Ad	138	C	O4'-C1'-C2'	-9.18	96.62	105.80
84	Aa	1661	G	N1-C6-O6	9.18	125.41	119.90
2	Ae	16	U	O4'-C1'-N1	9.18	115.54	108.20
1	Ad	871	G	O4'-C1'-N9	9.18	115.54	108.20
1	Ad	1172	G	N9-C1'-C2'	9.18	125.93	114.00
86	Ab	15	C	O4'-C1'-N1	9.18	115.54	108.20
84	Aa	904	G	N1-C6-O6	9.17	125.40	119.90
84	Aa	1774	G	P-O3'-C3'	9.17	130.71	119.70
84	Aa	317	G	N1-C6-O6	9.17	125.40	119.90
84	Aa	1133	A	N1-C6-N6	9.17	124.10	118.60
84	Aa	1820	C	O4'-C1'-N1	9.17	115.54	108.20
84	Aa	1931	G	N1-C6-O6	9.17	125.40	119.90
84	Aa	2182	G	N1-C6-O6	9.17	125.40	119.90
84	Aa	2314	G	O4'-C1'-N9	9.17	115.53	108.20
84	Aa	424	G	N1-C6-O6	9.16	125.40	119.90
86	Ab	49	A	C5-C6-N6	-9.16	116.37	123.70
1	Ad	1045	G	C1'-O4'-C4'	-9.16	102.57	109.90
84	Aa	2538	G	O4'-C1'-N9	9.16	115.53	108.20
84	Aa	1698	C	O4'-C1'-N1	9.16	115.53	108.20
84	Aa	1637	G	N1-C6-O6	9.15	125.39	119.90
84	Aa	2352	G	N1-C6-O6	9.15	125.39	119.90
84	Aa	2527	G	C5-C6-O6	-9.15	123.11	128.60
84	Aa	2700	A	N1-C6-N6	9.15	124.09	118.60
85	Ac	15	G	N1-C6-O6	9.15	125.39	119.90
85	Ac	56	G	N1-C6-O6	9.15	125.39	119.90
84	Aa	2092	C	O3'-P-O5'	-9.15	86.61	104.00
84	Aa	1708	C	O4'-C1'-N1	9.15	115.52	108.20
84	Aa	1556	G	N1-C6-O6	9.14	125.39	119.90
84	Aa	2752	G	N1-C6-O6	9.14	125.39	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	Ab	65	G	C5-C6-O6	-9.14	123.11	128.60
1	Ad	535	C	N1-C1'-C2'	9.14	125.88	114.00
1	Ad	1058	G	O4'-C1'-N9	9.14	115.51	108.20
84	Aa	2196	G	N1-C6-O6	9.14	125.38	119.90
84	Aa	210	G	C5-C6-O6	-9.14	123.12	128.60
84	Aa	3379	C	O4'-C1'-N1	9.14	115.51	108.20
86	Ab	69	A	N1-C6-N6	9.14	124.08	118.60
1	Ad	1500	A	N9-C1'-C2'	-9.13	101.95	112.00
84	Aa	2746	G	N1-C6-O6	9.14	125.38	119.90
84	Aa	1818	C	O4'-C1'-N1	9.13	115.51	108.20
84	Aa	3286	G	N1-C6-O6	9.13	125.38	119.90
1	Ad	611	G	O4'-C1'-N9	9.13	115.50	108.20
2	Ae	50	G	N9-C1'-C2'	-9.13	101.96	112.00
84	Aa	2800	C	O4'-C1'-N1	9.13	115.50	108.20
84	Aa	2818	G	N1-C6-O6	9.13	125.38	119.90
84	Aa	390	G	N1-C6-O6	9.13	125.38	119.90
84	Aa	1935	G	N1-C6-O6	9.13	125.38	119.90
84	Aa	2122	C	O4'-C1'-N1	9.13	115.50	108.20
84	Aa	2834	C	O4'-C1'-N1	9.13	115.50	108.20
84	Aa	265	G	N1-C6-O6	9.12	125.38	119.90
84	Aa	1710	G	N1-C6-O6	9.13	125.38	119.90
84	Aa	3040	G	N1-C6-O6	9.13	125.38	119.90
84	Aa	1356	G	N1-C6-O6	9.12	125.37	119.90
84	Aa	2086	A	C4-N9-C1'	9.12	142.72	126.30
84	Aa	3147	G	N1-C6-O6	9.12	125.37	119.90
1	Ad	824	U	O4'-C1'-N1	9.12	115.50	108.20
1	Ad	1484	U	O4'-C1'-N1	9.12	115.50	108.20
84	Aa	2789	G	N1-C6-O6	9.12	125.37	119.90
84	Aa	583	C	O4'-C1'-N1	9.12	115.49	108.20
84	Aa	2401	A	N1-C6-N6	9.12	124.07	118.60
1	Ad	914	U	C1'-O4'-C4'	-9.11	102.61	109.90
84	Aa	1693	A	N1-C6-N6	9.11	124.07	118.60
85	Ac	96	A	N1-C6-N6	9.11	124.07	118.60
84	Aa	90	G	N1-C6-O6	9.11	125.36	119.90
84	Aa	1076	G	N1-C6-O6	9.11	125.36	119.90
84	Aa	2390	G	N1-C6-O6	9.11	125.36	119.90
84	Aa	3264	C	O4'-C1'-N1	9.11	115.49	108.20
84	Aa	1167	G	N1-C6-O6	9.11	125.36	119.90
84	Aa	943	G	N1-C6-O6	9.10	125.36	119.90
84	Aa	2370	G	N1-C6-O6	9.10	125.36	119.90
84	Aa	3281	G	N1-C6-O6	9.10	125.36	119.90
84	Aa	1810	G	N1-C6-O6	9.10	125.36	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3156	G	N1-C6-O6	9.10	125.36	119.90
1	Ad	1280	U	O4'-C1'-N1	9.10	115.48	108.20
84	Aa	2073	U	P-O3'-C3'	-9.10	108.78	119.70
84	Aa	2076	C	O4'-C1'-N1	9.10	115.48	108.20
84	Aa	1058	A	N1-C6-N6	9.10	124.06	118.60
84	Aa	270	G	N1-C6-O6	9.09	125.36	119.90
1	Ad	1372	C	P-O3'-C3'	9.09	130.61	119.70
84	Aa	771	G	N1-C6-O6	9.09	125.35	119.90
84	Aa	3129	G	N1-C6-O6	9.09	125.35	119.90
84	Aa	2537	G	O4'-C1'-N9	9.09	115.47	108.20
1	Ad	41	A	O4'-C1'-C2'	-9.09	96.71	105.80
1	Ad	96	G	C1'-O4'-C4'	-9.09	102.63	109.90
1	Ad	843	G	O4'-C1'-N9	-9.09	100.93	108.20
84	Aa	3177	A	O4'-C1'-N9	9.09	115.47	108.20
1	Ad	1116	G	O4'-C1'-N9	9.09	115.47	108.20
84	Aa	1465	A	N1-C6-N6	9.09	124.05	118.60
84	Aa	760	C	O4'-C1'-N1	9.08	115.46	108.20
84	Aa	1233	G	N1-C6-O6	9.08	125.35	119.90
84	Aa	1729	G	N1-C6-O6	9.08	125.35	119.90
2	Ae	74	C	O4'-C1'-C2'	-9.08	96.72	105.80
84	Aa	1413	C	O4'-C1'-N1	9.08	115.46	108.20
84	Aa	2245	G	N1-C6-O6	9.07	125.34	119.90
84	Aa	638	G	N1-C6-O6	9.07	125.34	119.90
84	Aa	1561	U	O3'-P-O5'	9.07	121.23	104.00
84	Aa	3279	G	N1-C6-O6	9.07	125.34	119.90
84	Aa	1776	G	N1-C6-O6	9.07	125.34	119.90
1	Ad	1530	G	P-O3'-C3'	9.07	130.58	119.70
84	Aa	310	C	O4'-C1'-N1	9.07	115.45	108.20
84	Aa	1189	G	N1-C6-O6	9.07	125.34	119.90
84	Aa	1144	C	O4'-C1'-N1	9.06	115.45	108.20
84	Aa	1645	G	N1-C6-O6	9.06	125.34	119.90
1	Ad	633	U	O4'-C1'-N1	9.06	115.45	108.20
84	Aa	3001	G	N1-C6-O6	9.06	125.34	119.90
84	Aa	508	G	N1-C6-O6	9.06	125.33	119.90
84	Aa	1553	C	O4'-C1'-N1	9.06	115.44	108.20
84	Aa	1379	G	N1-C6-O6	9.05	125.33	119.90
84	Aa	1625	G	N1-C6-O6	9.05	125.33	119.90
84	Aa	2903	G	N1-C6-O6	9.05	125.33	119.90
1	Ad	817	C	O4'-C1'-N1	9.05	115.44	108.20
1	Ad	255	U	O4'-C1'-N1	9.05	115.44	108.20
84	Aa	96	C	O4'-C1'-N1	9.05	115.44	108.20
84	Aa	2023	C	O4'-C1'-N1	9.05	115.44	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	231	C	O4'-C1'-N1	9.05	115.44	108.20
84	Aa	1794	A	N1-C6-N6	9.05	124.03	118.60
1	Ad	1390	A	O4'-C1'-N9	9.04	115.44	108.20
1	Ad	845	C	C5'-C4'-C3'	9.04	130.47	116.00
1	Ad	1447	C	N1-C1'-C2'	9.04	125.76	114.00
1	Ad	1449	U	O4'-C1'-N1	9.04	115.43	108.20
84	Aa	3322	A	N1-C6-N6	9.04	124.03	118.60
84	Aa	2594	A	N1-C6-N6	9.04	124.03	118.60
85	Ac	36	G	N1-C6-O6	9.04	125.33	119.90
84	Aa	342	A	N1-C6-N6	9.04	124.02	118.60
84	Aa	971	G	N1-C6-O6	9.04	125.32	119.90
84	Aa	1168	G	N1-C6-O6	9.04	125.32	119.90
84	Aa	2589	G	N1-C6-O6	9.04	125.32	119.90
1	Ad	737	G	C4'-C3'-O3'	9.04	131.07	113.00
84	Aa	19	C	O4'-C1'-N1	9.04	115.43	108.20
84	Aa	485	G	N1-C6-O6	9.04	125.32	119.90
84	Aa	1985	G	N1-C6-O6	9.03	125.32	119.90
84	Aa	763	G	N1-C6-O6	9.03	125.32	119.90
84	Aa	2522	C	O4'-C1'-N1	9.03	115.42	108.20
1	Ad	1740	G	O4'-C1'-N9	9.03	115.42	108.20
84	Aa	1467	G	N1-C6-O6	9.03	125.32	119.90
1	Ad	1591	A	O4'-C1'-N9	9.03	115.42	108.20
84	Aa	725	G	N1-C6-O6	9.03	125.31	119.90
84	Aa	1447	G	N1-C6-O6	9.03	125.31	119.90
84	Aa	2383	G	N1-C6-O6	9.03	125.31	119.90
1	Ad	862	U	O4'-C1'-N1	9.02	115.42	108.20
1	Ad	1047	G	C1'-O4'-C4'	-9.02	102.68	109.90
84	Aa	700	C	O4'-C1'-N1	9.02	115.42	108.20
1	Ad	1748	U	O4'-C1'-N1	9.02	115.42	108.20
84	Aa	1226	G	N1-C6-O6	9.02	125.31	119.90
1	Ad	1782	C	O4'-C1'-N1	9.02	115.42	108.20
84	Aa	1443	G	N1-C6-O6	9.02	125.31	119.90
84	Aa	1798	C	O4'-C1'-N1	9.02	115.42	108.20
84	Aa	2170	G	N1-C6-O6	9.02	125.31	119.90
84	Aa	2676	A	N1-C6-N6	9.02	124.01	118.60
1	Ad	41	A	O4'-C1'-N9	9.02	115.41	108.20
1	Ad	270	U	O4'-C1'-N1	9.02	115.41	108.20
84	Aa	742	G	N1-C6-O6	9.02	125.31	119.90
84	Aa	497	G	N1-C6-O6	9.02	125.31	119.90
84	Aa	3357	C	O4'-C1'-N1	9.02	115.41	108.20
1	Ad	1206	A	O4'-C1'-N9	-9.01	100.99	108.20
84	Aa	1276	C	O4'-C1'-N1	9.01	115.41	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1303	C	O4'-C1'-N1	9.01	115.41	108.20
84	Aa	1631	G	N1-C6-O6	9.01	125.31	119.90
84	Aa	956	G	N1-C6-O6	9.01	125.31	119.90
86	Ab	33	U	O4'-C1'-N1	9.01	115.41	108.20
84	Aa	1893	G	N1-C6-O6	9.01	125.30	119.90
84	Aa	2527	G	O4'-C1'-N9	9.01	115.41	108.20
84	Aa	3325	G	N1-C6-O6	9.01	125.30	119.90
84	Aa	12	G	P-O3'-C3'	9.00	130.50	119.70
84	Aa	1547	G	N1-C6-O6	9.00	125.30	119.90
84	Aa	1725	G	N1-C6-O6	9.00	125.30	119.90
84	Aa	2855	G	N1-C6-O6	9.00	125.30	119.90
86	Ab	11	A	N1-C2-N3	9.00	133.80	129.30
84	Aa	998	G	N1-C6-O6	9.00	125.30	119.90
84	Aa	1909	G	N1-C6-O6	9.00	125.30	119.90
84	Aa	2085	A	P-O3'-C3'	-9.00	108.90	119.70
84	Aa	2656	C	O4'-C1'-N1	9.00	115.40	108.20
84	Aa	1583	G	N1-C6-O6	8.99	125.30	119.90
84	Aa	1950	G	O4'-C1'-N9	8.99	115.39	108.20
85	Ac	82	C	O4'-C1'-N1	8.99	115.39	108.20
86	Ab	58	G	C5-C6-O6	-8.99	123.20	128.60
84	Aa	890	G	N1-C6-O6	8.99	125.29	119.90
84	Aa	2825	G	N1-C6-O6	8.99	125.30	119.90
84	Aa	1	G	N1-C6-O6	8.99	125.29	119.90
1	Ad	824	U	P-O3'-C3'	8.98	130.48	119.70
1	Ad	1306	U	O4'-C1'-N1	8.98	115.39	108.20
1	Ad	1546	U	O4'-C1'-N1	8.98	115.39	108.20
84	Aa	17	G	N1-C6-O6	8.98	125.29	119.90
84	Aa	2236	U	O4'-C1'-N1	8.98	115.39	108.20
84	Aa	3331	G	N1-C6-O6	8.98	125.29	119.90
84	Aa	491	G	N1-C6-O6	8.98	125.29	119.90
84	Aa	2963	G	N1-C6-O6	8.98	125.29	119.90
84	Aa	927	G	N1-C6-O6	8.98	125.29	119.90
85	Ac	46	G	N1-C6-O6	8.98	125.29	119.90
1	Ad	1006	A	O4'-C1'-C2'	-8.98	96.82	105.80
1	Ad	1065	A	C1'-O4'-C4'	-8.98	102.72	109.90
84	Aa	3075	G	N1-C6-O6	8.97	125.28	119.90
84	Aa	1476	G	N1-C6-O6	8.97	125.28	119.90
84	Aa	1967	C	O4'-C1'-N1	8.97	115.38	108.20
84	Aa	803	G	N1-C6-O6	8.97	125.28	119.90
84	Aa	2213	G	N1-C6-O6	8.97	125.28	119.90
1	Ad	306	U	O4'-C1'-N1	8.96	115.37	108.20
84	Aa	1697	G	N1-C6-O6	8.96	125.28	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2204	U	P-O3'-C3'	8.96	130.46	119.70
84	Aa	522	C	O4'-C1'-N1	8.96	115.37	108.20
84	Aa	1619	G	N1-C6-O6	8.96	125.28	119.90
84	Aa	386	G	N1-C6-O6	8.96	125.28	119.90
84	Aa	2283	G	N1-C6-O6	8.96	125.28	119.90
85	Ac	120	G	N1-C6-O6	8.96	125.28	119.90
84	Aa	1978	G	N1-C6-O6	8.96	125.27	119.90
84	Aa	3029	G	N1-C6-O6	8.96	125.27	119.90
1	Ad	271	C	O4'-C1'-N1	8.96	115.36	108.20
84	Aa	584	G	N1-C6-O6	8.96	125.27	119.90
84	Aa	1001	A	N1-C6-N6	8.96	123.97	118.60
84	Aa	1764	G	N1-C6-O6	8.96	125.27	119.90
84	Aa	2428	G	N1-C6-O6	8.96	125.27	119.90
1	Ad	636	U	O4'-C1'-N1	8.95	115.36	108.20
84	Aa	1381	G	N1-C6-O6	8.95	125.27	119.90
84	Aa	1803	G	N1-C6-O6	8.95	125.27	119.90
84	Aa	2723	G	N1-C6-O6	8.95	125.27	119.90
1	Ad	1348	A	P-O3'-C3'	8.95	130.44	119.70
84	Aa	1601	G	N1-C6-O6	8.94	125.27	119.90
84	Aa	2086	A	C8-N9-C1'	-8.94	111.60	127.70
84	Aa	3367	C	O4'-C1'-N1	8.94	115.36	108.20
84	Aa	1143	G	N1-C6-O6	8.94	125.27	119.90
1	Ad	221	U	O4'-C1'-N1	8.94	115.35	108.20
84	Aa	2024	G	N1-C6-O6	8.94	125.26	119.90
84	Aa	176	A	N1-C6-N6	8.94	123.96	118.60
84	Aa	1716	G	C5-C6-O6	-8.94	123.24	128.60
84	Aa	3141	G	N1-C6-O6	8.94	125.26	119.90
84	Aa	3106	U	O4'-C1'-N1	8.94	115.35	108.20
1	Ad	174	C	O4'-C1'-N1	8.93	115.35	108.20
84	Aa	1621	G	N1-C6-O6	8.93	125.26	119.90
1	Ad	393	G	O4'-C1'-N9	8.93	115.35	108.20
1	Ad	1796	G	C1'-O4'-C4'	-8.93	102.75	109.90
84	Aa	1979	G	N1-C6-O6	8.93	125.26	119.90
85	Ac	114	G	N1-C6-O6	8.93	125.26	119.90
1	Ad	880	G	O4'-C1'-N9	8.93	115.34	108.20
1	Ad	1805	U	O4'-C1'-N1	8.93	115.34	108.20
84	Aa	97	G	N1-C6-O6	8.93	125.26	119.90
84	Aa	1411	G	N1-C6-O6	8.93	125.26	119.90
84	Aa	2657	C	O4'-C1'-N1	8.93	115.34	108.20
1	Ad	836	U	C3'-C2'-C1'	8.93	108.64	101.50
84	Aa	1898	G	N1-C6-O6	8.93	125.25	119.90
84	Aa	1915	G	N1-C6-O6	8.93	125.25	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	Ab	16	A	C5-C6-N1	-8.93	113.24	117.70
84	Aa	194	G	N1-C6-O6	8.92	125.25	119.90
84	Aa	3079	G	N1-C6-O6	8.92	125.25	119.90
84	Aa	986	G	N1-C6-O6	8.92	125.25	119.90
84	Aa	1348	G	N1-C6-O6	8.92	125.25	119.90
84	Aa	2189	G	N1-C6-O6	8.92	125.25	119.90
84	Aa	3188	G	C5-C6-O6	-8.92	123.25	128.60
84	Aa	3222	G	N1-C6-O6	8.92	125.25	119.90
1	Ad	518	G	P-O3'-C3'	8.92	130.40	119.70
84	Aa	542	G	P-O3'-C3'	8.92	130.40	119.70
84	Aa	2105	G	N1-C6-O6	8.92	125.25	119.90
1	Ad	500	G	O4'-C1'-N9	8.91	115.33	108.20
1	Ad	944	A	O4'-C1'-C2'	-8.91	96.89	105.80
1	Ad	1513	A	C1'-O4'-C4'	8.91	117.03	109.90
84	Aa	450	C	O4'-C1'-N1	8.91	115.33	108.20
84	Aa	867	G	N1-C6-O6	8.91	125.25	119.90
85	Ac	75	G	N1-C6-O6	8.91	125.25	119.90
85	Ac	149	U	O4'-C1'-N1	8.91	115.33	108.20
84	Aa	2874	A	N1-C6-N6	8.91	123.95	118.60
1	Ad	989	G	O4'-C1'-N9	8.91	115.33	108.20
84	Aa	1589	G	P-O3'-C3'	8.91	130.39	119.70
85	Ac	148	C	O4'-C1'-N1	8.91	115.33	108.20
86	Ab	23	A	N1-C6-N6	8.91	123.94	118.60
2	Ae	74	C	C3'-C2'-C1'	8.91	108.62	101.50
84	Aa	486	G	N1-C6-O6	8.91	125.24	119.90
84	Aa	1567	G	N1-C6-O6	8.91	125.24	119.90
84	Aa	2897	G	N1-C6-O6	8.91	125.24	119.90
1	Ad	1766	A	P-O3'-C3'	8.90	130.38	119.70
84	Aa	2731	G	N1-C6-O6	8.90	125.24	119.90
84	Aa	3282	G	C5-C6-O6	-8.90	123.26	128.60
1	Ad	517	U	O4'-C1'-N1	8.90	115.32	108.20
84	Aa	1823	C	O4'-C1'-N1	8.90	115.32	108.20
84	Aa	2487	A	O4'-C1'-N9	8.90	115.32	108.20
84	Aa	3264	C	P-O3'-C3'	8.90	130.38	119.70
84	Aa	1822	C	O4'-C1'-N1	8.90	115.32	108.20
84	Aa	894	G	N1-C6-O6	8.90	125.24	119.90
84	Aa	1990	A	P-O3'-C3'	8.90	130.38	119.70
84	Aa	2376	G	O4'-C1'-N9	8.90	115.32	108.20
84	Aa	3242	G	N1-C6-O6	8.90	125.24	119.90
84	Aa	434	C	O4'-C1'-N1	8.89	115.32	108.20
84	Aa	2601	G	N1-C6-O6	8.89	125.24	119.90
84	Aa	2651	G	N1-C6-O6	8.89	125.24	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2712	C	O4'-C1'-N1	8.89	115.31	108.20
86	Ab	119	C	O4'-C1'-N1	8.89	115.31	108.20
1	Ad	1343	C	O4'-C1'-N1	8.89	115.31	108.20
84	Aa	870	G	N1-C6-O6	8.89	125.23	119.90
84	Aa	2086	A	N9-C1'-C2'	8.89	125.56	114.00
84	Aa	590	C	P-O3'-C3'	-8.88	109.04	119.70
84	Aa	1329	G	N1-C6-O6	8.88	125.23	119.90
84	Aa	495	G	O5'-P-OP2	-8.88	97.71	105.70
84	Aa	1405	G	N1-C6-O6	8.88	125.23	119.90
84	Aa	2068	G	N1-C6-O6	8.88	125.23	119.90
84	Aa	3183	G	N1-C6-O6	8.88	125.23	119.90
84	Aa	1403	G	N1-C6-O6	8.88	125.23	119.90
84	Aa	1419	G	N1-C6-O6	8.88	125.23	119.90
84	Aa	2416	U	O4'-C1'-N1	8.88	115.31	108.20
1	Ad	38	C	O4'-C1'-N1	8.88	115.30	108.20
84	Aa	2300	G	N1-C6-O6	8.88	125.23	119.90
84	Aa	362	G	C5-C6-O6	-8.88	123.27	128.60
84	Aa	381	G	N1-C6-O6	8.88	125.23	119.90
84	Aa	2966	G	N1-C6-O6	8.87	125.22	119.90
84	Aa	3037	G	N1-C6-O6	8.87	125.22	119.90
84	Aa	3372	C	O4'-C1'-N1	8.87	115.30	108.20
1	Ad	1202	G	O4'-C1'-N9	8.87	115.30	108.20
84	Aa	3164	C	O4'-C1'-N1	8.87	115.30	108.20
85	Ac	25	G	N1-C6-O6	8.87	125.22	119.90
1	Ad	770	U	O4'-C1'-N1	8.87	115.30	108.20
84	Aa	1142	G	N1-C6-O6	8.87	125.22	119.90
85	Ac	31	G	N1-C6-O6	8.87	125.22	119.90
1	Ad	488	C	N1-C1'-C2'	8.87	125.53	114.00
84	Aa	1808	G	N1-C6-O6	8.87	125.22	119.90
84	Aa	2465	G	N1-C6-O6	8.87	125.22	119.90
84	Aa	2475	C	C4'-C3'-O3'	8.87	130.74	113.00
84	Aa	8	C	O4'-C1'-N1	8.87	115.29	108.20
1	Ad	1375	C	O4'-C1'-C2'	-8.86	96.94	105.80
1	Ad	281	U	O4'-C1'-C2'	-8.86	96.94	105.80
1	Ad	637	U	O4'-C1'-N1	8.86	115.29	108.20
84	Aa	796	C	O4'-C1'-N1	8.86	115.29	108.20
84	Aa	859	G	N1-C6-O6	8.86	125.22	119.90
84	Aa	1161	G	N1-C6-O6	8.86	125.21	119.90
84	Aa	2404	C	O4'-C1'-N1	8.86	115.29	108.20
84	Aa	937	G	N1-C6-O6	8.85	125.21	119.90
84	Aa	2466	G	N1-C6-O6	8.85	125.21	119.90
84	Aa	3066	G	N1-C6-O6	8.85	125.21	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3206	C	C2-N1-C1'	8.85	128.54	118.80
86	Ab	38	U	O4'-C1'-N1	8.85	115.28	108.20
1	Ad	422	G	O4'-C1'-N9	8.85	115.28	108.20
1	Ad	1651	U	P-O3'-C3'	8.85	130.32	119.70
84	Aa	1304	G	N1-C6-O6	8.85	125.21	119.90
84	Aa	1696	G	N1-C6-O6	8.85	125.21	119.90
1	Ad	1101	C	N1-C1'-C2'	8.85	125.50	114.00
1	Ad	1586	U	O4'-C1'-N1	8.85	115.28	108.20
84	Aa	787	G	N1-C6-O6	8.85	125.21	119.90
84	Aa	1785	G	N1-C6-O6	8.85	125.21	119.90
84	Aa	1587	G	N1-C6-O6	8.85	125.21	119.90
84	Aa	277	U	O4'-C1'-N1	8.84	115.27	108.20
84	Aa	1079	G	N1-C6-O6	8.84	125.21	119.90
84	Aa	1231	C	O4'-C1'-N1	8.84	115.27	108.20
84	Aa	2995	G	N1-C6-O6	8.84	125.21	119.90
85	Ac	76	C	O4'-C1'-N1	8.84	115.28	108.20
84	Aa	2802	G	N1-C6-O6	8.84	125.20	119.90
84	Aa	2886	C	O4'-C1'-N1	8.84	115.27	108.20
84	Aa	3210	G	N1-C6-O6	8.84	125.20	119.90
1	Ad	836	U	O4'-C1'-C2'	-8.84	96.96	105.80
1	Ad	1094	U	N1-C1'-C2'	8.84	125.49	114.00
84	Aa	714	G	N1-C6-O6	8.84	125.20	119.90
84	Aa	1289	G	N1-C6-O6	8.83	125.20	119.90
84	Aa	1960	C	O4'-C1'-N1	8.83	115.27	108.20
1	Ad	903	A	C3'-C2'-C1'	-8.83	94.44	101.50
84	Aa	2057	G	N1-C6-O6	8.83	125.20	119.90
84	Aa	44	A	N1-C6-N6	8.83	123.90	118.60
84	Aa	915	G	N1-C6-O6	8.83	125.20	119.90
84	Aa	1300	C	O4'-C1'-N1	8.83	115.26	108.20
84	Aa	1540	G	N1-C6-O6	8.83	125.20	119.90
86	Ab	62	U	O4'-C1'-N1	8.83	115.26	108.20
84	Aa	784	G	N1-C6-O6	8.82	125.19	119.90
84	Aa	2648	G	N1-C6-O6	8.82	125.19	119.90
1	Ad	196	G	N9-C1'-C2'	8.82	125.47	114.00
84	Aa	1936	G	N1-C6-O6	8.82	125.19	119.90
1	Ad	1	U	O4'-C1'-C2'	-8.82	96.98	105.80
84	Aa	206	C	O4'-C1'-N1	8.82	115.25	108.20
84	Aa	392	C	O4'-C1'-N1	8.82	115.26	108.20
84	Aa	2368	G	N1-C6-O6	8.82	125.19	119.90
84	Aa	3356	C	O4'-C1'-N1	8.82	115.25	108.20
84	Aa	602	G	N1-C6-O6	8.82	125.19	119.90
84	Aa	2664	G	N1-C6-O6	8.82	125.19	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2812	C	O4'-C1'-N1	8.82	115.25	108.20
84	Aa	1999	G	N1-C6-O6	8.81	125.19	119.90
84	Aa	2748	G	N1-C6-O6	8.81	125.19	119.90
1	Ad	1626	C	C3'-C2'-C1'	8.81	108.55	101.50
84	Aa	1402	G	N1-C6-O6	8.81	125.19	119.90
1	Ad	143	A	O4'-C1'-N9	8.81	115.25	108.20
84	Aa	1175	G	N1-C6-O6	8.81	125.19	119.90
84	Aa	22	G	N1-C6-O6	8.81	125.18	119.90
84	Aa	665	G	N1-C6-O6	8.81	125.18	119.90
84	Aa	2337	C	O4'-C1'-N1	8.81	115.25	108.20
84	Aa	2582	G	N1-C6-O6	8.81	125.18	119.90
84	Aa	431	G	C5-C6-O6	-8.80	123.32	128.60
84	Aa	1496	G	N1-C6-O6	8.81	125.18	119.90
84	Aa	2627	G	N1-C6-O6	8.80	125.18	119.90
84	Aa	2959	G	N1-C6-O6	8.80	125.18	119.90
84	Aa	3298	G	O4'-C1'-N9	8.80	115.24	108.20
1	Ad	806	U	O4'-C1'-N1	8.80	115.24	108.20
84	Aa	1417	G	N1-C6-O6	8.80	125.18	119.90
86	Ab	60	G	C5-C6-O6	-8.80	123.32	128.60
1	Ad	32	U	N1-C1'-C2'	-8.80	102.32	112.00
84	Aa	1800	G	N1-C6-O6	8.80	125.18	119.90
84	Aa	2462	G	N1-C6-O6	8.80	125.18	119.90
84	Aa	3314	G	N1-C6-O6	8.80	125.18	119.90
1	Ad	222	G	O4'-C1'-N9	8.79	115.24	108.20
84	Aa	413	G	N1-C6-O6	8.80	125.18	119.90
84	Aa	445	C	O4'-C1'-N1	8.79	115.24	108.20
84	Aa	663	G	N1-C6-O6	8.79	125.18	119.90
84	Aa	2205	G	N1-C6-O6	8.80	125.18	119.90
84	Aa	3230	G	N1-C6-O6	8.80	125.18	119.90
84	Aa	60	G	N1-C6-O6	8.79	125.18	119.90
84	Aa	881	G	N1-C6-O6	8.79	125.17	119.90
84	Aa	1474	U	O4'-C1'-N1	8.79	115.23	108.20
84	Aa	2235	G	N1-C6-O6	8.79	125.18	119.90
84	Aa	3369	G	N1-C6-O6	8.79	125.17	119.90
84	Aa	429	G	N1-C6-O6	8.79	125.17	119.90
84	Aa	729	G	N1-C6-O6	8.79	125.17	119.90
84	Aa	2274	A	N1-C6-N6	8.79	123.87	118.60
84	Aa	3171	C	O4'-C1'-N1	8.79	115.23	108.20
84	Aa	640	C	P-O3'-C3'	-8.79	109.16	119.70
1	Ad	1016	C	N1-C1'-C2'	8.78	125.42	114.00
84	Aa	3027	G	N1-C6-O6	8.78	125.17	119.90
84	Aa	272	G	N1-C6-O6	8.78	125.17	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1432	G	N1-C6-O6	8.78	125.17	119.90
84	Aa	1588	G	N1-C6-O6	8.78	125.17	119.90
84	Aa	3237	G	N1-C6-O6	8.78	125.17	119.90
84	Aa	1354	G	N1-C6-O6	8.78	125.17	119.90
84	Aa	1925	G	N1-C6-O6	8.78	125.17	119.90
84	Aa	2878	C	O4'-C1'-N1	8.78	115.22	108.20
84	Aa	3172	G	O4'-C1'-N9	8.78	115.22	108.20
1	Ad	79	A	N9-C1'-C2'	-8.78	102.34	112.00
84	Aa	703	G	N1-C6-O6	8.78	125.17	119.90
84	Aa	1258	C	O4'-C1'-N1	8.78	115.22	108.20
84	Aa	1483	G	N1-C6-O6	8.78	125.17	119.90
84	Aa	2256	G	N1-C6-O6	8.78	125.17	119.90
84	Aa	2413	G	N1-C6-O6	8.78	125.17	119.90
84	Aa	2740	C	O4'-C1'-N1	8.78	115.22	108.20
84	Aa	3150	G	N1-C6-O6	8.78	125.17	119.90
84	Aa	3224	C	O4'-C1'-N1	8.78	115.22	108.20
84	Aa	3186	G	N1-C6-O6	8.77	125.16	119.90
84	Aa	3335	G	N1-C6-O6	8.77	125.16	119.90
1	Ad	201	G	N9-C1'-C2'	8.77	125.40	114.00
1	Ad	1432	C	O4'-C1'-N1	8.77	115.22	108.20
1	Ad	1778	G	N9-C1'-C2'	8.77	125.40	114.00
84	Aa	245	C	O4'-C1'-N1	8.77	115.22	108.20
84	Aa	1271	U	O4'-C1'-N1	8.77	115.21	108.20
84	Aa	1988	G	N1-C6-O6	8.77	125.16	119.90
84	Aa	3332	G	N1-C6-O6	8.77	125.16	119.90
85	Ac	159	G	P-O3'-C3'	8.77	130.22	119.70
84	Aa	534	G	N1-C6-O6	8.77	125.16	119.90
84	Aa	2258	C	O4'-C1'-N1	8.76	115.21	108.20
84	Aa	3384	G	N1-C6-O6	8.76	125.16	119.90
85	Ac	28	C	O4'-C1'-N1	8.76	115.21	108.20
84	Aa	438	G	N1-C6-O6	8.76	125.16	119.90
84	Aa	2115	G	N1-C6-O6	8.76	125.16	119.90
84	Aa	2578	G	N1-C6-O6	8.76	125.16	119.90
1	Ad	999	G	O4'-C1'-C2'	8.76	115.48	107.60
1	Ad	1191	U	O4'-C1'-N1	8.76	115.21	108.20
84	Aa	1134	G	N1-C6-O6	8.76	125.15	119.90
84	Aa	1283	C	O4'-C1'-N1	8.76	115.20	108.20
84	Aa	1699	C	O4'-C1'-N1	8.76	115.20	108.20
84	Aa	2012	C	P-O3'-C3'	8.76	130.21	119.70
84	Aa	2617	G	N1-C6-O6	8.76	125.15	119.90
84	Aa	3311	C	O4'-C1'-N1	8.76	115.20	108.20
85	Ac	158	C	O4'-C1'-N1	8.76	115.20	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1407	G	N1-C6-O6	8.75	125.15	119.90
84	Aa	1500	C	O4'-C1'-N1	8.75	115.20	108.20
84	Aa	2541	A	N1-C6-N6	8.75	123.85	118.60
84	Aa	3062	G	N1-C6-O6	8.75	125.15	119.90
1	Ad	281	U	O3'-P-O5'	-8.75	87.38	104.00
84	Aa	754	G	N1-C6-O6	8.75	125.15	119.90
84	Aa	773	G	N1-C6-O6	8.75	125.15	119.90
84	Aa	2375	G	N1-C6-O6	8.75	125.15	119.90
84	Aa	2979	G	N1-C6-O6	8.75	125.15	119.90
84	Aa	3084	G	N1-C6-O6	8.75	125.15	119.90
84	Aa	3337	G	N1-C6-O6	8.75	125.15	119.90
84	Aa	1100	G	N1-C6-O6	8.75	125.15	119.90
84	Aa	2780	G	N1-C6-O6	8.75	125.15	119.90
1	Ad	898	U	O4'-C1'-N1	8.74	115.20	108.20
84	Aa	114	G	N1-C6-O6	8.74	125.15	119.90
84	Aa	1728	G	N1-C6-O6	8.74	125.14	119.90
84	Aa	2838	C	O4'-C1'-N1	8.74	115.19	108.20
86	Ab	5	G	N1-C6-O6	8.74	125.14	119.90
1	Ad	756	U	O4'-C1'-N1	8.74	115.19	108.20
1	Ad	796	U	O4'-C1'-N1	8.74	115.19	108.20
84	Aa	199	G	N1-C6-O6	8.74	125.14	119.90
84	Aa	669	G	N1-C6-O6	8.74	125.14	119.90
84	Aa	2667	C	O4'-C1'-N1	8.74	115.19	108.20
84	Aa	944	G	N1-C6-O6	8.74	125.14	119.90
1	Ad	1170	G	O4'-C1'-N9	8.74	115.19	108.20
84	Aa	136	C	O4'-C1'-N1	8.74	115.19	108.20
84	Aa	2598	A	N1-C6-N6	8.74	123.84	118.60
84	Aa	126	G	N1-C6-O6	8.74	125.14	119.90
84	Aa	1299	G	N1-C6-O6	8.74	125.14	119.90
1	Ad	1325	A	C1'-O4'-C4'	-8.73	102.91	109.90
84	Aa	57	G	N1-C6-O6	8.73	125.14	119.90
84	Aa	2075	C	O4'-C1'-N1	8.73	115.19	108.20
84	Aa	2021	G	N1-C6-O6	8.73	125.14	119.90
85	Ac	24	G	N1-C6-O6	8.73	125.14	119.90
1	Ad	1061	G	O4'-C1'-N9	8.73	115.19	108.20
1	Ad	1474	U	O4'-C1'-N1	8.73	115.19	108.20
84	Aa	82	C	O4'-C1'-N1	8.73	115.18	108.20
84	Aa	531	G	N1-C6-O6	8.73	125.14	119.90
84	Aa	1982	G	N1-C6-O6	8.73	125.14	119.90
84	Aa	2540	C	O4'-C1'-N1	8.73	115.18	108.20
84	Aa	2448	G	N1-C6-O6	8.73	125.14	119.90
84	Aa	2128	G	N1-C6-O6	8.72	125.14	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1986	G	N1-C6-O6	8.72	125.13	119.90
1	Ad	67	G	O4'-C1'-N9	-8.72	101.22	108.20
84	Aa	208	G	N1-C6-O6	8.72	125.13	119.90
84	Aa	212	G	N1-C6-O6	8.72	125.13	119.90
84	Aa	357	C	O4'-C1'-N1	8.72	115.18	108.20
84	Aa	1767	G	N1-C6-O6	8.72	125.13	119.90
84	Aa	2067	G	N1-C6-O6	8.72	125.13	119.90
84	Aa	2112	C	O4'-C1'-N1	8.72	115.18	108.20
84	Aa	2231	G	C5-C6-O6	-8.72	123.37	128.60
84	Aa	2968	G	N1-C6-O6	8.72	125.13	119.90
84	Aa	358	G	N1-C6-O6	8.72	125.13	119.90
84	Aa	431	G	P-O3'-C3'	8.72	130.16	119.70
84	Aa	87	A	C4-C5-C6	8.71	121.36	117.00
84	Aa	790	G	N1-C6-O6	8.71	125.13	119.90
1	Ad	392	G	O4'-C1'-N9	8.71	115.17	108.20
1	Ad	870	A	C1'-O4'-C4'	8.71	116.87	109.90
84	Aa	100	C	O4'-C1'-N1	8.71	115.17	108.20
84	Aa	1701	G	N1-C6-O6	8.71	125.13	119.90
84	Aa	3295	G	N1-C6-O6	8.71	125.13	119.90
85	Ac	159	G	N1-C6-O6	8.71	125.13	119.90
84	Aa	2034	G	N1-C6-O6	8.71	125.13	119.90
84	Aa	3103	G	N1-C6-O6	8.71	125.13	119.90
84	Aa	3274	G	N1-C6-O6	8.71	125.13	119.90
1	Ad	1776	A	O4'-C1'-N9	8.71	115.17	108.20
84	Aa	545	C	O4'-C1'-N1	8.71	115.17	108.20
84	Aa	1779	C	O4'-C1'-N1	8.71	115.17	108.20
84	Aa	2763	C	O4'-C1'-N1	8.71	115.17	108.20
86	Ab	118	C	O4'-C1'-N1	8.71	115.17	108.20
1	Ad	1593	U	N1-C1'-C2'	8.71	125.32	114.00
84	Aa	1085	G	N1-C6-O6	8.70	125.12	119.90
84	Aa	1119	G	N1-C6-O6	8.71	125.12	119.90
84	Aa	1572	C	O4'-C1'-N1	8.71	115.16	108.20
84	Aa	2493	C	O4'-C1'-N1	8.71	115.17	108.20
1	Ad	4	C	C1'-O4'-C4'	-8.70	102.94	109.90
84	Aa	493	G	N1-C6-O6	8.70	125.12	119.90
84	Aa	2728	C	O4'-C1'-N1	8.70	115.16	108.20
84	Aa	2860	U	O4'-C1'-N1	8.70	115.16	108.20
84	Aa	757	G	N1-C6-O6	8.70	125.12	119.90
85	Ac	39	G	N1-C6-O6	8.70	125.12	119.90
84	Aa	441	G	N1-C6-O6	8.70	125.12	119.90
84	Aa	1825	G	N1-C6-O6	8.70	125.12	119.90
84	Aa	2450	G	P-O3'-C3'	8.70	130.14	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	454	U	O4'-C1'-N1	8.70	115.16	108.20
1	Ad	811	U	O4'-C1'-N1	8.70	115.16	108.20
84	Aa	741	G	N1-C6-O6	8.70	125.12	119.90
84	Aa	2271	G	N1-C6-O6	8.70	125.12	119.90
84	Aa	1213	G	N1-C6-O6	8.69	125.11	119.90
84	Aa	3083	C	O4'-C1'-N1	8.69	115.15	108.20
84	Aa	3343	U	P-O3'-C3'	8.69	130.13	119.70
84	Aa	713	G	N1-C6-O6	8.69	125.11	119.90
84	Aa	782	G	N1-C6-O6	8.69	125.11	119.90
84	Aa	838	G	N1-C6-O6	8.69	125.11	119.90
84	Aa	1670	G	N1-C6-O6	8.69	125.11	119.90
84	Aa	2589	G	P-O3'-C3'	8.69	130.12	119.70
84	Aa	2735	G	N1-C6-O6	8.69	125.11	119.90
84	Aa	1268	G	N1-C6-O6	8.69	125.11	119.90
84	Aa	3023	G	N1-C6-O6	8.69	125.11	119.90
1	Ad	753	C	N1-C1'-C2'	8.69	125.29	114.00
84	Aa	1949	G	N1-C6-O6	8.69	125.11	119.90
84	Aa	1954	G	N1-C6-O6	8.69	125.11	119.90
84	Aa	3301	G	N1-C6-O6	8.69	125.11	119.90
84	Aa	3329	G	N1-C6-O6	8.69	125.11	119.90
1	Ad	787	C	C3'-C2'-C1'	-8.68	94.55	101.50
1	Ad	956	A	N9-C1'-C2'	-8.68	102.45	112.00
84	Aa	1404	G	N1-C6-O6	8.68	125.11	119.90
1	Ad	206	U	O4'-C1'-N1	8.68	115.15	108.20
84	Aa	1100	G	P-O3'-C3'	8.68	130.12	119.70
84	Aa	1941	G	N1-C6-O6	8.68	125.11	119.90
1	Ad	1363	G	C3'-C2'-C1'	-8.68	94.56	101.50
84	Aa	356	G	N1-C6-O6	8.68	125.11	119.90
84	Aa	720	G	N1-C6-O6	8.68	125.11	119.90
84	Aa	667	C	O4'-C1'-N1	8.68	115.14	108.20
84	Aa	988	G	N1-C6-O6	8.68	125.11	119.90
84	Aa	1723	C	O4'-C1'-N1	8.68	115.14	108.20
84	Aa	1804	G	N1-C6-O6	8.68	125.11	119.90
84	Aa	2048	C	O4'-C1'-N1	8.68	115.14	108.20
84	Aa	2865	G	N1-C6-O6	8.68	125.11	119.90
84	Aa	3172	G	N1-C6-O6	8.68	125.11	119.90
84	Aa	527	G	N1-C6-O6	8.67	125.10	119.90
84	Aa	745	G	N1-C6-O6	8.67	125.10	119.90
1	Ad	28	A	O4'-C1'-N9	8.67	115.14	108.20
84	Aa	200	G	N1-C6-O6	8.67	125.10	119.90
84	Aa	732	G	N1-C6-O6	8.67	125.10	119.90
84	Aa	802	G	N1-C6-O6	8.67	125.10	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1238	G	O5'-P-OP1	-8.67	97.90	105.70
84	Aa	1877	G	N1-C6-O6	8.67	125.10	119.90
84	Aa	2858	G	N1-C6-O6	8.67	125.10	119.90
84	Aa	2488	A	N1-C6-N6	8.67	123.80	118.60
1	Ad	136	U	O4'-C1'-N1	8.67	115.13	108.20
84	Aa	3245	G	N1-C6-O6	8.67	125.10	119.90
84	Aa	112	C	O4'-C1'-N1	8.67	115.13	108.20
84	Aa	3361	G	N1-C6-O6	8.67	125.10	119.90
84	Aa	535	G	N1-C6-O6	8.66	125.10	119.90
84	Aa	150	G	N1-C6-O6	8.66	125.10	119.90
84	Aa	1555	G	N1-C6-O6	8.66	125.10	119.90
84	Aa	1807	C	O4'-C1'-N1	8.66	115.13	108.20
84	Aa	2051	G	N1-C6-O6	8.66	125.10	119.90
1	Ad	165	U	N1-C1'-C2'	8.66	125.26	114.00
1	Ad	617	G	O4'-C1'-N9	-8.66	101.27	108.20
84	Aa	404	G	O4'-C1'-N9	8.66	115.13	108.20
84	Aa	1689	G	C5-C6-O6	-8.66	123.40	128.60
84	Aa	1759	C	O4'-C1'-N1	8.66	115.13	108.20
84	Aa	1826	G	N1-C6-O6	8.66	125.10	119.90
84	Aa	1980	C	O4'-C1'-N1	8.66	115.13	108.20
84	Aa	2031	G	N1-C6-O6	8.66	125.09	119.90
84	Aa	124	C	O4'-C1'-N1	8.66	115.12	108.20
84	Aa	142	G	N1-C6-O6	8.66	125.09	119.90
84	Aa	1340	G	N1-C6-O6	8.66	125.09	119.90
84	Aa	2703	G	N1-C6-O6	8.66	125.09	119.90
84	Aa	1974	C	O4'-C1'-N1	8.65	115.12	108.20
84	Aa	3167	G	O4'-C1'-N9	8.65	115.12	108.20
1	Ad	1029	U	O4'-C1'-C2'	-8.65	97.15	105.80
84	Aa	490	G	N1-C6-O6	8.65	125.09	119.90
84	Aa	1129	G	N1-C6-O6	8.65	125.09	119.90
84	Aa	2355	A	N1-C6-N6	8.65	123.79	118.60
1	Ad	1473	C	O4'-C1'-N1	8.65	115.12	108.20
84	Aa	613	G	N1-C6-O6	8.65	125.09	119.90
84	Aa	691	U	O4'-C1'-N1	8.65	115.12	108.20
84	Aa	1245	U	P-O5'-C5'	8.65	134.74	120.90
84	Aa	2050	G	N1-C6-O6	8.65	125.09	119.90
84	Aa	253	G	N1-C6-O6	8.64	125.09	119.90
84	Aa	254	G	N1-C6-O6	8.64	125.09	119.90
84	Aa	834	G	N1-C6-O6	8.64	125.09	119.90
84	Aa	2017	G	N1-C6-O6	8.64	125.09	119.90
84	Aa	2673	G	N1-C6-O6	8.64	125.09	119.90
84	Aa	2732	U	O4'-C1'-N1	8.64	115.11	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3077	C	O4'-C1'-N1	8.64	115.11	108.20
84	Aa	521	G	P-O3'-C3'	8.64	130.07	119.70
84	Aa	477	C	O4'-C1'-N1	8.64	115.11	108.20
84	Aa	2399	G	N1-C6-O6	8.64	125.08	119.90
84	Aa	2539	G	N1-C6-O6	8.64	125.08	119.90
1	Ad	438	G	O4'-C1'-N9	8.63	115.11	108.20
84	Aa	863	G	N1-C6-O6	8.63	125.08	119.90
84	Aa	2002	G	N1-C6-O6	8.63	125.08	119.90
84	Aa	3231	G	N1-C6-O6	8.63	125.08	119.90
86	Ab	57	C	N3-C4-N4	8.64	124.05	118.00
1	Ad	969	U	N1-C1'-C2'	8.63	125.22	114.00
84	Aa	567	G	C5-C6-O6	-8.63	123.42	128.60
84	Aa	1914	C	O4'-C1'-N1	8.63	115.11	108.20
84	Aa	2027	G	N1-C6-O6	8.63	125.08	119.90
84	Aa	2536	G	N1-C6-O6	8.63	125.08	119.90
84	Aa	748	C	O4'-C1'-N1	8.63	115.10	108.20
84	Aa	1544	G	N1-C6-O6	8.63	125.08	119.90
84	Aa	2411	G	N1-C6-O6	8.63	125.08	119.90
84	Aa	2924	G	N1-C6-O6	8.63	125.08	119.90
1	Ad	351	G	O4'-C1'-N9	8.63	115.10	108.20
84	Aa	1859	G	N1-C6-O6	8.62	125.08	119.90
84	Aa	2505	C	O4'-C1'-N1	8.62	115.10	108.20
1	Ad	1636	U	O4'-C1'-N1	8.62	115.10	108.20
2	Ae	8	U	O4'-C1'-N1	8.62	115.10	108.20
84	Aa	837	C	O4'-C1'-N1	8.62	115.10	108.20
84	Aa	1655	G	N1-C6-O6	8.62	125.07	119.90
84	Aa	2624	G	N1-C6-O6	8.62	125.07	119.90
84	Aa	512	G	O4'-C1'-N9	8.62	115.10	108.20
84	Aa	1263	A	O4'-C1'-N9	8.62	115.10	108.20
1	Ad	124	G	O4'-C1'-N9	8.62	115.09	108.20
1	Ad	748	C	P-O3'-C3'	8.62	130.04	119.70
84	Aa	2795	G	N1-C6-O6	8.62	125.07	119.90
84	Aa	3160	G	N1-C6-O6	8.62	125.07	119.90
84	Aa	3353	G	N1-C6-O6	8.62	125.07	119.90
85	Ac	156	C	O4'-C1'-N1	8.62	115.09	108.20
84	Aa	2441	G	N1-C6-O6	8.62	125.07	119.90
1	Ad	399	U	O4'-C1'-N1	8.62	115.09	108.20
84	Aa	351	G	N1-C6-O6	8.62	125.07	119.90
84	Aa	464	G	O4'-C1'-N9	8.62	115.09	108.20
84	Aa	685	G	N1-C6-O6	8.61	125.07	119.90
84	Aa	422	G	N1-C6-O6	8.61	125.07	119.90
84	Aa	469	U	O4'-C1'-N1	8.61	115.09	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	546	C	O4'-C1'-N1	8.61	115.09	108.20
84	Aa	3026	C	O4'-C1'-N1	8.61	115.09	108.20
84	Aa	914	C	O4'-C1'-N1	8.61	115.09	108.20
84	Aa	3349	C	O4'-C1'-N1	8.61	115.09	108.20
84	Aa	860	G	N1-C6-O6	8.61	125.07	119.90
84	Aa	3275	G	N1-C6-O6	8.61	125.07	119.90
84	Aa	625	G	C4'-C3'-O3'	8.61	130.22	113.00
84	Aa	2011	G	N1-C6-O6	8.61	125.06	119.90
84	Aa	2709	G	C5-C6-O6	-8.61	123.44	128.60
84	Aa	2811	C	O4'-C1'-N1	8.61	115.09	108.20
1	Ad	769	G	O4'-C1'-N9	8.61	115.08	108.20
1	Ad	1232	G	N9-C1'-C2'	8.61	125.19	114.00
68	Ch	74	TYR	CB-CG-CD1	-8.61	115.84	121.00
84	Aa	135	G	N1-C6-O6	8.61	125.06	119.90
84	Aa	680	G	N1-C6-O6	8.61	125.06	119.90
84	Aa	1361	G	C5-C6-O6	-8.61	123.44	128.60
84	Aa	1760	G	N1-C6-O6	8.61	125.06	119.90
84	Aa	1975	G	N1-C6-O6	8.61	125.06	119.90
84	Aa	2457	G	N1-C6-O6	8.61	125.06	119.90
84	Aa	3240	C	O4'-C1'-N1	8.61	115.08	108.20
84	Aa	1425	G	N1-C6-O6	8.60	125.06	119.90
84	Aa	1762	G	N1-C6-O6	8.60	125.06	119.90
86	Ab	5	G	O4'-C1'-N9	8.60	115.08	108.20
84	Aa	2459	U	P-O3'-C3'	8.60	130.02	119.70
84	Aa	2381	G	N1-C6-O6	8.60	125.06	119.90
84	Aa	2080	G	N1-C6-O6	8.60	125.06	119.90
1	Ad	1080	C	C1'-O4'-C4'	-8.60	103.02	109.90
84	Aa	1885	G	N1-C6-O6	8.60	125.06	119.90
85	Ac	115	C	O4'-C1'-N1	8.60	115.08	108.20
1	Ad	1545	A	C1'-O4'-C4'	8.59	116.78	109.90
84	Aa	1523	G	C5-C6-O6	-8.59	123.44	128.60
84	Aa	2543	G	N1-C6-O6	8.59	125.06	119.90
1	Ad	725	U	O4'-C1'-N1	8.59	115.07	108.20
3	Af	14	A	O4'-C1'-N9	8.59	115.07	108.20
84	Aa	543	C	C5'-C4'-O4'	8.59	119.41	109.10
84	Aa	831	G	N1-C6-O6	8.59	125.05	119.90
1	Ad	1163	C	O4'-C1'-C2'	-8.59	97.21	105.80
2	Ae	54	U	O4'-C1'-N1	8.59	115.07	108.20
84	Aa	975	G	N1-C6-O6	8.59	125.05	119.90
84	Aa	1371	G	N1-C6-O6	8.59	125.05	119.90
84	Aa	1696	G	O4'-C1'-N9	8.59	115.07	108.20
84	Aa	1717	G	C5-C6-O6	-8.59	123.45	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	366	G	N1-C6-O6	8.59	125.05	119.90
84	Aa	919	G	N1-C6-O6	8.59	125.05	119.90
84	Aa	2039	G	N1-C6-O6	8.59	125.05	119.90
84	Aa	2742	A	N1-C6-N6	8.59	123.75	118.60
84	Aa	600	G	N1-C6-O6	8.58	125.05	119.90
84	Aa	1510	G	N1-C6-O6	8.58	125.05	119.90
84	Aa	2118	G	N1-C6-O6	8.58	125.05	119.90
84	Aa	2985	C	O4'-C1'-N1	8.58	115.07	108.20
84	Aa	2066	G	N1-C6-O6	8.58	125.05	119.90
86	Ab	108	G	N3-C2-N2	8.58	125.91	119.90
84	Aa	3284	C	O4'-C1'-N1	8.58	115.06	108.20
1	Ad	1107	G	O4'-C1'-N9	8.58	115.06	108.20
84	Aa	396	G	N1-C6-O6	8.58	125.05	119.90
84	Aa	498	G	N1-C6-O6	8.58	125.05	119.90
84	Aa	1301	C	O4'-C1'-N1	8.58	115.06	108.20
84	Aa	2248	G	N1-C6-O6	8.58	125.05	119.90
84	Aa	1230	G	N1-C6-O6	8.57	125.05	119.90
1	Ad	434	G	O4'-C1'-N9	8.57	115.06	108.20
1	Ad	629	C	O4'-C1'-N1	8.57	115.06	108.20
84	Aa	1095	C	O4'-C1'-N1	8.57	115.06	108.20
84	Aa	2405	C	O4'-C1'-N1	8.57	115.06	108.20
84	Aa	1084	G	N1-C6-O6	8.57	125.04	119.90
85	Ac	142	G	N1-C6-O6	8.57	125.04	119.90
1	Ad	184	C	O4'-C1'-N1	8.57	115.06	108.20
84	Aa	2595	G	N1-C6-O6	8.57	125.04	119.90
84	Aa	3208	G	N1-C6-O6	8.57	125.04	119.90
1	Ad	966	U	N1-C1'-C2'	8.56	125.13	114.00
84	Aa	1736	C	O4'-C1'-N1	8.56	115.05	108.20
84	Aa	374	G	N1-C6-O6	8.56	125.04	119.90
84	Aa	1482	C	O4'-C1'-N1	8.56	115.05	108.20
84	Aa	1783	G	N1-C6-O6	8.56	125.04	119.90
84	Aa	2477	G	C4'-C3'-O3'	8.56	130.12	113.00
84	Aa	2914	G	N1-C6-O6	8.56	125.04	119.90
1	Ad	1466	A	N9-C1'-C2'	8.56	125.13	114.00
1	Ad	162	A	C1'-O4'-C4'	-8.56	103.05	109.90
84	Aa	85	G	N1-C6-O6	8.56	125.03	119.90
86	Ab	66	G	C5-C6-O6	-8.56	123.47	128.60
84	Aa	833	G	N1-C6-O6	8.55	125.03	119.90
84	Aa	1888	G	N1-C6-O6	8.56	125.03	119.90
84	Aa	2666	G	N1-C6-O6	8.55	125.03	119.90
84	Aa	5	G	N1-C6-O6	8.55	125.03	119.90
84	Aa	2608	G	N1-C6-O6	8.55	125.03	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	Ab	75	G	O4'-C1'-N9	8.55	115.04	108.20
84	Aa	2717	G	N1-C6-O6	8.55	125.03	119.90
84	Aa	3380	G	N3-C2-N2	8.55	125.89	119.90
84	Aa	2744	C	O4'-C1'-N1	8.55	115.04	108.20
84	Aa	1857	G	N1-C6-O6	8.55	125.03	119.90
84	Aa	2013	G	N1-C6-O6	8.55	125.03	119.90
84	Aa	2093	G	C5-C6-O6	-8.55	123.47	128.60
84	Aa	2109	G	N1-C6-O6	8.55	125.03	119.90
85	Ac	6	C	O4'-C1'-N1	8.55	115.04	108.20
1	Ad	290	C	C3'-C2'-C1'	8.55	108.34	101.50
1	Ad	1395	C	C1'-O4'-C4'	-8.55	103.06	109.90
84	Aa	880	C	O4'-C1'-N1	8.54	115.04	108.20
84	Aa	3277	C	O4'-C1'-N1	8.54	115.04	108.20
1	Ad	118	U	O4'-C1'-N1	8.54	115.03	108.20
84	Aa	3195	C	O4'-C1'-N1	8.54	115.03	108.20
84	Aa	1919	C	O4'-C1'-N1	8.54	115.03	108.20
1	Ad	218	G	C1'-O4'-C4'	-8.54	103.07	109.90
84	Aa	2315	G	N1-C6-O6	8.54	125.02	119.90
84	Aa	2920	G	N1-C6-O6	8.54	125.02	119.90
86	Ab	104	C	O4'-C1'-N1	8.54	115.03	108.20
84	Aa	1711	G	N1-C6-O6	8.54	125.02	119.90
84	Aa	2045	G	N1-C6-O6	8.54	125.02	119.90
85	Ac	133	C	O4'-C1'-N1	8.54	115.03	108.20
86	Ab	99	G	C5-C6-O6	-8.54	123.48	128.60
1	Ad	220	C	N1-C1'-C2'	8.54	125.10	114.00
1	Ad	1128	C	C3'-C2'-C1'	8.53	108.33	101.50
84	Aa	1420	G	N1-C6-O6	8.53	125.02	119.90
1	Ad	1425	G	O4'-C1'-N9	8.53	115.03	108.20
84	Aa	1067	G	N1-C6-O6	8.53	125.02	119.90
84	Aa	1726	G	N1-C6-O6	8.53	125.02	119.90
1	Ad	1065	A	O4'-C1'-C2'	8.53	115.28	107.60
85	Ac	3	A	N1-C6-N6	8.53	123.72	118.60
84	Aa	1335	C	O4'-C1'-N1	8.53	115.02	108.20
84	Aa	1966	C	O4'-C1'-N1	8.53	115.02	108.20
84	Aa	2229	G	N1-C6-O6	8.53	125.02	119.90
1	Ad	1706	G	O4'-C1'-N9	8.53	115.02	108.20
86	Ab	55	A	C4-C5-C6	8.53	121.26	117.00
1	Ad	266	C	O4'-C1'-N1	8.52	115.02	108.20
84	Aa	1751	G	N1-C6-O6	8.52	125.01	119.90
84	Aa	2417	G	N1-C6-O6	8.52	125.02	119.90
86	Ab	93	U	O4'-C1'-N1	8.52	115.02	108.20
84	Aa	160	G	N1-C6-O6	8.52	125.01	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1118	G	N1-C6-O6	8.52	125.01	119.90
84	Aa	2916	G	N1-C6-O6	8.52	125.01	119.90
84	Aa	2556	G	N1-C6-O6	8.52	125.01	119.90
1	Ad	1363	G	O4'-C1'-N9	8.52	115.01	108.20
84	Aa	179	G	N1-C6-O6	8.52	125.01	119.90
84	Aa	1771	G	N1-C6-O6	8.52	125.01	119.90
85	Ac	124	C	O4'-C1'-N1	8.52	115.02	108.20
84	Aa	557	C	O4'-C1'-N1	8.52	115.01	108.20
86	Ab	13	A	N1-C6-N6	8.52	123.71	118.60
1	Ad	386	C	O4'-C1'-N1	8.51	115.01	108.20
84	Aa	187	G	N1-C6-O6	8.51	125.01	119.90
84	Aa	876	C	O4'-C1'-N1	8.51	115.01	108.20
84	Aa	1968	C	O4'-C1'-N1	8.51	115.01	108.20
84	Aa	2029	G	N1-C6-O6	8.51	125.01	119.90
84	Aa	2038	G	N1-C6-O6	8.51	125.00	119.90
84	Aa	2102	C	O4'-C1'-N1	8.51	115.01	108.20
84	Aa	2207	C	O4'-C1'-N1	8.51	115.01	108.20
84	Aa	3252	G	N1-C6-O6	8.51	125.01	119.90
84	Aa	905	G	N1-C6-O6	8.51	125.00	119.90
84	Aa	345	G	N1-C6-O6	8.51	125.00	119.90
84	Aa	400	G	N1-C6-O6	8.51	125.00	119.90
84	Aa	1241	G	N1-C6-O6	8.51	125.00	119.90
84	Aa	1996	C	O4'-C1'-N1	8.51	115.01	108.20
84	Aa	2099	G	N1-C6-O6	8.51	125.00	119.90
84	Aa	756	C	O4'-C1'-N1	8.51	115.00	108.20
84	Aa	848	G	N1-C6-O6	8.51	125.00	119.90
84	Aa	1077	C	O4'-C1'-N1	8.51	115.00	108.20
84	Aa	1242	U	O4'-C1'-N1	8.51	115.00	108.20
84	Aa	1973	C	O4'-C1'-N1	8.51	115.00	108.20
84	Aa	3006	G	N1-C6-O6	8.51	125.00	119.90
1	Ad	1665	U	O4'-C1'-C2'	-8.50	97.30	105.80
84	Aa	1113	C	O4'-C1'-N1	8.50	115.00	108.20
84	Aa	1165	C	O4'-C1'-N1	8.50	115.00	108.20
84	Aa	1614	G	N1-C6-O6	8.50	125.00	119.90
85	Ac	47	U	O4'-C1'-N1	8.50	115.00	108.20
1	Ad	1131	G	O4'-C1'-N9	8.50	115.00	108.20
1	Ad	1729	A	O4'-C1'-N9	8.50	115.00	108.20
84	Aa	3089	G	N1-C6-O6	8.50	125.00	119.90
84	Aa	1491	G	N1-C6-O6	8.49	125.00	119.90
84	Aa	2402	G	N1-C6-O6	8.49	125.00	119.90
84	Aa	2563	G	N1-C6-O6	8.49	125.00	119.90
1	Ad	41	A	C1'-O4'-C4'	8.49	116.69	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	55	G	N1-C6-O6	8.49	125.00	119.90
84	Aa	805	C	O4'-C1'-N1	8.49	114.99	108.20
84	Aa	2689	U	O4'-C1'-N1	8.49	114.99	108.20
84	Aa	2754	G	N1-C6-O6	8.49	125.00	119.90
84	Aa	1608	C	O4'-C1'-N1	8.49	114.99	108.20
84	Aa	2526	G	N1-C6-O6	8.49	124.99	119.90
84	Aa	778	G	N1-C6-O6	8.49	124.99	119.90
84	Aa	109	G	N1-C6-O6	8.48	124.99	119.90
84	Aa	689	G	N1-C6-O6	8.48	124.99	119.90
84	Aa	2306	G	C5-C6-O6	-8.48	123.51	128.60
86	Ab	1	G	O4'-C1'-N9	8.48	114.99	108.20
1	Ad	599	G	O4'-C1'-N9	8.48	114.99	108.20
84	Aa	940	G	N1-C6-O6	8.48	124.99	119.90
84	Aa	2364	C	O4'-C1'-N1	8.48	114.99	108.20
84	Aa	1022	G	O4'-C1'-N9	8.48	114.98	108.20
84	Aa	2322	G	N1-C6-O6	8.48	124.99	119.90
84	Aa	2654	G	N1-C6-O6	8.48	124.99	119.90
1	Ad	968	A	C1'-O4'-C4'	8.48	116.68	109.90
84	Aa	619	C	O4'-C1'-N1	8.48	114.98	108.20
84	Aa	1666	C	O4'-C1'-N1	8.48	114.98	108.20
84	Aa	1895	G	N1-C6-O6	8.48	124.99	119.90
84	Aa	1210	G	N1-C6-O6	8.48	124.99	119.90
84	Aa	2714	U	O4'-C1'-N1	8.48	114.98	108.20
85	Ac	141	C	O4'-C1'-N1	8.48	114.98	108.20
84	Aa	446	C	O4'-C1'-N1	8.47	114.98	108.20
84	Aa	2025	C	O4'-C1'-N1	8.47	114.98	108.20
86	Ab	101	A	C4-C5-C6	8.47	121.24	117.00
1	Ad	782	G	C1'-O4'-C4'	-8.47	103.12	109.90
84	Aa	1186	C	O4'-C1'-N1	8.47	114.98	108.20
84	Aa	3234	G	N1-C6-O6	8.47	124.98	119.90
84	Aa	611	C	O4'-C1'-N1	8.47	114.98	108.20
84	Aa	2841	G	N1-C6-O6	8.47	124.98	119.90
85	Ac	16	G	N1-C6-O6	8.47	124.98	119.90
84	Aa	1977	C	O4'-C1'-N1	8.47	114.97	108.20
84	Aa	2486	G	N1-C6-O6	8.47	124.98	119.90
1	Ad	902	C	O4'-C1'-C2'	-8.47	97.33	105.80
84	Aa	230	G	N1-C6-O6	8.47	124.98	119.90
84	Aa	1997	G	N1-C6-O6	8.47	124.98	119.90
84	Aa	3109	G	N1-C6-O6	8.47	124.98	119.90
1	Ad	1380	A	O4'-C1'-N9	8.46	114.97	108.20
84	Aa	544	C	O4'-C1'-N1	8.46	114.97	108.20
84	Aa	829	G	N1-C6-O6	8.46	124.98	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1519	C	O4'-C1'-N1	8.47	114.97	108.20
1	Ad	1724	U	O4'-C1'-N1	8.46	114.97	108.20
84	Aa	256	G	N1-C6-O6	8.46	124.98	119.90
84	Aa	579	G	C5-C6-O6	-8.46	123.52	128.60
84	Aa	3090	C	O4'-C1'-N1	8.46	114.97	108.20
1	Ad	15	U	O4'-C1'-N1	8.46	114.97	108.20
84	Aa	195	G	N1-C6-O6	8.46	124.97	119.90
84	Aa	1665	G	N1-C6-O6	8.46	124.97	119.90
1	Ad	1646	G	C1'-O4'-C4'	-8.46	103.14	109.90
84	Aa	2642	G	N1-C6-O6	8.46	124.97	119.90
84	Aa	1318	C	O4'-C1'-N1	8.45	114.96	108.20
85	Ac	150	G	O4'-C1'-N9	8.46	114.96	108.20
1	Ad	839	G	P-O5'-C5'	8.45	134.42	120.90
1	Ad	1717	C	O4'-C1'-N1	8.45	114.96	108.20
84	Aa	1408	C	O4'-C1'-N1	8.45	114.96	108.20
84	Aa	20	G	N1-C6-O6	8.45	124.97	119.90
84	Aa	1686	U	O4'-C1'-N1	8.45	114.96	108.20
84	Aa	2622	G	N1-C6-O6	8.45	124.97	119.90
84	Aa	3100	C	P-O3'-C3'	8.45	129.84	119.70
84	Aa	1606	C	O4'-C1'-N1	8.45	114.96	108.20
1	Ad	328	U	O4'-C1'-N1	8.45	114.96	108.20
84	Aa	144	A	C5-C6-N6	-8.45	116.94	123.70
84	Aa	221	C	O4'-C1'-N1	8.45	114.96	108.20
84	Aa	707	G	N1-C6-O6	8.45	124.97	119.90
84	Aa	749	C	O4'-C1'-N1	8.45	114.96	108.20
84	Aa	882	U	O4'-C1'-N1	8.45	114.96	108.20
84	Aa	2108	C	O4'-C1'-N1	8.45	114.96	108.20
84	Aa	2173	G	O4'-C1'-N9	8.45	114.96	108.20
1	Ad	1155	G	C1'-O4'-C4'	-8.44	103.14	109.90
84	Aa	3198	C	O4'-C1'-N1	8.44	114.95	108.20
84	Aa	2033	C	O4'-C1'-N1	8.44	114.95	108.20
84	Aa	2086	A	C4'-C3'-O3'	-8.44	91.68	109.40
84	Aa	612	U	P-O3'-C3'	8.44	129.82	119.70
84	Aa	1281	C	O4'-C1'-N1	8.44	114.95	108.20
1	Ad	1492	G	N9-C1'-C2'	-8.44	102.72	112.00
84	Aa	2821	U	O4'-C1'-N1	8.43	114.95	108.20
84	Aa	471	C	O4'-C1'-N1	8.43	114.94	108.20
84	Aa	275	G	C5-C6-O6	-8.43	123.54	128.60
84	Aa	2535	C	O4'-C1'-N1	8.43	114.94	108.20
84	Aa	639	A	P-O3'-C3'	8.43	129.81	119.70
84	Aa	2555	G	N1-C6-O6	8.43	124.96	119.90
84	Aa	3053	G	N1-C6-O6	8.42	124.95	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1634	G	N1-C6-O6	8.42	124.95	119.90
84	Aa	2144	G	N1-C6-O6	8.42	124.95	119.90
85	Ac	144	C	O4'-C1'-N1	8.42	114.94	108.20
84	Aa	2940	G	N1-C6-O6	8.42	124.95	119.90
84	Aa	102	G	N1-C6-O6	8.42	124.95	119.90
84	Aa	1444	G	N1-C6-O6	8.42	124.95	119.90
1	Ad	296	A	N9-C1'-C2'	8.41	124.93	114.00
1	Ad	513	G	O4'-C1'-N9	8.41	114.93	108.20
1	Ad	733	U	O4'-C1'-N1	8.41	114.93	108.20
84	Aa	24	C	O4'-C1'-N1	8.41	114.93	108.20
1	Ad	1744	C	O4'-C1'-N1	8.40	114.92	108.20
1	Ad	88	C	O4'-C1'-N1	8.40	114.92	108.20
1	Ad	919	G	N9-C1'-C2'	-8.40	102.76	112.00
1	Ad	1103	U	O4'-C1'-N1	8.40	114.92	108.20
84	Aa	244	G	O4'-C1'-N9	8.40	114.92	108.20
84	Aa	1984	C	O4'-C1'-N1	8.40	114.92	108.20
1	Ad	1604	C	C3'-C2'-C1'	-8.40	94.78	101.50
84	Aa	52	G	N1-C6-O6	8.40	124.94	119.90
84	Aa	103	G	N1-C6-O6	8.40	124.94	119.90
84	Aa	1872	C	O4'-C1'-N1	8.40	114.92	108.20
84	Aa	1913	C	O4'-C1'-N1	8.40	114.92	108.20
1	Ad	407	G	O4'-C1'-N9	8.40	114.92	108.20
85	Ac	152	G	N1-C6-O6	8.40	124.94	119.90
1	Ad	224	C	O4'-C1'-N1	8.39	114.92	108.20
1	Ad	1593	U	C1'-O4'-C4'	-8.39	103.18	109.90
84	Aa	1150	G	N1-C6-O6	8.39	124.94	119.90
84	Aa	2299	C	O4'-C1'-N1	8.39	114.92	108.20
84	Aa	3350	C	O4'-C1'-N1	8.39	114.91	108.20
84	Aa	290	C	O4'-C1'-N1	8.39	114.91	108.20
1	Ad	1665	U	C1'-O4'-C4'	8.39	116.61	109.90
84	Aa	3203	G	N1-C6-O6	8.39	124.94	119.90
84	Aa	1415	G	C5-C6-O6	-8.39	123.57	128.60
84	Aa	1596	G	N1-C6-O6	8.39	124.93	119.90
1	Ad	164	C	N1-C1'-C2'	8.38	124.90	114.00
84	Aa	1562	A	C5'-C4'-C3'	8.38	129.42	116.00
84	Aa	2464	G	N1-C6-O6	8.38	124.93	119.90
1	Ad	1725	C	O4'-C1'-C2'	-8.38	97.42	105.80
84	Aa	1685	U	O4'-C1'-N1	8.38	114.91	108.20
84	Aa	3182	A	N1-C6-N6	8.38	123.63	118.60
84	Aa	1782	G	N1-C6-O6	8.38	124.93	119.90
84	Aa	3134	C	O4'-C1'-N1	8.38	114.90	108.20
1	Ad	571	A	O4'-C1'-N9	8.38	114.90	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2592	G	N1-C6-O6	8.38	124.93	119.90
1	Ad	1752	U	O4'-C1'-N1	8.38	114.90	108.20
84	Aa	1841	G	N1-C6-O6	8.38	124.93	119.90
86	Ab	23	A	C4-C5-C6	8.38	121.19	117.00
1	Ad	1466	A	C1'-O4'-C4'	-8.38	103.20	109.90
84	Aa	361	G	N1-C6-O6	8.38	124.93	119.90
84	Aa	305	G	N1-C6-O6	8.37	124.92	119.90
84	Aa	437	C	O4'-C1'-N1	8.37	114.90	108.20
86	Ab	51	G	C5-C6-O6	-8.38	123.58	128.60
1	Ad	5	U	O4'-C1'-N1	8.37	114.90	108.20
84	Aa	1422	G	N1-C6-O6	8.37	124.92	119.90
1	Ad	409	C	N1-C1'-C2'	8.37	124.88	114.00
84	Aa	3167	G	N1-C6-O6	8.37	124.92	119.90
86	Ab	118	C	N3-C4-N4	8.37	123.86	118.00
1	Ad	291	G	O4'-C1'-N9	8.37	114.89	108.20
84	Aa	80	C	O4'-C1'-N1	8.37	114.89	108.20
84	Aa	307	C	O4'-C1'-N1	8.37	114.89	108.20
84	Aa	2032	C	O4'-C1'-N1	8.37	114.89	108.20
84	Aa	2040	G	N1-C6-O6	8.37	124.92	119.90
1	Ad	67	G	C1'-O4'-C4'	-8.37	103.21	109.90
1	Ad	212	A	N9-C1'-C2'	8.37	124.87	114.00
1	Ad	1235	U	O4'-C1'-N1	8.37	114.89	108.20
84	Aa	162	G	N1-C6-O6	8.37	124.92	119.90
84	Aa	1042	C	O4'-C1'-N1	8.37	114.89	108.20
84	Aa	1573	G	N1-C6-O6	8.37	124.92	119.90
84	Aa	2246	G	C5-C6-O6	-8.37	123.58	128.60
84	Aa	2911	C	O4'-C1'-N1	8.37	114.89	108.20
84	Aa	2154	G	N1-C6-O6	8.36	124.92	119.90
84	Aa	3199	C	O4'-C1'-N1	8.37	114.89	108.20
1	Ad	249	G	N9-C1'-C2'	8.36	124.87	114.00
1	Ad	1622	A	C3'-C2'-C1'	-8.36	94.81	101.50
84	Aa	3054	G	N1-C6-O6	8.36	124.92	119.90
84	Aa	3063	C	O4'-C1'-N1	8.36	114.89	108.20
84	Aa	487	C	O4'-C1'-N1	8.36	114.89	108.20
84	Aa	575	C	C4'-C3'-O3'	-8.36	91.85	109.40
84	Aa	2628	C	O4'-C1'-N1	8.36	114.89	108.20
84	Aa	1323	G	N1-C6-O6	8.36	124.92	119.90
84	Aa	1632	G	N1-C6-O6	8.36	124.92	119.90
84	Aa	209	G	N1-C6-O6	8.36	124.91	119.90
84	Aa	433	C	O4'-C1'-N1	8.36	114.88	108.20
84	Aa	898	G	N1-C6-O6	8.35	124.91	119.90
1	Ad	570	C	N1-C1'-C2'	8.35	124.86	114.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	343	G	N1-C6-O6	8.35	124.91	119.90
84	Aa	3046	C	O4'-C1'-N1	8.35	114.88	108.20
84	Aa	1241	G	O4'-C1'-N9	8.35	114.88	108.20
84	Aa	2000	C	O4'-C1'-N1	8.35	114.88	108.20
84	Aa	470	G	N1-C6-O6	8.35	124.91	119.90
84	Aa	3317	G	C5-C6-O6	-8.35	123.59	128.60
84	Aa	3323	U	O4'-C1'-N1	8.35	114.88	108.20
1	Ad	1394	A	P-O3'-C3'	8.35	129.72	119.70
84	Aa	1628	G	N1-C6-O6	8.35	124.91	119.90
84	Aa	2597	C	O4'-C1'-N1	8.34	114.88	108.20
84	Aa	64	A	C5-C6-N6	-8.34	117.03	123.70
84	Aa	95	G	N1-C6-O6	8.34	124.91	119.90
84	Aa	2953	G	N1-C6-O6	8.34	124.91	119.90
84	Aa	516	C	O4'-C1'-N1	8.34	114.87	108.20
84	Aa	3068	U	O4'-C1'-N1	8.34	114.87	108.20
84	Aa	690	G	N1-C6-O6	8.34	124.90	119.90
84	Aa	3163	G	N1-C6-O6	8.34	124.90	119.90
1	Ad	1510	G	O4'-C1'-N9	8.34	114.87	108.20
2	Ae	31	C	C3'-C2'-C1'	8.34	108.17	101.50
84	Aa	835	G	N1-C6-O6	8.34	124.90	119.90
84	Aa	2511	U	C4'-C3'-O3'	-8.34	91.90	109.40
1	Ad	1321	C	N1-C1'-C2'	8.33	124.83	114.00
84	Aa	225	G	N1-C6-O6	8.33	124.90	119.90
84	Aa	2065	G	N1-C6-O6	8.33	124.90	119.90
85	Ac	118	C	O4'-C1'-N1	8.33	114.87	108.20
86	Ab	67	C	O4'-C1'-N1	8.33	114.87	108.20
84	Aa	1424	G	N1-C6-O6	8.33	124.90	119.90
84	Aa	165	C	O4'-C1'-N1	8.33	114.86	108.20
1	Ad	988	G	O4'-C1'-N9	8.33	114.86	108.20
84	Aa	902	U	O4'-C1'-N1	8.33	114.86	108.20
84	Aa	2163	G	O4'-C1'-N9	8.33	114.86	108.20
86	Ab	34	C	O4'-C1'-N1	8.33	114.86	108.20
84	Aa	704	G	C5-C6-O6	-8.32	123.61	128.60
84	Aa	872	G	N1-C6-O6	8.32	124.89	119.90
84	Aa	2097	C	O4'-C1'-N1	8.32	114.86	108.20
84	Aa	2786	G	N1-C6-O6	8.32	124.89	119.90
85	Ac	154	G	N1-C6-O6	8.32	124.89	119.90
84	Aa	3205	C	O4'-C1'-N1	8.32	114.86	108.20
1	Ad	1784	G	O4'-C1'-N9	8.32	114.86	108.20
84	Aa	3226	G	P-O3'-C3'	8.32	129.69	119.70
1	Ad	358	C	C3'-C2'-C1'	8.32	108.16	101.50
1	Ad	834	A	C1'-O4'-C4'	8.32	116.56	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	443	G	N1-C6-O6	8.32	124.89	119.90
84	Aa	448	G	N1-C6-O6	8.32	124.89	119.90
84	Aa	1029	C	O4'-C1'-N1	8.32	114.86	108.20
84	Aa	1571	A	O4'-C1'-N9	8.32	114.86	108.20
84	Aa	2621	G	N1-C6-O6	8.32	124.89	119.90
84	Aa	3211	C	O4'-C1'-N1	8.32	114.86	108.20
1	Ad	597	U	O4'-C1'-N1	8.32	114.85	108.20
84	Aa	1014	G	N1-C6-O6	8.32	124.89	119.90
84	Aa	1418	C	O4'-C1'-N1	8.32	114.86	108.20
84	Aa	3179	G	C5-C6-O6	-8.32	123.61	128.60
84	Aa	2426	C	O4'-C1'-N1	8.32	114.85	108.20
84	Aa	3060	G	N1-C6-O6	8.32	124.89	119.90
85	Ac	124	C	P-O3'-C3'	8.32	129.68	119.70
84	Aa	1016	G	N1-C6-O6	8.31	124.89	119.90
84	Aa	1741	G	C5-C6-O6	-8.31	123.61	128.60
84	Aa	2990	C	O4'-C1'-N1	8.31	114.85	108.20
84	Aa	3339	G	N1-C6-O6	8.31	124.89	119.90
86	Ab	31	G	C6-C5-N7	-8.31	125.41	130.40
1	Ad	962	G	O4'-C1'-N9	8.31	114.85	108.20
1	Ad	1599	C	C3'-C2'-C1'	8.31	108.15	101.50
84	Aa	1151	G	N1-C6-O6	8.31	124.89	119.90
84	Aa	2297	G	N1-C6-O6	8.31	124.89	119.90
84	Aa	827	C	O4'-C1'-N1	8.31	114.85	108.20
84	Aa	3126	U	O4'-C1'-N1	8.31	114.85	108.20
84	Aa	733	C	O4'-C1'-N1	8.31	114.85	108.20
84	Aa	1053	C	O4'-C1'-N1	8.31	114.85	108.20
84	Aa	2124	G	N1-C6-O6	8.31	124.89	119.90
84	Aa	478	G	N1-C6-O6	8.31	124.88	119.90
84	Aa	1989	G	N1-C6-O6	8.31	124.88	119.90
84	Aa	2460	A	N1-C6-N6	8.31	123.58	118.60
84	Aa	2511	U	C2'-C3'-O3'	8.31	127.78	109.50
84	Aa	3176	C	O4'-C1'-N1	8.31	114.85	108.20
84	Aa	3276	G	N1-C6-O6	8.31	124.88	119.90
84	Aa	761	C	O4'-C1'-N1	8.31	114.84	108.20
85	Ac	151	G	C5-C6-O6	-8.31	123.62	128.60
1	Ad	1608	A	N9-C1'-C2'	8.30	124.80	114.00
84	Aa	10	C	O4'-C1'-N1	8.30	114.84	108.20
84	Aa	832	C	O4'-C1'-N1	8.30	114.84	108.20
84	Aa	1742	G	N1-C6-O6	8.30	124.88	119.90
84	Aa	1864	G	N1-C6-O6	8.30	124.88	119.90
1	Ad	1045	G	O4'-C1'-N9	8.30	114.84	108.20
84	Aa	2049	C	O4'-C1'-N1	8.30	114.84	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2268	G	N1-C6-O6	8.30	124.88	119.90
84	Aa	2623	G	C5-C6-O6	-8.30	123.62	128.60
84	Aa	2378	U	O4'-C1'-N1	8.30	114.84	108.20
85	Ac	30	C	O4'-C1'-N1	8.30	114.84	108.20
84	Aa	40	G	N1-C6-O6	8.30	124.88	119.90
84	Aa	2328	C	O4'-C1'-N1	8.30	114.84	108.20
84	Aa	2940	G	O4'-C1'-N9	8.30	114.84	108.20
84	Aa	562	G	N1-C6-O6	8.29	124.88	119.90
84	Aa	585	A	O4'-C1'-N9	8.29	114.84	108.20
84	Aa	1612	C	O4'-C1'-N1	8.29	114.83	108.20
84	Aa	1824	C	O4'-C1'-N1	8.29	114.83	108.20
84	Aa	2474	A	O4'-C1'-N9	8.29	114.83	108.20
84	Aa	2588	G	C5-C6-O6	-8.29	123.63	128.60
1	Ad	152	G	O4'-C1'-N9	8.29	114.83	108.20
84	Aa	369	G	N1-C6-O6	8.29	124.87	119.90
84	Aa	751	C	O4'-C1'-N1	8.29	114.83	108.20
84	Aa	1994	C	O4'-C1'-N1	8.29	114.83	108.20
84	Aa	2546	C	O4'-C1'-N1	8.29	114.83	108.20
85	Ac	71	A	C5-C6-N6	-8.29	117.07	123.70
85	Ac	91	C	O4'-C1'-N1	8.29	114.83	108.20
84	Aa	14	U	O4'-C1'-N1	8.29	114.83	108.20
84	Aa	1552	C	O4'-C1'-N1	8.28	114.83	108.20
84	Aa	2864	U	O4'-C1'-N1	8.29	114.83	108.20
86	Ab	94	C	O4'-C1'-N1	8.29	114.83	108.20
84	Aa	3303	C	O4'-C1'-N1	8.28	114.83	108.20
85	Ac	138	G	N1-C6-O6	8.28	124.87	119.90
1	Ad	817	C	O4'-C1'-C2'	-8.28	97.52	105.80
1	Ad	1544	G	O4'-C1'-N9	8.28	114.83	108.20
84	Aa	2847	A	N1-C6-N6	8.28	123.57	118.60
84	Aa	3061	C	O4'-C1'-N1	8.28	114.83	108.20
84	Aa	3342	C	O4'-C1'-N1	8.28	114.83	108.20
85	Ac	126	A	O4'-C1'-N9	8.28	114.83	108.20
1	Ad	37	U	O4'-C1'-N1	8.28	114.82	108.20
84	Aa	633	C	O4'-C1'-N1	8.28	114.82	108.20
84	Aa	2196	G	C5-C6-O6	-8.28	123.63	128.60
85	Ac	153	C	O4'-C1'-N1	8.28	114.82	108.20
84	Aa	515	C	O4'-C1'-N1	8.28	114.82	108.20
84	Aa	1081	U	O4'-C1'-N1	8.28	114.82	108.20
84	Aa	2076	C	P-O3'-C3'	8.28	129.63	119.70
84	Aa	2927	C	O4'-C1'-N1	8.28	114.82	108.20
84	Aa	2985	C	C2-N1-C1'	8.28	127.90	118.80
1	Ad	1445	C	C3'-C2'-C1'	8.27	108.12	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1757	G	N1-C6-O6	8.27	124.86	119.90
84	Aa	993	A	O4'-C1'-N9	8.27	114.82	108.20
84	Aa	1759	C	C2-N1-C1'	8.27	127.90	118.80
1	Ad	1758	G	C1'-O4'-C4'	-8.27	103.29	109.90
84	Aa	2059	C	O4'-C1'-N1	8.27	114.81	108.20
84	Aa	2501	U	O4'-C1'-N1	8.27	114.82	108.20
85	Ac	134	G	C5-C6-O6	-8.27	123.64	128.60
84	Aa	1331	C	O4'-C1'-N1	8.27	114.81	108.20
84	Aa	1732	G	N1-C6-O6	8.27	124.86	119.90
85	Ac	50	C	O4'-C1'-N1	8.27	114.81	108.20
1	Ad	450	A	O4'-C1'-C2'	-8.26	97.54	105.80
84	Aa	331	G	N1-C6-O6	8.26	124.86	119.90
84	Aa	727	G	N1-C6-O6	8.26	124.86	119.90
84	Aa	2046	G	N1-C6-O6	8.26	124.86	119.90
1	Ad	1032	A	N9-C1'-C2'	-8.26	102.91	112.00
84	Aa	1434	G	N1-C6-O6	8.26	124.86	119.90
84	Aa	1495	G	N1-C6-O6	8.26	124.86	119.90
84	Aa	1706	C	O4'-C1'-N1	8.26	114.81	108.20
84	Aa	800	C	O4'-C1'-N1	8.26	114.81	108.20
84	Aa	3044	C	O4'-C1'-N1	8.26	114.81	108.20
1	Ad	509	A	O4'-C1'-C2'	-8.26	97.55	105.80
84	Aa	3187	C	O4'-C1'-N1	8.26	114.80	108.20
84	Aa	259	G	N1-C6-O6	8.25	124.85	119.90
84	Aa	1102	A	N1-C6-N6	8.25	123.55	118.60
84	Aa	592	U	O4'-C1'-N1	8.25	114.80	108.20
84	Aa	1245	U	O4'-C1'-N1	8.25	114.80	108.20
84	Aa	2010	G	N1-C6-O6	8.25	124.85	119.90
84	Aa	155	G	N1-C6-O6	8.25	124.85	119.90
84	Aa	354	C	O4'-C1'-N1	8.25	114.80	108.20
84	Aa	1350	G	N1-C6-O6	8.25	124.85	119.90
84	Aa	2814	C	O4'-C1'-N1	8.25	114.80	108.20
84	Aa	101	C	O4'-C1'-N1	8.25	114.80	108.20
84	Aa	147	G	N1-C6-O6	8.25	124.85	119.90
84	Aa	1449	A	O4'-C1'-N9	8.25	114.80	108.20
1	Ad	25	C	C3'-C2'-C1'	8.25	108.10	101.50
84	Aa	1656	C	O4'-C1'-N1	8.25	114.80	108.20
84	Aa	1930	G	N1-C6-O6	8.25	124.85	119.90
85	Ac	51	G	N1-C6-O6	8.25	124.85	119.90
84	Aa	1185	G	N1-C6-O6	8.24	124.85	119.90
84	Aa	1667	C	O4'-C1'-N1	8.24	114.79	108.20
84	Aa	3249	G	N1-C6-O6	8.24	124.84	119.90
1	Ad	615	U	O4'-C1'-N1	8.24	114.79	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1141	U	O4'-C1'-N1	8.24	114.79	108.20
1	Ad	1345	G	C1'-O4'-C4'	-8.24	103.31	109.90
84	Aa	1642	G	N1-C6-O6	8.24	124.84	119.90
84	Aa	1609	G	O4'-C1'-N9	8.23	114.79	108.20
84	Aa	1854	A	N1-C6-N6	8.23	123.54	118.60
84	Aa	9	C	O4'-C1'-N1	8.23	114.78	108.20
84	Aa	2028	C	O4'-C1'-N1	8.23	114.78	108.20
84	Aa	2393	G	N1-C6-O6	8.23	124.84	119.90
1	Ad	748	C	C3'-C2'-C1'	8.23	108.08	101.50
85	Ac	2	G	N1-C6-O6	8.23	124.84	119.90
84	Aa	1947	U	O4'-C1'-N1	8.22	114.78	108.20
85	Ac	103	G	N1-C6-O6	8.22	124.83	119.90
85	Ac	49	G	N1-C6-O6	8.22	124.83	119.90
85	Ac	58	G	N1-C6-O6	8.22	124.83	119.90
1	Ad	416	A	O4'-C1'-N9	8.22	114.78	108.20
84	Aa	737	C	O4'-C1'-N1	8.22	114.77	108.20
84	Aa	1856	G	C5-C6-O6	-8.22	123.67	128.60
84	Aa	2093	G	O4'-C1'-N9	8.22	114.77	108.20
84	Aa	2568	G	C5-C6-O6	-8.22	123.67	128.60
1	Ad	1664	U	P-O3'-C3'	8.22	129.56	119.70
84	Aa	505	G	N1-C6-O6	8.21	124.83	119.90
84	Aa	960	C	O4'-C1'-N1	8.21	114.77	108.20
84	Aa	1414	C	O4'-C1'-N1	8.21	114.77	108.20
1	Ad	6	G	O4'-C1'-N9	8.21	114.77	108.20
1	Ad	84	G	O4'-C1'-N9	8.21	114.77	108.20
1	Ad	277	G	C3'-C2'-C1'	8.21	108.07	101.50
1	Ad	1301	G	N9-C1'-C2'	-8.21	102.97	112.00
84	Aa	398	G	C5-C6-O6	-8.21	123.67	128.60
84	Aa	655	G	N1-C6-O6	8.21	124.83	119.90
1	Ad	1774	C	O4'-C1'-N1	8.21	114.77	108.20
84	Aa	201	G	N1-C6-O6	8.21	124.83	119.90
84	Aa	1636	C	O4'-C1'-N1	8.21	114.77	108.20
84	Aa	1659	G	N1-C6-O6	8.21	124.83	119.90
84	Aa	2103	U	O4'-C1'-N1	8.21	114.77	108.20
84	Aa	3210	G	O4'-C1'-N9	8.21	114.77	108.20
84	Aa	3319	G	N1-C6-O6	8.21	124.83	119.90
84	Aa	558	G	N1-C6-O6	8.21	124.83	119.90
85	Ac	45	C	O4'-C1'-N1	8.21	114.77	108.20
1	Ad	373	U	O4'-C1'-N1	8.21	114.76	108.20
1	Ad	1069	G	O4'-C1'-N9	8.21	114.76	108.20
84	Aa	549	G	C5-C6-O6	-8.21	123.68	128.60
84	Aa	569	C	O4'-C1'-N1	8.21	114.77	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2908	C	O4'-C1'-N1	8.21	114.77	108.20
84	Aa	1112	C	O4'-C1'-N1	8.20	114.76	108.20
86	Ab	104	C	N3-C4-N4	8.20	123.74	118.00
18	BN	128	TYR	CB-CG-CD1	-8.20	116.08	121.00
84	Aa	1784	C	O4'-C1'-N1	8.20	114.76	108.20
3	Af	13	A	O4'-C1'-N9	8.20	114.76	108.20
84	Aa	218	G	N1-C6-O6	8.20	124.82	119.90
84	Aa	1075	G	N1-C6-O6	8.20	124.82	119.90
84	Aa	1660	C	O4'-C1'-N1	8.20	114.76	108.20
84	Aa	1963	G	N1-C6-O6	8.20	124.82	119.90
1	Ad	244	C	O4'-C1'-N1	8.20	114.76	108.20
1	Ad	492	G	P-O5'-C5'	8.20	134.01	120.90
84	Aa	91	G	N1-C6-O6	8.20	124.82	119.90
84	Aa	2041	G	N1-C6-O6	8.20	124.82	119.90
84	Aa	2450	G	N1-C6-O6	8.19	124.82	119.90
85	Ac	65	G	C5-C6-O6	-8.19	123.68	128.60
1	Ad	366	G	O4'-C1'-N9	8.19	114.75	108.20
84	Aa	164	C	O4'-C1'-N1	8.19	114.75	108.20
84	Aa	532	G	N1-C6-O6	8.19	124.81	119.90
84	Aa	1369	G	N1-C6-O6	8.19	124.81	119.90
84	Aa	3387	U	O4'-C1'-N1	8.19	114.75	108.20
86	Ab	2	G	N1-C6-O6	8.19	124.82	119.90
86	Ab	7	G	N1-C6-O6	8.19	124.82	119.90
84	Aa	1378	G	N1-C6-O6	8.19	124.81	119.90
1	Ad	1731	A	C3'-C2'-C1'	8.19	108.05	101.50
84	Aa	267	G	N1-C6-O6	8.19	124.81	119.90
84	Aa	2857	U	O4'-C1'-N1	8.19	114.75	108.20
2	Ae	57	A	O4'-C1'-N9	8.19	114.75	108.20
84	Aa	523	C	O4'-C1'-N1	8.19	114.75	108.20
84	Aa	2342	C	O4'-C1'-N1	8.18	114.75	108.20
84	Aa	3324	U	O4'-C1'-N1	8.18	114.75	108.20
1	Ad	324	U	O4'-C1'-N1	8.18	114.74	108.20
84	Aa	536	C	O4'-C1'-N1	8.18	114.74	108.20
84	Aa	641	C	C5'-C4'-C3'	8.18	129.09	116.00
84	Aa	744	C	O4'-C1'-N1	8.18	114.75	108.20
84	Aa	2816	G	N1-C6-O6	8.18	124.81	119.90
84	Aa	1735	U	O4'-C1'-N1	8.18	114.74	108.20
84	Aa	1972	C	O4'-C1'-N1	8.18	114.74	108.20
84	Aa	3207	C	O4'-C1'-N1	8.18	114.74	108.20
84	Aa	3239	G	N1-C6-O6	8.18	124.81	119.90
86	Ab	8	A	C4-C5-C6	8.18	121.09	117.00
86	Ab	75	G	N1-C6-O6	8.18	124.81	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	75	U	O4'-C1'-N1	8.18	114.74	108.20
84	Aa	1346	C	O4'-C1'-N1	8.17	114.74	108.20
84	Aa	1566	C	O4'-C1'-N1	8.17	114.74	108.20
84	Aa	2019	G	N1-C6-O6	8.17	124.80	119.90
84	Aa	2388	C	O4'-C1'-N1	8.17	114.74	108.20
84	Aa	3118	C	O4'-C1'-N1	8.17	114.74	108.20
84	Aa	2183	A	C5-C6-N1	-8.17	113.61	117.70
84	Aa	2187	C	O4'-C1'-N1	8.17	114.74	108.20
1	Ad	1127	G	N9-C1'-C2'	-8.17	103.01	112.00
84	Aa	1889	G	C5-C6-O6	-8.17	123.70	128.60
1	Ad	532	U	O4'-C1'-N1	8.17	114.73	108.20
84	Aa	2891	C	O4'-C1'-N1	8.17	114.73	108.20
2	Ae	3	C	O4'-C1'-N1	8.17	114.73	108.20
84	Aa	946	U	O4'-C1'-N1	8.17	114.73	108.20
84	Aa	1965	C	O4'-C1'-N1	8.17	114.73	108.20
84	Aa	2222	C	O4'-C1'-N1	8.17	114.73	108.20
86	Ab	67	C	C5-C4-N4	-8.17	114.48	120.20
84	Aa	1284	C	O4'-C1'-N1	8.16	114.73	108.20
84	Aa	2332	C	O4'-C1'-N1	8.16	114.73	108.20
84	Aa	98	A	O4'-C1'-N9	8.16	114.73	108.20
84	Aa	140	C	O4'-C1'-N1	8.16	114.73	108.20
84	Aa	684	C	O4'-C1'-N1	8.16	114.73	108.20
84	Aa	1038	C	O4'-C1'-N1	8.16	114.73	108.20
84	Aa	642	C	C5'-C4'-C3'	8.16	129.05	116.00
84	Aa	1156	A	N1-C6-N6	8.16	123.50	118.60
84	Aa	2003	C	O4'-C1'-N1	8.16	114.73	108.20
84	Aa	588	G	N1-C6-O6	8.16	124.79	119.90
84	Aa	1690	C	O4'-C1'-N1	8.16	114.73	108.20
84	Aa	2869	C	O4'-C1'-N1	8.16	114.72	108.20
84	Aa	3263	C	P-O5'-C5'	8.16	133.95	120.90
1	Ad	73	A	O4'-C1'-N9	8.15	114.72	108.20
84	Aa	1096	C	O4'-C1'-N1	8.15	114.72	108.20
84	Aa	1259	C	O4'-C1'-N1	8.15	114.72	108.20
84	Aa	2148	U	O4'-C1'-N1	8.15	114.72	108.20
84	Aa	2672	C	O4'-C1'-N1	8.15	114.72	108.20
1	Ad	1072	U	O4'-C1'-N1	8.15	114.72	108.20
84	Aa	582	C	O4'-C1'-N1	8.15	114.72	108.20
84	Aa	1959	U	P-O5'-C5'	-8.15	107.86	120.90
84	Aa	2569	G	N1-C6-O6	8.15	124.79	119.90
1	Ad	617	G	N9-C1'-C2'	8.15	124.59	114.00
84	Aa	1174	G	N1-C6-O6	8.15	124.79	119.90
84	Aa	1848	G	N1-C6-O6	8.15	124.79	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	131	C	O4'-C1'-N1	8.15	114.72	108.20
84	Aa	1695	C	N3-C4-C5	-8.15	118.64	121.90
84	Aa	3192	G	N1-C6-O6	8.14	124.79	119.90
86	Ab	11	A	C5-C6-N6	-8.14	117.18	123.70
1	Ad	481	A	O4'-C1'-N9	8.14	114.71	108.20
84	Aa	3004	G	N1-C6-O6	8.14	124.78	119.90
1	Ad	1047	G	O4'-C1'-N9	8.14	114.71	108.20
84	Aa	913	G	N1-C6-O6	8.14	124.78	119.90
84	Aa	1962	C	O4'-C1'-N1	8.14	114.71	108.20
1	Ad	189	U	P-O3'-C3'	8.14	129.46	119.70
1	Ad	836	U	N1-C1'-C2'	8.13	124.58	114.00
84	Aa	3216	G	N1-C6-O6	8.14	124.78	119.90
85	Ac	94	C	O4'-C1'-N1	8.13	114.71	108.20
84	Aa	520	G	N1-C6-O6	8.13	124.78	119.90
84	Aa	2241	G	N1-C6-O6	8.13	124.78	119.90
84	Aa	2402	G	P-O3'-C3'	8.13	129.46	119.70
84	Aa	2588	G	O4'-C1'-N9	8.13	114.70	108.20
84	Aa	2807	G	N1-C6-O6	8.13	124.78	119.90
1	Ad	903	A	N9-C1'-C2'	-8.13	103.06	112.00
86	Ab	69	A	C5-C6-N1	-8.13	113.64	117.70
84	Aa	777	G	N1-C6-O6	8.12	124.77	119.90
84	Aa	3184	G	N1-C6-O6	8.12	124.77	119.90
86	Ab	100	A	C5-C6-N1	-8.12	113.64	117.70
1	Ad	1711	G	O4'-C1'-N9	8.12	114.70	108.20
84	Aa	631	C	O4'-C1'-N1	8.12	114.70	108.20
84	Aa	1448	U	O4'-C1'-N1	8.12	114.70	108.20
84	Aa	3316	C	O4'-C1'-N1	8.12	114.70	108.20
86	Ab	72	G	O4'-C1'-N9	8.12	114.70	108.20
1	Ad	913	U	O4'-C1'-N1	8.12	114.69	108.20
84	Aa	1522	G	N1-C6-O6	8.12	124.77	119.90
85	Ac	1	C	O4'-C1'-N1	8.12	114.70	108.20
1	Ad	1139	C	C3'-C2'-C1'	8.12	107.99	101.50
84	Aa	621	C	O4'-C1'-N1	8.12	114.69	108.20
84	Aa	710	C	O4'-C1'-N1	8.11	114.69	108.20
84	Aa	570	G	N1-C6-O6	8.11	124.77	119.90
84	Aa	1615	G	N1-C6-O6	8.11	124.77	119.90
84	Aa	278	U	O4'-C1'-N1	8.11	114.69	108.20
84	Aa	1672	G	N1-C6-O6	8.11	124.77	119.90
84	Aa	2484	G	N1-C6-O6	8.11	124.77	119.90
84	Aa	3082	G	N1-C6-O6	8.11	124.77	119.90
84	Aa	3365	U	O4'-C1'-N1	8.11	114.69	108.20
1	Ad	193	G	C1'-O4'-C4'	-8.11	103.41	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2880	G	N1-C6-O6	8.11	124.77	119.90
84	Aa	2161	G	N1-C6-O6	8.11	124.76	119.90
84	Aa	2380	G	N1-C6-O6	8.11	124.76	119.90
84	Aa	2734	C	O4'-C1'-N1	8.11	114.69	108.20
84	Aa	2838	C	C2-N1-C1'	8.11	127.72	118.80
84	Aa	1969	G	N1-C6-O6	8.10	124.76	119.90
84	Aa	204	G	N1-C6-O6	8.10	124.76	119.90
84	Aa	3138	C	O4'-C1'-N1	8.10	114.68	108.20
84	Aa	1240	G	N1-C6-O6	8.10	124.76	119.90
84	Aa	1453	G	N1-C6-O6	8.10	124.76	119.90
84	Aa	2615	U	O4'-C1'-N1	8.10	114.68	108.20
1	Ad	158	C	C1'-O4'-C4'	-8.10	103.42	109.90
1	Ad	1062	C	O4'-C1'-N1	8.10	114.68	108.20
84	Aa	404	G	N1-C6-O6	8.10	124.76	119.90
86	Ab	82	G	C4-C5-C6	8.10	123.66	118.80
1	Ad	254	A	N9-C1'-C2'	8.10	124.53	114.00
84	Aa	3278	G	C5-C6-O6	-8.10	123.74	128.60
1	Ad	452	C	O4'-C1'-N1	8.10	114.68	108.20
84	Aa	2688	G	N1-C6-O6	8.10	124.76	119.90
1	Ad	617	G	C1'-O4'-C4'	-8.09	103.42	109.90
1	Ad	1513	A	O4'-C1'-N9	8.09	114.67	108.20
84	Aa	781	C	O4'-C1'-N1	8.09	114.67	108.20
84	Aa	2753	C	O4'-C1'-N1	8.09	114.67	108.20
84	Aa	3375	G	C5-C6-O6	-8.09	123.74	128.60
84	Aa	415	G	N1-C6-O6	8.09	124.75	119.90
84	Aa	596	C	O4'-C1'-N1	8.09	114.67	108.20
84	Aa	643	G	N1-C6-O6	8.09	124.75	119.90
84	Aa	1609	G	N1-C6-O6	8.09	124.75	119.90
84	Aa	2026	C	O4'-C1'-N1	8.09	114.67	108.20
84	Aa	2585	C	O4'-C1'-N1	8.09	114.67	108.20
1	Ad	393	G	C1'-O4'-C4'	-8.09	103.43	109.90
84	Aa	251	G	N1-C6-O6	8.09	124.75	119.90
84	Aa	511	C	O4'-C1'-N1	8.09	114.67	108.20
84	Aa	1827	U	O4'-C1'-N1	8.09	114.67	108.20
84	Aa	2826	G	N1-C6-O6	8.09	124.75	119.90
84	Aa	2687	C	O4'-C1'-N1	8.09	114.67	108.20
84	Aa	2837	C	O4'-C1'-N1	8.09	114.67	108.20
84	Aa	2913	A	N1-C6-N6	8.09	123.45	118.60
85	Ac	9	G	N1-C6-O6	8.09	124.75	119.90
1	Ad	765	U	O4'-C1'-N1	8.08	114.67	108.20
84	Aa	260	U	O4'-C1'-N1	8.08	114.67	108.20
84	Aa	1261	C	N3-C4-N4	8.08	123.66	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	746	C	C5'-C4'-O4'	8.08	118.80	109.10
84	Aa	977	G	N1-C6-O6	8.08	124.75	119.90
1	Ad	8	U	O4'-C1'-N1	8.08	114.66	108.20
1	Ad	1080	C	N1-C1'-C2'	8.08	124.50	114.00
84	Aa	442	C	O4'-C1'-N1	8.08	114.66	108.20
84	Aa	645	C	O4'-C1'-N1	8.08	114.67	108.20
84	Aa	2324	G	N1-C6-O6	8.08	124.75	119.90
84	Aa	3269	C	O4'-C1'-N1	8.08	114.66	108.20
86	Ab	66	G	C4-C5-C6	8.08	123.65	118.80
84	Aa	2408	G	N1-C6-O6	8.08	124.75	119.90
1	Ad	231	U	O4'-C1'-N1	8.07	114.66	108.20
1	Ad	1721	A	O4'-C1'-N9	8.07	114.66	108.20
84	Aa	233	C	O4'-C1'-N1	8.07	114.66	108.20
84	Aa	1173	C	O4'-C1'-N1	8.07	114.66	108.20
84	Aa	2123	C	O4'-C1'-N1	8.07	114.66	108.20
1	Ad	1588	C	N1-C1'-C2'	8.07	124.49	114.00
1	Ad	1685	U	C1'-O4'-C4'	-8.07	103.44	109.90
84	Aa	504	U	O4'-C1'-N1	8.07	114.66	108.20
84	Aa	565	C	O4'-C1'-N1	8.07	114.66	108.20
84	Aa	621	C	C2-N1-C1'	8.07	127.68	118.80
84	Aa	701	U	O4'-C1'-N1	8.07	114.66	108.20
84	Aa	1603	U	O4'-C1'-N1	8.07	114.66	108.20
84	Aa	717	G	N1-C6-O6	8.07	124.74	119.90
84	Aa	1878	G	N1-C6-O6	8.07	124.74	119.90
84	Aa	2117	G	N1-C6-O6	8.07	124.74	119.90
84	Aa	31	U	O4'-C1'-N1	8.07	114.65	108.20
84	Aa	1734	G	N1-C6-O6	8.07	124.74	119.90
84	Aa	2987	C	O4'-C1'-N1	8.07	114.66	108.20
85	Ac	123	G	N1-C6-O6	8.07	124.74	119.90
84	Aa	335	G	N1-C6-O6	8.06	124.74	119.90
84	Aa	461	A	C4'-C3'-O3'	8.06	129.13	113.00
84	Aa	597	C	O4'-C1'-N1	8.06	114.65	108.20
84	Aa	788	G	N1-C6-O6	8.06	124.74	119.90
84	Aa	1209	G	N1-C6-O6	8.06	124.74	119.90
84	Aa	337	C	O4'-C1'-N1	8.06	114.65	108.20
84	Aa	911	G	C5-C6-O6	-8.06	123.76	128.60
84	Aa	3189	C	O4'-C1'-N1	8.06	114.65	108.20
84	Aa	298	G	N1-C6-O6	8.06	124.74	119.90
84	Aa	436	G	C5-C6-O6	-8.06	123.76	128.60
84	Aa	419	G	N1-C6-O6	8.06	124.73	119.90
84	Aa	3158	C	O4'-C1'-N1	8.06	114.65	108.20
1	Ad	840	U	O4'-C1'-N1	8.06	114.64	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	735	C	O4'-C1'-N1	8.06	114.65	108.20
84	Aa	2713	G	N1-C6-O6	8.06	124.73	119.90
84	Aa	495	G	N1-C6-O6	8.05	124.73	119.90
84	Aa	2614	U	O4'-C1'-N1	8.05	114.64	108.20
84	Aa	571	G	N1-C6-O6	8.05	124.73	119.90
84	Aa	2191	C	O4'-C1'-N1	8.05	114.64	108.20
1	Ad	1049	U	O4'-C1'-N1	8.05	114.64	108.20
1	Ad	1496	A	P-O3'-C3'	8.05	129.36	119.70
1	Ad	1521	G	O4'-C1'-N9	8.05	114.64	108.20
84	Aa	1714	A	P-O3'-C3'	8.05	129.36	119.70
1	Ad	238	G	C3'-C2'-C1'	8.05	107.94	101.50
84	Aa	3183	G	P-O5'-C5'	8.05	133.77	120.90
1	Ad	1482	U	O4'-C1'-N1	8.04	114.64	108.20
84	Aa	400	G	C5-C6-O6	-8.04	123.77	128.60
84	Aa	519	C	O4'-C1'-N1	8.04	114.63	108.20
84	Aa	3086	G	C5-C6-O6	-8.04	123.78	128.60
84	Aa	3145	G	N1-C6-O6	8.04	124.72	119.90
1	Ad	288	G	C1'-O4'-C4'	-8.04	103.47	109.90
84	Aa	138	G	C5-C6-O6	-8.04	123.78	128.60
84	Aa	961	C	O4'-C1'-N1	8.04	114.63	108.20
84	Aa	1311	G	O4'-C1'-N9	8.04	114.63	108.20
84	Aa	1345	U	O4'-C1'-N1	8.04	114.63	108.20
84	Aa	2551	U	O4'-C1'-N1	8.04	114.63	108.20
84	Aa	1068	A	P-O3'-C3'	8.03	129.34	119.70
84	Aa	1087	G	N1-C6-O6	8.04	124.72	119.90
84	Aa	2278	G	C5-C6-O6	-8.04	123.78	128.60
84	Aa	3154	G	N1-C6-O6	8.04	124.72	119.90
85	Ac	57	C	O4'-C1'-N1	8.03	114.63	108.20
1	Ad	468	A	N9-C1'-C2'	-8.03	103.16	112.00
84	Aa	1725	G	O4'-C1'-N9	8.03	114.63	108.20
84	Aa	2711	U	O4'-C1'-N1	8.03	114.63	108.20
84	Aa	2941	G	N1-C6-O6	8.03	124.72	119.90
84	Aa	1733	G	N1-C6-O6	8.03	124.72	119.90
84	Aa	2329	C	O4'-C1'-N1	8.03	114.62	108.20
85	Ac	154	G	O4'-C1'-N9	8.03	114.62	108.20
84	Aa	1579	C	O4'-C1'-N1	8.03	114.62	108.20
84	Aa	1646	U	O4'-C1'-N1	8.03	114.62	108.20
1	Ad	619	A	O4'-C1'-N9	8.03	114.62	108.20
1	Ad	1260	A	N9-C1'-C2'	8.03	124.44	114.00
1	Ad	1499	U	O4'-C1'-N1	8.03	114.62	108.20
84	Aa	1217	G	C5-C6-O6	-8.03	123.78	128.60
84	Aa	2363	G	C5-C6-O6	-8.03	123.78	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2919	G	C5-C6-O6	-8.03	123.78	128.60
85	Ac	66	G	N1-C6-O6	8.03	124.72	119.90
84	Aa	1961	C	O4'-C1'-N1	8.02	114.62	108.20
1	Ad	1571	G	O4'-C1'-N9	8.02	114.62	108.20
84	Aa	174	G	N1-C6-O6	8.02	124.71	119.90
84	Aa	281	G	C5-C6-O6	-8.02	123.79	128.60
84	Aa	1071	G	N1-C6-O6	8.02	124.71	119.90
1	Ad	1434	G	C1'-O4'-C4'	-8.02	103.48	109.90
84	Aa	2233	G	N1-C6-O6	8.02	124.71	119.90
1	Ad	9	U	O4'-C1'-N1	8.02	114.61	108.20
1	Ad	928	A	O4'-C1'-N9	8.02	114.61	108.20
84	Aa	492	G	C5'-C4'-O4'	8.02	118.72	109.10
84	Aa	533	G	N1-C6-O6	8.02	124.71	119.90
84	Aa	2848	U	O4'-C1'-N1	8.02	114.61	108.20
85	Ac	132	C	O4'-C1'-N1	8.02	114.61	108.20
1	Ad	808	G	O4'-C1'-N9	8.02	114.61	108.20
84	Aa	795	C	O4'-C1'-N1	8.02	114.61	108.20
84	Aa	2243	C	O4'-C1'-N1	8.02	114.61	108.20
84	Aa	2579	G	N1-C6-O6	8.02	124.71	119.90
84	Aa	3256	C	O4'-C1'-N1	8.02	114.61	108.20
84	Aa	1775	C	O4'-C1'-N1	8.01	114.61	108.20
84	Aa	1908	C	O4'-C1'-N1	8.01	114.61	108.20
1	Ad	98	C	N1-C1'-C2'	8.01	124.42	114.00
84	Aa	2513	U	C5'-C4'-O4'	-8.01	99.49	109.10
84	Aa	2843	G	N1-C6-O6	8.01	124.71	119.90
84	Aa	412	C	O4'-C1'-N1	8.01	114.61	108.20
84	Aa	1327	G	N1-C6-O6	8.01	124.71	119.90
84	Aa	250	C	O4'-C1'-N1	8.01	114.61	108.20
84	Aa	1383	G	O4'-C1'-N9	8.01	114.61	108.20
84	Aa	1649	G	C5-C6-O6	-8.01	123.80	128.60
84	Aa	2359	C	O4'-C1'-N1	8.01	114.61	108.20
1	Ad	1472	G	C1'-O4'-C4'	-8.01	103.50	109.90
84	Aa	2850	G	C5-C6-O6	-8.01	123.80	128.60
84	Aa	127	G	N1-C6-O6	8.00	124.70	119.90
84	Aa	49	U	O4'-C1'-N1	8.00	114.60	108.20
1	Ad	954	C	C1'-O4'-C4'	-8.00	103.50	109.90
84	Aa	63	G	N1-C6-O6	8.00	124.70	119.90
84	Aa	1821	G	N1-C6-O6	8.00	124.70	119.90
84	Aa	2549	C	O4'-C1'-N1	8.00	114.60	108.20
86	Ab	119	C	N3-C4-C5	-8.00	118.70	121.90
1	Ad	19	A	O4'-C1'-N9	8.00	114.60	108.20
84	Aa	1035	C	O4'-C1'-N1	8.00	114.60	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1923	G	C5-C6-O6	-8.00	123.80	128.60
86	Ab	19	A	O4'-C1'-N9	7.99	114.59	108.20
1	Ad	945	A	O4'-C1'-N9	7.99	114.59	108.20
84	Aa	1110	C	O4'-C1'-N1	7.99	114.59	108.20
84	Aa	1338	C	O4'-C1'-N1	7.99	114.59	108.20
84	Aa	1787	C	O4'-C1'-N1	7.99	114.59	108.20
84	Aa	2086	A	O4'-C1'-N9	7.99	114.59	108.20
84	Aa	801	G	N1-C6-O6	7.99	124.69	119.90
1	Ad	89	U	O4'-C1'-N1	7.99	114.59	108.20
1	Ad	1576	C	C1'-O4'-C4'	7.99	116.29	109.90
1	Ad	1263	C	N1-C1'-C2'	7.98	124.38	114.00
84	Aa	694	U	O4'-C1'-N1	7.98	114.59	108.20
84	Aa	852	C	O4'-C1'-N1	7.98	114.59	108.20
84	Aa	161	C	O4'-C1'-N1	7.98	114.58	108.20
84	Aa	750	G	N1-C6-O6	7.98	124.69	119.90
84	Aa	1647	C	O4'-C1'-N1	7.98	114.58	108.20
84	Aa	2015	G	N1-C6-O6	7.98	124.69	119.90
84	Aa	2873	G	C5-C6-O6	-7.98	123.81	128.60
84	Aa	1033	G	N1-C6-O6	7.98	124.69	119.90
84	Aa	1080	C	O4'-C1'-N1	7.98	114.58	108.20
84	Aa	1194	C	O4'-C1'-N1	7.98	114.58	108.20
84	Aa	1293	C	O4'-C1'-N1	7.98	114.58	108.20
84	Aa	1362	C	O4'-C1'-N1	7.98	114.58	108.20
84	Aa	712	A	N1-C6-N6	7.98	123.39	118.60
84	Aa	967	G	C5-C6-O6	-7.97	123.81	128.60
84	Aa	1111	U	O4'-C1'-N1	7.97	114.58	108.20
1	Ad	1011	C	O4'-C1'-N1	7.97	114.58	108.20
84	Aa	1511	C	O4'-C1'-N1	7.97	114.58	108.20
84	Aa	2625	C	O4'-C1'-N1	7.97	114.58	108.20
84	Aa	3132	U	O4'-C1'-N1	7.97	114.58	108.20
85	Ac	97	G	C5-C6-O6	-7.97	123.82	128.60
1	Ad	14	C	N1-C1'-C2'	7.97	124.36	114.00
84	Aa	2061	C	O4'-C1'-N1	7.97	114.57	108.20
84	Aa	3032	G	C5-C6-O6	-7.97	123.82	128.60
84	Aa	1743	C	O4'-C1'-N1	7.96	114.57	108.20
84	Aa	1752	C	O4'-C1'-N1	7.96	114.57	108.20
1	Ad	1653	G	O4'-C1'-N9	7.96	114.57	108.20
1	Ad	1766	A	C1'-O4'-C4'	7.96	116.27	109.90
1	Ad	1202	G	C1'-O4'-C4'	-7.96	103.53	109.90
84	Aa	1383	G	C5-C6-O6	-7.96	123.82	128.60
84	Aa	2759	C	O4'-C1'-N1	7.96	114.57	108.20
84	Aa	1796	A	C5-C6-N6	-7.96	117.33	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	Ac	137	G	N1-C6-O6	7.96	124.68	119.90
1	Ad	1731	A	O4'-C1'-C2'	-7.96	97.84	105.80
84	Aa	1486	G	C5-C6-O6	-7.96	123.83	128.60
84	Aa	2499	U	O4'-C1'-N1	7.96	114.57	108.20
84	Aa	604	C	O4'-C1'-N1	7.96	114.57	108.20
84	Aa	3020	C	O4'-C1'-N1	7.96	114.56	108.20
84	Aa	1160	G	N1-C6-O6	7.96	124.67	119.90
84	Aa	1983	U	O4'-C1'-N1	7.96	114.56	108.20
84	Aa	2618	G	N1-C6-O6	7.96	124.67	119.90
84	Aa	952	C	O4'-C1'-N1	7.95	114.56	108.20
84	Aa	3197	C	O4'-C1'-N1	7.95	114.56	108.20
85	Ac	116	G	N1-C6-O6	7.95	124.67	119.90
84	Aa	2419	C	O4'-C1'-N1	7.95	114.56	108.20
84	Aa	2715	U	O4'-C1'-N1	7.95	114.56	108.20
84	Aa	1549	A	N1-C6-N6	7.95	123.37	118.60
84	Aa	878	G	C5-C6-O6	-7.95	123.83	128.60
84	Aa	2272	C	O4'-C1'-N1	7.95	114.56	108.20
80	Cf	107	TYR	CB-CG-CD2	-7.94	116.23	121.00
84	Aa	2581	C	O4'-C1'-N1	7.94	114.56	108.20
84	Aa	3269	C	C2-N1-C1'	7.94	127.54	118.80
84	Aa	1901	G	C5-C6-O6	-7.94	123.83	128.60
84	Aa	2225	C	O4'-C1'-N1	7.94	114.55	108.20
84	Aa	3111	C	O4'-C1'-N1	7.94	114.55	108.20
1	Ad	1194	C	O4'-C1'-C2'	-7.94	97.86	105.80
1	Ad	269	A	O4'-C1'-N9	7.94	114.55	108.20
41	CA	69	TYR	N-CA-CB	7.94	124.89	110.60
84	Aa	3244	G	C5-C6-O6	-7.94	123.84	128.60
84	Aa	603	G	O4'-C1'-N9	7.94	114.55	108.20
84	Aa	1070	G	N1-C6-O6	7.94	124.66	119.90
84	Aa	1390	G	O4'-C1'-N9	7.94	114.55	108.20
84	Aa	3385	G	N1-C6-O6	7.94	124.66	119.90
84	Aa	604	C	N3-C4-C5	-7.93	118.73	121.90
84	Aa	2260	C	O4'-C1'-N1	7.93	114.55	108.20
84	Aa	3031	G	N1-C6-O6	7.93	124.66	119.90
84	Aa	2952	G	N1-C6-O6	7.93	124.66	119.90
1	Ad	937	A	C1'-O4'-C4'	7.93	116.24	109.90
84	Aa	822	U	O4'-C1'-N1	7.93	114.54	108.20
84	Aa	1352	G	C5-C6-O6	-7.93	123.84	128.60
84	Aa	1648	C	O4'-C1'-N1	7.93	114.54	108.20
84	Aa	3214	U	O4'-C1'-N1	7.93	114.54	108.20
84	Aa	2833	G	N1-C6-O6	7.93	124.66	119.90
84	Aa	347	A	N1-C6-N6	7.93	123.36	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	676	G	N1-C6-O6	7.93	124.66	119.90
84	Aa	1853	C	O4'-C1'-N1	7.93	114.54	108.20
84	Aa	2092	C	O4'-C4'-C3'	-7.93	96.07	104.00
84	Aa	2975	G	N1-C6-O6	7.93	124.66	119.90
84	Aa	3019	C	O4'-C1'-N1	7.93	114.54	108.20
86	Ab	48	G	N1-C6-O6	7.93	124.66	119.90
1	Ad	1113	G	O4'-C1'-C2'	7.92	114.73	107.60
84	Aa	580	C	P-O3'-C3'	7.92	129.21	119.70
84	Aa	1306	A	O4'-C1'-N9	7.92	114.54	108.20
84	Aa	1700	U	O4'-C1'-N1	7.92	114.54	108.20
84	Aa	2064	C	O4'-C1'-N1	7.92	114.54	108.20
84	Aa	2491	A	P-O3'-C3'	7.92	129.21	119.70
84	Aa	2523	G	C5-C6-O6	-7.92	123.85	128.60
84	Aa	3143	A	C5-C6-N6	-7.92	117.36	123.70
84	Aa	2177	U	N1-C1'-C2'	7.92	124.30	114.00
84	Aa	2531	G	N1-C6-O6	7.92	124.65	119.90
84	Aa	3223	C	O4'-C1'-N1	7.92	114.54	108.20
86	Ab	14	C	C5-C6-N1	7.92	124.96	121.00
86	Ab	44	C	C2-N3-C4	7.92	123.86	119.90
1	Ad	1056	A	O4'-C1'-N9	7.92	114.53	108.20
84	Aa	32	G	C5-C6-O6	-7.92	123.85	128.60
1	Ad	281	U	C3'-C2'-C1'	-7.92	95.17	101.50
84	Aa	311	G	N1-C6-O6	7.92	124.65	119.90
85	Ac	93	U	O4'-C1'-N1	7.92	114.53	108.20
1	Ad	821	G	C3'-C2'-C1'	7.91	107.83	101.50
84	Aa	75	G	N1-C6-O6	7.91	124.65	119.90
84	Aa	3119	C	O4'-C1'-N1	7.91	114.53	108.20
84	Aa	4	C	O4'-C1'-N1	7.91	114.53	108.20
84	Aa	489	C	O4'-C1'-N1	7.91	114.53	108.20
84	Aa	2463	U	O4'-C1'-N1	7.91	114.53	108.20
84	Aa	2992	G	C5-C6-O6	-7.91	123.85	128.60
1	Ad	1568	U	C1'-O4'-C4'	7.91	116.23	109.90
84	Aa	182	C	O4'-C1'-N1	7.91	114.53	108.20
84	Aa	2008	G	N1-C6-O6	7.91	124.64	119.90
84	Aa	2321	C	O4'-C1'-N1	7.91	114.53	108.20
84	Aa	2866	A	N1-C6-N6	7.91	123.34	118.60
84	Aa	205	C	O4'-C1'-N1	7.91	114.53	108.20
1	Ad	183	C	N1-C1'-C2'	7.90	124.27	114.00
1	Ad	1128	C	P-O5'-C5'	7.90	133.54	120.90
84	Aa	3	G	N1-C6-O6	7.90	124.64	119.90
84	Aa	1707	C	O4'-C1'-N1	7.90	114.52	108.20
84	Aa	2167	G	C5'-C4'-C3'	7.90	128.65	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	339	G	C1'-O4'-C4'	-7.90	103.58	109.90
84	Aa	1108	U	O4'-C1'-N1	7.90	114.52	108.20
84	Aa	1320	G	C5-C6-O6	-7.90	123.86	128.60
1	Ad	96	G	N9-C1'-C2'	7.90	124.27	114.00
1	Ad	1218	U	C1'-O4'-C4'	7.90	116.22	109.90
84	Aa	1580	C	O4'-C1'-N1	7.90	114.52	108.20
84	Aa	2851	C	O4'-C1'-N1	7.90	114.52	108.20
1	Ad	321	C	C3'-C2'-C1'	7.90	107.82	101.50
1	Ad	369	G	O4'-C1'-C2'	-7.90	97.90	105.80
84	Aa	3232	C	O4'-C1'-N1	7.90	114.52	108.20
1	Ad	834	A	O4'-C1'-C2'	-7.89	97.91	105.80
84	Aa	54	G	N1-C6-O6	7.89	124.64	119.90
1	Ad	565	G	O4'-C1'-N9	7.89	114.51	108.20
1	Ad	1594	A	O4'-C1'-C2'	-7.89	97.91	105.80
84	Aa	191	C	O4'-C1'-N1	7.89	114.51	108.20
84	Aa	521	G	C5-C6-O6	-7.89	123.86	128.60
84	Aa	1131	U	O4'-C1'-N1	7.89	114.51	108.20
84	Aa	1197	A	O4'-C1'-N9	7.89	114.51	108.20
84	Aa	1613	C	O4'-C1'-N1	7.89	114.51	108.20
84	Aa	2494	A	C5-C6-N6	-7.89	117.39	123.70
84	Aa	3373	C	O4'-C1'-N1	7.89	114.51	108.20
84	Aa	285	G	C5-C6-O6	-7.89	123.86	128.60
84	Aa	1287	C	O4'-C1'-N1	7.89	114.51	108.20
84	Aa	500	C	O4'-C1'-N1	7.89	114.51	108.20
84	Aa	1328	C	O4'-C1'-N1	7.89	114.51	108.20
84	Aa	3289	U	O4'-C1'-N1	7.89	114.51	108.20
1	Ad	825	U	O4'-C1'-N1	7.89	114.51	108.20
1	Ad	1205	G	N9-C1'-C2'	-7.89	103.33	112.00
1	Ad	1755	G	O4'-C1'-N9	7.89	114.51	108.20
84	Aa	2037	C	O4'-C1'-N1	7.89	114.51	108.20
84	Aa	3283	G	N1-C6-O6	7.89	124.63	119.90
1	Ad	535	C	C3'-C2'-C1'	7.88	107.81	101.50
84	Aa	543	C	C2'-C3'-O3'	7.88	126.85	109.50
84	Aa	587	A	C4-C5-C6	7.88	120.94	117.00
1	Ad	176	A	N9-C1'-C2'	-7.88	103.33	112.00
84	Aa	743	C	O4'-C1'-N1	7.88	114.50	108.20
84	Aa	1722	G	P-O3'-C3'	7.88	129.16	119.70
84	Aa	2534	G	N1-C6-O6	7.88	124.63	119.90
84	Aa	2870	U	O4'-C1'-N1	7.88	114.51	108.20
84	Aa	458	G	N1-C6-O6	7.88	124.63	119.90
1	Ad	743	G	O4'-C1'-C2'	-7.88	97.92	105.80
84	Aa	1246	G	N1-C6-O6	7.88	124.63	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	275	C	O4'-C1'-N1	7.88	114.50	108.20
84	Aa	640	C	C6-N1-C2	-7.88	117.15	120.30
84	Aa	856	G	N1-C6-O6	7.88	124.63	119.90
84	Aa	3246	U	O4'-C1'-N1	7.88	114.50	108.20
1	Ad	1397	A	C1'-O4'-C4'	-7.87	103.60	109.90
84	Aa	2773	G	N1-C6-O6	7.87	124.62	119.90
84	Aa	2997	C	O4'-C1'-N1	7.87	114.50	108.20
84	Aa	1017	G	C5-C6-O6	-7.87	123.88	128.60
84	Aa	1364	C	O4'-C1'-N1	7.87	114.50	108.20
84	Aa	2643	A	N1-C6-N6	7.87	123.32	118.60
84	Aa	3213	A	O4'-C1'-N9	7.87	114.50	108.20
1	Ad	14	C	C1'-O4'-C4'	-7.87	103.60	109.90
1	Ad	1096	A	C1'-O4'-C4'	-7.87	103.60	109.90
84	Aa	244	G	N1-C6-O6	7.87	124.62	119.90
84	Aa	334	A	C5-C6-N1	-7.87	113.77	117.70
84	Aa	1574	C	O4'-C1'-N1	7.87	114.49	108.20
84	Aa	2906	U	O4'-C1'-N1	7.87	114.49	108.20
2	Ae	74	C	P-O3'-C3'	7.87	129.14	119.70
84	Aa	391	U	O4'-C1'-N1	7.87	114.49	108.20
84	Aa	912	G	N1-C6-O6	7.87	124.62	119.90
1	Ad	214	A	C1'-O4'-C4'	7.86	116.19	109.90
1	Ad	1041	A	N9-C1'-C2'	-7.86	103.35	112.00
1	Ad	1645	C	O4'-C1'-C2'	-7.86	97.94	105.80
84	Aa	2859	C	O4'-C1'-N1	7.86	114.49	108.20
85	Ac	19	A	N1-C6-N6	7.86	123.32	118.60
1	Ad	784	C	C3'-C2'-C1'	-7.86	95.21	101.50
84	Aa	227	C	O4'-C1'-N1	7.86	114.49	108.20
84	Aa	584	G	P-O3'-C3'	7.86	129.13	119.70
84	Aa	1162	A	O4'-C1'-N9	7.86	114.49	108.20
84	Aa	1976	U	O4'-C1'-N1	7.86	114.49	108.20
84	Aa	1530	C	O4'-C1'-N1	7.86	114.48	108.20
84	Aa	2145	C	O4'-C1'-N1	7.86	114.48	108.20
84	Aa	2895	G	N1-C6-O6	7.86	124.61	119.90
84	Aa	3341	C	O4'-C1'-N1	7.86	114.48	108.20
84	Aa	2143	A	C5-C6-N6	-7.85	117.42	123.70
84	Aa	1382	C	O4'-C1'-N1	7.85	114.48	108.20
84	Aa	1939	C	O4'-C1'-N1	7.85	114.48	108.20
84	Aa	2391	C	O4'-C1'-N1	7.85	114.48	108.20
84	Aa	3200	A	P-O3'-C3'	7.85	129.12	119.70
85	Ac	95	G	C5-C6-O6	-7.85	123.89	128.60
84	Aa	985	C	O4'-C1'-N1	7.85	114.48	108.20
84	Aa	2354	G	C5-C6-O6	-7.85	123.89	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	628	G	O4'-C1'-N9	7.85	114.48	108.20
84	Aa	559	U	O4'-C1'-N1	7.85	114.48	108.20
84	Aa	767	U	O4'-C1'-N1	7.85	114.48	108.20
84	Aa	1687	C	O4'-C1'-N1	7.85	114.48	108.20
84	Aa	2018	C	O4'-C1'-N1	7.85	114.48	108.20
84	Aa	2692	G	C5-C6-O6	-7.85	123.89	128.60
84	Aa	3390	G	C5-C6-O6	-7.85	123.89	128.60
84	Aa	2610	G	C5-C6-O6	-7.85	123.89	128.60
84	Aa	3025	A	C5-C6-N6	-7.85	117.42	123.70
1	Ad	367	G	O4'-C1'-N9	7.84	114.47	108.20
84	Aa	34	G	N1-C6-O6	7.84	124.61	119.90
84	Aa	1105	G	C5-C6-O6	-7.84	123.89	128.60
84	Aa	2506	G	O4'-C1'-N9	7.84	114.47	108.20
85	Ac	117	C	O4'-C1'-N1	7.84	114.48	108.20
84	Aa	674	G	N1-C6-O6	7.84	124.61	119.90
84	Aa	2587	G	C5-C6-O6	-7.84	123.89	128.60
1	Ad	833	U	O4'-C1'-N1	7.84	114.47	108.20
1	Ad	1730	G	C3'-C2'-C1'	7.84	107.77	101.50
84	Aa	130	G	N1-C6-O6	7.84	124.61	119.90
84	Aa	280	G	O4'-C1'-N9	7.84	114.47	108.20
84	Aa	1023	G	N1-C6-O6	7.84	124.61	119.90
84	Aa	1641	G	C5-C6-O6	-7.84	123.89	128.60
84	Aa	3225	G	C5-C6-O6	-7.84	123.89	128.60
84	Aa	759	C	O4'-C1'-N1	7.84	114.47	108.20
1	Ad	1432	C	N1-C1'-C2'	-7.84	103.38	112.00
84	Aa	1541	G	N1-C6-O6	7.84	124.60	119.90
2	Ae	6	G	N9-C1'-C2'	7.84	124.19	114.00
84	Aa	27	C	O4'-C1'-N1	7.84	114.47	108.20
84	Aa	2056	C	P-O3'-C3'	7.84	129.10	119.70
84	Aa	2626	G	N1-C6-O6	7.84	124.60	119.90
86	Ab	100	A	C4-C5-C6	7.84	120.92	117.00
84	Aa	51	A	C5-C6-N6	-7.83	117.43	123.70
84	Aa	2343	U	O4'-C1'-N1	7.83	114.47	108.20
1	Ad	430	G	C1'-O4'-C4'	-7.83	103.63	109.90
1	Ad	1789	U	O4'-C1'-N1	7.83	114.47	108.20
84	Aa	2831	U	O4'-C1'-N1	7.83	114.47	108.20
1	Ad	1027	C	N1-C1'-C2'	7.83	124.18	114.00
1	Ad	1698	A	O4'-C1'-N9	7.83	114.47	108.20
84	Aa	2005	C	O4'-C1'-N1	7.83	114.47	108.20
84	Aa	3291	C	O4'-C1'-N1	7.83	114.47	108.20
84	Aa	1694	A	C5-C6-N6	-7.83	117.44	123.70
85	Ac	68	G	N1-C6-O6	7.83	124.60	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	Ab	44	C	N3-C4-N4	7.83	123.48	118.00
1	Ad	1700	G	C1'-O4'-C4'	-7.83	103.64	109.90
84	Aa	2605	G	C5-C6-O6	-7.83	123.90	128.60
1	Ad	205	U	O4'-C1'-N1	7.83	114.46	108.20
84	Aa	871	C	O4'-C1'-N1	7.83	114.46	108.20
84	Aa	1073	G	N1-C6-O6	7.83	124.59	119.90
84	Aa	1089	G	C5-C6-O6	-7.83	123.90	128.60
84	Aa	2456	G	N1-C6-O6	7.83	124.59	119.90
84	Aa	2567	C	O4'-C1'-N1	7.83	114.46	108.20
84	Aa	1616	G	N1-C6-O6	7.82	124.59	119.90
84	Aa	2384	G	C5-C6-O6	-7.82	123.91	128.60
1	Ad	834	A	P-O3'-C3'	7.82	129.09	119.70
84	Aa	203	C	O4'-C1'-N1	7.82	114.46	108.20
1	Ad	612	U	C1'-O4'-C4'	-7.82	103.64	109.90
84	Aa	373	A	P-O3'-C3'	7.82	129.08	119.70
84	Aa	2525	G	C5-C6-O6	-7.82	123.91	128.60
84	Aa	28	C	O4'-C1'-N1	7.82	114.45	108.20
84	Aa	3257	G	N1-C6-O6	7.82	124.59	119.90
86	Ab	87	G	C5-C6-O6	-7.82	123.91	128.60
1	Ad	563	C	N1-C1'-C2'	7.82	124.16	114.00
1	Ad	1436	U	O4'-C1'-N1	7.82	114.45	108.20
84	Aa	1037	U	O4'-C1'-N1	7.81	114.45	108.20
84	Aa	3366	C	O4'-C1'-N1	7.81	114.45	108.20
1	Ad	243	U	P-O3'-C3'	7.81	129.08	119.70
84	Aa	2410	U	O4'-C1'-N1	7.81	114.45	108.20
84	Aa	3217	G	N1-C6-O6	7.81	124.59	119.90
84	Aa	3251	C	O4'-C1'-N1	7.81	114.45	108.20
84	Aa	2440	U	O4'-C1'-N1	7.81	114.45	108.20
86	Ab	21	U	O4'-C1'-N1	7.81	114.45	108.20
84	Aa	335	G	O4'-C1'-N9	7.81	114.45	108.20
85	Ac	122	G	C5-C6-O6	-7.81	123.92	128.60
84	Aa	296	C	O4'-C1'-N1	7.81	114.44	108.20
84	Aa	1575	G	O4'-C1'-N9	7.81	114.44	108.20
1	Ad	895	U	O4'-C1'-N1	7.80	114.44	108.20
1	Ad	1746	U	O4'-C1'-N1	7.80	114.44	108.20
85	Ac	59	A	C4-C5-C6	7.80	120.90	117.00
1	Ad	1185	U	O4'-C1'-N1	7.80	114.44	108.20
1	Ad	1338	U	O4'-C1'-N1	7.80	114.44	108.20
1	Ad	1528	U	O4'-C1'-N1	7.80	114.44	108.20
84	Aa	806	C	O4'-C1'-N1	7.80	114.44	108.20
84	Aa	3340	G	N1-C6-O6	7.80	124.58	119.90
84	Aa	1953	C	O4'-C1'-N1	7.80	114.44	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	Ab	17	G	C5-C6-O6	-7.80	123.92	128.60
84	Aa	1121	C	O4'-C1'-N1	7.80	114.44	108.20
84	Aa	1937	C	O4'-C1'-N1	7.80	114.44	108.20
84	Aa	302	G	O4'-C1'-N9	7.80	114.44	108.20
84	Aa	828	U	O4'-C1'-N1	7.80	114.44	108.20
84	Aa	1237	G	C5-C6-O6	-7.80	123.92	128.60
84	Aa	3196	C	O4'-C1'-N1	7.80	114.44	108.20
84	Aa	3351	A	C5-C6-N6	-7.80	117.46	123.70
84	Aa	479	C	O4'-C1'-N1	7.79	114.44	108.20
84	Aa	2729	C	O4'-C1'-N1	7.79	114.44	108.20
1	Ad	814	C	N1-C1'-C2'	7.79	124.13	114.00
84	Aa	125	G	C5-C6-O6	-7.79	123.92	128.60
84	Aa	268	U	O4'-C1'-N1	7.79	114.44	108.20
84	Aa	1009	G	N1-C6-O6	7.79	124.58	119.90
1	Ad	743	G	P-O3'-C3'	7.79	129.05	119.70
1	Ad	1017	U	O4'-C1'-N1	7.79	114.43	108.20
84	Aa	561	G	N1-C6-O6	7.79	124.57	119.90
84	Aa	1772	G	C5-C6-O6	-7.79	123.93	128.60
84	Aa	2242	G	C5-C6-O6	-7.79	123.93	128.60
84	Aa	2339	U	O4'-C1'-N1	7.79	114.43	108.20
84	Aa	2779	G	N1-C6-O6	7.79	124.57	119.90
1	Ad	1511	A	C3'-C2'-C1'	-7.79	95.27	101.50
84	Aa	2164	G	N1-C6-O6	7.79	124.57	119.90
84	Aa	1754	C	O4'-C1'-N1	7.78	114.43	108.20
84	Aa	234	G	N1-C6-O6	7.78	124.57	119.90
84	Aa	2077	C	O4'-C1'-N1	7.78	114.42	108.20
84	Aa	3112	U	O4'-C1'-N1	7.78	114.43	108.20
84	Aa	2279	C	O4'-C1'-N1	7.78	114.42	108.20
84	Aa	2616	U	O4'-C1'-N1	7.78	114.42	108.20
84	Aa	2766	U	O4'-C1'-N1	7.78	114.42	108.20
84	Aa	1308	A	C5-C6-N6	-7.78	117.48	123.70
1	Ad	401	A	P-O5'-C5'	-7.77	108.47	120.90
1	Ad	1471	C	C1'-O4'-C4'	-7.77	103.68	109.90
84	Aa	355	C	O4'-C1'-N1	7.77	114.42	108.20
84	Aa	2167	G	C5'-C4'-O4'	7.77	118.43	109.10
1	Ad	1761	G	O4'-C1'-N9	7.77	114.41	108.20
84	Aa	874	U	O4'-C1'-N1	7.77	114.41	108.20
84	Aa	1135	C	O4'-C1'-N1	7.77	114.41	108.20
84	Aa	1749	G	N1-C6-O6	7.77	124.56	119.90
84	Aa	1057	A	C5-C6-N6	-7.77	117.49	123.70
84	Aa	1987	C	O4'-C1'-N1	7.76	114.41	108.20
85	Ac	131	G	C5-C6-O6	-7.76	123.94	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	Ab	97	G	N3-C2-N2	7.76	125.33	119.90
1	Ad	567	U	O4'-C1'-N1	7.76	114.41	108.20
84	Aa	257	C	O4'-C1'-N1	7.76	114.41	108.20
84	Aa	734	C	O4'-C1'-N1	7.76	114.41	108.20
84	Aa	1873	C	O4'-C1'-N1	7.76	114.41	108.20
84	Aa	3388	U	O4'-C1'-N1	7.76	114.41	108.20
85	Ac	26	C	O4'-C1'-N1	7.76	114.41	108.20
84	Aa	1951	C	O4'-C1'-N1	7.76	114.41	108.20
1	Ad	785	A	C1'-O4'-C4'	7.76	116.11	109.90
84	Aa	93	G	N1-C6-O6	7.76	124.55	119.90
84	Aa	497	G	C5-C6-O6	-7.76	123.95	128.60
84	Aa	2830	G	C5-C6-O6	-7.76	123.95	128.60
84	Aa	3345	G	C5-C6-O6	-7.76	123.95	128.60
84	Aa	2052	G	N1-C6-O6	7.75	124.55	119.90
1	Ad	435	C	O4'-C1'-N1	7.75	114.40	108.20
1	Ad	1574	U	N1-C1'-C2'	-7.75	103.47	112.00
84	Aa	522	C	C5'-C4'-O4'	7.75	118.40	109.10
84	Aa	594	C	O4'-C1'-N1	7.75	114.40	108.20
84	Aa	1582	C	O4'-C1'-N1	7.75	114.40	108.20
84	Aa	2696	C	O4'-C1'-N1	7.75	114.40	108.20
1	Ad	900	G	N9-C1'-C2'	7.75	124.08	114.00
1	Ad	1576	C	O4'-C1'-N1	7.75	114.40	108.20
84	Aa	673	U	O4'-C1'-N1	7.75	114.40	108.20
84	Aa	2962	C	O4'-C1'-N1	7.75	114.40	108.20
84	Aa	2988	U	O4'-C1'-N1	7.75	114.40	108.20
84	Aa	1740	U	O4'-C1'-N1	7.75	114.40	108.20
84	Aa	3247	C	O4'-C1'-N1	7.75	114.40	108.20
84	Aa	192	C	O4'-C1'-N1	7.75	114.40	108.20
84	Aa	229	G	N1-C6-O6	7.75	124.55	119.90
1	Ad	1450	A	C3'-C2'-C1'	7.75	107.70	101.50
1	Ad	1734	U	O4'-C1'-N1	7.75	114.40	108.20
85	Ac	147	C	O4'-C1'-N1	7.75	114.40	108.20
84	Aa	648	G	C5-C6-O6	-7.74	123.95	128.60
84	Aa	1219	C	O4'-C1'-N1	7.74	114.39	108.20
1	Ad	239	C	P-O3'-C3'	7.74	128.99	119.70
84	Aa	755	C	O4'-C1'-N1	7.74	114.39	108.20
84	Aa	2363	G	O4'-C1'-N9	7.74	114.39	108.20
85	Ac	108	C	O4'-C1'-N1	7.74	114.39	108.20
1	Ad	867	A	O4'-C1'-N9	7.74	114.39	108.20
84	Aa	1159	C	O4'-C1'-N1	7.74	114.39	108.20
84	Aa	2012	C	O4'-C1'-N1	7.74	114.39	108.20
84	Aa	2607	U	O4'-C1'-N1	7.74	114.39	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	803	G	O4'-C1'-N9	7.73	114.39	108.20
84	Aa	1082	U	O4'-C1'-N1	7.73	114.39	108.20
86	Ab	40	A	N1-C6-N6	7.73	123.24	118.60
1	Ad	1193	A	O4'-C1'-N9	7.73	114.38	108.20
84	Aa	654	C	O4'-C1'-N1	7.73	114.38	108.20
84	Aa	2406	C	O4'-C1'-N1	7.73	114.38	108.20
84	Aa	2425	U	O4'-C1'-N1	7.73	114.39	108.20
2	Ae	52	G	O4'-C1'-N9	7.73	114.38	108.20
84	Aa	1650	G	C5-C6-O6	-7.73	123.96	128.60
84	Aa	2201	G	N1-C6-O6	7.73	124.54	119.90
1	Ad	644	U	C1'-O4'-C4'	-7.73	103.72	109.90
84	Aa	2186	U	O4'-C1'-N1	7.73	114.38	108.20
84	Aa	3043	U	O4'-C1'-N1	7.73	114.38	108.20
84	Aa	2020	G	N1-C6-O6	7.73	124.53	119.90
85	Ac	55	U	O4'-C1'-N1	7.73	114.38	108.20
84	Aa	79	C	O4'-C1'-N1	7.72	114.38	108.20
84	Aa	939	A	C5-C6-N6	-7.72	117.52	123.70
84	Aa	1870	G	C5-C6-O6	-7.72	123.97	128.60
84	Aa	1778	C	O4'-C1'-N1	7.72	114.38	108.20
84	Aa	2508	U	O4'-C1'-N1	7.72	114.38	108.20
84	Aa	2669	C	O4'-C1'-N1	7.72	114.38	108.20
84	Aa	3233	C	O4'-C1'-N1	7.72	114.38	108.20
84	Aa	1388	C	O4'-C1'-N1	7.72	114.38	108.20
84	Aa	1828	C	O4'-C1'-N1	7.72	114.38	108.20
85	Ac	69	U	O4'-C1'-N1	7.72	114.38	108.20
1	Ad	1648	C	C3'-C2'-C1'	7.72	107.68	101.50
84	Aa	2340	G	N1-C6-O6	7.72	124.53	119.90
84	Aa	481	G	N1-C6-O6	7.72	124.53	119.90
84	Aa	1190	C	O4'-C1'-N1	7.72	114.37	108.20
84	Aa	2575	C	O4'-C1'-N1	7.72	114.37	108.20
1	Ad	829	G	O4'-C1'-N9	7.71	114.37	108.20
1	Ad	1046	G	C1'-O4'-C4'	-7.71	103.73	109.90
84	Aa	2394	G	N1-C6-O6	7.71	124.53	119.90
1	Ad	1610	C	O4'-C1'-N1	7.71	114.37	108.20
84	Aa	276	U	O4'-C1'-N1	7.71	114.37	108.20
84	Aa	1427	C	O4'-C1'-N1	7.71	114.37	108.20
84	Aa	1671	G	N1-C6-O6	7.71	124.53	119.90
85	Ac	5	U	O4'-C1'-N1	7.71	114.37	108.20
1	Ad	985	G	C3'-C2'-C1'	-7.71	95.33	101.50
1	Ad	1363	G	C1'-O4'-C4'	-7.71	103.73	109.90
84	Aa	1940	U	O4'-C1'-N1	7.71	114.37	108.20
84	Aa	2887	C	O4'-C1'-N1	7.71	114.37	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1764	G	O4'-C1'-N9	7.71	114.36	108.20
84	Aa	702	G	N1-C6-O6	7.71	124.52	119.90
86	Ab	98	G	O4'-C1'-N9	7.71	114.36	108.20
1	Ad	832	C	N1-C1'-C2'	7.70	124.01	114.00
84	Aa	2999	G	N1-C6-O6	7.70	124.52	119.90
1	Ad	595	A	O4'-C1'-N9	7.70	114.36	108.20
84	Aa	948	C	O4'-C1'-N1	7.70	114.36	108.20
1	Ad	507	G	C1'-O4'-C4'	-7.69	103.75	109.90
1	Ad	1350	C	O4'-C1'-C2'	7.69	114.52	107.60
84	Aa	699	C	O4'-C1'-N1	7.69	114.35	108.20
84	Aa	1466	U	O4'-C1'-N1	7.69	114.35	108.20
84	Aa	2057	G	O4'-C1'-N9	7.69	114.36	108.20
1	Ad	428	C	N1-C1'-C2'	7.69	124.00	114.00
1	Ad	785	A	O4'-C1'-C2'	-7.69	98.11	105.80
1	Ad	888	U	C1'-O4'-C4'	-7.69	103.75	109.90
84	Aa	978	C	O4'-C1'-N1	7.69	114.35	108.20
84	Aa	1400	C	O4'-C1'-N1	7.69	114.35	108.20
84	Aa	2507	U	O4'-C1'-N1	7.69	114.35	108.20
1	Ad	1275	G	O4'-C1'-N9	7.69	114.35	108.20
1	Ad	1781	U	O4'-C1'-N1	7.69	114.35	108.20
84	Aa	1562	A	C2'-C3'-O3'	-7.69	92.59	109.50
84	Aa	2422	U	O4'-C1'-N1	7.69	114.35	108.20
1	Ad	1601	A	O4'-C1'-N9	7.68	114.35	108.20
84	Aa	502	G	C5-C6-O6	-7.68	123.99	128.60
86	Ab	61	C	O4'-C1'-N1	7.68	114.35	108.20
86	Ab	102	G	O4'-C1'-N9	7.68	114.35	108.20
84	Aa	1145	G	O4'-C1'-N9	7.68	114.35	108.20
84	Aa	1611	G	N1-C6-O6	7.68	124.51	119.90
84	Aa	169	G	N1-C6-O6	7.68	124.51	119.90
1	Ad	869	U	C1'-O4'-C4'	7.68	116.04	109.90
84	Aa	169	G	O4'-C1'-N9	7.68	114.34	108.20
84	Aa	1120	G	N1-C6-O6	7.68	124.51	119.90
84	Aa	3085	C	O4'-C1'-N1	7.68	114.34	108.20
84	Aa	25	U	O4'-C1'-N1	7.68	114.34	108.20
1	Ad	511	U	P-O3'-C3'	7.68	128.91	119.70
84	Aa	2899	A	N1-C6-N6	7.68	123.21	118.60
1	Ad	482	A	O4'-C1'-N9	7.67	114.34	108.20
1	Ad	1795	U	O4'-C1'-N1	7.67	114.34	108.20
84	Aa	542	G	C5-C6-O6	-7.67	124.00	128.60
84	Aa	1737	C	O4'-C1'-N1	7.67	114.34	108.20
84	Aa	2910	C	O4'-C1'-N1	7.67	114.34	108.20
84	Aa	2985	C	C6-N1-C1'	-7.67	111.59	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1317	G	C5-C6-O6	-7.67	124.00	128.60
84	Aa	2538	G	C5-C6-O6	-7.67	124.00	128.60
84	Aa	2879	G	C5-C6-O6	-7.67	124.00	128.60
1	Ad	142	G	O4'-C1'-N9	7.67	114.33	108.20
84	Aa	175	G	N1-C6-O6	7.67	124.50	119.90
84	Aa	1663	G	C5-C6-O6	-7.67	124.00	128.60
84	Aa	2557	C	O4'-C1'-N1	7.67	114.33	108.20
84	Aa	1607	C	O4'-C1'-N1	7.67	114.33	108.20
84	Aa	1894	G	N1-C6-O6	7.67	124.50	119.90
1	Ad	1082	C	C1'-O4'-C4'	-7.66	103.77	109.90
1	Ad	1310	C	N1-C1'-C2'	7.66	123.96	114.00
84	Aa	291	C	O4'-C1'-N1	7.66	114.33	108.20
84	Aa	1265	G	O4'-C1'-N9	7.66	114.33	108.20
84	Aa	1840	C	O4'-C1'-N1	7.66	114.33	108.20
84	Aa	2110	G	N1-C6-O6	7.66	124.50	119.90
84	Aa	2445	U	O4'-C1'-N1	7.66	114.33	108.20
84	Aa	2544	C	O4'-C1'-N1	7.66	114.33	108.20
84	Aa	189	C	O4'-C1'-N1	7.66	114.33	108.20
84	Aa	696	A	O4'-C1'-N9	7.66	114.33	108.20
1	Ad	1112	G	C1'-O4'-C4'	-7.66	103.77	109.90
84	Aa	2702	G	N1-C6-O6	7.66	124.50	119.90
84	Aa	3079	G	O4'-C1'-N9	7.66	114.33	108.20
84	Aa	120	G	N1-C6-O6	7.66	124.50	119.90
84	Aa	1473	U	O4'-C1'-N1	7.66	114.33	108.20
1	Ad	999	G	N9-C1'-C2'	7.66	123.95	114.00
84	Aa	240	U	P-O3'-C3'	7.66	128.89	119.70
84	Aa	309	C	O4'-C1'-N1	7.66	114.32	108.20
84	Aa	2565	C	O4'-C1'-N1	7.66	114.33	108.20
84	Aa	2721	C	O4'-C1'-N1	7.66	114.33	108.20
84	Aa	2158	C	O4'-C1'-N1	7.65	114.32	108.20
84	Aa	3222	G	O4'-C1'-N9	7.65	114.32	108.20
1	Ad	94	A	O4'-C1'-N9	7.65	114.32	108.20
84	Aa	652	C	O4'-C1'-N1	7.65	114.32	108.20
84	Aa	1964	G	N1-C6-O6	7.65	124.49	119.90
84	Aa	892	U	O4'-C1'-N1	7.65	114.32	108.20
84	Aa	58	G	N1-C6-O6	7.65	124.49	119.90
84	Aa	1834	C	O4'-C1'-N1	7.65	114.32	108.20
84	Aa	2007	C	O4'-C1'-N1	7.65	114.32	108.20
85	Ac	72	A	O4'-C1'-N9	7.65	114.32	108.20
1	Ad	157	U	O4'-C1'-N1	7.65	114.32	108.20
1	Ad	1650	G	C1'-O4'-C4'	-7.65	103.78	109.90
84	Aa	300	C	O4'-C1'-N1	7.65	114.32	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2586	C	O4'-C1'-N1	7.65	114.32	108.20
1	Ad	950	U	O4'-C1'-N1	7.64	114.32	108.20
1	Ad	1219	C	C5'-C4'-O4'	7.64	118.27	109.10
2	Ae	30	G	N9-C1'-C2'	7.64	123.94	114.00
84	Aa	190	C	O4'-C1'-N1	7.64	114.32	108.20
84	Aa	352	U	O4'-C1'-N1	7.64	114.31	108.20
84	Aa	1351	C	O4'-C1'-N1	7.64	114.31	108.20
84	Aa	1715	C	O4'-C1'-N1	7.64	114.31	108.20
84	Aa	3321	C	O4'-C1'-N1	7.64	114.31	108.20
86	Ab	91	C	O4'-C1'-N1	7.64	114.31	108.20
84	Aa	167	C	O4'-C1'-N1	7.64	114.31	108.20
84	Aa	679	C	O4'-C1'-N1	7.64	114.31	108.20
84	Aa	1399	C	O4'-C1'-N1	7.64	114.31	108.20
84	Aa	1678	U	O4'-C1'-N1	7.64	114.31	108.20
84	Aa	1900	C	O4'-C1'-N1	7.64	114.31	108.20
84	Aa	2001	U	O4'-C1'-N1	7.64	114.31	108.20
1	Ad	302	C	O4'-C1'-N1	7.64	114.31	108.20
1	Ad	1629	U	O4'-C1'-N1	7.64	114.31	108.20
84	Aa	1768	U	O4'-C1'-N1	7.64	114.31	108.20
1	Ad	1654	C	C3'-C2'-C1'	7.63	107.61	101.50
84	Aa	2547	C	O4'-C1'-N1	7.63	114.31	108.20
84	Aa	2675	G	N1-C6-O6	7.63	124.48	119.90
84	Aa	3206	C	C6-N1-C1'	-7.63	111.64	120.80
1	Ad	494	G	O4'-C1'-N9	7.63	114.31	108.20
84	Aa	1902	G	C5-C6-O6	-7.63	124.02	128.60
84	Aa	1674	A	C5-C6-N1	-7.63	113.89	117.70
84	Aa	457	C	O4'-C1'-N1	7.63	114.30	108.20
84	Aa	492	G	N1-C6-O6	7.62	124.47	119.90
1	Ad	1027	C	C3'-C2'-C1'	7.62	107.60	101.50
84	Aa	264	C	O4'-C1'-N1	7.62	114.30	108.20
84	Aa	237	C	O4'-C1'-N1	7.62	114.30	108.20
86	Ab	16	A	C4-C5-C6	7.62	120.81	117.00
1	Ad	3	C	O4'-C1'-N1	7.62	114.30	108.20
1	Ad	80	C	C3'-C2'-C1'	7.62	107.59	101.50
84	Aa	1536	U	O4'-C1'-N1	7.62	114.29	108.20
84	Aa	2611	G	N1-C6-O6	7.62	124.47	119.90
1	Ad	1454	G	O4'-C1'-N9	7.62	114.29	108.20
84	Aa	308	U	O4'-C1'-N1	7.62	114.29	108.20
84	Aa	2125	A	C5-C6-N1	-7.62	113.89	117.70
1	Ad	338	G	O4'-C1'-N9	7.61	114.29	108.20
1	Ad	1645	C	C1'-O4'-C4'	7.61	115.99	109.90
84	Aa	2074	C	C5'-C4'-C3'	7.61	128.18	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	825	G	N1-C6-O6	7.61	124.47	119.90
84	Aa	847	G	N1-C6-O6	7.61	124.47	119.90
84	Aa	482	C	O4'-C1'-N1	7.61	114.29	108.20
84	Aa	630	C	O4'-C1'-N1	7.61	114.29	108.20
84	Aa	1214	U	O4'-C1'-N1	7.61	114.29	108.20
84	Aa	2092	C	C5'-C4'-C3'	7.61	128.18	116.00
84	Aa	2252	C	O4'-C1'-N1	7.61	114.29	108.20
84	Aa	1325	G	N1-C6-O6	7.61	124.47	119.90
84	Aa	3230	G	O4'-C1'-N9	7.61	114.29	108.20
1	Ad	155	A	C1'-O4'-C4'	7.61	115.99	109.90
1	Ad	395	A	O4'-C1'-N9	7.61	114.29	108.20
84	Aa	841	G	C5-C6-O6	-7.61	124.03	128.60
84	Aa	3105	U	O4'-C1'-N1	7.61	114.29	108.20
1	Ad	1286	U	O4'-C1'-N1	7.61	114.28	108.20
84	Aa	1426	C	O4'-C1'-N1	7.61	114.28	108.20
84	Aa	1533	U	O4'-C1'-N1	7.61	114.28	108.20
85	Ac	100	U	O4'-C1'-N1	7.60	114.28	108.20
84	Aa	1127	U	O4'-C1'-N1	7.60	114.28	108.20
84	Aa	1341	G	N1-C6-O6	7.60	124.46	119.90
84	Aa	1682	C	O4'-C1'-N1	7.60	114.28	108.20
84	Aa	3154	G	P-O3'-C3'	7.60	128.82	119.70
84	Aa	3163	G	O4'-C1'-N9	7.60	114.28	108.20
86	Ab	111	U	C5-C6-N1	7.60	126.50	122.70
1	Ad	1124	G	C1'-O4'-C4'	-7.60	103.82	109.90
1	Ad	1153	C	O4'-C1'-N1	7.60	114.28	108.20
84	Aa	261	C	O4'-C1'-N1	7.60	114.28	108.20
84	Aa	1806	C	O4'-C1'-N1	7.60	114.28	108.20
1	Ad	917	U	O4'-C1'-N1	7.60	114.28	108.20
84	Aa	263	A	O4'-C1'-N9	7.60	114.28	108.20
84	Aa	652	C	N3-C4-C5	-7.60	118.86	121.90
84	Aa	2767	C	O4'-C1'-N1	7.60	114.28	108.20
84	Aa	2827	C	O4'-C1'-N1	7.60	114.28	108.20
84	Aa	3183	G	O4'-C1'-N9	7.60	114.28	108.20
84	Aa	1746	G	C5-C6-O6	-7.60	124.04	128.60
84	Aa	2603	C	O4'-C1'-N1	7.60	114.28	108.20
85	Ac	27	U	O4'-C1'-N1	7.60	114.28	108.20
84	Aa	185	A	N1-C6-N6	7.59	123.16	118.60
84	Aa	495	G	O4'-C1'-N9	7.59	114.28	108.20
84	Aa	1992	U	O4'-C1'-N1	7.59	114.28	108.20
84	Aa	2259	U	O4'-C1'-N1	7.59	114.28	108.20
84	Aa	1068	A	N1-C6-N6	7.59	123.16	118.60
84	Aa	1118	G	C5-C6-O6	-7.59	124.05	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1594	G	C5-C6-O6	-7.59	124.05	128.60
86	Ab	68	G	C5-C6-N1	-7.59	107.70	111.50
1	Ad	1554	G	O4'-C1'-N9	7.59	114.27	108.20
84	Aa	3219	U	O4'-C1'-N1	7.59	114.27	108.20
84	Aa	512	G	C5-C6-O6	-7.59	124.05	128.60
84	Aa	2713	G	O4'-C1'-N9	7.58	114.27	108.20
84	Aa	3070	G	N1-C6-O6	7.58	124.45	119.90
84	Aa	3271	A	C5-C6-N6	-7.58	117.63	123.70
84	Aa	444	C	O4'-C1'-N1	7.58	114.27	108.20
84	Aa	2087	A	C5'-C4'-O4'	7.58	118.20	109.10
84	Aa	2609	G	N1-C6-O6	7.58	124.45	119.90
84	Aa	889	C	O4'-C1'-N1	7.58	114.26	108.20
1	Ad	73	A	C1'-O4'-C4'	7.58	115.96	109.90
1	Ad	1445	C	N1-C1'-C2'	7.58	123.85	114.00
2	Ae	30	G	C1'-O4'-C4'	-7.58	103.84	109.90
84	Aa	2701	G	O4'-C1'-N9	7.58	114.26	108.20
84	Aa	2269	U	O4'-C1'-N1	7.58	114.26	108.20
84	Aa	492	G	C2'-C3'-O3'	7.58	126.17	109.50
84	Aa	1461	U	O4'-C1'-N1	7.58	114.26	108.20
84	Aa	2806	A	N1-C6-N6	7.58	123.14	118.60
1	Ad	27	U	O4'-C1'-N1	7.57	114.26	108.20
1	Ad	1308	G	C1'-O4'-C4'	-7.57	103.84	109.90
68	Ch	74	TYR	CB-CG-CD2	7.57	125.54	121.00
84	Aa	1005	C	O4'-C1'-N1	7.57	114.26	108.20
84	Aa	2832	G	N1-C6-O6	7.57	124.44	119.90
84	Aa	1314	G	N1-C6-O6	7.57	124.44	119.90
84	Aa	1247	G	C5-C6-O6	-7.57	124.06	128.60
86	Ab	22	A	C4-C5-C6	7.57	120.78	117.00
84	Aa	1366	G	N1-C6-O6	7.57	124.44	119.90
84	Aa	3194	G	C5-C6-O6	-7.57	124.06	128.60
1	Ad	1178	C	C3'-C2'-C1'	7.57	107.55	101.50
84	Aa	1109	G	N1-C6-O6	7.57	124.44	119.90
1	Ad	860	A	C1'-O4'-C4'	7.57	115.95	109.90
1	Ad	873	G	O4'-C1'-N9	7.57	114.25	108.20
1	Ad	935	A	O4'-C1'-N9	7.57	114.25	108.20
84	Aa	717	G	O4'-C1'-N9	7.56	114.25	108.20
84	Aa	1122	C	O4'-C1'-N1	7.56	114.25	108.20
84	Aa	2576	C	O4'-C1'-N1	7.56	114.25	108.20
84	Aa	388	G	C5-C6-O6	-7.56	124.06	128.60
84	Aa	2974	G	N1-C6-O6	7.56	124.44	119.90
84	Aa	3263	C	C4'-C3'-O3'	7.56	128.12	113.00
1	Ad	290	C	O4'-C1'-C2'	-7.56	98.24	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	448	C	O4'-C1'-C2'	-7.56	98.24	105.80
84	Aa	2286	A	C5-C6-N6	-7.56	117.65	123.70
84	Aa	2823	C	O4'-C1'-N1	7.56	114.25	108.20
1	Ad	1051	G	O4'-C1'-N9	7.56	114.25	108.20
84	Aa	338	C	O4'-C1'-N1	7.56	114.25	108.20
84	Aa	1347	U	O4'-C1'-N1	7.56	114.25	108.20
1	Ad	1140	U	O4'-C1'-N1	7.55	114.24	108.20
84	Aa	215	U	O4'-C1'-N1	7.55	114.24	108.20
84	Aa	314	C	O4'-C1'-N1	7.55	114.24	108.20
84	Aa	595	C	O4'-C1'-N1	7.55	114.24	108.20
84	Aa	1141	U	O4'-C1'-N1	7.55	114.24	108.20
84	Aa	128	C	O4'-C1'-N1	7.55	114.24	108.20
84	Aa	726	C	O4'-C1'-N1	7.55	114.24	108.20
1	Ad	12	U	C1'-O4'-C4'	-7.55	103.86	109.90
84	Aa	2330	C	O4'-C1'-N1	7.55	114.24	108.20
84	Aa	3091	U	O4'-C1'-N1	7.55	114.24	108.20
84	Aa	983	U	O4'-C1'-N1	7.55	114.24	108.20
84	Aa	1581	C	O4'-C1'-N1	7.55	114.24	108.20
84	Aa	2421	C	O4'-C1'-N1	7.55	114.24	108.20
84	Aa	1039	G	N1-C6-O6	7.55	124.43	119.90
1	Ad	983	A	C1'-O4'-C4'	7.55	115.94	109.90
1	Ad	1453	U	O4'-C1'-N1	7.55	114.24	108.20
84	Aa	1440	C	O4'-C1'-N1	7.55	114.24	108.20
84	Aa	3300	C	O4'-C1'-N1	7.55	114.24	108.20
84	Aa	1995	U	P-O3'-C3'	7.54	128.75	119.70
84	Aa	3346	C	O4'-C1'-N1	7.54	114.24	108.20
80	Cf	107	TYR	CB-CG-CD1	7.54	125.53	121.00
84	Aa	2580	C	O4'-C1'-N1	7.54	114.23	108.20
84	Aa	2828	U	O4'-C1'-N1	7.54	114.23	108.20
85	Ac	99	C	O4'-C1'-N1	7.54	114.23	108.20
84	Aa	414	G	N1-C6-O6	7.54	124.42	119.90
84	Aa	1551	C	O4'-C1'-N1	7.54	114.23	108.20
84	Aa	2844	U	O4'-C1'-N1	7.54	114.23	108.20
1	Ad	1294	U	O4'-C1'-N1	7.54	114.23	108.20
1	Ad	1471	C	N1-C1'-C2'	7.54	123.80	114.00
2	Ae	25	U	O4'-C1'-N1	7.54	114.23	108.20
1	Ad	487	A	O4'-C1'-N9	7.54	114.23	108.20
84	Aa	215	U	P-O3'-C3'	7.54	128.74	119.70
84	Aa	824	U	O4'-C1'-N1	7.54	114.23	108.20
84	Aa	285	G	O4'-C1'-N9	7.53	114.23	108.20
84	Aa	618	G	C5-C6-O6	-7.53	124.08	128.60
84	Aa	716	A	C4-C5-C6	7.53	120.77	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1094	G	C5-C6-O6	-7.53	124.08	128.60
84	Aa	2498	C	O4'-C1'-N1	7.53	114.23	108.20
84	Aa	1145	G	C5-C6-O6	-7.53	124.08	128.60
84	Aa	2327	U	O4'-C1'-N1	7.53	114.23	108.20
1	Ad	1634	U	O4'-C1'-N1	7.53	114.22	108.20
84	Aa	753	G	C5-C6-O6	-7.53	124.08	128.60
1	Ad	552	G	O4'-C1'-N9	7.53	114.22	108.20
1	Ad	860	A	P-O3'-C3'	-7.53	110.67	119.70
84	Aa	2009	C	O4'-C1'-N1	7.53	114.22	108.20
84	Aa	2556	G	O4'-C1'-N9	7.53	114.22	108.20
84	Aa	632	C	O4'-C1'-N1	7.53	114.22	108.20
1	Ad	130	A	O4'-C1'-C2'	-7.52	98.28	105.80
1	Ad	1670	G	O4'-C1'-C2'	7.52	114.37	107.60
84	Aa	1928	A	C4-C5-C6	7.52	120.76	117.00
84	Aa	2782	G	N1-C6-O6	7.52	124.41	119.90
1	Ad	757	G	C1'-O4'-C4'	-7.52	103.89	109.90
1	Ad	953	G	O4'-C1'-N9	7.52	114.22	108.20
1	Ad	1801	A	C1'-O4'-C4'	7.52	115.92	109.90
84	Aa	1028	G	N1-C6-O6	7.52	124.41	119.90
84	Aa	1546	G	N1-C6-O6	7.52	124.41	119.90
85	Ac	10	G	N1-C6-O6	7.52	124.41	119.90
1	Ad	501	U	O4'-C1'-N1	7.52	114.21	108.20
84	Aa	938	U	O4'-C1'-N1	7.52	114.21	108.20
84	Aa	1363	C	O4'-C1'-N1	7.52	114.21	108.20
85	Ac	8	C	O4'-C1'-N1	7.52	114.21	108.20
84	Aa	372	A	O4'-C1'-N9	7.51	114.21	108.20
84	Aa	2577	G	C5-C6-O6	-7.51	124.09	128.60
84	Aa	3153	U	C2-N1-C1'	7.51	126.72	117.70
84	Aa	8	C	C2-N1-C1'	7.51	127.06	118.80
84	Aa	1146	A	C5-C6-N6	-7.51	117.69	123.70
84	Aa	2716	U	O4'-C1'-N1	7.51	114.21	108.20
84	Aa	3024	U	O4'-C1'-N1	7.51	114.21	108.20
84	Aa	2503	A	C4-C5-C6	7.51	120.75	117.00
84	Aa	3121	C	O4'-C1'-N1	7.51	114.21	108.20
84	Aa	2957	U	O4'-C1'-N1	7.51	114.21	108.20
1	Ad	394	G	C1'-O4'-C4'	7.51	115.91	109.90
1	Ad	990	G	O4'-C1'-N9	7.51	114.20	108.20
1	Ad	1145	G	O4'-C1'-N9	7.51	114.20	108.20
84	Aa	1326	C	O4'-C1'-N1	7.51	114.21	108.20
1	Ad	1512	C	N1-C1'-C2'	7.50	123.76	114.00
84	Aa	658	C	O4'-C1'-N1	7.50	114.20	108.20
84	Aa	896	C	O4'-C1'-N1	7.50	114.20	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2584	U	O4'-C1'-N1	7.50	114.20	108.20
1	Ad	1231	A	P-O3'-C3'	7.50	128.70	119.70
1	Ad	1500	A	C1'-O4'-C4'	7.50	115.90	109.90
84	Aa	617	C	O4'-C1'-N1	7.50	114.20	108.20
84	Aa	1409	G	N1-C6-O6	7.50	124.40	119.90
84	Aa	2178	G	C5'-C4'-O4'	7.50	118.10	109.10
84	Aa	2262	C	O4'-C1'-N1	7.50	114.20	108.20
84	Aa	1288	C	O4'-C1'-N1	7.50	114.20	108.20
1	Ad	114	U	C3'-C2'-C1'	7.50	107.50	101.50
1	Ad	755	U	N1-C1'-C2'	7.50	123.75	114.00
84	Aa	430	G	C5-C6-O6	-7.50	124.10	128.60
84	Aa	1130	G	C5-C6-O6	-7.50	124.10	128.60
84	Aa	2295	G	C5-C6-O6	-7.50	124.10	128.60
84	Aa	808	G	C5-C6-O6	-7.49	124.10	128.60
1	Ad	329	G	O4'-C1'-N9	7.49	114.19	108.20
84	Aa	449	G	C5-C6-O6	-7.49	124.11	128.60
84	Aa	1535	C	O4'-C1'-N1	7.49	114.19	108.20
1	Ad	23	G	O4'-C1'-N9	7.49	114.19	108.20
84	Aa	1865	C	O4'-C1'-N1	7.49	114.19	108.20
84	Aa	3290	C	O4'-C1'-N1	7.49	114.19	108.20
2	Ae	47	U	N1-C1'-C2'	-7.49	103.76	112.00
84	Aa	1850	C	O4'-C1'-N1	7.49	114.19	108.20
84	Aa	2157	C	O4'-C1'-N1	7.49	114.19	108.20
1	Ad	369	G	C1'-O4'-C4'	7.49	115.89	109.90
1	Ad	373	U	C5'-C4'-O4'	7.49	118.08	109.10
1	Ad	499	A	O4'-C1'-N9	7.49	114.19	108.20
84	Aa	375	G	C5-C6-O6	-7.49	124.11	128.60
84	Aa	464	G	N1-C6-O6	7.49	124.39	119.90
84	Aa	739	C	O4'-C1'-N1	7.49	114.19	108.20
84	Aa	2155	G	N1-C6-O6	7.49	124.39	119.90
84	Aa	339	G	C5-C6-O6	-7.48	124.11	128.60
84	Aa	1406	C	O4'-C1'-N1	7.48	114.19	108.20
84	Aa	2954	G	N1-C6-O6	7.48	124.39	119.90
84	Aa	1981	U	O4'-C1'-N1	7.48	114.19	108.20
1	Ad	648	C	O4'-C1'-C2'	-7.48	98.32	105.80
84	Aa	1428	G	C5-C6-O6	-7.48	124.11	128.60
84	Aa	3137	G	N1-C6-O6	7.48	124.39	119.90
84	Aa	815	G	N1-C6-O6	7.48	124.39	119.90
84	Aa	1503	G	N1-C6-O6	7.48	124.39	119.90
1	Ad	1063	U	O4'-C1'-N1	7.48	114.18	108.20
84	Aa	168	A	C5-C6-N6	-7.48	117.72	123.70
84	Aa	1344	A	O4'-C1'-N9	7.48	114.18	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1560	A	N9-C1'-C2'	7.48	123.72	114.00
84	Aa	2147	U	O4'-C1'-N1	7.48	114.18	108.20
84	Aa	1589	G	O4'-C1'-N9	7.48	114.18	108.20
1	Ad	1409	G	O4'-C1'-C2'	7.47	114.33	107.60
84	Aa	659	C	O4'-C1'-N1	7.47	114.18	108.20
84	Aa	1178	C	O4'-C1'-N1	7.47	114.18	108.20
84	Aa	2637	U	O4'-C1'-N1	7.47	114.18	108.20
84	Aa	3159	C	O4'-C1'-N1	7.47	114.18	108.20
86	Ab	56	G	N1-C6-O6	7.47	124.38	119.90
86	Ab	80	A	C4-C5-C6	7.47	120.74	117.00
84	Aa	1750	A	O4'-C1'-N9	7.47	114.18	108.20
84	Aa	895	U	O4'-C1'-N1	7.47	114.17	108.20
84	Aa	1211	G	C5-C6-O6	-7.47	124.12	128.60
84	Aa	3241	C	O4'-C1'-N1	7.47	114.17	108.20
1	Ad	146	A	C1'-O4'-C4'	7.47	115.88	109.90
84	Aa	2710	C	O4'-C1'-N1	7.47	114.17	108.20
84	Aa	425	G	C5-C6-O6	-7.47	124.12	128.60
84	Aa	628	C	O4'-C1'-N1	7.47	114.17	108.20
84	Aa	1125	U	O4'-C1'-N1	7.47	114.17	108.20
84	Aa	3218	C	O4'-C1'-N1	7.47	114.17	108.20
1	Ad	1192	G	O4'-C1'-N9	7.46	114.17	108.20
84	Aa	537	U	P-O3'-C3'	7.46	128.66	119.70
84	Aa	1695	C	O4'-C1'-N1	7.46	114.17	108.20
84	Aa	2557	C	C2-N1-C1'	7.46	127.01	118.80
1	Ad	1194	C	O4'-C1'-N1	7.46	114.17	108.20
1	Ad	67	G	C3'-C2'-C1'	7.46	107.47	101.50
1	Ad	1441	C	C3'-C2'-C1'	7.46	107.47	101.50
84	Aa	1249	A	O4'-C1'-N9	7.46	114.17	108.20
1	Ad	390	G	O4'-C1'-N9	7.46	114.17	108.20
1	Ad	1266	U	O4'-C1'-N1	7.46	114.17	108.20
84	Aa	1045	U	O4'-C1'-N1	7.46	114.17	108.20
84	Aa	3348	G	O4'-C1'-N9	7.46	114.17	108.20
84	Aa	2234	G	C5-C6-O6	-7.46	124.13	128.60
1	Ad	1082	C	N1-C1'-C2'	7.46	123.69	114.00
84	Aa	1489	G	N1-C6-O6	7.46	124.37	119.90
84	Aa	2195	U	O4'-C1'-N1	7.46	114.16	108.20
84	Aa	2519	U	O4'-C1'-N1	7.45	114.16	108.20
84	Aa	547	C	O4'-C1'-N1	7.45	114.16	108.20
1	Ad	93	A	C1'-O4'-C4'	7.45	115.86	109.90
84	Aa	1056	U	O4'-C1'-N1	7.45	114.16	108.20
84	Aa	2358	C	O4'-C1'-N1	7.45	114.16	108.20
84	Aa	2	C	O4'-C1'-N1	7.45	114.16	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	166	U	O4'-C1'-N1	7.45	114.16	108.20
84	Aa	344	C	O4'-C1'-N1	7.45	114.16	108.20
84	Aa	2167	G	C1'-O4'-C4'	-7.45	103.94	109.90
84	Aa	2717	G	O4'-C1'-N9	7.45	114.16	108.20
84	Aa	1164	G	N1-C6-O6	7.45	124.37	119.90
84	Aa	2521	C	O4'-C1'-N1	7.45	114.16	108.20
1	Ad	804	C	N1-C1'-C2'	7.45	123.68	114.00
1	Ad	819	U	O4'-C1'-N1	7.45	114.16	108.20
84	Aa	2063	U	O4'-C1'-N1	7.45	114.16	108.20
84	Aa	2072	U	O4'-C1'-N1	7.45	114.16	108.20
84	Aa	2600	U	O4'-C1'-N1	7.45	114.16	108.20
84	Aa	236	A	N1-C6-N6	7.44	123.07	118.60
1	Ad	1467	C	O4'-C1'-N1	7.44	114.15	108.20
84	Aa	893	C	O4'-C1'-N1	7.44	114.15	108.20
84	Aa	1781	C	O4'-C1'-N1	7.44	114.15	108.20
84	Aa	1218	U	O4'-C1'-N1	7.44	114.15	108.20
4	BY	41	SER	N-CA-CB	7.44	121.66	110.50
84	Aa	724	A	O4'-C1'-N9	7.44	114.15	108.20
1	Ad	163	G	O4'-C1'-N9	7.44	114.15	108.20
1	Ad	701	C	O4'-C1'-N1	7.44	114.15	108.20
84	Aa	518	G	N1-C6-O6	7.44	124.36	119.90
84	Aa	556	U	O4'-C1'-N1	7.44	114.15	108.20
84	Aa	2476	G	C4'-C3'-O3'	-7.44	93.78	109.40
1	Ad	1723	G	C1'-O4'-C4'	-7.43	103.95	109.90
84	Aa	1252	C	O4'-C1'-N1	7.43	114.15	108.20
84	Aa	2069	G	C5-C6-O6	-7.43	124.14	128.60
84	Aa	2824	U	O4'-C1'-N1	7.43	114.15	108.20
84	Aa	2965	C	O4'-C1'-N1	7.43	114.15	108.20
84	Aa	1063	G	N1-C6-O6	7.43	124.36	119.90
84	Aa	1657	C	O4'-C1'-N1	7.43	114.15	108.20
1	Ad	1234	A	O4'-C1'-N9	7.43	114.14	108.20
3	Af	18	C	C3'-C2'-C1'	7.43	107.44	101.50
84	Aa	1957	G	C2'-C3'-O3'	-7.43	93.15	109.50
84	Aa	2244	G	C5-C6-O6	-7.43	124.14	128.60
1	Ad	1334	G	C1'-O4'-C4'	7.43	115.84	109.90
84	Aa	297	G	C5-C6-O6	-7.43	124.14	128.60
84	Aa	641	C	N3-C4-C5	-7.43	118.93	121.90
84	Aa	930	C	O4'-C1'-N1	7.43	114.14	108.20
84	Aa	2237	A	C5-C6-N6	-7.43	117.76	123.70
84	Aa	1724	C	O4'-C1'-N1	7.43	114.14	108.20
1	Ad	1665	U	C3'-C2'-C1'	7.42	107.44	101.50
84	Aa	318	G	O4'-C1'-N9	7.42	114.14	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1481	C	O4'-C1'-N1	7.42	114.14	108.20
84	Aa	2236	U	P-O3'-C3'	7.42	128.61	119.70
86	Ab	57	C	C5-C4-N4	-7.42	115.00	120.20
86	Ab	80	A	C5-C6-N6	-7.42	117.76	123.70
84	Aa	900	C	O4'-C1'-N1	7.42	114.14	108.20
84	Aa	2454	U	O4'-C1'-N1	7.42	114.14	108.20
84	Aa	2842	C	O4'-C1'-N1	7.42	114.14	108.20
1	Ad	410	U	O4'-C1'-N1	7.42	114.14	108.20
1	Ad	1128	C	C5'-C4'-O4'	7.42	118.01	109.10
84	Aa	774	A	C5-C6-N6	-7.42	117.76	123.70
84	Aa	799	U	O4'-C1'-N1	7.42	114.14	108.20
84	Aa	1274	A	C4-C5-C6	7.42	120.71	117.00
84	Aa	2560	C	O4'-C1'-N1	7.42	114.14	108.20
84	Aa	2686	U	O4'-C1'-N1	7.42	114.14	108.20
84	Aa	1008	U	O4'-C1'-N1	7.42	114.14	108.20
84	Aa	1027	C	O4'-C1'-N1	7.42	114.14	108.20
84	Aa	1316	C	C6-N1-C2	-7.42	117.33	120.30
1	Ad	1168	A	C3'-C2'-C1'	7.42	107.44	101.50
84	Aa	133	G	C5-C6-O6	-7.42	124.15	128.60
84	Aa	506	U	O4'-C1'-N1	7.42	114.14	108.20
84	Aa	1311	G	N1-C6-O6	7.42	124.35	119.90
84	Aa	2768	C	O4'-C1'-N1	7.42	114.14	108.20
84	Aa	541	C	O4'-C1'-N1	7.42	114.13	108.20
84	Aa	3352	C	O4'-C1'-N1	7.42	114.13	108.20
84	Aa	328	G	C5-C6-O6	-7.42	124.15	128.60
84	Aa	353	A	C5-C6-N6	-7.41	117.77	123.70
84	Aa	408	U	O4'-C1'-N1	7.41	114.13	108.20
84	Aa	2261	U	O4'-C1'-N1	7.41	114.13	108.20
84	Aa	3161	C	O4'-C1'-N1	7.41	114.13	108.20
85	Ac	150	G	C5-C6-O6	-7.41	124.15	128.60
84	Aa	289	C	O4'-C1'-N1	7.41	114.13	108.20
86	Ab	71	A	N1-C6-N6	7.41	123.05	118.60
84	Aa	1633	C	O4'-C1'-N1	7.41	114.13	108.20
85	Ac	42	G	N1-C6-O6	7.41	124.35	119.90
1	Ad	215	A	N9-C1'-C2'	-7.41	103.85	112.00
84	Aa	3248	G	C5-C6-O6	-7.41	124.16	128.60
84	Aa	3202	G	N1-C6-O6	7.41	124.34	119.90
84	Aa	777	G	O4'-C1'-N9	7.41	114.12	108.20
84	Aa	2729	C	C2-N1-C1'	7.41	126.94	118.80
85	Ac	4	C	O4'-C1'-N1	7.41	114.12	108.20
84	Aa	1272	G	C5-C6-O6	-7.40	124.16	128.60
84	Aa	2071	U	O4'-C1'-N1	7.40	114.12	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2951	U	O4'-C1'-N1	7.40	114.12	108.20
84	Aa	3294	U	P-O3'-C3'	7.40	128.59	119.70
1	Ad	413	C	C3'-C2'-C1'	7.40	107.42	101.50
84	Aa	1201	C	O4'-C1'-N1	7.40	114.12	108.20
84	Aa	3138	C	N3-C4-C5	-7.40	118.94	121.90
84	Aa	1442	U	O4'-C1'-N1	7.40	114.12	108.20
84	Aa	2106	U	O4'-C1'-N1	7.40	114.12	108.20
84	Aa	2431	U	O4'-C1'-N1	7.40	114.12	108.20
84	Aa	92	C	O4'-C1'-N1	7.40	114.12	108.20
84	Aa	2211	G	N1-C6-O6	7.40	124.34	119.90
1	Ad	1196	C	C3'-C2'-C1'	7.40	107.42	101.50
84	Aa	3191	U	O4'-C1'-N1	7.40	114.12	108.20
1	Ad	1209	C	O4'-C1'-N1	7.40	114.12	108.20
84	Aa	1262	U	O4'-C1'-N1	7.39	114.11	108.20
84	Aa	1575	G	C5-C6-O6	-7.39	124.16	128.60
1	Ad	391	A	N9-C1'-C2'	-7.39	103.87	112.00
84	Aa	246	C	O4'-C1'-N1	7.39	114.11	108.20
84	Aa	1324	C	O4'-C1'-N1	7.39	114.11	108.20
84	Aa	1384	G	C5-C6-O6	-7.39	124.17	128.60
84	Aa	1760	G	O4'-C1'-N9	7.39	114.11	108.20
84	Aa	1935	G	C5-C6-O6	-7.39	124.17	128.60
84	Aa	2035	G	N1-C6-O6	7.39	124.34	119.90
86	Ab	3	A	C5-C6-N1	-7.39	114.00	117.70
1	Ad	1104	U	O4'-C1'-N1	7.39	114.11	108.20
1	Ad	1189	U	N1-C1'-C2'	7.39	123.61	114.00
84	Aa	1049	C	O4'-C1'-N1	7.39	114.11	108.20
84	Aa	867	G	C5-C6-O6	-7.39	124.17	128.60
84	Aa	2209	A	N1-C6-N6	7.39	123.03	118.60
1	Ad	105	A	O4'-C1'-C2'	-7.39	98.41	105.80
84	Aa	2194	G	C5-C6-O6	-7.39	124.17	128.60
84	Aa	3359	C	O4'-C1'-N1	7.39	114.11	108.20
1	Ad	1734	U	N1-C1'-C2'	7.38	123.60	114.00
2	Ae	10	G	C3'-C2'-C1'	7.38	107.41	101.50
84	Aa	2030	U	O4'-C1'-N1	7.38	114.11	108.20
84	Aa	2226	C	O4'-C1'-N1	7.38	114.11	108.20
84	Aa	2285	C	O4'-C1'-N1	7.38	114.11	108.20
84	Aa	2939	G	N1-C6-O6	7.38	124.33	119.90
1	Ad	900	G	C1'-O4'-C4'	-7.38	103.99	109.90
84	Aa	845	G	C5-C6-O6	-7.38	124.17	128.60
84	Aa	1620	U	O4'-C1'-N1	7.38	114.11	108.20
84	Aa	3177	A	C4-C5-C6	7.38	120.69	117.00
84	Aa	3228	C	O4'-C1'-N1	7.38	114.11	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1103	U	O4'-C1'-N1	7.38	114.11	108.20
84	Aa	1556	G	O4'-C1'-N9	7.38	114.11	108.20
84	Aa	1623	C	O4'-C1'-N1	7.38	114.11	108.20
1	Ad	1108	U	O4'-C1'-N1	7.38	114.10	108.20
84	Aa	318	G	C5-C6-O6	-7.38	124.17	128.60
1	Ad	148	C	O4'-C1'-N1	7.38	114.10	108.20
84	Aa	1952	U	O4'-C1'-N1	7.38	114.10	108.20
84	Aa	3133	C	O4'-C1'-N1	7.38	114.10	108.20
84	Aa	2785	U	O4'-C1'-N1	7.38	114.10	108.20
84	Aa	2845	U	O4'-C1'-N1	7.37	114.10	108.20
84	Aa	2970	G	C5-C6-O6	-7.37	124.18	128.60
84	Aa	3099	G	C5-C6-O6	-7.37	124.18	128.60
1	Ad	1193	A	C1'-O4'-C4'	7.37	115.80	109.90
84	Aa	2752	G	C5-C6-O6	-7.37	124.18	128.60
84	Aa	525	A	C5-C6-N6	-7.37	117.80	123.70
84	Aa	857	G	C5-C6-O6	-7.37	124.18	128.60
84	Aa	2175	A	O4'-C1'-N9	7.37	114.09	108.20
85	Ac	68	G	O4'-C1'-N9	7.37	114.09	108.20
1	Ad	271	C	O4'-C1'-C2'	-7.37	98.43	105.80
1	Ad	1669	U	O4'-C1'-N1	7.37	114.09	108.20
84	Aa	329	G	N1-C6-O6	7.37	124.32	119.90
84	Aa	465	C	O4'-C1'-N1	7.36	114.09	108.20
84	Aa	867	G	O4'-C1'-N9	7.36	114.09	108.20
84	Aa	1193	A	C4-C5-C6	7.36	120.68	117.00
84	Aa	2090	G	C5-C6-O6	-7.36	124.18	128.60
84	Aa	1062	G	N1-C6-O6	7.36	124.32	119.90
84	Aa	3185	G	C5-C6-O6	-7.36	124.18	128.60
84	Aa	1597	U	O4'-C1'-N1	7.36	114.09	108.20
84	Aa	1798	C	O3'-P-O5'	-7.36	90.02	104.00
84	Aa	1626	U	O4'-C1'-N1	7.36	114.09	108.20
84	Aa	2062	U	O4'-C1'-N1	7.36	114.09	108.20
84	Aa	110	C	O4'-C1'-N1	7.36	114.08	108.20
84	Aa	682	G	C5-C6-O6	-7.36	124.19	128.60
84	Aa	2221	U	O4'-C1'-N1	7.36	114.08	108.20
84	Aa	2636	U	O4'-C1'-N1	7.35	114.08	108.20
84	Aa	1604	U	P-O3'-C3'	7.35	128.52	119.70
84	Aa	1534	C	O4'-C1'-N1	7.35	114.08	108.20
1	Ad	834	A	C2'-C3'-O3'	7.35	125.67	109.50
84	Aa	514	G	C5-C6-O6	-7.35	124.19	128.60
84	Aa	991	C	O4'-C1'-N1	7.35	114.08	108.20
84	Aa	2771	U	O4'-C1'-N1	7.35	114.08	108.20
84	Aa	3267	U	O4'-C1'-N1	7.35	114.08	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	213	G	C5-C6-O6	-7.35	124.19	128.60
84	Aa	1462	C	O4'-C1'-N1	7.35	114.08	108.20
2	Ae	61	C	N1-C1'-C2'	7.35	123.55	114.00
1	Ad	348	A	O4'-C1'-N9	7.34	114.08	108.20
84	Aa	940	G	C5-C6-O6	-7.34	124.19	128.60
84	Aa	2571	C	O4'-C1'-N1	7.34	114.08	108.20
84	Aa	1137	G	N1-C6-O6	7.34	124.31	119.90
1	Ad	1409	G	N9-C1'-C2'	7.34	123.54	114.00
1	Ad	1735	C	C3'-C2'-C1'	7.34	107.37	101.50
84	Aa	302	G	C5-C6-O6	-7.34	124.20	128.60
84	Aa	1624	G	N1-C6-O6	7.34	124.30	119.90
84	Aa	1866	C	O4'-C1'-N1	7.34	114.07	108.20
84	Aa	2690	G	C5-C6-O6	-7.34	124.20	128.60
85	Ac	87	G	N1-C6-O6	7.34	124.30	119.90
86	Ab	58	G	N1-C6-O6	7.34	124.30	119.90
1	Ad	356	G	O4'-C1'-C2'	-7.34	98.46	105.80
1	Ad	1574	U	O4'-C1'-C2'	-7.34	98.46	105.80
84	Aa	647	U	O4'-C1'-N1	7.34	114.07	108.20
84	Aa	1505	G	N1-C6-O6	7.34	124.30	119.90
84	Aa	1774	G	C5-C6-O6	-7.34	124.20	128.60
1	Ad	1616	U	O4'-C1'-N1	7.34	114.07	108.20
84	Aa	1124	U	O4'-C1'-N1	7.34	114.07	108.20
84	Aa	1446	G	C5-C6-O6	-7.34	124.20	128.60
84	Aa	2303	C	O4'-C1'-N1	7.34	114.07	108.20
1	Ad	1685	U	N1-C1'-C2'	7.33	123.53	114.00
84	Aa	974	G	C5-C6-O6	-7.33	124.20	128.60
84	Aa	1093	U	O4'-C1'-N1	7.33	114.07	108.20
84	Aa	1216	G	O4'-C1'-N9	7.33	114.07	108.20
84	Aa	1270	G	N1-C6-O6	7.33	124.30	119.90
1	Ad	1514	G	O4'-C1'-N9	7.33	114.06	108.20
84	Aa	2652	G	C5-C6-O6	-7.33	124.20	128.60
84	Aa	2854	C	N3-C4-N4	7.33	123.13	118.00
84	Aa	766	C	O4'-C1'-N1	7.33	114.06	108.20
84	Aa	2168	C	O4'-C1'-N1	7.33	114.06	108.20
84	Aa	3035	C	O4'-C1'-N1	7.33	114.06	108.20
84	Aa	1337	C	O4'-C1'-N1	7.33	114.06	108.20
84	Aa	1890	C	O4'-C1'-N1	7.33	114.06	108.20
84	Aa	2326	U	O4'-C1'-N1	7.33	114.06	108.20
1	Ad	1255	U	O4'-C1'-C2'	-7.32	98.48	105.80
84	Aa	2060	C	O4'-C1'-N1	7.32	114.06	108.20
84	Aa	3036	C	O4'-C1'-N1	7.32	114.06	108.20
84	Aa	360	G	N1-C6-O6	7.32	124.29	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	695	G	C5-C6-O6	-7.32	124.21	128.60
84	Aa	2280	C	O4'-C1'-N1	7.32	114.06	108.20
84	Aa	636	C	O4'-C1'-N1	7.32	114.06	108.20
84	Aa	2682	A	O4'-C1'-N9	7.32	114.06	108.20
85	Ac	130	G	C5-C6-O6	-7.32	124.21	128.60
84	Aa	1390	G	C5-C6-O6	-7.32	124.21	128.60
84	Aa	2202	A	O4'-C1'-N9	7.32	114.06	108.20
1	Ad	1458	U	O4'-C1'-N1	7.32	114.05	108.20
84	Aa	177	C	O4'-C1'-N1	7.32	114.06	108.20
84	Aa	1205	C	O4'-C1'-N1	7.32	114.06	108.20
86	Ab	81	G	C5-C6-N1	-7.32	107.84	111.50
84	Aa	141	C	O4'-C1'-N1	7.32	114.05	108.20
84	Aa	716	A	O4'-C1'-N9	7.32	114.05	108.20
86	Ab	110	G	O4'-C1'-N9	7.32	114.05	108.20
84	Aa	773	G	C5-C6-O6	-7.31	124.21	128.60
84	Aa	2633	C	O4'-C1'-N1	7.31	114.05	108.20
84	Aa	2717	G	C5-C6-O6	-7.31	124.21	128.60
84	Aa	623	G	O4'-C1'-N9	7.31	114.05	108.20
84	Aa	3082	G	O4'-C1'-N9	7.31	114.05	108.20
1	Ad	349	U	N1-C1'-C2'	-7.31	103.96	112.00
84	Aa	1180	C	C2-N1-C1'	7.31	126.84	118.80
85	Ac	20	U	O4'-C1'-N1	7.31	114.05	108.20
84	Aa	87	A	C5-C6-N1	-7.31	114.05	117.70
84	Aa	106	G	N1-C6-O6	7.31	124.28	119.90
84	Aa	2809	U	O4'-C1'-N1	7.31	114.05	108.20
84	Aa	922	U	O4'-C1'-N1	7.30	114.04	108.20
84	Aa	1371	G	O4'-C1'-N9	7.30	114.04	108.20
84	Aa	2925	U	O4'-C1'-N1	7.30	114.04	108.20
1	Ad	398	C	N1-C1'-C2'	7.30	123.49	114.00
1	Ad	1801	A	O4'-C1'-N9	7.30	114.04	108.20
84	Aa	568	C	O4'-C1'-N1	7.30	114.04	108.20
84	Aa	662	G	C5-C6-O6	-7.30	124.22	128.60
1	Ad	1470	G	O4'-C1'-N9	7.30	114.04	108.20
84	Aa	3100	C	O4'-C1'-N1	7.30	114.04	108.20
86	Ab	80	A	C5-C6-N1	-7.30	114.05	117.70
84	Aa	826	C	O4'-C1'-N1	7.30	114.04	108.20
84	Aa	2106	U	P-O3'-C3'	7.30	128.46	119.70
84	Aa	2230	C	O4'-C1'-N1	7.30	114.04	108.20
85	Ac	4	C	N3-C4-N4	7.30	123.11	118.00
85	Ac	43	A	O4'-C1'-N9	7.30	114.04	108.20
84	Aa	468	U	O4'-C1'-N1	7.30	114.04	108.20
84	Aa	980	C	O4'-C1'-N1	7.30	114.04	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2934	C	O4'-C1'-N1	7.30	114.04	108.20
86	Ab	48	G	O4'-C1'-N9	7.30	114.04	108.20
84	Aa	139	U	O4'-C1'-N1	7.29	114.03	108.20
84	Aa	271	G	N1-C6-O6	7.29	124.28	119.90
84	Aa	1072	C	O4'-C1'-N1	7.29	114.03	108.20
1	Ad	1623	C	O4'-C1'-C2'	-7.29	98.51	105.80
1	Ad	1709	U	C3'-C2'-C1'	7.29	107.33	101.50
84	Aa	526	A	P-O3'-C3'	7.29	128.45	119.70
84	Aa	770	U	O4'-C1'-N1	7.29	114.03	108.20
84	Aa	3097	G	O4'-C1'-N9	7.29	114.03	108.20
86	Ab	55	A	C5-C6-N6	-7.29	117.87	123.70
84	Aa	36	U	O4'-C1'-N1	7.29	114.03	108.20
84	Aa	525	A	O4'-C1'-N9	7.29	114.03	108.20
84	Aa	1339	C	O4'-C1'-N1	7.29	114.03	108.20
84	Aa	517	G	C5-C6-O6	-7.29	124.23	128.60
84	Aa	1763	C	O4'-C1'-N1	7.29	114.03	108.20
84	Aa	2929	C	O4'-C1'-N1	7.29	114.03	108.20
1	Ad	996	G	O4'-C1'-N9	7.28	114.03	108.20
84	Aa	254	G	O4'-C1'-N9	7.28	114.03	108.20
84	Aa	2121	U	O4'-C1'-N1	7.28	114.03	108.20
84	Aa	267	G	O4'-C1'-N9	7.28	114.03	108.20
84	Aa	1701	G	C5-C6-O6	-7.28	124.23	128.60
84	Aa	2392	G	C5-C6-O6	-7.28	124.23	128.60
84	Aa	35	U	O4'-C1'-N1	7.28	114.02	108.20
84	Aa	2784	U	O4'-C1'-N1	7.28	114.03	108.20
84	Aa	3377	G	C5-C6-O6	-7.28	124.23	128.60
84	Aa	651	A	P-O3'-C3'	7.28	128.44	119.70
84	Aa	2035	G	O4'-C1'-N9	7.28	114.02	108.20
84	Aa	1504	U	O4'-C1'-N1	7.28	114.02	108.20
84	Aa	1565	G	P-O5'-C5'	7.28	132.54	120.90
84	Aa	1938	U	O4'-C1'-N1	7.28	114.02	108.20
84	Aa	2941	G	O4'-C1'-N9	7.28	114.02	108.20
84	Aa	72	A	O4'-C1'-N9	7.28	114.02	108.20
84	Aa	1722	G	N1-C6-O6	7.28	124.27	119.90
84	Aa	2267	G	N1-C6-O6	7.28	124.27	119.90
84	Aa	326	C	O4'-C1'-N1	7.27	114.02	108.20
84	Aa	773	G	O4'-C1'-N9	7.27	114.02	108.20
84	Aa	2316	A	O4'-C1'-N9	7.27	114.02	108.20
84	Aa	3318	G	C5-C6-O6	-7.27	124.24	128.60
84	Aa	2530	G	N1-C6-O6	7.27	124.26	119.90
1	Ad	137	A	P-O3'-C3'	7.27	128.42	119.70
84	Aa	3202	G	O4'-C1'-N9	7.27	114.02	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1149	C	O4'-C1'-N1	7.27	114.01	108.20
84	Aa	1327	G	O4'-C1'-N9	7.27	114.01	108.20
84	Aa	1431	G	C5-C6-O6	-7.27	124.24	128.60
84	Aa	1614	G	O4'-C1'-N9	7.27	114.01	108.20
86	Ab	31	G	O4'-C1'-N9	7.27	114.01	108.20
1	Ad	508	U	O4'-C1'-N1	7.26	114.01	108.20
1	Ad	1062	C	O4'-C1'-C2'	-7.26	98.53	105.80
84	Aa	1665	G	O4'-C1'-N9	7.26	114.01	108.20
86	Ab	66	G	C6-C5-N7	-7.26	126.04	130.40
1	Ad	826	C	O4'-C1'-C2'	-7.26	98.54	105.80
84	Aa	1827	U	C5'-C4'-C3'	-7.26	104.38	116.00
71	CB	265	TYR	CB-CG-CD2	-7.26	116.64	121.00
84	Aa	2966	G	C5-C6-O6	-7.26	124.24	128.60
1	Ad	1472	G	N9-C1'-C2'	7.26	123.44	114.00
84	Aa	1995	U	O4'-C1'-N1	7.26	114.01	108.20
84	Aa	2926	U	O4'-C1'-N1	7.26	114.01	108.20
85	Ac	29	U	O4'-C1'-N1	7.26	114.01	108.20
84	Aa	202	G	C5-C6-O6	-7.26	124.25	128.60
1	Ad	945	A	N9-C1'-C2'	-7.26	104.02	112.00
84	Aa	18	G	N1-C6-O6	7.26	124.25	119.90
84	Aa	463	G	N1-C6-O6	7.26	124.25	119.90
84	Aa	1385	C	O4'-C1'-N1	7.26	114.00	108.20
84	Aa	2481	C	O4'-C1'-N1	7.26	114.00	108.20
84	Aa	3059	C	C2-N1-C1'	7.26	126.78	118.80
1	Ad	1372	C	C3'-C2'-C1'	7.25	107.30	101.50
84	Aa	173	C	O4'-C1'-N1	7.25	114.00	108.20
84	Aa	807	C	O4'-C1'-N1	7.25	114.00	108.20
84	Aa	2394	G	O4'-C1'-N9	7.25	114.00	108.20
84	Aa	2116	G	N1-C6-O6	7.25	124.25	119.90
84	Aa	2379	U	O4'-C1'-N1	7.25	114.00	108.20
1	Ad	1464	G	C1'-O4'-C4'	-7.25	104.10	109.90
85	Ac	111	G	C5-C6-O6	-7.25	124.25	128.60
1	Ad	1250	C	C3'-C2'-C1'	-7.25	95.70	101.50
84	Aa	44	A	O4'-C1'-N9	7.25	114.00	108.20
84	Aa	614	C	O4'-C1'-N1	7.25	114.00	108.20
84	Aa	1171	U	O4'-C1'-N1	7.25	114.00	108.20
84	Aa	2356	A	C4-C5-C6	7.25	120.62	117.00
84	Aa	2727	U	O4'-C1'-N1	7.25	114.00	108.20
86	Ab	18	C	N3-C4-N4	7.25	123.07	118.00
84	Aa	794	G	C5-C6-O6	-7.25	124.25	128.60
84	Aa	865	U	O4'-C1'-N1	7.24	113.99	108.20
1	Ad	26	A	O4'-C1'-N9	7.24	113.99	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	744	G	O4'-C4'-C3'	-7.24	96.76	104.00
1	Ad	1523	A	O4'-C1'-C2'	-7.24	98.56	105.80
84	Aa	2335	U	O4'-C1'-N1	7.24	113.99	108.20
84	Aa	2775	C	O4'-C1'-N1	7.24	113.99	108.20
1	Ad	352	U	O4'-C1'-N1	7.24	113.99	108.20
1	Ad	1105	G	O4'-C1'-C2'	7.24	114.11	107.60
18	BN	128	TYR	CB-CG-CD2	7.24	125.34	121.00
84	Aa	1085	G	O4'-C1'-N9	7.24	113.99	108.20
84	Aa	2515	C	O4'-C1'-N1	7.24	113.99	108.20
84	Aa	2770	U	O4'-C1'-N1	7.24	113.99	108.20
1	Ad	1429	U	O4'-C1'-N1	7.24	113.99	108.20
1	Ad	1235	U	C3'-C2'-C1'	7.24	107.29	101.50
1	Ad	1339	C	O4'-C1'-N1	7.24	113.99	108.20
1	Ad	1647	C	C3'-C2'-C1'	7.24	107.29	101.50
84	Aa	747	A	C5-C6-N6	-7.24	117.91	123.70
84	Aa	1425	G	O4'-C1'-N9	7.24	113.99	108.20
84	Aa	403	U	O4'-C1'-N1	7.23	113.99	108.20
84	Aa	1090	C	O4'-C1'-N1	7.23	113.99	108.20
84	Aa	1187	G	N1-C6-O6	7.23	124.24	119.90
84	Aa	1198	G	N1-C6-O6	7.23	124.24	119.90
84	Aa	2492	C	O4'-C1'-N1	7.23	113.99	108.20
84	Aa	2959	G	C5-C6-O6	-7.23	124.26	128.60
84	Aa	90	G	C5-C6-O6	-7.23	124.26	128.60
84	Aa	816	G	C5-C6-O6	-7.23	124.26	128.60
84	Aa	1924	G	O4'-C1'-N9	7.23	113.98	108.20
84	Aa	2446	G	N1-C6-O6	7.23	124.24	119.90
84	Aa	2604	A	C5-C6-N1	-7.23	114.08	117.70
1	Ad	54	C	N1-C1'-C2'	-7.23	104.05	112.00
1	Ad	1391	G	C1'-O4'-C4'	-7.23	104.12	109.90
84	Aa	2708	A	C5-C6-N6	-7.23	117.92	123.70
84	Aa	3016	C	O4'-C1'-N1	7.23	113.98	108.20
1	Ad	1444	G	C3'-C2'-C1'	7.23	107.28	101.50
84	Aa	2293	U	O4'-C1'-N1	7.23	113.98	108.20
1	Ad	1121	A	O4'-C1'-N9	7.22	113.98	108.20
84	Aa	1652	G	C5-C6-O6	-7.22	124.27	128.60
84	Aa	2407	U	O4'-C1'-N1	7.22	113.98	108.20
42	CJ	70	TYR	CB-CG-CD2	7.22	125.33	121.00
84	Aa	45	U	O4'-C1'-N1	7.22	113.98	108.20
84	Aa	220	G	N1-C6-O6	7.22	124.23	119.90
84	Aa	333	G	N1-C6-O6	7.22	124.23	119.90
84	Aa	675	C	O4'-C1'-N1	7.22	113.98	108.20
84	Aa	2180	G	O4'-C1'-N9	7.22	113.98	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
71	CB	197	TYR	CB-CG-CD1	-7.22	116.67	121.00
1	Ad	412	C	C3'-C2'-C1'	7.22	107.28	101.50
84	Aa	379	U	O4'-C1'-N1	7.22	113.98	108.20
84	Aa	526	A	C5'-C4'-C3'	-7.22	104.45	116.00
84	Aa	270	G	O4'-C1'-N9	7.22	113.97	108.20
84	Aa	708	C	O4'-C1'-N1	7.22	113.97	108.20
84	Aa	1055	U	O4'-C1'-N1	7.22	113.97	108.20
1	Ad	123	U	O4'-C1'-N1	7.22	113.97	108.20
84	Aa	467	C	O4'-C1'-N1	7.22	113.97	108.20
84	Aa	603	G	N1-C6-O6	7.22	124.23	119.90
84	Aa	1510	G	O4'-C1'-N9	7.22	113.97	108.20
84	Aa	2376	G	C5-C6-O6	-7.22	124.27	128.60
84	Aa	2646	A	C5-C6-N6	-7.22	117.93	123.70
84	Aa	1469	G	C5-C6-O6	-7.21	124.27	128.60
86	Ab	7	G	O4'-C1'-N9	7.21	113.97	108.20
84	Aa	834	G	O4'-C1'-N9	7.21	113.97	108.20
84	Aa	2566	C	N3-C4-N4	7.21	123.05	118.00
1	Ad	334	G	O4'-C1'-C2'	7.21	114.09	107.60
84	Aa	836	G	C5'-C4'-C3'	-7.21	104.46	116.00
84	Aa	2289	U	O4'-C1'-N1	7.21	113.97	108.20
85	Ac	15	G	C5-C6-O6	-7.21	124.27	128.60
1	Ad	919	G	O4'-C1'-N9	7.21	113.97	108.20
1	Ad	1303	G	O4'-C1'-C2'	7.21	114.09	107.60
84	Aa	86	U	O4'-C1'-N1	7.21	113.97	108.20
84	Aa	2589	G	C5-C6-O6	-7.21	124.28	128.60
84	Aa	2779	G	P-O3'-C3'	7.21	128.35	119.70
84	Aa	1401	C	O4'-C1'-N1	7.21	113.97	108.20
84	Aa	1764	G	C5-C6-O6	-7.21	124.28	128.60
1	Ad	1218	U	O4'-C1'-N1	7.21	113.96	108.20
84	Aa	50	A	C5-C6-N6	-7.21	117.94	123.70
84	Aa	1060	U	O4'-C1'-N1	7.20	113.96	108.20
84	Aa	2483	A	C5-C6-N6	-7.20	117.94	123.70
84	Aa	2997	C	C2-N1-C1'	7.20	126.72	118.80
84	Aa	3255	U	O4'-C1'-N1	7.20	113.96	108.20
84	Aa	390	G	C5-C6-O6	-7.20	124.28	128.60
84	Aa	771	G	C5-C6-O6	-7.20	124.28	128.60
84	Aa	862	G	C5-C6-O6	-7.20	124.28	128.60
84	Aa	3389	C	O4'-C1'-N1	7.20	113.96	108.20
86	Ab	70	G	C5-C6-O6	-7.20	124.28	128.60
1	Ad	645	G	O4'-C1'-N9	7.20	113.96	108.20
84	Aa	989	U	O4'-C1'-N1	7.20	113.96	108.20
84	Aa	1018	C	O4'-C1'-N1	7.20	113.96	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	163	U	O4'-C1'-N1	7.20	113.96	108.20
84	Aa	971	G	C5-C6-O6	-7.20	124.28	128.60
84	Aa	3250	C	O4'-C1'-N1	7.20	113.96	108.20
86	Ab	14	C	N3-C4-C5	-7.20	119.02	121.90
84	Aa	514	G	O4'-C1'-N9	7.20	113.96	108.20
84	Aa	1484	A	N1-C6-N6	7.20	122.92	118.60
84	Aa	1718	U	O4'-C1'-N1	7.20	113.96	108.20
84	Aa	2557	C	N3-C4-C5	-7.20	119.02	121.90
84	Aa	3000	U	O4'-C1'-N1	7.20	113.96	108.20
85	Ac	32	C	O4'-C1'-N1	7.20	113.96	108.20
86	Ab	113	G	O4'-C1'-N9	7.20	113.96	108.20
84	Aa	3354	A	O4'-C1'-N9	7.19	113.96	108.20
84	Aa	496	U	O4'-C1'-N1	7.19	113.95	108.20
1	Ad	1672	U	O4'-C1'-N1	7.19	113.95	108.20
84	Aa	97	G	C5-C6-O6	-7.19	124.28	128.60
84	Aa	1241	G	C5-C6-O6	-7.19	124.29	128.60
84	Aa	1265	G	N1-C6-O6	7.19	124.21	119.90
84	Aa	1380	C	O4'-C1'-N1	7.19	113.95	108.20
84	Aa	2453	G	C5-C6-O6	-7.19	124.28	128.60
84	Aa	3067	G	C5-C6-O6	-7.19	124.29	128.60
84	Aa	566	G	C5-C6-O6	-7.19	124.29	128.60
1	Ad	865	U	O4'-C1'-N1	7.19	113.95	108.20
84	Aa	1074	C	O4'-C1'-N1	7.19	113.95	108.20
1	Ad	1330	A	O4'-C1'-N9	7.19	113.95	108.20
84	Aa	313	C	O4'-C1'-N1	7.19	113.95	108.20
84	Aa	1329	G	O4'-C1'-N9	7.19	113.95	108.20
84	Aa	994	U	O4'-C1'-N1	7.18	113.95	108.20
84	Aa	2955	U	O4'-C1'-N1	7.18	113.95	108.20
84	Aa	2427	C	O4'-C1'-N1	7.18	113.95	108.20
84	Aa	2972	C	O4'-C1'-N1	7.18	113.95	108.20
86	Ab	55	A	O4'-C1'-N9	7.18	113.95	108.20
84	Aa	1250	G	N1-C6-O6	7.18	124.21	119.90
84	Aa	1402	G	O4'-C1'-N9	7.18	113.94	108.20
84	Aa	2613	G	C5-C6-O6	-7.18	124.29	128.60
86	Ab	72	G	C8-N9-C4	-7.18	103.53	106.40
84	Aa	2725	U	O4'-C1'-N1	7.18	113.94	108.20
84	Aa	2915	U	O4'-C1'-N1	7.18	113.94	108.20
84	Aa	713	G	O4'-C1'-N9	7.18	113.94	108.20
84	Aa	934	C	O4'-C1'-N1	7.18	113.94	108.20
84	Aa	1935	G	O4'-C1'-N9	7.18	113.94	108.20
1	Ad	1316	A	N9-C1'-C2'	7.17	123.33	114.00
84	Aa	1142	G	O4'-C1'-N9	7.17	113.94	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2994	U	O4'-C1'-N1	7.17	113.94	108.20
1	Ad	426	G	O4'-C1'-N9	7.17	113.94	108.20
57	Ce	67	TYR	CB-CG-CD2	-7.17	116.70	121.00
84	Aa	432	G	C5-C6-O6	-7.17	124.30	128.60
84	Aa	1041	C	O4'-C1'-N1	7.17	113.94	108.20
84	Aa	1521	U	O4'-C1'-N1	7.17	113.94	108.20
1	Ad	406	C	C3'-C2'-C1'	7.17	107.24	101.50
1	Ad	1569	U	O4'-C1'-N1	7.17	113.93	108.20
1	Ad	1720	G	P-O3'-C3'	7.17	128.30	119.70
84	Aa	1697	G	O4'-C1'-N9	7.17	113.93	108.20
84	Aa	1705	A	O4'-C1'-N9	7.17	113.93	108.20
84	Aa	1975	G	O4'-C1'-N9	7.17	113.93	108.20
1	Ad	1554	G	O4'-C1'-C2'	7.17	114.05	107.60
84	Aa	410	G	O4'-C1'-N9	7.17	113.93	108.20
84	Aa	2004	U	O4'-C1'-N1	7.17	113.93	108.20
1	Ad	1796	G	N9-C1'-C2'	7.16	123.31	114.00
84	Aa	1034	U	O4'-C1'-N1	7.16	113.93	108.20
84	Aa	2245	G	C5-C6-O6	-7.16	124.30	128.60
84	Aa	3042	U	O4'-C1'-N1	7.16	113.93	108.20
84	Aa	41	C	O4'-C1'-N1	7.16	113.93	108.20
84	Aa	752	U	O4'-C1'-N1	7.16	113.93	108.20
86	Ab	44	C	O4'-C1'-N1	7.16	113.93	108.20
1	Ad	117	U	O4'-C1'-N1	7.16	113.93	108.20
1	Ad	149	G	O4'-C1'-N9	7.16	113.93	108.20
84	Aa	671	C	O4'-C1'-N1	7.16	113.93	108.20
84	Aa	891	U	O4'-C1'-N1	7.16	113.93	108.20
84	Aa	2253	U	O4'-C1'-N1	7.16	113.93	108.20
1	Ad	1663	A	O4'-C1'-C2'	-7.16	98.64	105.80
84	Aa	311	G	O4'-C1'-N9	7.16	113.93	108.20
84	Aa	1232	A	C5-C6-N1	-7.16	114.12	117.70
84	Aa	1433	U	O4'-C1'-N1	7.16	113.93	108.20
84	Aa	1959	U	O4'-C1'-N1	7.16	113.93	108.20
1	Ad	708	G	O4'-C1'-N9	-7.16	102.47	108.20
84	Aa	2395	G	N1-C6-O6	7.16	124.19	119.90
84	Aa	1676	A	C5-C6-N1	-7.16	114.12	117.70
84	Aa	2333	U	O4'-C1'-N1	7.16	113.92	108.20
1	Ad	475	A	P-O3'-C3'	7.15	128.28	119.70
84	Aa	581	G	C5-C6-O6	-7.15	124.31	128.60
84	Aa	1064	U	O4'-C1'-N1	7.15	113.92	108.20
84	Aa	1531	G	N1-C6-O6	7.15	124.19	119.90
84	Aa	2302	G	C5-C6-O6	-7.15	124.31	128.60
1	Ad	710	G	O4'-C1'-N9	7.15	113.92	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3108	U	O4'-C1'-N1	7.15	113.92	108.20
84	Aa	1302	C	O4'-C1'-N1	7.15	113.92	108.20
84	Aa	1315	G	N1-C6-O6	7.15	124.19	119.90
84	Aa	2070	C	O4'-C1'-N1	7.15	113.92	108.20
84	Aa	3304	U	O4'-C1'-N1	7.15	113.92	108.20
1	Ad	242	A	O4'-C1'-N9	7.15	113.92	108.20
84	Aa	2220	U	O4'-C1'-N1	7.15	113.92	108.20
1	Ad	800	U	C3'-C2'-C1'	-7.15	95.78	101.50
84	Aa	706	U	O4'-C1'-N1	7.15	113.92	108.20
84	Aa	1786	G	C5-C6-O6	-7.15	124.31	128.60
84	Aa	2945	G	C5-C6-O6	-7.15	124.31	128.60
1	Ad	164	C	O4'-C1'-N1	7.15	113.92	108.20
1	Ad	1260	A	C3'-C2'-C1'	7.15	107.22	101.50
84	Aa	2092	C	N3-C4-C5	-7.15	119.04	121.90
86	Ab	4	U	O4'-C1'-N1	7.15	113.92	108.20
84	Aa	2346	U	O4'-C1'-N1	7.14	113.92	108.20
84	Aa	2383	G	C5-C6-O6	-7.14	124.31	128.60
1	Ad	1194	C	P-O3'-C3'	7.14	128.27	119.70
84	Aa	720	G	C5-C6-O6	-7.14	124.31	128.60
1	Ad	31	C	C3'-C2'-C1'	7.14	107.21	101.50
1	Ad	339	G	N9-C1'-C2'	7.14	123.28	114.00
84	Aa	554	C	C4'-C3'-O3'	7.14	127.28	113.00
84	Aa	2868	C	O4'-C1'-N1	7.14	113.91	108.20
84	Aa	3225	G	O4'-C1'-N9	7.14	113.91	108.20
1	Ad	258	U	O4'-C1'-N1	7.14	113.91	108.20
1	Ad	593	C	O4'-C1'-N1	7.14	113.91	108.20
1	Ad	1042	C	N1-C1'-C2'	7.14	123.28	114.00
84	Aa	1265	G	C5-C6-O6	-7.14	124.32	128.60
1	Ad	308	U	O4'-C1'-N1	7.14	113.91	108.20
84	Aa	157	G	C5-C6-O6	-7.14	124.32	128.60
84	Aa	330	C	O4'-C1'-N1	7.14	113.91	108.20
84	Aa	2451	G	O4'-C1'-N9	7.14	113.91	108.20
84	Aa	1494	A	O4'-C1'-N9	7.13	113.91	108.20
84	Aa	2307	A	C5-C6-N1	-7.13	114.13	117.70
84	Aa	3301	G	C5-C6-O6	-7.13	124.32	128.60
1	Ad	1587	G	O4'-C1'-N9	7.13	113.91	108.20
84	Aa	1948	G	O4'-C1'-N9	7.13	113.91	108.20
84	Aa	570	G	O4'-C1'-N9	7.13	113.90	108.20
84	Aa	2077	C	P-O3'-C3'	7.13	128.25	119.70
84	Aa	2310	G	N1-C6-O6	7.13	124.18	119.90
84	Aa	2081	C	C2'-C3'-O3'	7.13	125.18	109.50
85	Ac	21	C	O4'-C1'-N1	7.13	113.90	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1248	A	C3'-C2'-C1'	-7.13	95.80	101.50
1	Ad	1725	C	O4'-C1'-N1	7.13	113.90	108.20
84	Aa	2905	A	C4-C5-C6	7.13	120.56	117.00
1	Ad	1528	U	C1'-O4'-C4'	7.12	115.60	109.90
84	Aa	2093	G	C4-N9-C1'	-7.12	117.24	126.50
84	Aa	2341	U	O4'-C1'-N1	7.12	113.90	108.20
84	Aa	3170	C	O4'-C1'-N1	7.12	113.90	108.20
84	Aa	223	C	O4'-C1'-N1	7.12	113.90	108.20
84	Aa	2836	G	N1-C6-O6	7.12	124.17	119.90
1	Ad	530	A	P-O3'-C3'	7.12	128.25	119.70
84	Aa	2856	U	O4'-C1'-N1	7.12	113.90	108.20
1	Ad	1647	C	O4'-C1'-C2'	-7.12	98.68	105.80
84	Aa	321	A	N1-C6-N6	7.12	122.87	118.60
84	Aa	931	C	O4'-C1'-N1	7.12	113.89	108.20
85	Ac	52	A	C5-C6-N6	-7.12	118.01	123.70
84	Aa	1948	G	N1-C6-O6	7.12	124.17	119.90
84	Aa	1439	U	O4'-C1'-N1	7.12	113.89	108.20
84	Aa	2242	G	O4'-C1'-N9	7.12	113.89	108.20
84	Aa	2528	U	O4'-C1'-N1	7.12	113.89	108.20
84	Aa	3029	G	O4'-C1'-N9	7.12	113.89	108.20
84	Aa	3333	C	C5'-C4'-O4'	7.12	117.64	109.10
84	Aa	1556	G	C5-C6-O6	-7.11	124.33	128.60
84	Aa	2801	A	C5-C6-N1	-7.11	114.14	117.70
84	Aa	3286	G	C5-C6-O6	-7.11	124.33	128.60
84	Aa	2434	G	N1-C6-O6	7.11	124.17	119.90
84	Aa	745	G	O4'-C1'-N9	7.11	113.89	108.20
84	Aa	1538	A	C5-C6-N6	-7.11	118.01	123.70
84	Aa	2385	A	C4-C5-C6	7.11	120.56	117.00
84	Aa	2611	G	O4'-C1'-N9	7.11	113.89	108.20
84	Aa	2701	G	C5-C6-O6	-7.11	124.33	128.60
1	Ad	99	U	N1-C1'-C2'	7.11	123.24	114.00
1	Ad	414	A	C1'-O4'-C4'	-7.11	104.21	109.90
1	Ad	716	A	C4'-C3'-O3'	-7.11	94.47	109.40
1	Ad	323	U	P-O3'-C3'	7.11	128.23	119.70
1	Ad	1184	C	C3'-C2'-C1'	7.11	107.19	101.50
3	Af	21	C	O4'-C1'-C2'	-7.11	98.69	105.80
84	Aa	456	G	C5-C6-O6	-7.11	124.34	128.60
84	Aa	1280	U	O4'-C1'-N1	7.11	113.89	108.20
1	Ad	391	A	C1'-O4'-C4'	7.10	115.58	109.90
84	Aa	765	U	O4'-C1'-N1	7.10	113.88	108.20
86	Ab	82	G	N3-C2-N2	7.10	124.87	119.90
84	Aa	513	C	C5'-C4'-O4'	7.10	117.62	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1372	U	O4'-C1'-N1	7.10	113.88	108.20
84	Aa	1650	G	O4'-C1'-N9	7.10	113.88	108.20
84	Aa	1662	G	C5-C6-O6	-7.10	124.34	128.60
1	Ad	480	U	N1-C1'-C2'	7.10	123.23	114.00
84	Aa	1350	G	O4'-C1'-N9	7.10	113.88	108.20
84	Aa	637	C	P-O3'-C3'	-7.10	111.19	119.70
1	Ad	1313	G	O4'-C1'-N9	7.09	113.88	108.20
84	Aa	1358	C	O4'-C1'-N1	7.09	113.88	108.20
84	Aa	2838	C	C6-N1-C1'	-7.09	112.29	120.80
84	Aa	3179	G	O4'-C1'-N9	7.09	113.88	108.20
86	Ab	73	U	C5'-C4'-O4'	7.09	117.61	109.10
84	Aa	455	U	O4'-C1'-N1	7.09	113.87	108.20
84	Aa	476	C	O4'-C1'-N1	7.09	113.87	108.20
84	Aa	1683	U	O4'-C1'-N1	7.09	113.87	108.20
84	Aa	2430	C	O4'-C1'-N1	7.09	113.87	108.20
84	Aa	2746	G	O4'-C1'-N9	7.09	113.87	108.20
1	Ad	151	A	O4'-C1'-N9	7.09	113.87	108.20
1	Ad	1154	G	O4'-C1'-C2'	7.09	113.98	107.60
84	Aa	293	A	O4'-C1'-N9	7.09	113.87	108.20
84	Aa	1088	A	C5-C6-N6	-7.09	118.03	123.70
84	Aa	1526	A	C4-C5-C6	7.09	120.54	117.00
84	Aa	3200	A	O4'-C1'-N9	7.09	113.87	108.20
86	Ab	71	A	C5-N7-C8	7.09	107.44	103.90
84	Aa	1924	G	N1-C6-O6	7.08	124.15	119.90
1	Ad	1172	G	C1'-O4'-C4'	-7.08	104.23	109.90
1	Ad	1376	A	C1'-O4'-C4'	7.08	115.57	109.90
84	Aa	1304	G	C5-C6-O6	-7.08	124.35	128.60
85	Ac	119	C	O4'-C1'-N1	7.08	113.87	108.20
84	Aa	440	U	O4'-C1'-N1	7.08	113.86	108.20
84	Aa	1526	A	C5-C6-N1	-7.08	114.16	117.70
84	Aa	3165	C	O4'-C1'-N1	7.08	113.86	108.20
1	Ad	76	U	O4'-C1'-N1	7.08	113.86	108.20
1	Ad	161	G	C3'-C2'-C1'	-7.08	95.84	101.50
84	Aa	510	C	O4'-C1'-N1	7.08	113.86	108.20
1	Ad	223	A	C1'-O4'-C4'	-7.08	104.24	109.90
84	Aa	644	U	O4'-C1'-N1	7.08	113.86	108.20
84	Aa	2078	G	C5-C6-O6	-7.08	124.35	128.60
84	Aa	3369	G	C5-C6-O6	-7.08	124.35	128.60
1	Ad	742	C	O4'-C1'-N1	7.08	113.86	108.20
1	Ad	1667	A	O4'-C1'-N9	7.08	113.86	108.20
1	Ad	792	U	N1-C1'-C2'	-7.08	104.22	112.00
84	Aa	921	C	O4'-C1'-N1	7.08	113.86	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1036	C	O4'-C1'-N1	7.08	113.86	108.20
84	Aa	1539	G	C5-C6-O6	-7.08	124.36	128.60
84	Aa	1649	G	O4'-C1'-N9	7.08	113.86	108.20
1	Ad	526	U	O4'-C1'-N1	7.07	113.86	108.20
84	Aa	666	U	O4'-C1'-N1	7.07	113.86	108.20
84	Aa	687	C	N3-C4-N4	7.07	122.95	118.00
84	Aa	2099	G	O4'-C1'-N9	7.07	113.86	108.20
84	Aa	377	C	O4'-C1'-N1	7.07	113.86	108.20
1	Ad	213	U	O4'-C1'-N1	7.07	113.86	108.20
1	Ad	535	C	C1'-O4'-C4'	-7.07	104.24	109.90
84	Aa	1025	G	C5-C6-O6	-7.07	124.36	128.60
84	Aa	1645	G	C5-C6-O6	-7.07	124.36	128.60
84	Aa	1729	G	C5-C6-O6	-7.07	124.36	128.60
1	Ad	1392	G	O4'-C1'-C2'	7.07	113.96	107.60
84	Aa	564	A	O4'-C1'-N9	7.07	113.85	108.20
84	Aa	574	C	O4'-C1'-N1	7.07	113.85	108.20
84	Aa	2903	G	C5-C6-O6	-7.07	124.36	128.60
1	Ad	1033	C	N1-C1'-C2'	7.06	123.18	114.00
86	Ab	78	C	C6-N1-C2	-7.06	117.47	120.30
86	Ab	86	G	O4'-C1'-N9	7.06	113.85	108.20
84	Aa	2340	G	O4'-C1'-N9	7.06	113.85	108.20
1	Ad	758	A	N9-C1'-C2'	7.06	123.18	114.00
58	Cj	84	ALA	N-CA-CB	7.06	119.98	110.10
84	Aa	247	C	O4'-C1'-N1	7.06	113.85	108.20
84	Aa	1909	G	C5-C6-O6	-7.06	124.36	128.60
84	Aa	2756	G	N3-C2-N2	7.06	124.84	119.90
84	Aa	2949	G	O4'-C1'-N9	7.06	113.85	108.20
1	Ad	430	G	O4'-C1'-C2'	7.06	113.95	107.60
1	Ad	1019	G	O4'-C1'-C2'	7.06	113.95	107.60
1	Ad	1235	U	O4'-C1'-C2'	-7.06	98.74	105.80
1	Ad	1255	U	N1-C1'-C2'	-7.06	104.24	112.00
84	Aa	299	G	N1-C6-O6	7.06	124.14	119.90
84	Aa	621	C	C6-N1-C1'	-7.06	112.33	120.80
84	Aa	1318	C	N3-C4-C5	-7.06	119.08	121.90
84	Aa	2444	U	O4'-C1'-N1	7.06	113.85	108.20
84	Aa	623	G	C5-C6-O6	-7.06	124.37	128.60
84	Aa	686	A	C5-C6-N6	-7.05	118.06	123.70
84	Aa	791	C	O4'-C1'-N1	7.05	113.84	108.20
84	Aa	3264	C	C2-N1-C1'	7.05	126.56	118.80
84	Aa	2336	C	O4'-C1'-N1	7.05	113.84	108.20
84	Aa	3237	G	C5-C6-O6	-7.05	124.37	128.60
84	Aa	818	G	N1-C6-O6	7.05	124.13	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1228	C	O4'-C1'-N1	7.05	113.84	108.20
84	Aa	1422	G	O4'-C1'-N9	7.05	113.84	108.20
84	Aa	2552	U	O4'-C1'-N1	7.05	113.84	108.20
84	Aa	870	G	C5-C6-O6	-7.05	124.37	128.60
84	Aa	1202	C	O4'-C1'-N1	7.05	113.84	108.20
84	Aa	1601	G	C5-C6-O6	-7.05	124.37	128.60
84	Aa	616	A	O4'-C1'-N9	7.05	113.84	108.20
1	Ad	1725	C	C1'-O4'-C4'	7.05	115.54	109.90
84	Aa	884	C	O4'-C1'-N1	7.05	113.84	108.20
84	Aa	1126	U	O4'-C1'-N1	7.05	113.84	108.20
84	Aa	2793	G	N1-C6-O6	7.04	124.13	119.90
1	Ad	176	A	O4'-C1'-C2'	-7.04	98.76	105.80
84	Aa	1513	C	O4'-C1'-N1	7.04	113.83	108.20
84	Aa	2163	G	C5'-C4'-O4'	-7.04	100.65	109.10
84	Aa	2635	G	N1-C6-O6	7.04	124.12	119.90
84	Aa	2932	A	C5-C6-N6	-7.04	118.07	123.70
84	Aa	1664	G	C5-C6-O6	-7.04	124.38	128.60
1	Ad	567	U	N1-C1'-C2'	7.04	123.15	114.00
84	Aa	1325	G	O4'-C1'-N9	7.04	113.83	108.20
84	Aa	1509	G	N1-C6-O6	7.04	124.12	119.90
84	Aa	1868	C	N3-C4-C5	-7.04	119.08	121.90
86	Ab	68	G	C5-C6-O6	-7.04	124.38	128.60
84	Aa	193	U	O4'-C1'-N1	7.04	113.83	108.20
84	Aa	1898	G	C5-C6-O6	-7.04	124.38	128.60
84	Aa	3075	G	C5-C6-O6	-7.04	124.38	128.60
1	Ad	1584	A	C3'-C2'-C1'	7.04	107.13	101.50
84	Aa	1282	A	O4'-C1'-N9	7.04	113.83	108.20
84	Aa	2536	G	C5-C6-O6	-7.04	124.38	128.60
1	Ad	478	A	O4'-C1'-N9	7.03	113.83	108.20
84	Aa	2602	U	O4'-C1'-N1	7.03	113.83	108.20
84	Aa	1563	G	P-O3'-C3'	7.03	128.14	119.70
84	Aa	2208	A	O4'-C1'-N9	7.03	113.83	108.20
84	Aa	3079	G	C5-C6-O6	-7.03	124.38	128.60
84	Aa	740	G	C5-C6-O6	-7.03	124.38	128.60
84	Aa	1851	U	O4'-C1'-N1	7.03	113.82	108.20
86	Ab	90	A	C4-C5-C6	7.03	120.52	117.00
84	Aa	992	U	O4'-C1'-N1	7.03	113.82	108.20
84	Aa	1258	C	N3-C4-C5	-7.03	119.09	121.90
84	Aa	1869	U	O4'-C1'-N1	7.03	113.82	108.20
86	Ab	17	G	N1-C6-O6	7.03	124.12	119.90
48	CD	183	PHE	CB-CG-CD1	-7.03	115.88	120.80
84	Aa	512	G	N3-C2-N2	7.03	124.82	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1989	G	O4'-C1'-N9	7.03	113.82	108.20
84	Aa	2524	U	O4'-C1'-N1	7.03	113.82	108.20
84	Aa	2570	U	O4'-C1'-N1	7.03	113.82	108.20
84	Aa	2573	U	O4'-C1'-N1	7.03	113.82	108.20
85	Ac	90	C	O4'-C1'-N1	7.03	113.82	108.20
84	Aa	956	G	C5-C6-O6	-7.02	124.39	128.60
84	Aa	2188	U	O4'-C1'-N1	7.02	113.82	108.20
1	Ad	351	G	C3'-C2'-C1'	-7.02	95.88	101.50
84	Aa	1669	C	O4'-C1'-N1	7.02	113.82	108.20
84	Aa	2088	C	N3-C4-C5	-7.02	119.09	121.90
84	Aa	2190	C	O4'-C1'-N1	7.02	113.82	108.20
84	Aa	3146	C	O4'-C1'-N1	7.02	113.82	108.20
1	Ad	1088	G	O4'-C1'-N9	7.02	113.82	108.20
84	Aa	951	C	O4'-C1'-N1	7.02	113.82	108.20
84	Aa	1079	G	C5-C6-O6	-7.02	124.39	128.60
84	Aa	1106	G	C5-C6-O6	-7.02	124.39	128.60
1	Ad	1	U	O4'-C1'-N1	7.02	113.81	108.20
84	Aa	239	C	O4'-C1'-N1	7.02	113.81	108.20
84	Aa	972	C	O4'-C1'-N1	7.02	113.82	108.20
84	Aa	1993	G	C5-C6-O6	-7.02	124.39	128.60
84	Aa	74	G	N1-C6-O6	7.02	124.11	119.90
84	Aa	1170	U	O4'-C1'-N1	7.02	113.81	108.20
84	Aa	1187	G	O4'-C1'-N9	7.02	113.81	108.20
84	Aa	1548	U	O4'-C1'-N1	7.02	113.81	108.20
84	Aa	1625	G	C5-C6-O6	-7.02	124.39	128.60
84	Aa	2080	G	O4'-C1'-N9	7.02	113.81	108.20
84	Aa	3274	G	O4'-C1'-N9	7.02	113.81	108.20
85	Ac	35	C	O4'-C1'-N1	7.02	113.81	108.20
85	Ac	78	G	N1-C6-O6	7.02	124.11	119.90
1	Ad	372	U	N1-C1'-C2'	-7.01	104.28	112.00
1	Ad	904	G	O4'-C1'-N9	7.01	113.81	108.20
84	Aa	539	C	O4'-C1'-N1	7.01	113.81	108.20
84	Aa	1528	G	N1-C6-O6	7.01	124.11	119.90
84	Aa	2964	U	O4'-C1'-N1	7.01	113.81	108.20
84	Aa	3279	G	C5-C6-O6	-7.01	124.39	128.60
84	Aa	181	G	N1-C6-O6	7.01	124.11	119.90
84	Aa	331	G	O4'-C1'-N9	7.01	113.81	108.20
84	Aa	507	C	N3-C4-N4	7.01	122.91	118.00
85	Ac	56	G	C5-C6-O6	-7.01	124.39	128.60
86	Ab	118	C	C2-N3-C4	7.01	123.41	119.90
1	Ad	1124	G	O4'-C1'-C2'	7.01	113.91	107.60
1	Ad	1489	A	O4'-C1'-N9	7.01	113.81	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1567	G	C3'-C2'-C1'	-7.01	95.89	101.50
84	Aa	678	G	C5-C6-O6	-7.01	124.39	128.60
84	Aa	815	G	O4'-C1'-N9	7.01	113.81	108.20
84	Aa	1991	U	O4'-C1'-N1	7.01	113.81	108.20
84	Aa	2151	G	C5-C6-O6	-7.01	124.39	128.60
84	Aa	2466	G	C5-C6-O6	-7.01	124.39	128.60
84	Aa	1730	U	O4'-C1'-N1	7.01	113.81	108.20
57	Ce	26	TYR	CB-CG-CD2	-7.01	116.80	121.00
84	Aa	835	G	C5-C6-O6	-7.01	124.40	128.60
84	Aa	2789	G	C5-C6-O6	-7.01	124.40	128.60
85	Ac	24	G	O4'-C1'-N9	7.01	113.81	108.20
86	Ab	36	C	O4'-C1'-N1	7.01	113.81	108.20
1	Ad	1164	C	O4'-C1'-N1	7.00	113.80	108.20
84	Aa	2726	U	O4'-C1'-N1	7.00	113.80	108.20
84	Aa	2956	U	O4'-C1'-N1	7.00	113.80	108.20
1	Ad	975	A	C3'-C2'-C1'	7.00	107.10	101.50
1	Ad	1080	C	C3'-C2'-C1'	7.00	107.10	101.50
84	Aa	152	C	O4'-C1'-N1	7.00	113.80	108.20
84	Aa	275	G	N3-C2-N2	7.00	124.80	119.90
85	Ac	7	U	O4'-C1'-N1	7.00	113.80	108.20
84	Aa	106	G	O4'-C1'-N9	7.00	113.80	108.20
84	Aa	3384	G	C5-C6-O6	-7.00	124.40	128.60
85	Ac	107	G	C5-C6-O6	-7.00	124.40	128.60
84	Aa	720	G	P-O3'-C3'	-7.00	111.30	119.70
1	Ad	179	A	C3'-C2'-C1'	7.00	107.10	101.50
84	Aa	2163	G	C5-C6-O6	-6.99	124.40	128.60
84	Aa	2206	U	O4'-C1'-N1	6.99	113.79	108.20
84	Aa	1722	G	O4'-C1'-N9	6.99	113.79	108.20
84	Aa	2947	G	N1-C6-O6	6.99	124.09	119.90
84	Aa	1445	U	O4'-C1'-N1	6.99	113.79	108.20
84	Aa	1853	C	N3-C4-C5	-6.99	119.10	121.90
1	Ad	1405	U	C5'-C4'-C3'	-6.99	104.82	116.00
84	Aa	627	G	C5-C6-O6	-6.99	124.41	128.60
84	Aa	2914	G	O4'-C1'-N9	6.99	113.79	108.20
84	Aa	1829	G	C5-C6-O6	-6.99	124.41	128.60
1	Ad	873	G	N9-C1'-C2'	-6.99	104.32	112.00
84	Aa	709	G	N1-C6-O6	6.99	124.09	119.90
84	Aa	787	G	O4'-C1'-N9	6.99	113.79	108.20
84	Aa	1076	G	O4'-C1'-N9	6.99	113.79	108.20
84	Aa	3180	U	O4'-C1'-N1	6.99	113.79	108.20
84	Aa	3014	U	O4'-C1'-N1	6.98	113.79	108.20
84	Aa	2398	A	O4'-C1'-N9	6.98	113.78	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	317	G	C5-C6-O6	-6.98	124.41	128.60
84	Aa	814	U	O4'-C1'-N1	6.98	113.78	108.20
84	Aa	2871	U	O4'-C1'-N1	6.98	113.78	108.20
84	Aa	3150	G	C5-C6-O6	-6.98	124.41	128.60
84	Aa	1119	G	C5-C6-O6	-6.98	124.41	128.60
86	Ab	82	G	N1-C2-N3	-6.98	119.71	123.90
1	Ad	1432	C	C1'-O4'-C4'	6.98	115.48	109.90
42	CJ	70	TYR	CB-CG-CD1	-6.98	116.81	121.00
84	Aa	51	A	C4-C5-C6	6.98	120.49	117.00
84	Aa	104	G	C5-C6-O6	-6.98	124.41	128.60
84	Aa	241	G	C5-C6-O6	-6.98	124.41	128.60
84	Aa	1387	G	N1-C6-O6	6.98	124.09	119.90
84	Aa	2439	A	O4'-C1'-N9	6.98	113.78	108.20
84	Aa	2510	U	O4'-C1'-N1	6.98	113.78	108.20
84	Aa	3378	U	O4'-C1'-N1	6.98	113.78	108.20
1	Ad	64	U	O4'-C1'-C2'	-6.98	98.82	105.80
84	Aa	728	G	N1-C6-O6	6.98	124.09	119.90
84	Aa	1368	U	O4'-C1'-N1	6.98	113.78	108.20
84	Aa	1622	G	C5-C6-O6	-6.98	124.41	128.60
84	Aa	2846	C	O4'-C1'-N1	6.98	113.78	108.20
1	Ad	1549	G	C3'-C2'-C1'	6.97	107.08	101.50
2	Ae	29	C	C3'-C2'-C1'	6.97	107.08	101.50
84	Aa	1091	C	O4'-C1'-N1	6.97	113.78	108.20
84	Aa	1673	A	C5-C6-N1	-6.97	114.21	117.70
84	Aa	1605	U	O4'-C1'-N1	6.97	113.78	108.20
84	Aa	3298	G	C5-C6-O6	-6.97	124.42	128.60
84	Aa	625	G	P-O3'-C3'	6.97	128.07	119.70
1	Ad	627	A	O4'-C1'-N9	6.97	113.78	108.20
1	Ad	746	A	C1'-O4'-C4'	6.97	115.47	109.90
1	Ad	1683	G	C3'-C2'-C1'	-6.97	95.92	101.50
84	Aa	279	G	N1-C6-O6	6.97	124.08	119.90
86	Ab	90	A	C5-C6-N1	-6.97	114.22	117.70
1	Ad	711	C	O4'-C1'-N1	6.97	113.78	108.20
84	Aa	350	A	O4'-C1'-N9	6.97	113.77	108.20
84	Aa	1563	G	C2'-C3'-O3'	6.97	124.85	113.70
84	Aa	3217	G	O4'-C1'-N9	6.97	113.77	108.20
86	Ab	10	C	N3-C4-N4	6.97	122.88	118.00
1	Ad	385	C	C3'-C2'-C1'	6.96	107.07	101.50
1	Ad	726	G	O4'-C1'-N9	6.96	113.77	108.20
84	Aa	1169	G	C5-C6-O6	-6.96	124.42	128.60
84	Aa	1630	C	O4'-C1'-N1	6.96	113.77	108.20
84	Aa	2169	U	O4'-C1'-N1	6.96	113.77	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3370	U	O4'-C1'-N1	6.96	113.77	108.20
84	Aa	3376	C	O4'-C1'-N1	6.96	113.77	108.20
1	Ad	751	U	O4'-C1'-N1	6.96	113.77	108.20
84	Aa	2255	U	O4'-C1'-N1	6.96	113.77	108.20
1	Ad	1735	C	N1-C1'-C2'	6.96	123.05	114.00
84	Aa	1085	G	C5-C6-O6	-6.96	124.42	128.60
84	Aa	1264	A	O4'-C1'-N9	6.96	113.77	108.20
84	Aa	2566	C	O4'-C1'-N1	6.96	113.77	108.20
84	Aa	1726	G	O4'-C1'-N9	6.96	113.77	108.20
84	Aa	2485	U	O4'-C1'-N1	6.96	113.77	108.20
84	Aa	3178	C	C2-N1-C1'	6.96	126.46	118.80
1	Ad	1787	G	C4'-C3'-C2'	-6.96	95.64	102.60
84	Aa	1577	A	O4'-C1'-N9	6.96	113.77	108.20
1	Ad	1512	C	O4'-C1'-N1	6.96	113.77	108.20
84	Aa	380	U	O4'-C1'-N1	6.96	113.77	108.20
84	Aa	750	G	O4'-C1'-N9	6.96	113.77	108.20
84	Aa	1288	C	N3-C4-C5	-6.96	119.12	121.90
84	Aa	2693	G	C5-C6-O6	-6.96	124.43	128.60
1	Ad	246	G	O4'-C1'-N9	6.96	113.76	108.20
84	Aa	425	G	O4'-C1'-N9	6.96	113.76	108.20
84	Aa	1884	U	O4'-C1'-N1	6.96	113.76	108.20
84	Aa	3334	A	C5-C6-N1	-6.96	114.22	117.70
86	Ab	82	G	C6-C5-N7	-6.96	126.23	130.40
84	Aa	1010	A	N1-C6-N6	6.95	122.77	118.60
84	Aa	1259	C	N3-C4-C5	-6.95	119.12	121.90
84	Aa	3177	A	C5-C6-N1	-6.95	114.22	117.70
84	Aa	3300	C	N3-C4-N4	6.95	122.87	118.00
84	Aa	2598	A	O4'-C1'-N9	6.95	113.76	108.20
1	Ad	54	C	C1'-O4'-C4'	6.95	115.46	109.90
1	Ad	173	G	O4'-C1'-N9	6.95	113.76	108.20
1	Ad	856	G	O4'-C1'-C2'	6.95	113.86	107.60
84	Aa	607	U	O4'-C1'-N1	6.95	113.76	108.20
84	Aa	1419	G	C5-C6-O6	-6.95	124.43	128.60
1	Ad	1538	C	C3'-C2'-C1'	6.95	107.06	101.50
84	Aa	1773	U	O4'-C1'-N1	6.95	113.76	108.20
84	Aa	1803	G	C5-C6-O6	-6.95	124.43	128.60
86	Ab	23	A	C5-C6-N1	-6.95	114.23	117.70
1	Ad	1202	G	C3'-C2'-C1'	-6.95	95.94	101.50
84	Aa	1104	C	O4'-C1'-N1	6.95	113.76	108.20
84	Aa	1719	U	O4'-C1'-N1	6.95	113.76	108.20
84	Aa	2599	U	O4'-C1'-N1	6.95	113.76	108.20
84	Aa	3281	G	C5-C6-O6	-6.95	124.43	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	Ab	47	C	N3-C4-N4	6.95	122.86	118.00
84	Aa	1234	G	N1-C6-O6	6.94	124.07	119.90
84	Aa	1618	U	O4'-C1'-N1	6.94	113.75	108.20
1	Ad	1448	U	C4'-C3'-C2'	-6.94	95.66	102.60
6	BK	87	VAL	CA-C-N	6.94	136.54	117.10
84	Aa	1261	C	O4'-C1'-N1	6.94	113.75	108.20
84	Aa	2695	A	O4'-C1'-N9	6.94	113.75	108.20
84	Aa	2808	U	O4'-C1'-N1	6.94	113.75	108.20
84	Aa	3142	C	O4'-C1'-N1	6.94	113.75	108.20
1	Ad	631	C	C1'-O4'-C4'	-6.94	104.35	109.90
84	Aa	252	A	N1-C6-N6	6.94	122.77	118.60
84	Aa	301	G	N1-C6-O6	6.94	124.06	119.90
84	Aa	428	G	N1-C6-O6	6.94	124.06	119.90
84	Aa	1847	G	C5-C6-O6	-6.94	124.44	128.60
84	Aa	1876	U	O4'-C1'-N1	6.94	113.75	108.20
84	Aa	2457	G	P-O3'-C3'	6.94	128.03	119.70
84	Aa	2659	A	C5-C6-N1	-6.94	114.23	117.70
84	Aa	2876	G	N1-C6-O6	6.94	124.06	119.90
1	Ad	1237	G	O4'-C1'-N9	6.94	113.75	108.20
84	Aa	150	G	C5-C6-O6	-6.94	124.44	128.60
84	Aa	1033	G	O4'-C1'-N9	6.94	113.75	108.20
84	Aa	3325	G	O4'-C1'-N9	6.94	113.75	108.20
85	Ac	136	G	C5-C6-O6	-6.94	124.44	128.60
1	Ad	1150	U	O4'-C1'-N1	6.94	113.75	108.20
52	CW	54	THR	N-CA-CB	6.94	123.48	110.30
1	Ad	914	U	N1-C1'-C2'	6.94	123.02	114.00
1	Ad	1062	C	C1'-O4'-C4'	6.93	115.45	109.90
1	Ad	1273	U	O4'-C1'-N1	6.93	113.75	108.20
1	Ad	1519	G	O4'-C1'-N9	6.93	113.75	108.20
85	Ac	70	G	C5-C6-O6	-6.93	124.44	128.60
84	Aa	1728	G	C5-C6-O6	-6.93	124.44	128.60
84	Aa	1348	G	O4'-C1'-N9	6.93	113.75	108.20
84	Aa	2022	U	O4'-C1'-N1	6.93	113.75	108.20
84	Aa	2142	A	C5-C6-N6	-6.93	118.16	123.70
84	Aa	745	G	C5-C6-O6	-6.93	124.44	128.60
84	Aa	2338	C	O4'-C1'-N1	6.93	113.74	108.20
84	Aa	2377	C	O4'-C1'-N1	6.93	113.74	108.20
84	Aa	2415	U	O4'-C1'-N1	6.93	113.74	108.20
84	Aa	2683	A	C4-C5-C6	6.93	120.46	117.00
1	Ad	229	G	O4'-C1'-N9	6.93	113.74	108.20
84	Aa	699	C	N3-C4-C5	-6.93	119.13	121.90
84	Aa	797	U	O4'-C1'-N1	6.93	113.74	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	977	G	O4'-C1'-N9	6.92	113.74	108.20
84	Aa	78	U	O4'-C1'-N1	6.92	113.74	108.20
84	Aa	2760	U	O4'-C1'-N1	6.92	113.74	108.20
84	Aa	653	A	O4'-C1'-N9	6.92	113.74	108.20
84	Aa	984	A	O4'-C1'-N9	6.92	113.74	108.20
84	Aa	3125	G	N1-C6-O6	6.92	124.05	119.90
1	Ad	1154	G	C1'-O4'-C4'	-6.92	104.36	109.90
84	Aa	1336	A	C4-C5-C6	6.92	120.46	117.00
84	Aa	1341	G	O4'-C1'-N9	6.92	113.74	108.20
84	Aa	3048	C	O4'-C1'-N1	6.92	113.74	108.20
84	Aa	289	C	N3-C4-C5	-6.92	119.13	121.90
84	Aa	1226	G	C5-C6-O6	-6.92	124.45	128.60
86	Ab	6	C	O4'-C1'-N1	6.92	113.74	108.20
84	Aa	622	U	O4'-C1'-N1	6.92	113.73	108.20
84	Aa	1801	G	C5-C6-O6	-6.92	124.45	128.60
84	Aa	15	C	N3-C4-N4	6.92	122.84	118.00
84	Aa	2961	C	O4'-C1'-N1	6.92	113.73	108.20
84	Aa	494	C	O4'-C1'-N1	6.92	113.73	108.20
84	Aa	1137	G	O4'-C1'-N9	6.91	113.73	108.20
84	Aa	1158	C	N3-C4-C5	-6.91	119.13	121.90
84	Aa	1430	C	O4'-C1'-N1	6.91	113.73	108.20
84	Aa	2568	G	O4'-C1'-N9	6.91	113.73	108.20
84	Aa	554	C	C2'-C3'-O3'	-6.91	94.29	109.50
84	Aa	1279	C	O4'-C1'-N1	6.91	113.73	108.20
1	Ad	1486	U	O4'-C1'-N1	6.91	113.73	108.20
84	Aa	381	G	C5-C6-O6	-6.91	124.45	128.60
84	Aa	2958	A	C5-C6-N6	-6.91	118.17	123.70
84	Aa	2997	C	C6-N1-C1'	-6.91	112.51	120.80
84	Aa	3215	U	O4'-C1'-N1	6.91	113.73	108.20
84	Aa	18	G	O4'-C1'-N9	6.91	113.73	108.20
84	Aa	638	G	C5-C6-O6	-6.91	124.45	128.60
84	Aa	2490	U	O4'-C1'-N1	6.91	113.73	108.20
86	Ab	55	A	C5-C6-N1	-6.91	114.25	117.70
84	Aa	487	C	N3-C4-N4	6.91	122.83	118.00
84	Aa	2818	G	C5-C6-O6	-6.91	124.46	128.60
1	Ad	179	A	C1'-O4'-C4'	-6.91	104.38	109.90
1	Ad	212	A	O4'-C1'-N9	-6.91	102.68	108.20
1	Ad	273	C	N1-C1'-C2'	6.91	122.98	114.00
84	Aa	1562	A	C5'-C4'-O4'	6.91	117.39	109.10
84	Aa	400	G	O4'-C1'-N9	6.90	113.72	108.20
84	Aa	3059	C	O4'-C1'-N1	6.90	113.72	108.20
1	Ad	1092	A	N9-C1'-C2'	6.90	122.97	114.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1658	G	C5-C6-O6	-6.90	124.46	128.60
1	Ad	729	C	O4'-C1'-N1	6.90	113.72	108.20
1	Ad	738	U	O4'-C1'-N1	6.90	113.72	108.20
1	Ad	1434	G	N9-C1'-C2'	6.90	122.97	114.00
84	Aa	3286	G	O4'-C1'-N9	6.90	113.72	108.20
85	Ac	120	G	C5-C6-O6	-6.90	124.46	128.60
1	Ad	473	C	N1-C1'-C2'	6.90	122.97	114.00
1	Ad	1327	C	C3'-C2'-C1'	6.90	107.02	101.50
84	Aa	640	C	O3'-P-O5'	6.90	117.11	104.00
84	Aa	1612	C	N3-C4-C5	-6.90	119.14	121.90
85	Ac	66	G	O4'-C1'-N9	6.90	113.72	108.20
1	Ad	295	C	C3'-C2'-C1'	6.90	107.02	101.50
84	Aa	22	G	C5-C6-O6	-6.90	124.46	128.60
84	Aa	1621	G	O4'-C1'-N9	6.90	113.72	108.20
84	Aa	3315	A	C5-C6-N6	-6.90	118.18	123.70
85	Ac	42	G	O4'-C1'-N9	6.90	113.72	108.20
86	Ab	77	A	C5-C6-N6	-6.90	118.18	123.70
1	Ad	1309	U	C1'-O4'-C4'	6.89	115.42	109.90
84	Aa	926	C	O4'-C1'-N1	6.89	113.72	108.20
1	Ad	883	G	O4'-C1'-N9	6.89	113.72	108.20
84	Aa	1677	G	O4'-C1'-N9	6.89	113.71	108.20
84	Aa	2564	G	N1-C6-O6	6.89	124.03	119.90
1	Ad	433	G	O4'-C1'-N9	6.89	113.71	108.20
84	Aa	284	U	O4'-C1'-N1	6.89	113.71	108.20
84	Aa	552	G	C4'-C3'-O3'	-6.89	94.93	109.40
84	Aa	1066	G	C5-C6-O6	-6.89	124.47	128.60
84	Aa	2202	A	C4-C5-C6	6.89	120.44	117.00
84	Aa	3243	C	N3-C4-N4	6.89	122.82	118.00
84	Aa	3361	G	C5-C6-O6	-6.89	124.47	128.60
84	Aa	151	U	O4'-C1'-N1	6.89	113.71	108.20
84	Aa	295	U	O4'-C1'-N1	6.89	113.71	108.20
84	Aa	883	G	N1-C6-O6	6.89	124.03	119.90
1	Ad	1445	C	O4'-C1'-C2'	-6.89	98.91	105.80
2	Ae	47	U	C1'-O4'-C4'	6.89	115.41	109.90
84	Aa	60	G	C5-C6-O6	-6.89	124.47	128.60
84	Aa	663	G	C5-C6-O6	-6.89	124.47	128.60
84	Aa	859	G	C5-C6-O6	-6.89	124.47	128.60
84	Aa	1644	A	C4-C5-C6	6.89	120.44	117.00
86	Ab	73	U	C1'-O4'-C4'	-6.88	104.39	109.90
1	Ad	1464	G	C3'-C2'-C1'	-6.88	96.00	101.50
84	Aa	563	C	O4'-C1'-N1	6.88	113.70	108.20
84	Aa	1189	G	C5-C6-O6	-6.88	124.47	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1904	A	C5-C6-N6	-6.88	118.19	123.70
1	Ad	881	G	O4'-C1'-N9	6.88	113.70	108.20
84	Aa	1432	G	C5-C6-O6	-6.88	124.47	128.60
84	Aa	3138	C	N3-C4-N4	6.88	122.82	118.00
84	Aa	787	G	C5-C6-O6	-6.88	124.47	128.60
84	Aa	3008	U	O4'-C1'-N1	6.88	113.70	108.20
84	Aa	3054	G	O4'-C1'-N9	6.88	113.70	108.20
84	Aa	3076	C	N3-C4-N4	6.88	122.81	118.00
84	Aa	531	G	O4'-C1'-N9	6.88	113.70	108.20
84	Aa	927	G	C5-C6-O6	-6.88	124.47	128.60
84	Aa	2170	G	C5-C6-O6	-6.88	124.47	128.60
1	Ad	1212	A	O4'-C1'-N9	6.88	113.70	108.20
1	Ad	1618	G	C1'-O4'-C4'	-6.88	104.40	109.90
73	CO	117	TYR	CB-CG-CD2	-6.88	116.88	121.00
1	Ad	273	C	O4'-C1'-N1	6.87	113.70	108.20
84	Aa	121	A	O4'-C1'-N9	6.87	113.70	108.20
84	Aa	1925	G	C5-C6-O6	-6.87	124.48	128.60
1	Ad	1152	A	C3'-C2'-C1'	6.87	107.00	101.50
84	Aa	485	G	O4'-C1'-N9	6.87	113.70	108.20
84	Aa	533	G	O4'-C1'-N9	6.87	113.70	108.20
84	Aa	2417	G	O4'-C1'-N9	6.87	113.70	108.20
84	Aa	3320	G	O4'-C1'-N9	6.87	113.70	108.20
85	Ac	159	G	C5-C6-O6	-6.87	124.48	128.60
84	Aa	763	G	C5-C6-O6	-6.87	124.48	128.60
84	Aa	1878	G	O4'-C1'-N9	6.87	113.69	108.20
84	Aa	2313	U	O4'-C1'-N1	6.87	113.70	108.20
84	Aa	2455	A	N1-C6-N6	6.87	122.72	118.60
84	Aa	2854	C	O4'-C1'-N1	6.87	113.69	108.20
84	Aa	1292	U	O4'-C1'-N1	6.87	113.69	108.20
84	Aa	3042	U	P-O5'-C5'	6.87	131.89	120.90
84	Aa	3027	G	C5-C6-O6	-6.87	124.48	128.60
84	Aa	3046	C	N3-C4-C5	-6.87	119.15	121.90
84	Aa	3236	A	O4'-C1'-N9	6.87	113.69	108.20
86	Ab	98	G	N1-C2-N3	-6.87	119.78	123.90
1	Ad	1613	G	C1'-O4'-C4'	-6.86	104.41	109.90
84	Aa	1537	A	O4'-C1'-N9	6.86	113.69	108.20
84	Aa	1912	U	O4'-C1'-N1	6.86	113.69	108.20
84	Aa	2089	A	O4'-C1'-N9	6.86	113.69	108.20
84	Aa	2149	G	N1-C6-O6	6.86	124.02	119.90
84	Aa	2438	A	O4'-C1'-N9	6.86	113.69	108.20
86	Ab	69	A	C4-C5-C6	6.86	120.43	117.00
84	Aa	692	U	O4'-C1'-N1	6.86	113.69	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	580	C	O4'-C1'-N1	6.86	113.69	108.20
84	Aa	962	C	O4'-C1'-N1	6.86	113.69	108.20
84	Aa	1116	G	C5-C6-O6	-6.86	124.48	128.60
84	Aa	2802	G	C5-C6-O6	-6.86	124.48	128.60
84	Aa	2986	C	O4'-C1'-N1	6.86	113.69	108.20
84	Aa	3208	G	C5-C6-O6	-6.86	124.48	128.60
84	Aa	3363	G	O4'-C1'-N9	6.86	113.69	108.20
1	Ad	1172	G	O4'-C1'-C2'	6.86	113.77	107.60
85	Ac	41	A	C5-C6-N6	-6.86	118.21	123.70
1	Ad	1677	U	O4'-C1'-N1	6.86	113.69	108.20
84	Aa	1810	G	C5-C6-O6	-6.86	124.48	128.60
1	Ad	848	C	O4'-C1'-C2'	-6.86	98.94	105.80
84	Aa	527	G	C5-C6-O6	-6.86	124.49	128.60
1	Ad	1190	U	N1-C1'-C2'	-6.85	104.46	112.00
1	Ad	1213	C	N1-C1'-C2'	6.85	122.91	114.00
2	Ae	9	A	O4'-C1'-C2'	-6.85	98.95	105.80
84	Aa	119	A	C5-C6-N6	-6.85	118.22	123.70
84	Aa	1134	G	C5-C6-O6	-6.85	124.49	128.60
84	Aa	1822	C	N3-C4-C5	-6.85	119.16	121.90
84	Aa	2115	G	C5-C6-O6	-6.85	124.49	128.60
84	Aa	2996	A	P-O3'-C3'	6.85	127.92	119.70
1	Ad	356	G	C3'-C2'-C1'	6.85	106.98	101.50
84	Aa	2292	U	O4'-C1'-N1	6.85	113.68	108.20
84	Aa	294	A	C4-C5-C6	6.85	120.42	117.00
84	Aa	964	C	O4'-C1'-N1	6.85	113.68	108.20
84	Aa	1411	G	O4'-C1'-N9	6.85	113.68	108.20
84	Aa	1524	G	C5-C6-O6	-6.85	124.49	128.60
84	Aa	2212	U	O4'-C1'-N1	6.85	113.68	108.20
86	Ab	47	C	O4'-C1'-N1	6.85	113.68	108.20
84	Aa	265	G	C5-C6-O6	-6.85	124.49	128.60
84	Aa	1545	G	O4'-C1'-N9	6.85	113.68	108.20
1	Ad	60	C	N1-C1'-C2'	6.85	122.90	114.00
84	Aa	2791	U	O4'-C1'-N1	6.85	113.68	108.20
1	Ad	719	C	C3'-C2'-C1'	6.84	106.97	101.50
1	Ad	1221	A	C1'-O4'-C4'	6.84	115.38	109.90
1	Ad	1582	G	O4'-C1'-C2'	-6.84	98.95	105.80
84	Aa	798	G	C5-C6-O6	-6.84	124.49	128.60
1	Ad	1112	G	O4'-C1'-C2'	6.84	113.76	107.60
1	Ad	25	C	P-O3'-C3'	6.84	127.91	119.70
84	Aa	3005	C	O4'-C1'-N1	6.84	113.67	108.20
1	Ad	706	U	C5'-C4'-C3'	-6.84	105.06	116.00
84	Aa	881	G	C5-C6-O6	-6.84	124.50	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1825	G	O4'-C1'-N9	6.84	113.67	108.20
84	Aa	2078	G	O4'-C1'-N9	6.84	113.67	108.20
1	Ad	1251	U	O4'-C1'-N1	6.84	113.67	108.20
84	Aa	2995	G	C5-C6-O6	-6.84	124.50	128.60
1	Ad	339	G	C3'-C2'-C1'	-6.84	96.03	101.50
1	Ad	1680	A	C3'-C2'-C1'	6.84	106.97	101.50
84	Aa	2348	U	O4'-C1'-N1	6.84	113.67	108.20
84	Aa	2652	G	O4'-C1'-N9	6.84	113.67	108.20
84	Aa	2949	G	N1-C6-O6	6.84	124.00	119.90
1	Ad	351	G	O4'-C1'-C2'	6.83	113.75	107.60
1	Ad	1324	U	O4'-C1'-N1	6.83	113.67	108.20
84	Aa	37	U	O4'-C1'-N1	6.83	113.67	108.20
84	Aa	1760	G	C5-C6-O6	-6.83	124.50	128.60
84	Aa	3144	U	O4'-C1'-N1	6.83	113.67	108.20
84	Aa	3245	G	C5-C6-O6	-6.83	124.50	128.60
1	Ad	728	C	N1-C1'-C2'	6.83	122.88	114.00
84	Aa	1871	G	N1-C6-O6	6.83	124.00	119.90
84	Aa	2832	G	O4'-C1'-N9	6.83	113.67	108.20
84	Aa	2898	A	C5-C6-N1	-6.83	114.28	117.70
84	Aa	3102	G	O4'-C1'-N9	6.83	113.67	108.20
1	Ad	262	U	N1-C1'-C2'	-6.83	104.49	112.00
1	Ad	929	A	O4'-C1'-N9	6.83	113.66	108.20
84	Aa	2171	A	C5-C6-N6	-6.83	118.24	123.70
86	Ab	74	A	C4-C5-C6	6.83	120.41	117.00
84	Aa	848	G	C5-C6-O6	-6.83	124.50	128.60
84	Aa	2246	G	O4'-C1'-N9	6.83	113.66	108.20
1	Ad	971	A	O4'-C1'-C2'	-6.83	98.97	105.80
84	Aa	355	C	N3-C4-C5	-6.83	119.17	121.90
84	Aa	1759	C	C6-N1-C1'	-6.83	112.61	120.80
84	Aa	1958	G	C5'-C4'-C3'	-6.83	105.08	116.00
84	Aa	2305	U	O4'-C1'-N1	6.83	113.66	108.20
84	Aa	2506	G	N1-C6-O6	6.83	124.00	119.90
1	Ad	416	A	C1'-O4'-C4'	6.82	115.36	109.90
1	Ad	1143	A	O4'-C1'-N9	6.82	113.66	108.20
84	Aa	241	G	O4'-C1'-N9	6.82	113.66	108.20
84	Aa	1161	G	C5-C6-O6	-6.82	124.51	128.60
84	Aa	2182	G	C5-C6-O6	-6.82	124.50	128.60
84	Aa	2640	A	C4-C5-C6	6.82	120.41	117.00
86	Ab	27	A	C5-N7-C8	6.82	107.31	103.90
86	Ab	75	G	C5-C6-O6	-6.82	124.51	128.60
1	Ad	1590	U	C1'-O4'-C4'	6.82	115.36	109.90
84	Aa	849	A	C5-C6-N6	-6.82	118.24	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1502	U	O4'-C1'-N1	6.82	113.66	108.20
84	Aa	2657	C	N3-C4-C5	-6.82	119.17	121.90
1	Ad	419	C	O4'-C1'-C2'	-6.82	98.98	105.80
1	Ad	1212	A	O4'-C1'-C2'	6.82	113.74	107.60
84	Aa	1100	G	C5-C6-O6	-6.82	124.51	128.60
84	Aa	1357	C	N3-C4-C5	-6.82	119.17	121.90
84	Aa	2735	G	C5-C6-O6	-6.82	124.51	128.60
85	Ac	83	C	O4'-C1'-N1	6.82	113.66	108.20
84	Aa	3348	G	N1-C6-O6	6.82	123.99	119.90
84	Aa	263	A	P-O3'-C3'	6.82	127.88	119.70
84	Aa	1260	G	C5-C6-O6	-6.82	124.51	128.60
86	Ab	50	A	C5-C6-N6	-6.82	118.25	123.70
1	Ad	804	C	O4'-C1'-N1	6.81	113.65	108.20
1	Ad	936	C	C3'-C2'-C1'	6.81	106.95	101.50
1	Ad	1593	U	O4'-C1'-C2'	6.81	113.73	107.60
48	CD	2	SER	N-CA-CB	6.81	120.72	110.50
84	Aa	1269	U	O4'-C1'-N1	6.81	113.65	108.20
86	Ab	58	G	N3-C2-N2	6.81	124.67	119.90
1	Ad	498	U	O4'-C1'-C2'	-6.81	98.99	105.80
84	Aa	683	U	O4'-C1'-N1	6.81	113.65	108.20
1	Ad	79	A	O4'-C4'-C3'	-6.81	97.19	104.00
84	Aa	123	U	O4'-C1'-N1	6.81	113.65	108.20
84	Aa	2867	U	O4'-C1'-N1	6.81	113.65	108.20
1	Ad	1193	A	O4'-C1'-C2'	-6.81	98.99	105.80
84	Aa	2582	G	C5-C6-O6	-6.81	124.52	128.60
84	Aa	3263	C	C5'-C4'-O4'	6.81	117.27	109.10
1	Ad	171	G	O4'-C1'-N9	6.81	113.64	108.20
1	Ad	762	A	O4'-C1'-C2'	-6.81	98.99	105.80
84	Aa	12	G	O4'-C1'-N9	6.81	113.64	108.20
84	Aa	1407	G	C5-C6-O6	-6.81	124.52	128.60
84	Aa	1547	G	C5-C6-O6	-6.81	124.52	128.60
1	Ad	59	G	O4'-C1'-N9	6.80	113.64	108.20
1	Ad	1434	G	O4'-C1'-C2'	6.80	113.72	107.60
84	Aa	800	C	N3-C4-C5	-6.80	119.18	121.90
84	Aa	998	G	C5-C6-O6	-6.80	124.52	128.60
84	Aa	1016	G	O4'-C1'-N9	6.80	113.64	108.20
85	Ac	72	A	C4-C5-C6	6.80	120.40	117.00
84	Aa	3103	G	O4'-C1'-N9	6.80	113.64	108.20
85	Ac	139	C	O4'-C1'-N1	6.80	113.64	108.20
84	Aa	76	A	C5-C6-N1	-6.80	114.30	117.70
84	Aa	808	G	P-O3'-C3'	6.80	127.86	119.70
84	Aa	1158	C	O4'-C1'-N1	6.80	113.64	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2745	C	O4'-C1'-N1	6.80	113.64	108.20
61	CM	6	PHE	CB-CG-CD2	6.80	125.56	120.80
84	Aa	170	C	N3-C4-N4	6.80	122.76	118.00
84	Aa	413	G	C5-C6-O6	-6.80	124.52	128.60
84	Aa	1561	U	P-O3'-C3'	6.80	127.86	119.70
84	Aa	1600	A	O4'-C1'-N9	6.80	113.64	108.20
84	Aa	2146	A	C5-C6-N1	-6.80	114.30	117.70
84	Aa	3184	G	O4'-C1'-N9	6.80	113.64	108.20
84	Aa	1762	G	C5-C6-O6	-6.80	124.52	128.60
84	Aa	2520	U	O4'-C1'-N1	6.80	113.64	108.20
1	Ad	351	G	N9-C1'-C2'	6.80	122.83	114.00
2	Ae	24	A	O4'-C1'-N9	6.80	113.64	108.20
84	Aa	1931	G	C5-C6-O6	-6.80	124.52	128.60
84	Aa	2948	A	C5-C6-N6	-6.80	118.26	123.70
1	Ad	214	A	P-O3'-C3'	6.79	127.85	119.70
1	Ad	442	A	O4'-C1'-N9	6.79	113.64	108.20
84	Aa	1525	U	O4'-C1'-N1	6.79	113.64	108.20
84	Aa	3001	G	C5-C6-O6	-6.79	124.52	128.60
1	Ad	1046	G	O4'-C1'-N9	6.79	113.63	108.20
84	Aa	3336	A	C5-C6-N6	-6.79	118.27	123.70
86	Ab	36	C	C6-N1-C2	-6.79	117.58	120.30
86	Ab	73	U	O4'-C1'-N1	6.79	113.63	108.20
84	Aa	1136	A	C5-C6-N6	-6.79	118.27	123.70
84	Aa	1216	G	N1-C6-O6	6.79	123.97	119.90
85	Ac	28	C	N3-C4-C5	-6.79	119.18	121.90
1	Ad	963	U	O4'-C1'-N1	6.79	113.63	108.20
84	Aa	3355	U	O4'-C1'-N1	6.79	113.63	108.20
86	Ab	11	A	C4-C5-C6	6.79	120.39	117.00
1	Ad	1184	C	N1-C1'-C2'	6.79	122.83	114.00
1	Ad	1698	A	P-O3'-C3'	6.79	127.85	119.70
84	Aa	429	G	C5-C6-O6	-6.79	124.53	128.60
84	Aa	1054	U	O4'-C1'-N1	6.79	113.63	108.20
84	Aa	2301	C	O4'-C1'-N1	6.79	113.63	108.20
84	Aa	3129	G	C5-C6-O6	-6.79	124.53	128.60
84	Aa	152	C	N3-C4-N4	6.79	122.75	118.00
84	Aa	2849	A	C5-C6-N6	-6.79	118.27	123.70
84	Aa	208	G	C5-C6-O6	-6.79	124.53	128.60
84	Aa	629	U	O4'-C1'-N1	6.79	113.63	108.20
84	Aa	1676	A	C4-C5-C6	6.79	120.39	117.00
84	Aa	2309	U	O4'-C1'-N1	6.79	113.63	108.20
84	Aa	2678	C	O4'-C1'-N1	6.79	113.63	108.20
84	Aa	2989	A	C5-C6-N6	-6.79	118.27	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3382	A	C4-C5-C6	6.79	120.39	117.00
86	Ab	81	G	O4'-C1'-N9	6.79	113.63	108.20
84	Aa	2663	U	O4'-C1'-N1	6.78	113.63	108.20
84	Aa	2935	A	C4-C5-C6	6.78	120.39	117.00
84	Aa	3183	G	C5-C6-O6	-6.78	124.53	128.60
1	Ad	1125	U	C1'-O4'-C4'	6.78	115.33	109.90
84	Aa	73	A	C5-C6-N1	-6.78	114.31	117.70
84	Aa	435	G	O4'-C1'-N9	6.78	113.62	108.20
84	Aa	776	G	O4'-C1'-N9	6.78	113.62	108.20
84	Aa	833	G	C5-C6-O6	-6.78	124.53	128.60
84	Aa	1508	C	O4'-C1'-N1	6.78	113.62	108.20
84	Aa	3353	G	C5-C6-O6	-6.78	124.53	128.60
1	Ad	1363	G	O4'-C1'-C2'	6.78	113.70	107.60
84	Aa	3273	C	O4'-C1'-N1	6.78	113.62	108.20
1	Ad	81	U	O4'-C1'-N1	6.78	113.62	108.20
1	Ad	1660	C	O4'-C1'-N1	6.78	113.62	108.20
84	Aa	207	U	O4'-C1'-N1	6.78	113.62	108.20
84	Aa	1480	G	C5-C6-O6	-6.78	124.53	128.60
84	Aa	1547	G	O4'-C1'-N9	6.78	113.62	108.20
84	Aa	2170	G	O4'-C1'-N9	6.78	113.62	108.20
86	Ab	7	G	C5-C6-N1	-6.78	108.11	111.50
84	Aa	324	U	O4'-C1'-N1	6.78	113.62	108.20
84	Aa	2331	A	C5-C6-N6	-6.78	118.28	123.70
84	Aa	2748	G	C5-C6-O6	-6.78	124.53	128.60
2	Ae	1	U	O4'-C1'-N1	6.77	113.62	108.20
84	Aa	480	C	O4'-C1'-N1	6.77	113.62	108.20
84	Aa	1299	G	O4'-C1'-N9	6.77	113.62	108.20
84	Aa	1352	G	O4'-C1'-N9	6.77	113.62	108.20
84	Aa	1675	G	O4'-C1'-N9	6.77	113.62	108.20
84	Aa	2140	C	O4'-C1'-N1	6.77	113.62	108.20
84	Aa	2375	G	C5-C6-O6	-6.77	124.54	128.60
1	Ad	1013	G	C1'-O4'-C4'	-6.77	104.48	109.90
71	CB	197	TYR	CB-CG-CD2	6.77	125.06	121.00
84	Aa	105	A	O4'-C1'-N9	6.77	113.62	108.20
1	Ad	524	A	C1'-O4'-C4'	-6.77	104.48	109.90
71	CB	118	PHE	CB-CG-CD1	6.77	125.54	120.80
84	Aa	17	G	C5-C6-O6	-6.77	124.54	128.60
84	Aa	85	G	C5-C6-O6	-6.77	124.54	128.60
84	Aa	156	A	C5-C6-N6	-6.77	118.28	123.70
84	Aa	368	U	O4'-C1'-N1	6.77	113.62	108.20
84	Aa	386	G	C5-C6-O6	-6.77	124.54	128.60
84	Aa	1374	G	C5-C6-O6	-6.77	124.54	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1573	G	O4'-C1'-N9	6.77	113.61	108.20
84	Aa	2393	G	O4'-C1'-N9	6.77	113.61	108.20
84	Aa	2943	A	C4-C5-C6	6.77	120.38	117.00
84	Aa	271	G	O4'-C1'-N9	6.77	113.61	108.20
84	Aa	2031	G	O4'-C1'-N9	6.77	113.61	108.20
84	Aa	2991	U	O4'-C1'-N1	6.77	113.61	108.20
1	Ad	1192	G	O4'-C1'-C2'	6.76	113.69	107.60
84	Aa	1918	A	C5-C6-N6	-6.76	118.29	123.70
84	Aa	2537	G	N1-C6-O6	6.76	123.96	119.90
84	Aa	3178	C	C6-N1-C1'	-6.76	112.68	120.80
84	Aa	1283	C	N3-C4-C5	-6.76	119.19	121.90
84	Aa	3015	U	O4'-C1'-N1	6.76	113.61	108.20
84	Aa	2574	A	O4'-C1'-N9	6.76	113.61	108.20
84	Aa	3076	C	O4'-C1'-N1	6.76	113.61	108.20
84	Aa	3210	G	C5-C6-O6	-6.76	124.54	128.60
1	Ad	922	U	N1-C1'-C2'	6.76	122.79	114.00
84	Aa	1180	C	C6-N1-C1'	-6.76	112.69	120.80
84	Aa	1588	G	C5-C6-O6	-6.76	124.54	128.60
84	Aa	2465	G	O4'-C1'-N9	6.76	113.61	108.20
84	Aa	2893	U	O4'-C1'-N1	6.76	113.61	108.20
84	Aa	719	U	O4'-C1'-N1	6.76	113.61	108.20
84	Aa	2563	G	C5-C6-O6	-6.76	124.55	128.60
1	Ad	1728	G	P-O5'-C5'	6.76	131.71	120.90
84	Aa	1688	U	O4'-C1'-N1	6.76	113.61	108.20
86	Ab	37	G	N3-C2-N2	6.76	124.63	119.90
1	Ad	264	G	O4'-C1'-N9	6.75	113.60	108.20
1	Ad	1013	G	O4'-C1'-N9	6.75	113.60	108.20
84	Aa	180	G	C5-C6-O6	-6.75	124.55	128.60
84	Aa	43	U	O4'-C1'-N1	6.75	113.60	108.20
84	Aa	594	C	N3-C4-C5	-6.75	119.20	121.90
84	Aa	600	G	C5-C6-O6	-6.75	124.55	128.60
84	Aa	986	G	O4'-C1'-N9	6.75	113.60	108.20
84	Aa	1004	C	N3-C4-C5	-6.75	119.20	121.90
84	Aa	1949	G	C5-C6-O6	-6.75	124.55	128.60
84	Aa	2370	G	C5-C6-O6	-6.75	124.55	128.60
84	Aa	2608	G	C5-C6-O6	-6.75	124.55	128.60
84	Aa	3315	A	O4'-C1'-N9	6.75	113.60	108.20
1	Ad	1596	G	O4'-C1'-C2'	-6.75	99.05	105.80
84	Aa	919	G	C5-C6-O6	-6.75	124.55	128.60
86	Ab	85	G	O4'-C1'-N9	6.75	113.60	108.20
1	Ad	789	C	P-O3'-C3'	6.75	127.80	119.70
84	Aa	1808	G	C5-C6-O6	-6.75	124.55	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3160	G	C5-C6-O6	-6.75	124.55	128.60
1	Ad	189	U	C1'-O4'-C4'	6.75	115.30	109.90
84	Aa	715	A	N1-C6-N6	6.75	122.65	118.60
84	Aa	2787	A	C5-C6-N6	-6.75	118.30	123.70
63	CU	92	TYR	CB-CG-CD2	-6.75	116.95	121.00
84	Aa	627	G	O4'-C1'-N9	6.75	113.60	108.20
84	Aa	1076	G	C5-C6-O6	-6.75	124.55	128.60
84	Aa	3182	A	O5'-C5'-C4'	6.75	124.52	111.70
84	Aa	736	U	O4'-C1'-N1	6.75	113.60	108.20
84	Aa	838	G	C5-C6-O6	-6.75	124.55	128.60
84	Aa	2731	G	C5-C6-O6	-6.75	124.55	128.60
84	Aa	2968	G	C5-C6-O6	-6.75	124.55	128.60
1	Ad	1547	G	O4'-C1'-N9	6.74	113.59	108.20
1	Ad	1687	G	N9-C1'-C2'	6.74	122.77	114.00
84	Aa	2420	U	O4'-C1'-N1	6.74	113.59	108.20
85	Ac	36	G	C5-C6-O6	-6.74	124.55	128.60
84	Aa	2555	G	C5-C6-O6	-6.74	124.56	128.60
1	Ad	79	A	O4'-C1'-C2'	-6.74	99.06	105.80
84	Aa	226	U	O4'-C1'-N1	6.74	113.59	108.20
84	Aa	1659	G	O4'-C1'-N9	6.74	113.59	108.20
84	Aa	2021	G	C5-C6-O6	-6.74	124.56	128.60
84	Aa	2266	A	C4-C5-C6	6.74	120.37	117.00
84	Aa	3064	U	O4'-C1'-N1	6.74	113.59	108.20
84	Aa	3102	G	C5-C6-O6	-6.74	124.56	128.60
84	Aa	610	G	C8-N9-C4	-6.74	103.70	106.40
84	Aa	901	U	O4'-C1'-N1	6.74	113.59	108.20
84	Aa	1604	U	O4'-C1'-N1	6.74	113.59	108.20
84	Aa	1933	U	O4'-C1'-N1	6.74	113.59	108.20
84	Aa	2526	G	O4'-C1'-N9	6.74	113.59	108.20
1	Ad	295	C	O4'-C1'-N1	6.74	113.59	108.20
1	Ad	789	C	O4'-C1'-C2'	-6.74	99.06	105.80
84	Aa	441	G	C5-C6-O6	-6.74	124.56	128.60
84	Aa	587	A	C5-C6-N6	-6.74	118.31	123.70
84	Aa	1375	G	C5-C6-O6	-6.74	124.56	128.60
86	Ab	16	A	C5-C6-N6	-6.74	118.31	123.70
84	Aa	1212	U	O4'-C1'-N1	6.73	113.59	108.20
84	Aa	2126	C	O4'-C1'-N1	6.73	113.59	108.20
84	Aa	2434	G	O4'-C1'-N9	6.73	113.59	108.20
84	Aa	57	G	C5-C6-O6	-6.73	124.56	128.60
84	Aa	214	G	N1-C6-O6	6.73	123.94	119.90
84	Aa	581	G	O4'-C1'-N9	6.73	113.59	108.20
84	Aa	2014	A	C4-C5-C6	6.73	120.37	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2217	A	C4-C5-C6	6.73	120.37	117.00
84	Aa	3204	G	C5-C6-O6	-6.73	124.56	128.60
84	Aa	508	G	C5-C6-O6	-6.73	124.56	128.60
84	Aa	703	G	C5-C6-O6	-6.73	124.56	128.60
84	Aa	1529	C	N3-C4-C5	-6.73	119.21	121.90
84	Aa	1583	G	C5-C6-O6	-6.73	124.56	128.60
84	Aa	2465	G	C5-C6-O6	-6.73	124.56	128.60
1	Ad	1809	U	O4'-C1'-N1	6.73	113.58	108.20
84	Aa	81	C	O4'-C1'-N1	6.73	113.58	108.20
84	Aa	949	C	N3-C4-C5	-6.73	119.21	121.90
84	Aa	2013	G	C5-C6-O6	-6.73	124.56	128.60
84	Aa	2758	C	N3-C4-C5	-6.73	119.21	121.90
84	Aa	3066	G	O4'-C1'-N9	6.73	113.58	108.20
84	Aa	404	G	C5-C6-O6	-6.73	124.56	128.60
84	Aa	1102	A	O4'-C1'-N9	6.73	113.58	108.20
84	Aa	1555	G	C5-C6-O6	-6.73	124.56	128.60
84	Aa	1567	G	O4'-C1'-N9	6.73	113.58	108.20
84	Aa	2624	G	C5-C6-O6	-6.73	124.56	128.60
84	Aa	2901	C	C2-N1-C1'	6.73	126.20	118.80
84	Aa	100	C	C2-N1-C1'	6.72	126.20	118.80
84	Aa	890	G	C5-C6-O6	-6.72	124.56	128.60
84	Aa	1550	A	O4'-C1'-N9	6.72	113.58	108.20
1	Ad	445	A	O4'-C1'-N9	6.72	113.58	108.20
1	Ad	956	A	O4'-C1'-C2'	-6.72	99.08	105.80
84	Aa	265	G	P-O3'-C3'	6.72	127.77	119.70
84	Aa	333	G	O4'-C1'-N9	6.72	113.58	108.20
84	Aa	349	A	C5-C6-N6	-6.72	118.32	123.70
84	Aa	2345	C	P-O5'-C5'	6.72	131.66	120.90
84	Aa	2621	G	O4'-C1'-N9	6.72	113.58	108.20
1	Ad	1589	C	N1-C1'-C2'	6.72	122.74	114.00
84	Aa	2653	U	O4'-C1'-N1	6.72	113.58	108.20
84	Aa	3325	G	C5-C6-O6	-6.72	124.57	128.60
84	Aa	1510	G	C5-C6-O6	-6.72	124.57	128.60
84	Aa	1681	U	O4'-C1'-N1	6.72	113.58	108.20
84	Aa	2741	G	C5-C6-O6	-6.72	124.57	128.60
86	Ab	10	C	N3-C4-C5	-6.72	119.21	121.90
84	Aa	1671	G	O4'-C1'-N9	6.72	113.58	108.20
84	Aa	2648	G	C5-C6-O6	-6.72	124.57	128.60
84	Aa	612	U	O4'-C1'-N1	6.72	113.57	108.20
84	Aa	1116	G	O4'-C1'-N9	6.72	113.57	108.20
84	Aa	1668	U	O4'-C1'-N1	6.72	113.57	108.20
84	Aa	2052	G	O4'-C1'-N9	6.72	113.57	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3021	U	O4'-C1'-N1	6.72	113.57	108.20
1	Ad	717	G	N9-C1'-C2'	6.71	122.73	114.00
1	Ad	868	A	O4'-C1'-C2'	-6.71	99.09	105.80
84	Aa	1544	G	C5-C6-O6	-6.71	124.57	128.60
84	Aa	2399	G	C5-C6-O6	-6.71	124.57	128.60
1	Ad	562	U	C1'-O4'-C4'	-6.71	104.53	109.90
84	Aa	1065	A	C5-C6-N1	-6.71	114.34	117.70
84	Aa	2413	G	C5-C6-O6	-6.71	124.57	128.60
84	Aa	2924	G	C5-C6-O6	-6.71	124.57	128.60
84	Aa	172	A	C5-C6-N1	-6.71	114.34	117.70
84	Aa	1844	U	O4'-C1'-N1	6.71	113.57	108.20
84	Aa	3320	G	N1-C6-O6	6.71	123.93	119.90
86	Ab	39	C	O4'-C1'-N1	6.71	113.57	108.20
84	Aa	601	G	C5-C6-O6	-6.71	124.57	128.60
1	Ad	955	C	C3'-C2'-C1'	6.71	106.87	101.50
1	Ad	1114	G	O4'-C1'-N9	6.71	113.57	108.20
84	Aa	2073	U	C2-N1-C1'	6.71	125.75	117.70
84	Aa	2539	G	C5-C6-O6	-6.71	124.58	128.60
84	Aa	2827	C	N3-C4-N4	6.71	122.69	118.00
84	Aa	3363	G	N1-C6-O6	6.71	123.92	119.90
1	Ad	1538	C	O4'-C1'-N1	6.71	113.56	108.20
84	Aa	68	U	O4'-C1'-N1	6.71	113.57	108.20
84	Aa	593	G	O4'-C1'-N9	6.71	113.56	108.20
84	Aa	1176	U	O4'-C1'-N1	6.71	113.56	108.20
84	Aa	3065	U	O4'-C1'-N1	6.71	113.56	108.20
1	Ad	225	G	N9-C1'-C2'	6.70	122.72	114.00
1	Ad	613	U	C3'-C2'-C1'	6.70	106.86	101.50
84	Aa	1084	G	C5-C6-O6	-6.70	124.58	128.60
84	Aa	2738	U	O4'-C1'-N1	6.70	113.56	108.20
84	Aa	3242	G	O4'-C1'-N9	6.70	113.56	108.20
84	Aa	1801	G	O4'-C1'-N9	6.70	113.56	108.20
2	Ae	35	U	N1-C1'-C2'	6.70	122.71	114.00
84	Aa	1177	G	N1-C6-O6	6.70	123.92	119.90
84	Aa	1765	G	N1-C6-O6	6.70	123.92	119.90
84	Aa	1893	G	O4'-C1'-N9	6.70	113.56	108.20
84	Aa	2482	A	O4'-C1'-N9	6.70	113.56	108.20
1	Ad	363	G	N9-C1'-C2'	-6.70	104.63	112.00
1	Ad	1472	G	C3'-C2'-C1'	-6.70	96.14	101.50
84	Aa	117	U	O4'-C1'-N1	6.70	113.56	108.20
84	Aa	304	A	C5-C6-N6	-6.70	118.34	123.70
84	Aa	2447	A	O4'-C1'-N9	6.70	113.56	108.20
84	Aa	2865	G	C5-C6-O6	-6.70	124.58	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3023	G	C5-C6-O6	-6.70	124.58	128.60
1	Ad	327	A	C3'-C2'-C1'	6.70	106.86	101.50
84	Aa	1545	G	C5-C6-O6	-6.70	124.58	128.60
84	Aa	1897	A	C4-C5-C6	6.70	120.35	117.00
86	Ab	7	G	N1-C2-N3	-6.70	119.88	123.90
1	Ad	968	A	N9-C1'-C2'	-6.70	104.64	112.00
84	Aa	601	G	O4'-C1'-N9	6.70	113.56	108.20
84	Aa	1344	A	C5-C6-N6	-6.70	118.34	123.70
84	Aa	2182	G	O4'-C1'-N9	6.70	113.56	108.20
84	Aa	995	C	O4'-C1'-N1	6.69	113.56	108.20
1	Ad	209	U	P-O3'-C3'	-6.69	111.67	119.70
84	Aa	548	G	C5-C6-O6	-6.69	124.58	128.60
84	Aa	1843	A	O4'-C1'-N9	6.69	113.55	108.20
84	Aa	1852	C	O4'-C1'-N1	6.69	113.55	108.20
84	Aa	2130	U	O4'-C1'-N1	6.69	113.55	108.20
84	Aa	3026	C	N3-C4-C5	-6.69	119.22	121.90
1	Ad	990	G	C1'-O4'-C4'	-6.69	104.55	109.90
1	Ad	1045	G	O4'-C1'-C2'	6.69	113.62	107.60
84	Aa	269	C	O4'-C1'-N1	6.69	113.55	108.20
84	Aa	1834	C	N3-C4-N4	6.69	122.68	118.00
84	Aa	1895	G	O4'-C1'-N9	6.69	113.55	108.20
84	Aa	2034	G	O4'-C1'-N9	6.69	113.55	108.20
84	Aa	965	A	C5-C6-N1	-6.69	114.36	117.70
84	Aa	1881	C	O4'-C1'-N1	6.69	113.55	108.20
1	Ad	551	U	O4'-C1'-N1	6.69	113.55	108.20
84	Aa	273	U	O4'-C1'-N1	6.69	113.55	108.20
84	Aa	1693	A	C4-C5-C6	6.69	120.34	117.00
84	Aa	179	G	O4'-C1'-N9	6.69	113.55	108.20
84	Aa	2459	U	O4'-C1'-N1	6.68	113.55	108.20
84	Aa	3117	G	C5-C6-O6	-6.68	124.59	128.60
86	Ab	10	C	O4'-C1'-N1	6.68	113.55	108.20
1	Ad	1618	G	O4'-C1'-C2'	6.68	113.61	107.60
84	Aa	1354	G	C5-C6-O6	-6.68	124.59	128.60
84	Aa	2039	G	O4'-C1'-N9	6.68	113.55	108.20
84	Aa	2091	U	O4'-C1'-N1	6.68	113.55	108.20
84	Aa	2530	G	O4'-C1'-N9	6.68	113.55	108.20
84	Aa	775	A	C5-C6-N6	-6.68	118.36	123.70
84	Aa	1724	C	N3-C4-C5	-6.68	119.23	121.90
84	Aa	3361	G	O4'-C1'-N9	6.68	113.54	108.20
84	Aa	1589	G	N1-C6-O6	6.68	123.91	119.90
84	Aa	1910	G	N1-C6-O6	6.68	123.91	119.90
1	Ad	287	C	N1-C1'-C2'	6.68	122.68	114.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1221	A	O4'-C1'-N9	6.68	113.54	108.20
84	Aa	929	A	C4-C5-C6	6.68	120.34	117.00
84	Aa	945	U	O4'-C1'-N1	6.68	113.54	108.20
84	Aa	1044	A	C5-C6-N1	-6.68	114.36	117.70
84	Aa	1148	G	N1-C6-O6	6.68	123.91	119.90
84	Aa	1411	G	C5-C6-O6	-6.68	124.59	128.60
84	Aa	2024	G	O4'-C1'-N9	6.68	113.54	108.20
33	BJ	8	TYR	N-CA-CB	6.67	122.61	110.60
84	Aa	780	U	O4'-C1'-N1	6.67	113.54	108.20
84	Aa	1624	G	P-O3'-C3'	6.67	127.71	119.70
84	Aa	3168	C	O4'-C1'-N1	6.67	113.54	108.20
1	Ad	721	U	C4'-C3'-O3'	6.67	126.35	113.00
1	Ad	1069	G	O4'-C1'-C2'	6.67	113.61	107.60
1	Ad	1199	C	N1-C1'-C2'	6.67	122.67	114.00
84	Aa	243	C	O4'-C1'-N1	6.67	113.54	108.20
84	Aa	1223	U	O4'-C1'-N1	6.67	113.54	108.20
86	Ab	2	G	O4'-C1'-N9	6.67	113.54	108.20
84	Aa	252	A	C5-C6-N1	-6.67	114.36	117.70
84	Aa	537	U	O4'-C1'-N1	6.67	113.54	108.20
84	Aa	1386	G	O4'-C1'-N9	6.67	113.54	108.20
84	Aa	1670	G	O4'-C1'-N9	6.67	113.54	108.20
84	Aa	1747	A	O4'-C1'-N9	6.67	113.54	108.20
84	Aa	1897	A	C5-C6-N1	-6.67	114.36	117.70
84	Aa	1920	U	O4'-C1'-N1	6.67	113.54	108.20
84	Aa	2896	C	O4'-C1'-N1	6.67	113.54	108.20
84	Aa	1268	G	C5-C6-O6	-6.67	124.60	128.60
84	Aa	2904	A	P-O3'-C3'	6.67	127.70	119.70
85	Ac	107	G	O4'-C1'-N9	6.67	113.54	108.20
84	Aa	942	U	O4'-C1'-N1	6.67	113.53	108.20
84	Aa	2245	G	O4'-C1'-N9	6.67	113.53	108.20
84	Aa	3242	G	C5-C6-O6	-6.67	124.60	128.60
1	Ad	1198	A	O4'-C1'-N9	6.67	113.53	108.20
84	Aa	279	G	O4'-C1'-N9	6.67	113.53	108.20
84	Aa	714	G	C5-C6-O6	-6.67	124.60	128.60
84	Aa	803	G	C5-C6-O6	-6.67	124.60	128.60
84	Aa	2406	C	N3-C4-C5	-6.67	119.23	121.90
84	Aa	3252	G	O4'-C1'-N9	6.67	113.53	108.20
84	Aa	144	A	C4-C5-C6	6.67	120.33	117.00
84	Aa	2543	G	C5-C6-O6	-6.67	124.60	128.60
84	Aa	2666	G	O4'-C1'-N9	6.67	113.53	108.20
1	Ad	1450	A	O4'-C1'-C2'	-6.66	99.14	105.80
84	Aa	1387	G	O4'-C1'-N9	6.66	113.53	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1710	G	C5-C6-O6	-6.66	124.60	128.60
84	Aa	2617	G	C5-C6-O6	-6.66	124.60	128.60
1	Ad	794	G	O4'-C1'-N9	6.66	113.53	108.20
84	Aa	509	G	C5-C6-O6	-6.66	124.60	128.60
79	CE	74	THR	N-CA-CB	6.66	122.96	110.30
84	Aa	1043	U	O4'-C1'-N1	6.66	113.53	108.20
84	Aa	3029	G	C5-C6-O6	-6.66	124.60	128.60
85	Ac	16	G	O4'-C1'-N9	6.66	113.53	108.20
84	Aa	1207	A	C5-C6-N6	-6.66	118.37	123.70
84	Aa	1615	G	O4'-C1'-N9	6.66	113.53	108.20
84	Aa	1621	G	C5-C6-O6	-6.66	124.60	128.60
84	Aa	1826	G	C5-C6-O6	-6.66	124.60	128.60
1	Ad	1406	U	N1-C1'-C2'	6.66	122.65	114.00
84	Aa	395	A	C5-C6-N6	-6.66	118.38	123.70
84	Aa	2931	C	O4'-C1'-N1	6.66	113.53	108.20
84	Aa	3010	G	C5-C6-O6	-6.66	124.61	128.60
1	Ad	744	G	C1'-O4'-C4'	6.66	115.22	109.90
84	Aa	56	A	C5-C6-N6	-6.66	118.38	123.70
84	Aa	2556	G	C5-C6-O6	-6.66	124.61	128.60
1	Ad	260	A	C1'-O4'-C4'	-6.65	104.58	109.90
72	CC	305	SER	N-CA-CB	6.65	120.48	110.50
84	Aa	256	G	C5-C6-O6	-6.65	124.61	128.60
84	Aa	426	A	C5-C6-N1	-6.65	114.37	117.70
84	Aa	776	G	N1-C6-O6	6.65	123.89	119.90
84	Aa	1071	G	O4'-C1'-N9	6.65	113.52	108.20
84	Aa	1330	A	O4'-C1'-N9	6.65	113.52	108.20
84	Aa	3299	A	C4-C5-C6	6.65	120.33	117.00
86	Ab	86	G	C5-C6-O6	-6.65	124.61	128.60
1	Ad	604	U	O4'-C1'-N1	6.65	113.52	108.20
1	Ad	773	U	C3'-C2'-C1'	-6.65	96.18	101.50
84	Aa	6	A	O4'-C1'-N9	6.65	113.52	108.20
84	Aa	730	A	O4'-C1'-N9	6.65	113.52	108.20
84	Aa	1476	G	C5-C6-O6	-6.65	124.61	128.60
84	Aa	2529	C	N3-C4-C5	-6.65	119.24	121.90
84	Aa	2706	A	C4-C5-C6	6.65	120.33	117.00
86	Ab	17	G	N1-C2-N3	-6.65	119.91	123.90
84	Aa	2630	A	C4-C5-C6	6.65	120.33	117.00
2	Ae	10	G	O4'-C1'-C2'	-6.65	99.15	105.80
84	Aa	501	U	O4'-C1'-N1	6.65	113.52	108.20
84	Aa	1435	C	O4'-C1'-N1	6.65	113.52	108.20
84	Aa	2943	A	C5-C6-N6	-6.65	118.38	123.70
84	Aa	3040	G	C5-C6-O6	-6.65	124.61	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3109	G	O4'-C1'-N9	6.65	113.52	108.20
84	Aa	1697	G	C5-C6-O6	-6.65	124.61	128.60
84	Aa	2833	G	O4'-C1'-N9	6.65	113.52	108.20
84	Aa	2979	G	C5-C6-O6	-6.65	124.61	128.60
1	Ad	1411	C	N1-C1'-C2'	6.64	122.64	114.00
84	Aa	402	U	O4'-C1'-N1	6.64	113.52	108.20
84	Aa	617	C	N3-C4-N4	6.64	122.65	118.00
84	Aa	898	G	C5-C6-O6	-6.64	124.61	128.60
84	Aa	2457	G	C5-C6-O6	-6.64	124.61	128.60
84	Aa	1046	U	O4'-C1'-N1	6.64	113.51	108.20
84	Aa	3173	A	O4'-C1'-N9	6.64	113.51	108.20
84	Aa	3383	C	O4'-C1'-N1	6.64	113.51	108.20
84	Aa	364	A	C4-C5-C6	6.64	120.32	117.00
84	Aa	580	C	N3-C4-C5	-6.64	119.25	121.90
84	Aa	1355	U	C2-N1-C1'	6.64	125.67	117.70
84	Aa	1496	G	C5-C6-O6	-6.64	124.62	128.60
84	Aa	3037	G	C5-C6-O6	-6.64	124.62	128.60
84	Aa	3076	C	P-O3'-C3'	6.64	127.67	119.70
84	Aa	906	U	O4'-C1'-N1	6.64	113.51	108.20
84	Aa	3174	C	O4'-C1'-N1	6.64	113.51	108.20
1	Ad	125	A	N9-C1'-C2'	6.64	122.63	114.00
84	Aa	438	G	C5-C6-O6	-6.64	124.62	128.60
84	Aa	1011	U	O4'-C1'-N1	6.64	113.51	108.20
84	Aa	1833	U	O4'-C1'-N1	6.64	113.51	108.20
84	Aa	3081	G	N1-C6-O6	6.64	123.88	119.90
84	Aa	224	C	O4'-C1'-N1	6.63	113.51	108.20
84	Aa	793	C	O4'-C1'-N1	6.63	113.51	108.20
84	Aa	1154	U	O4'-C1'-N1	6.63	113.51	108.20
84	Aa	1289	G	C5-C6-O6	-6.63	124.62	128.60
84	Aa	3294	U	O4'-C1'-N1	6.63	113.51	108.20
1	Ad	838	U	C5'-C4'-O4'	6.63	117.06	109.10
84	Aa	2491	A	C5-C6-N6	-6.63	118.39	123.70
1	Ad	127	G	O4'-C1'-C2'	-6.63	99.17	105.80
1	Ad	205	U	O4'-C1'-C2'	-6.63	99.17	105.80
1	Ad	1159	G	O4'-C1'-N9	6.63	113.50	108.20
1	Ad	1179	C	N1-C1'-C2'	6.63	122.62	114.00
84	Aa	414	G	O4'-C1'-N9	6.63	113.50	108.20
84	Aa	491	G	C5-C6-O6	-6.63	124.62	128.60
84	Aa	788	G	C5-C6-O6	-6.63	124.62	128.60
84	Aa	915	G	C5-C6-O6	-6.63	124.62	128.60
84	Aa	1739	G	O4'-C1'-N9	6.63	113.50	108.20
84	Aa	1747	A	C3'-C2'-C1'	-6.63	96.19	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3092	A	C4-C5-C6	6.63	120.31	117.00
1	Ad	1443	U	O4'-C1'-N1	6.63	113.50	108.20
1	Ad	1395	C	O4'-C1'-N1	6.63	113.50	108.20
84	Aa	127	G	O4'-C1'-N9	6.63	113.50	108.20
84	Aa	620	C	O4'-C1'-N1	6.63	113.50	108.20
84	Aa	839	A	C4-C5-C6	6.63	120.31	117.00
84	Aa	1503	G	O4'-C1'-N9	6.63	113.50	108.20
84	Aa	1887	A	C5-C6-N1	-6.63	114.39	117.70
84	Aa	1897	A	O4'-C1'-N9	6.63	113.50	108.20
1	Ad	1747	A	O4'-C1'-N9	6.63	113.50	108.20
84	Aa	409	U	O4'-C1'-N1	6.63	113.50	108.20
84	Aa	925	U	O4'-C1'-N1	6.63	113.50	108.20
84	Aa	1839	C	O4'-C1'-N1	6.63	113.50	108.20
84	Aa	1955	G	C5-C6-O6	-6.63	124.62	128.60
84	Aa	2758	C	O4'-C1'-N1	6.63	113.50	108.20
84	Aa	3226	G	N1-C6-O6	6.63	123.88	119.90
84	Aa	2264	U	O4'-C1'-N1	6.62	113.50	108.20
84	Aa	2390	G	C5-C6-O6	-6.62	124.62	128.60
84	Aa	3120	U	O4'-C1'-N1	6.62	113.50	108.20
1	Ad	438	G	N9-C1'-C2'	-6.62	104.71	112.00
84	Aa	195	G	C5-C6-O6	-6.62	124.62	128.60
84	Aa	286	C	O4'-C1'-N1	6.62	113.50	108.20
84	Aa	643	G	O4'-C1'-N9	6.62	113.50	108.20
84	Aa	973	U	O4'-C1'-N1	6.62	113.50	108.20
84	Aa	1528	G	O4'-C1'-N9	6.62	113.50	108.20
84	Aa	2595	G	C5-C6-O6	-6.62	124.63	128.60
84	Aa	3141	G	C5-C6-O6	-6.62	124.63	128.60
63	CU	84	TYR	CB-CG-CD2	-6.62	117.03	121.00
84	Aa	235	G	C5-C6-O6	-6.62	124.63	128.60
84	Aa	307	C	N3-C4-C5	-6.62	119.25	121.90
84	Aa	1112	C	N3-C4-C5	-6.62	119.25	121.90
84	Aa	1751	G	O4'-C1'-N9	6.62	113.50	108.20
84	Aa	2104	G	C5-C6-O6	-6.62	124.63	128.60
84	Aa	2486	G	C5-C6-O6	-6.62	124.63	128.60
84	Aa	2776	U	O4'-C1'-N1	6.62	113.50	108.20
84	Aa	345	G	C5-C6-O6	-6.62	124.63	128.60
84	Aa	1402	G	C5-C6-O6	-6.62	124.63	128.60
1	Ad	1273	U	N1-C1'-C2'	6.62	122.61	114.00
1	Ad	187	C	N1-C1'-C2'	6.62	122.60	114.00
1	Ad	474	A	O4'-C1'-C2'	-6.62	99.18	105.80
84	Aa	463	G	O4'-C1'-N9	6.62	113.49	108.20
1	Ad	856	G	C1'-O4'-C4'	-6.62	104.61	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
19	BL	116	PHE	CB-CG-CD1	6.62	125.43	120.80
84	Aa	1243	C	C5'-C4'-O4'	6.62	117.04	109.10
84	Aa	2017	G	O4'-C1'-N9	6.62	113.49	108.20
84	Aa	3018	A	C4-C5-C6	6.62	120.31	117.00
84	Aa	3264	C	C6-N1-C1'	-6.62	112.86	120.80
1	Ad	1110	C	N1-C1'-C2'	6.61	122.60	114.00
84	Aa	10	C	N3-C4-N4	6.61	122.63	118.00
84	Aa	1526	A	C5-C6-N6	-6.61	118.41	123.70
86	Ab	113	G	N1-C2-N3	-6.61	119.93	123.90
84	Aa	2095	C	C2'-C3'-O3'	6.61	124.28	113.70
84	Aa	2369	G	O4'-C1'-N9	6.61	113.49	108.20
1	Ad	59	G	O4'-C1'-C2'	6.61	113.55	107.60
74	Cp	14	TYR	CB-CG-CD2	-6.61	117.03	121.00
84	Aa	76	A	C4-C5-C6	6.61	120.31	117.00
84	Aa	417	G	O4'-C1'-N9	6.61	113.49	108.20
84	Aa	2172	C	P-O3'-C3'	6.61	127.63	119.70
84	Aa	2981	U	O4'-C1'-N1	6.61	113.49	108.20
84	Aa	3194	G	O4'-C1'-N9	6.61	113.49	108.20
85	Ac	39	G	C5-C6-O6	-6.61	124.63	128.60
84	Aa	503	U	O4'-C1'-N1	6.61	113.49	108.20
84	Aa	716	A	C5-C6-N6	-6.61	118.41	123.70
84	Aa	785	U	O4'-C1'-N1	6.61	113.49	108.20
84	Aa	3336	A	O4'-C1'-N9	6.61	113.49	108.20
1	Ad	94	A	O4'-C1'-C2'	6.61	113.55	107.60
2	Ae	34	G	N9-C1'-C2'	6.61	122.59	114.00
84	Aa	2307	A	C4-C5-C6	6.61	120.30	117.00
85	Ac	75	G	C5-C6-O6	-6.61	124.64	128.60
1	Ad	87	A	O4'-C1'-N9	6.61	113.48	108.20
1	Ad	1593	U	C3'-C2'-C1'	-6.61	96.22	101.50
84	Aa	650	A	O4'-C1'-N9	6.61	113.48	108.20
1	Ad	110	G	C1'-O4'-C4'	-6.60	104.62	109.90
84	Aa	2372	A	O4'-C1'-N9	6.60	113.48	108.20
84	Aa	2984	A	C5-C6-N6	-6.60	118.42	123.70
1	Ad	41	A	N9-C1'-C2'	-6.60	104.74	112.00
1	Ad	1012	C	O4'-C1'-N1	6.60	113.48	108.20
84	Aa	12	G	N3-C2-N2	6.60	124.52	119.90
84	Aa	974	G	O4'-C1'-N9	6.60	113.48	108.20
84	Aa	1467	G	C5-C6-O6	-6.60	124.64	128.60
84	Aa	3269	C	C6-N1-C1'	-6.60	112.88	120.80
84	Aa	3330	U	O4'-C1'-N1	6.60	113.48	108.20
84	Aa	13	G	N1-C6-O6	6.60	123.86	119.90
84	Aa	885	A	C4-C5-C6	6.60	120.30	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2642	G	C5-C6-O6	-6.60	124.64	128.60
86	Ab	50	A	C5-C6-N1	-6.60	114.40	117.70
84	Aa	2283	G	C5-C6-O6	-6.60	124.64	128.60
84	Aa	2531	G	O4'-C1'-N9	6.60	113.48	108.20
84	Aa	2777	U	O4'-C1'-N1	6.60	113.48	108.20
84	Aa	2820	U	O4'-C1'-N1	6.60	113.48	108.20
84	Aa	3070	G	N3-C2-N2	6.60	124.52	119.90
1	Ad	208	U	C4'-C3'-C2'	-6.60	96.00	102.60
1	Ad	449	A	C3'-C2'-C1'	6.60	106.78	101.50
1	Ad	1117	G	N9-C1'-C2'	6.60	122.58	114.00
1	Ad	1189	U	O4'-C1'-N1	6.60	113.48	108.20
86	Ab	26	C	N3-C4-C5	-6.60	119.26	121.90
84	Aa	351	G	C5-C6-O6	-6.59	124.64	128.60
84	Aa	1723	C	N3-C4-N4	6.59	122.62	118.00
84	Aa	2360	A	C4-C5-C6	6.59	120.30	117.00
1	Ad	1662	G	O4'-C1'-N9	6.59	113.47	108.20
84	Aa	122	A	C4-C5-C6	6.59	120.30	117.00
84	Aa	2038	G	C5-C6-O6	-6.59	124.64	128.60
84	Aa	2417	G	C5-C6-O6	-6.59	124.64	128.60
1	Ad	1189	U	P-O3'-C3'	6.59	127.61	119.70
84	Aa	1	G	O4'-C1'-N9	6.59	113.47	108.20
84	Aa	943	G	C5-C6-O6	-6.59	124.64	128.60
84	Aa	1792	G	N1-C6-O6	6.59	123.86	119.90
84	Aa	2020	G	O4'-C1'-N9	6.59	113.47	108.20
84	Aa	3364	A	O4'-C1'-N9	6.59	113.47	108.20
85	Ac	96	A	C5-C6-N1	-6.59	114.40	117.70
84	Aa	888	U	O4'-C1'-N1	6.59	113.47	108.20
84	Aa	1783	G	C5-C6-O6	-6.59	124.65	128.60
84	Aa	1837	A	O4'-C1'-N9	6.59	113.47	108.20
86	Ab	40	A	O4'-C1'-N9	6.59	113.47	108.20
84	Aa	378	U	O4'-C1'-N1	6.59	113.47	108.20
84	Aa	2117	G	O4'-C1'-N9	6.59	113.47	108.20
84	Aa	2187	C	N3-C4-C5	-6.59	119.27	121.90
84	Aa	2210	A	C4-C5-C6	6.59	120.29	117.00
84	Aa	61	A	C4-C5-C6	6.58	120.29	117.00
84	Aa	2141	A	C4-C5-C6	6.58	120.29	117.00
84	Aa	2697	A	C5-C6-N1	-6.58	114.41	117.70
84	Aa	3012	A	C5-C6-N6	-6.58	118.43	123.70
84	Aa	337	C	N3-C4-C5	-6.58	119.27	121.90
84	Aa	520	G	O4'-C1'-N9	6.58	113.47	108.20
84	Aa	3087	A	C4-C5-C6	6.58	120.29	117.00
84	Aa	1153	A	C4-C5-C6	6.58	120.29	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1506	A	O4'-C1'-N9	6.58	113.47	108.20
84	Aa	3362	A	C4-C5-C6	6.58	120.29	117.00
84	Aa	1596	G	O4'-C1'-N9	6.58	113.46	108.20
84	Aa	1985	G	C5-C6-O6	-6.58	124.65	128.60
84	Aa	2396	A	C5-C6-N6	-6.58	118.44	123.70
1	Ad	753	C	C1'-O4'-C4'	-6.58	104.64	109.90
1	Ad	1274	G	C1'-O4'-C4'	-6.58	104.64	109.90
57	Ce	26	TYR	CB-CG-CD1	6.58	124.95	121.00
84	Aa	1175	G	C5-C6-O6	-6.58	124.65	128.60
84	Aa	1540	G	O4'-C1'-N9	6.58	113.46	108.20
84	Aa	2601	G	C5-C6-O6	-6.58	124.65	128.60
84	Aa	2944	C	O4'-C1'-N1	6.58	113.46	108.20
84	Aa	3237	G	O4'-C1'-N9	6.58	113.46	108.20
84	Aa	396	G	C5-C6-O6	-6.58	124.65	128.60
84	Aa	430	G	O4'-C1'-N9	6.58	113.46	108.20
1	Ad	110	G	C3'-C2'-C1'	-6.58	96.24	101.50
1	Ad	250	A	C1'-O4'-C4'	6.58	115.16	109.90
1	Ad	648	C	C3'-C2'-C1'	6.58	106.76	101.50
84	Aa	94	A	C4-C5-C6	6.58	120.29	117.00
84	Aa	1166	C	O4'-C1'-N1	6.58	113.46	108.20
84	Aa	2229	G	C5-C6-O6	-6.58	124.65	128.60
84	Aa	2769	U	O4'-C1'-N1	6.58	113.46	108.20
1	Ad	256	G	O4'-C1'-C2'	-6.57	99.23	105.80
1	Ad	632	G	N9-C1'-C2'	6.57	122.55	114.00
1	Ad	750	U	O4'-C1'-N1	6.57	113.46	108.20
1	Ad	905	A	O4'-C1'-N9	6.57	113.46	108.20
84	Aa	424	G	C4-N9-C1'	6.57	135.05	126.50
84	Aa	1167	G	C5-C6-O6	-6.57	124.66	128.60
84	Aa	1224	A	C5-C6-N6	-6.57	118.44	123.70
84	Aa	1691	U	O4'-C1'-N1	6.57	113.46	108.20
84	Aa	1766	U	O4'-C1'-N1	6.57	113.46	108.20
84	Aa	1870	G	O4'-C1'-N9	6.57	113.46	108.20
84	Aa	2618	G	O4'-C1'-N9	6.57	113.46	108.20
84	Aa	2914	G	C5-C6-O6	-6.57	124.66	128.60
86	Ab	59	U	C5-C6-N1	6.57	125.99	122.70
84	Aa	685	G	O4'-C1'-N9	6.57	113.46	108.20
84	Aa	1163	A	C5-C6-N1	-6.57	114.41	117.70
84	Aa	1777	C	O4'-C1'-N1	6.57	113.46	108.20
85	Ac	88	A	C4-C5-C6	6.57	120.29	117.00
1	Ad	1702	G	C1'-O4'-C4'	-6.57	104.64	109.90
73	CO	117	TYR	CB-CG-CD1	6.57	124.94	121.00
84	Aa	21	G	O4'-C1'-N9	6.57	113.46	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	685	G	C5-C6-O6	-6.57	124.66	128.60
84	Aa	1266	G	N1-C6-O6	6.57	123.84	119.90
84	Aa	2606	G	O4'-C1'-N9	6.57	113.46	108.20
84	Aa	2700	A	C5-C6-N1	-6.57	114.42	117.70
84	Aa	2778	C	O4'-C1'-N1	6.57	113.46	108.20
35	BG	160	ASN	N-CA-CB	6.57	122.42	110.60
84	Aa	613	G	C5-C6-O6	-6.57	124.66	128.60
84	Aa	1371	G	C5-C6-O6	-6.57	124.66	128.60
84	Aa	2132	A	C5-C6-N1	-6.57	114.42	117.70
84	Aa	2989	A	C4-C5-C6	6.57	120.28	117.00
1	Ad	1615	G	C1'-O4'-C4'	-6.57	104.64	109.90
84	Aa	1091	C	N3-C4-N4	6.57	122.60	118.00
84	Aa	1267	A	O4'-C1'-N9	6.57	113.45	108.20
84	Aa	1454	C	O4'-C1'-N1	6.57	113.45	108.20
84	Aa	1916	U	O4'-C1'-N1	6.57	113.45	108.20
84	Aa	2146	A	O4'-C1'-N9	6.57	113.45	108.20
84	Aa	1600	A	C4-C5-C6	6.57	120.28	117.00
84	Aa	1982	G	C5-C6-O6	-6.57	124.66	128.60
84	Aa	2621	G	C5-C6-O6	-6.57	124.66	128.60
84	Aa	2894	U	O4'-C1'-N1	6.57	113.45	108.20
84	Aa	3253	C	N3-C4-C5	-6.57	119.27	121.90
84	Aa	729	G	O4'-C1'-N9	6.56	113.45	108.20
84	Aa	1595	G	C5-C6-O6	-6.56	124.66	128.60
84	Aa	3147	G	C5-C6-O6	-6.56	124.66	128.60
84	Aa	697	A	O4'-C1'-N9	6.56	113.45	108.20
84	Aa	2381	G	O4'-C1'-N9	6.56	113.45	108.20
84	Aa	532	G	O4'-C1'-N9	6.56	113.45	108.20
84	Aa	607	U	C5'-C4'-O4'	6.56	116.97	109.10
84	Aa	1420	G	C5-C6-O6	-6.56	124.66	128.60
84	Aa	1540	G	C5-C6-O6	-6.56	124.66	128.60
1	Ad	467	U	O4'-C1'-N1	6.56	113.45	108.20
1	Ad	720	U	P-O3'-C3'	-6.56	111.83	119.70
84	Aa	2066	G	C5-C6-O6	-6.56	124.66	128.60
84	Aa	2074	C	N3-C4-N4	6.56	122.59	118.00
84	Aa	1643	A	O4'-C1'-N9	6.56	113.44	108.20
84	Aa	1839	C	N3-C4-C5	-6.56	119.28	121.90
84	Aa	1979	G	O4'-C1'-N9	6.56	113.45	108.20
84	Aa	2877	U	O4'-C1'-N1	6.56	113.45	108.20
86	Ab	5	G	C5-C6-O6	-6.56	124.67	128.60
84	Aa	2665	A	O4'-C1'-N9	6.56	113.44	108.20
85	Ac	8	C	N3-C4-N4	6.56	122.59	118.00
84	Aa	707	G	C5-C6-O6	-6.55	124.67	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1531	G	O4'-C1'-N9	6.55	113.44	108.20
84	Aa	1725	G	C5-C6-O6	-6.55	124.67	128.60
85	Ac	108	C	N3-C4-C5	-6.55	119.28	121.90
86	Ab	29	C	O4'-C1'-N1	6.55	113.44	108.20
1	Ad	1304	A	C1'-O4'-C4'	6.55	115.14	109.90
1	Ad	1720	G	C1'-O4'-C4'	-6.55	104.66	109.90
84	Aa	1290	A	O4'-C1'-N9	6.55	113.44	108.20
84	Aa	1083	C	O4'-C1'-N1	6.55	113.44	108.20
84	Aa	1321	A	C5-C6-N1	-6.55	114.42	117.70
84	Aa	1786	G	O4'-C1'-N9	6.55	113.44	108.20
84	Aa	1845	C	N3-C4-N4	6.55	122.59	118.00
84	Aa	2664	G	O4'-C1'-N9	6.55	113.44	108.20
84	Aa	3154	G	O4'-C1'-N9	6.55	113.44	108.20
84	Aa	599	C	O4'-C1'-N1	6.55	113.44	108.20
84	Aa	1143	G	C5-C6-O6	-6.55	124.67	128.60
84	Aa	1679	U	O4'-C1'-N1	6.55	113.44	108.20
84	Aa	2165	A	C5-C6-N1	-6.55	114.42	117.70
84	Aa	2498	C	N3-C4-C5	-6.55	119.28	121.90
84	Aa	2939	G	O4'-C1'-N9	6.55	113.44	108.20
84	Aa	2723	G	C5-C6-O6	-6.55	124.67	128.60
84	Aa	3183	G	O4'-C4'-C3'	-6.55	97.45	104.00
1	Ad	1211	U	O4'-C1'-N1	6.55	113.44	108.20
84	Aa	234	G	O4'-C1'-N9	6.55	113.44	108.20
84	Aa	1318	C	N3-C4-N4	6.55	122.58	118.00
84	Aa	1579	C	N3-C4-C5	-6.55	119.28	121.90
84	Aa	1874	A	C4-C5-C6	6.55	120.27	117.00
84	Aa	2290	A	C4-C5-C6	6.55	120.27	117.00
84	Aa	3384	G	O4'-C1'-N9	6.55	113.44	108.20
84	Aa	904	G	C5-C6-O6	-6.54	124.67	128.60
84	Aa	713	G	C5-C6-O6	-6.54	124.67	128.60
84	Aa	1963	G	O4'-C1'-N9	6.54	113.44	108.20
84	Aa	2260	C	N3-C4-C5	-6.54	119.28	121.90
84	Aa	2311	A	C5-C6-N1	-6.54	114.43	117.70
84	Aa	3052	U	O4'-C1'-N1	6.54	113.44	108.20
84	Aa	909	A	C5-C6-N6	-6.54	118.47	123.70
84	Aa	1479	G	C5-C6-O6	-6.54	124.67	128.60
85	Ac	130	G	O4'-C1'-N9	6.54	113.43	108.20
86	Ab	100	A	N9-C4-C5	6.54	108.42	105.80
1	Ad	411	A	O4'-C1'-N9	6.54	113.43	108.20
1	Ad	749	G	O4'-C1'-C2'	6.54	113.49	107.60
2	Ae	36	C	C1'-O4'-C4'	-6.54	104.67	109.90
84	Aa	1452	A	C4-C5-C6	6.54	120.27	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2424	G	N3-C2-N2	6.54	124.48	119.90
84	Aa	3238	U	O4'-C1'-N1	6.54	113.43	108.20
1	Ad	913	U	N1-C1'-C2'	6.54	122.50	114.00
84	Aa	258	C	O4'-C1'-N1	6.54	113.43	108.20
84	Aa	1798	C	C5'-C4'-C3'	6.54	126.46	116.00
84	Aa	2982	U	O4'-C1'-N1	6.54	113.43	108.20
84	Aa	3216	G	O4'-C1'-N9	6.54	113.43	108.20
1	Ad	640	A	O4'-C1'-N9	6.54	113.43	108.20
1	Ad	642	C	N1-C1'-C2'	6.54	122.50	114.00
1	Ad	1594	A	O4'-C1'-N9	6.54	113.43	108.20
84	Aa	228	C	O4'-C1'-N1	6.54	113.43	108.20
84	Aa	1838	A	C5-C6-N6	-6.54	118.47	123.70
86	Ab	56	G	O4'-C1'-N9	6.54	113.43	108.20
86	Ab	92	C	C6-N1-C2	-6.54	117.69	120.30
84	Aa	2423	A	O4'-C1'-N9	6.53	113.43	108.20
84	Aa	2775	C	N3-C4-C5	-6.53	119.29	121.90
84	Aa	852	C	N3-C4-C5	-6.53	119.29	121.90
84	Aa	2933	C	N3-C4-N4	6.53	122.57	118.00
1	Ad	288	G	O4'-C1'-C2'	6.53	113.48	107.60
84	Aa	555	G	N1-C6-O6	6.53	123.82	119.90
84	Aa	1919	C	N3-C4-C5	-6.53	119.29	121.90
84	Aa	3005	C	N3-C4-C5	-6.53	119.29	121.90
84	Aa	3190	U	C2-N1-C1'	6.53	125.54	117.70
84	Aa	3182	A	P-O5'-C5'	-6.53	110.45	120.90
71	CB	123	CYS	N-CA-CB	6.53	122.35	110.60
84	Aa	255	C	O4'-C1'-N1	6.53	113.42	108.20
84	Aa	259	G	C5-C6-O6	-6.53	124.68	128.60
84	Aa	615	A	C5-C6-N1	-6.53	114.44	117.70
84	Aa	2677	A	C5-C6-N6	-6.53	118.48	123.70
84	Aa	2996	A	C4-C5-C6	6.53	120.26	117.00
1	Ad	1003	A	C3'-C2'-C1'	6.53	106.72	101.50
84	Aa	40	G	C5-C6-O6	-6.53	124.68	128.60
84	Aa	1488	G	N3-C2-N2	6.53	124.47	119.90
84	Aa	2409	U	O4'-C1'-N1	6.53	113.42	108.20
84	Aa	1993	G	O4'-C1'-N9	6.52	113.42	108.20
84	Aa	2746	G	C5-C6-O6	-6.52	124.69	128.60
1	Ad	448	C	N1-C1'-C2'	6.52	122.48	114.00
18	BN	81	ALA	N-CA-CB	6.52	119.23	110.10
84	Aa	1067	G	C5-C6-O6	-6.52	124.69	128.60
84	Aa	1261	C	C5-C4-N4	-6.52	115.63	120.20
84	Aa	2165	A	C5'-C4'-C3'	-6.52	105.56	116.00
84	Aa	2736	A	C5-C6-N1	-6.52	114.44	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1243	C	N3-C4-C5	-6.52	119.29	121.90
84	Aa	1941	G	O4'-C1'-N9	6.52	113.42	108.20
84	Aa	385	A	C5-C6-N1	-6.52	114.44	117.70
1	Ad	1809	U	P-O3'-C3'	6.52	127.52	119.70
84	Aa	266	A	C5-C6-N6	-6.52	118.49	123.70
84	Aa	970	A	O4'-C1'-N9	6.52	113.41	108.20
1	Ad	201	G	C1'-O4'-C4'	-6.52	104.69	109.90
1	Ad	546	U	P-O3'-C3'	6.52	127.52	119.70
84	Aa	59	A	O4'-C1'-N9	6.52	113.41	108.20
84	Aa	421	A	O4'-C1'-N9	6.52	113.41	108.20
84	Aa	1726	G	C5-C6-O6	-6.52	124.69	128.60
86	Ab	96	U	O4'-C1'-N1	6.52	113.41	108.20
1	Ad	1382	C	C1'-O4'-C4'	-6.51	104.69	109.90
84	Aa	1340	G	O4'-C1'-N9	6.51	113.41	108.20
84	Aa	1532	A	C5-C6-N1	-6.51	114.44	117.70
84	Aa	1936	G	C5-C6-O6	-6.51	124.69	128.60
84	Aa	1950	G	N1-C6-O6	6.51	123.81	119.90
86	Ab	18	C	O4'-C1'-N1	6.51	113.41	108.20
1	Ad	240	U	N1-C1'-C2'	6.51	122.47	114.00
84	Aa	1441	U	O4'-C1'-N1	6.51	113.41	108.20
84	Aa	67	C	O4'-C1'-N1	6.51	113.41	108.20
84	Aa	125	G	O4'-C1'-N9	6.51	113.41	108.20
84	Aa	263	A	C4-C5-C6	6.51	120.26	117.00
84	Aa	1308	A	C4-C5-C6	6.51	120.26	117.00
84	Aa	2086	A	P-O3'-C3'	6.51	127.51	119.70
84	Aa	2721	C	N3-C4-C5	-6.51	119.30	121.90
84	Aa	259	G	O4'-C1'-N9	6.51	113.41	108.20
84	Aa	1836	U	O4'-C1'-N1	6.51	113.41	108.20
84	Aa	1922	C	O4'-C1'-N1	6.51	113.41	108.20
84	Aa	2263	U	O4'-C1'-N1	6.51	113.41	108.20
84	Aa	2539	G	O4'-C1'-N9	6.51	113.41	108.20
84	Aa	3380	G	O4'-C1'-N9	6.51	113.41	108.20
1	Ad	55	A	N9-C1'-C2'	6.51	122.46	114.00
84	Aa	1855	A	C4-C5-C6	6.51	120.25	117.00
1	Ad	1679	A	C3'-C2'-C1'	6.51	106.70	101.50
84	Aa	972	C	N3-C4-C5	-6.51	119.30	121.90
84	Aa	1632	G	C5-C6-O6	-6.51	124.70	128.60
84	Aa	1849	U	O4'-C1'-N1	6.51	113.41	108.20
84	Aa	2029	G	C5-C6-O6	-6.51	124.70	128.60
84	Aa	2669	C	N3-C4-C5	-6.51	119.30	121.90
84	Aa	3143	A	O4'-C1'-N9	6.51	113.41	108.20
1	Ad	1624	G	C5'-C4'-O4'	6.50	116.91	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	BZ	108	ALA	N-CA-CB	6.50	119.21	110.10
84	Aa	941	C	O4'-C1'-N1	6.50	113.40	108.20
84	Aa	2173	G	C5-C6-O6	-6.50	124.70	128.60
84	Aa	1964	G	O4'-C1'-N9	6.50	113.40	108.20
84	Aa	2344	A	C5-C6-N1	-6.50	114.45	117.70
84	Aa	2756	G	N1-C6-O6	6.50	123.80	119.90
1	Ad	141	G	C3'-C2'-C1'	6.50	106.70	101.50
1	Ad	1058	G	C1'-O4'-C4'	-6.50	104.70	109.90
84	Aa	670	A	O4'-C1'-N9	6.50	113.40	108.20
84	Aa	988	G	C5-C6-O6	-6.50	124.70	128.60
84	Aa	2040	G	O4'-C1'-N9	6.50	113.40	108.20
84	Aa	2080	G	C5-C6-O6	-6.50	124.70	128.60
84	Aa	2855	G	C5-C6-O6	-6.50	124.70	128.60
84	Aa	155	G	C5-C6-O6	-6.50	124.70	128.60
84	Aa	2353	C	N3-C4-C5	-6.50	119.30	121.90
1	Ad	998	A	C1'-O4'-C4'	-6.50	104.70	109.90
2	Ae	41	G	O4'-C1'-N9	6.50	113.40	108.20
84	Aa	758	A	C5-C6-N1	-6.50	114.45	117.70
84	Aa	1506	A	C5-C6-N6	-6.50	118.50	123.70
2	Ae	5	U	C3'-C2'-C1'	6.50	106.70	101.50
84	Aa	1	G	C5-C6-O6	-6.50	124.70	128.60
84	Aa	290	C	N3-C4-C5	-6.50	119.30	121.90
84	Aa	1063	G	O4'-C1'-N9	6.50	113.40	108.20
84	Aa	1365	C	N3-C4-N4	6.50	122.55	118.00
84	Aa	2790	C	O4'-C1'-N1	6.50	113.40	108.20
1	Ad	1041	A	C1'-O4'-C4'	6.50	115.10	109.90
84	Aa	358	G	C5-C6-O6	-6.50	124.70	128.60
84	Aa	2315	G	C5-C6-O6	-6.50	124.70	128.60
1	Ad	850	G	O4'-C1'-N9	6.49	113.39	108.20
1	Ad	919	G	C1'-O4'-C4'	6.49	115.09	109.90
84	Aa	742	G	C5-C6-O6	-6.49	124.70	128.60
1	Ad	1036	U	O4'-C1'-C2'	-6.49	99.31	105.80
1	Ad	1467	C	C1'-O4'-C4'	-6.49	104.71	109.90
84	Aa	997	G	N1-C6-O6	6.49	123.80	119.90
84	Aa	1625	G	O4'-C1'-N9	6.49	113.39	108.20
84	Aa	2523	G	C5'-C4'-O4'	6.49	116.89	109.10
84	Aa	2724	A	O4'-C1'-N9	6.49	113.39	108.20
84	Aa	99	A	C4-C5-C6	6.49	120.25	117.00
84	Aa	744	C	N3-C4-C5	-6.49	119.31	121.90
84	Aa	1485	A	C4-C5-C6	6.49	120.24	117.00
84	Aa	1997	G	C5-C6-O6	-6.49	124.71	128.60
84	Aa	2368	G	C5-C6-O6	-6.49	124.71	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2426	C	N3-C4-C5	-6.49	119.30	121.90
84	Aa	2797	U	O4'-C1'-N1	6.49	113.39	108.20
84	Aa	3226	G	O4'-C1'-N9	6.49	113.39	108.20
86	Ab	22	A	O4'-C1'-N9	6.49	113.39	108.20
86	Ab	27	A	C5-C6-N6	-6.49	118.51	123.70
84	Aa	918	A	C4-C5-C6	6.49	120.24	117.00
84	Aa	2284	U	O4'-C1'-N1	6.49	113.39	108.20
1	Ad	922	U	C1'-O4'-C4'	-6.49	104.71	109.90
1	Ad	1268	G	O4'-C1'-N9	6.49	113.39	108.20
84	Aa	51	A	C5'-C4'-O4'	6.49	116.88	109.10
84	Aa	958	U	O4'-C1'-N1	6.49	113.39	108.20
84	Aa	1235	A	O4'-C1'-N9	6.49	113.39	108.20
84	Aa	1978	G	C5-C6-O6	-6.49	124.71	128.60
84	Aa	2591	G	N1-C6-O6	6.49	123.79	119.90
84	Aa	3152	C	P-O3'-C3'	6.49	127.48	119.70
84	Aa	184	C	O4'-C1'-N1	6.48	113.39	108.20
84	Aa	944	G	C5-C6-O6	-6.48	124.71	128.60
1	Ad	11	A	O4'-C1'-C2'	-6.48	99.32	105.80
1	Ad	1679	A	O4'-C1'-C2'	-6.48	99.32	105.80
72	CC	330	VAL	N-CA-C	-6.48	93.50	111.00
84	Aa	2365	C	O4'-C1'-N1	6.48	113.39	108.20
84	Aa	2977	U	O4'-C1'-N1	6.48	113.39	108.20
84	Aa	3006	G	C5-C6-O6	-6.48	124.71	128.60
1	Ad	714	C	O4'-C1'-N1	6.48	113.39	108.20
84	Aa	791	C	N3-C4-C5	-6.48	119.31	121.90
84	Aa	1377	G	C5-C6-O6	-6.48	124.71	128.60
84	Aa	1680	A	O4'-C1'-N9	6.48	113.39	108.20
86	Ab	92	C	C2-N3-C4	6.48	123.14	119.90
84	Aa	369	G	C5-C6-O6	-6.48	124.71	128.60
84	Aa	1308	A	P-O3'-C3'	-6.48	111.92	119.70
1	Ad	255	U	C1'-O4'-C4'	6.48	115.08	109.90
50	CP	167	ALA	N-CA-CB	6.48	119.17	110.10
84	Aa	253	G	C5-C6-O6	-6.48	124.71	128.60
84	Aa	1648	C	N3-C4-C5	-6.48	119.31	121.90
84	Aa	2898	A	O4'-C1'-N9	6.48	113.38	108.20
84	Aa	3193	C	O4'-C1'-N1	6.48	113.38	108.20
1	Ad	1259	G	C3'-C2'-C1'	-6.48	96.32	101.50
84	Aa	3231	G	O4'-C1'-N9	6.48	113.38	108.20
84	Aa	217	A	C5-C6-N6	-6.47	118.52	123.70
84	Aa	1617	A	C5-C6-N1	-6.47	114.46	117.70
84	Aa	1893	G	C5-C6-O6	-6.47	124.72	128.60
1	Ad	1180	U	O4'-C1'-N1	6.47	113.38	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1282	A	P-O3'-C3'	6.47	127.47	119.70
84	Aa	3128	A	O4'-C1'-N9	6.47	113.38	108.20
86	Ab	50	A	C4-C5-C6	6.47	120.24	117.00
1	Ad	245	C	C1'-O4'-C4'	-6.47	104.72	109.90
1	Ad	108	C	N1-C1'-C2'	6.47	122.41	114.00
84	Aa	1233	G	C5-C6-O6	-6.47	124.72	128.60
84	Aa	3084	G	C5-C6-O6	-6.47	124.72	128.60
84	Aa	3313	C	O4'-C1'-N1	6.47	113.38	108.20
85	Ac	67	U	O4'-C1'-N1	6.47	113.38	108.20
86	Ab	9	U	O4'-C1'-N1	6.47	113.38	108.20
86	Ab	50	A	O4'-C1'-N9	6.47	113.38	108.20
1	Ad	194	G	O4'-C1'-C2'	-6.47	99.33	105.80
84	Aa	1450	G	C5-C6-O6	-6.47	124.72	128.60
84	Aa	2065	G	C5-C6-O6	-6.47	124.72	128.60
84	Aa	2119	A	C5-C6-N1	-6.47	114.47	117.70
1	Ad	536	U	O4'-C1'-C2'	6.47	113.42	107.60
84	Aa	5	G	C5-C6-O6	-6.47	124.72	128.60
84	Aa	209	G	C5-C6-O6	-6.47	124.72	128.60
84	Aa	986	G	C5-C6-O6	-6.47	124.72	128.60
84	Aa	1142	G	C5-C6-O6	-6.47	124.72	128.60
84	Aa	1334	A	C5-C6-N6	-6.47	118.53	123.70
84	Aa	1841	G	C5-C6-O6	-6.47	124.72	128.60
84	Aa	2173	G	N1-C6-O6	6.47	123.78	119.90
85	Ac	89	A	C5-C6-N1	-6.47	114.47	117.70
85	Ac	102	U	O4'-C1'-N1	6.47	113.37	108.20
1	Ad	521	U	O4'-C1'-N1	6.46	113.37	108.20
84	Aa	923	A	N1-C6-N6	6.46	122.48	118.60
84	Aa	1655	G	C5-C6-O6	-6.46	124.72	128.60
84	Aa	2461	A	C4-C5-C6	6.46	120.23	117.00
1	Ad	926	G	C5'-C4'-O4'	6.46	116.86	109.10
1	Ad	1502	C	N1-C1'-C2'	6.46	122.40	114.00
84	Aa	1757	G	C5-C6-O6	-6.46	124.72	128.60
1	Ad	357	A	C3'-C2'-C1'	6.46	106.67	101.50
84	Aa	349	A	C5-C6-N1	-6.46	114.47	117.70
84	Aa	682	G	O4'-C1'-N9	6.46	113.37	108.20
84	Aa	1123	A	C5-C6-N1	-6.46	114.47	117.70
84	Aa	1356	G	C5-C6-O6	-6.46	124.72	128.60
84	Aa	1634	G	C5-C6-O6	-6.46	124.72	128.60
84	Aa	1895	G	C5-C6-O6	-6.46	124.72	128.60
84	Aa	3107	A	C4-C5-C6	6.46	120.23	117.00
85	Ac	2	G	O4'-C1'-N9	6.46	113.37	108.20
86	Ab	83	A	C5-C6-N6	-6.46	118.53	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	778	G	O4'-C1'-N9	6.46	113.37	108.20
84	Aa	991	C	N3-C4-N4	6.46	122.52	118.00
84	Aa	1353	A	C4-C5-C6	6.46	120.23	117.00
84	Aa	2688	G	O4'-C1'-N9	6.46	113.37	108.20
84	Aa	3276	G	O4'-C1'-N9	6.46	113.37	108.20
1	Ad	79	A	C1'-O4'-C4'	6.46	115.07	109.90
1	Ad	749	G	C1'-O4'-C4'	-6.46	104.73	109.90
1	Ad	1727	C	N1-C1'-C2'	6.46	122.39	114.00
1	Ad	1792	A	O4'-C1'-N9	6.46	113.37	108.20
2	Ae	7	A	P-O3'-C3'	6.46	127.45	119.70
84	Aa	478	G	C5-C6-O6	-6.46	124.72	128.60
84	Aa	1096	C	N3-C4-C5	-6.46	119.32	121.90
84	Aa	1883	A	C4-C5-C6	6.46	120.23	117.00
84	Aa	2024	G	C5-C6-O6	-6.46	124.72	128.60
84	Aa	2126	C	N3-C4-C5	-6.46	119.32	121.90
84	Aa	3017	A	C4-C5-C6	6.46	120.23	117.00
84	Aa	28	C	N3-C4-C5	-6.46	119.32	121.90
84	Aa	1758	U	O4'-C1'-N1	6.46	113.36	108.20
84	Aa	2050	G	O4'-C1'-N9	6.46	113.37	108.20
84	Aa	2138	A	C5-C6-N6	-6.46	118.53	123.70
84	Aa	2290	A	C5-C6-N1	-6.46	114.47	117.70
84	Aa	3156	G	C5-C6-O6	-6.46	124.73	128.60
85	Ac	140	A	C5-C6-N1	-6.46	114.47	117.70
86	Ab	42	A	N9-C4-C5	6.46	108.38	105.80
86	Ab	101	A	C8-N9-C4	-6.46	103.22	105.80
86	Ab	12	U	O4'-C1'-N1	6.46	113.36	108.20
1	Ad	139	U	N1-C1'-C2'	6.45	122.39	114.00
1	Ad	1806	C	C1'-O4'-C4'	6.45	115.06	109.90
84	Aa	84	A	O4'-C1'-N9	6.45	113.36	108.20
84	Aa	1188	C	P-O5'-C5'	6.45	131.22	120.90
84	Aa	2577	G	O4'-C1'-N9	6.45	113.36	108.20
84	Aa	2967	U	O4'-C1'-N1	6.45	113.36	108.20
1	Ad	398	C	O4'-C1'-N1	6.45	113.36	108.20
84	Aa	674	G	O4'-C1'-N9	6.45	113.36	108.20
84	Aa	2683	A	C5-C6-N1	-6.45	114.47	117.70
84	Aa	2729	C	C6-N1-C1'	-6.45	113.06	120.80
84	Aa	2922	U	O4'-C1'-N1	6.45	113.36	108.20
85	Ac	157	A	C5-C6-N6	-6.45	118.54	123.70
1	Ad	329	G	C1'-O4'-C4'	-6.45	104.74	109.90
73	CO	136	PRO	CB-CA-C	6.45	128.13	112.00
84	Aa	754	G	C5-C6-O6	-6.45	124.73	128.60
84	Aa	957	U	O4'-C1'-N1	6.45	113.36	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2105	G	C5-C6-O6	-6.45	124.73	128.60
84	Aa	3110	A	C4-C5-C6	6.45	120.23	117.00
84	Aa	3316	C	N3-C4-N4	6.45	122.52	118.00
85	Ac	16	G	C5-C6-O6	-6.45	124.73	128.60
1	Ad	1450	A	C1'-O4'-C4'	6.45	115.06	109.90
1	Ad	1530	G	O4'-C1'-C2'	-6.45	99.35	105.80
84	Aa	363	A	C4-C5-C6	6.45	120.22	117.00
84	Aa	486	G	C5-C6-O6	-6.45	124.73	128.60
84	Aa	518	G	O4'-C1'-N9	6.45	113.36	108.20
84	Aa	937	G	C5-C6-O6	-6.45	124.73	128.60
84	Aa	941	C	N3-C4-C5	-6.45	119.32	121.90
84	Aa	1397	A	C4-C5-C6	6.45	120.22	117.00
84	Aa	1802	A	O4'-C1'-N9	6.45	113.36	108.20
84	Aa	2461	A	C5-C6-N1	-6.45	114.48	117.70
84	Aa	3130	A	O4'-C1'-N9	6.45	113.36	108.20
84	Aa	3227	U	O4'-C1'-N1	6.45	113.36	108.20
1	Ad	166	A	C3'-C2'-C1'	6.45	106.66	101.50
1	Ad	1638	U	O4'-C1'-N1	6.45	113.36	108.20
84	Aa	2550	C	N3-C4-C5	-6.45	119.32	121.90
1	Ad	834	A	N9-C1'-C2'	-6.45	104.91	112.00
84	Aa	436	G	O4'-C1'-N9	6.45	113.36	108.20
84	Aa	1313	U	O4'-C1'-N1	6.45	113.36	108.20
84	Aa	1344	A	C4-C5-C6	6.45	120.22	117.00
84	Aa	1367	A	O4'-C1'-N9	6.45	113.36	108.20
84	Aa	1472	C	O4'-C1'-N1	6.45	113.36	108.20
84	Aa	2494	A	C4-C5-C6	6.45	120.22	117.00
1	Ad	1589	C	O4'-C1'-C2'	-6.44	99.36	105.80
84	Aa	1366	G	O4'-C1'-N9	6.44	113.36	108.20
84	Aa	2324	G	O4'-C1'-N9	6.44	113.36	108.20
86	Ab	42	A	C8-N9-C4	-6.44	103.22	105.80
84	Aa	1180	C	O4'-C1'-N1	6.44	113.35	108.20
84	Aa	1296	C	O4'-C1'-N1	6.44	113.36	108.20
84	Aa	1447	G	C5-C6-O6	-6.44	124.73	128.60
84	Aa	2002	G	O4'-C1'-N9	6.44	113.35	108.20
84	Aa	3041	A	C5-C6-N1	-6.44	114.48	117.70
84	Aa	3095	G	N1-C6-O6	6.44	123.77	119.90
1	Ad	1783	C	C1'-O4'-C4'	-6.44	104.75	109.90
72	CC	91	ALA	N-CA-CB	6.44	119.12	110.10
84	Aa	111	C	O4'-C1'-N1	6.44	113.35	108.20
84	Aa	505	G	C5-C6-O6	-6.44	124.73	128.60
84	Aa	966	G	C5-C6-O6	-6.44	124.74	128.60
84	Aa	1516	G	N1-C6-O6	6.44	123.76	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1006	A	C3'-C2'-C1'	6.44	106.65	101.50
1	Ad	1648	C	O4'-C1'-C2'	-6.44	99.36	105.80
2	Ae	30	G	O4'-C1'-C2'	6.44	113.39	107.60
84	Aa	2875	U	O4'-C1'-N1	6.44	113.35	108.20
1	Ad	21	U	O4'-C1'-N1	6.44	113.35	108.20
1	Ad	1186	U	C1'-O4'-C4'	6.44	115.05	109.90
57	Ce	67	TYR	CB-CG-CD1	6.44	124.86	121.00
84	Aa	1106	G	O4'-C1'-N9	6.44	113.35	108.20
84	Aa	1379	G	C5-C6-O6	-6.44	124.74	128.60
84	Aa	2235	G	C5-C6-O6	-6.44	124.74	128.60
86	Ab	88	U	O4'-C1'-N1	6.44	113.35	108.20
84	Aa	920	A	C5-C6-N6	-6.44	118.55	123.70
84	Aa	2414	C	O4'-C1'-N1	6.44	113.35	108.20
84	Aa	2518	A	C4-C5-C6	6.44	120.22	117.00
84	Aa	894	G	C5-C6-O6	-6.43	124.74	128.60
84	Aa	1181	A	C4-C5-C6	6.43	120.22	117.00
84	Aa	1913	C	N3-C4-C5	-6.43	119.33	121.90
84	Aa	2486	G	O4'-C1'-N9	6.43	113.35	108.20
84	Aa	2932	A	O4'-C1'-N9	6.43	113.35	108.20
1	Ad	1508	C	N1-C1'-C2'	6.43	122.36	114.00
84	Aa	908	U	O4'-C1'-N1	6.43	113.35	108.20
84	Aa	1707	C	N3-C4-C5	-6.43	119.33	121.90
84	Aa	2496	U	C4'-C3'-O3'	-6.43	95.89	109.40
84	Aa	3222	G	C5-C6-O6	-6.43	124.74	128.60
1	Ad	1305	U	O4'-C1'-N1	6.43	113.34	108.20
84	Aa	2960	A	C5-C6-N1	-6.43	114.48	117.70
84	Aa	3013	A	C5-C6-N1	-6.43	114.48	117.70
84	Aa	2239	A	C4-C5-C6	6.43	120.22	117.00
84	Aa	3220	A	C5-C6-N6	-6.43	118.56	123.70
1	Ad	1572	U	C1'-O4'-C4'	-6.43	104.76	109.90
84	Aa	65	A	C5-C6-N6	-6.43	118.56	123.70
84	Aa	3172	G	C5-C6-O6	-6.43	124.74	128.60
59	Cl	39	ALA	N-CA-CB	6.43	119.10	110.10
84	Aa	83	U	O4'-C1'-N1	6.43	113.34	108.20
84	Aa	177	C	N3-C4-C5	-6.43	119.33	121.90
84	Aa	959	U	O4'-C1'-N1	6.43	113.34	108.20
84	Aa	1070	G	O4'-C1'-N9	6.43	113.34	108.20
84	Aa	2723	G	O4'-C1'-N9	6.43	113.34	108.20
3	Af	12	A	O4'-C1'-C2'	6.42	113.38	107.60
84	Aa	232	C	O4'-C1'-N1	6.42	113.34	108.20
84	Aa	429	G	O4'-C1'-N9	6.42	113.34	108.20
84	Aa	598	U	O4'-C1'-N1	6.42	113.34	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1370	A	C4-C5-C6	6.42	120.21	117.00
84	Aa	1642	G	O4'-C1'-N9	6.42	113.34	108.20
84	Aa	1982	G	O4'-C1'-N9	6.42	113.34	108.20
84	Aa	2286	A	C4-C5-C6	6.42	120.21	117.00
84	Aa	2514	A	C5-C6-N1	-6.42	114.49	117.70
86	Ab	111	U	C6-N1-C2	-6.42	117.15	121.00
84	Aa	997	G	O4'-C1'-N9	6.42	113.34	108.20
84	Aa	1333	C	O4'-C1'-N1	6.42	113.34	108.20
84	Aa	1519	C	N3-C4-C5	-6.42	119.33	121.90
84	Aa	2388	C	N3-C4-N4	6.42	122.50	118.00
84	Aa	2930	C	O4'-C1'-N1	6.42	113.34	108.20
1	Ad	237	C	C1'-O4'-C4'	-6.42	104.76	109.90
84	Aa	639	A	C4-C5-C6	6.42	120.21	117.00
84	Aa	1386	G	C5-C6-O6	-6.42	124.75	128.60
84	Aa	2159	U	O4'-C1'-N1	6.42	113.34	108.20
85	Ac	17	A	O4'-C1'-N9	6.42	113.34	108.20
84	Aa	3322	A	C5-C6-N1	-6.42	114.49	117.70
85	Ac	142	G	C5-C6-O6	-6.42	124.75	128.60
1	Ad	314	C	O4'-C1'-N1	6.42	113.33	108.20
84	Aa	34	G	O4'-C1'-N9	6.42	113.33	108.20
84	Aa	294	A	C5-C6-N6	-6.42	118.57	123.70
84	Aa	746	C	C1'-O4'-C4'	-6.42	104.77	109.90
84	Aa	1443	G	C5-C6-O6	-6.42	124.75	128.60
84	Aa	2334	G	C5-C6-O6	-6.42	124.75	128.60
84	Aa	3332	G	O4'-C1'-N9	6.42	113.33	108.20
1	Ad	363	G	C3'-C2'-C1'	-6.42	96.37	101.50
84	Aa	593	G	C5-C6-O6	-6.42	124.75	128.60
84	Aa	2256	G	C5-C6-O6	-6.42	124.75	128.60
84	Aa	2679	A	C4-C5-C6	6.42	120.21	117.00
84	Aa	2817	G	O4'-C1'-N9	6.42	113.33	108.20
84	Aa	3044	C	N3-C4-C5	-6.42	119.33	121.90
1	Ad	474	A	C3'-C2'-C1'	6.42	106.63	101.50
1	Ad	196	G	C1'-O4'-C4'	-6.41	104.77	109.90
84	Aa	829	G	C5-C6-O6	-6.41	124.75	128.60
84	Aa	1661	G	C5-C6-O6	-6.41	124.75	128.60
84	Aa	3230	G	C5-C6-O6	-6.41	124.75	128.60
71	CB	265	TYR	CB-CG-CD1	6.41	124.85	121.00
84	Aa	2463	U	C5'-C4'-O4'	6.41	116.80	109.10
86	Ab	43	A	O4'-C1'-N9	6.41	113.33	108.20
86	Ab	68	G	O4'-C1'-N9	6.41	113.33	108.20
84	Aa	270	G	C5-C6-O6	-6.41	124.75	128.60
84	Aa	969	U	O4'-C1'-N1	6.41	113.33	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2057	G	C5-C6-O6	-6.41	124.75	128.60
84	Aa	2698	A	C5-C6-N1	-6.41	114.49	117.70
86	Ab	27	A	C4-C5-C6	6.41	120.20	117.00
84	Aa	1184	U	C5-C4-O4	-6.41	122.06	125.90
84	Aa	1954	G	O4'-C1'-N9	6.41	113.33	108.20
63	CU	92	TYR	CB-CG-CD1	6.41	124.84	121.00
84	Aa	1816	U	O4'-C1'-N1	6.41	113.33	108.20
84	Aa	1886	U	O4'-C1'-N1	6.41	113.33	108.20
85	Ac	126	A	C4-C5-C6	6.41	120.20	117.00
1	Ad	100	C	O4'-C1'-N1	6.41	113.33	108.20
1	Ad	151	A	C1'-O4'-C4'	6.41	115.03	109.90
1	Ad	1020	U	C3'-C2'-C1'	6.41	106.62	101.50
20	BT	51	TYR	CB-CG-CD2	6.41	124.84	121.00
84	Aa	1416	G	C5-C6-O6	-6.41	124.76	128.60
84	Aa	1595	G	O4'-C1'-N9	6.41	113.32	108.20
84	Aa	3057	A	C5-C6-N1	-6.41	114.50	117.70
86	Ab	68	G	C6-C5-N7	-6.41	126.56	130.40
45	CN	81	TYR	CB-CG-CD1	-6.40	117.16	121.00
1	Ad	529	A	P-O3'-C3'	6.40	127.38	119.70
1	Ad	60	C	P-O3'-C3'	-6.40	112.02	119.70
84	Aa	242	U	O4'-C1'-N1	6.40	113.32	108.20
84	Aa	288	G	P-O3'-C3'	6.40	127.38	119.70
84	Aa	733	C	N3-C4-C5	-6.40	119.34	121.90
84	Aa	1809	A	C4-C5-C6	6.40	120.20	117.00
84	Aa	2722	U	O4'-C1'-N1	6.40	113.32	108.20
84	Aa	657	A	C5-C6-N6	-6.40	118.58	123.70
84	Aa	1918	A	O4'-C1'-N9	6.40	113.32	108.20
84	Aa	2743	A	C4-C5-C6	6.40	120.20	117.00
1	Ad	1541	C	C3'-C2'-C1'	6.40	106.62	101.50
79	CE	124	TYR	CB-CG-CD2	-6.40	117.16	121.00
84	Aa	2076	C	N3-C4-C5	-6.40	119.34	121.90
84	Aa	2228	A	C4-C5-C6	6.40	120.20	117.00
84	Aa	2232	C	O4'-C1'-N1	6.40	113.32	108.20
84	Aa	2889	A	C4-C5-C6	6.40	120.20	117.00
84	Aa	3059	C	C6-N1-C1'	-6.40	113.12	120.80
1	Ad	593	C	O4'-C1'-C2'	-6.40	99.40	105.80
1	Ad	1409	G	C1'-O4'-C4'	-6.40	104.78	109.90
84	Aa	405	A	C4-C5-C6	6.40	120.20	117.00
84	Aa	1451	U	O4'-C1'-N1	6.40	113.32	108.20
85	Ac	103	G	C5-C6-O6	-6.40	124.76	128.60
84	Aa	250	C	P-O3'-C3'	6.39	127.37	119.70
84	Aa	729	G	C5-C6-O6	-6.39	124.76	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	947	C	O4'-C1'-N1	6.39	113.32	108.20
84	Aa	1447	G	O4'-C1'-N9	6.39	113.31	108.20
84	Aa	1865	C	N3-C4-C5	-6.39	119.34	121.90
84	Aa	1979	G	C5-C6-O6	-6.39	124.76	128.60
84	Aa	2134	U	O4'-C1'-N1	6.39	113.32	108.20
85	Ac	123	G	O4'-C1'-N9	6.39	113.32	108.20
84	Aa	2239	A	O4'-C1'-N9	6.39	113.31	108.20
84	Aa	1219	C	N3-C4-N4	6.39	122.47	118.00
84	Aa	1299	G	C5-C6-O6	-6.39	124.77	128.60
1	Ad	287	C	C3'-C2'-C1'	6.39	106.61	101.50
1	Ad	1073	C	C3'-C2'-C1'	6.39	106.61	101.50
84	Aa	239	C	N3-C4-N4	6.39	122.47	118.00
84	Aa	300	C	N3-C4-C5	-6.39	119.34	121.90
84	Aa	571	G	C5-C6-O6	-6.39	124.77	128.60
84	Aa	578	C	N3-C4-C5	-6.39	119.34	121.90
84	Aa	844	A	C5-C6-N1	-6.39	114.50	117.70
84	Aa	3284	C	N3-C4-N4	6.39	122.47	118.00
84	Aa	128	C	N3-C4-C5	-6.39	119.34	121.90
84	Aa	1032	C	O4'-C1'-N1	6.39	113.31	108.20
1	Ad	356	G	P-O3'-C3'	6.39	127.36	119.70
84	Aa	2319	A	C4-C5-C6	6.39	120.19	117.00
1	Ad	68	A	C1'-O4'-C4'	-6.38	104.79	109.90
1	Ad	651	G	N9-C1'-C2'	6.38	122.30	114.00
84	Aa	196	A	C4-C5-C6	6.38	120.19	117.00
84	Aa	978	C	N3-C4-C5	-6.38	119.35	121.90
84	Aa	1138	A	C5-C6-N1	-6.38	114.51	117.70
84	Aa	1757	G	O4'-C1'-N9	6.38	113.31	108.20
84	Aa	2050	G	C5-C6-O6	-6.38	124.77	128.60
84	Aa	2258	C	N3-C4-C5	-6.38	119.35	121.90
84	Aa	3303	C	N3-C4-C5	-6.38	119.35	121.90
1	Ad	483	C	N1-C1'-C2'	6.38	122.30	114.00
2	Ae	50	G	C1'-O4'-C4'	6.38	115.01	109.90
84	Aa	2909	A	C5-C6-N1	-6.38	114.51	117.70
85	Ac	2	G	C5-C6-O6	-6.38	124.77	128.60
1	Ad	350	G	O4'-C1'-N9	6.38	113.31	108.20
1	Ad	382	A	O4'-C1'-N9	6.38	113.31	108.20
18	BN	151	ALA	N-CA-CB	6.38	119.03	110.10
84	Aa	1348	G	C5-C6-O6	-6.38	124.77	128.60
84	Aa	2450	G	P-O5'-C5'	6.38	131.11	120.90
84	Aa	2472	U	O4'-C1'-N1	6.38	113.31	108.20
84	Aa	129	G	C5-C6-O6	-6.38	124.77	128.60
84	Aa	530	C	N3-C4-C5	-6.38	119.35	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1230	G	C5-C6-O6	-6.38	124.77	128.60
84	Aa	1901	G	O4'-C1'-N9	6.38	113.30	108.20
1	Ad	1538	C	O4'-C1'-C2'	-6.38	99.42	105.80
84	Aa	354	C	N3-C4-C5	-6.38	119.35	121.90
84	Aa	477	C	N3-C4-C5	-6.38	119.35	121.90
84	Aa	1631	G	C5-C6-O6	-6.38	124.77	128.60
84	Aa	2405	C	N3-C4-C5	-6.38	119.35	121.90
84	Aa	3121	C	N3-C4-N4	6.38	122.47	118.00
84	Aa	3139	U	O4'-C1'-N1	6.38	113.30	108.20
1	Ad	1314	U	O4'-C1'-N1	6.38	113.30	108.20
84	Aa	422	G	C5-C6-O6	-6.38	124.77	128.60
84	Aa	442	C	N3-C4-C5	-6.38	119.35	121.90
84	Aa	1341	G	N3-C2-N2	6.38	124.36	119.90
84	Aa	310	C	N3-C4-C5	-6.38	119.35	121.90
3	Af	17	A	N9-C1'-C2'	6.37	122.29	114.00
84	Aa	424	G	C5-C6-O6	-6.37	124.78	128.60
84	Aa	1281	C	N3-C4-C5	-6.37	119.35	121.90
84	Aa	2747	U	O4'-C1'-N1	6.37	113.30	108.20
1	Ad	227	G	O3'-P-O5'	6.37	116.11	104.00
1	Ad	236	U	O4'-C1'-N1	6.37	113.30	108.20
1	Ad	784	C	N1-C1'-C2'	-6.37	104.99	112.00
79	CE	50	PHE	CB-CG-CD1	6.37	125.26	120.80
84	Aa	254	G	C5-C6-O6	-6.37	124.78	128.60
84	Aa	282	A	C5-C6-N6	-6.37	118.60	123.70
84	Aa	868	A	C4-C5-C6	6.37	120.19	117.00
84	Aa	2168	C	P-O3'-C3'	6.37	127.35	119.70
84	Aa	2435	U	O4'-C1'-N1	6.37	113.30	108.20
84	Aa	2500	U	O4'-C1'-N1	6.37	113.30	108.20
84	Aa	3167	G	C5-C6-O6	-6.37	124.78	128.60
85	Ac	18	U	O4'-C1'-N1	6.37	113.30	108.20
84	Aa	322	A	C4-C5-C6	6.37	120.19	117.00
84	Aa	2695	A	C4-C5-C6	6.37	120.19	117.00
1	Ad	82	G	O4'-C1'-N9	6.37	113.29	108.20
84	Aa	356	G	C5-C6-O6	-6.37	124.78	128.60
84	Aa	811	A	C5-C6-N6	-6.37	118.61	123.70
84	Aa	817	U	O4'-C1'-N1	6.37	113.30	108.20
84	Aa	1297	U	O4'-C1'-N1	6.37	113.29	108.20
84	Aa	2139	A	C4-C5-C6	6.37	120.18	117.00
84	Aa	2164	G	O4'-C1'-N9	6.37	113.30	108.20
84	Aa	3084	G	O4'-C1'-N9	6.37	113.29	108.20
84	Aa	3333	C	C1'-O4'-C4'	-6.37	104.81	109.90
84	Aa	29	G	N3-C2-N2	6.37	124.36	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	130	G	O4'-C1'-N9	6.37	113.29	108.20
84	Aa	1855	A	C5-C6-N6	-6.37	118.61	123.70
84	Aa	584	G	C5-C6-O6	-6.37	124.78	128.60
84	Aa	1282	A	C4-C5-C6	6.37	120.18	117.00
84	Aa	1796	A	C4-C5-C6	6.37	120.18	117.00
84	Aa	2211	G	O4'-C1'-N9	6.37	113.29	108.20
84	Aa	2969	A	C5-C6-N1	-6.37	114.52	117.70
84	Aa	3140	A	O4'-C1'-N9	6.37	113.29	108.20
85	Ac	46	G	C5-C6-O6	-6.37	124.78	128.60
85	Ac	133	C	N3-C4-C5	-6.37	119.35	121.90
71	CB	55	HIS	CA-CB-CG	6.36	124.42	113.60
84	Aa	488	U	O4'-C1'-N1	6.36	113.29	108.20
84	Aa	855	U	O4'-C1'-N1	6.36	113.29	108.20
84	Aa	2626	G	O4'-C1'-N9	6.36	113.29	108.20
84	Aa	2719	U	O4'-C1'-N1	6.36	113.29	108.20
84	Aa	3062	G	C5-C6-O6	-6.36	124.78	128.60
84	Aa	3331	G	C5-C6-O6	-6.36	124.78	128.60
84	Aa	3335	G	C5-C6-O6	-6.36	124.78	128.60
84	Aa	2783	U	O4'-C1'-N1	6.36	113.29	108.20
1	Ad	1452	A	O4'-C1'-N9	6.36	113.29	108.20
84	Aa	1155	G	N1-C6-O6	6.36	123.72	119.90
84	Aa	1330	A	C4-C5-C6	6.36	120.18	117.00
84	Aa	1338	C	N3-C4-C5	-6.36	119.36	121.90
84	Aa	2976	U	O4'-C1'-N1	6.36	113.29	108.20
84	Aa	2984	A	C4-C5-C6	6.36	120.18	117.00
84	Aa	3062	G	O4'-C1'-N9	6.36	113.29	108.20
84	Aa	3197	C	N3-C4-C5	-6.36	119.36	121.90
84	Aa	3066	G	C5-C6-O6	-6.36	124.78	128.60
84	Aa	21	G	C5-C6-O6	-6.36	124.79	128.60
84	Aa	728	G	O4'-C1'-N9	6.36	113.29	108.20
84	Aa	2685	C	N3-C4-C5	-6.36	119.36	121.90
86	Ab	91	C	C2-N3-C4	6.36	123.08	119.90
1	Ad	381	G	C1'-O4'-C4'	-6.36	104.82	109.90
84	Aa	650	A	C4-C5-C6	6.36	120.18	117.00
1	Ad	235	C	C3'-C2'-C1'	6.35	106.58	101.50
84	Aa	2127	U	O4'-C1'-N1	6.35	113.28	108.20
84	Aa	3100	C	N3-C4-N4	6.35	122.45	118.00
84	Aa	987	A	C4-C5-C6	6.35	120.18	117.00
84	Aa	1824	C	N3-C4-C5	-6.35	119.36	121.90
84	Aa	1852	C	N3-C4-C5	-6.35	119.36	121.90
84	Aa	1915	G	C5-C6-O6	-6.35	124.79	128.60
84	Aa	2487	A	C4-C5-C6	6.35	120.18	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2853	A	C4-C5-C6	6.35	120.18	117.00
1	Ad	61	A	P-O5'-C5'	-6.35	110.74	120.90
84	Aa	665	G	C5-C6-O6	-6.35	124.79	128.60
84	Aa	3178	C	N3-C4-N4	6.35	122.44	118.00
1	Ad	1477	A	O4'-C1'-C2'	-6.35	99.45	105.80
33	BJ	85	TYR	CB-CG-CD2	-6.35	117.19	121.00
84	Aa	63	G	C5-C6-O6	-6.35	124.79	128.60
84	Aa	97	G	N3-C2-N2	6.35	124.34	119.90
84	Aa	684	C	N3-C4-C5	-6.35	119.36	121.90
84	Aa	771	G	C4'-C3'-C2'	6.35	108.95	102.60
84	Aa	784	G	C5-C6-O6	-6.35	124.79	128.60
84	Aa	1454	C	N3-C4-C5	-6.35	119.36	121.90
84	Aa	2247	A	C4-C5-C6	6.35	120.17	117.00
1	Ad	1210	U	O4'-C1'-C2'	-6.35	99.45	105.80
1	Ad	1295	G	O4'-C1'-N9	6.35	113.28	108.20
84	Aa	2885	U	O5'-P-OP1	-6.35	99.99	105.70
84	Aa	3135	A	C5-C6-N1	-6.35	114.53	117.70
85	Ac	40	A	C5-C6-N6	-6.35	118.62	123.70
1	Ad	1507	G	O4'-C1'-N9	6.34	113.28	108.20
84	Aa	319	C	N3-C4-N4	6.34	122.44	118.00
84	Aa	393	A	C4-C5-C6	6.34	120.17	117.00
84	Aa	1168	G	C5-C6-O6	-6.34	124.79	128.60
84	Aa	1501	A	C5-C6-N6	-6.34	118.62	123.70
84	Aa	2954	G	O4'-C1'-N9	6.34	113.28	108.20
85	Ac	137	G	O4'-C1'-N9	6.34	113.28	108.20
84	Aa	1385	C	N3-C4-C5	-6.34	119.36	121.90
84	Aa	1490	A	C5-C6-N6	-6.34	118.63	123.70
84	Aa	1518	A	C4-C5-C6	6.34	120.17	117.00
84	Aa	2199	C	N3-C4-C5	-6.34	119.36	121.90
1	Ad	1187	A	N9-C1'-C2'	6.34	122.24	114.00
2	Ae	58	U	C1'-O4'-C4'	6.34	114.97	109.90
84	Aa	2045	G	O4'-C1'-N9	6.34	113.27	108.20
84	Aa	2389	A	C4-C5-C6	6.34	120.17	117.00
85	Ac	154	G	C5-C6-O6	-6.34	124.80	128.60
84	Aa	70	A	C5-C6-N1	-6.34	114.53	117.70
84	Aa	747	A	C4-C5-C6	6.34	120.17	117.00
84	Aa	960	C	N3-C4-C5	-6.34	119.36	121.90
84	Aa	1426	C	N3-C4-C5	-6.34	119.36	121.90
84	Aa	2645	A	C4-C5-C6	6.34	120.17	117.00
1	Ad	594	C	O4'-C1'-N1	6.34	113.27	108.20
48	CD	260	GLU	N-CA-CB	6.34	122.01	110.60
84	Aa	2681	A	C4-C5-C6	6.34	120.17	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	412	C	N3-C4-N4	6.34	122.44	118.00
84	Aa	1457	A	O4'-C1'-N9	6.34	113.27	108.20
84	Aa	1647	C	N3-C4-C5	-6.34	119.37	121.90
84	Aa	2051	G	O4'-C1'-N9	6.34	113.27	108.20
84	Aa	2128	G	C5-C6-O6	-6.34	124.80	128.60
84	Aa	426	A	C4-C5-C6	6.33	120.17	117.00
84	Aa	1363	C	N3-C4-C5	-6.33	119.37	121.90
84	Aa	2606	G	N1-C6-O6	6.33	123.70	119.90
1	Ad	1693	C	O4'-C1'-N1	6.33	113.27	108.20
84	Aa	8	C	C6-N1-C1'	-6.33	113.20	120.80
84	Aa	131	C	N3-C4-C5	-6.33	119.37	121.90
84	Aa	213	G	O4'-C1'-N9	6.33	113.27	108.20
84	Aa	561	G	O4'-C1'-N9	6.33	113.27	108.20
84	Aa	1741	G	O4'-C1'-N9	6.33	113.27	108.20
84	Aa	1823	C	N3-C4-C5	-6.33	119.37	121.90
84	Aa	2613	G	O4'-C1'-N9	6.33	113.27	108.20
84	Aa	2694	A	O4'-C1'-N9	6.33	113.27	108.20
85	Ac	57	C	N3-C4-C5	-6.33	119.37	121.90
1	Ad	1508	C	O4'-C1'-N1	6.33	113.27	108.20
84	Aa	114	G	C5-C6-O6	-6.33	124.80	128.60
84	Aa	190	C	N3-C4-C5	-6.33	119.37	121.90
84	Aa	841	G	O4'-C1'-N9	6.33	113.27	108.20
84	Aa	1696	G	C5-C6-O6	-6.33	124.80	128.60
84	Aa	1929	A	C4-C5-C6	6.33	120.17	117.00
84	Aa	2046	G	O4'-C1'-N9	6.33	113.27	108.20
84	Aa	2132	A	C4-C5-C6	6.33	120.17	117.00
84	Aa	2191	C	N3-C4-C5	-6.33	119.37	121.90
85	Ac	106	C	O4'-C1'-N1	6.33	113.26	108.20
84	Aa	292	A	C4-C5-C6	6.33	120.17	117.00
84	Aa	1312	A	C5-C6-N1	-6.33	114.53	117.70
84	Aa	3053	G	O4'-C1'-N9	6.33	113.26	108.20
84	Aa	316	A	O4'-C1'-N9	6.33	113.26	108.20
84	Aa	1129	G	C5-C6-O6	-6.33	124.80	128.60
84	Aa	1653	A	C5-C6-N6	-6.33	118.64	123.70
84	Aa	1713	A	C4-C5-C6	6.33	120.16	117.00
84	Aa	1204	A	C4-C5-C6	6.33	120.16	117.00
84	Aa	1367	A	C5-C6-N6	-6.33	118.64	123.70
84	Aa	1750	A	C4-C5-C6	6.33	120.16	117.00
84	Aa	2997	C	P-O3'-C3'	6.33	127.29	119.70
1	Ad	602	U	O4'-C1'-N1	6.33	113.26	108.20
84	Aa	1049	C	N3-C4-C5	-6.33	119.37	121.90
84	Aa	1295	A	C5-C6-N1	-6.33	114.54	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2096	U	P-O5'-C5'	-6.33	110.78	120.90
84	Aa	2534	G	O4'-C1'-N9	6.33	113.26	108.20
84	Aa	2849	A	C4-C5-C6	6.33	120.16	117.00
84	Aa	1227	A	C4-C5-C6	6.32	120.16	117.00
84	Aa	2189	G	C5-C6-O6	-6.32	124.81	128.60
84	Aa	2412	A	O4'-C1'-N9	6.32	113.26	108.20
84	Aa	2942	A	C4-C5-C6	6.32	120.16	117.00
85	Ac	110	A	C4-C5-C6	6.32	120.16	117.00
1	Ad	531	A	O4'-C1'-N9	6.32	113.26	108.20
84	Aa	297	G	O4'-C1'-N9	6.32	113.26	108.20
84	Aa	538	C	O4'-C1'-N1	6.32	113.26	108.20
84	Aa	1159	C	N3-C4-C5	-6.32	119.37	121.90
84	Aa	1417	G	C5-C6-O6	-6.32	124.81	128.60
84	Aa	2021	G	O4'-C1'-N9	6.32	113.26	108.20
84	Aa	2300	G	C5-C6-O6	-6.32	124.81	128.60
1	Ad	227	G	O4'-C1'-N9	6.32	113.26	108.20
84	Aa	286	C	N3-C4-N4	6.32	122.42	118.00
84	Aa	716	A	C5-C6-N1	-6.32	114.54	117.70
84	Aa	726	C	N3-C4-C5	-6.32	119.37	121.90
84	Aa	1298	A	C5-C6-N1	-6.32	114.54	117.70
84	Aa	1776	G	C5-C6-O6	-6.32	124.81	128.60
84	Aa	419	G	C5-C6-O6	-6.32	124.81	128.60
84	Aa	3168	C	N3-C4-C5	-6.32	119.37	121.90
1	Ad	244	C	C3'-C2'-C1'	6.32	106.56	101.50
1	Ad	1137	A	C3'-C2'-C1'	6.32	106.55	101.50
84	Aa	792	A	C5-C6-N6	-6.32	118.65	123.70
84	Aa	1023	G	O4'-C1'-N9	6.32	113.25	108.20
84	Aa	1664	G	O4'-C1'-N9	6.32	113.25	108.20
84	Aa	2077	C	N3-C4-C5	-6.32	119.37	121.90
84	Aa	2111	A	C4-C5-C6	6.32	120.16	117.00
84	Aa	2300	G	O4'-C1'-N9	6.32	113.25	108.20
84	Aa	2428	G	C5-C6-O6	-6.32	124.81	128.60
84	Aa	3136	A	C4-C5-C6	6.32	120.16	117.00
1	Ad	558	C	C3'-C2'-C1'	6.32	106.55	101.50
84	Aa	1135	C	N3-C4-C5	-6.32	119.37	121.90
84	Aa	2336	C	N3-C4-C5	-6.32	119.37	121.90
84	Aa	3053	G	C5-C6-O6	-6.32	124.81	128.60
1	Ad	728	C	C3'-C2'-C1'	6.31	106.55	101.50
84	Aa	258	C	N3-C4-C5	-6.31	119.37	121.90
84	Aa	1086	U	O4'-C1'-N1	6.31	113.25	108.20
84	Aa	2297	G	O4'-C1'-N9	6.31	113.25	108.20
84	Aa	2534	G	C5-C6-O6	-6.31	124.81	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
17	BS	83	PHE	CB-CG-CD2	6.31	125.22	120.80
84	Aa	1191	U	O4'-C1'-N1	6.31	113.25	108.20
84	Aa	1463	A	C5-C6-N6	-6.31	118.65	123.70
84	Aa	1553	C	N3-C4-C5	-6.31	119.38	121.90
84	Aa	490	G	C5-C6-O6	-6.31	124.81	128.60
84	Aa	507	C	O4'-C1'-N1	6.31	113.25	108.20
84	Aa	1013	A	C5-C6-N1	-6.31	114.55	117.70
84	Aa	2651	G	C5-C6-O6	-6.31	124.81	128.60
84	Aa	2657	C	P-O3'-C3'	6.31	127.27	119.70
84	Aa	550	C	C4'-C3'-O3'	6.31	125.62	113.00
84	Aa	680	G	C5-C6-O6	-6.31	124.81	128.60
84	Aa	1403	G	C5-C6-O6	-6.31	124.81	128.60
84	Aa	1576	C	O4'-C1'-N1	6.31	113.25	108.20
84	Aa	1613	C	N3-C4-C5	-6.31	119.38	121.90
84	Aa	2474	A	C4-C5-C6	6.31	120.16	117.00
84	Aa	2585	C	N3-C4-C5	-6.31	119.38	121.90
84	Aa	2668	U	O4'-C1'-N1	6.31	113.25	108.20
84	Aa	3187	C	N3-C4-N4	6.31	122.42	118.00
1	Ad	344	U	O4'-C1'-C2'	-6.31	99.49	105.80
45	CN	4	TYR	CB-CG-CD1	-6.31	117.22	121.00
84	Aa	141	C	N3-C4-N4	6.31	122.42	118.00
84	Aa	786	U	O4'-C1'-N1	6.31	113.25	108.20
84	Aa	1015	A	O4'-C1'-N9	6.31	113.25	108.20
84	Aa	1310	G	N1-C6-O6	6.31	123.69	119.90
84	Aa	1859	G	C5-C6-O6	-6.31	124.82	128.60
84	Aa	2291	A	C5-C6-N1	-6.31	114.55	117.70
84	Aa	2780	G	C5-C6-O6	-6.31	124.82	128.60
84	Aa	3229	C	O4'-C1'-N1	6.31	113.25	108.20
84	Aa	3319	G	O4'-C1'-N9	6.31	113.25	108.20
84	Aa	2973	A	O4'-C1'-N9	6.31	113.25	108.20
84	Aa	2983	U	O4'-C1'-N1	6.31	113.25	108.20
1	Ad	1783	C	N1-C1'-C2'	6.30	122.20	114.00
84	Aa	838	G	O4'-C1'-N9	6.30	113.24	108.20
84	Aa	1188	C	N3-C4-C5	-6.30	119.38	121.90
84	Aa	2509	A	C5-C6-N1	-6.30	114.55	117.70
84	Aa	3136	A	C5-C6-N1	-6.30	114.55	117.70
84	Aa	3281	G	O4'-C1'-N9	6.30	113.24	108.20
84	Aa	2017	G	C5-C6-O6	-6.30	124.82	128.60
84	Aa	2561	A	C5-C6-N6	-6.30	118.66	123.70
1	Ad	614	G	P-O3'-C3'	6.30	127.26	119.70
2	Ae	46	A	O4'-C1'-N9	6.30	113.24	108.20
41	CA	67	PHE	N-CA-CB	6.30	121.94	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	456	G	O4'-C1'-N9	6.30	113.24	108.20
84	Aa	621	C	N3-C4-N4	6.30	122.41	118.00
84	Aa	1332	C	O4'-C1'-N1	6.30	113.24	108.20
84	Aa	1662	G	O4'-C1'-N9	6.30	113.24	108.20
84	Aa	2027	G	O4'-C1'-N9	6.30	113.24	108.20
84	Aa	2199	C	O4'-C1'-N1	6.30	113.24	108.20
84	Aa	2307	A	O4'-C1'-N9	6.30	113.24	108.20
84	Aa	2311	A	O4'-C1'-N9	6.30	113.24	108.20
84	Aa	3247	C	N3-C4-C5	-6.30	119.38	121.90
85	Ac	125	C	O4'-C1'-N1	6.30	113.24	108.20
85	Ac	157	A	O4'-C1'-N9	6.30	113.24	108.20
84	Aa	428	G	O4'-C1'-N9	6.30	113.24	108.20
84	Aa	1675	G	C5-C6-O6	-6.30	124.82	128.60
86	Ab	85	G	N1-C6-O6	6.30	123.68	119.90
84	Aa	1637	G	C5-C6-O6	-6.30	124.82	128.60
84	Aa	2162	C	O4'-C4'-C3'	-6.30	97.70	104.00
1	Ad	353	G	O4'-C1'-N9	6.30	113.24	108.20
48	CD	262	ALA	N-CA-CB	6.30	118.92	110.10
84	Aa	875	A	O4'-C1'-N9	6.30	113.24	108.20
84	Aa	1936	G	O4'-C1'-N9	6.30	113.24	108.20
84	Aa	2257	A	O4'-C1'-N9	6.30	113.24	108.20
84	Aa	2294	A	C4-C5-C6	6.30	120.15	117.00
84	Aa	2345	C	N3-C4-C5	-6.30	119.38	121.90
84	Aa	121	A	C4-C5-C6	6.29	120.15	117.00
84	Aa	1340	G	N3-C2-N2	6.29	124.31	119.90
84	Aa	2730	A	C4-C5-C6	6.29	120.15	117.00
84	Aa	2985	C	N3-C4-C5	-6.29	119.38	121.90
1	Ad	1237	G	O4'-C1'-C2'	6.29	113.26	107.60
84	Aa	401	C	N3-C4-C5	-6.29	119.38	121.90
84	Aa	415	G	C5-C6-O6	-6.29	124.82	128.60
84	Aa	573	A	C5-C6-N6	-6.29	118.67	123.70
84	Aa	886	A	C4-C5-C6	6.29	120.15	117.00
84	Aa	2053	A	C4-C5-C6	6.29	120.15	117.00
84	Aa	2144	G	C5-C6-O6	-6.29	124.82	128.60
84	Aa	2798	G	N1-C6-O6	6.29	123.68	119.90
84	Aa	3124	A	C4-C5-C6	6.29	120.15	117.00
1	Ad	1232	G	C1'-O4'-C4'	-6.29	104.87	109.90
84	Aa	8	C	N3-C4-C5	-6.29	119.38	121.90
84	Aa	761	C	N3-C4-C5	-6.29	119.38	121.90
84	Aa	782	G	C5-C6-O6	-6.29	124.83	128.60
84	Aa	2332	C	N3-C4-C5	-6.29	119.38	121.90
84	Aa	2907	U	O4'-C1'-N1	6.29	113.23	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3287	A	O4'-C1'-N9	6.29	113.23	108.20
84	Aa	1427	C	N3-C4-C5	-6.29	119.38	121.90
84	Aa	2122	C	N3-C4-C5	-6.29	119.38	121.90
84	Aa	3263	C	C5'-C4'-C3'	6.29	126.06	116.00
84	Aa	124	C	N3-C4-C5	-6.29	119.38	121.90
84	Aa	535	G	C5-C6-O6	-6.29	124.83	128.60
84	Aa	719	U	P-O5'-C5'	6.29	130.96	120.90
84	Aa	1695	C	C2-N3-C4	6.29	123.04	119.90
84	Aa	2114	A	C4-C5-C6	6.29	120.14	117.00
84	Aa	2311	A	C5-C6-N6	-6.29	118.67	123.70
84	Aa	3166	C	O4'-C1'-N1	6.29	113.23	108.20
86	Ab	108	G	C4-C5-N7	-6.29	108.28	110.80
1	Ad	901	U	O4'-C1'-N1	6.29	113.23	108.20
84	Aa	1264	A	C4-C5-C6	6.29	120.14	117.00
84	Aa	2357	A	C5-C6-N6	-6.29	118.67	123.70
84	Aa	3089	G	C5-C6-O6	-6.29	124.83	128.60
1	Ad	252	U	O4'-C1'-C2'	-6.29	99.51	105.80
1	Ad	561	G	O4'-C1'-N9	-6.29	103.17	108.20
1	Ad	744	G	P-O5'-C5'	6.29	130.96	120.90
1	Ad	1112	G	N9-C1'-C2'	6.29	122.17	114.00
46	Ca	52	TYR	CB-CG-CD2	-6.29	117.23	121.00
84	Aa	700	C	N3-C4-C5	-6.29	119.39	121.90
84	Aa	830	A	C5-C6-N1	-6.29	114.56	117.70
84	Aa	1729	G	O4'-C1'-N9	6.29	113.23	108.20
84	Aa	1999	G	C5-C6-O6	-6.29	124.83	128.60
84	Aa	2427	C	N3-C4-C5	-6.29	119.39	121.90
84	Aa	2923	U	O4'-C1'-N1	6.29	113.23	108.20
84	Aa	3049	A	C5-C6-N6	-6.29	118.67	123.70
85	Ac	58	G	O4'-C1'-N9	6.29	113.23	108.20
1	Ad	1354	C	O4'-C1'-N1	6.28	113.23	108.20
1	Ad	1524	A	P-O3'-C3'	6.28	127.24	119.70
50	CP	4	TYR	CB-CG-CD2	-6.28	117.23	121.00
84	Aa	138	G	O4'-C1'-N9	6.28	113.23	108.20
84	Aa	359	A	C5-C6-N1	-6.28	114.56	117.70
84	Aa	1057	A	C4-C5-C6	6.28	120.14	117.00
84	Aa	1436	A	C4-C5-C6	6.28	120.14	117.00
84	Aa	1481	C	N3-C4-C5	-6.28	119.39	121.90
84	Aa	2412	A	C5-C6-N1	-6.28	114.56	117.70
84	Aa	2548	U	O4'-C1'-N1	6.28	113.23	108.20
84	Aa	2990	C	N3-C4-C5	-6.28	119.39	121.90
85	Ac	152	G	C5-C6-O6	-6.28	124.83	128.60
84	Aa	47	A	O4'-C1'-N9	6.28	113.23	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1162	A	C4-C5-C6	6.28	120.14	117.00
84	Aa	1206	A	C4-C5-C6	6.28	120.14	117.00
84	Aa	1501	A	C4-C5-C6	6.28	120.14	117.00
84	Aa	1986	G	O4'-C1'-N9	6.28	113.23	108.20
1	Ad	318	C	C3'-C2'-C1'	6.28	106.52	101.50
84	Aa	20	G	O4'-C1'-N9	6.28	113.22	108.20
84	Aa	1222	U	O4'-C1'-N1	6.28	113.22	108.20
84	Aa	1267	A	N1-C6-N6	6.28	122.37	118.60
84	Aa	2892	A	O4'-C1'-N9	6.28	113.22	108.20
1	Ad	1292	G	O4'-C1'-N9	6.28	113.22	108.20
84	Aa	218	G	O4'-C1'-N9	6.28	113.22	108.20
84	Aa	1619	G	C5-C6-O6	-6.28	124.83	128.60
84	Aa	3391	U	O4'-C1'-N1	6.28	113.22	108.20
1	Ad	976	A	O4'-C1'-N9	6.28	113.22	108.20
84	Aa	466	U	O4'-C1'-N1	6.28	113.22	108.20
84	Aa	485	G	C5-C6-O6	-6.28	124.83	128.60
84	Aa	2751	A	C5-C6-N6	-6.28	118.68	123.70
84	Aa	2781	A	C5-C6-N1	-6.28	114.56	117.70
84	Aa	3195	C	N3-C4-N4	6.28	122.39	118.00
1	Ad	162	A	O4'-C1'-N9	6.28	113.22	108.20
84	Aa	1139	A	O4'-C1'-N9	6.28	113.22	108.20
84	Aa	1477	A	C5-C6-N1	-6.28	114.56	117.70
84	Aa	1495	G	O4'-C1'-N9	6.28	113.22	108.20
84	Aa	1657	C	N3-C4-C5	-6.28	119.39	121.90
84	Aa	3266	U	O4'-C1'-N1	6.28	113.22	108.20
85	Ac	13	A	C5-C6-N1	-6.28	114.56	117.70
85	Ac	60	U	O4'-C1'-N1	6.28	113.22	108.20
85	Ac	114	G	O4'-C1'-N9	6.28	113.22	108.20
84	Aa	510	C	N3-C4-C5	-6.27	119.39	121.90
84	Aa	968	A	C4-C5-C6	6.27	120.14	117.00
84	Aa	1567	G	C5-C6-O6	-6.27	124.84	128.60
84	Aa	2735	G	O4'-C1'-N9	6.27	113.22	108.20
84	Aa	2395	G	O4'-C1'-N9	6.27	113.22	108.20
84	Aa	2562	A	C4-C5-C6	6.27	120.14	117.00
84	Aa	2795	G	C5-C6-O6	-6.27	124.84	128.60
84	Aa	723	G	O3'-P-O5'	6.27	115.92	104.00
84	Aa	3107	A	O4'-C1'-N9	6.27	113.22	108.20
1	Ad	931	A	O4'-C1'-N9	6.27	113.22	108.20
84	Aa	143	A	C4-C5-C6	6.27	120.13	117.00
84	Aa	751	C	N3-C4-C5	-6.27	119.39	121.90
84	Aa	835	G	C5'-C4'-C3'	6.27	126.03	116.00
1	Ad	719	C	O4'-C1'-C2'	-6.27	99.53	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	73	A	C4-C5-C6	6.27	120.13	117.00
84	Aa	1278	A	C4-C5-C6	6.27	120.13	117.00
84	Aa	2113	A	C4-C5-C6	6.27	120.13	117.00
84	Aa	2354	G	O4'-C1'-N9	6.27	113.22	108.20
84	Aa	2687	C	N3-C4-C5	-6.27	119.39	121.90
84	Aa	3055	U	O4'-C1'-N1	6.27	113.22	108.20
84	Aa	3133	C	N3-C4-C5	-6.27	119.39	121.90
84	Aa	3299	A	C5-C6-N1	-6.27	114.57	117.70
84	Aa	3337	G	C5-C6-O6	-6.27	124.84	128.60
1	Ad	1421	U	C1'-O4'-C4'	-6.27	104.89	109.90
84	Aa	2254	A	C4-C5-C6	6.27	120.13	117.00
84	Aa	2493	C	N3-C4-C5	-6.27	119.39	121.90
1	Ad	788	G	O4'-C1'-N9	-6.26	103.19	108.20
1	Ad	888	U	C3'-C2'-C1'	-6.26	96.49	101.50
84	Aa	77	U	O4'-C1'-N1	6.26	113.21	108.20
84	Aa	132	U	O4'-C1'-N1	6.26	113.21	108.20
84	Aa	401	C	O4'-C1'-N1	6.26	113.21	108.20
84	Aa	508	G	O4'-C1'-N9	6.26	113.21	108.20
84	Aa	640	C	N3-C4-C5	-6.26	119.39	121.90
84	Aa	949	C	N3-C4-N4	6.26	122.39	118.00
84	Aa	1039	G	O4'-C1'-N9	6.26	113.21	108.20
84	Aa	1806	C	N3-C4-C5	-6.26	119.39	121.90
84	Aa	1985	G	O4'-C1'-N9	6.26	113.21	108.20
84	Aa	3152	C	O4'-C1'-N1	6.26	113.21	108.20
85	Ac	51	G	C5-C6-O6	-6.26	124.84	128.60
1	Ad	1217	G	O4'-C1'-N9	6.26	113.21	108.20
84	Aa	489	C	N3-C4-N4	6.26	122.38	118.00
84	Aa	931	C	N3-C4-C5	-6.26	119.39	121.90
84	Aa	2034	G	C5-C6-O6	-6.26	124.84	128.60
84	Aa	2316	A	C4-C5-C6	6.26	120.13	117.00
63	CU	84	TYR	CB-CG-CD1	6.26	124.76	121.00
71	CB	118	PHE	CB-CG-CD2	-6.26	116.42	120.80
84	Aa	420	A	C4-C5-C6	6.26	120.13	117.00
84	Aa	795	C	N3-C4-C5	-6.26	119.39	121.90
84	Aa	912	G	O4'-C1'-N9	6.26	113.21	108.20
84	Aa	929	A	C5-C6-N6	-6.26	118.69	123.70
84	Aa	1880	A	O4'-C1'-N9	6.26	113.21	108.20
84	Aa	3314	G	C5-C6-O6	-6.26	124.84	128.60
1	Ad	174	C	O4'-C1'-C2'	-6.26	99.54	105.80
1	Ad	417	U	O4'-C1'-N1	6.26	113.21	108.20
1	Ad	1732	A	O4'-C1'-C2'	-6.26	99.54	105.80
84	Aa	287	A	C5-C6-N1	-6.26	114.57	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	864	C	N3-C4-C5	-6.26	119.40	121.90
84	Aa	866	C	N3-C4-C5	-6.26	119.40	121.90
84	Aa	1000	A	C4-C5-C6	6.26	120.13	117.00
84	Aa	2157	C	N3-C4-C5	-6.26	119.40	121.90
84	Aa	2372	A	C5-C6-N6	-6.26	118.69	123.70
84	Aa	2800	C	N3-C4-C5	-6.26	119.40	121.90
1	Ad	975	A	O4'-C1'-C2'	-6.26	99.54	105.80
1	Ad	1123	G	O4'-C1'-N9	6.26	113.21	108.20
1	Ad	368	A	O4'-C1'-N9	6.26	113.20	108.20
84	Aa	212	G	C5-C6-O6	-6.26	124.85	128.60
84	Aa	962	C	P-O3'-C3'	6.26	127.21	119.70
84	Aa	1405	G	C5-C6-O6	-6.26	124.85	128.60
84	Aa	2358	C	N3-C4-C5	-6.26	119.40	121.90
84	Aa	2804	A	C4-C5-C6	6.26	120.13	117.00
1	Ad	94	A	C1'-O4'-C4'	-6.25	104.90	109.90
1	Ad	957	A	O4'-C1'-N9	6.25	113.20	108.20
1	Ad	1157	A	O4'-C1'-N9	6.25	113.20	108.20
84	Aa	536	C	N3-C4-C5	-6.25	119.40	121.90
84	Aa	733	C	N3-C4-N4	6.25	122.38	118.00
84	Aa	1783	G	O4'-C1'-N9	6.25	113.20	108.20
84	Aa	1932	A	C5-C6-N1	-6.25	114.57	117.70
84	Aa	2213	G	C5-C6-O6	-6.25	124.85	128.60
84	Aa	2325	A	C5-C6-N1	-6.25	114.57	117.70
1	Ad	1722	C	O4'-C1'-N1	6.25	113.20	108.20
16	BO	72	ALA	N-CA-CB	6.25	118.85	110.10
84	Aa	836	G	C5-C6-O6	-6.25	124.85	128.60
84	Aa	2008	G	O4'-C1'-N9	6.25	113.20	108.20
84	Aa	2742	A	C5-C6-N1	-6.25	114.57	117.70
84	Aa	3347	U	O4'-C1'-N1	6.25	113.20	108.20
85	Ac	11	C	N3-C4-N4	6.25	122.38	118.00
84	Aa	614	C	N3-C4-N4	6.25	122.38	118.00
84	Aa	1724	C	N3-C4-N4	6.25	122.38	118.00
84	Aa	2256	G	O4'-C1'-N9	6.25	113.20	108.20
85	Ac	24	G	C5-C6-O6	-6.25	124.85	128.60
84	Aa	980	C	N3-C4-C5	-6.25	119.40	121.90
84	Aa	1539	G	O4'-C1'-N9	6.25	113.20	108.20
84	Aa	1799	C	N3-C4-N4	6.25	122.37	118.00
84	Aa	2627	G	C5-C6-O6	-6.25	124.85	128.60
28	BA	43	TYR	CB-CG-CD2	-6.25	117.25	121.00
84	Aa	1804	G	C5-C6-O6	-6.25	124.85	128.60
84	Aa	1843	A	C5-C6-N6	-6.25	118.70	123.70
85	Ac	43	A	C5-C6-N1	-6.25	114.58	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	498	G	C5-C6-O6	-6.25	124.85	128.60
84	Aa	587	A	C5-C6-N1	-6.25	114.58	117.70
84	Aa	2609	G	C5-C6-O6	-6.25	124.85	128.60
84	Aa	3111	C	N3-C4-C5	-6.25	119.40	121.90
1	Ad	842	G	C3'-C2'-C1'	-6.24	96.51	101.50
1	Ad	1525	U	O4'-C1'-N1	6.24	113.19	108.20
84	Aa	454	A	C5-C6-N1	-6.24	114.58	117.70
84	Aa	1644	A	C5-C6-N1	-6.24	114.58	117.70
84	Aa	2108	C	N3-C4-C5	-6.24	119.40	121.90
84	Aa	2638	A	C5-C6-N6	-6.24	118.70	123.70
85	Ac	31	G	C5-C6-O6	-6.24	124.85	128.60
15	BU	3	ALA	N-CA-CB	6.24	118.84	110.10
84	Aa	583	C	N3-C4-C5	-6.24	119.40	121.90
84	Aa	661	A	C5-C6-N1	-6.24	114.58	117.70
84	Aa	1101	A	C5-C6-N1	-6.24	114.58	117.70
84	Aa	1989	G	C5-C6-O6	-6.24	124.86	128.60
84	Aa	2418	A	C4-C5-C6	6.24	120.12	117.00
84	Aa	2712	C	N3-C4-C5	-6.24	119.40	121.90
84	Aa	2898	A	C4-C5-C6	6.24	120.12	117.00
1	Ad	397	C	O4'-C1'-C2'	-6.24	99.56	105.80
84	Aa	50	A	C4-C5-C6	6.24	120.12	117.00
84	Aa	1072	C	N3-C4-N4	6.24	122.37	118.00
84	Aa	1554	C	N3-C4-C5	-6.24	119.40	121.90
84	Aa	2051	G	C5-C6-O6	-6.24	124.86	128.60
84	Aa	2401	A	C4-C5-C6	6.24	120.12	117.00
85	Ac	59	A	C5-C6-N1	-6.24	114.58	117.70
1	Ad	1074	C	C3'-C2'-C1'	6.24	106.49	101.50
1	Ad	1465	C	O4'-C1'-C2'	-6.24	99.56	105.80
38	CT	19	PHE	CB-CG-CD1	6.24	125.17	120.80
84	Aa	642	C	C2-N1-C1'	6.24	125.66	118.80
84	Aa	1200	A	C4-C5-C6	6.24	120.12	117.00
84	Aa	2144	G	O4'-C1'-N9	6.24	113.19	108.20
84	Aa	2619	C	N3-C4-C5	-6.24	119.40	121.90
85	Ac	92	A	C4-C5-C6	6.24	120.12	117.00
41	CA	76	PHE	CB-CG-CD1	6.24	125.17	120.80
84	Aa	2291	A	C4-C5-C6	6.24	120.12	117.00
84	Aa	2765	A	C4-C5-C6	6.24	120.12	117.00
84	Aa	3382	A	C5-C6-N6	-6.24	118.71	123.70
1	Ad	1787	G	O4'-C1'-N9	6.24	113.19	108.20
84	Aa	1172	A	C5-C6-N6	-6.24	118.71	123.70
84	Aa	2011	G	C5-C6-O6	-6.24	124.86	128.60
84	Aa	2720	U	O4'-C1'-N1	6.24	113.19	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2882	U	O4'-C1'-N1	6.24	113.19	108.20
84	Aa	2905	A	C5-C6-N6	-6.24	118.71	123.70
84	Aa	3203	G	O4'-C1'-N9	6.24	113.19	108.20
84	Aa	1640	A	C4-C5-C6	6.23	120.12	117.00
84	Aa	2109	G	C5-C6-O6	-6.23	124.86	128.60
1	Ad	1461	G	C1'-O4'-C4'	-6.23	104.92	109.90
84	Aa	1704	A	C5-C6-N6	-6.23	118.71	123.70
84	Aa	2956	U	P-O3'-C3'	6.23	127.18	119.70
84	Aa	3057	A	C5-C6-N6	-6.23	118.71	123.70
86	Ab	7	G	C6-N1-C2	6.23	128.84	125.10
84	Aa	1279	C	N3-C4-C5	-6.23	119.41	121.90
84	Aa	1572	C	N3-C4-C5	-6.23	119.41	121.90
84	Aa	2129	U	O4'-C1'-N1	6.23	113.18	108.20
84	Aa	2265	A	O4'-C1'-N9	6.23	113.19	108.20
84	Aa	2996	A	C5-C6-N6	-6.23	118.72	123.70
84	Aa	214	G	O4'-C1'-N9	6.23	113.18	108.20
84	Aa	1281	C	P-O3'-C3'	6.23	127.17	119.70
84	Aa	1779	C	N3-C4-C5	-6.23	119.41	121.90
84	Aa	1860	A	O4'-C1'-N9	6.23	113.18	108.20
1	Ad	429	A	O4'-C1'-N9	6.23	113.18	108.20
1	Ad	510	A	C1'-O4'-C4'	6.23	114.88	109.90
1	Ad	951	U	O4'-C1'-N1	6.23	113.18	108.20
84	Aa	419	G	O4'-C1'-N9	6.23	113.18	108.20
84	Aa	486	G	O4'-C1'-N9	6.23	113.18	108.20
84	Aa	718	C	C5'-C4'-O4'	6.23	116.57	109.10
84	Aa	853	U	O4'-C1'-N1	6.23	113.18	108.20
84	Aa	1705	A	C5-C6-N1	-6.23	114.59	117.70
84	Aa	1880	A	C4-C5-C6	6.23	120.11	117.00
84	Aa	2774	A	C4-C5-C6	6.23	120.11	117.00
84	Aa	2918	U	O4'-C1'-N1	6.23	113.18	108.20
86	Ab	1	G	N1-C6-O6	6.23	123.64	119.90
84	Aa	357	C	N3-C4-N4	6.23	122.36	118.00
84	Aa	657	A	C4-C5-C6	6.23	120.11	117.00
84	Aa	985	C	N3-C4-N4	6.23	122.36	118.00
84	Aa	1075	G	C5-C6-O6	-6.23	124.86	128.60
84	Aa	2070	C	N3-C4-C5	-6.23	119.41	121.90
84	Aa	3090	C	N3-C4-C5	-6.23	119.41	121.90
1	Ad	277	G	N9-C1'-C2'	6.22	122.09	114.00
84	Aa	109	G	C5-C6-O6	-6.22	124.86	128.60
84	Aa	1000	A	C5-C6-N6	-6.22	118.72	123.70
84	Aa	1714	A	C4-C5-C6	6.22	120.11	117.00
84	Aa	1805	A	O4'-C1'-N9	6.22	113.18	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2098	A	C5-C6-N6	-6.22	118.72	123.70
84	Aa	2484	G	C5-C6-O6	-6.22	124.86	128.60
84	Aa	2670	A	C5-C6-N6	-6.22	118.72	123.70
1	Ad	223	A	N9-C1'-C2'	6.22	122.09	114.00
1	Ad	1503	C	N1-C1'-C2'	6.22	122.09	114.00
1	Ad	1581	A	C3'-C2'-C1'	6.22	106.48	101.50
84	Aa	1644	A	C5-C6-N6	-6.22	118.72	123.70
84	Aa	3201	A	C4-C5-C6	6.22	120.11	117.00
1	Ad	1543	U	C3'-C2'-C1'	6.22	106.48	101.50
84	Aa	2483	A	O4'-C1'-N9	6.22	113.18	108.20
1	Ad	1501	G	P-O3'-C3'	6.22	127.16	119.70
1	Ad	1670	G	O4'-C1'-N9	6.22	113.17	108.20
84	Aa	1294	A	O4'-C1'-N9	6.22	113.17	108.20
84	Aa	1928	A	C5-C6-N1	-6.22	114.59	117.70
84	Aa	2542	U	O4'-C1'-N1	6.22	113.17	108.20
84	Aa	2676	A	C5-C6-N1	-6.22	114.59	117.70
84	Aa	2859	C	N3-C4-C5	-6.22	119.41	121.90
84	Aa	3333	C	N3-C4-C5	-6.22	119.41	121.90
84	Aa	568	C	N3-C4-C5	-6.22	119.41	121.90
84	Aa	842	C	N3-C4-C5	-6.22	119.41	121.90
84	Aa	1157	A	C5-C6-N1	-6.22	114.59	117.70
84	Aa	1282	A	C5-C6-N6	-6.22	118.73	123.70
1	Ad	1303	G	P-O3'-C3'	6.22	127.16	119.70
84	Aa	640	C	C5'-C4'-O4'	6.22	116.56	109.10
84	Aa	1381	G	C5-C6-O6	-6.22	124.87	128.60
84	Aa	1819	A	O4'-C1'-N9	6.22	113.17	108.20
84	Aa	1902	G	O4'-C1'-N9	6.22	113.17	108.20
84	Aa	2205	G	C5-C6-O6	-6.22	124.87	128.60
84	Aa	2448	G	C5-C6-O6	-6.22	124.87	128.60
84	Aa	2645	A	C5-C6-N6	-6.22	118.73	123.70
84	Aa	2679	A	C5-C6-N6	-6.22	118.73	123.70
84	Aa	262	A	C4-C5-C6	6.21	120.11	117.00
84	Aa	1089	G	O4'-C1'-N9	6.21	113.17	108.20
84	Aa	1097	A	C4-C5-C6	6.21	120.11	117.00
84	Aa	1754	C	N3-C4-C5	-6.21	119.41	121.90
84	Aa	1927	A	C5-C6-N6	-6.21	118.73	123.70
84	Aa	2087	A	C1'-O4'-C4'	-6.21	104.93	109.90
84	Aa	2899	A	C5-C6-N1	-6.21	114.59	117.70
84	Aa	3003	C	O4'-C1'-N1	6.21	113.17	108.20
85	Ac	58	G	C5-C6-O6	-6.21	124.87	128.60
1	Ad	750	U	N1-C1'-C2'	6.21	122.08	114.00
84	Aa	721	A	C5-C6-N6	-6.21	118.73	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1114	A	O4'-C1'-N9	6.21	113.17	108.20
84	Aa	1273	U	C2-N1-C1'	6.21	125.16	117.70
1	Ad	1687	G	P-O3'-C3'	6.21	127.16	119.70
84	Aa	1036	C	N3-C4-C5	-6.21	119.42	121.90
84	Aa	1713	A	O4'-C1'-N9	6.21	113.17	108.20
84	Aa	2350	C	O4'-C1'-N1	6.21	113.17	108.20
84	Aa	3295	G	C5-C6-O6	-6.21	124.87	128.60
84	Aa	3332	G	C5-C6-O6	-6.21	124.87	128.60
84	Aa	433	C	N3-C4-C5	-6.21	119.42	121.90
84	Aa	513	C	P-O3'-C3'	6.21	127.15	119.70
84	Aa	1149	C	N3-C4-C5	-6.21	119.42	121.90
84	Aa	1978	G	O4'-C1'-N9	6.21	113.17	108.20
84	Aa	48	A	C4-C5-C6	6.21	120.10	117.00
84	Aa	586	A	C4-C5-C6	6.21	120.10	117.00
84	Aa	1814	C	N3-C4-N4	6.21	122.35	118.00
84	Aa	2268	G	C5-C6-O6	-6.21	124.87	128.60
84	Aa	2319	A	O4'-C1'-N9	6.21	113.17	108.20
84	Aa	2402	G	C5-C6-O6	-6.21	124.88	128.60
1	Ad	1163	C	C3'-C2'-C1'	6.21	106.47	101.50
84	Aa	834	G	C5-C6-O6	-6.21	124.88	128.60
84	Aa	887	A	C4-C5-C6	6.21	120.10	117.00
84	Aa	2436	G	C2'-C3'-O3'	-6.21	95.84	109.50
84	Aa	2667	C	N3-C4-C5	-6.21	119.42	121.90
84	Aa	3254	C	N3-C4-C5	-6.21	119.42	121.90
86	Ab	49	A	O4'-C1'-N9	6.21	113.17	108.20
1	Ad	303	A	O4'-C1'-C2'	-6.21	99.59	105.80
1	Ad	402	G	C3'-C2'-C1'	6.21	106.46	101.50
84	Aa	1224	A	C4-C5-C6	6.21	120.10	117.00
84	Aa	1930	G	O4'-C1'-N9	6.21	113.16	108.20
84	Aa	2572	U	O4'-C1'-N1	6.21	113.16	108.20
84	Aa	385	A	C4-C5-C6	6.20	120.10	117.00
84	Aa	1117	U	O4'-C1'-N1	6.20	113.16	108.20
84	Aa	1850	C	N3-C4-C5	-6.20	119.42	121.90
84	Aa	2411	G	C5-C6-O6	-6.20	124.88	128.60
84	Aa	2810	A	C5-C6-N6	-6.20	118.74	123.70
84	Aa	219	A	C4-C5-C6	6.20	120.10	117.00
84	Aa	1403	G	O4'-C1'-N9	6.20	113.16	108.20
1	Ad	24	U	O4'-C1'-N1	6.20	113.16	108.20
1	Ad	1755	G	C3'-C2'-C1'	-6.20	96.54	101.50
10	Bg	342	SER	N-CA-CB	6.20	119.80	110.50
84	Aa	516	C	N3-C4-C5	-6.20	119.42	121.90
84	Aa	663	G	O4'-C1'-N9	6.20	113.16	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1113	C	N3-C4-C5	-6.20	119.42	121.90
84	Aa	1892	A	C4-C5-C6	6.20	120.10	117.00
84	Aa	3232	C	N3-C4-C5	-6.20	119.42	121.90
84	Aa	3244	G	O4'-C1'-N9	6.20	113.16	108.20
84	Aa	3356	C	N3-C4-C5	-6.20	119.42	121.90
85	Ac	148	C	N3-C4-N4	6.20	122.34	118.00
1	Ad	27	U	C5'-C4'-O4'	6.20	116.54	109.10
1	Ad	357	A	C1'-O4'-C4'	6.20	114.86	109.90
1	Ad	1487	U	O4'-C1'-N1	6.20	113.16	108.20
84	Aa	186	A	C4-C5-C6	6.20	120.10	117.00
84	Aa	724	A	C5-C6-N1	-6.20	114.60	117.70
84	Aa	951	C	N3-C4-C5	-6.20	119.42	121.90
84	Aa	1030	A	C4-C5-C6	6.20	120.10	117.00
84	Aa	1285	U	O4'-C1'-N1	6.20	113.16	108.20
84	Aa	1514	U	O4'-C1'-N1	6.20	113.16	108.20
84	Aa	1517	C	C2-N1-C1'	6.20	125.62	118.80
84	Aa	2418	A	C5-C6-N1	-6.20	114.60	117.70
84	Aa	2869	C	N3-C4-C5	-6.20	119.42	121.90
85	Ac	157	A	C4-C5-C6	6.20	120.10	117.00
1	Ad	954	C	N1-C1'-C2'	6.20	122.06	114.00
2	Ae	20	C	O4'-C1'-N1	6.20	113.16	108.20
84	Aa	1246	G	C5-C6-O6	-6.20	124.88	128.60
1	Ad	934	A	P-O3'-C3'	6.20	127.13	119.70
1	Ad	1360	G	O4'-C1'-N9	6.20	113.16	108.20
84	Aa	32	G	O4'-C1'-N9	6.20	113.16	108.20
84	Aa	289	C	N3-C4-N4	6.20	122.34	118.00
84	Aa	1225	A	C5-C6-N6	-6.20	118.74	123.70
84	Aa	1840	C	N3-C4-C5	-6.20	119.42	121.90
84	Aa	2887	C	N3-C4-C5	-6.20	119.42	121.90
84	Aa	3135	A	C4-C5-C6	6.20	120.10	117.00
1	Ad	498	U	N1-C1'-C2'	-6.19	105.19	112.00
84	Aa	2904	A	C5-C6-N6	-6.19	118.74	123.70
84	Aa	850	A	C4-C5-C6	6.19	120.10	117.00
84	Aa	1687	C	N3-C4-C5	-6.19	119.42	121.90
84	Aa	2079	A	C5-C6-N1	-6.19	114.60	117.70
84	Aa	2912	A	C4-C5-C6	6.19	120.10	117.00
84	Aa	2978	A	C4-C5-C6	6.19	120.10	117.00
84	Aa	3220	A	C4-C5-C6	6.19	120.10	117.00
1	Ad	899	A	P-O3'-C3'	6.19	127.13	119.70
1	Ad	1663	A	C3'-C2'-C1'	6.19	106.45	101.50
84	Aa	199	G	C5-C6-O6	-6.19	124.89	128.60
84	Aa	645	C	N3-C4-C5	-6.19	119.42	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1405	G	O4'-C1'-N9	6.19	113.15	108.20
84	Aa	1733	G	O4'-C1'-N9	6.19	113.15	108.20
84	Aa	3386	A	C5-C6-N1	-6.19	114.61	117.70
86	Ab	118	C	N3-C4-C5	-6.19	119.42	121.90
84	Aa	968	A	C5-C6-N1	-6.19	114.61	117.70
84	Aa	1284	C	N3-C4-C5	-6.19	119.42	121.90
1	Ad	470	U	O4'-C1'-N1	6.19	113.15	108.20
84	Aa	98	A	C4-C5-C6	6.19	120.09	117.00
84	Aa	1095	C	N3-C4-C5	-6.19	119.42	121.90
84	Aa	1868	C	O4'-C1'-N1	6.19	113.15	108.20
84	Aa	3115	A	O4'-C1'-N9	6.19	113.15	108.20
1	Ad	1127	G	O4'-C1'-N9	6.19	113.15	108.20
61	CM	6	PHE	CB-CG-CD1	-6.19	116.47	120.80
84	Aa	1232	A	C4-C5-C6	6.19	120.09	117.00
84	Aa	1863	A	C4-C5-C6	6.19	120.09	117.00
84	Aa	3274	G	C5-C6-O6	-6.19	124.89	128.60
84	Aa	348	C	N3-C4-N4	6.18	122.33	118.00
84	Aa	802	G	O4'-C1'-N9	6.18	113.15	108.20
84	Aa	1068	A	C4-C5-C6	6.18	120.09	117.00
84	Aa	1818	C	N3-C4-C5	-6.18	119.43	121.90
84	Aa	1861	A	C4-C5-C6	6.18	120.09	117.00
84	Aa	1892	A	C5-C6-N1	-6.18	114.61	117.70
84	Aa	1986	G	C5-C6-O6	-6.18	124.89	128.60
84	Aa	2963	G	C5-C6-O6	-6.18	124.89	128.60
84	Aa	3386	A	O4'-C1'-N9	6.18	113.15	108.20
1	Ad	943	G	O4'-C1'-N9	6.18	113.15	108.20
84	Aa	88	A	C5-C6-N1	-6.18	114.61	117.70
84	Aa	262	A	C5-C6-N6	-6.18	118.75	123.70
84	Aa	304	A	C4-C5-C6	6.18	120.09	117.00
84	Aa	1917	A	C4-C5-C6	6.18	120.09	117.00
84	Aa	1929	A	C5-C6-N6	-6.18	118.75	123.70
84	Aa	2480	G	P-O5'-C5'	6.18	130.79	120.90
84	Aa	2921	A	C5-C6-N1	-6.18	114.61	117.70
84	Aa	849	A	C4-C5-C6	6.18	120.09	117.00
84	Aa	3009	A	C4-C5-C6	6.18	120.09	117.00
1	Ad	201	G	O4'-C1'-C2'	6.18	113.16	107.60
1	Ad	715	U	P-O5'-C5'	6.18	130.79	120.90
84	Aa	122	A	C5-C6-N1	-6.18	114.61	117.70
84	Aa	995	C	N3-C4-C5	-6.18	119.43	121.90
84	Aa	1478	A	C5-C6-N1	-6.18	114.61	117.70
84	Aa	1791	U	O4'-C1'-N1	6.18	113.14	108.20
84	Aa	3384	G	P-O3'-C3'	6.18	127.11	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	Ab	81	G	C5-C6-O6	-6.18	124.89	128.60
84	Aa	192	C	N3-C4-C5	-6.18	119.43	121.90
85	Ac	31	G	O4'-C1'-N9	6.18	113.14	108.20
1	Ad	1690	U	N1-C1'-C2'	-6.18	105.20	112.00
84	Aa	137	C	N3-C4-C5	-6.18	119.43	121.90
84	Aa	1031	A	C4-C5-C6	6.18	120.09	117.00
84	Aa	1249	A	C4-C5-C6	6.18	120.09	117.00
84	Aa	1523	G	O4'-C1'-N9	6.18	113.14	108.20
84	Aa	2015	G	O4'-C1'-N9	6.18	113.14	108.20
1	Ad	902	C	C3'-C2'-C1'	6.17	106.44	101.50
1	Ad	1206	A	C3'-C2'-C1'	6.17	106.44	101.50
84	Aa	127	G	C5-C6-O6	-6.17	124.89	128.60
84	Aa	634	A	C5-C6-N1	-6.17	114.61	117.70
84	Aa	1107	G	O4'-C1'-N9	6.17	113.14	108.20
84	Aa	1888	G	C5-C6-O6	-6.17	124.89	128.60
86	Ab	20	C	C2-N3-C4	6.17	122.99	119.90
1	Ad	541	G	O4'-C1'-N9	6.17	113.14	108.20
1	Ad	835	U	C4'-C3'-C2'	-6.17	96.43	102.60
84	Aa	7	C	O4'-C1'-N1	6.17	113.14	108.20
84	Aa	39	A	C5-C6-N6	-6.17	118.76	123.70
84	Aa	1359	A	C5-C6-N6	-6.17	118.76	123.70
84	Aa	1400	C	N3-C4-C5	-6.17	119.43	121.90
1	Ad	225	G	C1'-O4'-C4'	-6.17	104.96	109.90
1	Ad	1346	C	N1-C1'-C2'	6.17	122.02	114.00
1	Ad	1392	G	C1'-O4'-C4'	-6.17	104.96	109.90
84	Aa	160	G	C5-C6-O6	-6.17	124.90	128.60
84	Aa	228	C	N3-C4-C5	-6.17	119.43	121.90
84	Aa	448	G	O4'-C1'-N9	6.17	113.14	108.20
84	Aa	585	A	C4-C5-C6	6.17	120.08	117.00
84	Aa	687	C	O4'-C1'-N1	6.17	113.14	108.20
84	Aa	903	G	N1-C6-O6	6.17	123.60	119.90
84	Aa	3302	A	C4-C5-C6	6.17	120.09	117.00
1	Ad	1461	G	O4'-C1'-C2'	6.17	113.15	107.60
84	Aa	2068	G	C5-C6-O6	-6.17	124.90	128.60
84	Aa	2933	C	O4'-C1'-N1	6.17	113.14	108.20
85	Ac	139	C	N3-C4-C5	-6.17	119.43	121.90
84	Aa	424	G	C8-N9-C1'	-6.17	118.98	127.00
84	Aa	1790	A	C4-C5-C6	6.17	120.08	117.00
84	Aa	3089	G	O4'-C1'-N9	6.17	113.14	108.20
84	Aa	3277	C	C5'-C4'-O4'	6.17	116.50	109.10
1	Ad	630	U	O4'-C1'-N1	6.17	113.13	108.20
84	Aa	596	C	N3-C4-C5	-6.17	119.43	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	812	G	N1-C6-O6	6.17	123.60	119.90
84	Aa	840	A	O4'-C1'-N9	6.17	113.13	108.20
84	Aa	1326	C	N3-C4-C5	-6.17	119.43	121.90
84	Aa	2250	A	C4-C5-C6	6.17	120.08	117.00
84	Aa	2482	A	C5-C6-N6	-6.17	118.77	123.70
84	Aa	2659	A	C4-C5-C6	6.17	120.08	117.00
84	Aa	3302	A	C5-C6-N1	-6.17	114.62	117.70
84	Aa	1201	C	N3-C4-N4	6.17	122.31	118.00
1	Ad	1547	G	C3'-C2'-C1'	6.16	106.43	101.50
84	Aa	454	A	C4-C5-C6	6.16	120.08	117.00
84	Aa	1463	A	C5-C6-N1	-6.16	114.62	117.70
84	Aa	1794	A	C5-C6-N1	-6.16	114.62	117.70
84	Aa	1899	U	O4'-C1'-N1	6.16	113.13	108.20
84	Aa	2664	G	C5-C6-O6	-6.16	124.90	128.60
84	Aa	2673	G	O4'-C1'-N9	6.16	113.13	108.20
84	Aa	2737	A	C4-C5-C6	6.16	120.08	117.00
84	Aa	2858	G	C5-C6-O6	-6.16	124.90	128.60
84	Aa	3074	A	C4-C5-C6	6.16	120.08	117.00
1	Ad	292	A	C3'-C2'-C1'	6.16	106.43	101.50
1	Ad	1135	G	O4'-C1'-N9	6.16	113.13	108.20
84	Aa	519	C	N3-C4-C5	-6.16	119.44	121.90
84	Aa	1001	A	O4'-C1'-N9	6.16	113.13	108.20
84	Aa	1984	C	N3-C4-C5	-6.16	119.44	121.90
84	Aa	1988	G	C5-C6-O6	-6.16	124.90	128.60
84	Aa	2092	C	C6-N1-C2	-6.16	117.83	120.30
84	Aa	2708	A	C5-C6-N1	-6.16	114.62	117.70
86	Ab	25	G	N1-C6-O6	6.16	123.60	119.90
1	Ad	1682	U	O4'-C1'-C2'	-6.16	99.64	105.80
1	Ad	1697	G	O4'-C1'-N9	6.16	113.13	108.20
1	Ad	1728	G	OP1-P-OP2	-6.16	110.36	119.60
84	Aa	470	G	O4'-C1'-N9	6.16	113.13	108.20
84	Aa	632	C	N3-C4-C5	-6.16	119.44	121.90
84	Aa	879	A	C4-C5-C6	6.16	120.08	117.00
84	Aa	1732	G	O4'-C1'-N9	6.16	113.13	108.20
84	Aa	2917	U	O4'-C1'-N1	6.16	113.13	108.20
84	Aa	2930	C	N3-C4-C5	-6.16	119.44	121.90
84	Aa	3292	U	O4'-C1'-N1	6.16	113.13	108.20
86	Ab	5	G	N1-C2-N3	-6.16	120.20	123.90
1	Ad	847	U	C1'-O4'-C4'	-6.16	104.97	109.90
1	Ad	848	C	C3'-C2'-C1'	6.16	106.43	101.50
84	Aa	126	G	O4'-C1'-N9	6.16	113.13	108.20
84	Aa	1255	A	C4-C5-C6	6.16	120.08	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1398	A	C5-C6-N6	-6.16	118.77	123.70
84	Aa	1456	A	C5-C6-N1	-6.16	114.62	117.70
84	Aa	3269	C	N3-C4-N4	6.16	122.31	118.00
1	Ad	27	U	C3'-C2'-C1'	6.16	106.43	101.50
84	Aa	531	G	C5-C6-O6	-6.16	124.91	128.60
84	Aa	1012	U	O4'-C1'-N1	6.16	113.12	108.20
84	Aa	3003	C	N3-C4-C5	-6.16	119.44	121.90
85	Ac	73	U	O4'-C1'-N1	6.16	113.13	108.20
1	Ad	47	A	N9-C1'-C2'	6.16	122.00	114.00
1	Ad	79	A	C5'-C4'-O4'	6.16	116.49	109.10
84	Aa	1234	G	O4'-C1'-N9	6.16	113.12	108.20
84	Aa	1699	C	N3-C4-C5	-6.16	119.44	121.90
84	Aa	1805	A	C4-C5-C6	6.16	120.08	117.00
84	Aa	2037	C	N3-C4-N4	6.16	122.31	118.00
84	Aa	2458	A	C4-C5-C6	6.16	120.08	117.00
84	Aa	2852	G	N1-C6-O6	6.16	123.59	119.90
84	Aa	2987	C	N3-C4-C5	-6.16	119.44	121.90
1	Ad	228	G	O4'-C1'-C2'	-6.15	99.65	105.80
84	Aa	104	G	O4'-C1'-N9	6.15	113.12	108.20
84	Aa	56	A	C4-C5-C6	6.15	120.08	117.00
84	Aa	322	A	C5-C6-N6	-6.15	118.78	123.70
84	Aa	1336	A	C5-C6-N1	-6.15	114.62	117.70
84	Aa	1373	A	C4-C5-C6	6.15	120.08	117.00
84	Aa	1690	C	N3-C4-C5	-6.15	119.44	121.90
84	Aa	2013	G	O4'-C1'-N9	6.15	113.12	108.20
84	Aa	2993	A	C5-C6-N1	-6.15	114.62	117.70
1	Ad	17	C	C3'-C2'-C1'	6.15	106.42	101.50
1	Ad	1732	A	O4'-C1'-N9	6.15	113.12	108.20
47	CQ	53	PHE	CB-CG-CD2	-6.15	116.49	120.80
84	Aa	255	C	N3-C4-N4	6.15	122.31	118.00
84	Aa	365	A	C4-C5-C6	6.15	120.08	117.00
84	Aa	382	A	C4-C5-C6	6.15	120.08	117.00
84	Aa	749	C	N3-C4-C5	-6.15	119.44	121.90
84	Aa	933	U	O4'-C1'-N1	6.15	113.12	108.20
84	Aa	1246	G	O4'-C1'-N9	6.15	113.12	108.20
84	Aa	1666	C	N3-C4-C5	-6.15	119.44	121.90
84	Aa	2093	G	N3-C2-N2	6.15	124.20	119.90
84	Aa	660	A	C5-C6-N1	-6.15	114.62	117.70
84	Aa	2154	G	N3-C2-N2	6.15	124.20	119.90
1	Ad	1745	U	O4'-C1'-C2'	-6.15	99.65	105.80
84	Aa	225	G	C5-C6-O6	-6.15	124.91	128.60
84	Aa	981	A	C4-C5-C6	6.15	120.07	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2826	G	C5-C6-O6	-6.15	124.91	128.60
84	Aa	2069	G	O4'-C1'-N9	6.15	113.12	108.20
84	Aa	2331	A	C4-C5-C6	6.15	120.07	117.00
84	Aa	3072	A	C4-C5-C6	6.15	120.07	117.00
1	Ad	700	C	O4'-C1'-C2'	-6.14	99.66	105.80
84	Aa	970	A	C5-C6-N1	-6.14	114.63	117.70
84	Aa	1225	A	C4-C5-C6	6.14	120.07	117.00
84	Aa	1301	C	N3-C4-N4	6.14	122.30	118.00
84	Aa	1843	A	C4-C5-C6	6.14	120.07	117.00
84	Aa	2066	G	O4'-C1'-N9	6.14	113.11	108.20
84	Aa	2662	A	O4'-C1'-N9	6.14	113.12	108.20
84	Aa	2825	G	C5-C6-O6	-6.14	124.91	128.60
84	Aa	3205	C	N3-C4-N4	6.14	122.30	118.00
63	CU	107	ALA	N-CA-CB	6.14	118.70	110.10
84	Aa	162	G	C5-C6-O6	-6.14	124.91	128.60
84	Aa	370	A	O4'-C1'-N9	6.14	113.11	108.20
84	Aa	371	A	C4-C5-C6	6.14	120.07	117.00
84	Aa	1149	C	N3-C4-N4	6.14	122.30	118.00
84	Aa	1812	A	C5-C6-N1	-6.14	114.63	117.70
84	Aa	2549	C	N3-C4-C5	-6.14	119.44	121.90
84	Aa	3051	U	O4'-C1'-N1	6.14	113.11	108.20
86	Ab	14	C	C6-N1-C2	-6.14	117.84	120.30
84	Aa	1240	G	C5-C6-O6	-6.14	124.92	128.60
1	Ad	1165	A	O4'-C1'-N9	6.14	113.11	108.20
84	Aa	669	G	O4'-C1'-N9	6.14	113.11	108.20
84	Aa	790	G	C5-C6-O6	-6.14	124.92	128.60
84	Aa	1815	G	N1-C6-O6	6.14	123.58	119.90
84	Aa	2656	C	N3-C4-C5	-6.14	119.44	121.90
84	Aa	3073	A	C4-C5-C6	6.14	120.07	117.00
63	CU	57	GLY	C-N-CA	6.14	137.04	121.70
84	Aa	94	A	C5-C6-N1	-6.14	114.63	117.70
84	Aa	864	C	O4'-C1'-N1	6.14	113.11	108.20
84	Aa	2561	A	C4-C5-C6	6.14	120.07	117.00
84	Aa	3005	C	C5'-C4'-C3'	-6.14	106.18	116.00
1	Ad	1162	A	O4'-C1'-C2'	-6.14	99.66	105.80
9	BX	60	GLN	CB-CA-C	6.14	122.67	110.40
84	Aa	1040	A	C5-C6-N6	-6.14	118.79	123.70
84	Aa	1293	C	N3-C4-C5	-6.14	119.44	121.90
84	Aa	1401	C	N3-C4-C5	-6.14	119.44	121.90
84	Aa	1456	A	C5-C6-N6	-6.14	118.79	123.70
84	Aa	1532	A	C4-C5-C6	6.14	120.07	117.00
84	Aa	1965	C	N3-C4-N4	6.14	122.30	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2098	A	O4'-C1'-N9	6.14	113.11	108.20
1	Ad	822	G	C5'-C4'-O4'	6.13	116.46	109.10
1	Ad	932	C	N1-C1'-C2'	6.13	121.98	114.00
1	Ad	1342	C	N1-C1'-C2'	6.13	121.98	114.00
84	Aa	65	A	C4-C5-C6	6.13	120.07	117.00
84	Aa	481	G	O4'-C1'-N9	6.13	113.11	108.20
84	Aa	657	A	C5-C6-N1	-6.13	114.63	117.70
84	Aa	1543	A	C4-C5-C6	6.13	120.07	117.00
84	Aa	1807	C	N3-C4-C5	-6.13	119.45	121.90
84	Aa	1828	C	N3-C4-C5	-6.13	119.45	121.90
84	Aa	2366	A	C5-C6-N1	-6.13	114.63	117.70
84	Aa	2647	C	N3-C4-C5	-6.13	119.45	121.90
84	Aa	2886	C	N3-C4-C5	-6.13	119.45	121.90
84	Aa	115	C	P-O3'-C3'	6.13	127.06	119.70
84	Aa	660	A	O4'-C1'-N9	6.13	113.11	108.20
84	Aa	711	A	C5-C6-N1	-6.13	114.63	117.70
84	Aa	1328	C	N3-C4-C5	-6.13	119.45	121.90
84	Aa	2357	A	C4-C5-C6	6.13	120.07	117.00
84	Aa	2835	A	O4'-C1'-N9	6.13	113.11	108.20
84	Aa	3047	A	C5-C6-N1	-6.13	114.63	117.70
84	Aa	3109	G	C5-C6-O6	-6.13	124.92	128.60
1	Ad	291	G	C3'-C2'-C1'	-6.13	96.59	101.50
84	Aa	376	A	C4-C5-C6	6.13	120.07	117.00
84	Aa	802	G	C5-C6-O6	-6.13	124.92	128.60
84	Aa	1075	G	O4'-C1'-N9	6.13	113.11	108.20
84	Aa	1831	A	P-O3'-C3'	6.13	127.06	119.70
84	Aa	2092	C	C2-N1-C1'	-6.13	112.05	118.80
84	Aa	2580	C	N3-C4-C5	-6.13	119.45	121.90
84	Aa	2739	A	O4'-C1'-N9	6.13	113.11	108.20
84	Aa	2904	A	O4'-C1'-N9	6.13	113.11	108.20
1	Ad	146	A	O4'-C1'-C2'	-6.13	99.67	105.80
84	Aa	158	A	C4-C5-C6	6.13	120.06	117.00
84	Aa	170	C	O4'-C1'-N1	6.13	113.10	108.20
84	Aa	366	G	C5-C6-O6	-6.13	124.92	128.60
84	Aa	543	C	O5'-C5'-C4'	-6.13	100.05	111.70
84	Aa	1607	C	N3-C4-C5	-6.13	119.45	121.90
84	Aa	1798	C	C1'-O4'-C4'	-6.13	105.00	109.90
84	Aa	3095	G	O4'-C1'-N9	6.13	113.10	108.20
84	Aa	3154	G	C5-C6-O6	-6.13	124.92	128.60
1	Ad	185	G	O4'-C1'-N9	6.13	113.10	108.20
1	Ad	641	C	O4'-C1'-N1	6.13	113.10	108.20
84	Aa	305	G	O4'-C1'-N9	6.13	113.10	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1670	G	C5-C6-O6	-6.13	124.92	128.60
84	Aa	1744	C	C5'-C4'-O4'	6.13	116.45	109.10
84	Aa	2167	G	P-O3'-C3'	6.13	127.05	119.70
84	Aa	2397	A	C5-C6-N1	-6.13	114.64	117.70
84	Aa	2413	G	O4'-C1'-N9	6.13	113.10	108.20
84	Aa	3368	A	O4'-C1'-N9	6.13	113.10	108.20
84	Aa	376	A	O4'-C1'-N9	6.13	113.10	108.20
84	Aa	383	A	C4-C5-C6	6.13	120.06	117.00
84	Aa	717	G	C5-C6-O6	-6.13	124.92	128.60
84	Aa	1538	A	C4-C5-C6	6.13	120.06	117.00
84	Aa	2878	C	N3-C4-C5	-6.13	119.45	121.90
84	Aa	1026	A	O4'-C1'-N9	6.12	113.10	108.20
84	Aa	2442	A	O4'-C1'-N9	6.12	113.10	108.20
84	Aa	280	G	N1-C6-O6	6.12	123.57	119.90
84	Aa	932	A	O4'-C1'-N9	6.12	113.10	108.20
84	Aa	1123	A	C4-C5-C6	6.12	120.06	117.00
84	Aa	1367	A	C4-C5-C6	6.12	120.06	117.00
84	Aa	1413	C	N3-C4-C5	-6.12	119.45	121.90
84	Aa	2232	C	N3-C4-C5	-6.12	119.45	121.90
85	Ac	25	G	C5-C6-O6	-6.12	124.92	128.60
85	Ac	48	A	C5-C6-N1	-6.12	114.64	117.70
86	Ab	38	U	N3-C4-O4	6.12	123.69	119.40
1	Ad	152	G	C3'-C2'-C1'	-6.12	96.60	101.50
84	Aa	470	G	C5-C6-O6	-6.12	124.93	128.60
84	Aa	588	G	O4'-C1'-N9	6.12	113.10	108.20
84	Aa	842	C	N3-C4-N4	6.12	122.28	118.00
84	Aa	987	A	N1-C6-N6	6.12	122.27	118.60
84	Aa	1133	A	C4-C5-C6	6.12	120.06	117.00
84	Aa	1969	G	P-O3'-C3'	6.12	127.05	119.70
85	Ac	64	U	O4'-C1'-N1	6.12	113.10	108.20
1	Ad	96	G	O4'-C1'-C2'	6.12	113.11	107.60
84	Aa	252	A	C4-C5-C6	6.12	120.06	117.00
84	Aa	920	A	C4-C5-C6	6.12	120.06	117.00
84	Aa	2089	A	C4-C5-C6	6.12	120.06	117.00
84	Aa	2948	A	C4-C5-C6	6.12	120.06	117.00
84	Aa	3016	C	N3-C4-C5	-6.12	119.45	121.90
1	Ad	791	C	N1-C1'-C2'	6.12	121.95	114.00
1	Ad	1424	G	O4'-C1'-N9	6.12	113.10	108.20
56	Cd	33	PHE	CB-CG-CD1	6.12	125.08	120.80
84	Aa	711	A	C4-C5-C6	6.12	120.06	117.00
84	Aa	1161	G	O4'-C1'-N9	6.12	113.09	108.20
84	Aa	1475	U	O4'-C1'-N1	6.12	113.09	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1980	C	N3-C4-C5	-6.12	119.45	121.90
84	Aa	2744	C	N3-C4-N4	6.12	122.28	118.00
84	Aa	369	G	O4'-C1'-N9	6.12	113.09	108.20
84	Aa	665	G	O4'-C1'-N9	6.12	113.09	108.20
84	Aa	1490	A	C4-C5-C6	6.12	120.06	117.00
84	Aa	2084	G	O3'-P-O5'	-6.12	92.38	104.00
84	Aa	2843	G	C5-C6-O6	-6.12	124.93	128.60
85	Ac	54	A	C5-C6-N6	-6.12	118.81	123.70
1	Ad	296	A	O4'-C1'-N9	-6.12	103.31	108.20
1	Ad	298	C	C1'-O4'-C4'	-6.12	105.01	109.90
1	Ad	1555	A	O4'-C1'-N9	6.12	113.09	108.20
84	Aa	1004	C	O4'-C1'-N1	6.12	113.09	108.20
84	Aa	1711	G	C5-C6-O6	-6.12	124.93	128.60
84	Aa	1738	A	O4'-C1'-N9	6.12	113.09	108.20
84	Aa	1967	C	N3-C4-C5	-6.12	119.45	121.90
84	Aa	2367	A	C5-C6-N1	-6.12	114.64	117.70
84	Aa	2397	A	C4-C5-C6	6.12	120.06	117.00
84	Aa	2514	A	O4'-C1'-N9	6.12	113.09	108.20
84	Aa	2618	G	C5-C6-O6	-6.12	124.93	128.60
84	Aa	3277	C	N3-C4-C5	-6.12	119.45	121.90
86	Ab	105	C	C2-N3-C4	6.12	122.96	119.90
1	Ad	425	A	O4'-C1'-N9	6.11	113.09	108.20
1	Ad	967	C	C1'-O4'-C4'	-6.11	105.01	109.90
14	BQ	102	TYR	CB-CG-CD2	-6.11	117.33	121.00
25	Bd	47	ALA	N-CA-CB	6.11	118.66	110.10
84	Aa	1389	C	N3-C4-C5	-6.11	119.45	121.90
84	Aa	1793	A	C4-C5-C6	6.11	120.06	117.00
84	Aa	1888	G	O4'-C1'-N9	6.11	113.09	108.20
84	Aa	2479	C	O5'-C5'-C4'	-6.11	100.08	111.70
84	Aa	2750	A	C4-C5-C6	6.11	120.06	117.00
84	Aa	2889	A	C5-C6-N6	-6.11	118.81	123.70
84	Aa	3236	A	C4-C5-C6	6.11	120.06	117.00
2	Ae	17	G	C3'-C2'-C1'	6.11	106.39	101.50
84	Aa	602	G	C5-C6-O6	-6.11	124.93	128.60
84	Aa	2910	C	N3-C4-C5	-6.11	119.45	121.90
1	Ad	1120	U	N1-C1'-C2'	6.11	121.94	114.00
1	Ad	1576	C	C5'-C4'-C3'	-6.11	106.22	116.00
84	Aa	642	C	O4'-C1'-N1	6.11	113.09	108.20
84	Aa	1061	A	C5-C6-N6	-6.11	118.81	123.70
84	Aa	2366	A	C4-C5-C6	6.11	120.06	117.00
84	Aa	2371	A	C4-C5-C6	6.11	120.06	117.00
84	Aa	2512	U	P-O5'-C5'	-6.11	111.12	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3150	G	O4'-C1'-N9	6.11	113.09	108.20
84	Aa	3213	A	C5-C6-N6	-6.11	118.81	123.70
84	Aa	3270	C	C2-N1-C1'	6.11	125.52	118.80
85	Ac	35	C	N3-C4-C5	-6.11	119.46	121.90
85	Ac	117	C	N3-C4-C5	-6.11	119.45	121.90
84	Aa	1333	C	N3-C4-N4	6.11	122.28	118.00
84	Aa	1746	G	P-O5'-C5'	-6.11	111.13	120.90
84	Aa	2271	G	C5-C6-O6	-6.11	124.94	128.60
84	Aa	3158	C	N3-C4-C5	-6.11	119.46	121.90
51	CX	140	TYR	CB-CG-CD1	-6.11	117.33	121.00
84	Aa	498	G	P-O3'-C3'	6.11	127.03	119.70
84	Aa	1305	A	C4-C5-C6	6.11	120.05	117.00
84	Aa	1316	C	N3-C4-C5	-6.11	119.46	121.90
84	Aa	2432	U	O4'-C1'-N1	6.11	113.08	108.20
84	Aa	2482	A	C4-C5-C6	6.11	120.05	117.00
84	Aa	3342	C	N3-C4-C5	-6.11	119.46	121.90
46	Ca	52	TYR	CB-CG-CD1	6.11	124.66	121.00
84	Aa	498	G	O4'-C1'-N9	6.11	113.08	108.20
84	Aa	573	A	C4-C5-C6	6.11	120.05	117.00
84	Aa	2730	A	C5-C6-N1	-6.11	114.65	117.70
84	Aa	3071	A	C4-C5-C6	6.11	120.05	117.00
84	Aa	3231	G	C5-C6-O6	-6.11	124.94	128.60
84	Aa	3338	U	O4'-C1'-N1	6.11	113.08	108.20
84	Aa	1704	A	C4-C5-C6	6.10	120.05	117.00
84	Aa	2755	U	O4'-C1'-N1	6.10	113.08	108.20
84	Aa	1438	A	C5-C6-N6	-6.10	118.82	123.70
86	Ab	29	C	N3-C4-C5	-6.10	119.46	121.90
84	Aa	1500	C	N3-C4-C5	-6.10	119.46	121.90
84	Aa	3349	C	N3-C4-C5	-6.10	119.46	121.90
84	Aa	722	C	O4'-C1'-N1	6.10	113.08	108.20
84	Aa	725	G	C5-C6-O6	-6.10	124.94	128.60
84	Aa	1194	C	N3-C4-N4	6.10	122.27	118.00
84	Aa	2238	A	C5-C6-N1	-6.10	114.65	117.70
84	Aa	2962	C	N3-C4-C5	-6.10	119.46	121.90
84	Aa	3156	G	O4'-C1'-N9	6.10	113.08	108.20
1	Ad	609	A	O4'-C1'-N9	6.10	113.08	108.20
19	BL	116	PHE	CB-CG-CD2	-6.10	116.53	120.80
84	Aa	293	A	C4-C5-C6	6.10	120.05	117.00
84	Aa	332	A	C5-C6-N6	-6.10	118.82	123.70
84	Aa	443	G	O4'-C1'-N9	6.10	113.08	108.20
84	Aa	2010	G	C5-C6-O6	-6.10	124.94	128.60
84	Aa	2101	A	C4-C5-C6	6.10	120.05	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2400	A	C4-C5-C6	6.10	120.05	117.00
84	Aa	2484	G	O4'-C1'-N9	6.10	113.08	108.20
84	Aa	2803	A	C5-C6-N1	-6.10	114.65	117.70
84	Aa	2806	A	C5-C6-N1	-6.10	114.65	117.70
84	Aa	3280	U	O4'-C1'-N1	6.10	113.08	108.20
1	Ad	621	U	N1-C1'-C2'	6.10	121.92	114.00
1	Ad	1652	C	C1'-O4'-C4'	-6.10	105.02	109.90
84	Aa	20	G	C5-C6-O6	-6.10	124.94	128.60
84	Aa	2023	C	N3-C4-C5	-6.10	119.46	121.90
84	Aa	2786	G	C5-C6-O6	-6.10	124.94	128.60
1	Ad	838	U	O4'-C1'-N1	6.09	113.08	108.20
63	CU	106	ALA	N-CA-CB	6.09	118.63	110.10
84	Aa	628	C	N3-C4-N4	6.09	122.27	118.00
84	Aa	1435	C	N3-C4-C5	-6.09	119.46	121.90
84	Aa	1535	C	N3-C4-C5	-6.09	119.46	121.90
84	Aa	1663	G	O4'-C1'-N9	6.09	113.08	108.20
84	Aa	2879	G	O4'-C1'-N9	6.09	113.08	108.20
1	Ad	581	G	O4'-C1'-N9	6.09	113.07	108.20
84	Aa	534	G	C5-C6-O6	-6.09	124.94	128.60
84	Aa	3088	A	C4-C5-C6	6.09	120.05	117.00
84	Aa	3092	A	C5-C6-N1	-6.09	114.65	117.70
1	Ad	605	A	O4'-C1'-N9	6.09	113.07	108.20
1	Ad	1513	A	N9-C1'-C2'	-6.09	105.30	112.00
84	Aa	54	G	O4'-C1'-N9	6.09	113.07	108.20
84	Aa	493	G	O4'-C1'-N9	6.09	113.07	108.20
84	Aa	1846	A	C4-C5-C6	6.09	120.05	117.00
84	Aa	2165	A	C5-C6-N6	-6.09	118.83	123.70
84	Aa	2172	C	N3-C4-C5	-6.09	119.46	121.90
84	Aa	2678	C	N3-C4-N4	6.09	122.26	118.00
84	Aa	3041	A	C4-C5-C6	6.09	120.05	117.00
84	Aa	3103	G	C5-C6-O6	-6.09	124.94	128.60
84	Aa	3146	C	N3-C4-C5	-6.09	119.46	121.90
84	Aa	3241	C	N3-C4-C5	-6.09	119.46	121.90
85	Ac	153	C	N3-C4-N4	6.09	122.26	118.00
84	Aa	507	C	N3-C4-C5	-6.09	119.46	121.90
84	Aa	721	A	C4-C5-C6	6.09	120.05	117.00
84	Aa	1424	G	O4'-C1'-N9	6.09	113.07	108.20
84	Aa	1763	C	N3-C4-N4	6.09	122.26	118.00
84	Aa	2081	C	N3-C4-N4	6.09	122.26	118.00
84	Aa	1153	A	C5-C6-N6	-6.09	118.83	123.70
84	Aa	1291	A	C4-C5-C6	6.09	120.04	117.00
85	Ac	54	A	C4-C5-C6	6.09	120.04	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	489	C	O4'-C1'-N1	6.09	113.07	108.20
1	Ad	607	U	O4'-C1'-N1	6.09	113.07	108.20
39	CZ	77	PHE	CB-CG-CD2	-6.09	116.54	120.80
84	Aa	991	C	N3-C4-C5	-6.09	119.47	121.90
84	Aa	1248	A	C5-C6-N1	-6.09	114.66	117.70
84	Aa	1769	C	N3-C4-C5	-6.09	119.47	121.90
84	Aa	1996	C	N3-C4-C5	-6.09	119.47	121.90
84	Aa	2082	A	C4-C5-C6	6.09	120.04	117.00
84	Aa	2622	G	C5-C6-O6	-6.09	124.95	128.60
84	Aa	3371	C	N3-C4-C5	-6.09	119.47	121.90
84	Aa	458	G	O4'-C1'-N9	6.08	113.07	108.20
84	Aa	1295	A	O4'-C1'-N9	6.08	113.07	108.20
84	Aa	2360	A	O4'-C1'-N9	6.08	113.07	108.20
84	Aa	2430	C	N3-C4-C5	-6.08	119.47	121.90
84	Aa	2466	G	O4'-C1'-N9	6.08	113.07	108.20
84	Aa	3166	C	N3-C4-C5	-6.08	119.47	121.90
1	Ad	537	U	N1-C1'-C2'	6.08	121.91	114.00
84	Aa	126	G	C5-C6-O6	-6.08	124.95	128.60
84	Aa	1463	A	C4-C5-C6	6.08	120.04	117.00
84	Aa	1464	A	O4'-C1'-N9	6.08	113.07	108.20
84	Aa	1939	C	N3-C4-C5	-6.08	119.47	121.90
84	Aa	2190	C	N3-C4-N4	6.08	122.26	118.00
84	Aa	3221	A	C5-C6-N6	-6.08	118.83	123.70
84	Aa	3386	A	C4-C5-C6	6.08	120.04	117.00
86	Ab	24	G	C6-C5-N7	-6.08	126.75	130.40
1	Ad	475	A	O4'-C1'-N9	6.08	113.07	108.20
1	Ad	528	U	O4'-C1'-N1	6.08	113.06	108.20
49	CR	55	GLN	N-CA-CB	6.08	121.55	110.60
84	Aa	88	A	C4-C5-C6	6.08	120.04	117.00
84	Aa	279	G	N3-C2-N2	6.08	124.16	119.90
84	Aa	905	G	C5-C6-O6	-6.08	124.95	128.60
84	Aa	2897	G	C5-C6-O6	-6.08	124.95	128.60
84	Aa	2928	A	C5-C6-N1	-6.08	114.66	117.70
84	Aa	3049	A	C4-C5-C6	6.08	120.04	117.00
2	Ae	45	G	C3'-C2'-C1'	6.08	106.36	101.50
79	CE	39	ALA	N-CA-CB	6.08	118.61	110.10
84	Aa	439	A	C4-C5-C6	6.08	120.04	117.00
84	Aa	541	C	N3-C4-N4	6.08	122.26	118.00
84	Aa	2377	C	C1'-O4'-C4'	6.08	114.76	109.90
84	Aa	2773	G	O4'-C1'-N9	6.08	113.06	108.20
1	Ad	1699	C	O4'-C1'-C2'	-6.08	99.72	105.80
1	Ad	1741	A	C5'-C4'-O4'	6.08	116.39	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	33	A	C4-C5-C6	6.08	120.04	117.00
84	Aa	642	C	C6-N1-C2	-6.08	117.87	120.30
84	Aa	725	G	O4'-C1'-N9	6.08	113.06	108.20
84	Aa	904	G	O4'-C1'-N9	6.08	113.06	108.20
84	Aa	1251	U	O4'-C1'-N1	6.08	113.06	108.20
84	Aa	1665	G	C5-C6-O6	-6.08	124.95	128.60
84	Aa	1812	A	O4'-C1'-N9	6.08	113.06	108.20
84	Aa	1975	G	C5-C6-O6	-6.08	124.95	128.60
84	Aa	2266	A	C5-C6-N1	-6.08	114.66	117.70
84	Aa	2276	A	C5-C6-N6	-6.08	118.84	123.70
84	Aa	2380	G	C5-C6-O6	-6.08	124.95	128.60
84	Aa	3329	G	O4'-C1'-N9	6.08	113.06	108.20
85	Ac	145	U	O4'-C1'-N1	6.08	113.06	108.20
85	Ac	155	U	O4'-C1'-N1	6.08	113.06	108.20
86	Ab	18	C	C2-N3-C4	6.08	122.94	119.90
1	Ad	815	A	C4'-C3'-C2'	-6.08	96.52	102.60
84	Aa	55	G	O4'-C1'-N9	6.08	113.06	108.20
84	Aa	110	C	N3-C4-C5	-6.08	119.47	121.90
84	Aa	187	G	O4'-C1'-N9	6.08	113.06	108.20
84	Aa	732	G	C5-C6-O6	-6.08	124.95	128.60
84	Aa	778	G	C5-C6-O6	-6.08	124.95	128.60
84	Aa	1346	C	N3-C4-C5	-6.08	119.47	121.90
84	Aa	1826	G	O4'-C1'-N9	6.08	113.06	108.20
84	Aa	2093	G	C8-N9-C1'	6.08	134.90	127.00
1	Ad	760	G	O4'-C1'-C2'	6.08	113.07	107.60
1	Ad	825	U	P-O5'-C5'	6.08	130.62	120.90
84	Aa	194	G	C5-C6-O6	-6.08	124.95	128.60
84	Aa	1619	G	O4'-C1'-N9	6.08	113.06	108.20
84	Aa	2163	G	N3-C2-N2	6.08	124.15	119.90
84	Aa	2651	G	O4'-C1'-N9	6.08	113.06	108.20
84	Aa	2666	G	C5-C6-O6	-6.08	124.95	128.60
85	Ac	11	C	N3-C4-C5	-6.08	119.47	121.90
84	Aa	394	A	C4-C5-C6	6.07	120.04	117.00
84	Aa	671	C	N3-C4-C5	-6.07	119.47	121.90
84	Aa	827	C	N3-C4-C5	-6.07	119.47	121.90
84	Aa	1658	G	O4'-C1'-N9	6.07	113.06	108.20
84	Aa	2193	A	O4'-C1'-N9	6.07	113.06	108.20
84	Aa	2908	C	N3-C4-C5	-6.07	119.47	121.90
1	Ad	1733	G	N9-C1'-C2'	6.07	121.89	114.00
84	Aa	2161	G	C5-C6-O6	-6.07	124.96	128.60
84	Aa	2920	G	C5'-C4'-C3'	-6.07	106.28	116.00
84	Aa	3123	A	C4-C5-C6	6.07	120.04	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1194	C	C1'-O4'-C4'	6.07	114.76	109.90
1	Ad	1726	G	O4'-C1'-N9	6.07	113.06	108.20
84	Aa	427	U	O4'-C1'-N1	6.07	113.06	108.20
84	Aa	1115	A	O4'-C1'-N9	6.07	113.06	108.20
84	Aa	1175	G	N3-C2-N2	6.07	124.15	119.90
84	Aa	1953	C	N3-C4-C5	-6.07	119.47	121.90
84	Aa	248	C	N3-C4-C5	-6.07	119.47	121.90
84	Aa	1050	A	N1-C6-N6	6.07	122.24	118.60
84	Aa	2521	C	N3-C4-C5	-6.07	119.47	121.90
84	Aa	2950	C	N3-C4-C5	-6.07	119.47	121.90
85	Ac	40	A	O4'-C1'-N9	6.07	113.06	108.20
86	Ab	111	U	O4'-C1'-N1	6.07	113.06	108.20
78	CL	25	PHE	CB-CG-CD1	6.07	125.05	120.80
84	Aa	103	G	C5-C6-O6	-6.07	124.96	128.60
84	Aa	686	A	C4-C5-C6	6.07	120.03	117.00
84	Aa	1590	A	C5-C6-N6	-6.07	118.85	123.70
84	Aa	2778	C	N3-C4-C5	-6.07	119.47	121.90
85	Ac	110	A	O4'-C1'-N9	6.07	113.05	108.20
1	Ad	958	G	O4'-C1'-N9	6.07	113.05	108.20
1	Ad	1225	A	C3'-C2'-C1'	6.07	106.35	101.50
1	Ad	1597	C	C3'-C2'-C1'	6.07	106.35	101.50
84	Aa	205	C	N3-C4-C5	-6.07	119.47	121.90
84	Aa	616	A	C5-C6-N1	-6.07	114.67	117.70
84	Aa	710	C	N3-C4-C5	-6.07	119.47	121.90
84	Aa	1186	C	N3-C4-C5	-6.07	119.47	121.90
84	Aa	1253	G	O4'-C1'-N9	6.07	113.05	108.20
84	Aa	1376	A	C5-C6-N1	-6.07	114.67	117.70
84	Aa	1795	A	C4-C5-C6	6.07	120.03	117.00
84	Aa	3034	A	C4-C5-C6	6.07	120.03	117.00
84	Aa	3251	C	N3-C4-N4	6.07	122.25	118.00
84	Aa	1456	A	C4-C5-C6	6.06	120.03	117.00
84	Aa	1818	C	N3-C4-N4	6.06	122.25	118.00
1	Ad	637	U	O4'-C1'-C2'	-6.06	99.74	105.80
36	BH	17	SER	N-CA-CB	6.06	119.59	110.50
84	Aa	523	C	N3-C4-N4	6.06	122.24	118.00
84	Aa	529	C	N3-C4-N4	6.06	122.24	118.00
84	Aa	1376	A	C4-C5-C6	6.06	120.03	117.00
84	Aa	1388	C	N3-C4-C5	-6.06	119.47	121.90
84	Aa	2624	G	O4'-C1'-N9	6.06	113.05	108.20
84	Aa	2779	G	C5-C6-O6	-6.06	124.96	128.60
84	Aa	3240	C	N3-C4-C5	-6.06	119.47	121.90
84	Aa	3310	A	C4-C5-C6	6.06	120.03	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1728	G	O4'-C1'-N9	6.06	113.05	108.20
84	Aa	1925	G	O4'-C1'-N9	6.06	113.05	108.20
84	Aa	2429	A	C5-C6-N1	-6.06	114.67	117.70
1	Ad	11	A	C3'-C2'-C1'	6.06	106.35	101.50
1	Ad	470	U	N1-C1'-C2'	6.06	121.88	114.00
1	Ad	914	U	P-O5'-C5'	6.06	130.59	120.90
84	Aa	12	G	C2-N3-C4	6.06	114.93	111.90
84	Aa	377	C	N3-C4-N4	6.06	122.24	118.00
84	Aa	595	C	N3-C4-C5	-6.06	119.48	121.90
84	Aa	830	A	C4-C5-C6	6.06	120.03	117.00
84	Aa	879	A	C5-C6-N1	-6.06	114.67	117.70
84	Aa	1040	A	C4-C5-C6	6.06	120.03	117.00
84	Aa	1253	G	C5-C6-O6	-6.06	124.96	128.60
84	Aa	2201	G	O4'-C1'-N9	6.06	113.05	108.20
84	Aa	2578	G	C5-C6-O6	-6.06	124.97	128.60
84	Aa	2838	C	N3-C4-C5	-6.06	119.48	121.90
84	Aa	2866	A	C4-C5-C6	6.06	120.03	117.00
84	Aa	3033	A	C4-C5-C6	6.06	120.03	117.00
84	Aa	3176	C	N3-C4-C5	-6.06	119.48	121.90
85	Ac	126	A	C5-C6-N6	-6.06	118.85	123.70
1	Ad	889	C	O4'-C1'-N1	6.06	113.05	108.20
2	Ae	60	C	N1-C1'-C2'	6.06	121.88	114.00
84	Aa	738	A	C4-C5-C6	6.06	120.03	117.00
84	Aa	1015	A	C5-C6-N1	-6.06	114.67	117.70
84	Aa	1359	A	C4-C5-C6	6.06	120.03	117.00
84	Aa	1364	C	N3-C4-C5	-6.06	119.48	121.90
84	Aa	1751	G	C5-C6-O6	-6.06	124.97	128.60
84	Aa	1752	C	N3-C4-C5	-6.06	119.48	121.90
84	Aa	2364	C	N3-C4-N4	6.06	122.24	118.00
84	Aa	2954	G	C5-C6-O6	-6.06	124.97	128.60
84	Aa	3239	G	O4'-C1'-N9	6.06	113.05	108.20
84	Aa	688	G	N1-C6-O6	6.06	123.53	119.90
84	Aa	820	A	C5-C6-N6	-6.06	118.86	123.70
84	Aa	1337	C	N3-C4-C5	-6.06	119.48	121.90
84	Aa	2097	C	N3-C4-C5	-6.06	119.48	121.90
84	Aa	2753	C	N3-C4-C5	-6.06	119.48	121.90
85	Ac	32	C	N3-C4-C5	-6.06	119.48	121.90
1	Ad	1059	U	N1-C1'-C2'	6.05	121.87	114.00
1	Ad	1700	G	N9-C1'-C2'	6.05	121.87	114.00
84	Aa	528	C	C4'-C3'-O3'	6.05	125.11	113.00
84	Aa	1041	C	N3-C4-C5	-6.05	119.48	121.90
84	Aa	2398	A	C5-C6-N1	-6.05	114.67	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2462	G	C5-C6-O6	-6.05	124.97	128.60
84	Aa	2557	C	C6-N1-C1'	-6.05	113.53	120.80
84	Aa	2751	A	C5-C6-N1	-6.05	114.67	117.70
84	Aa	2813	A	C4-C5-C6	6.05	120.03	117.00
85	Ac	30	C	N3-C4-C5	-6.05	119.48	121.90
2	Ae	17	G	N9-C1'-C2'	6.05	121.87	114.00
84	Aa	563	C	C2'-C3'-O3'	6.05	123.38	113.70
84	Aa	734	C	N3-C4-C5	-6.05	119.48	121.90
84	Aa	953	G	N3-C2-N2	6.05	124.14	119.90
84	Aa	2240	C	N3-C4-N4	6.05	122.24	118.00
84	Aa	3123	A	O4'-C1'-N9	6.05	113.04	108.20
84	Aa	3306	A	C4-C5-C6	6.05	120.03	117.00
1	Ad	257	A	O4'-C1'-N9	-6.05	103.36	108.20
84	Aa	513	C	N3-C4-N4	6.05	122.23	118.00
84	Aa	1819	A	C4-C5-C6	6.05	120.03	117.00
84	Aa	1831	A	C5-C6-N1	-6.05	114.67	117.70
84	Aa	2509	A	O4'-C1'-N9	6.05	113.04	108.20
84	Aa	2805	A	C5-C6-N1	-6.05	114.67	117.70
84	Aa	315	A	C4-C5-C6	6.05	120.02	117.00
84	Aa	1002	A	C5-C6-N6	-6.05	118.86	123.70
84	Aa	2328	C	N3-C4-C5	-6.05	119.48	121.90
85	Ac	12	A	C5-C6-N6	-6.05	118.86	123.70
84	Aa	142	G	C5-C6-O6	-6.05	124.97	128.60
84	Aa	1404	G	O4'-C1'-N9	6.05	113.04	108.20
84	Aa	1484	A	C5-C6-N1	-6.05	114.68	117.70
84	Aa	1492	A	C5-C6-N6	-6.05	118.86	123.70
84	Aa	2038	G	O4'-C1'-N9	6.05	113.04	108.20
84	Aa	2503	A	C5-C6-N1	-6.05	114.68	117.70
84	Aa	2522	C	N3-C4-C5	-6.05	119.48	121.90
84	Aa	2640	A	C5-C6-N6	-6.05	118.86	123.70
84	Aa	3147	G	O4'-C1'-N9	6.05	113.04	108.20
86	Ab	99	G	C5-C6-N1	-6.05	108.48	111.50
84	Aa	1879	A	O4'-C1'-N9	6.04	113.04	108.20
84	Aa	2086	A	O5'-C5'-C4'	-6.04	100.21	111.70
84	Aa	2737	A	C5-C6-N1	-6.04	114.68	117.70
1	Ad	594	C	N1-C1'-C2'	6.04	121.86	114.00
1	Ad	1627	C	C3'-C2'-C1'	6.04	106.33	101.50
84	Aa	231	C	N3-C4-C5	-6.04	119.48	121.90
84	Aa	251	G	C5-C6-O6	-6.04	124.97	128.60
84	Aa	586	A	P-O3'-C3'	6.04	126.95	119.70
84	Aa	1028	G	C5-C6-O6	-6.04	124.97	128.60
84	Aa	1139	A	C5-C6-N1	-6.04	114.68	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1152	G	N1-C6-O6	6.04	123.53	119.90
84	Aa	1832	C	N3-C4-C5	-6.04	119.48	121.90
84	Aa	2180	G	N1-C6-O6	6.04	123.53	119.90
84	Aa	2697	A	O4'-C1'-N9	6.04	113.03	108.20
84	Aa	2840	A	O4'-C1'-N9	6.04	113.03	108.20
85	Ac	77	A	O4'-C1'-N9	6.04	113.03	108.20
85	Ac	118	C	N3-C4-C5	-6.04	119.48	121.90
1	Ad	357	A	O4'-C1'-C2'	-6.04	99.76	105.80
1	Ad	1502	C	C3'-C2'-C1'	6.04	106.33	101.50
47	CQ	123	PHE	CB-CG-CD1	6.04	125.03	120.80
84	Aa	849	A	C5-C6-N1	-6.04	114.68	117.70
84	Aa	854	C	N3-C4-C5	-6.04	119.48	121.90
84	Aa	955	A	C4-C5-C6	6.04	120.02	117.00
84	Aa	1182	A	C5-C6-N1	-6.04	114.68	117.70
84	Aa	1362	C	N3-C4-C5	-6.04	119.48	121.90
84	Aa	1998	A	O4'-C1'-N9	6.04	113.03	108.20
84	Aa	2032	C	N3-C4-C5	-6.04	119.48	121.90
84	Aa	2781	A	C4-C5-C6	6.04	120.02	117.00
84	Aa	3136	A	O4'-C1'-N9	6.04	113.03	108.20
1	Ad	1127	G	C1'-O4'-C4'	6.04	114.73	109.90
2	Ae	17	G	C5'-C4'-O4'	6.04	116.35	109.10
84	Aa	57	G	O4'-C1'-N9	6.04	113.03	108.20
84	Aa	187	G	C5-C6-O6	-6.04	124.98	128.60
84	Aa	2389	A	C5-C6-N1	-6.04	114.68	117.70
84	Aa	2754	G	C5-C6-O6	-6.04	124.98	128.60
85	Ac	81	U	O4'-C1'-N1	6.04	113.03	108.20
1	Ad	1803	G	O4'-C1'-N9	-6.04	103.37	108.20
14	BQ	149	ARG	N-CA-CB	6.04	121.47	110.60
84	Aa	534	G	O4'-C1'-N9	6.04	113.03	108.20
84	Aa	2019	G	O4'-C1'-N9	6.04	113.03	108.20
84	Aa	2251	A	C4-C5-C6	6.04	120.02	117.00
1	Ad	1210	U	C1'-O4'-C4'	6.04	114.73	109.90
1	Ad	1736	C	C3'-C2'-C1'	6.04	106.33	101.50
41	CA	76	PHE	CB-CG-CD2	-6.04	116.57	120.80
84	Aa	990	U	O4'-C1'-N1	6.04	113.03	108.20
84	Aa	1468	A	O4'-C1'-N9	6.04	113.03	108.20
84	Aa	2670	A	C4-C5-C6	6.04	120.02	117.00
84	Aa	3310	A	O4'-C1'-N9	6.04	113.03	108.20
84	Aa	33	A	C5-C6-N6	-6.04	118.87	123.70
84	Aa	471	C	N3-C4-C5	-6.04	119.49	121.90
84	Aa	2396	A	C4-C5-C6	6.04	120.02	117.00
84	Aa	2424	G	O4'-C1'-N9	6.04	113.03	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2695	A	C5-C6-N6	-6.04	118.87	123.70
84	Aa	2952	G	C5-C6-O6	-6.04	124.98	128.60
84	Aa	3265	C	N3-C4-N4	6.04	122.22	118.00
84	Aa	454	A	C5-C6-N6	-6.03	118.87	123.70
84	Aa	1340	G	C5-C6-O6	-6.03	124.98	128.60
84	Aa	1422	G	C5-C6-O6	-6.03	124.98	128.60
84	Aa	2562	A	C5-C6-N6	-6.03	118.87	123.70
1	Ad	1493	A	O4'-C1'-N9	6.03	113.03	108.20
84	Aa	544	C	N3-C4-C5	-6.03	119.49	121.90
84	Aa	2393	G	C5-C6-O6	-6.03	124.98	128.60
84	Aa	2806	A	C4-C5-C6	6.03	120.02	117.00
86	Ab	119	C	N3-C4-N4	6.03	122.22	118.00
1	Ad	455	G	O4'-C1'-N9	6.03	113.02	108.20
1	Ad	795	A	C3'-C2'-C1'	6.03	106.33	101.50
84	Aa	27	C	N3-C4-C5	-6.03	119.49	121.90
84	Aa	359	A	C4-C5-C6	6.03	120.02	117.00
84	Aa	672	A	C5-C6-N1	-6.03	114.69	117.70
84	Aa	1298	A	O4'-C1'-N9	6.03	113.03	108.20
84	Aa	1656	C	N3-C4-C5	-6.03	119.49	121.90
84	Aa	2285	C	N3-C4-C5	-6.03	119.49	121.90
84	Aa	2342	C	N3-C4-C5	-6.03	119.49	121.90
84	Aa	3383	C	N3-C4-N4	6.03	122.22	118.00
85	Ac	114	G	C5-C6-O6	-6.03	124.98	128.60
84	Aa	343	G	C5-C6-O6	-6.03	124.98	128.60
84	Aa	831	G	C5-C6-O6	-6.03	124.98	128.60
1	Ad	378	U	O4'-C1'-N1	6.03	113.02	108.20
1	Ad	1312	G	C1'-O4'-C4'	-6.03	105.08	109.90
84	Aa	444	C	N3-C4-C5	-6.03	119.49	121.90
84	Aa	579	G	N3-C2-N2	6.03	124.12	119.90
84	Aa	586	A	C5-C6-N6	-6.03	118.88	123.70
84	Aa	1221	A	C4-C5-C6	6.03	120.01	117.00
84	Aa	2696	C	N3-C4-C5	-6.03	119.49	121.90
86	Ab	70	G	C8-N9-C4	-6.03	103.99	106.40
1	Ad	1124	G	O4'-C1'-N9	6.03	113.02	108.20
84	Aa	89	C	N3-C4-C5	-6.03	119.49	121.90
84	Aa	179	G	C5-C6-O6	-6.03	124.98	128.60
84	Aa	312	U	O4'-C1'-N1	6.03	113.02	108.20
84	Aa	798	G	O4'-C1'-N9	6.03	113.02	108.20
84	Aa	1767	G	C5-C6-O6	-6.03	124.98	128.60
84	Aa	2884	U	O4'-C1'-N1	6.03	113.02	108.20
84	Aa	3114	A	C4-C5-C6	6.03	120.01	117.00
85	Ac	14	C	O4'-C1'-N1	6.03	113.02	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	Ab	58	G	O4'-C1'-N9	6.03	113.02	108.20
1	Ad	771	G	O4'-C1'-N9	6.02	113.02	108.20
1	Ad	1077	C	C3'-C2'-C1'	6.02	106.32	101.50
84	Aa	1727	A	C4-C5-C6	6.02	120.01	117.00
84	Aa	1873	C	N3-C4-C5	-6.02	119.49	121.90
86	Ab	120	C	N3-C4-N4	6.02	122.22	118.00
1	Ad	196	G	O4'-C1'-N9	6.02	113.02	108.20
1	Ad	792	U	C1'-O4'-C4'	6.02	114.72	109.90
9	BX	40	PHE	CB-CG-CD1	6.02	125.02	120.80
84	Aa	305	G	C5-C6-O6	-6.02	124.99	128.60
84	Aa	332	A	C5-C6-N1	-6.02	114.69	117.70
84	Aa	1946	C	N3-C4-C5	-6.02	119.49	121.90
84	Aa	2436	G	P-O3'-C3'	6.02	126.93	119.70
84	Aa	2905	A	C5-C6-N1	-6.02	114.69	117.70
85	Ac	50	C	N3-C4-C5	-6.02	119.49	121.90
85	Ac	131	G	O4'-C1'-N9	6.02	113.02	108.20
84	Aa	1777	C	N3-C4-C5	-6.02	119.49	121.90
84	Aa	2441	G	C5-C6-O6	-6.02	124.99	128.60
84	Aa	690	G	C5-C6-O6	-6.02	124.99	128.60
84	Aa	975	G	O4'-C1'-N9	6.02	113.02	108.20
84	Aa	1144	C	N3-C4-C5	-6.02	119.49	121.90
84	Aa	1507	A	C4-C5-C6	6.02	120.01	117.00
84	Aa	2514	A	C4-C5-C6	6.02	120.01	117.00
84	Aa	2902	A	C5-C6-N1	-6.02	114.69	117.70
84	Aa	3123	A	C5-C6-N1	-6.02	114.69	117.70
84	Aa	3224	C	N3-C4-C5	-6.02	119.49	121.90
1	Ad	853	U	O4'-C1'-N1	6.02	113.01	108.20
2	Ae	69	G	N9-C1'-C2'	6.02	121.82	114.00
17	BS	98	VAL	N-CA-C	-6.02	94.75	111.00
84	Aa	168	A	C4-C5-C6	6.02	120.01	117.00
84	Aa	1635	A	C4-C5-C6	6.02	120.01	117.00
84	Aa	2100	A	C4-C5-C6	6.02	120.01	117.00
84	Aa	2788	A	C5-C6-N6	-6.02	118.89	123.70
85	Ac	87	G	O4'-C1'-N9	6.02	113.01	108.20
1	Ad	533	C	P-O5'-C5'	-6.02	111.28	120.90
84	Aa	515	C	N3-C4-C5	-6.02	119.49	121.90
1	Ad	619	A	N9-C1'-C2'	-6.01	105.38	112.00
1	Ad	1054	G	O4'-C1'-N9	6.01	113.01	108.20
84	Aa	920	A	O4'-C1'-N9	6.01	113.01	108.20
84	Aa	1155	G	O4'-C1'-N9	6.01	113.01	108.20
84	Aa	2275	A	C4-C5-C6	6.01	120.01	117.00
84	Aa	2380	G	O4'-C1'-N9	6.01	113.01	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	393	G	N9-C1'-C2'	6.01	121.82	114.00
1	Ad	1520	G	C1'-O4'-C4'	-6.01	105.09	109.90
84	Aa	875	A	C4-C5-C6	6.01	120.01	117.00
84	Aa	876	C	N3-C4-C5	-6.01	119.50	121.90
84	Aa	1941	G	C5-C6-O6	-6.01	124.99	128.60
1	Ad	1181	G	O4'-C1'-N9	6.01	113.01	108.20
1	Ad	1778	G	O4'-C1'-N9	-6.01	103.39	108.20
84	Aa	697	A	C5-C6-N6	-6.01	118.89	123.70
84	Aa	1221	A	C5-C6-N6	-6.01	118.89	123.70
84	Aa	1470	A	C5-C6-N6	-6.01	118.89	123.70
84	Aa	1486	G	N3-C2-N2	6.01	124.11	119.90
84	Aa	1596	G	C5-C6-O6	-6.01	124.99	128.60
84	Aa	1954	G	C5-C6-O6	-6.01	124.99	128.60
84	Aa	2118	G	C5-C6-O6	-6.01	124.99	128.60
84	Aa	2175	A	C4-C5-C6	6.01	120.00	117.00
84	Aa	2222	C	N3-C4-C5	-6.01	119.50	121.90
84	Aa	2276	A	O4'-C1'-N9	6.01	113.01	108.20
84	Aa	2351	A	C5-C6-N6	-6.01	118.89	123.70
84	Aa	2381	G	C5-C6-O6	-6.01	124.99	128.60
84	Aa	2805	A	C4-C5-C6	6.01	120.01	117.00
84	Aa	3275	G	C5-C6-O6	-6.01	124.99	128.60
85	Ac	91	C	N3-C4-C5	-6.01	119.50	121.90
1	Ad	1331	C	O4'-C1'-N1	6.01	113.01	108.20
1	Ad	1526	C	O4'-C1'-N1	6.01	113.01	108.20
3	Af	16	G	N9-C1'-C2'	-6.01	105.39	112.00
84	Aa	361	G	O4'-C1'-N9	6.01	113.01	108.20
84	Aa	639	A	C5-C6-N1	-6.01	114.69	117.70
84	Aa	650	A	C5-C6-N1	-6.01	114.69	117.70
84	Aa	1552	C	N3-C4-C5	-6.01	119.50	121.90
84	Aa	1723	C	N3-C4-C5	-6.01	119.50	121.90
84	Aa	1944	G	C5'-C4'-C3'	6.01	125.61	116.00
84	Aa	2010	G	O4'-C1'-N9	6.01	113.01	108.20
84	Aa	2638	A	C4-C5-C6	6.01	120.00	117.00
84	Aa	2863	U	O4'-C1'-N1	6.01	113.01	108.20
84	Aa	3009	A	C5-C6-N1	-6.01	114.69	117.70
84	Aa	3077	C	N3-C4-C5	-6.01	119.50	121.90
85	Ac	115	C	N3-C4-C5	-6.01	119.50	121.90
1	Ad	451	U	O4'-C1'-N1	6.01	113.01	108.20
84	Aa	1274	A	C5-C6-N1	-6.01	114.70	117.70
84	Aa	1894	G	O4'-C1'-N9	6.01	113.01	108.20
84	Aa	2027	G	C5-C6-O6	-6.01	125.00	128.60
84	Aa	2142	A	C4-C5-C6	6.01	120.00	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2276	A	C4-C5-C6	6.01	120.00	117.00
84	Aa	374	G	C5-C6-O6	-6.01	125.00	128.60
84	Aa	482	C	N3-C4-N4	6.01	122.20	118.00
84	Aa	570	G	C5-C6-O6	-6.01	125.00	128.60
84	Aa	1629	A	C5-C6-N1	-6.01	114.70	117.70
84	Aa	2270	A	O4'-C1'-N9	6.01	113.00	108.20
1	Ad	164	C	C3'-C2'-C1'	6.00	106.30	101.50
84	Aa	961	C	N3-C4-C5	-6.00	119.50	121.90
84	Aa	1749	G	O3'-P-O5'	-6.00	92.59	104.00
84	Aa	2149	G	C4-N9-C1'	6.00	134.31	126.50
84	Aa	2587	G	N1-C2-N3	-6.00	120.30	123.90
84	Aa	3218	C	N3-C4-C5	-6.00	119.50	121.90
1	Ad	367	G	C3'-C2'-C1'	-6.00	96.70	101.50
84	Aa	432	G	O4'-C1'-N9	6.00	113.00	108.20
84	Aa	903	G	O4'-C1'-N9	6.00	113.00	108.20
84	Aa	1296	C	N3-C4-C5	-6.00	119.50	121.90
84	Aa	1609	G	C5-C6-O6	-6.00	125.00	128.60
84	Aa	1812	A	C4-C5-C6	6.00	120.00	117.00
84	Aa	2369	G	N1-C6-O6	6.00	123.50	119.90
84	Aa	2579	G	C5-C6-O6	-6.00	125.00	128.60
84	Aa	2835	A	C4-C5-C6	6.00	120.00	117.00
85	Ac	51	G	O4'-C1'-N9	6.00	113.00	108.20
1	Ad	193	G	N9-C1'-C2'	6.00	121.80	114.00
84	Aa	93	G	C5-C6-O6	-6.00	125.00	128.60
84	Aa	670	A	C5-C6-N1	-6.00	114.70	117.70
84	Aa	2412	A	C4-C5-C6	6.00	120.00	117.00
84	Aa	2429	A	C4-C5-C6	6.00	120.00	117.00
84	Aa	474	G	O5'-C5'-C4'	6.00	123.10	111.70
17	BS	100	SER	N-CA-CB	6.00	119.50	110.50
84	Aa	675	C	N3-C4-C5	-6.00	119.50	121.90
84	Aa	872	G	O4'-C1'-N9	6.00	113.00	108.20
84	Aa	2039	G	C5-C6-O6	-6.00	125.00	128.60
84	Aa	2533	A	C4-C5-C6	6.00	120.00	117.00
84	Aa	2750	A	C5-C6-N1	-6.00	114.70	117.70
84	Aa	2822	A	C4-C5-C6	6.00	120.00	117.00
84	Aa	3186	G	C5-C6-O6	-6.00	125.00	128.60
1	Ad	86	A	N9-C1'-C2'	6.00	121.80	114.00
1	Ad	782	G	N9-C1'-C2'	6.00	121.80	114.00
1	Ad	1475	A	O4'-C1'-C2'	-6.00	99.80	105.80
84	Aa	1438	A	C4-C5-C6	6.00	120.00	117.00
84	Aa	2019	G	C5-C6-O6	-6.00	125.00	128.60
84	Aa	2100	A	C5-C6-N1	-6.00	114.70	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2139	A	C5-C6-N1	-6.00	114.70	117.70
84	Aa	2706	A	C5-C6-N1	-6.00	114.70	117.70
84	Aa	2790	C	N3-C4-N4	6.00	122.20	118.00
84	Aa	3372	C	N3-C4-C5	-6.00	119.50	121.90
84	Aa	916	A	C4-C5-C6	6.00	120.00	117.00
84	Aa	2643	A	O4'-C1'-N9	6.00	113.00	108.20
84	Aa	3335	G	N3-C2-N2	6.00	124.10	119.90
1	Ad	318	C	C5'-C4'-O4'	5.99	116.29	109.10
84	Aa	156	A	C4-C5-C6	5.99	120.00	117.00
84	Aa	411	C	O4'-C1'-N1	5.99	113.00	108.20
84	Aa	511	C	N3-C4-C5	-5.99	119.50	121.90
84	Aa	1317	G	O4'-C1'-N9	5.99	113.00	108.20
84	Aa	1749	G	C5-C6-O6	-5.99	125.00	128.60
84	Aa	1877	G	C5-C6-O6	-5.99	125.00	128.60
84	Aa	1960	C	N3-C4-C5	-5.99	119.50	121.90
84	Aa	2274	A	C4-C5-C6	5.99	120.00	117.00
84	Aa	2632	U	O4'-C1'-N1	5.99	112.99	108.20
84	Aa	3131	A	C4-C5-C6	5.99	120.00	117.00
1	Ad	1016	C	C3'-C2'-C1'	5.99	106.29	101.50
84	Aa	206	C	N3-C4-C5	-5.99	119.50	121.90
84	Aa	810	A	C4-C5-C6	5.99	120.00	117.00
84	Aa	1659	G	C5-C6-O6	-5.99	125.00	128.60
86	Ab	105	C	N3-C4-N4	5.99	122.19	118.00
1	Ad	569	C	O4'-C1'-C2'	-5.99	99.81	105.80
84	Aa	81	C	N3-C4-C5	-5.99	119.50	121.90
84	Aa	119	A	C4-C5-C6	5.99	120.00	117.00
84	Aa	1846	A	O4'-C1'-N9	5.99	112.99	108.20
84	Aa	2079	A	O4'-C1'-N9	5.99	112.99	108.20
84	Aa	2338	C	N3-C4-N4	5.99	122.19	118.00
84	Aa	3350	C	N3-C4-C5	-5.99	119.50	121.90
36	BH	117	ARG	N-CA-CB	5.99	121.38	110.60
63	CU	100	ASP	N-CA-CB	5.99	121.38	110.60
84	Aa	69	U	O4'-C1'-N1	5.99	112.99	108.20
84	Aa	420	A	O4'-C1'-N9	5.99	112.99	108.20
84	Aa	800	C	C2-N3-C4	5.99	122.89	119.90
84	Aa	900	C	N3-C4-C5	-5.99	119.50	121.90
84	Aa	1210	G	C5-C6-O6	-5.99	125.01	128.60
84	Aa	1825	G	C5-C6-O6	-5.99	125.01	128.60
84	Aa	2152	A	C5-C6-N6	-5.99	118.91	123.70
84	Aa	2455	A	C5-C6-N1	-5.99	114.71	117.70
84	Aa	2773	G	C5-C6-O6	-5.99	125.01	128.60
84	Aa	3157	C	N3-C4-C5	-5.99	119.50	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	113	A	C5-C6-N6	-5.99	118.91	123.70
84	Aa	696	A	C4-C5-C6	5.99	119.99	117.00
84	Aa	887	A	C5-C6-N6	-5.99	118.91	123.70
84	Aa	1213	G	C5-C6-O6	-5.99	125.01	128.60
84	Aa	1602	A	C5-C6-N6	-5.99	118.91	123.70
84	Aa	2502	U	O3'-P-O5'	-5.99	92.63	104.00
84	Aa	2801	A	C4-C5-C6	5.99	119.99	117.00
84	Aa	16	A	C5-C6-N1	-5.99	114.71	117.70
84	Aa	2745	C	N3-C4-C5	-5.99	119.51	121.90
1	Ad	762	A	C3'-C2'-C1'	5.98	106.29	101.50
84	Aa	917	A	C4-C5-C6	5.98	119.99	117.00
84	Aa	1087	G	C5-C6-O6	-5.98	125.01	128.60
84	Aa	2029	G	O4'-C1'-N9	5.98	112.99	108.20
84	Aa	2047	A	C4-C5-C6	5.98	119.99	117.00
84	Aa	3223	C	N3-C4-C5	-5.98	119.51	121.90
1	Ad	614	G	P-O5'-C5'	-5.98	111.33	120.90
1	Ad	1305	U	O4'-C1'-C2'	-5.98	99.82	105.80
13	BF	148	TYR	CB-CG-CD2	-5.98	117.41	121.00
84	Aa	705	A	C5-C6-N6	-5.98	118.91	123.70
84	Aa	890	G	O4'-C1'-N9	5.98	112.99	108.20
1	Ad	614	G	O4'-C1'-N9	5.98	112.98	108.20
23	Bc	8	ALA	N-CA-CB	5.98	118.47	110.10
84	Aa	218	G	C5-C6-O6	-5.98	125.01	128.60
84	Aa	1546	G	O4'-C1'-N9	5.98	112.98	108.20
84	Aa	2681	A	O4'-C1'-N9	5.98	112.98	108.20
84	Aa	72	A	C4-C5-C6	5.98	119.99	117.00
84	Aa	948	C	N3-C4-C5	-5.98	119.51	121.90
84	Aa	1307	A	C4-C5-C6	5.98	119.99	117.00
84	Aa	1444	G	O4'-C1'-N9	5.98	112.98	108.20
84	Aa	2031	G	C5-C6-O6	-5.98	125.01	128.60
2	Ae	72	G	P-O3'-C3'	-5.98	112.53	119.70
84	Aa	1923	G	O4'-C1'-N9	5.98	112.98	108.20
84	Aa	2145	C	N3-C4-C5	-5.98	119.51	121.90
84	Aa	2224	A	C5-C6-N1	-5.98	114.71	117.70
84	Aa	2439	A	C5-C6-N6	-5.98	118.92	123.70
84	Aa	3044	C	N3-C4-N4	5.98	122.18	118.00
84	Aa	3045	A	C5-C6-N6	-5.98	118.92	123.70
84	Aa	3114	A	C5-C6-N6	-5.98	118.92	123.70
85	Ac	45	C	N3-C4-C5	-5.98	119.51	121.90
1	Ad	755	U	O4'-C1'-N1	5.98	112.98	108.20
84	Aa	499	A	C5-C6-N6	-5.98	118.92	123.70
84	Aa	1747	A	C4-C5-C6	5.98	119.99	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2098	A	C4-C5-C6	5.98	119.99	117.00
85	Ac	41	A	O4'-C1'-N9	5.98	112.98	108.20
1	Ad	868	A	N9-C1'-C2'	-5.97	105.43	112.00
1	Ad	1090	G	O4'-C1'-N9	5.97	112.98	108.20
84	Aa	522	C	N3-C4-C5	-5.97	119.51	121.90
84	Aa	653	A	C4-C5-C6	5.97	119.99	117.00
84	Aa	792	A	C4-C5-C6	5.97	119.99	117.00
84	Aa	1306	A	C4-C5-C6	5.97	119.99	117.00
84	Aa	1630	C	N3-C4-C5	-5.97	119.51	121.90
84	Aa	1650	G	N3-C2-N2	5.97	124.08	119.90
84	Aa	2505	C	N3-C4-N4	5.97	122.18	118.00
84	Aa	2152	A	C4-C5-C6	5.97	119.99	117.00
84	Aa	3327	A	C4-C5-C6	5.97	119.99	117.00
1	Ad	949	A	C3'-C2'-C1'	5.97	106.28	101.50
2	Ae	28	G	O4'-C1'-C2'	5.97	112.97	107.60
1	Ad	244	C	O4'-C1'-C2'	-5.97	99.83	105.80
84	Aa	387	A	C5-C6-N1	-5.97	114.72	117.70
84	Aa	697	A	C5-C6-N1	-5.97	114.72	117.70
84	Aa	1199	A	C5-C6-N1	-5.97	114.72	117.70
84	Aa	1373	A	O4'-C1'-N9	5.97	112.98	108.20
84	Aa	1484	A	P-O3'-C3'	5.97	126.86	119.70
84	Aa	1970	A	C4-C5-C6	5.97	119.98	117.00
84	Aa	2107	A	C4-C5-C6	5.97	119.98	117.00
84	Aa	2312	A	C4-C5-C6	5.97	119.98	117.00
1	Ad	741	C	O4'-C1'-N1	5.97	112.97	108.20
84	Aa	66	A	C5-C6-N1	-5.97	114.72	117.70
84	Aa	1968	C	N3-C4-C5	-5.97	119.51	121.90
84	Aa	2067	G	C5-C6-O6	-5.97	125.02	128.60
84	Aa	3328	A	C4-C5-C6	5.97	119.98	117.00
1	Ad	238	G	O4'-C1'-C2'	-5.97	99.83	105.80
51	CX	140	TYR	CB-CG-CD2	5.97	124.58	121.00
84	Aa	62	A	C4-C5-C6	5.97	119.98	117.00
84	Aa	754	G	O4'-C1'-N9	5.97	112.97	108.20
84	Aa	1048	U	O4'-C1'-N1	5.97	112.97	108.20
84	Aa	2120	A	C4-C5-C6	5.97	119.98	117.00
84	Aa	3013	A	C4-C5-C6	5.97	119.98	117.00
84	Aa	3101	C	N3-C4-N4	5.97	122.18	118.00
84	Aa	3249	G	O4'-C1'-N9	5.97	112.97	108.20
1	Ad	1574	U	C1'-O4'-C4'	5.96	114.67	109.90
84	Aa	50	A	O4'-C1'-N9	5.96	112.97	108.20
84	Aa	1393	G	N1-C6-O6	5.96	123.48	119.90
84	Aa	1698	C	N3-C4-C5	-5.96	119.52	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1706	C	N3-C4-C5	-5.96	119.52	121.90
84	Aa	1879	A	C5-C6-N1	-5.96	114.72	117.70
84	Aa	1972	C	N3-C4-C5	-5.96	119.51	121.90
84	Aa	2059	C	N3-C4-C5	-5.96	119.51	121.90
84	Aa	2165	A	C4-C5-C6	5.96	119.98	117.00
84	Aa	2337	C	N3-C4-C5	-5.96	119.51	121.90
84	Aa	2943	A	O4'-C1'-N9	5.96	112.97	108.20
84	Aa	3288	A	C5-C6-N1	-5.96	114.72	117.70
84	Aa	3310	A	C5-C6-N6	-5.96	118.93	123.70
84	Aa	3340	G	O4'-C1'-N9	5.96	112.97	108.20
1	Ad	61	A	O4'-C4'-C3'	-5.96	98.04	104.00
1	Ad	70	C	O4'-C1'-N1	5.96	112.97	108.20
2	Ae	43	C	O4'-C1'-N1	5.96	112.97	108.20
84	Aa	743	C	N3-C4-C5	-5.96	119.52	121.90
84	Aa	1566	C	N3-C4-C5	-5.96	119.52	121.90
85	Ac	56	G	O4'-C1'-N9	5.96	112.97	108.20
84	Aa	178	C	N3-C4-C5	-5.96	119.52	121.90
84	Aa	543	C	N3-C4-N4	5.96	122.17	118.00
84	Aa	655	G	C5-C6-O6	-5.96	125.02	128.60
84	Aa	813	A	C5-C6-N1	-5.96	114.72	117.70
84	Aa	987	A	O4'-C1'-N9	5.96	112.97	108.20
84	Aa	1450	G	C4'-C3'-C2'	-5.96	96.64	102.60
84	Aa	1499	C	O4'-C1'-N1	5.96	112.97	108.20
84	Aa	2156	U	O4'-C1'-N1	5.96	112.97	108.20
84	Aa	2351	A	C5-C6-N1	-5.96	114.72	117.70
84	Aa	564	A	C5-C6-N1	-5.96	114.72	117.70
84	Aa	656	G	O4'-C1'-N9	5.96	112.97	108.20
84	Aa	1136	A	C4-C5-C6	5.96	119.98	117.00
84	Aa	1365	C	O4'-C1'-N1	5.96	112.97	108.20
85	Ac	14	C	N3-C4-N4	5.96	122.17	118.00
1	Ad	788	G	P-O5'-C5'	-5.96	111.37	120.90
39	CZ	77	PHE	CB-CG-CD1	5.96	124.97	120.80
47	CQ	53	PHE	CB-CG-CD1	5.96	124.97	120.80
84	Aa	47	A	C5-C6-N6	-5.96	118.93	123.70
84	Aa	646	U	O4'-C1'-N1	5.96	112.97	108.20
84	Aa	1114	A	C5-C6-N6	-5.96	118.93	123.70
84	Aa	1378	G	O4'-C1'-N9	5.96	112.97	108.20
84	Aa	2641	A	C4-C5-C6	5.96	119.98	117.00
84	Aa	3173	A	C4-C5-C6	5.96	119.98	117.00
84	Aa	3352	C	N3-C4-C5	-5.96	119.52	121.90
1	Ad	1587	G	C1'-O4'-C4'	-5.96	105.14	109.90
84	Aa	1073	G	O4'-C1'-N9	5.96	112.97	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1882	A	O4'-C1'-N9	5.96	112.97	108.20
84	Aa	2047	A	C5-C6-N1	-5.96	114.72	117.70
84	Aa	2227	A	C4-C5-C6	5.96	119.98	117.00
84	Aa	3028	A	C4-C5-C6	5.96	119.98	117.00
84	Aa	3381	C	N3-C4-N4	5.96	122.17	118.00
85	Ac	119	C	N3-C4-C5	-5.96	119.52	121.90
1	Ad	237	C	N1-C1'-C2'	5.96	121.74	114.00
84	Aa	541	C	N3-C4-C5	-5.96	119.52	121.90
84	Aa	944	G	O4'-C1'-N9	5.96	112.96	108.20
84	Aa	2296	U	O4'-C1'-N1	5.96	112.96	108.20
84	Aa	2662	A	C4-C5-C6	5.96	119.98	117.00
84	Aa	2792	A	C4-C5-C6	5.96	119.98	117.00
84	Aa	3357	C	N3-C4-C5	-5.96	119.52	121.90
1	Ad	1035	A	C3'-C2'-C1'	5.95	106.26	101.50
1	Ad	1163	C	O4'-C1'-N1	-5.95	103.44	108.20
41	CA	69	TYR	CB-CG-CD1	5.95	124.57	121.00
84	Aa	1254	A	C4-C5-C6	5.95	119.98	117.00
84	Aa	1315	G	O4'-C1'-N9	5.95	112.96	108.20
84	Aa	1785	G	C5-C6-O6	-5.95	125.03	128.60
84	Aa	2428	G	O4'-C1'-N9	5.95	112.96	108.20
84	Aa	2665	A	C4-C5-C6	5.95	119.98	117.00
84	Aa	2921	A	C4-C5-C6	5.95	119.98	117.00
84	Aa	3104	A	C4-C5-C6	5.95	119.98	117.00
84	Aa	3331	G	O4'-C1'-N9	5.95	112.96	108.20
85	Ac	92	A	C5-C6-N1	-5.95	114.72	117.70
84	Aa	1684	U	P-O3'-C3'	-5.95	112.56	119.70
84	Aa	2002	G	C5-C6-O6	-5.95	125.03	128.60
84	Aa	2110	G	O4'-C1'-N9	5.95	112.96	108.20
85	Ac	88	A	O4'-C1'-N9	5.95	112.96	108.20
1	Ad	153	U	C3'-C2'-C1'	5.95	106.26	101.50
1	Ad	885	C	C3'-C2'-C1'	5.95	106.26	101.50
1	Ad	1179	C	C3'-C2'-C1'	5.95	106.26	101.50
84	Aa	704	G	N3-C2-N2	5.95	124.07	119.90
84	Aa	782	G	O4'-C1'-N9	5.95	112.96	108.20
84	Aa	1288	C	N3-C4-N4	5.95	122.17	118.00
84	Aa	1459	A	C4-C5-C6	5.95	119.97	117.00
84	Aa	1785	G	O4'-C1'-N9	5.95	112.96	108.20
84	Aa	2350	C	N3-C4-C5	-5.95	119.52	121.90
84	Aa	3297	A	C4-C5-C6	5.95	119.98	117.00
85	Ac	33	A	C5-C6-N6	-5.95	118.94	123.70
84	Aa	1183	C	N3-C4-C5	-5.95	119.52	121.90
84	Aa	2354	G	N3-C2-N2	5.95	124.06	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2998	A	C5-C6-N1	-5.95	114.73	117.70
84	Aa	1819	A	C5-C6-N6	-5.95	118.94	123.70
1	Ad	1231	A	C3'-C2'-C1'	5.95	106.26	101.50
2	Ae	2	C	O4'-C1'-N1	5.95	112.96	108.20
84	Aa	111	C	P-O3'-C3'	5.95	126.84	119.70
84	Aa	143	A	C5-C6-N1	-5.95	114.73	117.70
84	Aa	836	G	C8-N9-C4	-5.95	104.02	106.40
84	Aa	1944	G	O3'-P-O5'	-5.95	92.70	104.00
84	Aa	2193	A	C4-C5-C6	5.95	119.97	117.00
84	Aa	2456	G	O4'-C1'-N9	5.95	112.96	108.20
84	Aa	2993	A	C4-C5-C6	5.95	119.97	117.00
84	Aa	3019	C	N3-C4-C5	-5.95	119.52	121.90
1	Ad	904	G	O4'-C1'-C2'	5.94	112.95	107.60
84	Aa	1731	A	C5-C6-N1	-5.94	114.73	117.70
84	Aa	416	A	C4-C5-C6	5.94	119.97	117.00
84	Aa	547	C	N3-C4-C5	-5.94	119.52	121.90
84	Aa	660	A	C4-C5-C6	5.94	119.97	117.00
84	Aa	936	A	C4-C5-C6	5.94	119.97	117.00
84	Aa	2674	A	C4-C5-C6	5.94	119.97	117.00
84	Aa	2697	A	C4-C5-C6	5.94	119.97	117.00
84	Aa	2788	A	C5-C6-N1	-5.94	114.73	117.70
85	Ac	12	A	C5-C6-N1	-5.94	114.73	117.70
2	Ae	2	C	N1-C1'-C2'	5.94	121.72	114.00
84	Aa	947	C	N3-C4-C5	-5.94	119.52	121.90
84	Aa	993	A	C5-C6-N1	-5.94	114.73	117.70
84	Aa	1010	A	C5-C6-N1	-5.94	114.73	117.70
84	Aa	1459	A	C5-C6-N1	-5.94	114.73	117.70
84	Aa	2161	G	O4'-C1'-N9	5.94	112.95	108.20
84	Aa	2227	A	O4'-C1'-N9	5.94	112.95	108.20
84	Aa	2682	A	C4-C5-C6	5.94	119.97	117.00
84	Aa	2812	C	N3-C4-C5	-5.94	119.52	121.90
84	Aa	2815	A	C5-C6-N1	-5.94	114.73	117.70
84	Aa	3336	A	C4-C5-C6	5.94	119.97	117.00
1	Ad	807	G	O4'-C1'-N9	5.94	112.95	108.20
84	Aa	327	A	C4-C5-C6	5.94	119.97	117.00
84	Aa	1576	C	N3-C4-N4	5.94	122.16	118.00
1	Ad	562	U	O4'-C1'-C2'	5.94	112.94	107.60
1	Ad	1524	A	C3'-C2'-C1'	-5.94	96.75	101.50
84	Aa	1499	C	N3-C4-C5	-5.94	119.53	121.90
84	Aa	1673	A	C4-C5-C6	5.94	119.97	117.00
84	Aa	1864	G	C5-C6-O6	-5.94	125.04	128.60
84	Aa	2006	A	C4-C5-C6	5.94	119.97	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2352	G	C5-C6-O6	-5.94	125.04	128.60
1	Ad	633	U	P-O5'-C5'	-5.94	111.40	120.90
1	Ad	1545	A	O4'-C1'-N9	5.94	112.95	108.20
84	Aa	75	G	C5-C6-O6	-5.94	125.04	128.60
84	Aa	2208	A	C4-C5-C6	5.94	119.97	117.00
84	Aa	2772	A	C4-C5-C6	5.94	119.97	117.00
85	Ac	75	G	O4'-C1'-N9	5.94	112.95	108.20
1	Ad	971	A	C3'-C2'-C1'	5.93	106.25	101.50
1	Ad	1480	G	O4'-C1'-N9	5.93	112.95	108.20
84	Aa	261	C	N3-C4-C5	-5.93	119.53	121.90
84	Aa	2087	A	O4'-C1'-C2'	5.93	112.94	107.60
84	Aa	3083	C	N3-C4-C5	-5.93	119.53	121.90
85	Ac	120	G	O4'-C1'-N9	5.93	112.95	108.20
1	Ad	139	U	P-O5'-C5'	5.93	130.39	120.90
1	Ad	184	C	N1-C1'-C2'	5.93	121.71	114.00
1	Ad	626	A	P-O3'-C3'	5.93	126.82	119.70
84	Aa	517	G	O4'-C1'-N9	5.93	112.94	108.20
84	Aa	557	C	N3-C4-C5	-5.93	119.53	121.90
84	Aa	630	C	N3-C4-N4	5.93	122.15	118.00
84	Aa	1192	A	C4-C5-C6	5.93	119.97	117.00
84	Aa	1294	A	C5-C6-N1	-5.93	114.73	117.70
84	Aa	1395	A	C4-C5-C6	5.93	119.97	117.00
84	Aa	1896	A	N1-C6-N6	5.93	122.16	118.60
84	Aa	1999	G	O4'-C1'-N9	5.93	112.95	108.20
84	Aa	2503	A	O4'-C1'-N9	5.93	112.95	108.20
84	Aa	2737	A	O4'-C1'-N9	5.93	112.94	108.20
84	Aa	2810	A	C4-C5-C6	5.93	119.97	117.00
84	Aa	2841	G	C5-C6-O6	-5.93	125.04	128.60
84	Aa	2997	C	N3-C4-N4	5.93	122.15	118.00
56	Cd	33	PHE	CB-CG-CD2	-5.93	116.65	120.80
84	Aa	1058	A	C4-C5-C6	5.93	119.97	117.00
84	Aa	1229	A	C4-C5-C6	5.93	119.97	117.00
84	Aa	2003	C	N3-C4-C5	-5.93	119.53	121.90
85	Ac	1	C	N3-C4-N4	5.93	122.15	118.00
1	Ad	493	C	C3'-C2'-C1'	5.93	106.24	101.50
1	Ad	1733	G	O4'-C1'-N9	5.93	112.94	108.20
84	Aa	174	G	C5-C6-O6	-5.93	125.04	128.60
84	Aa	319	C	N3-C4-C5	-5.93	119.53	121.90
84	Aa	1192	A	C5-C6-N1	-5.93	114.73	117.70
84	Aa	1534	C	N3-C4-C5	-5.93	119.53	121.90
84	Aa	1750	A	C5-C6-N1	-5.93	114.73	117.70
84	Aa	2377	C	N3-C4-C5	-5.93	119.53	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2700	A	C4-C5-C6	5.93	119.96	117.00
84	Aa	2855	G	O4'-C1'-N9	5.93	112.94	108.20
84	Aa	3153	U	C6-N1-C1'	-5.93	112.90	121.20
79	CE	205	ARG	N-CA-CB	5.93	121.27	110.60
84	Aa	928	A	O4'-C1'-N9	5.93	112.94	108.20
84	Aa	2248	G	C5-C6-O6	-5.93	125.04	128.60
84	Aa	2357	A	O4'-C1'-N9	5.93	112.94	108.20
84	Aa	3057	A	C4-C5-C6	5.93	119.96	117.00
85	Ac	76	C	N3-C4-C5	-5.93	119.53	121.90
1	Ad	215	A	C1'-O4'-C4'	5.93	114.64	109.90
84	Aa	1274	A	C5-C6-N6	-5.93	118.96	123.70
84	Aa	1339	C	N3-C4-C5	-5.93	119.53	121.90
84	Aa	1493	A	C5-C6-N1	-5.93	114.74	117.70
84	Aa	1977	C	N3-C4-C5	-5.93	119.53	121.90
84	Aa	2197	C	O4'-C1'-N1	5.93	112.94	108.20
84	Aa	2273	C	N3-C4-N4	5.93	122.15	118.00
84	Aa	2368	G	O4'-C1'-N9	5.93	112.94	108.20
84	Aa	2594	A	C4-C5-C6	5.93	119.96	117.00
84	Aa	3115	A	C4-C5-C6	5.93	119.96	117.00
84	Aa	3148	A	C5-C6-N6	-5.93	118.96	123.70
2	Ae	75	A	O4'-C1'-N9	5.92	112.94	108.20
84	Aa	224	C	N3-C4-N4	5.92	122.15	118.00
84	Aa	387	A	C4-C5-C6	5.92	119.96	117.00
84	Aa	2065	G	O4'-C1'-N9	5.92	112.94	108.20
84	Aa	2385	A	C5-C6-N1	-5.92	114.74	117.70
84	Aa	2385	A	P-O5'-C5'	5.92	130.38	120.90
84	Aa	2958	A	O4'-C1'-N9	5.92	112.94	108.20
84	Aa	3007	A	C4-C5-C6	5.92	119.96	117.00
85	Ac	62	C	N3-C4-C5	-5.92	119.53	121.90
85	Ac	106	C	N3-C4-C5	-5.92	119.53	121.90
84	Aa	103	G	O4'-C1'-N9	5.92	112.94	108.20
84	Aa	407	A	P-O5'-C5'	5.92	130.38	120.90
84	Aa	3028	A	O4'-C1'-N9	5.92	112.94	108.20
1	Ad	541	G	P-O3'-C3'	5.92	126.81	119.70
84	Aa	26	A	C4-C5-C6	5.92	119.96	117.00
84	Aa	939	A	C4-C5-C6	5.92	119.96	117.00
84	Aa	1455	A	C4-C5-C6	5.92	119.96	117.00
84	Aa	2449	A	C4-C5-C6	5.92	119.96	117.00
84	Aa	2840	A	C5-C6-N6	-5.92	118.96	123.70
84	Aa	3012	A	C4-C5-C6	5.92	119.96	117.00
84	Aa	3329	G	C5-C6-O6	-5.92	125.05	128.60
84	Aa	272	G	C5-C6-O6	-5.92	125.05	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	783	A	C4-C5-C6	5.92	119.96	117.00
84	Aa	1044	A	O4'-C1'-N9	5.92	112.94	108.20
84	Aa	1484	A	C4-C5-C6	5.92	119.96	117.00
1	Ad	1629	U	N1-C1'-C2'	5.92	121.69	114.00
84	Aa	23	A	C4-C5-C6	5.92	119.96	117.00
84	Aa	183	C	O4'-C1'-N1	5.92	112.93	108.20
84	Aa	314	C	N3-C4-C5	-5.92	119.53	121.90
84	Aa	412	C	N3-C4-C5	-5.92	119.53	121.90
84	Aa	494	C	N3-C4-C5	-5.92	119.53	121.90
84	Aa	558	G	O4'-C1'-N9	5.92	112.94	108.20
84	Aa	996	A	C5-C6-N1	-5.92	114.74	117.70
84	Aa	1009	G	O4'-C1'-N9	5.92	112.93	108.20
84	Aa	1355	U	N1-C1'-C2'	5.92	121.69	114.00
84	Aa	1382	C	N3-C4-N4	5.92	122.14	118.00
84	Aa	2751	A	C4-C5-C6	5.92	119.96	117.00
84	Aa	3134	C	N3-C4-C5	-5.92	119.53	121.90
1	Ad	371	A	P-O5'-C5'	-5.92	111.43	120.90
84	Aa	175	G	O4'-C1'-N9	5.92	112.93	108.20
84	Aa	1406	C	N3-C4-C5	-5.92	119.53	121.90
84	Aa	1410	A	C4-C5-C6	5.92	119.96	117.00
84	Aa	1951	C	N3-C4-N4	5.92	122.14	118.00
84	Aa	2048	C	N3-C4-C5	-5.92	119.53	121.90
84	Aa	2294	A	C5-C6-N6	-5.92	118.97	123.70
84	Aa	2359	C	N3-C4-C5	-5.92	119.53	121.90
84	Aa	2601	G	O4'-C1'-N9	5.92	112.93	108.20
84	Aa	2829	U	O4'-C1'-N1	5.92	112.93	108.20
84	Aa	3063	C	N3-C4-C5	-5.92	119.53	121.90
85	Ac	160	C	N3-C4-N4	5.92	122.14	118.00
84	Aa	1452	A	C5-C6-N1	-5.92	114.74	117.70
84	Aa	2532	A	C5-C6-N1	-5.92	114.74	117.70
45	CN	30	TYR	CB-CG-CD2	-5.91	117.45	121.00
84	Aa	91	G	C5-C6-O6	-5.91	125.05	128.60
84	Aa	301	G	O4'-C1'-N9	5.91	112.93	108.20
84	Aa	574	C	N3-C4-N4	5.91	122.14	118.00
84	Aa	830	A	C5-C6-N6	-5.91	118.97	123.70
84	Aa	1114	A	C4-C5-C6	5.91	119.96	117.00
84	Aa	1512	A	C4-C5-C6	5.91	119.96	117.00
84	Aa	2016	A	O4'-C1'-N9	5.91	112.93	108.20
84	Aa	2935	A	C5-C6-N6	-5.91	118.97	123.70
84	Aa	3086	G	O4'-C1'-N9	5.91	112.93	108.20
84	Aa	3306	A	C5-C6-N1	-5.91	114.74	117.70
1	Ad	1284	C	O4'-C1'-N1	5.91	112.93	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	732	G	O4'-C1'-N9	5.91	112.93	108.20
84	Aa	921	C	N3-C4-C5	-5.91	119.53	121.90
84	Aa	1027	C	N3-C4-C5	-5.91	119.53	121.90
84	Aa	3069	U	O4'-C1'-N1	5.91	112.93	108.20
84	Aa	3228	C	N3-C4-C5	-5.91	119.53	121.90
84	Aa	3358	A	C5-C6-N6	-5.91	118.97	123.70
1	Ad	93	A	O4'-C1'-C2'	-5.91	99.89	105.80
84	Aa	886	A	C5-C6-N6	-5.91	118.97	123.70
84	Aa	1042	C	N3-C4-C5	-5.91	119.54	121.90
84	Aa	1180	C	N3-C4-N4	5.91	122.14	118.00
84	Aa	1835	A	C5-C6-N6	-5.91	118.97	123.70
84	Aa	2026	C	N3-C4-C5	-5.91	119.54	121.90
84	Aa	3339	G	O4'-C1'-N9	5.91	112.93	108.20
1	Ad	1229	C	C3'-C2'-C1'	5.91	106.23	101.50
1	Ad	1732	A	C1'-O4'-C4'	5.91	114.63	109.90
84	Aa	217	A	C4-C5-C6	5.91	119.95	117.00
84	Aa	323	A	C5-C6-N1	-5.91	114.75	117.70
84	Aa	346	A	C4-C5-C6	5.91	119.95	117.00
84	Aa	480	C	N3-C4-N4	5.91	122.14	118.00
84	Aa	748	C	N3-C4-C5	-5.91	119.54	121.90
84	Aa	1052	A	C5-C6-N6	-5.91	118.97	123.70
84	Aa	2694	A	C4-C5-C6	5.91	119.95	117.00
84	Aa	2944	C	N3-C4-N4	5.91	122.14	118.00
84	Aa	3050	A	O4'-C1'-N9	5.91	112.93	108.20
85	Ac	37	A	C4-C5-C6	5.91	119.95	117.00
48	CD	125	GLU	N-CA-CB	5.91	121.23	110.60
84	Aa	230	G	O4'-C1'-N9	5.91	112.92	108.20
84	Aa	121	A	C5-C6-N1	-5.91	114.75	117.70
84	Aa	313	C	N3-C4-C5	-5.91	119.54	121.90
84	Aa	847	G	O4'-C1'-N9	5.91	112.92	108.20
84	Aa	2370	G	O4'-C1'-N9	5.91	112.92	108.20
84	Aa	3287	A	C5-C6-N1	-5.91	114.75	117.70
1	Ad	892	A	C1'-O4'-C4'	5.90	114.62	109.90
84	Aa	1464	A	C4-C5-C6	5.90	119.95	117.00
1	Ad	487	A	O4'-C1'-C2'	5.90	112.91	107.60
1	Ad	749	G	C3'-C2'-C1'	-5.90	96.78	101.50
84	Aa	711	A	O4'-C1'-N9	5.90	112.92	108.20
84	Aa	924	A	C5-C6-N6	-5.90	118.98	123.70
84	Aa	953	G	C5-C6-O6	-5.90	125.06	128.60
84	Aa	955	A	C5-C6-N6	-5.90	118.98	123.70
84	Aa	1770	C	N3-C4-C5	-5.90	119.54	121.90
84	Aa	2364	C	N3-C4-C5	-5.90	119.54	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2423	A	C4-C5-C6	5.90	119.95	117.00
85	Ac	43	A	C4-C5-C6	5.90	119.95	117.00
1	Ad	136	U	N1-C1'-C2'	5.90	121.67	114.00
1	Ad	716	A	C2'-C3'-O3'	5.90	123.14	113.70
84	Aa	244	G	C5-C6-O6	-5.90	125.06	128.60
84	Aa	360	G	O4'-C1'-N9	5.90	112.92	108.20
84	Aa	1136	A	O4'-C1'-N9	5.90	112.92	108.20
84	Aa	1283	C	N3-C4-N4	5.90	122.13	118.00
84	Aa	2110	G	C5-C6-O6	-5.90	125.06	128.60
1	Ad	85	A	O4'-C1'-C2'	-5.90	99.90	105.80
84	Aa	2012	C	N3-C4-C5	-5.90	119.54	121.90
84	Aa	2631	A	C5-C6-N1	-5.90	114.75	117.70
84	Aa	2643	A	C5-C6-N1	-5.90	114.75	117.70
1	Ad	1285	G	O4'-C1'-N9	5.90	112.92	108.20
1	Ad	1408	G	C2'-C3'-O3'	5.90	123.14	113.70
84	Aa	410	G	N3-C2-N2	5.90	124.03	119.90
84	Aa	758	A	O4'-C1'-N9	5.90	112.92	108.20
84	Aa	1074	C	N3-C4-C5	-5.90	119.54	121.90
84	Aa	1098	U	O4'-C1'-N1	5.90	112.92	108.20
84	Aa	1674	A	C4-C5-C6	5.90	119.95	117.00
84	Aa	2011	G	O4'-C1'-N9	5.90	112.92	108.20
84	Aa	2055	U	O4'-C1'-N1	5.90	112.92	108.20
84	Aa	2655	U	O4'-C1'-N1	5.90	112.92	108.20
84	Aa	2839	A	C4-C5-C6	5.90	119.95	117.00
84	Aa	2998	A	C4-C5-C6	5.90	119.95	117.00
84	Aa	3030	A	O4'-C1'-N9	5.90	112.92	108.20
84	Aa	3305	U	O4'-C1'-N1	5.90	112.92	108.20
84	Aa	161	C	N3-C4-N4	5.90	122.13	118.00
84	Aa	1987	C	N3-C4-C5	-5.90	119.54	121.90
84	Aa	2795	G	O4'-C1'-N9	5.90	112.92	108.20
84	Aa	3234	G	C5-C6-O6	-5.90	125.06	128.60
1	Ad	1368	C	C3'-C2'-C1'	-5.89	96.78	101.50
84	Aa	325	A	C4-C5-C6	5.89	119.95	117.00
84	Aa	529	C	O4'-C1'-N1	5.89	112.92	108.20
84	Aa	658	C	N3-C4-C5	-5.89	119.54	121.90
84	Aa	1649	G	N3-C2-N2	5.89	124.03	119.90
84	Aa	2590	C	N3-C4-N4	5.89	122.13	118.00
84	Aa	2665	A	C5-C6-N1	-5.89	114.75	117.70
84	Aa	3373	C	N3-C4-C5	-5.89	119.54	121.90
1	Ad	1345	G	O4'-C1'-C2'	5.89	112.90	107.60
9	BX	128	SER	N-CA-CB	5.89	119.34	110.50
71	CB	120	LYS	N-CA-C	-5.89	95.09	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	790	G	O4'-C1'-N9	5.89	112.91	108.20
84	Aa	1640	A	C5-C6-N6	-5.89	118.99	123.70
84	Aa	1882	A	C4-C5-C6	5.89	119.95	117.00
84	Aa	2014	A	C5-C6-N1	-5.89	114.75	117.70
84	Aa	2593	A	O4'-C1'-N9	5.89	112.91	108.20
84	Aa	2667	C	C2-N3-C4	5.89	122.85	119.90
84	Aa	2934	C	N3-C4-N4	5.89	122.12	118.00
84	Aa	3211	C	N3-C4-N4	5.89	122.12	118.00
1	Ad	1342	C	O4'-C1'-N1	5.89	112.91	108.20
84	Aa	46	A	C4-C5-C6	5.89	119.94	117.00
84	Aa	2386	A	C5-C6-N1	-5.89	114.75	117.70
85	Ac	124	C	N3-C4-C5	-5.89	119.54	121.90
1	Ad	1047	G	O4'-C1'-C2'	5.89	112.90	107.60
84	Aa	140	C	N3-C4-C5	-5.89	119.54	121.90
84	Aa	631	C	N3-C4-N4	5.89	122.12	118.00
84	Aa	856	G	O4'-C1'-N9	5.89	112.91	108.20
84	Aa	1300	C	N3-C4-C5	-5.89	119.54	121.90
84	Aa	1905	A	O4'-C1'-N9	5.89	112.91	108.20
84	Aa	2060	C	N3-C4-C5	-5.89	119.54	121.90
84	Aa	2123	C	N3-C4-C5	-5.89	119.54	121.90
84	Aa	2532	A	C4-C5-C6	5.89	119.94	117.00
84	Aa	373	A	C5-C6-N6	-5.89	118.99	123.70
84	Aa	672	A	C5-C6-N6	-5.89	118.99	123.70
84	Aa	2281	U	O4'-C1'-N1	5.89	112.91	108.20
84	Aa	2759	C	N3-C4-C5	-5.89	119.55	121.90
84	Aa	491	G	O4'-C1'-N9	5.89	112.91	108.20
84	Aa	1440	C	N3-C4-C5	-5.89	119.55	121.90
84	Aa	2349	C	N3-C4-C5	-5.89	119.55	121.90
84	Aa	2846	C	N3-C4-N4	5.89	122.12	118.00
85	Ac	87	G	C5-C6-O6	-5.89	125.07	128.60
86	Ab	30	G	N1-C6-O6	5.89	123.43	119.90
1	Ad	1012	C	C3'-C2'-C1'	5.88	106.21	101.50
1	Ad	1023	C	C3'-C2'-C1'	5.88	106.21	101.50
1	Ad	1042	C	O4'-C1'-C2'	-5.88	99.92	105.80
84	Aa	1580	C	N3-C4-C5	-5.88	119.55	121.90
84	Aa	2090	G	O4'-C1'-N9	5.88	112.91	108.20
84	Aa	2794	A	C5-C6-N1	-5.88	114.76	117.70
1	Ad	1710	C	C3'-C2'-C1'	5.88	106.21	101.50
20	BT	51	TYR	CB-CG-CD1	-5.88	117.47	121.00
84	Aa	1263	A	C4-C5-C6	5.88	119.94	117.00
84	Aa	2351	A	C4-C5-C6	5.88	119.94	117.00
84	Aa	2718	A	C4-C5-C6	5.88	119.94	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
6	BK	64	TYR	CB-CG-CD2	-5.88	117.47	121.00
41	CA	68	ARG	N-CA-CB	5.88	121.19	110.60
84	Aa	1035	C	N3-C4-C5	-5.88	119.55	121.90
84	Aa	842	C	O4'-C1'-N1	5.88	112.90	108.20
84	Aa	869	A	C5-C6-N6	-5.88	119.00	123.70
84	Aa	2762	U	O4'-C1'-N1	5.88	112.90	108.20
84	Aa	2813	A	O4'-C1'-N9	5.88	112.90	108.20
1	Ad	535	C	P-O5'-C5'	5.88	130.31	120.90
1	Ad	1705	C	C1'-O4'-C4'	-5.88	105.20	109.90
84	Aa	9	C	N3-C4-C5	-5.88	119.55	121.90
84	Aa	846	A	C5-C6-N1	-5.88	114.76	117.70
84	Aa	2045	G	C5-C6-O6	-5.88	125.07	128.60
84	Aa	3321	C	N3-C4-N4	5.88	122.11	118.00
1	Ad	1670	G	C1'-O4'-C4'	-5.88	105.20	109.90
25	Bd	53	ILE	C-N-CA	5.88	136.39	121.70
84	Aa	6	A	C5-C6-N6	-5.88	119.00	123.70
84	Aa	1002	A	C4-C5-C6	5.88	119.94	117.00
84	Aa	1866	C	N3-C4-C5	-5.88	119.55	121.90
84	Aa	2102	C	N3-C4-N4	5.88	122.11	118.00
1	Ad	413	C	O4'-C1'-C2'	-5.88	99.92	105.80
84	Aa	46	A	C5-C6-N6	-5.88	119.00	123.70
84	Aa	99	A	C5-C6-N6	-5.88	119.00	123.70
84	Aa	1731	A	C4-C5-C6	5.88	119.94	117.00
84	Aa	2491	A	C4-C5-C6	5.88	119.94	117.00
84	Aa	2581	C	N3-C4-C5	-5.88	119.55	121.90
84	Aa	2969	A	C4-C5-C6	5.88	119.94	117.00
84	Aa	3189	C	N3-C4-C5	-5.88	119.55	121.90
85	Ac	109	A	C4-C5-C6	5.88	119.94	117.00
1	Ad	152	G	C1'-O4'-C4'	-5.87	105.20	109.90
1	Ad	1299	G	C1'-O4'-C4'	5.87	114.60	109.90
1	Ad	1624	G	O4'-C1'-C2'	5.87	112.89	107.60
84	Aa	95	G	C5-C6-O6	-5.87	125.08	128.60
84	Aa	610	G	N1-C6-O6	5.87	123.42	119.90
84	Aa	730	A	C5-C6-N6	-5.87	119.00	123.70
84	Aa	1278	A	O4'-C1'-N9	5.87	112.90	108.20
84	Aa	1582	C	N3-C4-N4	5.87	122.11	118.00
84	Aa	1623	C	N3-C4-N4	5.87	122.11	118.00
84	Aa	1802	A	C4-C5-C6	5.87	119.94	117.00
84	Aa	2630	A	C5-C6-N6	-5.87	119.00	123.70
84	Aa	2851	C	N3-C4-C5	-5.87	119.55	121.90
84	Aa	2942	A	O4'-C1'-N9	5.87	112.90	108.20
84	Aa	3085	C	N3-C4-C5	-5.87	119.55	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	Ac	12	A	P-O5'-C5'	-5.87	111.50	120.90
1	Ad	1143	A	C3'-C2'-C1'	5.87	106.20	101.50
84	Aa	320	U	O4'-C1'-N1	5.87	112.90	108.20
84	Aa	566	G	C4'-C3'-C2'	-5.87	96.73	102.60
84	Aa	615	A	O4'-C1'-N9	5.87	112.90	108.20
84	Aa	1110	C	N3-C4-C5	-5.87	119.55	121.90
84	Aa	1568	A	C4-C5-C6	5.87	119.94	117.00
84	Aa	1788	C	N3-C4-C5	-5.87	119.55	121.90
84	Aa	2101	A	C5-C6-N1	-5.87	114.76	117.70
84	Aa	2724	A	C4-C5-C6	5.87	119.94	117.00
84	Aa	2996	A	C5-C6-N1	-5.87	114.77	117.70
84	Aa	3221	A	C4-C5-C6	5.87	119.94	117.00
84	Aa	3235	A	C4-C5-C6	5.87	119.94	117.00
85	Ac	79	A	C4-C5-C6	5.87	119.94	117.00
1	Ad	941	G	O4'-C1'-N9	5.87	112.90	108.20
1	Ad	1109	U	P-O5'-C5'	5.87	130.29	120.90
84	Aa	861	A	C4-C5-C6	5.87	119.94	117.00
84	Aa	3271	A	C4-C5-C6	5.87	119.94	117.00
86	Ab	30	G	C8-N9-C4	-5.87	104.05	106.40
1	Ad	326	G	P-O3'-C3'	5.87	126.74	119.70
84	Aa	434	C	N3-C4-C5	-5.87	119.55	121.90
84	Aa	474	G	O3'-P-O5'	-5.87	92.85	104.00
84	Aa	1097	A	C5-C6-N6	-5.87	119.01	123.70
84	Aa	1629	A	C4-C5-C6	5.87	119.93	117.00
84	Aa	2503	A	P-O5'-C5'	5.87	130.29	120.90
84	Aa	2692	G	C4-N9-C1'	5.87	134.13	126.50
84	Aa	2749	A	C4-C5-C6	5.87	119.93	117.00
84	Aa	3228	C	N3-C4-N4	5.87	122.11	118.00
84	Aa	3291	C	N3-C4-N4	5.87	122.11	118.00
84	Aa	121	A	C5-C6-N6	-5.87	119.01	123.70
84	Aa	200	G	C5-C6-O6	-5.87	125.08	128.60
84	Aa	1007	A	C4-C5-C6	5.87	119.93	117.00
85	Ac	144	C	N3-C4-N4	5.87	122.11	118.00
1	Ad	203	A	C1'-O4'-C4'	-5.87	105.21	109.90
48	CD	245	ALA	N-CA-CB	5.87	118.31	110.10
84	Aa	212	G	N3-C2-N2	5.87	124.01	119.90
84	Aa	811	A	O4'-C1'-N9	5.87	112.89	108.20
84	Aa	2329	C	N3-C4-C5	-5.87	119.55	121.90
84	Aa	2671	A	C4-C5-C6	5.87	119.93	117.00
86	Ab	7	G	C6-C5-N7	-5.87	126.88	130.40
1	Ad	263	C	O4'-C1'-C2'	-5.86	99.94	105.80
84	Aa	956	G	O4'-C1'-N9	5.86	112.89	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1398	A	C4-C5-C6	5.86	119.93	117.00
84	Aa	1600	A	C5-C6-N1	-5.86	114.77	117.70
84	Aa	2054	A	C4-C5-C6	5.86	119.93	117.00
84	Aa	2713	G	C5-C6-O6	-5.86	125.08	128.60
84	Aa	62	A	C5-C6-N6	-5.86	119.01	123.70
84	Aa	149	A	C4-C5-C6	5.86	119.93	117.00
84	Aa	349	A	C4-C5-C6	5.86	119.93	117.00
84	Aa	569	C	N3-C4-C5	-5.86	119.56	121.90
84	Aa	1590	A	C4-C5-C6	5.86	119.93	117.00
84	Aa	2374	G	C5-C6-O6	-5.86	125.08	128.60
84	Aa	2375	G	O4'-C1'-N9	5.86	112.89	108.20
85	Ac	19	A	O4'-C1'-N9	5.86	112.89	108.20
85	Ac	125	C	N3-C4-C5	-5.86	119.56	121.90
62	CS	6	PHE	N-CA-CB	5.86	121.15	110.60
84	Aa	367	A	C4-C5-C6	5.86	119.93	117.00
84	Aa	897	U	O4'-C1'-N1	5.86	112.89	108.20
84	Aa	2009	C	N3-C4-C5	-5.86	119.56	121.90
84	Aa	2803	A	C4-C5-C6	5.86	119.93	117.00
84	Aa	3003	C	N3-C4-N4	5.86	122.10	118.00
85	Ac	9	G	C5-C6-O6	-5.86	125.08	128.60
84	Aa	1797	U	C5'-C4'-O4'	5.86	116.13	109.10
84	Aa	2586	C	N3-C4-C5	-5.86	119.56	121.90
1	Ad	827	C	O4'-C1'-N1	5.86	112.89	108.20
84	Aa	134	U	O4'-C1'-N1	5.86	112.89	108.20
84	Aa	1582	C	N3-C4-C5	-5.86	119.56	121.90
84	Aa	1951	C	N3-C4-C5	-5.86	119.56	121.90
84	Aa	2116	G	O4'-C1'-N9	5.86	112.89	108.20
84	Aa	2733	A	C4-C5-C6	5.86	119.93	117.00
84	Aa	2912	A	C5-C6-N6	-5.86	119.01	123.70
84	Aa	2974	G	O4'-C1'-N9	5.86	112.89	108.20
84	Aa	3116	C	O4'-C1'-N1	5.86	112.89	108.20
1	Ad	1237	G	C1'-O4'-C4'	-5.86	105.22	109.90
1	Ad	1794	C	N1-C1'-C2'	5.86	121.61	114.00
84	Aa	309	C	N3-C4-C5	-5.86	119.56	121.90
84	Aa	707	G	O4'-C1'-N9	5.86	112.88	108.20
84	Aa	1157	A	C4-C5-C6	5.86	119.93	117.00
84	Aa	1392	U	O4'-C1'-N1	5.86	112.89	108.20
84	Aa	1487	A	O4'-C1'-N9	5.86	112.88	108.20
84	Aa	2223	A	C4-C5-C6	5.86	119.93	117.00
84	Aa	2356	A	C5-C6-N1	-5.86	114.77	117.70
84	Aa	3119	C	N3-C4-C5	-5.86	119.56	121.90
85	Ac	50	C	N3-C4-N4	5.86	122.10	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1412	A	C1'-O4'-C4'	-5.85	105.22	109.90
70	Cq	140	PHE	CB-CG-CD1	5.85	124.90	120.80
84	Aa	100	C	C6-N1-C1'	-5.85	113.78	120.80
84	Aa	221	C	N3-C4-C5	-5.85	119.56	121.90
85	Ac	38	U	O4'-C1'-N1	5.85	112.88	108.20
85	Ac	144	C	N3-C4-C5	-5.85	119.56	121.90
1	Ad	1182	C	O4'-C1'-N1	5.85	112.88	108.20
84	Aa	741	G	C5-C6-O6	-5.85	125.09	128.60
84	Aa	1178	C	N3-C4-C5	-5.85	119.56	121.90
84	Aa	1516	G	O4'-C1'-N9	5.85	112.88	108.20
84	Aa	2077	C	N3-C4-N4	5.85	122.10	118.00
84	Aa	2227	A	C5-C6-N6	-5.85	119.02	123.70
84	Aa	2443	C	N3-C4-C5	-5.85	119.56	121.90
84	Aa	2683	A	N1-C6-N6	5.85	122.11	118.60
84	Aa	2708	A	C4-C5-C6	5.85	119.93	117.00
1	Ad	1391	G	O4'-C1'-C2'	5.85	112.87	107.60
84	Aa	1128	U	O4'-C1'-N1	5.85	112.88	108.20
84	Aa	1969	G	C5-C6-O6	-5.85	125.09	128.60
84	Aa	2053	A	C5-C6-N1	-5.85	114.77	117.70
84	Aa	2650	A	C5-C6-N1	-5.85	114.78	117.70
1	Ad	805	A	O4'-C1'-N9	5.85	112.88	108.20
82	Cb	39	PHE	N-CA-CB	5.85	121.13	110.60
84	Aa	438	G	O4'-C1'-N9	5.85	112.88	108.20
84	Aa	854	C	O4'-C1'-N1	5.85	112.88	108.20
84	Aa	952	C	N3-C4-C5	-5.85	119.56	121.90
84	Aa	1109	G	O4'-C1'-N9	5.85	112.88	108.20
84	Aa	1308	A	O4'-C1'-N9	5.85	112.88	108.20
84	Aa	2124	G	C5-C6-O6	-5.85	125.09	128.60
84	Aa	2400	A	C5-C6-N6	-5.85	119.02	123.70
84	Aa	2739	A	C5-C6-N1	-5.85	114.78	117.70
84	Aa	2883	C	N3-C4-C5	-5.85	119.56	121.90
14	BQ	102	TYR	CB-CG-CD1	5.85	124.51	121.00
35	BG	28	PHE	CB-CG-CD1	5.85	124.89	120.80
84	Aa	355	C	C5'-C4'-C3'	-5.85	106.64	116.00
84	Aa	634	A	C5'-C4'-C3'	-5.85	106.64	116.00
84	Aa	1819	A	C5-C6-N1	-5.85	114.78	117.70
84	Aa	2099	G	C5-C6-O6	-5.85	125.09	128.60
85	Ac	121	A	C5-C6-N6	-5.85	119.02	123.70
1	Ad	530	A	C3'-C2'-C1'	5.85	106.18	101.50
84	Aa	249	A	C4-C5-C6	5.85	119.92	117.00
84	Aa	366	G	O4'-C1'-N9	5.85	112.88	108.20
84	Aa	1324	C	N3-C4-C5	-5.85	119.56	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1494	A	C4-C5-C6	5.85	119.92	117.00
84	Aa	1660	C	N3-C4-C5	-5.85	119.56	121.90
84	Aa	2209	A	O4'-C1'-N9	5.85	112.88	108.20
84	Aa	3164	C	N3-C4-C5	-5.85	119.56	121.90
84	Aa	489	C	N3-C4-C5	-5.84	119.56	121.90
84	Aa	634	A	O4'-C1'-N9	5.84	112.88	108.20
84	Aa	984	A	C4-C5-C6	5.84	119.92	117.00
84	Aa	1373	A	C5-C6-N6	-5.84	119.03	123.70
84	Aa	1961	C	N3-C4-C5	-5.84	119.56	121.90
84	Aa	2214	A	C5-C6-N1	-5.84	114.78	117.70
84	Aa	2631	A	C4-C5-C6	5.84	119.92	117.00
84	Aa	2971	A	O4'-C1'-N9	5.84	112.88	108.20
85	Ac	146	G	O4'-C1'-N9	5.84	112.88	108.20
86	Ab	17	G	O4'-C1'-N9	5.84	112.88	108.20
1	Ad	1075	G	O4'-C1'-N9	5.84	112.87	108.20
84	Aa	148	U	O4'-C1'-N1	5.84	112.87	108.20
84	Aa	323	A	O4'-C1'-N9	5.84	112.87	108.20
84	Aa	2150	C	C5'-C4'-O4'	5.84	116.11	109.10
84	Aa	3203	G	C5-C6-O6	-5.84	125.09	128.60
85	Ac	138	G	O4'-C1'-N9	5.84	112.87	108.20
1	Ad	1492	G	C1'-O4'-C4'	5.84	114.57	109.90
84	Aa	257	C	N3-C4-C5	-5.84	119.56	121.90
84	Aa	662	G	O4'-C1'-N9	5.84	112.87	108.20
84	Aa	1704	A	C5-C6-N1	-5.84	114.78	117.70
84	Aa	1781	C	N3-C4-C5	-5.84	119.56	121.90
84	Aa	2082	A	C5-C6-N1	-5.84	114.78	117.70
84	Aa	2411	G	O4'-C1'-N9	5.84	112.87	108.20
1	Ad	432	A	C5'-C4'-O4'	5.84	116.11	109.10
1	Ad	1544	G	P-O3'-C3'	5.84	126.71	119.70
84	Aa	2627	G	O4'-C1'-N9	5.84	112.87	108.20
84	Aa	3011	U	O4'-C1'-N1	5.84	112.87	108.20
84	Aa	832	C	N3-C4-N4	5.84	122.09	118.00
84	Aa	1365	C	N3-C4-C5	-5.84	119.56	121.90
84	Aa	1629	A	O4'-C1'-N9	5.84	112.87	108.20
84	Aa	2591	G	O4'-C1'-N9	5.84	112.87	108.20
84	Aa	3210	G	P-O3'-C3'	5.84	126.70	119.70
84	Aa	3283	G	O4'-C1'-N9	5.84	112.87	108.20
1	Ad	202	C	O4'-C1'-N1	5.84	112.87	108.20
1	Ad	526	U	N1-C1'-C2'	5.84	121.59	114.00
1	Ad	861	A	O4'-C1'-N9	-5.84	103.53	108.20
1	Ad	969	U	C3'-C2'-C1'	5.84	106.17	101.50
1	Ad	1386	U	C1'-O4'-C4'	-5.84	105.23	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	129	G	O4'-C1'-N9	5.84	112.87	108.20
84	Aa	212	G	O4'-C1'-N9	5.84	112.87	108.20
84	Aa	739	C	N3-C4-N4	5.84	122.09	118.00
84	Aa	1904	A	C4-C5-C6	5.84	119.92	117.00
84	Aa	1905	A	C5-C6-N1	-5.84	114.78	117.70
84	Aa	1927	A	C4-C5-C6	5.84	119.92	117.00
84	Aa	2183	A	C4-C5-C6	5.84	119.92	117.00
84	Aa	2203	A	C4-C5-C6	5.84	119.92	117.00
84	Aa	2567	C	N3-C4-N4	5.84	122.09	118.00
84	Aa	2826	G	O4'-C1'-N9	5.84	112.87	108.20
85	Ac	26	C	N3-C4-C5	-5.84	119.56	121.90
84	Aa	26	A	C5-C6-N1	-5.83	114.78	117.70
84	Aa	709	G	P-O3'-C3'	5.83	126.70	119.70
84	Aa	2016	A	C4-C5-C6	5.83	119.92	117.00
84	Aa	3220	A	C5-C6-N1	-5.83	114.78	117.70
1	Ad	288	G	O4'-C1'-N9	5.83	112.87	108.20
1	Ad	1342	C	C1'-O4'-C4'	-5.83	105.23	109.90
84	Aa	111	C	N3-C4-C5	-5.83	119.57	121.90
84	Aa	1573	G	C5-C6-O6	-5.83	125.10	128.60
84	Aa	2674	A	C5-C6-N1	-5.83	114.78	117.70
84	Aa	3201	A	C5-C6-N6	-5.83	119.03	123.70
1	Ad	135	C	N1-C1'-C2'	5.83	121.58	114.00
1	Ad	854	C	O4'-C1'-N1	5.83	112.86	108.20
1	Ad	993	C	C3'-C2'-C1'	5.83	106.16	101.50
3	Af	15	A	O4'-C1'-N9	5.83	112.86	108.20
84	Aa	191	C	N3-C4-C5	-5.83	119.57	121.90
84	Aa	913	G	O4'-C1'-N9	5.83	112.86	108.20
84	Aa	1911	A	C5-C6-N1	-5.83	114.78	117.70
84	Aa	2139	A	C5-C6-N6	-5.83	119.03	123.70
84	Aa	2272	C	N3-C4-C5	-5.83	119.57	121.90
84	Aa	2421	C	N3-C4-C5	-5.83	119.57	121.90
84	Aa	2813	A	C5-C6-N1	-5.83	114.78	117.70
84	Aa	2837	C	N3-C4-C5	-5.83	119.57	121.90
86	Ab	74	A	C5-C6-N1	-5.83	114.78	117.70
1	Ad	58	U	O4'-C1'-N1	5.83	112.86	108.20
84	Aa	2733	A	C5-C6-N6	-5.83	119.04	123.70
84	Aa	196	A	C5-C6-N6	-5.83	119.04	123.70
84	Aa	1570	C	N3-C4-N4	5.83	122.08	118.00
84	Aa	1599	A	C5-C6-N6	-5.83	119.04	123.70
84	Aa	2120	A	C5-C6-N1	-5.83	114.78	117.70
84	Aa	2311	A	C4-C5-C6	5.83	119.91	117.00
84	Aa	2892	A	C5-C6-N1	-5.83	114.78	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3071	A	C5-C6-N1	-5.83	114.79	117.70
84	Aa	3334	A	O4'-C1'-N9	5.83	112.86	108.20
84	Aa	3374	C	C2-N1-C1'	5.83	125.21	118.80
86	Ab	108	G	N1-C2-N3	-5.83	120.40	123.90
1	Ad	1647	C	C1'-O4'-C4'	5.83	114.56	109.90
84	Aa	2140	C	N3-C4-C5	-5.83	119.57	121.90
84	Aa	2322	G	C5-C6-O6	-5.83	125.10	128.60
84	Aa	2344	A	C4-C5-C6	5.83	119.91	117.00
85	Ac	147	C	N3-C4-C5	-5.83	119.57	121.90
1	Ad	214	A	O4'-C1'-N9	5.83	112.86	108.20
1	Ad	1159	G	C3'-C2'-C1'	-5.83	96.84	101.50
84	Aa	1610	A	C5-C6-N1	-5.83	114.79	117.70
84	Aa	1974	C	N3-C4-C5	-5.83	119.57	121.90
84	Aa	2312	A	C5-C6-N1	-5.83	114.79	117.70
84	Aa	2704	U	O4'-C1'-N1	5.83	112.86	108.20
84	Aa	2895	G	O4'-C1'-N9	5.83	112.86	108.20
84	Aa	3045	A	C4-C5-C6	5.83	119.91	117.00
84	Aa	3078	A	C5-C6-N6	-5.83	119.04	123.70
84	Aa	569	C	N3-C4-N4	5.82	122.08	118.00
84	Aa	735	C	N3-C4-C5	-5.82	119.57	121.90
84	Aa	1129	G	O4'-C1'-N9	5.82	112.86	108.20
84	Aa	1483	G	C5-C6-O6	-5.82	125.11	128.60
84	Aa	2233	G	O4'-C1'-N9	5.82	112.86	108.20
84	Aa	2265	A	C4-C5-C6	5.82	119.91	117.00
84	Aa	2901	C	N3-C4-C5	-5.82	119.57	121.90
84	Aa	3174	C	N3-C4-C5	-5.82	119.57	121.90
84	Aa	3175	C	N3-C4-C5	-5.82	119.57	121.90
84	Aa	290	C	N3-C4-N4	5.82	122.08	118.00
84	Aa	912	G	C5-C6-O6	-5.82	125.11	128.60
84	Aa	1359	A	O4'-C1'-N9	5.82	112.86	108.20
84	Aa	2557	C	N3-C4-N4	5.82	122.08	118.00
84	Aa	2960	A	C4-C5-C6	5.82	119.91	117.00
84	Aa	3252	G	C5-C6-O6	-5.82	125.11	128.60
1	Ad	1414	G	O4'-C1'-N9	5.82	112.86	108.20
1	Ad	1528	U	O4'-C1'-C2'	-5.82	99.98	105.80
84	Aa	1330	A	C5-C6-N1	-5.82	114.79	117.70
84	Aa	1574	C	N3-C4-C5	-5.82	119.57	121.90
84	Aa	1753	A	C5-C6-N1	-5.82	114.79	117.70
84	Aa	2772	A	O4'-C1'-N9	5.82	112.86	108.20
85	Ac	71	A	C4-C5-C6	5.82	119.91	117.00
84	Aa	6	A	C4-C5-C6	5.82	119.91	117.00
84	Aa	447	C	N3-C4-C5	-5.82	119.57	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	628	C	N3-C4-C5	-5.82	119.57	121.90
84	Aa	1494	A	C5-C6-N6	-5.82	119.05	123.70
84	Aa	2761	A	C4-C5-C6	5.82	119.91	117.00
84	Aa	3159	C	N3-C4-C5	-5.82	119.57	121.90
84	Aa	144	A	C5-C6-N1	-5.82	114.79	117.70
84	Aa	642	C	P-O5'-C5'	5.82	130.21	120.90
84	Aa	1512	A	C5-C6-N6	-5.82	119.05	123.70
84	Aa	1651	A	C5-C6-N1	-5.82	114.79	117.70
84	Aa	2634	U	O4'-C1'-N1	5.82	112.85	108.20
85	Ac	44	A	C4-C5-C6	5.82	119.91	117.00
85	Ac	140	A	O4'-C1'-N9	5.82	112.85	108.20
86	Ab	11	A	C8-N9-C4	-5.82	103.47	105.80
1	Ad	507	G	O4'-C1'-N9	5.82	112.85	108.20
84	Aa	55	G	C5-C6-O6	-5.82	125.11	128.60
84	Aa	448	G	C5-C6-O6	-5.82	125.11	128.60
84	Aa	1323	G	C5-C6-O6	-5.82	125.11	128.60
84	Aa	1391	A	O4'-C1'-N9	5.82	112.85	108.20
84	Aa	1708	C	N3-C4-C5	-5.82	119.57	121.90
84	Aa	1973	C	N3-C4-C5	-5.82	119.57	121.90
84	Aa	2304	A	C4-C5-C6	5.82	119.91	117.00
84	Aa	2367	A	O4'-C1'-N9	5.82	112.85	108.20
84	Aa	3249	G	C5-C6-O6	-5.82	125.11	128.60
1	Ad	247	A	C5'-C4'-O4'	5.81	116.08	109.10
84	Aa	550	C	O3'-P-O5'	-5.81	92.95	104.00
84	Aa	971	G	O4'-C1'-N9	5.81	112.85	108.20
84	Aa	1396	A	C4-C5-C6	5.81	119.91	117.00
84	Aa	2016	A	C5-C6-N1	-5.81	114.79	117.70
84	Aa	2385	A	C5-C6-N6	-5.81	119.05	123.70
84	Aa	2554	U	O4'-C1'-N1	5.81	112.85	108.20
1	Ad	11	A	C1'-O4'-C4'	5.81	114.55	109.90
1	Ad	415	C	C5'-C4'-O4'	5.81	116.07	109.10
1	Ad	618	C	O4'-C1'-N1	5.81	112.85	108.20
1	Ad	1097	A	O4'-C1'-N9	5.81	112.85	108.20
1	Ad	1615	G	N9-C1'-C2'	5.81	121.56	114.00
24	BW	128	PHE	CB-CG-CD2	-5.81	116.73	120.80
84	Aa	672	A	C4-C5-C6	5.81	119.91	117.00
84	Aa	730	A	C4-C5-C6	5.81	119.91	117.00
84	Aa	1115	A	C4-C5-C6	5.81	119.91	117.00
84	Aa	1295	A	C4-C5-C6	5.81	119.91	117.00
84	Aa	1424	G	C5-C6-O6	-5.81	125.11	128.60
84	Aa	2113	A	C5-C6-N1	-5.81	114.79	117.70
86	Ab	64	G	C4-C5-C6	5.81	122.29	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	Ab	98	G	N3-C2-N2	5.81	123.97	119.90
1	Ad	1386	U	O4'-C1'-N1	5.81	112.85	108.20
48	CD	291	SER	N-CA-CB	5.81	119.22	110.50
84	Aa	851	A	C4-C5-C6	5.81	119.91	117.00
84	Aa	2178	G	P-O3'-C3'	5.81	126.67	119.70
84	Aa	2214	A	C4-C5-C6	5.81	119.91	117.00
84	Aa	2931	C	N3-C4-C5	-5.81	119.58	121.90
1	Ad	1497	U	C3'-C2'-C1'	-5.81	96.85	101.50
38	CT	19	PHE	CB-CG-CD2	-5.81	116.73	120.80
84	Aa	211	A	C5-C6-N6	-5.81	119.05	123.70
84	Aa	509	G	O4'-C1'-N9	5.81	112.85	108.20
84	Aa	532	G	C5-C6-O6	-5.81	125.11	128.60
84	Aa	607	U	N1-C1'-C2'	5.81	121.55	114.00
84	Aa	756	C	N3-C4-C5	-5.81	119.58	121.90
84	Aa	1617	A	C4-C5-C6	5.81	119.91	117.00
84	Aa	2872	C	N3-C4-N4	5.81	122.07	118.00
85	Ac	122	G	O4'-C1'-N9	5.81	112.85	108.20
1	Ad	1034	G	O4'-C1'-N9	5.81	112.85	108.20
1	Ad	1258	U	O4'-C1'-N1	5.81	112.85	108.20
84	Aa	17	G	O4'-C1'-N9	5.81	112.84	108.20
84	Aa	135	G	C5-C6-O6	-5.81	125.11	128.60
84	Aa	272	G	O4'-C1'-N9	5.81	112.85	108.20
84	Aa	823	A	C4-C5-C6	5.81	119.90	117.00
84	Aa	978	C	C2-N3-C4	5.81	122.80	119.90
84	Aa	1370	A	C5-C6-N6	-5.81	119.05	123.70
84	Aa	3060	G	C5-C6-O6	-5.81	125.11	128.60
84	Aa	3136	A	C5-C6-N6	-5.81	119.05	123.70
84	Aa	3210	G	C4-N9-C1'	5.81	134.05	126.50
48	CD	183	PHE	CB-CG-CD2	5.81	124.86	120.80
84	Aa	1256	A	C4-C5-C6	5.81	119.90	117.00
84	Aa	2025	C	N3-C4-C5	-5.81	119.58	121.90
84	Aa	2483	A	C4-C5-C6	5.81	119.90	117.00
84	Aa	3213	A	C4-C5-C6	5.81	119.90	117.00
1	Ad	310	U	C5'-C4'-O4'	5.80	116.06	109.10
1	Ad	493	C	O4'-C1'-N1	5.80	112.84	108.20
1	Ad	627	A	P-O3'-C3'	5.80	126.67	119.70
1	Ad	1388	A	N9-C1'-C2'	-5.80	105.62	112.00
48	CD	7	PHE	CB-CG-CD1	5.80	124.86	120.80
84	Aa	539	C	N3-C4-N4	5.80	122.06	118.00
84	Aa	610	G	P-O5'-C5'	5.80	130.19	120.90
84	Aa	715	A	C5-C6-N1	-5.80	114.80	117.70
84	Aa	1335	C	N3-C4-C5	-5.80	119.58	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1636	C	N3-C4-C5	-5.80	119.58	121.90
84	Aa	2574	A	C4-C5-C6	5.80	119.90	117.00
84	Aa	2734	C	N3-C4-N4	5.80	122.06	118.00
84	Aa	2835	A	C5-C6-N1	-5.80	114.80	117.70
84	Aa	2892	A	C4-C5-C6	5.80	119.90	117.00
84	Aa	2946	U	O4'-C1'-N1	5.80	112.84	108.20
84	Aa	3308	A	C4-C5-C6	5.80	119.90	117.00
85	Ac	77	A	C4-C5-C6	5.80	119.90	117.00
86	Ab	61	C	N3-C4-C5	-5.80	119.58	121.90
1	Ad	197	G	P-O3'-C3'	5.80	126.66	119.70
84	Aa	70	A	P-O3'-C3'	5.80	126.66	119.70
84	Aa	1380	C	N3-C4-C5	-5.80	119.58	121.90
84	Aa	1520	A	C4-C5-C6	5.80	119.90	117.00
84	Aa	2657	C	C2-N3-C4	5.80	122.80	119.90
84	Aa	3142	C	N3-C4-N4	5.80	122.06	118.00
24	BW	15	TYR	CB-CG-CD2	-5.80	117.52	121.00
84	Aa	875	A	C5-C6-N1	-5.80	114.80	117.70
84	Aa	2112	C	N3-C4-N4	5.80	122.06	118.00
84	Aa	2460	A	C5-C6-N1	-5.80	114.80	117.70
84	Aa	2736	A	C4-C5-C6	5.80	119.90	117.00
85	Ac	66	G	C5-C6-O6	-5.80	125.12	128.60
1	Ad	36	C	C3'-C2'-C1'	5.80	106.14	101.50
1	Ad	1214	C	P-O3'-C3'	-5.80	112.74	119.70
84	Aa	61	A	C5-C6-N6	-5.80	119.06	123.70
84	Aa	158	A	C5-C6-N1	-5.80	114.80	117.70
84	Aa	1156	A	C4-C5-C6	5.80	119.90	117.00
84	Aa	2386	A	C4-C5-C6	5.80	119.90	117.00
84	Aa	2629	C	N3-C4-N4	5.80	122.06	118.00
84	Aa	2658	U	O4'-C1'-N1	5.80	112.84	108.20
84	Aa	2662	A	C5-C6-N1	-5.80	114.80	117.70
84	Aa	3022	A	C4-C5-C6	5.80	119.90	117.00
1	Ad	1052	G	O4'-C1'-N9	5.80	112.84	108.20
78	CL	64	LYS	N-CA-CB	5.80	121.04	110.60
84	Aa	1932	A	O4'-C1'-N9	5.80	112.84	108.20
84	Aa	2273	C	N3-C4-C5	-5.80	119.58	121.90
85	Ac	106	C	N3-C4-N4	5.80	122.06	118.00
86	Ab	48	G	N3-C2-N2	5.80	123.96	119.90
1	Ad	966	U	C1'-O4'-C4'	-5.80	105.26	109.90
84	Aa	80	C	N3-C4-C5	-5.80	119.58	121.90
84	Aa	171	G	O4'-C1'-N9	5.80	112.84	108.20
84	Aa	555	G	O4'-C1'-N9	5.80	112.84	108.20
84	Aa	1643	A	C4-C5-C6	5.80	119.90	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2903	G	O4'-C1'-N9	5.80	112.84	108.20
84	Aa	2932	A	C5-C6-N1	-5.80	114.80	117.70
84	Aa	2934	C	P-O3'-C3'	5.80	126.66	119.70
84	Aa	3275	G	O4'-C1'-N9	5.80	112.84	108.20
1	Ad	83	U	N1-C1'-C2'	-5.79	105.62	112.00
1	Ad	547	C	P-O3'-C3'	5.79	126.65	119.70
84	Aa	793	C	N3-C4-C5	-5.79	119.58	121.90
84	Aa	1006	A	C4-C5-C6	5.79	119.90	117.00
84	Aa	1026	A	C5-C6-N1	-5.79	114.80	117.70
1	Ad	539	A	C1'-O4'-C4'	-5.79	105.27	109.90
84	Aa	480	C	N3-C4-C5	-5.79	119.58	121.90
84	Aa	920	A	C5-C6-N1	-5.79	114.80	117.70
84	Aa	3257	G	C5-C6-O6	-5.79	125.12	128.60
85	Ac	109	A	O4'-C1'-N9	5.79	112.84	108.20
85	Ac	158	C	N3-C4-N4	5.79	122.06	118.00
1	Ad	1283	C	O4'-C1'-N1	5.79	112.83	108.20
84	Aa	372	A	C4-C5-C6	5.79	119.90	117.00
84	Aa	493	G	C5-C6-O6	-5.79	125.12	128.60
84	Aa	708	C	N3-C4-N4	5.79	122.06	118.00
84	Aa	1061	A	C4-C5-C6	5.79	119.90	117.00
84	Aa	2671	A	C5-C6-N1	-5.79	114.80	117.70
84	Aa	2998	A	O4'-C1'-N9	5.79	112.83	108.20
1	Ad	104	A	P-O3'-C3'	5.79	126.65	119.70
1	Ad	496	A	P-O3'-C3'	-5.79	112.75	119.70
72	CC	336	TYR	CB-CG-CD2	5.79	124.47	121.00
84	Aa	1550	A	C4-C5-C6	5.79	119.89	117.00
86	Ab	82	G	C5-C6-N1	-5.79	108.61	111.50
1	Ad	1233	G	C3'-C2'-C1'	5.79	106.13	101.50
84	Aa	58	G	O4'-C1'-N9	5.79	112.83	108.20
84	Aa	693	C	N3-C4-C5	-5.79	119.58	121.90
84	Aa	954	A	C4-C5-C6	5.79	119.89	117.00
84	Aa	995	C	N3-C4-N4	5.79	122.05	118.00
84	Aa	1875	A	O4'-C1'-N9	5.79	112.83	108.20
84	Aa	2847	A	O4'-C1'-N9	5.79	112.83	108.20
84	Aa	3095	G	N3-C2-N2	5.79	123.95	119.90
86	Ab	68	G	C4-C5-C6	5.79	122.27	118.80
1	Ad	714	C	C3'-C2'-C1'	5.79	106.13	101.50
84	Aa	265	G	N3-C2-N2	5.79	123.95	119.90
84	Aa	492	G	C4'-C3'-C2'	5.79	108.39	102.60
84	Aa	907	A	C4-C5-C6	5.79	119.89	117.00
1	Ad	574	A	C5'-C4'-O4'	5.79	116.04	109.10
1	Ad	1705	C	C3'-C2'-C1'	5.79	106.13	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	267	G	C5-C6-O6	-5.79	125.13	128.60
84	Aa	393	A	C5-C6-N6	-5.79	119.07	123.70
84	Aa	1468	A	C4-C5-C6	5.79	119.89	117.00
84	Aa	1753	A	C5-C6-N6	-5.79	119.07	123.70
84	Aa	2678	C	N3-C4-C5	-5.79	119.59	121.90
84	Aa	2836	G	N3-C2-N2	5.79	123.95	119.90
84	Aa	2850	G	O4'-C1'-N9	5.79	112.83	108.20
84	Aa	3250	C	N3-C4-C5	-5.79	119.59	121.90
84	Aa	3251	C	N3-C4-C5	-5.79	119.59	121.90
1	Ad	131	C	C4'-C3'-C2'	-5.78	96.82	102.60
1	Ad	1192	G	C1'-O4'-C4'	-5.78	105.27	109.90
1	Ad	1755	G	C1'-O4'-C4'	-5.78	105.27	109.90
84	Aa	181	G	O4'-C1'-N9	5.78	112.83	108.20
84	Aa	2487	A	C5-C6-N1	-5.78	114.81	117.70
84	Aa	2672	C	N3-C4-C5	-5.78	119.59	121.90
84	Aa	3358	A	C4-C5-C6	5.78	119.89	117.00
84	Aa	932	A	C5-C6-N1	-5.78	114.81	117.70
84	Aa	2961	C	N3-C4-C5	-5.78	119.59	121.90
85	Ac	110	A	C5-C6-N6	-5.78	119.08	123.70
1	Ad	285	G	N9-C1'-C2'	5.78	121.51	114.00
1	Ad	1573	C	O4'-C1'-N1	5.78	112.82	108.20
61	CM	89	TRP	N-CA-CB	5.78	121.00	110.60
84	Aa	224	C	N3-C4-C5	-5.78	119.59	121.90
84	Aa	1087	G	O4'-C1'-N9	5.78	112.82	108.20
84	Aa	1132	A	C4-C5-C6	5.78	119.89	117.00
84	Aa	1581	C	N3-C4-C5	-5.78	119.59	121.90
84	Aa	1855	A	C5-C6-N1	-5.78	114.81	117.70
84	Aa	2177	U	C2'-C3'-O3'	5.78	122.95	113.70
84	Aa	2509	A	C4-C5-C6	5.78	119.89	117.00
84	Aa	2822	A	C5-C6-N1	-5.78	114.81	117.70
84	Aa	3100	C	N3-C4-C5	-5.78	119.59	121.90
1	Ad	344	U	C1'-O4'-C4'	5.78	114.52	109.90
1	Ad	1597	C	O4'-C1'-N1	5.78	112.82	108.20
84	Aa	1890	C	N3-C4-C5	-5.78	119.59	121.90
84	Aa	1943	G	P-O3'-C3'	-5.78	112.77	119.70
84	Aa	2237	A	C4-C5-C6	5.78	119.89	117.00
84	Aa	2321	C	N3-C4-C5	-5.78	119.59	121.90
86	Ab	75	G	C3'-C2'-C1'	-5.78	96.88	101.50
1	Ad	860	A	P-O5'-C5'	-5.78	111.66	120.90
1	Ad	949	A	O4'-C1'-C2'	-5.78	100.02	105.80
1	Ad	1254	U	C1'-O4'-C4'	5.78	114.52	109.90
41	CA	67	PHE	CB-CG-CD1	-5.78	116.76	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	120	G	O4'-C1'-N9	5.78	112.82	108.20
84	Aa	1418	C	N3-C4-N4	5.78	122.04	118.00
84	Aa	1599	A	C4-C5-C6	5.78	119.89	117.00
84	Aa	1875	A	C4-C5-C6	5.78	119.89	117.00
84	Aa	2119	A	C4-C5-C6	5.78	119.89	117.00
84	Aa	2250	A	C5-C6-N1	-5.78	114.81	117.70
84	Aa	2456	G	C5-C6-O6	-5.78	125.13	128.60
84	Aa	2792	A	C5-C6-N1	-5.78	114.81	117.70
84	Aa	565	C	C2-N3-C4	5.78	122.79	119.90
84	Aa	805	C	N3-C4-C5	-5.78	119.59	121.90
84	Aa	1203	C	N3-C4-C5	-5.78	119.59	121.90
84	Aa	1208	A	C4-C5-C6	5.78	119.89	117.00
84	Aa	1778	C	N3-C4-C5	-5.78	119.59	121.90
84	Aa	2757	G	N1-C6-O6	5.78	123.36	119.90
84	Aa	3017	A	C5-C6-N6	-5.78	119.08	123.70
84	Aa	3212	C	N3-C4-N4	5.78	122.04	118.00
84	Aa	3358	A	C5-C6-N1	-5.78	114.81	117.70
1	Ad	10	G	O4'-C1'-N9	5.77	112.82	108.20
1	Ad	312	C	C3'-C2'-C1'	5.77	106.12	101.50
84	Aa	2288	C	O4'-C1'-N1	5.77	112.82	108.20
84	Aa	2400	A	O4'-C1'-N9	5.77	112.82	108.20
85	Ac	135	A	C4-C5-C6	5.77	119.89	117.00
84	Aa	26	A	O4'-C1'-N9	5.77	112.82	108.20
84	Aa	232	C	N3-C4-C5	-5.77	119.59	121.90
84	Aa	354	C	N3-C4-N4	5.77	122.04	118.00
84	Aa	2210	A	C5-C6-N6	-5.77	119.08	123.70
84	Aa	2347	A	O4'-C1'-N9	5.77	112.82	108.20
84	Aa	2847	A	C4-C5-C6	5.77	119.89	117.00
84	Aa	2880	G	O4'-C1'-N9	5.77	112.82	108.20
86	Ab	65	G	N1-C2-N3	-5.77	120.44	123.90
84	Aa	160	G	O4'-C1'-N9	5.77	112.82	108.20
84	Aa	189	C	N3-C4-C5	-5.77	119.59	121.90
84	Aa	1016	G	C5-C6-O6	-5.77	125.14	128.60
84	Aa	1291	A	C5-C6-N1	-5.77	114.81	117.70
84	Aa	1492	A	C4-C5-C6	5.77	119.89	117.00
84	Aa	1635	A	C5-C6-N1	-5.77	114.81	117.70
84	Aa	2896	C	N3-C4-N4	5.77	122.04	118.00
84	Aa	3140	A	C5-C6-N1	-5.77	114.81	117.70
1	Ad	586	U	O4'-C1'-N1	5.77	112.81	108.20
2	Ae	68	C	N1-C1'-C2'	5.77	121.50	114.00
84	Aa	197	A	C5-C6-N6	-5.77	119.08	123.70
84	Aa	1490	A	C5-C6-N1	-5.77	114.81	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2973	A	C4-C5-C6	5.77	119.89	117.00
84	Aa	3206	C	N3-C4-C5	-5.77	119.59	121.90
84	Aa	3302	A	O4'-C1'-N9	5.77	112.82	108.20
86	Ab	34	C	C2-N3-C4	5.77	122.78	119.90
1	Ad	206	U	P-O3'-C3'	5.77	126.62	119.70
1	Ad	961	U	O4'-C1'-N1	5.77	112.81	108.20
34	BC	146	ASN	N-CA-CB	5.77	120.98	110.60
84	Aa	545	C	N3-C4-C5	-5.77	119.59	121.90
84	Aa	871	C	N3-C4-C5	-5.77	119.59	121.90
84	Aa	1061	A	C5-C6-N1	-5.77	114.82	117.70
84	Aa	1361	G	O4'-C1'-N9	5.77	112.81	108.20
84	Aa	1472	C	N3-C4-C5	-5.77	119.59	121.90
84	Aa	1755	A	C5-C6-N6	-5.77	119.08	123.70
84	Aa	2623	G	O4'-C1'-N9	5.77	112.81	108.20
84	Aa	3272	A	C5-C6-N6	-5.77	119.09	123.70
84	Aa	3288	A	O4'-C1'-N9	5.77	112.81	108.20
1	Ad	1132	G	O4'-C1'-N9	5.77	112.81	108.20
1	Ad	1341	G	P-O3'-C3'	5.77	126.62	119.70
84	Aa	147	G	C5-C6-O6	-5.77	125.14	128.60
84	Aa	873	A	C4-C5-C6	5.77	119.88	117.00
84	Aa	1305	A	C5-C6-N6	-5.77	119.09	123.70
84	Aa	2092	C	C3'-C2'-C1'	-5.77	96.89	101.50
84	Aa	2244	G	O4'-C1'-N9	5.77	112.81	108.20
1	Ad	646	G	O4'-C1'-C2'	5.76	112.79	107.60
1	Ad	1352	A	O4'-C1'-N9	5.76	112.81	108.20
1	Ad	1369	C	N1-C1'-C2'	5.76	121.49	114.00
1	Ad	1502	C	O4'-C1'-C2'	-5.76	100.03	105.80
1	Ad	1507	G	C1'-O4'-C4'	-5.76	105.29	109.90
84	Aa	821	C	O4'-C1'-N1	5.76	112.81	108.20
84	Aa	2243	C	N3-C4-C5	-5.76	119.59	121.90
84	Aa	826	C	N3-C4-C5	-5.76	119.59	121.90
84	Aa	2238	A	C4-C5-C6	5.76	119.88	117.00
84	Aa	2442	A	C4-C5-C6	5.76	119.88	117.00
1	Ad	965	U	O4'-C1'-C2'	-5.76	100.04	105.80
1	Ad	1005	C	C5'-C4'-O4'	5.76	116.01	109.10
48	CD	122	GLY	N-CA-C	-5.76	98.70	113.10
84	Aa	3	G	C5-C6-O6	-5.76	125.14	128.60
84	Aa	1460	U	O4'-C1'-N1	5.76	112.81	108.20
84	Aa	1491	G	C5-C6-O6	-5.76	125.14	128.60
84	Aa	1577	A	C4-C5-C6	5.76	119.88	117.00
84	Aa	2439	A	P-O5'-C5'	-5.76	111.68	120.90
84	Aa	2928	A	C4-C5-C6	5.76	119.88	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2932	A	C4-C5-C6	5.76	119.88	117.00
84	Aa	3199	C	N3-C4-C5	-5.76	119.60	121.90
1	Ad	1687	G	C3'-C2'-C1'	5.76	106.11	101.50
2	Ae	10	G	C1'-O4'-C4'	5.76	114.51	109.90
56	Cd	87	ARG	N-CA-CB	5.76	120.97	110.60
84	Aa	73	A	C5-C6-N6	-5.76	119.09	123.70
84	Aa	976	A	O4'-C1'-N9	5.76	112.81	108.20
84	Aa	1059	A	O4'-C1'-N9	5.76	112.81	108.20
84	Aa	1885	G	C5-C6-O6	-5.76	125.14	128.60
84	Aa	2571	C	N3-C4-C5	-5.76	119.60	121.90
84	Aa	3104	A	C5-C6-N6	-5.76	119.09	123.70
84	Aa	38	A	O4'-C1'-N9	5.76	112.81	108.20
84	Aa	917	A	C5-C6-N6	-5.76	119.09	123.70
84	Aa	1477	A	C4-C5-C6	5.76	119.88	117.00
84	Aa	1622	G	O4'-C1'-N9	5.76	112.81	108.20
84	Aa	2699	A	C4-C5-C6	5.76	119.88	117.00
84	Aa	2834	C	N3-C4-C5	-5.76	119.60	121.90
1	Ad	1250	C	O4'-C1'-C2'	5.76	112.78	107.60
84	Aa	93	G	O4'-C1'-N9	5.76	112.81	108.20
84	Aa	654	C	N3-C4-C5	-5.76	119.60	121.90
84	Aa	905	G	O4'-C1'-N9	5.76	112.81	108.20
84	Aa	1190	C	N3-C4-C5	-5.76	119.60	121.90
84	Aa	1412	C	N3-C4-C5	-5.76	119.60	121.90
84	Aa	1848	G	O4'-C1'-N9	5.76	112.81	108.20
84	Aa	1887	A	O4'-C1'-N9	5.76	112.81	108.20
84	Aa	1926	A	C5-C6-N6	-5.76	119.09	123.70
84	Aa	2137	A	C4-C5-C6	5.76	119.88	117.00
85	Ac	11	C	C4'-C3'-C2'	-5.76	96.84	102.60
85	Ac	104	A	C4-C5-C6	5.76	119.88	117.00
33	BJ	85	TYR	CB-CG-CD1	5.75	124.45	121.00
71	CB	351	SER	N-CA-CB	5.75	119.13	110.50
84	Aa	1062	G	O4'-C1'-N9	5.75	112.80	108.20
84	Aa	2041	G	O4'-C1'-N9	5.75	112.80	108.20
84	Aa	2081	C	O4'-C1'-N1	5.75	112.80	108.20
84	Aa	2125	A	C4-C5-C6	5.75	119.88	117.00
84	Aa	3037	G	O4'-C1'-N9	5.75	112.80	108.20
86	Ab	90	A	O4'-C1'-N9	5.75	112.80	108.20
1	Ad	1091	A	N9-C1'-C2'	5.75	121.48	114.00
84	Aa	659	C	N3-C4-C5	-5.75	119.60	121.90
84	Aa	1838	A	C4-C5-C6	5.75	119.88	117.00
84	Aa	1908	C	N3-C4-N4	5.75	122.03	118.00
84	Aa	2365	C	N3-C4-C5	-5.75	119.60	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2794	A	C4-C5-C6	5.75	119.88	117.00
84	Aa	2799	U	O4'-C1'-N1	5.75	112.80	108.20
84	Aa	3165	C	N3-C4-C5	-5.75	119.60	121.90
84	Aa	3235	A	C5-C6-N6	-5.75	119.10	123.70
84	Aa	3360	U	O4'-C1'-N1	5.75	112.80	108.20
85	Ac	96	A	C4-C5-C6	5.75	119.88	117.00
34	BC	235	PHE	N-CA-CB	5.75	120.95	110.60
84	Aa	886	A	C5-C6-N1	-5.75	114.83	117.70
84	Aa	1256	A	C5-C6-N6	-5.75	119.10	123.70
84	Aa	1748	A	C5-C6-N6	-5.75	119.10	123.70
84	Aa	1771	G	C5-C6-O6	-5.75	125.15	128.60
84	Aa	3368	A	C4-C5-C6	5.75	119.88	117.00
86	Ab	3	A	C5-C6-N6	-5.75	119.10	123.70
1	Ad	1155	G	N9-C1'-C2'	5.75	121.47	114.00
84	Aa	1114	A	C5-C6-N1	-5.75	114.83	117.70
84	Aa	1450	G	C5'-C4'-O4'	5.75	116.00	109.10
84	Aa	2786	G	O4'-C1'-N9	5.75	112.80	108.20
1	Ad	419	C	O4'-C1'-N1	5.75	112.80	108.20
1	Ad	1802	G	O4'-C1'-C2'	5.75	112.77	107.60
11	BD	184	ILE	N-CA-CB	5.75	124.02	110.80
14	BQ	135	PHE	CB-CG-CD2	5.75	124.82	120.80
33	BJ	9	SER	N-CA-CB	5.75	119.12	110.50
84	Aa	338	C	N3-C4-C5	-5.75	119.60	121.90
84	Aa	383	A	C5-C6-N6	-5.75	119.10	123.70
84	Aa	760	C	N3-C4-C5	-5.75	119.60	121.90
84	Aa	873	A	C5-C6-N6	-5.75	119.10	123.70
84	Aa	1882	A	C5-C6-N6	-5.75	119.10	123.70
84	Aa	2215	A	C4-C5-C6	5.75	119.88	117.00
84	Aa	3193	C	N3-C4-N4	5.75	122.02	118.00
1	Ad	559	A	O4'-C1'-C2'	-5.75	100.05	105.80
84	Aa	818	G	C5-C6-O6	-5.75	125.15	128.60
84	Aa	1378	G	C5-C6-O6	-5.75	125.15	128.60
84	Aa	2124	G	O4'-C1'-N9	5.75	112.80	108.20
84	Aa	2223	A	C5-C6-N1	-5.75	114.83	117.70
84	Aa	2598	A	C4-C5-C6	5.75	119.87	117.00
84	Aa	19	C	O5'-P-OP2	-5.75	100.53	105.70
84	Aa	976	A	C5-C6-N1	-5.75	114.83	117.70
84	Aa	1872	C	N3-C4-N4	5.75	122.02	118.00
84	Aa	2280	C	N3-C4-N4	5.75	122.02	118.00
1	Ad	518	G	O4'-C1'-N9	5.74	112.80	108.20
1	Ad	936	C	O4'-C1'-C2'	-5.74	100.06	105.80
84	Aa	216	G	O4'-C1'-N9	5.74	112.80	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1259	C	N3-C4-N4	5.74	122.02	118.00
84	Aa	1404	G	C5-C6-O6	-5.74	125.15	128.60
84	Aa	1628	G	C5-C6-O6	-5.74	125.15	128.60
84	Aa	2596	A	C4-C5-C6	5.74	119.87	117.00
84	Aa	3101	C	N3-C4-C5	-5.74	119.60	121.90
84	Aa	970	A	C4-C5-C6	5.74	119.87	117.00
84	Aa	1414	C	N3-C4-C5	-5.74	119.60	121.90
84	Aa	1694	A	C4-C5-C6	5.74	119.87	117.00
84	Aa	2140	C	N3-C4-N4	5.74	122.02	118.00
1	Ad	1074	C	O4'-C1'-C2'	-5.74	100.06	105.80
84	Aa	120	G	C5-C6-O6	-5.74	125.16	128.60
84	Aa	164	C	N3-C4-N4	5.74	122.02	118.00
84	Aa	246	C	N3-C4-C5	-5.74	119.60	121.90
84	Aa	813	A	O4'-C1'-N9	5.74	112.79	108.20
84	Aa	2000	C	N3-C4-C5	-5.74	119.60	121.90
84	Aa	2018	C	N3-C4-C5	-5.74	119.60	121.90
84	Aa	2033	C	N3-C4-C5	-5.74	119.60	121.90
84	Aa	2049	C	N3-C4-C5	-5.74	119.60	121.90
84	Aa	2288	C	N3-C4-C5	-5.74	119.60	121.90
84	Aa	2442	A	C5-C6-N1	-5.74	114.83	117.70
84	Aa	2593	A	C5-C6-N1	-5.74	114.83	117.70
84	Aa	2971	A	C4-C5-C6	5.74	119.87	117.00
1	Ad	217	A	P-O5'-C5'	5.74	130.08	120.90
84	Aa	233	C	N3-C4-C5	-5.74	119.60	121.90
84	Aa	715	A	O4'-C1'-N9	5.74	112.79	108.20
84	Aa	1018	C	N3-C4-N4	5.74	122.02	118.00
84	Aa	1667	C	N3-C4-C5	-5.74	119.61	121.90
84	Aa	2107	A	C5-C6-N6	-5.74	119.11	123.70
84	Aa	2133	A	C4-C5-C6	5.74	119.87	117.00
84	Aa	2223	A	O4'-C1'-N9	5.74	112.79	108.20
84	Aa	2676	A	C4-C5-C6	5.74	119.87	117.00
84	Aa	2920	G	C5-C6-O6	-5.74	125.16	128.60
84	Aa	3359	C	N3-C4-C5	-5.74	119.61	121.90
1	Ad	1083	C	C3'-C2'-C1'	5.74	106.09	101.50
1	Ad	1220	C	C3'-C2'-C1'	5.74	106.09	101.50
48	CD	238	SER	N-CA-CB	5.74	119.11	110.50
84	Aa	1994	C	N3-C4-C5	-5.74	119.61	121.90
84	Aa	2971	A	C5-C6-N1	-5.74	114.83	117.70
84	Aa	3375	G	O4'-C1'-N9	5.74	112.79	108.20
1	Ad	1159	G	P-O3'-C3'	-5.74	112.82	119.70
1	Ad	1699	C	C3'-C2'-C1'	5.74	106.09	101.50
84	Aa	2168	C	N3-C4-C5	-5.74	119.61	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2788	A	C4-C5-C6	5.74	119.87	117.00
84	Aa	2928	A	O4'-C1'-N9	5.74	112.79	108.20
84	Aa	363	A	O4'-C1'-N9	5.73	112.79	108.20
84	Aa	505	G	O4'-C1'-N9	5.73	112.79	108.20
84	Aa	1354	G	O4'-C1'-N9	5.73	112.79	108.20
86	Ab	34	C	N3-C4-C5	-5.73	119.61	121.90
1	Ad	554	A	O4'-C1'-N9	5.73	112.79	108.20
1	Ad	910	A	O4'-C1'-N9	5.73	112.79	108.20
84	Aa	200	G	O4'-C1'-N9	5.73	112.78	108.20
84	Aa	1066	G	O4'-C1'-N9	5.73	112.79	108.20
84	Aa	1206	A	O4'-C1'-N9	5.73	112.78	108.20
84	Aa	2215	A	C5-C6-N1	-5.73	114.83	117.70
84	Aa	2224	A	C4-C5-C6	5.73	119.87	117.00
84	Aa	2619	C	N3-C4-N4	5.73	122.01	118.00
84	Aa	2757	G	O4'-C1'-N9	5.73	112.79	108.20
1	Ad	1193	A	C3'-C2'-C1'	5.73	106.08	101.50
1	Ad	1451	G	N9-C1'-C2'	5.73	121.45	114.00
84	Aa	293	A	C5-C6-N1	-5.73	114.83	117.70
84	Aa	1193	A	C5-C6-N6	-5.73	119.11	123.70
84	Aa	1835	A	C4-C5-C6	5.73	119.86	117.00
84	Aa	1857	G	C5-C6-O6	-5.73	125.16	128.60
84	Aa	2088	C	C5'-C4'-C3'	5.73	125.17	116.00
84	Aa	2218	A	C5-C6-N6	-5.73	119.12	123.70
84	Aa	3306	A	C5-C6-N6	-5.73	119.12	123.70
84	Aa	3389	C	N3-C4-C5	-5.73	119.61	121.90
1	Ad	281	U	C4'-C3'-C2'	-5.73	96.87	102.60
85	Ac	89	A	C4-C5-C6	5.73	119.86	117.00
1	Ad	827	C	O4'-C1'-C2'	-5.73	100.07	105.80
84	Aa	346	A	O4'-C1'-N9	5.73	112.78	108.20
84	Aa	1508	C	N3-C4-C5	-5.73	119.61	121.90
84	Aa	1782	G	O4'-C1'-N9	5.73	112.78	108.20
84	Aa	2603	C	N3-C4-C5	-5.73	119.61	121.90
84	Aa	2673	G	C5-C6-O6	-5.73	125.16	128.60
84	Aa	3162	C	N3-C4-N4	5.73	122.01	118.00
1	Ad	64	U	C4'-C3'-C2'	-5.73	96.87	102.60
84	Aa	16	A	C4-C5-C6	5.73	119.86	117.00
84	Aa	1290	A	C5-C6-N1	-5.73	114.84	117.70
1	Ad	1125	U	O4'-C1'-C2'	-5.72	100.08	105.80
1	Ad	1591	A	O4'-C1'-C2'	-5.72	100.08	105.80
84	Aa	848	G	O4'-C1'-N9	5.72	112.78	108.20
86	Ab	2	G	N9-C4-C5	-5.72	103.11	105.40
1	Ad	174	C	C5'-C4'-O4'	5.72	115.97	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	760	G	O4'-C1'-N9	5.72	112.78	108.20
84	Aa	813	A	C4-C5-C6	5.72	119.86	117.00
84	Aa	1588	G	O4'-C1'-N9	5.72	112.78	108.20
84	Aa	1969	G	O4'-C1'-N9	5.72	112.78	108.20
84	Aa	2790	C	N3-C4-C5	-5.72	119.61	121.90
84	Aa	2881	C	N3-C4-C5	-5.72	119.61	121.90
84	Aa	3206	C	N3-C4-N4	5.72	122.01	118.00
84	Aa	3322	A	C4-C5-C6	5.72	119.86	117.00
85	Ac	129	C	N3-C4-C5	-5.72	119.61	121.90
84	Aa	1444	G	C5-C6-O6	-5.72	125.17	128.60
84	Aa	2504	A	C4-C5-C6	5.72	119.86	117.00
84	Aa	2992	G	O4'-C1'-N9	5.72	112.78	108.20
84	Aa	3087	A	C5-C6-N6	-5.72	119.12	123.70
1	Ad	232	C	C3'-C2'-C1'	5.72	106.08	101.50
1	Ad	491	G	N9-C1'-C2'	-5.72	105.71	112.00
84	Aa	1375	G	O4'-C1'-N9	5.72	112.78	108.20
84	Aa	2043	A	C4'-C3'-O3'	5.72	124.44	113.00
84	Aa	2694	A	C5-C6-N6	-5.72	119.12	123.70
84	Aa	3061	C	N3-C4-C5	-5.72	119.61	121.90
84	Aa	757	G	C5-C6-O6	-5.72	125.17	128.60
84	Aa	1342	C	N3-C4-C5	-5.72	119.61	121.90
84	Aa	1584	A	C4-C5-C6	5.72	119.86	117.00
84	Aa	2133	A	O4'-C1'-N9	5.72	112.77	108.20
84	Aa	3059	C	N3-C4-N4	5.72	122.00	118.00
84	Aa	3293	U	O4'-C1'-N1	5.72	112.78	108.20
1	Ad	488	C	O4'-C1'-C2'	-5.72	100.08	105.80
1	Ad	1488	C	C3'-C2'-C1'	5.72	106.07	101.50
84	Aa	159	G	O4'-C1'-N9	5.72	112.77	108.20
84	Aa	1208	A	O4'-C1'-N9	5.72	112.77	108.20
84	Aa	1511	C	N3-C4-C5	-5.72	119.61	121.90
84	Aa	1571	A	C4-C5-C6	5.72	119.86	117.00
84	Aa	2941	G	C5-C6-O6	-5.72	125.17	128.60
85	Ac	44	A	C5-C6-N6	-5.72	119.13	123.70
1	Ad	490	G	N9-C1'-C2'	5.71	121.43	114.00
1	Ad	978	A	O4'-C1'-N9	5.71	112.77	108.20
84	Aa	136	C	N3-C4-C5	-5.71	119.61	121.90
84	Aa	1236	C	N3-C4-C5	-5.71	119.61	121.90
84	Aa	2325	A	C4-C5-C6	5.71	119.86	117.00
84	Aa	2612	A	C5-C6-N1	-5.71	114.84	117.70
85	Ac	65	G	P-O5'-C5'	-5.71	111.76	120.90
1	Ad	250	A	O4'-C1'-C2'	-5.71	100.09	105.80
1	Ad	372	U	C1'-O4'-C4'	5.71	114.47	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	325	A	O4'-C1'-N9	5.71	112.77	108.20
84	Aa	667	C	N3-C4-C5	-5.71	119.61	121.90
84	Aa	1693	A	C5-C6-N1	-5.71	114.84	117.70
86	Ab	3	A	C4-C5-C6	5.71	119.86	117.00
1	Ad	397	C	C3'-C2'-C1'	5.71	106.07	101.50
1	Ad	431	C	C3'-C2'-C1'	5.71	106.07	101.50
1	Ad	561	G	C3'-C2'-C1'	5.71	106.07	101.50
84	Aa	105	A	C5-C6-N1	-5.71	114.84	117.70
84	Aa	604	C	C6-N1-C2	-5.71	118.02	120.30
84	Aa	1374	G	O4'-C1'-N9	5.71	112.77	108.20
84	Aa	2742	A	C4-C5-C6	5.71	119.86	117.00
85	Ac	156	C	N3-C4-C5	-5.71	119.61	121.90
1	Ad	742	C	P-O3'-C3'	5.71	126.55	119.70
84	Aa	2963	G	O4'-C1'-N9	5.71	112.77	108.20
85	Ac	123	G	C5-C6-O6	-5.71	125.17	128.60
86	Ab	42	A	O4'-C1'-N9	5.71	112.77	108.20
84	Aa	755	C	N3-C4-C5	-5.71	119.62	121.90
84	Aa	928	A	C4-C5-C6	5.71	119.86	117.00
84	Aa	977	G	C5-C6-O6	-5.71	125.17	128.60
84	Aa	1591	A	C4-C5-C6	5.71	119.85	117.00
84	Aa	1767	G	O4'-C1'-N9	5.71	112.77	108.20
84	Aa	2767	C	N3-C4-N4	5.71	122.00	118.00
84	Aa	2792	A	O4'-C1'-N9	5.71	112.77	108.20
84	Aa	2916	G	C5-C6-O6	-5.71	125.17	128.60
84	Aa	3362	A	O4'-C1'-N9	5.71	112.77	108.20
1	Ad	405	A	O4'-C1'-C2'	-5.71	100.09	105.80
1	Ad	1088	G	C4'-C3'-C2'	-5.71	96.89	102.60
84	Aa	546	C	N3-C4-C5	-5.71	119.62	121.90
84	Aa	939	A	C5-C6-N1	-5.71	114.85	117.70
84	Aa	1122	C	N3-C4-C5	-5.71	119.62	121.90
84	Aa	1862	C	N3-C4-N4	5.71	122.00	118.00
84	Aa	2401	A	C5-C6-N1	-5.71	114.85	117.70
84	Aa	2449	A	C5-C6-N1	-5.71	114.85	117.70
84	Aa	2488	A	C4-C5-C6	5.71	119.85	117.00
84	Aa	2929	C	N3-C4-N4	5.71	121.99	118.00
84	Aa	3114	A	C5'-C4'-O4'	5.71	115.95	109.10
84	Aa	16	A	O4'-C1'-N9	5.71	112.76	108.20
84	Aa	738	A	C5-C6-N6	-5.71	119.14	123.70
84	Aa	2301	C	C2-N1-C1'	5.71	125.08	118.80
84	Aa	2640	A	C5-C6-N1	-5.71	114.85	117.70
84	Aa	2753	C	C5'-C4'-C3'	5.71	125.13	116.00
1	Ad	637	U	C1'-O4'-C4'	5.70	114.46	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1116	G	C3'-C2'-C1'	-5.70	96.94	101.50
1	Ad	1221	A	P-O3'-C3'	5.70	126.54	119.70
1	Ad	1665	U	O4'-C1'-N1	5.70	112.76	108.20
50	CP	31	GLU	N-CA-CB	5.70	120.86	110.60
84	Aa	670	A	C4-C5-C6	5.70	119.85	117.00
84	Aa	1101	A	C4-C5-C6	5.70	119.85	117.00
84	Aa	1150	G	C5-C6-O6	-5.70	125.18	128.60
84	Aa	1905	A	C4-C5-C6	5.70	119.85	117.00
84	Aa	1962	C	N3-C4-C5	-5.70	119.62	121.90
84	Aa	2361	C	N3-C4-C5	-5.70	119.62	121.90
84	Aa	3154	G	C1'-O4'-C4'	-5.70	105.34	109.90
84	Aa	3290	C	N3-C4-C5	-5.70	119.62	121.90
84	Aa	1151	G	C5-C6-O6	-5.70	125.18	128.60
84	Aa	1477	A	O4'-C1'-N9	5.70	112.76	108.20
84	Aa	2032	C	N3-C4-N4	5.70	121.99	118.00
84	Aa	2056	C	N3-C4-N4	5.70	121.99	118.00
1	Ad	1097	A	O4'-C1'-C2'	5.70	112.73	107.60
45	CN	30	TYR	CB-CG-CD1	5.70	124.42	121.00
84	Aa	201	G	O4'-C1'-N9	5.70	112.76	108.20
84	Aa	1123	A	C5-C6-N6	-5.70	119.14	123.70
84	Aa	1610	A	C4-C5-C6	5.70	119.85	117.00
84	Aa	1863	A	C5-C6-N6	-5.70	119.14	123.70
84	Aa	2080	G	P-O3'-C3'	5.70	126.54	119.70
84	Aa	2230	C	N3-C4-C5	-5.70	119.62	121.90
84	Aa	2356	A	C5-C6-N6	-5.70	119.14	123.70
84	Aa	2978	A	C5-C6-N6	-5.70	119.14	123.70
1	Ad	575	G	C1'-O4'-C4'	5.70	114.46	109.90
84	Aa	7	C	N3-C4-C5	-5.70	119.62	121.90
84	Aa	394	A	C5-C6-N6	-5.70	119.14	123.70
84	Aa	450	C	N3-C4-C5	-5.70	119.62	121.90
84	Aa	766	C	N3-C4-C5	-5.70	119.62	121.90
84	Aa	873	A	O4'-C1'-N9	5.70	112.76	108.20
84	Aa	1227	A	C5-C6-N1	-5.70	114.85	117.70
84	Aa	1409	G	O4'-C1'-N9	5.70	112.76	108.20
84	Aa	1906	A	C4-C5-C6	5.70	119.85	117.00
84	Aa	3209	U	P-O3'-C3'	5.70	126.54	119.70
1	Ad	1600	G	C3'-C2'-C1'	5.70	106.06	101.50
84	Aa	1217	G	O4'-C1'-N9	5.70	112.76	108.20
84	Aa	1501	A	O4'-C1'-N9	5.70	112.76	108.20
84	Aa	1907	A	C4-C5-C6	5.70	119.85	117.00
84	Aa	1930	G	C5-C6-O6	-5.70	125.18	128.60
84	Aa	3110	A	C5-C6-N1	-5.70	114.85	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3149	C	N3-C4-C5	-5.70	119.62	121.90
86	Ab	50	A	C5-N7-C8	5.70	106.75	103.90
1	Ad	1426	C	O4'-C1'-C2'	-5.70	100.10	105.80
17	BS	83	PHE	CB-CG-CD1	-5.70	116.81	120.80
42	CJ	1	MET	C-N-CA	5.70	135.94	121.70
78	CL	48	ARG	N-CA-CB	5.70	120.85	110.60
84	Aa	102	G	C5-C6-O6	-5.70	125.18	128.60
84	Aa	420	A	C5-C6-N1	-5.70	114.85	117.70
84	Aa	617	C	N3-C4-C5	-5.70	119.62	121.90
84	Aa	1392	U	P-O5'-C5'	-5.70	111.79	120.90
84	Aa	2079	A	C4-C5-C6	5.70	119.85	117.00
84	Aa	2594	A	C5-C6-N1	-5.70	114.85	117.70
84	Aa	292	A	C5-C6-N1	-5.69	114.85	117.70
84	Aa	1267	A	C4-C5-C6	5.69	119.85	117.00
84	Aa	1395	A	C5-C6-N6	-5.69	119.14	123.70
84	Aa	1837	A	C4-C5-C6	5.69	119.85	117.00
84	Aa	3027	G	O4'-C1'-N9	5.69	112.75	108.20
84	Aa	3034	A	C5-C6-N6	-5.69	119.14	123.70
84	Aa	3288	A	C4-C5-C6	5.69	119.85	117.00
84	Aa	257	C	N3-C4-N4	5.69	121.98	118.00
84	Aa	377	C	N3-C4-C5	-5.69	119.62	121.90
84	Aa	389	A	C4-C5-C6	5.69	119.85	117.00
84	Aa	679	C	N3-C4-C5	-5.69	119.62	121.90
84	Aa	709	G	C5-C6-O6	-5.69	125.18	128.60
84	Aa	1146	A	P-O5'-C5'	-5.69	111.79	120.90
84	Aa	1197	A	C4-C5-C6	5.69	119.85	117.00
84	Aa	1485	A	C5-C6-N6	-5.69	119.15	123.70
84	Aa	1715	C	N3-C4-C5	-5.69	119.62	121.90
84	Aa	1842	C	N3-C4-N4	5.69	121.98	118.00
84	Aa	1904	A	O4'-C1'-N9	5.69	112.75	108.20
84	Aa	2639	A	C4-C5-C6	5.69	119.85	117.00
84	Aa	2804	A	C5-C6-N6	-5.69	119.14	123.70
61	CM	17	TYR	N-CA-CB	5.69	120.84	110.60
84	Aa	331	G	C5-C6-O6	-5.69	125.19	128.60
84	Aa	615	A	C4-C5-C6	5.69	119.85	117.00
84	Aa	690	G	O4'-C1'-N9	5.69	112.75	108.20
84	Aa	1038	C	N3-C4-C5	-5.69	119.62	121.90
84	Aa	1434	G	C5-C6-O6	-5.69	125.19	128.60
84	Aa	2239	A	C5-C6-N1	-5.69	114.86	117.70
84	Aa	3054	G	C5-C6-O6	-5.69	125.19	128.60
85	Ac	121	A	C4-C5-C6	5.69	119.84	117.00
84	Aa	702	G	O4'-C1'-N9	5.69	112.75	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1593	C	N3-C4-C5	-5.69	119.62	121.90
84	Aa	2817	G	N1-C6-O6	5.69	123.31	119.90
1	Ad	71	C	O4'-C1'-C2'	-5.69	100.11	105.80
84	Aa	269	C	N3-C4-N4	5.69	121.98	118.00
84	Aa	416	A	O4'-C1'-N9	5.69	112.75	108.20
84	Aa	812	G	O4'-C1'-N9	5.69	112.75	108.20
84	Aa	907	A	C5-C6-N1	-5.69	114.86	117.70
84	Aa	1799	C	N3-C4-C5	-5.69	119.62	121.90
84	Aa	1914	C	N3-C4-C5	-5.69	119.62	121.90
84	Aa	2743	A	C5-C6-N1	-5.69	114.86	117.70
84	Aa	2815	A	C4-C5-C6	5.69	119.84	117.00
84	Aa	2975	G	C5-C6-O6	-5.69	125.19	128.60
84	Aa	3187	C	N3-C4-C5	-5.69	119.62	121.90
85	Ac	49	G	O4'-C1'-N9	5.69	112.75	108.20
84	Aa	337	C	N3-C4-N4	5.69	121.98	118.00
84	Aa	1136	A	C5-C6-N1	-5.69	114.86	117.70
84	Aa	2224	A	O4'-C1'-N9	5.69	112.75	108.20
84	Aa	2671	A	O4'-C1'-N9	5.69	112.75	108.20
84	Aa	2819	A	C5-C6-N1	-5.69	114.86	117.70
84	Aa	3385	G	C5-C6-O6	-5.69	125.19	128.60
84	Aa	1509	G	C5-C6-O6	-5.68	125.19	128.60
84	Aa	2089	A	C5-C6-N6	-5.68	119.15	123.70
84	Aa	2190	C	N3-C4-C5	-5.68	119.63	121.90
1	Ad	912	A	O4'-C1'-N9	5.68	112.75	108.20
1	Ad	1374	G	O4'-C1'-N9	5.68	112.75	108.20
84	Aa	228	C	N3-C4-N4	5.68	121.98	118.00
84	Aa	294	A	O4'-C1'-N9	5.68	112.75	108.20
84	Aa	3045	A	C5-C6-N1	-5.68	114.86	117.70
86	Ab	74	A	O4'-C1'-N9	5.68	112.75	108.20
84	Aa	210	G	O4'-C1'-N9	5.68	112.75	108.20
84	Aa	2564	G	O4'-C1'-N9	5.68	112.75	108.20
84	Aa	2575	C	N3-C4-N4	5.68	121.98	118.00
84	Aa	3264	C	N3-C4-C5	-5.68	119.63	121.90
1	Ad	801	U	C3'-C2'-C1'	-5.68	96.96	101.50
84	Aa	889	C	N3-C4-C5	-5.68	119.63	121.90
84	Aa	1249	A	C5-C6-N1	-5.68	114.86	117.70
84	Aa	1953	C	N3-C4-N4	5.68	121.98	118.00
84	Aa	1990	A	C4-C5-C6	5.68	119.84	117.00
84	Aa	2592	G	C5-C6-O6	-5.68	125.19	128.60
84	Aa	2703	G	C5-C6-O6	-5.68	125.19	128.60
84	Aa	2718	A	C5-C6-N6	-5.68	119.16	123.70
84	Aa	2868	C	N3-C4-N4	5.68	121.97	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3272	A	C4-C5-C6	5.68	119.84	117.00
50	CP	4	TYR	CB-CG-CD1	5.68	124.41	121.00
84	Aa	22	G	O4'-C1'-N9	5.68	112.74	108.20
84	Aa	1789	C	N3-C4-N4	5.68	121.97	118.00
84	Aa	3364	A	C5-C6-N1	-5.68	114.86	117.70
84	Aa	24	C	N3-C4-C5	-5.68	119.63	121.90
84	Aa	3067	G	O4'-C1'-N9	5.68	112.74	108.20
2	Ae	47	U	O4'-C1'-C2'	-5.67	100.13	105.80
71	CB	373	ARG	N-CA-CB	5.67	120.82	110.60
84	Aa	364	A	C5-C6-N6	-5.67	119.16	123.70
84	Aa	449	G	O4'-C1'-N9	5.67	112.74	108.20
84	Aa	893	C	N3-C4-C5	-5.67	119.63	121.90
84	Aa	1118	G	C4-N9-C1'	5.67	133.88	126.50
84	Aa	1134	G	O4'-C1'-N9	5.67	112.74	108.20
84	Aa	1793	A	C5-C6-N6	-5.67	119.16	123.70
84	Aa	1883	A	O4'-C1'-N9	5.67	112.74	108.20
84	Aa	2217	A	C5-C6-N6	-5.67	119.16	123.70
85	Ac	140	A	C4-C5-C6	5.67	119.84	117.00
84	Aa	235	G	O4'-C1'-N9	5.67	112.74	108.20
84	Aa	361	G	C5-C6-O6	-5.67	125.20	128.60
84	Aa	363	A	C5-C6-N1	-5.67	114.86	117.70
84	Aa	2175	A	C5-C6-N1	-5.67	114.86	117.70
84	Aa	2489	A	C5-C6-N6	-5.67	119.16	123.70
84	Aa	2655	U	C5'-C4'-C3'	-5.67	106.92	116.00
84	Aa	3078	A	C4-C5-C6	5.67	119.84	117.00
1	Ad	1598	G	C5'-C4'-O4'	5.67	115.91	109.10
84	Aa	999	U	O4'-C1'-N1	5.67	112.74	108.20
84	Aa	1872	C	N3-C4-C5	-5.67	119.63	121.90
85	Ac	52	A	C4-C5-C6	5.67	119.84	117.00
86	Ab	15	C	N3-C4-C5	-5.67	119.63	121.90
84	Aa	1570	C	N3-C4-C5	-5.67	119.63	121.90
84	Aa	2650	A	C4-C5-C6	5.67	119.83	117.00
84	Aa	2761	A	C5-C6-N6	-5.67	119.16	123.70
1	Ad	815	A	O4'-C1'-N9	5.67	112.73	108.20
84	Aa	211	A	O4'-C1'-N9	5.67	112.73	108.20
84	Aa	698	A	C4-C5-C6	5.67	119.83	117.00
84	Aa	1156	A	C5-C6-N1	-5.67	114.86	117.70
84	Aa	1254	A	O4'-C1'-N9	5.67	112.73	108.20
84	Aa	2320	A	O4'-C1'-N9	5.67	112.73	108.20
84	Aa	3152	C	N3-C4-N4	5.67	121.97	118.00
1	Ad	847	U	C5'-C4'-O4'	5.67	115.90	109.10
84	Aa	578	C	O4'-C1'-N1	5.67	112.73	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	661	A	O4'-C1'-N9	5.67	112.73	108.20
84	Aa	975	G	C5-C6-O6	-5.67	125.20	128.60
84	Aa	1926	A	C4-C5-C6	5.67	119.83	117.00
84	Aa	2214	A	C5-C6-N6	-5.67	119.17	123.70
84	Aa	2724	A	C5-C6-N1	-5.67	114.87	117.70
84	Aa	2807	G	C5-C6-O6	-5.67	125.20	128.60
84	Aa	2942	A	C5-C6-N1	-5.67	114.87	117.70
84	Aa	3122	U	P-O3'-C3'	5.67	126.50	119.70
1	Ad	1226	U	C1'-O4'-C4'	5.67	114.43	109.90
84	Aa	437	C	N3-C4-C5	-5.67	119.63	121.90
84	Aa	758	A	C4-C5-C6	5.67	119.83	117.00
84	Aa	883	G	O4'-C1'-N9	5.67	112.73	108.20
1	Ad	289	G	O4'-C1'-C2'	5.66	112.70	107.60
1	Ad	936	C	N1-C1'-C2'	5.66	121.36	114.00
1	Ad	1753	U	N1-C1'-C2'	-5.66	105.77	112.00
84	Aa	71	C	N3-C4-N4	5.66	121.96	118.00
84	Aa	393	A	O4'-C1'-N9	5.66	112.73	108.20
84	Aa	1332	C	N3-C4-C5	-5.66	119.64	121.90
84	Aa	1397	A	C5-C6-N1	-5.66	114.87	117.70
84	Aa	1496	G	O4'-C1'-N9	5.66	112.73	108.20
84	Aa	1543	A	C5-C6-N1	-5.66	114.87	117.70
84	Aa	1883	A	C5-C6-N1	-5.66	114.87	117.70
84	Aa	2247	A	C5-C6-N6	-5.66	119.17	123.70
84	Aa	2458	A	O4'-C1'-N9	5.66	112.73	108.20
85	Ac	21	C	N3-C4-C5	-5.66	119.63	121.90
84	Aa	1306	A	C5-C6-N6	-5.66	119.17	123.70
84	Aa	1587	G	C5-C6-O6	-5.66	125.20	128.60
1	Ad	548	C	O4'-C1'-N1	5.66	112.73	108.20
84	Aa	563	C	N3-C4-N4	5.66	121.96	118.00
84	Aa	1501	A	C5-C6-N1	-5.66	114.87	117.70
84	Aa	1906	A	O4'-C1'-N9	5.66	112.73	108.20
84	Aa	1918	A	C4-C5-C6	5.66	119.83	117.00
84	Aa	1971	A	O4'-C1'-N9	5.66	112.73	108.20
1	Ad	297	U	C1'-O4'-C4'	-5.66	105.37	109.90
84	Aa	1643	A	C5-C6-N6	-5.66	119.17	123.70
84	Aa	1794	A	O4'-C1'-N9	5.66	112.73	108.20
84	Aa	1800	G	C5-C6-O6	-5.66	125.20	128.60
84	Aa	2347	A	C4-C5-C6	5.66	119.83	117.00
84	Aa	2518	A	C5-C6-N6	-5.66	119.17	123.70
84	Aa	2899	A	O4'-C1'-N9	5.66	112.73	108.20
84	Aa	2975	G	O4'-C1'-N9	5.66	112.73	108.20
85	Ac	61	A	C4-C5-C6	5.66	119.83	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1727	A	C5-C6-N6	-5.66	119.17	123.70
84	Aa	1971	A	C4-C5-C6	5.66	119.83	117.00
84	Aa	2330	C	N3-C4-C5	-5.66	119.64	121.90
84	Aa	2367	A	C4-C5-C6	5.66	119.83	117.00
84	Aa	3162	C	N3-C4-C5	-5.66	119.64	121.90
1	Ad	1768	U	C1'-O4'-C4'	-5.66	105.38	109.90
84	Aa	66	A	C4-C5-C6	5.66	119.83	117.00
84	Aa	371	A	C5-C6-N1	-5.66	114.87	117.70
84	Aa	413	G	O4'-C1'-N9	5.66	112.72	108.20
84	Aa	907	A	O4'-C1'-N9	5.66	112.72	108.20
84	Aa	1001	A	C4-C5-C6	5.66	119.83	117.00
84	Aa	1026	A	C4-C5-C6	5.66	119.83	117.00
84	Aa	1255	A	C5-C6-N1	-5.66	114.87	117.70
84	Aa	1318	C	C5-C6-N1	5.66	123.83	121.00
1	Ad	1688	G	C1'-O4'-C4'	5.65	114.42	109.90
84	Aa	39	A	C4-C5-C6	5.65	119.83	117.00
84	Aa	1576	C	N3-C4-C5	-5.65	119.64	121.90
84	Aa	2427	C	N3-C4-N4	5.65	121.96	118.00
1	Ad	1258	U	C5'-C4'-O4'	5.65	115.88	109.10
79	CE	124	TYR	CB-CG-CD1	5.65	124.39	121.00
84	Aa	203	C	N3-C4-C5	-5.65	119.64	121.90
84	Aa	465	C	N3-C4-C5	-5.65	119.64	121.90
84	Aa	2462	G	C5'-C4'-C3'	5.65	125.05	116.00
84	Aa	2574	A	C5-C6-N6	-5.65	119.18	123.70
84	Aa	2700	A	O4'-C1'-N9	5.65	112.72	108.20
84	Aa	3104	A	C5-C6-N1	-5.65	114.87	117.70
84	Aa	3196	C	N3-C4-C5	-5.65	119.64	121.90
1	Ad	108	C	C1'-O4'-C4'	-5.65	105.38	109.90
1	Ad	336	U	C1'-O4'-C4'	5.65	114.42	109.90
1	Ad	1614	C	C1'-O4'-C4'	-5.65	105.38	109.90
84	Aa	1562	A	O5'-C5'-C4'	-5.65	100.97	111.70
84	Aa	1879	A	C4-C5-C6	5.65	119.83	117.00
84	Aa	1949	G	O4'-C1'-N9	5.65	112.72	108.20
84	Aa	2058	C	N3-C4-C5	-5.65	119.64	121.90
84	Aa	2217	A	C5-C6-N1	-5.65	114.88	117.70
84	Aa	2228	A	C5-C6-N6	-5.65	119.18	123.70
84	Aa	2819	A	C4-C5-C6	5.65	119.83	117.00
84	Aa	3362	A	C5-C6-N1	-5.65	114.88	117.70
85	Ac	109	A	C5-C6-N6	-5.65	119.18	123.70
55	Cc	102	SER	N-CA-CB	5.65	118.97	110.50
84	Aa	1248	A	C4-C5-C6	5.65	119.83	117.00
84	Aa	2114	A	C5-C6-N6	-5.65	119.18	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2372	A	C4-C5-C6	5.65	119.82	117.00
84	Aa	3020	C	N3-C4-C5	-5.65	119.64	121.90
1	Ad	595	A	C1'-O4'-C4'	5.65	114.42	109.90
1	Ad	624	A	O4'-C1'-N9	5.65	112.72	108.20
79	CE	50	PHE	CB-CG-CD2	-5.65	116.85	120.80
84	Aa	1321	A	C4-C5-C6	5.65	119.82	117.00
84	Aa	1642	G	C5-C6-O6	-5.65	125.21	128.60
84	Aa	1680	A	C4-C5-C6	5.65	119.82	117.00
84	Aa	1775	C	N3-C4-N4	5.65	121.95	118.00
84	Aa	2272	C	N3-C4-N4	5.65	121.95	118.00
84	Aa	2805	A	C5-C6-N6	-5.65	119.18	123.70
84	Aa	3107	A	C5-C6-N6	-5.65	119.18	123.70
85	Ac	14	C	N3-C4-C5	-5.65	119.64	121.90
85	Ac	111	G	O4'-C1'-N9	5.65	112.72	108.20
1	Ad	158	C	C4'-C3'-C2'	-5.65	96.95	102.60
1	Ad	1294	U	O4'-C1'-C2'	-5.65	100.15	105.80
1	Ad	1768	U	O4'-C1'-N1	5.65	112.72	108.20
1	Ad	1801	A	O4'-C1'-C2'	-5.65	100.15	105.80
1	Ad	216	A	O4'-C1'-C2'	5.64	112.68	107.60
1	Ad	979	A	O4'-C1'-N9	5.64	112.72	108.20
84	Aa	112	C	N3-C4-N4	5.64	121.95	118.00
84	Aa	579	G	O4'-C1'-N9	5.64	112.72	108.20
84	Aa	788	G	O4'-C1'-N9	5.64	112.72	108.20
84	Aa	846	A	C4-C5-C6	5.64	119.82	117.00
84	Aa	1100	G	O4'-C1'-N9	5.64	112.72	108.20
84	Aa	1204	A	C5-C6-N6	-5.64	119.19	123.70
84	Aa	1304	G	O4'-C1'-N9	5.64	112.72	108.20
84	Aa	1633	C	N3-C4-N4	5.64	121.95	118.00
84	Aa	2014	A	C5-C6-N6	-5.64	119.18	123.70
84	Aa	2724	A	C5-C6-N6	-5.64	119.18	123.70
84	Aa	2814	C	N3-C4-C5	-5.64	119.64	121.90
84	Aa	3346	C	N3-C4-N4	5.64	121.95	118.00
1	Ad	923	U	C3'-C2'-C1'	5.64	106.01	101.50
84	Aa	219	A	C5-C6-N6	-5.64	119.19	123.70
84	Aa	863	G	C5-C6-O6	-5.64	125.22	128.60
84	Aa	1917	A	O4'-C1'-N9	5.64	112.71	108.20
84	Aa	2909	A	O4'-C1'-N9	5.64	112.72	108.20
84	Aa	3269	C	N3-C4-C5	-5.64	119.64	121.90
86	Ab	94	C	C6-N1-C2	5.64	122.56	120.30
48	CD	199	ILE	N-CA-CB	5.64	123.78	110.80
84	Aa	558	G	C5-C6-O6	-5.64	125.22	128.60
84	Aa	886	A	O4'-C1'-N9	5.64	112.71	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1318	C	C6-N1-C2	-5.64	118.04	120.30
1	Ad	205	U	N1-C1'-C2'	-5.64	105.80	112.00
84	Aa	108	A	P-O3'-C3'	5.64	126.47	119.70
84	Aa	923	A	C4-C5-C6	5.64	119.82	117.00
84	Aa	1015	A	C4-C5-C6	5.64	119.82	117.00
84	Aa	2371	A	C5-C6-N1	-5.64	114.88	117.70
84	Aa	2492	C	N3-C4-N4	5.64	121.95	118.00
86	Ab	36	C	N3-C4-N4	5.64	121.95	118.00
1	Ad	820	A	O4'-C1'-C2'	5.64	112.67	107.60
84	Aa	2304	A	O4'-C1'-N9	5.64	112.71	108.20
84	Aa	2598	A	C5-C6-N1	-5.64	114.88	117.70
85	Ac	97	G	O4'-C1'-N9	5.64	112.71	108.20
1	Ad	203	A	O4'-C1'-N9	5.64	112.71	108.20
1	Ad	329	G	C4'-C3'-C2'	-5.64	96.96	102.60
84	Aa	560	C	N3-C4-N4	5.64	121.95	118.00
84	Aa	631	C	N3-C4-C5	-5.64	119.64	121.90
84	Aa	1084	G	O4'-C1'-N9	5.64	112.71	108.20
84	Aa	1160	G	C5-C6-O6	-5.64	125.22	128.60
84	Aa	1551	C	N3-C4-N4	5.64	121.95	118.00
84	Aa	2088	C	O4'-C4'-C3'	-5.64	98.36	104.00
84	Aa	2853	A	C5-C6-N6	-5.64	119.19	123.70
84	Aa	3022	A	C5-C6-N6	-5.64	119.19	123.70
1	Ad	465	G	C3'-C2'-C1'	5.63	106.01	101.50
1	Ad	507	G	O4'-C1'-C2'	5.63	112.67	107.60
1	Ad	1675	G	N9-C1'-C2'	-5.63	105.80	112.00
84	Aa	446	C	N3-C4-C5	-5.63	119.65	121.90
84	Aa	502	G	O4'-C1'-N9	5.63	112.71	108.20
84	Aa	563	C	N3-C4-C5	-5.63	119.65	121.90
84	Aa	1109	G	C5-C6-O6	-5.63	125.22	128.60
84	Aa	1329	G	C5-C6-O6	-5.63	125.22	128.60
84	Aa	1534	C	N3-C4-N4	5.63	121.94	118.00
84	Aa	1714	A	C5-C6-N1	-5.63	114.88	117.70
84	Aa	1934	U	O4'-C1'-N1	5.63	112.71	108.20
84	Aa	2756	G	O4'-C1'-N9	5.63	112.71	108.20
84	Aa	1860	A	C4-C5-C6	5.63	119.82	117.00
84	Aa	1861	A	O4'-C1'-N9	5.63	112.71	108.20
84	Aa	3034	A	O4'-C1'-N9	5.63	112.71	108.20
1	Ad	732	G	N9-C1'-C2'	-5.63	105.81	112.00
1	Ad	1159	G	C5'-C4'-O4'	5.63	115.86	109.10
84	Aa	633	C	N3-C4-N4	5.63	121.94	118.00
84	Aa	1182	A	C4-C5-C6	5.63	119.82	117.00
84	Aa	1206	A	C5-C6-N1	-5.63	114.88	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1221	A	C5-C6-N1	-5.63	114.89	117.70
84	Aa	1784	C	N3-C4-C5	-5.63	119.65	121.90
84	Aa	3077	C	N3-C4-N4	5.63	121.94	118.00
86	Ab	78	C	C5-C6-N1	5.63	123.82	121.00
1	Ad	276	G	C4'-C3'-C2'	-5.63	96.97	102.60
84	Aa	10	C	N3-C4-C5	-5.63	119.65	121.90
84	Aa	60	G	O4'-C1'-N9	5.63	112.70	108.20
84	Aa	291	C	N3-C4-C5	-5.63	119.65	121.90
84	Aa	880	C	N3-C4-N4	5.63	121.94	118.00
84	Aa	1732	G	C5-C6-O6	-5.63	125.22	128.60
84	Aa	1900	C	N3-C4-N4	5.63	121.94	118.00
84	Aa	2223	A	C5-C6-N6	-5.63	119.20	123.70
24	BW	15	TYR	CB-CG-CD1	5.63	124.38	121.00
84	Aa	222	C	O4'-C1'-N1	5.63	112.70	108.20
84	Aa	317	G	O4'-C1'-N9	5.63	112.70	108.20
84	Aa	669	G	C5-C6-O6	-5.63	125.22	128.60
84	Aa	919	G	O4'-C1'-N9	5.63	112.70	108.20
84	Aa	2040	G	C5-C6-O6	-5.63	125.22	128.60
84	Aa	2741	G	O4'-C1'-N9	5.63	112.70	108.20
84	Aa	3327	A	O4'-C1'-N9	5.63	112.70	108.20
1	Ad	141	G	N9-C1'-C2'	5.63	121.32	114.00
22	BZ	34	LYS	N-CA-CB	5.63	120.73	110.60
84	Aa	1079	G	N3-C2-N2	5.63	123.84	119.90
84	Aa	1542	A	O4'-C1'-N9	5.63	112.70	108.20
84	Aa	2173	G	C4-N9-C1'	-5.63	119.19	126.50
84	Aa	2203	A	C5-C6-N6	-5.63	119.20	123.70
84	Aa	2629	C	N3-C4-C5	-5.63	119.65	121.90
84	Aa	2740	C	N3-C4-N4	5.63	121.94	118.00
84	Aa	3079	G	P-O3'-C3'	5.63	126.45	119.70
84	Aa	3128	A	C5-C6-N1	-5.63	114.89	117.70
80	Cf	1	MET	CG-SD-CE	-5.62	91.20	100.20
84	Aa	651	A	C4-C5-C6	5.62	119.81	117.00
84	Aa	2058	C	N3-C4-N4	5.62	121.94	118.00
85	Ac	129	C	O4'-C1'-N1	5.62	112.70	108.20
41	CA	119	HIS	N-CA-CB	5.62	120.72	110.60
84	Aa	2743	A	O4'-C1'-N9	5.62	112.70	108.20
85	Ac	33	A	C5-C6-N1	-5.62	114.89	117.70
1	Ad	249	G	C1'-O4'-C4'	-5.62	105.40	109.90
1	Ad	448	C	C1'-O4'-C4'	5.62	114.40	109.90
1	Ad	1053	C	O4'-C1'-N1	5.62	112.70	108.20
1	Ad	1272	G	O4'-C1'-N9	5.62	112.70	108.20
1	Ad	1569	U	C1'-O4'-C4'	5.62	114.40	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1785	U	O4'-C1'-N1	5.62	112.70	108.20
84	Aa	230	G	C5-C6-O6	-5.62	125.23	128.60
84	Aa	232	C	N3-C4-N4	5.62	121.94	118.00
84	Aa	687	C	N3-C4-C5	-5.62	119.65	121.90
84	Aa	1051	A	C4-C5-C6	5.62	119.81	117.00
84	Aa	1099	G	N1-C6-O6	5.62	123.27	119.90
84	Aa	1742	G	C5-C6-O6	-5.62	125.23	128.60
84	Aa	1846	A	C5-C6-N6	-5.62	119.20	123.70
84	Aa	2306	G	O4'-C1'-N9	5.62	112.70	108.20
84	Aa	3141	G	O4'-C1'-N9	5.62	112.70	108.20
84	Aa	3339	G	C5-C6-O6	-5.62	125.23	128.60
86	Ab	1	G	C4-C5-C6	5.62	122.17	118.80
1	Ad	187	C	O4'-C1'-C2'	-5.62	100.18	105.80
60	Co	59	HIS	N-CA-CB	5.62	120.72	110.60
84	Aa	58	G	C5-C6-O6	-5.62	125.23	128.60
84	Aa	79	C	N3-C4-C5	-5.62	119.65	121.90
84	Aa	2414	C	N3-C4-C5	-5.62	119.65	121.90
84	Aa	3173	A	C5-C6-N6	-5.62	119.20	123.70
1	Ad	488	C	P-O3'-C3'	5.62	126.44	119.70
84	Aa	681	A	C5-C6-N6	-5.62	119.20	123.70
84	Aa	2239	A	C5-C6-N6	-5.62	119.20	123.70
84	Aa	2633	C	N3-C4-C5	-5.62	119.65	121.90
84	Aa	3171	C	C2-N1-C1'	5.62	124.98	118.80
85	Ac	154	G	P-O5'-C5'	5.62	129.89	120.90
84	Aa	298	G	O4'-C1'-N9	5.62	112.69	108.20
84	Aa	1182	A	O4'-C1'-N9	5.62	112.69	108.20
84	Aa	1651	A	C5-C6-N6	-5.62	119.21	123.70
84	Aa	1932	A	C4-C5-C6	5.62	119.81	117.00
84	Aa	2254	A	O4'-C1'-N9	5.62	112.69	108.20
84	Aa	2294	A	C5-C6-N1	-5.62	114.89	117.70
1	Ad	1780	U	C1'-O4'-C4'	5.62	114.39	109.90
70	Cq	140	PHE	CB-CG-CD2	-5.62	116.87	120.80
84	Aa	1065	A	C4-C5-C6	5.62	119.81	117.00
84	Aa	1274	A	O4'-C1'-N9	5.62	112.69	108.20
84	Aa	2750	A	O4'-C1'-N9	5.62	112.69	108.20
84	Aa	2945	G	O4'-C1'-N9	5.62	112.69	108.20
86	Ab	74	A	C5-C6-N6	-5.62	119.21	123.70
1	Ad	970	U	O4'-C1'-N1	5.61	112.69	108.20
1	Ad	1013	G	O4'-C1'-C2'	5.61	112.65	107.60
71	CB	353	LEU	N-CA-C	-5.61	95.84	111.00
84	Aa	198	A	C4-C5-C6	5.61	119.81	117.00
84	Aa	238	C	N3-C4-N4	5.61	121.93	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	854	C	N3-C4-N4	5.61	121.93	118.00
84	Aa	913	G	C5-C6-O6	-5.61	125.23	128.60
84	Aa	1028	G	O4'-C1'-N9	5.61	112.69	108.20
84	Aa	1143	G	O4'-C1'-N9	5.61	112.69	108.20
84	Aa	1260	G	O4'-C1'-N9	5.61	112.69	108.20
84	Aa	1385	C	N3-C4-N4	5.61	121.93	118.00
84	Aa	2119	A	O4'-C1'-N9	5.61	112.69	108.20
84	Aa	2297	G	C5-C6-O6	-5.61	125.23	128.60
84	Aa	3268	C	O4'-C1'-N1	5.61	112.69	108.20
85	Ac	90	C	N3-C4-N4	5.61	121.93	118.00
84	Aa	696	A	C5-C6-N6	-5.61	119.21	123.70
84	Aa	1104	C	N3-C4-N4	5.61	121.93	118.00
84	Aa	1343	C	O4'-C1'-N1	5.61	112.69	108.20
84	Aa	1511	C	N3-C4-N4	5.61	121.93	118.00
84	Aa	2436	G	O4'-C1'-N9	5.61	112.69	108.20
1	Ad	523	C	C3'-C2'-C1'	5.61	105.99	101.50
1	Ad	639	G	O4'-C1'-N9	5.61	112.69	108.20
84	Aa	382	A	C5-C6-N6	-5.61	119.21	123.70
84	Aa	928	A	C5-C6-N6	-5.61	119.21	123.70
84	Aa	1174	G	N3-C2-N2	5.61	123.83	119.90
84	Aa	1436	A	C5-C6-N1	-5.61	114.89	117.70
84	Aa	1937	C	N3-C4-N4	5.61	121.93	118.00
84	Aa	2219	A	O4'-C1'-N9	5.61	112.69	108.20
84	Aa	2800	C	N3-C4-N4	5.61	121.93	118.00
84	Aa	3018	A	C5-C6-N1	-5.61	114.89	117.70
84	Aa	3018	A	C5-C6-N6	-5.61	119.21	123.70
84	Aa	3169	C	N3-C4-N4	5.61	121.93	118.00
84	Aa	3276	G	C5-C6-O6	-5.61	125.23	128.60
1	Ad	1005	C	N1-C1'-C2'	-5.61	105.83	112.00
1	Ad	1385	C	O3'-P-O5'	-5.61	93.34	104.00
1	Ad	1516	C	N1-C1'-C2'	5.61	121.29	114.00
84	Aa	34	G	C5-C6-O6	-5.61	125.23	128.60
84	Aa	1462	C	N3-C4-C5	-5.61	119.66	121.90
85	Ac	158	C	N3-C4-C5	-5.61	119.66	121.90
84	Aa	714	G	O4'-C1'-N9	5.61	112.69	108.20
84	Aa	1146	A	C4-C5-C6	5.61	119.80	117.00
84	Aa	1443	G	O4'-C1'-N9	5.61	112.69	108.20
84	Aa	1772	G	O4'-C1'-N9	5.61	112.69	108.20
84	Aa	1809	A	C5-C6-N6	-5.61	119.21	123.70
84	Aa	3110	A	C5-C6-N6	-5.61	119.21	123.70
86	Ab	97	G	C8-N9-C4	-5.61	104.16	106.40
84	Aa	253	G	N1-C2-N3	-5.61	120.54	123.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1252	C	N3-C4-C5	-5.61	119.66	121.90
84	Aa	1353	A	C5-C6-N6	-5.61	119.22	123.70
84	Aa	2589	G	O4'-C1'-N9	5.61	112.68	108.20
84	Aa	3056	C	O4'-C1'-N1	5.61	112.69	108.20
84	Aa	3074	A	C5-C6-N6	-5.61	119.22	123.70
85	Ac	33	A	C4-C5-C6	5.61	119.80	117.00
1	Ad	923	U	P-O3'-C3'	5.60	126.42	119.70
22	BZ	18	SER	C-N-CA	5.60	134.07	122.30
84	Aa	741	G	O4'-C1'-N9	5.60	112.68	108.20
84	Aa	862	G	O4'-C1'-N9	5.60	112.68	108.20
84	Aa	1820	C	N3-C4-N4	5.60	121.92	118.00
85	Ac	88	A	C5-C6-N1	-5.60	114.90	117.70
86	Ab	31	G	C5-C6-N1	-5.60	108.70	111.50
1	Ad	283	G	P-O3'-C3'	5.60	126.42	119.70
37	CG	233	VAL	N-CA-CB	5.60	123.83	111.50
84	Aa	411	C	N3-C4-C5	-5.60	119.66	121.90
84	Aa	457	C	N3-C4-N4	5.60	121.92	118.00
84	Aa	608	G	P-O5'-C5'	5.60	129.86	120.90
84	Aa	633	C	N3-C4-C5	-5.60	119.66	121.90
84	Aa	899	A	C4-C5-C6	5.60	119.80	117.00
84	Aa	1813	C	N3-C4-C5	-5.60	119.66	121.90
84	Aa	2458	A	C5-C6-N6	-5.60	119.22	123.70
86	Ab	41	G	N3-C2-N2	5.60	123.82	119.90
1	Ad	333	G	O4'-C1'-C2'	5.60	112.64	107.60
23	Bc	2	ASP	N-CA-CB	5.60	120.68	110.60
84	Aa	23	A	C5-C6-N6	-5.60	119.22	123.70
84	Aa	99	A	C5-C6-N1	-5.60	114.90	117.70
84	Aa	229	G	N3-C2-N2	5.60	123.82	119.90
84	Aa	574	C	N3-C4-C5	-5.60	119.66	121.90
84	Aa	1353	A	O4'-C1'-N9	5.60	112.68	108.20
84	Aa	1527	A	C5-C6-N6	-5.60	119.22	123.70
56	Cd	102	THR	N-CA-CB	5.60	120.94	110.30
84	Aa	1200	A	C5-C6-N6	-5.60	119.22	123.70
84	Aa	1790	A	O4'-C1'-N9	5.60	112.68	108.20
84	Aa	2141	A	C5-C6-N1	-5.60	114.90	117.70
84	Aa	2533	A	C5-C6-N6	-5.60	119.22	123.70
84	Aa	3232	C	N3-C4-N4	5.60	121.92	118.00
84	Aa	3353	G	O4'-C1'-N9	5.60	112.68	108.20
84	Aa	3382	A	O4'-C1'-N9	5.60	112.68	108.20
1	Ad	71	C	O4'-C1'-N1	5.60	112.68	108.20
24	BW	65	LEU	N-CA-CB	5.60	121.59	110.40
25	Bd	33	LYS	N-CA-CB	5.60	120.67	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	306	A	C5-C6-N1	-5.60	114.90	117.70
84	Aa	467	C	N3-C4-N4	5.60	121.92	118.00
84	Aa	1112	C	C2-N3-C4	5.60	122.70	119.90
84	Aa	1788	C	O4'-C1'-N1	5.60	112.68	108.20
84	Aa	2390	G	O4'-C1'-N9	5.60	112.68	108.20
84	Aa	2544	C	N3-C4-N4	5.60	121.92	118.00
84	Aa	2938	A	C4-C5-C6	5.60	119.80	117.00
84	Aa	738	A	O4'-C1'-N9	5.60	112.68	108.20
84	Aa	1962	C	N3-C4-N4	5.60	121.92	118.00
84	Aa	2230	C	N3-C4-N4	5.60	121.92	118.00
84	Aa	3034	A	C5-C6-N1	-5.60	114.90	117.70
1	Ad	462	G	O4'-C1'-C2'	5.59	112.64	107.60
1	Ad	1707	G	O4'-C1'-N9	5.59	112.68	108.20
71	CB	70	LYS	N-CA-CB	5.59	120.67	110.60
84	Aa	173	C	N3-C4-C5	-5.59	119.66	121.90
84	Aa	1485	A	P-O5'-C5'	-5.59	111.95	120.90
84	Aa	1988	G	O4'-C1'-N9	5.59	112.68	108.20
85	Ac	132	C	N3-C4-C5	-5.59	119.66	121.90
2	Ae	28	G	C4'-C3'-C2'	-5.59	97.01	102.60
84	Aa	1044	A	C4-C5-C6	5.59	119.80	117.00
84	Aa	2504	A	C5-C6-N1	-5.59	114.90	117.70
85	Ac	53	A	C4-C5-C6	5.59	119.80	117.00
86	Ab	77	A	O4'-C1'-N9	5.59	112.67	108.20
86	Ab	104	C	N3-C4-C5	-5.59	119.66	121.90
84	Aa	619	C	N3-C4-N4	5.59	121.91	118.00
84	Aa	676	G	C5-C6-O6	-5.59	125.25	128.60
84	Aa	914	C	N3-C4-N4	5.59	121.91	118.00
84	Aa	1321	A	O4'-C1'-N9	5.59	112.67	108.20
84	Aa	1889	G	O4'-C1'-N9	5.59	112.67	108.20
84	Aa	2005	C	N3-C4-N4	5.59	121.91	118.00
84	Aa	2056	C	N3-C4-C5	-5.59	119.66	121.90
84	Aa	2298	A	C5-C6-N1	-5.59	114.91	117.70
84	Aa	3073	A	C5-C6-N6	-5.59	119.23	123.70
1	Ad	757	G	O4'-C1'-C2'	5.59	112.63	107.60
1	Ad	1772	A	O4'-C1'-C2'	-5.59	100.21	105.80
71	CB	61	GLU	N-CA-CB	5.59	120.66	110.60
84	Aa	245	C	N3-C4-C5	-5.59	119.66	121.90
84	Aa	304	A	C5-C6-N1	-5.59	114.91	117.70
84	Aa	384	A	C4-C5-C6	5.59	119.80	117.00
84	Aa	1270	G	C5-C6-O6	-5.59	125.25	128.60
84	Aa	1324	C	N3-C4-N4	5.59	121.91	118.00
84	Aa	1880	A	C5-C6-N1	-5.59	114.91	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2151	G	P-O3'-C3'	5.59	126.41	119.70
84	Aa	2546	C	N3-C4-N4	5.59	121.91	118.00
84	Aa	2898	A	C5-C6-N6	-5.59	119.23	123.70
84	Aa	2993	A	C5-C6-N6	-5.59	119.23	123.70
1	Ad	1252	C	N1-C1'-C2'	5.59	121.27	114.00
1	Ad	1418	G	O4'-C1'-C2'	-5.59	100.21	105.80
75	CI	119	PHE	N-CA-CB	5.59	120.66	110.60
84	Aa	543	C	N3-C4-C5	-5.59	119.67	121.90
86	Ab	16	A	O4'-C1'-N9	5.59	112.67	108.20
1	Ad	42	G	O4'-C1'-N9	5.59	112.67	108.20
1	Ad	174	C	N1-C1'-C2'	-5.59	105.86	112.00
1	Ad	1345	G	O4'-C1'-N9	5.59	112.67	108.20
84	Aa	186	A	C5-C6-N6	-5.59	119.23	123.70
84	Aa	405	A	C5-C6-N1	-5.59	114.91	117.70
84	Aa	490	G	O4'-C1'-N9	5.59	112.67	108.20
84	Aa	520	G	C5-C6-O6	-5.59	125.25	128.60
84	Aa	562	G	C5-C6-O6	-5.59	125.25	128.60
84	Aa	695	G	O4'-C1'-N9	5.59	112.67	108.20
84	Aa	703	G	O4'-C1'-N9	5.59	112.67	108.20
84	Aa	860	G	C5-C6-O6	-5.59	125.25	128.60
84	Aa	1003	G	O4'-C1'-N9	5.59	112.67	108.20
84	Aa	2036	C	N3-C4-C5	-5.59	119.67	121.90
84	Aa	2193	A	C5-C6-N6	-5.59	119.23	123.70
84	Aa	2201	G	C5-C6-O6	-5.58	125.25	128.60
84	Aa	2545	C	O4'-C1'-N1	5.58	112.67	108.20
14	BQ	32	ARG	N-CA-CB	5.58	120.65	110.60
84	Aa	384	A	O4'-C1'-N9	5.58	112.67	108.20
84	Aa	688	G	N3-C2-N2	5.58	123.81	119.90
84	Aa	801	G	C5-C6-O6	-5.58	125.25	128.60
84	Aa	1072	C	N3-C4-C5	-5.58	119.67	121.90
84	Aa	1418	C	N3-C4-C5	-5.58	119.67	121.90
84	Aa	1906	A	C5-C6-N6	-5.58	119.23	123.70
84	Aa	2208	A	C5-C6-N1	-5.58	114.91	117.70
1	Ad	287	C	O4'-C1'-C2'	-5.58	100.22	105.80
1	Ad	455	G	C3'-C2'-C1'	-5.58	97.03	101.50
1	Ad	1311	U	N1-C1'-C2'	-5.58	105.86	112.00
1	Ad	1616	U	P-O3'-C3'	5.58	126.40	119.70
41	CA	67	PHE	CB-CG-CD2	5.58	124.71	120.80
84	Aa	568	C	N3-C4-N4	5.58	121.91	118.00
84	Aa	1254	A	C5-C6-N6	-5.58	119.23	123.70
84	Aa	1784	C	N3-C4-N4	5.58	121.91	118.00
84	Aa	2101	A	C5-C6-N6	-5.58	119.23	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2360	A	C5-C6-N1	-5.58	114.91	117.70
84	Aa	2904	A	C4-C5-C6	5.58	119.79	117.00
85	Ac	143	C	N3-C4-C5	-5.58	119.67	121.90
84	Aa	122	A	O4'-C1'-N9	5.58	112.66	108.20
84	Aa	2439	A	C5-C6-N1	-5.58	114.91	117.70
86	Ab	72	G	C4-N9-C1'	5.58	133.75	126.50
1	Ad	76	U	C2'-C3'-O3'	5.58	122.62	113.70
1	Ad	1301	G	C1'-O4'-C4'	5.58	114.36	109.90
84	Aa	303	U	O4'-C1'-N1	5.58	112.66	108.20
84	Aa	705	A	C4-C5-C6	5.58	119.79	117.00
84	Aa	1820	C	N3-C4-C5	-5.58	119.67	121.90
85	Ac	95	G	O4'-C1'-N9	5.58	112.66	108.20
1	Ad	926	G	C1'-O4'-C4'	-5.58	105.44	109.90
1	Ad	1132	G	P-O5'-C5'	5.58	129.82	120.90
84	Aa	19	C	N3-C4-C5	-5.58	119.67	121.90
84	Aa	61	A	C5-C6-N1	-5.58	114.91	117.70
84	Aa	108	A	C4-C5-C6	5.58	119.79	117.00
84	Aa	364	A	C5-C6-N1	-5.58	114.91	117.70
84	Aa	1410	A	C5-C6-N6	-5.58	119.24	123.70
84	Aa	2371	A	C5-C6-N6	-5.58	119.24	123.70
84	Aa	2571	C	N3-C4-N4	5.58	121.90	118.00
84	Aa	2973	A	C5-C6-N6	-5.58	119.24	123.70
84	Aa	3207	C	N3-C4-N4	5.58	121.90	118.00
84	Aa	3268	C	N3-C4-N4	5.58	121.90	118.00
84	Aa	3311	C	N3-C4-C5	-5.58	119.67	121.90
1	Ad	798	C	O4'-C1'-N1	5.57	112.66	108.20
84	Aa	106	G	C5-C6-O6	-5.57	125.26	128.60
84	Aa	513	C	N3-C4-C5	-5.57	119.67	121.90
84	Aa	1107	G	P-O5'-C5'	5.57	129.82	120.90
84	Aa	1350	G	C5-C6-O6	-5.57	125.26	128.60
84	Aa	1606	C	N3-C4-C5	-5.57	119.67	121.90
84	Aa	1922	C	N3-C4-N4	5.57	121.90	118.00
84	Aa	2515	C	N3-C4-C5	-5.57	119.67	121.90
84	Aa	2639	A	C5-C6-N1	-5.57	114.91	117.70
84	Aa	3207	C	N3-C4-C5	-5.57	119.67	121.90
1	Ad	242	A	P-O3'-C3'	5.57	126.39	119.70
1	Ad	393	G	O4'-C1'-C2'	5.57	112.61	107.60
84	Aa	453	U	C5'-C4'-C3'	5.57	124.92	116.00
84	Aa	881	G	O4'-C1'-N9	5.57	112.66	108.20
84	Aa	916	A	C5-C6-N1	-5.57	114.91	117.70
84	Aa	1090	C	N3-C4-N4	5.57	121.90	118.00
84	Aa	1351	C	N3-C4-C5	-5.57	119.67	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1790	A	C5-C6-N6	-5.57	119.24	123.70
84	Aa	3240	C	N3-C4-N4	5.57	121.90	118.00
1	Ad	63	G	C5'-C4'-C3'	5.57	124.91	116.00
1	Ad	135	C	C3'-C2'-C1'	5.57	105.96	101.50
1	Ad	1181	G	C1'-O4'-C4'	-5.57	105.44	109.90
1	Ad	1731	A	C4'-C3'-C2'	-5.57	97.03	102.60
7	BM	96	SER	N-CA-CB	5.57	118.86	110.50
84	Aa	274	U	O4'-C1'-N1	5.57	112.66	108.20
84	Aa	1410	A	C5-C6-N1	-5.57	114.92	117.70
84	Aa	2883	C	O4'-C1'-N1	5.57	112.66	108.20
84	Aa	3155	C	N3-C4-C5	-5.57	119.67	121.90
85	Ac	40	A	C4-C5-C6	5.57	119.78	117.00
1	Ad	1527	U	O4'-C1'-N1	5.57	112.66	108.20
84	Aa	831	G	O4'-C1'-N9	5.57	112.66	108.20
84	Aa	846	A	C5-C6-N6	-5.57	119.25	123.70
84	Aa	2562	A	C5-C6-N1	-5.57	114.92	117.70
84	Aa	2810	A	C5-C6-N1	-5.57	114.92	117.70
1	Ad	1378	C	N1-C1'-C2'	5.57	121.24	114.00
9	BX	40	PHE	CB-CG-CD2	-5.57	116.90	120.80
84	Aa	79	C	N3-C4-N4	5.57	121.90	118.00
84	Aa	1782	G	C5-C6-O6	-5.57	125.26	128.60
84	Aa	2936	A	C5-C6-N6	-5.57	119.25	123.70
84	Aa	3127	C	N3-C4-C5	-5.57	119.67	121.90
84	Aa	3128	A	C4-C5-C6	5.57	119.78	117.00
84	Aa	3148	A	C4-C5-C6	5.57	119.78	117.00
1	Ad	30	G	C1'-O4'-C4'	-5.57	105.45	109.90
1	Ad	311	G	O4'-C1'-N9	5.57	112.65	108.20
1	Ad	646	G	C1'-O4'-C4'	-5.57	105.45	109.90
10	Bg	98	SER	N-CA-CB	5.57	118.85	110.50
78	CL	25	PHE	CB-CG-CD2	-5.57	116.90	120.80
84	Aa	567	G	N3-C2-N2	5.57	123.80	119.90
84	Aa	2068	G	O4'-C1'-N9	5.57	112.65	108.20
84	Aa	2354	G	C4-C5-C6	5.57	122.14	118.80
84	Aa	3257	G	O4'-C1'-N9	5.57	112.65	108.20
84	Aa	3367	C	N3-C4-N4	5.57	121.90	118.00
1	Ad	358	C	O4'-C1'-C2'	-5.56	100.24	105.80
84	Aa	292	A	C5-C6-N6	-5.56	119.25	123.70
84	Aa	1052	A	O4'-C1'-N9	5.56	112.65	108.20
1	Ad	886	A	O4'-C1'-N9	5.56	112.65	108.20
48	CD	236	MET	N-CA-CB	5.56	120.61	110.60
63	CU	111	ARG	N-CA-CB	5.56	120.61	110.60
84	Aa	183	C	N3-C4-C5	-5.56	119.67	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	186	A	C5-C6-N1	-5.56	114.92	117.70
84	Aa	548	G	O4'-C1'-N9	5.56	112.65	108.20
84	Aa	612	U	C4'-C3'-C2'	5.56	108.16	102.60
84	Aa	894	G	O4'-C1'-N9	5.56	112.65	108.20
84	Aa	1875	A	C5-C6-N6	-5.56	119.25	123.70
84	Aa	2020	G	C5-C6-O6	-5.56	125.26	128.60
84	Aa	2547	C	N3-C4-N4	5.56	121.89	118.00
84	Aa	2628	C	N3-C4-C5	-5.56	119.67	121.90
84	Aa	3163	G	C5-C6-O6	-5.56	125.26	128.60
85	Ac	84	C	N3-C4-N4	5.56	121.89	118.00
85	Ac	110	A	C5-C6-N1	-5.56	114.92	117.70
84	Aa	112	C	N3-C4-C5	-5.56	119.68	121.90
84	Aa	884	C	N3-C4-N4	5.56	121.89	118.00
84	Aa	1715	C	N3-C4-N4	5.56	121.89	118.00
84	Aa	2593	A	C4-C5-C6	5.56	119.78	117.00
1	Ad	78	A	C5'-C4'-O4'	5.56	115.77	109.10
84	Aa	149	A	C5-C6-N6	-5.56	119.25	123.70
84	Aa	860	G	O4'-C1'-N9	5.56	112.65	108.20
84	Aa	1431	G	O4'-C1'-N9	5.56	112.65	108.20
84	Aa	1881	C	N3-C4-N4	5.56	121.89	118.00
84	Aa	1994	C	N3-C4-N4	5.56	121.89	118.00
86	Ab	2	G	N7-C8-N9	5.56	115.88	113.10
86	Ab	85	G	C6-C5-N7	-5.56	127.06	130.40
1	Ad	591	C	N1-C1'-C2'	5.56	121.22	114.00
69	CF	151	TYR	CB-CG-CD1	-5.56	117.67	121.00
84	Aa	445	C	N3-C4-C5	-5.56	119.68	121.90
84	Aa	1614	G	C5-C6-O6	-5.56	125.27	128.60
84	Aa	1669	C	N3-C4-C5	-5.56	119.68	121.90
84	Aa	1705	A	C4-C5-C6	5.56	119.78	117.00
84	Aa	1747	A	C5'-C4'-O4'	5.56	115.77	109.10
84	Aa	1885	G	O4'-C1'-N9	5.56	112.65	108.20
84	Aa	2604	A	O4'-C1'-N9	5.56	112.65	108.20
84	Aa	3159	C	N3-C4-N4	5.56	121.89	118.00
84	Aa	3171	C	N3-C4-N4	5.56	121.89	118.00
84	Aa	3341	C	N3-C4-C5	-5.56	119.68	121.90
85	Ac	104	A	C5-C6-N6	-5.56	119.25	123.70
84	Aa	1640	A	C5-C6-N1	-5.56	114.92	117.70
84	Aa	1641	G	O4'-C1'-N9	5.56	112.64	108.20
84	Aa	1917	A	C5-C6-N1	-5.56	114.92	117.70
84	Aa	1970	A	C5-C6-N6	-5.56	119.25	123.70
84	Aa	2028	C	N3-C4-C5	-5.56	119.68	121.90
84	Aa	2098	A	C5-C6-N1	-5.56	114.92	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2837	C	N3-C4-N4	5.56	121.89	118.00
1	Ad	59	G	C1'-O4'-C4'	-5.55	105.46	109.90
1	Ad	453	C	C3'-C2'-C1'	5.55	105.94	101.50
84	Aa	4	C	N3-C4-N4	5.55	121.89	118.00
84	Aa	256	G	O4'-C1'-N9	5.55	112.64	108.20
84	Aa	614	C	N3-C4-C5	-5.55	119.68	121.90
84	Aa	1030	A	C5-C6-N6	-5.55	119.26	123.70
84	Aa	1302	C	N3-C4-C5	-5.55	119.68	121.90
84	Aa	1307	A	C5-C6-N6	-5.55	119.26	123.70
84	Aa	1482	C	N3-C4-C5	-5.55	119.68	121.90
84	Aa	1692	U	O4'-C1'-N1	5.55	112.64	108.20
84	Aa	1702	C	N3-C4-N4	5.55	121.89	118.00
84	Aa	2252	C	N3-C4-N4	5.55	121.89	118.00
84	Aa	2630	A	C5-C6-N1	-5.55	114.92	117.70
84	Aa	3210	G	C5'-C4'-O4'	5.55	115.77	109.10
85	Ac	63	C	N3-C4-N4	5.55	121.89	118.00
1	Ad	403	A	O4'-C1'-N9	5.55	112.64	108.20
85	Ac	19	A	C4-C5-C6	5.55	119.78	117.00
86	Ab	48	G	C5-C6-O6	-5.55	125.27	128.60
1	Ad	193	G	O4'-C1'-N9	5.55	112.64	108.20
1	Ad	610	A	O4'-C1'-N9	5.55	112.64	108.20
84	Aa	97	G	O4'-C1'-N9	5.55	112.64	108.20
84	Aa	110	C	N3-C4-N4	5.55	121.89	118.00
84	Aa	560	C	N3-C4-C5	-5.55	119.68	121.90
84	Aa	576	C	O4'-C1'-N1	5.55	112.64	108.20
84	Aa	722	C	N3-C4-C5	-5.55	119.68	121.90
84	Aa	1343	C	N3-C4-N4	5.55	121.89	118.00
84	Aa	2779	G	O4'-C1'-N9	5.55	112.64	108.20
84	Aa	2944	C	N3-C4-C5	-5.55	119.68	121.90
1	Ad	1426	C	O4'-C1'-N1	5.55	112.64	108.20
6	BK	64	TYR	CB-CG-CD1	5.55	124.33	121.00
84	Aa	1025	G	O4'-C1'-N9	5.55	112.64	108.20
84	Aa	1574	C	N3-C4-N4	5.55	121.89	118.00
84	Aa	3023	G	O4'-C1'-N9	5.55	112.64	108.20
86	Ab	71	A	C4-C5-N7	-5.55	107.92	110.70
1	Ad	1160	G	C5'-C4'-O4'	5.55	115.76	109.10
84	Aa	580	C	N3-C4-N4	5.55	121.88	118.00
84	Aa	1031	A	C5-C6-N1	-5.55	114.93	117.70
84	Aa	1624	G	O4'-C1'-N9	5.55	112.64	108.20
84	Aa	1753	A	C4-C5-C6	5.55	119.77	117.00
85	Ac	17	A	C4-C5-C6	5.55	119.77	117.00
1	Ad	298	C	N1-C1'-C2'	5.55	121.21	114.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	507	G	C3'-C2'-C1'	-5.55	97.06	101.50
1	Ad	1793	C	O4'-C1'-C2'	-5.55	100.25	105.80
72	CC	336	TYR	CB-CG-CD1	-5.55	117.67	121.00
84	Aa	4	C	N3-C4-C5	-5.55	119.68	121.90
84	Aa	325	A	C5-C6-N6	-5.55	119.26	123.70
84	Aa	1336	A	C5-C6-N6	-5.55	119.26	123.70
84	Aa	1568	A	C5-C6-N6	-5.55	119.26	123.70
84	Aa	1731	A	C5-C6-N6	-5.55	119.26	123.70
84	Aa	1787	C	N3-C4-C5	-5.55	119.68	121.90
84	Aa	1987	C	N3-C4-N4	5.55	121.88	118.00
84	Aa	2018	C	N3-C4-N4	5.55	121.88	118.00
84	Aa	2303	C	N3-C4-C5	-5.55	119.68	121.90
84	Aa	3201	A	C5-C6-N1	-5.55	114.93	117.70
84	Aa	2219	A	C5-C6-N1	-5.54	114.93	117.70
84	Aa	2249	U	O4'-C1'-N1	5.54	112.64	108.20
84	Aa	3107	A	C5-C6-N1	-5.54	114.93	117.70
86	Ab	89	G	N1-C6-O6	5.54	123.23	119.90
84	Aa	1156	A	O4'-C1'-N9	5.54	112.63	108.20
84	Aa	1738	A	C4-C5-C6	5.54	119.77	117.00
84	Aa	2046	G	C5-C6-O6	-5.54	125.27	128.60
84	Aa	2408	G	C5-C6-O6	-5.54	125.27	128.60
84	Aa	2839	A	C5-C6-N6	-5.54	119.27	123.70
84	Aa	3056	C	N3-C4-C5	-5.54	119.68	121.90
84	Aa	3327	A	C5-C6-N6	-5.54	119.27	123.70
85	Ac	13	A	C4-C5-C6	5.54	119.77	117.00
86	Ab	35	C	O4'-C1'-N1	5.54	112.63	108.20
84	Aa	1745	G	O4'-C1'-N9	5.54	112.63	108.20
84	Aa	2447	A	C4-C5-C6	5.54	119.77	117.00
84	Aa	2647	C	O4'-C1'-N1	5.54	112.63	108.20
84	Aa	2901	C	C6-N1-C1'	-5.54	114.15	120.80
84	Aa	3169	C	N3-C4-C5	-5.54	119.68	121.90
86	Ab	22	A	C5-C6-N1	-5.54	114.93	117.70
1	Ad	1032	A	C1'-O4'-C4'	5.54	114.33	109.90
1	Ad	1792	A	P-O3'-C3'	5.54	126.35	119.70
84	Aa	348	C	N3-C4-C5	-5.54	119.68	121.90
84	Aa	976	A	C4-C5-C6	5.54	119.77	117.00
84	Aa	1307	A	C5-C6-N1	-5.54	114.93	117.70
84	Aa	2262	C	N3-C4-C5	-5.54	119.68	121.90
84	Aa	2517	U	O4'-C1'-N1	5.54	112.63	108.20
84	Aa	2565	C	N3-C4-N4	5.54	121.88	118.00
1	Ad	599	G	C5'-C4'-O4'	5.54	115.75	109.10
84	Aa	98	A	C5-C6-N6	-5.54	119.27	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	837	C	N3-C4-C5	-5.54	119.69	121.90
84	Aa	839	A	C5-C6-N1	-5.54	114.93	117.70
84	Aa	876	C	N3-C4-N4	5.54	121.88	118.00
84	Aa	1094	G	O4'-C1'-N9	5.54	112.63	108.20
84	Aa	1138	A	C4-C5-C6	5.54	119.77	117.00
84	Aa	1279	C	N3-C4-N4	5.54	121.88	118.00
84	Aa	1343	C	N3-C4-C5	-5.54	119.69	121.90
84	Aa	2947	G	C5-C6-O6	-5.54	125.28	128.60
1	Ad	1444	G	N9-C1'-C2'	5.54	121.20	114.00
73	CO	69	THR	N-CA-CB	5.54	120.82	110.30
84	Aa	698	A	C5-C6-N1	-5.54	114.93	117.70
84	Aa	868	A	C5-C6-N1	-5.54	114.93	117.70
84	Aa	1493	A	C4-C5-C6	5.54	119.77	117.00
84	Aa	1493	A	O4'-C1'-N9	5.54	112.63	108.20
84	Aa	1775	C	N3-C4-C5	-5.54	119.69	121.90
84	Aa	1831	A	C5-C6-N6	-5.54	119.27	123.70
84	Aa	1958	G	C5'-C4'-O4'	5.54	115.74	109.10
84	Aa	2104	G	O4'-C1'-N9	5.54	112.63	108.20
84	Aa	3155	C	O4'-C1'-N1	5.54	112.63	108.20
84	Aa	3245	G	O4'-C1'-N9	5.54	112.63	108.20
84	Aa	3272	A	O4'-C1'-N9	5.54	112.63	108.20
1	Ad	573	C	C3'-C2'-C1'	5.54	105.93	101.50
48	CD	219	PHE	CB-CG-CD1	5.54	124.67	120.80
62	CS	152	PRO	CA-N-CD	-5.54	103.75	111.50
84	Aa	21	G	P-O3'-C3'	-5.54	113.06	119.70
84	Aa	439	A	C5-C6-N6	-5.54	119.27	123.70
84	Aa	640	C	N3-C4-N4	5.54	121.88	118.00
84	Aa	781	C	N3-C4-C5	-5.54	119.69	121.90
84	Aa	872	G	C5-C6-O6	-5.54	125.28	128.60
84	Aa	1464	A	C5-C6-N1	-5.54	114.93	117.70
84	Aa	2547	C	N3-C4-C5	-5.54	119.69	121.90
84	Aa	2936	A	O4'-C1'-N9	5.54	112.63	108.20
86	Ab	92	C	C4-C5-C6	5.54	120.17	117.40
6	BK	54	TYR	CB-CG-CD1	-5.53	117.68	121.00
64	Ci	56	TYR	CB-CG-CD2	-5.53	117.68	121.00
84	Aa	289	C	C5-C6-N1	5.53	123.77	121.00
84	Aa	806	C	N3-C4-C5	-5.53	119.69	121.90
84	Aa	843	C	N3-C4-C5	-5.53	119.69	121.90
84	Aa	1013	A	C4-C5-C6	5.53	119.77	117.00
84	Aa	1249	A	C5-C6-N6	-5.53	119.27	123.70
84	Aa	1264	A	C5-C6-N1	-5.53	114.93	117.70
84	Aa	1080	C	N3-C4-C5	-5.53	119.69	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1468	A	C5-C6-N6	-5.53	119.27	123.70
84	Aa	2064	C	N3-C4-N4	5.53	121.87	118.00
84	Aa	2110	G	P-O5'-C5'	-5.53	112.05	120.90
84	Aa	2160	C	N3-C4-N4	5.53	121.87	118.00
85	Ac	159	G	O4'-C1'-N9	5.53	112.62	108.20
1	Ad	159	U	C1'-O4'-C4'	-5.53	105.47	109.90
1	Ad	799	A	O4'-C1'-N9	5.53	112.62	108.20
1	Ad	1745	U	C1'-O4'-C4'	5.53	114.33	109.90
84	Aa	393	A	C5-C6-N1	-5.53	114.94	117.70
84	Aa	1494	A	C5-C6-N1	-5.53	114.94	117.70
84	Aa	1517	C	N3-C4-C5	-5.53	119.69	121.90
84	Aa	1922	C	N3-C4-C5	-5.53	119.69	121.90
84	Aa	2301	C	N3-C4-N4	5.53	121.87	118.00
84	Aa	3092	A	C5-C6-N6	-5.53	119.28	123.70
84	Aa	3104	A	O4'-C1'-N9	5.53	112.62	108.20
84	Aa	3178	C	P-O3'-C3'	5.53	126.34	119.70
1	Ad	462	G	C1'-O4'-C4'	-5.53	105.48	109.90
1	Ad	1431	A	O4'-C1'-N9	5.53	112.62	108.20
2	Ae	22	G	C1'-O4'-C4'	-5.53	105.48	109.90
84	Aa	1033	G	C5-C6-O6	-5.53	125.28	128.60
84	Aa	2804	A	C5-C6-N1	-5.53	114.94	117.70
1	Ad	147	C	C3'-C2'-C1'	5.53	105.92	101.50
1	Ad	369	G	N9-C1'-C2'	-5.53	105.92	112.00
1	Ad	1439	G	N9-C1'-C2'	5.53	121.19	114.00
84	Aa	611	C	N3-C4-N4	5.53	121.87	118.00
84	Aa	759	C	N3-C4-C5	-5.53	119.69	121.90
84	Aa	783	A	C5-C6-N6	-5.53	119.28	123.70
84	Aa	1276	C	N3-C4-C5	-5.53	119.69	121.90
84	Aa	1455	A	C5-C6-N6	-5.53	119.28	123.70
84	Aa	1635	A	O4'-C1'-N9	5.53	112.62	108.20
84	Aa	2391	C	N3-C4-N4	5.53	121.87	118.00
84	Aa	2710	C	N3-C4-C5	-5.53	119.69	121.90
84	Aa	3233	C	N3-C4-C5	-5.53	119.69	121.90
84	Aa	111	C	N3-C4-N4	5.53	121.87	118.00
84	Aa	1148	G	C5-C6-O6	-5.53	125.28	128.60
84	Aa	1275	A	C5-C6-N6	-5.53	119.28	123.70
84	Aa	1415	G	O4'-C1'-N9	5.53	112.62	108.20
84	Aa	1805	A	C5-C6-N1	-5.53	114.94	117.70
84	Aa	1990	A	O4'-C1'-N9	5.53	112.62	108.20
84	Aa	2728	C	N3-C4-C5	-5.53	119.69	121.90
84	Aa	3041	A	C5-C6-N6	-5.53	119.28	123.70
85	Ac	152	G	O4'-C1'-N9	5.53	112.62	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1564	A	O3'-P-O5'	-5.52	93.50	104.00
50	CP	140	TYR	CB-CG-CD1	5.52	124.31	121.00
84	Aa	2207	C	N3-C4-C5	-5.52	119.69	121.90
84	Aa	2698	A	C4-C5-C6	5.52	119.76	117.00
84	Aa	327	A	C5-C6-N6	-5.52	119.28	123.70
84	Aa	376	A	C5-C6-N6	-5.52	119.28	123.70
84	Aa	656	G	N1-C6-O6	5.52	123.21	119.90
84	Aa	1053	C	N3-C4-N4	5.52	121.86	118.00
84	Aa	1308	A	C5-C6-N1	-5.52	114.94	117.70
84	Aa	1522	G	O4'-C1'-N9	5.52	112.62	108.20
84	Aa	1802	A	C5-C6-N1	-5.52	114.94	117.70
84	Aa	2935	A	C5-C6-N1	-5.52	114.94	117.70
86	Ab	31	G	C4-C5-N7	5.52	113.01	110.80
42	CJ	113	ASP	N-CA-CB	5.52	120.54	110.60
84	Aa	639	A	C5-C6-N6	-5.52	119.28	123.70
84	Aa	879	A	C5-C6-N6	-5.52	119.28	123.70
85	Ac	98	U	O4'-C1'-N1	5.52	112.62	108.20
85	Ac	134	G	O4'-C1'-N9	5.52	112.62	108.20
1	Ad	236	U	N1-C1'-C2'	5.52	121.18	114.00
84	Aa	250	C	N3-C4-N4	5.52	121.86	118.00
84	Aa	1276	C	N3-C4-N4	5.52	121.86	118.00
84	Aa	1657	C	N3-C4-N4	5.52	121.86	118.00
84	Aa	1721	A	C4'-C3'-C2'	-5.52	97.08	102.60
84	Aa	1738	A	C5-C6-N6	-5.52	119.28	123.70
84	Aa	2331	A	O4'-C1'-N9	5.52	112.62	108.20
84	Aa	2460	A	C4-C5-C6	5.52	119.76	117.00
84	Aa	2487	A	C5-C6-N6	-5.52	119.28	123.70
84	Aa	3382	A	C5-C6-N1	-5.52	114.94	117.70
1	Ad	200	C	C3'-C2'-C1'	5.52	105.91	101.50
84	Aa	2160	C	N3-C4-C5	-5.52	119.69	121.90
84	Aa	2352	G	N3-C2-N2	5.52	123.76	119.90
84	Aa	2526	G	C5-C6-O6	-5.52	125.29	128.60
84	Aa	2774	A	C5-C6-N6	-5.52	119.29	123.70
84	Aa	1166	C	N3-C4-C5	-5.52	119.69	121.90
84	Aa	2114	A	C5-C6-N1	-5.52	114.94	117.70
1	Ad	1361	G	C4'-C3'-C2'	-5.51	97.08	102.60
42	CJ	58	SER	N-CA-CB	5.51	118.77	110.50
48	CD	187	GLU	N-CA-CB	5.51	120.53	110.60
84	Aa	385	A	C5-C6-N6	-5.51	119.29	123.70
84	Aa	416	A	C5-C6-N6	-5.51	119.29	123.70
84	Aa	1102	A	C4-C5-C6	5.51	119.76	117.00
84	Aa	1335	C	N3-C4-N4	5.51	121.86	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1399	C	N3-C4-C5	-5.51	119.69	121.90
84	Aa	1417	G	O4'-C1'-N9	5.51	112.61	108.20
84	Aa	2089	A	C5-C6-N1	-5.51	114.94	117.70
84	Aa	2832	G	C5-C6-O6	-5.51	125.29	128.60
85	Ac	83	C	N3-C4-N4	5.51	121.86	118.00
1	Ad	1713	C	O4'-C1'-N1	5.51	112.61	108.20
84	Aa	499	A	C4-C5-C6	5.51	119.76	117.00
84	Aa	1542	A	C5-C6-N6	-5.51	119.29	123.70
84	Aa	2174	C	N3-C4-N4	5.51	121.86	118.00
84	Aa	2940	G	C5-C6-O6	-5.51	125.29	128.60
63	CU	34	LYS	N-CA-CB	5.51	120.52	110.60
84	Aa	95	G	P-O5'-C5'	5.51	129.72	120.90
84	Aa	1911	A	C4-C5-C6	5.51	119.76	117.00
84	Aa	2226	C	N3-C4-C5	-5.51	119.69	121.90
84	Aa	2251	A	C5-C6-N6	-5.51	119.29	123.70
84	Aa	2559	C	N3-C4-C5	-5.51	119.69	121.90
85	Ac	44	A	O4'-C1'-N9	5.51	112.61	108.20
85	Ac	126	A	C5-C6-N1	-5.51	114.94	117.70
1	Ad	165	U	O4'-C1'-N1	5.51	112.61	108.20
1	Ad	1409	G	C3'-C2'-C1'	-5.51	97.09	101.50
84	Aa	421	A	C4-C5-C6	5.51	119.75	117.00
84	Aa	564	A	C4-C5-C6	5.51	119.75	117.00
84	Aa	869	A	O4'-C1'-N9	5.51	112.61	108.20
84	Aa	1520	A	C5-C6-N6	-5.51	119.29	123.70
84	Aa	2681	A	C5-C6-N6	-5.51	119.29	123.70
84	Aa	3028	A	C5-C6-N6	-5.51	119.29	123.70
84	Aa	3170	C	N3-C4-C5	-5.51	119.70	121.90
10	Bg	202	SER	N-CA-CB	5.51	118.76	110.50
84	Aa	2641	A	O4'-C1'-N9	5.51	112.61	108.20
84	Aa	3073	A	C5-C6-N1	-5.51	114.95	117.70
1	Ad	360	G	C3'-C2'-C1'	-5.51	97.09	101.50
1	Ad	1388	A	O4'-C1'-C2'	-5.51	100.29	105.80
50	CP	140	TYR	CB-CG-CD2	-5.51	117.70	121.00
84	Aa	82	C	N3-C4-N4	5.51	121.86	118.00
84	Aa	643	G	C5-C6-O6	-5.51	125.30	128.60
84	Aa	839	A	C5-C6-N6	-5.51	119.30	123.70
84	Aa	1455	A	O4'-C1'-N9	5.51	112.61	108.20
84	Aa	2111	A	C5-C6-N1	-5.51	114.95	117.70
6	BK	82	LEU	CA-C-N	5.50	132.51	117.10
84	Aa	1059	A	C5-C6-N6	-5.50	119.30	123.70
84	Aa	2740	C	N3-C4-C5	-5.50	119.70	121.90
84	Aa	2912	A	C5-C6-N1	-5.50	114.95	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	Ab	20	C	C6-N1-C2	5.50	122.50	120.30
1	Ad	315	U	O4'-C1'-C2'	-5.50	100.30	105.80
84	Aa	582	C	N3-C4-C5	-5.50	119.70	121.90
84	Aa	914	C	N3-C4-C5	-5.50	119.70	121.90
84	Aa	1181	A	C5-C6-N1	-5.50	114.95	117.70
84	Aa	2158	C	N3-C4-C5	-5.50	119.70	121.90
84	Aa	2254	A	C5-C6-N1	-5.50	114.95	117.70
84	Aa	2545	C	N3-C4-C5	-5.50	119.70	121.90
85	Ac	105	A	C5-C6-N1	-5.50	114.95	117.70
1	Ad	600	C	O4'-C1'-N1	5.50	112.60	108.20
84	Aa	661	A	C4-C5-C6	5.50	119.75	117.00
84	Aa	769	C	N3-C4-C5	-5.50	119.70	121.90
84	Aa	1721	A	C5-C6-N1	-5.50	114.95	117.70
84	Aa	2082	A	C5-C6-N6	-5.50	119.30	123.70
84	Aa	2361	C	N3-C4-N4	5.50	121.85	118.00
84	Aa	2749	A	C5-C6-N1	-5.50	114.95	117.70
84	Aa	2949	G	N3-C2-N2	5.50	123.75	119.90
84	Aa	3088	A	C5-C6-N6	-5.50	119.30	123.70
84	Aa	3137	G	C5-C6-O6	-5.50	125.30	128.60
85	Ac	124	C	N3-C4-N4	5.50	121.85	118.00
86	Ab	66	G	O4'-C1'-N9	5.50	112.60	108.20
1	Ad	1442	A	C1'-O4'-C4'	-5.50	105.50	109.90
84	Aa	243	C	N3-C4-C5	-5.50	119.70	121.90
84	Aa	2323	A	C4-C5-C6	5.50	119.75	117.00
84	Aa	2359	C	N3-C4-N4	5.50	121.85	118.00
1	Ad	191	U	C4'-C3'-C2'	-5.50	97.10	102.60
1	Ad	462	G	O4'-C1'-N9	5.50	112.60	108.20
1	Ad	1044	A	P-O3'-C3'	5.50	126.30	119.70
1	Ad	1428	A	O4'-C1'-N9	5.50	112.60	108.20
30	BB	49	SER	N-CA-CB	5.50	118.75	110.50
84	Aa	108	A	O4'-C1'-N9	5.50	112.60	108.20
84	Aa	1471	A	C4-C5-C6	5.50	119.75	117.00
84	Aa	2025	C	N3-C4-N4	5.50	121.85	118.00
84	Aa	2650	A	O4'-C1'-N9	5.50	112.60	108.20
84	Aa	2899	A	C4-C5-C6	5.50	119.75	117.00
84	Aa	3200	A	C4-C5-C6	5.50	119.75	117.00
1	Ad	536	U	C1'-O4'-C4'	-5.50	105.50	109.90
84	Aa	306	A	O4'-C1'-N9	5.50	112.60	108.20
84	Aa	588	G	C5-C6-O6	-5.50	125.30	128.60
84	Aa	1512	A	C5-C6-N1	-5.50	114.95	117.70
84	Aa	1632	G	O4'-C1'-N9	5.50	112.60	108.20
84	Aa	2075	C	N3-C4-C5	-5.50	119.70	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2374	G	C5-C6-N1	-5.50	108.75	111.50
84	Aa	2540	C	N3-C4-C5	-5.50	119.70	121.90
84	Aa	2733	A	O4'-C1'-N9	5.50	112.60	108.20
84	Aa	3161	C	N3-C4-C5	-5.50	119.70	121.90
84	Aa	2566	C	N3-C4-C5	-5.50	119.70	121.90
84	Aa	3115	A	C5-C6-N6	-5.50	119.30	123.70
84	Aa	3297	A	O4'-C1'-N9	5.50	112.60	108.20
84	Aa	833	G	O4'-C1'-N9	5.49	112.59	108.20
84	Aa	954	A	C5-C6-N1	-5.49	114.95	117.70
84	Aa	1023	G	C5-C6-O6	-5.49	125.30	128.60
84	Aa	1554	C	O4'-C1'-N1	5.49	112.59	108.20
84	Aa	1963	G	C5-C6-O6	-5.49	125.30	128.60
84	Aa	158	A	C5-C6-N6	-5.49	119.31	123.70
84	Aa	217	A	C5-C6-N1	-5.49	114.95	117.70
84	Aa	365	A	C5-C6-N6	-5.49	119.31	123.70
1	Ad	546	U	C3'-C2'-C1'	-5.49	97.11	101.50
1	Ad	1111	C	O4'-C1'-N1	5.49	112.59	108.20
84	Aa	136	C	N3-C4-N4	5.49	121.84	118.00
84	Aa	952	C	N3-C4-N4	5.49	121.84	118.00
84	Aa	1425	G	C5-C6-O6	-5.49	125.31	128.60
84	Aa	1429	U	O4'-C1'-N1	5.49	112.59	108.20
84	Aa	2304	A	C5-C6-N1	-5.49	114.95	117.70
84	Aa	2458	A	C5-C6-N1	-5.49	114.95	117.70
84	Aa	3072	A	C5-C6-N6	-5.49	119.31	123.70
84	Aa	3075	G	O4'-C1'-N9	5.49	112.59	108.20
1	Ad	960	A	O4'-C1'-N9	5.49	112.59	108.20
1	Ad	1796	G	O4'-C1'-C2'	5.49	112.54	107.60
36	BH	135	GLU	N-CA-CB	5.49	120.48	110.60
37	CG	47	PHE	CB-CG-CD1	5.49	124.64	120.80
72	CC	345	THR	N-CA-CB	5.49	120.73	110.30
84	Aa	1520	A	C5-C6-N1	-5.49	114.95	117.70
84	Aa	1842	C	N3-C4-C5	-5.49	119.70	121.90
84	Aa	2265	A	C5-C6-N1	-5.49	114.95	117.70
84	Aa	2527	G	C4-C5-C6	5.49	122.09	118.80
1	Ad	13	C	O4'-C1'-N1	5.49	112.59	108.20
84	Aa	1199	A	C4-C5-C6	5.49	119.74	117.00
84	Aa	1358	C	N3-C4-N4	5.49	121.84	118.00
84	Aa	1887	A	C4-C5-C6	5.49	119.74	117.00
84	Aa	2772	A	C5-C6-N1	-5.49	114.96	117.70
85	Ac	53	A	C5-C6-N6	-5.49	119.31	123.70
85	Ac	135	A	C5-C6-N1	-5.49	114.96	117.70
84	Aa	264	C	N3-C4-N4	5.49	121.84	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	525	A	C4-C5-C6	5.49	119.74	117.00
84	Aa	742	G	O4'-C1'-N9	5.49	112.59	108.20
84	Aa	870	G	O4'-C1'-N9	5.49	112.59	108.20
84	Aa	1382	C	N3-C4-C5	-5.49	119.71	121.90
84	Aa	1858	U	O4'-C1'-N1	5.49	112.59	108.20
84	Aa	1861	A	C5-C6-N1	-5.49	114.96	117.70
84	Aa	884	C	N3-C4-C5	-5.48	119.71	121.90
84	Aa	1586	A	C5-C6-N6	-5.48	119.31	123.70
84	Aa	2659	A	O4'-C1'-N9	5.48	112.59	108.20
84	Aa	2927	C	N3-C4-N4	5.48	121.84	118.00
84	Aa	173	C	N3-C4-N4	5.48	121.84	118.00
84	Aa	330	C	N3-C4-C5	-5.48	119.71	121.90
84	Aa	640	C	P-O5'-C5'	5.48	129.67	120.90
84	Aa	782	G	O3'-P-O5'	-5.48	93.58	104.00
84	Aa	1565	G	O5'-C5'-C4'	-5.48	101.28	111.70
84	Aa	1739	G	N1-C6-O6	5.48	123.19	119.90
84	Aa	2969	A	C5-C6-N6	-5.48	119.31	123.70
84	Aa	3198	C	N3-C4-C5	-5.48	119.71	121.90
85	Ac	116	G	O4'-C1'-N9	5.48	112.59	108.20
1	Ad	178	A	O4'-C1'-C2'	-5.48	100.32	105.80
1	Ad	869	U	O4'-C1'-C2'	-5.48	100.32	105.80
1	Ad	1506	G	O4'-C1'-N9	5.48	112.58	108.20
84	Aa	1467	G	O4'-C1'-N9	5.48	112.58	108.20
84	Aa	1637	G	O4'-C1'-N9	5.48	112.58	108.20
84	Aa	2319	A	C5-C6-N6	-5.48	119.31	123.70
84	Aa	2402	G	O4'-C1'-N9	5.48	112.58	108.20
84	Aa	2971	A	C5-C6-N6	-5.48	119.32	123.70
84	Aa	2979	G	O4'-C1'-N9	5.48	112.58	108.20
84	Aa	3087	A	O4'-C1'-N9	5.48	112.58	108.20
85	Ac	23	C	N3-C4-C5	-5.48	119.71	121.90
1	Ad	1360	G	P-O3'-C3'	5.48	126.28	119.70
84	Aa	1073	G	C5-C6-O6	-5.48	125.31	128.60
84	Aa	1167	G	O4'-C1'-N9	5.48	112.58	108.20
84	Aa	1874	A	O4'-C1'-N9	5.48	112.58	108.20
84	Aa	2210	A	C5-C6-N1	-5.48	114.96	117.70
84	Aa	2279	C	N3-C4-N4	5.48	121.83	118.00
84	Aa	2840	A	C4-C5-C6	5.48	119.74	117.00
86	Ab	102	G	N1-C6-O6	5.48	123.19	119.90
48	CD	7	PHE	CB-CG-CD2	-5.48	116.97	120.80
84	Aa	321	A	C4-C5-C6	5.48	119.74	117.00
84	Aa	722	C	N3-C4-N4	5.48	121.83	118.00
84	Aa	917	A	O4'-C1'-N9	5.48	112.58	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1287	C	N3-C4-C5	-5.48	119.71	121.90
84	Aa	1495	G	C5-C6-O6	-5.48	125.31	128.60
84	Aa	3351	A	C4-C5-C6	5.48	119.74	117.00
1	Ad	839	G	C1'-O4'-C4'	-5.47	105.52	109.90
84	Aa	1172	A	C4-C5-C6	5.47	119.74	117.00
84	Aa	1499	C	N3-C4-N4	5.47	121.83	118.00
84	Aa	1713	A	C5-C6-N6	-5.47	119.32	123.70
84	Aa	2117	G	C5-C6-O6	-5.47	125.31	128.60
84	Aa	2362	A	C4-C5-C6	5.47	119.74	117.00
84	Aa	2384	G	O4'-C1'-N9	5.47	112.58	108.20
84	Aa	2705	A	C4-C5-C6	5.47	119.74	117.00
84	Aa	2768	C	N3-C4-C5	-5.47	119.71	121.90
85	Ac	54	A	O4'-C1'-N9	5.47	112.58	108.20
1	Ad	487	A	C3'-C2'-C1'	-5.47	97.12	101.50
29	BR	1	MET	C-N-CA	5.47	133.79	122.30
46	Ca	8	ASN	N-CA-C	-5.47	96.23	111.00
48	CD	119	GLU	N-CA-CB	5.47	120.45	110.60
84	Aa	59	A	C4-C5-C6	5.47	119.74	117.00
84	Aa	96	C	N3-C4-C5	-5.47	119.71	121.90
84	Aa	415	G	O4'-C1'-N9	5.47	112.58	108.20
84	Aa	476	C	N3-C4-C5	-5.47	119.71	121.90
84	Aa	481	G	C5-C6-O6	-5.47	125.32	128.60
84	Aa	1039	G	C5-C6-O6	-5.47	125.32	128.60
84	Aa	1275	A	O4'-C1'-N9	5.47	112.58	108.20
84	Aa	1854	A	C5-C6-N1	-5.47	114.96	117.70
84	Aa	1944	G	O4'-C4'-C3'	5.47	110.48	106.10
84	Aa	2391	C	N3-C4-C5	-5.47	119.71	121.90
84	Aa	2679	A	C5-C6-N1	-5.47	114.96	117.70
84	Aa	3389	C	N3-C4-N4	5.47	121.83	118.00
86	Ab	22	A	C5-C6-N6	-5.47	119.32	123.70
86	Ab	95	U	O4'-C1'-N1	5.47	112.58	108.20
84	Aa	258	C	N3-C4-N4	5.47	121.83	118.00
84	Aa	454	A	O4'-C1'-N9	5.47	112.58	108.20
84	Aa	723	G	C4'-C3'-C2'	-5.47	97.13	102.60
84	Aa	2433	U	O4'-C1'-N1	5.47	112.58	108.20
84	Aa	2823	C	N3-C4-N4	5.47	121.83	118.00
84	Aa	2853	A	C5-C6-N1	-5.47	114.97	117.70
86	Ab	91	C	N3-C4-C5	-5.47	119.71	121.90
1	Ad	1580	G	C3'-C2'-C1'	-5.47	97.12	101.50
84	Aa	499	A	C5-C6-N1	-5.47	114.97	117.70
84	Aa	1105	G	P-O3'-C3'	5.47	126.26	119.70
84	Aa	1476	G	N3-C2-N2	5.47	123.73	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1563	G	C5'-C4'-O4'	-5.47	102.54	109.10
84	Aa	1712	A	C4-C5-C6	5.47	119.73	117.00
84	Aa	3076	C	N3-C4-C5	-5.47	119.71	121.90
84	Aa	3155	C	N3-C4-N4	5.47	121.83	118.00
86	Ab	37	G	O4'-C1'-N9	5.47	112.58	108.20
86	Ab	43	A	C4-C5-C6	5.47	119.73	117.00
84	Aa	1185	G	O4'-C1'-N9	5.47	112.58	108.20
84	Aa	1795	A	C5-C6-N1	-5.47	114.97	117.70
84	Aa	2901	C	N3-C4-N4	5.47	121.83	118.00
1	Ad	717	G	C2'-C3'-O3'	5.47	122.45	113.70
1	Ad	826	C	O4'-C1'-N1	5.47	112.57	108.20
71	CB	121	ASN	N-CA-CB	5.47	120.44	110.60
84	Aa	101	C	N3-C4-C5	-5.47	119.71	121.90
84	Aa	810	A	O4'-C1'-N9	5.47	112.57	108.20
84	Aa	2138	A	C4-C5-C6	5.47	119.73	117.00
84	Aa	2654	G	C5-C6-O6	-5.47	125.32	128.60
85	Ac	119	C	N3-C4-N4	5.47	121.83	118.00
1	Ad	724	U	C1'-O4'-C4'	5.46	114.27	109.90
1	Ad	1232	G	C5'-C4'-O4'	5.46	115.66	109.10
84	Aa	12	G	C4'-C3'-C2'	-5.46	97.14	102.60
84	Aa	677	U	O4'-C1'-N1	5.46	112.57	108.20
84	Aa	1174	G	C5-C6-O6	-5.46	125.32	128.60
84	Aa	1225	A	C5-C6-N1	-5.46	114.97	117.70
84	Aa	1409	G	C5-C6-O6	-5.46	125.32	128.60
84	Aa	2389	A	C5-C6-N6	-5.46	119.33	123.70
84	Aa	2493	C	N3-C4-N4	5.46	121.83	118.00
84	Aa	3036	C	N3-C4-C5	-5.46	119.71	121.90
85	Ac	49	G	C5-C6-O6	-5.46	125.32	128.60
84	Aa	353	A	P-O5'-C5'	-5.46	112.16	120.90
1	Ad	706	U	O4'-C1'-N1	5.46	112.57	108.20
1	Ad	1092	A	O4'-C1'-N9	5.46	112.57	108.20
84	Aa	1709	U	O4'-C1'-N1	5.46	112.57	108.20
84	Aa	2662	A	C5-C6-N6	-5.46	119.33	123.70
85	Ac	105	A	C5-C6-N6	-5.46	119.33	123.70
84	Aa	2105	G	O4'-C1'-N9	5.46	112.57	108.20
84	Aa	2174	C	N3-C4-C5	-5.46	119.72	121.90
84	Aa	3182	A	C4'-C3'-C2'	-5.46	97.14	102.60
1	Ad	333	G	C3'-C2'-C1'	-5.46	97.13	101.50
84	Aa	1185	G	C5-C6-O6	-5.46	125.33	128.60
84	Aa	1195	C	N3-C4-N4	5.46	121.82	118.00
84	Aa	1802	A	C5-C6-N6	-5.46	119.33	123.70
84	Aa	2298	A	C4-C5-C6	5.46	119.73	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2807	G	O4'-C1'-N9	5.46	112.57	108.20
84	Aa	2972	C	N3-C4-C5	-5.46	119.72	121.90
84	Aa	3030	A	C4-C5-C6	5.46	119.73	117.00
85	Ac	80	A	C4-C5-C6	5.46	119.73	117.00
1	Ad	468	A	C5'-C4'-O4'	5.46	115.65	109.10
1	Ad	1316	A	C1'-O4'-C4'	-5.46	105.53	109.90
1	Ad	1515	G	O4'-C1'-N9	5.46	112.57	108.20
35	BG	28	PHE	CB-CG-CD2	-5.46	116.98	120.80
84	Aa	1	G	C4-N9-C1'	5.46	133.59	126.50
84	Aa	109	G	O4'-C1'-N9	5.46	112.57	108.20
84	Aa	223	C	N3-C4-N4	5.46	121.82	118.00
84	Aa	573	A	C5-C6-N1	-5.46	114.97	117.70
84	Aa	619	C	N3-C4-C5	-5.46	119.72	121.90
84	Aa	823	A	O4'-C1'-N9	5.46	112.57	108.20
84	Aa	972	C	N3-C4-N4	5.46	121.82	118.00
84	Aa	1200	A	C5-C6-N1	-5.46	114.97	117.70
84	Aa	1235	A	C4-C5-C6	5.46	119.73	117.00
84	Aa	1854	A	C4-C5-C6	5.46	119.73	117.00
84	Aa	1915	G	O4'-C1'-N9	5.46	112.56	108.20
84	Aa	2228	A	C5-C6-N1	-5.46	114.97	117.70
84	Aa	2933	C	C5'-C4'-O4'	5.46	115.65	109.10
84	Aa	330	C	N3-C4-N4	5.46	121.82	118.00
84	Aa	943	G	N3-C2-N2	5.46	123.72	119.90
84	Aa	2999	G	N3-C2-N2	5.46	123.72	119.90
1	Ad	768	A	OP1-P-OP2	-5.45	111.42	119.60
84	Aa	249	A	C5-C6-N1	-5.45	114.97	117.70
84	Aa	620	C	N3-C4-C5	-5.45	119.72	121.90
84	Aa	1438	A	O4'-C1'-N9	5.45	112.56	108.20
84	Aa	1878	G	C5-C6-O6	-5.45	125.33	128.60
84	Aa	2026	C	N3-C4-N4	5.45	121.82	118.00
84	Aa	2149	G	C8-N9-C1'	-5.45	119.91	127.00
84	Aa	2450	G	C5-C6-O6	-5.45	125.33	128.60
84	Aa	2985	C	C5'-C4'-O4'	5.45	115.64	109.10
84	Aa	3087	A	C5-C6-N1	-5.45	114.97	117.70
85	Ac	23	C	N3-C4-N4	5.45	121.82	118.00
1	Ad	267	G	O4'-C1'-N9	5.45	112.56	108.20
84	Aa	162	G	O4'-C1'-N9	5.45	112.56	108.20
84	Aa	1744	C	P-O3'-C3'	-5.45	113.16	119.70
84	Aa	2158	C	N3-C4-N4	5.45	121.82	118.00
84	Aa	2185	U	C5'-C4'-C3'	-5.45	107.28	116.00
1	Ad	934	A	C3'-C2'-C1'	5.45	105.86	101.50
84	Aa	199	G	O4'-C1'-N9	5.45	112.56	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	712	A	C4-C5-C6	5.45	119.72	117.00
84	Aa	889	C	N3-C4-N4	5.45	121.81	118.00
84	Aa	985	C	N3-C4-C5	-5.45	119.72	121.90
84	Aa	2503	A	C5-C6-N6	-5.45	119.34	123.70
84	Aa	2660	A	O4'-C1'-N9	5.45	112.56	108.20
84	Aa	3082	G	C5-C6-O6	-5.45	125.33	128.60
84	Aa	3341	C	N3-C4-N4	5.45	121.81	118.00
1	Ad	922	U	O4'-C1'-N1	5.45	112.56	108.20
1	Ad	1226	U	O4'-C1'-C2'	-5.45	100.35	105.80
84	Aa	689	G	C5-C6-O6	-5.45	125.33	128.60
84	Aa	777	G	C5-C6-O6	-5.45	125.33	128.60
84	Aa	1789	C	N3-C4-C5	-5.45	119.72	121.90
84	Aa	2789	G	C2'-C3'-O3'	5.45	122.42	113.70
84	Aa	2885	U	C4'-C3'-C2'	-5.45	97.15	102.60
86	Ab	51	G	C6-C5-N7	-5.45	127.13	130.40
1	Ad	140	C	O4'-C1'-C2'	-5.45	100.35	105.80
1	Ad	1512	C	C3'-C2'-C1'	5.45	105.86	101.50
84	Aa	332	A	C4-C5-C6	5.45	119.72	117.00
84	Aa	769	C	N3-C4-N4	5.45	121.81	118.00
84	Aa	1157	A	C5-C6-N6	-5.45	119.34	123.70
84	Aa	2033	C	N3-C4-N4	5.45	121.81	118.00
1	Ad	1365	C	O4'-C1'-N1	5.45	112.56	108.20
28	BA	43	TYR	CB-CG-CD1	5.45	124.27	121.00
84	Aa	803	G	O4'-C1'-N9	5.45	112.56	108.20
84	Aa	1139	A	C4-C5-C6	5.45	119.72	117.00
84	Aa	1160	G	O4'-C1'-N9	5.45	112.56	108.20
84	Aa	1752	C	N3-C4-N4	5.45	121.81	118.00
84	Aa	2625	C	N3-C4-N4	5.45	121.81	118.00
84	Aa	3140	A	C4-C5-C6	5.45	119.72	117.00
84	Aa	3251	C	P-O3'-C3'	5.45	126.23	119.70
85	Ac	79	A	C5-C6-N6	-5.45	119.34	123.70
1	Ad	29	U	O4'-C1'-C2'	-5.44	100.36	105.80
1	Ad	179	A	C4'-C3'-C2'	5.44	108.04	102.60
84	Aa	1990	A	C5-C6-N6	-5.44	119.34	123.70
85	Ac	61	A	O4'-C1'-N9	5.44	112.56	108.20
1	Ad	100	C	C1'-O4'-C4'	5.44	114.25	109.90
1	Ad	1618	G	N9-C1'-C2'	5.44	121.07	114.00
1	Ad	1776	A	C3'-C2'-C1'	-5.44	97.14	101.50
73	CO	136	PRO	CA-N-CD	-5.44	103.88	111.50
84	Aa	24	C	N3-C4-N4	5.44	121.81	118.00
84	Aa	640	C	O4'-C1'-N1	5.44	112.55	108.20
84	Aa	1162	A	C5-C6-N6	-5.44	119.35	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1520	A	O4'-C1'-N9	5.44	112.55	108.20
84	Aa	1862	C	N3-C4-C5	-5.44	119.72	121.90
84	Aa	2184	U	O4'-C1'-N1	5.44	112.55	108.20
48	CD	180	PHE	CB-CG-CD1	5.44	124.61	120.80
84	Aa	5	G	C4-N9-C1'	5.44	133.57	126.50
84	Aa	727	G	C5-C6-O6	-5.44	125.34	128.60
84	Aa	1291	A	C5-C6-N6	-5.44	119.35	123.70
84	Aa	1480	G	O4'-C1'-N9	5.44	112.55	108.20
84	Aa	1744	C	N3-C4-C5	-5.44	119.72	121.90
85	Ac	81	U	P-O3'-C3'	5.44	126.23	119.70
84	Aa	418	G	O4'-C1'-N9	5.44	112.55	108.20
84	Aa	1998	A	C4-C5-C6	5.44	119.72	117.00
1	Ad	450	A	C3'-C2'-C1'	5.44	105.85	101.50
22	BZ	89	ALA	N-CA-CB	5.44	117.71	110.10
70	Cq	52	SER	N-CA-CB	5.44	118.66	110.50
84	Aa	2	C	N3-C4-C5	-5.44	119.72	121.90
84	Aa	237	C	N3-C4-C5	-5.44	119.72	121.90
84	Aa	450	C	N3-C4-N4	5.44	121.81	118.00
84	Aa	949	C	C5-C6-N1	5.44	123.72	121.00
84	Aa	2225	C	N3-C4-N4	5.44	121.81	118.00
84	Aa	2373	C	C5'-C4'-O4'	-5.44	102.58	109.10
84	Aa	3007	A	C5-C6-N6	-5.44	119.35	123.70
84	Aa	3192	G	C5-C6-O6	-5.44	125.34	128.60
84	Aa	3205	C	N3-C4-C5	-5.44	119.72	121.90
84	Aa	3287	A	C4-C5-C6	5.44	119.72	117.00
86	Ab	29	C	C5-C6-N1	5.44	123.72	121.00
1	Ad	1234	A	C1'-O4'-C4'	5.44	114.25	109.90
70	Cq	9	GLU	N-CA-C	-5.44	96.32	111.00
84	Aa	3350	C	N3-C4-N4	5.44	121.81	118.00
85	Ac	62	C	P-O5'-C5'	-5.44	112.20	120.90
1	Ad	879	C	N1-C1'-C2'	5.43	121.06	114.00
1	Ad	1507	G	C3'-C2'-C1'	-5.43	97.15	101.50
84	Aa	262	A	C5-C6-N1	-5.43	114.98	117.70
84	Aa	342	A	O4'-C1'-N9	5.43	112.55	108.20
84	Aa	898	G	C4-N9-C1'	5.43	133.57	126.50
84	Aa	1229	A	C5-C6-N6	-5.43	119.35	123.70
84	Aa	2950	C	N3-C4-N4	5.43	121.80	118.00
84	Aa	3212	C	N3-C4-C5	-5.43	119.73	121.90
84	Aa	479	C	N3-C4-C5	-5.43	119.73	121.90
84	Aa	792	A	O4'-C1'-N9	5.43	112.55	108.20
84	Aa	1907	A	C5-C6-N1	-5.43	114.98	117.70
84	Aa	2061	C	N3-C4-N4	5.43	121.80	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2397	A	C5-C6-N6	-5.43	119.36	123.70
84	Aa	3094	C	N3-C4-N4	5.43	121.80	118.00
84	Aa	3319	G	C5-C6-O6	-5.43	125.34	128.60
84	Aa	3366	C	N3-C4-C5	-5.43	119.73	121.90
69	CF	77	PHE	CB-CG-CD1	5.43	124.60	120.80
84	Aa	712	A	O4'-C1'-N9	5.43	112.55	108.20
84	Aa	883	G	C5-C6-O6	-5.43	125.34	128.60
84	Aa	1088	A	C5-C6-N1	-5.43	114.98	117.70
84	Aa	2481	C	N3-C4-N4	5.43	121.80	118.00
84	Aa	2660	A	C5-C6-N1	-5.43	114.98	117.70
1	Ad	58	U	P-O5'-C5'	5.43	129.59	120.90
84	Aa	343	G	O4'-C1'-N9	5.43	112.54	108.20
84	Aa	1312	A	C4-C5-C6	5.43	119.72	117.00
84	Aa	1591	A	C5-C6-N6	-5.43	119.36	123.70
84	Aa	1761	C	N3-C4-C5	-5.43	119.73	121.90
1	Ad	1799	G	O4'-C1'-N9	5.43	112.54	108.20
84	Aa	1206	A	C5-C6-N6	-5.43	119.36	123.70
84	Aa	1850	C	N3-C4-N4	5.43	121.80	118.00
84	Aa	2048	C	N3-C4-N4	5.43	121.80	118.00
84	Aa	3070	G	C5-C6-O6	-5.43	125.34	128.60
84	Aa	84	A	C4-C5-C6	5.43	119.71	117.00
84	Aa	1228	C	N3-C4-C5	-5.43	119.73	121.90
84	Aa	1369	G	C5-C6-O6	-5.43	125.34	128.60
84	Aa	1790	A	C5-C6-N1	-5.43	114.99	117.70
84	Aa	3017	A	C5-C6-N1	-5.43	114.99	117.70
84	Aa	3272	A	C5-C6-N1	-5.43	114.99	117.70
85	Ac	10	G	O4'-C1'-N9	5.43	112.54	108.20
1	Ad	969	U	P-O3'-C3'	5.42	126.21	119.70
1	Ad	1133	C	O4'-C1'-N1	5.42	112.54	108.20
1	Ad	1752	U	C3'-C2'-C1'	-5.42	97.16	101.50
84	Aa	130	G	C5-C6-O6	-5.42	125.34	128.60
84	Aa	373	A	C5-C6-N1	-5.42	114.99	117.70
84	Aa	603	G	P-O5'-C5'	5.42	129.58	120.90
84	Aa	1013	A	C5-C6-N6	-5.42	119.36	123.70
84	Aa	1278	A	C5-C6-N1	-5.42	114.99	117.70
84	Aa	1513	C	N3-C4-C5	-5.42	119.73	121.90
84	Aa	2674	A	C5-C6-N6	-5.42	119.36	123.70
84	Aa	3263	C	O4'-C4'-C3'	5.42	110.44	106.10
1	Ad	281	U	O4'-C4'-C3'	-5.42	98.58	104.00
1	Ad	1010	A	O4'-C1'-N9	5.42	112.54	108.20
84	Aa	586	A	C5-C6-N1	-5.42	114.99	117.70
84	Aa	850	A	C5-C6-N6	-5.42	119.36	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1353	A	C5-C6-N1	-5.42	114.99	117.70
84	Aa	2400	A	C5-C6-N1	-5.42	114.99	117.70
84	Aa	3088	A	C5-C6-N1	-5.42	114.99	117.70
84	Aa	3088	A	O4'-C1'-N9	5.42	112.54	108.20
84	Aa	3114	A	C5-C6-N1	-5.42	114.99	117.70
84	Aa	3148	A	C5-C6-N1	-5.42	114.99	117.70
84	Aa	477	C	N3-C4-N4	5.42	121.80	118.00
84	Aa	771	G	N1-C2-N3	-5.42	120.65	123.90
84	Aa	1911	A	O4'-C1'-N9	5.42	112.54	108.20
84	Aa	2028	C	N3-C4-N4	5.42	121.80	118.00
84	Aa	2533	A	O4'-C1'-N9	5.42	112.54	108.20
84	Aa	3033	A	O4'-C1'-N9	5.42	112.54	108.20
85	Ac	34	U	O4'-C1'-N1	5.42	112.54	108.20
1	Ad	1177	G	C1'-O4'-C4'	-5.42	105.56	109.90
84	Aa	114	G	P-O3'-C3'	5.42	126.20	119.70
84	Aa	932	A	C4-C5-C6	5.42	119.71	117.00
84	Aa	1370	A	C5-C6-N1	-5.42	114.99	117.70
84	Aa	2457	G	O4'-C1'-N9	5.42	112.54	108.20
1	Ad	61	A	C4'-C3'-O3'	-5.42	98.02	109.40
1	Ad	238	G	O4'-C1'-N9	-5.42	103.86	108.20
1	Ad	262	U	O4'-C1'-C2'	-5.42	100.38	105.80
84	Aa	1277	A	C4-C5-C6	5.42	119.71	117.00
84	Aa	1490	A	O4'-C1'-N9	5.42	112.53	108.20
84	Aa	1937	C	N3-C4-C5	-5.42	119.73	121.90
84	Aa	2603	C	N3-C4-N4	5.42	121.79	118.00
84	Aa	2681	A	C5-C6-N1	-5.42	114.99	117.70
84	Aa	2961	C	N3-C4-N4	5.42	121.79	118.00
84	Aa	3248	G	O4'-C1'-N9	5.42	112.53	108.20
1	Ad	144	U	P-O3'-C3'	-5.42	113.20	119.70
1	Ad	1355	U	N1-C1'-C2'	5.42	121.04	114.00
1	Ad	1790	G	C1'-O4'-C4'	-5.42	105.57	109.90
21	BP	70	ARG	N-CA-CB	5.42	120.35	110.60
84	Aa	710	C	N3-C4-N4	5.42	121.79	118.00
84	Aa	857	G	O4'-C1'-N9	5.42	112.53	108.20
84	Aa	1231	C	N3-C4-N4	5.42	121.79	118.00
84	Aa	1568	A	C5-C6-N1	-5.42	114.99	117.70
84	Aa	1713	A	C5-C6-N1	-5.42	114.99	117.70
84	Aa	1917	A	C5-C6-N6	-5.42	119.37	123.70
84	Aa	1977	C	N3-C4-N4	5.42	121.79	118.00
84	Aa	2612	A	C4-C5-C6	5.42	119.71	117.00
84	Aa	227	C	N3-C4-C5	-5.42	119.73	121.90
84	Aa	1397	A	O4'-C1'-N9	5.42	112.53	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1507	A	C5-C6-N6	-5.42	119.37	123.70
84	Aa	1516	G	N3-C2-N2	5.42	123.69	119.90
84	Aa	2136	A	C5-C6-N6	-5.42	119.37	123.70
84	Aa	2635	G	O4'-C1'-N9	5.42	112.53	108.20
86	Ab	88	U	C5-C6-N1	5.42	125.41	122.70
1	Ad	902	C	P-O3'-C3'	5.41	126.20	119.70
1	Ad	968	A	C3'-C2'-C1'	-5.41	97.17	101.50
1	Ad	1620	C	C3'-C2'-C1'	5.41	105.83	101.50
84	Aa	2	C	N3-C4-N4	5.41	121.79	118.00
84	Aa	72	A	C5-C6-N1	-5.41	114.99	117.70
84	Aa	197	A	C4-C5-C6	5.41	119.71	117.00
84	Aa	201	G	C5-C6-O6	-5.41	125.35	128.60
84	Aa	298	G	C5-C6-O6	-5.41	125.35	128.60
84	Aa	316	A	C4-C5-C6	5.41	119.71	117.00
84	Aa	917	A	C5-C6-N1	-5.41	114.99	117.70
84	Aa	1272	G	O4'-C1'-N9	5.41	112.53	108.20
84	Aa	1471	A	O4'-C1'-N9	5.41	112.53	108.20
84	Aa	1594	G	O4'-C1'-N9	5.41	112.53	108.20
84	Aa	1743	C	N3-C4-C5	-5.41	119.73	121.90
84	Aa	1882	A	C5-C6-N1	-5.41	114.99	117.70
84	Aa	2575	C	N3-C4-C5	-5.41	119.73	121.90
84	Aa	2822	A	O4'-C1'-N9	5.41	112.53	108.20
84	Aa	3110	A	O4'-C1'-N9	5.41	112.53	108.20
84	Aa	3321	C	N3-C4-C5	-5.41	119.73	121.90
1	Ad	1465	C	C5'-C4'-O4'	5.41	115.59	109.10
84	Aa	1029	C	N3-C4-C5	-5.41	119.73	121.90
84	Aa	2558	U	C2-N1-C1'	5.41	124.19	117.70
84	Aa	3364	A	C4-C5-C6	5.41	119.71	117.00
84	Aa	3374	C	O4'-C1'-N1	5.41	112.53	108.20
1	Ad	1234	A	O4'-C1'-C2'	-5.41	100.39	105.80
84	Aa	420	A	C5-C6-N6	-5.41	119.37	123.70
84	Aa	1465	A	O4'-C1'-N9	5.41	112.53	108.20
84	Aa	1837	A	C5-C6-N1	-5.41	115.00	117.70
84	Aa	2081	C	N3-C4-C5	-5.41	119.74	121.90
84	Aa	2924	G	O4'-C1'-N9	5.41	112.53	108.20
84	Aa	3025	A	C4-C5-C6	5.41	119.71	117.00
1	Ad	315	U	C4'-C3'-C2'	-5.41	97.19	102.60
1	Ad	573	C	O4'-C1'-C2'	-5.41	100.39	105.80
1	Ad	1347	U	O4'-C1'-N1	5.41	112.53	108.20
84	Aa	708	C	N3-C4-C5	-5.41	119.74	121.90
84	Aa	1571	A	C5-C6-N6	-5.41	119.37	123.70
84	Aa	2202	A	C5-C6-N1	-5.41	115.00	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2279	C	N3-C4-C5	-5.41	119.74	121.90
84	Aa	3059	C	N3-C4-C5	-5.41	119.74	121.90
85	Ac	53	A	O4'-C1'-N9	5.41	112.53	108.20
85	Ac	146	G	N1-C2-N3	-5.41	120.66	123.90
86	Ab	95	U	N3-C4-O4	5.41	123.19	119.40
1	Ad	317	U	C3'-C2'-C1'	5.41	105.83	101.50
1	Ad	1641	A	O4'-C1'-N9	5.41	112.53	108.20
84	Aa	533	G	C5-C6-O6	-5.41	125.36	128.60
1	Ad	311	G	P-O5'-C5'	-5.41	112.25	120.90
84	Aa	383	A	C5-C6-N1	-5.41	115.00	117.70
84	Aa	1002	A	O4'-C1'-N9	5.41	112.52	108.20
84	Aa	1312	A	O4'-C1'-N9	5.41	112.52	108.20
84	Aa	2006	A	C5-C6-N6	-5.41	119.38	123.70
84	Aa	2052	G	C5-C6-O6	-5.41	125.36	128.60
84	Aa	2299	C	N3-C4-C5	-5.41	119.74	121.90
84	Aa	2730	A	C5-C6-N6	-5.41	119.38	123.70
84	Aa	2768	C	N3-C4-N4	5.41	121.78	118.00
1	Ad	1327	C	N1-C1'-C2'	5.40	121.03	114.00
84	Aa	222	C	N3-C4-C5	-5.40	119.74	121.90
84	Aa	414	G	C5-C6-O6	-5.40	125.36	128.60
84	Aa	664	A	O4'-C1'-N9	5.40	112.52	108.20
84	Aa	1005	C	N3-C4-N4	5.40	121.78	118.00
84	Aa	3039	U	O4'-C1'-N1	5.40	112.52	108.20
86	Ab	18	C	N3-C4-C5	-5.40	119.74	121.90
86	Ab	56	G	N1-C2-N3	-5.40	120.66	123.90
84	Aa	213	G	N3-C2-N2	5.40	123.68	119.90
84	Aa	720	G	C5'-C4'-O4'	5.40	115.58	109.10
84	Aa	820	A	C4-C5-C6	5.40	119.70	117.00
84	Aa	2260	C	C2-N3-C4	5.40	122.60	119.90
84	Aa	2388	C	N3-C4-C5	-5.40	119.74	121.90
84	Aa	2595	G	O4'-C1'-N9	5.40	112.52	108.20
84	Aa	2596	A	O4'-C1'-N9	5.40	112.52	108.20
84	Aa	3048	C	N3-C4-C5	-5.40	119.74	121.90
85	Ac	101	U	O4'-C1'-N1	5.40	112.52	108.20
1	Ad	289	G	C1'-O4'-C4'	-5.40	105.58	109.90
1	Ad	1543	U	N1-C1'-C2'	5.40	121.02	114.00
1	Ad	1553	A	N9-C1'-C2'	5.40	121.02	114.00
1	Ad	1751	U	N1-C1'-C2'	5.40	121.02	114.00
84	Aa	224	C	C4'-C3'-C2'	-5.40	97.20	102.60
84	Aa	1209	G	C5-C6-O6	-5.40	125.36	128.60
84	Aa	2365	C	N3-C4-N4	5.40	121.78	118.00
84	Aa	2729	C	N3-C4-C5	-5.40	119.74	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	39	A	C5-C6-N1	-5.40	115.00	117.70
84	Aa	523	C	N3-C4-C5	-5.40	119.74	121.90
84	Aa	1831	A	P-O5'-C5'	5.40	129.54	120.90
84	Aa	2086	A	O4'-C1'-C2'	5.40	112.46	107.60
1	Ad	792	U	O4'-C1'-C2'	-5.40	100.40	105.80
1	Ad	1758	G	N9-C1'-C2'	5.40	121.02	114.00
53	CY	8	THR	N-CA-CB	5.40	120.56	110.30
84	Aa	219	A	C5-C6-N1	-5.40	115.00	117.70
84	Aa	823	A	C5-C6-N1	-5.40	115.00	117.70
84	Aa	1174	G	O4'-C1'-N9	5.40	112.52	108.20
84	Aa	1864	G	O4'-C1'-N9	5.40	112.52	108.20
84	Aa	1932	A	C5-C6-N6	-5.40	119.38	123.70
84	Aa	2301	C	N3-C4-C5	-5.40	119.74	121.90
84	Aa	2643	A	C4-C5-C6	5.40	119.70	117.00
84	Aa	2767	C	N3-C4-C5	-5.40	119.74	121.90
84	Aa	2836	G	O4'-C1'-N9	5.40	112.52	108.20
1	Ad	1792	A	C3'-C2'-C1'	5.40	105.82	101.50
84	Aa	176	A	C4-C5-C6	5.40	119.70	117.00
84	Aa	539	C	N3-C4-C5	-5.40	119.74	121.90
84	Aa	731	G	O4'-C1'-N9	5.40	112.52	108.20
84	Aa	1543	A	C5-C6-N6	-5.40	119.38	123.70
84	Aa	1586	A	O4'-C1'-N9	5.40	112.52	108.20
64	Ci	38	LYS	N-CA-CB	5.39	120.31	110.60
84	Aa	578	C	P-O3'-C3'	5.39	126.17	119.70
84	Aa	651	A	C5-C6-N1	-5.39	115.00	117.70
84	Aa	2449	A	C5-C6-N6	-5.39	119.38	123.70
84	Aa	2464	G	C5-C6-O6	-5.39	125.36	128.60
84	Aa	2739	A	P-O5'-C5'	-5.39	112.27	120.90
85	Ac	104	A	C5-C6-N1	-5.39	115.00	117.70
84	Aa	19	C	N3-C4-N4	5.39	121.78	118.00
84	Aa	157	G	O4'-C1'-N9	5.39	112.51	108.20
84	Aa	196	A	C5-C6-N1	-5.39	115.00	117.70
84	Aa	616	A	C4-C5-C6	5.39	119.70	117.00
84	Aa	981	A	C5-C6-N1	-5.39	115.00	117.70
84	Aa	1063	G	C5-C6-O6	-5.39	125.36	128.60
84	Aa	1090	C	N3-C4-C5	-5.39	119.74	121.90
84	Aa	1344	A	C5-C6-N1	-5.39	115.00	117.70
84	Aa	1837	A	C5-C6-N6	-5.39	119.39	123.70
84	Aa	2816	G	C5-C6-O6	-5.39	125.36	128.60
84	Aa	2874	A	O4'-C1'-N9	5.39	112.51	108.20
84	Aa	3299	A	O4'-C1'-N9	5.39	112.51	108.20
84	Aa	3328	A	O4'-C1'-N9	5.39	112.51	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	Ac	113	U	O4'-C1'-N1	5.39	112.51	108.20
85	Ac	129	C	N3-C4-N4	5.39	121.78	118.00
84	Aa	1970	A	O4'-C1'-N9	5.39	112.51	108.20
84	Aa	3142	C	N3-C4-C5	-5.39	119.74	121.90
86	Ab	1	G	C6-C5-N7	-5.39	127.17	130.40
1	Ad	365	C	N1-C1'-C2'	5.39	121.01	114.00
1	Ad	555	G	C1'-O4'-C4'	-5.39	105.59	109.90
1	Ad	1305	U	C1'-O4'-C4'	5.39	114.21	109.90
1	Ad	1632	C	P-O3'-C3'	5.39	126.17	119.70
84	Aa	426	A	C5-C6-N6	-5.39	119.39	123.70
84	Aa	1010	A	C4-C5-C6	5.39	119.69	117.00
84	Aa	1162	A	C5-C6-N1	-5.39	115.00	117.70
84	Aa	1602	A	C4-C5-C6	5.39	119.69	117.00
84	Aa	1761	C	N3-C4-N4	5.39	121.77	118.00
84	Aa	2882	U	P-O3'-C3'	5.39	126.17	119.70
1	Ad	1069	G	C1'-O4'-C4'	-5.39	105.59	109.90
84	Aa	70	A	C4-C5-C6	5.39	119.69	117.00
84	Aa	2960	A	O4'-C1'-N9	5.39	112.51	108.20
1	Ad	1015	C	O4'-C1'-N1	5.39	112.51	108.20
1	Ad	1665	U	P-O3'-C3'	5.39	126.16	119.70
84	Aa	582	C	N3-C4-N4	5.39	121.77	118.00
84	Aa	918	A	C5-C6-N1	-5.39	115.01	117.70
84	Aa	1337	C	N3-C4-N4	5.39	121.77	118.00
84	Aa	1537	A	C4-C5-C6	5.39	119.69	117.00
84	Aa	1584	A	C5-C6-N1	-5.39	115.01	117.70
84	Aa	1797	U	C5'-C4'-C3'	-5.39	107.38	116.00
84	Aa	2070	C	N3-C4-N4	5.39	121.77	118.00
84	Aa	2430	C	N3-C4-N4	5.39	121.77	118.00
84	Aa	2596	A	C5-C6-N6	-5.39	119.39	123.70
84	Aa	2710	C	N3-C4-N4	5.39	121.77	118.00
84	Aa	2758	C	C2-N3-C4	5.39	122.59	119.90
84	Aa	3334	A	C5'-C4'-C3'	-5.39	107.38	116.00
1	Ad	346	C	C3'-C2'-C1'	5.38	105.81	101.50
1	Ad	1400	G	O4'-C1'-N9	5.38	112.51	108.20
84	Aa	389	A	C5-C6-N1	-5.38	115.01	117.70
84	Aa	1029	C	N3-C4-N4	5.38	121.77	118.00
84	Aa	1097	A	C5-C6-N1	-5.38	115.01	117.70
84	Aa	1780	C	N3-C4-N4	5.38	121.77	118.00
84	Aa	2041	G	C5-C6-O6	-5.38	125.37	128.60
84	Aa	2605	G	O4'-C1'-N9	5.38	112.51	108.20
84	Aa	2952	G	O4'-C1'-N9	5.38	112.51	108.20
84	Aa	3004	G	C5-C6-O6	-5.38	125.37	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3376	C	N3-C4-N4	5.38	121.77	118.00
1	Ad	1063	U	P-O3'-C3'	5.38	126.16	119.70
1	Ad	1074	C	O4'-C1'-N1	5.38	112.50	108.20
84	Aa	825	G	C5-C6-O6	-5.38	125.37	128.60
84	Aa	887	A	C5-C6-N1	-5.38	115.01	117.70
84	Aa	2021	G	P-O3'-C3'	5.38	126.16	119.70
84	Aa	2404	C	N3-C4-N4	5.38	121.77	118.00
84	Aa	2694	A	C5-C6-N1	-5.38	115.01	117.70
84	Aa	2765	A	C5-C6-N1	-5.38	115.01	117.70
84	Aa	2909	A	C4-C5-C6	5.38	119.69	117.00
84	Aa	2986	C	N3-C4-C5	-5.38	119.75	121.90
85	Ac	99	C	N3-C4-N4	5.38	121.77	118.00
86	Ab	101	A	C5-C6-N1	-5.38	115.01	117.70
1	Ad	844	C	O4'-C1'-C2'	-5.38	100.42	105.80
84	Aa	2781	A	O4'-C1'-N9	5.38	112.50	108.20
84	Aa	3193	C	N3-C4-C5	-5.38	119.75	121.90
1	Ad	82	G	C1'-O4'-C4'	-5.38	105.60	109.90
1	Ad	980	C	O4'-C1'-N1	5.38	112.50	108.20
24	BW	128	PHE	CB-CG-CD1	5.38	124.56	120.80
84	Aa	98	A	C5-C6-N1	-5.38	115.01	117.70
84	Aa	323	A	C4-C5-C6	5.38	119.69	117.00
84	Aa	1195	C	N3-C4-C5	-5.38	119.75	121.90
84	Aa	1759	C	N3-C4-C5	-5.38	119.75	121.90
84	Aa	2545	C	N3-C4-N4	5.38	121.77	118.00
84	Aa	2641	A	C5-C6-N6	-5.38	119.40	123.70
1	Ad	719	C	O4'-C1'-N1	5.38	112.50	108.20
1	Ad	1137	A	O4'-C1'-C2'	-5.38	100.42	105.80
1	Ad	1207	A	O4'-C1'-N9	-5.38	103.90	108.20
1	Ad	1655	U	C1'-O4'-C4'	5.38	114.20	109.90
84	Aa	474	G	C2'-C3'-O3'	-5.38	97.67	109.50
84	Aa	1331	C	N3-C4-N4	5.38	121.76	118.00
84	Aa	1717	G	O4'-C1'-N9	5.38	112.50	108.20
84	Aa	2308	A	C4-C5-C6	5.38	119.69	117.00
84	Aa	2372	A	C5-C6-N1	-5.38	115.01	117.70
84	Aa	3103	G	N3-C2-N2	5.38	123.66	119.90
84	Aa	3270	C	O4'-C1'-N1	5.38	112.50	108.20
86	Ab	118	C	C5-C4-N4	-5.38	116.44	120.20
84	Aa	447	C	N3-C4-N4	5.38	121.76	118.00
84	Aa	996	A	C4-C5-C6	5.38	119.69	117.00
84	Aa	2429	A	O4'-C1'-N9	5.38	112.50	108.20
1	Ad	253	C	N1-C1'-C2'	5.37	120.99	114.00
1	Ad	385	C	C1'-O4'-C4'	-5.37	105.60	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	319	C	O4'-C1'-N1	5.37	112.50	108.20
84	Aa	1552	C	N3-C4-N4	5.37	121.76	118.00
84	Aa	1841	G	O4'-C1'-N9	5.37	112.50	108.20
84	Aa	2000	C	N3-C4-N4	5.37	121.76	118.00
84	Aa	2005	C	N3-C4-C5	-5.37	119.75	121.90
84	Aa	2318	U	O4'-C1'-N1	5.37	112.50	108.20
84	Aa	3098	U	O4'-C1'-N1	5.37	112.50	108.20
84	Aa	3116	C	N3-C4-C5	-5.37	119.75	121.90
85	Ac	156	C	N3-C4-N4	5.37	121.76	118.00
86	Ab	43	A	N1-C2-N3	5.37	131.99	129.30
1	Ad	164	C	C4'-C3'-C2'	-5.37	97.23	102.60
1	Ad	227	G	P-O3'-C3'	5.37	126.15	119.70
1	Ad	323	U	C3'-C2'-C1'	5.37	105.80	101.50
84	Aa	113	A	C4-C5-C6	5.37	119.69	117.00
84	Aa	338	C	N3-C4-N4	5.37	121.76	118.00
84	Aa	376	A	C5-C6-N1	-5.37	115.01	117.70
84	Aa	636	C	N3-C4-C5	-5.37	119.75	121.90
84	Aa	698	A	C5-C6-N6	-5.37	119.40	123.70
84	Aa	821	C	N3-C4-N4	5.37	121.76	118.00
84	Aa	1518	A	C5-C6-N1	-5.37	115.01	117.70
84	Aa	1583	G	N3-C2-N2	5.37	123.66	119.90
84	Aa	1795	A	O4'-C1'-N9	5.37	112.50	108.20
84	Aa	2100	A	C5-C6-N6	-5.37	119.40	123.70
84	Aa	2436	G	C4'-C3'-O3'	5.37	123.74	113.00
84	Aa	2451	G	N1-C6-O6	5.37	123.12	119.90
84	Aa	2576	C	N3-C4-C5	-5.37	119.75	121.90
84	Aa	2815	A	O4'-C1'-N9	5.37	112.50	108.20
84	Aa	2891	C	N3-C4-C5	-5.37	119.75	121.90
86	Ab	90	A	N7-C8-N9	5.37	116.49	113.80
1	Ad	1203	G	C5'-C4'-O4'	5.37	115.54	109.10
3	Af	21	C	C1'-O4'-C4'	5.37	114.20	109.90
84	Aa	1367	A	C5-C6-N1	-5.37	115.02	117.70
1	Ad	153	U	O4'-C1'-C2'	-5.37	100.43	105.80
84	Aa	347	A	C4-C5-C6	5.37	119.68	117.00
84	Aa	518	G	C5-C6-O6	-5.37	125.38	128.60
84	Aa	926	C	N3-C4-C5	-5.37	119.75	121.90
84	Aa	1030	A	C5-C6-N1	-5.37	115.02	117.70
84	Aa	1723	C	C4'-C3'-C2'	-5.37	97.23	102.60
84	Aa	2729	C	N3-C4-N4	5.37	121.76	118.00
85	Ac	42	G	C5-C6-O6	-5.37	125.38	128.60
85	Ac	105	A	O4'-C1'-N9	5.37	112.50	108.20
84	Aa	618	G	O4'-C1'-N9	5.37	112.49	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1071	G	C5-C6-O6	-5.37	125.38	128.60
84	Aa	2515	C	N3-C4-N4	5.37	121.76	118.00
86	Ab	51	G	N1-C2-N3	-5.37	120.68	123.90
1	Ad	134	G	O4'-C1'-N9	-5.37	103.91	108.20
1	Ad	761	A	C1'-O4'-C4'	-5.37	105.61	109.90
78	CL	156	ILE	N-CA-CB	5.37	123.14	110.80
84	Aa	54	G	C5-C6-O6	-5.37	125.38	128.60
84	Aa	238	C	N3-C4-C5	-5.37	119.75	121.90
84	Aa	826	C	N3-C4-N4	5.37	121.75	118.00
84	Aa	993	A	C4-C5-C6	5.37	119.68	117.00
84	Aa	1377	G	O4'-C1'-N9	5.37	112.49	108.20
84	Aa	1743	C	N3-C4-N4	5.37	121.76	118.00
84	Aa	2257	A	C4-C5-C6	5.37	119.68	117.00
84	Aa	2529	C	C5'-C4'-O4'	5.37	115.54	109.10
84	Aa	3346	C	N3-C4-C5	-5.37	119.75	121.90
85	Ac	84	C	N3-C4-C5	-5.37	119.75	121.90
1	Ad	542	A	N9-C1'-C2'	-5.36	106.10	112.00
2	Ae	44	A	C4'-C3'-C2'	-5.36	97.24	102.60
84	Aa	1254	A	C5-C6-N1	-5.36	115.02	117.70
84	Aa	1394	C	N3-C4-N4	5.36	121.75	118.00
84	Aa	1843	A	C5-C6-N1	-5.36	115.02	117.70
84	Aa	2266	A	C5-C6-N6	-5.36	119.41	123.70
84	Aa	2388	C	C2-N3-C4	5.36	122.58	119.90
84	Aa	3229	C	N3-C4-C5	-5.36	119.75	121.90
86	Ab	95	U	C5-C4-O4	-5.36	122.68	125.90
1	Ad	466	G	O4'-C1'-N9	5.36	112.49	108.20
1	Ad	852	A	O4'-C1'-N9	5.36	112.49	108.20
13	BF	41	HIS	N-CA-CB	5.36	120.25	110.60
84	Aa	371	A	N1-C6-N6	5.36	121.82	118.60
84	Aa	2900	G	N3-C2-N2	5.36	123.65	119.90
84	Aa	3239	G	C5-C6-O6	-5.36	125.38	128.60
1	Ad	282	C	P-O5'-C5'	-5.36	112.32	120.90
1	Ad	965	U	O4'-C1'-N1	5.36	112.49	108.20
1	Ad	1742	A	O4'-C1'-N9	5.36	112.49	108.20
84	Aa	1579	C	C2-N3-C4	5.36	122.58	119.90
84	Aa	1610	A	C5-C6-N6	-5.36	119.41	123.70
84	Aa	1906	A	C5-C6-N1	-5.36	115.02	117.70
84	Aa	2527	G	C6-C5-N7	-5.36	127.18	130.40
84	Aa	2578	G	N3-C2-N2	5.36	123.65	119.90
84	Aa	2774	A	C5-C6-N1	-5.36	115.02	117.70
84	Aa	2810	A	O4'-C1'-N9	5.36	112.49	108.20
84	Aa	1391	A	C5-C6-N1	-5.36	115.02	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1616	G	O4'-C1'-N9	5.36	112.49	108.20
84	Aa	1813	C	N3-C4-N4	5.36	121.75	118.00
84	Aa	2007	C	N3-C4-N4	5.36	121.75	118.00
84	Aa	2765	A	C5-C6-N6	-5.36	119.41	123.70
1	Ad	361	G	O4'-C1'-N9	5.36	112.49	108.20
1	Ad	1769	C	O4'-C1'-C2'	-5.36	100.44	105.80
84	Aa	1040	A	O4'-C1'-N9	5.36	112.48	108.20
84	Aa	1067	G	O4'-C1'-N9	5.36	112.49	108.20
84	Aa	1080	C	N3-C4-N4	5.36	121.75	118.00
84	Aa	1205	C	N3-C4-N4	5.36	121.75	118.00
84	Aa	2141	A	O4'-C1'-N9	5.36	112.48	108.20
84	Aa	2698	A	O4'-C1'-N9	5.36	112.49	108.20
84	Aa	2938	A	C5-C6-N1	-5.36	115.02	117.70
84	Aa	3129	G	O4'-C1'-N9	5.36	112.48	108.20
84	Aa	3211	C	N3-C4-C5	-5.36	119.76	121.90
84	Aa	3264	C	N3-C4-N4	5.36	121.75	118.00
1	Ad	403	A	C1'-O4'-C4'	5.36	114.18	109.90
1	Ad	457	C	C3'-C2'-C1'	-5.36	97.22	101.50
1	Ad	1355	U	O4'-C1'-N1	5.36	112.48	108.20
2	Ae	19	U	O4'-C1'-N1	-5.36	103.92	108.20
84	Aa	239	C	N3-C4-C5	-5.36	119.76	121.90
84	Aa	943	G	O4'-C1'-N9	5.36	112.48	108.20
84	Aa	1238	G	N1-C6-O6	5.36	123.11	119.90
84	Aa	1333	C	N3-C4-C5	-5.36	119.76	121.90
84	Aa	1334	A	C5-C6-N1	-5.36	115.02	117.70
84	Aa	1891	A	C5-C6-N1	-5.36	115.02	117.70
84	Aa	1971	A	C5-C6-N6	-5.36	119.42	123.70
84	Aa	2006	A	C5-C6-N1	-5.36	115.02	117.70
84	Aa	2386	A	C5-C6-N6	-5.36	119.42	123.70
84	Aa	2474	A	C5-C6-N6	-5.36	119.42	123.70
84	Aa	3130	A	C4-C5-C6	5.36	119.68	117.00
1	Ad	72	A	C5'-C4'-O4'	5.35	115.53	109.10
1	Ad	596	A	C3'-C2'-C1'	5.35	105.78	101.50
2	Ae	66	C	O4'-C1'-N1	5.35	112.48	108.20
84	Aa	434	C	N3-C4-N4	5.35	121.75	118.00
84	Aa	493	G	O3'-P-O5'	5.35	114.17	104.00
84	Aa	1311	G	C5-C6-O6	-5.35	125.39	128.60
84	Aa	1734	G	O4'-C1'-N9	5.35	112.48	108.20
84	Aa	2015	G	C5-C6-O6	-5.35	125.39	128.60
84	Aa	2543	G	O4'-C1'-N9	5.35	112.48	108.20
1	Ad	760	G	C1'-O4'-C4'	-5.35	105.62	109.90
84	Aa	167	C	C4'-C3'-C2'	5.35	107.95	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	198	A	C5-C6-N1	-5.35	115.02	117.70
84	Aa	296	C	N3-C4-N4	5.35	121.75	118.00
84	Aa	1241	G	C5'-C4'-C3'	5.35	124.56	116.00
84	Aa	1734	G	C5-C6-O6	-5.35	125.39	128.60
84	Aa	1852	C	N3-C4-N4	5.35	121.75	118.00
84	Aa	2101	A	O4'-C1'-N9	5.35	112.48	108.20
1	Ad	1359	C	P-O3'-C3'	5.35	126.12	119.70
24	BW	97	ARG	N-CA-CB	5.35	120.23	110.60
84	Aa	1209	G	O4'-C1'-N9	5.35	112.48	108.20
84	Aa	2016	A	C5-C6-N6	-5.35	119.42	123.70
84	Aa	2745	C	N3-C4-N4	5.35	121.75	118.00
86	Ab	46	C	O4'-C1'-N1	5.35	112.48	108.20
1	Ad	1164	C	C3'-C2'-C1'	5.35	105.78	101.50
1	Ad	1239	C	C3'-C2'-C1'	5.35	105.78	101.50
1	Ad	1503	C	C3'-C2'-C1'	5.35	105.78	101.50
84	Aa	779	U	O4'-C1'-N1	5.35	112.48	108.20
84	Aa	3028	A	C5-C6-N1	-5.35	115.03	117.70
84	Aa	3173	A	C5-C6-N1	-5.35	115.03	117.70
84	Aa	3216	G	C5-C6-O6	-5.35	125.39	128.60
84	Aa	3256	C	N3-C4-C5	-5.35	119.76	121.90
86	Ab	59	U	O4'-C1'-N1	5.35	112.48	108.20
1	Ad	164	C	C1'-O4'-C4'	-5.35	105.62	109.90
1	Ad	1288	C	O4'-C1'-N1	5.35	112.48	108.20
41	CA	40	TYR	CB-CG-CD2	-5.35	117.79	121.00
48	CD	289	ASN	N-CA-CB	5.35	120.22	110.60
84	Aa	1314	G	N3-C2-N2	5.35	123.64	119.90
84	Aa	1363	C	C2-N3-C4	5.35	122.57	119.90
84	Aa	1500	C	N3-C4-N4	5.35	121.74	118.00
84	Aa	1509	G	N3-C2-N2	5.35	123.64	119.90
84	Aa	2604	A	C4-C5-C6	5.35	119.67	117.00
84	Aa	3296	C	N3-C4-N4	5.35	121.74	118.00
86	Ab	75	G	P-O3'-C3'	5.35	126.12	119.70
1	Ad	926	G	C3'-C2'-C1'	-5.35	97.22	101.50
1	Ad	1007	G	C1'-O4'-C4'	-5.35	105.62	109.90
84	Aa	143	A	C5-C6-N6	-5.35	119.42	123.70
84	Aa	482	C	N3-C4-C5	-5.35	119.76	121.90
84	Aa	1868	C	C2-N3-C4	5.35	122.57	119.90
84	Aa	1945	A	P-O5'-C5'	-5.35	112.35	120.90
84	Aa	3182	A	C5'-C4'-C3'	5.35	124.55	116.00
1	Ad	886	A	C5'-C4'-O4'	5.34	115.51	109.10
30	BB	134	MET	CG-SD-CE	-5.34	91.65	100.20
84	Aa	738	A	C5-C6-N1	-5.34	115.03	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1831	A	C4-C5-C6	5.34	119.67	117.00
84	Aa	2473	C	N3-C4-N4	5.34	121.74	118.00
84	Aa	3322	A	O4'-C1'-N9	5.34	112.48	108.20
1	Ad	153	U	O4'-C1'-N1	5.34	112.47	108.20
84	Aa	2178	G	C2'-C3'-O3'	-5.34	97.74	109.50
84	Aa	2749	A	C5-C6-N6	-5.34	119.43	123.70
1	Ad	844	C	C5'-C4'-O4'	5.34	115.51	109.10
49	CR	188	SER	N-CA-CB	5.34	118.51	110.50
84	Aa	392	C	N3-C4-C5	-5.34	119.76	121.90
84	Aa	885	A	C5-C6-N1	-5.34	115.03	117.70
84	Aa	1018	C	N3-C4-C5	-5.34	119.76	121.90
84	Aa	1153	A	C5-C6-N1	-5.34	115.03	117.70
84	Aa	1264	A	C5'-C4'-O4'	5.34	115.51	109.10
84	Aa	2083	U	C5'-C4'-C3'	5.34	124.55	116.00
84	Aa	2118	G	O4'-C1'-N9	5.34	112.47	108.20
84	Aa	2439	A	C4-C5-C6	5.34	119.67	117.00
84	Aa	2576	C	N3-C4-N4	5.34	121.74	118.00
84	Aa	2620	U	O4'-C1'-N1	5.34	112.47	108.20
84	Aa	3033	A	C5-C6-N6	-5.34	119.43	123.70
2	Ae	53	U	O4'-C1'-N1	5.34	112.47	108.20
84	Aa	6	A	C5-C6-N1	-5.34	115.03	117.70
84	Aa	439	A	C5-C6-N1	-5.34	115.03	117.70
84	Aa	785	U	P-O3'-C3'	5.34	126.11	119.70
84	Aa	981	A	C5-C6-N6	-5.34	119.43	123.70
84	Aa	1104	C	N3-C4-C5	-5.34	119.76	121.90
84	Aa	2054	A	C5-C6-N1	-5.34	115.03	117.70
84	Aa	2481	C	N3-C4-C5	-5.34	119.76	121.90
84	Aa	2611	G	C5-C6-O6	-5.34	125.40	128.60
84	Aa	3038	U	O4'-C1'-N1	5.34	112.47	108.20
1	Ad	255	U	O4'-C1'-C2'	-5.34	100.46	105.80
84	Aa	33	A	C5-C6-N1	-5.34	115.03	117.70
84	Aa	1255	A	C5-C6-N6	-5.34	119.43	123.70
84	Aa	1780	C	N3-C4-C5	-5.34	119.77	121.90
86	Ab	68	G	C6-N1-C2	5.34	128.30	125.10
1	Ad	831	C	C1'-O4'-C4'	5.34	114.17	109.90
1	Ad	973	U	N1-C1'-C2'	5.34	120.94	114.00
1	Ad	1115	G	C3'-C2'-C1'	5.34	105.77	101.50
1	Ad	1203	G	P-O3'-C3'	-5.34	113.30	119.70
84	Aa	150	G	P-O5'-C5'	-5.34	112.36	120.90
84	Aa	180	G	O4'-C1'-N9	5.34	112.47	108.20
84	Aa	679	C	N3-C4-N4	5.34	121.73	118.00
84	Aa	844	A	C4-C5-C6	5.34	119.67	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1364	C	N3-C4-N4	5.34	121.74	118.00
84	Aa	1607	C	C2-N3-C4	5.34	122.57	119.90
84	Aa	2699	A	C5-C6-N6	-5.34	119.43	123.70
84	Aa	3074	A	C5-C6-N1	-5.34	115.03	117.70
84	Aa	114	G	C4'-C3'-C2'	-5.33	97.27	102.60
84	Aa	642	C	O5'-C5'-C4'	5.33	121.84	111.70
84	Aa	727	G	O4'-C1'-N9	5.33	112.47	108.20
84	Aa	850	A	C5-C6-N1	-5.33	115.03	117.70
84	Aa	467	C	N3-C4-C5	-5.33	119.77	121.90
84	Aa	558	G	P-O5'-C5'	-5.33	112.37	120.90
84	Aa	737	C	N3-C4-C5	-5.33	119.77	121.90
84	Aa	744	C	C2-N3-C4	5.33	122.57	119.90
84	Aa	1541	G	C5-C6-O6	-5.33	125.40	128.60
84	Aa	2197	C	N3-C4-N4	5.33	121.73	118.00
84	Aa	2320	A	C4-C5-C6	5.33	119.67	117.00
84	Aa	3236	A	C5-C6-N6	-5.33	119.43	123.70
84	Aa	3284	C	N3-C4-C5	-5.33	119.77	121.90
1	Ad	381	G	O4'-C1'-C2'	5.33	112.40	107.60
1	Ad	1133	C	N1-C1'-C2'	5.33	120.93	114.00
20	BT	56	TYR	CB-CG-CD2	-5.33	117.80	121.00
84	Aa	195	G	O4'-C1'-N9	5.33	112.47	108.20
84	Aa	442	C	N3-C4-N4	5.33	121.73	118.00
84	Aa	599	C	N3-C4-C5	-5.33	119.77	121.90
84	Aa	775	A	C4-C5-C6	5.33	119.67	117.00
84	Aa	789	A	C5-C6-N6	-5.33	119.44	123.70
84	Aa	1027	C	N3-C4-N4	5.33	121.73	118.00
84	Aa	1727	A	O4'-C1'-N9	5.33	112.47	108.20
84	Aa	1794	A	C4-C5-C6	5.33	119.67	117.00
84	Aa	2132	A	C4'-C3'-C2'	-5.33	97.27	102.60
84	Aa	2252	C	N3-C4-C5	-5.33	119.77	121.90
84	Aa	2705	A	C5-C6-N1	-5.33	115.03	117.70
84	Aa	2883	C	N3-C4-N4	5.33	121.73	118.00
85	Ac	103	G	O4'-C1'-N9	5.33	112.47	108.20
86	Ab	28	U	C5-C4-O4	-5.33	122.70	125.90
84	Aa	1303	C	N3-C4-C5	-5.33	119.77	121.90
84	Aa	2532	A	O4'-C1'-N9	5.33	112.46	108.20
84	Aa	3012	A	C5-C6-N1	-5.33	115.03	117.70
1	Ad	180	A	O4'-C1'-C2'	-5.33	100.47	105.80
1	Ad	821	G	C1'-O4'-C4'	-5.33	105.64	109.90
1	Ad	1522	U	O4'-C1'-C2'	5.33	112.40	107.60
1	Ad	1606	U	O4'-C1'-N1	5.33	112.46	108.20
84	Aa	306	A	C4-C5-C6	5.33	119.66	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	783	A	C5-C6-N1	-5.33	115.04	117.70
84	Aa	1255	A	O4'-C1'-N9	5.33	112.46	108.20
84	Aa	1263	A	C5-C6-N1	-5.33	115.04	117.70
84	Aa	1676	A	O4'-C1'-N9	5.33	112.46	108.20
84	Aa	2789	G	C5'-C4'-C3'	-5.33	107.47	116.00
86	Ab	113	G	N1-C6-O6	5.33	123.10	119.90
1	Ad	1186	U	O4'-C1'-C2'	-5.33	100.47	105.80
1	Ad	1456	U	N1-C1'-C2'	5.33	120.92	114.00
84	Aa	1227	A	C5-C6-N6	-5.33	119.44	123.70
84	Aa	2102	C	N3-C4-C5	-5.33	119.77	121.90
84	Aa	3073	A	O4'-C1'-N9	5.33	112.46	108.20
84	Aa	3093	C	N3-C4-N4	5.33	121.73	118.00
84	Aa	3168	C	C6-N1-C2	-5.33	118.17	120.30
84	Aa	3278	G	P-O3'-C3'	5.33	126.09	119.70
1	Ad	41	A	C3'-C2'-C1'	5.33	105.76	101.50
1	Ad	229	G	P-O3'-C3'	5.33	126.09	119.70
1	Ad	1161	C	C3'-C2'-C1'	5.33	105.76	101.50
84	Aa	699	C	N3-C4-N4	5.33	121.73	118.00
84	Aa	1485	A	C5-C6-N1	-5.33	115.04	117.70
84	Aa	1580	C	N3-C4-N4	5.33	121.73	118.00
84	Aa	2706	A	C5-C6-N6	-5.33	119.44	123.70
84	Aa	2880	G	C5-C6-O6	-5.33	125.41	128.60
1	Ad	1650	G	O4'-C1'-C2'	5.32	112.39	107.60
84	Aa	326	C	N3-C4-C5	-5.32	119.77	121.90
84	Aa	680	G	O4'-C1'-N9	5.32	112.46	108.20
84	Aa	681	A	C4-C5-C6	5.32	119.66	117.00
84	Aa	1486	G	C6-C5-N7	-5.32	127.20	130.40
84	Aa	1633	C	N3-C4-C5	-5.32	119.77	121.90
84	Aa	2780	G	O4'-C1'-N9	5.32	112.46	108.20
84	Aa	2978	A	C5-C6-N1	-5.32	115.04	117.70
1	Ad	1109	U	N1-C1'-C2'	5.32	120.92	114.00
86	Ab	25	G	C5-C6-O6	-5.32	125.41	128.60
1	Ad	94	A	C5'-C4'-C3'	-5.32	107.49	116.00
1	Ad	194	G	P-O3'-C3'	5.32	126.08	119.70
1	Ad	261	C	C1'-O4'-C4'	-5.32	105.64	109.90
84	Aa	1551	C	N3-C4-C5	-5.32	119.77	121.90
84	Aa	1565	G	C5'-C4'-C3'	5.32	124.51	116.00
84	Aa	2044	C	C5'-C4'-C3'	-5.32	107.49	116.00
84	Aa	2455	A	O4'-C1'-N9	5.32	112.46	108.20
84	Aa	3152	C	C3'-C2'-C1'	5.32	105.76	101.50
84	Aa	819	A	C4-C5-C6	5.32	119.66	117.00
84	Aa	869	A	C4-C5-C6	5.32	119.66	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1457	A	C5-C6-N6	-5.32	119.44	123.70
84	Aa	1860	A	C5-C6-N1	-5.32	115.04	117.70
84	Aa	2137	A	C5-C6-N1	-5.32	115.04	117.70
85	Ac	25	G	O4'-C1'-N9	5.32	112.45	108.20
86	Ab	19	A	C5-C6-N6	-5.32	119.44	123.70
1	Ad	1281	G	N9-C1'-C2'	5.32	120.91	114.00
84	Aa	880	C	N3-C4-C5	-5.32	119.77	121.90
84	Aa	1208	A	C5-C6-N1	-5.32	115.04	117.70
84	Aa	1455	A	C5-C6-N1	-5.32	115.04	117.70
84	Aa	1806	C	C5'-C4'-O4'	5.32	115.48	109.10
84	Aa	2054	A	C5-C6-N6	-5.32	119.45	123.70
84	Aa	2061	C	N3-C4-C5	-5.32	119.77	121.90
84	Aa	2111	A	C5-C6-N6	-5.32	119.45	123.70
84	Aa	2150	C	N3-C4-N4	5.32	121.72	118.00
84	Aa	2874	A	C4-C5-C6	5.32	119.66	117.00
85	Ac	142	G	O4'-C1'-N9	5.32	112.45	108.20
1	Ad	1252	C	O4'-C1'-N1	5.32	112.45	108.20
1	Ad	1401	C	N1-C1'-C2'	5.32	120.91	114.00
47	CQ	161	SER	N-CA-C	-5.32	96.65	111.00
60	Co	48	SER	N-CA-CB	5.32	118.47	110.50
84	Aa	33	A	O4'-C1'-N9	5.32	112.45	108.20
84	Aa	316	A	C5-C6-N1	-5.32	115.04	117.70
84	Aa	763	G	O4'-C1'-N9	5.32	112.45	108.20
84	Aa	936	A	C5-C6-N1	-5.32	115.04	117.70
84	Aa	1763	C	N3-C4-C5	-5.32	119.77	121.90
84	Aa	2691	U	O4'-C1'-N1	5.32	112.45	108.20
84	Aa	3050	A	C5-C6-N6	-5.32	119.45	123.70
84	Aa	3121	C	N3-C4-C5	-5.32	119.77	121.90
84	Aa	3169	C	O4'-C1'-N1	5.32	112.45	108.20
84	Aa	202	G	O4'-C1'-N9	5.31	112.45	108.20
84	Aa	347	A	C5-C6-N1	-5.31	115.04	117.70
84	Aa	500	C	N3-C4-N4	5.31	121.72	118.00
84	Aa	1970	A	C5-C6-N1	-5.31	115.04	117.70
84	Aa	2801	A	O4'-C1'-N9	5.31	112.45	108.20
84	Aa	3134	C	N3-C4-N4	5.31	121.72	118.00
1	Ad	1091	A	C1'-O4'-C4'	-5.31	105.65	109.90
46	Ca	9	ARG	N-CA-CB	5.31	120.16	110.60
84	Aa	329	G	O4'-C1'-N9	5.31	112.45	108.20
84	Aa	1120	G	O4'-C1'-N9	5.31	112.45	108.20
84	Aa	1192	A	C5-C6-N6	-5.31	119.45	123.70
84	Aa	1856	G	O4'-C1'-N9	5.31	112.45	108.20
84	Aa	3311	C	N3-C4-N4	5.31	121.72	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	Ae	19	U	C3'-C2'-C1'	5.31	105.75	101.50
47	CQ	123	PHE	CB-CG-CD2	-5.31	117.08	120.80
84	Aa	1645	G	O4'-C1'-N9	5.31	112.45	108.20
84	Aa	1861	A	C5-C6-N6	-5.31	119.45	123.70
84	Aa	3253	C	N3-C4-N4	5.31	121.72	118.00
85	Ac	128	C	N3-C4-C5	-5.31	119.78	121.90
1	Ad	865	U	N1-C1'-C2'	5.31	120.90	114.00
1	Ad	1173	U	O4'-C1'-N1	5.31	112.45	108.20
84	Aa	286	C	C4'-C3'-C2'	-5.31	97.29	102.60
84	Aa	365	A	C5-C6-N1	-5.31	115.05	117.70
84	Aa	794	G	O4'-C1'-N9	5.31	112.45	108.20
84	Aa	806	C	N3-C4-N4	5.31	121.72	118.00
84	Aa	1051	A	C5-C6-N6	-5.31	119.45	123.70
84	Aa	1278	A	C5-C6-N6	-5.31	119.45	123.70
84	Aa	1396	A	O4'-C1'-N9	5.31	112.45	108.20
84	Aa	1590	A	C5-C6-N1	-5.31	115.05	117.70
84	Aa	3011	U	C5'-C4'-C3'	5.31	124.50	116.00
2	Ae	74	C	N1-C1'-C2'	-5.31	106.16	112.00
84	Aa	705	A	C5-C6-N1	-5.31	115.05	117.70
84	Aa	1896	A	O4'-C1'-N9	5.31	112.44	108.20
84	Aa	2092	C	C4'-C3'-C2'	5.31	107.91	102.60
84	Aa	3033	A	C5-C6-N1	-5.31	115.05	117.70
84	Aa	3165	C	N3-C4-N4	5.31	121.72	118.00
84	Aa	1306	A	C5-C6-N1	-5.31	115.05	117.70
84	Aa	1865	C	N3-C4-N4	5.31	121.71	118.00
84	Aa	2035	G	C5-C6-O6	-5.31	125.42	128.60
84	Aa	2037	C	N3-C4-C5	-5.31	119.78	121.90
1	Ad	167	A	N9-C1'-C2'	-5.30	106.17	112.00
1	Ad	941	G	P-O3'-C3'	5.30	126.07	119.70
1	Ad	995	C	O4'-C1'-N1	5.30	112.44	108.20
1	Ad	1511	A	O4'-C1'-C2'	5.30	112.37	107.60
1	Ad	1701	G	C3'-C2'-C1'	-5.30	97.26	101.50
53	CY	10	SER	N-CA-CB	5.30	118.45	110.50
84	Aa	11	A	O4'-C1'-N9	5.30	112.44	108.20
84	Aa	249	A	C5-C6-N6	-5.30	119.46	123.70
84	Aa	363	A	C5-C6-N6	-5.30	119.46	123.70
84	Aa	1233	G	O4'-C1'-N9	5.30	112.44	108.20
84	Aa	1290	A	C4-C5-C6	5.30	119.65	117.00
84	Aa	1961	C	N3-C4-N4	5.30	121.71	118.00
84	Aa	2059	C	N3-C4-N4	5.30	121.71	118.00
84	Aa	2167	G	C6-C5-N7	-5.30	127.22	130.40
84	Aa	2533	A	C5-C6-N1	-5.30	115.05	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2839	A	C5-C6-N1	-5.30	115.05	117.70
84	Aa	3184	G	C5-C6-O6	-5.30	125.42	128.60
85	Ac	21	C	N3-C4-N4	5.30	121.71	118.00
85	Ac	48	A	C4-C5-C6	5.30	119.65	117.00
85	Ac	53	A	C5-C6-N1	-5.30	115.05	117.70
86	Ab	3	A	O4'-C1'-N9	5.30	112.44	108.20
86	Ab	7	G	N3-C2-N2	5.30	123.61	119.90
84	Aa	296	C	N3-C4-C5	-5.30	119.78	121.90
84	Aa	535	G	O4'-C1'-N9	5.30	112.44	108.20
84	Aa	2462	G	O4'-C4'-C3'	-5.30	98.70	104.00
84	Aa	2902	A	C4-C5-C6	5.30	119.65	117.00
84	Aa	336	A	C5-C6-N1	-5.30	115.05	117.70
84	Aa	661	A	C5-C6-N6	-5.30	119.46	123.70
84	Aa	859	G	O4'-C1'-N9	5.30	112.44	108.20
84	Aa	1517	C	C6-N1-C1'	-5.30	114.44	120.80
84	Aa	1875	A	C5-C6-N1	-5.30	115.05	117.70
84	Aa	2146	A	C4-C5-C6	5.30	119.65	117.00
84	Aa	2325	A	O4'-C1'-N9	5.30	112.44	108.20
84	Aa	2774	A	O4'-C1'-N9	5.30	112.44	108.20
84	Aa	2896	C	N3-C4-C5	-5.30	119.78	121.90
84	Aa	3022	A	C5-C6-N1	-5.30	115.05	117.70
8	Bf	56	PHE	CB-CG-CD2	5.30	124.51	120.80
84	Aa	416	A	C5-C6-N1	-5.30	115.05	117.70
84	Aa	632	C	N3-C4-N4	5.30	121.71	118.00
84	Aa	924	A	C4-C5-C6	5.30	119.65	117.00
84	Aa	1207	A	O4'-C1'-N9	5.30	112.44	108.20
84	Aa	1550	A	C5-C6-N6	-5.30	119.46	123.70
84	Aa	2135	U	O4'-C1'-N1	5.30	112.44	108.20
84	Aa	2240	C	O4'-C1'-N1	5.30	112.44	108.20
84	Aa	2298	A	C5-C6-N6	-5.30	119.46	123.70
84	Aa	804	A	C4-C5-C6	5.30	119.65	117.00
84	Aa	1006	A	O4'-C1'-N9	5.30	112.44	108.20
84	Aa	2441	G	O4'-C1'-N9	5.30	112.44	108.20
84	Aa	2474	A	C5-C6-N1	-5.30	115.05	117.70
1	Ad	721	U	P-O5'-C5'	-5.30	112.42	120.90
1	Ad	746	A	N9-C1'-C2'	-5.30	106.17	112.00
1	Ad	1270	U	C1'-O4'-C4'	-5.30	105.66	109.90
2	Ae	70	G	O4'-C1'-N9	5.30	112.44	108.20
11	BD	78	ASN	N-CA-CB	5.30	120.13	110.60
84	Aa	12	G	C5-C6-O6	-5.30	125.42	128.60
84	Aa	167	C	N3-C4-C5	-5.30	119.78	121.90
84	Aa	410	G	N1-C6-O6	5.30	123.08	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	720	G	O4'-C1'-N9	5.30	112.44	108.20
84	Aa	819	A	C5-C6-N1	-5.30	115.05	117.70
84	Aa	928	A	C5-C6-N1	-5.30	115.05	117.70
84	Aa	949	C	C2-N3-C4	5.30	122.55	119.90
84	Aa	1036	C	C2-N3-C4	5.30	122.55	119.90
84	Aa	1267	A	C5-C6-N1	-5.30	115.05	117.70
84	Aa	1323	G	O4'-C1'-N9	5.30	112.44	108.20
84	Aa	1608	C	N3-C4-N4	5.30	121.71	118.00
84	Aa	2141	A	C5-C6-N6	-5.30	119.46	123.70
84	Aa	2399	G	O4'-C1'-N9	5.30	112.44	108.20
84	Aa	3162	C	P-O3'-C3'	5.30	126.06	119.70
84	Aa	3256	C	N3-C4-N4	5.30	121.71	118.00
84	Aa	3336	A	C1'-O4'-C4'	-5.30	105.66	109.90
84	Aa	597	C	N3-C4-N4	5.29	121.71	118.00
84	Aa	1430	C	N3-C4-C5	-5.29	119.78	121.90
84	Aa	1846	A	C5-C6-N1	-5.29	115.05	117.70
84	Aa	3072	A	C5-C6-N1	-5.29	115.05	117.70
84	Aa	2064	C	N3-C4-C5	-5.29	119.78	121.90
84	Aa	2137	A	C5-C6-N6	-5.29	119.47	123.70
84	Aa	2185	U	O4'-C1'-N1	5.29	112.44	108.20
84	Aa	2561	A	C5-C6-N1	-5.29	115.05	117.70
84	Aa	3078	A	C5-C6-N1	-5.29	115.05	117.70
1	Ad	1079	G	N9-C1'-C2'	5.29	120.88	114.00
1	Ad	1299	G	O4'-C4'-C3'	-5.29	98.71	104.00
2	Ae	5	U	O4'-C1'-C2'	-5.29	100.51	105.80
84	Aa	143	A	O4'-C1'-N9	5.29	112.43	108.20
84	Aa	825	G	O4'-C1'-N9	5.29	112.43	108.20
84	Aa	1172	A	O4'-C1'-N9	5.29	112.43	108.20
84	Aa	1800	G	P-O3'-C3'	5.29	126.05	119.70
84	Aa	1860	A	C5-C6-N6	-5.29	119.47	123.70
84	Aa	2885	U	O4'-C1'-N1	5.29	112.43	108.20
84	Aa	3186	G	O4'-C1'-N9	5.29	112.43	108.20
85	Ac	94	C	N3-C4-N4	5.29	121.70	118.00
3	Af	20	U	N1-C1'-C2'	5.29	120.88	114.00
84	Aa	1177	G	O4'-C1'-N9	5.29	112.43	108.20
84	Aa	1577	A	C5-C6-N1	-5.29	115.06	117.70
84	Aa	3166	C	N3-C4-N4	5.29	121.70	118.00
84	Aa	3168	C	N3-C4-N4	5.29	121.70	118.00
85	Ac	22	U	O4'-C1'-N1	5.29	112.43	108.20
86	Ab	5	G	N3-C2-N2	5.29	123.60	119.90
1	Ad	281	U	C5'-C4'-O4'	-5.29	102.75	109.10
1	Ad	1185	U	N1-C1'-C2'	5.29	120.87	114.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	Ae	57	A	O4'-C1'-C2'	-5.29	100.51	105.80
84	Aa	1351	C	C2-N3-C4	5.29	122.54	119.90
84	Aa	1438	A	C5-C6-N1	-5.29	115.06	117.70
84	Aa	1512	A	O4'-C1'-N9	5.29	112.43	108.20
84	Aa	1928	A	C5-C6-N6	-5.29	119.47	123.70
84	Aa	2120	A	O4'-C1'-N9	5.29	112.43	108.20
84	Aa	2815	A	C5'-C4'-C3'	5.29	124.46	116.00
84	Aa	3270	C	C6-N1-C1'	-5.29	114.45	120.80
84	Aa	3307	A	C5-C6-N1	-5.29	115.06	117.70
85	Ac	153	C	N3-C4-C5	-5.29	119.78	121.90
86	Ab	38	U	C5-C4-O4	-5.29	122.73	125.90
1	Ad	236	U	P-O3'-C3'	5.29	126.05	119.70
84	Aa	177	C	C2-N3-C4	5.29	122.54	119.90
84	Aa	381	G	O4'-C1'-N9	5.29	112.43	108.20
84	Aa	759	C	N3-C4-N4	5.29	121.70	118.00
84	Aa	1119	G	O4'-C1'-N9	5.29	112.43	108.20
84	Aa	2104	G	N1-C2-N3	-5.29	120.73	123.90
84	Aa	2316	A	C5-C6-N1	-5.29	115.06	117.70
84	Aa	52	G	C5'-C4'-C3'	-5.29	107.54	116.00
84	Aa	149	A	O4'-C1'-N9	5.29	112.43	108.20
84	Aa	407	A	C2'-C3'-O3'	5.29	122.16	113.70
84	Aa	861	A	C5-C6-N1	-5.29	115.06	117.70
84	Aa	885	A	C5-C6-N6	-5.29	119.47	123.70
84	Aa	966	G	O4'-C1'-N9	5.29	112.43	108.20
84	Aa	1360	U	P-O3'-C3'	5.29	126.04	119.70
84	Aa	1726	G	C1'-O4'-C4'	-5.29	105.67	109.90
84	Aa	2449	A	O4'-C1'-N9	5.29	112.43	108.20
85	Ac	139	C	N3-C4-N4	5.29	121.70	118.00
86	Ab	11	A	N3-C4-C5	-5.29	123.10	126.80
1	Ad	1657	C	C5'-C4'-O4'	5.28	115.44	109.10
84	Aa	636	C	N3-C4-N4	5.28	121.70	118.00
84	Aa	664	A	C4-C5-C6	5.28	119.64	117.00
84	Aa	1032	C	N3-C4-N4	5.28	121.70	118.00
84	Aa	2275	A	C5-C6-N6	-5.28	119.47	123.70
84	Aa	2323	A	C5-C6-N1	-5.28	115.06	117.70
84	Aa	2763	C	N3-C4-N4	5.28	121.70	118.00
84	Aa	3351	A	O4'-C1'-N9	5.28	112.43	108.20
85	Ac	12	A	C4-C5-C6	5.28	119.64	117.00
1	Ad	17	C	N1-C1'-C2'	5.28	120.87	114.00
84	Aa	322	A	C5-C6-N1	-5.28	115.06	117.70
84	Aa	1341	G	C5-C6-O6	-5.28	125.43	128.60
84	Aa	2229	G	O4'-C1'-N9	5.28	112.42	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	3	C	C1'-O4'-C4'	5.28	114.12	109.90
1	Ad	1355	U	C3'-C2'-C1'	5.28	105.72	101.50
84	Aa	97	G	P-O5'-C5'	-5.28	112.45	120.90
84	Aa	737	C	N3-C4-N4	5.28	121.70	118.00
84	Aa	861	A	C5-C6-N6	-5.28	119.47	123.70
84	Aa	2214	A	O4'-C1'-N9	5.28	112.42	108.20
84	Aa	2772	A	C5-C6-N6	-5.28	119.47	123.70
84	Aa	3241	C	N3-C4-N4	5.28	121.70	118.00
1	Ad	1311	U	O4'-C1'-C2'	-5.28	100.52	105.80
1	Ad	1478	C	O4'-C1'-C2'	-5.28	100.52	105.80
54	Cr	93	ARG	N-CA-CB	5.28	120.10	110.60
84	Aa	74	G	N3-C2-N2	5.28	123.59	119.90
84	Aa	1173	C	N3-C4-C5	-5.28	119.79	121.90
84	Aa	1302	C	N3-C4-N4	5.28	121.69	118.00
84	Aa	30	C	N3-C4-C5	-5.28	119.79	121.90
84	Aa	96	C	N3-C4-N4	5.28	121.69	118.00
84	Aa	345	G	O4'-C1'-N9	5.28	112.42	108.20
84	Aa	443	G	C5-C6-O6	-5.28	125.43	128.60
84	Aa	1755	A	C5-C6-N1	-5.28	115.06	117.70
84	Aa	1848	G	C5-C6-O6	-5.28	125.43	128.60
84	Aa	3152	C	N3-C4-C5	-5.28	119.79	121.90
86	Ab	27	A	N7-C8-N9	-5.28	111.16	113.80
1	Ad	843	G	P-O5'-C5'	5.28	129.34	120.90
2	Ae	27	G	C3'-C2'-C1'	-5.28	97.28	101.50
84	Aa	48	A	C5-C6-N6	-5.28	119.48	123.70
84	Aa	397	A	C5-C6-N1	-5.28	115.06	117.70
84	Aa	1002	A	C5-C6-N1	-5.28	115.06	117.70
84	Aa	1041	C	N3-C4-N4	5.28	121.69	118.00
84	Aa	1135	C	N3-C4-N4	5.28	121.69	118.00
84	Aa	1591	A	C5-C6-N1	-5.28	115.06	117.70
84	Aa	1736	C	N3-C4-C5	-5.28	119.79	121.90
84	Aa	2222	C	N3-C4-N4	5.28	121.69	118.00
84	Aa	2560	C	N3-C4-C5	-5.28	119.79	121.90
84	Aa	2718	A	C5-C6-N1	-5.28	115.06	117.70
84	Aa	2778	C	N3-C4-N4	5.28	121.69	118.00
84	Aa	2911	C	N3-C4-N4	5.28	121.69	118.00
1	Ad	73	A	O4'-C1'-C2'	-5.27	100.53	105.80
1	Ad	153	U	N1-C1'-C2'	-5.27	106.20	112.00
1	Ad	238	G	C5'-C4'-O4'	5.27	115.43	109.10
84	Aa	8	C	N3-C4-N4	5.27	121.69	118.00
84	Aa	1213	G	O4'-C1'-N9	5.27	112.42	108.20
84	Aa	1595	G	N3-C2-N2	5.27	123.59	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2492	C	N3-C4-C5	-5.27	119.79	121.90
84	Aa	2504	A	O4'-C1'-N9	5.27	112.42	108.20
84	Aa	2891	C	C2-N3-C4	5.27	122.54	119.90
86	Ab	103	U	N3-C4-O4	5.27	123.09	119.40
1	Ad	322	U	C3'-C2'-C1'	5.27	105.72	101.50
1	Ad	1808	U	C3'-C2'-C1'	5.27	105.72	101.50
51	CX	50	LYS	N-CA-CB	5.27	120.09	110.60
84	Aa	684	C	C2-N3-C4	5.27	122.54	119.90
84	Aa	896	C	N3-C4-N4	5.27	121.69	118.00
84	Aa	948	C	N3-C4-N4	5.27	121.69	118.00
84	Aa	1014	G	C5-C6-O6	-5.27	125.44	128.60
84	Aa	1097	A	O4'-C1'-N9	5.27	112.42	108.20
84	Aa	1908	C	N3-C4-C5	-5.27	119.79	121.90
84	Aa	2202	A	C5-C6-N6	-5.27	119.48	123.70
84	Aa	2434	G	C5-C6-O6	-5.27	125.44	128.60
84	Aa	3151	C	O4'-C1'-N1	5.27	112.42	108.20
84	Aa	3308	A	C5-C6-N1	-5.27	115.06	117.70
84	Aa	810	A	C5-C6-N1	-5.27	115.06	117.70
84	Aa	2324	G	C5-C6-O6	-5.27	125.44	128.60
84	Aa	2688	G	C5-C6-O6	-5.27	125.44	128.60
84	Aa	3362	A	C5-C6-N6	-5.27	119.48	123.70
1	Ad	18	C	O4'-C1'-N1	5.27	112.42	108.20
1	Ad	1038	C	C3'-C2'-C1'	5.27	105.72	101.50
1	Ad	1763	A	N9-C1'-C2'	-5.27	106.20	112.00
84	Aa	1518	A	C5-C6-N6	-5.27	119.48	123.70
84	Aa	2225	C	N3-C4-C5	-5.27	119.79	121.90
84	Aa	2966	G	O4'-C1'-N9	5.27	112.42	108.20
84	Aa	3115	A	C5-C6-N1	-5.27	115.06	117.70
86	Ab	99	G	N3-C2-N2	5.27	123.59	119.90
84	Aa	1358	C	N3-C4-C5	-5.27	119.79	121.90
84	Aa	1399	C	N3-C4-N4	5.27	121.69	118.00
85	Ac	128	C	N3-C4-N4	5.27	121.69	118.00
85	Ac	148	C	N3-C4-C5	-5.27	119.79	121.90
1	Ad	646	G	O4'-C1'-N9	5.27	112.41	108.20
84	Aa	220	G	O4'-C1'-N9	5.27	112.41	108.20
84	Aa	1675	G	C3'-C2'-C1'	-5.27	97.29	101.50
84	Aa	2660	A	C4-C5-C6	5.27	119.63	117.00
85	Ac	160	C	N3-C4-C5	-5.27	119.79	121.90
84	Aa	327	A	C5-C6-N1	-5.26	115.07	117.70
84	Aa	1219	C	N3-C4-C5	-5.26	119.79	121.90
84	Aa	1373	A	C5-C6-N1	-5.26	115.07	117.70
84	Aa	1468	A	C5-C6-N1	-5.26	115.07	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2765	A	O4'-C1'-N9	5.26	112.41	108.20
84	Aa	3147	G	N3-C2-N2	5.26	123.58	119.90
86	Ab	29	C	N3-C4-N4	5.26	121.69	118.00
1	Ad	1086	A	C3'-C2'-C1'	5.26	105.71	101.50
28	BA	108	THR	N-CA-CB	5.26	120.30	110.30
84	Aa	1088	A	C4-C5-C6	5.26	119.63	117.00
86	Ab	39	C	N3-C4-N4	5.26	121.68	118.00
34	BC	39	THR	N-CA-CB	5.26	120.30	110.30
84	Aa	197	A	C5-C6-N1	-5.26	115.07	117.70
84	Aa	250	C	N3-C4-C5	-5.26	119.80	121.90
84	Aa	529	C	C6-N1-C2	-5.26	118.20	120.30
84	Aa	1507	A	O4'-C1'-N9	5.26	112.41	108.20
84	Aa	2985	C	N3-C4-N4	5.26	121.68	118.00
84	Aa	3316	C	N3-C4-C5	-5.26	119.80	121.90
85	Ac	94	C	N3-C4-C5	-5.26	119.80	121.90
1	Ad	388	G	N9-C1'-C2'	5.26	120.84	114.00
1	Ad	1065	A	C3'-C2'-C1'	-5.26	97.29	101.50
65	CK	95	LYS	C-N-CA	5.26	134.85	121.70
84	Aa	165	C	N3-C4-N4	5.26	121.68	118.00
84	Aa	424	G	N3-C2-N2	5.26	123.58	119.90
84	Aa	764	A	O4'-C1'-N9	5.26	112.41	108.20
84	Aa	1682	C	N3-C4-N4	5.26	121.68	118.00
84	Aa	2075	C	N3-C4-N4	5.26	121.68	118.00
84	Aa	2943	A	C5-C6-N1	-5.26	115.07	117.70
86	Ab	87	G	C6-C5-N7	-5.26	127.24	130.40
1	Ad	877	G	O4'-C1'-N9	5.26	112.41	108.20
84	Aa	48	A	C5-C6-N1	-5.26	115.07	117.70
84	Aa	2275	A	C5-C6-N1	-5.26	115.07	117.70
1	Ad	230	C	O4'-C1'-C2'	-5.26	100.54	105.80
1	Ad	1314	U	O4'-C4'-C3'	-5.26	98.74	104.00
37	CG	47	PHE	CB-CG-CD2	-5.26	117.12	120.80
45	CN	4	TYR	CB-CG-CD2	5.26	124.15	121.00
84	Aa	392	C	N3-C4-N4	5.26	121.68	118.00
84	Aa	594	C	C2-N3-C4	5.26	122.53	119.90
84	Aa	724	A	C5'-C4'-C3'	5.26	124.41	116.00
84	Aa	840	A	C4-C5-C6	5.26	119.63	117.00
84	Aa	1301	C	N3-C4-C5	-5.26	119.80	121.90
84	Aa	1707	C	N3-C4-N4	5.26	121.68	118.00
84	Aa	1874	A	C5-C6-N1	-5.26	115.07	117.70
84	Aa	1900	C	N3-C4-C5	-5.26	119.80	121.90
84	Aa	2362	A	O4'-C1'-N9	5.26	112.41	108.20
84	Aa	2398	A	C4-C5-C6	5.26	119.63	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2868	C	P-O3'-C3'	5.26	126.01	119.70
84	Aa	3151	C	N3-C4-C5	-5.26	119.80	121.90
84	Aa	3170	C	N3-C4-N4	5.26	121.68	118.00
84	Aa	3312	G	O4'-C1'-N9	5.26	112.41	108.20
1	Ad	439	C	C3'-C2'-C1'	5.25	105.70	101.50
84	Aa	165	C	N3-C4-C5	-5.25	119.80	121.90
84	Aa	344	C	N3-C4-C5	-5.25	119.80	121.90
84	Aa	2173	G	C8-N9-C1'	5.25	133.83	127.00
84	Aa	3268	C	N3-C4-C5	-5.25	119.80	121.90
1	Ad	1693	C	C3'-C2'-C1'	5.25	105.70	101.50
84	Aa	318	G	N3-C2-N2	5.25	123.58	119.90
84	Aa	697	A	C4-C5-C6	5.25	119.63	117.00
84	Aa	2544	C	N3-C4-C5	-5.25	119.80	121.90
84	Aa	3067	G	C5'-C4'-O4'	5.25	115.40	109.10
84	Aa	3096	U	O4'-C1'-N1	5.25	112.40	108.20
1	Ad	831	C	O4'-C1'-C2'	-5.25	100.55	105.80
1	Ad	1448	U	C3'-C2'-C1'	5.25	105.70	101.50
1	Ad	1517	C	N1-C1'-C2'	5.25	120.83	114.00
84	Aa	113	A	O4'-C1'-N9	5.25	112.40	108.20
84	Aa	585	A	C5-C6-N1	-5.25	115.08	117.70
84	Aa	686	A	C5-C6-N1	-5.25	115.07	117.70
84	Aa	774	A	C4-C5-C6	5.25	119.62	117.00
84	Aa	1655	G	O4'-C1'-N9	5.25	112.40	108.20
84	Aa	2023	C	N3-C4-N4	5.25	121.68	118.00
84	Aa	2182	G	P-O3'-C3'	5.25	126.00	119.70
84	Aa	2197	C	N3-C4-C5	-5.25	119.80	121.90
84	Aa	3071	A	C5-C6-N6	-5.25	119.50	123.70
84	Aa	3171	C	C6-N1-C1'	-5.25	114.50	120.80
85	Ac	116	G	C5-C6-O6	-5.25	125.45	128.60
1	Ad	974	C	C3'-C2'-C1'	5.25	105.70	101.50
84	Aa	1499	C	C6-N1-C2	-5.25	118.20	120.30
84	Aa	3006	G	O4'-C1'-N9	5.25	112.40	108.20
72	CC	90	ARG	N-CA-CB	5.25	120.05	110.60
84	Aa	382	A	C5-C6-N1	-5.25	115.08	117.70
84	Aa	1291	A	O4'-C1'-N9	5.25	112.40	108.20
84	Aa	1607	C	N3-C4-N4	5.25	121.67	118.00
84	Aa	2113	A	C5-C6-N6	-5.25	119.50	123.70
84	Aa	2414	C	N3-C4-N4	5.25	121.67	118.00
84	Aa	2642	G	O4'-C1'-N9	5.25	112.40	108.20
84	Aa	3356	C	N3-C4-N4	5.25	121.67	118.00
84	Aa	3374	C	N3-C4-N4	5.25	121.67	118.00
1	Ad	1188	A	C1'-O4'-C4'	5.25	114.10	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	124	C	N3-C4-N4	5.25	121.67	118.00
84	Aa	340	A	C4-C5-C6	5.25	119.62	117.00
84	Aa	1316	C	C2-N1-C1'	5.25	124.57	118.80
84	Aa	1704	A	O4'-C1'-N9	5.25	112.40	108.20
84	Aa	1733	G	C5-C6-O6	-5.25	125.45	128.60
84	Aa	2248	G	O4'-C1'-N9	5.25	112.40	108.20
1	Ad	1806	C	C5'-C4'-O4'	5.25	115.39	109.10
2	Ae	58	U	P-O5'-C5'	5.25	129.29	120.90
53	CY	73	TYR	CB-CG-CD1	-5.25	117.85	121.00
84	Aa	82	C	N3-C4-C5	-5.25	119.80	121.90
84	Aa	119	A	C5-C6-N1	-5.25	115.08	117.70
84	Aa	620	C	N3-C4-N4	5.25	121.67	118.00
84	Aa	964	C	N3-C4-N4	5.25	121.67	118.00
84	Aa	2574	A	C5-C6-N1	-5.25	115.08	117.70
86	Ab	104	C	C5-C4-N4	-5.25	116.53	120.20
1	Ad	25	C	C1'-O4'-C4'	-5.24	105.70	109.90
1	Ad	104	A	O4'-C1'-C2'	-5.24	100.56	105.80
2	Ae	44	A	C1'-O4'-C4'	5.24	114.09	109.90
84	Aa	1196	U	O4'-C1'-N1	5.24	112.39	108.20
84	Aa	1391	A	C5-C6-N6	-5.24	119.51	123.70
84	Aa	1587	G	O4'-C1'-N9	5.24	112.39	108.20
84	Aa	3109	G	N3-C2-N2	5.24	123.57	119.90
1	Ad	927	A	C4'-C3'-C2'	-5.24	97.36	102.60
1	Ad	1263	C	P-O3'-C3'	5.24	125.99	119.70
1	Ad	1410	C	C3'-C2'-C1'	5.24	105.69	101.50
1	Ad	1627	C	N1-C1'-C2'	5.24	120.81	114.00
84	Aa	72	A	C5-C6-N6	-5.24	119.51	123.70
84	Aa	458	G	C5-C6-O6	-5.24	125.45	128.60
84	Aa	497	G	O4'-C1'-N9	5.24	112.39	108.20
84	Aa	962	C	N3-C4-N4	5.24	121.67	118.00
84	Aa	968	A	C5-C6-N6	-5.24	119.51	123.70
84	Aa	1421	A	C4-C5-C6	5.24	119.62	117.00
84	Aa	1550	A	C5-C6-N1	-5.24	115.08	117.70
15	BU	11	PRO	C-N-CA	5.24	134.80	121.70
84	Aa	585	A	C5-C6-N6	-5.24	119.51	123.70
84	Aa	678	G	O4'-C1'-N9	5.24	112.39	108.20
84	Aa	1047	C	N3-C4-C5	-5.24	119.80	121.90
84	Aa	1052	A	C4-C5-C6	5.24	119.62	117.00
84	Aa	3290	C	N3-C4-N4	5.24	121.67	118.00
1	Ad	269	A	C3'-C2'-C1'	-5.24	97.31	101.50
1	Ad	560	A	O4'-C1'-N9	5.24	112.39	108.20
84	Aa	12	G	N1-C6-O6	5.24	123.04	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	353	A	C4-C5-C6	5.24	119.62	117.00
84	Aa	426	A	O4'-C1'-N9	5.24	112.39	108.20
84	Aa	486	G	N3-C2-N2	5.24	123.57	119.90
84	Aa	2648	G	O4'-C1'-N9	5.24	112.39	108.20
84	Aa	2931	C	N3-C4-N4	5.24	121.67	118.00
84	Aa	3061	C	N3-C4-N4	5.24	121.67	118.00
72	CC	110	LYS	N-CA-CB	5.24	120.03	110.60
84	Aa	2277	U	O4'-C1'-N1	5.24	112.39	108.20
84	Aa	2362	A	C5-C6-N6	-5.24	119.51	123.70
84	Aa	2531	G	C5-C6-O6	-5.24	125.46	128.60
84	Aa	2958	A	C4-C5-C6	5.24	119.62	117.00
1	Ad	138	C	O4'-C1'-N1	5.24	112.39	108.20
1	Ad	700	C	O4'-C1'-N1	5.24	112.39	108.20
84	Aa	464	G	P-O3'-C3'	5.24	125.98	119.70
84	Aa	693	C	N3-C4-N4	5.24	121.67	118.00
84	Aa	962	C	N3-C4-C5	-5.24	119.81	121.90
84	Aa	1568	A	O4'-C1'-N9	5.24	112.39	108.20
84	Aa	1697	G	N3-C2-N2	5.24	123.56	119.90
84	Aa	2133	A	C5-C6-N6	-5.24	119.51	123.70
84	Aa	2496	U	C2'-C3'-O3'	5.24	122.08	113.70
84	Aa	2874	A	C5-C6-N1	-5.24	115.08	117.70
84	Aa	3047	A	C4-C5-C6	5.24	119.62	117.00
85	Ac	36	G	O4'-C1'-N9	5.24	112.39	108.20
86	Ab	120	C	N1-C2-O2	5.24	122.04	118.90
1	Ad	401	A	O4'-C1'-N9	5.23	112.39	108.20
84	Aa	2227	A	C5-C6-N1	-5.23	115.08	117.70
1	Ad	991	G	C4'-C3'-C2'	-5.23	97.37	102.60
1	Ad	1210	U	P-O3'-C3'	5.23	125.98	119.70
37	CG	222	PHE	CB-CG-CD2	5.23	124.46	120.80
69	CF	77	PHE	CB-CG-CD2	-5.23	117.14	120.80
84	Aa	142	G	O4'-C1'-N9	5.23	112.39	108.20
84	Aa	527	G	O4'-C1'-N9	5.23	112.39	108.20
84	Aa	1874	A	C5-C6-N6	-5.23	119.51	123.70
84	Aa	1883	A	C5-C6-N6	-5.23	119.52	123.70
84	Aa	1958	G	N9-C1'-C2'	-5.23	106.25	112.00
84	Aa	2498	C	N3-C4-N4	5.23	121.66	118.00
84	Aa	2597	C	N3-C4-C5	-5.23	119.81	121.90
84	Aa	2695	A	C5-C6-N1	-5.23	115.08	117.70
1	Ad	776	A	C1'-O4'-C4'	5.23	114.08	109.90
1	Ad	1676	G	O4'-C1'-N9	5.23	112.39	108.20
7	BM	117	GLU	N-CA-CB	5.23	120.01	110.60
64	Ci	95	SER	N-CA-C	-5.23	96.88	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	965	A	C4-C5-C6	5.23	119.62	117.00
1	Ad	1494	G	P-O3'-C3'	5.23	125.97	119.70
84	Aa	38	A	C4-C5-C6	5.23	119.61	117.00
84	Aa	1905	A	C5-C6-N6	-5.23	119.52	123.70
84	Aa	2247	A	C5-C6-N1	-5.23	115.08	117.70
84	Aa	3190	U	C6-N1-C1'	-5.23	113.88	121.20
1	Ad	1256	C	O4'-C1'-N1	5.23	112.38	108.20
84	Aa	326	C	N3-C4-N4	5.23	121.66	118.00
84	Aa	562	G	O4'-C1'-N9	5.23	112.38	108.20
84	Aa	1139	A	C5-C6-N6	-5.23	119.52	123.70
84	Aa	1423	C	N3-C4-C5	-5.23	119.81	121.90
84	Aa	2007	C	N3-C4-C5	-5.23	119.81	121.90
84	Aa	2429	A	C5-C6-N6	-5.23	119.52	123.70
1	Ad	100	C	N1-C1'-C2'	-5.23	106.25	112.00
84	Aa	512	G	N1-C2-N3	-5.23	120.76	123.90
84	Aa	1179	C	N3-C4-C5	-5.23	119.81	121.90
84	Aa	1458	U	O4'-C1'-N1	5.23	112.38	108.20
84	Aa	1744	C	N3-C4-N4	5.23	121.66	118.00
84	Aa	2262	C	N3-C4-N4	5.23	121.66	118.00
84	Aa	3374	C	N3-C4-C5	-5.23	119.81	121.90
86	Ab	74	A	P-O3'-C3'	5.23	125.97	119.70
84	Aa	967	G	C5'-C4'-C3'	-5.22	107.64	116.00
84	Aa	1007	A	O4'-C1'-N9	5.22	112.38	108.20
84	Aa	1440	C	N3-C4-N4	5.22	121.66	118.00
84	Aa	1615	G	C5-C6-O6	-5.22	125.47	128.60
84	Aa	1795	A	C5-C6-N6	-5.22	119.52	123.70
84	Aa	2209	A	C4-C5-C6	5.22	119.61	117.00
84	Aa	2565	C	N3-C4-C5	-5.22	119.81	121.90
84	Aa	3307	A	C4-C5-C6	5.22	119.61	117.00
86	Ab	29	C	C2'-C3'-O3'	5.22	122.06	113.70
1	Ad	420	A	O4'-C1'-N9	5.22	112.38	108.20
1	Ad	545	A	P-O5'-C5'	-5.22	112.54	120.90
1	Ad	649	C	O4'-C1'-N1	5.22	112.38	108.20
1	Ad	1672	U	N1-C1'-C2'	5.22	120.79	114.00
84	Aa	510	C	N3-C4-N4	5.22	121.66	118.00
84	Aa	750	G	C5-C6-O6	-5.22	125.47	128.60
84	Aa	979	C	N3-C4-N4	5.22	121.66	118.00
84	Aa	1258	C	P-O5'-C5'	-5.22	112.55	120.90
84	Aa	1376	A	O4'-C1'-N9	5.22	112.38	108.20
84	Aa	1408	C	N3-C4-N4	5.22	121.66	118.00
84	Aa	2590	C	N3-C4-C5	-5.22	119.81	121.90
1	Ad	906	G	O4'-C1'-C2'	5.22	112.30	107.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1594	A	C5'-C4'-O4'	5.22	115.36	109.10
84	Aa	981	A	O4'-C1'-N9	5.22	112.38	108.20
84	Aa	1571	A	C5-C6-N1	-5.22	115.09	117.70
84	Aa	1880	A	C5-C6-N6	-5.22	119.52	123.70
84	Aa	1950	G	O3'-P-O5'	5.22	113.92	104.00
84	Aa	2473	C	N3-C4-C5	-5.22	119.81	121.90
84	Aa	3334	A	C4-C5-C6	5.22	119.61	117.00
1	Ad	162	A	O4'-C1'-C2'	5.22	112.30	107.60
1	Ad	521	U	C4'-C3'-C2'	-5.22	97.38	102.60
1	Ad	534	C	O4'-C1'-N1	5.22	112.38	108.20
3	Af	16	G	C4'-C3'-C2'	-5.22	97.38	102.60
69	CF	110	LEU	N-CA-C	-5.22	96.91	111.00
84	Aa	721	A	P-O5'-C5'	5.22	129.25	120.90
84	Aa	1010	A	O4'-C1'-N9	5.22	112.38	108.20
84	Aa	1737	C	N3-C4-C5	-5.22	119.81	121.90
84	Aa	2953	G	C5-C6-O6	-5.22	125.47	128.60
1	Ad	1184	C	P-O3'-C3'	5.22	125.96	119.70
84	Aa	791	C	N3-C4-N4	5.22	121.65	118.00
84	Aa	1578	U	O4'-C1'-N1	5.22	112.38	108.20
84	Aa	5	G	N3-C2-N2	5.22	123.55	119.90
84	Aa	565	C	C4'-C3'-C2'	-5.22	97.38	102.60
84	Aa	597	C	N3-C4-C5	-5.22	119.81	121.90
84	Aa	1395	A	C5-C6-N1	-5.22	115.09	117.70
84	Aa	1407	G	O4'-C1'-N9	5.22	112.37	108.20
84	Aa	1759	C	N3-C4-N4	5.22	121.65	118.00
84	Aa	2216	G	C6-C5-N7	-5.22	127.27	130.40
84	Aa	2233	G	C5-C6-O6	-5.22	125.47	128.60
84	Aa	2291	A	O4'-C1'-N9	5.22	112.37	108.20
84	Aa	2362	A	C5-C6-N1	-5.22	115.09	117.70
84	Aa	2865	G	O4'-C1'-N9	5.22	112.37	108.20
84	Aa	2902	A	O4'-C1'-N9	5.22	112.37	108.20
84	Aa	424	G	C6-C5-N7	-5.21	127.27	130.40
84	Aa	454	A	P-O5'-C5'	5.21	129.24	120.90
84	Aa	1507	A	C5-C6-N1	-5.21	115.09	117.70
84	Aa	1566	C	N3-C4-N4	5.21	121.65	118.00
84	Aa	2178	G	N9-C1'-C2'	-5.21	106.26	112.00
84	Aa	2524	U	C4'-C3'-C2'	-5.21	97.39	102.60
84	Aa	2641	A	C5-C6-N1	-5.21	115.09	117.70
84	Aa	3195	C	N3-C4-C5	-5.21	119.81	121.90
84	Aa	1581	C	N3-C4-N4	5.21	121.65	118.00
84	Aa	2707	A	C5-C6-N6	-5.21	119.53	123.70
84	Aa	3124	A	C5-C6-N1	-5.21	115.09	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	716	A	O3'-P-O5'	5.21	113.90	104.00
1	Ad	1035	A	O4'-C1'-C2'	-5.21	100.59	105.80
1	Ad	1167	C	O4'-C1'-N1	5.21	112.37	108.20
18	BN	129	TYR	CB-CG-CD2	-5.21	117.87	121.00
30	BB	188	PHE	CB-CG-CD2	-5.21	117.15	120.80
84	Aa	237	C	N3-C4-N4	5.21	121.65	118.00
84	Aa	964	C	N3-C4-C5	-5.21	119.82	121.90
84	Aa	1720	C	N3-C4-C5	-5.21	119.81	121.90
84	Aa	2086	A	O3'-P-O5'	5.21	113.90	104.00
84	Aa	2207	C	N3-C4-N4	5.21	121.65	118.00
84	Aa	2546	C	N3-C4-C5	-5.21	119.82	121.90
2	Ae	37	G	O4'-C1'-C2'	5.21	112.29	107.60
40	Cz	167	ALA	N-CA-CB	5.21	117.39	110.10
84	Aa	851	A	C5-C6-N1	-5.21	115.09	117.70
84	Aa	2107	A	C5-C6-N1	-5.21	115.09	117.70
86	Ab	53	U	C5-C6-N1	5.21	125.31	122.70
1	Ad	269	A	P-O5'-C5'	5.21	129.23	120.90
1	Ad	839	G	N9-C1'-C2'	5.21	120.77	114.00
1	Ad	857	A	O4'-C1'-N9	5.21	112.37	108.20
1	Ad	1624	G	C1'-O4'-C4'	-5.21	105.73	109.90
84	Aa	128	C	C2-N3-C4	5.21	122.50	119.90
84	Aa	140	C	N3-C4-N4	5.21	121.65	118.00
84	Aa	795	C	N3-C4-N4	5.21	121.65	118.00
84	Aa	1007	A	C5-C6-N6	-5.21	119.53	123.70
84	Aa	1305	A	C5-C6-N1	-5.21	115.10	117.70
84	Aa	1599	A	P-O5'-C5'	-5.21	112.57	120.90
84	Aa	2286	A	C5'-C4'-C3'	5.21	124.33	116.00
84	Aa	2373	C	N3-C4-C5	-5.21	119.82	121.90
84	Aa	3349	C	N3-C4-N4	5.21	121.65	118.00
85	Ac	80	A	C5-C6-N6	-5.21	119.53	123.70
1	Ad	983	A	O4'-C1'-N9	5.21	112.36	108.20
1	Ad	1442	A	C5'-C4'-O4'	5.21	115.35	109.10
1	Ad	1492	G	C3'-C2'-C1'	-5.21	97.33	101.50
49	CR	159	PHE	CB-CG-CD1	5.21	124.44	120.80
84	Aa	386	G	O4'-C1'-N9	5.21	112.36	108.20
84	Aa	1457	A	C4-C5-C6	5.21	119.60	117.00
84	Aa	1537	A	C5-C6-N6	-5.21	119.53	123.70
2	Ae	21	A	P-O5'-C5'	5.21	129.23	120.90
19	BL	54	TYR	CB-CG-CD2	-5.21	117.88	121.00
84	Aa	189	C	N3-C4-N4	5.21	121.64	118.00
84	Aa	1491	G	O4'-C1'-N9	5.21	112.36	108.20
84	Aa	3217	G	C5-C6-O6	-5.21	125.48	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	329	G	P-O3'-C3'	5.20	125.94	119.70
1	Ad	547	C	C4'-C3'-C2'	-5.20	97.40	102.60
1	Ad	933	G	O4'-C1'-N9	5.20	112.36	108.20
1	Ad	1609	G	O4'-C1'-C2'	5.20	112.28	107.60
84	Aa	1020	U	O4'-C1'-N1	5.20	112.36	108.20
84	Aa	1359	A	C5-C6-N1	-5.20	115.10	117.70
84	Aa	1599	A	C5-C6-N1	-5.20	115.10	117.70
84	Aa	2344	A	C5-C6-N6	-5.20	119.54	123.70
84	Aa	3116	C	N3-C4-N4	5.20	121.64	118.00
84	Aa	3297	A	C5-C6-N6	-5.20	119.54	123.70
85	Ac	123	G	N3-C2-N2	5.20	123.54	119.90
64	Ci	42	PHE	N-CA-CB	5.20	119.96	110.60
84	Aa	2446	G	O4'-C1'-N9	5.20	112.36	108.20
84	Aa	2786	G	N3-C2-N2	5.20	123.54	119.90
84	Aa	2916	G	O4'-C1'-N9	5.20	112.36	108.20
85	Ac	108	C	C2-N3-C4	5.20	122.50	119.90
1	Ad	823	A	N9-C1'-C2'	5.20	120.76	114.00
1	Ad	1071	C	C5'-C4'-O4'	5.20	115.34	109.10
84	Aa	383	A	O4'-C1'-N9	5.20	112.36	108.20
84	Aa	739	C	N3-C4-C5	-5.20	119.82	121.90
84	Aa	1330	A	C5-C6-N6	-5.20	119.54	123.70
84	Aa	2047	A	O4'-C1'-N9	5.20	112.36	108.20
84	Aa	2646	A	C4-C5-C6	5.20	119.60	117.00
1	Ad	489	C	O4'-C1'-C2'	-5.20	100.60	105.80
1	Ad	578	G	C3'-C2'-C1'	-5.20	97.34	101.50
1	Ad	1110	C	C1'-O4'-C4'	-5.20	105.74	109.90
84	Aa	1179	C	O4'-C1'-N1	5.20	112.36	108.20
84	Aa	1369	G	N3-C2-N2	5.20	123.54	119.90
84	Aa	1859	G	O4'-C1'-N9	5.20	112.36	108.20
84	Aa	2635	G	N3-C2-N2	5.20	123.54	119.90
84	Aa	2665	A	C5-C6-N6	-5.20	119.54	123.70
84	Aa	2736	A	O4'-C1'-N9	5.20	112.36	108.20
84	Aa	2761	A	C5-C6-N1	-5.20	115.10	117.70
1	Ad	1698	A	N9-C1'-C2'	5.20	120.76	114.00
63	CU	87	TYR	CB-CG-CD1	5.20	124.12	121.00
84	Aa	1199	A	O4'-C1'-N9	5.20	112.36	108.20
84	Aa	1736	C	N3-C4-N4	5.20	121.64	118.00
84	Aa	2093	G	N1-C2-N3	-5.20	120.78	123.90
84	Aa	2794	A	C5-C6-N6	-5.20	119.54	123.70
84	Aa	3125	G	O4'-C1'-N9	5.20	112.36	108.20
85	Ac	72	A	C5'-C4'-O4'	5.20	115.34	109.10
1	Ad	82	G	O4'-C1'-C2'	5.20	112.28	107.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	700	C	N1-C1'-C2'	5.20	120.75	114.00
1	Ad	1645	C	C5'-C4'-O4'	5.20	115.33	109.10
1	Ad	1771	U	C4'-C3'-C2'	5.20	107.80	102.60
84	Aa	609	C	C5'-C4'-O4'	5.20	115.34	109.10
84	Aa	1809	A	C5-C6-N1	-5.20	115.10	117.70
84	Aa	2131	U	O4'-C1'-N1	5.20	112.36	108.20
84	Aa	2251	A	O4'-C1'-N9	5.20	112.36	108.20
84	Aa	2715	U	C5'-C4'-C3'	-5.20	107.69	116.00
1	Ad	1147	A	O4'-C1'-N9	5.19	112.36	108.20
84	Aa	774	A	O4'-C1'-N9	5.19	112.36	108.20
84	Aa	811	A	C4-C5-C6	5.19	119.60	117.00
84	Aa	1261	C	P-O5'-C5'	5.19	129.21	120.90
84	Aa	3381	C	N3-C4-C5	-5.19	119.82	121.90
85	Ac	138	G	C5-C6-O6	-5.19	125.48	128.60
35	BG	159	VAL	C-N-CA	5.19	134.68	121.70
84	Aa	2179	U	C5'-C4'-O4'	5.19	115.33	109.10
84	Aa	2283	G	O4'-C1'-N9	5.19	112.35	108.20
86	Ab	69	A	C6-C5-N7	-5.19	128.66	132.30
1	Ad	581	G	C1'-O4'-C4'	-5.19	105.75	109.90
1	Ad	982	A	O4'-C1'-N9	5.19	112.35	108.20
2	Ae	60	C	O4'-C1'-N1	5.19	112.35	108.20
84	Aa	395	A	C4-C5-C6	5.19	119.60	117.00
84	Aa	653	A	C5-C6-N6	-5.19	119.55	123.70
84	Aa	878	G	O4'-C1'-N9	5.19	112.35	108.20
84	Aa	929	A	C5-C6-N1	-5.19	115.11	117.70
84	Aa	1379	G	O4'-C1'-N9	5.19	112.35	108.20
84	Aa	2142	A	C5-C6-N1	-5.19	115.10	117.70
84	Aa	2316	A	C5-C6-N6	-5.19	119.55	123.70
84	Aa	3148	A	O4'-C1'-N9	5.19	112.35	108.20
84	Aa	3354	A	C4-C5-C6	5.19	119.59	117.00
86	Ab	70	G	C4-C5-C6	5.19	121.92	118.80
86	Ab	89	G	C5-C6-O6	-5.19	125.49	128.60
84	Aa	567	G	C6-C5-N7	-5.19	127.29	130.40
84	Aa	1115	A	C5-C6-N6	-5.19	119.55	123.70
84	Aa	1327	G	C5-C6-O6	-5.19	125.49	128.60
84	Aa	1847	G	O4'-C1'-N9	5.19	112.35	108.20
84	Aa	3299	A	C5-C6-N6	-5.19	119.55	123.70
1	Ad	1001	C	O4'-C1'-C2'	-5.19	100.61	105.80
1	Ad	1097	A	N9-C1'-C2'	5.19	120.74	114.00
70	Cq	14	TYR	CB-CG-CD2	-5.19	117.89	121.00
82	Cb	25	LYS	N-CA-CB	5.19	119.94	110.60
84	Aa	94	A	C5-C6-N6	-5.19	119.55	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	827	C	N3-C4-N4	5.19	121.63	118.00
84	Aa	1284	C	N3-C4-N4	5.19	121.63	118.00
84	Aa	1594	G	N3-C2-N2	5.19	123.53	119.90
84	Aa	1793	A	O4'-C1'-N9	5.19	112.35	108.20
84	Aa	2677	A	O4'-C1'-N9	5.19	112.35	108.20
85	Ac	13	A	C5-C6-N6	-5.19	119.55	123.70
85	Ac	61	A	C5-C6-N6	-5.19	119.55	123.70
86	Ab	44	C	P-O5'-C5'	-5.19	112.60	120.90
1	Ad	72	A	C3'-C2'-C1'	5.19	105.65	101.50
1	Ad	428	C	C1'-O4'-C4'	-5.19	105.75	109.90
84	Aa	435	G	N1-C6-O6	5.19	123.01	119.90
84	Aa	547	C	N3-C4-N4	5.19	121.63	118.00
84	Aa	2203	A	C5-C6-N1	-5.19	115.11	117.70
84	Aa	2210	A	O4'-C1'-N9	5.19	112.35	108.20
84	Aa	3327	A	C5-C6-N1	-5.19	115.11	117.70
85	Ac	3	A	C4-C5-C6	5.19	119.59	117.00
1	Ad	813	A	C1'-O4'-C4'	5.18	114.05	109.90
1	Ad	965	U	P-O5'-C5'	-5.18	112.60	120.90
1	Ad	1137	A	O4'-C1'-N9	5.18	112.35	108.20
84	Aa	1293	C	C2-N3-C4	5.18	122.49	119.90
84	Aa	2707	A	C4-C5-C6	5.18	119.59	117.00
84	Aa	2744	C	N3-C4-C5	-5.18	119.83	121.90
86	Ab	75	G	C5'-C4'-O4'	5.18	115.32	109.10
1	Ad	874	A	O4'-C1'-C2'	-5.18	100.62	105.80
1	Ad	1556	U	C1'-O4'-C4'	5.18	114.05	109.90
30	BB	188	PHE	CB-CG-CD1	5.18	124.43	120.80
84	Aa	23	A	C5-C6-N1	-5.18	115.11	117.70
84	Aa	334	A	C4-C5-C6	5.18	119.59	117.00
84	Aa	899	A	C5-C6-N6	-5.18	119.55	123.70
84	Aa	1244	A	O4'-C1'-N9	5.18	112.35	108.20
84	Aa	1397	A	C5-C6-N6	-5.18	119.55	123.70
84	Aa	1787	C	N3-C4-N4	5.18	121.63	118.00
84	Aa	1965	C	N3-C4-C5	-5.18	119.83	121.90
84	Aa	2529	C	C2-N3-C4	5.18	122.49	119.90
84	Aa	2625	C	N3-C4-C5	-5.18	119.83	121.90
84	Aa	2900	G	N1-C6-O6	5.18	123.01	119.90
84	Aa	3124	A	O4'-C1'-N9	5.18	112.35	108.20
85	Ac	92	A	C5-C6-N6	-5.18	119.55	123.70
85	Ac	137	G	C5-C6-O6	-5.18	125.49	128.60
84	Aa	1137	G	C5-C6-O6	-5.18	125.49	128.60
84	Aa	1584	A	C5-C6-N6	-5.18	119.56	123.70
1	Ad	517	U	C5'-C4'-O4'	5.18	115.32	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1538	C	C1'-O4'-C4'	5.18	114.04	109.90
1	Ad	1584	A	O4'-C1'-C2'	-5.18	100.62	105.80
84	Aa	789	A	C4-C5-C6	5.18	119.59	117.00
84	Aa	804	A	C5-C6-N6	-5.18	119.56	123.70
84	Aa	871	C	N3-C4-N4	5.18	121.62	118.00
84	Aa	1282	A	C5-C6-N1	-5.18	115.11	117.70
84	Aa	1356	G	O4'-C1'-N9	5.18	112.34	108.20
84	Aa	1926	A	O4'-C1'-N9	5.18	112.34	108.20
84	Aa	2193	A	C5-C6-N1	-5.18	115.11	117.70
85	Ac	80	A	O4'-C1'-N9	5.18	112.34	108.20
85	Ac	125	C	N3-C4-N4	5.18	121.62	118.00
84	Aa	227	C	N3-C4-N4	5.18	121.62	118.00
84	Aa	1030	A	O4'-C1'-N9	5.18	112.34	108.20
84	Aa	1850	C	C2-N3-C4	5.18	122.49	119.90
1	Ad	177	C	O4'-C1'-C2'	-5.18	100.62	105.80
1	Ad	260	A	N9-C1'-C2'	5.18	120.73	114.00
1	Ad	1538	C	P-O5'-C5'	5.18	129.18	120.90
1	Ad	1699	C	O4'-C1'-N1	5.18	112.34	108.20
2	Ae	56	A	C3'-C2'-C1'	5.18	105.64	101.50
78	CL	66	ASN	N-CA-CB	5.18	119.92	110.60
84	Aa	149	A	C5-C6-N1	-5.18	115.11	117.70
84	Aa	672	A	O4'-C1'-N9	5.18	112.34	108.20
84	Aa	1513	C	N3-C4-N4	5.18	121.62	118.00
84	Aa	2388	C	C5'-C4'-C3'	5.18	124.28	116.00
84	Aa	2649	C	N3-C4-C5	-5.18	119.83	121.90
1	Ad	511	U	C1'-O4'-C4'	5.17	114.04	109.90
84	Aa	382	A	O4'-C1'-N9	5.17	112.34	108.20
84	Aa	1140	C	N3-C4-N4	5.17	121.62	118.00
84	Aa	2179	U	C4'-C3'-O3'	-5.17	98.53	109.40
84	Aa	2280	C	N3-C4-C5	-5.17	119.83	121.90
84	Aa	2356	A	O4'-C1'-N9	5.17	112.34	108.20
84	Aa	3367	C	N3-C4-C5	-5.17	119.83	121.90
86	Ab	78	C	O4'-C1'-N1	5.17	112.34	108.20
1	Ad	409	C	O4'-C1'-N1	5.17	112.34	108.20
1	Ad	774	C	C3'-C2'-C1'	5.17	105.64	101.50
1	Ad	961	U	C3'-C2'-C1'	5.17	105.64	101.50
84	Aa	67	C	N3-C4-N4	5.17	121.62	118.00
84	Aa	813	A	C5-C6-N6	-5.17	119.56	123.70
1	Ad	29	U	C1'-O4'-C4'	5.17	114.04	109.90
84	Aa	1654	C	O4'-C1'-N1	5.17	112.34	108.20
84	Aa	1898	G	O4'-C1'-N9	5.17	112.34	108.20
84	Aa	2770	U	C5'-C4'-C3'	-5.17	107.73	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2869	C	N3-C4-N4	5.17	121.62	118.00
84	Aa	3099	G	O4'-C1'-N9	5.17	112.34	108.20
84	Aa	3123	A	C5-C6-N6	-5.17	119.56	123.70
1	Ad	1808	U	O4'-C1'-C2'	-5.17	100.63	105.80
22	BZ	18	SER	N-CA-CB	5.17	118.25	110.50
84	Aa	1482	C	N3-C4-N4	5.17	121.62	118.00
84	Aa	2401	A	O4'-C1'-N9	5.17	112.34	108.20
84	Aa	3196	C	N3-C4-N4	5.17	121.62	118.00
1	Ad	222	G	OP1-P-OP2	-5.17	111.85	119.60
1	Ad	576	C	C3'-C2'-C1'	5.17	105.64	101.50
1	Ad	1497	U	C5'-C4'-O4'	5.17	115.30	109.10
1	Ad	1798	G	C5'-C4'-O4'	5.17	115.30	109.10
84	Aa	565	C	N3-C4-C5	-5.17	119.83	121.90
84	Aa	630	C	N3-C4-C5	-5.17	119.83	121.90
84	Aa	655	G	O4'-C1'-N9	5.17	112.33	108.20
84	Aa	1230	G	O4'-C1'-N9	5.17	112.33	108.20
84	Aa	2013	G	C5'-C4'-O4'	5.17	115.30	109.10
84	Aa	2541	A	C4-C5-C6	5.17	119.58	117.00
84	Aa	2674	A	O4'-C1'-N9	5.17	112.33	108.20
84	Aa	2815	A	C5-C6-N6	-5.17	119.56	123.70
84	Aa	2869	C	C2-N3-C4	5.17	122.48	119.90
84	Aa	3048	C	N3-C4-N4	5.17	121.62	118.00
86	Ab	56	G	C6-C5-N7	-5.17	127.30	130.40
86	Ab	82	G	N1-C6-O6	5.17	123.00	119.90
1	Ad	1793	C	O4'-C1'-N1	5.17	112.33	108.20
11	BD	120	TYR	CB-CG-CD1	-5.17	117.90	121.00
84	Aa	1334	A	C4-C5-C6	5.17	119.58	117.00
84	Aa	2304	A	C5-C6-N6	-5.17	119.57	123.70
84	Aa	2699	A	C5-C6-N1	-5.17	115.12	117.70
84	Aa	2841	G	O4'-C1'-N9	5.17	112.33	108.20
84	Aa	3220	A	O4'-C1'-N9	5.17	112.33	108.20
85	Ac	135	A	C5-C6-N6	-5.17	119.57	123.70
1	Ad	231	U	N1-C1'-C2'	5.17	120.72	114.00
1	Ad	265	A	C4'-C3'-C2'	-5.17	97.44	102.60
1	Ad	473	C	C3'-C2'-C1'	5.17	105.63	101.50
84	Aa	105	A	C4-C5-C6	5.17	119.58	117.00
85	Ac	129	C	C6-N1-C2	-5.17	118.23	120.30
1	Ad	1191	U	N1-C1'-C2'	5.16	120.71	114.00
1	Ad	1280	U	O4'-C1'-C2'	-5.16	100.64	105.80
84	Aa	236	A	C4-C5-C6	5.16	119.58	117.00
84	Aa	282	A	O5'-C5'-C4'	5.16	121.51	111.70
84	Aa	724	A	C4-C5-C6	5.16	119.58	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1506	A	C4-C5-C6	5.16	119.58	117.00
84	Aa	3221	A	C5-C6-N1	-5.16	115.12	117.70
84	Aa	3333	C	N3-C4-N4	5.16	121.61	118.00
86	Ab	50	A	N1-C2-N3	5.16	131.88	129.30
84	Aa	101	C	N3-C4-N4	5.16	121.61	118.00
84	Aa	263	A	C5-C6-N1	-5.16	115.12	117.70
84	Aa	638	G	P-O5'-C5'	5.16	129.16	120.90
84	Aa	299	G	O4'-C1'-N9	5.16	112.33	108.20
84	Aa	1469	G	O4'-C1'-N9	5.16	112.33	108.20
84	Aa	2270	A	C5-C6-N6	-5.16	119.57	123.70
84	Aa	3094	C	N3-C4-C5	-5.16	119.84	121.90
1	Ad	451	U	N1-C1'-C2'	5.16	120.70	114.00
1	Ad	485	A	N9-C1'-C2'	5.16	120.71	114.00
1	Ad	1326	A	O4'-C1'-N9	5.16	112.33	108.20
84	Aa	94	A	O4'-C1'-N9	5.16	112.33	108.20
84	Aa	519	C	N3-C4-N4	5.16	121.61	118.00
84	Aa	1395	A	O4'-C1'-N9	5.16	112.33	108.20
84	Aa	1424	G	N3-C2-N2	5.16	123.51	119.90
84	Aa	2288	C	N3-C4-N4	5.16	121.61	118.00
84	Aa	3321	C	P-O3'-C3'	5.16	125.89	119.70
84	Aa	2709	G	O4'-C1'-N9	5.16	112.33	108.20
84	Aa	3332	G	P-O3'-C3'	5.16	125.89	119.70
1	Ad	758	A	C1'-O4'-C4'	-5.16	105.78	109.90
84	Aa	294	A	C5-C6-N1	-5.16	115.12	117.70
84	Aa	1092	G	O4'-C1'-N9	5.16	112.33	108.20
84	Aa	1140	C	N3-C4-C5	-5.16	119.84	121.90
84	Aa	1585	A	C4-C5-C6	5.16	119.58	117.00
84	Aa	1638	U	O4'-C1'-N1	5.16	112.32	108.20
84	Aa	1907	A	O4'-C1'-N9	5.16	112.33	108.20
84	Aa	2754	G	O4'-C1'-N9	5.16	112.32	108.20
84	Aa	2868	C	N3-C4-C5	-5.16	119.84	121.90
86	Ab	48	G	C2-N3-C4	-5.16	109.32	111.90
84	Aa	832	C	N3-C4-C5	-5.15	119.84	121.90
84	Aa	1450	G	C4-C5-C6	5.15	121.89	118.80
84	Aa	350	A	C5-C6-N1	-5.15	115.12	117.70
84	Aa	771	G	O4'-C1'-N9	5.15	112.32	108.20
84	Aa	1731	A	O4'-C1'-N9	5.15	112.32	108.20
84	Aa	1913	C	N3-C4-N4	5.15	121.61	118.00
84	Aa	2282	C	N3-C4-C5	-5.15	119.84	121.90
84	Aa	3279	G	O4'-C1'-N9	5.15	112.32	108.20
85	Ac	6	C	N3-C4-C5	-5.15	119.84	121.90
85	Ac	41	A	C4-C5-C6	5.15	119.58	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	938	A	O4'-C1'-C2'	5.15	112.24	107.60
1	Ad	1626	C	O4'-C1'-C2'	-5.15	100.65	105.80
84	Aa	1256	A	C5-C6-N1	-5.15	115.12	117.70
84	Aa	2386	A	O4'-C1'-N9	5.15	112.32	108.20
84	Aa	3127	C	N3-C4-N4	5.15	121.61	118.00
85	Ac	77	A	C5-C6-N6	-5.15	119.58	123.70
85	Ac	99	C	N3-C4-C5	-5.15	119.84	121.90
84	Aa	311	G	C5-C6-O6	-5.15	125.51	128.60
84	Aa	1560	A	C5'-C4'-O4'	5.15	115.28	109.10
84	Aa	1894	G	C5-C6-O6	-5.15	125.51	128.60
85	Ac	132	C	N3-C4-N4	5.15	121.61	118.00
1	Ad	506	G	C1'-O4'-C4'	-5.15	105.78	109.90
1	Ad	593	C	C3'-C2'-C1'	5.15	105.62	101.50
2	Ae	61	C	O4'-C1'-N1	5.15	112.32	108.20
84	Aa	11	A	C4-C5-C6	5.15	119.57	117.00
84	Aa	1224	A	C5-C6-N1	-5.15	115.13	117.70
84	Aa	1401	C	C2-N3-C4	5.15	122.47	119.90
84	Aa	1435	C	N3-C4-N4	5.15	121.60	118.00
84	Aa	1660	C	N3-C4-N4	5.15	121.60	118.00
84	Aa	1771	G	N3-C2-N2	5.15	123.50	119.90
84	Aa	1998	A	C5-C6-N1	-5.15	115.13	117.70
84	Aa	2125	A	C5-C6-N6	-5.15	119.58	123.70
84	Aa	2251	A	C5-C6-N1	-5.15	115.13	117.70
84	Aa	2374	G	O4'-C1'-N9	5.15	112.32	108.20
84	Aa	2849	A	O4'-C1'-N9	5.15	112.32	108.20
42	CJ	63	ARG	N-CA-CB	5.15	119.86	110.60
84	Aa	1019	A	O4'-C1'-N9	5.15	112.32	108.20
84	Aa	2290	A	C5-C6-N6	-5.15	119.58	123.70
1	Ad	8	U	O4'-C1'-C2'	-5.14	100.66	105.80
1	Ad	1548	G	C3'-C2'-C1'	-5.14	97.38	101.50
1	Ad	1716	C	N1-C1'-C2'	5.14	120.69	114.00
84	Aa	660	A	C5-C6-N6	-5.14	119.58	123.70
84	Aa	873	A	C5-C6-N1	-5.14	115.13	117.70
84	Aa	2235	G	N3-C2-N2	5.14	123.50	119.90
84	Aa	3172	G	C4-N9-C1'	5.14	133.19	126.50
84	Aa	3209	U	O4'-C1'-N1	5.14	112.31	108.20
86	Ab	26	C	N3-C4-N4	5.14	121.60	118.00
2	Ae	58	U	N1-C1'-C2'	-5.14	106.34	112.00
71	CB	2	SER	N-CA-C	-5.14	97.11	111.00
84	Aa	396	G	O4'-C1'-N9	5.14	112.31	108.20
84	Aa	955	A	C5-C6-N1	-5.14	115.13	117.70
84	Aa	1083	C	N3-C4-C5	-5.14	119.84	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2434	G	P-O3'-C3'	-5.14	113.53	119.70
84	Aa	3317	G	N3-C2-N2	5.14	123.50	119.90
84	Aa	384	A	C5-C6-N1	-5.14	115.13	117.70
1	Ad	405	A	O4'-C1'-N9	5.14	112.31	108.20
63	CU	108	ASN	N-CA-CB	5.14	119.85	110.60
84	Aa	500	C	N3-C4-C5	-5.14	119.84	121.90
84	Aa	1193	A	C8-N9-C4	-5.14	103.75	105.80
84	Aa	1235	A	C5-C6-N1	-5.14	115.13	117.70
84	Aa	1400	C	N3-C4-N4	5.14	121.60	118.00
84	Aa	1838	A	C5-C6-N1	-5.14	115.13	117.70
84	Aa	2569	G	C5-C6-O6	-5.14	125.52	128.60
84	Aa	2814	C	N3-C4-N4	5.14	121.60	118.00
84	Aa	3313	C	N3-C4-C5	-5.14	119.84	121.90
1	Ad	66	U	C2'-C3'-O3'	5.14	121.92	113.70
84	Aa	1960	C	N3-C4-N4	5.14	121.60	118.00
84	Aa	2358	C	N3-C4-N4	5.14	121.60	118.00
84	Aa	2834	C	N3-C4-N4	5.14	121.60	118.00
1	Ad	400	G	P-O3'-C3'	-5.14	113.53	119.70
1	Ad	1105	G	C1'-O4'-C4'	-5.14	105.79	109.90
16	BO	66	ASP	N-CA-CB	5.14	119.84	110.60
84	Aa	554	C	C5'-C4'-O4'	5.14	115.26	109.10
84	Aa	963	U	O4'-C1'-N1	5.14	112.31	108.20
84	Aa	1235	A	C5-C6-N6	-5.14	119.59	123.70
84	Aa	1560	A	O5'-P-OP1	-5.14	101.08	105.70
84	Aa	1927	A	C5-C6-N1	-5.14	115.13	117.70
85	Ac	83	C	N3-C4-C5	-5.14	119.84	121.90
86	Ab	47	C	C5-C4-N4	-5.14	116.61	120.20
1	Ad	802	A	O4'-C1'-N9	5.13	112.31	108.20
25	Bd	14	TYR	N-CA-CB	5.13	119.84	110.60
60	Co	102	THR	N-CA-CB	5.13	120.06	110.30
84	Aa	186	A	O4'-C1'-N9	5.13	112.31	108.20
84	Aa	674	G	C5-C6-O6	-5.13	125.52	128.60
84	Aa	2073	U	C6-N1-C1'	-5.13	114.01	121.20
84	Aa	2276	A	C5-C6-N1	-5.13	115.13	117.70
84	Aa	2804	A	O4'-C1'-N9	5.13	112.31	108.20
1	Ad	45	U	N1-C1'-C2'	5.13	120.67	114.00
84	Aa	341	U	O4'-C1'-N1	5.13	112.31	108.20
84	Aa	1059	A	C4-C5-C6	5.13	119.57	117.00
84	Aa	1132	A	O4'-C1'-N9	5.13	112.31	108.20
84	Aa	1151	G	O4'-C1'-N9	5.13	112.31	108.20
84	Aa	1946	C	N3-C4-N4	5.13	121.59	118.00
84	Aa	3177	A	C5-C6-N6	-5.13	119.59	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	Ac	131	G	N3-C2-N2	5.13	123.49	119.90
1	Ad	515	A	C3'-C2'-C1'	5.13	105.61	101.50
1	Ad	642	C	C3'-C2'-C1'	5.13	105.61	101.50
1	Ad	732	G	C3'-C2'-C1'	-5.13	97.39	101.50
2	Ae	53	U	O4'-C1'-C2'	-5.13	100.67	105.80
84	Aa	62	A	O4'-C1'-N9	5.13	112.31	108.20
84	Aa	724	A	P-O5'-C5'	5.13	129.11	120.90
84	Aa	1275	A	C4-C5-C6	5.13	119.56	117.00
84	Aa	1370	A	O4'-C1'-N9	5.13	112.31	108.20
84	Aa	1462	C	N3-C4-N4	5.13	121.59	118.00
84	Aa	2009	C	N3-C4-N4	5.13	121.59	118.00
84	Aa	2152	A	C5-C6-N1	-5.13	115.13	117.70
84	Aa	2752	G	O4'-C1'-N9	5.13	112.31	108.20
84	Aa	3259	A	C4'-C3'-O3'	-5.13	98.63	109.40
84	Aa	1077	C	N3-C4-N4	5.13	121.59	118.00
84	Aa	2929	C	N3-C4-C5	-5.13	119.85	121.90
85	Ac	109	A	C5-C6-N1	-5.13	115.14	117.70
1	Ad	34	G	N9-C1'-C2'	5.13	120.67	114.00
1	Ad	838	U	O4'-C4'-C3'	-5.13	98.87	104.00
14	BQ	135	PHE	CB-CG-CD1	-5.13	117.21	120.80
84	Aa	1239	U	O4'-C1'-N1	5.13	112.30	108.20
84	Aa	1298	A	C4-C5-C6	5.13	119.56	117.00
84	Aa	1487	A	C5-C6-N1	-5.13	115.14	117.70
84	Aa	3188	G	O4'-C1'-N9	5.13	112.30	108.20
84	Aa	3376	C	N3-C4-C5	-5.13	119.85	121.90
84	Aa	283	A	C4-C5-C6	5.13	119.56	117.00
84	Aa	910	G	N1-C6-O6	5.13	122.98	119.90
84	Aa	916	A	O4'-C1'-N9	5.13	112.30	108.20
84	Aa	960	C	C2-N3-C4	5.13	122.46	119.90
84	Aa	2241	G	C5-C6-O6	-5.13	125.52	128.60
84	Aa	2282	C	N3-C4-N4	5.13	121.59	118.00
84	Aa	251	G	N3-C2-N2	5.12	123.49	119.90
84	Aa	584	G	O4'-C1'-N9	5.12	112.30	108.20
84	Aa	1398	A	C5-C6-N1	-5.12	115.14	117.70
84	Aa	1651	A	C4-C5-C6	5.12	119.56	117.00
84	Aa	1742	G	O4'-C1'-N9	5.12	112.30	108.20
1	Ad	426	G	P-O3'-C3'	5.12	125.85	119.70
84	Aa	29	G	O4'-C1'-N9	5.12	112.30	108.20
84	Aa	561	G	C5-C6-O6	-5.12	125.53	128.60
84	Aa	980	C	N3-C4-N4	5.12	121.59	118.00
84	Aa	1651	A	O4'-C1'-N9	5.12	112.30	108.20
84	Aa	1723	C	C5'-C4'-O4'	5.12	115.25	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1984	C	N3-C4-N4	5.12	121.59	118.00
84	Aa	2049	C	N3-C4-N4	5.12	121.59	118.00
84	Aa	2287	U	O4'-C1'-N1	5.12	112.30	108.20
84	Aa	2347	A	C5-C6-N1	-5.12	115.14	117.70
85	Ac	61	A	C5-C6-N1	-5.12	115.14	117.70
85	Ac	86	U	O4'-C1'-N1	5.12	112.30	108.20
1	Ad	288	G	C4'-C3'-C2'	-5.12	97.48	102.60
1	Ad	946	A	O4'-C1'-N9	5.12	112.30	108.20
2	Ae	59	U	O4'-C1'-C2'	-5.12	100.68	105.80
15	BU	81	GLU	N-CA-CB	5.12	119.82	110.60
84	Aa	1057	A	C5-C6-N1	-5.12	115.14	117.70
84	Aa	2895	G	C5-C6-O6	-5.12	125.53	128.60
85	Ac	44	A	C5-C6-N1	-5.12	115.14	117.70
85	Ac	59	A	C5-N7-C8	5.12	106.46	103.90
1	Ad	1359	C	O4'-C1'-N1	5.12	112.30	108.20
84	Aa	340	A	C5-C6-N6	-5.12	119.61	123.70
84	Aa	918	A	C5-C6-N6	-5.12	119.60	123.70
84	Aa	1608	C	N3-C4-C5	-5.12	119.85	121.90
84	Aa	1892	A	C8-N9-C4	-5.12	103.75	105.80
84	Aa	3297	A	C5-C6-N1	-5.12	115.14	117.70
1	Ad	114	U	C1'-O4'-C4'	-5.12	105.81	109.90
1	Ad	1576	C	C3'-C2'-C1'	5.12	105.59	101.50
84	Aa	2070	C	P-O5'-C5'	5.12	129.09	120.90
84	Aa	2904	A	C5-C6-N1	-5.12	115.14	117.70
84	Aa	3315	A	C4-C5-C6	5.12	119.56	117.00
1	Ad	550	U	O4'-C1'-N1	5.12	112.29	108.20
1	Ad	631	C	N1-C1'-C2'	5.12	120.65	114.00
1	Ad	1151	G	O4'-C1'-C2'	5.12	112.20	107.60
3	Af	21	C	C3'-C2'-C1'	5.12	105.59	101.50
62	CS	72	THR	N-CA-CB	5.12	120.02	110.30
84	Aa	27	C	N3-C4-N4	5.12	121.58	118.00
84	Aa	437	C	N3-C4-N4	5.12	121.58	118.00
84	Aa	1391	A	C4-C5-C6	5.12	119.56	117.00
84	Aa	2559	C	N3-C4-N4	5.12	121.58	118.00
1	Ad	1274	G	C5'-C4'-O4'	5.11	115.23	109.10
64	Ci	95	SER	N-CA-CB	5.11	118.17	110.50
84	Aa	550	C	C2'-C3'-O3'	-5.11	98.25	109.50
84	Aa	595	C	N3-C4-N4	5.11	121.58	118.00
84	Aa	1323	G	N3-C2-N2	5.11	123.48	119.90
84	Aa	2226	C	N3-C4-N4	5.11	121.58	118.00
84	Aa	2270	A	C4-C5-C6	5.11	119.56	117.00
1	Ad	858	G	N9-C1'-C2'	5.11	120.64	114.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1310	G	C4-N9-C1'	5.11	133.15	126.50
84	Aa	1738	A	C5-C6-N1	-5.11	115.14	117.70
84	Aa	2451	G	N3-C2-N2	5.11	123.48	119.90
84	Aa	2733	A	C5-C6-N1	-5.11	115.14	117.70
84	Aa	3004	G	O4'-C1'-N9	5.11	112.29	108.20
84	Aa	3171	C	N3-C4-C5	-5.11	119.86	121.90
1	Ad	458	A	P-O3'-C3'	5.11	125.83	119.70
1	Ad	837	G	O4'-C1'-N9	5.11	112.29	108.20
1	Ad	1066	U	C3'-C2'-C1'	-5.11	97.41	101.50
84	Aa	59	A	C5-C6-N1	-5.11	115.14	117.70
84	Aa	298	G	N3-C2-N2	5.11	123.48	119.90
84	Aa	325	A	C5-C6-N1	-5.11	115.14	117.70
84	Aa	1005	C	N3-C4-C5	-5.11	119.86	121.90
84	Aa	1464	A	C5-C6-N6	-5.11	119.61	123.70
84	Aa	1807	C	N3-C4-N4	5.11	121.58	118.00
84	Aa	2677	A	C4-C5-C6	5.11	119.56	117.00
1	Ad	1143	A	O4'-C1'-C2'	-5.11	100.69	105.80
1	Ad	1735	C	O4'-C1'-C2'	-5.11	100.69	105.80
84	Aa	493	G	P-O3'-C3'	-5.11	113.57	119.70
84	Aa	495	G	C5-C6-O6	-5.11	125.53	128.60
84	Aa	1032	C	N3-C4-C5	-5.11	119.86	121.90
84	Aa	2377	C	N3-C4-N4	5.11	121.58	118.00
84	Aa	2434	G	C4'-C3'-C2'	-5.11	97.49	102.60
1	Ad	93	A	C3'-C2'-C1'	5.11	105.59	101.50
1	Ad	1616	U	C1'-O4'-C4'	5.11	113.99	109.90
84	Aa	651	A	C5-C6-N6	-5.11	119.61	123.70
84	Aa	1077	C	N3-C4-C5	-5.11	119.86	121.90
84	Aa	1396	A	C5-C6-N1	-5.11	115.15	117.70
84	Aa	1804	G	O4'-C1'-N9	5.11	112.29	108.20
84	Aa	1866	C	C2-N3-C4	5.11	122.45	119.90
84	Aa	2082	A	O4'-C1'-N9	5.11	112.29	108.20
1	Ad	169	A	C3'-C2'-C1'	5.11	105.58	101.50
1	Ad	1134	U	C1'-O4'-C4'	5.11	113.98	109.90
84	Aa	247	C	N3-C4-N4	5.11	121.57	118.00
84	Aa	764	A	C4-C5-C6	5.11	119.55	117.00
84	Aa	1109	G	N1-C2-N3	-5.11	120.84	123.90
84	Aa	1760	G	P-O3'-C3'	-5.11	113.57	119.70
84	Aa	1779	C	C2-N3-C4	5.11	122.45	119.90
84	Aa	1966	C	N3-C4-C5	-5.11	119.86	121.90
84	Aa	2518	A	C5-C6-N1	-5.11	115.15	117.70
84	Aa	2645	A	C5-C6-N1	-5.11	115.15	117.70
84	Aa	2986	C	N3-C4-N4	5.11	121.57	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3066	G	N3-C2-N2	5.11	123.47	119.90
84	Aa	3310	A	C5-C6-N1	-5.11	115.15	117.70
85	Ac	3	A	C5-C6-N1	-5.11	115.15	117.70
85	Ac	4	C	N3-C4-C5	-5.11	119.86	121.90
2	Ae	67	G	C4'-C3'-C2'	-5.10	97.50	102.60
84	Aa	1356	G	C4-C5-C6	5.10	121.86	118.80
84	Aa	1427	C	C2-N3-C4	5.10	122.45	119.90
84	Aa	565	C	N3-C4-N4	5.10	121.57	118.00
84	Aa	792	A	C5-C6-N1	-5.10	115.15	117.70
84	Aa	1248	A	O4'-C1'-N9	5.10	112.28	108.20
84	Aa	1739	G	N3-C2-N2	5.10	123.47	119.90
84	Aa	1835	A	C5-C6-N1	-5.10	115.15	117.70
84	Aa	3085	C	N3-C4-N4	5.10	121.57	118.00
49	CR	159	PHE	CB-CG-CD2	-5.10	117.23	120.80
84	Aa	941	C	N3-C4-N4	5.10	121.57	118.00
84	Aa	1805	A	C5-C6-N6	-5.10	119.62	123.70
86	Ab	44	C	C5'-C4'-O4'	5.10	115.22	109.10
1	Ad	747	U	N1-C1'-C2'	5.10	120.63	114.00
11	BD	107	TYR	CB-CG-CD1	-5.10	117.94	121.00
84	Aa	87	A	C8-N9-C4	-5.10	103.76	105.80
84	Aa	172	A	C4-C5-C6	5.10	119.55	117.00
84	Aa	175	G	C5-C6-O6	-5.10	125.54	128.60
84	Aa	452	G	C5'-C4'-O4'	5.10	115.22	109.10
84	Aa	513	C	O4'-C1'-N1	5.10	112.28	108.20
84	Aa	793	C	N3-C4-N4	5.10	121.57	118.00
84	Aa	1437	G	O4'-C1'-N9	5.10	112.28	108.20
84	Aa	2008	G	C5-C6-O6	-5.10	125.54	128.60
84	Aa	2852	G	O4'-C1'-N9	5.10	112.28	108.20
1	Ad	340	G	O4'-C1'-N9	5.10	112.28	108.20
1	Ad	1408	G	O3'-P-O5'	5.10	113.68	104.00
28	BA	36	PHE	CB-CG-CD1	5.10	124.37	120.80
84	Aa	255	C	N3-C4-C5	-5.10	119.86	121.90
84	Aa	397	A	C4-C5-C6	5.10	119.55	117.00
84	Aa	399	U	O4'-C1'-N1	5.10	112.28	108.20
84	Aa	1197	A	C5-C6-N6	-5.10	119.62	123.70
84	Aa	2286	A	C5-C6-N1	-5.10	115.15	117.70
84	Aa	2816	G	O4'-C1'-N9	5.10	112.28	108.20
84	Aa	2913	A	C5-C6-N1	-5.10	115.15	117.70
84	Aa	3160	G	O4'-C1'-N9	5.10	112.28	108.20
84	Aa	650	A	C3'-C2'-C1'	5.10	105.58	101.50
1	Ad	226	C	O4'-C1'-N1	5.09	112.28	108.20
84	Aa	796	C	N3-C4-N4	5.09	121.57	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	1394	C	N3-C4-C5	-5.09	119.86	121.90
84	Aa	1727	A	C5-C6-N1	-5.09	115.15	117.70
84	Aa	2699	A	O4'-C1'-N9	5.09	112.28	108.20
1	Ad	700	C	C3'-C2'-C1'	5.09	105.58	101.50
48	CD	144	ALA	N-CA-CB	5.09	117.23	110.10
84	Aa	1538	A	C5-C6-N1	-5.09	115.15	117.70
84	Aa	1793	A	C5-C6-N1	-5.09	115.15	117.70
1	Ad	776	A	O4'-C1'-N9	5.09	112.27	108.20
84	Aa	479	C	N3-C4-N4	5.09	121.56	118.00
84	Aa	753	G	O4'-C1'-N9	5.09	112.27	108.20
84	Aa	1637	G	N3-C2-N2	5.09	123.47	119.90
84	Aa	1871	G	O4'-C1'-N9	5.09	112.27	108.20
84	Aa	2247	A	O4'-C1'-N9	5.09	112.27	108.20
85	Ac	17	A	C5-C6-N6	-5.09	119.63	123.70
86	Ab	32	A	C5-C6-N1	-5.09	115.15	117.70
1	Ad	100	C	O4'-C1'-C2'	-5.09	100.71	105.80
1	Ad	207	A	N9-C1'-C2'	-5.09	106.40	112.00
1	Ad	307	U	O4'-C1'-N1	5.09	112.27	108.20
1	Ad	954	C	C3'-C2'-C1'	5.09	105.57	101.50
6	BK	54	TYR	CB-CG-CD2	5.09	124.05	121.00
84	Aa	457	C	N3-C4-C5	-5.09	119.86	121.90
84	Aa	552	G	C2'-C3'-O3'	5.09	121.84	113.70
84	Aa	1355	U	C6-N1-C1'	-5.09	114.07	121.20
84	Aa	1383	G	N1-C2-N3	-5.09	120.85	123.90
84	Aa	1693	A	C5-N7-C8	5.09	106.44	103.90
84	Aa	2529	C	C5'-C4'-C3'	-5.09	107.86	116.00
1	Ad	1701	G	O4'-C1'-N9	5.09	112.27	108.20
1	Ad	851	G	O4'-C1'-C2'	5.09	112.18	107.60
1	Ad	1395	C	O4'-C1'-C2'	5.09	112.18	107.60
1	Ad	1701	G	O4'-C1'-C2'	5.09	112.18	107.60
84	Aa	203	C	N3-C4-N4	5.09	121.56	118.00
84	Aa	512	G	C5'-C4'-C3'	5.09	124.14	116.00
84	Aa	564	A	C5-C6-N6	-5.09	119.63	123.70
84	Aa	984	A	C5-C6-N6	-5.09	119.63	123.70
84	Aa	1236	C	N3-C4-N4	5.09	121.56	118.00
84	Aa	1966	C	N3-C4-N4	5.09	121.56	118.00
84	Aa	2782	G	O4'-C1'-N9	5.09	112.27	108.20
84	Aa	3031	G	O4'-C1'-N9	5.09	112.27	108.20
1	Ad	592	U	O4'-C1'-N1	5.08	112.27	108.20
22	BZ	26	LYS	N-CA-CB	5.08	119.75	110.60
84	Aa	421	A	C5-C6-N1	-5.08	115.16	117.70
1	Ad	176	A	P-O3'-C3'	5.08	125.80	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	621	U	C1'-O4'-C4'	-5.08	105.83	109.90
1	Ad	784	C	O4'-C1'-C2'	5.08	112.17	107.60
1	Ad	1155	G	O4'-C1'-C2'	5.08	112.18	107.60
84	Aa	46	A	C5-C6-N1	-5.08	115.16	117.70
84	Aa	282	A	C4-C5-C6	5.08	119.54	117.00
84	Aa	1000	A	C5-C6-N1	-5.08	115.16	117.70
84	Aa	1310	G	N3-C2-N2	5.08	123.46	119.90
84	Aa	1526	A	O4'-C1'-N9	5.08	112.27	108.20
84	Aa	1702	C	N3-C4-C5	-5.08	119.87	121.90
84	Aa	2241	G	C5'-C4'-C3'	-5.08	107.87	116.00
2	Ae	9	A	O4'-C1'-N9	-5.08	104.14	108.20
2	Ae	72	G	C3'-C2'-C1'	-5.08	97.44	101.50
84	Aa	1393	G	O4'-C1'-N9	5.08	112.27	108.20
84	Aa	1421	A	C5-C6-N1	-5.08	115.16	117.70
84	Aa	2512	U	O4'-C4'-C3'	5.08	110.17	106.10
84	Aa	2833	G	C5-C6-O6	-5.08	125.55	128.60
84	Aa	3265	C	N3-C4-C5	-5.08	119.87	121.90
1	Ad	93	A	N9-C1'-C2'	-5.08	106.41	112.00
1	Ad	542	A	O4'-C1'-N9	5.08	112.26	108.20
1	Ad	1752	U	C1'-O4'-C4'	-5.08	105.84	109.90
84	Aa	211	A	C4-C5-C6	5.08	119.54	117.00
1	Ad	718	C	C4'-C3'-O3'	5.08	123.16	113.00
1	Ad	1487	U	C1'-O4'-C4'	5.08	113.96	109.90
1	Ad	1633	C	O4'-C1'-N1	5.08	112.26	108.20
84	Aa	287	A	C4-C5-C6	5.08	119.54	117.00
84	Aa	1671	G	C5-C6-O6	-5.08	125.55	128.60
84	Aa	2647	C	N3-C4-N4	5.08	121.56	118.00
84	Aa	3049	A	C5-C6-N1	-5.08	115.16	117.70
1	Ad	1554	G	C3'-C2'-C1'	-5.08	97.44	101.50
84	Aa	764	A	C5-C6-N1	-5.08	115.16	117.70
84	Aa	1969	G	N3-C2-N2	5.08	123.45	119.90
84	Aa	3036	C	N3-C4-N4	5.08	121.55	118.00
1	Ad	948	C	N1-C1'-C2'	5.08	120.60	114.00
1	Ad	1549	G	O4'-C1'-N9	-5.08	104.14	108.20
84	Aa	3	G	O4'-C1'-N9	5.08	112.26	108.20
84	Aa	279	G	C5-C6-O6	-5.08	125.55	128.60
84	Aa	1546	G	C5'-C4'-O4'	5.08	115.19	109.10
84	Aa	1712	A	C5-C6-N6	-5.08	119.64	123.70
84	Aa	2092	C	C5'-C4'-O4'	-5.08	103.01	109.10
84	Aa	2527	G	N3-C2-N2	5.08	123.45	119.90
84	Aa	2638	A	C5-C6-N1	-5.08	115.16	117.70
84	Aa	3252	G	C5'-C4'-C3'	-5.08	107.88	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	Ad	1664	U	C1'-O4'-C4'	5.07	113.96	109.90
13	BF	148	TYR	CB-CG-CD1	5.07	124.04	121.00
37	CG	222	PHE	CB-CG-CD1	-5.07	117.25	120.80
42	CJ	2	SER	N-CA-C	5.07	124.70	111.00
84	Aa	289	C	C6-N1-C2	-5.07	118.27	120.30
84	Aa	356	G	O4'-C1'-N9	5.07	112.26	108.20
84	Aa	2330	C	N3-C4-N4	5.07	121.55	118.00
1	Ad	960	A	O4'-C1'-C2'	-5.07	100.73	105.80
1	Ad	1390	A	C3'-C2'-C1'	-5.07	97.44	101.50
84	Aa	84	A	C5-C6-N6	-5.07	119.64	123.70
84	Aa	264	C	N3-C4-C5	-5.07	119.87	121.90
84	Aa	315	A	C5-C6-N1	-5.07	115.17	117.70
84	Aa	705	A	O4'-C1'-N9	5.07	112.26	108.20
84	Aa	1040	A	C5-C6-N1	-5.07	115.17	117.70
84	Aa	1808	G	O4'-C1'-N9	5.07	112.26	108.20
84	Aa	1839	C	N3-C4-N4	5.07	121.55	118.00
84	Aa	2382	C	N3-C4-C5	-5.07	119.87	121.90
84	Aa	2675	G	O4'-C1'-N9	5.07	112.26	108.20
84	Aa	2745	C	C2-N3-C4	5.07	122.44	119.90
84	Aa	2992	G	N1-C2-N3	-5.07	120.86	123.90
84	Aa	3060	G	N3-C2-N2	5.07	123.45	119.90
84	Aa	3118	C	N3-C4-N4	5.07	121.55	118.00
86	Ab	81	G	C5-N7-C8	5.07	106.83	104.30
79	CE	26	TRP	N-CA-CB	5.07	119.72	110.60
84	Aa	975	G	N3-C2-N2	5.07	123.45	119.90
84	Aa	2274	A	O4'-C1'-N9	5.07	112.25	108.20
84	Aa	3296	C	N3-C4-C5	-5.07	119.87	121.90
84	Aa	1434	G	O4'-C1'-N9	5.07	112.25	108.20
84	Aa	2204	U	O4'-C1'-N1	5.07	112.25	108.20
84	Aa	3035	C	N3-C4-C5	-5.07	119.87	121.90
84	Aa	3166	C	C2-N3-C4	5.07	122.43	119.90
84	Aa	3273	C	N3-C4-C5	-5.07	119.87	121.90
85	Ac	6	C	N3-C4-N4	5.07	121.55	118.00
22	BZ	28	TRP	N-CA-CB	5.07	119.72	110.60
84	Aa	493	G	C1'-O4'-C4'	-5.07	105.85	109.90
84	Aa	932	A	C5-C6-N6	-5.07	119.65	123.70
84	Aa	1207	A	C4-C5-C6	5.07	119.53	117.00
84	Aa	2450	G	O4'-C1'-N9	5.07	112.25	108.20
84	Aa	3010	G	N3-C2-N2	5.07	123.45	119.90
84	Aa	3131	A	C5-C6-N6	-5.07	119.65	123.70
84	Aa	3236	A	C5-C6-N1	-5.07	115.17	117.70
84	Aa	405	A	C5-C6-N6	-5.06	119.65	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	463	G	C5-C6-O6	-5.06	125.56	128.60
84	Aa	1682	C	N3-C4-C5	-5.06	119.87	121.90
84	Aa	141	C	N3-C4-C5	-5.06	119.88	121.90
84	Aa	1079	G	C5'-C4'-O4'	5.06	115.17	109.10
84	Aa	1661	G	O4'-C1'-N9	5.06	112.25	108.20
84	Aa	1190	C	N3-C4-N4	5.06	121.54	118.00
86	Ab	79	A	C4-C5-N7	-5.06	108.17	110.70
84	Aa	15	C	N3-C4-C5	-5.06	119.88	121.90
84	Aa	97	G	C6-C5-N7	-5.06	127.36	130.40
84	Aa	115	C	C1'-O4'-C4'	-5.06	105.85	109.90
84	Aa	169	G	C5-C6-O6	-5.06	125.56	128.60
84	Aa	1443	G	N3-C2-N2	5.06	123.44	119.90
84	Aa	2446	G	C5-C6-O6	-5.06	125.56	128.60
1	Ad	176	A	C3'-C2'-C1'	5.06	105.55	101.50
17	BS	83	PHE	N-CA-CB	5.06	119.70	110.60
84	Aa	572	U	O4'-C1'-N1	5.06	112.25	108.20
84	Aa	594	C	N3-C4-N4	5.06	121.54	118.00
84	Aa	796	C	N3-C4-C5	-5.06	119.88	121.90
84	Aa	1588	G	C5'-C4'-C3'	5.06	124.09	116.00
84	Aa	2074	C	C5'-C4'-O4'	-5.06	103.03	109.10
84	Aa	2303	C	N3-C4-N4	5.06	121.54	118.00
1	Ad	1443	U	N1-C1'-C2'	5.06	120.57	114.00
1	Ad	1169	G	C5'-C4'-O4'	5.05	115.17	109.10
84	Aa	478	G	O4'-C1'-N9	5.05	112.24	108.20
84	Aa	645	C	C2-N3-C4	5.05	122.43	119.90
84	Aa	1457	A	C5-C6-N1	-5.05	115.17	117.70
84	Aa	2295	G	O4'-C1'-N9	5.05	112.24	108.20
84	Aa	2357	A	C5-C6-N1	-5.05	115.17	117.70
1	Ad	838	U	C3'-C2'-C1'	5.05	105.54	101.50
84	Aa	350	A	C5-C6-N6	-5.05	119.66	123.70
1	Ad	182	C	C3'-C2'-C1'	5.05	105.54	101.50
84	Aa	161	C	N3-C4-C5	-5.05	119.88	121.90
84	Aa	1318	C	P-O5'-C5'	5.05	128.98	120.90
84	Aa	2557	C	C6-N1-C2	-5.05	118.28	120.30
84	Aa	2617	G	O4'-C1'-N9	5.05	112.24	108.20
84	Aa	3229	C	N3-C4-N4	5.05	121.54	118.00
85	Ac	85	G	C6-C5-N7	-5.05	127.37	130.40
86	Ab	3	A	C6-N1-C2	5.05	121.63	118.60
1	Ad	529	A	O4'-C1'-N9	5.05	112.24	108.20
1	Ad	1151	G	C5'-C4'-O4'	5.05	115.16	109.10
84	Aa	156	A	O4'-C1'-N9	5.05	112.24	108.20
84	Aa	2482	A	C5-C6-N1	-5.05	115.18	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2586	C	N3-C4-N4	5.05	121.53	118.00
1	Ad	780	A	O4'-C1'-C2'	5.05	112.14	107.60
84	Aa	64	A	O4'-C1'-N9	5.05	112.24	108.20
84	Aa	858	U	O4'-C1'-N1	5.05	112.24	108.20
84	Aa	2835	A	C5-C6-N6	-5.05	119.66	123.70
86	Ab	54	A	C5-C6-N6	-5.05	119.66	123.70
1	Ad	299	A	C1'-O4'-C4'	-5.05	105.86	109.90
1	Ad	421	A	C3'-C2'-C1'	5.05	105.54	101.50
1	Ad	1071	C	C1'-O4'-C4'	-5.05	105.86	109.90
84	Aa	52	G	C5-C6-O6	-5.05	125.57	128.60
84	Aa	124	C	C2-N3-C4	5.05	122.42	119.90
84	Aa	715	A	C4-C5-C6	5.05	119.52	117.00
84	Aa	1365	C	C2-N3-C4	5.05	122.42	119.90
84	Aa	2124	G	N3-C2-N2	5.05	123.43	119.90
84	Aa	2198	U	C5'-C4'-C3'	-5.05	107.92	116.00
84	Aa	2948	A	C5-C6-N1	-5.05	115.18	117.70
1	Ad	1268	G	C3'-C2'-C1'	5.04	105.54	101.50
2	Ae	1	U	P-O3'-C3'	5.04	125.75	119.70
86	Ab	115	A	O4'-C1'-N9	5.04	112.24	108.20
1	Ad	334	G	C1'-O4'-C4'	-5.04	105.86	109.90
1	Ad	1368	C	P-O3'-C3'	5.04	125.75	119.70
84	Aa	182	C	N3-C4-C5	-5.04	119.88	121.90
84	Aa	394	A	C5-C6-N1	-5.04	115.18	117.70
84	Aa	417	G	N1-C6-O6	5.04	122.93	119.90
84	Aa	1051	A	C5-C6-N1	-5.04	115.18	117.70
84	Aa	2014	A	P-O3'-C3'	5.04	125.75	119.70
84	Aa	2086	A	C4'-C3'-C2'	5.04	107.64	102.60
84	Aa	3308	A	C5-C6-N6	-5.04	119.67	123.70
1	Ad	838	U	P-O3'-C3'	5.04	125.75	119.70
84	Aa	1478	A	C4-C5-C6	5.04	119.52	117.00
84	Aa	1753	A	O4'-C1'-N9	5.04	112.23	108.20
84	Aa	3278	G	N3-C2-N2	5.04	123.43	119.90
1	Ad	735	G	N9-C1'-C2'	5.04	120.55	114.00
84	Aa	2152	A	C2'-C3'-O3'	5.04	121.76	113.70
1	Ad	239	C	O4'-C1'-N1	5.04	112.23	108.20
1	Ad	934	A	N9-C1'-C2'	5.04	120.55	114.00
84	Aa	61	A	O4'-C1'-N9	5.04	112.23	108.20
84	Aa	1050	A	C5-C6-N1	-5.04	115.18	117.70
84	Aa	1239	U	P-O3'-C3'	5.04	125.75	119.70
84	Aa	1376	A	C5-C6-N6	-5.04	119.67	123.70
84	Aa	1929	A	C5-C6-N1	-5.04	115.18	117.70
84	Aa	1974	C	N3-C4-N4	5.04	121.53	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2646	A	C5-C6-N1	-5.04	115.18	117.70
84	Aa	2690	G	O4'-C1'-N9	5.04	112.23	108.20
84	Aa	3374	C	C6-N1-C1'	-5.04	114.75	120.80
86	Ab	8	A	O4'-C1'-N9	5.04	112.23	108.20
1	Ad	774	C	N1-C1'-C2'	5.04	120.55	114.00
1	Ad	961	U	O4'-C1'-C2'	-5.04	100.76	105.80
1	Ad	1040	G	C1'-O4'-C4'	-5.04	105.87	109.90
48	CD	219	PHE	CB-CG-CD2	-5.04	117.27	120.80
84	Aa	135	G	O4'-C1'-N9	5.04	112.23	108.20
84	Aa	164	C	N3-C4-C5	-5.04	119.89	121.90
84	Aa	3368	A	C5-C6-N6	-5.04	119.67	123.70
1	Ad	1024	A	O4'-C1'-N9	5.04	112.23	108.20
84	Aa	319	C	C2-N3-C4	5.04	122.42	119.90
84	Aa	1369	G	O4'-C1'-N9	5.04	112.23	108.20
84	Aa	1631	G	N3-C2-N2	5.04	123.42	119.90
84	Aa	2164	G	C5-C6-O6	-5.04	125.58	128.60
84	Aa	2941	G	N3-C2-N2	5.04	123.42	119.90
85	Ac	9	G	N1-C2-N3	-5.04	120.88	123.90
1	Ad	219	G	O4'-C1'-C2'	-5.03	100.77	105.80
1	Ad	651	G	C1'-O4'-C4'	-5.03	105.87	109.90
1	Ad	1465	C	O4'-C1'-N1	5.03	112.23	108.20
48	CD	280	SER	N-CA-CB	5.03	118.05	110.50
84	Aa	309	C	N3-C4-N4	5.03	121.52	118.00
84	Aa	441	G	O4'-C1'-N9	5.03	112.23	108.20
84	Aa	526	A	C5-C6-N1	-5.03	115.18	117.70
84	Aa	573	A	O4'-C1'-N9	5.03	112.23	108.20
84	Aa	860	G	C4-C5-C6	5.03	121.82	118.80
84	Aa	984	A	C5-C6-N1	-5.03	115.18	117.70
84	Aa	1157	A	O4'-C1'-N9	5.03	112.23	108.20
84	Aa	2120	A	C5-C6-N6	-5.03	119.67	123.70
84	Aa	2266	A	O4'-C1'-N9	5.03	112.23	108.20
84	Aa	2540	C	N3-C4-N4	5.03	121.52	118.00
1	Ad	279	C	C1'-O4'-C4'	5.03	113.93	109.90
84	Aa	2853	A	O4'-C1'-N9	5.03	112.22	108.20
84	Aa	3290	C	P-O3'-C3'	5.03	125.74	119.70
1	Ad	30	G	O4'-C1'-C2'	5.03	112.13	107.60
1	Ad	446	C	C3'-C2'-C1'	5.03	105.52	101.50
84	Aa	333	G	C8-N9-C1'	5.03	133.54	127.00
84	Aa	934	C	N3-C4-C5	-5.03	119.89	121.90
84	Aa	996	A	O4'-C1'-N9	5.03	112.22	108.20
84	Aa	2165	A	O4'-C1'-N9	5.03	112.22	108.20
84	Aa	2489	A	C4-C5-C6	5.03	119.52	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2518	A	O4'-C1'-N9	5.03	112.22	108.20
84	Aa	2842	C	N3-C4-C5	-5.03	119.89	121.90
84	Aa	1042	C	N3-C4-N4	5.03	121.52	118.00
84	Aa	3182	A	O4'-C4'-C3'	-5.03	98.97	104.00
84	Aa	232	C	C5'-C4'-O4'	5.03	115.13	109.10
84	Aa	389	A	O4'-C1'-N9	5.03	112.22	108.20
84	Aa	721	A	C5-C6-N1	-5.03	115.19	117.70
84	Aa	1465	A	C4-C5-C6	5.03	119.51	117.00
84	Aa	1490	A	P-O3'-C3'	5.03	125.73	119.70
84	Aa	1509	G	O4'-C1'-N9	5.03	112.22	108.20
84	Aa	1968	C	N3-C4-N4	5.03	121.52	118.00
84	Aa	2622	G	N3-C2-N2	5.03	123.42	119.90
84	Aa	3139	U	P-O3'-C3'	5.03	125.73	119.70
1	Ad	78	A	O4'-C1'-N9	5.03	112.22	108.20
1	Ad	1466	A	P-O3'-C3'	5.03	125.73	119.70
84	Aa	300	C	C2-N3-C4	5.03	122.41	119.90
84	Aa	492	G	N3-C2-N2	5.03	123.42	119.90
84	Aa	3283	G	C5-C6-O6	-5.03	125.58	128.60
85	Ac	78	G	O4'-C1'-N9	5.03	112.22	108.20
85	Ac	90	C	N3-C4-C5	-5.03	119.89	121.90
86	Ab	83	A	C4-C5-C6	5.03	119.51	117.00
1	Ad	403	A	N9-C1'-C2'	-5.02	106.47	112.00
1	Ad	1343	C	C5'-C4'-C3'	-5.02	107.96	116.00
84	Aa	553	C	P-O3'-C3'	5.02	125.73	119.70
84	Aa	2205	G	N3-C2-N2	5.02	123.42	119.90
85	Ac	74	U	O4'-C1'-N1	5.02	112.22	108.20
1	Ad	619	A	C1'-O4'-C4'	5.02	113.92	109.90
1	Ad	1652	C	O4'-C1'-N1	5.02	112.22	108.20
84	Aa	1330	A	C4'-C3'-C2'	-5.02	97.58	102.60
84	Aa	1522	G	C5-C6-O6	-5.02	125.59	128.60
84	Aa	2858	G	O4'-C1'-N9	5.02	112.22	108.20
85	Ac	54	A	C5-C6-N1	-5.02	115.19	117.70
1	Ad	1280	U	C1'-O4'-C4'	5.02	113.92	109.90
2	Ae	4	G	C1'-O4'-C4'	-5.02	105.88	109.90
84	Aa	544	C	N3-C4-N4	5.02	121.52	118.00
84	Aa	1389	C	N3-C4-N4	5.02	121.52	118.00
84	Aa	2872	C	N3-C4-C5	-5.02	119.89	121.90
1	Ad	601	G	O4'-C1'-N9	5.02	112.22	108.20
21	BP	71	LYS	N-CA-CB	5.02	119.63	110.60
84	Aa	1006	A	C5-C6-N1	-5.02	115.19	117.70
84	Aa	1070	G	C5-C6-O6	-5.02	125.59	128.60
84	Aa	1967	C	N3-C4-N4	5.02	121.51	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	2866	A	O4'-C1'-N9	5.02	112.22	108.20
85	Ac	15	G	N3-C2-N2	5.02	123.41	119.90
1	Ad	1488	C	O4'-C1'-C2'	-5.02	100.78	105.80
73	CO	134	LEU	C-N-CA	5.02	134.24	121.70
84	Aa	951	C	N3-C4-N4	5.02	121.51	118.00
84	Aa	1452	A	O4'-C1'-N9	5.02	112.22	108.20
84	Aa	1479	G	N3-C2-N2	5.02	123.41	119.90
84	Aa	1643	A	C5-C6-N1	-5.02	115.19	117.70
84	Aa	1781	C	N3-C4-N4	5.02	121.51	118.00
84	Aa	1806	C	N3-C4-N4	5.02	121.51	118.00
84	Aa	2216	G	N1-C2-N3	-5.02	120.89	123.90
84	Aa	2502	U	P-O5'-C5'	5.02	128.93	120.90
84	Aa	2889	A	C5-C6-N1	-5.02	115.19	117.70
84	Aa	3114	A	O4'-C1'-N9	5.02	112.21	108.20
84	Aa	3275	G	N3-C2-N2	5.02	123.41	119.90
1	Ad	1405	U	C5'-C4'-O4'	5.02	115.12	109.10
84	Aa	204	G	O4'-C1'-N9	5.02	112.21	108.20
84	Aa	924	A	C5-C6-N1	-5.02	115.19	117.70
84	Aa	2208	A	C5-C6-N6	-5.02	119.69	123.70
1	Ad	1715	C	N1-C1'-C2'	5.01	120.52	114.00
2	Ae	64	G	C1'-O4'-C4'	-5.01	105.89	109.90
53	CY	9	SER	N-CA-CB	5.01	118.02	110.50
84	Aa	621	C	N3-C4-C5	-5.01	119.89	121.90
84	Aa	2267	G	C5-C6-O6	-5.01	125.59	128.60
86	Ab	101	A	C6-C5-N7	-5.01	128.79	132.30
84	Aa	328	G	O4'-C1'-N9	5.01	112.21	108.20
84	Aa	3056	C	N3-C4-N4	5.01	121.51	118.00
84	Aa	1017	G	O4'-C1'-N9	5.01	112.21	108.20
84	Aa	1454	C	C2-N3-C4	5.01	122.41	119.90
84	Aa	2285	C	N3-C4-N4	5.01	121.51	118.00
84	Aa	2387	U	O4'-C1'-N1	5.01	112.21	108.20
84	Aa	2919	G	N3-C2-N2	5.01	123.41	119.90
1	Ad	649	C	N1-C1'-C2'	5.01	120.51	114.00
84	Aa	397	A	O4'-C1'-N9	5.01	112.21	108.20
84	Aa	801	G	O4'-C1'-N9	5.01	112.21	108.20
84	Aa	1018	C	P-O3'-C3'	5.01	125.71	119.70
84	Aa	1065	A	O4'-C1'-N9	5.01	112.21	108.20
84	Aa	1264	A	C5-C6-N6	-5.01	119.69	123.70
84	Aa	1311	G	C2'-C3'-O3'	5.01	121.72	113.70
84	Aa	1877	G	O4'-C1'-N9	5.01	112.21	108.20
84	Aa	2088	C	C4'-C3'-C2'	-5.01	97.59	102.60
84	Aa	2505	C	N3-C4-C5	-5.01	119.90	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
84	Aa	3030	A	C5-C6-N6	-5.01	119.69	123.70
1	Ad	559	A	C1'-O4'-C4'	5.01	113.91	109.90
45	CN	53	TYR	CB-CG-CD1	-5.01	118.00	121.00
84	Aa	204	G	C5-C6-O6	-5.01	125.60	128.60
84	Aa	820	A	O4'-C1'-N9	5.01	112.21	108.20
84	Aa	1024	G	O4'-C1'-N9	5.01	112.21	108.20
84	Aa	1130	G	O4'-C1'-N9	5.01	112.20	108.20
84	Aa	2301	C	C6-N1-C1'	-5.01	114.79	120.80
84	Aa	2649	C	N3-C4-N4	5.01	121.50	118.00
84	Aa	3072	A	O4'-C1'-N9	5.01	112.21	108.20
84	Aa	3203	G	N3-C2-N2	5.01	123.40	119.90
84	Aa	3337	G	O4'-C1'-N9	5.01	112.20	108.20
84	Aa	789	A	C5-C6-N1	-5.00	115.20	117.70
84	Aa	1675	G	C8-N9-C1'	5.00	133.51	127.00
84	Aa	1774	G	N3-C2-N2	5.00	123.40	119.90
86	Ab	64	G	N1-C6-O6	5.00	122.90	119.90
1	Ad	19	A	O4'-C1'-C2'	-5.00	100.80	105.80
1	Ad	1349	A	N9-C1'-C2'	5.00	120.51	114.00
2	Ae	26	G	O4'-C1'-N9	5.00	112.20	108.20
84	Aa	348	C	O4'-C1'-N1	5.00	112.20	108.20
84	Aa	1722	G	C5-C6-O6	-5.00	125.60	128.60
1	Ad	56	U	P-O3'-C3'	5.00	125.70	119.70
1	Ad	341	G	O4'-C1'-N9	-5.00	104.20	108.20
1	Ad	550	U	C3'-C2'-C1'	5.00	105.50	101.50
1	Ad	783	C	O4'-C1'-N1	5.00	112.20	108.20
84	Aa	245	C	N3-C4-N4	5.00	121.50	118.00
84	Aa	617	C	C5-C6-N1	5.00	123.50	121.00
84	Aa	980	C	C2-N3-C4	5.00	122.40	119.90
84	Aa	2143	A	C4-C5-C6	5.00	119.50	117.00
84	Aa	2185	U	C5-C6-N1	5.00	125.20	122.70
84	Aa	2959	G	O4'-C1'-N9	5.00	112.20	108.20

There are no chirality outliers.

All (486) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
84	Aa	1004	C	Sidechain
84	Aa	1010	A	Sidechain
84	Aa	1019	A	Sidechain
84	Aa	1022	G	Sidechain
84	Aa	1028	G	Sidechain
84	Aa	1051	A	Sidechain

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>Group</b>
84	Aa	106	G	Sidechain
84	Aa	1073	G	Sidechain
84	Aa	108	A	Sidechain
84	Aa	1083	C	Sidechain
84	Aa	11	A	Sidechain
84	Aa	1102	A	Sidechain
84	Aa	1109	G	Sidechain
84	Aa	111	C	Sidechain
84	Aa	1116	G	Sidechain
84	Aa	1118	G	Sidechain
84	Aa	1119	G	Sidechain
84	Aa	114	G	Sidechain
84	Aa	1150	G	Sidechain
84	Aa	1156	A	Sidechain
84	Aa	1157	A	Sidechain
84	Aa	1163	A	Sidechain
84	Aa	1164	G	Sidechain
84	Aa	1167	G	Sidechain
84	Aa	1177	G	Sidechain
84	Aa	1184	U	Sidechain
84	Aa	1187	G	Sidechain
84	Aa	120	G	Sidechain
84	Aa	1216	G	Sidechain
84	Aa	1230	G	Sidechain
84	Aa	1237	G	Sidechain
84	Aa	1241	G	Sidechain
84	Aa	1247	G	Sidechain
84	Aa	126	G	Sidechain
84	Aa	1262	U	Sidechain
84	Aa	1267	A	Sidechain
84	Aa	1275	A	Sidechain
84	Aa	1297	U	Sidechain
84	Aa	1298	A	Sidechain
84	Aa	13	G	Sidechain
84	Aa	1308	A	Sidechain
84	Aa	1312	A	Sidechain
84	Aa	1313	U	Sidechain
84	Aa	1320	G	Sidechain
84	Aa	1322	A	Sidechain
84	Aa	1323	G	Sidechain
84	Aa	1335	C	Sidechain
84	Aa	1341	G	Sidechain

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>Group</b>
84	Aa	1354	G	Sidechain
84	Aa	1409	G	Sidechain
84	Aa	144	A	Sidechain
84	Aa	1449	A	Sidechain
84	Aa	1457	A	Sidechain
84	Aa	147	G	Sidechain
84	Aa	1476	G	Sidechain
84	Aa	1482	C	Sidechain
84	Aa	1486	G	Sidechain
84	Aa	1489	G	Sidechain
84	Aa	1528	G	Sidechain
84	Aa	1531	G	Sidechain
84	Aa	1538	A	Sidechain
84	Aa	1542	A	Sidechain
84	Aa	158	A	Sidechain
84	Aa	1586	A	Sidechain
84	Aa	159	G	Sidechain
84	Aa	1594	G	Sidechain
84	Aa	1598	U	Sidechain
84	Aa	1601	G	Sidechain
84	Aa	1609	G	Sidechain
84	Aa	1611	G	Sidechain
84	Aa	1618	U	Sidechain
84	Aa	1619	G	Sidechain
84	Aa	1634	G	Sidechain
84	Aa	1635	A	Sidechain
84	Aa	1638	U	Sidechain
84	Aa	1653	A	Sidechain
84	Aa	1671	G	Sidechain
84	Aa	1672	G	Sidechain
84	Aa	1678	U	Sidechain
84	Aa	1689	G	Sidechain
84	Aa	1711	G	Sidechain
84	Aa	1716	G	Sidechain
84	Aa	1721	A	Sidechain
84	Aa	1723	C	Sidechain
84	Aa	1725	G	Sidechain
84	Aa	1728	G	Sidechain
84	Aa	1746	G	Sidechain
84	Aa	1747	A	Sidechain
84	Aa	1749	G	Sidechain
84	Aa	1750	A	Sidechain

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>Group</b>
84	Aa	1756	C	Sidechain
84	Aa	176	A	Sidechain
84	Aa	1767	G	Sidechain
84	Aa	18	G	Sidechain
84	Aa	1804	G	Sidechain
84	Aa	1816	U	Sidechain
84	Aa	1821	G	Sidechain
84	Aa	1842	C	Sidechain
84	Aa	1853	C	Sidechain
84	Aa	1857	G	Sidechain
84	Aa	1859	G	Sidechain
84	Aa	186	A	Sidechain
84	Aa	188	U	Sidechain
84	Aa	1892	A	Sidechain
84	Aa	1903	C	Sidechain
84	Aa	1910	G	Sidechain
84	Aa	1912	U	Sidechain
84	Aa	1921	U	Sidechain
84	Aa	1923	G	Sidechain
84	Aa	1935	G	Sidechain
84	Aa	20	G	Sidechain
84	Aa	2075	C	Sidechain
84	Aa	2077	C	Sidechain
84	Aa	2088	C	Sidechain
84	Aa	2092	C	Sidechain
84	Aa	2093	G	Sidechain
84	Aa	2104	G	Sidechain
84	Aa	2113	A	Sidechain
84	Aa	2132	A	Sidechain
84	Aa	214	G	Sidechain
84	Aa	2149	G	Sidechain
84	Aa	2150	C	Sidechain
84	Aa	2167	G	Sidechain
84	Aa	2171	A	Sidechain
84	Aa	218	G	Sidechain
84	Aa	2189	G	Sidechain
84	Aa	2192	C	Sidechain
84	Aa	2218	A	Sidechain
84	Aa	2242	G	Sidechain
84	Aa	2269	U	Sidechain
84	Aa	2270	A	Sidechain
84	Aa	2290	A	Sidechain

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>Group</b>
84	Aa	2303	C	Sidechain
84	Aa	2348	U	Sidechain
84	Aa	235	G	Sidechain
84	Aa	2352	G	Sidechain
84	Aa	2354	G	Sidechain
84	Aa	2375	G	Sidechain
84	Aa	2376	G	Sidechain
84	Aa	2380	G	Sidechain
84	Aa	2381	G	Sidechain
84	Aa	2416	U	Sidechain
84	Aa	2424	G	Sidechain
84	Aa	244	G	Sidechain
84	Aa	2444	U	Sidechain
84	Aa	2451	G	Sidechain
84	Aa	2491	A	Sidechain
84	Aa	2502	U	Sidechain
84	Aa	2506	G	Sidechain
84	Aa	2508	U	Sidechain
84	Aa	2526	G	Sidechain
84	Aa	2528	U	Sidechain
84	Aa	2529	C	Sidechain
84	Aa	2537	G	Sidechain
84	Aa	2538	G	Sidechain
84	Aa	2539	G	Sidechain
84	Aa	2542	U	Sidechain
84	Aa	2543	G	Sidechain
84	Aa	2552	U	Sidechain
84	Aa	2557	C	Sidechain
84	Aa	2558	U	Sidechain
84	Aa	2583	A	Sidechain
84	Aa	2588	G	Sidechain
84	Aa	2591	G	Sidechain
84	Aa	2610	G	Sidechain
84	Aa	2623	G	Sidechain
84	Aa	2627	G	Sidechain
84	Aa	2639	A	Sidechain
84	Aa	2648	G	Sidechain
84	Aa	265	G	Sidechain
84	Aa	2666	G	Sidechain
84	Aa	2673	G	Sidechain
84	Aa	2677	A	Sidechain
84	Aa	2697	A	Sidechain

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>Group</b>
84	Aa	2709	G	Sidechain
84	Aa	2732	U	Sidechain
84	Aa	2733	A	Sidechain
84	Aa	2751	A	Sidechain
84	Aa	2789	G	Sidechain
84	Aa	279	G	Sidechain
84	Aa	2801	A	Sidechain
84	Aa	2807	G	Sidechain
84	Aa	2815	A	Sidechain
84	Aa	2820	U	Sidechain
84	Aa	2843	G	Sidechain
84	Aa	2877	U	Sidechain
84	Aa	2882	U	Sidechain
84	Aa	2888	U	Sidechain
84	Aa	2918	U	Sidechain
84	Aa	2932	A	Sidechain
84	Aa	2935	A	Sidechain
84	Aa	2936	A	Sidechain
84	Aa	2966	G	Sidechain
84	Aa	2967	U	Sidechain
84	Aa	2968	G	Sidechain
84	Aa	2992	G	Sidechain
84	Aa	3002	U	Sidechain
84	Aa	3006	G	Sidechain
84	Aa	3007	A	Sidechain
84	Aa	3012	A	Sidechain
84	Aa	302	G	Sidechain
84	Aa	3035	C	Sidechain
84	Aa	3057	A	Sidechain
84	Aa	3067	G	Sidechain
84	Aa	3097	G	Sidechain
84	Aa	3123	A	Sidechain
84	Aa	3125	G	Sidechain
84	Aa	3137	G	Sidechain
84	Aa	3143	A	Sidechain
84	Aa	3171	C	Sidechain
84	Aa	3179	G	Sidechain
84	Aa	318	G	Sidechain
84	Aa	3204	G	Sidechain
84	Aa	3208	G	Sidechain
84	Aa	321	A	Sidechain
84	Aa	3219	U	Sidechain

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>Group</b>
84	Aa	3220	A	Sidechain
84	Aa	3222	G	Sidechain
84	Aa	3242	G	Sidechain
84	Aa	3248	G	Sidechain
84	Aa	3252	G	Sidechain
84	Aa	3264	C	Sidechain
84	Aa	3267	U	Sidechain
84	Aa	3269	C	Sidechain
84	Aa	3271	A	Sidechain
84	Aa	3276	G	Sidechain
84	Aa	3286	G	Sidechain
84	Aa	3294	U	Sidechain
84	Aa	3295	G	Sidechain
84	Aa	3309	U	Sidechain
84	Aa	3312	G	Sidechain
84	Aa	3314	G	Sidechain
84	Aa	3315	A	Sidechain
84	Aa	3320	G	Sidechain
84	Aa	3334	A	Sidechain
84	Aa	3340	G	Sidechain
84	Aa	3345	G	Sidechain
84	Aa	3361	G	Sidechain
84	Aa	3377	G	Sidechain
84	Aa	3379	C	Sidechain
84	Aa	3380	G	Sidechain
84	Aa	339	G	Sidechain
84	Aa	342	A	Sidechain
84	Aa	352	U	Sidechain
84	Aa	370	A	Sidechain
84	Aa	372	A	Sidechain
84	Aa	373	A	Sidechain
84	Aa	404	G	Sidechain
84	Aa	413	G	Sidechain
84	Aa	424	G	Sidechain
84	Aa	425	G	Sidechain
84	Aa	431	G	Sidechain
84	Aa	436	G	Sidechain
84	Aa	492	G	Sidechain
84	Aa	493	G	Sidechain
84	Aa	497	G	Sidechain
84	Aa	522	C	Sidechain
84	Aa	526	A	Sidechain

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>Group</b>
84	Aa	527	G	Sidechain
84	Aa	53	C	Sidechain
84	Aa	566	G	Sidechain
84	Aa	602	G	Sidechain
84	Aa	603	G	Sidechain
84	Aa	604	C	Sidechain
84	Aa	62	A	Sidechain
84	Aa	647	U	Sidechain
84	Aa	666	U	Sidechain
84	Aa	669	G	Sidechain
84	Aa	67	C	Sidechain
84	Aa	676	G	Sidechain
84	Aa	688	G	Sidechain
84	Aa	693	C	Sidechain
84	Aa	713	G	Sidechain
84	Aa	723	G	Sidechain
84	Aa	732	G	Sidechain
84	Aa	736	U	Sidechain
84	Aa	746	C	Sidechain
84	Aa	75	G	Sidechain
84	Aa	757	G	Sidechain
84	Aa	763	G	Sidechain
84	Aa	765	U	Sidechain
84	Aa	768	U	Sidechain
84	Aa	771	G	Sidechain
84	Aa	772	U	Sidechain
84	Aa	773	G	Sidechain
84	Aa	775	A	Sidechain
84	Aa	779	U	Sidechain
84	Aa	783	A	Sidechain
84	Aa	787	G	Sidechain
84	Aa	801	G	Sidechain
84	Aa	808	G	Sidechain
84	Aa	818	G	Sidechain
84	Aa	825	G	Sidechain
84	Aa	835	G	Sidechain
84	Aa	838	G	Sidechain
84	Aa	844	A	Sidechain
84	Aa	849	A	Sidechain
84	Aa	85	G	Sidechain
84	Aa	858	U	Sidechain
84	Aa	861	A	Sidechain

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>Group</b>
84	Aa	877	U	Sidechain
84	Aa	886	A	Sidechain
84	Aa	911	G	Sidechain
84	Aa	93	G	Sidechain
84	Aa	936	A	Sidechain
84	Aa	966	G	Sidechain
84	Aa	97	G	Sidechain
84	Aa	977	G	Sidechain
84	Aa	997	G	Sidechain
86	Ab	104	C	Sidechain
86	Ab	117	U	Sidechain
86	Ab	17	G	Sidechain
86	Ab	30	G	Sidechain
86	Ab	33	U	Sidechain
86	Ab	37	G	Sidechain
86	Ab	40	A	Sidechain
86	Ab	42	A	Sidechain
86	Ab	46	C	Sidechain
86	Ab	56	G	Sidechain
86	Ab	61	C	Sidechain
86	Ab	62	U	Sidechain
86	Ab	79	A	Sidechain
86	Ab	83	A	Sidechain
86	Ab	88	U	Sidechain
86	Ab	9	U	Sidechain
86	Ab	90	A	Sidechain
86	Ab	96	U	Sidechain
86	Ab	97	G	Sidechain
85	Ac	100	U	Sidechain
85	Ac	105	A	Sidechain
85	Ac	146	G	Sidechain
85	Ac	147	C	Sidechain
85	Ac	149	U	Sidechain
85	Ac	16	G	Sidechain
85	Ac	22	U	Sidechain
85	Ac	34	U	Sidechain
85	Ac	49	G	Sidechain
85	Ac	51	G	Sidechain
85	Ac	53	A	Sidechain
85	Ac	60	U	Sidechain
85	Ac	64	U	Sidechain
85	Ac	69	U	Sidechain

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>Group</b>
85	Ac	70	G	Sidechain
85	Ac	74	U	Sidechain
85	Ac	87	G	Sidechain
85	Ac	9	G	Sidechain
30	BB	157	GLN	Peptide
11	BD	112	GLY	Peptide
11	BD	211	HIS	Peptide
11	BD	212	PRO	Peptide
12	BE	132	GLY	Peptide
12	BE	240	LYS	Peptide
13	BF	138	SER	Peptide
13	BF	44	TYR	Sidechain
35	BG	182	ARG	Peptide
36	BH	105	PRO	Peptide
36	BH	114	PRO	Peptide
36	BH	117	ARG	Sidechain
36	BH	184	PHE	Peptide
6	BK	22	TYR	Sidechain
6	BK	83	PRO	Peptide
6	BK	86	ILE	Peptide
6	BK	87	VAL	Peptide
19	BL	98	TYR	Sidechain
7	BM	92	CYS	Peptide
14	BQ	131	GLU	Peptide
29	BR	2	GLY	Peptide
17	BS	142	ARG	Sidechain
20	BT	37	VAL	Peptide
20	BT	46	LYS	Peptide
20	BT	91	ARG	Sidechain
20	BT	92	PRO	Peptide
15	BU	10	PRO	Peptide
15	BU	78	PRO	Peptide
24	BW	62	VAL	Peptide
4	BY	48	LYS	Peptide
32	Ba	87	ARG	Sidechain
26	Bb	29	SER	Peptide
26	Bb	3	LEU	Peptide
23	Bc	21	GLY	Peptide
25	Bd	13	ASN	Peptide
8	Bf	52	GLY	Peptide
10	Bg	301	VAL	Peptide
41	CA	122	ASP	Peptide

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>Group</b>
41	CA	244	GLY	Peptide
41	CA	63	PHE	Peptide
41	CA	70	LYS	Peptide
71	CB	117	ARG	Peptide
71	CB	119	TYR	Sidechain
71	CB	259	HIS	Sidechain
71	CB	291	SER	Peptide
71	CB	303	ASP	Peptide
71	CB	349	GLN	Peptide
71	CB	357	GLU	Peptide
71	CB	365	THR	Peptide
71	CB	42	HIS	Peptide
72	CC	201	ARG	Sidechain
72	CC	215	TYR	Sidechain
72	CC	24	SER	Peptide
72	CC	95	ALA	Peptide
48	CD	114	ARG	Peptide
48	CD	121	GLU	Peptide
48	CD	139	ARG	Peptide
48	CD	187	GLU	Peptide
48	CD	201	GLY	Peptide
48	CD	21	GLN	Peptide
48	CD	217	GLU	Peptide
48	CD	218	LYS	Peptide
48	CD	235	GLY	Peptide
48	CD	238	SER	Peptide
48	CD	244	HIS	Peptide
48	CD	257	THR	Peptide
48	CD	261	PRO	Peptide
79	CE	20	TYR	Sidechain
79	CE	27	ALA	Peptide
79	CE	36	LEU	Peptide
79	CE	48	PRO	Peptide
79	CE	49	LYS	Peptide
79	CE	51	TYR	Peptide
79	CE	71	LEU	Peptide
69	CF	204	LEU	Peptide
69	CF	35	GLU	Peptide
69	CF	77	PHE	Peptide
37	CG	50	TRP	Peptide
37	CG	74	ASN	Peptide
43	CH	168	ASN	Peptide

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>Group</b>
43	CH	169	LYS	Peptide
43	CH	183	LYS	Peptide
75	CI	110	ARG	Peptide
75	CI	172	GLY	Peptide
75	CI	29	PRO	Peptide
75	CI	88	ARG	Sidechain
42	CJ	2	SER	Peptide
42	CJ	9	ASN	Peptide
65	CK	75	VAL	Peptide
65	CK	76	PRO	Peptide
78	CL	12	HIS	Peptide
78	CL	13	PHE	Peptide,Sidechain
78	CL	20	TYR	Sidechain
78	CL	65	TYR	Sidechain
61	CM	17	TYR	Sidechain
61	CM	18	GLY	Peptide
61	CM	40	ALA	Peptide
61	CM	6	PHE	Peptide
45	CN	81	TYR	Sidechain
73	CO	115	PRO	Peptide
73	CO	125	ILE	Peptide
73	CO	152	TRP	Peptide
73	CO	70	LYS	Peptide
50	CP	3	LYS	Peptide
47	CQ	14	THR	Peptide
47	CQ	156	PRO	Peptide
47	CQ	157	GLY	Peptide
47	CQ	158	VAL	Peptide
47	CQ	16	ARG	Peptide
47	CQ	59	ARG	Sidechain
49	CR	73	GLY	Peptide
49	CR	84	THR	Peptide
49	CR	89	LEU	Peptide
62	CS	139	ASP	Peptide
62	CS	151	PHE	Peptide
62	CS	82	ARG	Sidechain
63	CU	100	ASP	Peptide
63	CU	55	LYS	Peptide
51	CX	133	TYR	Sidechain
46	Ca	103	TYR	Sidechain
46	Ca	116	ARG	Peptide
46	Ca	137	GLY	Peptide

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Mol	Chain	Res	Type	Group
46	Ca	39	HIS	Peptide
46	Ca	5	PHE	Peptide
46	Ca	60	TYR	Peptide
46	Ca	8	ASN	Peptide
55	Cc	52	CYS	Peptide
57	Ce	13	LYS	Peptide
80	Cf	20	TYR	Sidechain
80	Cf	4	ARG	Sidechain
83	Cg	1	MET	Peptide
68	Ch	90	ARG	Peptide
58	Cj	88	LYS	Peptide
59	Cl	3	SER	Peptide
60	Co	32	LYS	Peptide
60	Co	87	ARG	Sidechain
74	Cp	35	SER	Peptide
70	Cq	7	LYS	Peptide
70	Cq	9	GLU	Peptide

## 5.2 Too-close contacts [i](#)

Due to software issues we are unable to calculate clashes - this section is therefore empty.

## 5.3 Torsion angles [i](#)

### 5.3.1 Protein backbone [i](#)

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
4	BY	136/138 (99%)	118 (87%)	8 (6%)	10 (7%)	1	13
5	BI	64/220 (29%)	61 (95%)	2 (3%)	1 (2%)	9	43
6	BK	94/183 (51%)	66 (70%)	17 (18%)	11 (12%)	0	6
7	BM	121/171 (71%)	84 (69%)	20 (16%)	17 (14%)	0	4
8	Bf	69/155 (44%)	46 (67%)	10 (14%)	13 (19%)	0	2

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
9	BX	140/142 (99%)	124 (89%)	11 (8%)	5 (4%)	3	25
10	Bg	378/380 (100%)	334 (88%)	26 (7%)	18 (5%)	2	21
11	BD	206/208 (99%)	125 (61%)	34 (16%)	47 (23%)	0	1
12	BE	198/265 (75%)	173 (87%)	16 (8%)	9 (4%)	2	22
13	BF	189/191 (99%)	162 (86%)	20 (11%)	7 (4%)	3	25
14	BQ	124/149 (83%)	93 (75%)	15 (12%)	16 (13%)	0	5
15	BU	126/128 (98%)	102 (81%)	14 (11%)	10 (8%)	1	12
16	BO	117/151 (78%)	91 (78%)	12 (10%)	14 (12%)	0	6
17	BS	150/152 (99%)	109 (73%)	16 (11%)	25 (17%)	0	3
18	BN	119/151 (79%)	92 (77%)	14 (12%)	13 (11%)	0	8
19	BL	83/160 (52%)	61 (74%)	16 (19%)	6 (7%)	1	14
20	BT	144/146 (99%)	123 (85%)	13 (9%)	8 (6%)	2	18
21	BP	89/154 (58%)	69 (78%)	12 (14%)	8 (9%)	1	11
22	BZ	98/108 (91%)	75 (76%)	10 (10%)	13 (13%)	0	4
23	Bc	56/65 (86%)	40 (71%)	5 (9%)	11 (20%)	0	2
24	BW	128/130 (98%)	101 (79%)	16 (12%)	11 (9%)	1	11
25	Bd	46/56 (82%)	29 (63%)	6 (13%)	11 (24%)	0	1
26	Bb	84/86 (98%)	75 (89%)	6 (7%)	3 (4%)	3	25
27	Be	58/62 (94%)	49 (84%)	5 (9%)	4 (7%)	1	14
28	BA	195/260 (75%)	176 (90%)	10 (5%)	9 (5%)	2	21
29	BR	114/141 (81%)	89 (78%)	15 (13%)	10 (9%)	1	11
30	BB	209/262 (80%)	153 (73%)	31 (15%)	25 (12%)	0	6
31	BV	74/82 (90%)	62 (84%)	9 (12%)	3 (4%)	3	22
32	Ba	91/133 (68%)	65 (71%)	13 (14%)	13 (14%)	0	4
33	BJ	185/195 (95%)	162 (88%)	16 (9%)	7 (4%)	3	24
34	BC	212/263 (81%)	189 (89%)	16 (8%)	7 (3%)	4	26
35	BG	227/245 (93%)	211 (93%)	10 (4%)	6 (3%)	5	31
36	BH	182/189 (96%)	154 (85%)	10 (6%)	18 (10%)	0	9
37	CG	235/257 (91%)	205 (87%)	24 (10%)	6 (3%)	5	31
38	CT	158/164 (96%)	137 (87%)	6 (4%)	15 (10%)	0	10
39	CZ	134/136 (98%)	123 (92%)	10 (8%)	1 (1%)	22	62

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
40	Cz	214/216 (99%)	197 (92%)	9 (4%)	8 (4%)	3	25
41	CA	253/261 (97%)	219 (87%)	19 (8%)	15 (6%)	1	17
42	CJ	168/180 (93%)	132 (79%)	14 (8%)	22 (13%)	0	4
43	CH	188/190 (99%)	167 (89%)	16 (8%)	5 (3%)	5	31
44	CV	138/140 (99%)	124 (90%)	7 (5%)	7 (5%)	2	19
45	CN	192/200 (96%)	168 (88%)	18 (9%)	6 (3%)	4	27
46	Ca	142/144 (99%)	101 (71%)	24 (17%)	17 (12%)	0	6
47	CQ	161/188 (86%)	127 (79%)	18 (11%)	16 (10%)	0	9
48	CD	302/304 (99%)	213 (70%)	35 (12%)	54 (18%)	0	3
49	CR	187/209 (90%)	163 (87%)	14 (8%)	10 (5%)	2	19
50	CP	169/171 (99%)	140 (83%)	12 (7%)	17 (10%)	0	9
51	CX	120/152 (79%)	100 (83%)	17 (14%)	3 (2%)	5	32
52	CW	73/162 (45%)	55 (75%)	12 (16%)	6 (8%)	1	12
53	CY	128/150 (85%)	114 (89%)	8 (6%)	6 (5%)	2	21
54	Cr	71/147 (48%)	49 (69%)	13 (18%)	9 (13%)	0	5
55	Cc	110/112 (98%)	96 (87%)	10 (9%)	4 (4%)	3	25
56	Cd	118/123 (96%)	98 (83%)	8 (7%)	12 (10%)	0	8
57	Ce	131/133 (98%)	113 (86%)	10 (8%)	8 (6%)	1	16
58	Cj	92/94 (98%)	58 (63%)	19 (21%)	15 (16%)	0	3
59	Cl	49/51 (96%)	36 (74%)	8 (16%)	5 (10%)	0	8
60	Co	103/105 (98%)	76 (74%)	13 (13%)	14 (14%)	0	4
61	CM	132/134 (98%)	101 (76%)	14 (11%)	17 (13%)	0	5
62	CS	165/178 (93%)	122 (74%)	20 (12%)	23 (14%)	0	4
63	CU	106/130 (82%)	76 (72%)	13 (12%)	17 (16%)	0	3
64	Ci	75/112 (67%)	59 (79%)	5 (7%)	11 (15%)	0	4
65	CK	126/166 (76%)	94 (75%)	17 (14%)	15 (12%)	0	6
66	Cu	56/110 (51%)	54 (96%)	1 (2%)	1 (2%)	8	40
66	Cv	56/110 (51%)	53 (95%)	2 (4%)	1 (2%)	8	40
67	Cs	57/113 (50%)	54 (95%)	3 (5%)	0	100	100
67	Ct	57/113 (50%)	54 (95%)	3 (5%)	0	100	100
68	Ch	122/124 (98%)	103 (84%)	11 (9%)	8 (7%)	1	15

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
69	CF	242/244 (99%)	217 (90%)	16 (7%)	9 (4%)	3	25
70	Cq	260/319 (82%)	233 (90%)	15 (6%)	12 (5%)	2	21
71	CB	387/389 (100%)	307 (79%)	43 (11%)	37 (10%)	0	9
72	CC	368/405 (91%)	311 (84%)	27 (7%)	30 (8%)	1	12
73	CO	204/206 (99%)	179 (88%)	14 (7%)	11 (5%)	2	19
74	Cp	90/92 (98%)	81 (90%)	7 (8%)	2 (2%)	6	35
75	CI	182/224 (81%)	147 (81%)	24 (13%)	11 (6%)	1	16
76	Cn	23/25 (92%)	21 (91%)	1 (4%)	1 (4%)	2	22
77	Cm	50/53 (94%)	46 (92%)	3 (6%)	1 (2%)	7	37
78	CL	206/208 (99%)	168 (82%)	13 (6%)	25 (12%)	0	5
79	CE	217/219 (99%)	177 (82%)	14 (6%)	26 (12%)	0	6
80	Cf	109/111 (98%)	103 (94%)	5 (5%)	1 (1%)	17	56
81	Ck	67/69 (97%)	63 (94%)	2 (3%)	2 (3%)	4	28
82	Cb	56/60 (93%)	48 (86%)	4 (7%)	4 (7%)	1	14
83	Cg	108/119 (91%)	96 (89%)	8 (7%)	4 (4%)	3	25
All	All	11663/13543 (86%)	9641 (83%)	1083 (9%)	939 (8%)	2	12

All (939) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
4	BY	2	ALA
4	BY	39	ASN
4	BY	41	SER
4	BY	46	LYS
4	BY	49	LEU
4	BY	68	THR
6	BK	82	LEU
6	BK	87	VAL
6	BK	88	PRO
6	BK	89	ALA
7	BM	79	VAL
7	BM	81	SER
7	BM	96	SER
8	Bf	26	VAL
9	BX	60	GLN
9	BX	128	SER
10	Bg	2	ALA

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
10	Bg	149	VAL
10	Bg	216	LEU
10	Bg	342	SER
10	Bg	343	HIS
11	BD	32	ASP
11	BD	48	ILE
11	BD	90	LYS
11	BD	96	LEU
11	BD	99	ILE
11	BD	111	GLY
11	BD	125	PHE
11	BD	126	VAL
11	BD	131	ALA
11	BD	156	TYR
11	BD	162	GLN
11	BD	183	GLY
11	BD	184	ILE
11	BD	197	LYS
11	BD	200	PRO
11	BD	213	PRO
11	BD	215	GLU
11	BD	216	GLU
11	BD	217	ASN
11	BD	218	GLU
12	BE	58	TYR
12	BE	153	ILE
12	BE	154	ILE
13	BF	41	HIS
13	BF	57	PHE
13	BF	63	PRO
13	BF	77	ARG
14	BQ	45	ILE
14	BQ	76	ARG
14	BQ	90	ALA
14	BQ	91	ILE
14	BQ	142	ALA
15	BU	3	ALA
15	BU	7	ALA
15	BU	9	ALA
15	BU	12	MET
15	BU	81	GLU
16	BO	66	ASP

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
16	BO	72	ALA
16	BO	97	ARG
16	BO	134	VAL
16	BO	148	ARG
17	BS	5	ALA
17	BS	8	GLU
17	BS	9	PHE
17	BS	17	ASN
17	BS	74	PRO
17	BS	80	PRO
17	BS	87	LYS
17	BS	99	VAL
17	BS	100	SER
17	BS	149	SER
18	BN	42	LYS
18	BN	57	GLN
18	BN	69	SER
18	BN	81	ALA
18	BN	86	GLU
18	BN	109	LYS
18	BN	137	PRO
19	BL	55	ILE
19	BL	109	PRO
19	BL	110	ALA
19	BL	120	GLU
20	BT	34	PRO
20	BT	40	VAL
21	BP	66	ILE
21	BP	70	ARG
21	BP	71	LYS
21	BP	139	ARG
22	BZ	18	SER
22	BZ	19	GLY
22	BZ	24	LYS
22	BZ	26	LYS
22	BZ	28	TRP
22	BZ	33	GLN
23	Bc	2	ASP
23	Bc	16	ARG
23	Bc	17	THR
23	Bc	24	THR
24	BW	4	VAL

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
24	BW	52	PHE
24	BW	53	VAL
24	BW	65	LEU
24	BW	97	ARG
25	Bd	11	PRO
25	Bd	14	TYR
25	Bd	54	LYS
26	Bb	64	VAL
27	Be	57	PRO
28	BA	10	ARG
28	BA	100	ALA
29	BR	2	GLY
29	BR	88	LYS
29	BR	93	VAL
29	BR	101	GLU
29	BR	115	PRO
30	BB	49	SER
30	BB	113	LEU
30	BB	148	ASN
30	BB	179	CYS
30	BB	182	LYS
30	BB	206	PRO
31	BV	22	ARG
32	Ba	45	VAL
32	Ba	63	VAL
32	Ba	86	VAL
33	BJ	5	PRO
33	BJ	8	TYR
34	BC	146	ASN
35	BG	20	ASP
35	BG	87	TYR
35	BG	160	ASN
36	BH	17	SER
36	BH	34	ASN
36	BH	104	PRO
36	BH	105	PRO
36	BH	106	LYS
36	BH	118	THR
37	CG	48	VAL
37	CG	225	VAL
37	CG	233	VAL
38	CT	8	ARG

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
38	CT	70	ARG
38	CT	141	VAL
38	CT	151	PRO
40	Cz	150	THR
40	Cz	167	ALA
40	Cz	197	TRP
40	Cz	198	GLN
41	CA	28	ARG
41	CA	67	PHE
41	CA	68	ARG
41	CA	119	HIS
42	CJ	2	SER
42	CJ	3	THR
42	CJ	7	GLN
42	CJ	58	SER
42	CJ	94	LEU
42	CJ	113	ASP
43	CH	140	LYS
43	CH	185	THR
43	CH	186	ILE
44	CV	13	LYS
44	CV	36	ASN
45	CN	122	ASN
45	CN	185	ARG
46	Ca	9	ARG
46	Ca	10	LYS
46	Ca	15	VAL
46	Ca	22	ILE
46	Ca	28	HIS
46	Ca	109	LYS
46	Ca	128	VAL
47	CQ	13	ARG
47	CQ	20	LYS
47	CQ	156	PRO
47	CQ	159	PRO
48	CD	2	SER
48	CD	16	TYR
48	CD	74	ILE
48	CD	116	LEU
48	CD	138	GLU
48	CD	188	LYS
48	CD	191	ASP

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
48	CD	199	ILE
48	CD	204	VAL
48	CD	218	LYS
48	CD	236	MET
48	CD	239	LEU
48	CD	260	GLU
48	CD	262	ALA
48	CD	284	ARG
48	CD	285	LEU
48	CD	289	ASN
48	CD	290	SER
48	CD	291	SER
49	CR	55	GLN
49	CR	56	LYS
49	CR	59	SER
49	CR	188	SER
50	CP	112	THR
50	CP	167	ALA
51	CX	34	LYS
51	CX	46	LYS
51	CX	50	LYS
52	CW	54	THR
52	CW	72	LYS
53	CY	8	THR
53	CY	9	SER
53	CY	10	SER
54	Cr	110	ASN
55	Cc	102	SER
56	Cd	8	ALA
56	Cd	27	ARG
56	Cd	87	ARG
56	Cd	101	VAL
56	Cd	102	THR
56	Cd	119	VAL
57	Ce	13	LYS
57	Ce	131	GLU
58	Cj	4	GLY
58	Cj	41	ALA
58	Cj	51	VAL
58	Cj	84	ALA
58	Cj	90	ALA
59	Cl	4	HIS

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
59	Cl	34	THR
59	Cl	38	ASN
60	Co	48	SER
60	Co	97	LYS
61	CM	6	PHE
61	CM	7	VAL
61	CM	17	TYR
61	CM	23	ARG
61	CM	66	PRO
61	CM	80	VAL
61	CM	89	TRP
61	CM	102	LEU
62	CS	4	PHE
62	CS	6	PHE
62	CS	17	PRO
62	CS	23	HIS
62	CS	35	ASN
62	CS	54	LYS
62	CS	70	ASN
62	CS	71	PRO
62	CS	73	THR
62	CS	87	THR
62	CS	117	ARG
62	CS	138	ARG
62	CS	151	PHE
62	CS	152	PRO
62	CS	161	PRO
63	CU	31	VAL
63	CU	32	GLU
63	CU	34	LYS
63	CU	58	ASN
63	CU	79	ALA
63	CU	100	ASP
63	CU	101	TRP
63	CU	104	VAL
63	CU	105	ILE
63	CU	106	ALA
63	CU	111	ARG
64	Ci	38	LYS
64	Ci	40	VAL
64	Ci	95	SER
65	CK	15	VAL

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
65	CK	32	ILE
65	CK	37	LEU
65	CK	76	PRO
65	CK	95	LYS
65	CK	96	VAL
65	CK	132	GLU
65	CK	133	ILE
66	Cu	62	VAL
66	Cv	62	VAL
68	Ch	117	GLN
68	Ch	119	LYS
69	CF	109	ARG
69	CF	159	ASN
69	CF	205	TRP
70	Cq	63	ARG
70	Cq	208	ASP
70	Cq	212	ASP
71	CB	4	ARG
71	CB	60	VAL
71	CB	61	GLU
71	CB	63	PRO
71	CB	123	CYS
71	CB	129	ALA
71	CB	292	GLY
71	CB	296	HIS
71	CB	335	PRO
71	CB	350	THR
71	CB	351	SER
71	CB	358	ILE
72	CC	3	THR
72	CC	17	ASP
72	CC	21	ASP
72	CC	24	SER
72	CC	62	ALA
72	CC	90	ARG
72	CC	107	ALA
72	CC	108	PRO
72	CC	110	LYS
72	CC	202	ASN
72	CC	305	SER
72	CC	348	GLU
72	CC	349	ALA

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
73	CO	11	ARG
73	CO	71	PRO
73	CO	126	PRO
73	CO	132	LEU
73	CO	135	GLN
73	CO	136	PRO
75	CI	18	PRO
75	CI	39	LYS
75	CI	113	THR
75	CI	115	MET
78	CL	13	PHE
78	CL	21	VAL
78	CL	46	PHE
78	CL	48	ARG
78	CL	62	THR
78	CL	64	LYS
78	CL	66	ASN
78	CL	127	PRO
78	CL	154	MET
78	CL	156	ILE
79	CE	26	TRP
79	CE	28	ILE
79	CE	37	PRO
79	CE	39	ALA
79	CE	40	GLU
79	CE	41	LYS
79	CE	44	ALA
79	CE	51	TYR
79	CE	52	PRO
79	CE	73	SER
79	CE	74	THR
79	CE	159	GLU
79	CE	166	ASP
80	Cf	7	GLN
82	Cb	21	ILE
82	Cb	39	PHE
83	Cg	11	HIS
83	Cg	65	PRO
6	BK	53	GLU
6	BK	61	TRP
7	BM	115	GLY
7	BM	117	GLU

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
8	Bf	13	LYS
8	Bf	16	LYS
8	Bf	19	HIS
9	BX	86	ASN
10	Bg	202	SER
10	Bg	203	GLY
10	Bg	271	GLY
11	BD	36	GLY
11	BD	63	GLY
11	BD	80	LEU
11	BD	81	GLU
11	BD	91	VAL
11	BD	98	ALA
11	BD	113	LEU
11	BD	140	GLY
11	BD	152	PHE
11	BD	195	LYS
12	BE	53	LYS
12	BE	163	ASP
12	BE	217	GLN
13	BF	178	LYS
14	BQ	26	SER
14	BQ	36	LYS
14	BQ	138	ARG
15	BU	13	LYS
15	BU	30	ARG
15	BU	58	LYS
16	BO	75	LEU
16	BO	94	ILE
16	BO	138	SER
17	BS	15	VAL
17	BS	75	ARG
17	BS	91	LYS
17	BS	95	PHE
17	BS	151	LYS
18	BN	149	LEU
19	BL	121	GLY
20	BT	45	PHE
21	BP	116	ILE
22	BZ	11	PRO
22	BZ	34	LYS
22	BZ	89	ALA

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
23	Bc	19	SER
23	Bc	25	GLN
24	BW	23	ARG
24	BW	50	PHE
24	BW	92	ARG
24	BW	96	SER
25	Bd	19	ARG
25	Bd	20	VAL
25	Bd	33	LYS
25	Bd	53	ILE
28	BA	32	LYS
28	BA	71	ALA
29	BR	86	PRO
29	BR	95	GLU
29	BR	112	ALA
30	BB	93	GLY
30	BB	177	SER
30	BB	221	PRO
31	BV	7	GLN
32	Ba	10	ARG
32	Ba	19	LYS
32	Ba	62	TYR
33	BJ	123	SER
33	BJ	136	ILE
33	BJ	152	VAL
34	BC	148	ILE
36	BH	67	TYR
36	BH	77	HIS
36	BH	111	VAL
36	BH	135	GLU
37	CG	121	GLU
38	CT	5	HIS
38	CT	10	ARG
38	CT	159	ASP
40	Cz	73	VAL
40	Cz	98	LEU
40	Cz	134	PRO
41	CA	34	PHE
42	CJ	33	THR
42	CJ	63	ARG
42	CJ	88	VAL
42	CJ	108	ILE

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
42	CJ	124	ILE
42	CJ	127	MET
44	CV	5	GLY
44	CV	9	SER
44	CV	12	ASN
45	CN	2	GLY
45	CN	56	LYS
46	Ca	7	LYS
46	Ca	8	ASN
46	Ca	11	LYS
46	Ca	124	LEU
47	CQ	94	GLU
47	CQ	98	MET
47	CQ	160	HIS
48	CD	3	LEU
48	CD	14	HIS
48	CD	59	LYS
48	CD	73	ASP
48	CD	89	LEU
48	CD	117	ASP
48	CD	119	GLU
48	CD	125	GLU
48	CD	144	ALA
48	CD	183	PHE
48	CD	187	GLU
48	CD	200	TYR
48	CD	203	HIS
48	CD	238	SER
48	CD	248	ARG
48	CD	249	ALA
49	CR	54	PRO
50	CP	2	VAL
50	CP	9	ASN
50	CP	31	GLU
50	CP	38	LYS
50	CP	68	GLY
50	CP	70	THR
50	CP	74	LYS
50	CP	128	ARG
50	CP	166	ILE
50	CP	169	ARG
52	CW	63	LYS

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
54	Cr	90	SER
54	Cr	91	VAL
56	Cd	2	SER
57	Ce	120	VAL
58	Cj	76	SER
58	Cj	79	ARG
58	Cj	87	ARG
59	Cl	39	ALA
60	Co	34	SER
60	Co	37	ALA
61	CM	3	PHE
61	CM	32	ASP
62	CS	52	LYS
63	CU	37	GLU
64	Ci	42	PHE
64	Ci	66	VAL
64	Ci	94	SER
65	CK	94	LYS
65	CK	137	CYS
69	CF	36	LYS
70	Cq	52	SER
71	CB	2	SER
71	CB	124	LYS
71	CB	126	LYS
71	CB	131	THR
71	CB	215	ASP
71	CB	301	GLU
71	CB	303	ASP
71	CB	359	LYS
71	CB	373	ARG
71	CB	374	PHE
72	CC	19	ALA
72	CC	91	ALA
72	CC	151	VAL
72	CC	208	ARG
72	CC	345	THR
74	Cp	6	LYS
74	Cp	18	TYR
75	CI	99	ILE
75	CI	110	ARG
75	CI	118	ALA
78	CL	12	HIS

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
78	CL	49	PRO
78	CL	67	MET
78	CL	70	ARG
78	CL	130	ALA
79	CE	12	LYS
79	CE	54	ASP
79	CE	75	ILE
79	CE	172	ASN
79	CE	205	ARG
79	CE	209	ARG
81	Ck	20	ALA
82	Cb	25	LYS
83	Cg	10	ARG
4	BY	16	LYS
5	BI	161	GLN
6	BK	34	GLN
7	BM	10	GLU
7	BM	44	LYS
7	BM	72	HIS
7	BM	89	ALA
8	Bf	12	PRO
8	Bf	30	TYR
10	Bg	98	SER
10	Bg	150	SER
10	Bg	151	ARG
10	Bg	180	GLN
10	Bg	301	VAL
10	Bg	366	LYS
11	BD	62	LYS
11	BD	78	ASN
11	BD	93	ASN
11	BD	173	ARG
11	BD	211	HIS
12	BE	94	LYS
12	BE	213	ALA
14	BQ	32	ARG
14	BQ	81	THR
14	BQ	92	ALA
14	BQ	118	TYR
14	BQ	146	LYS
16	BO	93	HIS
16	BO	109	PRO

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
16	BO	149	ARG
17	BS	49	ASP
17	BS	140	GLY
18	BN	78	HIS
18	BN	87	ASP
20	BT	11	ASP
22	BZ	101	ILE
23	Bc	8	ALA
24	BW	95	PRO
25	Bd	15	GLY
25	Bd	47	ALA
26	Bb	19	LEU
26	Bb	65	LEU
28	BA	44	LYS
30	BB	55	LYS
32	Ba	46	GLU
32	Ba	82	HIS
36	BH	57	ASN
36	BH	89	SER
36	BH	132	TYR
37	CG	153	VAL
37	CG	232	GLY
38	CT	80	VAL
39	CZ	103	THR
41	CA	11	GLY
41	CA	71	HIS
41	CA	245	ARG
42	CJ	65	GLU
42	CJ	89	LYS
42	CJ	145	ARG
46	Ca	66	ASN
46	Ca	114	PRO
47	CQ	11	ASN
47	CQ	119	GLU
47	CQ	142	PRO
48	CD	23	LYS
48	CD	52	LYS
48	CD	55	PHE
48	CD	132	TYR
48	CD	141	PRO
48	CD	197	LYS
48	CD	212	ALA

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
48	CD	261	PRO
48	CD	287	ALA
48	CD	288	LEU
48	CD	292	ALA
48	CD	293	GLY
50	CP	8	ALA
50	CP	63	TYR
50	CP	79	ASN
50	CP	168	ALA
54	Cr	87	TYR
54	Cr	93	ARG
56	Cd	4	LYS
56	Cd	11	ARG
56	Cd	26	LYS
58	Cj	26	SER
58	Cj	77	ASN
59	Cl	47	THR
60	Co	14	ASN
60	Co	56	PRO
60	Co	76	SER
60	Co	91	PHE
60	Co	94	GLY
60	Co	98	LYS
60	Co	102	THR
61	CM	31	VAL
61	CM	88	SER
61	CM	105	PHE
62	CS	139	ASP
62	CS	162	THR
63	CU	29	LYS
63	CU	97	ASN
63	CU	107	ALA
63	CU	108	ASN
64	Ci	39	ARG
64	Ci	80	LEU
65	CK	134	LEU
68	Ch	99	ASP
68	Ch	113	VAL
69	CF	35	GLU
69	CF	62	LYS
69	CF	167	ASN
70	Cq	73	THR

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
70	Cq	209	LEU
70	Cq	210	THR
70	Cq	211	GLU
71	CB	40	PRO
71	CB	103	ASN
71	CB	137	TYR
71	CB	158	THR
71	CB	262	ARG
71	CB	375	GLN
71	CB	386	ARG
72	CC	4	GLN
72	CC	330	VAL
73	CO	42	ARG
73	CO	69	THR
76	Cn	24	SER
78	CL	22	LYS
78	CL	128	ARG
78	CL	134	LYS
78	CL	142	GLU
78	CL	148	GLN
78	CL	152	ASP
79	CE	157	LYS
4	BY	11	THR
4	BY	48	LYS
6	BK	30	ALA
6	BK	60	SER
6	BK	64	TYR
7	BM	30	GLY
7	BM	94	ILE
7	BM	106	CYS
7	BM	107	SER
8	Bf	14	LYS
8	Bf	21	LYS
8	Bf	34	ASP
8	Bf	72	THR
9	BX	64	ALA
11	BD	30	ALA
11	BD	61	GLU
11	BD	64	ARG
11	BD	71	SER
11	BD	172	VAL
12	BE	149	TYR

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
13	BF	56	ARG
15	BU	25	VAL
16	BO	53	VAL
16	BO	73	ALA
16	BO	142	LYS
17	BS	6	GLY
17	BS	7	GLU
17	BS	79	VAL
17	BS	83	PHE
17	BS	94	ARG
17	BS	139	THR
18	BN	41	ALA
18	BN	58	HIS
19	BL	61	PHE
20	BT	48	LEU
22	BZ	13	SER
22	BZ	96	HIS
23	Bc	3	THR
23	Bc	4	GLN
25	Bd	16	ALA
28	BA	108	THR
29	BR	100	LYS
30	BB	35	PRO
30	BB	79	GLN
30	BB	82	ARG
30	BB	176	ALA
31	BV	59	ARG
32	Ba	36	ILE
32	Ba	64	LEU
32	Ba	65	PRO
34	BC	35	TRP
34	BC	36	VAL
34	BC	106	ASP
34	BC	150	GLN
34	BC	235	PHE
35	BG	69	THR
36	BH	102	VAL
36	BH	141	VAL
36	BH	157	PRO
38	CT	124	GLU
38	CT	125	VAL
41	CA	38	ASN

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
41	CA	115	ASN
41	CA	127	ALA
42	CJ	6	LYS
42	CJ	10	PRO
42	CJ	32	LEU
42	CJ	74	ARG
43	CH	49	GLU
43	CH	126	ASP
44	CV	6	ARG
45	CN	78	GLY
45	CN	148	ILE
46	Ca	4	ARG
46	Ca	108	GLY
46	Ca	117	PRO
46	Ca	134	LYS
47	CQ	78	ASN
47	CQ	139	LEU
47	CQ	148	ALA
47	CQ	157	GLY
47	CQ	158	VAL
48	CD	17	PHE
48	CD	90	GLU
48	CD	118	GLN
48	CD	245	ALA
48	CD	259	LYS
52	CW	66	HIS
53	CY	4	ASN
53	CY	21	ALA
55	Cc	31	TYR
55	Cc	89	TYR
57	Ce	46	LYS
57	Ce	90	ASN
57	Ce	132	ASP
58	Cj	10	LYS
58	Cj	37	CYS
58	Cj	39	TYR
58	Cj	61	THR
60	Co	35	LEU
61	CM	65	VAL
61	CM	67	LYS
62	CS	15	GLY
62	CS	16	LEU

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
62	CS	19	PRO
62	CS	86	ARG
64	Ci	41	HIS
68	Ch	80	ALA
69	CF	206	PRO
70	Cq	12	VAL
70	Cq	74	GLY
71	CB	204	LYS
71	CB	205	GLU
71	CB	270	ALA
72	CC	22	ASN
72	CC	87	GLY
72	CC	154	LEU
72	CC	331	LEU
72	CC	346	LEU
72	CC	389	SER
75	CI	84	ALA
75	CI	85	PHE
75	CI	93	PRO
78	CL	65	TYR
81	Ck	26	LYS
82	Cb	23	LYS
4	BY	45	LEU
7	BM	91	LEU
7	BM	113	ASP
8	Bf	43	ARG
9	BX	9	ALA
10	Bg	111	VAL
17	BS	137	LYS
17	BS	150	LYS
20	BT	53	PRO
21	BP	117	LYS
21	BP	118	PRO
23	Bc	57	SER
24	BW	79	PHE
27	Be	4	VAL
27	Be	48	VAL
28	BA	163	ILE
29	BR	92	GLU
30	BB	38	PHE
30	BB	54	THR
30	BB	58	SER

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
30	BB	62	LYS
30	BB	63	HIS
30	BB	64	ARG
30	BB	210	VAL
30	BB	213	ARG
33	BJ	142	ILE
35	BG	154	ASP
36	BH	113	ARG
41	CA	66	PRO
41	CA	69	TYR
41	CA	140	ASN
42	CJ	9	ASN
47	CQ	149	VAL
48	CD	140	ARG
48	CD	294	ALA
49	CR	185	PRO
50	CP	162	PRO
52	CW	45	ARG
53	CY	49	ILE
54	Cr	72	LEU
54	Cr	108	SER
56	Cd	90	GLU
57	Ce	66	HIS
60	Co	59	HIS
60	Co	80	TYR
61	CM	78	ALA
61	CM	79	ASP
63	CU	55	LYS
65	CK	75	VAL
65	CK	78	ALA
65	CK	91	ARG
68	Ch	45	LEU
68	Ch	90	ARG
69	CF	160	LYS
70	Cq	145	ASN
71	CB	70	LYS
71	CB	111	SER
72	CC	18	MET
72	CC	109	THR
72	CC	343	MET
73	CO	131	VAL
73	CO	152	TRP

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
73	CO	193	LYS
77	Cm	2	ILE
78	CL	60	CYS
79	CE	29	LYS
79	CE	69	THR
7	BM	95	ASP
8	Bf	11	LYS
10	Bg	237	ILE
10	Bg	340	GLN
11	BD	31	GLU
11	BD	84	VAL
11	BD	194	PRO
11	BD	206	ASP
11	BD	212	PRO
14	BQ	47	PRO
20	BT	90	SER
21	BP	63	MET
22	BZ	44	ASP
23	Bc	34	GLN
25	Bd	38	CYS
28	BA	193	ILE
30	BB	37	VAL
30	BB	207	LEU
33	BJ	4	ALA
35	BG	159	VAL
38	CT	43	LYS
41	CA	73	LYS
42	CJ	25	VAL
44	CV	49	LEU
49	CR	91	THR
49	CR	137	VAL
52	CW	71	LYS
54	Cr	67	ASP
57	Ce	121	THR
62	CS	166	LYS
64	Ci	81	GLY
65	CK	87	LYS
70	Cq	136	SER
71	CB	69	LYS
71	CB	242	PRO
71	CB	297	GLU
78	CL	126	PHE

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
79	CE	11	ILE
14	BQ	31	GLY
27	Be	56	GLY
32	Ba	84	VAL
38	CT	123	GLY
49	CR	57	ILE
79	CE	45	ILE
79	CE	117	ILE
83	Cg	2	VAL
13	BF	137	ILE
14	BQ	46	ARG
18	BN	66	VAL
38	CT	84	ILE
38	CT	146	ILE
49	CR	73	GLY
54	Cr	61	GLN
7	BM	103	VAL
10	Bg	127	GLY
11	BD	95	GLY
20	BT	33	LEU
28	BA	122	GLU
32	Ba	75	VAL
40	Cz	55	PRO
42	CJ	60	GLY
64	Ci	43	VAL
68	Ch	97	SER
78	CL	2	VAL
6	BK	83	PRO
15	BU	66	PRO
30	BB	215	VAL
36	BH	156	ASP
38	CT	126	ILE
56	Cd	16	VAL
75	CI	42	GLY
8	Bf	15	ILE
11	BD	202	THR
55	Cc	79	VAL
58	Cj	38	GLY
72	CC	386	ILE
79	CE	36	LEU

### 5.3.2 Protein sidechains [i](#)

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
4	BY	116/116 (100%)	110 (95%)	6 (5%)	23	49
5	BI	56/179 (31%)	53 (95%)	3 (5%)	22	49
6	BK	90/146 (62%)	88 (98%)	2 (2%)	52	71
7	BM	101/142 (71%)	101 (100%)	0	100	100
8	Bf	62/135 (46%)	60 (97%)	2 (3%)	39	61
9	BX	113/113 (100%)	111 (98%)	2 (2%)	59	77
10	Bg	323/323 (100%)	310 (96%)	13 (4%)	31	56
11	BD	175/175 (100%)	170 (97%)	5 (3%)	42	64
12	BE	176/225 (78%)	172 (98%)	4 (2%)	50	70
13	BF	159/159 (100%)	153 (96%)	6 (4%)	33	57
14	BQ	103/120 (86%)	97 (94%)	6 (6%)	20	46
15	BU	113/113 (100%)	107 (95%)	6 (5%)	22	49
16	BO	94/120 (78%)	90 (96%)	4 (4%)	29	54
17	BS	133/133 (100%)	125 (94%)	8 (6%)	19	46
18	BN	106/130 (82%)	101 (95%)	5 (5%)	26	52
19	BL	74/135 (55%)	70 (95%)	4 (5%)	22	49
20	BT	121/121 (100%)	115 (95%)	6 (5%)	24	50
21	BP	77/130 (59%)	71 (92%)	6 (8%)	12	38
22	BZ	87/93 (94%)	84 (97%)	3 (3%)	37	60
23	Bc	52/58 (90%)	48 (92%)	4 (8%)	13	39
24	BW	113/113 (100%)	109 (96%)	4 (4%)	36	60
25	Bd	40/47 (85%)	39 (98%)	1 (2%)	47	68
26	Bb	78/78 (100%)	78 (100%)	0	100	100
27	Be	47/49 (96%)	46 (98%)	1 (2%)	53	72
28	BA	161/204 (79%)	153 (95%)	8 (5%)	24	50
29	BR	105/127 (83%)	103 (98%)	2 (2%)	57	75

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
30	BB	188/226 (83%)	186 (99%)	2 (1%)	73	84
31	BV	63/68 (93%)	59 (94%)	4 (6%)	18	44
32	Ba	80/107 (75%)	78 (98%)	2 (2%)	47	68
33	BJ	160/167 (96%)	157 (98%)	3 (2%)	57	75
34	BC	182/211 (86%)	177 (97%)	5 (3%)	44	65
35	BG	201/210 (96%)	192 (96%)	9 (4%)	27	53
36	BH	164/168 (98%)	157 (96%)	7 (4%)	29	54
37	CG	205/220 (93%)	194 (95%)	11 (5%)	22	49
38	CT	139/141 (99%)	135 (97%)	4 (3%)	42	64
39	CZ	113/113 (100%)	109 (96%)	4 (4%)	36	60
40	Cz	192/192 (100%)	182 (95%)	10 (5%)	23	49
41	CA	195/199 (98%)	184 (94%)	11 (6%)	21	47
42	CJ	149/157 (95%)	139 (93%)	10 (7%)	16	42
43	CH	164/164 (100%)	158 (96%)	6 (4%)	34	58
44	CV	109/109 (100%)	106 (97%)	3 (3%)	43	65
45	CN	167/173 (96%)	161 (96%)	6 (4%)	35	59
46	Ca	110/110 (100%)	101 (92%)	9 (8%)	11	36
47	CQ	138/160 (86%)	132 (96%)	6 (4%)	29	54
48	CD	251/251 (100%)	234 (93%)	17 (7%)	16	42
49	CR	166/183 (91%)	154 (93%)	12 (7%)	14	41
50	CP	144/144 (100%)	139 (96%)	5 (4%)	36	60
51	CX	109/130 (84%)	102 (94%)	7 (6%)	17	44
52	CW	66/133 (50%)	65 (98%)	1 (2%)	65	80
53	CY	115/128 (90%)	110 (96%)	5 (4%)	29	54
54	Cr	64/131 (49%)	61 (95%)	3 (5%)	26	52
55	Cc	98/98 (100%)	94 (96%)	4 (4%)	30	55
56	Cd	103/106 (97%)	98 (95%)	5 (5%)	25	51
57	Ce	122/122 (100%)	116 (95%)	6 (5%)	25	51
58	Cj	77/77 (100%)	74 (96%)	3 (4%)	32	57
59	Cl	48/48 (100%)	47 (98%)	1 (2%)	53	72
60	Co	94/94 (100%)	87 (93%)	7 (7%)	13	40

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
61	CM	116/116 (100%)	110 (95%)	6 (5%)	23	49
62	CS	153/163 (94%)	145 (95%)	8 (5%)	23	49
63	CU	94/106 (89%)	86 (92%)	8 (8%)	10	35
64	Ci	62/92 (67%)	59 (95%)	3 (5%)	25	52
65	CK	105/139 (76%)	98 (93%)	7 (7%)	16	42
66	Cu	46/77 (60%)	45 (98%)	1 (2%)	52	71
66	Cv	46/77 (60%)	46 (100%)	0	100	100
67	Cs	48/82 (58%)	47 (98%)	1 (2%)	53	72
67	Ct	48/82 (58%)	46 (96%)	2 (4%)	30	54
68	Ch	109/109 (100%)	103 (94%)	6 (6%)	21	48
69	CF	206/206 (100%)	199 (97%)	7 (3%)	37	60
70	Cq	222/265 (84%)	216 (97%)	6 (3%)	44	65
71	CB	335/335 (100%)	315 (94%)	20 (6%)	19	46
72	CC	302/329 (92%)	286 (95%)	16 (5%)	22	49
73	CO	173/173 (100%)	160 (92%)	13 (8%)	13	39
74	Cp	73/73 (100%)	72 (99%)	1 (1%)	67	80
75	CI	156/183 (85%)	152 (97%)	4 (3%)	46	67
76	Cn	24/24 (100%)	23 (96%)	1 (4%)	30	54
77	Cm	47/48 (98%)	46 (98%)	1 (2%)	53	72
78	CL	175/175 (100%)	166 (95%)	9 (5%)	24	50
79	CE	185/185 (100%)	169 (91%)	16 (9%)	10	34
80	Cf	96/96 (100%)	93 (97%)	3 (3%)	40	62
81	Ck	63/63 (100%)	58 (92%)	5 (8%)	12	38
82	Cb	51/53 (96%)	51 (100%)	0	100	100
83	Cg	98/107 (92%)	91 (93%)	7 (7%)	14	41
All	All	10084/11382 (89%)	9634 (96%)	450 (4%)	31	53

All (450) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
4	BY	21	ARG
4	BY	37	ARG
4	BY	47	GLU

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
4	BY	95	TYR
4	BY	123	ILE
4	BY	127	LYS
5	BI	151	ASN
5	BI	201	LYS
5	BI	209	LYS
6	BK	54	TYR
6	BK	92	LYS
8	Bf	59	ASN
8	Bf	65	TYR
9	BX	60	GLN
9	BX	141	ARG
10	Bg	15	THR
10	Bg	52	ASN
10	Bg	62	LEU
10	Bg	78	LYS
10	Bg	107	HIS
10	Bg	112	MET
10	Bg	128	LEU
10	Bg	143	ARG
10	Bg	166	VAL
10	Bg	230	VAL
10	Bg	246	HIS
10	Bg	255	VAL
10	Bg	263	ARG
11	BD	40	ARG
11	BD	91	VAL
11	BD	99	ILE
11	BD	124	ARG
11	BD	207	LEU
12	BE	80	LYS
12	BE	149	TYR
12	BE	168	LYS
12	BE	208	ILE
13	BF	10	GLN
13	BF	51	ARG
13	BF	58	ARG
13	BF	113	ILE
13	BF	132	ARG
13	BF	178	LYS
14	BQ	53	LYS
14	BQ	66	PHE

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
14	BQ	129	ARG
14	BQ	135	PHE
14	BQ	143	ARG
14	BQ	149	ARG
15	BU	12	MET
15	BU	30	ARG
15	BU	75	ARG
15	BU	86	TRP
15	BU	87	ASP
15	BU	92	ARG
16	BO	49	ARG
16	BO	65	ARG
16	BO	67	GLU
16	BO	87	LEU
17	BS	25	LYS
17	BS	36	VAL
17	BS	55	ARG
17	BS	78	LYS
17	BS	79	VAL
17	BS	108	ARG
17	BS	135	HIS
17	BS	136	THR
18	BN	70	LYS
18	BN	83	GLU
18	BN	108	ASP
18	BN	117	LEU
18	BN	124	ARG
19	BL	86	ILE
19	BL	96	LYS
19	BL	100	ARG
19	BL	116	PHE
20	BT	34	PRO
20	BT	46	LYS
20	BT	51	TYR
20	BT	91	ARG
20	BT	93	PRO
20	BT	106	ILE
21	BP	59	LYS
21	BP	60	ARG
21	BP	99	ILE
21	BP	116	ILE
21	BP	124	TYR

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
21	BP	139	ARG
22	BZ	39	ASN
22	BZ	68	GLU
22	BZ	104	ARG
23	Bc	1	MET
23	Bc	16	ARG
23	Bc	41	ASN
23	Bc	47	ARG
24	BW	22	LYS
24	BW	51	GLU
24	BW	69	LEU
24	BW	113	HIS
25	Bd	55	TYR
27	Be	26	LYS
28	BA	37	GLN
28	BA	40	ARG
28	BA	52	ILE
28	BA	68	VAL
28	BA	76	GLN
28	BA	108	THR
28	BA	126	LEU
28	BA	159	ARG
29	BR	85	VAL
29	BR	105	MET
30	BB	105	PHE
30	BB	231	VAL
31	BV	20	THR
31	BV	41	GLU
31	BV	63	ASP
31	BV	72	TRP
32	Ba	45	VAL
32	Ba	70	LYS
33	BJ	18	ARG
33	BJ	23	LYS
33	BJ	70	ARG
34	BC	90	THR
34	BC	113	LEU
34	BC	148	ILE
34	BC	196	VAL
34	BC	216	ASP
35	BG	4	ASN
35	BG	7	ASN

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
35	BG	32	ILE
35	BG	44	GLU
35	BG	74	ARG
35	BG	87	TYR
35	BG	130	GLU
35	BG	144	ARG
35	BG	228	ARG
36	BH	32	ASN
36	BH	36	GLU
36	BH	117	ARG
36	BH	130	VAL
36	BH	158	LYS
36	BH	167	LEU
36	BH	182	VAL
37	CG	61	ARG
37	CG	68	LYS
37	CG	82	LYS
37	CG	126	ILE
37	CG	132	LEU
37	CG	177	LYS
37	CG	189	LYS
37	CG	193	VAL
37	CG	214	ILE
37	CG	217	ASN
37	CG	221	LYS
38	CT	27	LEU
38	CT	30	TYR
38	CT	60	ARG
38	CT	63	ARG
39	CZ	29	PHE
39	CZ	34	ARG
39	CZ	84	ARG
39	CZ	136	PHE
40	Cz	9	VAL
40	Cz	33	LEU
40	Cz	46	LYS
40	Cz	61	LYS
40	Cz	102	LEU
40	Cz	129	LYS
40	Cz	140	GLN
40	Cz	182	ILE
40	Cz	206	LYS

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
40	Cz	211	LYS
41	CA	1	MET
41	CA	5	ILE
41	CA	30	ARG
41	CA	37	ARG
41	CA	69	TYR
41	CA	74	GLU
41	CA	158	VAL
41	CA	177	LYS
41	CA	193	ARG
41	CA	227	ARG
41	CA	247	ARG
42	CJ	2	SER
42	CJ	31	ARG
42	CJ	34	ARG
42	CJ	51	LYS
42	CJ	70	TYR
42	CJ	82	LEU
42	CJ	89	LYS
42	CJ	93	LEU
42	CJ	94	LEU
42	CJ	142	ARG
43	CH	39	LYS
43	CH	41	LEU
43	CH	70	ARG
43	CH	141	ASP
43	CH	143	LEU
43	CH	185	THR
44	CV	39	ILE
44	CV	62	MET
44	CV	92	ASP
45	CN	97	ASN
45	CN	116	LEU
45	CN	123	GLU
45	CN	160	GLU
45	CN	175	ARG
45	CN	180	THR
46	Ca	21	ARG
46	Ca	26	ARG
46	Ca	41	HIS
46	Ca	47	LYS
46	Ca	62	HIS

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
46	Ca	111	MET
46	Ca	121	LYS
46	Ca	123	LYS
46	Ca	142	LEU
47	CQ	4	ASP
47	CQ	56	LYS
47	CQ	77	LYS
47	CQ	91	ARG
47	CQ	139	LEU
47	CQ	156	PRO
48	CD	19	ARG
48	CD	23	LYS
48	CD	52	LYS
48	CD	61	ILE
48	CD	142	PHE
48	CD	158	ARG
48	CD	190	LEU
48	CD	200	TYR
48	CD	203	HIS
48	CD	208	MET
48	CD	219	PHE
48	CD	236	MET
48	CD	241	LYS
48	CD	242	LYS
48	CD	261	PRO
48	CD	281	LEU
48	CD	285	LEU
49	CR	1	MET
49	CR	6	LEU
49	CR	9	ARG
49	CR	56	LYS
49	CR	58	HIS
49	CR	84	THR
49	CR	86	GLU
49	CR	91	THR
49	CR	131	MET
49	CR	135	LYS
49	CR	151	ARG
49	CR	153	LYS
50	CP	4	TYR
50	CP	16	LYS
50	CP	22	LEU

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
50	CP	61	ARG
50	CP	81	GLN
51	CX	46	LYS
51	CX	47	THR
51	CX	48	LEU
51	CX	66	LYS
51	CX	80	GLU
51	CX	98	LEU
51	CX	149	ILE
52	CW	45	ARG
53	CY	1	MET
53	CY	27	ARG
53	CY	86	ARG
53	CY	114	ARG
53	CY	129	LYS
54	Cr	87	TYR
54	Cr	92	MET
54	Cr	115	ASP
55	Cc	14	ILE
55	Cc	57	LYS
55	Cc	91	VAL
55	Cc	104	ILE
56	Cd	45	ARG
56	Cd	63	LEU
56	Cd	85	ARG
56	Cd	102	THR
56	Cd	105	GLU
57	Ce	24	ASP
57	Ce	44	ARG
57	Ce	91	ARG
57	Ce	120	VAL
57	Ce	121	THR
57	Ce	126	ARG
58	Cj	1	MET
58	Cj	14	LYS
58	Cj	30	GLN
59	Cl	23	ILE
60	Co	2	VAL
60	Co	16	GLU
60	Co	53	GLN
60	Co	61	LYS
60	Co	78	LYS

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
60	Co	83	HIS
60	Co	102	THR
61	CM	5	ARG
61	CM	6	PHE
61	CM	8	GLU
61	CM	42	ASP
61	CM	63	LYS
61	CM	102	LEU
62	CS	38	ARG
62	CS	55	LYS
62	CS	69	LYS
62	CS	71	PRO
62	CS	75	LYS
62	CS	115	ARG
62	CS	119	ARG
62	CS	161	PRO
63	CU	41	LEU
63	CU	48	ARG
63	CU	55	LYS
63	CU	85	LEU
63	CU	95	LYS
63	CU	101	TRP
63	CU	102	LEU
63	CU	114	TYR
64	Ci	88	LYS
64	Ci	96	VAL
64	Ci	99	LYS
65	CK	16	ARG
65	CK	32	ILE
65	CK	41	LYS
65	CK	42	ILE
65	CK	55	LYS
65	CK	58	ARG
65	CK	76	PRO
66	Cu	10	LEU
67	Cs	8	LEU
67	Ct	1	MET
67	Ct	8	LEU
68	Ch	7	LYS
68	Ch	16	LYS
68	Ch	35	ILE
68	Ch	37	LYS

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
68	Ch	75	LYS
68	Ch	90	ARG
69	CF	68	LYS
69	CF	69	ARG
69	CF	72	ARG
69	CF	115	ASN
69	CF	143	LEU
69	CF	144	LYS
69	CF	190	ILE
70	Cq	59	ASN
70	Cq	92	ILE
70	Cq	98	LEU
70	Cq	155	VAL
70	Cq	209	LEU
70	Cq	210	THR
71	CB	4	ARG
71	CB	20	LYS
71	CB	30	LYS
71	CB	39	LYS
71	CB	55	HIS
71	CB	60	VAL
71	CB	94	LYS
71	CB	101	THR
71	CB	118	PHE
71	CB	123	CYS
71	CB	162	VAL
71	CB	169	ARG
71	CB	182	MET
71	CB	201	PHE
71	CB	244	LYS
71	CB	259	HIS
71	CB	306	GLU
71	CB	327	MET
71	CB	347	LEU
71	CB	348	LYS
72	CC	8	LEU
72	CC	20	THR
72	CC	53	ARG
72	CC	60	ARG
72	CC	108	PRO
72	CC	109	THR
72	CC	154	LEU

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
72	CC	200	MET
72	CC	215	TYR
72	CC	233	ASP
72	CC	313	GLU
72	CC	319	LYS
72	CC	345	THR
72	CC	346	LEU
72	CC	352	ILE
72	CC	357	GLU
73	CO	11	ARG
73	CO	37	ARG
73	CO	44	GLU
73	CO	54	ARG
73	CO	57	MET
73	CO	66	ARG
73	CO	76	ILE
73	CO	126	PRO
73	CO	134	LEU
73	CO	152	TRP
73	CO	181	LYS
73	CO	185	LYS
73	CO	193	LYS
74	Cp	73	THR
75	CI	57	LYS
75	CI	65	LEU
75	CI	141	LYS
75	CI	179	GLU
76	Cn	25	LYS
77	Cm	23	CYS
78	CL	10	ASN
78	CL	12	HIS
78	CL	54	LEU
78	CL	63	LEU
78	CL	102	LYS
78	CL	120	LYS
78	CL	148	GLN
78	CL	183	ARG
78	CL	204	LYS
79	CE	22	ARG
79	CE	36	LEU
79	CE	38	LYS
79	CE	50	PHE

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
79	CE	51	TYR
79	CE	52	PRO
79	CE	75	ILE
79	CE	92	ARG
79	CE	118	ARG
79	CE	149	ARG
79	CE	152	LYS
79	CE	161	GLU
79	CE	180	ASP
79	CE	184	ILE
79	CE	209	ARG
79	CE	217	MET
80	Cf	4	ARG
80	Cf	5	GLN
80	Cf	9	VAL
81	Ck	1	MET
81	Ck	19	ASP
81	Ck	33	LYS
81	Ck	56	LEU
81	Ck	58	GLN
83	Cg	9	LYS
83	Cg	32	TYR
83	Cg	50	LYS
83	Cg	51	ILE
83	Cg	63	LYS
83	Cg	66	ARG
83	Cg	72	ARG

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (129) such sidechains are listed below:

<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
4	BY	39	ASN
5	BI	151	ASN
5	BI	152	HIS
5	BI	189	GLN
6	BK	32	HIS
9	BX	20	GLN
9	BX	60	GLN
9	BX	109	HIS
10	Bg	63	GLN
10	Bg	217	ASN
10	Bg	219	ASN

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
10	Bg	282	HIS
10	Bg	343	HIS
10	Bg	367	ASN
11	BD	162	GLN
13	BF	91	HIS
15	BU	84	ASN
17	BS	42	ASN
17	BS	72	HIS
17	BS	120	HIS
17	BS	125	HIS
18	BN	58	HIS
18	BN	101	HIS
19	BL	77	HIS
19	BL	105	HIS
19	BL	123	HIS
20	BT	15	HIS
21	BP	112	ASN
22	BZ	99	GLN
24	BW	42	GLN
24	BW	44	HIS
24	BW	113	HIS
25	Bd	28	HIS
26	Bb	51	HIS
29	BR	56	HIS
30	BB	99	ASN
30	BB	232	HIS
31	BV	38	HIS
31	BV	73	GLN
32	Ba	7	ASN
32	Ba	11	ASN
32	Ba	73	HIS
32	Ba	80	HIS
32	Ba	82	HIS
33	BJ	125	HIS
33	BJ	178	ASN
35	BG	34	GLN
35	BG	59	GLN
35	BG	160	ASN
37	CG	57	GLN
37	CG	134	HIS
38	CT	54	HIS
38	CT	58	HIS

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
38	CT	90	HIS
40	Cz	75	GLN
41	CA	65	HIS
41	CA	139	HIS
41	CA	187	HIS
41	CA	209	HIS
41	CA	211	HIS
41	CA	216	HIS
41	CA	218	HIS
41	CA	221	HIS
41	CA	233	GLN
42	CJ	17	GLN
42	CJ	153	HIS
43	CH	40	HIS
45	CN	87	GLN
45	CN	95	GLN
45	CN	97	ASN
45	CN	182	HIS
46	Ca	19	HIS
46	Ca	69	HIS
48	CD	14	HIS
48	CD	43	GLN
48	CD	157	ASN
49	CR	58	HIS
49	CR	121	HIS
49	CR	143	HIS
50	CP	9	ASN
50	CP	25	HIS
50	CP	54	HIS
50	CP	56	GLN
50	CP	117	HIS
50	CP	121	ASN
50	CP	146	HIS
51	CX	43	HIS
51	CX	69	GLN
51	CX	89	ASN
51	CX	121	ASN
52	CW	61	HIS
53	CY	99	HIS
57	Ce	22	HIS
58	Cj	28	HIS
58	Cj	94	ASN

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Mol	Chain	Res	Type
59	Cl	20	ASN
62	CS	132	HIS
64	Ci	83	HIS
68	Ch	46	ASN
68	Ch	66	GLN
68	Ch	117	GLN
70	Cq	40	GLN
70	Cq	59	ASN
71	CB	121	ASN
71	CB	165	HIS
71	CB	259	HIS
71	CB	276	HIS
71	CB	282	ASN
72	CC	64	HIS
72	CC	120	ASN
72	CC	146	HIS
72	CC	250	HIS
73	CO	18	HIS
73	CO	138	HIS
73	CO	145	GLN
75	CI	14	ASN
78	CL	99	HIS
79	CE	18	HIS
79	CE	21	HIS
79	CE	67	HIS
79	CE	100	GLN
80	Cf	26	ASN
80	Cf	27	GLN
82	Cb	6	ASN
82	Cb	10	HIS
82	Cb	17	HIS
82	Cb	42	ASN
82	Cb	49	HIS
83	Cg	11	HIS

### 5.3.3 RNA [i](#)

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	Ad	1760/1810 (97%)	458 (26%)	0
2	Ae	74/75 (98%)	19 (25%)	0
3	Af	10/11 (90%)	2 (20%)	0

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Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
84	Aa	3389/3391 (99%)	748 (22%)	0
85	Ac	159/160 (99%)	35 (22%)	0
86	Ab	119/120 (99%)	23 (19%)	0
All	All	5511/5567 (98%)	1285 (23%)	0

All (1285) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	Ad	4	C
1	Ad	8	U
1	Ad	16	G
1	Ad	25	C
1	Ad	26	A
1	Ad	27	U
1	Ad	34	G
1	Ad	46	A
1	Ad	47	A
1	Ad	50	C
1	Ad	55	A
1	Ad	56	U
1	Ad	57	G
1	Ad	58	U
1	Ad	59	G
1	Ad	60	C
1	Ad	65	A
1	Ad	68	A
1	Ad	72	A
1	Ad	73	A
1	Ad	75	U
1	Ad	76	U
1	Ad	77	G
1	Ad	78	A
1	Ad	79	A
1	Ad	80	C
1	Ad	81	U
1	Ad	103	U
1	Ad	105	A
1	Ad	112	U
1	Ad	115	A
1	Ad	127	G
1	Ad	128	G
1	Ad	132	G

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
1	Ad	133	U
1	Ad	134	G
1	Ad	135	C
1	Ad	136	U
1	Ad	137	A
1	Ad	138	C
1	Ad	139	U
1	Ad	140	C
1	Ad	142	G
1	Ad	143	A
1	Ad	144	U
1	Ad	151	A
1	Ad	157	U
1	Ad	158	C
1	Ad	164	C
1	Ad	175	A
1	Ad	176	A
1	Ad	177	C
1	Ad	179	A
1	Ad	183	C
1	Ad	184	C
1	Ad	185	G
1	Ad	186	A
1	Ad	187	C
1	Ad	189	U
1	Ad	190	C
1	Ad	191	U
1	Ad	192	G
1	Ad	193	G
1	Ad	194	G
1	Ad	195	A
1	Ad	198	G
1	Ad	203	A
1	Ad	209	U
1	Ad	212	A
1	Ad	215	A
1	Ad	216	A
1	Ad	220	C
1	Ad	222	G
1	Ad	223	A
1	Ad	224	C
1	Ad	225	G

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
1	Ad	226	C
1	Ad	229	G
1	Ad	230	C
1	Ad	231	U
1	Ad	235	C
1	Ad	236	U
1	Ad	237	C
1	Ad	238	G
1	Ad	239	C
1	Ad	240	U
1	Ad	241	G
1	Ad	242	A
1	Ad	243	U
1	Ad	244	C
1	Ad	245	C
1	Ad	251	U
1	Ad	252	U
1	Ad	253	C
1	Ad	263	C
1	Ad	264	G
1	Ad	265	A
1	Ad	268	G
1	Ad	269	A
1	Ad	270	U
1	Ad	271	C
1	Ad	272	G
1	Ad	277	G
1	Ad	278	C
1	Ad	279	C
1	Ad	282	C
1	Ad	283	G
1	Ad	284	U
1	Ad	285	G
1	Ad	292	A
1	Ad	303	A
1	Ad	318	C
1	Ad	320	A
1	Ad	324	U
1	Ad	337	A
1	Ad	341	G
1	Ad	342	C
1	Ad	345	A

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
1	Ad	352	U
1	Ad	356	G
1	Ad	364	A
1	Ad	365	C
1	Ad	373	U
1	Ad	384	U
1	Ad	403	A
1	Ad	405	A
1	Ad	406	C
1	Ad	408	G
1	Ad	415	C
1	Ad	420	A
1	Ad	421	A
1	Ad	422	G
1	Ad	428	C
1	Ad	429	A
1	Ad	430	G
1	Ad	432	A
1	Ad	438	G
1	Ad	443	U
1	Ad	448	C
1	Ad	450	A
1	Ad	452	C
1	Ad	458	A
1	Ad	474	A
1	Ad	479	A
1	Ad	481	A
1	Ad	488	C
1	Ad	489	C
1	Ad	490	G
1	Ad	491	G
1	Ad	492	G
1	Ad	498	U
1	Ad	500	G
1	Ad	501	U
1	Ad	502	G
1	Ad	503	U
1	Ad	506	G
1	Ad	507	G
1	Ad	508	U
1	Ad	509	A
1	Ad	510	A

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
1	Ad	512	U
1	Ad	514	G
1	Ad	515	A
1	Ad	517	U
1	Ad	519	A
1	Ad	520	G
1	Ad	523	C
1	Ad	529	A
1	Ad	531	A
1	Ad	535	C
1	Ad	536	U
1	Ad	545	A
1	Ad	547	C
1	Ad	548	C
1	Ad	549	A
1	Ad	552	G
1	Ad	560	A
1	Ad	561	G
1	Ad	562	U
1	Ad	569	C
1	Ad	572	G
1	Ad	574	A
1	Ad	579	C
1	Ad	584	A
1	Ad	589	A
1	Ad	598	A
1	Ad	599	G
1	Ad	601	G
1	Ad	610	A
1	Ad	611	G
1	Ad	613	U
1	Ad	615	U
1	Ad	623	A
1	Ad	626	A
1	Ad	628	G
1	Ad	634	A
1	Ad	642	C
1	Ad	643	U
1	Ad	644	U
1	Ad	705	A
1	Ad	708	G
1	Ad	722	A

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
1	Ad	723	A
1	Ad	732	G
1	Ad	733	U
1	Ad	744	G
1	Ad	745	C
1	Ad	749	G
1	Ad	760	G
1	Ad	761	A
1	Ad	762	A
1	Ad	771	G
1	Ad	772	C
1	Ad	780	A
1	Ad	781	A
1	Ad	784	C
1	Ad	789	C
1	Ad	790	U
1	Ad	791	C
1	Ad	793	G
1	Ad	795	A
1	Ad	800	U
1	Ad	801	U
1	Ad	812	A
1	Ad	816	U
1	Ad	817	C
1	Ad	818	A
1	Ad	821	G
1	Ad	822	G
1	Ad	824	U
1	Ad	825	U
1	Ad	826	C
1	Ad	828	G
1	Ad	829	G
1	Ad	834	A
1	Ad	835	U
1	Ad	836	U
1	Ad	838	U
1	Ad	839	G
1	Ad	842	G
1	Ad	843	G
1	Ad	845	C
1	Ad	851	G
1	Ad	854	C

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
1	Ad	857	A
1	Ad	859	U
1	Ad	867	A
1	Ad	868	A
1	Ad	878	U
1	Ad	881	G
1	Ad	903	A
1	Ad	917	U
1	Ad	918	G
1	Ad	919	G
1	Ad	926	G
1	Ad	933	G
1	Ad	934	A
1	Ad	935	A
1	Ad	937	A
1	Ad	938	A
1	Ad	940	U
1	Ad	947	G
1	Ad	949	A
1	Ad	956	A
1	Ad	964	U
1	Ad	965	U
1	Ad	966	U
1	Ad	971	A
1	Ad	973	U
1	Ad	987	U
1	Ad	997	A
1	Ad	998	A
1	Ad	1000	A
1	Ad	1002	G
1	Ad	1009	U
1	Ad	1010	A
1	Ad	1025	A
1	Ad	1026	C
1	Ad	1031	A
1	Ad	1033	C
1	Ad	1044	A
1	Ad	1045	G
1	Ad	1057	U
1	Ad	1058	G
1	Ad	1064	U
1	Ad	1077	C

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
1	Ad	1079	G
1	Ad	1084	U
1	Ad	1087	U
1	Ad	1089	A
1	Ad	1091	A
1	Ad	1096	A
1	Ad	1097	A
1	Ad	1101	C
1	Ad	1103	U
1	Ad	1105	G
1	Ad	1109	U
1	Ad	1114	G
1	Ad	1116	G
1	Ad	1128	C
1	Ad	1143	A
1	Ad	1144	A
1	Ad	1151	G
1	Ad	1154	G
1	Ad	1156	A
1	Ad	1157	A
1	Ad	1160	G
1	Ad	1162	A
1	Ad	1163	C
1	Ad	1165	A
1	Ad	1169	G
1	Ad	1172	G
1	Ad	1184	C
1	Ad	1189	U
1	Ad	1192	G
1	Ad	1195	U
1	Ad	1197	A
1	Ad	1198	A
1	Ad	1200	A
1	Ad	1201	C
1	Ad	1203	G
1	Ad	1204	G
1	Ad	1205	G
1	Ad	1206	A
1	Ad	1211	U
1	Ad	1221	A
1	Ad	1222	G
1	Ad	1232	G

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
1	Ad	1233	G
1	Ad	1247	G
1	Ad	1248	A
1	Ad	1249	G
1	Ad	1254	U
1	Ad	1255	U
1	Ad	1258	U
1	Ad	1260	A
1	Ad	1261	U
1	Ad	1262	U
1	Ad	1264	U
1	Ad	1273	U
1	Ad	1292	G
1	Ad	1305	U
1	Ad	1318	U
1	Ad	1319	U
1	Ad	1325	A
1	Ad	1326	A
1	Ad	1344	U
1	Ad	1345	G
1	Ad	1348	A
1	Ad	1349	A
1	Ad	1354	C
1	Ad	1358	G
1	Ad	1359	C
1	Ad	1366	A
1	Ad	1369	C
1	Ad	1373	C
1	Ad	1376	A
1	Ad	1377	G
1	Ad	1381	G
1	Ad	1382	C
1	Ad	1388	A
1	Ad	1394	A
1	Ad	1395	C
1	Ad	1396	U
1	Ad	1404	U
1	Ad	1405	U
1	Ad	1408	G
1	Ad	1409	G
1	Ad	1418	G
1	Ad	1419	U

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
1	Ad	1421	U
1	Ad	1433	A
1	Ad	1434	G
1	Ad	1437	C
1	Ad	1452	A
1	Ad	1454	G
1	Ad	1463	C
1	Ad	1464	G
1	Ad	1465	C
1	Ad	1466	A
1	Ad	1468	G
1	Ad	1477	A
1	Ad	1479	U
1	Ad	1480	G
1	Ad	1481	A
1	Ad	1484	U
1	Ad	1488	C
1	Ad	1494	G
1	Ad	1496	A
1	Ad	1497	U
1	Ad	1499	U
1	Ad	1500	A
1	Ad	1501	G
1	Ad	1502	C
1	Ad	1507	G
1	Ad	1508	C
1	Ad	1514	G
1	Ad	1522	U
1	Ad	1524	A
1	Ad	1526	C
1	Ad	1529	G
1	Ad	1531	G
1	Ad	1542	G
1	Ad	1543	U
1	Ad	1544	G
1	Ad	1545	A
1	Ad	1546	U
1	Ad	1547	G
1	Ad	1548	G
1	Ad	1565	U
1	Ad	1567	G
1	Ad	1577	A

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
1	Ad	1581	A
1	Ad	1582	G
1	Ad	1591	A
1	Ad	1592	G
1	Ad	1598	G
1	Ad	1609	G
1	Ad	1624	G
1	Ad	1627	C
1	Ad	1632	C
1	Ad	1639	A
1	Ad	1642	C
1	Ad	1643	A
1	Ad	1652	C
1	Ad	1664	U
1	Ad	1665	U
1	Ad	1666	G
1	Ad	1691	C
1	Ad	1692	G
1	Ad	1694	G
1	Ad	1698	A
1	Ad	1699	C
1	Ad	1708	U
1	Ad	1725	C
1	Ad	1726	G
1	Ad	1728	G
1	Ad	1737	A
1	Ad	1739	U
1	Ad	1764	G
1	Ad	1765	A
1	Ad	1766	A
1	Ad	1767	G
1	Ad	1769	C
1	Ad	1770	G
1	Ad	1771	U
1	Ad	1772	A
1	Ad	1776	A
1	Ad	1779	U
1	Ad	1780	U
1	Ad	1783	C
1	Ad	1793	C
1	Ad	1799	G
1	Ad	1802	G

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
1	Ad	1804	A
1	Ad	1806	C
1	Ad	1807	A
1	Ad	1809	U
2	Ae	8	U
2	Ae	17	G
2	Ae	19	U
2	Ae	20	C
2	Ae	21	A
2	Ae	22	G
2	Ae	33	U
2	Ae	37	G
2	Ae	38	C
2	Ae	41	G
2	Ae	42	C
2	Ae	45	G
2	Ae	47	U
2	Ae	51	G
2	Ae	60	C
2	Ae	68	C
2	Ae	72	G
2	Ae	74	C
2	Ae	75	A
3	Af	13	A
3	Af	14	A
84	Aa	2	C
84	Aa	3	G
84	Aa	6	A
84	Aa	12	G
84	Aa	13	G
84	Aa	15	C
84	Aa	25	U
84	Aa	39	A
84	Aa	41	C
84	Aa	48	A
84	Aa	58	G
84	Aa	59	A
84	Aa	64	A
84	Aa	65	A
84	Aa	73	A
84	Aa	74	G
84	Aa	75	G

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	84	A
84	Aa	85	G
84	Aa	91	G
84	Aa	92	C
84	Aa	98	A
84	Aa	108	A
84	Aa	109	G
84	Aa	112	C
84	Aa	115	C
84	Aa	116	U
84	Aa	121	A
84	Aa	134	U
84	Aa	135	G
84	Aa	153	U
84	Aa	155	G
84	Aa	156	A
84	Aa	159	G
84	Aa	164	C
84	Aa	167	C
84	Aa	168	A
84	Aa	171	G
84	Aa	180	G
84	Aa	188	U
84	Aa	189	C
84	Aa	190	C
84	Aa	198	A
84	Aa	208	G
84	Aa	212	G
84	Aa	216	G
84	Aa	217	A
84	Aa	232	C
84	Aa	233	C
84	Aa	236	A
84	Aa	238	C
84	Aa	239	C
84	Aa	241	G
84	Aa	242	U
84	Aa	243	C
84	Aa	247	C
84	Aa	248	C
84	Aa	249	A
84	Aa	250	C

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	251	G
84	Aa	263	A
84	Aa	267	G
84	Aa	281	G
84	Aa	284	U
84	Aa	293	A
84	Aa	296	C
84	Aa	305	G
84	Aa	321	A
84	Aa	327	A
84	Aa	336	A
84	Aa	337	C
84	Aa	347	A
84	Aa	348	C
84	Aa	349	A
84	Aa	368	U
84	Aa	370	A
84	Aa	371	A
84	Aa	374	G
84	Aa	393	A
84	Aa	395	A
84	Aa	396	G
84	Aa	397	A
84	Aa	399	U
84	Aa	400	G
84	Aa	401	C
84	Aa	404	G
84	Aa	419	G
84	Aa	421	A
84	Aa	422	G
84	Aa	424	G
84	Aa	432	G
84	Aa	435	G
84	Aa	438	G
84	Aa	440	U
84	Aa	441	G
84	Aa	464	G
84	Aa	465	C
84	Aa	466	U
84	Aa	467	C
84	Aa	469	U
84	Aa	479	C

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	482	C
84	Aa	488	U
84	Aa	489	C
84	Aa	492	G
84	Aa	493	G
84	Aa	499	A
84	Aa	500	C
84	Aa	507	C
84	Aa	521	G
84	Aa	522	C
84	Aa	523	C
84	Aa	524	A
84	Aa	543	C
84	Aa	544	C
84	Aa	549	G
84	Aa	550	C
84	Aa	555	G
84	Aa	564	A
84	Aa	571	G
84	Aa	572	U
84	Aa	573	A
84	Aa	574	C
84	Aa	575	C
84	Aa	581	G
84	Aa	585	A
84	Aa	588	G
84	Aa	598	U
84	Aa	601	G
84	Aa	612	U
84	Aa	613	G
84	Aa	621	C
84	Aa	623	G
84	Aa	639	A
84	Aa	640	C
84	Aa	642	C
84	Aa	651	A
84	Aa	652	C
84	Aa	653	A
84	Aa	660	A
84	Aa	664	A
84	Aa	681	A
84	Aa	685	G

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	697	A
84	Aa	703	G
84	Aa	709	G
84	Aa	712	A
84	Aa	716	A
84	Aa	718	C
84	Aa	719	U
84	Aa	720	G
84	Aa	722	C
84	Aa	723	G
84	Aa	724	A
84	Aa	729	G
84	Aa	736	U
84	Aa	746	C
84	Aa	747	A
84	Aa	761	C
84	Aa	767	U
84	Aa	768	U
84	Aa	769	C
84	Aa	770	U
84	Aa	779	U
84	Aa	784	G
84	Aa	787	G
84	Aa	788	G
84	Aa	804	A
84	Aa	809	A
84	Aa	810	A
84	Aa	820	A
84	Aa	840	A
84	Aa	852	C
84	Aa	864	C
84	Aa	877	U
84	Aa	882	U
84	Aa	886	A
84	Aa	899	A
84	Aa	900	C
84	Aa	910	G
84	Aa	911	G
84	Aa	917	A
84	Aa	919	G
84	Aa	920	A
84	Aa	923	A

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	924	A
84	Aa	926	C
84	Aa	928	A
84	Aa	937	G
84	Aa	940	G
84	Aa	947	C
84	Aa	950	U
84	Aa	962	C
84	Aa	963	U
84	Aa	965	A
84	Aa	977	G
84	Aa	982	U
84	Aa	983	U
84	Aa	984	A
84	Aa	985	C
84	Aa	986	G
84	Aa	997	G
84	Aa	998	G
84	Aa	1005	C
84	Aa	1006	A
84	Aa	1007	A
84	Aa	1010	A
84	Aa	1014	G
84	Aa	1018	C
84	Aa	1019	A
84	Aa	1020	U
84	Aa	1022	G
84	Aa	1024	G
84	Aa	1025	G
84	Aa	1028	G
84	Aa	1033	G
84	Aa	1036	C
84	Aa	1040	A
84	Aa	1041	C
84	Aa	1051	A
84	Aa	1053	C
84	Aa	1056	U
84	Aa	1061	A
84	Aa	1068	A
84	Aa	1069	U
84	Aa	1075	G
84	Aa	1076	G

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	1085	G
84	Aa	1086	U
84	Aa	1087	G
84	Aa	1097	A
84	Aa	1098	U
84	Aa	1099	G
84	Aa	1101	A
84	Aa	1107	G
84	Aa	1120	G
84	Aa	1134	G
84	Aa	1146	A
84	Aa	1147	U
84	Aa	1156	A
84	Aa	1162	A
84	Aa	1183	C
84	Aa	1184	U
84	Aa	1185	G
84	Aa	1188	C
84	Aa	1194	C
84	Aa	1196	U
84	Aa	1205	C
84	Aa	1206	A
84	Aa	1213	G
84	Aa	1217	G
84	Aa	1220	G
84	Aa	1222	U
84	Aa	1226	G
84	Aa	1229	A
84	Aa	1231	C
84	Aa	1236	C
84	Aa	1237	G
84	Aa	1240	G
84	Aa	1241	G
84	Aa	1242	U
84	Aa	1243	C
84	Aa	1245	U
84	Aa	1246	G
84	Aa	1247	G
84	Aa	1248	A
84	Aa	1249	A
84	Aa	1250	G
84	Aa	1252	C

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	1253	G
84	Aa	1255	A
84	Aa	1257	U
84	Aa	1262	U
84	Aa	1264	A
84	Aa	1266	G
84	Aa	1267	A
84	Aa	1268	G
84	Aa	1269	U
84	Aa	1270	G
84	Aa	1271	U
84	Aa	1273	U
84	Aa	1274	A
84	Aa	1275	A
84	Aa	1276	C
84	Aa	1281	C
84	Aa	1282	A
84	Aa	1283	C
84	Aa	1285	U
84	Aa	1289	G
84	Aa	1291	A
84	Aa	1296	C
84	Aa	1309	U
84	Aa	1311	G
84	Aa	1312	A
84	Aa	1313	U
84	Aa	1317	G
84	Aa	1329	G
84	Aa	1334	A
84	Aa	1349	G
84	Aa	1352	G
84	Aa	1355	U
84	Aa	1356	G
84	Aa	1357	C
84	Aa	1360	U
84	Aa	1361	G
84	Aa	1365	C
84	Aa	1402	G
84	Aa	1403	G
84	Aa	1404	G
84	Aa	1417	G
84	Aa	1421	A

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	1422	G
84	Aa	1431	G
84	Aa	1436	A
84	Aa	1437	G
84	Aa	1440	C
84	Aa	1446	G
84	Aa	1449	A
84	Aa	1452	A
84	Aa	1453	G
84	Aa	1455	A
84	Aa	1481	C
84	Aa	1484	A
84	Aa	1488	G
84	Aa	1491	G
84	Aa	1511	C
84	Aa	1526	A
84	Aa	1529	C
84	Aa	1530	C
84	Aa	1531	G
84	Aa	1542	A
84	Aa	1545	G
84	Aa	1546	G
84	Aa	1550	A
84	Aa	1554	C
84	Aa	1556	G
84	Aa	1566	C
84	Aa	1567	G
84	Aa	1568	A
84	Aa	1570	C
84	Aa	1572	C
84	Aa	1577	A
84	Aa	1578	U
84	Aa	1584	A
84	Aa	1586	A
84	Aa	1602	A
84	Aa	1605	U
84	Aa	1618	U
84	Aa	1625	G
84	Aa	1640	A
84	Aa	1642	G
84	Aa	1652	G
84	Aa	1654	C

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	1680	A
84	Aa	1703	C
84	Aa	1714	A
84	Aa	1715	C
84	Aa	1720	C
84	Aa	1721	A
84	Aa	1722	G
84	Aa	1723	C
84	Aa	1724	C
84	Aa	1726	G
84	Aa	1727	A
84	Aa	1728	G
84	Aa	1729	G
84	Aa	1734	G
84	Aa	1740	U
84	Aa	1748	A
84	Aa	1749	G
84	Aa	1758	U
84	Aa	1760	G
84	Aa	1761	C
84	Aa	1762	G
84	Aa	1766	U
84	Aa	1775	C
84	Aa	1776	G
84	Aa	1777	C
84	Aa	1793	A
84	Aa	1806	C
84	Aa	1808	G
84	Aa	1809	A
84	Aa	1810	G
84	Aa	1812	A
84	Aa	1813	C
84	Aa	1815	G
84	Aa	1816	U
84	Aa	1817	U
84	Aa	1830	U
84	Aa	1831	A
84	Aa	1835	A
84	Aa	1836	U
84	Aa	1837	A
84	Aa	1838	A
84	Aa	1842	C

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	1845	C
84	Aa	1846	A
84	Aa	1851	U
84	Aa	1862	C
84	Aa	1872	C
84	Aa	1874	A
84	Aa	1875	A
84	Aa	1876	U
84	Aa	1882	A
84	Aa	1897	A
84	Aa	1901	G
84	Aa	1902	G
84	Aa	1926	A
84	Aa	1938	U
84	Aa	1951	C
84	Aa	1970	A
84	Aa	1991	U
84	Aa	1996	C
84	Aa	1997	G
84	Aa	1999	G
84	Aa	2003	C
84	Aa	2004	U
84	Aa	2006	A
84	Aa	2007	C
84	Aa	2008	G
84	Aa	2012	C
84	Aa	2013	G
84	Aa	2015	G
84	Aa	2021	G
84	Aa	2042	G
84	Aa	2054	A
84	Aa	2056	C
84	Aa	2057	G
84	Aa	2058	C
84	Aa	2071	U
84	Aa	2073	U
84	Aa	2075	C
84	Aa	2077	C
84	Aa	2081	C
84	Aa	2082	A
84	Aa	2084	G
84	Aa	2088	C

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	2101	A
84	Aa	2107	A
84	Aa	2108	C
84	Aa	2115	G
84	Aa	2116	G
84	Aa	2125	A
84	Aa	2134	U
84	Aa	2150	C
84	Aa	2151	G
84	Aa	2152	A
84	Aa	2153	U
84	Aa	2154	G
84	Aa	2160	C
84	Aa	2161	G
84	Aa	2162	C
84	Aa	2163	G
84	Aa	2167	G
84	Aa	2168	C
84	Aa	2170	G
84	Aa	2183	A
84	Aa	2188	U
84	Aa	2196	G
84	Aa	2200	U
84	Aa	2203	A
84	Aa	2205	G
84	Aa	2223	A
84	Aa	2239	A
84	Aa	2244	G
84	Aa	2245	G
84	Aa	2247	A
84	Aa	2248	G
84	Aa	2250	A
84	Aa	2251	A
84	Aa	2267	G
84	Aa	2268	G
84	Aa	2276	A
84	Aa	2277	U
84	Aa	2278	G
84	Aa	2279	C
84	Aa	2283	G
84	Aa	2287	U
84	Aa	2302	G

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	2303	C
84	Aa	2304	A
84	Aa	2305	U
84	Aa	2308	A
84	Aa	2309	U
84	Aa	2310	G
84	Aa	2314	G
84	Aa	2315	G
84	Aa	2317	U
84	Aa	2319	A
84	Aa	2335	U
84	Aa	2372	A
84	Aa	2373	C
84	Aa	2374	G
84	Aa	2384	G
84	Aa	2385	A
84	Aa	2387	U
84	Aa	2392	G
84	Aa	2396	A
84	Aa	2401	A
84	Aa	2402	G
84	Aa	2403	A
84	Aa	2405	C
84	Aa	2410	U
84	Aa	2443	C
84	Aa	2445	U
84	Aa	2450	G
84	Aa	2451	G
84	Aa	2452	U
84	Aa	2453	G
84	Aa	2454	U
84	Aa	2458	A
84	Aa	2460	A
84	Aa	2461	A
84	Aa	2462	G
84	Aa	2465	G
84	Aa	2467	A
84	Aa	2473	C
84	Aa	2474	A
84	Aa	2481	C
84	Aa	2483	A
84	Aa	2485	U

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	2490	U
84	Aa	2491	A
84	Aa	2492	C
84	Aa	2493	C
84	Aa	2494	A
84	Aa	2498	C
84	Aa	2499	U
84	Aa	2501	U
84	Aa	2502	U
84	Aa	2503	A
84	Aa	2504	A
84	Aa	2505	C
84	Aa	2506	G
84	Aa	2510	U
84	Aa	2511	U
84	Aa	2515	C
84	Aa	2516	U
84	Aa	2517	U
84	Aa	2518	A
84	Aa	2524	U
84	Aa	2526	G
84	Aa	2528	U
84	Aa	2529	C
84	Aa	2532	A
84	Aa	2534	G
84	Aa	2535	C
84	Aa	2536	G
84	Aa	2537	G
84	Aa	2539	G
84	Aa	2542	U
84	Aa	2543	G
84	Aa	2546	C
84	Aa	2547	C
84	Aa	2548	U
84	Aa	2549	C
84	Aa	2550	C
84	Aa	2552	U
84	Aa	2553	U
84	Aa	2559	C
84	Aa	2565	C
84	Aa	2566	C
84	Aa	2573	U

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	2574	A
84	Aa	2579	G
84	Aa	2584	U
84	Aa	2585	C
84	Aa	2588	G
84	Aa	2590	C
84	Aa	2596	A
84	Aa	2597	C
84	Aa	2609	G
84	Aa	2610	G
84	Aa	2617	G
84	Aa	2629	C
84	Aa	2655	U
84	Aa	2659	A
84	Aa	2675	G
84	Aa	2677	A
84	Aa	2680	G
84	Aa	2681	A
84	Aa	2684	U
84	Aa	2692	G
84	Aa	2693	G
84	Aa	2694	A
84	Aa	2696	C
84	Aa	2697	A
84	Aa	2699	A
84	Aa	2702	G
84	Aa	2708	A
84	Aa	2717	G
84	Aa	2731	G
84	Aa	2732	U
84	Aa	2755	U
84	Aa	2756	G
84	Aa	2765	A
84	Aa	2774	A
84	Aa	2779	G
84	Aa	2780	G
84	Aa	2781	A
84	Aa	2798	G
84	Aa	2801	A
84	Aa	2802	G
84	Aa	2803	A
84	Aa	2804	A

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	2812	C
84	Aa	2818	G
84	Aa	2819	A
84	Aa	2820	U
84	Aa	2831	U
84	Aa	2844	U
84	Aa	2847	A
84	Aa	2851	C
84	Aa	2869	C
84	Aa	2873	G
84	Aa	2874	A
84	Aa	2875	U
84	Aa	2877	U
84	Aa	2880	G
84	Aa	2881	C
84	Aa	2889	A
84	Aa	2891	C
84	Aa	2898	A
84	Aa	2899	A
84	Aa	2900	G
84	Aa	2901	C
84	Aa	2916	G
84	Aa	2925	U
84	Aa	2929	C
84	Aa	2937	U
84	Aa	2938	A
84	Aa	2939	G
84	Aa	2944	C
84	Aa	2949	G
84	Aa	2953	G
84	Aa	2957	U
84	Aa	2959	G
84	Aa	2973	A
84	Aa	2985	C
84	Aa	2992	G
84	Aa	2994	U
84	Aa	2997	C
84	Aa	2998	A
84	Aa	3013	A
84	Aa	3050	A
84	Aa	3058	U
84	Aa	3059	C

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	3060	G
84	Aa	3079	G
84	Aa	3080	U
84	Aa	3081	G
84	Aa	3087	A
84	Aa	3093	C
84	Aa	3114	A
84	Aa	3120	U
84	Aa	3123	A
84	Aa	3129	G
84	Aa	3131	A
84	Aa	3132	U
84	Aa	3140	A
84	Aa	3143	A
84	Aa	3144	U
84	Aa	3152	C
84	Aa	3153	U
84	Aa	3154	G
84	Aa	3155	C
84	Aa	3162	C
84	Aa	3163	G
84	Aa	3166	C
84	Aa	3167	G
84	Aa	3168	C
84	Aa	3169	C
84	Aa	3170	C
84	Aa	3171	C
84	Aa	3172	G
84	Aa	3174	C
84	Aa	3176	C
84	Aa	3177	A
84	Aa	3178	C
84	Aa	3182	A
84	Aa	3190	U
84	Aa	3191	U
84	Aa	3192	G
84	Aa	3193	C
84	Aa	3201	A
84	Aa	3202	G
84	Aa	3208	G
84	Aa	3209	U
84	Aa	3210	G

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	3211	C
84	Aa	3212	C
84	Aa	3213	A
84	Aa	3222	G
84	Aa	3227	U
84	Aa	3230	G
84	Aa	3231	G
84	Aa	3234	G
84	Aa	3235	A
84	Aa	3236	A
84	Aa	3237	G
84	Aa	3239	G
84	Aa	3245	G
84	Aa	3251	C
84	Aa	3252	G
84	Aa	3264	C
84	Aa	3265	C
84	Aa	3266	U
84	Aa	3268	C
84	Aa	3271	A
84	Aa	3273	C
84	Aa	3274	G
84	Aa	3278	G
84	Aa	3279	G
84	Aa	3281	G
84	Aa	3286	G
84	Aa	3287	A
84	Aa	3295	G
84	Aa	3296	C
84	Aa	3305	U
84	Aa	3308	A
84	Aa	3309	U
84	Aa	3310	A
84	Aa	3320	G
84	Aa	3322	A
84	Aa	3324	U
84	Aa	3328	A
84	Aa	3333	C
84	Aa	3334	A
84	Aa	3337	G
84	Aa	3339	G
84	Aa	3340	G

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
84	Aa	3341	C
84	Aa	3342	C
84	Aa	3343	U
84	Aa	3344	U
84	Aa	3345	G
84	Aa	3346	C
84	Aa	3347	U
84	Aa	3348	G
84	Aa	3361	G
84	Aa	3367	C
84	Aa	3370	U
84	Aa	3374	C
84	Aa	3381	C
84	Aa	3382	A
84	Aa	3383	C
84	Aa	3385	G
84	Aa	3391	U
85	Ac	23	C
85	Ac	34	U
85	Ac	47	U
85	Ac	48	A
85	Ac	49	G
85	Ac	52	A
85	Ac	59	A
85	Ac	62	C
85	Ac	63	C
85	Ac	73	U
85	Ac	80	A
85	Ac	81	U
85	Ac	82	C
85	Ac	83	C
85	Ac	85	G
85	Ac	86	U
85	Ac	87	G
85	Ac	90	C
85	Ac	92	A
85	Ac	93	U
85	Ac	95	G
85	Ac	104	A
85	Ac	105	A
85	Ac	106	C
85	Ac	111	G

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
85	Ac	113	U
85	Ac	125	C
85	Ac	128	C
85	Ac	129	C
85	Ac	130	G
85	Ac	140	A
85	Ac	150	G
85	Ac	155	U
85	Ac	159	G
85	Ac	160	C
86	Ab	11	A
86	Ab	13	A
86	Ab	14	C
86	Ab	22	A
86	Ab	26	C
86	Ab	41	G
86	Ab	42	A
86	Ab	48	G
86	Ab	49	A
86	Ab	50	A
86	Ab	52	U
86	Ab	53	U
86	Ab	63	U
86	Ab	64	G
86	Ab	73	U
86	Ab	75	G
86	Ab	93	U
86	Ab	100	A
86	Ab	101	A
86	Ab	108	G
86	Ab	110	G
86	Ab	113	G
86	Ab	119	C

There are no RNA pucker outliers to report.

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

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

## 5.5 Carbohydrates [i](#)

There are no monosaccharides in this entry.

## 5.6 Ligand geometry [i](#)

There are no ligands in this entry.

## 5.7 Other polymers [i](#)

There are no such residues in this entry.

## 5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

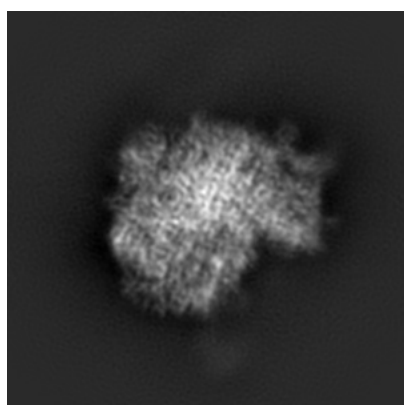
## 6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-1780. These allow visual inspection of the internal detail of the map and identification of artifacts.

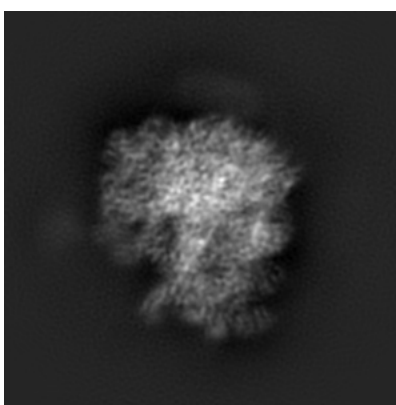
No raw map or half-maps were deposited for this entry and therefore no images, graphs, etc. pertaining to the raw map can be shown.

### 6.1 Orthogonal projections [i](#)

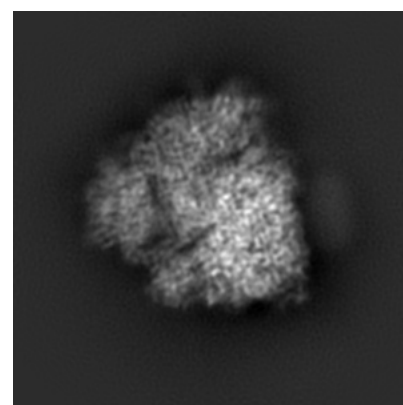
#### 6.1.1 Primary map



X



Y

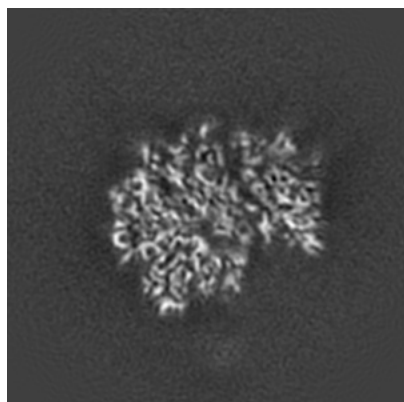


Z

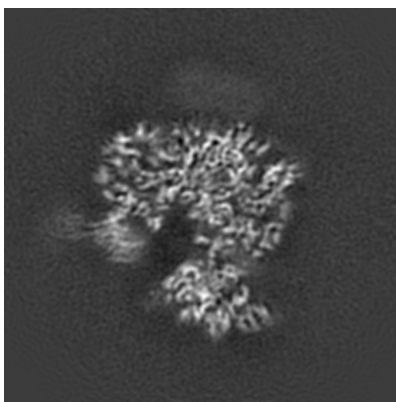
The images above show the map projected in three orthogonal directions.

### 6.2 Central slices [i](#)

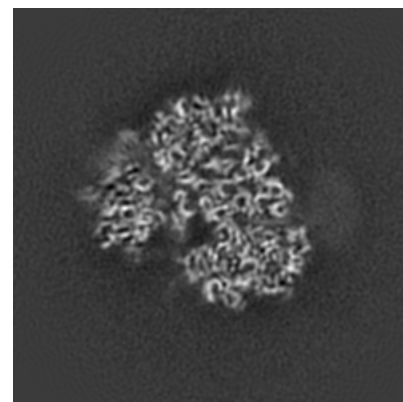
#### 6.2.1 Primary map



X Index: 184



Y Index: 184

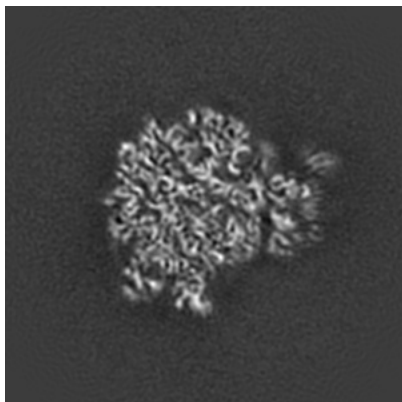


Z Index: 184

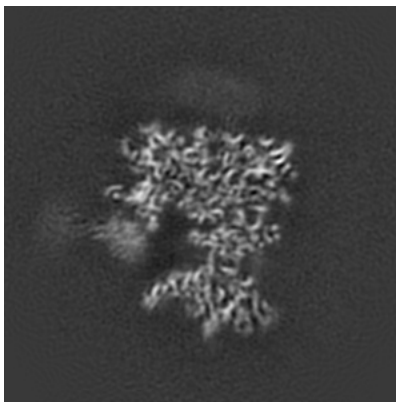
The images above show central slices of the map in three orthogonal directions.

## 6.3 Largest variance slices [i](#)

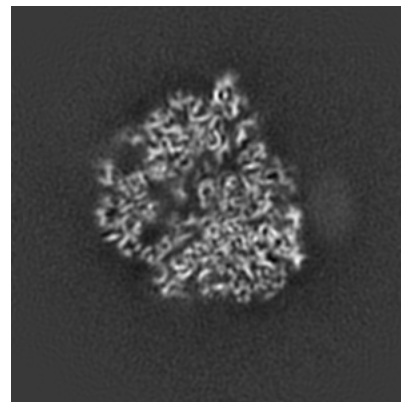
### 6.3.1 Primary map



X Index: 213



Y Index: 193

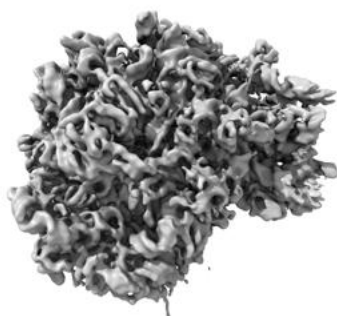


Z Index: 173

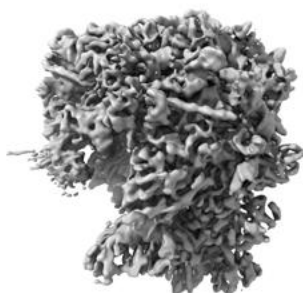
The images above show the largest variance slices of the map in three orthogonal directions.

## 6.4 Orthogonal surface views [i](#)

### 6.4.1 Primary map



X



Y



Z

The images above show the 3D surface view of the map at the recommended contour level 0.11. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

## 6.5 Mask visualisation

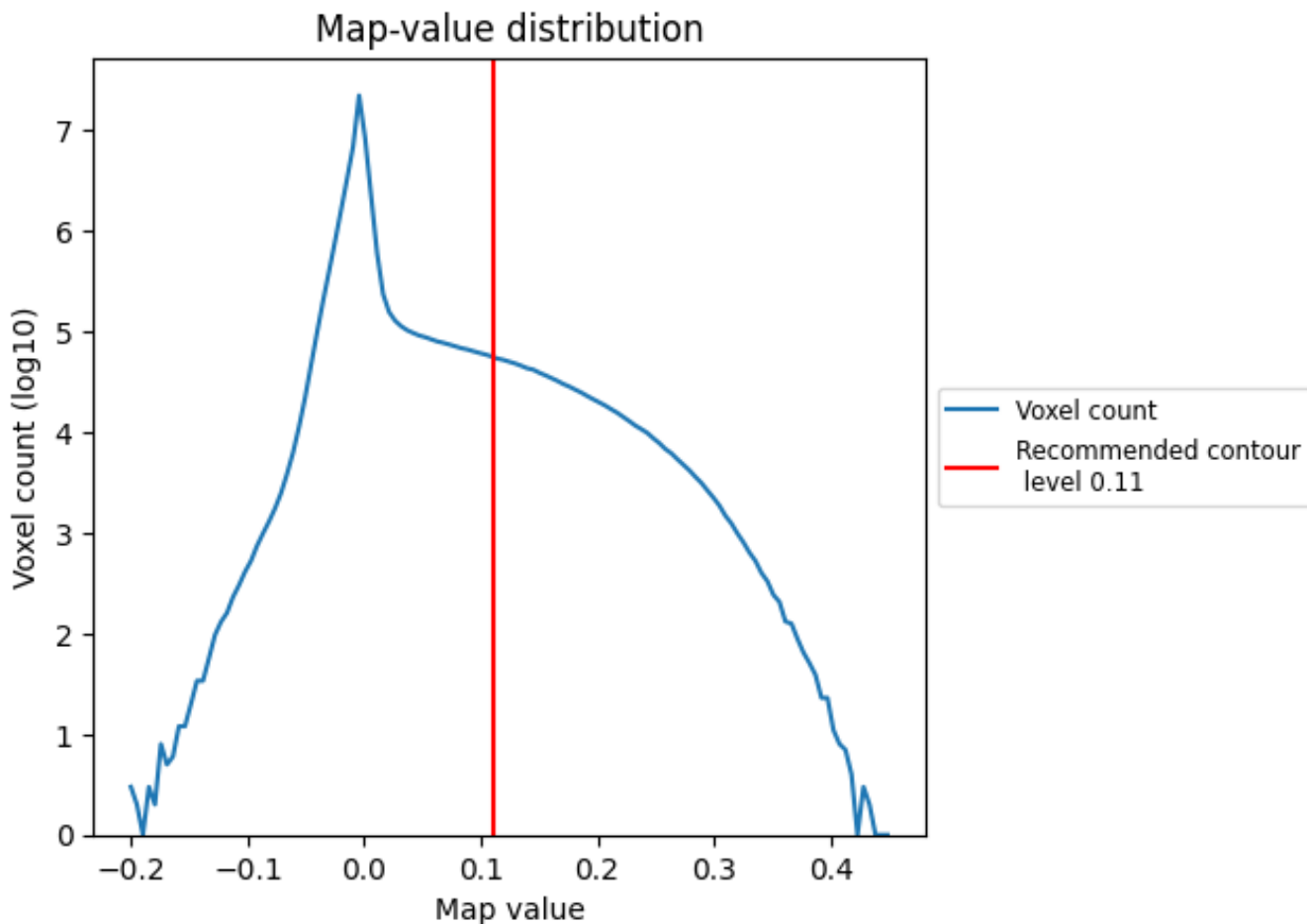
This section was not generated. No masks/segmentation were deposited.



## 7 Map analysis [i](#)

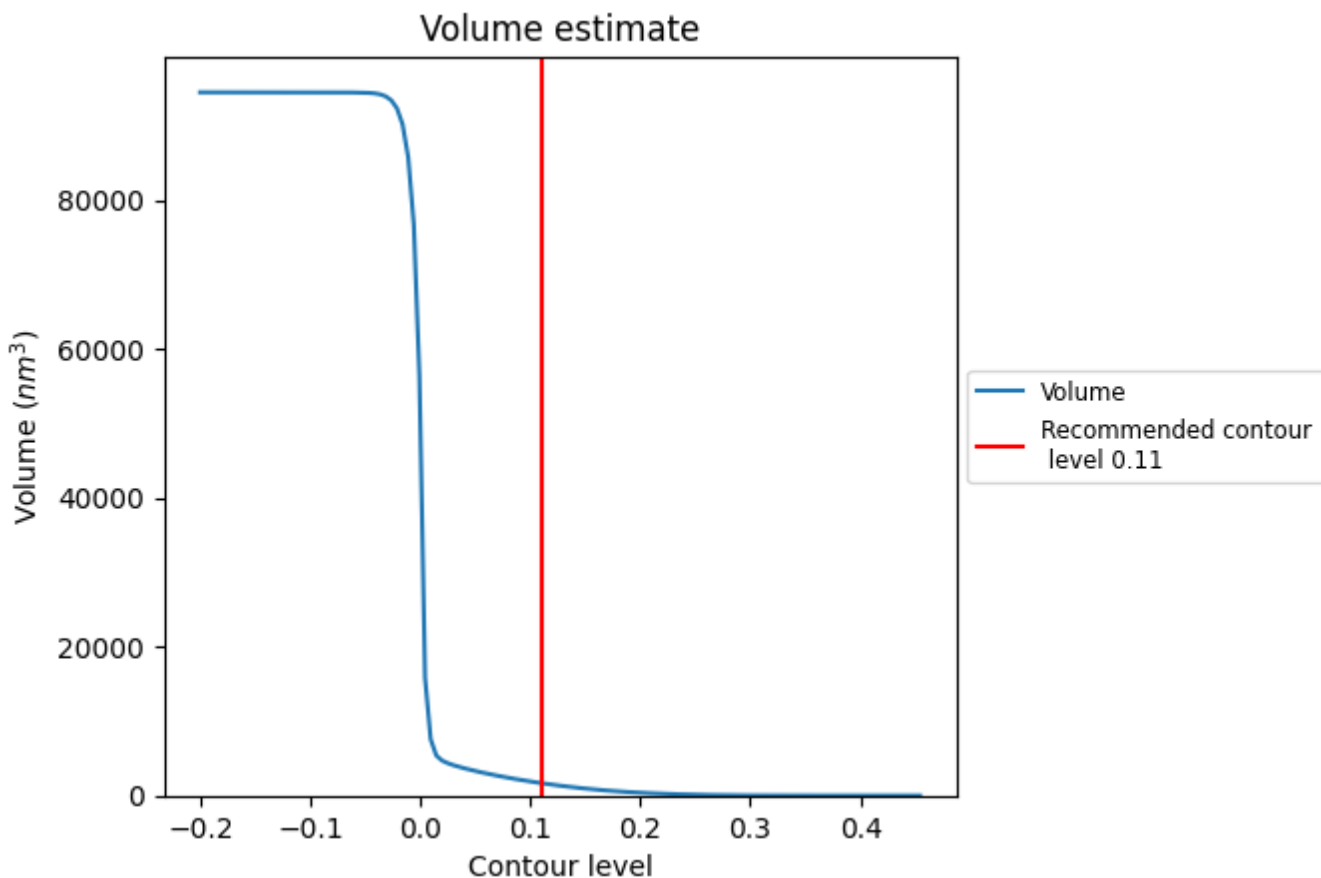
This section contains the results of statistical analysis of the map.

### 7.1 Map-value distribution [i](#)



The map-value distribution is plotted in 128 intervals along the x-axis. The y-axis is logarithmic. A spike in this graph at zero usually indicates that the volume has been masked.

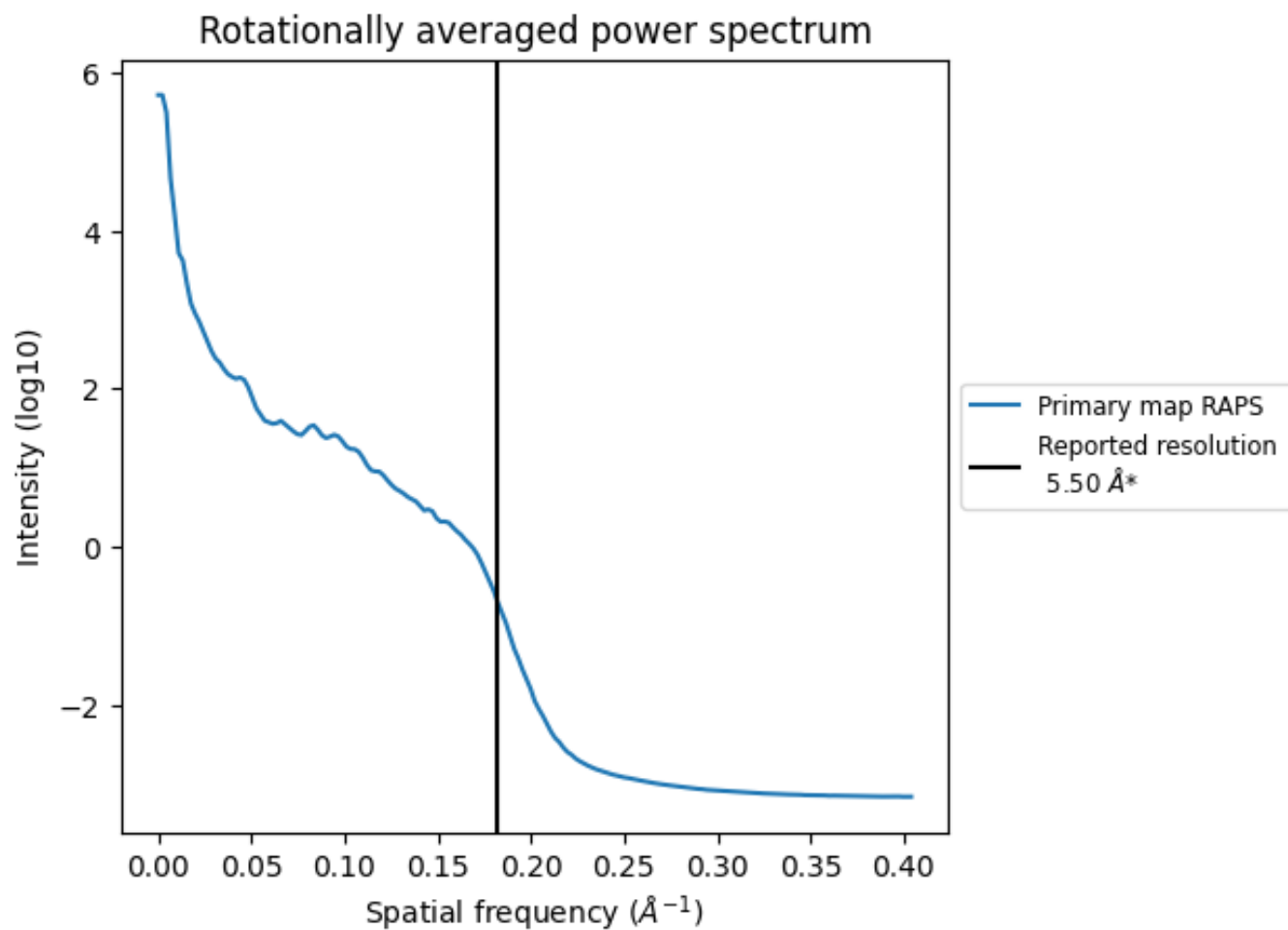
## 7.2 Volume estimate [i](#)



The volume at the recommended contour level is 1668 nm<sup>3</sup>; this corresponds to an approximate mass of 1506 kDa.

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

### 7.3 Rotationally averaged power spectrum [i](#)



\*Reported resolution corresponds to spatial frequency of  $0.182 \text{\AA}^{-1}$

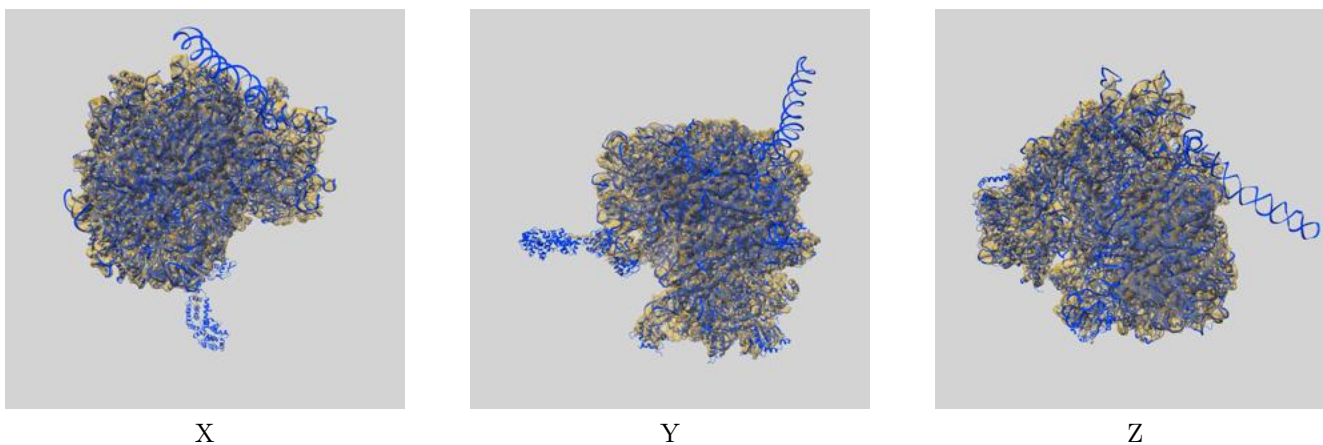
## 8 Fourier-Shell correlation

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

## 9 Map-model fit [i](#)

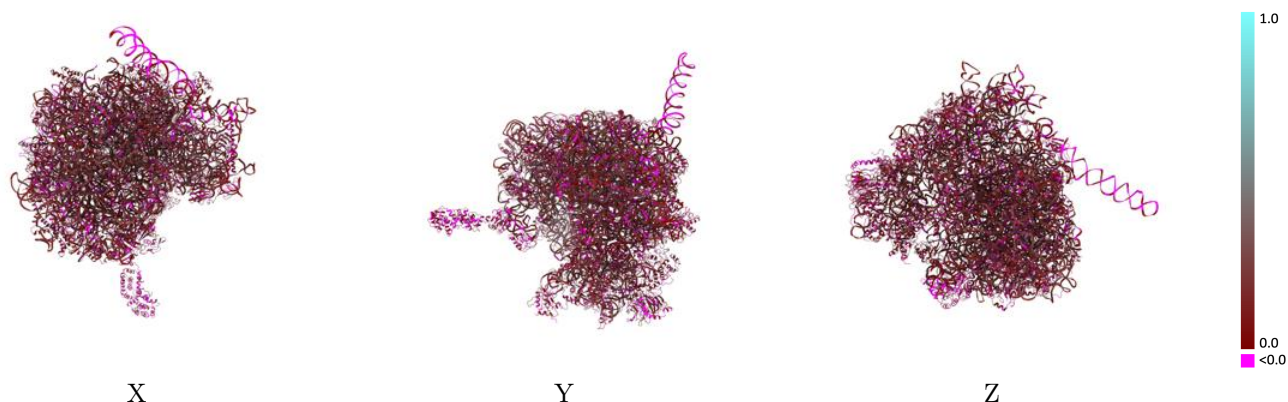
This section contains information regarding the fit between EMDB map EMD-1780 and PDB model 4V7E. Per-residue inclusion information can be found in section 3 on page 20.

### 9.1 Map-model overlay [i](#)



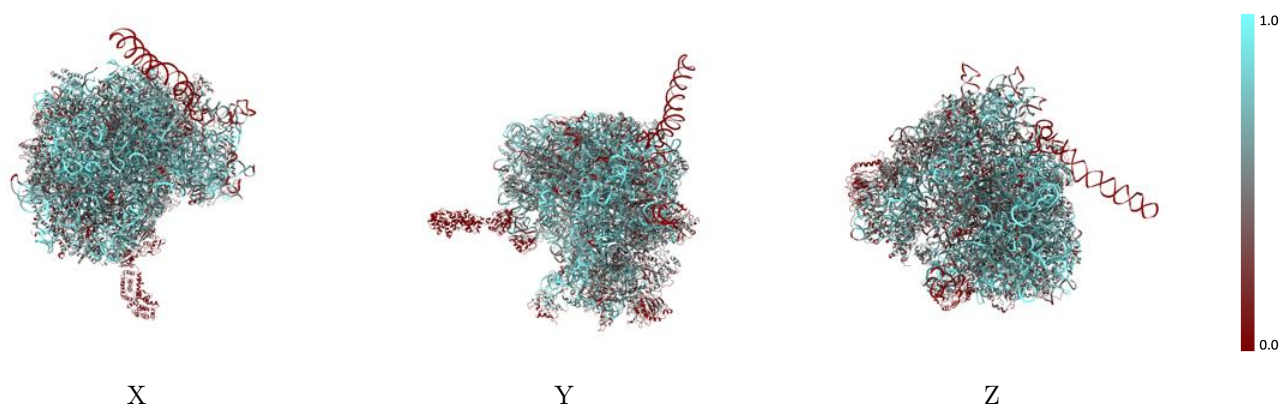
The images above show the 3D surface view of the map at the recommended contour level 0.11 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



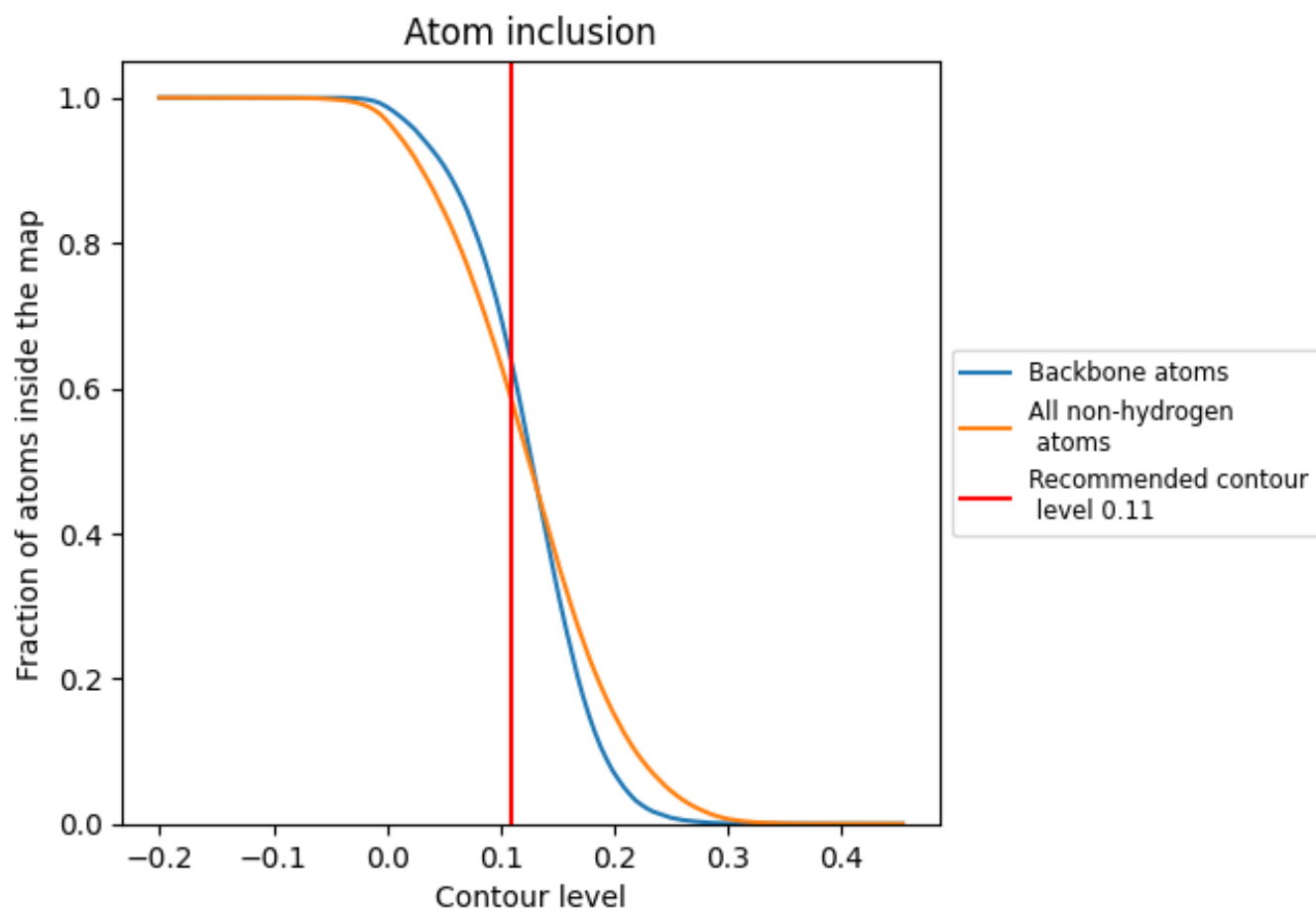
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions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

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



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (0.11).







































































## 9.4 Atom inclusion [i](#)



At the recommended contour level, 63% of all backbone atoms, 58% of all non-hydrogen atoms, are inside the map.

## 9.5 Map-model fit summary

The table lists the average atom inclusion at the recommended contour level (0.11) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.5826	 0.1550
Aa	 0.7377	 0.1940
Ab	 0.8528	 0.2080
Ac	 0.7934	 0.2070
Ad	 0.7396	 0.1960
Ae	 0.4915	 0.1860
Af	 0.1293	 0.0800
BA	 0.3804	 0.1240
BB	 0.4210	 0.1260
BC	 0.3646	 0.1210
BD	 0.3293	 0.1190
BE	 0.4311	 0.1040
BF	 0.4542	 0.1280
BG	 0.4302	 0.1120
BH	 0.3666	 0.1230
BI	 0.4139	 0.1110
BJ	 0.4894	 0.1170
BK	 0.4180	 0.0950
BL	 0.3153	 0.1070
BM	 0.2871	 0.0870
BN	 0.3866	 0.0980
BO	 0.3839	 0.1130
BP	 0.4370	 0.1110
BQ	 0.3975	 0.0950
BR	 0.3454	 0.1130
BS	 0.4000	 0.1130
BT	 0.4459	 0.1070
BU	 0.3075	 0.0920
BV	 0.3601	 0.1050
BW	 0.3448	 0.0840
BX	 0.4157	 0.1190
BY	 0.4148	 0.0740
BZ	 0.3447	 0.0990
Ba	 0.4599	 0.1350
Bb	 0.4000	 0.1200



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Chain	Atom inclusion	Q-score
Bc	0.3077	0.0630
Bd	0.3736	0.0680
Be	0.3031	0.0730
Bf	0.3569	0.0730
Bg	0.2151	0.0850
CA	0.3965	0.1150
CB	0.4292	0.1080
CC	0.4029	0.1160
CD	0.4762	0.1050
CE	0.3678	0.0960
CF	0.4640	0.1200
CG	0.4751	0.1180
CH	0.4758	0.1140
CI	0.4448	0.1210
CJ	0.4656	0.1300
CK	0.0583	0.0530
CL	0.4417	0.1020
CM	0.4739	0.1230
CN	0.4819	0.1090
CO	0.4495	0.1070
CP	0.4274	0.1150
CQ	0.4246	0.1150
CR	0.4299	0.1140
CS	0.4637	0.1040
CT	0.3952	0.1130
CU	0.3175	0.0650
CV	0.2652	0.1300
CW	0.2757	0.1230
CX	0.3930	0.1120
CY	0.5448	0.1270
CZ	0.5066	0.1270
Ca	0.4074	0.0920
Cb	0.3613	0.0940
Cc	0.4303	0.1090
Cd	0.4082	0.0930
Ce	0.3661	0.1060
Cf	0.4198	0.0960
Cg	0.4448	0.1210
Ch	0.4536	0.0910
Ci	0.4105	0.0870
Cj	0.4672	0.1040
Ck	0.4709	0.1140

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Chain	Atom inclusion	Q-score
Cl	 0.4312	 0.1180
Cm	 0.5290	 0.1580
Cn	 0.2831	 -0.0170
Co	 0.3911	 0.0940
Cp	 0.3991	 0.1240
Cq	 0.0737	 0.0580
Cr	 0.4893	 0.1140
Cs	 0.0137	 0.0590
Ct	 0.0000	 0.0300
Cu	 0.0000	 0.0430
Cv	 0.0000	 0.0280
Cz	 0.0260	 0.0410