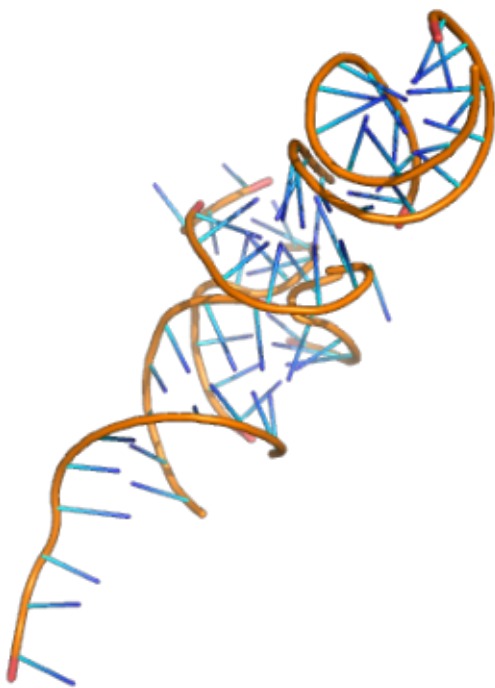


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angle value	angle type	residue 1	residue 2	chain	structure id	z -score	standered deviation	mean
140.96	alpha	('', 6, '')	('', 5, '')	<Chain id=A>	281_36031_1C0A_B	2.37	62.420	-71.120
145.40	alpha	('', 13, '')	('', 12, '')	<Chain id=A>	281_36031_1C0A_B	2.30	62.420	-71.120
142.81	alpha	('', 14, '')	('', 13, '')	<Chain id=A>	281_36031_1C0A_B	2.34	62.420	-71.120
161.27	alpha	('', 53, '')	('', 52, '')	<Chain id=A>	281_36031_1C0A_B	2.04	62.420	-71.120
160.26	alpha	('', 60, '')	('', 59, '')	<Chain id=A>	281_36031_1C0A_B	2.06	62.420	-71.120
-172.99	gamma	('', 13, '')	('', 13, '')	<Chain id=A>	281_36031_1C0A_B	2.40	50.440	65.790
-170.62	gamma	('', 14, '')	('', 14, '')	<Chain id=A>	281_36031_1C0A_B	2.45	50.440	65.790
168.58	gamma	('', 53, '')	('', 53, '')	<Chain id=A>	281_36031_1C0A_B	2.04	50.440	65.790
168.31	gamma	('', 75, '')	('', 75, '')	<Chain id=A>	281_36031_1C0A_B	2.03	50.440	65.790
158.90	epsilon	('', 48, '')	('', 49, '')	<Chain id=A>	281_36031_1C0A_B	2.31	26.930	-138.950
144.15	epsilon	('', 61, '')	('', 62, '')	<Chain id=A>	281_36031_1C0A_B	2.86	26.930	-138.950
135.21	zeta	('', 17, '')	('', 18, '')	<Chain id=A>	281_36031_1C0A_B	3.51	41.730	-78.480
105.22	zeta	('', 48, '')	('', 49, '')	<Chain id=A>	281_36031_1C0A_B	4.22	41.730	-78.480
179.53	zeta	('', 49, '')	('', 50, '')	<Chain id=A>	281_36031_1C0A_B	2.44	41.730	-78.480
81.94	zeta	('', 61, '')	('', 62, '')	<Chain id=A>	281_36031_1C0A_B	3.84	41.730	-78.480
-88.43	chi purine	('', 9, '')	('', 9, '')	<Chain id=A>	281_36031_1C0A_B	2.31	31.310	-160.640
-89.08	chi purine	('', 18, '')	('', 18, '')	<Chain id=A>	281_36031_1C0A_B	2.29	31.310	-160.640
-89.38	chi purine	('', 19, '')	('', 19, '')	<Chain id=A>	281_36031_1C0A_B	2.28	31.310	-160.640

distance value	residue	chain	structure id	atom1, atom2	z-score	standered deviation	mean
1.12	(' ', 6, ' ')	<Chain id=A>	281_36031_1C0A_B	O3',P	-4.00	0.122	1.608
1.99	(' ', 19, ' ')	<Chain id=A>	281_36031_1C0A_B	O3',P	3.14	0.122	1.608
1.90	(' ', 23, ' ')	<Chain id=A>	281_36031_1C0A_B	O3',P	2.40	0.122	1.608
2.11	(' ', 49, ' ')	<Chain id=A>	281_36031_1C0A_B	O3',P	4.14	0.122	1.608
1.89	(' ', 59, ' ')	<Chain id=A>	281_36031_1C0A_B	O3',P	2.30	0.122	1.608
1.93	(' ', 60, ' ')	<Chain id=A>	281_36031_1C0A_B	O3',P	2.61	0.122	1.608