



Full wwPDB X-ray Structure Validation Report ⓘ

Mar 28, 2026 – 10:03 PM UTC

PDB ID : 4U4Z / pdb_00004u4z
Title : Crystal structure of Phyllanthoside bound to the yeast 80S ribosome
Authors : Garreau de Loubresse, N.; Prokhorova, I.; Yusupova, G.; Yusupov, M.
Deposited on : 2014-07-24
Resolution : 3.10 Å(reported)

This is a Full wwPDB X-ray Structure Validation Report for a publicly released PDB entry.

We welcome your comments at validation@mail.wwpdb.org

A user guide is available at

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

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

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

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

MolProbity : **FAILED**
Mogul : 2022.3.0, CSD as543be (2022)
Xtriage (Phenix) : 2.0
EDS : 3.0
Buster-report : wwPDB partial adaption of 1.1.7 (2018)
Percentile statistics : 20250101.v01 (using entries in the PDB archive January 1st 2025)
CCP4 : 9.0.010 (Gargrove)
Density-Fitness : 1.0.12
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.49

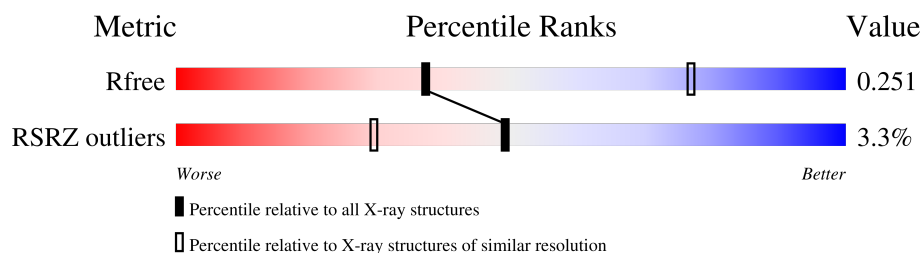
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 3.10 Å.

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



Metric	Whole archive (#Entries)	Similar resolution (#Entries, resolution range(Å))
R_{free}	180053	1456 (3.10-3.10)
RSRZ outliers	180081	1456 (3.10-3.10)

MolProbity failed to run properly - the sequence quality summary graphics cannot be shown.

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

Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
85	MG	1	3668	-	-	-	X
85	MG	1	3717	-	-	-	X
85	MG	1	3815	-	-	-	X
85	MG	1	3825	-	-	-	X
85	MG	2	1909	-	-	-	X
85	MG	2	1913	-	-	-	X
85	MG	2	1915	-	-	-	X
85	MG	2	1924	-	-	-	X
85	MG	2	1928	-	-	-	X
85	MG	2	1949	-	-	-	X
85	MG	2	1952	-	-	-	X
85	MG	2	1958	-	-	-	X
85	MG	2	1961	-	-	-	X
85	MG	2	1967	-	-	-	X
85	MG	2	1974	-	-	-	X
85	MG	2	1977	-	-	-	X
85	MG	2	1985	-	-	-	X
85	MG	2	1988	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
85	MG	2	1993	-	-	-	X
85	MG	2	2001	-	-	-	X
85	MG	2	2018	-	-	-	X
85	MG	5	3467	-	-	-	X
85	MG	5	3666	-	-	-	X
85	MG	5	3687	-	-	-	X
85	MG	5	3882	-	-	-	X
85	MG	6	1909	-	-	-	X
85	MG	6	1911	-	-	-	X
85	MG	6	1923	-	-	-	X
85	MG	6	1939	-	-	-	X
85	MG	6	1989	-	-	-	X
85	MG	6	1997	-	-	-	X
85	MG	6	2039	-	-	-	X
85	MG	D0	201	-	-	-	X

2 Entry composition

There are 88 unique types of molecules in this entry. The entry contains 411276 atoms, of which 0 are hydrogens and 0 are deuteriums.

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
1	2	1750	Total	C	N	O	P	0	0	0
			37283	16668	6591	12274	1750			
1	6	1795	Total	C	N	O	P	0	0	0
			38238	17095	6758	12590	1795			

- Molecule 2 is a protein called 40S ribosomal protein S0-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
2	S0	206	Total	C	N	O	S	0	0	0
			1577	1014	278	283	2			
2	s0	206	Total	C	N	O	S	0	0	0
			1583	1017	281	283	2			

- Molecule 3 is a protein called 40S ribosomal protein S1-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
3	S1	214	Total	C	N	O	S	0	0	0
			1709	1084	310	311	4			
3	s1	216	Total	C	N	O	S	0	0	0
			1722	1091	312	315	4			

- Molecule 4 is a protein called 40S ribosomal protein S2.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
4	S2	217	Total	C	N	O	S	0	0	0
			1635	1047	289	297	2			
4	s2	217	Total	C	N	O	S	0	0	0
			1635	1047	289	297	2			

- Molecule 5 is a protein called 40S ribosomal protein S3.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
5	S3	223	Total	C	N	O	S	0	0	0
			1734	1101	313	314	6			
5	s3	223	Total	C	N	O	S	0	0	0
			1734	1101	313	314	6			

- Molecule 6 is a protein called 40S ribosomal protein S4-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
6	S4	260	Total	C	N	O	S	0	0	0
			2068	1316	389	360	3			
6	s4	260	Total	C	N	O	S	0	0	0
			2068	1316	389	360	3			

- Molecule 7 is a protein called 40S ribosomal protein S5.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
7	S5	206	Total	C	N	O	S	0	0	0
			1609	1007	300	299	3			
7	s5	206	Total	C	N	O	S	0	0	0
			1609	1007	300	299	3			

- Molecule 8 is a protein called 40S ribosomal protein S6-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
8	S6	226	Total	C	N	O	S	0	0	0
			1799	1129	346	321	3			
8	s6	218	Total	C	N	O	S	0	0	0
			1755	1102	337	313	3			

- Molecule 9 is a protein called 40S ribosomal protein S7-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
9	S7	184	Total	C	N	O		0	0	0
			1481	951	265	265				
9	s7	186	Total	C	N	O		0	0	0
			1491	957	267	267				

- Molecule 10 is a protein called 40S ribosomal protein S8-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	S8	188	Total	C	N	O	S	0	0	0
			1489	925	298	264	2			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	s8	188	Total	C	N	O	S	0	0	0
			1489	925	298	264	2			

- Molecule 11 is a protein called 40S ribosomal protein S9-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
11	S9	185	Total	C	N	O	S	0	0	0
			1494	943	289	261	1			
11	s9	185	Total	C	N	O	S	0	0	0
			1494	943	289	261	1			

- Molecule 12 is a protein called 40S ribosomal protein S10-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
12	C0	96	Total	C	N	O	S	0	0	0
			773	500	126	145	2			
12	c0	96	Total	C	N	O	S	0	0	0
			762	491	125	144	2			

There are 2 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
C0	89	ALA	GLY	conflict	UNP Q08745
c0	89	ALA	GLY	conflict	UNP Q08745

- Molecule 13 is a protein called 40S ribosomal protein S11-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
13	C1	155	Total	C	N	O	S	0	0	0
			1214	775	230	206	3			
13	c1	146	Total	C	N	O	S	0	0	0
			1168	747	221	197	3			

There are 2 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
C1	147	ALA	GLY	conflict	UNP P0CX47
c1	147	ALA	GLY	conflict	UNP P0CX47

- Molecule 14 is a protein called 40S ribosomal protein S12.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
14	C2	124	Total	C	N	O	S	0	0	0
			892	562	156	172	2			
14	c2	124	Total	C	N	O	S	0	0	0
			892	562	156	172	2			

There are 4 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
C2	104	ALA	GLY	conflict	UNP P48589
C2	110	ALA	GLY	conflict	UNP P48589
c2	104	ALA	GLY	conflict	UNP P48589
c2	110	ALA	GLY	conflict	UNP P48589

- Molecule 15 is a protein called 40S ribosomal protein S13.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
15	C3	150	Total	C	N	O	S	0	0	0
			1192	759	224	207	2			
15	c3	150	Total	C	N	O	S	0	0	0
			1192	759	224	207	2			

- Molecule 16 is a protein called 40S ribosomal protein S14-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
16	C4	127	Total	C	N	O	S	0	0	0
			891	545	182	163	1			
16	c4	128	Total	C	N	O	S	0	0	0
			949	582	188	176	3			

- Molecule 17 is a protein called 40S ribosomal protein S15.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
17	C5	124	Total	C	N	O	S	0	0	0
			977	622	182	166	7			
17	c5	135	Total	C	N	O	S	0	0	0
			1039	658	196	178	7			

There are 2 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
C5	137	SER	ARG	conflict	UNP Q01855
c5	137	SER	ARG	conflict	UNP Q01855

- Molecule 18 is a protein called 40S ribosomal protein S16-A.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
18	C6	141	Total	C	N	O	0	0	0
			1105	708	203	194			
18	c6	142	Total	C	N	O	0	0	0
			1111	711	204	196			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
19	C7	120	Total	C	N	O	S	0	0	0
			926	577	177	170	2			
19	c7	117	Total	C	N	O	S	0	0	0
			906	563	174	167	2			

- Molecule 20 is a protein called 40S ribosomal protein S18-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
20	C8	145	Total	C	N	O	S	0	0	0
			1192	743	237	210	2			
20	c8	145	Total	C	N	O	S	0	0	0
			1192	743	237	210	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
21	C9	143	Total	C	N	O	S	0	0	0
			1112	694	208	208	2			
21	c9	143	Total	C	N	O	S	0	0	0
			1112	694	208	208	2			

- Molecule 22 is a protein called 40S ribosomal protein S20.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
22	D0	107	Total	C	N	O	S	0	0	0
			855	539	156	159	1			
22	d0	110	Total	C	N	O	S	0	0	0
			882	554	161	166	1			

- Molecule 23 is a protein called 40S ribosomal protein S21-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
23	D1	87	Total	C	N	O	S	0	0	0
			684	420	125	137	2			
23	d1	87	Total	C	N	O	S	0	0	0
			684	420	125	137	2			

- Molecule 24 is a protein called 40S ribosomal protein S22-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
24	D2	129	Total	C	N	O	S	0	0	0
			1021	650	188	180	3			
24	d2	129	Total	C	N	O	S	0	0	0
			1021	650	188	180	3			

- Molecule 25 is a protein called 40S ribosomal protein S23-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
25	D3	144	Total	C	N	O	S	0	0	0
			1121	708	220	191	2			
25	d3	144	Total	C	N	O	S	0	0	0
			1121	708	220	191	2			

- Molecule 26 is a protein called 40S ribosomal protein S24-A.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
26	D4	134	Total	C	N	O	0	0	0
			1073	676	208	189			
26	d4	134	Total	C	N	O	0	0	0
			1073	676	208	189			

- Molecule 27 is a protein called 40S ribosomal protein S25-A.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
27	D5	70	Total	C	N	O	0	0	0
			563	360	104	99			
27	d5	69	Total	C	N	O	0	0	0
			558	357	103	98			

- Molecule 28 is a protein called 40S ribosomal protein S26-B.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
28	D6	97	Total	C	N	O	S	0	0	0
			769	475	160	129	5			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
28	d6	97	Total	C	N	O	S	0	0	0
			769	475	160	129	5			

- Molecule 29 is a protein called 40S ribosomal protein S27-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
29	D7	81	Total	C	N	O	S	0	0	0
			610	382	110	113	5			
29	d7	81	Total	C	N	O	S	0	0	0
			610	382	110	113	5			

- Molecule 30 is a protein called 40S ribosomal protein S28-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
30	D8	63	Total	C	N	O	S	0	0	0
			497	306	99	91	1			
30	d8	63	Total	C	N	O	S	0	0	0
			497	306	99	91	1			

- Molecule 31 is a protein called 40S ribosomal protein S29-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
31	D9	53	Total	C	N	O	S	0	0	0
			442	274	92	72	4			
31	d9	53	Total	C	N	O	S	0	0	0
			442	274	92	72	4			

- Molecule 32 is a protein called 40S ribosomal protein S30-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
32	E0	60	Total	C	N	O	S	0	0	0
			475	299	98	77	1			

- Molecule 33 is a protein called Ubiquitin-40S ribosomal protein S31.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
33	E1	71	Total	C	N	O	S	0	0	0
			566	362	106	94	4			
33	e1	76	Total	C	N	O	S	0	0	0
			608	388	117	99	4			

- Molecule 34 is a protein called Guanine nucleotide-binding protein subunit beta-like protein.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
34	SR	318	Total	C	N	O	S	0	0	0
			2441	1544	419	470	8			
34	sR	318	Total	C	N	O	S	0	0	0
			2442	1544	418	472	8			

- Molecule 35 is a protein called Suppressor protein STM1.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
35	SM	159	Total	C	N	O		0	0	0
			1104	652	221	231				
35	sM	104	Total	C	N	O		0	0	0
			679	402	140	137				

- Molecule 36 is a RNA chain called 25S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
36	1	3149	Total	C	N	O	P	0	0	0
			67355	30086	12142	21978	3149			
36	5	3150	Total	C	N	O	P	0	0	0
			67376	30095	12145	21987	3149			

- Molecule 37 is a RNA chain called 5S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
37	3	121	Total	C	N	O	P	0	0	0
			2579	1152	461	845	121			
37	7	121	Total	C	N	O	P	0	0	0
			2579	1152	461	845	121			

- Molecule 38 is a RNA chain called 5.8S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
38	4	158	Total	C	N	O	P	0	0	0
			3353	1500	586	1109	158			
38	8	158	Total	C	N	O	P	0	0	0
			3353	1500	586	1109	158			

- Molecule 39 is a protein called 60S ribosomal protein L2-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
39	L2	252	Total	C	N	O	S	0	0	0
			1914	1191	388	334	1			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
39	12	252	Total	C	N	O	S	0	0	0
			1912	1190	388	333	1			

- Molecule 40 is a protein called 60S ribosomal protein L3.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
40	L3	386	Total	C	N	O	S	0	0	0
			3075	1950	584	533	8			
40	l3	386	Total	C	N	O	S	0	0	0
			3075	1950	584	533	8			

- Molecule 41 is a protein called 60S ribosomal protein L4-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
41	L4	361	Total	C	N	O	S	0	0	0
			2748	1729	522	494	3			
41	l4	361	Total	C	N	O	S	0	0	0
			2748	1729	522	494	3			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
42	L5	296	Total	C	N	O	S	0	0	0
			2375	1501	414	458	2			
42	l5	294	Total	C	N	O	S	0	0	0
			2359	1489	412	456	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
43	L6	156	Total	C	N	O	S	0	0	0
			1239	800	222	216	1			
43	l6	157	Total	C	N	O	S	0	0	0
			1248	806	224	217	1			

- Molecule 44 is a protein called 60S ribosomal protein L7-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
44	L7	222	Total	C	N	O	S	0	0	0
			1784	1151	324	308	1			
44	l7	223	Total	C	N	O	S	0	0	0
			1791	1155	325	310	1			

- Molecule 45 is a protein called 60S ribosomal protein L8-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
45	L8	233	Total	C	N	O	S	0	0	0
			1804	1151	323	327	3			
45	l8	231	Total	C	N	O	S	0	0	0
			1763	1130	316	314	3			

- Molecule 46 is a protein called 60S ribosomal protein L9-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
46	L9	191	Total	C	N	O	S	0	0	0
			1518	963	274	277	4			
46	l9	191	Total	C	N	O	S	0	0	0
			1518	963	274	277	4			

- Molecule 47 is a protein called 60S ribosomal protein L10.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
47	M0	211	Total	C	N	O	S	0	0	0
			1705	1083	322	294	6			
47	m0	213	Total	C	N	O	S	0	0	0
			1722	1094	325	297	6			

- Molecule 48 is a protein called 60S ribosomal protein L11-B.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
48	M1	169	Total	C	N	O	S	0	0	0
			1353	847	253	249	4			
48	m1	169	Total	C	N	O	S	0	0	0
			1353	847	253	249	4			

- Molecule 49 is a protein called 60S ribosomal protein L13-A.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
49	M3	193	Total	C	N	O	0	0	0
			1543	962	315	266			
49	m3	194	Total	C	N	O	0	0	0
			1548	965	316	267			

- Molecule 50 is a protein called 60S ribosomal protein L14-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
50	M4	136	Total	C	N	O	S	0	0	0
			1053	675	199	177	2			
50	m4	137	Total	C	N	O	S	0	0	0
			1059	678	200	179	2			

- Molecule 51 is a protein called 60S ribosomal protein L15-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
51	M5	203	Total	C	N	O	S	0	0	0
			1720	1077	361	281	1			
51	m5	203	Total	C	N	O	S	0	0	0
			1720	1077	361	281	1			

- Molecule 52 is a protein called 60S ribosomal protein L16-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
52	M6	197	Total	C	N	O	S	0	0	0
			1555	1003	289	262	1			
52	m6	197	Total	C	N	O	S	0	0	0
			1555	1003	289	262	1			

- Molecule 53 is a protein called 60S ribosomal protein L17-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
53	M7	183	Total	C	N	O		0	0	0
			1420	882	281	257				
53	m7	155	Total	C	N	O		0	0	0
			1227	764	238	225				

- Molecule 54 is a protein called 60S ribosomal protein L18-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
54	M8	185	Total	C	N	O	S	0	0	0
			1441	908	290	241	2			
54	m8	185	Total	C	N	O	S	0	0	0
			1441	908	290	241	2			

- Molecule 55 is a protein called 60S ribosomal protein L19-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
55	M9	188	Total	C	N	O		0	0	0
			1521	935	326	260				

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Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
55	m9	188	Total	C	N	O	0	0	0
			1521	935	326	260			

- Molecule 56 is a protein called 60S ribosomal protein L20-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
56	N0	172	Total	C	N	O	S	0	0	0
			1445	930	267	244	4			
56	n0	172	Total	C	N	O	S	0	0	0
			1445	930	267	244	4			

- Molecule 57 is a protein called 60S ribosomal protein L21-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
57	N1	159	Total	C	N	O	S	0	0	0
			1276	805	246	221	4			
57	n1	159	Total	C	N	O	S	0	0	0
			1276	805	246	221	4			

- Molecule 58 is a protein called 60S ribosomal protein L22-A.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
58	N2	100	Total	C	N	O	0	0	0
			796	516	131	149			
58	n2	98	Total	C	N	O	0	0	0
			778	505	127	146			

- Molecule 59 is a protein called 60S ribosomal protein L23-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
59	N3	136	Total	C	N	O	S	0	0	0
			1003	628	189	179	7			
59	n3	136	Total	C	N	O	S	0	0	0
			1003	628	189	179	7			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
60	N4	98	Total	C	N	O	S	0	0	0
			699	443	137	118	1			
60	n4	135	Total	C	N	O	S	0	0	0
			1038	651	206	180	1			

- Molecule 61 is a protein called 60S ribosomal protein L25.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
61	N5	121	Total	C	N	O	S	0	0	0
			964	620	169	173	2			
61	n5	120	Total	C	N	O	S	0	0	0
			959	617	168	172	2			

- Molecule 62 is a protein called 60S ribosomal protein L26-A.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
62	N6	126	Total	C	N	O	0	0	0
			993	625	192	176			
62	n6	126	Total	C	N	O	0	0	0
			993	625	192	176			

- Molecule 63 is a protein called 60S ribosomal protein L27-A.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
63	N7	135	Total	C	N	O	0	0	0
			1092	710	202	180			
63	n7	135	Total	C	N	O	0	0	0
			1092	710	202	180			

- Molecule 64 is a protein called 60S ribosomal protein L28.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
64	N8	148	Total	C	N	O	S	0	0	0
			1173	749	231	190	3			
64	n8	148	Total	C	N	O	S	0	0	0
			1173	749	231	190	3			

- Molecule 65 is a protein called 60S ribosomal protein L29.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
65	N9	58	Total	C	N	O	0	0	0
			462	289	100	73			
65	n9	58	Total	C	N	O	0	0	0
			462	289	100	73			

- Molecule 66 is a protein called 60S ribosomal protein L30.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
66	O0	97	Total	C	N	O	S	0	0	0
			743	479	124	139	1			
66	o0	100	Total	C	N	O	S	0	0	0
			767	492	128	146	1			

- Molecule 67 is a protein called 60S ribosomal protein L31-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
67	O1	109	Total	C	N	O	S	0	0	0
			876	556	167	152	1			
67	o1	109	Total	C	N	O	S	0	0	0
			883	559	167	156	1			

- Molecule 68 is a protein called 60S ribosomal protein L32.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
68	O2	127	Total	C	N	O	S	0	0	0
			1020	647	205	167	1			
68	o2	127	Total	C	N	O	S	0	0	0
			1020	647	205	167	1			

- Molecule 69 is a protein called 60S ribosomal protein L33-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
69	O3	106	Total	C	N	O	S	0	0	0
			850	540	165	144	1			
69	o3	106	Total	C	N	O	S	0	0	0
			850	540	165	144	1			

- Molecule 70 is a protein called 60S ribosomal protein L34-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
70	O4	112	Total	C	N	O	S	0	0	0
			880	545	179	152	4			
70	o4	112	Total	C	N	O	S	0	0	0
			880	545	179	152	4			

There are 22 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
O4	110	GLU	-	expression tag	UNP P87262
O4	111	ALA	-	expression tag	UNP P87262
O4	112	ALA	-	expression tag	UNP P87262

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Chain	Residue	Modelled	Actual	Comment	Reference
O4	113	LYS	-	expression tag	UNP P87262
O4	114	SER	-	expression tag	UNP P87262
O4	115	GLU	-	expression tag	UNP P87262
O4	116	LYS	-	expression tag	UNP P87262
O4	117	LYS	-	expression tag	UNP P87262
O4	118	ALA	-	expression tag	UNP P87262
O4	119	LYS	-	expression tag	UNP P87262
O4	120	LYS	-	expression tag	UNP P87262
o4	110	GLU	-	expression tag	UNP P87262
o4	111	ALA	-	expression tag	UNP P87262
o4	112	ALA	-	expression tag	UNP P87262
o4	113	LYS	-	expression tag	UNP P87262
o4	114	SER	-	expression tag	UNP P87262
o4	115	GLU	-	expression tag	UNP P87262
o4	116	LYS	-	expression tag	UNP P87262
o4	117	LYS	-	expression tag	UNP P87262
o4	118	ALA	-	expression tag	UNP P87262
o4	119	LYS	-	expression tag	UNP P87262
o4	120	LYS	-	expression tag	UNP P87262

- Molecule 71 is a protein called 60S ribosomal protein L35-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
71	O5	119	Total	C	N	O	S	0	0	0
			969	615	186	167	1			
71	o5	119	Total	C	N	O	S	0	0	0
			965	612	185	167	1			

- Molecule 72 is a protein called 60S ribosomal protein L36-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
72	O6	99	Total	C	N	O	S	0	0	0
			771	481	156	132	2			
72	o6	99	Total	C	N	O	S	0	0	0
			770	481	156	131	2			

- Molecule 73 is a protein called 60S ribosomal protein L37-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
73	O7	87	Total	C	N	O	S	0	0	0
			681	414	148	114	5			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
73	o7	87	Total	C	N	O	S	0	0	0
			681	414	148	114	5			

- Molecule 74 is a protein called 60S ribosomal protein L38.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
74	O8	77	Total	C	N	O		0	0	0
			612	391	115	106				
74	o8	77	Total	C	N	O		0	0	0
			608	388	114	106				

- Molecule 75 is a protein called 60S ribosomal protein L39.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
75	O9	50	Total	C	N	O	S	0	0	0
			436	272	97	65	2			
75	o9	50	Total	C	N	O	S	0	0	0
			436	272	97	65	2			

- Molecule 76 is a protein called Ubiquitin-60S ribosomal protein L40.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
76	Q0	52	Total	C	N	O	S	0	0	0
			417	259	86	67	5			
76	q0	52	Total	C	N	O	S	0	0	0
			417	259	86	67	5			

- Molecule 77 is a protein called 60S ribosomal protein L41-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
77	Q1	25	Total	C	N	O	S	0	0	0
			233	142	63	27	1			
77	q1	25	Total	C	N	O	S	0	0	0
			233	142	63	27	1			

- Molecule 78 is a protein called 60S ribosomal protein L42-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
78	Q2	105	Total	C	N	O	S	0	0	0
			847	534	170	138	5			
78	q2	105	Total	C	N	O	S	0	0	0
			847	534	170	138	5			

- Molecule 79 is a protein called 60S ribosomal protein L43-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
79	Q3	91	Total	C	N	O	S	0	0	0
			694	429	138	121	6			
79	q3	91	Total	C	N	O	S	0	0	0
			694	429	138	121	6			

- Molecule 80 is a protein called 40S ribosomal protein S30-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
80	e0	62	Total	C	N	O	S	0	0	0
			491	309	101	80	1			

- Molecule 81 is a protein called Unknown protein m2.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
81	m2	150	Total	C	N	O	0	0	0
			750	450	150	150			

- Molecule 82 is a protein called 60S acidic ribosomal protein P0.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
82	p0	143	Total	C	N	O	S	0	0	0
			1076	686	192	195	3			

- Molecule 83 is a protein called Unknown protein p1.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
83	p1	47	Total	C	N	O	0	0	0
			235	141	47	47			

- Molecule 84 is a protein called Unknown protein p2.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
84	p2	46	Total	C	N	O	0	0	0
			230	138	46	46			

- Molecule 85 is MAGNESIUM ION (CCD ID: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
85	2	121	Total	Mg	0	0
			121	121		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
85	S2	2	Total 2	Mg 2	0	0
85	S4	1	Total 1	Mg 1	0	0
85	S6	1	Total 1	Mg 1	0	0
85	S8	1	Total 1	Mg 1	0	0
85	D0	1	Total 1	Mg 1	0	0
85	D3	1	Total 1	Mg 1	0	0
85	1	471	Total 471	Mg 471	0	0
85	3	13	Total 13	Mg 13	0	0
85	4	23	Total 23	Mg 23	0	0
85	L2	2	Total 2	Mg 2	0	0
85	L3	3	Total 3	Mg 3	0	0
85	L4	2	Total 2	Mg 2	0	0
85	L5	1	Total 1	Mg 1	0	0
85	L7	3	Total 3	Mg 3	0	0
85	L8	1	Total 1	Mg 1	0	0
85	M0	3	Total 3	Mg 3	0	0
85	M1	2	Total 2	Mg 2	0	0
85	M3	3	Total 3	Mg 3	0	0
85	M5	1	Total 1	Mg 1	0	0
85	M6	1	Total 1	Mg 1	0	0
85	M7	4	Total 4	Mg 4	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
85	M9	2	Total 2	Mg 2	0	0
85	N0	1	Total 1	Mg 1	0	0
85	N3	3	Total 3	Mg 3	0	0
85	N5	1	Total 1	Mg 1	0	0
85	N6	1	Total 1	Mg 1	0	0
85	N8	3	Total 3	Mg 3	0	0
85	O3	1	Total 1	Mg 1	0	0
85	O4	1	Total 1	Mg 1	0	0
85	O7	2	Total 2	Mg 2	0	0
85	Q2	1	Total 1	Mg 1	0	0
85	6	145	Total 145	Mg 145	0	0
85	s1	1	Total 1	Mg 1	0	0
85	s2	1	Total 1	Mg 1	0	0
85	s4	1	Total 1	Mg 1	0	0
85	s8	2	Total 2	Mg 2	0	0
85	c1	1	Total 1	Mg 1	0	0
85	c7	1	Total 1	Mg 1	0	0
85	c8	2	Total 2	Mg 2	0	0
85	c9	1	Total 1	Mg 1	0	0
85	d3	2	Total 2	Mg 2	0	0
85	d6	1	Total 1	Mg 1	0	0

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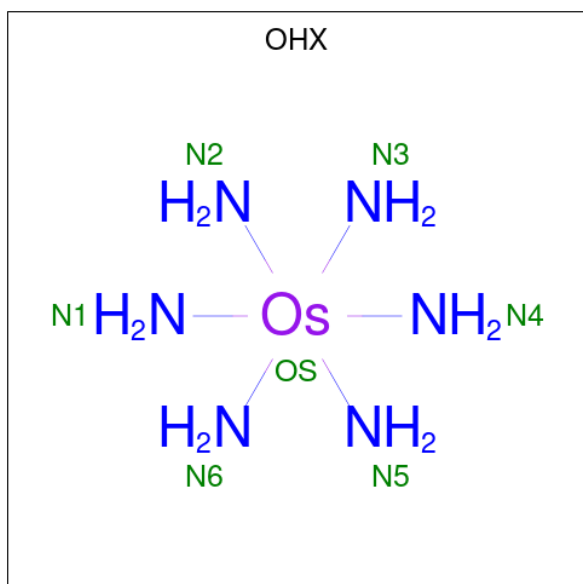
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
85	sM	2	Total 2	Mg 2	0	0
85	5	497	Total 497	Mg 497	0	0
85	7	16	Total 16	Mg 16	0	0
85	8	15	Total 15	Mg 15	0	0
85	12	2	Total 2	Mg 2	0	0
85	l3	3	Total 3	Mg 3	0	0
85	14	1	Total 1	Mg 1	0	0
85	15	2	Total 2	Mg 2	0	0
85	17	3	Total 3	Mg 3	0	0
85	18	1	Total 1	Mg 1	0	0
85	19	1	Total 1	Mg 1	0	0
85	m1	1	Total 1	Mg 1	0	0
85	m4	1	Total 1	Mg 1	0	0
85	m5	3	Total 3	Mg 3	0	0
85	m6	2	Total 2	Mg 2	0	0
85	m7	5	Total 5	Mg 5	0	0
85	n0	1	Total 1	Mg 1	0	0
85	n3	2	Total 2	Mg 2	0	0
85	n6	1	Total 1	Mg 1	0	0
85	n8	5	Total 5	Mg 5	0	0
85	o1	1	Total 1	Mg 1	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
85	o3	2	Total	Mg	0	0
			2	2		
85	o4	1	Total	Mg	0	0
			1	1		
85	o7	1	Total	Mg	0	0
			1	1		
85	q0	1	Total	Mg	0	0
			1	1		
85	q1	1	Total	Mg	0	0
			1	1		
85	q3	2	Total	Mg	0	0
			2	2		

- Molecule 86 is osmium (III) hexammine (CCD ID: OHX) (formula: $\text{H}_{12}\text{N}_6\text{Os}$).



Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
			7	6	1		
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			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	1	1	Total	N	Os	0	0
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			7	6	1		
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86	3	1	Total	N	Os	0	0
			7	6	1		
86	3	1	Total	N	Os	0	0
			7	6	1		
86	4	1	Total	N	Os	0	0
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86	4	1	Total	N	Os	0	0
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			7	6	1		
86	4	1	Total	N	Os	0	0
			7	6	1		
86	4	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	4	1	Total	N	Os	0	0
			7	6	1		
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86	4	1	Total	N	Os	0	0
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86	4	1	Total	N	Os	0	0
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86	4	1	Total	N	Os	0	0
			7	6	1		
86	4	1	Total	N	Os	0	0
			7	6	1		
86	4	1	Total	N	Os	0	0
			7	6	1		
86	L3	1	Total	N	Os	0	0
			7	6	1		
86	L3	1	Total	N	Os	0	0
			7	6	1		
86	L3	1	Total	N	Os	0	0
			7	6	1		
86	L4	1	Total	N	Os	0	0
			7	6	1		
86	M0	1	Total	N	Os	0	0
			7	6	1		
86	M5	1	Total	N	Os	0	0
			7	6	1		
86	M7	1	Total	N	Os	0	0
			7	6	1		
86	M7	1	Total	N	Os	0	0
			7	6	1		
86	M9	1	Total	N	Os	0	0
			7	6	1		
86	N1	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	N9	1	Total	N	Os	0	0
			7	6	1		
86	O1	1	Total	N	Os	0	0
			7	6	1		
86	O2	1	Total	N	Os	0	0
			7	6	1		
86	O3	1	Total	N	Os	0	0
			7	6	1		
86	O7	1	Total	N	Os	0	0
			7	6	1		
86	O7	1	Total	N	Os	0	0
			7	6	1		
86	Q2	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
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			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
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			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
			7	6	1		
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			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
			7	6	1		
86	6	1	Total	N	Os	0	0
			7	6	1		
86	s1	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	s1	1	Total 7	N 6	Os 1	0	0
86	s4	1	Total 7	N 6	Os 1	0	0
86	s8	1	Total 7	N 6	Os 1	0	0
86	c3	1	Total 7	N 6	Os 1	0	0
86	c5	1	Total 7	N 6	Os 1	0	0
86	c8	1	Total 7	N 6	Os 1	0	0
86	d4	1	Total 7	N 6	Os 1	0	0
86	d9	1	Total 7	N 6	Os 1	0	0
86	sR	1	Total 7	N 6	Os 1	0	0
86	5	1	Total 7	N 6	Os 1	0	0
86	5	1	Total 7	N 6	Os 1	0	0
86	5	1	Total 7	N 6	Os 1	0	0
86	5	1	Total 7	N 6	Os 1	0	0
86	5	1	Total 7	N 6	Os 1	0	0
86	5	1	Total 7	N 6	Os 1	0	0
86	5	1	Total 7	N 6	Os 1	0	0
86	5	1	Total 7	N 6	Os 1	0	0
86	5	1	Total 7	N 6	Os 1	0	0
86	5	1	Total 7	N 6	Os 1	0	0
86	5	1	Total 7	N 6	Os 1	0	0
86	5	1	Total 7	N 6	Os 1	0	0

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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			7	6	1		
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			7	6	1		
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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			7	6	1		
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			7	6	1		
86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	5	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	5	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	5	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	8	1	Total	N	Os	0	0
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86	8	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	13	1	Total	N	Os	0	0
			7	6	1		
86	14	1	Total	N	Os	0	0
			7	6	1		
86	14	1	Total	N	Os	0	0
			7	6	1		
86	15	1	Total	N	Os	0	0
			7	6	1		
86	15	1	Total	N	Os	0	0
			7	6	1		
86	15	1	Total	N	Os	0	0
			7	6	1		
86	15	1	Total	N	Os	0	0
			7	6	1		
86	19	1	Total	N	Os	0	0
			7	6	1		
86	m0	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	m0	1	Total	N	Os	0	0
			7	6	1		
86	m1	1	Total	N	Os	0	0
			7	6	1		
86	m4	1	Total	N	Os	0	0
			7	6	1		
86	m5	1	Total	N	Os	0	0
			7	6	1		
86	m5	1	Total	N	Os	0	0
			7	6	1		
86	m6	1	Total	N	Os	0	0
			7	6	1		
86	m7	1	Total	N	Os	0	0
			7	6	1		
86	m8	1	Total	N	Os	0	0
			7	6	1		
86	n3	1	Total	N	Os	0	0
			7	6	1		
86	n9	1	Total	N	Os	0	0
			7	6	1		
86	o2	1	Total	N	Os	0	0
			7	6	1		
86	o3	1	Total	N	Os	0	0
			7	6	1		
86	o7	1	Total	N	Os	0	0
			7	6	1		
86	q1	1	Total	N	Os	0	0
			7	6	1		
86	q2	1	Total	N	Os	0	0
			7	6	1		

- Molecule 87 is ZINC ION (CCD ID: ZN) (formula: Zn).

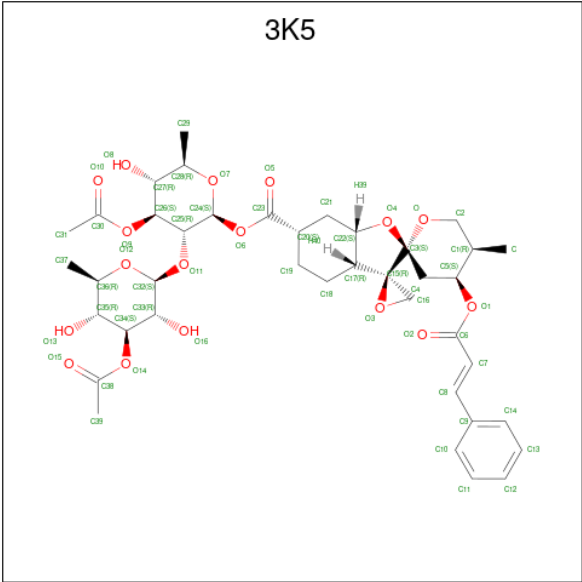
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
87	D6	1	Total	Zn	0	0
			1	1		
87	D7	1	Total	Zn	0	0
			1	1		
87	D9	1	Total	Zn	0	0
			1	1		
87	E1	1	Total	Zn	0	0
			1	1		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
87	O7	1	Total	Zn	0	0
			1	1		
87	Q0	1	Total	Zn	0	0
			1	1		
87	Q2	1	Total	Zn	0	0
			1	1		
87	Q3	1	Total	Zn	0	0
			1	1		
87	d6	1	Total	Zn	0	0
			1	1		
87	d7	1	Total	Zn	0	0
			1	1		
87	d9	1	Total	Zn	0	0
			1	1		
87	e1	1	Total	Zn	0	0
			1	1		
87	o7	1	Total	Zn	0	0
			1	1		
87	q0	1	Total	Zn	0	0
			1	1		
87	q2	1	Total	Zn	0	0
			1	1		
87	q3	1	Total	Zn	0	0
			1	1		

- Molecule 88 is 3-O-acetyl-2-O-(3-O-acetyl-6-deoxy-beta-D-glucopyranosyl)-6-deoxy-1-O-([(2R,2'S,3a'R,4'S,5''R,6'S,7a'S)-5''-methyl-4''-{[(2E)-3-phenylprop-2-enoyl]oxy}decahydrodispiro[oxirane-2,3'-[1]benzofuran-2',2''-pyran]-6'-yl]carbonyl)-beta-D-glucopyranose (CCD ID: 3K5) (formula: C₄₀H₅₂O₁₇).



Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
88	1	1	Total	C	O	0	0
			57	40	17		
88	5	1	Total	C	O	0	0
			57	40	17		

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

3 Data and refinement statistics

Property	Value	Source
Space group	P 1 21 1	Depositor
Cell constants a, b, c, α , β , γ	434.98Å 287.50Å 303.22Å 90.00° 98.85° 90.00°	Depositor
Resolution (Å)	299.60 – 3.10 299.60 – 3.10	Depositor EDS
% Data completeness (in resolution range)	100.0 (299.60-3.10) 99.9 (299.60-3.10)	Depositor EDS
R_{merge}	0.37	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.22 (at 3.07Å)	Xtriage
Refinement program	PHENIX (phenix.refine: dev_1702)	Depositor
R, R_{free}	0.202 , 0.252 0.203 , 0.251	Depositor DCC
R_{free} test set	26549 reflections (2.00%)	wwPDB-VP
Wilson B-factor (Å ²)	76.9	Xtriage
Anisotropy	0.178	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.39 , 130.8	EDS
L-test for twinning ²	$\langle L \rangle = 0.48$, $\langle L^2 \rangle = 0.30$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
F_o, F_c correlation	0.90	EDS
Total number of atoms	411276	wwPDB-VP
Average B, all atoms (Å ²)	76.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.54% of the height of the origin peak. No significant pseudotranslation is detected.*

¹Intensities estimated from amplitudes.

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

4 Model quality [i](#)

4.1 Standard geometry [i](#)

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

4.2 Too-close contacts [i](#)

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

4.3 Torsion angles [i](#)

4.3.1 Protein backbone [i](#)

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

4.3.2 Protein sidechains [i](#)

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

4.3.3 RNA [i](#)

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

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

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

4.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

4.6 Ligand geometry [i](#)

Of 2559 ligands modelled in this entry, 1424 are monoatomic - leaving 1135 for Mogul analysis.

In the following table, the Counts columns list the number of bonds (or angles) for which Mogul statistics could be retrieved, the number of bonds (or angles) that are observed in the model and the number of bonds (or angles) that are defined in the Chemical Component Dictionary. The Link column lists molecule types, if any, to which the group is linked. The Z score for a bond

length (or angle) is the number of standard deviations the observed value is removed from the expected value. A bond length (or angle) with $|Z| > 2$ is considered an outlier worth inspection. RMSZ is the root-mean-square of all Z scores of the bond lengths (or angles).

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	6	2059	-	0,6,6	-	-	-		
86	OHX	1	4003	-	0,6,6	-	-	-		
86	OHX	2	2051	-	0,6,6	-	-	-		
86	OHX	6	2074	-	0,6,6	-	-	-		
86	OHX	S8	302	-	0,6,6	-	-	-		
86	OHX	5	3960	-	0,6,6	-	-	-		
86	OHX	13	404	-	0,6,6	-	-	-		
86	OHX	5	4214	-	0,6,6	-	-	-		
86	OHX	8	225	-	0,6,6	-	-	-		
86	OHX	1	4213	-	0,6,6	-	-	-		
86	OHX	1	4207	-	0,6,6	-	-	-		
86	OHX	1	4115	-	0,6,6	-	-	-		
86	OHX	6	2082	-	0,6,6	-	-	-		
86	OHX	1	4028	-	0,6,6	-	-	-		
86	OHX	6	2105	-	0,6,6	-	-	-		
86	OHX	5	4141	-	0,6,6	-	-	-		
86	OHX	5	4080	-	0,6,6	-	-	-		
86	OHX	5	4149	-	0,6,6	-	-	-		
86	OHX	1	3893	-	0,6,6	-	-	-		
86	OHX	2	2135	-	0,6,6	-	-	-		
86	OHX	5	4165	-	0,6,6	-	-	-		
86	OHX	5	4040	-	0,6,6	-	-	-		
86	OHX	6	2122	-	0,6,6	-	-	-		
86	OHX	5	3963	-	0,6,6	-	-	-		
86	OHX	5	4100	-	0,6,6	-	-	-		
86	OHX	1	4195	-	0,6,6	-	-	-		
86	OHX	5	4228	-	0,6,6	-	-	-		
86	OHX	5	3906	-	0,6,6	-	-	-		
86	OHX	5	4126	-	0,6,6	-	-	-		
86	OHX	5	4130	-	0,6,6	-	-	-		
86	OHX	1	4008	-	0,6,6	-	-	-		
86	OHX	6	2196	-	0,6,6	-	-	-		
86	OHX	5	4180	-	0,6,6	-	-	-		
86	OHX	1	4154	-	0,6,6	-	-	-		
86	OHX	4	229	-	0,6,6	-	-	-		
86	OHX	2	2045	-	0,6,6	-	-	-		
86	OHX	3	222	-	0,6,6	-	-	-		
86	OHX	6	2144	-	0,6,6	-	-	-		
86	OHX	2	2060	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	1	4211	-	0,6,6	-	-	-		
86	OHX	2	2025	-	0,6,6	-	-	-		
86	OHX	5	3940	-	0,6,6	-	-	-		
86	OHX	q1	102	-	0,6,6	-	-	-		
86	OHX	1	4052	-	0,6,6	-	-	-		
86	OHX	1	4050	-	0,6,6	-	-	-		
86	OHX	1	4155	-	0,6,6	-	-	-		
86	OHX	1	4040	-	0,6,6	-	-	-		
86	OHX	6	2114	-	0,6,6	-	-	-		
86	OHX	5	4230	-	0,6,6	-	-	-		
86	OHX	5	4003	-	0,6,6	-	-	-		
86	OHX	8	226	-	0,6,6	-	-	-		
86	OHX	l3	405	-	0,6,6	-	-	-		
86	OHX	6	2125	-	0,6,6	-	-	-		
86	OHX	1	3931	-	0,6,6	-	-	-		
86	OHX	1	4180	-	0,6,6	-	-	-		
86	OHX	L3	406	-	0,6,6	-	-	-		
86	OHX	5	4057	-	0,6,6	-	-	-		
86	OHX	2	2034	-	0,6,6	-	-	-		
86	OHX	1	4048	-	0,6,6	-	-	-		
86	OHX	5	3924	-	0,6,6	-	-	-		
86	OHX	2	2155	-	0,6,6	-	-	-		
86	OHX	6	2187	-	0,6,6	-	-	-		
86	OHX	1	3884	-	0,6,6	-	-	-		
86	OHX	5	4218	-	0,6,6	-	-	-		
86	OHX	6	2198	-	0,6,6	-	-	-		
86	OHX	1	4184	-	0,6,6	-	-	-		
86	OHX	5	3901	-	0,6,6	-	-	-		
86	OHX	5	4039	-	0,6,6	-	-	-		
86	OHX	7	222	-	0,6,6	-	-	-		
86	OHX	5	3975	-	0,6,6	-	-	-		
86	OHX	6	2186	-	0,6,6	-	-	-		
86	OHX	5	4084	-	0,6,6	-	-	-		
86	OHX	1	4044	-	0,6,6	-	-	-		
86	OHX	5	4070	-	0,6,6	-	-	-		
86	OHX	6	2055	-	0,6,6	-	-	-		
86	OHX	5	4113	-	0,6,6	-	-	-		
86	OHX	2	2030	-	0,6,6	-	-	-		
86	OHX	6	2048	-	0,6,6	-	-	-		
86	OHX	1	3894	-	0,6,6	-	-	-		
86	OHX	6	2095	-	0,6,6	-	-	-		
86	OHX	5	4222	-	0,6,6	-	-	-		
86	OHX	1	4168	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	1	4047	-	0,6,6	-	-	-		
86	OHX	1	4090	-	0,6,6	-	-	-		
86	OHX	m5	305	-	0,6,6	-	-	-		
86	OHX	6	2172	-	0,6,6	-	-	-		
86	OHX	1	3919	-	0,6,6	-	-	-		
86	OHX	5	3925	-	0,6,6	-	-	-		
86	OHX	1	4020	-	0,6,6	-	-	-		
86	OHX	1	4216	-	0,6,6	-	-	-		
86	OHX	2	2110	-	0,6,6	-	-	-		
86	OHX	2	2062	-	0,6,6	-	-	-		
86	OHX	6	2134	-	0,6,6	-	-	-		
86	OHX	5	4140	-	0,6,6	-	-	-		
86	OHX	1	3956	-	0,6,6	-	-	-		
86	OHX	1	3904	-	0,6,6	-	-	-		
86	OHX	6	2174	-	0,6,6	-	-	-		
86	OHX	6	2177	-	0,6,6	-	-	-		
86	OHX	1	3889	-	0,6,6	-	-	-		
86	OHX	5	4092	-	0,6,6	-	-	-		
86	OHX	6	2119	-	0,6,6	-	-	-		
86	OHX	1	3954	-	0,6,6	-	-	-		
86	OHX	1	3897	-	0,6,6	-	-	-		
86	OHX	5	3930	-	0,6,6	-	-	-		
86	OHX	6	2190	-	0,6,6	-	-	-		
86	OHX	1	3914	-	0,6,6	-	-	-		
86	OHX	2	2023	-	0,6,6	-	-	-		
86	OHX	1	3945	-	0,6,6	-	-	-		
86	OHX	6	2176	-	0,6,6	-	-	-		
86	OHX	1	4124	-	0,6,6	-	-	-		
86	OHX	2	2079	-	0,6,6	-	-	-		
86	OHX	2	2114	-	0,6,6	-	-	-		
86	OHX	5	3923	-	0,6,6	-	-	-		
86	OHX	5	4043	-	0,6,6	-	-	-		
86	OHX	5	4050	-	0,6,6	-	-	-		
86	OHX	5	4242	-	0,6,6	-	-	-		
86	OHX	2	2061	-	0,6,6	-	-	-		
86	OHX	2	2058	-	0,6,6	-	-	-		
86	OHX	1	3983	-	0,6,6	-	-	-		
86	OHX	1	3998	-	0,6,6	-	-	-		
86	OHX	1	3930	-	0,6,6	-	-	-		
86	OHX	1	3934	-	0,6,6	-	-	-		
86	OHX	1	4055	-	0,6,6	-	-	-		
86	OHX	6	2200	-	0,6,6	-	-	-		
86	OHX	2	2040	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	M9	203	-	0,6,6	-	-	-		
86	OHX	1	4013	-	0,6,6	-	-	-		
86	OHX	1	4033	-	0,6,6	-	-	-		
86	OHX	2	2106	-	0,6,6	-	-	-		
86	OHX	5	4169	-	0,6,6	-	-	-		
86	OHX	1	3917	-	0,6,6	-	-	-		
86	OHX	5	3973	-	0,6,6	-	-	-		
86	OHX	5	4089	-	0,6,6	-	-	-		
86	OHX	6	2115	-	0,6,6	-	-	-		
86	OHX	6	2153	-	0,6,6	-	-	-		
86	OHX	5	4076	-	0,6,6	-	-	-		
86	OHX	5	4108	-	0,6,6	-	-	-		
86	OHX	5	4047	-	0,6,6	-	-	-		
86	OHX	5	3964	-	0,6,6	-	-	-		
86	OHX	6	2137	-	0,6,6	-	-	-		
86	OHX	5	4176	-	0,6,6	-	-	-		
86	OHX	1	3990	-	0,6,6	-	-	-		
86	OHX	M5	302	-	0,6,6	-	-	-		
86	OHX	1	4158	-	0,6,6	-	-	-		
86	OHX	5	4066	-	0,6,6	-	-	-		
86	OHX	2	2161	-	0,6,6	-	-	-		
86	OHX	1	4114	-	0,6,6	-	-	-		
86	OHX	1	4134	-	0,6,6	-	-	-		
86	OHX	5	4001	-	0,6,6	-	-	-		
86	OHX	l5	305	-	0,6,6	-	-	-		
86	OHX	5	4121	-	0,6,6	-	-	-		
86	OHX	6	2173	-	0,6,6	-	-	-		
86	OHX	6	2180	-	0,6,6	-	-	-		
86	OHX	L3	405	-	0,6,6	-	-	-		
86	OHX	N1	201	-	0,6,6	-	-	-		
86	OHX	2	2050	-	0,6,6	-	-	-		
86	OHX	1	3872	-	0,6,6	-	-	-		
86	OHX	5	4189	-	0,6,6	-	-	-		
86	OHX	2	2107	-	0,6,6	-	-	-		
86	OHX	2	2032	-	0,6,6	-	-	-		
86	OHX	2	2069	-	0,6,6	-	-	-		
86	OHX	5	3998	-	0,6,6	-	-	-		
86	OHX	5	4017	-	0,6,6	-	-	-		
86	OHX	6	2178	-	0,6,6	-	-	-		
86	OHX	1	3962	-	0,6,6	-	-	-		
86	OHX	5	4011	-	0,6,6	-	-	-		
86	OHX	1	3976	-	0,6,6	-	-	-		
86	OHX	5	4106	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	5	4135	-	0,6,6	-	-	-		
86	OHX	D3	202	-	0,6,6	-	-	-		
86	OHX	1	3901	-	0,6,6	-	-	-		
86	OHX	6	2118	-	0,6,6	-	-	-		
86	OHX	1	3874	-	0,6,6	-	-	-		
86	OHX	6	2096	-	0,6,6	-	-	-		
86	OHX	5	4028	-	0,6,6	-	-	-		
86	OHX	5	3912	-	0,6,6	-	-	-		
86	OHX	n3	203	-	0,6,6	-	-	-		
86	OHX	1	3991	-	0,6,6	-	-	-		
86	OHX	1	4084	-	0,6,6	-	-	-		
86	OHX	2	2074	-	0,6,6	-	-	-		
86	OHX	1	4120	-	0,6,6	-	-	-		
86	OHX	1	4131	-	0,6,6	-	-	-		
86	OHX	1	4182	-	0,6,6	-	-	-		
86	OHX	1	4189	-	0,6,6	-	-	-		
86	OHX	2	2097	-	0,6,6	-	-	-		
86	OHX	1	4178	-	0,6,6	-	-	-		
86	OHX	6	2145	-	0,6,6	-	-	-		
86	OHX	6	2168	-	0,6,6	-	-	-		
86	OHX	1	3961	-	0,6,6	-	-	-		
86	OHX	1	4101	-	0,6,6	-	-	-		
86	OHX	6	2171	-	0,6,6	-	-	-		
86	OHX	5	4209	-	0,6,6	-	-	-		
86	OHX	1	3895	-	0,6,6	-	-	-		
86	OHX	5	4123	-	0,6,6	-	-	-		
86	OHX	8	218	-	0,6,6	-	-	-		
86	OHX	5	4012	-	0,6,6	-	-	-		
86	OHX	1	4073	-	0,6,6	-	-	-		
86	OHX	M7	205	-	0,6,6	-	-	-		
86	OHX	1	4140	-	0,6,6	-	-	-		
86	OHX	1	3965	-	0,6,6	-	-	-		
86	OHX	5	3915	-	0,6,6	-	-	-		
86	OHX	5	4205	-	0,6,6	-	-	-		
86	OHX	1	4002	-	0,6,6	-	-	-		
86	OHX	5	4194	-	0,6,6	-	-	-		
86	OHX	1	4006	-	0,6,6	-	-	-		
86	OHX	1	3939	-	0,6,6	-	-	-		
86	OHX	2	2028	-	0,6,6	-	-	-		
86	OHX	5	4013	-	0,6,6	-	-	-		
86	OHX	8	229	-	0,6,6	-	-	-		
86	OHX	1	4177	-	0,6,6	-	-	-		
86	OHX	1	4063	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	1	4141	-	0,6,6	-	-	-		
86	OHX	1	4102	-	0,6,6	-	-	-		
86	OHX	1	4148	-	0,6,6	-	-	-		
86	OHX	1	3938	-	0,6,6	-	-	-		
86	OHX	4	225	-	0,6,6	-	-	-		
86	OHX	5	4079	-	0,6,6	-	-	-		
86	OHX	4	228	-	0,6,6	-	-	-		
86	OHX	5	4103	-	0,6,6	-	-	-		
86	OHX	14	403	-	0,6,6	-	-	-		
86	OHX	6	2085	-	0,6,6	-	-	-		
86	OHX	5	4131	-	0,6,6	-	-	-		
86	OHX	5	4022	-	0,6,6	-	-	-		
86	OHX	1	4193	-	0,6,6	-	-	-		
86	OHX	5	4142	-	0,6,6	-	-	-		
86	OHX	5	3989	-	0,6,6	-	-	-		
86	OHX	5	4221	-	0,6,6	-	-	-		
86	OHX	5	4232	-	0,6,6	-	-	-		
86	OHX	7	218	-	0,6,6	-	-	-		
86	OHX	3	219	-	0,6,6	-	-	-		
86	OHX	5	3962	-	0,6,6	-	-	-		
86	OHX	5	4200	-	0,6,6	-	-	-		
86	OHX	5	4024	-	0,6,6	-	-	-		
86	OHX	2	2177	-	0,6,6	-	-	-		
86	OHX	2	2154	-	0,6,6	-	-	-		
86	OHX	5	4077	-	0,6,6	-	-	-		
86	OHX	5	4212	-	0,6,6	-	-	-		
86	OHX	2	2048	-	0,6,6	-	-	-		
86	OHX	1	4133	-	0,6,6	-	-	-		
86	OHX	1	4061	-	0,6,6	-	-	-		
86	OHX	5	3993	-	0,6,6	-	-	-		
86	OHX	6	2064	-	0,6,6	-	-	-		
86	OHX	6	2136	-	0,6,6	-	-	-		
86	OHX	5	4038	-	0,6,6	-	-	-		
86	OHX	5	4093	-	0,6,6	-	-	-		
86	OHX	5	4107	-	0,6,6	-	-	-		
86	OHX	1	4142	-	0,6,6	-	-	-		
86	OHX	2	2131	-	0,6,6	-	-	-		
86	OHX	5	3957	-	0,6,6	-	-	-		
86	OHX	1	4171	-	0,6,6	-	-	-		
86	OHX	5	4027	-	0,6,6	-	-	-		
86	OHX	5	4102	-	0,6,6	-	-	-		
86	OHX	5	4148	-	0,6,6	-	-	-		
86	OHX	5	4223	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	5	4088	-	0,6,6	-	-	-		
86	OHX	6	2188	-	0,6,6	-	-	-		
86	OHX	m8	201	-	0,6,6	-	-	-		
86	OHX	5	3920	-	0,6,6	-	-	-		
86	OHX	1	3986	-	0,6,6	-	-	-		
86	OHX	2	2171	-	0,6,6	-	-	-		
86	OHX	Q2	503	-	0,6,6	-	-	-		
86	OHX	6	2157	-	0,6,6	-	-	-		
86	OHX	1	4039	-	0,6,6	-	-	-		
86	OHX	5	3978	-	0,6,6	-	-	-		
86	OHX	1	4188	-	0,6,6	-	-	-		
86	OHX	1	4026	-	0,6,6	-	-	-		
86	OHX	4	234	-	0,6,6	-	-	-		
86	OHX	6	2201	-	0,6,6	-	-	-		
86	OHX	4	226	-	0,6,6	-	-	-		
86	OHX	5	3935	-	0,6,6	-	-	-		
86	OHX	2	2036	-	0,6,6	-	-	-		
86	OHX	1	3875	-	0,6,6	-	-	-		
86	OHX	1	3911	-	0,6,6	-	-	-		
86	OHX	5	3959	-	0,6,6	-	-	-		
86	OHX	5	3990	-	0,6,6	-	-	-		
86	OHX	5	4037	-	0,6,6	-	-	-		
86	OHX	2	2044	-	0,6,6	-	-	-		
86	OHX	6	2166	-	0,6,6	-	-	-		
86	OHX	5	4072	-	0,6,6	-	-	-		
86	OHX	1	4042	-	0,6,6	-	-	-		
86	OHX	O1	201	-	0,6,6	-	-	-		
86	OHX	5	4041	-	0,6,6	-	-	-		
86	OHX	5	3994	-	0,6,6	-	-	-		
86	OHX	1	4135	-	0,6,6	-	-	-		
86	OHX	5	3911	-	0,6,6	-	-	-		
86	OHX	6	2103	-	0,6,6	-	-	-		
86	OHX	5	3997	-	0,6,6	-	-	-		
86	OHX	6	2120	-	0,6,6	-	-	-		
86	OHX	1	3950	-	0,6,6	-	-	-		
86	OHX	2	2173	-	0,6,6	-	-	-		
86	OHX	5	3914	-	0,6,6	-	-	-		
86	OHX	c5	201	-	0,6,6	-	-	-		
86	OHX	5	3932	-	0,6,6	-	-	-		
86	OHX	5	4014	-	0,6,6	-	-	-		
86	OHX	6	2150	-	0,6,6	-	-	-		
86	OHX	1	4161	-	0,6,6	-	-	-		
86	OHX	5	3909	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	5	3946	-	0,6,6	-	-	-		
86	OHX	2	2026	-	0,6,6	-	-	-		
86	OHX	5	4235	-	0,6,6	-	-	-		
86	OHX	2	2140	-	0,6,6	-	-	-		
86	OHX	2	2098	-	0,6,6	-	-	-		
86	OHX	5	4204	-	0,6,6	-	-	-		
86	OHX	1	4031	-	0,6,6	-	-	-		
86	OHX	4	239	-	0,6,6	-	-	-		
86	OHX	6	2047	-	0,6,6	-	-	-		
86	OHX	6	2051	-	0,6,6	-	-	-		
86	OHX	6	2185	-	0,6,6	-	-	-		
86	OHX	5	4158	-	0,6,6	-	-	-		
86	OHX	1	3942	-	0,6,6	-	-	-		
86	OHX	6	2199	-	0,6,6	-	-	-		
86	OHX	6	2061	-	0,6,6	-	-	-		
86	OHX	5	4119	-	0,6,6	-	-	-		
86	OHX	C8	201	-	0,6,6	-	-	-		
86	OHX	5	4097	-	0,6,6	-	-	-		
86	OHX	2	2130	-	0,6,6	-	-	-		
86	OHX	6	2135	-	0,6,6	-	-	-		
86	OHX	1	4156	-	0,6,6	-	-	-		
86	OHX	19	202	-	0,6,6	-	-	-		
86	OHX	n9	101	-	0,6,6	-	-	-		
86	OHX	1	3960	-	0,6,6	-	-	-		
86	OHX	1	4103	-	0,6,6	-	-	-		
86	OHX	6	2139	-	0,6,6	-	-	-		
86	OHX	6	2158	-	0,6,6	-	-	-		
86	OHX	2	2178	-	0,6,6	-	-	-		
86	OHX	1	4175	-	0,6,6	-	-	-		
86	OHX	1	4029	-	0,6,6	-	-	-		
86	OHX	6	2110	-	0,6,6	-	-	-		
86	OHX	6	2197	-	0,6,6	-	-	-		
86	OHX	1	4022	-	0,6,6	-	-	-		
86	OHX	5	3976	-	0,6,6	-	-	-		
86	OHX	2	2159	-	0,6,6	-	-	-		
86	OHX	6	2091	-	0,6,6	-	-	-		
86	OHX	1	4192	-	0,6,6	-	-	-		
86	OHX	1	4116	-	0,6,6	-	-	-		
86	OHX	5	4184	-	0,6,6	-	-	-		
86	OHX	2	2142	-	0,6,6	-	-	-		
86	OHX	1	3903	-	0,6,6	-	-	-		
86	OHX	6	2088	-	0,6,6	-	-	-		
86	OHX	1	4200	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	5	4032	-	0,6,6	-	-	-		
86	OHX	5	3933	-	0,6,6	-	-	-		
86	OHX	5	3981	-	0,6,6	-	-	-		
86	OHX	7	225	-	0,6,6	-	-	-		
86	OHX	1	4007	-	0,6,6	-	-	-		
86	OHX	5	3967	-	0,6,6	-	-	-		
86	OHX	1	4092	-	0,6,6	-	-	-		
86	OHX	1	3993	-	0,6,6	-	-	-		
86	OHX	1	4079	-	0,6,6	-	-	-		
86	OHX	1	4111	-	0,6,6	-	-	-		
86	OHX	6	2147	-	0,6,6	-	-	-		
86	OHX	5	4210	-	0,6,6	-	-	-		
86	OHX	7	227	-	0,6,6	-	-	-		
86	OHX	5	4065	-	0,6,6	-	-	-		
86	OHX	5	4074	-	0,6,6	-	-	-		
86	OHX	1	4144	-	0,6,6	-	-	-		
86	OHX	1	3891	-	0,6,6	-	-	-		
86	OHX	1	4166	-	0,6,6	-	-	-		
86	OHX	1	3948	-	0,6,6	-	-	-		
86	OHX	1	3963	-	0,6,6	-	-	-		
86	OHX	1	4100	-	0,6,6	-	-	-		
86	OHX	2	2088	-	0,6,6	-	-	-		
86	OHX	2	2120	-	0,6,6	-	-	-		
86	OHX	2	2066	-	0,6,6	-	-	-		
86	OHX	1	4110	-	0,6,6	-	-	-		
86	OHX	6	2077	-	0,6,6	-	-	-		
86	OHX	6	2189	-	0,6,6	-	-	-		
86	OHX	4	224	-	0,6,6	-	-	-		
86	OHX	5	3965	-	0,6,6	-	-	-		
86	OHX	2	2094	-	0,6,6	-	-	-		
86	OHX	5	4030	-	0,6,6	-	-	-		
86	OHX	5	4156	-	0,6,6	-	-	-		
86	OHX	1	3918	-	0,6,6	-	-	-		
88	3K5	5	4249	-	63,63,63	0.65	1 (1%)	84,95,95	1.37	12 (14%)
86	OHX	5	4112	-	0,6,6	-	-	-		
86	OHX	1	3920	-	0,6,6	-	-	-		
86	OHX	5	4182	-	0,6,6	-	-	-		
86	OHX	5	4153	-	0,6,6	-	-	-		
86	OHX	1	4210	-	0,6,6	-	-	-		
86	OHX	1	3878	-	0,6,6	-	-	-		
86	OHX	6	2151	-	0,6,6	-	-	-		
86	OHX	1	4138	-	0,6,6	-	-	-		
86	OHX	1	4202	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	6	2131	-	0,6,6	-	-	-		
86	OHX	2	2083	-	0,6,6	-	-	-		
86	OHX	O3	202	-	0,6,6	-	-	-		
86	OHX	6	2156	-	0,6,6	-	-	-		
86	OHX	1	4128	-	0,6,6	-	-	-		
86	OHX	5	4179	-	0,6,6	-	-	-		
86	OHX	1	3987	-	0,6,6	-	-	-		
86	OHX	5	3936	-	0,6,6	-	-	-		
86	OHX	5	4191	-	0,6,6	-	-	-		
86	OHX	1	3906	-	0,6,6	-	-	-		
86	OHX	2	2164	-	0,6,6	-	-	-		
86	OHX	6	2167	-	0,6,6	-	-	-		
86	OHX	8	228	-	0,6,6	-	-	-		
86	OHX	6	2062	-	0,6,6	-	-	-		
86	OHX	1	3885	-	0,6,6	-	-	-		
86	OHX	1	3964	-	0,6,6	-	-	-		
86	OHX	5	4110	-	0,6,6	-	-	-		
86	OHX	1	3940	-	0,6,6	-	-	-		
86	OHX	1	4205	-	0,6,6	-	-	-		
86	OHX	1	4125	-	0,6,6	-	-	-		
86	OHX	6	2057	-	0,6,6	-	-	-		
86	OHX	5	4056	-	0,6,6	-	-	-		
86	OHX	2	2096	-	0,6,6	-	-	-		
86	OHX	1	3925	-	0,6,6	-	-	-		
86	OHX	1	3880	-	0,6,6	-	-	-		
86	OHX	1	4045	-	0,6,6	-	-	-		
86	OHX	6	2069	-	0,6,6	-	-	-		
86	OHX	6	2102	-	0,6,6	-	-	-		
86	OHX	6	2124	-	0,6,6	-	-	-		
86	OHX	1	3910	-	0,6,6	-	-	-		
86	OHX	6	2128	-	0,6,6	-	-	-		
86	OHX	6	2165	-	0,6,6	-	-	-		
86	OHX	1	4035	-	0,6,6	-	-	-		
86	OHX	6	2193	-	0,6,6	-	-	-		
86	OHX	5	3918	-	0,6,6	-	-	-		
86	OHX	5	4029	-	0,6,6	-	-	-		
86	OHX	2	2100	-	0,6,6	-	-	-		
86	OHX	6	2127	-	0,6,6	-	-	-		
86	OHX	1	3992	-	0,6,6	-	-	-		
86	OHX	D9	102	-	0,6,6	-	-	-		
86	OHX	2	2081	-	0,6,6	-	-	-		
86	OHX	5	4049	-	0,6,6	-	-	-		
86	OHX	5	4069	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	5	4133	-	0,6,6	-	-	-		
86	OHX	8	219	-	0,6,6	-	-	-		
86	OHX	2	2075	-	0,6,6	-	-	-		
86	OHX	1	4157	-	0,6,6	-	-	-		
86	OHX	m5	304	-	0,6,6	-	-	-		
86	OHX	1	4095	-	0,6,6	-	-	-		
86	OHX	7	226	-	0,6,6	-	-	-		
86	OHX	5	4019	-	0,6,6	-	-	-		
86	OHX	5	4144	-	0,6,6	-	-	-		
86	OHX	1	4132	-	0,6,6	-	-	-		
86	OHX	1	4015	-	0,6,6	-	-	-		
86	OHX	5	3947	-	0,6,6	-	-	-		
86	OHX	5	3928	-	0,6,6	-	-	-		
86	OHX	1	4018	-	0,6,6	-	-	-		
86	OHX	1	3899	-	0,6,6	-	-	-		
86	OHX	1	3888	-	0,6,6	-	-	-		
86	OHX	1	4023	-	0,6,6	-	-	-		
86	OHX	5	3910	-	0,6,6	-	-	-		
86	OHX	5	4060	-	0,6,6	-	-	-		
86	OHX	5	3902	-	0,6,6	-	-	-		
86	OHX	m6	203	-	0,6,6	-	-	-		
86	OHX	1	4170	-	0,6,6	-	-	-		
86	OHX	2	2033	-	0,6,6	-	-	-		
86	OHX	2	2109	-	0,6,6	-	-	-		
86	OHX	1	4059	-	0,6,6	-	-	-		
86	OHX	1	3967	-	0,6,6	-	-	-		
86	OHX	2	2054	-	0,6,6	-	-	-		
86	OHX	2	2147	-	0,6,6	-	-	-		
86	OHX	5	4163	-	0,6,6	-	-	-		
86	OHX	6	2205	-	0,6,6	-	-	-		
86	OHX	1	4057	-	0,6,6	-	-	-		
86	OHX	6	2191	-	0,6,6	-	-	-		
86	OHX	1	4174	-	0,6,6	-	-	-		
86	OHX	5	4005	-	0,6,6	-	-	-		
86	OHX	o2	201	-	0,6,6	-	-	-		
86	OHX	2	2090	-	0,6,6	-	-	-		
86	OHX	2	2080	-	0,6,6	-	-	-		
86	OHX	5	3916	-	0,6,6	-	-	-		
86	OHX	6	2143	-	0,6,6	-	-	-		
86	OHX	6	2070	-	0,6,6	-	-	-		
86	OHX	1	4081	-	0,6,6	-	-	-		
86	OHX	1	4153	-	0,6,6	-	-	-		
86	OHX	2	2165	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	6	2086	-	0,6,6	-	-	-		
86	OHX	8	220	-	0,6,6	-	-	-		
86	OHX	5	3977	-	0,6,6	-	-	-		
86	OHX	5	4166	-	0,6,6	-	-	-		
86	OHX	5	4075	-	0,6,6	-	-	-		
86	OHX	5	4008	-	0,6,6	-	-	-		
86	OHX	1	4071	-	0,6,6	-	-	-		
86	OHX	2	2158	-	0,6,6	-	-	-		
86	OHX	5	4225	-	0,6,6	-	-	-		
86	OHX	6	2126	-	0,6,6	-	-	-		
86	OHX	1	4099	-	0,6,6	-	-	-		
86	OHX	5	3969	-	0,6,6	-	-	-		
86	OHX	6	2160	-	0,6,6	-	-	-		
86	OHX	6	2149	-	0,6,6	-	-	-		
86	OHX	6	2146	-	0,6,6	-	-	-		
86	OHX	1	4093	-	0,6,6	-	-	-		
86	OHX	d9	102	-	0,6,6	-	-	-		
86	OHX	2	2118	-	0,6,6	-	-	-		
86	OHX	5	4234	-	0,6,6	-	-	-		
86	OHX	6	2079	-	0,6,6	-	-	-		
86	OHX	1	3887	-	0,6,6	-	-	-		
86	OHX	1	4054	-	0,6,6	-	-	-		
86	OHX	2	2049	-	0,6,6	-	-	-		
86	OHX	2	2055	-	0,6,6	-	-	-		
86	OHX	5	4081	-	0,6,6	-	-	-		
86	OHX	1	4117	-	0,6,6	-	-	-		
86	OHX	6	2152	-	0,6,6	-	-	-		
86	OHX	2	2095	-	0,6,6	-	-	-		
86	OHX	1	3881	-	0,6,6	-	-	-		
86	OHX	1	4088	-	0,6,6	-	-	-		
86	OHX	1	4024	-	0,6,6	-	-	-		
86	OHX	1	4097	-	0,6,6	-	-	-		
86	OHX	1	3952	-	0,6,6	-	-	-		
86	OHX	1	4126	-	0,6,6	-	-	-		
86	OHX	1	4219	-	0,6,6	-	-	-		
86	OHX	1	4077	-	0,6,6	-	-	-		
86	OHX	6	2087	-	0,6,6	-	-	-		
86	OHX	1	3985	-	0,6,6	-	-	-		
86	OHX	2	2072	-	0,6,6	-	-	-		
86	OHX	5	4237	-	0,6,6	-	-	-		
86	OHX	5	4117	-	0,6,6	-	-	-		
86	OHX	3	223	-	0,6,6	-	-	-		
86	OHX	1	4083	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	1	4183	-	0,6,6	-	-	-		
86	OHX	O2	201	-	0,6,6	-	-	-		
86	OHX	5	3903	-	0,6,6	-	-	-		
86	OHX	5	3904	-	0,6,6	-	-	-		
86	OHX	5	4064	-	0,6,6	-	-	-		
86	OHX	1	4011	-	0,6,6	-	-	-		
86	OHX	1	3879	-	0,6,6	-	-	-		
86	OHX	5	3972	-	0,6,6	-	-	-		
86	OHX	2	2082	-	0,6,6	-	-	-		
86	OHX	5	4143	-	0,6,6	-	-	-		
86	OHX	6	2163	-	0,6,6	-	-	-		
86	OHX	6	2117	-	0,6,6	-	-	-		
86	OHX	5	3919	-	0,6,6	-	-	-		
86	OHX	1	4080	-	0,6,6	-	-	-		
86	OHX	5	3970	-	0,6,6	-	-	-		
86	OHX	5	3992	-	0,6,6	-	-	-		
86	OHX	6	2046	-	0,6,6	-	-	-		
86	OHX	5	4188	-	0,6,6	-	-	-		
86	OHX	1	4160	-	0,6,6	-	-	-		
86	OHX	5	4009	-	0,6,6	-	-	-		
86	OHX	5	3941	-	0,6,6	-	-	-		
86	OHX	1	3927	-	0,6,6	-	-	-		
86	OHX	1	4130	-	0,6,6	-	-	-		
86	OHX	6	2084	-	0,6,6	-	-	-		
86	OHX	5	4000	-	0,6,6	-	-	-		
86	OHX	1	4094	-	0,6,6	-	-	-		
86	OHX	5	3907	-	0,6,6	-	-	-		
86	OHX	5	4021	-	0,6,6	-	-	-		
86	OHX	5	4115	-	0,6,6	-	-	-		
86	OHX	6	2133	-	0,6,6	-	-	-		
86	OHX	5	4167	-	0,6,6	-	-	-		
86	OHX	5	4183	-	0,6,6	-	-	-		
86	OHX	2	2126	-	0,6,6	-	-	-		
86	OHX	1	4043	-	0,6,6	-	-	-		
86	OHX	1	4181	-	0,6,6	-	-	-		
86	OHX	1	4163	-	0,6,6	-	-	-		
86	OHX	1	4034	-	0,6,6	-	-	-		
86	OHX	1	4119	-	0,6,6	-	-	-		
86	OHX	1	4215	-	0,6,6	-	-	-		
86	OHX	5	4096	-	0,6,6	-	-	-		
86	OHX	s8	303	-	0,6,6	-	-	-		
86	OHX	2	2125	-	0,6,6	-	-	-		
86	OHX	2	2078	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	1	3892	-	0,6,6	-	-	-		
86	OHX	2	2053	-	0,6,6	-	-	-		
86	OHX	2	2136	-	0,6,6	-	-	-		
86	OHX	4	230	-	0,6,6	-	-	-		
86	OHX	5	3996	-	0,6,6	-	-	-		
86	OHX	2	2073	-	0,6,6	-	-	-		
86	OHX	1	3909	-	0,6,6	-	-	-		
86	OHX	2	2141	-	0,6,6	-	-	-		
86	OHX	5	3934	-	0,6,6	-	-	-		
86	OHX	6	2192	-	0,6,6	-	-	-		
86	OHX	5	4033	-	0,6,6	-	-	-		
86	OHX	2	2059	-	0,6,6	-	-	-		
86	OHX	5	3917	-	0,6,6	-	-	-		
86	OHX	1	4027	-	0,6,6	-	-	-		
86	OHX	5	4061	-	0,6,6	-	-	-		
86	OHX	8	230	-	0,6,6	-	-	-		
86	OHX	5	3952	-	0,6,6	-	-	-		
86	OHX	1	3989	-	0,6,6	-	-	-		
86	OHX	1	3937	-	0,6,6	-	-	-		
86	OHX	6	2116	-	0,6,6	-	-	-		
86	OHX	1	4143	-	0,6,6	-	-	-		
86	OHX	5	4016	-	0,6,6	-	-	-		
86	OHX	1	4078	-	0,6,6	-	-	-		
86	OHX	1	4194	-	0,6,6	-	-	-		
86	OHX	5	4062	-	0,6,6	-	-	-		
86	OHX	2	2122	-	0,6,6	-	-	-		
86	OHX	3	217	-	0,6,6	-	-	-		
86	OHX	5	4146	-	0,6,6	-	-	-		
86	OHX	6	2089	-	0,6,6	-	-	-		
86	OHX	1	3935	-	0,6,6	-	-	-		
86	OHX	5	3937	-	0,6,6	-	-	-		
86	OHX	2	2029	-	0,6,6	-	-	-		
86	OHX	5	3945	-	0,6,6	-	-	-		
86	OHX	2	2144	-	0,6,6	-	-	-		
86	OHX	3	221	-	0,6,6	-	-	-		
86	OHX	6	2099	-	0,6,6	-	-	-		
86	OHX	4	227	-	0,6,6	-	-	-		
86	OHX	6	2132	-	0,6,6	-	-	-		
86	OHX	5	4109	-	0,6,6	-	-	-		
86	OHX	5	4243	-	0,6,6	-	-	-		
86	OHX	8	222	-	0,6,6	-	-	-		
86	OHX	1	4032	-	0,6,6	-	-	-		
86	OHX	1	3969	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	6	2164	-	0,6,6	-	-	-		
86	OHX	O7	104	-	0,6,6	-	-	-		
86	OHX	6	2112	-	0,6,6	-	-	-		
86	OHX	2	2102	-	0,6,6	-	-	-		
86	OHX	15	303	-	0,6,6	-	-	-		
86	OHX	6	2050	-	0,6,6	-	-	-		
86	OHX	5	4094	-	0,6,6	-	-	-		
86	OHX	1	4037	-	0,6,6	-	-	-		
86	OHX	5	4192	-	0,6,6	-	-	-		
86	OHX	2	2150	-	0,6,6	-	-	-		
86	OHX	1	4098	-	0,6,6	-	-	-		
86	OHX	1	3932	-	0,6,6	-	-	-		
86	OHX	1	4041	-	0,6,6	-	-	-		
86	OHX	1	3984	-	0,6,6	-	-	-		
86	OHX	7	219	-	0,6,6	-	-	-		
86	OHX	5	3948	-	0,6,6	-	-	-		
86	OHX	5	4127	-	0,6,6	-	-	-		
86	OHX	1	4204	-	0,6,6	-	-	-		
86	OHX	2	2099	-	0,6,6	-	-	-		
86	OHX	1	4010	-	0,6,6	-	-	-		
86	OHX	6	2100	-	0,6,6	-	-	-		
86	OHX	1	4107	-	0,6,6	-	-	-		
86	OHX	6	2182	-	0,6,6	-	-	-		
86	OHX	5	3958	-	0,6,6	-	-	-		
86	OHX	7	221	-	0,6,6	-	-	-		
86	OHX	2	2162	-	0,6,6	-	-	-		
86	OHX	2	2152	-	0,6,6	-	-	-		
86	OHX	1	3896	-	0,6,6	-	-	-		
86	OHX	2	2149	-	0,6,6	-	-	-		
86	OHX	1	3929	-	0,6,6	-	-	-		
86	OHX	1	4091	-	0,6,6	-	-	-		
86	OHX	1	4068	-	0,6,6	-	-	-		
86	OHX	6	2063	-	0,6,6	-	-	-		
86	OHX	5	4138	-	0,6,6	-	-	-		
86	OHX	1	4089	-	0,6,6	-	-	-		
86	OHX	5	4006	-	0,6,6	-	-	-		
86	OHX	5	4150	-	0,6,6	-	-	-		
86	OHX	6	2113	-	0,6,6	-	-	-		
86	OHX	5	4122	-	0,6,6	-	-	-		
86	OHX	1	4046	-	0,6,6	-	-	-		
86	OHX	5	3951	-	0,6,6	-	-	-		
86	OHX	2	2169	-	0,6,6	-	-	-		
86	OHX	1	4108	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	6	2129	-	0,6,6	-	-	-		
86	OHX	6	2094	-	0,6,6	-	-	-		
86	OHX	5	3968	-	0,6,6	-	-	-		
86	OHX	5	4063	-	0,6,6	-	-	-		
86	OHX	2	2172	-	0,6,6	-	-	-		
86	OHX	5	4147	-	0,6,6	-	-	-		
86	OHX	1	3936	-	0,6,6	-	-	-		
86	OHX	6	2179	-	0,6,6	-	-	-		
86	OHX	2	2116	-	0,6,6	-	-	-		
86	OHX	6	2093	-	0,6,6	-	-	-		
86	OHX	2	2134	-	0,6,6	-	-	-		
86	OHX	5	4219	-	0,6,6	-	-	-		
86	OHX	5	3955	-	0,6,6	-	-	-		
86	OHX	5	4190	-	0,6,6	-	-	-		
86	OHX	5	4087	-	0,6,6	-	-	-		
86	OHX	2	2119	-	0,6,6	-	-	-		
86	OHX	8	232	-	0,6,6	-	-	-		
86	OHX	5	4233	-	0,6,6	-	-	-		
86	OHX	6	2104	-	0,6,6	-	-	-		
86	OHX	2	2042	-	0,6,6	-	-	-		
86	OHX	2	2123	-	0,6,6	-	-	-		
86	OHX	1	3900	-	0,6,6	-	-	-		
86	OHX	1	3997	-	0,6,6	-	-	-		
86	OHX	5	4035	-	0,6,6	-	-	-		
86	OHX	5	4137	-	0,6,6	-	-	-		
86	OHX	5	3926	-	0,6,6	-	-	-		
86	OHX	8	224	-	0,6,6	-	-	-		
86	OHX	5	4187	-	0,6,6	-	-	-		
86	OHX	1	4074	-	0,6,6	-	-	-		
86	OHX	6	2054	-	0,6,6	-	-	-		
86	OHX	c8	203	-	0,6,6	-	-	-		
86	OHX	2	2132	-	0,6,6	-	-	-		
86	OHX	7	220	-	0,6,6	-	-	-		
86	OHX	5	4157	-	0,6,6	-	-	-		
86	OHX	s1	302	-	0,6,6	-	-	-		
86	OHX	5	4051	-	0,6,6	-	-	-		
86	OHX	1	3902	-	0,6,6	-	-	-		
86	OHX	1	4030	-	0,6,6	-	-	-		
86	OHX	2	2168	-	0,6,6	-	-	-		
86	OHX	5	3939	-	0,6,6	-	-	-		
86	OHX	C3	201	-	0,6,6	-	-	-		
86	OHX	6	2161	-	0,6,6	-	-	-		
86	OHX	2	2175	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	5	4154	-	0,6,6	-	-	-		
86	OHX	SR	401	-	0,6,6	-	-	-		
86	OHX	1	4146	-	0,6,6	-	-	-		
86	OHX	s1	303	-	0,6,6	-	-	-		
86	OHX	1	4086	-	0,6,6	-	-	-		
86	OHX	5	4227	-	0,6,6	-	-	-		
86	OHX	1	4169	-	0,6,6	-	-	-		
86	OHX	2	2153	-	0,6,6	-	-	-		
86	OHX	5	4213	-	0,6,6	-	-	-		
86	OHX	1	4104	-	0,6,6	-	-	-		
86	OHX	5	4059	-	0,6,6	-	-	-		
86	OHX	5	4181	-	0,6,6	-	-	-		
86	OHX	2	2137	-	0,6,6	-	-	-		
86	OHX	6	2075	-	0,6,6	-	-	-		
86	OHX	1	4159	-	0,6,6	-	-	-		
86	OHX	6	2204	-	0,6,6	-	-	-		
86	OHX	5	3943	-	0,6,6	-	-	-		
86	OHX	5	4042	-	0,6,6	-	-	-		
86	OHX	3	216	-	0,6,6	-	-	-		
86	OHX	2	2113	-	0,6,6	-	-	-		
86	OHX	5	3922	-	0,6,6	-	-	-		
86	OHX	5	4104	-	0,6,6	-	-	-		
86	OHX	6	2155	-	0,6,6	-	-	-		
86	OHX	1	4058	-	0,6,6	-	-	-		
86	OHX	1	4152	-	0,6,6	-	-	-		
86	OHX	1	4151	-	0,6,6	-	-	-		
86	OHX	2	2179	-	0,6,6	-	-	-		
86	OHX	l5	304	-	0,6,6	-	-	-		
86	OHX	2	2176	-	0,6,6	-	-	-		
86	OHX	5	4208	-	0,6,6	-	-	-		
86	OHX	1	4191	-	0,6,6	-	-	-		
86	OHX	6	2060	-	0,6,6	-	-	-		
86	OHX	5	3929	-	0,6,6	-	-	-		
86	OHX	2	2071	-	0,6,6	-	-	-		
86	OHX	3	218	-	0,6,6	-	-	-		
86	OHX	2	2128	-	0,6,6	-	-	-		
86	OHX	2	2160	-	0,6,6	-	-	-		
86	OHX	1	4067	-	0,6,6	-	-	-		
86	OHX	5	4203	-	0,6,6	-	-	-		
86	OHX	5	3938	-	0,6,6	-	-	-		
86	OHX	1	4049	-	0,6,6	-	-	-		
86	OHX	2	2104	-	0,6,6	-	-	-		
86	OHX	1	4069	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	3	224	-	0,6,6	-	-	-		
86	OHX	2	2064	-	0,6,6	-	-	-		
86	OHX	2	2024	-	0,6,6	-	-	-		
86	OHX	2	2077	-	0,6,6	-	-	-		
86	OHX	1	3877	-	0,6,6	-	-	-		
86	OHX	5	4114	-	0,6,6	-	-	-		
86	OHX	5	3942	-	0,6,6	-	-	-		
86	OHX	2	2145	-	0,6,6	-	-	-		
86	OHX	5	4198	-	0,6,6	-	-	-		
86	OHX	1	3913	-	0,6,6	-	-	-		
86	OHX	1	4019	-	0,6,6	-	-	-		
86	OHX	1	4208	-	0,6,6	-	-	-		
86	OHX	6	2203	-	0,6,6	-	-	-		
86	OHX	1	3957	-	0,6,6	-	-	-		
86	OHX	1	3947	-	0,6,6	-	-	-		
86	OHX	6	2130	-	0,6,6	-	-	-		
86	OHX	5	3954	-	0,6,6	-	-	-		
86	OHX	1	3979	-	0,6,6	-	-	-		
86	OHX	1	3928	-	0,6,6	-	-	-		
86	OHX	6	2138	-	0,6,6	-	-	-		
86	OHX	5	3980	-	0,6,6	-	-	-		
86	OHX	6	2183	-	0,6,6	-	-	-		
86	OHX	5	3953	-	0,6,6	-	-	-		
86	OHX	1	3959	-	0,6,6	-	-	-		
86	OHX	5	4044	-	0,6,6	-	-	-		
86	OHX	1	4206	-	0,6,6	-	-	-		
86	OHX	5	4067	-	0,6,6	-	-	-		
86	OHX	1	4179	-	0,6,6	-	-	-		
86	OHX	1	4145	-	0,6,6	-	-	-		
86	OHX	1	3883	-	0,6,6	-	-	-		
86	OHX	5	4164	-	0,6,6	-	-	-		
86	OHX	4	235	-	0,6,6	-	-	-		
86	OHX	5	4162	-	0,6,6	-	-	-		
86	OHX	6	2169	-	0,6,6	-	-	-		
86	OHX	5	4078	-	0,6,6	-	-	-		
86	OHX	5	4168	-	0,6,6	-	-	-		
86	OHX	2	2056	-	0,6,6	-	-	-		
86	OHX	2	2076	-	0,6,6	-	-	-		
86	OHX	8	227	-	0,6,6	-	-	-		
86	OHX	2	2057	-	0,6,6	-	-	-		
86	OHX	1	3926	-	0,6,6	-	-	-		
86	OHX	4	238	-	0,6,6	-	-	-		
86	OHX	5	4139	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	2	2085	-	0,6,6	-	-	-		
86	OHX	2	2133	-	0,6,6	-	-	-		
86	OHX	1	3916	-	0,6,6	-	-	-		
86	OHX	2	2143	-	0,6,6	-	-	-		
86	OHX	5	3921	-	0,6,6	-	-	-		
86	OHX	6	2106	-	0,6,6	-	-	-		
86	OHX	2	2111	-	0,6,6	-	-	-		
86	OHX	2	2070	-	0,6,6	-	-	-		
86	OHX	6	2148	-	0,6,6	-	-	-		
86	OHX	6	2195	-	0,6,6	-	-	-		
86	OHX	1	3972	-	0,6,6	-	-	-		
86	OHX	1	4053	-	0,6,6	-	-	-		
86	OHX	5	4018	-	0,6,6	-	-	-		
86	OHX	5	4229	-	0,6,6	-	-	-		
86	OHX	15	306	-	0,6,6	-	-	-		
86	OHX	1	4113	-	0,6,6	-	-	-		
86	OHX	5	4023	-	0,6,6	-	-	-		
86	OHX	5	3966	-	0,6,6	-	-	-		
86	OHX	5	4136	-	0,6,6	-	-	-		
86	OHX	1	3977	-	0,6,6	-	-	-		
86	OHX	1	3946	-	0,6,6	-	-	-		
86	OHX	1	4005	-	0,6,6	-	-	-		
86	OHX	5	4224	-	0,6,6	-	-	-		
86	OHX	6	2170	-	0,6,6	-	-	-		
86	OHX	5	3927	-	0,6,6	-	-	-		
86	OHX	3	214	-	0,6,6	-	-	-		
86	OHX	5	4173	-	0,6,6	-	-	-		
86	OHX	5	3988	-	0,6,6	-	-	-		
86	OHX	O7	105	-	0,6,6	-	-	-		
86	OHX	5	4239	-	0,6,6	-	-	-		
86	OHX	14	402	-	0,6,6	-	-	-		
86	OHX	1	4056	-	0,6,6	-	-	-		
86	OHX	2	2148	-	0,6,6	-	-	-		
86	OHX	6	2092	-	0,6,6	-	-	-		
86	OHX	6	2109	-	0,6,6	-	-	-		
86	OHX	1	3912	-	0,6,6	-	-	-		
86	OHX	1	4165	-	0,6,6	-	-	-		
86	OHX	1	4085	-	0,6,6	-	-	-		
86	OHX	5	3944	-	0,6,6	-	-	-		
86	OHX	1	4065	-	0,6,6	-	-	-		
86	OHX	5	4025	-	0,6,6	-	-	-		
86	OHX	5	3987	-	0,6,6	-	-	-		
86	OHX	2	2139	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	1	3966	-	0,6,6	-	-	-		
86	OHX	1	4075	-	0,6,6	-	-	-		
86	OHX	5	4206	-	0,6,6	-	-	-		
86	OHX	1	4123	-	0,6,6	-	-	-		
86	OHX	2	2157	-	0,6,6	-	-	-		
86	OHX	1	4186	-	0,6,6	-	-	-		
86	OHX	5	3931	-	0,6,6	-	-	-		
86	OHX	5	3908	-	0,6,6	-	-	-		
86	OHX	1	4012	-	0,6,6	-	-	-		
86	OHX	2	2091	-	0,6,6	-	-	-		
86	OHX	2	2087	-	0,6,6	-	-	-		
86	OHX	1	3876	-	0,6,6	-	-	-		
86	OHX	5	3956	-	0,6,6	-	-	-		
86	OHX	5	4125	-	0,6,6	-	-	-		
86	OHX	5	4246	-	0,6,6	-	-	-		
86	OHX	5	4244	-	0,6,6	-	-	-		
86	OHX	6	2056	-	0,6,6	-	-	-		
86	OHX	1	4076	-	0,6,6	-	-	-		
86	OHX	1	3971	-	0,6,6	-	-	-		
86	OHX	2	2089	-	0,6,6	-	-	-		
86	OHX	1	3981	-	0,6,6	-	-	-		
86	OHX	1	3982	-	0,6,6	-	-	-		
86	OHX	5	4197	-	0,6,6	-	-	-		
86	OHX	5	4245	-	0,6,6	-	-	-		
86	OHX	5	4101	-	0,6,6	-	-	-		
86	OHX	8	231	-	0,6,6	-	-	-		
86	OHX	5	3995	-	0,6,6	-	-	-		
86	OHX	6	2111	-	0,6,6	-	-	-		
86	OHX	5	4116	-	0,6,6	-	-	-		
86	OHX	5	4186	-	0,6,6	-	-	-		
86	OHX	C5	201	-	0,6,6	-	-	-		
86	OHX	5	4155	-	0,6,6	-	-	-		
86	OHX	1	4009	-	0,6,6	-	-	-		
86	OHX	1	4062	-	0,6,6	-	-	-		
86	OHX	5	4220	-	0,6,6	-	-	-		
86	OHX	6	2202	-	0,6,6	-	-	-		
86	OHX	6	2162	-	0,6,6	-	-	-		
86	OHX	sR	401	-	0,6,6	-	-	-		
86	OHX	1	4000	-	0,6,6	-	-	-		
86	OHX	2	2047	-	0,6,6	-	-	-		
86	OHX	1	3873	-	0,6,6	-	-	-		
86	OHX	1	4001	-	0,6,6	-	-	-		
86	OHX	1	4167	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	1	3907	-	0,6,6	-	-	-		
86	OHX	6	2067	-	0,6,6	-	-	-		
86	OHX	1	3905	-	0,6,6	-	-	-		
86	OHX	M7	206	-	0,6,6	-	-	-		
86	OHX	4	237	-	0,6,6	-	-	-		
86	OHX	5	4004	-	0,6,6	-	-	-		
86	OHX	2	2156	-	0,6,6	-	-	-		
86	OHX	5	4238	-	0,6,6	-	-	-		
86	OHX	5	4085	-	0,6,6	-	-	-		
86	OHX	5	4052	-	0,6,6	-	-	-		
86	OHX	5	3979	-	0,6,6	-	-	-		
86	OHX	1	4082	-	0,6,6	-	-	-		
86	OHX	s4	302	-	0,6,6	-	-	-		
86	OHX	2	2174	-	0,6,6	-	-	-		
86	OHX	5	3971	-	0,6,6	-	-	-		
86	OHX	1	3951	-	0,6,6	-	-	-		
86	OHX	1	3941	-	0,6,6	-	-	-		
86	OHX	1	4139	-	0,6,6	-	-	-		
86	OHX	6	2154	-	0,6,6	-	-	-		
86	OHX	1	4121	-	0,6,6	-	-	-		
86	OHX	5	3999	-	0,6,6	-	-	-		
86	OHX	o3	203	-	0,6,6	-	-	-		
86	OHX	5	3983	-	0,6,6	-	-	-		
86	OHX	5	4090	-	0,6,6	-	-	-		
86	OHX	2	2138	-	0,6,6	-	-	-		
86	OHX	1	4185	-	0,6,6	-	-	-		
86	OHX	5	4055	-	0,6,6	-	-	-		
86	OHX	L3	404	-	0,6,6	-	-	-		
86	OHX	2	2037	-	0,6,6	-	-	-		
86	OHX	6	2123	-	0,6,6	-	-	-		
86	OHX	5	4207	-	0,6,6	-	-	-		
86	OHX	5	4058	-	0,6,6	-	-	-		
86	OHX	2	2146	-	0,6,6	-	-	-		
86	OHX	2	2127	-	0,6,6	-	-	-		
86	OHX	1	4096	-	0,6,6	-	-	-		
86	OHX	4	232	-	0,6,6	-	-	-		
86	OHX	7	223	-	0,6,6	-	-	-		
86	OHX	6	2121	-	0,6,6	-	-	-		
86	OHX	2	2163	-	0,6,6	-	-	-		
86	OHX	1	3999	-	0,6,6	-	-	-		
86	OHX	6	2083	-	0,6,6	-	-	-		
86	OHX	5	4036	-	0,6,6	-	-	-		
86	OHX	5	4195	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	5	4159	-	0,6,6	-	-	-		
86	OHX	6	2080	-	0,6,6	-	-	-		
86	OHX	2	2167	-	0,6,6	-	-	-		
86	OHX	5	4071	-	0,6,6	-	-	-		
86	OHX	M0	304	-	0,6,6	-	-	-		
86	OHX	5	4082	-	0,6,6	-	-	-		
86	OHX	1	3944	-	0,6,6	-	-	-		
86	OHX	6	2098	-	0,6,6	-	-	-		
86	OHX	8	217	-	0,6,6	-	-	-		
86	OHX	c3	201	-	0,6,6	-	-	-		
86	OHX	1	4209	-	0,6,6	-	-	-		
86	OHX	5	4026	-	0,6,6	-	-	-		
86	OHX	1	4127	-	0,6,6	-	-	-		
86	OHX	5	3913	-	0,6,6	-	-	-		
86	OHX	2	2108	-	0,6,6	-	-	-		
86	OHX	d4	201	-	0,6,6	-	-	-		
86	OHX	2	2041	-	0,6,6	-	-	-		
86	OHX	6	2053	-	0,6,6	-	-	-		
86	OHX	6	2181	-	0,6,6	-	-	-		
86	OHX	1	3955	-	0,6,6	-	-	-		
86	OHX	1	4172	-	0,6,6	-	-	-		
86	OHX	1	4087	-	0,6,6	-	-	-		
86	OHX	1	4176	-	0,6,6	-	-	-		
86	OHX	5	4095	-	0,6,6	-	-	-		
86	OHX	2	2105	-	0,6,6	-	-	-		
86	OHX	5	4124	-	0,6,6	-	-	-		
86	OHX	5	3950	-	0,6,6	-	-	-		
86	OHX	2	2043	-	0,6,6	-	-	-		
86	OHX	1	4197	-	0,6,6	-	-	-		
86	OHX	5	4054	-	0,6,6	-	-	-		
86	OHX	5	4215	-	0,6,6	-	-	-		
86	OHX	6	2140	-	0,6,6	-	-	-		
86	OHX	2	2068	-	0,6,6	-	-	-		
86	OHX	1	4109	-	0,6,6	-	-	-		
86	OHX	1	4051	-	0,6,6	-	-	-		
86	OHX	8	216	-	0,6,6	-	-	-		
86	OHX	6	2058	-	0,6,6	-	-	-		
86	OHX	1	4038	-	0,6,6	-	-	-		
86	OHX	N9	101	-	0,6,6	-	-	-		
86	OHX	2	2124	-	0,6,6	-	-	-		
86	OHX	5	4170	-	0,6,6	-	-	-		
86	OHX	5	4241	-	0,6,6	-	-	-		
86	OHX	5	4217	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	2	2166	-	0,6,6	-	-	-		
86	OHX	1	4220	-	0,6,6	-	-	-		
86	OHX	5	4134	-	0,6,6	-	-	-		
86	OHX	2	2086	-	0,6,6	-	-	-		
86	OHX	5	4020	-	0,6,6	-	-	-		
86	OHX	5	4091	-	0,6,6	-	-	-		
86	OHX	5	4174	-	0,6,6	-	-	-		
86	OHX	5	4048	-	0,6,6	-	-	-		
86	OHX	1	4147	-	0,6,6	-	-	-		
86	OHX	2	2103	-	0,6,6	-	-	-		
86	OHX	1	4004	-	0,6,6	-	-	-		
86	OHX	2	2035	-	0,6,6	-	-	-		
86	OHX	1	3882	-	0,6,6	-	-	-		
86	OHX	7	217	-	0,6,6	-	-	-		
86	OHX	2	2151	-	0,6,6	-	-	-		
86	OHX	1	3886	-	0,6,6	-	-	-		
86	OHX	5	4128	-	0,6,6	-	-	-		
86	OHX	5	4196	-	0,6,6	-	-	-		
86	OHX	5	4201	-	0,6,6	-	-	-		
86	OHX	1	4017	-	0,6,6	-	-	-		
86	OHX	5	3991	-	0,6,6	-	-	-		
86	OHX	1	4106	-	0,6,6	-	-	-		
86	OHX	1	4150	-	0,6,6	-	-	-		
86	OHX	1	4190	-	0,6,6	-	-	-		
86	OHX	1	4122	-	0,6,6	-	-	-		
86	OHX	1	3994	-	0,6,6	-	-	-		
86	OHX	5	3961	-	0,6,6	-	-	-		
86	OHX	1	4137	-	0,6,6	-	-	-		
86	OHX	5	3974	-	0,6,6	-	-	-		
86	OHX	1	3968	-	0,6,6	-	-	-		
86	OHX	1	3978	-	0,6,6	-	-	-		
86	OHX	5	4083	-	0,6,6	-	-	-		
86	OHX	m4	202	-	0,6,6	-	-	-		
86	OHX	5	4099	-	0,6,6	-	-	-		
86	OHX	5	4118	-	0,6,6	-	-	-		
86	OHX	5	3900	-	0,6,6	-	-	-		
86	OHX	5	3898	-	0,6,6	-	-	-		
86	OHX	1	4014	-	0,6,6	-	-	-		
86	OHX	5	4175	-	0,6,6	-	-	-		
86	OHX	5	4151	-	0,6,6	-	-	-		
86	OHX	3	220	-	0,6,6	-	-	-		
86	OHX	6	2068	-	0,6,6	-	-	-		
86	OHX	5	4231	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	4	233	-	0,6,6	-	-	-		
86	OHX	1	3890	-	0,6,6	-	-	-		
86	OHX	m7	206	-	0,6,6	-	-	-		
86	OHX	5	4073	-	0,6,6	-	-	-		
86	OHX	5	3905	-	0,6,6	-	-	-		
86	OHX	6	2073	-	0,6,6	-	-	-		
86	OHX	1	4118	-	0,6,6	-	-	-		
86	OHX	6	2141	-	0,6,6	-	-	-		
86	OHX	5	4226	-	0,6,6	-	-	-		
86	OHX	5	4002	-	0,6,6	-	-	-		
86	OHX	1	4212	-	0,6,6	-	-	-		
86	OHX	2	2022	-	0,6,6	-	-	-		
86	OHX	6	2101	-	0,6,6	-	-	-		
86	OHX	m1	202	-	0,6,6	-	-	-		
86	OHX	1	3953	-	0,6,6	-	-	-		
86	OHX	L4	403	-	0,6,6	-	-	-		
86	OHX	6	2049	-	0,6,6	-	-	-		
86	OHX	2	2084	-	0,6,6	-	-	-		
86	OHX	1	4036	-	0,6,6	-	-	-		
86	OHX	5	4105	-	0,6,6	-	-	-		
86	OHX	1	3975	-	0,6,6	-	-	-		
86	OHX	2	2052	-	0,6,6	-	-	-		
86	OHX	2	2039	-	0,6,6	-	-	-		
86	OHX	2	2027	-	0,6,6	-	-	-		
86	OHX	1	4214	-	0,6,6	-	-	-		
86	OHX	5	4185	-	0,6,6	-	-	-		
86	OHX	1	3988	-	0,6,6	-	-	-		
86	OHX	2	2101	-	0,6,6	-	-	-		
86	OHX	5	3982	-	0,6,6	-	-	-		
86	OHX	5	4193	-	0,6,6	-	-	-		
86	OHX	7	224	-	0,6,6	-	-	-		
86	OHX	5	4129	-	0,6,6	-	-	-		
86	OHX	6	2108	-	0,6,6	-	-	-		
86	OHX	5	4160	-	0,6,6	-	-	-		
86	OHX	l3	406	-	0,6,6	-	-	-		
86	OHX	1	4203	-	0,6,6	-	-	-		
86	OHX	5	4152	-	0,6,6	-	-	-		
86	OHX	5	3984	-	0,6,6	-	-	-		
86	OHX	2	2129	-	0,6,6	-	-	-		
86	OHX	1	3949	-	0,6,6	-	-	-		
86	OHX	5	3986	-	0,6,6	-	-	-		
86	OHX	1	3923	-	0,6,6	-	-	-		
86	OHX	1	4105	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	1	4218	-	0,6,6	-	-	-		
86	OHX	6	2076	-	0,6,6	-	-	-		
86	OHX	1	4021	-	0,6,6	-	-	-		
86	OHX	6	2081	-	0,6,6	-	-	-		
86	OHX	6	2107	-	0,6,6	-	-	-		
86	OHX	6	2142	-	0,6,6	-	-	-		
86	OHX	5	4010	-	0,6,6	-	-	-		
86	OHX	5	4034	-	0,6,6	-	-	-		
86	OHX	5	4199	-	0,6,6	-	-	-		
86	OHX	5	4015	-	0,6,6	-	-	-		
86	OHX	2	2115	-	0,6,6	-	-	-		
86	OHX	1	4196	-	0,6,6	-	-	-		
86	OHX	1	4201	-	0,6,6	-	-	-		
86	OHX	1	3915	-	0,6,6	-	-	-		
86	OHX	1	3973	-	0,6,6	-	-	-		
86	OHX	5	4132	-	0,6,6	-	-	-		
86	OHX	5	4240	-	0,6,6	-	-	-		
86	OHX	q2	502	-	0,6,6	-	-	-		
86	OHX	2	2117	-	0,6,6	-	-	-		
86	OHX	6	2097	-	0,6,6	-	-	-		
86	OHX	1	4016	-	0,6,6	-	-	-		
86	OHX	5	4177	-	0,6,6	-	-	-		
86	OHX	5	4120	-	0,6,6	-	-	-		
86	OHX	5	4172	-	0,6,6	-	-	-		
86	OHX	6	2066	-	0,6,6	-	-	-		
86	OHX	5	4247	-	0,6,6	-	-	-		
86	OHX	5	4211	-	0,6,6	-	-	-		
86	OHX	1	4198	-	0,6,6	-	-	-		
86	OHX	2	2046	-	0,6,6	-	-	-		
86	OHX	5	4236	-	0,6,6	-	-	-		
86	OHX	1	4112	-	0,6,6	-	-	-		
86	OHX	1	3924	-	0,6,6	-	-	-		
86	OHX	2	2065	-	0,6,6	-	-	-		
86	OHX	1	3943	-	0,6,6	-	-	-		
86	OHX	1	3908	-	0,6,6	-	-	-		
86	OHX	1	3974	-	0,6,6	-	-	-		
86	OHX	1	4199	-	0,6,6	-	-	-		
86	OHX	1	3922	-	0,6,6	-	-	-		
86	OHX	6	2078	-	0,6,6	-	-	-		
86	OHX	5	4161	-	0,6,6	-	-	-		
86	OHX	5	3949	-	0,6,6	-	-	-		
86	OHX	8	221	-	0,6,6	-	-	-		
86	OHX	1	3980	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	5	4007	-	0,6,6	-	-	-		
86	OHX	5	4031	-	0,6,6	-	-	-		
86	OHX	2	2067	-	0,6,6	-	-	-		
86	OHX	6	2052	-	0,6,6	-	-	-		
86	OHX	5	4111	-	0,6,6	-	-	-		
86	OHX	1	4060	-	0,6,6	-	-	-		
86	OHX	m0	301	-	0,6,6	-	-	-		
86	OHX	6	2090	-	0,6,6	-	-	-		
86	OHX	6	2194	-	0,6,6	-	-	-		
86	OHX	2	2112	-	0,6,6	-	-	-		
86	OHX	o7	503	-	0,6,6	-	-	-		
86	OHX	4	236	-	0,6,6	-	-	-		
86	OHX	5	3899	-	0,6,6	-	-	-		
86	OHX	1	4066	-	0,6,6	-	-	-		
86	OHX	2	2121	-	0,6,6	-	-	-		
86	OHX	1	3995	-	0,6,6	-	-	-		
86	OHX	1	3958	-	0,6,6	-	-	-		
86	OHX	6	2071	-	0,6,6	-	-	-		
86	OHX	6	2184	-	0,6,6	-	-	-		
86	OHX	2	2170	-	0,6,6	-	-	-		
86	OHX	5	4171	-	0,6,6	-	-	-		
86	OHX	8	223	-	0,6,6	-	-	-		
86	OHX	1	3996	-	0,6,6	-	-	-		
86	OHX	6	2072	-	0,6,6	-	-	-		
86	OHX	2	2092	-	0,6,6	-	-	-		
86	OHX	1	3933	-	0,6,6	-	-	-		
86	OHX	1	4136	-	0,6,6	-	-	-		
86	OHX	1	3898	-	0,6,6	-	-	-		
86	OHX	1	4149	-	0,6,6	-	-	-		
86	OHX	2	2038	-	0,6,6	-	-	-		
86	OHX	1	4072	-	0,6,6	-	-	-		
86	OHX	1	4164	-	0,6,6	-	-	-		
86	OHX	3	215	-	0,6,6	-	-	-		
86	OHX	2	2063	-	0,6,6	-	-	-		
86	OHX	5	4098	-	0,6,6	-	-	-		
86	OHX	5	4145	-	0,6,6	-	-	-		
86	OHX	5	4202	-	0,6,6	-	-	-		
86	OHX	5	4216	-	0,6,6	-	-	-		
86	OHX	6	2065	-	0,6,6	-	-	-		
86	OHX	1	4025	-	0,6,6	-	-	-		
86	OHX	5	4086	-	0,6,6	-	-	-		
86	OHX	1	4187	-	0,6,6	-	-	-		
86	OHX	5	4053	-	0,6,6	-	-	-		

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	5	4178	-	0,6,6	-	-	-		
86	OHX	6	2175	-	0,6,6	-	-	-		
86	OHX	m0	302	-	0,6,6	-	-	-		
86	OHX	1	4064	-	0,6,6	-	-	-		
86	OHX	5	4248	-	0,6,6	-	-	-		
86	OHX	4	231	-	0,6,6	-	-	-		
86	OHX	5	4068	-	0,6,6	-	-	-		
86	OHX	1	4173	-	0,6,6	-	-	-		
86	OHX	2	2093	-	0,6,6	-	-	-		
86	OHX	2	2031	-	0,6,6	-	-	-		
86	OHX	1	4162	-	0,6,6	-	-	-		
88	3K5	1	4221	-	63,63,63	1.24	1 (1%)	84,95,95	1.51	11 (13%)
86	OHX	1	3921	-	0,6,6	-	-	-		
86	OHX	1	4070	-	0,6,6	-	-	-		
86	OHX	1	4129	-	0,6,6	-	-	-		
86	OHX	1	4217	-	0,6,6	-	-	-		
86	OHX	5	3985	-	0,6,6	-	-	-		
86	OHX	6	2159	-	0,6,6	-	-	-		
86	OHX	1	3970	-	0,6,6	-	-	-		
86	OHX	5	4045	-	0,6,6	-	-	-		
86	OHX	5	4046	-	0,6,6	-	-	-		

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

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
88	3K5	1	4221	-	-	2/29/121/121	0/7/7/7
88	3K5	5	4249	-	-	7/29/121/121	0/7/7/7

All (2) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
88	1	4221	3K5	O3-C15	9.20	1.60	1.44
88	5	4249	3K5	O3-C15	-3.61	1.38	1.44

All (23) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
88	1	4221	3K5	O3-C16-C15	7.94	67.35	59.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
88	5	4249	3K5	O14-C34-C35	5.55	120.52	107.76
88	1	4221	3K5	O3-C15-C16	-4.51	55.88	59.74
88	5	4249	3K5	O3-C15-C16	3.96	63.14	59.74
88	5	4249	3K5	C3-C15-C17	3.90	111.30	103.30
88	1	4221	3K5	O14-C34-C35	3.52	115.86	107.76
88	5	4249	3K5	C4-C3-C15	-3.26	108.80	114.34
88	1	4221	3K5	C16-O3-C15	-3.21	56.77	60.89
88	1	4221	3K5	C15-C17-C22	-3.19	95.72	102.89
88	5	4249	3K5	C2-O-C3	2.85	115.34	113.51
88	5	4249	3K5	O3-C16-C15	-2.68	56.68	59.37
88	1	4221	3K5	O9-C26-C27	2.63	113.81	107.76
88	1	4221	3K5	O14-C34-C33	-2.54	101.93	107.76
88	1	4221	3K5	O7-C24-O6	-2.51	101.25	107.90
88	5	4249	3K5	C34-C35-C36	-2.44	104.82	110.12
88	5	4249	3K5	O-C3-C15	2.33	111.87	107.32
88	5	4249	3K5	C8-C7-C6	-2.10	116.27	122.28
88	5	4249	3K5	O14-C34-C33	-2.10	102.94	107.76
88	1	4221	3K5	O13-C35-C34	2.09	115.30	109.94
88	5	4249	3K5	O11-C25-C26	2.06	112.64	107.42
88	1	4221	3K5	C32-O12-C36	2.05	117.14	113.63
88	5	4249	3K5	O1-C5-C1	2.03	110.10	107.23
88	1	4221	3K5	C26-C27-C28	-2.01	105.76	110.12

There are no chirality outliers.

All (9) torsion outliers are listed below:

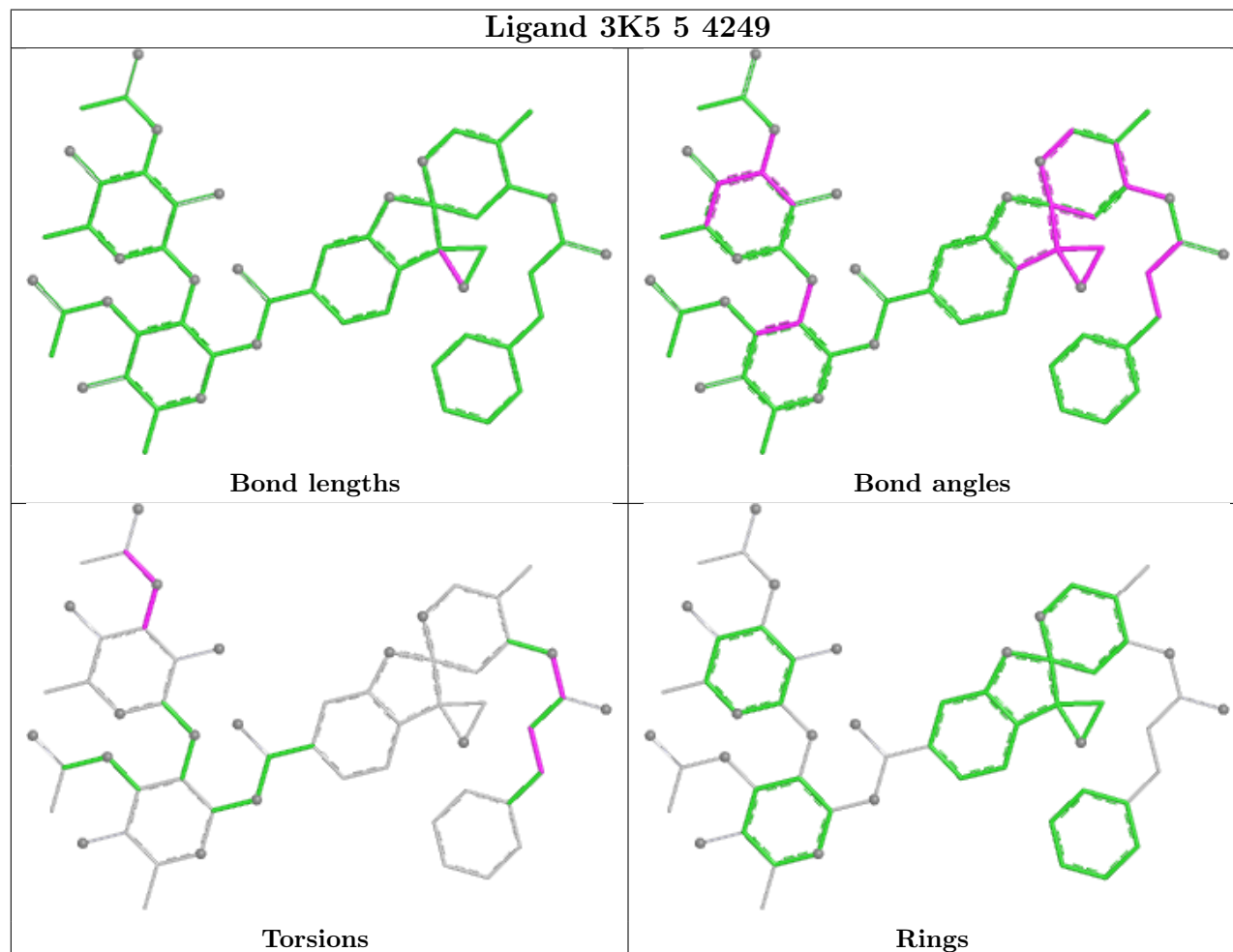
Mol	Chain	Res	Type	Atoms
88	5	4249	3K5	C6-C7-C8-C9
88	5	4249	3K5	C35-C34-O14-C38
88	5	4249	3K5	C39-C38-O14-C34
88	5	4249	3K5	C33-C34-O14-C38
88	5	4249	3K5	O15-C38-O14-C34
88	5	4249	3K5	O2-C6-O1-C5
88	1	4221	3K5	C19-C20-C23-O6
88	1	4221	3K5	C31-C30-O9-C26
88	5	4249	3K5	C7-C6-O1-C5

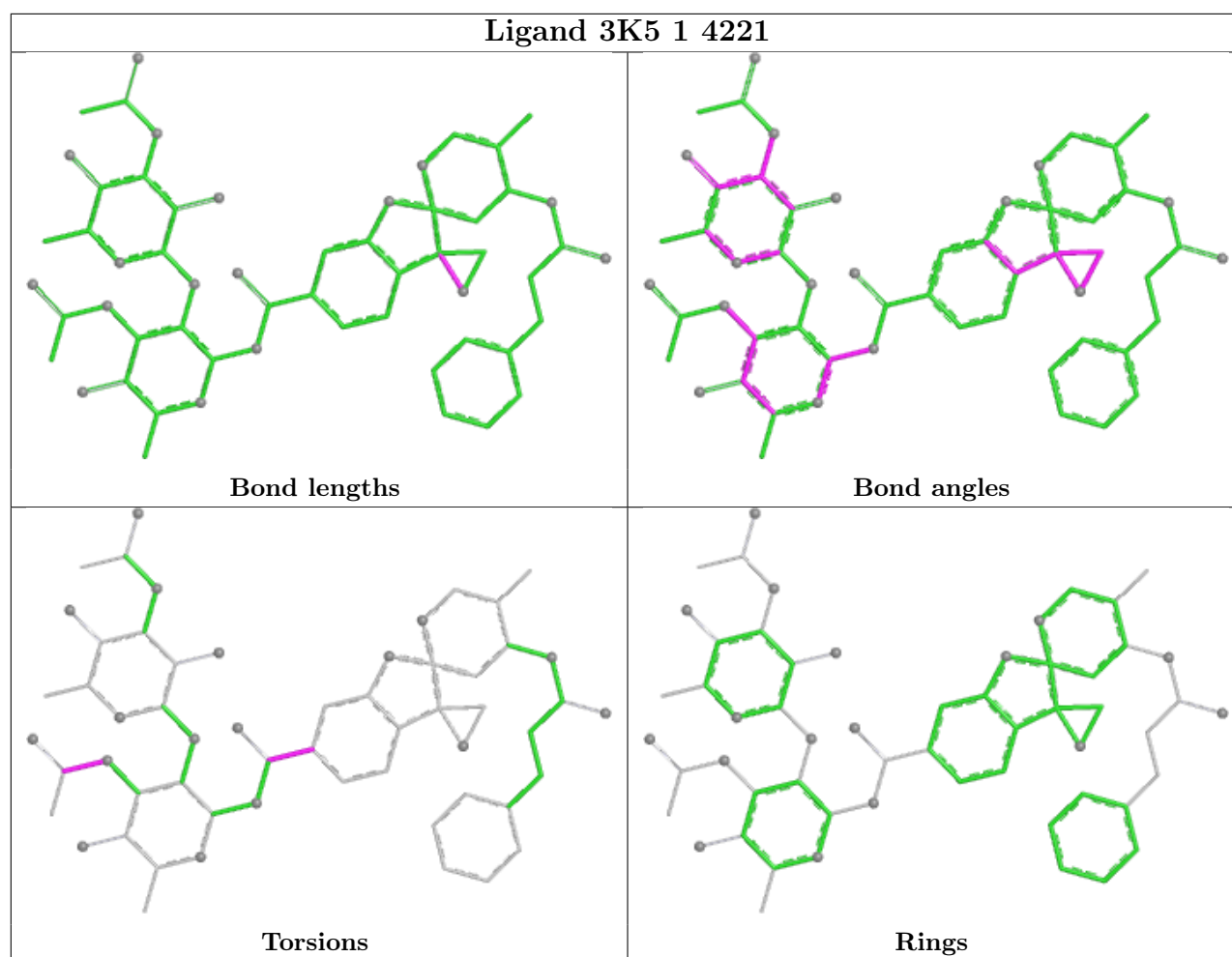
There are no ring outliers.

No monomer is involved in short contacts.

The following is a two-dimensional graphical depiction of Mogul quality analysis of bond lengths, bond angles, torsion angles, and ring geometry for all instances of the Ligand of Interest. In

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





4.7 Other polymers [i](#)

There are no such residues in this entry.

4.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

5 Fit of model and data ⓘ

5.1 Protein, DNA and RNA chains ⓘ

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

Mol	Chain	Analysed	<RSRZ>	#RSRZ > 2			OWAB(Å ²)	Q < 0.9
1	2	1750/1800 (97%)	0.37	76 (4%)	40	22	56, 92, 171, 265	0
1	6	1795/1800 (99%)	0.26	66 (3%)	45	25	43, 76, 180, 301	0
2	S0	206/251 (82%)	0.77	13 (6%)	26	14	94, 110, 125, 156	0
2	s0	206/251 (82%)	0.48	6 (2%)	53	32	73, 93, 110, 119	0
3	S1	214/254 (84%)	0.92	14 (6%)	25	13	100, 132, 158, 166	0
3	s1	216/254 (85%)	0.23	4 (1%)	66	45	66, 83, 109, 119	0
4	S2	217/253 (85%)	0.41	6 (2%)	55	34	72, 89, 106, 122	0
4	s2	217/253 (85%)	0.31	10 (4%)	37	20	56, 74, 98, 119	0
5	S3	223/239 (93%)	0.49	11 (4%)	35	18	79, 93, 131, 147	0
5	s3	223/239 (93%)	0.77	15 (6%)	24	13	74, 105, 131, 146	0
6	S4	260/260 (100%)	0.45	7 (2%)	56	35	68, 90, 104, 129	0
6	s4	260/260 (100%)	0.17	5 (1%)	66	45	54, 73, 92, 130	0
7	S5	206/224 (91%)	0.91	18 (8%)	16	9	97, 118, 136, 149	0
7	s5	206/224 (91%)	0.61	12 (5%)	29	16	70, 94, 125, 139	0
8	S6	226/236 (95%)	0.72	13 (5%)	29	16	65, 105, 126, 151	0
8	s6	218/236 (92%)	0.28	2 (0%)	81	63	53, 80, 105, 133	0
9	S7	184/189 (97%)	0.75	10 (5%)	31	17	89, 117, 148, 159	0
9	s7	186/189 (98%)	0.68	13 (6%)	22	12	67, 103, 136, 144	0
10	S8	188/200 (94%)	0.43	8 (4%)	40	22	58, 75, 115, 136	0
10	s8	188/200 (94%)	0.34	7 (3%)	45	25	47, 65, 114, 129	0
11	S9	185/196 (94%)	0.60	7 (3%)	44	25	83, 101, 136, 162	0
11	s9	185/196 (94%)	0.42	8 (4%)	40	22	63, 78, 117, 150	0
12	C0	96/105 (91%)	0.71	7 (7%)	21	11	82, 106, 140, 160	0
12	c0	96/105 (91%)	1.10	14 (14%)	6	3	97, 131, 158, 190	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
13	C1	155/155 (100%)	0.54	13 (8%) 17 9	60, 74, 134, 145	0
13	c1	146/155 (94%)	0.39	9 (6%) 26 14	47, 63, 99, 116	0
14	C2	124/142 (87%)	1.15	20 (16%) 4 2	129, 146, 167, 187	0
14	c2	124/142 (87%)	1.38	21 (16%) 4 2	172, 191, 208, 219	0
15	C3	150/150 (100%)	0.37	1 (0%) 84 68	69, 89, 108, 116	0
15	c3	150/150 (100%)	0.42	6 (4%) 42 23	57, 72, 93, 111	0
16	C4	127/136 (93%)	0.97	13 (10%) 12 6	71, 130, 147, 150	0
16	c4	128/136 (94%)	0.54	6 (4%) 36 19	53, 84, 94, 102	0
17	C5	124/141 (87%)	0.58	6 (4%) 35 19	79, 97, 136, 160	0
17	c5	135/141 (95%)	0.77	13 (9%) 13 7	76, 101, 136, 181	0
18	C6	141/142 (99%)	0.65	6 (4%) 40 22	83, 108, 115, 118	0
18	c6	142/142 (100%)	0.56	9 (6%) 26 14	66, 87, 106, 134	0
19	C7	120/136 (88%)	0.92	15 (12%) 8 5	92, 107, 138, 141	0
19	c7	117/136 (86%)	0.79	12 (10%) 12 6	78, 95, 122, 132	0
20	C8	145/145 (100%)	0.59	5 (3%) 48 28	80, 106, 132, 141	0
20	c8	145/145 (100%)	0.43	5 (3%) 48 28	72, 87, 117, 136	0
21	C9	143/143 (100%)	0.68	6 (4%) 40 22	90, 107, 127, 142	0
21	c9	143/143 (100%)	0.30	1 (0%) 84 68	66, 79, 100, 120	0
22	D0	107/120 (89%)	0.76	7 (6%) 25 13	73, 112, 152, 158	0
22	d0	110/120 (91%)	0.86	8 (7%) 21 11	68, 109, 156, 199	0
23	D1	87/87 (100%)	0.57	3 (3%) 48 28	92, 98, 116, 132	0
23	d1	87/87 (100%)	0.07	0 100 100	67, 78, 102, 117	0
24	D2	129/129 (100%)	0.39	2 (1%) 70 49	70, 83, 90, 100	0
24	d2	129/129 (100%)	-0.02	1 (0%) 82 65	52, 63, 71, 82	0
25	D3	144/144 (100%)	0.45	9 (6%) 26 14	61, 67, 79, 97	0
25	d3	144/144 (100%)	-0.02	3 (2%) 63 42	45, 52, 66, 79	0
26	D4	134/134 (100%)	0.75	8 (5%) 27 15	79, 109, 128, 138	0
26	d4	134/134 (100%)	0.25	6 (4%) 38 20	60, 83, 101, 127	0
27	D5	70/107 (65%)	0.79	5 (7%) 22 12	113, 129, 139, 142	0
27	d5	69/107 (64%)	0.44	1 (1%) 73 53	85, 111, 128, 134	0
28	D6	97/97 (100%)	0.98	11 (11%) 10 5	79, 96, 149, 155	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
28	d6	97/97 (100%)	0.41	3 (3%) 51 30	57, 73, 100, 109	0
29	D7	81/81 (100%)	0.71	3 (3%) 45 25	84, 103, 146, 152	0
29	d7	81/81 (100%)	0.63	5 (6%) 26 14	66, 85, 132, 137	0
30	D8	63/66 (95%)	1.01	7 (11%) 10 5	108, 129, 143, 159	0
30	d8	63/66 (95%)	0.56	2 (3%) 50 30	86, 105, 128, 135	0
31	D9	53/55 (96%)	0.64	5 (9%) 14 8	74, 82, 101, 114	0
31	d9	53/55 (96%)	0.84	4 (7%) 20 11	75, 82, 126, 139	0
32	E0	60/60 (100%)	0.55	2 (3%) 49 29	64, 98, 141, 153	0
33	E1	71/76 (93%)	1.26	11 (15%) 5 2	103, 126, 142, 150	0
33	e1	76/76 (100%)	1.61	19 (25%) 2 1	129, 159, 171, 175	0
34	SR	318/318 (100%)	0.68	18 (5%) 29 16	69, 115, 139, 157	0
34	sR	318/318 (100%)	0.79	21 (6%) 24 13	94, 116, 137, 155	0
35	SM	159/273 (58%)	0.73	19 (11%) 9 5	64, 92, 142, 149	0
35	sM	104/273 (38%)	0.84	11 (10%) 11 6	64, 101, 189, 194	0
36	1	3149/3396 (92%)	-0.10	73 (2%) 61 39	29, 53, 142, 278	0
36	5	3150/3396 (92%)	-0.19	53 (1%) 69 48	29, 52, 126, 288	0
37	3	121/121 (100%)	-0.29	82 65	44, 71, 87, 93	0
37	7	121/121 (100%)	-0.39	82 65	36, 56, 69, 78	0
38	4	158/158 (100%)	-0.29	75 56	35, 54, 96, 136	0
38	8	158/158 (100%)	-0.29	50 30	40, 63, 102, 133	0
39	L2	252/253 (99%)	-0.10	82 65	34, 49, 69, 79	0
39	l2	252/253 (99%)	0.09	46 26	35, 54, 74, 87	0
40	L3	386/386 (100%)	0.13	63 42	35, 59, 74, 110	0
40	l3	386/386 (100%)	-0.13	61 39	28, 42, 57, 96	0
41	L4	361/361 (100%)	-0.23	90 80	31, 46, 65, 78	0
41	l4	361/361 (100%)	0.01	69 48	35, 52, 72, 86	0
42	L5	296/296 (100%)	0.19	73 53	53, 78, 98, 128	0
42	l5	294/296 (99%)	-0.02	84 68	44, 59, 88, 136	0
43	L6	156/175 (89%)	-0.17	0 100 100	41, 49, 73, 101	0
43	l6	157/175 (89%)	-0.10	75 56	43, 53, 75, 90	0
44	L7	222/243 (91%)	-0.28	1 (0%) 87 73	34, 42, 75, 124	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2			OWAB(Å ²)	Q<0.9
44	l7	223/243 (91%)	-0.27	0	100	100	31, 41, 84, 129	0
45	L8	233/255 (91%)	0.30	6 (2%)	57	36	56, 71, 108, 122	0
45	l8	231/255 (90%)	0.44	10 (4%)	40	22	69, 83, 110, 117	0
46	L9	191/191 (100%)	0.13	2 (1%)	79	61	52, 67, 83, 103	0
46	l9	191/191 (100%)	-0.10	3 (1%)	70	49	38, 48, 71, 103	0
47	M0	211/220 (95%)	0.05	7 (3%)	49	29	39, 56, 94, 130	0
47	m0	213/220 (96%)	-0.08	3 (1%)	73	53	34, 53, 83, 102	0
48	M1	169/173 (97%)	0.24	2 (1%)	76	58	61, 82, 95, 103	0
48	m1	169/173 (97%)	0.22	3 (1%)	67	47	46, 65, 80, 92	0
49	M3	193/198 (97%)	-0.16	1 (0%)	87	73	34, 54, 102, 130	0
49	m3	194/198 (97%)	0.10	3 (1%)	72	52	41, 65, 108, 136	0
50	M4	136/137 (99%)	-0.18	0	100	100	44, 54, 68, 79	0
50	m4	137/137 (100%)	-0.21	1 (0%)	84	68	38, 45, 65, 77	0
51	M5	203/203 (100%)	-0.08	3 (1%)	72	52	32, 47, 59, 64	0
51	m5	203/203 (100%)	0.11	2 (0%)	79	61	40, 58, 70, 75	0
52	M6	197/198 (99%)	-0.14	1 (0%)	87	73	35, 44, 67, 72	0
52	m6	197/198 (99%)	-0.29	1 (0%)	87	73	29, 34, 62, 70	0
53	M7	183/183 (100%)	-0.05	7 (3%)	44	25	39, 46, 120, 144	0
53	m7	155/183 (84%)	-0.33	0	100	100	32, 41, 54, 89	0
54	M8	185/185 (100%)	-0.20	1 (0%)	87	73	36, 45, 61, 81	0
54	m8	185/185 (100%)	-0.15	1 (0%)	87	73	40, 51, 61, 67	0
55	M9	188/188 (100%)	0.40	12 (6%)	25	14	53, 70, 157, 171	0
55	m9	188/188 (100%)	0.26	5 (2%)	56	35	47, 60, 130, 144	0
56	N0	172/172 (100%)	-0.16	2 (1%)	76	58	44, 52, 66, 74	0
56	n0	172/172 (100%)	-0.22	2 (1%)	76	58	34, 42, 54, 67	0
57	N1	159/159 (100%)	0.02	1 (0%)	85	70	40, 51, 95, 103	0
57	n1	159/159 (100%)	-0.02	3 (1%)	66	45	37, 45, 89, 96	0
58	N2	100/120 (83%)	0.69	7 (7%)	22	12	83, 102, 119, 132	0
58	n2	98/120 (81%)	0.22	1 (1%)	79	61	72, 87, 102, 107	0
59	N3	136/136 (100%)	-0.00	2 (1%)	72	52	41, 53, 70, 87	0
59	n3	136/136 (100%)	-0.17	2 (1%)	72	52	30, 41, 59, 63	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
60	N4	98/155 (63%)	0.77	17 (17%) 4 2	52, 68, 162, 167	0
60	n4	135/155 (87%)	0.59	10 (7%) 20 11	41, 90, 134, 156	0
61	N5	121/141 (85%)	0.16	2 (1%) 69 48	49, 62, 83, 105	0
61	n5	120/141 (85%)	0.13	0 100 100	50, 65, 86, 93	0
62	N6	126/126 (100%)	-0.21	2 (1%) 70 49	40, 55, 68, 81	0
62	n6	126/126 (100%)	0.07	3 (2%) 59 38	47, 61, 80, 91	0
63	N7	135/135 (100%)	0.19	2 (1%) 72 52	70, 85, 101, 114	0
63	n7	135/135 (100%)	0.43	4 (2%) 52 31	75, 92, 116, 128	0
64	N8	148/148 (100%)	-0.10	1 (0%) 84 68	27, 46, 74, 86	0
64	n8	148/148 (100%)	-0.06	1 (0%) 84 68	32, 53, 76, 81	0
65	N9	58/58 (100%)	0.46	1 (1%) 69 48	36, 56, 107, 121	0
65	n9	58/58 (100%)	0.08	2 (3%) 48 28	35, 57, 89, 106	0
66	O0	97/104 (93%)	0.13	0 100 100	68, 80, 107, 120	0
66	o0	100/104 (96%)	0.34	3 (3%) 52 31	72, 83, 115, 126	0
67	O1	109/112 (97%)	0.17	2 (1%) 67 47	50, 64, 107, 121	0
67	o1	109/112 (97%)	0.01	2 (1%) 67 47	40, 53, 96, 123	0
68	O2	127/129 (98%)	-0.09	3 (2%) 59 38	28, 43, 56, 71	0
68	o2	127/129 (98%)	0.08	4 (3%) 51 30	30, 48, 65, 86	0
69	O3	106/106 (100%)	-0.30	0 100 100	35, 42, 65, 77	0
69	o3	106/106 (100%)	-0.28	1 (0%) 81 63	33, 41, 70, 85	0
70	O4	112/119 (94%)	0.22	1 (0%) 81 63	49, 67, 112, 129	0
70	o4	112/119 (94%)	0.22	1 (0%) 81 63	49, 69, 114, 127	0
71	O5	119/119 (100%)	-0.00	1 (0%) 82 65	45, 65, 73, 76	0
71	o5	119/119 (100%)	0.03	1 (0%) 82 65	54, 70, 82, 89	0
72	O6	99/99 (100%)	0.22	1 (1%) 79 61	53, 65, 101, 123	0
72	o6	99/99 (100%)	0.16	1 (1%) 79 61	58, 73, 99, 120	0
73	O7	87/87 (100%)	0.01	5 (5%) 29 16	35, 43, 75, 102	0
73	o7	87/87 (100%)	0.05	1 (1%) 78 59	39, 46, 82, 120	0
74	O8	77/77 (100%)	0.29	1 (1%) 75 56	73, 89, 113, 124	0
74	o8	77/77 (100%)	0.29	1 (1%) 75 56	73, 90, 105, 109	0
75	O9	50/50 (100%)	-0.02	1 (2%) 65 44	44, 50, 58, 61	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
75	o9	50/50 (100%)	0.14	1 (2%) 65 44	50, 54, 64, 75	0
76	Q0	52/52 (100%)	0.02	0 100 100	49, 59, 84, 94	0
76	q0	52/52 (100%)	-0.29	1 (1%) 66 45	34, 40, 54, 65	0
77	Q1	25/25 (100%)	0.12	0 100 100	54, 59, 63, 73	0
77	q1	25/25 (100%)	-0.16	0 100 100	44, 48, 63, 70	0
78	Q2	105/105 (100%)	0.05	2 (1%) 66 45	36, 54, 84, 116	0
78	q2	105/105 (100%)	-0.04	2 (1%) 66 45	43, 55, 84, 115	0
79	Q3	91/91 (100%)	-0.06	1 (1%) 78 59	44, 54, 74, 87	0
79	q3	91/91 (100%)	-0.03	1 (1%) 78 59	41, 55, 71, 84	0
80	e0	62/62 (100%)	0.32	2 (3%) 50 30	58, 81, 120, 139	0
81	m2	0/160	-	-	-	-
82	p0	143/311 (45%)	0.94	14 (9%) 13 7	86, 105, 187, 197	0
83	p1	0/47	-	-	-	-
84	p2	0/46	-	-	-	-
All	All	33063/35344 (93%)	0.19	1085 (3%) 49 29	27, 69, 136, 301	0

All (1085) RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
7	S5	152	GLY	7.2
36	5	2873	U	6.9
4	s2	90	THR	6.7
36	1	1349	G	6.5
60	N4	86	SER	5.8
56	N0	1	MET	5.6
36	5	1352	A	5.6
7	S5	37	GLN	5.5
36	5	3275	U	5.5
1	6	194	U	5.3
60	N4	76	VAL	5.3
30	D8	21	SER	5.3
36	1	1351	U	5.2
33	E1	124	PRO	5.2
36	1	1352	A	5.2
17	C5	104	GLN	5.2
36	5	2404	A	5.1
11	s9	2	PRO	5.1
1	6	320	U	5.1

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Mol	Chain	Res	Type	RSRZ
14	c2	123	VAL	5.1
36	5	2874	G	5.0
67	o1	82	GLU	5.0
36	5	1103	A	4.9
73	O7	88	ALA	4.9
4	s2	91	ARG	4.9
60	N4	89	LEU	4.9
1	6	678	A	4.9
33	e1	124	PRO	4.9
5	s3	145	ALA	4.8
1	2	320	U	4.7
49	m3	195	ALA	4.7
56	n0	2	ALA	4.7
58	N2	89	LEU	4.7
36	1	2874	G	4.7
9	s7	108	GLN	4.7
9	S7	145	GLY	4.7
17	c5	4	ALA	4.7
20	c8	17	LEU	4.6
1	6	718	U	4.6
1	6	794	U	4.5
36	5	1350	A	4.5
11	S9	2	PRO	4.5
45	L8	35	GLY	4.5
31	D9	23	VAL	4.4
1	6	754	A	4.4
13	c1	3	THR	4.3
1	2	238	U	4.3
1	2	260	U	4.3
36	5	1349	G	4.3
36	1	1103	A	4.3
29	D7	38	PRO	4.3
36	1	1350	A	4.3
47	M0	218	ALA	4.3
55	M9	189	ALA	4.3
1	2	310	C	4.2
47	M0	221	ALA	4.2
1	2	656	G	4.2
36	5	3276	G	4.2
34	sR	136	ILE	4.2
36	1	2873	U	4.2
36	1	2401	A	4.2

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Mol	Chain	Res	Type	RSRZ
33	e1	125	THR	4.2
22	D0	19	ILE	4.1
25	D3	3	LYS	4.1
7	S5	151	GLY	4.1
2	S0	40	ALA	4.1
16	C4	15	GLY	4.1
73	o7	88	ALA	4.1
7	S5	25	LEU	4.1
17	c5	11	VAL	4.1
19	C7	100	LEU	4.1
72	o6	11	LEU	4.1
60	N4	75	THR	4.1
2	s0	110	TYR	4.0
10	S8	21	PHE	4.0
82	p0	184	GLY	4.0
68	O2	26	HIS	4.0
1	2	1663	G	4.0
13	c1	2	SER	4.0
21	C9	5	SER	4.0
2	s0	40	ALA	4.0
20	c8	18	LEU	4.0
14	C2	88	LEU	4.0
47	m0	111	LEU	4.0
11	s9	8	TYR	4.0
13	c1	5	LEU	4.0
60	n4	97	LYS	4.0
6	s4	261	LEU	3.9
26	D4	135	ASP	3.9
3	S1	54	LEU	3.9
16	C4	102	LEU	3.9
9	s7	109	VAL	3.9
14	c2	30	VAL	3.9
28	D6	85	ARG	3.9
56	n0	1	MET	3.9
36	1	1581	C	3.9
38	8	81	U	3.9
7	S5	36	ALA	3.9
35	sM	119	ALA	3.9
31	d9	38	ILE	3.9
1	2	718	U	3.8
4	s2	92	ALA	3.8
26	D4	134	ALA	3.8

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Mol	Chain	Res	Type	RSRZ
18	c6	19	VAL	3.8
13	C1	152	GLN	3.8
36	1	2404	A	3.8
1	6	679	U	3.8
1	2	913	G	3.8
34	sR	121	MET	3.8
48	m1	116	TYR	3.8
60	N4	72	SER	3.8
31	d9	7	TRP	3.8
1	6	321	C	3.8
36	5	439	C	3.8
1	6	1150	G	3.8
53	M7	161	ALA	3.8
11	s9	5	PRO	3.8
14	C2	117	GLY	3.8
7	s5	79	ASN	3.7
11	S9	5	PRO	3.7
60	N4	81	PRO	3.7
17	c5	8	LYS	3.7
7	s5	37	GLN	3.7
36	1	1016	C	3.7
45	l8	124	ASP	3.7
1	6	1768	G	3.7
12	c0	98	THR	3.7
7	s5	24	VAL	3.7
25	D3	45	GLY	3.7
37	7	73	C	3.7
17	c5	7	ALA	3.7
8	S6	73	ILE	3.7
19	C7	126	ALA	3.7
13	C1	3	THR	3.7
1	2	321	C	3.7
26	d4	18	LEU	3.7
60	n4	68	ALA	3.6
1	2	1370	U	3.6
7	S5	222	LYS	3.6
25	D3	24	TRP	3.6
47	M0	219	ALA	3.6
39	l2	19	HIS	3.6
35	SM	9	GLY	3.6
36	5	2401	A	3.6
10	S8	8	ARG	3.6

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Mol	Chain	Res	Type	RSRZ
19	c7	97	ASN	3.6
33	e1	85	TYR	3.6
12	c0	13	GLN	3.6
10	S8	2	GLY	3.6
1	6	677	G	3.6
18	C6	92	TYR	3.6
33	e1	102	VAL	3.6
34	sR	141	LEU	3.6
1	6	738	G	3.5
33	e1	145	HIS	3.5
6	S4	25	GLY	3.5
14	C2	136	ILE	3.5
36	1	3079	U	3.5
80	e0	62	VAL	3.5
18	c6	3	ALA	3.5
36	1	2522	G	3.5
11	s9	3	ARG	3.5
9	s7	2	SER	3.5
16	c4	49	LYS	3.5
33	e1	84	VAL	3.5
62	N6	127	GLU	3.5
14	C2	130	THR	3.5
73	O7	87	SER	3.5
34	sR	162	ALA	3.5
36	1	1095	U	3.5
36	1	1353	U	3.5
23	D1	10	GLU	3.5
9	S7	105	THR	3.5
14	c2	128	ALA	3.5
3	S1	140	ILE	3.5
1	2	1130	G	3.5
41	l4	65	TRP	3.5
1	2	1362	U	3.5
12	c0	2	LEU	3.4
1	6	1157	A	3.4
12	c0	79	TYR	3.4
55	m9	78	TYR	3.4
1	2	1657	U	3.4
3	S1	46	THR	3.4
2	S0	23	HIS	3.4
53	M7	182	ILE	3.4
33	e1	77	GLY	3.4

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Mol	Chain	Res	Type	RSRZ
1	2	194	U	3.4
1	6	1129	U	3.4
22	D0	120	SER	3.4
60	N4	88	ASP	3.4
82	p0	203	ASP	3.4
33	e1	152	ALA	3.4
36	1	1583	A	3.4
9	s7	101	LYS	3.4
1	2	558	U	3.4
1	2	1666	U	3.4
9	s7	106	SER	3.4
30	D8	44	VAL	3.4
58	N2	27	VAL	3.4
42	l5	270	LYS	3.4
2	S0	207	PRO	3.4
34	sR	72	THR	3.4
14	c2	105	LYS	3.3
1	2	597	G	3.3
7	S5	33	VAL	3.3
20	C8	22	VAL	3.3
34	sR	107	LYS	3.3
82	p0	192	ASP	3.3
11	S9	6	ARG	3.3
40	l3	146	ARG	3.3
17	c5	138	PHE	3.3
5	s3	41	VAL	3.3
31	D9	9	SER	3.3
40	l3	363	SER	3.3
82	p0	200	SER	3.3
1	2	261	U	3.3
1	2	715	U	3.3
19	C7	11	ARG	3.3
1	2	716	C	3.3
33	e1	103	LEU	3.3
11	S9	181	ALA	3.3
22	D0	66	SER	3.3
82	p0	199	SER	3.3
12	c0	95	ARG	3.3
45	l8	254	ASP	3.3
1	6	1663	G	3.3
33	e1	147	VAL	3.3
1	2	1151	A	3.3

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Mol	Chain	Res	Type	RSRZ
1	6	192	U	3.3
36	1	2205	U	3.3
36	5	620	U	3.3
36	5	1356	U	3.3
38	4	82	U	3.3
5	s3	144	ALA	3.3
9	s7	12	ALA	3.3
47	m0	221	ALA	3.3
63	n7	124	ALA	3.3
12	c0	97	PRO	3.2
30	D8	15	VAL	3.2
13	c1	145	ALA	3.2
34	SR	78	ALA	3.2
36	1	1356	U	3.2
9	S7	108	GLN	3.2
19	c7	96	SER	3.2
22	D0	31	VAL	3.2
36	1	2400	G	3.2
38	8	80	A	3.2
35	SM	134	ASP	3.2
18	C6	3	ALA	3.2
65	N9	2	ALA	3.2
9	S7	109	VAL	3.2
31	d9	23	VAL	3.2
34	sR	104	VAL	3.2
36	5	1353	U	3.2
39	l2	248	GLY	3.2
12	c0	94	GLU	3.2
36	1	1196	C	3.2
5	s3	216	PRO	3.2
12	C0	13	GLN	3.2
33	e1	78	LYS	3.1
35	SM	11	ASP	3.1
16	C4	95	GLY	3.1
2	S0	18	LEU	3.1
3	s1	110	LEU	3.1
13	C1	153	PHE	3.1
36	5	3154	C	3.1
55	m9	85	ARG	3.1
7	s5	155	ALA	3.1
40	L3	94	GLU	3.1
34	SR	96	THR	3.1

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Mol	Chain	Res	Type	RSRZ
36	1	2772	C	3.1
1	2	1665	U	3.1
35	sM	156	ASN	3.1
38	4	81	U	3.1
10	s8	19	ALA	3.1
17	c5	133	ALA	3.1
16	c4	62	LEU	3.1
33	E1	87	THR	3.1
55	M9	78	TYR	3.1
33	E1	151	ASN	3.1
1	2	793	A	3.1
36	5	440	A	3.1
8	S6	158	ILE	3.1
36	5	3278	C	3.0
9	s7	52	ALA	3.0
12	c0	76	LEU	3.0
3	S1	93	GLY	3.0
74	O8	29	LYS	3.0
13	C1	2	SER	3.0
21	C9	36	ILE	3.0
14	C2	107	ASP	3.0
18	c6	142	TYR	3.0
14	C2	103	LEU	3.0
25	D3	33	LEU	3.0
1	2	1664	C	3.0
22	d0	70	THR	3.0
5	s3	217	ILE	3.0
36	5	3155	U	3.0
13	c1	144	ALA	3.0
63	n7	2	ALA	3.0
15	c3	104	ARG	3.0
7	S5	75	GLY	3.0
3	S1	25	THR	3.0
4	s2	208	GLU	3.0
36	5	1017	C	3.0
5	s3	43	PRO	3.0
18	c6	121	SER	3.0
59	N3	2	SER	3.0
7	S5	209	TYR	3.0
16	c4	60	ALA	3.0
28	d6	81	ALA	3.0
14	c2	131	ASP	3.0

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Mol	Chain	Res	Type	RSRZ
60	n4	117	LYS	3.0
2	S0	50	VAL	3.0
36	5	243	G	3.0
39	l2	186	PHE	3.0
14	c2	126	TRP	3.0
19	C7	89	SER	3.0
36	1	765	C	3.0
4	s2	93	GLY	3.0
7	S5	24	VAL	3.0
19	C7	110	VAL	3.0
47	M0	220	GLN	3.0
33	e1	123	ASN	3.0
45	l8	24	ASN	3.0
30	D8	53	ILE	3.0
41	l4	62	ALA	2.9
36	1	2139	A	2.9
36	1	2207	A	2.9
31	d9	6	VAL	2.9
35	SM	153	ASP	2.9
1	2	658	C	2.9
1	6	1285	U	2.9
1	6	1491	U	2.9
36	5	1569	U	2.9
82	p0	5	ARG	2.9
6	S4	60	GLU	2.9
13	C1	4	GLU	2.9
14	C2	141	SER	2.9
1	6	676	G	2.9
1	6	793	A	2.9
14	C2	62	LEU	2.9
5	S3	148	LYS	2.9
19	c7	35	CYS	2.9
82	p0	219	THR	2.9
1	6	1624	C	2.9
36	1	1574	C	2.9
1	2	494	U	2.9
1	6	662	U	2.9
1	6	1742	U	2.9
48	M1	96	PHE	2.9
15	c3	106	ARG	2.9
40	l3	140	ASP	2.9
9	s7	184	GLU	2.9

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Mol	Chain	Res	Type	RSRZ
47	m0	195	ALA	2.9
60	N4	68	ALA	2.9
33	E1	150	VAL	2.9
36	1	134	U	2.9
58	N2	108	TYR	2.9
73	O7	2	GLY	2.9
5	S3	179	GLN	2.9
29	d7	26	GLN	2.9
62	n6	120	GLN	2.9
18	C6	5	PRO	2.9
42	L5	291	ALA	2.9
16	C4	37	GLU	2.9
26	D4	46	GLU	2.9
28	D6	84	VAL	2.9
11	S9	3	ARG	2.9
1	6	1662	G	2.9
33	E1	103	LEU	2.9
17	c5	137	SER	2.8
1	2	1739	C	2.8
1	6	229	U	2.8
36	5	1567	U	2.8
46	l9	190	ASP	2.8
17	c5	109	PRO	2.8
14	C2	115	VAL	2.8
8	S6	54	GLY	2.8
10	s8	17	LYS	2.8
3	S1	229	MET	2.8
9	s7	187	SER	2.8
16	c4	50	ALA	2.8
36	5	3277	U	2.8
45	l8	25	PRO	2.8
19	c7	95	ARG	2.8
36	1	247	C	2.8
60	N4	92	GLU	2.8
20	c8	3	LEU	2.8
70	o4	81	CYS	2.8
5	S3	88	ALA	2.8
34	SR	102	ARG	2.8
15	c3	83	GLU	2.8
17	c5	9	LYS	2.8
27	D5	97	LYS	2.8
36	1	440	A	2.8

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Mol	Chain	Res	Type	RSRZ
5	S3	87	TYR	2.8
35	sM	66	ALA	2.8
14	c2	37	VAL	2.8
16	C4	67	VAL	2.8
35	SM	151	LEU	2.8
5	s3	225	TYR	2.8
36	1	1355	A	2.8
13	C1	151	LYS	2.8
1	2	1718	G	2.8
1	6	651	G	2.8
63	n7	13	VAL	2.8
14	c2	59	LEU	2.8
26	D4	2	SER	2.8
27	D5	41	ILE	2.8
7	s5	153	GLY	2.7
8	S6	80	ASN	2.7
39	l2	194	ASN	2.7
13	C1	146	ALA	2.7
49	m3	193	ALA	2.7
1	2	719	U	2.7
1	6	1152	A	2.7
6	S4	261	LEU	2.7
7	s5	152	GLY	2.7
33	e1	127	GLY	2.7
36	5	244	G	2.7
40	l3	244	ARG	2.7
8	S6	194	LYS	2.7
14	C2	85	LYS	2.7
26	d4	134	ALA	2.7
9	S7	100	PRO	2.7
5	S3	50	ILE	2.7
7	S5	150	GLY	2.7
35	SM	85	SER	2.7
22	D0	77	LYS	2.7
36	5	1582	C	2.7
40	l3	386	ASP	2.7
12	c0	89	ALA	2.7
13	c1	146	ALA	2.7
22	d0	118	VAL	2.7
40	L3	387	LEU	2.7
48	M1	163	PHE	2.7
2	s0	117	GLU	2.7

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Mol	Chain	Res	Type	RSRZ
62	n6	127	GLU	2.7
33	e1	80	ARG	2.7
60	N4	69	LYS	2.7
60	n4	95	SER	2.7
1	2	1577	A	2.7
36	5	2971	A	2.7
34	SR	27	ALA	2.7
35	SM	175	ASP	2.7
29	d7	24	LEU	2.7
32	E0	50	VAL	2.7
62	n6	126	LEU	2.7
35	sM	124	GLN	2.7
36	1	648	C	2.7
36	1	2360	C	2.7
48	m1	8	PRO	2.7
52	M6	80	PHE	2.7
24	D2	68	ARG	2.7
60	N4	73	ARG	2.7
30	d8	24	GLY	2.7
34	sR	62	LYS	2.7
34	SR	12	THR	2.7
36	1	2206	G	2.7
7	S5	154	ALA	2.7
12	C0	89	ALA	2.7
1	2	1129	U	2.7
5	s3	218	LEU	2.7
55	M9	184	LEU	2.7
14	C2	24	ILE	2.6
26	d4	106	GLN	2.6
36	1	1025	A	2.6
11	s9	10	LYS	2.6
16	c4	92	LYS	2.6
59	n3	3	GLY	2.6
70	O4	78	GLY	2.6
1	6	1537	C	2.6
1	6	1625	C	2.6
36	5	175	C	2.6
36	5	1016	C	2.6
4	S2	90	THR	2.6
33	E1	152	ALA	2.6
53	M7	165	VAL	2.6
2	S0	32	HIS	2.6

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Mol	Chain	Res	Type	RSRZ
1	2	279	G	2.6
36	5	437	G	2.6
82	p0	31	ASP	2.6
14	C2	137	MET	2.6
36	5	2875	U	2.6
36	1	621	A	2.6
78	q2	17	CYS	2.6
34	sR	167	VAL	2.6
61	N5	23	ALA	2.6
79	q3	2	ALA	2.6
18	c6	2	SER	2.6
36	1	1762	C	2.6
3	S1	55	LYS	2.6
13	c1	92	HIS	2.6
14	C2	89	ILE	2.6
39	L2	251	LYS	2.6
8	S6	181	PRO	2.6
45	l8	227	ASP	2.6
36	1	2875	U	2.6
36	5	1351	U	2.6
11	S9	130	THR	2.6
14	c2	45	LEU	2.6
13	C1	154	ALA	2.6
14	c2	86	VAL	2.6
34	sR	83	ALA	2.6
1	2	1357	A	2.6
1	2	1425	A	2.6
1	6	755	A	2.6
27	D5	88	ILE	2.6
3	S1	142	PHE	2.6
35	SM	152	GLN	2.6
51	M5	138	GLN	2.6
11	S9	38	ASN	2.6
39	L2	13	GLY	2.6
3	S1	65	VAL	2.6
13	C1	147	ALA	2.6
14	c2	130	THR	2.6
34	sR	25	THR	2.6
57	N1	121	ALA	2.6
17	C5	52	LYS	2.6
68	o2	16	LYS	2.6
9	s7	102	PRO	2.6

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Mol	Chain	Res	Type	RSRZ
1	6	1081	A	2.6
6	s4	260	GLY	2.6
7	s5	151	GLY	2.6
45	L8	37	GLY	2.6
2	S0	41	ARG	2.6
22	d0	97	VAL	2.6
36	1	2374	C	2.6
33	e1	101	ALA	2.6
2	S0	170	ILE	2.5
5	s3	115	ILE	2.5
1	6	663	U	2.5
36	5	249	U	2.5
19	c7	118	PRO	2.5
19	C7	125	SER	2.5
39	l2	14	SER	2.5
10	S8	20	GLN	2.5
12	C0	96	ASN	2.5
23	D1	21	ASN	2.5
34	SR	62	LYS	2.5
45	L8	156	ASP	2.5
46	l9	107	ASP	2.5
63	n7	59	ALA	2.5
9	S7	8	ILE	2.5
34	sR	134	TRP	2.5
47	M0	102	MET	2.5
1	2	1082	C	2.5
1	6	1000	C	2.5
34	sR	250	TYR	2.5
21	C9	85	SER	2.5
28	D6	9	GLY	2.5
34	SR	9	LEU	2.5
59	n3	2	SER	2.5
12	c0	93	GLN	2.5
68	o2	128	LEU	2.5
7	S5	179	ALA	2.5
12	c0	84	GLU	2.5
45	l8	125	ALA	2.5
79	Q3	92	ALA	2.5
10	S8	152	ILE	2.5
39	l2	15	ILE	2.5
43	l6	130	ILE	2.5
34	sR	61	PHE	2.5

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Mol	Chain	Res	Type	RSRZ
36	1	1879	A	2.5
36	5	100	A	2.5
36	1	1878	G	2.5
3	S1	47	LEU	2.5
63	N7	5	LEU	2.5
8	s6	217	SER	2.5
16	C4	99	GLN	2.5
20	c8	2	SER	2.5
42	l5	4	GLN	2.5
15	C3	61	THR	2.5
16	C4	93	THR	2.5
29	d7	66	PRO	2.5
9	S7	41	LEU	2.5
55	M9	85	ARG	2.5
1	2	1131	A	2.5
47	M0	114	GLY	2.5
4	S2	63	VAL	2.5
57	n1	126	VAL	2.5
36	1	1547	G	2.5
34	SR	136	ILE	2.5
1	6	239	C	2.5
67	o1	81	GLU	2.5
57	n1	79	MET	2.5
6	S4	57	ASN	2.5
1	2	1057	U	2.5
1	6	232	U	2.5
1	6	665	U	2.5
14	c2	78	LEU	2.5
20	C8	17	LEU	2.5
21	C9	87	GLY	2.5
14	C2	116	VAL	2.5
40	l3	139	GLN	2.5
19	C7	120	SER	2.4
31	D9	8	PHE	2.4
10	s8	200	LYS	2.4
19	c7	3	ARG	2.4
1	2	704	C	2.4
1	6	720	G	2.4
1	6	1159	C	2.4
25	d3	34	LEU	2.4
78	Q2	104	LEU	2.4
34	SR	79	TYR	2.4

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Mol	Chain	Res	Type	RSRZ
36	5	1764	U	2.4
22	d0	95	ALA	2.4
29	d7	5	GLN	2.4
53	M7	184	ALA	2.4
56	N0	138	GLN	2.4
60	n4	121	ALA	2.4
31	D9	50	ILE	2.4
34	sR	103	PHE	2.4
60	N4	80	ARG	2.4
30	D8	16	LEU	2.4
58	n2	89	LEU	2.4
1	2	1371	A	2.4
36	1	2971	A	2.4
2	S0	29	VAL	2.4
3	S1	225	VAL	2.4
1	2	1625	C	2.4
1	6	1447	C	2.4
1	2	1059	U	2.4
1	2	1108	G	2.4
7	s5	29	ILE	2.4
28	D6	63	ALA	2.4
36	5	1581	C	2.4
36	5	2507	C	2.4
71	o5	120	ALA	2.4
36	1	1028	U	2.4
26	d4	128	LYS	2.4
30	D8	22	ARG	2.4
34	SR	117	LYS	2.4
54	m8	11	LYS	2.4
8	s6	156	PHE	2.4
39	l2	16	PHE	2.4
35	SM	160	ASN	2.4
44	L7	209	ASN	2.4
6	S4	70	VAL	2.4
28	D6	80	HIS	2.4
39	l2	213	GLY	2.4
1	6	1371	A	2.4
18	c6	29	ILE	2.4
19	c7	90	ALA	2.4
40	L3	75	ALA	2.4
17	C5	12	PHE	2.4
1	2	313	U	2.4

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Mol	Chain	Res	Type	RSRZ
1	2	600	U	2.4
1	2	794	U	2.4
5	S3	21	LEU	2.4
22	d0	34	LEU	2.4
60	n4	66	GLU	2.4
19	C7	112	SER	2.4
10	S8	81	VAL	2.4
12	C0	90	THR	2.4
19	c7	8	THR	2.4
35	SM	19	VAL	2.4
4	s2	163	GLY	2.4
14	c2	111	ASN	2.4
6	s4	69	HIS	2.4
2	s0	111	ILE	2.4
7	S5	29	ILE	2.4
23	D1	23	ILE	2.4
25	D3	29	TYR	2.4
26	D4	130	ALA	2.4
55	m9	189	ALA	2.4
4	s2	97	ARG	2.4
35	SM	88	ARG	2.4
46	L9	31	ARG	2.4
8	S6	190	GLN	2.4
29	D7	32	PHE	2.4
9	s7	16	LEU	2.4
14	C2	45	LEU	2.4
60	N4	87	LEU	2.4
1	2	359	A	2.4
1	6	623	A	2.4
5	s3	128	GLU	2.4
38	8	79	A	2.4
60	n4	65	GLU	2.4
1	2	657	U	2.4
1	6	1661	U	2.4
36	1	1555	U	2.4
82	p0	80	VAL	2.4
32	E0	48	THR	2.3
37	3	73	C	2.3
25	d3	21	ASN	2.3
82	p0	23	LYS	2.3
82	p0	90	ASN	2.3
30	d8	65	ARG	2.3

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Mol	Chain	Res	Type	RSRZ
75	O9	45	ARG	2.3
34	sR	316	MET	2.3
12	C0	93	GLN	2.3
35	sM	158	GLN	2.3
39	l2	246	LEU	2.3
45	l8	77	GLN	2.3
66	o0	104	LEU	2.3
60	N4	66	GLU	2.3
67	O1	82	GLU	2.3
28	d6	58	VAL	2.3
14	C2	126	TRP	2.3
11	s9	6	ARG	2.3
19	C7	106	THR	2.3
33	E1	143	LYS	2.3
35	sM	67	GLY	2.3
49	M3	130	GLY	2.3
7	s5	154	ALA	2.3
60	n4	122	ALA	2.3
75	o9	51	ILE	2.3
1	2	1423	U	2.3
1	6	1708	U	2.3
20	C8	44	ASN	2.3
36	1	246	U	2.3
9	S7	153	LEU	2.3
36	1	1579	C	2.3
40	L3	226	PHE	2.3
55	M9	185	LEU	2.3
41	l4	63	GLU	2.3
28	D6	86	VAL	2.3
36	1	1576	G	2.3
60	n4	67	VAL	2.3
10	s8	135	LYS	2.3
17	c5	52	LYS	2.3
35	SM	21	PRO	2.3
35	sM	167	PRO	2.3
2	s0	44	GLY	2.3
34	SR	11	GLY	2.3
34	SR	115	ILE	2.3
51	M5	142	ILE	2.3
13	C1	145	ALA	2.3
14	C2	118	ALA	2.3
34	sR	106	HIS	2.3

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Mol	Chain	Res	Type	RSRZ
15	c3	149	LEU	2.3
28	D6	53	LEU	2.3
1	2	754	A	2.3
36	5	1355	A	2.3
78	q2	106	PHE	2.3
19	c7	103	ASP	2.3
1	2	1363	U	2.3
36	1	3155	U	2.3
36	5	250	U	2.3
1	2	726	C	2.3
1	6	1128	C	2.3
1	6	1623	C	2.3
17	c5	126	VAL	2.3
7	s5	148	ARG	2.3
10	s8	8	ARG	2.3
16	C4	94	PRO	2.3
40	L3	245	GLY	2.3
40	l3	245	GLY	2.3
34	SR	230	ALA	2.3
14	c2	34	THR	2.3
16	C4	11	SER	2.3
19	c7	25	THR	2.3
14	c2	36	LEU	2.3
20	C8	66	LEU	2.3
35	SM	20	LEU	2.3
1	6	624	G	2.3
33	e1	148	TYR	2.3
26	D4	60	PHE	2.3
35	sM	157	GLN	2.3
13	c1	29	LYS	2.3
1	2	881	A	2.3
28	D6	58	VAL	2.3
36	1	2996	U	2.3
36	1	3275	U	2.3
35	sM	40	PRO	2.3
3	S1	63	GLY	2.3
7	S5	105	GLY	2.3
11	s9	4	ALA	2.3
12	C0	19	GLY	2.3
53	M7	5	GLY	2.3
18	c6	17	THR	2.3
5	s3	195	SER	2.3

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Mol	Chain	Res	Type	RSRZ
15	c3	90	TYR	2.3
40	L3	154	TYR	2.3
40	l3	21	ARG	2.3
67	O1	79	ARG	2.3
4	s2	165	VAL	2.3
1	2	1737	G	2.2
1	6	1153	G	2.2
36	5	1953	G	2.2
36	5	2871	G	2.2
3	s1	60	ALA	2.2
8	S6	192	ALA	2.2
11	s9	165	GLY	2.2
28	D6	98	PRO	2.2
14	C2	128	ALA	2.2
15	c3	15	ALA	2.2
42	L5	287	ALA	2.2
55	m9	178	ALA	2.2
68	O2	2	ALA	2.2
1	2	133	U	2.2
1	2	1741	U	2.2
1	6	359	A	2.2
1	6	1800	A	2.2
26	d4	17	LEU	2.2
35	sM	155	LEU	2.2
36	1	245	U	2.2
36	1	2954	U	2.2
55	M9	138	LEU	2.2
1	2	233	C	2.2
18	c6	114	ARG	2.2
72	O6	56	ARG	2.2
25	D3	27	ASN	2.2
30	D8	7	VAL	2.2
35	SM	156	ASN	2.2
40	L3	93	VAL	2.2
45	L8	38	GLN	2.2
16	C4	83	ILE	2.2
8	S6	193	LEU	2.2
9	s7	58	LEU	2.2
29	D7	41	LEU	2.2
51	M5	88	GLY	2.2
68	o2	127	ALA	2.2
4	S2	224	PHE	2.2

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Mol	Chain	Res	Type	RSRZ
12	c0	67	THR	2.2
14	c2	53	THR	2.2
34	SR	99	THR	2.2
36	1	880	G	2.2
36	5	176	G	2.2
36	5	812	G	2.2
1	6	193	U	2.2
21	C9	80	TYR	2.2
51	m5	188	ARG	2.2
1	6	1719	A	2.2
22	d0	120	SER	2.2
25	D3	145	SER	2.2
36	1	647	A	2.2
36	5	3347	A	2.2
58	N2	72	SER	2.2
5	S3	222	VAL	2.2
22	d0	12	GLN	2.2
19	C7	101	ASN	2.2
55	M9	142	ILE	2.2
76	q0	77	ILE	2.2
14	c2	107	ASP	2.2
14	C2	104	ALA	2.2
41	l4	352	ALA	2.2
64	n8	149	ALA	2.2
73	O7	86	ALA	2.2
80	e0	56	MET	2.2
22	d0	57	ARG	2.2
3	S1	114	VAL	2.2
60	N4	95	SER	2.2
1	2	598	U	2.2
4	s2	89	GLN	2.2
1	2	1635	A	2.2
1	2	1662	G	2.2
1	6	228	G	2.2
1	6	1767	G	2.2
2	S0	24	LEU	2.2
12	C0	2	LEU	2.2
33	e1	100	LEU	2.2
34	SR	23	LEU	2.2
34	sR	32	LEU	2.2
57	n1	86	GLU	2.2
33	E1	128	ALA	2.2

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Mol	Chain	Res	Type	RSRZ
34	SR	95	ALA	2.2
45	l8	154	ALA	2.2
51	m5	39	ALA	2.2
55	M9	131	ALA	2.2
4	S2	88	LYS	2.2
16	C4	66	ASP	2.2
55	m9	153	LYS	2.2
60	N4	77	LYS	2.2
45	L8	57	ARG	2.2
68	O2	46	PHE	2.2
36	1	1192	C	2.2
36	1	1582	C	2.2
36	5	1562	C	2.2
38	8	84	C	2.2
55	M9	141	HIS	2.2
65	n9	27	TYR	2.2
66	o0	67	VAL	2.2
4	S2	248	SER	2.2
26	d4	69	SER	2.2
27	D5	65	LEU	2.2
17	c5	106	GLU	2.2
26	D4	39	GLU	2.2
49	m3	194	GLU	2.2
71	O5	8	GLU	2.2
9	S7	101	LYS	2.2
1	6	231	U	2.2
19	c7	98	GLY	2.2
41	l4	79	GLY	2.2
53	M7	164	LYS	2.2
5	s3	134	CYS	2.2
58	N2	107	PHE	2.2
78	Q2	106	PHE	2.2
1	2	1667	A	2.2
1	2	1118	G	2.2
1	2	1122	G	2.2
34	sR	294	TRP	2.2
36	1	2169	G	2.2
41	L4	311	HIS	2.2
1	6	1148	C	2.1
36	1	1556	C	2.1
4	s2	199	GLN	2.1
18	C6	28	LEU	2.1

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Mol	Chain	Res	Type	RSRZ
18	c6	44	LEU	2.1
19	C7	73	LEU	2.1
40	L3	241	LYS	2.1
35	SM	15	ALA	2.1
7	S5	153	GLY	2.1
34	SR	276	PRO	2.1
41	l4	80	GLY	2.1
64	N8	37	GLY	2.1
19	C7	123	ASN	2.1
52	m6	65	ASN	2.1
1	2	137	U	2.1
1	2	1361	U	2.1
5	S3	224	ASP	2.1
7	S5	149	VAL	2.1
1	6	1712	A	2.1
5	S3	225	TYR	2.1
19	C7	38	ILE	2.1
5	s3	176	LEU	2.1
8	S6	68	LEU	2.1
27	d5	97	LYS	2.1
45	l8	171	LYS	2.1
48	m1	174	LYS	2.1
1	6	739	G	2.1
3	S1	230	ALA	2.1
10	S8	22	ARG	2.1
36	1	2418	G	2.1
27	D5	36	ALA	2.1
33	E1	110	ALA	2.1
8	S6	33	GLY	2.1
54	M8	74	GLU	2.1
1	6	1215	C	2.1
36	5	1918	C	2.1
68	o2	17	PHE	2.1
33	e1	114	VAL	2.1
61	N5	26	VAL	2.1
1	6	1214	U	2.1
8	S6	226	ILE	2.1
16	C4	33	LEU	2.1
16	C4	89	THR	2.1
34	sR	137	LYS	2.1
36	1	979	U	2.1
36	5	981	U	2.1

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Mol	Chain	Res	Type	RSRZ
45	L8	231	LYS	2.1
60	N4	90	ILE	2.1
5	S3	40	ARG	2.1
29	d7	17	ARG	2.1
59	N3	32	ARG	2.1
1	2	217	A	2.1
1	6	1707	A	2.1
7	s5	22	PRO	2.1
35	SM	13	GLU	2.1
35	SM	167	PRO	2.1
35	sM	120	GLU	2.1
36	1	1881	A	2.1
36	5	361	A	2.1
2	S0	7	PHE	2.1
13	C1	156	PHE	2.1
13	c1	9	SER	2.1
13	C1	150	ASN	2.1
1	6	1288	G	2.1
14	c2	60	VAL	2.1
36	5	170	G	2.1
46	l9	75	VAL	2.1
53	M7	166	VAL	2.1
36	1	1017	C	2.1
36	5	1578	C	2.1
2	S0	54	TRP	2.1
25	d3	31	LYS	2.1
2	s0	24	LEU	2.1
14	c2	136	ILE	2.1
19	C7	46	LEU	2.1
22	D0	103	ILE	2.1
35	SM	14	ASP	2.1
58	N2	38	ILE	2.1
66	o0	14	LEU	2.1
17	C5	10	ARG	2.1
58	N2	94	ARG	2.1
17	C5	28	MET	2.1
14	c2	92	ALA	2.1
34	SR	162	ALA	2.1
40	l3	267	ALA	2.1
1	2	132	U	2.1
1	6	670	U	2.1
1	6	1626	U	2.1

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Mol	Chain	Res	Type	RSRZ
36	1	1455	U	2.1
36	1	1584	U	2.1
5	s3	220	PRO	2.1
6	S4	24	SER	2.1
55	M9	59	SER	2.1
55	M9	94	VAL	2.1
1	2	369	A	2.1
1	2	538	A	2.1
12	c0	96	ASN	2.1
13	C1	155	LYS	2.1
17	C5	59	LYS	2.1
65	n9	22	LYS	2.1
3	s1	217	LEU	2.1
6	S4	64	ILE	2.1
6	s4	90	ILE	2.1
9	s7	107	ARG	2.1
24	d2	26	LEU	2.1
18	C6	21	HIS	2.1
2	S0	72	ASP	2.1
1	2	239	C	2.1
1	6	717	C	2.1
1	6	1634	C	2.1
28	D6	6	ALA	2.1
36	1	439	C	2.1
36	1	918	C	2.1
36	1	2362	C	2.1
38	8	83	C	2.1
36	1	1952	G	2.1
10	S8	50	GLY	2.1
14	C2	102	GLY	2.1
45	l8	123	GLN	2.1
25	D3	64	PRO	2.1
43	l6	129	GLU	2.1
1	2	792	U	2.0
8	S6	41	VAL	2.0
9	S7	144	VAL	2.0
33	E1	116	LYS	2.0
36	1	249	U	2.0
36	1	1820	U	2.0
36	1	3156	U	2.0
36	5	2996	U	2.0
50	m4	8	LYS	2.0

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Mol	Chain	Res	Type	RSRZ
55	M9	177	VAL	2.0
7	s5	96	SER	2.0
82	p0	93	LEU	2.0
22	D0	84	MET	2.0
28	d6	80	HIS	2.0
5	s3	3	ALA	2.0
17	c5	54	ALA	2.0
19	c7	39	ALA	2.0
34	sR	135	THR	2.0
36	1	1893	A	2.0
42	L5	288	ALA	2.0
74	o8	2	ALA	2.0
6	s4	109	PHE	2.0
46	L9	85	GLY	2.0
20	C8	82	PRO	2.0
19	C7	75	GLU	2.0
1	2	717	C	2.0
1	6	1739	C	2.0
20	c8	27	LYS	2.0
35	SM	83	LYS	2.0
5	S3	138	VAL	2.0
14	c2	121	VAL	2.0
36	1	641	C	2.0
1	2	714	G	2.0
1	2	1112	G	2.0
1	2	1114	G	2.0
36	5	813	G	2.0
7	S5	40	ILE	2.0
82	p0	201	ILE	2.0
1	2	74	U	2.0
24	D2	70	ASN	2.0
36	5	246	U	2.0
31	D9	7	TRP	2.0
3	s1	126	THR	2.0
12	c0	90	THR	2.0
21	c9	136	ALA	2.0
42	L5	292	ALA	2.0
10	s8	71	GLY	2.0
25	D3	107	PHE	2.0
10	s8	20	GLN	2.0
16	c4	65	GLN	2.0
18	C6	132	LYS	2.0

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Mol	Chain	Res	Type	RSRZ
28	D6	60	PRO	2.0
47	M0	113	GLN	2.0
60	n4	116	LYS	2.0
73	O7	10	LYS	2.0
1	2	135	A	2.0
4	S2	205	ARG	2.0
21	C9	71	VAL	2.0
33	E1	102	VAL	2.0
36	1	2208	A	2.0
62	N6	88	GLU	2.0
82	p0	208	GLU	2.0
69	o3	60	ARG	2.0
26	D4	40	LEU	2.0
33	e1	117	LEU	2.0
63	N7	42	LEU	2.0
36	5	3285	C	2.0

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

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

5.3 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.4 Ligands [i](#)

In the following table, the Atoms column lists the number of modelled atoms in the group and the number defined in the chemical component dictionary. The B-factors column lists the minimum, median, 95th percentile and maximum values of B factors of atoms in the group. The column labelled 'Q< 0.9' lists the number of atoms with occupancy less than 0.9.

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(Å ²)	Q<0.9
85	MG	1	3815	1/1	-0.43	0.40	193,193,193,193	0
85	MG	1	3825	1/1	-0.19	0.43	121,121,121,121	0
85	MG	6	2013	1/1	0.06	0.34	150,150,150,150	0
85	MG	6	1923	1/1	0.10	0.43	103,103,103,103	0
85	MG	2	1993	1/1	0.11	0.47	114,114,114,114	0
85	MG	2	1967	1/1	0.24	0.50	121,121,121,121	0
85	MG	2	2001	1/1	0.26	0.46	109,109,109,109	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	1	3406	1/1	0.29	0.38	105,105,105,105	0
85	MG	1	3860	1/1	0.31	0.32	99,99,99,99	0
85	MG	2	1958	1/1	0.44	0.54	101,101,101,101	0
85	MG	1	3653	1/1	0.47	0.36	101,101,101,101	0
85	MG	2	1962	1/1	0.47	0.26	143,143,143,143	0
85	MG	6	2039	1/1	0.47	0.41	91,91,91,91	0
85	MG	6	1909	1/1	0.50	0.41	108,108,108,108	0
85	MG	2	1928	1/1	0.50	0.47	86,86,86,86	0
85	MG	2	1985	1/1	0.52	0.43	106,106,106,106	0
85	MG	5	3886	1/1	0.52	0.39	93,93,93,93	0
85	MG	2	1952	1/1	0.53	0.45	99,99,99,99	0
85	MG	5	3882	1/1	0.54	0.41	86,86,86,86	0
85	MG	1	3636	1/1	0.55	0.37	80,80,80,80	0
85	MG	2	1951	1/1	0.56	0.35	99,99,99,99	0
85	MG	2	1915	1/1	0.57	0.47	74,74,74,74	0
85	MG	6	2029	1/1	0.59	0.29	105,105,105,105	0
85	MG	2	1977	1/1	0.59	0.49	101,101,101,101	0
85	MG	5	3775	1/1	0.60	0.31	101,101,101,101	0
85	MG	5	3805	1/1	0.60	0.16	155,155,155,155	0
85	MG	2	1953	1/1	0.61	0.31	105,105,105,105	0
85	MG	5	3467	1/1	0.61	0.41	87,87,87,87	0
85	MG	6	1966	1/1	0.61	0.28	85,85,85,85	0
85	MG	2	1987	1/1	0.62	0.25	85,85,85,85	0
85	MG	6	1997	1/1	0.62	0.40	97,97,97,97	0
85	MG	2	2021	1/1	0.63	0.39	116,116,116,116	0
85	MG	5	3687	1/1	0.63	0.45	69,69,69,69	0
85	MG	6	2043	1/1	0.64	0.16	54,54,54,54	0
85	MG	2	2017	1/1	0.64	0.36	80,80,80,80	0
85	MG	5	3792	1/1	0.64	0.35	93,93,93,93	0
85	MG	2	1976	1/1	0.65	0.33	93,93,93,93	0
85	MG	6	2019	1/1	0.65	0.29	108,108,108,108	0
86	OHX	1	4192	7/7	0.65	0.20	190,190,190,190	0
85	MG	5	3781	1/1	0.66	0.36	80,80,80,80	0
85	MG	5	3653	1/1	0.66	0.27	120,120,120,120	0
85	MG	2	1949	1/1	0.68	0.49	88,88,88,88	0
85	MG	6	1989	1/1	0.68	0.42	78,78,78,78	0
85	MG	2	1950	1/1	0.69	0.37	99,99,99,99	0
85	MG	5	3421	1/1	0.69	0.29	96,96,96,96	0
86	OHX	1	4195	7/7	0.69	0.26	145,145,145,145	0
85	MG	o1	201	1/1	0.70	0.17	99,99,99,99	0
85	MG	2	1965	1/1	0.70	0.32	87,87,87,87	0
85	MG	3	211	1/1	0.70	0.17	75,75,75,75	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	5	3680	1/1	0.71	0.26	94,94,94,94	0
85	MG	5	3822	1/1	0.71	0.24	99,99,99,99	0
85	MG	6	1970	1/1	0.71	0.34	73,73,73,73	0
85	MG	5	3743	1/1	0.71	0.28	36,36,36,36	0
85	MG	6	1985	1/1	0.71	0.16	71,71,71,71	0
85	MG	6	1939	1/1	0.71	0.42	90,90,90,90	0
85	MG	2	1988	1/1	0.71	0.40	64,64,64,64	0
86	OHX	15	306	7/7	0.71	0.23	153,153,153,153	0
85	MG	1	3760	1/1	0.72	0.26	100,100,100,100	0
85	MG	2	1901	1/1	0.72	0.37	81,81,81,81	0
85	MG	2	1971	1/1	0.72	0.35	85,85,85,85	0
85	MG	1	3737	1/1	0.72	0.22	83,83,83,83	0
85	MG	6	1962	1/1	0.72	0.34	92,92,92,92	0
85	MG	3	209	1/1	0.72	0.39	62,62,62,62	0
86	OHX	5	4232	7/7	0.72	0.21	162,162,162,162	0
86	OHX	5	4234	7/7	0.72	0.21	165,165,165,165	0
85	MG	5	3666	1/1	0.72	0.41	65,65,65,65	0
86	OHX	2	2162	7/7	0.73	0.17	181,181,181,181	0
85	MG	2	1974	1/1	0.73	0.42	80,80,80,80	0
85	MG	5	3499	1/1	0.73	0.25	56,56,56,56	0
86	OHX	6	2192	7/7	0.73	0.20	176,176,176,176	0
86	OHX	5	4219	7/7	0.73	0.19	173,173,173,173	0
85	MG	1	3476	1/1	0.73	0.32	85,85,85,85	0
85	MG	6	1983	1/1	0.73	0.36	84,84,84,84	0
85	MG	2	1989	1/1	0.73	0.33	105,105,105,105	0
86	OHX	1	4147	7/7	0.74	0.25	133,133,133,133	0
85	MG	2	1914	1/1	0.74	0.38	71,71,71,71	0
85	MG	5	3807	1/1	0.74	0.26	97,97,97,97	0
85	MG	2	1945	1/1	0.74	0.28	84,84,84,84	0
85	MG	M1	201	1/1	0.74	0.18	74,74,74,74	0
86	OHX	5	4226	7/7	0.74	0.29	132,132,132,132	0
85	MG	2	1968	1/1	0.74	0.30	83,83,83,83	0
85	MG	2	1926	1/1	0.74	0.34	99,99,99,99	0
85	MG	6	1932	1/1	0.74	0.34	63,63,63,63	0
85	MG	6	2034	1/1	0.75	0.40	87,87,87,87	0
85	MG	1	3593	1/1	0.75	0.20	63,63,63,63	0
85	MG	6	1964	1/1	0.75	0.23	73,73,73,73	0
85	MG	2	1939	1/1	0.75	0.30	71,71,71,71	0
85	MG	2	1979	1/1	0.75	0.37	72,72,72,72	0
85	MG	6	1974	1/1	0.75	0.35	73,73,73,73	0
85	MG	1	3725	1/1	0.75	0.13	47,47,47,47	0
85	MG	2	1904	1/1	0.75	0.37	77,77,77,77	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	6	1987	1/1	0.75	0.24	86,86,86,86	0
85	MG	2	1913	1/1	0.75	0.43	84,84,84,84	0
85	MG	1	3479	1/1	0.75	0.19	42,42,42,42	0
85	MG	6	1934	1/1	0.75	0.22	61,61,61,61	0
85	MG	1	3582	1/1	0.75	0.21	47,47,47,47	0
85	MG	5	3787	1/1	0.75	0.38	87,87,87,87	0
85	MG	6	1960	1/1	0.75	0.38	79,79,79,79	0
85	MG	5	3803	1/1	0.75	0.19	65,65,65,65	0
85	MG	2	1924	1/1	0.76	0.45	87,87,87,87	0
86	OHX	1	4181	7/7	0.76	0.18	163,163,163,163	0
85	MG	1	3481	1/1	0.76	0.27	78,78,78,78	0
85	MG	6	1951	1/1	0.76	0.35	63,63,63,63	0
85	MG	4	222	1/1	0.76	0.39	76,76,76,76	0
85	MG	D0	201	1/1	0.76	0.43	77,77,77,77	0
85	MG	2	1933	1/1	0.76	0.36	77,77,77,77	0
85	MG	7	204	1/1	0.76	0.37	68,68,68,68	0
85	MG	1	3420	1/1	0.76	0.32	82,82,82,82	0
85	MG	2	1938	1/1	0.76	0.35	72,72,72,72	0
85	MG	15	302	1/1	0.77	0.20	67,67,67,67	0
85	MG	4	204	1/1	0.77	0.29	71,71,71,71	0
85	MG	5	3636	1/1	0.77	0.36	79,79,79,79	0
85	MG	1	3668	1/1	0.77	0.50	75,75,75,75	0
85	MG	1	3717	1/1	0.77	0.42	85,85,85,85	0
85	MG	2	2018	1/1	0.77	0.45	67,67,67,67	0
85	MG	5	3813	1/1	0.77	0.14	88,88,88,88	0
85	MG	3	208	1/1	0.77	0.21	82,82,82,82	0
86	OHX	6	2205	7/7	0.77	0.20	157,157,157,157	0
85	MG	2	1937	1/1	0.77	0.39	63,63,63,63	0
85	MG	2	1960	1/1	0.77	0.28	60,60,60,60	0
85	MG	5	3887	1/1	0.77	0.25	64,64,64,64	0
85	MG	5	3490	1/1	0.77	0.13	33,33,33,33	0
86	OHX	5	4238	7/7	0.77	0.14	238,238,238,238	0
85	MG	7	210	1/1	0.77	0.19	81,81,81,81	0
85	MG	5	3756	1/1	0.78	0.09	53,53,53,53	0
85	MG	3	201	1/1	0.78	0.30	73,73,73,73	0
86	OHX	2	2159	7/7	0.78	0.23	148,148,148,148	0
85	MG	2	1957	1/1	0.78	0.39	81,81,81,81	0
86	OHX	2	2177	7/7	0.78	0.16	188,188,188,188	0
85	MG	6	1933	1/1	0.78	0.35	77,77,77,77	0
85	MG	1	3408	1/1	0.78	0.24	45,45,45,45	0
85	MG	5	3470	1/1	0.78	0.23	110,110,110,110	0
85	MG	2	1909	1/1	0.78	0.41	72,72,72,72	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	6	2185	7/7	0.78	0.21	145,145,145,145	0
85	MG	1	3466	1/1	0.78	0.12	48,48,48,48	0
85	MG	2	1930	1/1	0.78	0.23	67,67,67,67	0
85	MG	L5	301	1/1	0.78	0.28	68,68,68,68	0
85	MG	1	3666	1/1	0.78	0.18	54,54,54,54	0
85	MG	6	1965	1/1	0.78	0.40	86,86,86,86	0
85	MG	6	2030	1/1	0.78	0.28	74,74,74,74	0
86	OHX	5	4236	7/7	0.78	0.18	158,158,158,158	0
85	MG	5	3710	1/1	0.78	0.38	77,77,77,77	0
86	OHX	14	403	7/7	0.78	0.19	154,154,154,154	0
85	MG	2	2016	1/1	0.78	0.40	81,81,81,81	0
85	MG	6	1904	1/1	0.79	0.32	75,75,75,75	0
86	OHX	1	4177	7/7	0.79	0.22	148,148,148,148	0
85	MG	6	1959	1/1	0.79	0.31	44,44,44,44	0
86	OHX	1	4183	7/7	0.79	0.19	159,159,159,159	0
85	MG	c9	201	1/1	0.79	0.29	74,74,74,74	0
85	MG	2	2000	1/1	0.79	0.17	104,104,104,104	0
86	OHX	1	4200	7/7	0.79	0.19	154,154,154,154	0
85	MG	6	1911	1/1	0.79	0.40	87,87,87,87	0
86	OHX	6	2189	7/7	0.79	0.19	153,153,153,153	0
85	MG	5	3892	1/1	0.79	0.31	69,69,69,69	0
85	MG	1	3674	1/1	0.79	0.35	68,68,68,68	0
86	OHX	5	4193	7/7	0.79	0.17	156,156,156,156	0
85	MG	1	3703	1/1	0.79	0.26	43,43,43,43	0
85	MG	2	1907	1/1	0.79	0.40	62,62,62,62	0
85	MG	5	3571	1/1	0.79	0.32	35,35,35,35	0
85	MG	6	1967	1/1	0.79	0.26	71,71,71,71	0
85	MG	1	3500	1/1	0.79	0.25	73,73,73,73	0
86	OHX	2	2163	7/7	0.79	0.22	154,154,154,154	0
86	OHX	5	4244	7/7	0.79	0.21	147,147,147,147	0
85	MG	2	1961	1/1	0.79	0.41	77,77,77,77	0
86	OHX	1	4132	7/7	0.79	0.18	151,151,151,151	0
85	MG	5	3476	1/1	0.80	0.27	81,81,81,81	0
85	MG	2	1998	1/1	0.80	0.35	74,74,74,74	0
85	MG	5	3493	1/1	0.80	0.30	62,62,62,62	0
85	MG	6	2028	1/1	0.80	0.42	83,83,83,83	0
85	MG	1	3863	1/1	0.80	0.30	46,46,46,46	0
85	MG	5	3607	1/1	0.80	0.18	45,45,45,45	0
85	MG	2	1969	1/1	0.80	0.24	73,73,73,73	0
85	MG	6	1950	1/1	0.80	0.40	72,72,72,72	0
85	MG	5	3859	1/1	0.80	0.20	93,93,93,93	0
86	OHX	4	239	7/7	0.80	0.22	144,144,144,144	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	6	2170	7/7	0.80	0.17	163,163,163,163	0
85	MG	5	3658	1/1	0.80	0.19	58,58,58,58	0
86	OHX	6	2186	7/7	0.80	0.15	181,181,181,181	0
85	MG	M3	202	1/1	0.80	0.28	86,86,86,86	0
86	OHX	6	2191	7/7	0.80	0.19	150,150,150,150	0
85	MG	5	3676	1/1	0.80	0.26	76,76,76,76	0
86	OHX	6	2194	7/7	0.80	0.14	192,192,192,192	0
85	MG	3	202	1/1	0.80	0.17	51,51,51,51	0
85	MG	5	3893	1/1	0.80	0.12	125,125,125,125	0
86	OHX	5	4201	7/7	0.80	0.20	146,146,146,146	0
85	MG	2	1970	1/1	0.80	0.39	71,71,71,71	0
86	OHX	5	4223	7/7	0.80	0.18	156,156,156,156	0
86	OHX	5	4225	7/7	0.80	0.17	167,167,167,167	0
85	MG	5	3695	1/1	0.80	0.23	55,55,55,55	0
86	OHX	5	4228	7/7	0.80	0.17	168,168,168,168	0
85	MG	7	215	1/1	0.80	0.22	62,62,62,62	0
85	MG	8	211	1/1	0.80	0.20	63,63,63,63	0
85	MG	2	2013	1/1	0.80	0.28	62,62,62,62	0
85	MG	5	3729	1/1	0.80	0.19	100,100,100,100	0
85	MG	5	3450	1/1	0.80	0.21	61,61,61,61	0
86	OHX	5	4245	7/7	0.80	0.21	139,139,139,139	0
85	MG	1	3700	1/1	0.80	0.14	51,51,51,51	0
85	MG	1	3640	1/1	0.80	0.27	64,64,64,64	0
86	OHX	1	4202	7/7	0.81	0.16	173,173,173,173	0
86	OHX	1	4208	7/7	0.81	0.22	150,150,150,150	0
85	MG	1	3430	1/1	0.81	0.39	49,49,49,49	0
85	MG	7	213	1/1	0.81	0.13	73,73,73,73	0
85	MG	6	2045	1/1	0.81	0.37	79,79,79,79	0
85	MG	5	3619	1/1	0.81	0.23	53,53,53,53	0
85	MG	1	3531	1/1	0.81	0.25	63,63,63,63	0
85	MG	5	3415	1/1	0.81	0.17	59,59,59,59	0
86	OHX	2	2158	7/7	0.81	0.17	167,167,167,167	0
85	MG	1	3848	1/1	0.81	0.35	59,59,59,59	0
86	OHX	6	2201	7/7	0.81	0.18	167,167,167,167	0
85	MG	2	1990	1/1	0.81	0.22	102,102,102,102	0
86	OHX	s1	303	7/7	0.81	0.17	168,168,168,168	0
86	OHX	5	4162	7/7	0.81	0.16	187,187,187,187	0
86	OHX	5	4185	7/7	0.81	0.16	168,168,168,168	0
85	MG	2	1947	1/1	0.81	0.31	92,92,92,92	0
86	OHX	2	2174	7/7	0.81	0.18	178,178,178,178	0
86	OHX	5	4204	7/7	0.81	0.21	154,154,154,154	0
85	MG	2	2008	1/1	0.81	0.39	77,77,77,77	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	5	3846	1/1	0.81	0.13	38,38,38,38	0
86	OHX	1	4146	7/7	0.81	0.23	151,151,151,151	0
85	MG	2	1981	1/1	0.81	0.30	77,77,77,77	0
86	OHX	1	4175	7/7	0.81	0.17	167,167,167,167	0
86	OHX	1	4176	7/7	0.81	0.15	185,185,185,185	0
85	MG	5	3477	1/1	0.81	0.32	39,39,39,39	0
85	MG	5	3480	1/1	0.81	0.32	67,67,67,67	0
86	OHX	5	4237	7/7	0.81	0.16	191,191,191,191	0
85	MG	6	1982	1/1	0.81	0.22	82,82,82,82	0
85	MG	1	3494	1/1	0.81	0.38	79,79,79,79	0
86	OHX	1	4193	7/7	0.81	0.19	148,148,148,148	0
86	OHX	14	402	7/7	0.81	0.16	168,168,168,168	0
85	MG	5	3746	1/1	0.81	0.25	61,61,61,61	0
85	MG	1	3496	1/1	0.81	0.19	53,53,53,53	0
86	OHX	m4	202	7/7	0.81	0.11	223,223,223,223	0
86	OHX	1	4100	7/7	0.82	0.16	156,156,156,156	0
85	MG	6	1922	1/1	0.82	0.31	70,70,70,70	0
85	MG	c7	201	1/1	0.82	0.28	72,72,72,72	0
85	MG	2	1940	1/1	0.82	0.17	71,71,71,71	0
85	MG	2	2005	1/1	0.82	0.43	78,78,78,78	0
85	MG	6	2011	1/1	0.82	0.16	52,52,52,52	0
86	OHX	5	4176	7/7	0.82	0.17	157,157,157,157	0
85	MG	5	3782	1/1	0.82	0.26	97,97,97,97	0
85	MG	2	1963	1/1	0.82	0.25	99,99,99,99	0
85	MG	5	3465	1/1	0.82	0.11	38,38,38,38	0
86	OHX	1	4186	7/7	0.82	0.21	141,141,141,141	0
86	OHX	5	4216	7/7	0.82	0.20	143,143,143,143	0
86	OHX	5	4218	7/7	0.82	0.19	150,150,150,150	0
85	MG	5	3664	1/1	0.82	0.27	36,36,36,36	0
85	MG	1	3667	1/1	0.82	0.17	46,46,46,46	0
85	MG	n8	201	1/1	0.82	0.17	40,40,40,40	0
85	MG	6	1935	1/1	0.82	0.30	78,78,78,78	0
85	MG	6	1938	1/1	0.82	0.43	67,67,67,67	0
85	MG	5	3815	1/1	0.82	0.21	64,64,64,64	0
86	OHX	1	4211	7/7	0.82	0.21	139,139,139,139	0
85	MG	1	3411	1/1	0.82	0.27	50,50,50,50	0
86	OHX	6	2148	7/7	0.82	0.19	149,149,149,149	0
86	OHX	6	2156	7/7	0.82	0.15	181,181,181,181	0
85	MG	1	3774	1/1	0.82	0.22	92,92,92,92	0
85	MG	2	1931	1/1	0.82	0.49	62,62,62,62	0
86	OHX	2	2176	7/7	0.82	0.14	187,187,187,187	0
86	OHX	6	2187	7/7	0.82	0.18	156,156,156,156	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	1	3427	1/1	0.82	0.14	63,63,63,63	0
86	OHX	1	4067	7/7	0.82	0.21	136,136,136,136	0
85	MG	S8	301	1/1	0.83	0.19	57,57,57,57	0
85	MG	1	3421	1/1	0.83	0.38	76,76,76,76	0
86	OHX	2	2099	7/7	0.83	0.18	156,156,156,156	0
86	OHX	2	2115	7/7	0.83	0.19	153,153,153,153	0
86	OHX	2	2148	7/7	0.83	0.15	173,173,173,173	0
85	MG	6	2000	1/1	0.83	0.26	77,77,77,77	0
86	OHX	6	2188	7/7	0.83	0.15	171,171,171,171	0
85	MG	5	3488	1/1	0.83	0.26	55,55,55,55	0
85	MG	3	203	1/1	0.83	0.20	94,94,94,94	0
85	MG	2	1943	1/1	0.83	0.30	69,69,69,69	0
85	MG	2	1964	1/1	0.83	0.26	63,63,63,63	0
85	MG	1	3654	1/1	0.83	0.34	70,70,70,70	0
85	MG	3	212	1/1	0.83	0.36	57,57,57,57	0
85	MG	5	3810	1/1	0.83	0.17	83,83,83,83	0
85	MG	3	213	1/1	0.83	0.22	61,61,61,61	0
86	OHX	1	4116	7/7	0.83	0.19	145,145,145,145	0
85	MG	6	2031	1/1	0.83	0.17	60,60,60,60	0
85	MG	1	3761	1/1	0.83	0.24	51,51,51,51	0
85	MG	1	3445	1/1	0.83	0.32	63,63,63,63	0
86	OHX	1	4174	7/7	0.83	0.21	126,126,126,126	0
85	MG	5	3853	1/1	0.83	0.20	56,56,56,56	0
86	OHX	5	4217	7/7	0.83	0.13	181,181,181,181	0
85	MG	1	3778	1/1	0.83	0.16	59,59,59,59	0
85	MG	5	3876	1/1	0.83	0.31	46,46,46,46	0
86	OHX	5	4221	7/7	0.83	0.23	136,136,136,136	0
85	MG	1	3508	1/1	0.83	0.30	32,32,32,32	0
85	MG	2	2009	1/1	0.83	0.31	66,66,66,66	0
85	MG	5	3678	1/1	0.83	0.18	45,45,45,45	0
85	MG	6	1902	1/1	0.83	0.18	59,59,59,59	0
85	MG	1	3833	1/1	0.83	0.15	50,50,50,50	0
86	OHX	1	4194	7/7	0.83	0.21	140,140,140,140	0
85	MG	5	3688	1/1	0.83	0.41	76,76,76,76	0
86	OHX	1	4198	7/7	0.83	0.15	158,158,158,158	0
85	MG	2	2003	1/1	0.83	0.19	71,71,71,71	0
86	OHX	1	4201	7/7	0.83	0.17	147,147,147,147	0
85	MG	1	3859	1/1	0.83	0.26	70,70,70,70	0
85	MG	1	3689	1/1	0.83	0.23	86,86,86,86	0
85	MG	1	3585	1/1	0.83	0.38	50,50,50,50	0
85	MG	1	3866	1/1	0.83	0.37	70,70,70,70	0
86	OHX	m1	202	7/7	0.83	0.17	160,160,160,160	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	L4	403	7/7	0.83	0.18	143,143,143,143	0
85	MG	o4	201	1/1	0.84	0.36	72,72,72,72	0
85	MG	6	1957	1/1	0.84	0.32	52,52,52,52	0
86	OHX	M7	205	7/7	0.84	0.27	118,118,118,118	0
86	OHX	6	2136	7/7	0.84	0.18	137,137,137,137	0
85	MG	1	3528	1/1	0.84	0.28	36,36,36,36	0
86	OHX	2	2136	7/7	0.84	0.15	175,175,175,175	0
85	MG	5	3487	1/1	0.84	0.25	51,51,51,51	0
86	OHX	6	2179	7/7	0.84	0.17	147,147,147,147	0
86	OHX	2	2152	7/7	0.84	0.14	182,182,182,182	0
86	OHX	2	2157	7/7	0.84	0.10	260,260,260,260	0
85	MG	N3	202	1/1	0.84	0.22	68,68,68,68	0
85	MG	1	3529	1/1	0.84	0.32	46,46,46,46	0
85	MG	1	3444	1/1	0.84	0.18	82,82,82,82	0
86	OHX	6	2190	7/7	0.84	0.18	147,147,147,147	0
85	MG	1	3765	1/1	0.84	0.22	64,64,64,64	0
85	MG	5	3506	1/1	0.84	0.29	46,46,46,46	0
85	MG	1	3561	1/1	0.84	0.30	42,42,42,42	0
85	MG	5	3583	1/1	0.84	0.30	49,49,49,49	0
85	MG	2	1975	1/1	0.84	0.26	62,62,62,62	0
85	MG	5	3612	1/1	0.84	0.11	38,38,38,38	0
86	OHX	5	4139	7/7	0.84	0.18	135,135,135,135	0
86	OHX	5	4151	7/7	0.84	0.23	131,131,131,131	0
86	OHX	1	4105	7/7	0.84	0.15	170,170,170,170	0
85	MG	5	3821	1/1	0.84	0.22	45,45,45,45	0
85	MG	6	2038	1/1	0.84	0.31	59,59,59,59	0
85	MG	5	3625	1/1	0.84	0.27	44,44,44,44	0
86	OHX	5	4199	7/7	0.84	0.16	145,145,145,145	0
85	MG	1	3782	1/1	0.84	0.18	77,77,77,77	0
86	OHX	5	4203	7/7	0.84	0.19	152,152,152,152	0
86	OHX	1	4152	7/7	0.84	0.19	141,141,141,141	0
86	OHX	1	4170	7/7	0.84	0.20	132,132,132,132	0
86	OHX	1	4171	7/7	0.84	0.10	215,215,215,215	0
85	MG	2	1929	1/1	0.84	0.38	69,69,69,69	0
85	MG	5	3860	1/1	0.84	0.15	62,62,62,62	0
85	MG	6	2044	1/1	0.84	0.36	76,76,76,76	0
85	MG	1	3498	1/1	0.84	0.27	44,44,44,44	0
85	MG	1	3826	1/1	0.84	0.20	54,54,54,54	0
85	MG	1	3609	1/1	0.84	0.31	65,65,65,65	0
86	OHX	5	4227	7/7	0.84	0.12	191,191,191,191	0
85	MG	1	3634	1/1	0.84	0.27	72,72,72,72	0
85	MG	L3	403	1/1	0.84	0.19	40,40,40,40	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	5	3684	1/1	0.84	0.38	78,78,78,78	0
85	MG	5	3431	1/1	0.84	0.22	69,69,69,69	0
85	MG	5	3432	1/1	0.84	0.15	44,44,44,44	0
85	MG	2	1996	1/1	0.84	0.13	85,85,85,85	0
85	MG	2	1935	1/1	0.84	0.36	55,55,55,55	0
85	MG	6	2002	1/1	0.84	0.18	97,97,97,97	0
86	OHX	7	227	7/7	0.84	0.16	150,150,150,150	0
85	MG	m7	204	1/1	0.84	0.14	39,39,39,39	0
86	OHX	1	4206	7/7	0.84	0.19	134,134,134,134	0
86	OHX	l5	305	7/7	0.84	0.18	151,151,151,151	0
85	MG	6	2004	1/1	0.84	0.27	60,60,60,60	0
85	MG	6	2009	1/1	0.84	0.30	65,65,65,65	0
86	OHX	1	4213	7/7	0.84	0.20	134,134,134,134	0
85	MG	7	216	1/1	0.85	0.23	57,57,57,57	0
85	MG	2	1995	1/1	0.85	0.22	92,92,92,92	0
85	MG	s2	301	1/1	0.85	0.12	55,55,55,55	0
85	MG	1	3472	1/1	0.85	0.15	43,43,43,43	0
85	MG	4	212	1/1	0.85	0.27	57,57,57,57	0
85	MG	6	1991	1/1	0.85	0.25	55,55,55,55	0
85	MG	5	3633	1/1	0.85	0.41	82,82,82,82	0
86	OHX	1	4185	7/7	0.85	0.20	143,143,143,143	0
85	MG	1	3845	1/1	0.85	0.21	39,39,39,39	0
85	MG	1	3728	1/1	0.85	0.09	52,52,52,52	0
85	MG	1	3564	1/1	0.85	0.22	51,51,51,51	0
86	OHX	5	4184	7/7	0.85	0.23	124,124,124,124	0
86	OHX	2	2144	7/7	0.85	0.17	153,153,153,153	0
86	OHX	2	2147	7/7	0.85	0.21	127,127,127,127	0
85	MG	1	3740	1/1	0.85	0.27	63,63,63,63	0
85	MG	1	3575	1/1	0.85	0.29	43,43,43,43	0
85	MG	5	3668	1/1	0.85	0.16	45,45,45,45	0
85	MG	5	3832	1/1	0.85	0.20	70,70,70,70	0
86	OHX	5	4215	7/7	0.85	0.16	154,154,154,154	0
86	OHX	1	4203	7/7	0.85	0.20	149,149,149,149	0
85	MG	2	1921	1/1	0.85	0.38	55,55,55,55	0
86	OHX	1	4207	7/7	0.85	0.19	147,147,147,147	0
85	MG	Q2	502	1/1	0.85	0.20	69,69,69,69	0
86	OHX	5	4220	7/7	0.85	0.16	151,151,151,151	0
85	MG	2	1959	1/1	0.85	0.16	63,63,63,63	0
85	MG	1	3516	1/1	0.85	0.35	39,39,39,39	0
86	OHX	4	237	7/7	0.85	0.17	150,150,150,150	0
85	MG	6	1907	1/1	0.85	0.29	68,68,68,68	0
85	MG	1	3517	1/1	0.85	0.31	43,43,43,43	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	2	1986	1/1	0.85	0.32	79,79,79,79	0
86	OHX	M7	206	7/7	0.85	0.19	145,145,145,145	0
86	OHX	M9	203	7/7	0.85	0.13	181,181,181,181	0
85	MG	5	3708	1/1	0.85	0.24	52,52,52,52	0
85	MG	6	1917	1/1	0.85	0.37	69,69,69,69	0
85	MG	1	3807	1/1	0.85	0.16	63,63,63,63	0
86	OHX	5	4242	7/7	0.85	0.17	165,165,165,165	0
86	OHX	5	4243	7/7	0.85	0.20	148,148,148,148	0
85	MG	5	3894	1/1	0.85	0.28	57,57,57,57	0
86	OHX	1	4143	7/7	0.85	0.17	151,151,151,151	0
86	OHX	6	2181	7/7	0.85	0.20	141,141,141,141	0
86	OHX	8	228	7/7	0.85	0.17	136,136,136,136	0
86	OHX	6	2184	7/7	0.85	0.17	147,147,147,147	0
85	MG	6	1978	1/1	0.85	0.23	69,69,69,69	0
85	MG	5	3504	1/1	0.85	0.22	45,45,45,45	0
85	MG	2	1910	1/1	0.85	0.29	57,57,57,57	0
86	OHX	1	4166	7/7	0.85	0.14	170,170,170,170	0
85	MG	1	3710	1/1	0.85	0.18	44,44,44,44	0
87	ZN	d7	101	1/1	0.85	0.15	161,161,161,161	0
85	MG	2	1922	1/1	0.86	0.38	66,66,66,66	0
85	MG	sM	301	1/1	0.86	0.16	46,46,46,46	0
85	MG	5	3403	1/1	0.86	0.24	49,49,49,49	0
85	MG	2	2004	1/1	0.86	0.18	68,68,68,68	0
85	MG	5	3420	1/1	0.86	0.22	69,69,69,69	0
85	MG	q1	101	1/1	0.86	0.24	44,44,44,44	0
85	MG	1	3632	1/1	0.86	0.10	38,38,38,38	0
85	MG	5	3424	1/1	0.86	0.24	62,62,62,62	0
86	OHX	6	2106	7/7	0.86	0.17	134,134,134,134	0
86	OHX	2	2133	7/7	0.86	0.17	148,148,148,148	0
85	MG	5	3426	1/1	0.86	0.20	41,41,41,41	0
85	MG	2	1942	1/1	0.86	0.25	74,74,74,74	0
86	OHX	6	2165	7/7	0.86	0.17	156,156,156,156	0
85	MG	1	3811	1/1	0.86	0.21	59,59,59,59	0
86	OHX	6	2175	7/7	0.86	0.15	165,165,165,165	0
85	MG	5	3440	1/1	0.86	0.41	62,62,62,62	0
85	MG	5	3718	1/1	0.86	0.21	65,65,65,65	0
85	MG	6	1927	1/1	0.86	0.32	71,71,71,71	0
85	MG	1	3635	1/1	0.86	0.24	63,63,63,63	0
85	MG	6	1994	1/1	0.86	0.31	71,71,71,71	0
86	OHX	2	2160	7/7	0.86	0.15	163,163,163,163	0
85	MG	5	3749	1/1	0.86	0.32	39,39,39,39	0
85	MG	1	3708	1/1	0.86	0.29	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(Å ²)	Q<0.9
86	OHX	2	2164	7/7	0.86	0.13	181,181,181,181	0
86	OHX	2	2170	7/7	0.86	0.16	153,153,153,153	0
86	OHX	2	2173	7/7	0.86	0.15	170,170,170,170	0
85	MG	6	1998	1/1	0.86	0.12	96,96,96,96	0
86	OHX	6	2197	7/7	0.86	0.12	180,180,180,180	0
85	MG	2	1948	1/1	0.86	0.17	62,62,62,62	0
86	OHX	6	2204	7/7	0.86	0.19	154,154,154,154	0
85	MG	1	3832	1/1	0.86	0.23	52,52,52,52	0
86	OHX	2	2179	7/7	0.86	0.18	168,168,168,168	0
86	OHX	s8	303	7/7	0.86	0.14	172,172,172,172	0
86	OHX	5	4137	7/7	0.86	0.15	147,147,147,147	0
85	MG	5	3481	1/1	0.86	0.24	61,61,61,61	0
86	OHX	1	4099	7/7	0.86	0.18	135,135,135,135	0
85	MG	5	3485	1/1	0.86	0.24	67,67,67,67	0
85	MG	1	3637	1/1	0.86	0.27	74,74,74,74	0
86	OHX	5	4180	7/7	0.86	0.17	138,138,138,138	0
85	MG	1	3431	1/1	0.86	0.20	52,52,52,52	0
85	MG	L8	301	1/1	0.86	0.22	50,50,50,50	0
86	OHX	5	4192	7/7	0.86	0.21	140,140,140,140	0
86	OHX	1	4134	7/7	0.86	0.14	174,174,174,174	0
85	MG	1	3641	1/1	0.86	0.15	70,70,70,70	0
85	MG	5	3495	1/1	0.86	0.15	45,45,45,45	0
85	MG	6	1953	1/1	0.86	0.32	51,51,51,51	0
86	OHX	1	4148	7/7	0.86	0.14	147,147,147,147	0
86	OHX	5	4207	7/7	0.86	0.18	142,142,142,142	0
86	OHX	5	4208	7/7	0.86	0.21	122,122,122,122	0
85	MG	6	1954	1/1	0.86	0.33	40,40,40,40	0
86	OHX	1	4165	7/7	0.86	0.15	156,156,156,156	0
85	MG	2	1912	1/1	0.86	0.26	68,68,68,68	0
85	MG	M9	201	1/1	0.86	0.12	67,67,67,67	0
85	MG	5	3573	1/1	0.86	0.29	40,40,40,40	0
86	OHX	1	4172	7/7	0.86	0.24	122,122,122,122	0
86	OHX	1	4173	7/7	0.86	0.19	129,129,129,129	0
85	MG	M9	202	1/1	0.86	0.23	71,71,71,71	0
85	MG	5	3596	1/1	0.86	0.38	41,41,41,41	0
85	MG	N0	201	1/1	0.86	0.37	49,49,49,49	0
85	MG	5	3873	1/1	0.86	0.27	48,48,48,48	0
85	MG	6	1963	1/1	0.86	0.22	57,57,57,57	0
86	OHX	1	4182	7/7	0.86	0.22	130,130,130,130	0
85	MG	5	3616	1/1	0.86	0.21	51,51,51,51	0
85	MG	1	3491	1/1	0.86	0.10	54,54,54,54	0
85	MG	1	3655	1/1	0.86	0.20	47,47,47,47	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	1	4188	7/7	0.86	0.18	143,143,143,143	0
86	OHX	5	4240	7/7	0.86	0.15	166,166,166,166	0
86	OHX	1	4191	7/7	0.86	0.17	159,159,159,159	0
85	MG	5	3632	1/1	0.86	0.11	45,45,45,45	0
85	MG	1	3522	1/1	0.86	0.20	77,77,77,77	0
85	MG	1	3590	1/1	0.86	0.41	42,42,42,42	0
86	OHX	7	225	7/7	0.86	0.17	154,154,154,154	0
85	MG	7	201	1/1	0.86	0.34	42,42,42,42	0
85	MG	5	3639	1/1	0.86	0.26	53,53,53,53	0
85	MG	7	207	1/1	0.86	0.16	61,61,61,61	0
85	MG	5	3640	1/1	0.86	0.19	62,62,62,62	0
85	MG	1	3773	1/1	0.86	0.33	65,65,65,65	0
85	MG	6	1908	1/1	0.86	0.19	50,50,50,50	0
85	MG	c8	202	1/1	0.86	0.23	63,63,63,63	0
85	MG	8	207	1/1	0.86	0.21	70,70,70,70	0
85	MG	5	3665	1/1	0.86	0.29	56,56,56,56	0
86	OHX	6	2144	7/7	0.87	0.16	145,145,145,145	0
86	OHX	2	2171	7/7	0.87	0.16	157,157,157,157	0
86	OHX	6	2150	7/7	0.87	0.16	145,145,145,145	0
85	MG	5	3626	1/1	0.87	0.24	46,46,46,46	0
86	OHX	6	2160	7/7	0.87	0.18	138,138,138,138	0
85	MG	5	3422	1/1	0.87	0.28	39,39,39,39	0
85	MG	6	2005	1/1	0.87	0.09	59,59,59,59	0
85	MG	5	3847	1/1	0.87	0.13	56,56,56,56	0
86	OHX	6	2178	7/7	0.87	0.20	139,139,139,139	0
85	MG	6	2008	1/1	0.87	0.14	57,57,57,57	0
86	OHX	6	2180	7/7	0.87	0.14	164,164,164,164	0
85	MG	5	3854	1/1	0.87	0.19	84,84,84,84	0
86	OHX	6	2182	7/7	0.87	0.21	147,147,147,147	0
86	OHX	1	4082	7/7	0.87	0.21	129,129,129,129	0
86	OHX	1	4085	7/7	0.87	0.20	125,125,125,125	0
86	OHX	1	4097	7/7	0.87	0.14	156,156,156,156	0
85	MG	1	3401	1/1	0.87	0.31	43,43,43,43	0
85	MG	2	1973	1/1	0.87	0.30	74,74,74,74	0
85	MG	5	3435	1/1	0.87	0.24	75,75,75,75	0
86	OHX	1	4114	7/7	0.87	0.16	147,147,147,147	0
85	MG	5	3439	1/1	0.87	0.18	59,59,59,59	0
86	OHX	1	4130	7/7	0.87	0.21	128,128,128,128	0
85	MG	2	2019	1/1	0.87	0.36	71,71,71,71	0
86	OHX	6	2196	7/7	0.87	0.14	171,171,171,171	0
85	MG	1	3612	1/1	0.87	0.10	45,45,45,45	0
86	OHX	6	2200	7/7	0.87	0.18	152,152,152,152	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	5	3456	1/1	0.87	0.22	51,51,51,51	0
86	OHX	1	4144	7/7	0.87	0.25	123,123,123,123	0
85	MG	5	3888	1/1	0.87	0.24	45,45,45,45	0
85	MG	6	2024	1/1	0.87	0.14	65,65,65,65	0
85	MG	5	3672	1/1	0.87	0.09	42,42,42,42	0
86	OHX	c3	201	7/7	0.87	0.14	159,159,159,159	0
86	OHX	5	4096	7/7	0.87	0.19	127,127,127,127	0
86	OHX	5	4136	7/7	0.87	0.22	134,134,134,134	0
86	OHX	1	4151	7/7	0.87	0.16	159,159,159,159	0
86	OHX	5	4138	7/7	0.87	0.19	134,134,134,134	0
85	MG	6	2026	1/1	0.87	0.24	86,86,86,86	0
86	OHX	1	4163	7/7	0.87	0.15	155,155,155,155	0
86	OHX	5	4153	7/7	0.87	0.17	142,142,142,142	0
86	OHX	5	4158	7/7	0.87	0.18	132,132,132,132	0
85	MG	L7	302	1/1	0.87	0.15	51,51,51,51	0
85	MG	5	3473	1/1	0.87	0.26	41,41,41,41	0
85	MG	1	3788	1/1	0.87	0.21	52,52,52,52	0
85	MG	7	209	1/1	0.87	0.27	45,45,45,45	0
85	MG	5	3686	1/1	0.87	0.08	36,36,36,36	0
85	MG	6	1924	1/1	0.87	0.33	38,38,38,38	0
85	MG	1	3789	1/1	0.87	0.14	46,46,46,46	0
86	OHX	5	4194	7/7	0.87	0.19	129,129,129,129	0
85	MG	M1	202	1/1	0.87	0.22	78,78,78,78	0
85	MG	8	206	1/1	0.87	0.21	50,50,50,50	0
85	MG	5	3707	1/1	0.87	0.14	60,60,60,60	0
86	OHX	1	4179	7/7	0.87	0.13	169,169,169,169	0
85	MG	6	2037	1/1	0.87	0.28	53,53,53,53	0
85	MG	1	3660	1/1	0.87	0.19	48,48,48,48	0
85	MG	m1	201	1/1	0.87	0.23	62,62,62,62	0
85	MG	6	1979	1/1	0.87	0.29	72,72,72,72	0
85	MG	5	3720	1/1	0.87	0.21	61,61,61,61	0
85	MG	6	2040	1/1	0.87	0.40	77,77,77,77	0
86	OHX	1	4190	7/7	0.87	0.21	143,143,143,143	0
85	MG	M7	201	1/1	0.87	0.20	36,36,36,36	0
85	MG	1	3480	1/1	0.87	0.18	81,81,81,81	0
86	OHX	5	4222	7/7	0.87	0.18	154,154,154,154	0
86	OHX	2	2092	7/7	0.87	0.15	155,155,155,155	0
85	MG	1	3510	1/1	0.87	0.27	53,53,53,53	0
85	MG	1	3739	1/1	0.87	0.11	39,39,39,39	0
86	OHX	1	4196	7/7	0.87	0.15	149,149,149,149	0
85	MG	5	3771	1/1	0.87	0.12	106,106,106,106	0
86	OHX	5	4229	7/7	0.87	0.21	130,130,130,130	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	5	4230	7/7	0.87	0.14	171,171,171,171	0
86	OHX	2	2134	7/7	0.87	0.16	142,142,142,142	0
85	MG	6	1940	1/1	0.87	0.23	54,54,54,54	0
85	MG	5	3545	1/1	0.87	0.37	71,71,71,71	0
86	OHX	2	2146	7/7	0.87	0.12	189,189,189,189	0
85	MG	5	3553	1/1	0.87	0.12	41,41,41,41	0
85	MG	c8	201	1/1	0.87	0.32	73,73,73,73	0
86	OHX	2	2150	7/7	0.87	0.12	197,197,197,197	0
85	MG	6	1948	1/1	0.87	0.29	52,52,52,52	0
86	OHX	1	4212	7/7	0.87	0.18	141,141,141,141	0
86	OHX	2	2155	7/7	0.87	0.10	244,244,244,244	0
86	OHX	5	4246	7/7	0.87	0.17	158,158,158,158	0
86	OHX	1	4219	7/7	0.87	0.20	140,140,140,140	0
86	OHX	3	222	7/7	0.87	0.15	172,172,172,172	0
85	MG	5	3795	1/1	0.87	0.20	63,63,63,63	0
86	OHX	8	231	7/7	0.87	0.18	133,133,133,133	0
86	OHX	8	232	7/7	0.87	0.18	142,142,142,142	0
85	MG	2	1991	1/1	0.87	0.15	65,65,65,65	0
85	MG	N3	203	1/1	0.87	0.14	54,54,54,54	0
85	MG	2	1978	1/1	0.87	0.30	60,60,60,60	0
85	MG	5	3405	1/1	0.87	0.21	32,32,32,32	0
86	OHX	m0	302	7/7	0.87	0.18	136,136,136,136	0
85	MG	1	3584	1/1	0.87	0.27	45,45,45,45	0
85	MG	4	201	1/1	0.87	0.23	49,49,49,49	0
86	OHX	m8	201	7/7	0.87	0.18	145,145,145,145	0
85	MG	2	1905	1/1	0.87	0.38	66,66,66,66	0
86	OHX	6	2167	7/7	0.88	0.12	180,180,180,180	0
86	OHX	6	2169	7/7	0.88	0.14	167,167,167,167	0
85	MG	5	3556	1/1	0.88	0.29	44,44,44,44	0
86	OHX	6	2173	7/7	0.88	0.18	153,153,153,153	0
85	MG	1	3742	1/1	0.88	0.33	70,70,70,70	0
85	MG	1	3754	1/1	0.88	0.24	50,50,50,50	0
86	OHX	S8	302	7/7	0.88	0.12	175,175,175,175	0
85	MG	1	3639	1/1	0.88	0.25	61,61,61,61	0
86	OHX	1	4068	7/7	0.88	0.19	131,131,131,131	0
85	MG	5	3591	1/1	0.88	0.26	35,35,35,35	0
85	MG	5	3844	1/1	0.88	0.26	58,58,58,58	0
85	MG	1	3537	1/1	0.88	0.29	45,45,45,45	0
85	MG	4	213	1/1	0.88	0.23	53,53,53,53	0
85	MG	5	3848	1/1	0.88	0.21	50,50,50,50	0
85	MG	1	3548	1/1	0.88	0.16	49,49,49,49	0
85	MG	1	3646	1/1	0.88	0.26	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	1	3652	1/1	0.88	0.21	70,70,70,70	0
86	OHX	1	4123	7/7	0.88	0.17	138,138,138,138	0
85	MG	5	3624	1/1	0.88	0.13	58,58,58,58	0
86	OHX	6	2193	7/7	0.88	0.14	169,169,169,169	0
85	MG	1	3549	1/1	0.88	0.27	51,51,51,51	0
86	OHX	6	2195	7/7	0.88	0.12	193,193,193,193	0
85	MG	1	3780	1/1	0.88	0.15	58,58,58,58	0
86	OHX	1	4139	7/7	0.88	0.19	122,122,122,122	0
85	MG	5	3630	1/1	0.88	0.19	65,65,65,65	0
85	MG	5	3884	1/1	0.88	0.23	63,63,63,63	0
85	MG	1	3490	1/1	0.88	0.25	57,57,57,57	0
85	MG	2	1983	1/1	0.88	0.15	63,63,63,63	0
85	MG	2	1908	1/1	0.88	0.25	74,74,74,74	0
85	MG	5	3638	1/1	0.88	0.25	46,46,46,46	0
85	MG	6	1968	1/1	0.88	0.21	60,60,60,60	0
86	OHX	d4	201	7/7	0.88	0.13	157,157,157,157	0
86	OHX	d9	102	7/7	0.88	0.14	169,169,169,169	0
86	OHX	1	4154	7/7	0.88	0.15	145,145,145,145	0
86	OHX	5	4106	7/7	0.88	0.19	131,131,131,131	0
86	OHX	5	4117	7/7	0.88	0.17	133,133,133,133	0
86	OHX	1	4155	7/7	0.88	0.14	157,157,157,157	0
86	OHX	1	4158	7/7	0.88	0.16	154,154,154,154	0
85	MG	1	3404	1/1	0.88	0.35	57,57,57,57	0
85	MG	5	3895	1/1	0.88	0.28	69,69,69,69	0
86	OHX	5	4150	7/7	0.88	0.21	120,120,120,120	0
85	MG	5	3649	1/1	0.88	0.28	38,38,38,38	0
86	OHX	1	4167	7/7	0.88	0.18	147,147,147,147	0
86	OHX	5	4154	7/7	0.88	0.16	146,146,146,146	0
86	OHX	5	4156	7/7	0.88	0.20	120,120,120,120	0
86	OHX	5	4157	7/7	0.88	0.19	134,134,134,134	0
85	MG	5	3411	1/1	0.88	0.32	41,41,41,41	0
85	MG	1	3441	1/1	0.88	0.14	45,45,45,45	0
86	OHX	5	4166	7/7	0.88	0.19	136,136,136,136	0
86	OHX	5	4168	7/7	0.88	0.14	173,173,173,173	0
85	MG	5	3659	1/1	0.88	0.20	46,46,46,46	0
85	MG	6	1976	1/1	0.88	0.23	51,51,51,51	0
85	MG	1	3812	1/1	0.88	0.18	60,60,60,60	0
85	MG	2	1925	1/1	0.88	0.34	67,67,67,67	0
86	OHX	5	4190	7/7	0.88	0.18	130,130,130,130	0
85	MG	1	3671	1/1	0.88	0.18	80,80,80,80	0
85	MG	2	2010	1/1	0.88	0.35	72,72,72,72	0
85	MG	1	3592	1/1	0.88	0.44	55,55,55,55	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	6	1986	1/1	0.88	0.10	48,48,48,48	0
86	OHX	5	4200	7/7	0.88	0.21	138,138,138,138	0
85	MG	1	3690	1/1	0.88	0.23	57,57,57,57	0
86	OHX	5	4202	7/7	0.88	0.18	141,141,141,141	0
85	MG	1	3693	1/1	0.88	0.09	37,37,37,37	0
85	MG	2	2012	1/1	0.88	0.28	69,69,69,69	0
85	MG	6	1993	1/1	0.88	0.11	46,46,46,46	0
85	MG	n8	203	1/1	0.88	0.27	46,46,46,46	0
86	OHX	5	4209	7/7	0.88	0.15	148,148,148,148	0
85	MG	1	3849	1/1	0.88	0.20	50,50,50,50	0
85	MG	5	3693	1/1	0.88	0.24	50,50,50,50	0
85	MG	1	3852	1/1	0.88	0.32	63,63,63,63	0
85	MG	5	3704	1/1	0.88	0.15	61,61,61,61	0
85	MG	1	3854	1/1	0.88	0.18	49,49,49,49	0
85	MG	6	1914	1/1	0.88	0.33	76,76,76,76	0
86	OHX	2	2119	7/7	0.88	0.17	144,144,144,144	0
86	OHX	2	2126	7/7	0.88	0.16	141,141,141,141	0
86	OHX	2	2132	7/7	0.88	0.13	162,162,162,162	0
85	MG	1	3414	1/1	0.88	0.32	57,57,57,57	0
85	MG	6	2003	1/1	0.88	0.28	72,72,72,72	0
85	MG	1	3707	1/1	0.88	0.24	49,49,49,49	0
85	MG	1	3418	1/1	0.88	0.24	44,44,44,44	0
85	MG	1	3620	1/1	0.88	0.23	59,59,59,59	0
85	MG	1	3868	1/1	0.88	0.13	110,110,110,110	0
86	OHX	1	4209	7/7	0.88	0.16	139,139,139,139	0
86	OHX	5	4233	7/7	0.88	0.20	138,138,138,138	0
86	OHX	1	4210	7/7	0.88	0.17	139,139,139,139	0
86	OHX	5	4235	7/7	0.88	0.15	151,151,151,151	0
85	MG	1	3871	1/1	0.88	0.25	69,69,69,69	0
85	MG	1	3626	1/1	0.88	0.35	72,72,72,72	0
85	MG	5	3763	1/1	0.88	0.20	73,73,73,73	0
86	OHX	1	4216	7/7	0.88	0.19	146,146,146,146	0
86	OHX	1	4218	7/7	0.88	0.12	175,175,175,175	0
86	OHX	2	2153	7/7	0.88	0.14	158,158,158,158	0
86	OHX	2	2154	7/7	0.88	0.16	152,152,152,152	0
86	OHX	4	236	7/7	0.88	0.19	123,123,123,123	0
85	MG	1	3478	1/1	0.88	0.17	51,51,51,51	0
85	MG	1	3419	1/1	0.88	0.19	55,55,55,55	0
85	MG	S6	301	1/1	0.88	0.25	93,93,93,93	0
86	OHX	8	227	7/7	0.88	0.17	150,150,150,150	0
85	MG	2	1920	1/1	0.88	0.27	63,63,63,63	0
85	MG	1	3536	1/1	0.88	0.33	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	2	2161	7/7	0.88	0.14	171,171,171,171	0
85	MG	6	1945	1/1	0.88	0.36	65,65,65,65	0
86	OHX	6	2133	7/7	0.88	0.15	166,166,166,166	0
85	MG	5	3529	1/1	0.88	0.47	58,58,58,58	0
85	MG	5	3799	1/1	0.88	0.23	78,78,78,78	0
86	OHX	19	202	7/7	0.88	0.17	132,132,132,132	0
85	MG	5	3535	1/1	0.88	0.20	53,53,53,53	0
85	MG	1	3741	1/1	0.88	0.13	35,35,35,35	0
86	OHX	2	2172	7/7	0.88	0.15	157,157,157,157	0
86	OHX	m7	206	7/7	0.88	0.20	125,125,125,125	0
85	MG	5	3552	1/1	0.88	0.23	37,37,37,37	0
85	MG	6	1949	1/1	0.88	0.23	47,47,47,47	0
85	MG	M7	203	1/1	0.89	0.12	36,36,36,36	0
85	MG	5	3724	1/1	0.89	0.19	51,51,51,51	0
86	OHX	1	4136	7/7	0.89	0.14	166,166,166,166	0
85	MG	1	3525	1/1	0.89	0.13	43,43,43,43	0
86	OHX	1	4142	7/7	0.89	0.17	135,135,135,135	0
85	MG	5	3738	1/1	0.89	0.21	49,49,49,49	0
85	MG	5	3595	1/1	0.89	0.32	40,40,40,40	0
86	OHX	1	4145	7/7	0.89	0.17	134,134,134,134	0
85	MG	5	3745	1/1	0.89	0.28	46,46,46,46	0
85	MG	1	3451	1/1	0.89	0.28	46,46,46,46	0
85	MG	8	212	1/1	0.89	0.26	43,43,43,43	0
86	OHX	1	4150	7/7	0.89	0.14	161,161,161,161	0
85	MG	8	214	1/1	0.89	0.27	57,57,57,57	0
85	MG	12	302	1/1	0.89	0.10	45,45,45,45	0
86	OHX	1	4153	7/7	0.89	0.14	149,149,149,149	0
85	MG	1	3456	1/1	0.89	0.23	55,55,55,55	0
85	MG	2	1902	1/1	0.89	0.23	48,48,48,48	0
85	MG	m6	201	1/1	0.89	0.15	77,77,77,77	0
85	MG	5	3758	1/1	0.89	0.07	48,48,48,48	0
86	OHX	5	4085	7/7	0.89	0.23	119,119,119,119	0
85	MG	5	3762	1/1	0.89	0.17	54,54,54,54	0
85	MG	1	3534	1/1	0.89	0.14	38,38,38,38	0
86	OHX	5	4108	7/7	0.89	0.15	146,146,146,146	0
85	MG	5	3768	1/1	0.89	0.07	43,43,43,43	0
86	OHX	5	4135	7/7	0.89	0.15	137,137,137,137	0
86	OHX	1	4169	7/7	0.89	0.13	169,169,169,169	0
85	MG	O4	201	1/1	0.89	0.20	63,63,63,63	0
85	MG	5	3621	1/1	0.89	0.13	46,46,46,46	0
86	OHX	2	2090	7/7	0.89	0.15	146,146,146,146	0
86	OHX	5	4145	7/7	0.89	0.18	134,134,134,134	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	1	3801	1/1	0.89	0.19	61,61,61,61	0
85	MG	1	3804	1/1	0.89	0.21	56,56,56,56	0
85	MG	5	3784	1/1	0.89	0.21	62,62,62,62	0
85	MG	5	3785	1/1	0.89	0.14	34,34,34,34	0
86	OHX	5	4155	7/7	0.89	0.17	141,141,141,141	0
85	MG	1	3721	1/1	0.89	0.17	79,79,79,79	0
85	MG	5	3790	1/1	0.89	0.11	40,40,40,40	0
86	OHX	1	4180	7/7	0.89	0.14	155,155,155,155	0
85	MG	2	1911	1/1	0.89	0.41	66,66,66,66	0
85	MG	5	3458	1/1	0.89	0.20	43,43,43,43	0
86	OHX	2	2135	7/7	0.89	0.16	144,144,144,144	0
86	OHX	5	4174	7/7	0.89	0.16	139,139,139,139	0
85	MG	2	1916	1/1	0.89	0.29	51,51,51,51	0
86	OHX	5	4179	7/7	0.89	0.14	153,153,153,153	0
86	OHX	2	2137	7/7	0.89	0.14	146,146,146,146	0
86	OHX	5	4181	7/7	0.89	0.18	126,126,126,126	0
86	OHX	5	4183	7/7	0.89	0.18	135,135,135,135	0
86	OHX	2	2138	7/7	0.89	0.15	177,177,177,177	0
86	OHX	2	2139	7/7	0.89	0.14	170,170,170,170	0
86	OHX	5	4189	7/7	0.89	0.19	133,133,133,133	0
86	OHX	2	2143	7/7	0.89	0.12	177,177,177,177	0
85	MG	6	2027	1/1	0.89	0.30	59,59,59,59	0
85	MG	5	3637	1/1	0.89	0.17	56,56,56,56	0
85	MG	5	3469	1/1	0.89	0.14	41,41,41,41	0
85	MG	1	3610	1/1	0.89	0.11	46,46,46,46	0
85	MG	1	3738	1/1	0.89	0.19	65,65,65,65	0
85	MG	5	3648	1/1	0.89	0.13	54,54,54,54	0
85	MG	1	3546	1/1	0.89	0.21	47,47,47,47	0
85	MG	6	1915	1/1	0.89	0.41	62,62,62,62	0
85	MG	6	1916	1/1	0.89	0.27	55,55,55,55	0
86	OHX	2	2156	7/7	0.89	0.20	125,125,125,125	0
86	OHX	1	4205	7/7	0.89	0.18	133,133,133,133	0
85	MG	5	3842	1/1	0.89	0.18	59,59,59,59	0
86	OHX	5	4210	7/7	0.89	0.17	148,148,148,148	0
86	OHX	5	4211	7/7	0.89	0.20	130,130,130,130	0
85	MG	6	1971	1/1	0.89	0.20	52,52,52,52	0
85	MG	5	3663	1/1	0.89	0.13	49,49,49,49	0
85	MG	1	3830	1/1	0.89	0.14	45,45,45,45	0
85	MG	6	1921	1/1	0.89	0.33	55,55,55,55	0
85	MG	6	1977	1/1	0.89	0.17	78,78,78,78	0
85	MG	2	1917	1/1	0.89	0.39	59,59,59,59	0
85	MG	S2	302	1/1	0.89	0.38	76,76,76,76	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	1	4214	7/7	0.89	0.18	138,138,138,138	0
85	MG	4	218	1/1	0.89	0.26	60,60,60,60	0
86	OHX	5	4224	7/7	0.89	0.16	139,139,139,139	0
85	MG	5	3867	1/1	0.89	0.10	45,45,45,45	0
85	MG	5	3870	1/1	0.89	0.34	43,43,43,43	0
86	OHX	1	4220	7/7	0.89	0.18	158,158,158,158	0
85	MG	5	3496	1/1	0.89	0.23	28,28,28,28	0
85	MG	5	3874	1/1	0.89	0.32	56,56,56,56	0
85	MG	1	3836	1/1	0.89	0.21	56,56,56,56	0
85	MG	1	3840	1/1	0.89	0.25	49,49,49,49	0
85	MG	1	3551	1/1	0.89	0.25	44,44,44,44	0
85	MG	1	3555	1/1	0.89	0.21	37,37,37,37	0
86	OHX	D3	202	7/7	0.89	0.16	148,148,148,148	0
86	OHX	1	4033	7/7	0.89	0.18	126,126,126,126	0
86	OHX	1	4066	7/7	0.89	0.13	156,156,156,156	0
85	MG	2	1999	1/1	0.89	0.23	83,83,83,83	0
85	MG	5	3543	1/1	0.89	0.35	35,35,35,35	0
86	OHX	5	4241	7/7	0.89	0.18	142,142,142,142	0
86	OHX	1	4074	7/7	0.89	0.18	126,126,126,126	0
86	OHX	1	4077	7/7	0.89	0.24	114,114,114,114	0
85	MG	5	3889	1/1	0.89	0.11	38,38,38,38	0
86	OHX	6	2153	7/7	0.89	0.14	156,156,156,156	0
85	MG	2	2015	1/1	0.89	0.37	74,74,74,74	0
86	OHX	1	4086	7/7	0.89	0.14	150,150,150,150	0
85	MG	5	3701	1/1	0.89	0.20	40,40,40,40	0
85	MG	1	3691	1/1	0.89	0.12	41,41,41,41	0
85	MG	1	3565	1/1	0.89	0.22	40,40,40,40	0
86	OHX	1	4102	7/7	0.89	0.14	152,152,152,152	0
85	MG	6	1941	1/1	0.89	0.14	34,34,34,34	0
86	OHX	6	2174	7/7	0.89	0.20	135,135,135,135	0
85	MG	2	2006	1/1	0.89	0.41	52,52,52,52	0
85	MG	7	205	1/1	0.89	0.31	28,28,28,28	0
86	OHX	1	4117	7/7	0.89	0.23	123,123,123,123	0
86	OHX	1	4118	7/7	0.89	0.12	186,186,186,186	0
85	MG	5	3716	1/1	0.89	0.20	56,56,56,56	0
86	OHX	1	4124	7/7	0.89	0.19	122,122,122,122	0
86	OHX	1	4125	7/7	0.89	0.17	138,138,138,138	0
86	OHX	1	4126	7/7	0.89	0.14	152,152,152,152	0
86	OHX	1	4128	7/7	0.89	0.16	147,147,147,147	0
85	MG	6	1999	1/1	0.89	0.09	54,54,54,54	0
85	MG	5	3448	1/1	0.90	0.10	36,36,36,36	0
86	OHX	1	4107	7/7	0.90	0.16	144,144,144,144	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	6	1990	1/1	0.90	0.08	53,53,53,53	0
85	MG	1	3545	1/1	0.90	0.19	35,35,35,35	0
85	MG	6	1992	1/1	0.90	0.18	66,66,66,66	0
85	MG	1	3755	1/1	0.90	0.23	65,65,65,65	0
86	OHX	1	4120	7/7	0.90	0.17	133,133,133,133	0
85	MG	5	3466	1/1	0.90	0.26	63,63,63,63	0
85	MG	5	3682	1/1	0.90	0.23	46,46,46,46	0
85	MG	6	1918	1/1	0.90	0.30	41,41,41,41	0
85	MG	6	1919	1/1	0.90	0.25	54,54,54,54	0
85	MG	7	214	1/1	0.90	0.14	53,53,53,53	0
85	MG	1	3469	1/1	0.90	0.32	46,46,46,46	0
86	OHX	1	4131	7/7	0.90	0.14	153,153,153,153	0
85	MG	1	3678	1/1	0.90	0.30	66,66,66,66	0
85	MG	8	205	1/1	0.90	0.12	41,41,41,41	0
86	OHX	1	4135	7/7	0.90	0.14	159,159,159,159	0
85	MG	1	3763	1/1	0.90	0.09	52,52,52,52	0
86	OHX	6	2199	7/7	0.90	0.18	147,147,147,147	0
86	OHX	1	4137	7/7	0.90	0.17	126,126,126,126	0
86	OHX	1	4138	7/7	0.90	0.18	141,141,141,141	0
85	MG	6	2001	1/1	0.90	0.14	75,75,75,75	0
85	MG	8	209	1/1	0.90	0.10	61,61,61,61	0
85	MG	8	210	1/1	0.90	0.12	44,44,44,44	0
85	MG	1	3683	1/1	0.90	0.20	67,67,67,67	0
85	MG	1	3771	1/1	0.90	0.12	56,56,56,56	0
85	MG	5	3483	1/1	0.90	0.26	58,58,58,58	0
85	MG	8	215	1/1	0.90	0.25	38,38,38,38	0
85	MG	6	1928	1/1	0.90	0.27	62,62,62,62	0
86	OHX	5	4093	7/7	0.90	0.14	156,156,156,156	0
85	MG	l5	301	1/1	0.90	0.26	65,65,65,65	0
86	OHX	5	4105	7/7	0.90	0.15	135,135,135,135	0
85	MG	6	1929	1/1	0.90	0.22	60,60,60,60	0
85	MG	6	1931	1/1	0.90	0.22	47,47,47,47	0
86	OHX	5	4116	7/7	0.90	0.17	124,124,124,124	0
85	MG	m4	201	1/1	0.90	0.11	46,46,46,46	0
85	MG	1	3772	1/1	0.90	0.16	57,57,57,57	0
85	MG	1	3547	1/1	0.90	0.30	66,66,66,66	0
85	MG	1	3470	1/1	0.90	0.26	54,54,54,54	0
85	MG	5	3725	1/1	0.90	0.18	41,41,41,41	0
86	OHX	1	4164	7/7	0.90	0.16	138,138,138,138	0
86	OHX	5	4144	7/7	0.90	0.19	122,122,122,122	0
85	MG	6	2018	1/1	0.90	0.16	49,49,49,49	0
85	MG	o3	202	1/1	0.90	0.25	37,37,37,37	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	5	3733	1/1	0.90	0.28	60,60,60,60	0
86	OHX	1	4168	7/7	0.90	0.13	152,152,152,152	0
85	MG	5	3736	1/1	0.90	0.13	47,47,47,47	0
85	MG	1	3503	1/1	0.90	0.40	45,45,45,45	0
85	MG	6	2022	1/1	0.90	0.16	62,62,62,62	0
85	MG	1	3504	1/1	0.90	0.25	29,29,29,29	0
86	OHX	2	2107	7/7	0.90	0.15	141,141,141,141	0
86	OHX	5	4161	7/7	0.90	0.15	148,148,148,148	0
86	OHX	2	2111	7/7	0.90	0.12	170,170,170,170	0
86	OHX	5	4164	7/7	0.90	0.17	132,132,132,132	0
85	MG	1	3413	1/1	0.90	0.26	42,42,42,42	0
85	MG	1	3701	1/1	0.90	0.11	64,64,64,64	0
86	OHX	5	4173	7/7	0.90	0.16	141,141,141,141	0
85	MG	5	3536	1/1	0.90	0.21	41,41,41,41	0
85	MG	5	3538	1/1	0.90	0.36	48,48,48,48	0
86	OHX	5	4178	7/7	0.90	0.16	142,142,142,142	0
85	MG	1	3702	1/1	0.90	0.33	66,66,66,66	0
85	MG	1	3794	1/1	0.90	0.14	46,46,46,46	0
85	MG	5	3547	1/1	0.90	0.41	55,55,55,55	0
86	OHX	5	4182	7/7	0.90	0.18	130,130,130,130	0
85	MG	1	3798	1/1	0.90	0.18	82,82,82,82	0
85	MG	4	219	1/1	0.90	0.21	48,48,48,48	0
85	MG	2	2002	1/1	0.90	0.19	85,85,85,85	0
86	OHX	5	4186	7/7	0.90	0.18	131,131,131,131	0
86	OHX	1	4187	7/7	0.90	0.20	113,113,113,113	0
85	MG	5	3557	1/1	0.90	0.18	46,46,46,46	0
86	OHX	1	4189	7/7	0.90	0.14	153,153,153,153	0
86	OHX	2	2141	7/7	0.90	0.15	143,143,143,143	0
85	MG	5	3561	1/1	0.90	0.31	38,38,38,38	0
85	MG	6	2035	1/1	0.90	0.17	62,62,62,62	0
85	MG	1	3439	1/1	0.90	0.21	52,52,52,52	0
85	MG	5	3575	1/1	0.90	0.40	33,33,33,33	0
85	MG	5	3579	1/1	0.90	0.39	48,48,48,48	0
85	MG	1	3416	1/1	0.90	0.10	54,54,54,54	0
85	MG	2	1944	1/1	0.90	0.19	69,69,69,69	0
86	OHX	5	4205	7/7	0.90	0.14	153,153,153,153	0
85	MG	6	1956	1/1	0.90	0.42	57,57,57,57	0
85	MG	1	3645	1/1	0.90	0.21	45,45,45,45	0
85	MG	5	3601	1/1	0.90	0.19	47,47,47,47	0
85	MG	M0	303	1/1	0.90	0.20	38,38,38,38	0
85	MG	1	3814	1/1	0.90	0.18	44,44,44,44	0
86	OHX	5	4212	7/7	0.90	0.10	207,207,207,207	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	1	3720	1/1	0.90	0.10	51,51,51,51	0
85	MG	1	3524	1/1	0.90	0.21	27,27,27,27	0
85	MG	1	3405	1/1	0.90	0.25	66,66,66,66	0
85	MG	1	3829	1/1	0.90	0.09	53,53,53,53	0
85	MG	5	3834	1/1	0.90	0.09	42,42,42,42	0
85	MG	1	3727	1/1	0.90	0.10	57,57,57,57	0
85	MG	1	3483	1/1	0.90	0.21	34,34,34,34	0
86	OHX	2	2167	7/7	0.90	0.18	132,132,132,132	0
86	OHX	2	2168	7/7	0.90	0.14	163,163,163,163	0
85	MG	5	3628	1/1	0.90	0.22	62,62,62,62	0
86	OHX	1	4217	7/7	0.90	0.20	132,132,132,132	0
85	MG	5	3401	1/1	0.90	0.15	61,61,61,61	0
85	MG	1	3733	1/1	0.90	0.21	35,35,35,35	0
85	MG	5	3849	1/1	0.90	0.21	61,61,61,61	0
85	MG	1	3734	1/1	0.90	0.18	40,40,40,40	0
86	OHX	2	2175	7/7	0.90	0.15	145,145,145,145	0
85	MG	5	3410	1/1	0.90	0.25	48,48,48,48	0
86	OHX	4	238	7/7	0.90	0.16	149,149,149,149	0
85	MG	5	3858	1/1	0.90	0.12	49,49,49,49	0
86	OHX	L3	406	7/7	0.90	0.13	164,164,164,164	0
86	OHX	2	2178	7/7	0.90	0.19	154,154,154,154	0
85	MG	2	1994	1/1	0.90	0.34	63,63,63,63	0
85	MG	N5	201	1/1	0.90	0.25	69,69,69,69	0
85	MG	1	3843	1/1	0.90	0.31	50,50,50,50	0
86	OHX	D9	102	7/7	0.90	0.16	149,149,149,149	0
85	MG	5	3868	1/1	0.90	0.14	40,40,40,40	0
86	OHX	1	4041	7/7	0.90	0.17	133,133,133,133	0
85	MG	2	1936	1/1	0.90	0.31	56,56,56,56	0
85	MG	5	3642	1/1	0.90	0.20	41,41,41,41	0
85	MG	1	3493	1/1	0.90	0.14	72,72,72,72	0
86	OHX	5	4247	7/7	0.90	0.15	145,145,145,145	0
86	OHX	5	4248	7/7	0.90	0.14	169,169,169,169	0
86	OHX	6	2152	7/7	0.90	0.14	161,161,161,161	0
86	OHX	1	4072	7/7	0.90	0.16	136,136,136,136	0
86	OHX	6	2154	7/7	0.90	0.13	143,143,143,143	0
85	MG	1	3608	1/1	0.90	0.19	70,70,70,70	0
86	OHX	6	2158	7/7	0.90	0.14	146,146,146,146	0
85	MG	1	3850	1/1	0.90	0.23	60,60,60,60	0
86	OHX	6	2164	7/7	0.90	0.11	192,192,192,192	0
85	MG	5	3657	1/1	0.90	0.27	68,68,68,68	0
85	MG	1	3460	1/1	0.90	0.14	57,57,57,57	0
85	MG	1	3853	1/1	0.90	0.24	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	1	4092	7/7	0.90	0.18	132,132,132,132	0
86	OHX	6	2172	7/7	0.90	0.13	153,153,153,153	0
85	MG	5	3660	1/1	0.90	0.18	47,47,47,47	0
85	MG	2	1941	1/1	0.90	0.35	77,77,77,77	0
85	MG	1	3743	1/1	0.90	0.21	57,57,57,57	0
86	OHX	6	2176	7/7	0.90	0.20	112,112,112,112	0
87	ZN	D7	101	1/1	0.90	0.14	167,167,167,167	0
85	MG	1	3745	1/1	0.90	0.29	63,63,63,63	0
88	3K5	5	4249	57/57	0.90	0.14	52,52,52,52	0
85	MG	1	3499	1/1	0.91	0.37	34,34,34,34	0
85	MG	5	3551	1/1	0.91	0.37	51,51,51,51	0
85	MG	1	3767	1/1	0.91	0.07	50,50,50,50	0
85	MG	1	3428	1/1	0.91	0.26	42,42,42,42	0
86	OHX	1	4127	7/7	0.91	0.21	115,115,115,115	0
85	MG	5	3735	1/1	0.91	0.32	69,69,69,69	0
85	MG	5	3555	1/1	0.91	0.21	37,37,37,37	0
85	MG	l3	403	1/1	0.91	0.31	36,36,36,36	0
85	MG	1	3409	1/1	0.91	0.27	33,33,33,33	0
86	OHX	1	4133	7/7	0.91	0.16	131,131,131,131	0
85	MG	5	3739	1/1	0.91	0.37	65,65,65,65	0
85	MG	2	2007	1/1	0.91	0.20	51,51,51,51	0
85	MG	5	3560	1/1	0.91	0.27	34,34,34,34	0
85	MG	m5	303	1/1	0.91	0.17	98,98,98,98	0
86	OHX	6	2198	7/7	0.91	0.15	149,149,149,149	0
85	MG	1	3438	1/1	0.91	0.26	33,33,33,33	0
85	MG	m7	201	1/1	0.91	0.26	37,37,37,37	0
86	OHX	1	4140	7/7	0.91	0.17	123,123,123,123	0
85	MG	m7	202	1/1	0.91	0.16	36,36,36,36	0
85	MG	5	3565	1/1	0.91	0.35	32,32,32,32	0
85	MG	n0	201	1/1	0.91	0.16	46,46,46,46	0
85	MG	1	3553	1/1	0.91	0.39	36,36,36,36	0
85	MG	5	3406	1/1	0.91	0.14	42,42,42,42	0
85	MG	2	1954	1/1	0.91	0.20	68,68,68,68	0
85	MG	1	3556	1/1	0.91	0.25	34,34,34,34	0
86	OHX	1	4149	7/7	0.91	0.17	126,126,126,126	0
85	MG	5	3764	1/1	0.91	0.20	44,44,44,44	0
85	MG	5	3767	1/1	0.91	0.09	65,65,65,65	0
86	OHX	5	4097	7/7	0.91	0.15	130,130,130,130	0
86	OHX	2	2080	7/7	0.91	0.15	143,143,143,143	0
85	MG	1	3787	1/1	0.91	0.13	56,56,56,56	0
85	MG	5	3589	1/1	0.91	0.43	56,56,56,56	0
86	OHX	5	4115	7/7	0.91	0.14	130,130,130,130	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	5	3416	1/1	0.91	0.10	42,42,42,42	0
85	MG	5	3779	1/1	0.91	0.26	57,57,57,57	0
86	OHX	5	4121	7/7	0.91	0.12	157,157,157,157	0
86	OHX	5	4125	7/7	0.91	0.15	133,133,133,133	0
86	OHX	5	4126	7/7	0.91	0.13	155,155,155,155	0
86	OHX	5	4134	7/7	0.91	0.18	135,135,135,135	0
85	MG	1	3514	1/1	0.91	0.27	26,26,26,26	0
85	MG	2	1932	1/1	0.91	0.31	61,61,61,61	0
86	OHX	2	2118	7/7	0.91	0.13	154,154,154,154	0
85	MG	1	3790	1/1	0.91	0.29	71,71,71,71	0
86	OHX	2	2121	7/7	0.91	0.13	149,149,149,149	0
86	OHX	5	4141	7/7	0.91	0.16	126,126,126,126	0
86	OHX	5	4142	7/7	0.91	0.17	139,139,139,139	0
86	OHX	2	2123	7/7	0.91	0.15	154,154,154,154	0
86	OHX	2	2125	7/7	0.91	0.14	143,143,143,143	0
86	OHX	5	4148	7/7	0.91	0.20	126,126,126,126	0
86	OHX	5	4149	7/7	0.91	0.14	152,152,152,152	0
85	MG	1	3712	1/1	0.91	0.08	54,54,54,54	0
86	OHX	2	2128	7/7	0.91	0.11	201,201,201,201	0
86	OHX	2	2131	7/7	0.91	0.13	165,165,165,165	0
85	MG	5	3609	1/1	0.91	0.23	33,33,33,33	0
85	MG	1	3443	1/1	0.91	0.28	30,30,30,30	0
85	MG	1	3719	1/1	0.91	0.10	45,45,45,45	0
85	MG	5	3794	1/1	0.91	0.12	49,49,49,49	0
85	MG	1	3519	1/1	0.91	0.30	31,31,31,31	0
86	OHX	5	4159	7/7	0.91	0.17	121,121,121,121	0
85	MG	5	3796	1/1	0.91	0.16	60,60,60,60	0
85	MG	5	3433	1/1	0.91	0.23	43,43,43,43	0
86	OHX	5	4163	7/7	0.91	0.16	145,145,145,145	0
85	MG	1	3649	1/1	0.91	0.10	53,53,53,53	0
85	MG	4	215	1/1	0.91	0.23	59,59,59,59	0
86	OHX	5	4167	7/7	0.91	0.09	197,197,197,197	0
85	MG	1	3723	1/1	0.91	0.40	60,60,60,60	0
86	OHX	5	4172	7/7	0.91	0.17	149,149,149,149	0
86	OHX	1	4184	7/7	0.91	0.07	245,245,245,245	0
85	MG	1	3650	1/1	0.91	0.07	32,32,32,32	0
86	OHX	5	4175	7/7	0.91	0.20	119,119,119,119	0
85	MG	1	3813	1/1	0.91	0.13	39,39,39,39	0
85	MG	D3	201	1/1	0.91	0.10	65,65,65,65	0
85	MG	2	1919	1/1	0.91	0.41	71,71,71,71	0
85	MG	5	3460	1/1	0.91	0.17	33,33,33,33	0
85	MG	5	3827	1/1	0.91	0.17	40,40,40,40	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	1	3822	1/1	0.91	0.18	50,50,50,50	0
85	MG	1	3730	1/1	0.91	0.12	64,64,64,64	0
85	MG	6	1946	1/1	0.91	0.29	53,53,53,53	0
85	MG	1	3486	1/1	0.91	0.33	44,44,44,44	0
85	MG	1	3828	1/1	0.91	0.15	45,45,45,45	0
86	OHX	5	4188	7/7	0.91	0.15	132,132,132,132	0
85	MG	5	3647	1/1	0.91	0.13	43,43,43,43	0
85	MG	2	1934	1/1	0.91	0.34	61,61,61,61	0
85	MG	1	3656	1/1	0.91	0.28	35,35,35,35	0
85	MG	5	3650	1/1	0.91	0.30	62,62,62,62	0
85	MG	1	3453	1/1	0.91	0.15	39,39,39,39	0
85	MG	5	3855	1/1	0.91	0.12	65,65,65,65	0
86	OHX	1	4204	7/7	0.91	0.15	142,142,142,142	0
85	MG	5	3856	1/1	0.91	0.18	52,52,52,52	0
85	MG	5	3857	1/1	0.91	0.21	68,68,68,68	0
85	MG	2	1984	1/1	0.91	0.15	69,69,69,69	0
85	MG	1	3594	1/1	0.91	0.27	30,30,30,30	0
85	MG	1	3459	1/1	0.91	0.19	43,43,43,43	0
86	OHX	5	4206	7/7	0.91	0.20	131,131,131,131	0
85	MG	1	3495	1/1	0.91	0.19	45,45,45,45	0
85	MG	5	3486	1/1	0.91	0.12	48,48,48,48	0
85	MG	S2	301	1/1	0.91	0.29	48,48,48,48	0
85	MG	1	3675	1/1	0.91	0.26	43,43,43,43	0
85	MG	2	2014	1/1	0.91	0.34	52,52,52,52	0
86	OHX	1	4215	7/7	0.91	0.19	131,131,131,131	0
86	OHX	5	4213	7/7	0.91	0.16	149,149,149,149	0
85	MG	N6	201	1/1	0.91	0.16	48,48,48,48	0
85	MG	5	3880	1/1	0.91	0.15	39,39,39,39	0
85	MG	O3	201	1/1	0.91	0.09	40,40,40,40	0
85	MG	1	3681	1/1	0.91	0.09	42,42,42,42	0
86	OHX	C3	201	7/7	0.91	0.12	167,167,167,167	0
86	OHX	3	221	7/7	0.91	0.18	130,130,130,130	0
86	OHX	C5	201	7/7	0.91	0.11	176,176,176,176	0
86	OHX	3	224	7/7	0.91	0.15	145,145,145,145	0
85	MG	1	3682	1/1	0.91	0.11	62,62,62,62	0
85	MG	5	3679	1/1	0.91	0.08	40,40,40,40	0
85	MG	5	3500	1/1	0.91	0.23	36,36,36,36	0
85	MG	5	3501	1/1	0.91	0.21	40,40,40,40	0
85	MG	5	3502	1/1	0.91	0.22	29,29,29,29	0
85	MG	1	3613	1/1	0.91	0.26	45,45,45,45	0
85	MG	6	2041	1/1	0.91	0.26	69,69,69,69	0
86	OHX	1	4071	7/7	0.91	0.18	120,120,120,120	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	5	3516	1/1	0.91	0.20	40,40,40,40	0
86	OHX	6	2104	7/7	0.91	0.16	127,127,127,127	0
85	MG	5	3689	1/1	0.91	0.10	46,46,46,46	0
86	OHX	6	2116	7/7	0.91	0.14	144,144,144,144	0
86	OHX	6	2132	7/7	0.91	0.18	132,132,132,132	0
86	OHX	1	4075	7/7	0.91	0.14	137,137,137,137	0
85	MG	5	3692	1/1	0.91	0.29	43,43,43,43	0
86	OHX	6	2138	7/7	0.91	0.15	125,125,125,125	0
86	OHX	1	4081	7/7	0.91	0.13	140,140,140,140	0
85	MG	5	3517	1/1	0.91	0.23	32,32,32,32	0
86	OHX	1	4084	7/7	0.91	0.18	131,131,131,131	0
85	MG	7	206	1/1	0.91	0.12	48,48,48,48	0
85	MG	5	3521	1/1	0.91	0.36	34,34,34,34	0
86	OHX	1	4087	7/7	0.91	0.17	136,136,136,136	0
86	OHX	1	4091	7/7	0.91	0.16	132,132,132,132	0
85	MG	6	1969	1/1	0.91	0.11	64,64,64,64	0
86	OHX	1	4094	7/7	0.91	0.14	140,140,140,140	0
86	OHX	6	2163	7/7	0.91	0.15	139,139,139,139	0
85	MG	5	3534	1/1	0.91	0.28	35,35,35,35	0
85	MG	1	3616	1/1	0.91	0.11	46,46,46,46	0
85	MG	6	1906	1/1	0.91	0.24	48,48,48,48	0
85	MG	s1	301	1/1	0.91	0.19	75,75,75,75	0
86	OHX	l3	406	7/7	0.91	0.14	142,142,142,142	0
86	OHX	1	4104	7/7	0.91	0.16	128,128,128,128	0
85	MG	5	3713	1/1	0.91	0.10	43,43,43,43	0
85	MG	8	202	1/1	0.91	0.28	38,38,38,38	0
86	OHX	1	4109	7/7	0.91	0.17	130,130,130,130	0
86	OHX	1	4112	7/7	0.91	0.15	143,143,143,143	0
86	OHX	1	4113	7/7	0.91	0.16	128,128,128,128	0
86	OHX	6	2177	7/7	0.91	0.20	129,129,129,129	0
85	MG	8	203	1/1	0.91	0.21	52,52,52,52	0
85	MG	5	3714	1/1	0.91	0.11	72,72,72,72	0
85	MG	5	3542	1/1	0.91	0.26	30,30,30,30	0
85	MG	6	1973	1/1	0.91	0.22	59,59,59,59	0
85	MG	1	3855	1/1	0.91	0.23	53,53,53,53	0
88	3K5	1	4221	57/57	0.91	0.12	44,44,44,44	0
86	OHX	6	2183	7/7	0.91	0.16	142,142,142,142	0
85	MG	1	3471	1/1	0.92	0.19	49,49,49,49	0
85	MG	5	3824	1/1	0.92	0.12	47,47,47,47	0
85	MG	d3	201	1/1	0.92	0.26	55,55,55,55	0
85	MG	5	3510	1/1	0.92	0.31	41,41,41,41	0
86	OHX	6	2202	7/7	0.92	0.14	155,155,155,155	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	5	3661	1/1	0.92	0.21	59,59,59,59	0
85	MG	d3	202	1/1	0.92	0.24	55,55,55,55	0
85	MG	d6	102	1/1	0.92	0.16	60,60,60,60	0
85	MG	5	3518	1/1	0.92	0.18	40,40,40,40	0
85	MG	1	3831	1/1	0.92	0.12	59,59,59,59	0
86	OHX	1	4159	7/7	0.92	0.17	138,138,138,138	0
85	MG	5	3523	1/1	0.92	0.22	36,36,36,36	0
86	OHX	5	4014	7/7	0.92	0.14	149,149,149,149	0
86	OHX	5	4023	7/7	0.92	0.16	109,109,109,109	0
85	MG	5	3526	1/1	0.92	0.27	30,30,30,30	0
85	MG	5	3527	1/1	0.92	0.21	49,49,49,49	0
86	OHX	5	4095	7/7	0.92	0.14	138,138,138,138	0
86	OHX	2	2142	7/7	0.92	0.18	134,134,134,134	0
85	MG	2	1923	1/1	0.92	0.26	61,61,61,61	0
86	OHX	5	4098	7/7	0.92	0.13	149,149,149,149	0
86	OHX	5	4099	7/7	0.92	0.17	110,110,110,110	0
86	OHX	5	4100	7/7	0.92	0.11	162,162,162,162	0
86	OHX	5	4103	7/7	0.92	0.18	119,119,119,119	0
85	MG	4	223	1/1	0.92	0.24	64,64,64,64	0
85	MG	1	3768	1/1	0.92	0.18	52,52,52,52	0
85	MG	1	3606	1/1	0.92	0.09	59,59,59,59	0
86	OHX	5	4113	7/7	0.92	0.12	153,153,153,153	0
85	MG	1	3452	1/1	0.92	0.26	44,44,44,44	0
86	OHX	2	2149	7/7	0.92	0.12	176,176,176,176	0
85	MG	5	3685	1/1	0.92	0.10	35,35,35,35	0
86	OHX	2	2151	7/7	0.92	0.14	152,152,152,152	0
85	MG	6	1996	1/1	0.92	0.23	57,57,57,57	0
85	MG	5	3862	1/1	0.92	0.08	45,45,45,45	0
86	OHX	5	4130	7/7	0.92	0.17	127,127,127,127	0
85	MG	5	3865	1/1	0.92	0.27	58,58,58,58	0
85	MG	5	3866	1/1	0.92	0.07	49,49,49,49	0
85	MG	1	3842	1/1	0.92	0.20	46,46,46,46	0
85	MG	6	1936	1/1	0.92	0.30	46,46,46,46	0
85	MG	M0	301	1/1	0.92	0.20	44,44,44,44	0
85	MG	5	3871	1/1	0.92	0.22	39,39,39,39	0
85	MG	5	3549	1/1	0.92	0.28	54,54,54,54	0
85	MG	2	1972	1/1	0.92	0.18	72,72,72,72	0
85	MG	2	1966	1/1	0.92	0.37	61,61,61,61	0
85	MG	5	3697	1/1	0.92	0.21	69,69,69,69	0
86	OHX	5	4147	7/7	0.92	0.14	151,151,151,151	0
85	MG	1	3552	1/1	0.92	0.30	36,36,36,36	0
86	OHX	2	2166	7/7	0.92	0.11	168,168,168,168	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	5	3702	1/1	0.92	0.16	62,62,62,62	0
85	MG	1	3779	1/1	0.92	0.14	62,62,62,62	0
85	MG	5	3706	1/1	0.92	0.09	51,51,51,51	0
85	MG	1	3724	1/1	0.92	0.29	55,55,55,55	0
85	MG	1	3457	1/1	0.92	0.37	53,53,53,53	0
85	MG	1	3615	1/1	0.92	0.17	40,40,40,40	0
85	MG	5	3711	1/1	0.92	0.10	46,46,46,46	0
86	OHX	1	4197	7/7	0.92	0.13	151,151,151,151	0
85	MG	5	3434	1/1	0.92	0.24	53,53,53,53	0
86	OHX	5	4160	7/7	0.92	0.19	118,118,118,118	0
85	MG	1	3458	1/1	0.92	0.20	23,23,23,23	0
85	MG	5	3896	1/1	0.92	0.16	87,87,87,87	0
85	MG	2	1992	1/1	0.92	0.16	68,68,68,68	0
85	MG	7	202	1/1	0.92	0.20	31,31,31,31	0
86	OHX	5	4165	7/7	0.92	0.11	158,158,158,158	0
85	MG	7	203	1/1	0.92	0.28	58,58,58,58	0
85	MG	6	2012	1/1	0.92	0.17	68,68,68,68	0
85	MG	5	3719	1/1	0.92	0.09	48,48,48,48	0
86	OHX	5	4169	7/7	0.92	0.22	109,109,109,109	0
86	OHX	5	4171	7/7	0.92	0.16	129,129,129,129	0
85	MG	5	3443	1/1	0.92	0.23	33,33,33,33	0
85	MG	5	3446	1/1	0.92	0.16	35,35,35,35	0
86	OHX	1	3958	7/7	0.92	0.16	105,105,105,105	0
86	OHX	1	3990	7/7	0.92	0.17	109,109,109,109	0
86	OHX	1	4009	7/7	0.92	0.16	114,114,114,114	0
86	OHX	1	4022	7/7	0.92	0.11	159,159,159,159	0
85	MG	1	3731	1/1	0.92	0.32	41,41,41,41	0
85	MG	1	3557	1/1	0.92	0.33	39,39,39,39	0
86	OHX	1	4054	7/7	0.92	0.17	125,125,125,125	0
86	OHX	1	4062	7/7	0.92	0.17	122,122,122,122	0
86	OHX	1	4065	7/7	0.92	0.11	156,156,156,156	0
85	MG	5	3451	1/1	0.92	0.22	57,57,57,57	0
85	MG	5	3593	1/1	0.92	0.24	29,29,29,29	0
85	MG	1	3558	1/1	0.92	0.14	57,57,57,57	0
86	OHX	3	220	7/7	0.92	0.12	148,148,148,148	0
85	MG	1	3633	1/1	0.92	0.09	37,37,37,37	0
85	MG	1	3803	1/1	0.92	0.08	31,31,31,31	0
86	OHX	5	4191	7/7	0.92	0.12	176,176,176,176	0
86	OHX	3	223	7/7	0.92	0.18	136,136,136,136	0
85	MG	1	3532	1/1	0.92	0.22	29,29,29,29	0
85	MG	1	3806	1/1	0.92	0.16	55,55,55,55	0
86	OHX	1	4076	7/7	0.92	0.13	152,152,152,152	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	2	1927	1/1	0.92	0.27	57,57,57,57	0
86	OHX	1	4080	7/7	0.92	0.15	133,133,133,133	0
85	MG	5	3613	1/1	0.92	0.18	51,51,51,51	0
85	MG	5	3468	1/1	0.92	0.18	35,35,35,35	0
86	OHX	1	4083	7/7	0.92	0.16	128,128,128,128	0
85	MG	5	3757	1/1	0.92	0.21	62,62,62,62	0
85	MG	6	1903	1/1	0.92	0.30	47,47,47,47	0
86	OHX	O3	202	7/7	0.92	0.17	124,124,124,124	0
86	OHX	6	2099	7/7	0.92	0.14	162,162,162,162	0
85	MG	1	3415	1/1	0.92	0.22	38,38,38,38	0
85	MG	8	213	1/1	0.92	0.21	59,59,59,59	0
86	OHX	1	4088	7/7	0.92	0.12	145,145,145,145	0
86	OHX	6	2124	7/7	0.92	0.16	138,138,138,138	0
86	OHX	6	2127	7/7	0.92	0.15	129,129,129,129	0
86	OHX	6	2131	7/7	0.92	0.12	148,148,148,148	0
86	OHX	1	4089	7/7	0.92	0.14	136,136,136,136	0
85	MG	5	3622	1/1	0.92	0.31	52,52,52,52	0
86	OHX	6	2134	7/7	0.92	0.14	135,135,135,135	0
86	OHX	6	2135	7/7	0.92	0.12	149,149,149,149	0
85	MG	1	3572	1/1	0.92	0.29	46,46,46,46	0
85	MG	12	301	1/1	0.92	0.23	46,46,46,46	0
86	OHX	6	2141	7/7	0.92	0.12	164,164,164,164	0
85	MG	5	3475	1/1	0.92	0.18	51,51,51,51	0
85	MG	6	2033	1/1	0.92	0.24	60,60,60,60	0
85	MG	5	3769	1/1	0.92	0.13	44,44,44,44	0
85	MG	1	3407	1/1	0.92	0.23	44,44,44,44	0
85	MG	17	303	1/1	0.92	0.20	47,47,47,47	0
85	MG	5	3772	1/1	0.92	0.12	68,68,68,68	0
86	OHX	1	4106	7/7	0.92	0.14	141,141,141,141	0
85	MG	1	3578	1/1	0.92	0.18	32,32,32,32	0
85	MG	6	2036	1/1	0.92	0.22	68,68,68,68	0
86	OHX	6	2161	7/7	0.92	0.15	133,133,133,133	0
86	OHX	6	2162	7/7	0.92	0.17	125,125,125,125	0
85	MG	1	3539	1/1	0.92	0.22	41,41,41,41	0
85	MG	5	3634	1/1	0.92	0.11	47,47,47,47	0
85	MG	5	3635	1/1	0.92	0.12	40,40,40,40	0
86	OHX	6	2166	7/7	0.92	0.12	152,152,152,152	0
86	OHX	1	4115	7/7	0.92	0.19	119,119,119,119	0
85	MG	1	3644	1/1	0.92	0.07	49,49,49,49	0
85	MG	5	3786	1/1	0.92	0.07	31,31,31,31	0
86	OHX	6	2171	7/7	0.92	0.20	121,121,121,121	0
85	MG	1	3540	1/1	0.92	0.26	26,26,26,26	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	4	202	1/1	0.92	0.29	49,49,49,49	0
86	OHX	1	4121	7/7	0.92	0.14	143,143,143,143	0
85	MG	n8	204	1/1	0.92	0.12	42,42,42,42	0
85	MG	1	3541	1/1	0.92	0.26	60,60,60,60	0
85	MG	5	3793	1/1	0.92	0.13	46,46,46,46	0
85	MG	4	205	1/1	0.92	0.29	43,43,43,43	0
85	MG	4	207	1/1	0.92	0.28	34,34,34,34	0
86	OHX	2	2074	7/7	0.92	0.14	151,151,151,151	0
86	OHX	8	229	7/7	0.92	0.13	144,144,144,144	0
86	OHX	8	230	7/7	0.92	0.15	139,139,139,139	0
85	MG	5	3644	1/1	0.92	0.21	67,67,67,67	0
86	OHX	2	2087	7/7	0.92	0.14	140,140,140,140	0
85	MG	5	3646	1/1	0.92	0.26	50,50,50,50	0
85	MG	4	211	1/1	0.92	0.15	54,54,54,54	0
85	MG	1	3827	1/1	0.92	0.12	61,61,61,61	0
86	OHX	l5	303	7/7	0.92	0.12	146,146,146,146	0
86	OHX	l5	304	7/7	0.92	0.15	144,144,144,144	0
86	OHX	2	2102	7/7	0.92	0.14	146,146,146,146	0
86	OHX	2	2104	7/7	0.92	0.14	128,128,128,128	0
86	OHX	2	2105	7/7	0.92	0.14	143,143,143,143	0
85	MG	5	3806	1/1	0.92	0.11	45,45,45,45	0
85	MG	1	3544	1/1	0.92	0.27	38,38,38,38	0
85	MG	5	3808	1/1	0.92	0.10	51,51,51,51	0
85	MG	5	3809	1/1	0.92	0.17	41,41,41,41	0
85	MG	4	214	1/1	0.92	0.10	43,43,43,43	0
85	MG	1	3446	1/1	0.92	0.19	52,52,52,52	0
87	ZN	Q2	501	1/1	0.92	0.24	95,95,95,95	0
85	MG	5	3656	1/1	0.92	0.13	43,43,43,43	0
87	ZN	q2	501	1/1	0.92	0.20	93,93,93,93	0
86	OHX	2	2124	7/7	0.92	0.14	136,136,136,136	0
85	MG	6	1925	1/1	0.92	0.25	51,51,51,51	0
86	OHX	5	4072	7/7	0.93	0.14	131,131,131,131	0
86	OHX	5	4084	7/7	0.93	0.17	122,122,122,122	0
85	MG	1	3604	1/1	0.93	0.19	43,43,43,43	0
86	OHX	5	4088	7/7	0.93	0.15	121,121,121,121	0
86	OHX	5	4089	7/7	0.93	0.16	124,124,124,124	0
86	OHX	2	2165	7/7	0.93	0.11	171,171,171,171	0
86	OHX	5	4094	7/7	0.93	0.15	128,128,128,128	0
85	MG	1	3735	1/1	0.93	0.12	48,48,48,48	0
85	MG	1	3605	1/1	0.93	0.17	45,45,45,45	0
85	MG	5	3891	1/1	0.93	0.06	92,92,92,92	0
86	OHX	2	2169	7/7	0.93	0.13	145,145,145,145	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	1	3805	1/1	0.93	0.13	35,35,35,35	0
85	MG	5	3427	1/1	0.93	0.17	42,42,42,42	0
86	OHX	5	4101	7/7	0.93	0.13	134,134,134,134	0
86	OHX	5	4102	7/7	0.93	0.18	110,110,110,110	0
85	MG	1	3638	1/1	0.93	0.10	43,43,43,43	0
86	OHX	5	4104	7/7	0.93	0.15	123,123,123,123	0
85	MG	5	3717	1/1	0.93	0.19	52,52,52,52	0
85	MG	6	1995	1/1	0.93	0.09	57,57,57,57	0
85	MG	5	3569	1/1	0.93	0.22	28,28,28,28	0
86	OHX	5	4112	7/7	0.93	0.16	126,126,126,126	0
85	MG	1	3684	1/1	0.93	0.07	45,45,45,45	0
85	MG	1	3450	1/1	0.93	0.11	49,49,49,49	0
85	MG	1	3436	1/1	0.93	0.11	50,50,50,50	0
85	MG	5	3577	1/1	0.93	0.23	57,57,57,57	0
85	MG	1	3429	1/1	0.93	0.17	56,56,56,56	0
85	MG	5	3581	1/1	0.93	0.26	34,34,34,34	0
85	MG	4	208	1/1	0.93	0.28	38,38,38,38	0
86	OHX	5	4128	7/7	0.93	0.13	122,122,122,122	0
86	OHX	5	4129	7/7	0.93	0.10	184,184,184,184	0
85	MG	5	3584	1/1	0.93	0.25	45,45,45,45	0
86	OHX	5	4131	7/7	0.93	0.14	135,135,135,135	0
85	MG	4	209	1/1	0.93	0.23	31,31,31,31	0
85	MG	5	3445	1/1	0.93	0.20	30,30,30,30	0
86	OHX	1	3971	7/7	0.93	0.12	121,121,121,121	0
85	MG	5	3744	1/1	0.93	0.13	36,36,36,36	0
86	OHX	1	4002	7/7	0.93	0.14	167,167,167,167	0
85	MG	6	1930	1/1	0.93	0.39	61,61,61,61	0
86	OHX	1	4017	7/7	0.93	0.14	136,136,136,136	0
85	MG	1	3579	1/1	0.93	0.23	28,28,28,28	0
86	OHX	1	4026	7/7	0.93	0.14	128,128,128,128	0
85	MG	5	3748	1/1	0.93	0.11	66,66,66,66	0
86	OHX	5	4146	7/7	0.93	0.15	132,132,132,132	0
86	OHX	1	4037	7/7	0.93	0.14	133,133,133,133	0
85	MG	1	3611	1/1	0.93	0.10	44,44,44,44	0
86	OHX	1	4042	7/7	0.93	0.13	149,149,149,149	0
85	MG	5	3752	1/1	0.93	0.16	44,44,44,44	0
85	MG	1	3820	1/1	0.93	0.10	43,43,43,43	0
85	MG	5	3603	1/1	0.93	0.11	62,62,62,62	0
85	MG	5	3605	1/1	0.93	0.08	42,42,42,42	0
85	MG	5	3759	1/1	0.93	0.18	62,62,62,62	0
85	MG	5	3760	1/1	0.93	0.14	61,61,61,61	0
85	MG	1	3748	1/1	0.93	0.09	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	1	3749	1/1	0.93	0.14	60,60,60,60	0
85	MG	4	216	1/1	0.93	0.12	48,48,48,48	0
85	MG	6	1937	1/1	0.93	0.15	43,43,43,43	0
85	MG	1	3752	1/1	0.93	0.13	61,61,61,61	0
85	MG	1	3580	1/1	0.93	0.34	38,38,38,38	0
86	OHX	1	4078	7/7	0.93	0.12	129,129,129,129	0
86	OHX	1	4079	7/7	0.93	0.15	127,127,127,127	0
85	MG	14	401	1/1	0.93	0.06	43,43,43,43	0
85	MG	1	3648	1/1	0.93	0.15	43,43,43,43	0
85	MG	1	3756	1/1	0.93	0.20	47,47,47,47	0
85	MG	5	3773	1/1	0.93	0.17	70,70,70,70	0
85	MG	6	1943	1/1	0.93	0.34	64,64,64,64	0
85	MG	5	3472	1/1	0.93	0.20	57,57,57,57	0
86	OHX	M0	304	7/7	0.93	0.14	120,120,120,120	0
85	MG	2	2011	1/1	0.93	0.18	58,58,58,58	0
85	MG	1	3520	1/1	0.93	0.16	42,42,42,42	0
85	MG	5	3629	1/1	0.93	0.21	57,57,57,57	0
85	MG	1	3651	1/1	0.93	0.08	51,51,51,51	0
86	OHX	5	4177	7/7	0.93	0.11	159,159,159,159	0
86	OHX	6	2094	7/7	0.93	0.14	133,133,133,133	0
85	MG	1	3535	1/1	0.93	0.28	32,32,32,32	0
85	MG	5	3478	1/1	0.93	0.17	33,33,33,33	0
85	MG	n3	201	1/1	0.93	0.25	27,27,27,27	0
85	MG	5	3479	1/1	0.93	0.29	68,68,68,68	0
86	OHX	6	2118	7/7	0.93	0.13	144,144,144,144	0
85	MG	n8	202	1/1	0.93	0.08	32,32,32,32	0
85	MG	1	3835	1/1	0.93	0.14	30,30,30,30	0
86	OHX	1	4101	7/7	0.93	0.11	151,151,151,151	0
85	MG	1	3711	1/1	0.93	0.05	61,61,61,61	0
85	MG	5	3482	1/1	0.93	0.20	39,39,39,39	0
85	MG	1	3839	1/1	0.93	0.20	25,25,25,25	0
85	MG	1	3506	1/1	0.93	0.17	42,42,42,42	0
85	MG	1	3841	1/1	0.93	0.23	33,33,33,33	0
86	OHX	1	4108	7/7	0.93	0.14	129,129,129,129	0
86	OHX	2	2067	7/7	0.93	0.12	159,159,159,159	0
86	OHX	5	4196	7/7	0.93	0.15	125,125,125,125	0
86	OHX	5	4198	7/7	0.93	0.16	128,128,128,128	0
86	OHX	6	2143	7/7	0.93	0.12	148,148,148,148	0
85	MG	M3	203	1/1	0.93	0.10	35,35,35,35	0
86	OHX	6	2147	7/7	0.93	0.17	117,117,117,117	0
85	MG	6	1958	1/1	0.93	0.32	53,53,53,53	0
85	MG	1	3770	1/1	0.93	0.12	45,45,45,45	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	6	2151	7/7	0.93	0.18	121,121,121,121	0
85	MG	1	3714	1/1	0.93	0.10	45,45,45,45	0
85	MG	1	3715	1/1	0.93	0.16	61,61,61,61	0
85	MG	1	3621	1/1	0.93	0.24	70,70,70,70	0
86	OHX	2	2100	7/7	0.93	0.12	150,150,150,150	0
86	OHX	6	2157	7/7	0.93	0.17	152,152,152,152	0
85	MG	6	2042	1/1	0.93	0.23	53,53,53,53	0
86	OHX	2	2103	7/7	0.93	0.09	202,202,202,202	0
85	MG	5	3811	1/1	0.93	0.11	42,42,42,42	0
85	MG	1	3622	1/1	0.93	0.09	37,37,37,37	0
85	MG	5	3654	1/1	0.93	0.28	72,72,72,72	0
85	MG	1	3777	1/1	0.93	0.11	49,49,49,49	0
86	OHX	2	2113	7/7	0.93	0.11	167,167,167,167	0
85	MG	1	3624	1/1	0.93	0.33	58,58,58,58	0
86	OHX	1	4129	7/7	0.93	0.11	152,152,152,152	0
86	OHX	6	2168	7/7	0.93	0.18	128,128,128,128	0
86	OHX	2	2117	7/7	0.93	0.10	166,166,166,166	0
85	MG	5	3823	1/1	0.93	0.21	65,65,65,65	0
85	MG	5	3503	1/1	0.93	0.17	53,53,53,53	0
85	MG	1	3625	1/1	0.93	0.14	47,47,47,47	0
85	MG	5	3505	1/1	0.93	0.30	34,34,34,34	0
85	MG	1	3665	1/1	0.93	0.08	50,50,50,50	0
85	MG	5	3835	1/1	0.93	0.07	48,48,48,48	0
85	MG	5	3837	1/1	0.93	0.16	37,37,37,37	0
85	MG	5	3840	1/1	0.93	0.16	78,78,78,78	0
86	OHX	2	2130	7/7	0.93	0.16	140,140,140,140	0
85	MG	5	3507	1/1	0.93	0.20	36,36,36,36	0
85	MG	5	3843	1/1	0.93	0.20	50,50,50,50	0
85	MG	s8	301	1/1	0.93	0.14	53,53,53,53	0
85	MG	5	3845	1/1	0.93	0.14	42,42,42,42	0
85	MG	5	3514	1/1	0.93	0.29	57,57,57,57	0
85	MG	N8	203	1/1	0.93	0.28	43,43,43,43	0
85	MG	1	3563	1/1	0.93	0.23	27,27,27,27	0
86	OHX	5	4239	7/7	0.93	0.19	110,110,110,110	0
85	MG	5	3671	1/1	0.93	0.11	31,31,31,31	0
85	MG	1	3783	1/1	0.93	0.11	57,57,57,57	0
85	MG	5	3519	1/1	0.93	0.25	27,27,27,27	0
85	MG	1	3785	1/1	0.93	0.17	63,63,63,63	0
85	MG	1	3627	1/1	0.93	0.16	41,41,41,41	0
85	MG	1	3631	1/1	0.93	0.12	40,40,40,40	0
85	MG	1	3464	1/1	0.93	0.21	28,28,28,28	0
85	MG	1	3870	1/1	0.93	0.32	57,57,57,57	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	1	4157	7/7	0.93	0.14	140,140,140,140	0
86	OHX	7	224	7/7	0.93	0.13	140,140,140,140	0
85	MG	5	3530	1/1	0.93	0.17	32,32,32,32	0
86	OHX	7	226	7/7	0.93	0.17	119,119,119,119	0
85	MG	S4	301	1/1	0.93	0.18	71,71,71,71	0
86	OHX	1	4160	7/7	0.93	0.17	138,138,138,138	0
85	MG	1	3792	1/1	0.93	0.06	49,49,49,49	0
85	MG	5	3404	1/1	0.93	0.08	48,48,48,48	0
85	MG	1	3598	1/1	0.93	0.26	43,43,43,43	0
85	MG	5	3540	1/1	0.93	0.33	46,46,46,46	0
85	MG	6	1984	1/1	0.93	0.07	49,49,49,49	0
85	MG	1	3797	1/1	0.93	0.14	73,73,73,73	0
85	MG	5	3872	1/1	0.93	0.37	51,51,51,51	0
85	MG	3	206	1/1	0.93	0.30	34,34,34,34	0
86	OHX	s4	302	7/7	0.93	0.13	155,155,155,155	0
85	MG	5	3698	1/1	0.93	0.13	46,46,46,46	0
85	MG	5	3414	1/1	0.93	0.25	31,31,31,31	0
86	OHX	c5	201	7/7	0.93	0.10	172,172,172,172	0
85	MG	3	207	1/1	0.93	0.33	64,64,64,64	0
85	MG	5	3550	1/1	0.93	0.36	43,43,43,43	0
86	OHX	5	4009	7/7	0.93	0.14	116,116,116,116	0
85	MG	6	1988	1/1	0.93	0.30	73,73,73,73	0
86	OHX	5	4022	7/7	0.93	0.17	112,112,112,112	0
85	MG	1	3600	1/1	0.93	0.23	41,41,41,41	0
86	OHX	5	4033	7/7	0.93	0.11	125,125,125,125	0
86	OHX	5	4038	7/7	0.93	0.13	133,133,133,133	0
86	OHX	5	4057	7/7	0.93	0.16	120,120,120,120	0
86	OHX	5	4059	7/7	0.93	0.12	130,130,130,130	0
86	OHX	5	4064	7/7	0.93	0.15	116,116,116,116	0
86	OHX	5	4067	7/7	0.93	0.15	134,134,134,134	0
85	MG	M7	204	1/1	0.94	0.07	38,38,38,38	0
86	OHX	sR	401	7/7	0.94	0.10	167,167,167,167	0
86	OHX	5	3997	7/7	0.94	0.13	114,114,114,114	0
86	OHX	2	2122	7/7	0.94	0.12	148,148,148,148	0
85	MG	5	3670	1/1	0.94	0.15	32,32,32,32	0
85	MG	1	3692	1/1	0.94	0.26	51,51,51,51	0
85	MG	5	3829	1/1	0.94	0.30	54,54,54,54	0
86	OHX	5	4025	7/7	0.94	0.15	101,101,101,101	0
86	OHX	5	4032	7/7	0.94	0.10	136,136,136,136	0
85	MG	5	3830	1/1	0.94	0.16	76,76,76,76	0
86	OHX	5	4037	7/7	0.94	0.12	159,159,159,159	0
85	MG	1	3425	1/1	0.94	0.16	50,50,50,50	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	5	4039	7/7	0.94	0.14	123,123,123,123	0
85	MG	5	3674	1/1	0.94	0.19	54,54,54,54	0
86	OHX	5	4058	7/7	0.94	0.13	128,128,128,128	0
86	OHX	1	4156	7/7	0.94	0.14	118,118,118,118	0
86	OHX	5	4061	7/7	0.94	0.14	122,122,122,122	0
85	MG	5	3436	1/1	0.94	0.18	32,32,32,32	0
86	OHX	5	4066	7/7	0.94	0.14	140,140,140,140	0
85	MG	6	2015	1/1	0.94	0.11	71,71,71,71	0
86	OHX	5	4068	7/7	0.94	0.14	122,122,122,122	0
86	OHX	5	4069	7/7	0.94	0.13	121,121,121,121	0
86	OHX	5	4071	7/7	0.94	0.17	121,121,121,121	0
85	MG	5	3548	1/1	0.94	0.33	50,50,50,50	0
86	OHX	5	4082	7/7	0.94	0.14	132,132,132,132	0
85	MG	1	3695	1/1	0.94	0.27	47,47,47,47	0
85	MG	6	1952	1/1	0.94	0.42	60,60,60,60	0
85	MG	5	3683	1/1	0.94	0.08	42,42,42,42	0
85	MG	6	2020	1/1	0.94	0.29	89,89,89,89	0
85	MG	N3	201	1/1	0.94	0.23	37,37,37,37	0
85	MG	1	3867	1/1	0.94	0.08	56,56,56,56	0
85	MG	5	3554	1/1	0.94	0.33	43,43,43,43	0
85	MG	1	3802	1/1	0.94	0.38	46,46,46,46	0
85	MG	5	3850	1/1	0.94	0.14	52,52,52,52	0
85	MG	1	3697	1/1	0.94	0.18	55,55,55,55	0
85	MG	5	3690	1/1	0.94	0.09	43,43,43,43	0
85	MG	5	3452	1/1	0.94	0.18	32,32,32,32	0
85	MG	5	3559	1/1	0.94	0.16	52,52,52,52	0
85	MG	1	3477	1/1	0.94	0.20	36,36,36,36	0
85	MG	1	3505	1/1	0.94	0.26	43,43,43,43	0
85	MG	5	3459	1/1	0.94	0.17	72,72,72,72	0
85	MG	5	3699	1/1	0.94	0.21	56,56,56,56	0
85	MG	2	1946	1/1	0.94	0.12	65,65,65,65	0
85	MG	5	3864	1/1	0.94	0.23	45,45,45,45	0
86	OHX	5	4109	7/7	0.94	0.15	128,128,128,128	0
86	OHX	5	4110	7/7	0.94	0.19	111,111,111,111	0
86	OHX	5	4111	7/7	0.94	0.16	119,119,119,119	0
85	MG	5	3464	1/1	0.94	0.25	54,54,54,54	0
85	MG	5	3703	1/1	0.94	0.09	40,40,40,40	0
85	MG	5	3572	1/1	0.94	0.29	30,30,30,30	0
85	MG	5	3705	1/1	0.94	0.30	67,67,67,67	0
85	MG	1	3581	1/1	0.94	0.31	43,43,43,43	0
86	OHX	5	4118	7/7	0.94	0.11	151,151,151,151	0
86	OHX	5	4119	7/7	0.94	0.13	144,144,144,144	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	5	4120	7/7	0.94	0.12	138,138,138,138	0
85	MG	6	2032	1/1	0.94	0.07	51,51,51,51	0
86	OHX	5	4124	7/7	0.94	0.14	134,134,134,134	0
85	MG	1	3808	1/1	0.94	0.15	89,89,89,89	0
85	MG	5	3709	1/1	0.94	0.22	48,48,48,48	0
85	MG	5	3578	1/1	0.94	0.21	30,30,30,30	0
85	MG	6	1901	1/1	0.94	0.27	44,44,44,44	0
85	MG	5	3712	1/1	0.94	0.18	89,89,89,89	0
85	MG	5	3881	1/1	0.94	0.25	49,49,49,49	0
85	MG	1	3704	1/1	0.94	0.11	47,47,47,47	0
85	MG	5	3883	1/1	0.94	0.19	36,36,36,36	0
85	MG	1	3706	1/1	0.94	0.25	53,53,53,53	0
85	MG	5	3715	1/1	0.94	0.17	47,47,47,47	0
85	MG	1	3617	1/1	0.94	0.08	67,67,67,67	0
86	OHX	1	4199	7/7	0.94	0.12	176,176,176,176	0
85	MG	5	3587	1/1	0.94	0.32	28,28,28,28	0
85	MG	1	3467	1/1	0.94	0.18	62,62,62,62	0
85	MG	5	3890	1/1	0.94	0.08	29,29,29,29	0
85	MG	5	3590	1/1	0.94	0.29	31,31,31,31	0
85	MG	5	3474	1/1	0.94	0.24	52,52,52,52	0
85	MG	5	3721	1/1	0.94	0.13	59,59,59,59	0
85	MG	5	3723	1/1	0.94	0.12	38,38,38,38	0
85	MG	1	3583	1/1	0.94	0.22	36,36,36,36	0
85	MG	5	3594	1/1	0.94	0.21	32,32,32,32	0
85	MG	5	3726	1/1	0.94	0.14	43,43,43,43	0
85	MG	1	3762	1/1	0.94	0.08	36,36,36,36	0
85	MG	5	3732	1/1	0.94	0.09	45,45,45,45	0
85	MG	1	3509	1/1	0.94	0.22	23,23,23,23	0
86	OHX	1	3934	7/7	0.94	0.16	107,107,107,107	0
85	MG	5	3598	1/1	0.94	0.34	35,35,35,35	0
86	OHX	1	3968	7/7	0.94	0.15	104,104,104,104	0
85	MG	5	3599	1/1	0.94	0.39	30,30,30,30	0
86	OHX	1	3988	7/7	0.94	0.13	112,112,112,112	0
85	MG	5	3737	1/1	0.94	0.16	51,51,51,51	0
85	MG	1	3433	1/1	0.94	0.21	44,44,44,44	0
86	OHX	1	4006	7/7	0.94	0.14	122,122,122,122	0
85	MG	6	1913	1/1	0.94	0.24	37,37,37,37	0
86	OHX	1	4010	7/7	0.94	0.13	128,128,128,128	0
86	OHX	1	4014	7/7	0.94	0.14	129,129,129,129	0
86	OHX	1	4015	7/7	0.94	0.12	131,131,131,131	0
85	MG	7	211	1/1	0.94	0.17	37,37,37,37	0
86	OHX	4	233	7/7	0.94	0.15	119,119,119,119	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	4	234	7/7	0.94	0.11	147,147,147,147	0
86	OHX	1	4021	7/7	0.94	0.13	119,119,119,119	0
85	MG	7	212	1/1	0.94	0.24	66,66,66,66	0
86	OHX	1	4024	7/7	0.94	0.12	141,141,141,141	0
85	MG	5	3740	1/1	0.94	0.20	43,43,43,43	0
86	OHX	1	4031	7/7	0.94	0.11	134,134,134,134	0
85	MG	1	3587	1/1	0.94	0.34	38,38,38,38	0
85	MG	5	3606	1/1	0.94	0.10	31,31,31,31	0
85	MG	1	3554	1/1	0.94	0.32	53,53,53,53	0
85	MG	1	3769	1/1	0.94	0.16	49,49,49,49	0
85	MG	1	3661	1/1	0.94	0.17	48,48,48,48	0
86	OHX	1	4056	7/7	0.94	0.12	141,141,141,141	0
86	OHX	6	2076	7/7	0.94	0.15	117,117,117,117	0
86	OHX	1	4061	7/7	0.94	0.16	105,105,105,105	0
85	MG	6	1980	1/1	0.94	0.09	52,52,52,52	0
86	OHX	6	2102	7/7	0.94	0.14	123,123,123,123	0
86	OHX	5	4187	7/7	0.94	0.12	144,144,144,144	0
86	OHX	1	4064	7/7	0.94	0.13	129,129,129,129	0
85	MG	c1	201	1/1	0.94	0.26	49,49,49,49	0
86	OHX	6	2107	7/7	0.94	0.14	119,119,119,119	0
86	OHX	6	2110	7/7	0.94	0.13	134,134,134,134	0
86	OHX	6	2111	7/7	0.94	0.12	129,129,129,129	0
85	MG	5	3753	1/1	0.94	0.27	52,52,52,52	0
85	MG	1	3591	1/1	0.94	0.24	33,33,33,33	0
86	OHX	6	2120	7/7	0.94	0.14	119,119,119,119	0
86	OHX	5	4197	7/7	0.94	0.13	129,129,129,129	0
86	OHX	6	2121	7/7	0.94	0.11	146,146,146,146	0
86	OHX	6	2123	7/7	0.94	0.14	111,111,111,111	0
85	MG	1	3629	1/1	0.94	0.12	47,47,47,47	0
85	MG	1	3417	1/1	0.94	0.16	36,36,36,36	0
85	MG	1	3515	1/1	0.94	0.21	41,41,41,41	0
86	OHX	1	4073	7/7	0.94	0.15	131,131,131,131	0
85	MG	5	3494	1/1	0.94	0.18	45,45,45,45	0
85	MG	2	1906	1/1	0.94	0.26	58,58,58,58	0
85	MG	5	3627	1/1	0.94	0.19	33,33,33,33	0
85	MG	1	3597	1/1	0.94	0.30	27,27,27,27	0
85	MG	5	3765	1/1	0.94	0.16	40,40,40,40	0
86	OHX	6	2140	7/7	0.94	0.13	138,138,138,138	0
85	MG	l3	401	1/1	0.94	0.33	26,26,26,26	0
85	MG	5	3766	1/1	0.94	0.18	44,44,44,44	0
85	MG	5	3497	1/1	0.94	0.19	39,39,39,39	0
86	OHX	6	2146	7/7	0.94	0.12	136,136,136,136	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	1	3837	1/1	0.94	0.07	38,38,38,38	0
85	MG	1	3462	1/1	0.94	0.22	24,24,24,24	0
85	MG	17	301	1/1	0.94	0.08	39,39,39,39	0
85	MG	17	302	1/1	0.94	0.12	43,43,43,43	0
85	MG	sM	302	1/1	0.94	0.10	50,50,50,50	0
85	MG	1	3559	1/1	0.94	0.23	52,52,52,52	0
85	MG	1	3729	1/1	0.94	0.13	47,47,47,47	0
85	MG	m5	301	1/1	0.94	0.08	39,39,39,39	0
85	MG	m5	302	1/1	0.94	0.14	56,56,56,56	0
85	MG	1	3680	1/1	0.94	0.20	51,51,51,51	0
85	MG	5	3777	1/1	0.94	0.17	32,32,32,32	0
86	OHX	1	4096	7/7	0.94	0.15	125,125,125,125	0
85	MG	L3	401	1/1	0.94	0.10	75,75,75,75	0
85	MG	1	3518	1/1	0.94	0.26	35,35,35,35	0
85	MG	m7	203	1/1	0.94	0.10	57,57,57,57	0
85	MG	5	3407	1/1	0.94	0.16	41,41,41,41	0
86	OHX	5	4231	7/7	0.94	0.15	114,114,114,114	0
85	MG	5	3409	1/1	0.94	0.26	44,44,44,44	0
86	OHX	1	4103	7/7	0.94	0.16	119,119,119,119	0
85	MG	5	3511	1/1	0.94	0.27	32,32,32,32	0
85	MG	n6	201	1/1	0.94	0.26	53,53,53,53	0
85	MG	5	3513	1/1	0.94	0.12	34,34,34,34	0
85	MG	5	3645	1/1	0.94	0.07	43,43,43,43	0
85	MG	1	3786	1/1	0.94	0.12	48,48,48,48	0
85	MG	5	3791	1/1	0.94	0.09	56,56,56,56	0
86	OHX	1	4111	7/7	0.94	0.15	123,123,123,123	0
85	MG	1	3488	1/1	0.94	0.23	44,44,44,44	0
85	MG	1	3542	1/1	0.94	0.19	31,31,31,31	0
85	MG	1	3501	1/1	0.94	0.21	72,72,72,72	0
85	MG	o7	502	1/1	0.94	0.22	38,38,38,38	0
85	MG	M0	302	1/1	0.94	0.13	55,55,55,55	0
85	MG	q3	502	1/1	0.94	0.17	62,62,62,62	0
86	OHX	2	2039	7/7	0.94	0.16	106,106,106,106	0
86	OHX	1	4119	7/7	0.94	0.15	127,127,127,127	0
86	OHX	2	2062	7/7	0.94	0.11	136,136,136,136	0
85	MG	5	3652	1/1	0.94	0.15	43,43,43,43	0
86	OHX	1	4122	7/7	0.94	0.14	136,136,136,136	0
85	MG	5	3417	1/1	0.94	0.23	30,30,30,30	0
86	OHX	2	2075	7/7	0.94	0.15	135,135,135,135	0
86	OHX	2	2079	7/7	0.94	0.10	167,167,167,167	0
85	MG	5	3800	1/1	0.94	0.09	41,41,41,41	0
85	MG	5	3802	1/1	0.94	0.13	46,46,46,46	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	5	3522	1/1	0.94	0.15	36,36,36,36	0
85	MG	5	3804	1/1	0.94	0.08	52,52,52,52	0
86	OHX	2	2097	7/7	0.94	0.10	161,161,161,161	0
85	MG	5	3419	1/1	0.94	0.11	38,38,38,38	0
85	MG	5	3525	1/1	0.94	0.25	37,37,37,37	0
85	MG	1	3569	1/1	0.94	0.23	33,33,33,33	0
85	MG	1	3791	1/1	0.94	0.37	42,42,42,42	0
85	MG	1	3643	1/1	0.94	0.14	43,43,43,43	0
85	MG	1	3502	1/1	0.94	0.26	22,22,22,22	0
85	MG	5	3662	1/1	0.94	0.18	33,33,33,33	0
86	OHX	2	2108	7/7	0.94	0.10	163,163,163,163	0
85	MG	6	1942	1/1	0.94	0.27	39,39,39,39	0
86	OHX	6	2203	7/7	0.94	0.09	197,197,197,197	0
85	MG	5	3814	1/1	0.94	0.17	62,62,62,62	0
86	OHX	1	4141	7/7	0.94	0.13	114,114,114,114	0
85	MG	1	3856	1/1	0.94	0.18	50,50,50,50	0
86	OHX	2	2116	7/7	0.94	0.13	150,150,150,150	0
85	MG	5	3816	1/1	0.94	0.14	47,47,47,47	0
85	MG	1	3857	1/1	0.94	0.28	64,64,64,64	0
85	MG	1	3796	1/1	0.94	0.26	29,29,29,29	0
86	OHX	2	2120	7/7	0.94	0.10	155,155,155,155	0
85	MG	1	3821	1/1	0.95	0.13	52,52,52,52	0
85	MG	5	3562	1/1	0.95	0.25	30,30,30,30	0
85	MG	1	3672	1/1	0.95	0.09	54,54,54,54	0
86	OHX	1	4012	7/7	0.95	0.16	116,116,116,116	0
85	MG	5	3568	1/1	0.95	0.25	31,31,31,31	0
86	OHX	5	4073	7/7	0.95	0.11	163,163,163,163	0
86	OHX	5	4075	7/7	0.95	0.13	135,135,135,135	0
86	OHX	5	4078	7/7	0.95	0.15	108,108,108,108	0
86	OHX	5	4079	7/7	0.95	0.12	113,113,113,113	0
86	OHX	5	4080	7/7	0.95	0.10	143,143,143,143	0
86	OHX	5	4081	7/7	0.95	0.13	120,120,120,120	0
85	MG	3	210	1/1	0.95	0.29	64,64,64,64	0
86	OHX	5	4083	7/7	0.95	0.11	134,134,134,134	0
85	MG	1	3484	1/1	0.95	0.17	55,55,55,55	0
86	OHX	1	4018	7/7	0.95	0.12	145,145,145,145	0
86	OHX	5	4086	7/7	0.95	0.17	109,109,109,109	0
86	OHX	5	4087	7/7	0.95	0.14	116,116,116,116	0
86	OHX	1	4020	7/7	0.95	0.12	127,127,127,127	0
85	MG	5	3812	1/1	0.95	0.07	45,45,45,45	0
86	OHX	5	4091	7/7	0.95	0.11	135,135,135,135	0
86	OHX	5	4092	7/7	0.95	0.12	134,134,134,134	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	5	3471	1/1	0.95	0.22	39,39,39,39	0
86	OHX	1	4023	7/7	0.95	0.12	125,125,125,125	0
85	MG	1	3485	1/1	0.95	0.09	46,46,46,46	0
85	MG	5	3574	1/1	0.95	0.23	33,33,33,33	0
86	OHX	1	4028	7/7	0.95	0.10	147,147,147,147	0
86	OHX	1	4029	7/7	0.95	0.11	144,144,144,144	0
85	MG	1	3412	1/1	0.95	0.10	44,44,44,44	0
85	MG	1	3454	1/1	0.95	0.18	52,52,52,52	0
86	OHX	1	4034	7/7	0.95	0.12	134,134,134,134	0
85	MG	6	1975	1/1	0.95	0.12	48,48,48,48	0
86	OHX	1	4039	7/7	0.95	0.13	117,117,117,117	0
85	MG	5	3694	1/1	0.95	0.14	47,47,47,47	0
85	MG	1	3489	1/1	0.95	0.14	39,39,39,39	0
86	OHX	1	4044	7/7	0.95	0.13	129,129,129,129	0
86	OHX	1	4049	7/7	0.95	0.14	114,114,114,114	0
86	OHX	1	4050	7/7	0.95	0.14	116,116,116,116	0
85	MG	5	3580	1/1	0.95	0.21	35,35,35,35	0
86	OHX	1	4055	7/7	0.95	0.14	121,121,121,121	0
85	MG	n8	205	1/1	0.95	0.11	40,40,40,40	0
86	OHX	3	219	7/7	0.95	0.12	135,135,135,135	0
86	OHX	5	4114	7/7	0.95	0.16	112,112,112,112	0
86	OHX	1	4058	7/7	0.95	0.11	142,142,142,142	0
86	OHX	1	4059	7/7	0.95	0.17	115,115,115,115	0
85	MG	5	3828	1/1	0.95	0.20	29,29,29,29	0
85	MG	s4	301	1/1	0.95	0.23	55,55,55,55	0
85	MG	6	1910	1/1	0.95	0.25	51,51,51,51	0
86	OHX	4	232	7/7	0.95	0.11	122,122,122,122	0
85	MG	5	3831	1/1	0.95	0.09	56,56,56,56	0
86	OHX	5	4123	7/7	0.95	0.10	144,144,144,144	0
85	MG	1	3775	1/1	0.95	0.09	41,41,41,41	0
85	MG	5	3586	1/1	0.95	0.30	28,28,28,28	0
85	MG	q3	503	1/1	0.95	0.11	74,74,74,74	0
86	OHX	5	4127	7/7	0.95	0.10	142,142,142,142	0
86	OHX	1	4070	7/7	0.95	0.09	156,156,156,156	0
85	MG	1	3474	1/1	0.95	0.24	31,31,31,31	0
86	OHX	L3	404	7/7	0.95	0.13	121,121,121,121	0
86	OHX	2	2052	7/7	0.95	0.11	135,135,135,135	0
86	OHX	5	4132	7/7	0.95	0.14	127,127,127,127	0
86	OHX	5	4133	7/7	0.95	0.12	140,140,140,140	0
86	OHX	2	2053	7/7	0.95	0.11	136,136,136,136	0
85	MG	5	3588	1/1	0.95	0.23	29,29,29,29	0
85	MG	5	3839	1/1	0.95	0.10	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	4	206	1/1	0.95	0.21	32,32,32,32	0
85	MG	1	3475	1/1	0.95	0.20	27,27,27,27	0
86	OHX	2	2076	7/7	0.95	0.12	127,127,127,127	0
86	OHX	5	4140	7/7	0.95	0.13	138,138,138,138	0
86	OHX	6	2063	7/7	0.95	0.13	100,100,100,100	0
86	OHX	2	2077	7/7	0.95	0.10	139,139,139,139	0
86	OHX	5	4143	7/7	0.95	0.15	125,125,125,125	0
86	OHX	2	2078	7/7	0.95	0.14	131,131,131,131	0
85	MG	1	3463	1/1	0.95	0.19	28,28,28,28	0
85	MG	5	3592	1/1	0.95	0.23	44,44,44,44	0
86	OHX	2	2084	7/7	0.95	0.10	145,145,145,145	0
86	OHX	2	2085	7/7	0.95	0.13	121,121,121,121	0
85	MG	1	3686	1/1	0.95	0.12	49,49,49,49	0
85	MG	4	210	1/1	0.95	0.12	44,44,44,44	0
85	MG	1	3688	1/1	0.95	0.09	41,41,41,41	0
86	OHX	6	2114	7/7	0.95	0.11	132,132,132,132	0
86	OHX	6	2115	7/7	0.95	0.11	133,133,133,133	0
86	OHX	2	2096	7/7	0.95	0.09	175,175,175,175	0
85	MG	1	3455	1/1	0.95	0.28	39,39,39,39	0
85	MG	1	3838	1/1	0.95	0.22	37,37,37,37	0
85	MG	5	3491	1/1	0.95	0.29	49,49,49,49	0
85	MG	5	3852	1/1	0.95	0.10	59,59,59,59	0
86	OHX	1	4095	7/7	0.95	0.09	159,159,159,159	0
86	OHX	6	2125	7/7	0.95	0.13	138,138,138,138	0
86	OHX	6	2126	7/7	0.95	0.19	109,109,109,109	0
85	MG	5	3600	1/1	0.95	0.14	43,43,43,43	0
85	MG	1	3440	1/1	0.95	0.24	38,38,38,38	0
86	OHX	1	4098	7/7	0.95	0.10	145,145,145,145	0
85	MG	5	3602	1/1	0.95	0.07	47,47,47,47	0
85	MG	1	3589	1/1	0.95	0.22	34,34,34,34	0
85	MG	5	3604	1/1	0.95	0.18	37,37,37,37	0
85	MG	1	3538	1/1	0.95	0.38	35,35,35,35	0
86	OHX	2	2112	7/7	0.95	0.12	140,140,140,140	0
86	OHX	6	2139	7/7	0.95	0.13	143,143,143,143	0
85	MG	6	1926	1/1	0.95	0.24	46,46,46,46	0
85	MG	5	3722	1/1	0.95	0.13	48,48,48,48	0
85	MG	1	3423	1/1	0.95	0.07	36,36,36,36	0
85	MG	5	3498	1/1	0.95	0.25	34,34,34,34	0
85	MG	5	3610	1/1	0.95	0.24	33,33,33,33	0
85	MG	1	3623	1/1	0.95	0.20	45,45,45,45	0
86	OHX	1	4110	7/7	0.95	0.09	143,143,143,143	0
85	MG	5	3727	1/1	0.95	0.10	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	1	3696	1/1	0.95	0.13	50,50,50,50	0
85	MG	5	3730	1/1	0.95	0.23	33,33,33,33	0
85	MG	5	3614	1/1	0.95	0.06	35,35,35,35	0
85	MG	1	3846	1/1	0.95	0.25	42,42,42,42	0
86	OHX	6	2155	7/7	0.95	0.14	117,117,117,117	0
85	MG	5	3734	1/1	0.95	0.06	45,45,45,45	0
85	MG	1	3847	1/1	0.95	0.23	30,30,30,30	0
86	OHX	2	2127	7/7	0.95	0.11	154,154,154,154	0
86	OHX	6	2159	7/7	0.95	0.14	123,123,123,123	0
85	MG	5	3620	1/1	0.95	0.10	47,47,47,47	0
86	OHX	2	2129	7/7	0.95	0.13	127,127,127,127	0
85	MG	5	3878	1/1	0.95	0.20	24,24,24,24	0
85	MG	1	3497	1/1	0.95	0.19	34,34,34,34	0
85	MG	L4	401	1/1	0.95	0.09	67,67,67,67	0
85	MG	1	3468	1/1	0.95	0.10	53,53,53,53	0
85	MG	1	3793	1/1	0.95	0.17	57,57,57,57	0
85	MG	5	3742	1/1	0.95	0.11	39,39,39,39	0
85	MG	5	3885	1/1	0.95	0.27	64,64,64,64	0
85	MG	5	3418	1/1	0.95	0.31	27,27,27,27	0
85	MG	5	3509	1/1	0.95	0.31	26,26,26,26	0
85	MG	1	3437	1/1	0.95	0.21	45,45,45,45	0
86	OHX	2	2140	7/7	0.95	0.10	172,172,172,172	0
85	MG	1	3744	1/1	0.95	0.11	55,55,55,55	0
85	MG	5	3512	1/1	0.95	0.22	29,29,29,29	0
85	MG	1	3595	1/1	0.95	0.21	24,24,24,24	0
85	MG	1	3658	1/1	0.95	0.17	36,36,36,36	0
86	OHX	2	2145	7/7	0.95	0.12	128,128,128,128	0
85	MG	5	3515	1/1	0.95	0.28	31,31,31,31	0
85	MG	1	3659	1/1	0.95	0.11	40,40,40,40	0
85	MG	5	3425	1/1	0.95	0.19	41,41,41,41	0
85	MG	1	3596	1/1	0.95	0.28	27,27,27,27	0
85	MG	1	3482	1/1	0.95	0.26	44,44,44,44	0
86	OHX	5	4214	7/7	0.95	0.13	110,110,110,110	0
85	MG	1	3664	1/1	0.95	0.14	44,44,44,44	0
85	MG	M6	201	1/1	0.95	0.07	46,46,46,46	0
85	MG	1	3862	1/1	0.95	0.14	81,81,81,81	0
85	MG	M7	202	1/1	0.95	0.20	37,37,37,37	0
85	MG	1	3709	1/1	0.95	0.17	61,61,61,61	0
85	MG	1	3757	1/1	0.95	0.13	62,62,62,62	0
85	MG	7	208	1/1	0.95	0.33	52,52,52,52	0
85	MG	5	3437	1/1	0.95	0.11	35,35,35,35	0
85	MG	5	3438	1/1	0.95	0.19	47,47,47,47	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(Å ²)	Q<0.9
85	MG	5	3531	1/1	0.95	0.15	33,33,33,33	0
85	MG	5	3533	1/1	0.95	0.19	39,39,39,39	0
85	MG	6	2021	1/1	0.95	0.25	64,64,64,64	0
85	MG	1	3566	1/1	0.95	0.11	37,37,37,37	0
85	MG	1	3568	1/1	0.95	0.23	29,29,29,29	0
85	MG	5	3776	1/1	0.95	0.07	41,41,41,41	0
85	MG	8	201	1/1	0.95	0.17	36,36,36,36	0
85	MG	1	3869	1/1	0.95	0.22	54,54,54,54	0
85	MG	5	3778	1/1	0.95	0.22	71,71,71,71	0
85	MG	5	3539	1/1	0.95	0.19	35,35,35,35	0
86	OHX	1	4162	7/7	0.95	0.12	118,118,118,118	0
85	MG	1	3601	1/1	0.95	0.16	29,29,29,29	0
85	MG	5	3447	1/1	0.95	0.21	49,49,49,49	0
85	MG	6	1955	1/1	0.95	0.36	46,46,46,46	0
85	MG	1	3432	1/1	0.95	0.18	54,54,54,54	0
85	MG	1	3764	1/1	0.95	0.21	46,46,46,46	0
85	MG	1	3669	1/1	0.95	0.10	54,54,54,54	0
85	MG	5	3454	1/1	0.95	0.05	42,42,42,42	0
85	MG	5	3455	1/1	0.95	0.25	38,38,38,38	0
86	OHX	c8	203	7/7	0.95	0.10	157,157,157,157	0
85	MG	1	3523	1/1	0.95	0.27	38,38,38,38	0
85	MG	5	3667	1/1	0.95	0.16	60,60,60,60	0
85	MG	N8	202	1/1	0.95	0.17	52,52,52,52	0
86	OHX	5	3992	7/7	0.95	0.11	129,129,129,129	0
86	OHX	5	3993	7/7	0.95	0.13	113,113,113,113	0
85	MG	3	204	1/1	0.95	0.39	56,56,56,56	0
86	OHX	5	4000	7/7	0.95	0.12	120,120,120,120	0
85	MG	l3	402	1/1	0.95	0.33	31,31,31,31	0
85	MG	3	205	1/1	0.95	0.20	45,45,45,45	0
86	OHX	8	223	7/7	0.95	0.15	120,120,120,120	0
86	OHX	8	226	7/7	0.95	0.13	132,132,132,132	0
86	OHX	5	4019	7/7	0.95	0.12	113,113,113,113	0
85	MG	5	3462	1/1	0.95	0.09	41,41,41,41	0
86	OHX	1	3926	7/7	0.95	0.15	118,118,118,118	0
85	MG	5	3463	1/1	0.95	0.23	30,30,30,30	0
86	OHX	5	4026	7/7	0.95	0.13	122,122,122,122	0
86	OHX	5	4028	7/7	0.95	0.14	118,118,118,118	0
86	OHX	5	4031	7/7	0.95	0.12	126,126,126,126	0
86	OHX	1	3957	7/7	0.95	0.12	115,115,115,115	0
85	MG	5	3801	1/1	0.95	0.07	33,33,33,33	0
85	MG	1	3819	1/1	0.95	0.08	56,56,56,56	0
85	MG	5	3677	1/1	0.95	0.24	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	1	3982	7/7	0.95	0.25	96,96,96,96	0
86	OHX	5	4044	7/7	0.95	0.11	122,122,122,122	0
86	OHX	5	4047	7/7	0.95	0.13	114,114,114,114	0
86	OHX	5	4049	7/7	0.95	0.15	106,106,106,106	0
86	OHX	5	4050	7/7	0.95	0.13	116,116,116,116	0
86	OHX	5	4052	7/7	0.95	0.13	115,115,115,115	0
86	OHX	5	4053	7/7	0.95	0.11	131,131,131,131	0
86	OHX	5	4054	7/7	0.95	0.11	138,138,138,138	0
86	OHX	1	3983	7/7	0.95	0.31	98,98,98,98	0
87	ZN	E1	501	1/1	0.95	0.05	122,122,122,122	0
86	OHX	1	3984	7/7	0.95	0.12	112,112,112,112	0
86	OHX	1	3985	7/7	0.95	0.13	115,115,115,115	0
87	ZN	e1	501	1/1	0.95	0.06	167,167,167,167	0
85	MG	O7	103	1/1	0.95	0.14	44,44,44,44	0
85	MG	l8	301	1/1	0.95	0.32	70,70,70,70	0
85	MG	1	3718	1/1	0.95	0.07	34,34,34,34	0
85	MG	5	3492	1/1	0.96	0.08	48,48,48,48	0
85	MG	1	3599	1/1	0.96	0.29	23,23,23,23	0
85	MG	1	3511	1/1	0.96	0.20	34,34,34,34	0
85	MG	6	2010	1/1	0.96	0.12	49,49,49,49	0
85	MG	1	3732	1/1	0.96	0.18	75,75,75,75	0
85	MG	5	3780	1/1	0.96	0.08	54,54,54,54	0
85	MG	1	3628	1/1	0.96	0.12	40,40,40,40	0
86	OHX	L3	405	7/7	0.96	0.12	119,119,119,119	0
86	OHX	5	4090	7/7	0.96	0.12	121,121,121,121	0
85	MG	5	3681	1/1	0.96	0.22	43,43,43,43	0
85	MG	1	3657	1/1	0.96	0.29	50,50,50,50	0
85	MG	6	2014	1/1	0.96	0.06	41,41,41,41	0
86	OHX	M5	302	7/7	0.96	0.12	120,120,120,120	0
85	MG	5	3423	1/1	0.96	0.12	46,46,46,46	0
85	MG	1	3781	1/1	0.96	0.25	59,59,59,59	0
85	MG	5	3788	1/1	0.96	0.12	47,47,47,47	0
86	OHX	O1	201	7/7	0.96	0.10	118,118,118,118	0
85	MG	6	2016	1/1	0.96	0.08	43,43,43,43	0
86	OHX	Q2	503	7/7	0.96	0.21	95,95,95,95	0
85	MG	6	2017	1/1	0.96	0.24	46,46,46,46	0
86	OHX	6	2072	7/7	0.96	0.11	133,133,133,133	0
85	MG	1	3550	1/1	0.96	0.18	39,39,39,39	0
86	OHX	6	2084	7/7	0.96	0.11	131,131,131,131	0
86	OHX	6	2085	7/7	0.96	0.12	110,110,110,110	0
86	OHX	6	2090	7/7	0.96	0.14	122,122,122,122	0
85	MG	5	3430	1/1	0.96	0.09	32,32,32,32	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	6	2097	7/7	0.96	0.11	127,127,127,127	0
85	MG	1	3512	1/1	0.96	0.27	43,43,43,43	0
85	MG	5	3691	1/1	0.96	0.10	54,54,54,54	0
85	MG	1	3784	1/1	0.96	0.14	35,35,35,35	0
85	MG	5	3508	1/1	0.96	0.30	37,37,37,37	0
85	MG	O7	102	1/1	0.96	0.17	42,42,42,42	0
86	OHX	6	2108	7/7	0.96	0.10	127,127,127,127	0
85	MG	1	3513	1/1	0.96	0.27	28,28,28,28	0
85	MG	5	3696	1/1	0.96	0.09	54,54,54,54	0
86	OHX	6	2113	7/7	0.96	0.12	133,133,133,133	0
85	MG	6	2023	1/1	0.96	0.20	47,47,47,47	0
85	MG	8	204	1/1	0.96	0.27	53,53,53,53	0
85	MG	4	203	1/1	0.96	0.28	52,52,52,52	0
86	OHX	5	4122	7/7	0.96	0.09	154,154,154,154	0
86	OHX	6	2117	7/7	0.96	0.12	117,117,117,117	0
85	MG	1	3698	1/1	0.96	0.11	41,41,41,41	0
85	MG	1	3533	1/1	0.96	0.25	41,41,41,41	0
85	MG	1	3662	1/1	0.96	0.11	43,43,43,43	0
86	OHX	6	2122	7/7	0.96	0.10	138,138,138,138	0
85	MG	5	3608	1/1	0.96	0.21	49,49,49,49	0
85	MG	1	3663	1/1	0.96	0.05	35,35,35,35	0
85	MG	5	3441	1/1	0.96	0.15	32,32,32,32	0
85	MG	5	3611	1/1	0.96	0.17	33,33,33,33	0
85	MG	5	3442	1/1	0.96	0.19	45,45,45,45	0
86	OHX	6	2129	7/7	0.96	0.10	141,141,141,141	0
85	MG	1	3487	1/1	0.96	0.35	49,49,49,49	0
85	MG	5	3444	1/1	0.96	0.15	39,39,39,39	0
85	MG	5	3615	1/1	0.96	0.19	41,41,41,41	0
85	MG	1	3449	1/1	0.96	0.11	36,36,36,36	0
85	MG	5	3819	1/1	0.96	0.13	65,65,65,65	0
85	MG	5	3617	1/1	0.96	0.08	44,44,44,44	0
85	MG	1	3422	1/1	0.96	0.21	38,38,38,38	0
85	MG	1	3844	1/1	0.96	0.28	37,37,37,37	0
85	MG	1	3442	1/1	0.96	0.06	47,47,47,47	0
85	MG	5	3826	1/1	0.96	0.10	43,43,43,43	0
85	MG	5	3449	1/1	0.96	0.20	45,45,45,45	0
85	MG	5	3623	1/1	0.96	0.09	41,41,41,41	0
86	OHX	6	2145	7/7	0.96	0.12	136,136,136,136	0
85	MG	2	1997	1/1	0.96	0.10	111,111,111,111	0
85	MG	1	3795	1/1	0.96	0.10	38,38,38,38	0
85	MG	6	1972	1/1	0.96	0.12	54,54,54,54	0
86	OHX	6	2149	7/7	0.96	0.14	113,113,113,113	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	5	3453	1/1	0.96	0.33	44,44,44,44	0
86	OHX	5	4152	7/7	0.96	0.13	123,123,123,123	0
85	MG	1	3461	1/1	0.96	0.21	33,33,33,33	0
85	MG	1	3753	1/1	0.96	0.09	53,53,53,53	0
85	MG	1	3614	1/1	0.96	0.21	34,34,34,34	0
85	MG	5	3838	1/1	0.96	0.18	42,42,42,42	0
85	MG	5	3631	1/1	0.96	0.07	42,42,42,42	0
85	MG	2	1956	1/1	0.96	0.23	63,63,63,63	0
85	MG	4	221	1/1	0.96	0.18	53,53,53,53	0
85	MG	5	3728	1/1	0.96	0.18	43,43,43,43	0
85	MG	1	3673	1/1	0.96	0.14	50,50,50,50	0
85	MG	5	3461	1/1	0.96	0.07	34,34,34,34	0
85	MG	5	3731	1/1	0.96	0.19	53,53,53,53	0
85	MG	6	1920	1/1	0.96	0.28	43,43,43,43	0
85	MG	5	3544	1/1	0.96	0.25	29,29,29,29	0
85	MG	1	3713	1/1	0.96	0.22	38,38,38,38	0
85	MG	L2	302	1/1	0.96	0.11	42,42,42,42	0
85	MG	1	3759	1/1	0.96	0.10	31,31,31,31	0
85	MG	o3	201	1/1	0.96	0.20	53,53,53,53	0
85	MG	5	3641	1/1	0.96	0.15	46,46,46,46	0
86	OHX	C8	201	7/7	0.96	0.11	120,120,120,120	0
85	MG	1	3473	1/1	0.96	0.13	39,39,39,39	0
85	MG	1	3507	1/1	0.96	0.30	35,35,35,35	0
86	OHX	1	3902	7/7	0.96	0.14	84,84,84,84	0
86	OHX	1	3916	7/7	0.96	0.18	87,87,87,87	0
85	MG	1	3858	1/1	0.96	0.11	45,45,45,45	0
85	MG	1	3676	1/1	0.96	0.18	29,29,29,29	0
86	OHX	1	3941	7/7	0.96	0.12	108,108,108,108	0
86	OHX	1	3943	7/7	0.96	0.12	101,101,101,101	0
86	OHX	1	3945	7/7	0.96	0.12	108,108,108,108	0
86	OHX	1	3951	7/7	0.96	0.13	95,95,95,95	0
86	OHX	1	3955	7/7	0.96	0.13	119,119,119,119	0
85	MG	L7	303	1/1	0.96	0.12	46,46,46,46	0
86	OHX	2	2036	7/7	0.96	0.13	124,124,124,124	0
86	OHX	1	3967	7/7	0.96	0.10	123,123,123,123	0
85	MG	1	3618	1/1	0.96	0.13	57,57,57,57	0
86	OHX	2	2042	7/7	0.96	0.14	118,118,118,118	0
86	OHX	1	3978	7/7	0.96	0.12	111,111,111,111	0
85	MG	1	3809	1/1	0.96	0.08	59,59,59,59	0
85	MG	1	3619	1/1	0.96	0.27	51,51,51,51	0
86	OHX	2	2054	7/7	0.96	0.11	121,121,121,121	0
86	OHX	2	2055	7/7	0.96	0.11	135,135,135,135	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	2	2056	7/7	0.96	0.09	141,141,141,141	0
85	MG	1	3864	1/1	0.96	0.20	81,81,81,81	0
86	OHX	1	3993	7/7	0.96	0.12	127,127,127,127	0
86	OHX	1	3994	7/7	0.96	0.12	111,111,111,111	0
86	OHX	1	3996	7/7	0.96	0.13	111,111,111,111	0
86	OHX	1	3998	7/7	0.96	0.11	133,133,133,133	0
86	OHX	1	4001	7/7	0.96	0.14	106,106,106,106	0
86	OHX	2	2065	7/7	0.96	0.11	142,142,142,142	0
86	OHX	1	4004	7/7	0.96	0.13	118,118,118,118	0
86	OHX	2	2066	7/7	0.96	0.09	147,147,147,147	0
86	OHX	1	4007	7/7	0.96	0.15	104,104,104,104	0
85	MG	1	3865	1/1	0.96	0.35	47,47,47,47	0
86	OHX	2	2068	7/7	0.96	0.12	118,118,118,118	0
86	OHX	1	4178	7/7	0.96	0.12	113,113,113,113	0
86	OHX	1	4011	7/7	0.96	0.12	123,123,123,123	0
86	OHX	2	2072	7/7	0.96	0.08	151,151,151,151	0
86	OHX	2	2073	7/7	0.96	0.11	122,122,122,122	0
85	MG	5	3751	1/1	0.96	0.06	56,56,56,56	0
86	OHX	1	4016	7/7	0.96	0.12	129,129,129,129	0
85	MG	1	3647	1/1	0.96	0.12	38,38,38,38	0
85	MG	M3	201	1/1	0.96	0.09	44,44,44,44	0
85	MG	5	3754	1/1	0.96	0.16	50,50,50,50	0
85	MG	1	3403	1/1	0.96	0.18	39,39,39,39	0
85	MG	5	3563	1/1	0.96	0.27	27,27,27,27	0
86	OHX	5	3975	7/7	0.96	0.12	102,102,102,102	0
86	OHX	5	3991	7/7	0.96	0.14	102,102,102,102	0
85	MG	5	3402	1/1	0.96	0.20	33,33,33,33	0
86	OHX	2	2081	7/7	0.96	0.09	158,158,158,158	0
86	OHX	2	2083	7/7	0.96	0.13	133,133,133,133	0
86	OHX	1	4027	7/7	0.96	0.11	120,120,120,120	0
86	OHX	5	4003	7/7	0.96	0.11	116,116,116,116	0
85	MG	1	3722	1/1	0.96	0.08	38,38,38,38	0
85	MG	5	3875	1/1	0.96	0.37	40,40,40,40	0
86	OHX	5	4017	7/7	0.96	0.12	106,106,106,106	0
86	OHX	5	4018	7/7	0.96	0.11	122,122,122,122	0
86	OHX	2	2086	7/7	0.96	0.11	130,130,130,130	0
86	OHX	5	4021	7/7	0.96	0.12	116,116,116,116	0
85	MG	2	1982	1/1	0.96	0.12	72,72,72,72	0
86	OHX	2	2088	7/7	0.96	0.12	120,120,120,120	0
86	OHX	1	4035	7/7	0.96	0.10	138,138,138,138	0
85	MG	5	3877	1/1	0.96	0.26	51,51,51,51	0
86	OHX	5	4027	7/7	0.96	0.11	140,140,140,140	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	5	3570	1/1	0.96	0.26	34,34,34,34	0
86	OHX	5	4030	7/7	0.96	0.11	116,116,116,116	0
86	OHX	2	2094	7/7	0.96	0.08	159,159,159,159	0
85	MG	5	3879	1/1	0.96	0.17	37,37,37,37	0
85	MG	1	3817	1/1	0.96	0.12	41,41,41,41	0
86	OHX	5	4034	7/7	0.96	0.09	136,136,136,136	0
86	OHX	5	4036	7/7	0.96	0.12	132,132,132,132	0
86	OHX	1	4045	7/7	0.96	0.12	118,118,118,118	0
86	OHX	1	4047	7/7	0.96	0.11	115,115,115,115	0
86	OHX	1	4048	7/7	0.96	0.15	110,110,110,110	0
86	OHX	5	4040	7/7	0.96	0.16	105,105,105,105	0
86	OHX	5	4041	7/7	0.96	0.11	128,128,128,128	0
86	OHX	5	4043	7/7	0.96	0.12	127,127,127,127	0
86	OHX	2	2098	7/7	0.96	0.13	126,126,126,126	0
86	OHX	5	4045	7/7	0.96	0.14	112,112,112,112	0
85	MG	1	3526	1/1	0.96	0.19	31,31,31,31	0
86	OHX	8	219	7/7	0.96	0.12	114,114,114,114	0
86	OHX	8	220	7/7	0.96	0.12	118,118,118,118	0
86	OHX	8	221	7/7	0.96	0.11	128,128,128,128	0
86	OHX	8	222	7/7	0.96	0.11	123,123,123,123	0
86	OHX	5	4048	7/7	0.96	0.14	111,111,111,111	0
86	OHX	8	224	7/7	0.96	0.10	126,126,126,126	0
86	OHX	1	4051	7/7	0.96	0.13	119,119,119,119	0
86	OHX	1	4052	7/7	0.96	0.11	128,128,128,128	0
86	OHX	1	4053	7/7	0.96	0.10	137,137,137,137	0
85	MG	5	3484	1/1	0.96	0.28	30,30,30,30	0
86	OHX	2	2101	7/7	0.96	0.10	145,145,145,145	0
86	OHX	5	4056	7/7	0.96	0.10	150,150,150,150	0
85	MG	1	3685	1/1	0.96	0.12	45,45,45,45	0
86	OHX	13	405	7/7	0.96	0.12	123,123,123,123	0
86	OHX	1	4057	7/7	0.96	0.08	158,158,158,158	0
85	MG	1	3527	1/1	0.96	0.24	26,26,26,26	0
86	OHX	5	4060	7/7	0.96	0.08	156,156,156,156	0
85	MG	1	3448	1/1	0.96	0.30	44,44,44,44	0
86	OHX	5	4063	7/7	0.96	0.10	134,134,134,134	0
85	MG	1	3823	1/1	0.96	0.21	47,47,47,47	0
86	OHX	5	4065	7/7	0.96	0.09	148,148,148,148	0
86	OHX	2	2106	7/7	0.96	0.11	122,122,122,122	0
86	OHX	m0	301	7/7	0.96	0.11	125,125,125,125	0
86	OHX	1	4063	7/7	0.96	0.08	182,182,182,182	0
86	OHX	3	215	7/7	0.96	0.13	115,115,115,115	0
86	OHX	3	217	7/7	0.96	0.10	125,125,125,125	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	5	4070	7/7	0.96	0.12	121,121,121,121	0
85	MG	5	3770	1/1	0.96	0.12	52,52,52,52	0
85	MG	1	3573	1/1	0.96	0.21	34,34,34,34	0
86	OHX	2	2109	7/7	0.96	0.09	145,145,145,145	0
86	OHX	2	2110	7/7	0.96	0.12	128,128,128,128	0
86	OHX	5	4076	7/7	0.96	0.11	120,120,120,120	0
85	MG	6	2007	1/1	0.96	0.08	48,48,48,48	0
86	OHX	1	4069	7/7	0.96	0.17	111,111,111,111	0
86	OHX	4	231	7/7	0.96	0.13	114,114,114,114	0
85	MG	5	3673	1/1	0.96	0.09	35,35,35,35	0
86	OHX	1	3954	7/7	0.97	0.10	128,128,128,128	0
86	OHX	1	4090	7/7	0.97	0.07	191,191,191,191	0
85	MG	5	3798	1/1	0.97	0.08	44,44,44,44	0
85	MG	5	3669	1/1	0.97	0.19	49,49,49,49	0
86	OHX	1	4093	7/7	0.97	0.25	98,98,98,98	0
85	MG	1	3747	1/1	0.97	0.08	40,40,40,40	0
86	OHX	1	3960	7/7	0.97	0.11	101,101,101,101	0
86	OHX	1	3962	7/7	0.97	0.18	99,99,99,99	0
85	MG	1	3586	1/1	0.97	0.28	51,51,51,51	0
86	OHX	2	2114	7/7	0.97	0.12	126,126,126,126	0
86	OHX	1	3970	7/7	0.97	0.10	108,108,108,108	0
85	MG	4	217	1/1	0.97	0.12	61,61,61,61	0
86	OHX	1	3972	7/7	0.97	0.12	104,104,104,104	0
86	OHX	1	3973	7/7	0.97	0.08	132,132,132,132	0
86	OHX	1	3974	7/7	0.97	0.10	107,107,107,107	0
86	OHX	1	3977	7/7	0.97	0.12	108,108,108,108	0
86	OHX	3	216	7/7	0.97	0.10	109,109,109,109	0
85	MG	1	3679	1/1	0.97	0.15	48,48,48,48	0
86	OHX	3	218	7/7	0.97	0.10	129,129,129,129	0
86	OHX	1	3979	7/7	0.97	0.08	127,127,127,127	0
86	OHX	1	3981	7/7	0.97	0.12	112,112,112,112	0
85	MG	1	3750	1/1	0.97	0.13	31,31,31,31	0
85	MG	1	3751	1/1	0.97	0.10	43,43,43,43	0
85	MG	1	3571	1/1	0.97	0.16	26,26,26,26	0
85	MG	1	3642	1/1	0.97	0.18	40,40,40,40	0
86	OHX	4	228	7/7	0.97	0.12	104,104,104,104	0
86	OHX	4	230	7/7	0.97	0.11	125,125,125,125	0
85	MG	n3	202	1/1	0.97	0.07	47,47,47,47	0
85	MG	L2	301	1/1	0.97	0.10	31,31,31,31	0
86	OHX	1	3991	7/7	0.97	0.09	122,122,122,122	0
86	OHX	1	3992	7/7	0.97	0.11	117,117,117,117	0
86	OHX	4	235	7/7	0.97	0.09	139,139,139,139	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	1	3705	1/1	0.97	0.07	44,44,44,44	0
85	MG	1	3607	1/1	0.97	0.37	51,51,51,51	0
85	MG	1	3426	1/1	0.97	0.13	29,29,29,29	0
85	MG	1	3410	1/1	0.97	0.14	27,27,27,27	0
86	OHX	1	4000	7/7	0.97	0.09	139,139,139,139	0
85	MG	L4	402	1/1	0.97	0.07	32,32,32,32	0
85	MG	6	1981	1/1	0.97	0.30	58,58,58,58	0
86	OHX	5	3927	7/7	0.97	0.16	86,86,86,86	0
86	OHX	5	3938	7/7	0.97	0.15	89,89,89,89	0
86	OHX	5	3942	7/7	0.97	0.12	88,88,88,88	0
86	OHX	5	3947	7/7	0.97	0.11	101,101,101,101	0
86	OHX	5	3953	7/7	0.97	0.12	99,99,99,99	0
86	OHX	5	3961	7/7	0.97	0.12	99,99,99,99	0
86	OHX	5	3962	7/7	0.97	0.20	85,85,85,85	0
86	OHX	5	3964	7/7	0.97	0.11	105,105,105,105	0
86	OHX	5	3965	7/7	0.97	0.16	100,100,100,100	0
86	OHX	5	3972	7/7	0.97	0.11	109,109,109,109	0
86	OHX	5	4170	7/7	0.97	0.28	98,98,98,98	0
86	OHX	5	3973	7/7	0.97	0.12	95,95,95,95	0
85	MG	5	3747	1/1	0.97	0.31	30,30,30,30	0
86	OHX	5	3979	7/7	0.97	0.16	91,91,91,91	0
86	OHX	5	3988	7/7	0.97	0.11	112,112,112,112	0
86	OHX	5	3990	7/7	0.97	0.12	108,108,108,108	0
85	MG	5	3408	1/1	0.97	0.12	32,32,32,32	0
85	MG	5	3817	1/1	0.97	0.04	39,39,39,39	0
86	OHX	1	4008	7/7	0.97	0.10	128,128,128,128	0
86	OHX	5	3994	7/7	0.97	0.22	100,100,100,100	0
85	MG	5	3567	1/1	0.97	0.29	48,48,48,48	0
86	OHX	5	3999	7/7	0.97	0.22	80,80,80,80	0
85	MG	1	3758	1/1	0.97	0.06	43,43,43,43	0
85	MG	6	2025	1/1	0.97	0.08	92,92,92,92	0
86	OHX	5	4008	7/7	0.97	0.11	112,112,112,112	0
86	OHX	O2	201	7/7	0.97	0.23	101,101,101,101	0
86	OHX	5	4010	7/7	0.97	0.09	149,149,149,149	0
86	OHX	5	4012	7/7	0.97	0.11	111,111,111,111	0
86	OHX	5	4013	7/7	0.97	0.11	111,111,111,111	0
85	MG	L7	301	1/1	0.97	0.12	40,40,40,40	0
86	OHX	5	4015	7/7	0.97	0.09	124,124,124,124	0
86	OHX	O7	105	7/7	0.97	0.11	103,103,103,103	0
86	OHX	1	4013	7/7	0.97	0.10	117,117,117,117	0
86	OHX	2	2028	7/7	0.97	0.12	106,106,106,106	0
86	OHX	6	2065	7/7	0.97	0.14	114,114,114,114	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	5	4195	7/7	0.97	0.19	91,91,91,91	0
85	MG	5	3897	1/1	0.97	0.13	46,46,46,46	0
86	OHX	6	2073	7/7	0.97	0.09	131,131,131,131	0
86	OHX	2	2037	7/7	0.97	0.10	107,107,107,107	0
86	OHX	6	2077	7/7	0.97	0.10	100,100,100,100	0
85	MG	5	3412	1/1	0.97	0.20	36,36,36,36	0
85	MG	5	3755	1/1	0.97	0.07	48,48,48,48	0
86	OHX	6	2088	7/7	0.97	0.09	132,132,132,132	0
86	OHX	6	2089	7/7	0.97	0.10	116,116,116,116	0
86	OHX	1	4019	7/7	0.97	0.10	125,125,125,125	0
86	OHX	6	2093	7/7	0.97	0.09	132,132,132,132	0
86	OHX	2	2048	7/7	0.97	0.10	125,125,125,125	0
86	OHX	5	4035	7/7	0.97	0.29	96,96,96,96	0
86	OHX	6	2095	7/7	0.97	0.09	129,129,129,129	0
86	OHX	2	2049	7/7	0.97	0.09	123,123,123,123	0
86	OHX	6	2098	7/7	0.97	0.09	156,156,156,156	0
86	OHX	2	2051	7/7	0.97	0.08	121,121,121,121	0
86	OHX	6	2101	7/7	0.97	0.08	118,118,118,118	0
85	MG	6	1905	1/1	0.97	0.29	54,54,54,54	0
86	OHX	5	4042	7/7	0.97	0.15	111,111,111,111	0
85	MG	6	1944	1/1	0.97	0.32	37,37,37,37	0
86	OHX	6	2105	7/7	0.97	0.12	119,119,119,119	0
86	OHX	1	4025	7/7	0.97	0.10	116,116,116,116	0
86	OHX	5	4046	7/7	0.97	0.09	129,129,129,129	0
85	MG	1	3574	1/1	0.97	0.27	32,32,32,32	0
85	MG	1	3434	1/1	0.97	0.12	35,35,35,35	0
86	OHX	6	2109	7/7	0.97	0.11	120,120,120,120	0
85	MG	6	1947	1/1	0.97	0.25	44,44,44,44	0
86	OHX	2	2061	7/7	0.97	0.11	131,131,131,131	0
86	OHX	6	2112	7/7	0.97	0.11	111,111,111,111	0
85	MG	1	3576	1/1	0.97	0.23	21,21,21,21	0
86	OHX	5	4055	7/7	0.97	0.09	119,119,119,119	0
86	OHX	1	4032	7/7	0.97	0.09	129,129,129,129	0
86	OHX	2	2064	7/7	0.97	0.12	118,118,118,118	0
85	MG	5	3833	1/1	0.97	0.07	40,40,40,40	0
85	MG	1	3851	1/1	0.97	0.30	36,36,36,36	0
86	OHX	1	4036	7/7	0.97	0.13	112,112,112,112	0
85	MG	1	3577	1/1	0.97	0.23	28,28,28,28	0
85	MG	5	3520	1/1	0.97	0.16	28,28,28,28	0
86	OHX	1	4040	7/7	0.97	0.07	157,157,157,157	0
86	OHX	2	2070	7/7	0.97	0.10	133,133,133,133	0
86	OHX	2	2071	7/7	0.97	0.10	128,128,128,128	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	1	4043	7/7	0.97	0.11	115,115,115,115	0
85	MG	1	3435	1/1	0.97	0.11	51,51,51,51	0
86	OHX	1	4161	7/7	0.97	0.12	112,112,112,112	0
86	OHX	6	2128	7/7	0.97	0.11	152,152,152,152	0
85	MG	2	1918	1/1	0.97	0.41	55,55,55,55	0
86	OHX	6	2130	7/7	0.97	0.10	127,127,127,127	0
86	OHX	1	4046	7/7	0.97	0.09	134,134,134,134	0
85	MG	5	3643	1/1	0.97	0.20	52,52,52,52	0
85	MG	1	3824	1/1	0.97	0.23	53,53,53,53	0
86	OHX	5	4077	7/7	0.97	0.12	111,111,111,111	0
85	MG	2	1955	1/1	0.97	0.23	56,56,56,56	0
85	MG	2	1980	1/1	0.97	0.23	56,56,56,56	0
86	OHX	7	218	7/7	0.97	0.11	96,96,96,96	0
86	OHX	7	223	7/7	0.97	0.10	111,111,111,111	0
85	MG	2	2020	1/1	0.97	0.10	86,86,86,86	0
86	OHX	6	2137	7/7	0.97	0.12	130,130,130,130	0
85	MG	5	3528	1/1	0.97	0.19	27,27,27,27	0
85	MG	5	3774	1/1	0.97	0.09	36,36,36,36	0
85	MG	5	3428	1/1	0.97	0.28	45,45,45,45	0
86	OHX	2	2082	7/7	0.97	0.09	139,139,139,139	0
86	OHX	6	2142	7/7	0.97	0.10	132,132,132,132	0
85	MG	5	3429	1/1	0.97	0.14	29,29,29,29	0
85	MG	8	208	1/1	0.97	0.07	55,55,55,55	0
85	MG	5	3651	1/1	0.97	0.31	47,47,47,47	0
86	OHX	8	225	7/7	0.97	0.08	144,144,144,144	0
85	MG	M5	301	1/1	0.97	0.18	38,38,38,38	0
86	OHX	1	4060	7/7	0.97	0.08	152,152,152,152	0
85	MG	1	3567	1/1	0.97	0.34	30,30,30,30	0
85	MG	1	3492	1/1	0.97	0.13	35,35,35,35	0
85	MG	5	3655	1/1	0.97	0.13	30,30,30,30	0
86	OHX	2	2091	7/7	0.97	0.15	120,120,120,120	0
85	MG	1	3465	1/1	0.97	0.12	41,41,41,41	0
85	MG	5	3783	1/1	0.97	0.06	84,84,84,84	0
86	OHX	2	2095	7/7	0.97	0.11	138,138,138,138	0
85	MG	6	1961	1/1	0.97	0.09	47,47,47,47	0
85	MG	5	3537	1/1	0.97	0.30	34,34,34,34	0
85	MG	1	3799	1/1	0.97	0.09	60,60,60,60	0
85	MG	1	3800	1/1	0.97	0.23	32,32,32,32	0
85	MG	5	3863	1/1	0.97	0.15	41,41,41,41	0
85	MG	1	3677	1/1	0.97	0.08	72,72,72,72	0
86	OHX	SR	401	7/7	0.97	0.08	171,171,171,171	0
85	MG	5	3541	1/1	0.97	0.13	27,27,27,27	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	6	2006	1/1	0.97	0.14	53,53,53,53	0
86	OHX	1	3917	7/7	0.97	0.10	97,97,97,97	0
86	OHX	1	3923	7/7	0.97	0.11	101,101,101,101	0
86	OHX	m5	305	7/7	0.97	0.11	128,128,128,128	0
86	OHX	m6	203	7/7	0.97	0.10	101,101,101,101	0
85	MG	s8	302	1/1	0.97	0.09	50,50,50,50	0
86	OHX	1	3927	7/7	0.97	0.18	96,96,96,96	0
86	OHX	o3	203	7/7	0.97	0.12	113,113,113,113	0
85	MG	5	3489	1/1	0.97	0.26	31,31,31,31	0
85	MG	5	3869	1/1	0.97	0.19	53,53,53,53	0
85	MG	1	3834	1/1	0.97	0.12	64,64,64,64	0
86	OHX	1	3944	7/7	0.97	0.14	99,99,99,99	0
85	MG	5	3546	1/1	0.97	0.11	37,37,37,37	0
86	OHX	1	3950	7/7	0.97	0.12	105,105,105,105	0
85	MG	1	3746	1/1	0.97	0.18	47,47,47,47	0
86	OHX	1	3952	7/7	0.97	0.10	117,117,117,117	0
86	OHX	6	2068	7/7	0.98	0.08	104,104,104,104	0
86	OHX	6	2069	7/7	0.98	0.14	92,92,92,92	0
86	OHX	6	2070	7/7	0.98	0.08	112,112,112,112	0
86	OHX	6	2071	7/7	0.98	0.14	95,95,95,95	0
86	OHX	2	2057	7/7	0.98	0.10	111,111,111,111	0
86	OHX	1	3924	7/7	0.98	0.09	100,100,100,100	0
86	OHX	6	2075	7/7	0.98	0.07	111,111,111,111	0
86	OHX	2	2058	7/7	0.98	0.08	125,125,125,125	0
86	OHX	2	2060	7/7	0.98	0.08	134,134,134,134	0
86	OHX	5	3924	7/7	0.98	0.13	76,76,76,76	0
86	OHX	6	2078	7/7	0.98	0.07	109,109,109,109	0
86	OHX	5	3930	7/7	0.98	0.12	89,89,89,89	0
86	OHX	6	2080	7/7	0.98	0.08	104,104,104,104	0
86	OHX	5	3941	7/7	0.98	0.10	85,85,85,85	0
86	OHX	6	2081	7/7	0.98	0.08	110,110,110,110	0
86	OHX	5	3944	7/7	0.98	0.09	79,79,79,79	0
86	OHX	6	2082	7/7	0.98	0.08	116,116,116,116	0
86	OHX	5	3948	7/7	0.98	0.10	86,86,86,86	0
86	OHX	5	3949	7/7	0.98	0.14	108,108,108,108	0
86	OHX	5	3950	7/7	0.98	0.10	105,105,105,105	0
86	OHX	6	2083	7/7	0.98	0.08	116,116,116,116	0
86	OHX	5	3954	7/7	0.98	0.10	93,93,93,93	0
86	OHX	5	3956	7/7	0.98	0.10	90,90,90,90	0
86	OHX	5	3958	7/7	0.98	0.10	78,78,78,78	0
86	OHX	1	3930	7/7	0.98	0.10	93,93,93,93	0
86	OHX	1	3931	7/7	0.98	0.15	83,83,83,83	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	6	2086	7/7	0.98	0.08	115,115,115,115	0
86	OHX	6	2087	7/7	0.98	0.08	125,125,125,125	0
86	OHX	5	3966	7/7	0.98	0.09	93,93,93,93	0
86	OHX	5	3967	7/7	0.98	0.09	102,102,102,102	0
86	OHX	5	3969	7/7	0.98	0.17	94,94,94,94	0
86	OHX	5	3970	7/7	0.98	0.13	84,84,84,84	0
86	OHX	5	3971	7/7	0.98	0.10	86,86,86,86	0
86	OHX	1	3932	7/7	0.98	0.11	109,109,109,109	0
85	MG	1	3630	1/1	0.98	0.28	67,67,67,67	0
86	OHX	1	3935	7/7	0.98	0.08	107,107,107,107	0
86	OHX	5	3977	7/7	0.98	0.10	96,96,96,96	0
86	OHX	5	3978	7/7	0.98	0.10	109,109,109,109	0
86	OHX	6	2091	7/7	0.98	0.08	116,116,116,116	0
86	OHX	5	3981	7/7	0.98	0.10	101,101,101,101	0
86	OHX	5	3982	7/7	0.98	0.17	98,98,98,98	0
86	OHX	5	3984	7/7	0.98	0.14	91,91,91,91	0
86	OHX	5	3985	7/7	0.98	0.08	102,102,102,102	0
86	OHX	5	3986	7/7	0.98	0.08	121,121,121,121	0
86	OHX	5	3987	7/7	0.98	0.11	103,103,103,103	0
86	OHX	6	2092	7/7	0.98	0.08	109,109,109,109	0
86	OHX	5	3989	7/7	0.98	0.13	97,97,97,97	0
86	OHX	1	3936	7/7	0.98	0.14	91,91,91,91	0
86	OHX	1	3938	7/7	0.98	0.11	101,101,101,101	0
86	OHX	1	3939	7/7	0.98	0.09	107,107,107,107	0
86	OHX	6	2096	7/7	0.98	0.07	157,157,157,157	0
86	OHX	1	3940	7/7	0.98	0.19	100,100,100,100	0
86	OHX	5	3995	7/7	0.98	0.07	116,116,116,116	0
86	OHX	5	3996	7/7	0.98	0.07	108,108,108,108	0
85	MG	1	3521	1/1	0.98	0.10	39,39,39,39	0
86	OHX	5	3998	7/7	0.98	0.09	118,118,118,118	0
86	OHX	1	3942	7/7	0.98	0.11	105,105,105,105	0
86	OHX	6	2100	7/7	0.98	0.08	159,159,159,159	0
86	OHX	5	4001	7/7	0.98	0.15	99,99,99,99	0
86	OHX	5	4002	7/7	0.98	0.11	108,108,108,108	0
86	OHX	2	2063	7/7	0.98	0.10	117,117,117,117	0
86	OHX	5	4004	7/7	0.98	0.07	128,128,128,128	0
86	OHX	5	4006	7/7	0.98	0.11	100,100,100,100	0
86	OHX	5	4007	7/7	0.98	0.08	112,112,112,112	0
85	MG	4	220	1/1	0.98	0.08	40,40,40,40	0
86	OHX	6	2103	7/7	0.98	0.08	130,130,130,130	0
85	MG	19	201	1/1	0.98	0.04	43,43,43,43	0
86	OHX	5	4011	7/7	0.98	0.14	98,98,98,98	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	1	3946	7/7	0.98	0.07	99,99,99,99	0
86	OHX	1	3947	7/7	0.98	0.09	102,102,102,102	0
86	OHX	1	3948	7/7	0.98	0.09	108,108,108,108	0
85	MG	5	3558	1/1	0.98	0.25	28,28,28,28	0
86	OHX	5	4016	7/7	0.98	0.10	110,110,110,110	0
85	MG	1	3776	1/1	0.98	0.10	72,72,72,72	0
85	MG	1	3816	1/1	0.98	0.16	56,56,56,56	0
86	OHX	1	3953	7/7	0.98	0.09	114,114,114,114	0
86	OHX	2	2069	7/7	0.98	0.07	132,132,132,132	0
85	MG	1	3694	1/1	0.98	0.05	41,41,41,41	0
85	MG	5	3700	1/1	0.98	0.28	48,48,48,48	0
86	OHX	5	4024	7/7	0.98	0.14	93,93,93,93	0
85	MG	1	3818	1/1	0.98	0.06	41,41,41,41	0
86	OHX	1	3959	7/7	0.98	0.09	100,100,100,100	0
85	MG	m6	202	1/1	0.98	0.06	36,36,36,36	0
86	OHX	1	3961	7/7	0.98	0.08	106,106,106,106	0
86	OHX	5	4029	7/7	0.98	0.12	109,109,109,109	0
86	OHX	6	2119	7/7	0.98	0.09	120,120,120,120	0
85	MG	5	3861	1/1	0.98	0.11	67,67,67,67	0
86	OHX	1	3963	7/7	0.98	0.16	82,82,82,82	0
86	OHX	1	3964	7/7	0.98	0.07	115,115,115,115	0
86	OHX	1	3965	7/7	0.98	0.07	110,110,110,110	0
86	OHX	1	3966	7/7	0.98	0.13	98,98,98,98	0
85	MG	1	3570	1/1	0.98	0.21	33,33,33,33	0
85	MG	5	3564	1/1	0.98	0.33	28,28,28,28	0
85	MG	1	3726	1/1	0.98	0.13	64,64,64,64	0
85	MG	m7	205	1/1	0.98	0.09	38,38,38,38	0
85	MG	5	3566	1/1	0.98	0.04	30,30,30,30	0
85	MG	L3	402	1/1	0.98	0.17	42,42,42,42	0
85	MG	6	1912	1/1	0.98	0.26	48,48,48,48	0
86	OHX	1	3975	7/7	0.98	0.07	110,110,110,110	0
86	OHX	1	3976	7/7	0.98	0.11	116,116,116,116	0
85	MG	1	3562	1/1	0.98	0.24	35,35,35,35	0
85	MG	5	3825	1/1	0.98	0.10	42,42,42,42	0
85	MG	1	3543	1/1	0.98	0.19	36,36,36,36	0
86	OHX	1	3980	7/7	0.98	0.09	108,108,108,108	0
85	MG	5	3457	1/1	0.98	0.15	31,31,31,31	0
85	MG	5	3675	1/1	0.98	0.27	31,31,31,31	0
85	MG	1	3424	1/1	0.98	0.27	44,44,44,44	0
85	MG	5	3750	1/1	0.98	0.26	51,51,51,51	0
86	OHX	2	2089	7/7	0.98	0.09	129,129,129,129	0
85	MG	5	3789	1/1	0.98	0.10	54,54,54,54	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	1	3989	7/7	0.98	0.16	101,101,101,101	0
85	MG	1	3602	1/1	0.98	0.21	33,33,33,33	0
85	MG	1	3603	1/1	0.98	0.06	36,36,36,36	0
86	OHX	2	2093	7/7	0.98	0.06	158,158,158,158	0
85	MG	1	3766	1/1	0.98	0.07	41,41,41,41	0
85	MG	5	3576	1/1	0.98	0.26	33,33,33,33	0
86	OHX	5	4062	7/7	0.98	0.08	119,119,119,119	0
86	OHX	1	3995	7/7	0.98	0.10	113,113,113,113	0
85	MG	5	3836	1/1	0.98	0.19	68,68,68,68	0
86	OHX	1	3997	7/7	0.98	0.08	137,137,137,137	0
85	MG	1	3716	1/1	0.98	0.08	54,54,54,54	0
86	OHX	1	3999	7/7	0.98	0.11	109,109,109,109	0
86	OHX	2	2024	7/7	0.98	0.14	90,90,90,90	0
86	OHX	2	2027	7/7	0.98	0.10	97,97,97,97	0
85	MG	1	3687	1/1	0.98	0.21	52,52,52,52	0
86	OHX	1	4003	7/7	0.98	0.07	151,151,151,151	0
86	OHX	3	214	7/7	0.98	0.09	101,101,101,101	0
86	OHX	2	2029	7/7	0.98	0.12	101,101,101,101	0
86	OHX	7	217	7/7	0.98	0.10	89,89,89,89	0
86	OHX	1	4005	7/7	0.98	0.19	95,95,95,95	0
86	OHX	7	219	7/7	0.98	0.09	105,105,105,105	0
86	OHX	7	220	7/7	0.98	0.09	104,104,104,104	0
86	OHX	7	222	7/7	0.98	0.08	106,106,106,106	0
86	OHX	2	2030	7/7	0.98	0.08	114,114,114,114	0
86	OHX	2	2032	7/7	0.98	0.12	111,111,111,111	0
86	OHX	2	2033	7/7	0.98	0.12	107,107,107,107	0
86	OHX	2	2034	7/7	0.98	0.11	104,104,104,104	0
85	MG	1	3402	1/1	0.98	0.32	55,55,55,55	0
86	OHX	8	218	7/7	0.98	0.06	114,114,114,114	0
85	MG	5	3797	1/1	0.98	0.12	44,44,44,44	0
86	OHX	2	2038	7/7	0.98	0.08	103,103,103,103	0
85	MG	5	3841	1/1	0.98	0.14	43,43,43,43	0
86	OHX	4	227	7/7	0.98	0.22	96,96,96,96	0
85	MG	1	3447	1/1	0.98	0.08	45,45,45,45	0
86	OHX	4	229	7/7	0.98	0.08	120,120,120,120	0
86	OHX	2	2044	7/7	0.98	0.07	106,106,106,106	0
86	OHX	2	2045	7/7	0.98	0.08	121,121,121,121	0
86	OHX	2	2046	7/7	0.98	0.06	129,129,129,129	0
86	OHX	2	2047	7/7	0.98	0.07	117,117,117,117	0
85	MG	5	3413	1/1	0.98	0.22	43,43,43,43	0
85	MG	5	3582	1/1	0.98	0.33	33,33,33,33	0
86	OHX	2	2050	7/7	0.98	0.08	109,109,109,109	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
85	MG	5	3618	1/1	0.98	0.16	34,34,34,34	0
86	OHX	l3	404	7/7	0.98	0.09	105,105,105,105	0
85	MG	1	3736	1/1	0.98	0.08	63,63,63,63	0
85	MG	1	3810	1/1	0.98	0.09	62,62,62,62	0
86	OHX	1	3876	7/7	0.98	0.16	63,63,63,63	0
86	OHX	1	3896	7/7	0.98	0.13	82,82,82,82	0
86	OHX	1	3901	7/7	0.98	0.16	85,85,85,85	0
85	MG	2	1903	1/1	0.98	0.16	40,40,40,40	0
86	OHX	1	3903	7/7	0.98	0.11	82,82,82,82	0
86	OHX	1	4030	7/7	0.98	0.09	117,117,117,117	0
86	OHX	1	3905	7/7	0.98	0.13	92,92,92,92	0
86	OHX	1	3906	7/7	0.98	0.10	91,91,91,91	0
86	OHX	1	3909	7/7	0.98	0.11	81,81,81,81	0
86	OHX	1	3911	7/7	0.98	0.09	88,88,88,88	0
86	OHX	5	4107	7/7	0.98	0.11	98,98,98,98	0
86	OHX	1	3914	7/7	0.98	0.11	91,91,91,91	0
85	MG	5	3524	1/1	0.98	0.37	43,43,43,43	0
86	OHX	O7	104	7/7	0.98	0.10	101,101,101,101	0
85	MG	1	3560	1/1	0.98	0.09	31,31,31,31	0
86	OHX	o2	201	7/7	0.98	0.12	103,103,103,103	0
86	OHX	1	3919	7/7	0.98	0.08	88,88,88,88	0
86	OHX	o7	503	7/7	0.98	0.09	108,108,108,108	0
86	OHX	q1	102	7/7	0.98	0.11	103,103,103,103	0
86	OHX	q2	502	7/7	0.98	0.15	98,98,98,98	0
86	OHX	6	2050	7/7	0.98	0.11	75,75,75,75	0
86	OHX	6	2054	7/7	0.98	0.12	87,87,87,87	0
86	OHX	6	2062	7/7	0.98	0.11	90,90,90,90	0
86	OHX	1	3921	7/7	0.98	0.08	105,105,105,105	0
86	OHX	6	2064	7/7	0.98	0.08	103,103,103,103	0
86	OHX	1	3922	7/7	0.98	0.09	90,90,90,90	0
86	OHX	6	2066	7/7	0.98	0.18	96,96,96,96	0
86	OHX	6	2067	7/7	0.98	0.13	96,96,96,96	0
86	OHX	5	3917	7/7	0.99	0.07	72,72,72,72	0
86	OHX	5	3920	7/7	0.99	0.08	71,71,71,71	0
86	OHX	5	3921	7/7	0.99	0.10	72,72,72,72	0
86	OHX	5	3922	7/7	0.99	0.07	69,69,69,69	0
86	OHX	5	4051	7/7	0.99	0.08	108,108,108,108	0
86	OHX	6	2057	7/7	0.99	0.07	92,92,92,92	0
86	OHX	5	3925	7/7	0.99	0.06	81,81,81,81	0
86	OHX	5	3926	7/7	0.99	0.06	78,78,78,78	0
86	OHX	6	2058	7/7	0.99	0.11	97,97,97,97	0
86	OHX	5	3928	7/7	0.99	0.06	69,69,69,69	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	5	3929	7/7	0.99	0.09	76,76,76,76	0
86	OHX	6	2059	7/7	0.99	0.06	86,86,86,86	0
86	OHX	5	3931	7/7	0.99	0.06	76,76,76,76	0
86	OHX	5	3932	7/7	0.99	0.09	82,82,82,82	0
86	OHX	5	3933	7/7	0.99	0.05	79,79,79,79	0
86	OHX	5	3934	7/7	0.99	0.06	75,75,75,75	0
86	OHX	5	3935	7/7	0.99	0.09	80,80,80,80	0
86	OHX	5	3936	7/7	0.99	0.11	84,84,84,84	0
86	OHX	5	3937	7/7	0.99	0.11	91,91,91,91	0
86	OHX	6	2060	7/7	0.99	0.08	86,86,86,86	0
86	OHX	5	3939	7/7	0.99	0.08	84,84,84,84	0
86	OHX	5	3940	7/7	0.99	0.07	94,94,94,94	0
86	OHX	6	2061	7/7	0.99	0.06	90,90,90,90	0
86	OHX	1	3907	7/7	0.99	0.07	72,72,72,72	0
86	OHX	5	3943	7/7	0.99	0.07	78,78,78,78	0
86	OHX	1	3908	7/7	0.99	0.06	85,85,85,85	0
86	OHX	5	3945	7/7	0.99	0.08	90,90,90,90	0
86	OHX	5	4074	7/7	0.99	0.09	102,102,102,102	0
86	OHX	5	3946	7/7	0.99	0.10	99,99,99,99	0
85	MG	5	3818	1/1	0.99	0.08	72,72,72,72	0
86	OHX	1	3910	7/7	0.99	0.09	80,80,80,80	0
85	MG	N8	201	1/1	0.99	0.19	33,33,33,33	0
86	OHX	1	3912	7/7	0.99	0.10	96,96,96,96	0
86	OHX	5	3951	7/7	0.99	0.07	79,79,79,79	0
86	OHX	5	3952	7/7	0.99	0.07	94,94,94,94	0
86	OHX	1	3969	7/7	0.99	0.13	78,78,78,78	0
86	OHX	1	3913	7/7	0.99	0.09	83,83,83,83	0
86	OHX	5	3955	7/7	0.99	0.09	86,86,86,86	0
85	MG	5	3820	1/1	0.99	0.07	59,59,59,59	0
86	OHX	5	3957	7/7	0.99	0.07	77,77,77,77	0
86	OHX	1	3915	7/7	0.99	0.08	79,79,79,79	0
86	OHX	5	3959	7/7	0.99	0.11	94,94,94,94	0
86	OHX	5	3960	7/7	0.99	0.06	93,93,93,93	0
85	MG	5	3741	1/1	0.99	0.05	56,56,56,56	0
86	OHX	2	2031	7/7	0.99	0.07	105,105,105,105	0
86	OHX	5	3963	7/7	0.99	0.10	97,97,97,97	0
86	OHX	6	2074	7/7	0.99	0.14	93,93,93,93	0
86	OHX	1	3918	7/7	0.99	0.06	88,88,88,88	0
85	MG	5	3597	1/1	0.99	0.27	22,22,22,22	0
86	OHX	1	3920	7/7	0.99	0.08	93,93,93,93	0
86	OHX	5	3968	7/7	0.99	0.07	100,100,100,100	0
85	MG	5	3761	1/1	0.99	0.09	45,45,45,45	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	6	2079	7/7	0.99	0.07	104,104,104,104	0
86	OHX	1	4038	7/7	0.99	0.07	105,105,105,105	0
85	MG	1	3699	1/1	0.99	0.18	47,47,47,47	0
86	OHX	2	2035	7/7	0.99	0.10	95,95,95,95	0
86	OHX	5	3974	7/7	0.99	0.05	104,104,104,104	0
86	OHX	2	2059	7/7	0.99	0.10	113,113,113,113	0
86	OHX	5	3976	7/7	0.99	0.12	85,85,85,85	0
86	OHX	1	3925	7/7	0.99	0.08	89,89,89,89	0
85	MG	1	3588	1/1	0.99	0.25	28,28,28,28	0
85	MG	5	3585	1/1	0.99	0.30	28,28,28,28	0
86	OHX	5	3980	7/7	0.99	0.14	89,89,89,89	0
86	OHX	1	3928	7/7	0.99	0.07	95,95,95,95	0
86	OHX	1	3986	7/7	0.99	0.13	82,82,82,82	0
86	OHX	5	3983	7/7	0.99	0.09	96,96,96,96	0
86	OHX	1	3987	7/7	0.99	0.07	117,117,117,117	0
86	OHX	1	3929	7/7	0.99	0.09	84,84,84,84	0
85	MG	q0	202	1/1	0.99	0.06	47,47,47,47	0
85	MG	5	3851	1/1	0.99	0.10	49,49,49,49	0
86	OHX	2	2040	7/7	0.99	0.07	103,103,103,103	0
86	OHX	4	224	7/7	0.99	0.10	61,61,61,61	0
86	OHX	4	226	7/7	0.99	0.10	82,82,82,82	0
86	OHX	1	3933	7/7	0.99	0.07	90,90,90,90	0
86	OHX	2	2041	7/7	0.99	0.06	104,104,104,104	0
85	MG	1	3670	1/1	0.99	0.16	41,41,41,41	0
86	OHX	2	2043	7/7	0.99	0.07	107,107,107,107	0
86	OHX	7	221	7/7	0.99	0.12	101,101,101,101	0
86	OHX	1	3937	7/7	0.99	0.09	92,92,92,92	0
86	OHX	1	3877	7/7	0.99	0.08	55,55,55,55	0
86	OHX	1	3878	7/7	0.99	0.13	60,60,60,60	0
86	OHX	1	3879	7/7	0.99	0.08	58,58,58,58	0
86	OHX	1	3880	7/7	0.99	0.13	66,66,66,66	0
86	OHX	1	3881	7/7	0.99	0.08	61,61,61,61	0
86	OHX	8	217	7/7	0.99	0.09	62,62,62,62	0
86	OHX	1	3882	7/7	0.99	0.08	67,67,67,67	0
86	OHX	1	3885	7/7	0.99	0.08	70,70,70,70	0
86	OHX	1	3886	7/7	0.99	0.12	66,66,66,66	0
86	OHX	1	3888	7/7	0.99	0.09	70,70,70,70	0
86	OHX	5	4005	7/7	0.99	0.09	78,78,78,78	0
86	OHX	1	3889	7/7	0.99	0.10	71,71,71,71	0
86	OHX	1	3891	7/7	0.99	0.08	74,74,74,74	0
86	OHX	1	3949	7/7	0.99	0.07	98,98,98,98	0
86	OHX	1	3892	7/7	0.99	0.08	74,74,74,74	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	1	3894	7/7	0.99	0.11	82,82,82,82	0
85	MG	5	3532	1/1	0.99	0.30	23,23,23,23	0
86	OHX	1	3897	7/7	0.99	0.08	75,75,75,75	0
86	OHX	1	3899	7/7	0.99	0.10	77,77,77,77	0
86	OHX	N1	201	7/7	0.99	0.07	70,70,70,70	0
86	OHX	N9	101	7/7	0.99	0.10	68,68,68,68	0
86	OHX	1	3900	7/7	0.99	0.10	75,75,75,75	0
86	OHX	1	3956	7/7	0.99	0.07	109,109,109,109	0
86	OHX	2	2022	7/7	0.99	0.14	78,78,78,78	0
86	OHX	2	2023	7/7	0.99	0.12	87,87,87,87	0
86	OHX	5	4020	7/7	0.99	0.07	116,116,116,116	0
85	MG	1	3861	1/1	0.99	0.07	26,26,26,26	0
86	OHX	1	3904	7/7	0.99	0.08	80,80,80,80	0
86	OHX	6	2046	7/7	0.99	0.11	61,61,61,61	0
86	OHX	s1	302	7/7	0.99	0.10	85,85,85,85	0
86	OHX	6	2047	7/7	0.99	0.13	77,77,77,77	0
86	OHX	6	2048	7/7	0.99	0.11	68,68,68,68	0
86	OHX	6	2049	7/7	0.99	0.09	74,74,74,74	0
86	OHX	2	2025	7/7	0.99	0.10	91,91,91,91	0
86	OHX	6	2052	7/7	0.99	0.10	79,79,79,79	0
86	OHX	6	2053	7/7	0.99	0.12	82,82,82,82	0
86	OHX	2	2026	7/7	0.99	0.06	81,81,81,81	0
86	OHX	6	2055	7/7	0.99	0.10	78,78,78,78	0
86	OHX	6	2056	7/7	0.99	0.08	79,79,79,79	0
86	OHX	n3	203	7/7	0.99	0.10	91,91,91,91	0
86	OHX	n9	101	7/7	0.99	0.08	70,70,70,70	0
86	OHX	5	3898	7/7	0.99	0.12	51,51,51,51	0
86	OHX	5	3901	7/7	0.99	0.11	54,54,54,54	0
86	OHX	5	3902	7/7	0.99	0.10	56,56,56,56	0
86	OHX	5	3904	7/7	0.99	0.10	62,62,62,62	0
86	OHX	5	3905	7/7	0.99	0.10	64,64,64,64	0
86	OHX	5	3906	7/7	0.99	0.11	60,60,60,60	0
86	OHX	5	3908	7/7	0.99	0.15	73,73,73,73	0
86	OHX	5	3909	7/7	0.99	0.10	64,64,64,64	0
87	ZN	d6	101	1/1	0.99	0.02	64,64,64,64	0
86	OHX	5	3911	7/7	0.99	0.10	64,64,64,64	0
86	OHX	5	3912	7/7	0.99	0.10	68,68,68,68	0
86	OHX	5	3913	7/7	0.99	0.09	63,63,63,63	0
86	OHX	5	3914	7/7	0.99	0.08	73,73,73,73	0
86	OHX	5	3915	7/7	0.99	0.06	68,68,68,68	0
86	OHX	1	3872	7/7	1.00	0.10	47,47,47,47	0
86	OHX	5	3918	7/7	1.00	0.07	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
86	OHX	5	3919	7/7	1.00	0.06	67,67,67,67	0
86	OHX	1	3883	7/7	1.00	0.07	60,60,60,60	0
86	OHX	m5	304	7/7	1.00	0.08	67,67,67,67	0
86	OHX	5	3899	7/7	1.00	0.12	48,48,48,48	0
86	OHX	5	3900	7/7	1.00	0.11	49,49,49,49	0
86	OHX	5	3923	7/7	1.00	0.05	65,65,65,65	0
86	OHX	1	3893	7/7	1.00	0.07	75,75,75,75	0
86	OHX	1	3884	7/7	1.00	0.05	64,64,64,64	0
86	OHX	5	3903	7/7	1.00	0.12	60,60,60,60	0
86	OHX	6	2051	7/7	1.00	0.07	72,72,72,72	0
86	OHX	1	3895	7/7	1.00	0.06	75,75,75,75	0
86	OHX	1	3873	7/7	1.00	0.10	53,53,53,53	0
86	OHX	5	3907	7/7	1.00	0.06	54,54,54,54	0
86	OHX	4	225	7/7	1.00	0.09	66,66,66,66	0
87	ZN	D6	500	1/1	1.00	0.02	86,86,86,86	0
86	OHX	1	3874	7/7	1.00	0.09	52,52,52,52	0
87	ZN	D9	101	1/1	1.00	0.04	81,81,81,81	0
86	OHX	5	3910	7/7	1.00	0.06	61,61,61,61	0
87	ZN	O7	101	1/1	1.00	0.01	41,41,41,41	0
87	ZN	Q0	500	1/1	1.00	0.02	52,52,52,52	0
86	OHX	1	3898	7/7	1.00	0.07	74,74,74,74	0
87	ZN	Q3	501	1/1	1.00	0.02	65,65,65,65	0
86	OHX	1	3887	7/7	1.00	0.06	70,70,70,70	0
86	OHX	1	3875	7/7	1.00	0.09	57,57,57,57	0
87	ZN	d9	101	1/1	1.00	0.03	81,81,81,81	0
85	MG	1	3530	1/1	1.00	0.31	34,34,34,34	0
87	ZN	o7	501	1/1	1.00	0.03	47,47,47,47	0
87	ZN	q0	201	1/1	1.00	0.01	34,34,34,34	0
86	OHX	8	216	7/7	1.00	0.08	61,61,61,61	0
87	ZN	q3	501	1/1	1.00	0.02	73,73,73,73	0
86	OHX	1	3890	7/7	1.00	0.04	65,65,65,65	0
86	OHX	5	3916	7/7	1.00	0.06	67,67,67,67	0

5.5 Other polymers [i](#)

There are no such residues in this entry.