



## Full wwPDB EM Validation Report ⓘ

Mar 19, 2026 – 09:26 PM UTC

PDB ID : 9HZY / pdb\_00009hzy  
EMDB ID : EMD-52538  
Title : 290 A SynPspA rod after incubation with EPL  
Authors : Hudina, E.; Junglas, B.; Sachse, C.  
Deposited on : 2025-01-14  
Resolution : 4.70 Å(reported)

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

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

A user guide is available at

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

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

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

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

MolProbity : 4-5-2 with Phenix2.0  
Percentile statistics : 20250101.v01 (using entries in the PDB archive January 1st 2025)  
Ideal geometry (proteins) : Engh & Huber (2001)  
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)  
Validation Pipeline (wwPDB-VP) : 2.49

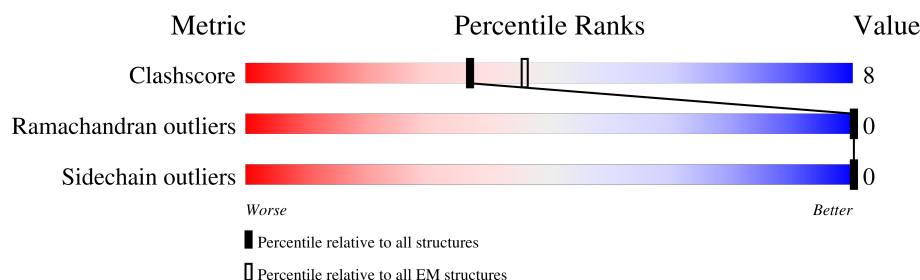
# 1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

*ELECTRON MICROSCOPY*

The reported resolution of this entry is 4.70 Å.

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
















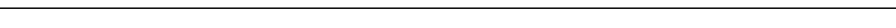











Metric	Whole archive (#Entries)	EM structures (#Entries)
Clashscore	229148	23984
Ramachandran outliers	224038	23583
Sidechain outliers	223484	23102

The table below summarises the geometric issues observed across the polymeric chains and their fit to the map. The red, orange, yellow and green segments of the bar indicate the fraction of residues that contain outliers for  $\geq 3$ , 2, 1 and 0 types of geometric quality criteria respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions  $\leq 5\%$ .

Mol	Chain	Length	Quality of chain
1	0	246	73% 15% 12%
1	1	246	72% 16% 12%
1	2	246	72% 17% 12%
1	3	246	72% 16% 12%
1	4	246	72% 16% 12%
1	5	246	73% 15% 12%
1	6	246	73% 15% 12%
1	7	246	75% 13% 12%
1	A	246	75% 13% 12%














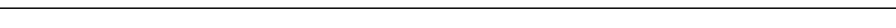











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Mol	Chain	Length	Quality of chain
1	B	246	
1	C	246	
1	D	246	
1	E	246	
1	F	246	
1	G	246	
1	H	246	
1	I	246	
1	J	246	
1	K	246	
1	L	246	
1	M	246	
1	N	246	
1	O	246	
1	P	246	
1	Q	246	
1	R	246	
1	S	246	
1	T	246	
1	U	246	
1	V	246	
1	W	246	
1	X	246	
1	Y	246	
1	Z	246	

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Mol	Chain	Length	Quality of chain
1	a	246	
1	b	246	
1	c	246	
1	d	246	
1	e	246	
1	f	246	
1	g	246	
1	h	246	
1	i	246	
1	j	246	
1	k	246	
1	l	246	
1	m	246	
1	n	246	
1	o	246	
1	p	246	
1	q	246	
1	r	246	
1	s	246	
1	t	246	
1	u	246	
1	v	246	
1	w	246	
1	x	246	
1	y	246	

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Mol	Chain	Length	Quality of chain
1	z	246	<div><div></div><div>72%</div><div>16%</div><div>12%</div></div>

## 2 Entry composition

There is only 1 type of molecule in this entry. The entry contains 212760 atoms, of which 107220 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a protein called Chloroplast membrane-associated 30 kD protein.

Mol	Chain	Residues	Atoms						AltConf	Trace
1	0	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	1	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	2	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	3	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	4	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	5	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	6	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	7	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	A	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	B	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	C	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	D	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	E	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	F	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	G	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	H	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		
1	I	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		

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Mol	Chain	Residues	Atoms						AltConf	Trace
1	J	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	K	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	L	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	M	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	N	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	O	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	P	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	Q	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	R	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	S	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	T	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	U	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	V	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	W	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	X	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	Y	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	Z	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	a	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	b	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	c	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	d	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0

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Mol	Chain	Residues	Atoms						AltConf	Trace
1	e	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	f	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	g	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	h	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	i	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	j	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	k	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	l	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	m	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	n	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	o	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	p	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	q	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	r	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	s	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	t	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	u	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	v	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	w	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	x	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0
1	y	217	Total 3546	C 1093	H 1787	N 325	O 338	S 3	0	0

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Mol	Chain	Residues	Atoms						AltConf	Trace
1	z	217	Total	C	H	N	O	S	0	0
			3546	1093	1787	325	338	3		

There are 1380 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
0	-22	MET	-	initiating methionine	UNP P74717
0	-21	GLY	-	expression tag	UNP P74717
0	-20	SER	-	expression tag	UNP P74717
0	-19	SER	-	expression tag	UNP P74717
0	-18	HIS	-	expression tag	UNP P74717
0	-17	HIS	-	expression tag	UNP P74717
0	-16	HIS	-	expression tag	UNP P74717
0	-15	HIS	-	expression tag	UNP P74717
0	-14	HIS	-	expression tag	UNP P74717
0	-13	HIS	-	expression tag	UNP P74717
0	-12	SER	-	expression tag	UNP P74717
0	-11	SER	-	expression tag	UNP P74717
0	-10	SER	-	expression tag	UNP P74717
0	-9	ALA	-	expression tag	UNP P74717
0	-8	ALA	-	expression tag	UNP P74717
0	-7	LEU	-	expression tag	UNP P74717
0	-6	GLU	-	expression tag	UNP P74717
0	-5	VAL	-	expression tag	UNP P74717
0	-4	LEU	-	expression tag	UNP P74717
0	-3	PHE	-	expression tag	UNP P74717
0	-2	GLN	-	expression tag	UNP P74717
0	-1	GLY	-	expression tag	UNP P74717
0	0	PRO	-	expression tag	UNP P74717
1	-22	MET	-	initiating methionine	UNP P74717
1	-21	GLY	-	expression tag	UNP P74717
1	-20	SER	-	expression tag	UNP P74717
1	-19	SER	-	expression tag	UNP P74717
1	-18	HIS	-	expression tag	UNP P74717
1	-17	HIS	-	expression tag	UNP P74717
1	-16	HIS	-	expression tag	UNP P74717
1	-15	HIS	-	expression tag	UNP P74717
1	-14	HIS	-	expression tag	UNP P74717
1	-13	HIS	-	expression tag	UNP P74717
1	-12	SER	-	expression tag	UNP P74717
1	-11	SER	-	expression tag	UNP P74717
1	-10	SER	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
1	-9	ALA	-	expression tag	UNP P74717
1	-8	ALA	-	expression tag	UNP P74717
1	-7	LEU	-	expression tag	UNP P74717
1	-6	GLU	-	expression tag	UNP P74717
1	-5	VAL	-	expression tag	UNP P74717
1	-4	LEU	-	expression tag	UNP P74717
1	-3	PHE	-	expression tag	UNP P74717
1	-2	GLN	-	expression tag	UNP P74717
1	-1	GLY	-	expression tag	UNP P74717
1	0	PRO	-	expression tag	UNP P74717
2	-22	MET	-	initiating methionine	UNP P74717
2	-21	GLY	-	expression tag	UNP P74717
2	-20	SER	-	expression tag	UNP P74717
2	-19	SER	-	expression tag	UNP P74717
2	-18	HIS	-	expression tag	UNP P74717
2	-17	HIS	-	expression tag	UNP P74717
2	-16	HIS	-	expression tag	UNP P74717
2	-15	HIS	-	expression tag	UNP P74717
2	-14	HIS	-	expression tag	UNP P74717
2	-13	HIS	-	expression tag	UNP P74717
2	-12	SER	-	expression tag	UNP P74717
2	-11	SER	-	expression tag	UNP P74717
2	-10	SER	-	expression tag	UNP P74717
2	-9	ALA	-	expression tag	UNP P74717
2	-8	ALA	-	expression tag	UNP P74717
2	-7	LEU	-	expression tag	UNP P74717
2	-6	GLU	-	expression tag	UNP P74717
2	-5	VAL	-	expression tag	UNP P74717
2	-4	LEU	-	expression tag	UNP P74717
2	-3	PHE	-	expression tag	UNP P74717
2	-2	GLN	-	expression tag	UNP P74717
2	-1	GLY	-	expression tag	UNP P74717
2	0	PRO	-	expression tag	UNP P74717
3	-22	MET	-	initiating methionine	UNP P74717
3	-21	GLY	-	expression tag	UNP P74717
3	-20	SER	-	expression tag	UNP P74717
3	-19	SER	-	expression tag	UNP P74717
3	-18	HIS	-	expression tag	UNP P74717
3	-17	HIS	-	expression tag	UNP P74717
3	-16	HIS	-	expression tag	UNP P74717
3	-15	HIS	-	expression tag	UNP P74717
3	-14	HIS	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
3	-13	HIS	-	expression tag	UNP P74717
3	-12	SER	-	expression tag	UNP P74717
3	-11	SER	-	expression tag	UNP P74717
3	-10	SER	-	expression tag	UNP P74717
3	-9	ALA	-	expression tag	UNP P74717
3	-8	ALA	-	expression tag	UNP P74717
3	-7	LEU	-	expression tag	UNP P74717
3	-6	GLU	-	expression tag	UNP P74717
3	-5	VAL	-	expression tag	UNP P74717
3	-4	LEU	-	expression tag	UNP P74717
3	-3	PHE	-	expression tag	UNP P74717
3	-2	GLN	-	expression tag	UNP P74717
3	-1	GLY	-	expression tag	UNP P74717
3	0	PRO	-	expression tag	UNP P74717
4	-22	MET	-	initiating methionine	UNP P74717
4	-21	GLY	-	expression tag	UNP P74717
4	-20	SER	-	expression tag	UNP P74717
4	-19	SER	-	expression tag	UNP P74717
4	-18	HIS	-	expression tag	UNP P74717
4	-17	HIS	-	expression tag	UNP P74717
4	-16	HIS	-	expression tag	UNP P74717
4	-15	HIS	-	expression tag	UNP P74717
4	-14	HIS	-	expression tag	UNP P74717
4	-13	HIS	-	expression tag	UNP P74717
4	-12	SER	-	expression tag	UNP P74717
4	-11	SER	-	expression tag	UNP P74717
4	-10	SER	-	expression tag	UNP P74717
4	-9	ALA	-	expression tag	UNP P74717
4	-8	ALA	-	expression tag	UNP P74717
4	-7	LEU	-	expression tag	UNP P74717
4	-6	GLU	-	expression tag	UNP P74717
4	-5	VAL	-	expression tag	UNP P74717
4	-4	LEU	-	expression tag	UNP P74717
4	-3	PHE	-	expression tag	UNP P74717
4	-2	GLN	-	expression tag	UNP P74717
4	-1	GLY	-	expression tag	UNP P74717
4	0	PRO	-	expression tag	UNP P74717
5	-22	MET	-	initiating methionine	UNP P74717
5	-21	GLY	-	expression tag	UNP P74717
5	-20	SER	-	expression tag	UNP P74717
5	-19	SER	-	expression tag	UNP P74717
5	-18	HIS	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
5	-17	HIS	-	expression tag	UNP P74717
5	-16	HIS	-	expression tag	UNP P74717
5	-15	HIS	-	expression tag	UNP P74717
5	-14	HIS	-	expression tag	UNP P74717
5	-13	HIS	-	expression tag	UNP P74717
5	-12	SER	-	expression tag	UNP P74717
5	-11	SER	-	expression tag	UNP P74717
5	-10	SER	-	expression tag	UNP P74717
5	-9	ALA	-	expression tag	UNP P74717
5	-8	ALA	-	expression tag	UNP P74717
5	-7	LEU	-	expression tag	UNP P74717
5	-6	GLU	-	expression tag	UNP P74717
5	-5	VAL	-	expression tag	UNP P74717
5	-4	LEU	-	expression tag	UNP P74717
5	-3	PHE	-	expression tag	UNP P74717
5	-2	GLN	-	expression tag	UNP P74717
5	-1	GLY	-	expression tag	UNP P74717
5	0	PRO	-	expression tag	UNP P74717
6	-22	MET	-	initiating methionine	UNP P74717
6	-21	GLY	-	expression tag	UNP P74717
6	-20	SER	-	expression tag	UNP P74717
6	-19	SER	-	expression tag	UNP P74717
6	-18	HIS	-	expression tag	UNP P74717
6	-17	HIS	-	expression tag	UNP P74717
6	-16	HIS	-	expression tag	UNP P74717
6	-15	HIS	-	expression tag	UNP P74717
6	-14	HIS	-	expression tag	UNP P74717
6	-13	HIS	-	expression tag	UNP P74717
6	-12	SER	-	expression tag	UNP P74717
6	-11	SER	-	expression tag	UNP P74717
6	-10	SER	-	expression tag	UNP P74717
6	-9	ALA	-	expression tag	UNP P74717
6	-8	ALA	-	expression tag	UNP P74717
6	-7	LEU	-	expression tag	UNP P74717
6	-6	GLU	-	expression tag	UNP P74717
6	-5	VAL	-	expression tag	UNP P74717
6	-4	LEU	-	expression tag	UNP P74717
6	-3	PHE	-	expression tag	UNP P74717
6	-2	GLN	-	expression tag	UNP P74717
6	-1	GLY	-	expression tag	UNP P74717
6	0	PRO	-	expression tag	UNP P74717
7	-22	MET	-	initiating methionine	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
7	-21	GLY	-	expression tag	UNP P74717
7	-20	SER	-	expression tag	UNP P74717
7	-19	SER	-	expression tag	UNP P74717
7	-18	HIS	-	expression tag	UNP P74717
7	-17	HIS	-	expression tag	UNP P74717
7	-16	HIS	-	expression tag	UNP P74717
7	-15	HIS	-	expression tag	UNP P74717
7	-14	HIS	-	expression tag	UNP P74717
7	-13	HIS	-	expression tag	UNP P74717
7	-12	SER	-	expression tag	UNP P74717
7	-11	SER	-	expression tag	UNP P74717
7	-10	SER	-	expression tag	UNP P74717
7	-9	ALA	-	expression tag	UNP P74717
7	-8	ALA	-	expression tag	UNP P74717
7	-7	LEU	-	expression tag	UNP P74717
7	-6	GLU	-	expression tag	UNP P74717
7	-5	VAL	-	expression tag	UNP P74717
7	-4	LEU	-	expression tag	UNP P74717
7	-3	PHE	-	expression tag	UNP P74717
7	-2	GLN	-	expression tag	UNP P74717
7	-1	GLY	-	expression tag	UNP P74717
7	0	PRO	-	expression tag	UNP P74717
A	-22	MET	-	initiating methionine	UNP P74717
A	-21	GLY	-	expression tag	UNP P74717
A	-20	SER	-	expression tag	UNP P74717
A	-19	SER	-	expression tag	UNP P74717
A	-18	HIS	-	expression tag	UNP P74717
A	-17	HIS	-	expression tag	UNP P74717
A	-16	HIS	-	expression tag	UNP P74717
A	-15	HIS	-	expression tag	UNP P74717
A	-14	HIS	-	expression tag	UNP P74717
A	-13	HIS	-	expression tag	UNP P74717
A	-12	SER	-	expression tag	UNP P74717
A	-11	SER	-	expression tag	UNP P74717
A	-10	SER	-	expression tag	UNP P74717
A	-9	ALA	-	expression tag	UNP P74717
A	-8	ALA	-	expression tag	UNP P74717
A	-7	LEU	-	expression tag	UNP P74717
A	-6	GLU	-	expression tag	UNP P74717
A	-5	VAL	-	expression tag	UNP P74717
A	-4	LEU	-	expression tag	UNP P74717
A	-3	PHE	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
A	-2	GLN	-	expression tag	UNP P74717
A	-1	GLY	-	expression tag	UNP P74717
A	0	PRO	-	expression tag	UNP P74717
B	-22	MET	-	initiating methionine	UNP P74717
B	-21	GLY	-	expression tag	UNP P74717
B	-20	SER	-	expression tag	UNP P74717
B	-19	SER	-	expression tag	UNP P74717
B	-18	HIS	-	expression tag	UNP P74717
B	-17	HIS	-	expression tag	UNP P74717
B	-16	HIS	-	expression tag	UNP P74717
B	-15	HIS	-	expression tag	UNP P74717
B	-14	HIS	-	expression tag	UNP P74717
B	-13	HIS	-	expression tag	UNP P74717
B	-12	SER	-	expression tag	UNP P74717
B	-11	SER	-	expression tag	UNP P74717
B	-10	SER	-	expression tag	UNP P74717
B	-9	ALA	-	expression tag	UNP P74717
B	-8	ALA	-	expression tag	UNP P74717
B	-7	LEU	-	expression tag	UNP P74717
B	-6	GLU	-	expression tag	UNP P74717
B	-5	VAL	-	expression tag	UNP P74717
B	-4	LEU	-	expression tag	UNP P74717
B	-3	PHE	-	expression tag	UNP P74717
B	-2	GLN	-	expression tag	UNP P74717
B	-1	GLY	-	expression tag	UNP P74717
B	0	PRO	-	expression tag	UNP P74717
C	-22	MET	-	initiating methionine	UNP P74717
C	-21	GLY	-	expression tag	UNP P74717
C	-20	SER	-	expression tag	UNP P74717
C	-19	SER	-	expression tag	UNP P74717
C	-18	HIS	-	expression tag	UNP P74717
C	-17	HIS	-	expression tag	UNP P74717
C	-16	HIS	-	expression tag	UNP P74717
C	-15	HIS	-	expression tag	UNP P74717
C	-14	HIS	-	expression tag	UNP P74717
C	-13	HIS	-	expression tag	UNP P74717
C	-12	SER	-	expression tag	UNP P74717
C	-11	SER	-	expression tag	UNP P74717
C	-10	SER	-	expression tag	UNP P74717
C	-9	ALA	-	expression tag	UNP P74717
C	-8	ALA	-	expression tag	UNP P74717
C	-7	LEU	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
C	-6	GLU	-	expression tag	UNP P74717
C	-5	VAL	-	expression tag	UNP P74717
C	-4	LEU	-	expression tag	UNP P74717
C	-3	PHE	-	expression tag	UNP P74717
C	-2	GLN	-	expression tag	UNP P74717
C	-1	GLY	-	expression tag	UNP P74717
C	0	PRO	-	expression tag	UNP P74717
D	-22	MET	-	initiating methionine	UNP P74717
D	-21	GLY	-	expression tag	UNP P74717
D	-20	SER	-	expression tag	UNP P74717
D	-19	SER	-	expression tag	UNP P74717
D	-18	HIS	-	expression tag	UNP P74717
D	-17	HIS	-	expression tag	UNP P74717
D	-16	HIS	-	expression tag	UNP P74717
D	-15	HIS	-	expression tag	UNP P74717
D	-14	HIS	-	expression tag	UNP P74717
D	-13	HIS	-	expression tag	UNP P74717
D	-12	SER	-	expression tag	UNP P74717
D	-11	SER	-	expression tag	UNP P74717
D	-10	SER	-	expression tag	UNP P74717
D	-9	ALA	-	expression tag	UNP P74717
D	-8	ALA	-	expression tag	UNP P74717
D	-7	LEU	-	expression tag	UNP P74717
D	-6	GLU	-	expression tag	UNP P74717
D	-5	VAL	-	expression tag	UNP P74717
D	-4	LEU	-	expression tag	UNP P74717
D	-3	PHE	-	expression tag	UNP P74717
D	-2	GLN	-	expression tag	UNP P74717
D	-1	GLY	-	expression tag	UNP P74717
D	0	PRO	-	expression tag	UNP P74717
E	-22	MET	-	initiating methionine	UNP P74717
E	-21	GLY	-	expression tag	UNP P74717
E	-20	SER	-	expression tag	UNP P74717
E	-19	SER	-	expression tag	UNP P74717
E	-18	HIS	-	expression tag	UNP P74717
E	-17	HIS	-	expression tag	UNP P74717
E	-16	HIS	-	expression tag	UNP P74717
E	-15	HIS	-	expression tag	UNP P74717
E	-14	HIS	-	expression tag	UNP P74717
E	-13	HIS	-	expression tag	UNP P74717
E	-12	SER	-	expression tag	UNP P74717
E	-11	SER	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
E	-10	SER	-	expression tag	UNP P74717
E	-9	ALA	-	expression tag	UNP P74717
E	-8	ALA	-	expression tag	UNP P74717
E	-7	LEU	-	expression tag	UNP P74717
E	-6	GLU	-	expression tag	UNP P74717
E	-5	VAL	-	expression tag	UNP P74717
E	-4	LEU	-	expression tag	UNP P74717
E	-3	PHE	-	expression tag	UNP P74717
E	-2	GLN	-	expression tag	UNP P74717
E	-1	GLY	-	expression tag	UNP P74717
E	0	PRO	-	expression tag	UNP P74717
F	-22	MET	-	initiating methionine	UNP P74717
F	-21	GLY	-	expression tag	UNP P74717
F	-20	SER	-	expression tag	UNP P74717
F	-19	SER	-	expression tag	UNP P74717
F	-18	HIS	-	expression tag	UNP P74717
F	-17	HIS	-	expression tag	UNP P74717
F	-16	HIS	-	expression tag	UNP P74717
F	-15	HIS	-	expression tag	UNP P74717
F	-14	HIS	-	expression tag	UNP P74717
F	-13	HIS	-	expression tag	UNP P74717
F	-12	SER	-	expression tag	UNP P74717
F	-11	SER	-	expression tag	UNP P74717
F	-10	SER	-	expression tag	UNP P74717
F	-9	ALA	-	expression tag	UNP P74717
F	-8	ALA	-	expression tag	UNP P74717
F	-7	LEU	-	expression tag	UNP P74717
F	-6	GLU	-	expression tag	UNP P74717
F	-5	VAL	-	expression tag	UNP P74717
F	-4	LEU	-	expression tag	UNP P74717
F	-3	PHE	-	expression tag	UNP P74717
F	-2	GLN	-	expression tag	UNP P74717
F	-1	GLY	-	expression tag	UNP P74717
F	0	PRO	-	expression tag	UNP P74717
G	-22	MET	-	initiating methionine	UNP P74717
G	-21	GLY	-	expression tag	UNP P74717
G	-20	SER	-	expression tag	UNP P74717
G	-19	SER	-	expression tag	UNP P74717
G	-18	HIS	-	expression tag	UNP P74717
G	-17	HIS	-	expression tag	UNP P74717
G	-16	HIS	-	expression tag	UNP P74717
G	-15	HIS	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
G	-14	HIS	-	expression tag	UNP P74717
G	-13	HIS	-	expression tag	UNP P74717
G	-12	SER	-	expression tag	UNP P74717
G	-11	SER	-	expression tag	UNP P74717
G	-10	SER	-	expression tag	UNP P74717
G	-9	ALA	-	expression tag	UNP P74717
G	-8	ALA	-	expression tag	UNP P74717
G	-7	LEU	-	expression tag	UNP P74717
G	-6	GLU	-	expression tag	UNP P74717
G	-5	VAL	-	expression tag	UNP P74717
G	-4	LEU	-	expression tag	UNP P74717
G	-3	PHE	-	expression tag	UNP P74717
G	-2	GLN	-	expression tag	UNP P74717
G	-1	GLY	-	expression tag	UNP P74717
G	0	PRO	-	expression tag	UNP P74717
H	-22	MET	-	initiating methionine	UNP P74717
H	-21	GLY	-	expression tag	UNP P74717
H	-20	SER	-	expression tag	UNP P74717
H	-19	SER	-	expression tag	UNP P74717
H	-18	HIS	-	expression tag	UNP P74717
H	-17	HIS	-	expression tag	UNP P74717
H	-16	HIS	-	expression tag	UNP P74717
H	-15	HIS	-	expression tag	UNP P74717
H	-14	HIS	-	expression tag	UNP P74717
H	-13	HIS	-	expression tag	UNP P74717
H	-12	SER	-	expression tag	UNP P74717
H	-11	SER	-	expression tag	UNP P74717
H	-10	SER	-	expression tag	UNP P74717
H	-9	ALA	-	expression tag	UNP P74717
H	-8	ALA	-	expression tag	UNP P74717
H	-7	LEU	-	expression tag	UNP P74717
H	-6	GLU	-	expression tag	UNP P74717
H	-5	VAL	-	expression tag	UNP P74717
H	-4	LEU	-	expression tag	UNP P74717
H	-3	PHE	-	expression tag	UNP P74717
H	-2	GLN	-	expression tag	UNP P74717
H	-1	GLY	-	expression tag	UNP P74717
H	0	PRO	-	expression tag	UNP P74717
I	-22	MET	-	initiating methionine	UNP P74717
I	-21	GLY	-	expression tag	UNP P74717
I	-20	SER	-	expression tag	UNP P74717
I	-19	SER	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
I	-18	HIS	-	expression tag	UNP P74717
I	-17	HIS	-	expression tag	UNP P74717
I	-16	HIS	-	expression tag	UNP P74717
I	-15	HIS	-	expression tag	UNP P74717
I	-14	HIS	-	expression tag	UNP P74717
I	-13	HIS	-	expression tag	UNP P74717
I	-12	SER	-	expression tag	UNP P74717
I	-11	SER	-	expression tag	UNP P74717
I	-10	SER	-	expression tag	UNP P74717
I	-9	ALA	-	expression tag	UNP P74717
I	-8	ALA	-	expression tag	UNP P74717
I	-7	LEU	-	expression tag	UNP P74717
I	-6	GLU	-	expression tag	UNP P74717
I	-5	VAL	-	expression tag	UNP P74717
I	-4	LEU	-	expression tag	UNP P74717
I	-3	PHE	-	expression tag	UNP P74717
I	-2	GLN	-	expression tag	UNP P74717
I	-1	GLY	-	expression tag	UNP P74717
I	0	PRO	-	expression tag	UNP P74717
J	-22	MET	-	initiating methionine	UNP P74717
J	-21	GLY	-	expression tag	UNP P74717
J	-20	SER	-	expression tag	UNP P74717
J	-19	SER	-	expression tag	UNP P74717
J	-18	HIS	-	expression tag	UNP P74717
J	-17	HIS	-	expression tag	UNP P74717
J	-16	HIS	-	expression tag	UNP P74717
J	-15	HIS	-	expression tag	UNP P74717
J	-14	HIS	-	expression tag	UNP P74717
J	-13	HIS	-	expression tag	UNP P74717
J	-12	SER	-	expression tag	UNP P74717
J	-11	SER	-	expression tag	UNP P74717
J	-10	SER	-	expression tag	UNP P74717
J	-9	ALA	-	expression tag	UNP P74717
J	-8	ALA	-	expression tag	UNP P74717
J	-7	LEU	-	expression tag	UNP P74717
J	-6	GLU	-	expression tag	UNP P74717
J	-5	VAL	-	expression tag	UNP P74717
J	-4	LEU	-	expression tag	UNP P74717
J	-3	PHE	-	expression tag	UNP P74717
J	-2	GLN	-	expression tag	UNP P74717
J	-1	GLY	-	expression tag	UNP P74717
J	0	PRO	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
K	-22	MET	-	initiating methionine	UNP P74717
K	-21	GLY	-	expression tag	UNP P74717
K	-20	SER	-	expression tag	UNP P74717
K	-19	SER	-	expression tag	UNP P74717
K	-18	HIS	-	expression tag	UNP P74717
K	-17	HIS	-	expression tag	UNP P74717
K	-16	HIS	-	expression tag	UNP P74717
K	-15	HIS	-	expression tag	UNP P74717
K	-14	HIS	-	expression tag	UNP P74717
K	-13	HIS	-	expression tag	UNP P74717
K	-12	SER	-	expression tag	UNP P74717
K	-11	SER	-	expression tag	UNP P74717
K	-10	SER	-	expression tag	UNP P74717
K	-9	ALA	-	expression tag	UNP P74717
K	-8	ALA	-	expression tag	UNP P74717
K	-7	LEU	-	expression tag	UNP P74717
K	-6	GLU	-	expression tag	UNP P74717
K	-5	VAL	-	expression tag	UNP P74717
K	-4	LEU	-	expression tag	UNP P74717
K	-3	PHE	-	expression tag	UNP P74717
K	-2	GLN	-	expression tag	UNP P74717
K	-1	GLY	-	expression tag	UNP P74717
K	0	PRO	-	expression tag	UNP P74717
L	-22	MET	-	initiating methionine	UNP P74717
L	-21	GLY	-	expression tag	UNP P74717
L	-20	SER	-	expression tag	UNP P74717
L	-19	SER	-	expression tag	UNP P74717
L	-18	HIS	-	expression tag	UNP P74717
L	-17	HIS	-	expression tag	UNP P74717
L	-16	HIS	-	expression tag	UNP P74717
L	-15	HIS	-	expression tag	UNP P74717
L	-14	HIS	-	expression tag	UNP P74717
L	-13	HIS	-	expression tag	UNP P74717
L	-12	SER	-	expression tag	UNP P74717
L	-11	SER	-	expression tag	UNP P74717
L	-10	SER	-	expression tag	UNP P74717
L	-9	ALA	-	expression tag	UNP P74717
L	-8	ALA	-	expression tag	UNP P74717
L	-7	LEU	-	expression tag	UNP P74717
L	-6	GLU	-	expression tag	UNP P74717
L	-5	VAL	-	expression tag	UNP P74717
L	-4	LEU	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
L	-3	PHE	-	expression tag	UNP P74717
L	-2	GLN	-	expression tag	UNP P74717
L	-1	GLY	-	expression tag	UNP P74717
L	0	PRO	-	expression tag	UNP P74717
M	-22	MET	-	initiating methionine	UNP P74717
M	-21	GLY	-	expression tag	UNP P74717
M	-20	SER	-	expression tag	UNP P74717
M	-19	SER	-	expression tag	UNP P74717
M	-18	HIS	-	expression tag	UNP P74717
M	-17	HIS	-	expression tag	UNP P74717
M	-16	HIS	-	expression tag	UNP P74717
M	-15	HIS	-	expression tag	UNP P74717
M	-14	HIS	-	expression tag	UNP P74717
M	-13	HIS	-	expression tag	UNP P74717
M	-12	SER	-	expression tag	UNP P74717
M	-11	SER	-	expression tag	UNP P74717
M	-10	SER	-	expression tag	UNP P74717
M	-9	ALA	-	expression tag	UNP P74717
M	-8	ALA	-	expression tag	UNP P74717
M	-7	LEU	-	expression tag	UNP P74717
M	-6	GLU	-	expression tag	UNP P74717
M	-5	VAL	-	expression tag	UNP P74717
M	-4	LEU	-	expression tag	UNP P74717
M	-3	PHE	-	expression tag	UNP P74717
M	-2	GLN	-	expression tag	UNP P74717
M	-1	GLY	-	expression tag	UNP P74717
M	0	PRO	-	expression tag	UNP P74717
N	-22	MET	-	initiating methionine	UNP P74717
N	-21	GLY	-	expression tag	UNP P74717
N	-20	SER	-	expression tag	UNP P74717
N	-19	SER	-	expression tag	UNP P74717
N	-18	HIS	-	expression tag	UNP P74717
N	-17	HIS	-	expression tag	UNP P74717
N	-16	HIS	-	expression tag	UNP P74717
N	-15	HIS	-	expression tag	UNP P74717
N	-14	HIS	-	expression tag	UNP P74717
N	-13	HIS	-	expression tag	UNP P74717
N	-12	SER	-	expression tag	UNP P74717
N	-11	SER	-	expression tag	UNP P74717
N	-10	SER	-	expression tag	UNP P74717
N	-9	ALA	-	expression tag	UNP P74717
N	-8	ALA	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
N	-7	LEU	-	expression tag	UNP P74717
N	-6	GLU	-	expression tag	UNP P74717
N	-5	VAL	-	expression tag	UNP P74717
N	-4	LEU	-	expression tag	UNP P74717
N	-3	PHE	-	expression tag	UNP P74717
N	-2	GLN	-	expression tag	UNP P74717
N	-1	GLY	-	expression tag	UNP P74717
N	0	PRO	-	expression tag	UNP P74717
O	-22	MET	-	initiating methionine	UNP P74717
O	-21	GLY	-	expression tag	UNP P74717
O	-20	SER	-	expression tag	UNP P74717
O	-19	SER	-	expression tag	UNP P74717
O	-18	HIS	-	expression tag	UNP P74717
O	-17	HIS	-	expression tag	UNP P74717
O	-16	HIS	-	expression tag	UNP P74717
O	-15	HIS	-	expression tag	UNP P74717
O	-14	HIS	-	expression tag	UNP P74717
O	-13	HIS	-	expression tag	UNP P74717
O	-12	SER	-	expression tag	UNP P74717
O	-11	SER	-	expression tag	UNP P74717
O	-10	SER	-	expression tag	UNP P74717
O	-9	ALA	-	expression tag	UNP P74717
O	-8	ALA	-	expression tag	UNP P74717
O	-7	LEU	-	expression tag	UNP P74717
O	-6	GLU	-	expression tag	UNP P74717
O	-5	VAL	-	expression tag	UNP P74717
O	-4	LEU	-	expression tag	UNP P74717
O	-3	PHE	-	expression tag	UNP P74717
O	-2	GLN	-	expression tag	UNP P74717
O	-1	GLY	-	expression tag	UNP P74717
O	0	PRO	-	expression tag	UNP P74717
P	-22	MET	-	initiating methionine	UNP P74717
P	-21	GLY	-	expression tag	UNP P74717
P	-20	SER	-	expression tag	UNP P74717
P	-19	SER	-	expression tag	UNP P74717
P	-18	HIS	-	expression tag	UNP P74717
P	-17	HIS	-	expression tag	UNP P74717
P	-16	HIS	-	expression tag	UNP P74717
P	-15	HIS	-	expression tag	UNP P74717
P	-14	HIS	-	expression tag	UNP P74717
P	-13	HIS	-	expression tag	UNP P74717
P	-12	SER	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
P	-11	SER	-	expression tag	UNP P74717
P	-10	SER	-	expression tag	UNP P74717
P	-9	ALA	-	expression tag	UNP P74717
P	-8	ALA	-	expression tag	UNP P74717
P	-7	LEU	-	expression tag	UNP P74717
P	-6	GLU	-	expression tag	UNP P74717
P	-5	VAL	-	expression tag	UNP P74717
P	-4	LEU	-	expression tag	UNP P74717
P	-3	PHE	-	expression tag	UNP P74717
P	-2	GLN	-	expression tag	UNP P74717
P	-1	GLY	-	expression tag	UNP P74717
P	0	PRO	-	expression tag	UNP P74717
Q	-22	MET	-	initiating methionine	UNP P74717
Q	-21	GLY	-	expression tag	UNP P74717
Q	-20	SER	-	expression tag	UNP P74717
Q	-19	SER	-	expression tag	UNP P74717
Q	-18	HIS	-	expression tag	UNP P74717
Q	-17	HIS	-	expression tag	UNP P74717
Q	-16	HIS	-	expression tag	UNP P74717
Q	-15	HIS	-	expression tag	UNP P74717
Q	-14	HIS	-	expression tag	UNP P74717
Q	-13	HIS	-	expression tag	UNP P74717
Q	-12	SER	-	expression tag	UNP P74717
Q	-11	SER	-	expression tag	UNP P74717
Q	-10	SER	-	expression tag	UNP P74717
Q	-9	ALA	-	expression tag	UNP P74717
Q	-8	ALA	-	expression tag	UNP P74717
Q	-7	LEU	-	expression tag	UNP P74717
Q	-6	GLU	-	expression tag	UNP P74717
Q	-5	VAL	-	expression tag	UNP P74717
Q	-4	LEU	-	expression tag	UNP P74717
Q	-3	PHE	-	expression tag	UNP P74717
Q	-2	GLN	-	expression tag	UNP P74717
Q	-1	GLY	-	expression tag	UNP P74717
Q	0	PRO	-	expression tag	UNP P74717
R	-22	MET	-	initiating methionine	UNP P74717
R	-21	GLY	-	expression tag	UNP P74717
R	-20	SER	-	expression tag	UNP P74717
R	-19	SER	-	expression tag	UNP P74717
R	-18	HIS	-	expression tag	UNP P74717
R	-17	HIS	-	expression tag	UNP P74717
R	-16	HIS	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
R	-15	HIS	-	expression tag	UNP P74717
R	-14	HIS	-	expression tag	UNP P74717
R	-13	HIS	-	expression tag	UNP P74717
R	-12	SER	-	expression tag	UNP P74717
R	-11	SER	-	expression tag	UNP P74717
R	-10	SER	-	expression tag	UNP P74717
R	-9	ALA	-	expression tag	UNP P74717
R	-8	ALA	-	expression tag	UNP P74717
R	-7	LEU	-	expression tag	UNP P74717
R	-6	GLU	-	expression tag	UNP P74717
R	-5	VAL	-	expression tag	UNP P74717
R	-4	LEU	-	expression tag	UNP P74717
R	-3	PHE	-	expression tag	UNP P74717
R	-2	GLN	-	expression tag	UNP P74717
R	-1	GLY	-	expression tag	UNP P74717
R	0	PRO	-	expression tag	UNP P74717
S	-22	MET	-	initiating methionine	UNP P74717
S	-21	GLY	-	expression tag	UNP P74717
S	-20	SER	-	expression tag	UNP P74717
S	-19	SER	-	expression tag	UNP P74717
S	-18	HIS	-	expression tag	UNP P74717
S	-17	HIS	-	expression tag	UNP P74717
S	-16	HIS	-	expression tag	UNP P74717
S	-15	HIS	-	expression tag	UNP P74717
S	-14	HIS	-	expression tag	UNP P74717
S	-13	HIS	-	expression tag	UNP P74717
S	-12	SER	-	expression tag	UNP P74717
S	-11	SER	-	expression tag	UNP P74717
S	-10	SER	-	expression tag	UNP P74717
S	-9	ALA	-	expression tag	UNP P74717
S	-8	ALA	-	expression tag	UNP P74717
S	-7	LEU	-	expression tag	UNP P74717
S	-6	GLU	-	expression tag	UNP P74717
S	-5	VAL	-	expression tag	UNP P74717
S	-4	LEU	-	expression tag	UNP P74717
S	-3	PHE	-	expression tag	UNP P74717
S	-2	GLN	-	expression tag	UNP P74717
S	-1	GLY	-	expression tag	UNP P74717
S	0	PRO	-	expression tag	UNP P74717
T	-22	MET	-	initiating methionine	UNP P74717
T	-21	GLY	-	expression tag	UNP P74717
T	-20	SER	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
T	-19	SER	-	expression tag	UNP P74717
T	-18	HIS	-	expression tag	UNP P74717
T	-17	HIS	-	expression tag	UNP P74717
T	-16	HIS	-	expression tag	UNP P74717
T	-15	HIS	-	expression tag	UNP P74717
T	-14	HIS	-	expression tag	UNP P74717
T	-13	HIS	-	expression tag	UNP P74717
T	-12	SER	-	expression tag	UNP P74717
T	-11	SER	-	expression tag	UNP P74717
T	-10	SER	-	expression tag	UNP P74717
T	-9	ALA	-	expression tag	UNP P74717
T	-8	ALA	-	expression tag	UNP P74717
T	-7	LEU	-	expression tag	UNP P74717
T	-6	GLU	-	expression tag	UNP P74717
T	-5	VAL	-	expression tag	UNP P74717
T	-4	LEU	-	expression tag	UNP P74717
T	-3	PHE	-	expression tag	UNP P74717
T	-2	GLN	-	expression tag	UNP P74717
T	-1	GLY	-	expression tag	UNP P74717
T	0	PRO	-	expression tag	UNP P74717
U	-22	MET	-	initiating methionine	UNP P74717
U	-21	GLY	-	expression tag	UNP P74717
U	-20	SER	-	expression tag	UNP P74717
U	-19	SER	-	expression tag	UNP P74717
U	-18	HIS	-	expression tag	UNP P74717
U	-17	HIS	-	expression tag	UNP P74717
U	-16	HIS	-	expression tag	UNP P74717
U	-15	HIS	-	expression tag	UNP P74717
U	-14	HIS	-	expression tag	UNP P74717
U	-13	HIS	-	expression tag	UNP P74717
U	-12	SER	-	expression tag	UNP P74717
U	-11	SER	-	expression tag	UNP P74717
U	-10	SER	-	expression tag	UNP P74717
U	-9	ALA	-	expression tag	UNP P74717
U	-8	ALA	-	expression tag	UNP P74717
U	-7	LEU	-	expression tag	UNP P74717
U	-6	GLU	-	expression tag	UNP P74717
U	-5	VAL	-	expression tag	UNP P74717
U	-4	LEU	-	expression tag	UNP P74717
U	-3	PHE	-	expression tag	UNP P74717
U	-2	GLN	-	expression tag	UNP P74717
U	-1	GLY	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
U	0	PRO	-	expression tag	UNP P74717
V	-22	MET	-	initiating methionine	UNP P74717
V	-21	GLY	-	expression tag	UNP P74717
V	-20	SER	-	expression tag	UNP P74717
V	-19	SER	-	expression tag	UNP P74717
V	-18	HIS	-	expression tag	UNP P74717
V	-17	HIS	-	expression tag	UNP P74717
V	-16	HIS	-	expression tag	UNP P74717
V	-15	HIS	-	expression tag	UNP P74717
V	-14	HIS	-	expression tag	UNP P74717
V	-13	HIS	-	expression tag	UNP P74717
V	-12	SER	-	expression tag	UNP P74717
V	-11	SER	-	expression tag	UNP P74717
V	-10	SER	-	expression tag	UNP P74717
V	-9	ALA	-	expression tag	UNP P74717
V	-8	ALA	-	expression tag	UNP P74717
V	-7	LEU	-	expression tag	UNP P74717
V	-6	GLU	-	expression tag	UNP P74717
V	-5	VAL	-	expression tag	UNP P74717
V	-4	LEU	-	expression tag	UNP P74717
V	-3	PHE	-	expression tag	UNP P74717
V	-2	GLN	-	expression tag	UNP P74717
V	-1	GLY	-	expression tag	UNP P74717
V	0	PRO	-	expression tag	UNP P74717
W	-22	MET	-	initiating methionine	UNP P74717
W	-21	GLY	-	expression tag	UNP P74717
W	-20	SER	-	expression tag	UNP P74717
W	-19	SER	-	expression tag	UNP P74717
W	-18	HIS	-	expression tag	UNP P74717
W	-17	HIS	-	expression tag	UNP P74717
W	-16	HIS	-	expression tag	UNP P74717
W	-15	HIS	-	expression tag	UNP P74717
W	-14	HIS	-	expression tag	UNP P74717
W	-13	HIS	-	expression tag	UNP P74717
W	-12	SER	-	expression tag	UNP P74717
W	-11	SER	-	expression tag	UNP P74717
W	-10	SER	-	expression tag	UNP P74717
W	-9	ALA	-	expression tag	UNP P74717
W	-8	ALA	-	expression tag	UNP P74717
W	-7	LEU	-	expression tag	UNP P74717
W	-6	GLU	-	expression tag	UNP P74717
W	-5	VAL	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
W	-4	LEU	-	expression tag	UNP P74717
W	-3	PHE	-	expression tag	UNP P74717
W	-2	GLN	-	expression tag	UNP P74717
W	-1	GLY	-	expression tag	UNP P74717
W	0	PRO	-	expression tag	UNP P74717
X	-22	MET	-	initiating methionine	UNP P74717
X	-21	GLY	-	expression tag	UNP P74717
X	-20	SER	-	expression tag	UNP P74717
X	-19	SER	-	expression tag	UNP P74717
X	-18	HIS	-	expression tag	UNP P74717
X	-17	HIS	-	expression tag	UNP P74717
X	-16	HIS	-	expression tag	UNP P74717
X	-15	HIS	-	expression tag	UNP P74717
X	-14	HIS	-	expression tag	UNP P74717
X	-13	HIS	-	expression tag	UNP P74717
X	-12	SER	-	expression tag	UNP P74717
X	-11	SER	-	expression tag	UNP P74717
X	-10	SER	-	expression tag	UNP P74717
X	-9	ALA	-	expression tag	UNP P74717
X	-8	ALA	-	expression tag	UNP P74717
X	-7	LEU	-	expression tag	UNP P74717
X	-6	GLU	-	expression tag	UNP P74717
X	-5	VAL	-	expression tag	UNP P74717
X	-4	LEU	-	expression tag	UNP P74717
X	-3	PHE	-	expression tag	UNP P74717
X	-2	GLN	-	expression tag	UNP P74717
X	-1	GLY	-	expression tag	UNP P74717
X	0	PRO	-	expression tag	UNP P74717
Y	-22	MET	-	initiating methionine	UNP P74717
Y	-21	GLY	-	expression tag	UNP P74717
Y	-20	SER	-	expression tag	UNP P74717
Y	-19	SER	-	expression tag	UNP P74717
Y	-18	HIS	-	expression tag	UNP P74717
Y	-17	HIS	-	expression tag	UNP P74717
Y	-16	HIS	-	expression tag	UNP P74717
Y	-15	HIS	-	expression tag	UNP P74717
Y	-14	HIS	-	expression tag	UNP P74717
Y	-13	HIS	-	expression tag	UNP P74717
Y	-12	SER	-	expression tag	UNP P74717
Y	-11	SER	-	expression tag	UNP P74717
Y	-10	SER	-	expression tag	UNP P74717
Y	-9	ALA	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
Y	-8	ALA	-	expression tag	UNP P74717
Y	-7	LEU	-	expression tag	UNP P74717
Y	-6	GLU	-	expression tag	UNP P74717
Y	-5	VAL	-	expression tag	UNP P74717
Y	-4	LEU	-	expression tag	UNP P74717
Y	-3	PHE	-	expression tag	UNP P74717
Y	-2	GLN	-	expression tag	UNP P74717
Y	-1	GLY	-	expression tag	UNP P74717
Y	0	PRO	-	expression tag	UNP P74717
Z	-22	MET	-	initiating methionine	UNP P74717
Z	-21	GLY	-	expression tag	UNP P74717
Z	-20	SER	-	expression tag	UNP P74717
Z	-19	SER	-	expression tag	UNP P74717
Z	-18	HIS	-	expression tag	UNP P74717
Z	-17	HIS	-	expression tag	UNP P74717
Z	-16	HIS	-	expression tag	UNP P74717
Z	-15	HIS	-	expression tag	UNP P74717
Z	-14	HIS	-	expression tag	UNP P74717
Z	-13	HIS	-	expression tag	UNP P74717
Z	-12	SER	-	expression tag	UNP P74717
Z	-11	SER	-	expression tag	UNP P74717
Z	-10	SER	-	expression tag	UNP P74717
Z	-9	ALA	-	expression tag	UNP P74717
Z	-8	ALA	-	expression tag	UNP P74717
Z	-7	LEU	-	expression tag	UNP P74717
Z	-6	GLU	-	expression tag	UNP P74717
Z	-5	VAL	-	expression tag	UNP P74717
Z	-4	LEU	-	expression tag	UNP P74717
Z	-3	PHE	-	expression tag	UNP P74717
Z	-2	GLN	-	expression tag	UNP P74717
Z	-1	GLY	-	expression tag	UNP P74717
Z	0	PRO	-	expression tag	UNP P74717
a	-22	MET	-	initiating methionine	UNP P74717
a	-21	GLY	-	expression tag	UNP P74717
a	-20	SER	-	expression tag	UNP P74717
a	-19	SER	-	expression tag	UNP P74717
a	-18	HIS	-	expression tag	UNP P74717
a	-17	HIS	-	expression tag	UNP P74717
a	-16	HIS	-	expression tag	UNP P74717
a	-15	HIS	-	expression tag	UNP P74717
a	-14	HIS	-	expression tag	UNP P74717
a	-13	HIS	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
a	-12	SER	-	expression tag	UNP P74717
a	-11	SER	-	expression tag	UNP P74717
a	-10	SER	-	expression tag	UNP P74717
a	-9	ALA	-	expression tag	UNP P74717
a	-8	ALA	-	expression tag	UNP P74717
a	-7	LEU	-	expression tag	UNP P74717
a	-6	GLU	-	expression tag	UNP P74717
a	-5	VAL	-	expression tag	UNP P74717
a	-4	LEU	-	expression tag	UNP P74717
a	-3	PHE	-	expression tag	UNP P74717
a	-2	GLN	-	expression tag	UNP P74717
a	-1	GLY	-	expression tag	UNP P74717
a	0	PRO	-	expression tag	UNP P74717
b	-22	MET	-	initiating methionine	UNP P74717
b	-21	GLY	-	expression tag	UNP P74717
b	-20	SER	-	expression tag	UNP P74717
b	-19	SER	-	expression tag	UNP P74717
b	-18	HIS	-	expression tag	UNP P74717
b	-17	HIS	-	expression tag	UNP P74717
b	-16	HIS	-	expression tag	UNP P74717
b	-15	HIS	-	expression tag	UNP P74717
b	-14	HIS	-	expression tag	UNP P74717
b	-13	HIS	-	expression tag	UNP P74717
b	-12	SER	-	expression tag	UNP P74717
b	-11	SER	-	expression tag	UNP P74717
b	-10	SER	-	expression tag	UNP P74717
b	-9	ALA	-	expression tag	UNP P74717
b	-8	ALA	-	expression tag	UNP P74717
b	-7	LEU	-	expression tag	UNP P74717
b	-6	GLU	-	expression tag	UNP P74717
b	-5	VAL	-	expression tag	UNP P74717
b	-4	LEU	-	expression tag	UNP P74717
b	-3	PHE	-	expression tag	UNP P74717
b	-2	GLN	-	expression tag	UNP P74717
b	-1	GLY	-	expression tag	UNP P74717
b	0	PRO	-	expression tag	UNP P74717
c	-22	MET	-	initiating methionine	UNP P74717
c	-21	GLY	-	expression tag	UNP P74717
c	-20	SER	-	expression tag	UNP P74717
c	-19	SER	-	expression tag	UNP P74717
c	-18	HIS	-	expression tag	UNP P74717
c	-17	HIS	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
c	-16	HIS	-	expression tag	UNP P74717
c	-15	HIS	-	expression tag	UNP P74717
c	-14	HIS	-	expression tag	UNP P74717
c	-13	HIS	-	expression tag	UNP P74717
c	-12	SER	-	expression tag	UNP P74717
c	-11	SER	-	expression tag	UNP P74717
c	-10	SER	-	expression tag	UNP P74717
c	-9	ALA	-	expression tag	UNP P74717
c	-8	ALA	-	expression tag	UNP P74717
c	-7	LEU	-	expression tag	UNP P74717
c	-6	GLU	-	expression tag	UNP P74717
c	-5	VAL	-	expression tag	UNP P74717
c	-4	LEU	-	expression tag	UNP P74717
c	-3	PHE	-	expression tag	UNP P74717
c	-2	GLN	-	expression tag	UNP P74717
c	-1	GLY	-	expression tag	UNP P74717
c	0	PRO	-	expression tag	UNP P74717
d	-22	MET	-	initiating methionine	UNP P74717
d	-21	GLY	-	expression tag	UNP P74717
d	-20	SER	-	expression tag	UNP P74717
d	-19	SER	-	expression tag	UNP P74717
d	-18	HIS	-	expression tag	UNP P74717
d	-17	HIS	-	expression tag	UNP P74717
d	-16	HIS	-	expression tag	UNP P74717
d	-15	HIS	-	expression tag	UNP P74717
d	-14	HIS	-	expression tag	UNP P74717
d	-13	HIS	-	expression tag	UNP P74717
d	-12	SER	-	expression tag	UNP P74717
d	-11	SER	-	expression tag	UNP P74717
d	-10	SER	-	expression tag	UNP P74717
d	-9	ALA	-	expression tag	UNP P74717
d	-8	ALA	-	expression tag	UNP P74717
d	-7	LEU	-	expression tag	UNP P74717
d	-6	GLU	-	expression tag	UNP P74717
d	-5	VAL	-	expression tag	UNP P74717
d	-4	LEU	-	expression tag	UNP P74717
d	-3	PHE	-	expression tag	UNP P74717
d	-2	GLN	-	expression tag	UNP P74717
d	-1	GLY	-	expression tag	UNP P74717
d	0	PRO	-	expression tag	UNP P74717
e	-22	MET	-	initiating methionine	UNP P74717
e	-21	GLY	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
e	-20	SER	-	expression tag	UNP P74717
e	-19	SER	-	expression tag	UNP P74717
e	-18	HIS	-	expression tag	UNP P74717
e	-17	HIS	-	expression tag	UNP P74717
e	-16	HIS	-	expression tag	UNP P74717
e	-15	HIS	-	expression tag	UNP P74717
e	-14	HIS	-	expression tag	UNP P74717
e	-13	HIS	-	expression tag	UNP P74717
e	-12	SER	-	expression tag	UNP P74717
e	-11	SER	-	expression tag	UNP P74717
e	-10	SER	-	expression tag	UNP P74717
e	-9	ALA	-	expression tag	UNP P74717
e	-8	ALA	-	expression tag	UNP P74717
e	-7	LEU	-	expression tag	UNP P74717
e	-6	GLU	-	expression tag	UNP P74717
e	-5	VAL	-	expression tag	UNP P74717
e	-4	LEU	-	expression tag	UNP P74717
e	-3	PHE	-	expression tag	UNP P74717
e	-2	GLN	-	expression tag	UNP P74717
e	-1	GLY	-	expression tag	UNP P74717
e	0	PRO	-	expression tag	UNP P74717
f	-22	MET	-	initiating methionine	UNP P74717
f	-21	GLY	-	expression tag	UNP P74717
f	-20	SER	-	expression tag	UNP P74717
f	-19	SER	-	expression tag	UNP P74717
f	-18	HIS	-	expression tag	UNP P74717
f	-17	HIS	-	expression tag	UNP P74717
f	-16	HIS	-	expression tag	UNP P74717
f	-15	HIS	-	expression tag	UNP P74717
f	-14	HIS	-	expression tag	UNP P74717
f	-13	HIS	-	expression tag	UNP P74717
f	-12	SER	-	expression tag	UNP P74717
f	-11	SER	-	expression tag	UNP P74717
f	-10	SER	-	expression tag	UNP P74717
f	-9	ALA	-	expression tag	UNP P74717
f	-8	ALA	-	expression tag	UNP P74717
f	-7	LEU	-	expression tag	UNP P74717
f	-6	GLU	-	expression tag	UNP P74717
f	-5	VAL	-	expression tag	UNP P74717
f	-4	LEU	-	expression tag	UNP P74717
f	-3	PHE	-	expression tag	UNP P74717
f	-2	GLN	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
f	-1	GLY	-	expression tag	UNP P74717
f	0	PRO	-	expression tag	UNP P74717
g	-22	MET	-	initiating methionine	UNP P74717
g	-21	GLY	-	expression tag	UNP P74717
g	-20	SER	-	expression tag	UNP P74717
g	-19	SER	-	expression tag	UNP P74717
g	-18	HIS	-	expression tag	UNP P74717
g	-17	HIS	-	expression tag	UNP P74717
g	-16	HIS	-	expression tag	UNP P74717
g	-15	HIS	-	expression tag	UNP P74717
g	-14	HIS	-	expression tag	UNP P74717
g	-13	HIS	-	expression tag	UNP P74717
g	-12	SER	-	expression tag	UNP P74717
g	-11	SER	-	expression tag	UNP P74717
g	-10	SER	-	expression tag	UNP P74717
g	-9	ALA	-	expression tag	UNP P74717
g	-8	ALA	-	expression tag	UNP P74717
g	-7	LEU	-	expression tag	UNP P74717
g	-6	GLU	-	expression tag	UNP P74717
g	-5	VAL	-	expression tag	UNP P74717
g	-4	LEU	-	expression tag	UNP P74717
g	-3	PHE	-	expression tag	UNP P74717
g	-2	GLN	-	expression tag	UNP P74717
g	-1	GLY	-	expression tag	UNP P74717
g	0	PRO	-	expression tag	UNP P74717
h	-22	MET	-	initiating methionine	UNP P74717
h	-21	GLY	-	expression tag	UNP P74717
h	-20	SER	-	expression tag	UNP P74717
h	-19	SER	-	expression tag	UNP P74717
h	-18	HIS	-	expression tag	UNP P74717
h	-17	HIS	-	expression tag	UNP P74717
h	-16	HIS	-	expression tag	UNP P74717
h	-15	HIS	-	expression tag	UNP P74717
h	-14	HIS	-	expression tag	UNP P74717
h	-13	HIS	-	expression tag	UNP P74717
h	-12	SER	-	expression tag	UNP P74717
h	-11	SER	-	expression tag	UNP P74717
h	-10	SER	-	expression tag	UNP P74717
h	-9	ALA	-	expression tag	UNP P74717
h	-8	ALA	-	expression tag	UNP P74717
h	-7	LEU	-	expression tag	UNP P74717
h	-6	GLU	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
h	-5	VAL	-	expression tag	UNP P74717
h	-4	LEU	-	expression tag	UNP P74717
h	-3	PHE	-	expression tag	UNP P74717
h	-2	GLN	-	expression tag	UNP P74717
h	-1	GLY	-	expression tag	UNP P74717
h	0	PRO	-	expression tag	UNP P74717
i	-22	MET	-	initiating methionine	UNP P74717
i	-21	GLY	-	expression tag	UNP P74717
i	-20	SER	-	expression tag	UNP P74717
i	-19	SER	-	expression tag	UNP P74717
i	-18	HIS	-	expression tag	UNP P74717
i	-17	HIS	-	expression tag	UNP P74717
i	-16	HIS	-	expression tag	UNP P74717
i	-15	HIS	-	expression tag	UNP P74717
i	-14	HIS	-	expression tag	UNP P74717
i	-13	HIS	-	expression tag	UNP P74717
i	-12	SER	-	expression tag	UNP P74717
i	-11	SER	-	expression tag	UNP P74717
i	-10	SER	-	expression tag	UNP P74717
i	-9	ALA	-	expression tag	UNP P74717
i	-8	ALA	-	expression tag	UNP P74717
i	-7	LEU	-	expression tag	UNP P74717
i	-6	GLU	-	expression tag	UNP P74717
i	-5	VAL	-	expression tag	UNP P74717
i	-4	LEU	-	expression tag	UNP P74717
i	-3	PHE	-	expression tag	UNP P74717
i	-2	GLN	-	expression tag	UNP P74717
i	-1	GLY	-	expression tag	UNP P74717
i	0	PRO	-	expression tag	UNP P74717
j	-22	MET	-	initiating methionine	UNP P74717
j	-21	GLY	-	expression tag	UNP P74717
j	-20	SER	-	expression tag	UNP P74717
j	-19	SER	-	expression tag	UNP P74717
j	-18	HIS	-	expression tag	UNP P74717
j	-17	HIS	-	expression tag	UNP P74717
j	-16	HIS	-	expression tag	UNP P74717
j	-15	HIS	-	expression tag	UNP P74717
j	-14	HIS	-	expression tag	UNP P74717
j	-13	HIS	-	expression tag	UNP P74717
j	-12	SER	-	expression tag	UNP P74717
j	-11	SER	-	expression tag	UNP P74717
j	-10	SER	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
j	-9	ALA	-	expression tag	UNP P74717
j	-8	ALA	-	expression tag	UNP P74717
j	-7	LEU	-	expression tag	UNP P74717
j	-6	GLU	-	expression tag	UNP P74717
j	-5	VAL	-	expression tag	UNP P74717
j	-4	LEU	-	expression tag	UNP P74717
j	-3	PHE	-	expression tag	UNP P74717
j	-2	GLN	-	expression tag	UNP P74717
j	-1	GLY	-	expression tag	UNP P74717
j	0	PRO	-	expression tag	UNP P74717
k	-22	MET	-	initiating methionine	UNP P74717
k	-21	GLY	-	expression tag	UNP P74717
k	-20	SER	-	expression tag	UNP P74717
k	-19	SER	-	expression tag	UNP P74717
k	-18	HIS	-	expression tag	UNP P74717
k	-17	HIS	-	expression tag	UNP P74717
k	-16	HIS	-	expression tag	UNP P74717
k	-15	HIS	-	expression tag	UNP P74717
k	-14	HIS	-	expression tag	UNP P74717
k	-13	HIS	-	expression tag	UNP P74717
k	-12	SER	-	expression tag	UNP P74717
k	-11	SER	-	expression tag	UNP P74717
k	-10	SER	-	expression tag	UNP P74717
k	-9	ALA	-	expression tag	UNP P74717
k	-8	ALA	-	expression tag	UNP P74717
k	-7	LEU	-	expression tag	UNP P74717
k	-6	GLU	-	expression tag	UNP P74717
k	-5	VAL	-	expression tag	UNP P74717
k	-4	LEU	-	expression tag	UNP P74717
k	-3	PHE	-	expression tag	UNP P74717
k	-2	GLN	-	expression tag	UNP P74717
k	-1	GLY	-	expression tag	UNP P74717
k	0	PRO	-	expression tag	UNP P74717
l	-22	MET	-	initiating methionine	UNP P74717
l	-21	GLY	-	expression tag	UNP P74717
l	-20	SER	-	expression tag	UNP P74717
l	-19	SER	-	expression tag	UNP P74717
l	-18	HIS	-	expression tag	UNP P74717
l	-17	HIS	-	expression tag	UNP P74717
l	-16	HIS	-	expression tag	UNP P74717
l	-15	HIS	-	expression tag	UNP P74717
l	-14	HIS	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
l	-13	HIS	-	expression tag	UNP P74717
l	-12	SER	-	expression tag	UNP P74717
l	-11	SER	-	expression tag	UNP P74717
l	-10	SER	-	expression tag	UNP P74717
l	-9	ALA	-	expression tag	UNP P74717
l	-8	ALA	-	expression tag	UNP P74717
l	-7	LEU	-	expression tag	UNP P74717
l	-6	GLU	-	expression tag	UNP P74717
l	-5	VAL	-	expression tag	UNP P74717
l	-4	LEU	-	expression tag	UNP P74717
l	-3	PHE	-	expression tag	UNP P74717
l	-2	GLN	-	expression tag	UNP P74717
l	-1	GLY	-	expression tag	UNP P74717
l	0	PRO	-	expression tag	UNP P74717
m	-22	MET	-	initiating methionine	UNP P74717
m	-21	GLY	-	expression tag	UNP P74717
m	-20	SER	-	expression tag	UNP P74717
m	-19	SER	-	expression tag	UNP P74717
m	-18	HIS	-	expression tag	UNP P74717
m	-17	HIS	-	expression tag	UNP P74717
m	-16	HIS	-	expression tag	UNP P74717
m	-15	HIS	-	expression tag	UNP P74717
m	-14	HIS	-	expression tag	UNP P74717
m	-13	HIS	-	expression tag	UNP P74717
m	-12	SER	-	expression tag	UNP P74717
m	-11	SER	-	expression tag	UNP P74717
m	-10	SER	-	expression tag	UNP P74717
m	-9	ALA	-	expression tag	UNP P74717
m	-8	ALA	-	expression tag	UNP P74717
m	-7	LEU	-	expression tag	UNP P74717
m	-6	GLU	-	expression tag	UNP P74717
m	-5	VAL	-	expression tag	UNP P74717
m	-4	LEU	-	expression tag	UNP P74717
m	-3	PHE	-	expression tag	UNP P74717
m	-2	GLN	-	expression tag	UNP P74717
m	-1	GLY	-	expression tag	UNP P74717
m	0	PRO	-	expression tag	UNP P74717
n	-22	MET	-	initiating methionine	UNP P74717
n	-21	GLY	-	expression tag	UNP P74717
n	-20	SER	-	expression tag	UNP P74717
n	-19	SER	-	expression tag	UNP P74717
n	-18	HIS	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
n	-17	HIS	-	expression tag	UNP P74717
n	-16	HIS	-	expression tag	UNP P74717
n	-15	HIS	-	expression tag	UNP P74717
n	-14	HIS	-	expression tag	UNP P74717
n	-13	HIS	-	expression tag	UNP P74717
n	-12	SER	-	expression tag	UNP P74717
n	-11	SER	-	expression tag	UNP P74717
n	-10	SER	-	expression tag	UNP P74717
n	-9	ALA	-	expression tag	UNP P74717
n	-8	ALA	-	expression tag	UNP P74717
n	-7	LEU	-	expression tag	UNP P74717
n	-6	GLU	-	expression tag	UNP P74717
n	-5	VAL	-	expression tag	UNP P74717
n	-4	LEU	-	expression tag	UNP P74717
n	-3	PHE	-	expression tag	UNP P74717
n	-2	GLN	-	expression tag	UNP P74717
n	-1	GLY	-	expression tag	UNP P74717
n	0	PRO	-	expression tag	UNP P74717
o	-22	MET	-	initiating methionine	UNP P74717
o	-21	GLY	-	expression tag	UNP P74717
o	-20	SER	-	expression tag	UNP P74717
o	-19	SER	-	expression tag	UNP P74717
o	-18	HIS	-	expression tag	UNP P74717
o	-17	HIS	-	expression tag	UNP P74717
o	-16	HIS	-	expression tag	UNP P74717
o	-15	HIS	-	expression tag	UNP P74717
o	-14	HIS	-	expression tag	UNP P74717
o	-13	HIS	-	expression tag	UNP P74717
o	-12	SER	-	expression tag	UNP P74717
o	-11	SER	-	expression tag	UNP P74717
o	-10	SER	-	expression tag	UNP P74717
o	-9	ALA	-	expression tag	UNP P74717
o	-8	ALA	-	expression tag	UNP P74717
o	-7	LEU	-	expression tag	UNP P74717
o	-6	GLU	-	expression tag	UNP P74717
o	-5	VAL	-	expression tag	UNP P74717
o	-4	LEU	-	expression tag	UNP P74717
o	-3	PHE	-	expression tag	UNP P74717
o	-2	GLN	-	expression tag	UNP P74717
o	-1	GLY	-	expression tag	UNP P74717
o	0	PRO	-	expression tag	UNP P74717
p	-22	MET	-	initiating methionine	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
p	-21	GLY	-	expression tag	UNP P74717
p	-20	SER	-	expression tag	UNP P74717
p	-19	SER	-	expression tag	UNP P74717
p	-18	HIS	-	expression tag	UNP P74717
p	-17	HIS	-	expression tag	UNP P74717
p	-16	HIS	-	expression tag	UNP P74717
p	-15	HIS	-	expression tag	UNP P74717
p	-14	HIS	-	expression tag	UNP P74717
p	-13	HIS	-	expression tag	UNP P74717
p	-12	SER	-	expression tag	UNP P74717
p	-11	SER	-	expression tag	UNP P74717
p	-10	SER	-	expression tag	UNP P74717
p	-9	ALA	-	expression tag	UNP P74717
p	-8	ALA	-	expression tag	UNP P74717
p	-7	LEU	-	expression tag	UNP P74717
p	-6	GLU	-	expression tag	UNP P74717
p	-5	VAL	-	expression tag	UNP P74717
p	-4	LEU	-	expression tag	UNP P74717
p	-3	PHE	-	expression tag	UNP P74717
p	-2	GLN	-	expression tag	UNP P74717
p	-1	GLY	-	expression tag	UNP P74717
p	0	PRO	-	expression tag	UNP P74717
q	-22	MET	-	initiating methionine	UNP P74717
q	-21	GLY	-	expression tag	UNP P74717
q	-20	SER	-	expression tag	UNP P74717
q	-19	SER	-	expression tag	UNP P74717
q	-18	HIS	-	expression tag	UNP P74717
q	-17	HIS	-	expression tag	UNP P74717
q	-16	HIS	-	expression tag	UNP P74717
q	-15	HIS	-	expression tag	UNP P74717
q	-14	HIS	-	expression tag	UNP P74717
q	-13	HIS	-	expression tag	UNP P74717
q	-12	SER	-	expression tag	UNP P74717
q	-11	SER	-	expression tag	UNP P74717
q	-10	SER	-	expression tag	UNP P74717
q	-9	ALA	-	expression tag	UNP P74717
q	-8	ALA	-	expression tag	UNP P74717
q	-7	LEU	-	expression tag	UNP P74717
q	-6	GLU	-	expression tag	UNP P74717
q	-5	VAL	-	expression tag	UNP P74717
q	-4	LEU	-	expression tag	UNP P74717
q	-3	PHE	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
q	-2	GLN	-	expression tag	UNP P74717
q	-1	GLY	-	expression tag	UNP P74717
q	0	PRO	-	expression tag	UNP P74717
r	-22	MET	-	initiating methionine	UNP P74717
r	-21	GLY	-	expression tag	UNP P74717
r	-20	SER	-	expression tag	UNP P74717
r	-19	SER	-	expression tag	UNP P74717
r	-18	HIS	-	expression tag	UNP P74717
r	-17	HIS	-	expression tag	UNP P74717
r	-16	HIS	-	expression tag	UNP P74717
r	-15	HIS	-	expression tag	UNP P74717
r	-14	HIS	-	expression tag	UNP P74717
r	-13	HIS	-	expression tag	UNP P74717
r	-12	SER	-	expression tag	UNP P74717
r	-11	SER	-	expression tag	UNP P74717
r	-10	SER	-	expression tag	UNP P74717
r	-9	ALA	-	expression tag	UNP P74717
r	-8	ALA	-	expression tag	UNP P74717
r	-7	LEU	-	expression tag	UNP P74717
r	-6	GLU	-	expression tag	UNP P74717
r	-5	VAL	-	expression tag	UNP P74717
r	-4	LEU	-	expression tag	UNP P74717
r	-3	PHE	-	expression tag	UNP P74717
r	-2	GLN	-	expression tag	UNP P74717
r	-1	GLY	-	expression tag	UNP P74717
r	0	PRO	-	expression tag	UNP P74717
s	-22	MET	-	initiating methionine	UNP P74717
s	-21	GLY	-	expression tag	UNP P74717
s	-20	SER	-	expression tag	UNP P74717
s	-19	SER	-	expression tag	UNP P74717
s	-18	HIS	-	expression tag	UNP P74717
s	-17	HIS	-	expression tag	UNP P74717
s	-16	HIS	-	expression tag	UNP P74717
s	-15	HIS	-	expression tag	UNP P74717
s	-14	HIS	-	expression tag	UNP P74717
s	-13	HIS	-	expression tag	UNP P74717
s	-12	SER	-	expression tag	UNP P74717
s	-11	SER	-	expression tag	UNP P74717
s	-10	SER	-	expression tag	UNP P74717
s	-9	ALA	-	expression tag	UNP P74717
s	-8	ALA	-	expression tag	UNP P74717
s	-7	LEU	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
s	-6	GLU	-	expression tag	UNP P74717
s	-5	VAL	-	expression tag	UNP P74717
s	-4	LEU	-	expression tag	UNP P74717
s	-3	PHE	-	expression tag	UNP P74717
s	-2	GLN	-	expression tag	UNP P74717
s	-1	GLY	-	expression tag	UNP P74717
s	0	PRO	-	expression tag	UNP P74717
t	-22	MET	-	initiating methionine	UNP P74717
t	-21	GLY	-	expression tag	UNP P74717
t	-20	SER	-	expression tag	UNP P74717
t	-19	SER	-	expression tag	UNP P74717
t	-18	HIS	-	expression tag	UNP P74717
t	-17	HIS	-	expression tag	UNP P74717
t	-16	HIS	-	expression tag	UNP P74717
t	-15	HIS	-	expression tag	UNP P74717
t	-14	HIS	-	expression tag	UNP P74717
t	-13	HIS	-	expression tag	UNP P74717
t	-12	SER	-	expression tag	UNP P74717
t	-11	SER	-	expression tag	UNP P74717
t	-10	SER	-	expression tag	UNP P74717
t	-9	ALA	-	expression tag	UNP P74717
t	-8	ALA	-	expression tag	UNP P74717
t	-7	LEU	-	expression tag	UNP P74717
t	-6	GLU	-	expression tag	UNP P74717
t	-5	VAL	-	expression tag	UNP P74717
t	-4	LEU	-	expression tag	UNP P74717
t	-3	PHE	-	expression tag	UNP P74717
t	-2	GLN	-	expression tag	UNP P74717
t	-1	GLY	-	expression tag	UNP P74717
t	0	PRO	-	expression tag	UNP P74717
u	-22	MET	-	initiating methionine	UNP P74717
u	-21	GLY	-	expression tag	UNP P74717
u	-20	SER	-	expression tag	UNP P74717
u	-19	SER	-	expression tag	UNP P74717
u	-18	HIS	-	expression tag	UNP P74717
u	-17	HIS	-	expression tag	UNP P74717
u	-16	HIS	-	expression tag	UNP P74717
u	-15	HIS	-	expression tag	UNP P74717
u	-14	HIS	-	expression tag	UNP P74717
u	-13	HIS	-	expression tag	UNP P74717
u	-12	SER	-	expression tag	UNP P74717
u	-11	SER	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
u	-10	SER	-	expression tag	UNP P74717
u	-9	ALA	-	expression tag	UNP P74717
u	-8	ALA	-	expression tag	UNP P74717
u	-7	LEU	-	expression tag	UNP P74717
u	-6	GLU	-	expression tag	UNP P74717
u	-5	VAL	-	expression tag	UNP P74717
u	-4	LEU	-	expression tag	UNP P74717
u	-3	PHE	-	expression tag	UNP P74717
u	-2	GLN	-	expression tag	UNP P74717
u	-1	GLY	-	expression tag	UNP P74717
u	0	PRO	-	expression tag	UNP P74717
v	-22	MET	-	initiating methionine	UNP P74717
v	-21	GLY	-	expression tag	UNP P74717
v	-20	SER	-	expression tag	UNP P74717
v	-19	SER	-	expression tag	UNP P74717
v	-18	HIS	-	expression tag	UNP P74717
v	-17	HIS	-	expression tag	UNP P74717
v	-16	HIS	-	expression tag	UNP P74717
v	-15	HIS	-	expression tag	UNP P74717
v	-14	HIS	-	expression tag	UNP P74717
v	-13	HIS	-	expression tag	UNP P74717
v	-12	SER	-	expression tag	UNP P74717
v	-11	SER	-	expression tag	UNP P74717
v	-10	SER	-	expression tag	UNP P74717
v	-9	ALA	-	expression tag	UNP P74717
v	-8	ALA	-	expression tag	UNP P74717
v	-7	LEU	-	expression tag	UNP P74717
v	-6	GLU	-	expression tag	UNP P74717
v	-5	VAL	-	expression tag	UNP P74717
v	-4	LEU	-	expression tag	UNP P74717
v	-3	PHE	-	expression tag	UNP P74717
v	-2	GLN	-	expression tag	UNP P74717
v	-1	GLY	-	expression tag	UNP P74717
v	0	PRO	-	expression tag	UNP P74717
w	-22	MET	-	initiating methionine	UNP P74717
w	-21	GLY	-	expression tag	UNP P74717
w	-20	SER	-	expression tag	UNP P74717
w	-19	SER	-	expression tag	UNP P74717
w	-18	HIS	-	expression tag	UNP P74717
w	-17	HIS	-	expression tag	UNP P74717
w	-16	HIS	-	expression tag	UNP P74717
w	-15	HIS	-	expression tag	UNP P74717

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Chain	Residue	Modelled	Actual	Comment	Reference
w	-14	HIS	-	expression tag	UNP P74717
w	-13	HIS	-	expression tag	UNP P74717
w	-12	SER	-	expression tag	UNP P74717
w	-11	SER	-	expression tag	UNP P74717
w	-10	SER	-	expression tag	UNP P74717
w	-9	ALA	-	expression tag	UNP P74717
w	-8	ALA	-	expression tag	UNP P74717
w	-7	LEU	-	expression tag	UNP P74717
w	-6	GLU	-	expression tag	UNP P74717
w	-5	VAL	-	expression tag	UNP P74717
w	-4	LEU	-	expression tag	UNP P74717
w	-3	PHE	-	expression tag	UNP P74717
w	-2	GLN	-	expression tag	UNP P74717
w	-1	GLY	-	expression tag	UNP P74717
w	0	PRO	-	expression tag	UNP P74717
x	-22	MET	-	initiating methionine	UNP P74717
x	-21	GLY	-	expression tag	UNP P74717
x	-20	SER	-	expression tag	UNP P74717
x	-19	SER	-	expression tag	UNP P74717
x	-18	HIS	-	expression tag	UNP P74717
x	-17	HIS	-	expression tag	UNP P74717
x	-16	HIS	-	expression tag	UNP P74717
x	-15	HIS	-	expression tag	UNP P74717
x	-14	HIS	-	expression tag	UNP P74717
x	-13	HIS	-	expression tag	UNP P74717
x	-12	SER	-	expression tag	UNP P74717
x	-11	SER	-	expression tag	UNP P74717
x	-10	SER	-	expression tag	UNP P74717
x	-9	ALA	-	expression tag	UNP P74717
x	-8	ALA	-	expression tag	UNP P74717
x	-7	LEU	-	expression tag	UNP P74717
x	-6	GLU	-	expression tag	UNP P74717
x	-5	VAL	-	expression tag	UNP P74717
x	-4	LEU	-	expression tag	UNP P74717
x	-3	PHE	-	expression tag	UNP P74717
x	-2	GLN	-	expression tag	UNP P74717
x	-1	GLY	-	expression tag	UNP P74717
x	0	PRO	-	expression tag	UNP P74717
y	-22	MET	-	initiating methionine	UNP P74717
y	-21	GLY	-	expression tag	UNP P74717
y	-20	SER	-	expression tag	UNP P74717
y	-19	SER	-	expression tag	UNP P74717

*Continued on next page...*

*Continued from previous page...*

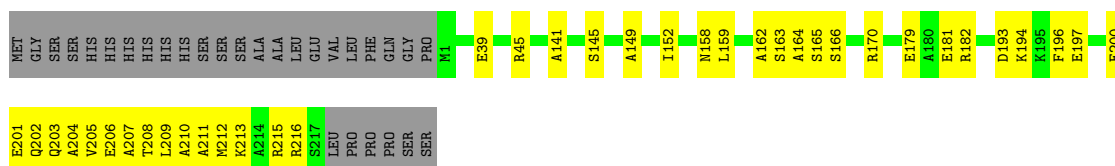
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y	-16	HIS	-	expression tag	UNP P74717
y	-15	HIS	-	expression tag	UNP P74717
y	-14	HIS	-	expression tag	UNP P74717
y	-13	HIS	-	expression tag	UNP P74717
y	-12	SER	-	expression tag	UNP P74717
y	-11	SER	-	expression tag	UNP P74717
y	-10	SER	-	expression tag	UNP P74717
y	-9	ALA	-	expression tag	UNP P74717
y	-8	ALA	-	expression tag	UNP P74717
y	-7	LEU	-	expression tag	UNP P74717
y	-6	GLU	-	expression tag	UNP P74717
y	-5	VAL	-	expression tag	UNP P74717
y	-4	LEU	-	expression tag	UNP P74717
y	-3	PHE	-	expression tag	UNP P74717
y	-2	GLN	-	expression tag	UNP P74717
y	-1	GLY	-	expression tag	UNP P74717
y	0	PRO	-	expression tag	UNP P74717
z	-22	MET	-	initiating methionine	UNP P74717
z	-21	GLY	-	expression tag	UNP P74717
z	-20	SER	-	expression tag	UNP P74717
z	-19	SER	-	expression tag	UNP P74717
z	-18	HIS	-	expression tag	UNP P74717
z	-17	HIS	-	expression tag	UNP P74717
z	-16	HIS	-	expression tag	UNP P74717
z	-15	HIS	-	expression tag	UNP P74717
z	-14	HIS	-	expression tag	UNP P74717
z	-13	HIS	-	expression tag	UNP P74717
z	-12	SER	-	expression tag	UNP P74717
z	-11	SER	-	expression tag	UNP P74717
z	-10	SER	-	expression tag	UNP P74717
z	-9	ALA	-	expression tag	UNP P74717
z	-8	ALA	-	expression tag	UNP P74717
z	-7	LEU	-	expression tag	UNP P74717
z	-6	GLU	-	expression tag	UNP P74717
z	-5	VAL	-	expression tag	UNP P74717
z	-4	LEU	-	expression tag	UNP P74717
z	-3	PHE	-	expression tag	UNP P74717
z	-2	GLN	-	expression tag	UNP P74717
z	-1	GLY	-	expression tag	UNP P74717
z	0	PRO	-	expression tag	UNP P74717

### 3 Residue-property plots

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

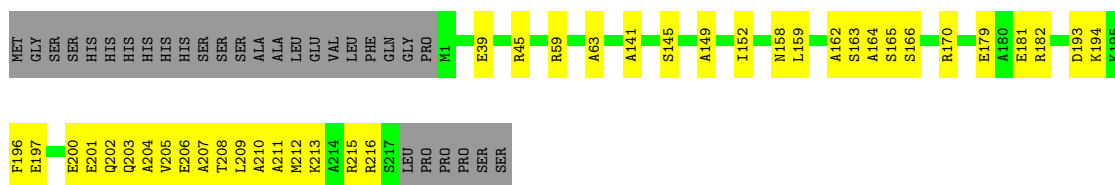
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain 0: 



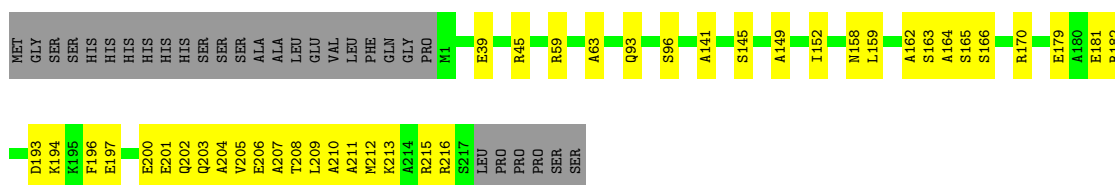
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain 1: 



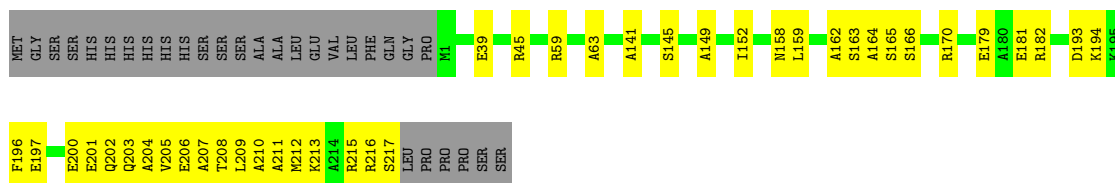
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain 2: 



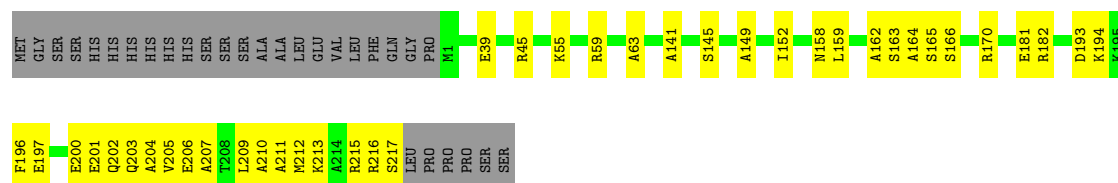
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain 3: 



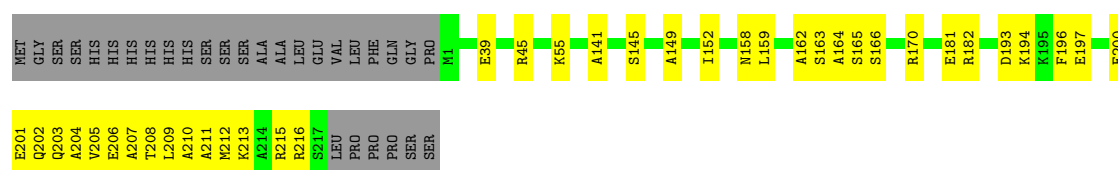
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain 4:  72% 16% 12%



- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain 5:  73% 15% 12%




- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain 6:  73% 15% 12%




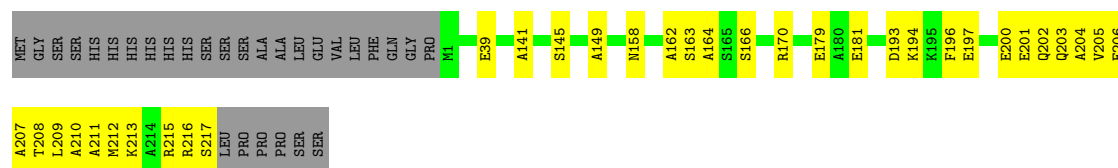
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain 7:  75% 13% 12%



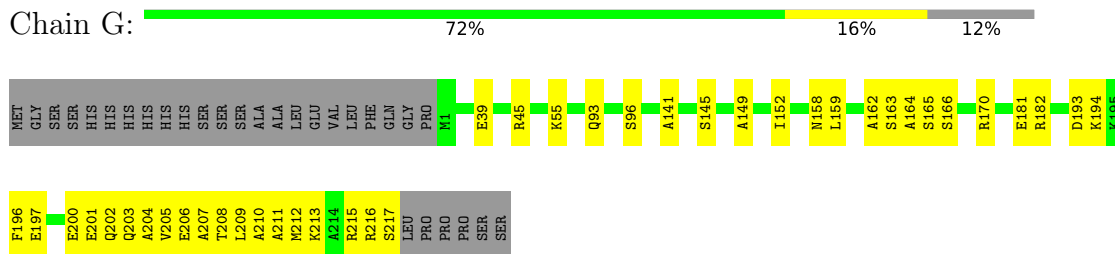
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain A:  75% 13% 12%

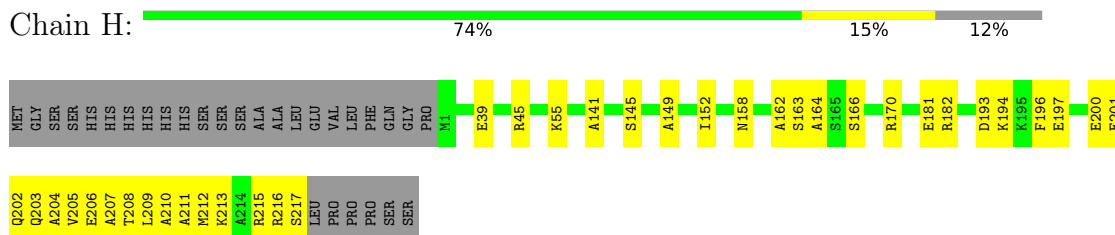


- Molecule 1: Chloroplast membrane-associated 30 kD protein

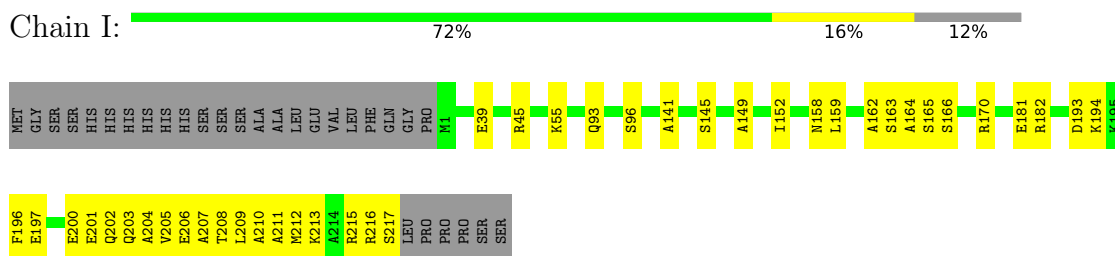




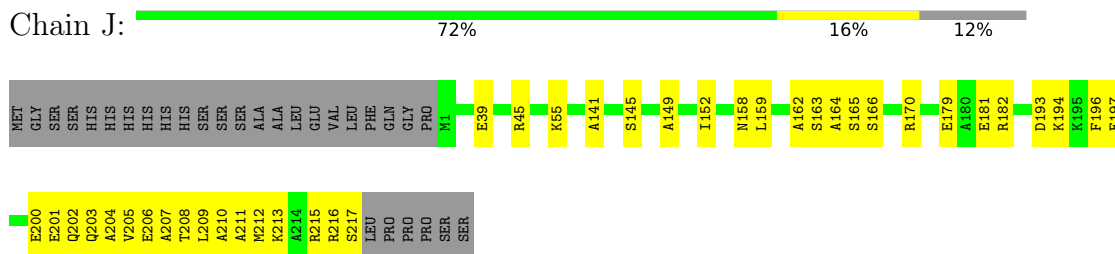
- Molecule 1: Chloroplast membrane-associated 30 kD protein



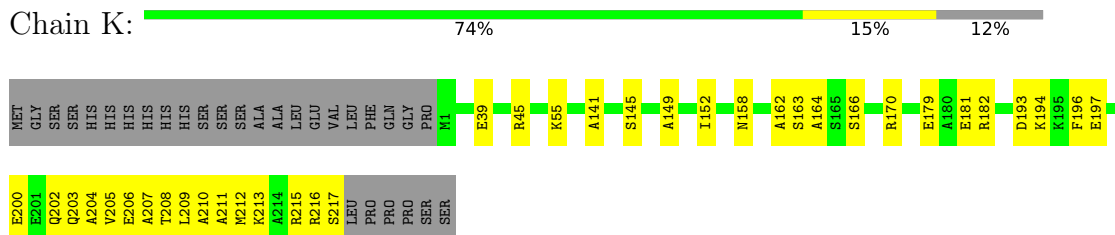
- Molecule 1: Chloroplast membrane-associated 30 kD protein



- Molecule 1: Chloroplast membrane-associated 30 kD protein



- Molecule 1: Chloroplast membrane-associated 30 kD protein



- Molecule 1: Chloroplast membrane-associated 30 kD protein

E200	ME1
E201	GLY
Q202	SER
Q203	SER
A204	H1S
V205	H1S
A206	H1S
A207	H1S
T208	H1S
L209	SER
A210	SER
A211	SER
M212	H1S
K213	ALA
R214	LEU
R215	GLU
R216	VAL
S217	LEU
PRO	PHE
PRO	GLN
PRO	GLY
PRO	PRO
SER	M1
SER	E39
	R45
	K55
	A141
	S145
	A149
	I152
	N158
	L159
	A162
	S163
	A164
	S165
	S166
	R170
	E179
	A180
	E181
	R182
	D193
	K194
	K195
	F196
	F197

- |      |      |
|------|------|
| E200 | ME1  |
| E201 | GLY  |
| Q202 | SER  |
| Q203 | SER  |
| A204 | HIS  |
| V205 | HIS  |
| E206 | HIS  |
| A207 | HIS  |
| T208 | HIS  |
| A210 | SER  |
| A211 | SER  |
| M212 | ALA  |
| K213 | ALA  |
| A214 | LEU  |
| R215 | GLU  |
| R216 | VAL  |
| S217 | LEU  |
| LEU  | PHE  |
| PRO  | GLN  |
| PRO  | GLY  |
| PRO  | PRO  |
| SER  | M1   |
| SER  | E99  |
|      | R45  |
|      | K55  |
|      | A141 |
|      | S145 |
|      | A149 |
|      | I152 |
|      | N158 |
|      | L159 |
|      | A162 |
|      | S163 |
|      | A164 |
|      | S165 |
|      | S166 |
|      | R170 |
|      | E179 |
|      | A180 |
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|      | R182 |
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|      | K195 |
|      | F196 |
|      | F197 |

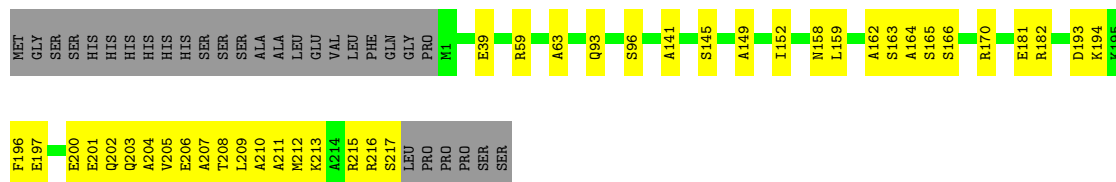
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|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|-----|-----|-----|-----|-----|-----|-----|----|-----|-----|-----|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|
| Q202 | Q203 | A204 | V205 | E206 | A207 | T208 | L209 | A210 | A211 | M212 | K213 | A214 | R215 | T216 | S217 | LEU | PRO | PRO | PRO | SER | SER |     |    |     |     |     |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |
| ME1  | GLY  | SER  | SER  | HIS  | HIS  | HIS  | HIS  | HIS  | HIS  | SER  | SER  | ALA  | ALA  | ALA  | LEU  | GLU | VAL | LEU | PHE | GLN | GLY | PRO | M1 | E39 | R45 | K55 | A141 | S145 | A149 | I152 | N158 | L159 | A162 | S163 | A164 | S165 | S166 | R170 | R182 | D193 | K194 | K195 | F196 | E197 | E200 | E204 |

- |      |      |      |      |      |      |      |      |      |      |      |      |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |    |     |     |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |
|------|------|------|------|------|------|------|------|------|------|------|------|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|----|-----|-----|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|
| E206 | E207 | T208 | L209 | A210 | A211 | M212 | K213 | A214 | R215 | R216 | S217 | LEU | PRO | PRO | PRO | PRO | SER | SER | SER | SER | HIS | HIS | HIS | HIS | HIS | HIS | SER | SER | ALA | ALA | LEU | GLU | VAL | LEU | PHE | GLN | GLY | PRO | M1 | R45 | K55 | A141 | S145 | I152 | N158 | L159 | A162 | S163 | A164 | S165 | S166 | R170 | R182 | D193 | K194 | K195 | F196 | E197 | E200 | E201 | Q202 | Q203 | A204 | Y205 |
|------|------|------|------|------|------|------|------|------|------|------|------|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|----|-----|-----|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|

- |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |     |     |     |     |     |     |     |     |     |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |
|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|
| E200 | E201 | Q202 | Q203 | A204 | V205 | A206 | A207 | T208 | H209 | L209 | A210 | A211 | M212 | K213 | A214 | R215 | R216 | S217 | LEU | PRO | PRO | PRO | PRO | SER | SER |     |     |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |      |
| MET  | GLY  | SER  | SER  | HIS  | HIS  | HIS  | HIS  | HIS  | HIS  | SER  | SER  | ALA  | ALA  | LEU  | GLU  | VAL  | LEU  | PHE  | GLN | GLY | PRO | M1  | E39 | R59 | A63 | Q93 | S96 | A141 | S145 | A149 | N158 | L159 | A162 | S163 | A164 | S165 | S166 | R170 | E181 | D193 | K194 | K196 | F196 | E197 |

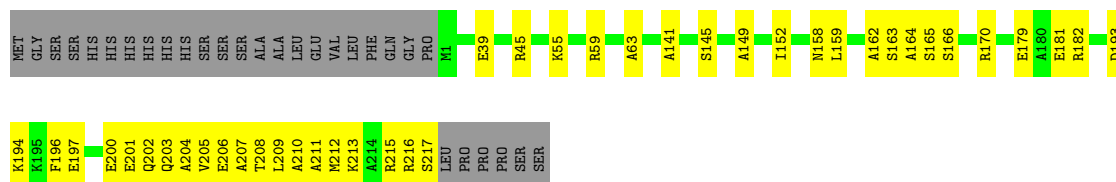
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Chain Q:  72% 16% 12%



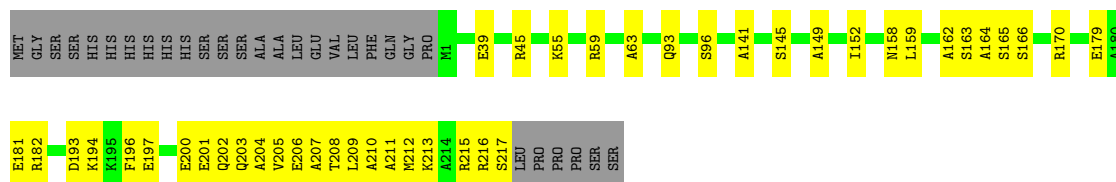
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain R:  72% 17% 12%



- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain S:  71% 17% 12%



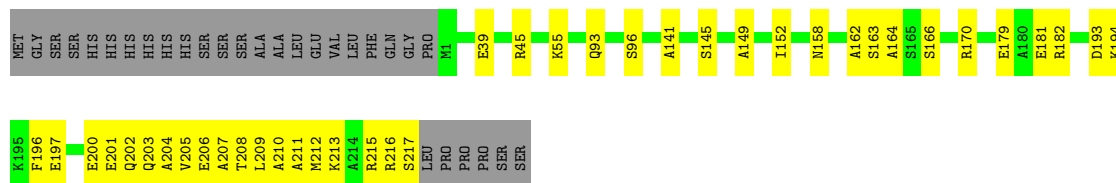
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain T:  74% 15% 12%



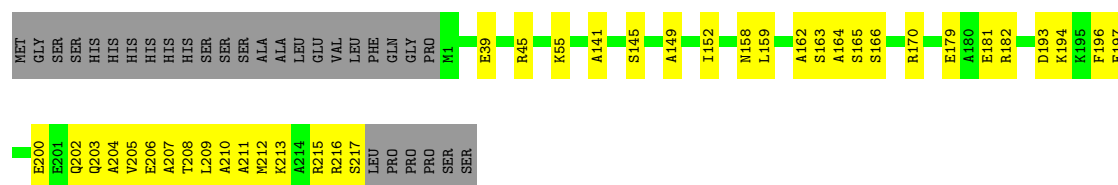
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain U:  72% 16% 12%



- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain V:  73% 15% 12%



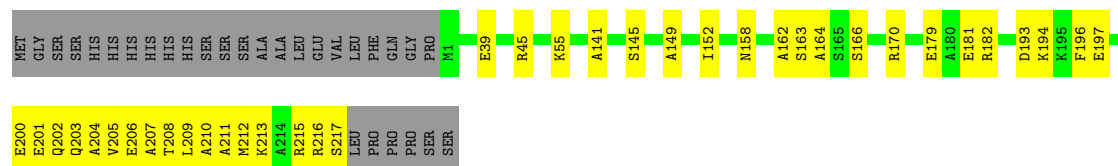
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain W:  74% 15% 12%



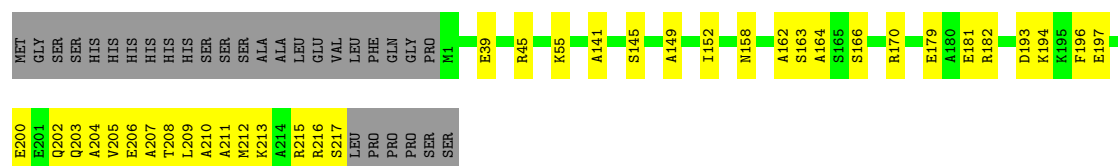
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain X:  73% 15% 12%



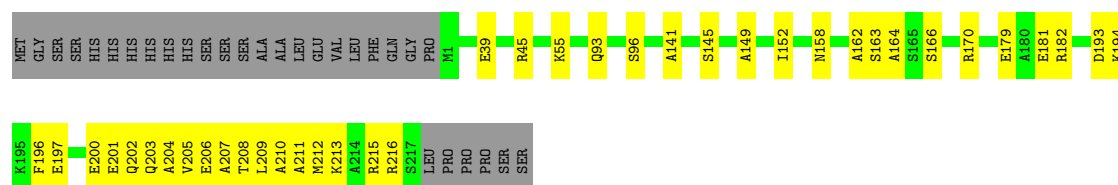
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain Y:  74% 15% 12%



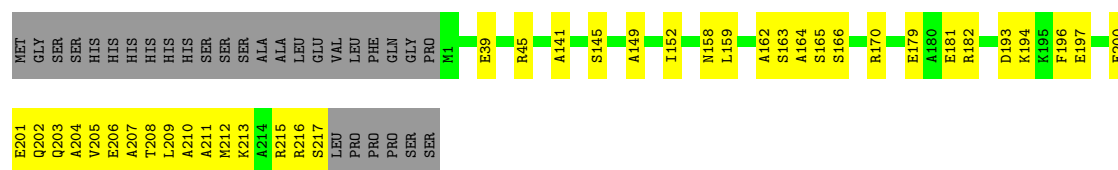
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain Z:  73% 15% 12%



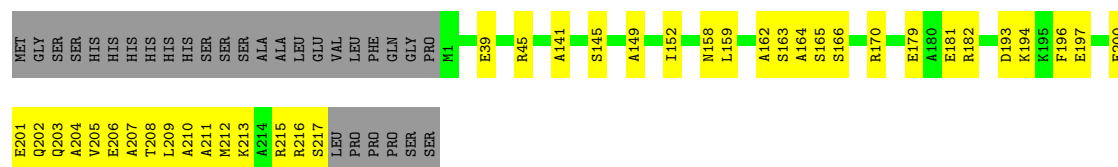
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain a: 



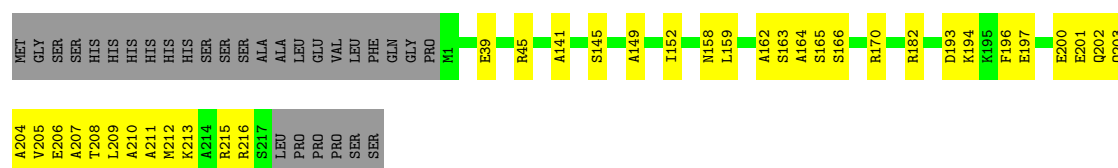
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain b: 




- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain c: 



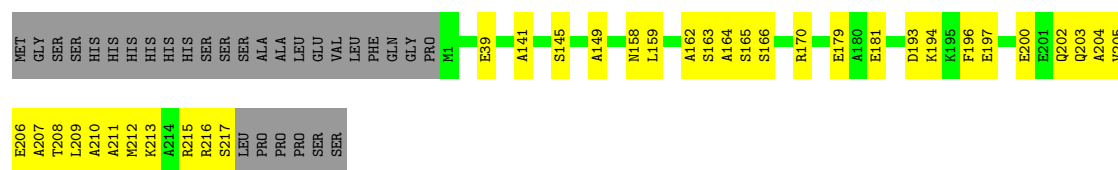
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain d: 



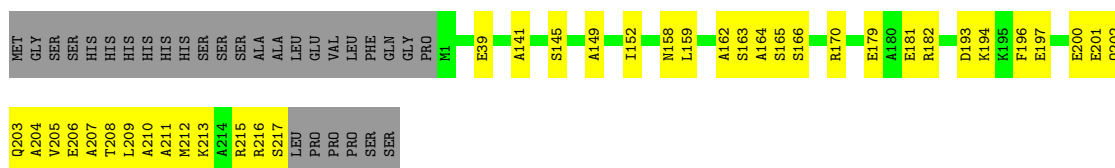
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain e: 

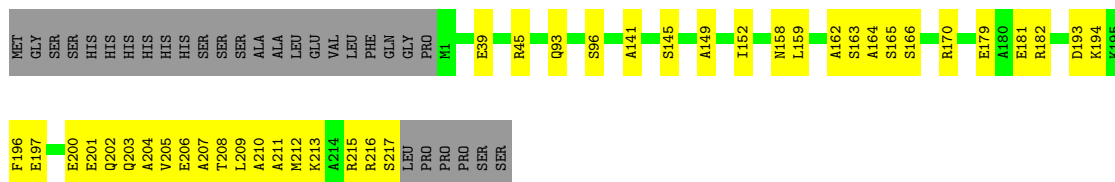


- Molecule 1: Chloroplast membrane-associated 30 kD protein

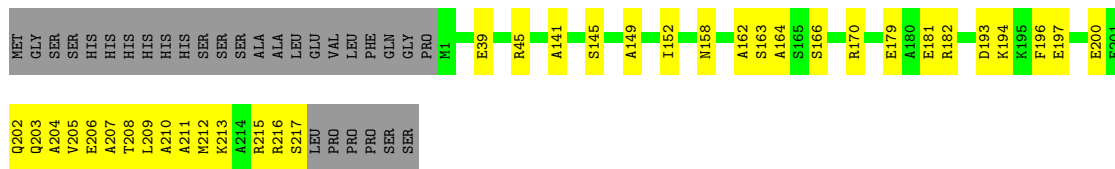
Chain f: 



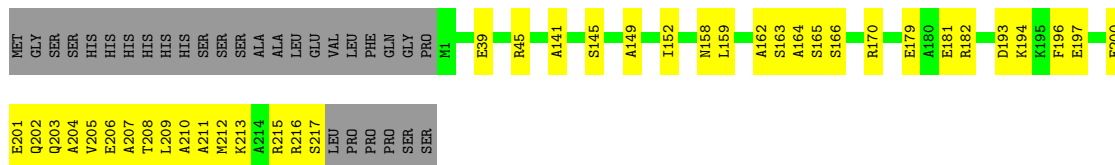
- Molecule 1: Chloroplast membrane-associated 30 kD protein



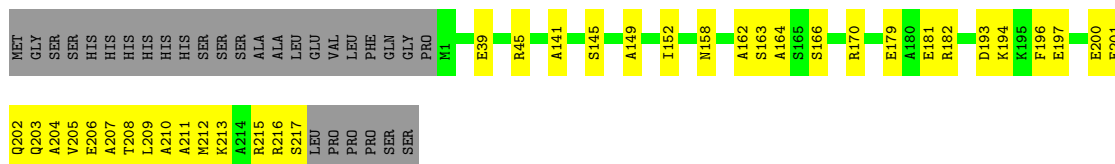
- Molecule 1: Chloroplast membrane-associated 30 kD protein



- Molecule 1: Chloroplast membrane-associated 30 kD protein

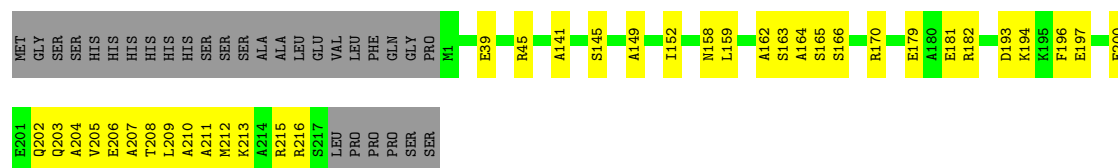


- Molecule 1: Chloroplast membrane-associated 30 kD protein

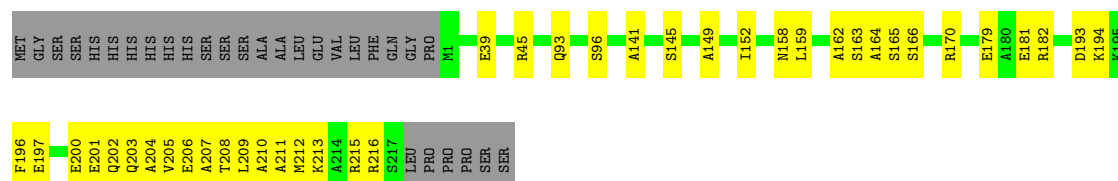


- Molecule 1: Chloroplast membrane-associated 30 kD protein

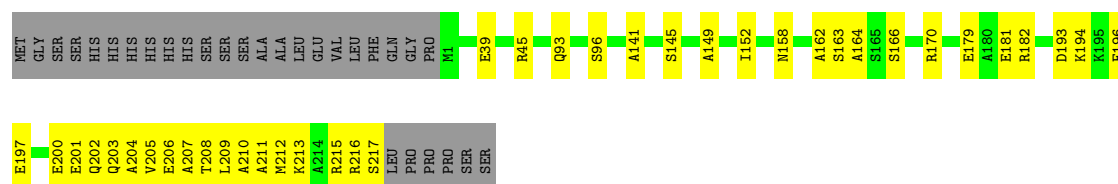




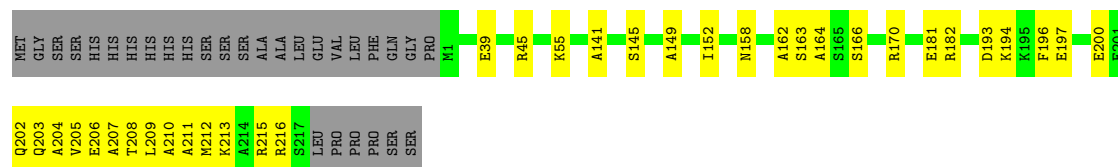
- Molecule 1: Chloroplast membrane-associated 30 kD protein



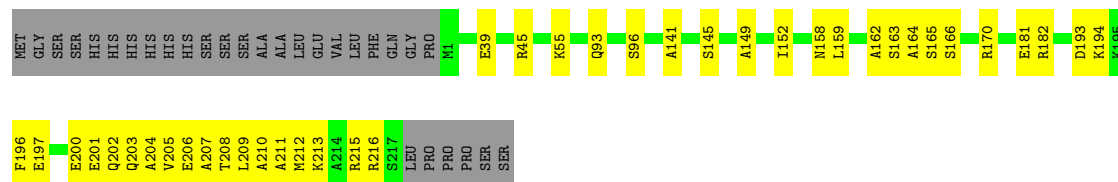
- Molecule 1: Chloroplast membrane-associated 30 kD protein



- Molecule 1: Chloroplast membrane-associated 30 kD protein



- Molecule 1: Chloroplast membrane-associated 30 kD protein



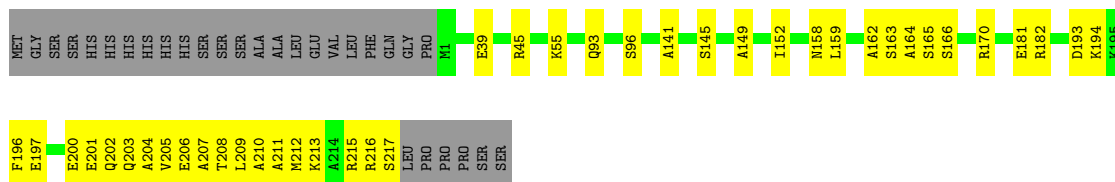
- Molecule 1: Chloroplast membrane-associated 30 kD protein





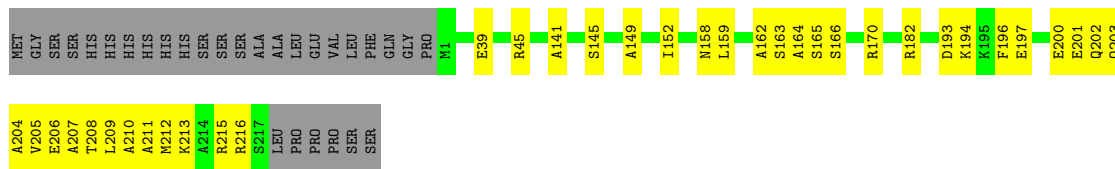
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain q: 72% 16% 12%



- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain r: 74% 14% 12%



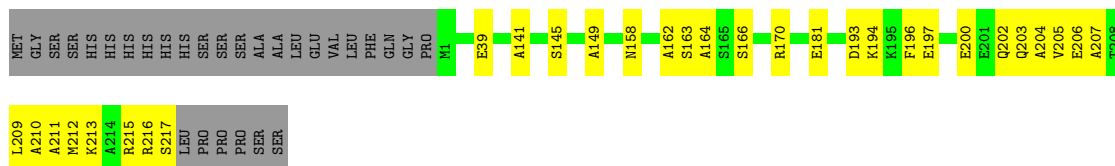
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain s: 74% 14% 12%



- Molecule 1: Chloroplast membrane-associated 30 kD protein

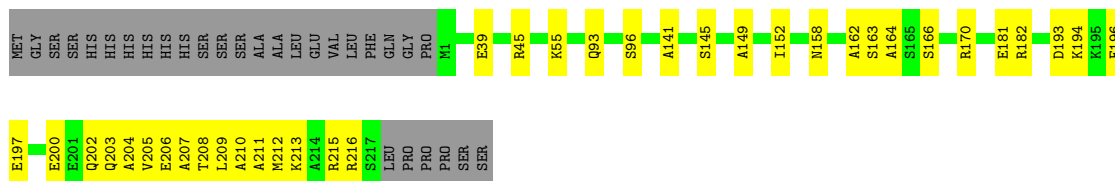
Chain t: 76% 12% 12%



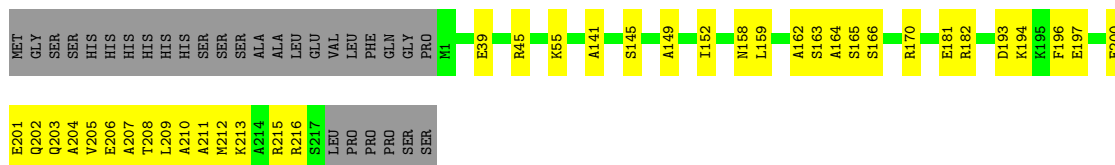
- Molecule 1: Chloroplast membrane-associated 30 kD protein

Chain u: 74% 15% 12%

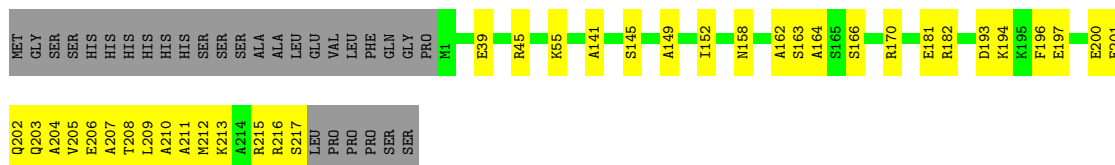
- Molecule 1: Chloroplast membrane-associated 30 kD protein



- Molecule 1: Chloroplast membrane-associated 30 kD protein



- Molecule 1: Chloroplast membrane-associated 30 kD protein



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## 4 Experimental information

Property	Value	Source
EM reconstruction method	HELICAL	Depositor
Imposed symmetry	HELICAL, twist=21.2°, rise=6.85 Å, axial sym=C4	Depositor
Number of segments used	113665	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	FEI TALOS ARCTICA	Depositor
Voltage (kV)	200	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	44	Depositor
Minimum defocus (nm)	1000	Depositor
Maximum defocus (nm)	3500	Depositor
Magnification	Not provided	
Image detector	GATAN K3 BIOQUANTUM (6k x 4k)	Depositor

## 5 Model quality [i](#)

### 5.1 Standard geometry [i](#)

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

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	$\# Z  > 5$	RMSZ	$\# Z  > 5$
1	0	0.08	0/1779	0.19	0/2388
1	1	0.08	0/1779	0.19	0/2388
1	2	0.08	0/1779	0.19	0/2388
1	3	0.08	0/1779	0.19	0/2388
1	4	0.08	0/1779	0.19	0/2388
1	5	0.08	0/1779	0.19	0/2388
1	6	0.08	0/1779	0.19	0/2388
1	7	0.08	0/1779	0.19	0/2388
1	A	0.08	0/1779	0.19	0/2388
1	B	0.08	0/1779	0.19	0/2388
1	C	0.08	0/1779	0.19	0/2388
1	D	0.08	0/1779	0.19	0/2388
1	E	0.08	0/1779	0.19	0/2388
1	F	0.08	0/1779	0.19	0/2388
1	G	0.08	0/1779	0.19	0/2388
1	H	0.08	0/1779	0.19	0/2388
1	I	0.08	0/1779	0.19	0/2388
1	J	0.08	0/1779	0.19	0/2388
1	K	0.08	0/1779	0.19	0/2388
1	L	0.08	0/1779	0.19	0/2388
1	M	0.08	0/1779	0.19	0/2388
1	N	0.08	0/1779	0.19	0/2388
1	O	0.08	0/1779	0.19	0/2388
1	P	0.08	0/1779	0.19	0/2388
1	Q	0.08	0/1779	0.19	0/2388
1	R	0.08	0/1779	0.19	0/2388
1	S	0.08	0/1779	0.19	0/2388
1	T	0.08	0/1779	0.19	0/2388
1	U	0.07	0/1779	0.19	0/2388
1	V	0.07	0/1779	0.19	0/2388
1	W	0.08	0/1779	0.19	0/2388
1	X	0.08	0/1779	0.19	0/2388
1	Y	0.08	0/1779	0.19	0/2388
1	Z	0.08	0/1779	0.19	0/2388

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
1	a	0.08	0/1779	0.19	0/2388
1	b	0.08	0/1779	0.19	0/2388
1	c	0.08	0/1779	0.19	0/2388
1	d	0.08	0/1779	0.19	0/2388
1	e	0.08	0/1779	0.19	0/2388
1	f	0.08	0/1779	0.19	0/2388
1	g	0.08	0/1779	0.19	0/2388
1	h	0.08	0/1779	0.19	0/2388
1	i	0.08	0/1779	0.19	0/2388
1	j	0.08	0/1779	0.19	0/2388
1	k	0.08	0/1779	0.19	0/2388
1	l	0.08	0/1779	0.19	0/2388
1	m	0.08	0/1779	0.19	0/2388
1	n	0.08	0/1779	0.19	0/2388
1	o	0.08	0/1779	0.19	0/2388
1	p	0.08	0/1779	0.19	0/2388
1	q	0.07	0/1779	0.19	0/2388
1	r	0.08	0/1779	0.19	0/2388
1	s	0.08	0/1779	0.19	0/2388
1	t	0.08	0/1779	0.19	0/2388
1	u	0.08	0/1779	0.19	0/2388
1	v	0.08	0/1779	0.19	0/2388
1	w	0.08	0/1779	0.19	0/2388
1	x	0.08	0/1779	0.19	0/2388
1	y	0.07	0/1779	0.19	0/2388
1	z	0.08	0/1779	0.19	0/2388
All	All	0.08	0/106740	0.19	0/143280

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

There are no planarity outliers.

## 5.2 Too-close contacts

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	0	1759	1787	1790	30	0
1	1	1759	1787	1790	31	0
1	2	1759	1787	1790	32	0
1	3	1759	1787	1790	32	0
1	4	1759	1787	1790	32	0
1	5	1759	1787	1790	30	0
1	6	1759	1787	1790	30	0
1	7	1759	1787	1790	27	0
1	A	1759	1787	1790	27	0
1	B	1759	1787	1790	30	0
1	C	1759	1787	1790	32	0
1	D	1759	1787	1790	31	0
1	E	1759	1787	1790	31	0
1	F	1759	1787	1790	32	0
1	G	1759	1787	1790	32	0
1	H	1759	1787	1790	31	0
1	I	1759	1787	1790	32	0
1	J	1759	1787	1790	32	0
1	K	1759	1787	1790	30	0
1	L	1759	1787	1790	32	0
1	M	1759	1787	1790	32	0
1	N	1759	1787	1790	29	0
1	O	1759	1787	1790	29	0
1	P	1759	1787	1790	28	0
1	Q	1759	1787	1790	31	0
1	R	1759	1787	1790	33	0
1	S	1759	1787	1790	34	0
1	T	1759	1787	1790	30	0
1	U	1759	1787	1790	32	0
1	V	1759	1787	1790	31	0
1	W	1759	1787	1790	31	0
1	X	1759	1787	1790	32	0
1	Y	1759	1787	1790	31	0
1	Z	1759	1787	1790	31	0
1	a	1759	1787	1790	31	0
1	b	1759	1787	1790	31	0
1	c	1759	1787	1790	28	0
1	d	1759	1787	1790	26	0
1	e	1759	1787	1790	27	0
1	f	1759	1787	1790	30	0
1	g	1759	1787	1790	32	0
1	h	1759	1787	1790	29	0
1	i	1759	1787	1790	32	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	j	1759	1787	1790	31	0
1	k	1759	1787	1790	30	0
1	l	1759	1787	1790	31	0
1	m	1759	1787	1790	31	0
1	n	1759	1787	1790	28	0
1	o	1759	1787	1790	31	0
1	p	1759	1787	1790	31	0
1	q	1759	1787	1790	32	0
1	r	1759	1787	1790	28	0
1	s	1759	1787	1790	27	0
1	t	1759	1787	1790	25	0
1	u	1759	1787	1790	30	0
1	v	1759	1787	1790	29	0
1	w	1759	1787	1790	30	0
1	x	1759	1787	1790	30	0
1	y	1759	1787	1790	32	0
1	z	1759	1787	1790	31	0
All	All	105540	107220	107400	1626	0

The all-atom clashscore is defined as the number of clashes found per 1000 atoms (including hydrogen atoms). The all-atom clashscore for this structure is 8.

All (1626) close contacts within the same asymmetric unit are listed below, sorted by their clash magnitude.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:V:149:ALA:O	1:W:182:ARG:NH2	2.18	0.77
1:X:206:GLU:O	1:X:210:ALA:N	2.19	0.76
1:h:206:GLU:O	1:h:210:ALA:N	2.19	0.76
1:N:206:GLU:O	1:N:210:ALA:N	2.19	0.76
1:K:206:GLU:O	1:K:210:ALA:N	2.19	0.76
1:u:206:GLU:O	1:u:210:ALA:N	2.19	0.76
1:U:206:GLU:O	1:U:210:ALA:N	2.19	0.76
1:g:206:GLU:O	1:g:210:ALA:N	2.19	0.76
1:k:206:GLU:O	1:k:210:ALA:N	2.19	0.76
1:7:206:GLU:O	1:7:210:ALA:N	2.19	0.76
1:J:206:GLU:O	1:J:210:ALA:N	2.19	0.76
1:W:206:GLU:O	1:W:210:ALA:N	2.19	0.76
1:0:206:GLU:O	1:0:210:ALA:N	2.19	0.76
1:O:206:GLU:O	1:O:210:ALA:N	2.19	0.76
1:Y:206:GLU:O	1:Y:210:ALA:N	2.19	0.76
1:e:206:GLU:O	1:e:210:ALA:N	2.19	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1:206:GLU:O	1:1:210:ALA:N	2.19	0.75
1:M:206:GLU:O	1:M:210:ALA:N	2.19	0.75
1:T:206:GLU:O	1:T:210:ALA:N	2.19	0.75
1:a:206:GLU:O	1:a:210:ALA:N	2.19	0.75
1:i:206:GLU:O	1:i:210:ALA:N	2.19	0.75
1:q:206:GLU:O	1:q:210:ALA:N	2.19	0.75
1:r:206:GLU:O	1:r:210:ALA:N	2.19	0.75
1:t:206:GLU:O	1:t:210:ALA:N	2.19	0.75
1:v:206:GLU:O	1:v:210:ALA:N	2.19	0.75
1:z:206:GLU:O	1:z:210:ALA:N	2.19	0.75
1:5:206:GLU:O	1:5:210:ALA:N	2.19	0.75
1:B:206:GLU:O	1:B:210:ALA:N	2.19	0.75
1:C:206:GLU:O	1:C:210:ALA:N	2.19	0.75
1:L:206:GLU:O	1:L:210:ALA:N	2.19	0.75
1:l:206:GLU:O	1:l:210:ALA:N	2.19	0.75
1:V:206:GLU:O	1:V:210:ALA:N	2.19	0.75
1:j:206:GLU:O	1:j:210:ALA:N	2.19	0.75
1:s:206:GLU:O	1:s:210:ALA:N	2.19	0.75
1:2:206:GLU:O	1:2:210:ALA:N	2.19	0.75
1:D:206:GLU:O	1:D:210:ALA:N	2.19	0.75
1:H:206:GLU:O	1:H:210:ALA:N	2.19	0.75
1:Q:206:GLU:O	1:Q:210:ALA:N	2.19	0.75
1:p:206:GLU:O	1:p:210:ALA:N	2.19	0.75
1:P:206:GLU:O	1:P:210:ALA:N	2.19	0.75
1:x:206:GLU:O	1:x:210:ALA:N	2.19	0.75
1:4:206:GLU:O	1:4:210:ALA:N	2.19	0.75
1:A:206:GLU:O	1:A:210:ALA:N	2.19	0.75
1:Z:206:GLU:O	1:Z:210:ALA:N	2.19	0.75
1:b:206:GLU:O	1:b:210:ALA:N	2.19	0.75
1:y:206:GLU:O	1:y:210:ALA:N	2.19	0.75
1:R:206:GLU:O	1:R:210:ALA:N	2.19	0.75
1:f:206:GLU:O	1:f:210:ALA:N	2.19	0.75
1:d:206:GLU:O	1:d:210:ALA:N	2.19	0.74
1:6:206:GLU:O	1:6:210:ALA:N	2.19	0.74
1:c:206:GLU:O	1:c:210:ALA:N	2.19	0.74
1:n:206:GLU:O	1:n:210:ALA:N	2.19	0.74
1:E:206:GLU:O	1:E:210:ALA:N	2.19	0.74
1:G:206:GLU:O	1:G:210:ALA:N	2.19	0.74
1:3:206:GLU:O	1:3:210:ALA:N	2.19	0.74
1:w:206:GLU:O	1:w:210:ALA:N	2.19	0.74
1:I:206:GLU:O	1:I:210:ALA:N	2.19	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:S:206:GLU:O	1:S:210:ALA:N	2.19	0.74
1:o:206:GLU:O	1:o:210:ALA:N	2.19	0.74
1:m:206:GLU:O	1:m:210:ALA:N	2.19	0.74
1:F:206:GLU:O	1:F:210:ALA:N	2.19	0.74
1:4:149:ALA:O	1:5:182:ARG:NH2	2.22	0.73
1:E:149:ALA:O	1:F:182:ARG:NH2	2.22	0.73
1:F:149:ALA:O	1:G:182:ARG:NH2	2.22	0.73
1:L:149:ALA:O	1:M:182:ARG:NH2	2.22	0.73
1:M:149:ALA:O	1:N:182:ARG:NH2	2.22	0.73
1:R:149:ALA:O	1:S:182:ARG:NH2	2.22	0.73
1:S:149:ALA:O	1:T:182:ARG:NH2	2.22	0.73
1:h:149:ALA:O	1:i:182:ARG:NH2	2.22	0.73
1:i:149:ALA:O	1:j:182:ARG:NH2	2.22	0.73
1:3:149:ALA:O	1:4:182:ARG:NH2	2.22	0.73
1:5:149:ALA:O	1:6:182:ARG:NH2	2.22	0.73
1:D:149:ALA:O	1:E:182:ARG:NH2	2.22	0.73
1:N:149:ALA:O	1:O:182:ARG:NH2	2.22	0.73
1:g:149:ALA:O	1:h:182:ARG:NH2	2.22	0.73
1:p:149:ALA:O	1:q:182:ARG:NH2	2.22	0.73
1:q:149:ALA:O	1:r:182:ARG:NH2	2.22	0.73
1:r:149:ALA:O	1:s:182:ARG:NH2	2.22	0.73
1:C:149:ALA:O	1:D:182:ARG:NH2	2.22	0.72
1:G:149:ALA:O	1:H:182:ARG:NH2	2.22	0.72
1:K:149:ALA:O	1:L:182:ARG:NH2	2.22	0.72
1:Q:149:ALA:O	1:R:182:ARG:NH2	2.22	0.72
1:X:149:ALA:O	1:Y:182:ARG:NH2	2.22	0.72
1:l:149:ALA:O	1:m:182:ARG:NH2	2.22	0.72
1:m:149:ALA:O	1:n:182:ARG:NH2	2.22	0.72
1:t:149:ALA:O	1:u:182:ARG:NH2	2.22	0.72
1:A:149:ALA:O	1:B:182:ARG:NH2	2.22	0.72
1:B:149:ALA:O	1:C:182:ARG:NH2	2.22	0.72
1:F:193:ASP:O	1:F:197:GLU:N	2.23	0.72
1:P:193:ASP:O	1:P:197:GLU:N	2.23	0.72
1:T:149:ALA:O	1:U:182:ARG:NH2	2.22	0.72
1:W:149:ALA:O	1:X:182:ARG:NH2	2.22	0.72
1:Y:149:ALA:O	1:Z:182:ARG:NH2	2.22	0.72
1:f:149:ALA:O	1:g:182:ARG:NH2	2.22	0.72
1:k:149:ALA:O	1:l:182:ARG:NH2	2.22	0.72
1:n:149:ALA:O	1:o:182:ARG:NH2	2.22	0.72
1:o:149:ALA:O	1:p:182:ARG:NH2	2.22	0.72
1:q:193:ASP:O	1:q:197:GLU:N	2.22	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:u:149:ALA:O	1:v:182:ARG:NH2	2.22	0.72
1:3:193:ASP:O	1:3:197:GLU:N	2.23	0.72
1:H:149:ALA:O	1:I:182:ARG:NH2	2.22	0.72
1:j:149:ALA:O	1:k:182:ARG:NH2	2.22	0.72
1:o:193:ASP:O	1:o:197:GLU:N	2.23	0.72
1:y:193:ASP:O	1:y:197:GLU:N	2.22	0.72
1:0:193:ASP:O	1:0:197:GLU:N	2.22	0.72
1:2:149:ALA:O	1:3:182:ARG:NH2	2.22	0.72
1:6:149:ALA:O	1:7:182:ARG:NH2	2.22	0.72
1:J:149:ALA:O	1:K:182:ARG:NH2	2.22	0.72
1:Z:149:ALA:O	1:a:182:ARG:NH2	2.22	0.72
1:v:149:ALA:O	1:w:182:ARG:NH2	2.22	0.72
1:A:193:ASP:O	1:A:197:GLU:N	2.23	0.72
1:a:149:ALA:O	1:b:182:ARG:NH2	2.22	0.72
1:e:149:ALA:O	1:f:182:ARG:NH2	2.22	0.72
1:C:193:ASP:O	1:C:197:GLU:N	2.23	0.72
1:w:149:ALA:O	1:x:182:ARG:NH2	2.22	0.72
1:I:149:ALA:O	1:J:182:ARG:NH2	2.22	0.72
1:P:149:ALA:O	1:Q:182:ARG:NH2	2.22	0.72
1:S:193:ASP:O	1:S:197:GLU:N	2.23	0.72
1:b:193:ASP:O	1:b:197:GLU:N	2.23	0.72
1:b:149:ALA:O	1:c:182:ARG:NH2	2.22	0.72
1:H:193:ASP:O	1:H:197:GLU:N	2.22	0.72
1:I:193:ASP:O	1:I:197:GLU:N	2.23	0.72
1:R:193:ASP:O	1:R:197:GLU:N	2.23	0.72
1:d:193:ASP:O	1:d:197:GLU:N	2.23	0.72
1:l:193:ASP:O	1:l:197:GLU:N	2.22	0.72
1:s:193:ASP:O	1:s:197:GLU:N	2.23	0.72
1:x:149:ALA:O	1:y:182:ARG:NH2	2.22	0.72
1:1:149:ALA:O	1:2:182:ARG:NH2	2.22	0.71
1:2:193:ASP:O	1:2:197:GLU:N	2.23	0.71
1:5:193:ASP:O	1:5:197:GLU:N	2.23	0.71
1:n:193:ASP:O	1:n:197:GLU:N	2.23	0.71
1:6:193:ASP:O	1:6:197:GLU:N	2.23	0.71
1:c:149:ALA:O	1:d:182:ARG:NH2	2.22	0.71
1:y:149:ALA:O	1:z:182:ARG:NH2	2.22	0.71
1:E:193:ASP:O	1:E:197:GLU:N	2.23	0.71
1:e:193:ASP:O	1:e:197:GLU:N	2.23	0.71
1:v:193:ASP:O	1:v:197:GLU:N	2.23	0.71
1:0:149:ALA:O	1:1:182:ARG:NH2	2.22	0.71
1:0:182:ARG:NH2	1:z:149:ALA:O	2.22	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:x:193:ASP:O	1:x:197:GLU:N	2.23	0.71
1:O:193:ASP:O	1:O:197:GLU:N	2.22	0.71
1:V:193:ASP:O	1:V:197:GLU:N	2.22	0.71
1:f:193:ASP:O	1:f:197:GLU:N	2.23	0.71
1:U:193:ASP:O	1:U:197:GLU:N	2.23	0.71
1:L:193:ASP:O	1:L:197:GLU:N	2.23	0.71
1:Y:193:ASP:O	1:Y:197:GLU:N	2.23	0.71
1:K:193:ASP:O	1:K:197:GLU:N	2.23	0.71
1:X:193:ASP:O	1:X:197:GLU:N	2.23	0.71
1:i:193:ASP:O	1:i:197:GLU:N	2.23	0.71
1:a:193:ASP:O	1:a:197:GLU:N	2.23	0.70
1:p:193:ASP:O	1:p:197:GLU:N	2.23	0.70
1:Z:193:ASP:O	1:Z:197:GLU:N	2.23	0.70
1:k:193:ASP:O	1:k:197:GLU:N	2.22	0.70
1:U:149:ALA:O	1:V:182:ARG:NH2	2.24	0.70
1:j:193:ASP:O	1:j:197:GLU:N	2.23	0.70
1:u:193:ASP:O	1:u:197:GLU:N	2.23	0.70
1:z:193:ASP:O	1:z:197:GLU:N	2.22	0.70
1:h:193:ASP:O	1:h:197:GLU:N	2.23	0.70
1:J:193:ASP:O	1:J:197:GLU:N	2.23	0.70
1:N:193:ASP:O	1:N:197:GLU:N	2.23	0.70
1:G:193:ASP:O	1:G:197:GLU:N	2.22	0.70
1:4:193:ASP:O	1:4:197:GLU:N	2.23	0.70
1:2:166:SER:O	1:2:170:ARG:N	2.25	0.70
1:B:193:ASP:O	1:B:197:GLU:N	2.23	0.70
1:P:166:SER:O	1:P:170:ARG:N	2.25	0.70
1:Q:166:SER:O	1:Q:170:ARG:N	2.25	0.70
1:1:166:SER:O	1:1:170:ARG:N	2.25	0.69
1:Q:193:ASP:O	1:Q:197:GLU:N	2.23	0.69
1:T:193:ASP:O	1:T:197:GLU:N	2.23	0.69
1:t:193:ASP:O	1:t:197:GLU:N	2.23	0.69
1:3:166:SER:O	1:3:170:ARG:N	2.25	0.69
1:C:166:SER:O	1:C:170:ARG:N	2.25	0.69
1:H:166:SER:O	1:H:170:ARG:N	2.25	0.69
1:R:166:SER:O	1:R:170:ARG:N	2.25	0.69
1:p:166:SER:O	1:p:170:ARG:N	2.25	0.69
1:D:166:SER:O	1:D:170:ARG:N	2.25	0.69
1:Y:166:SER:O	1:Y:170:ARG:N	2.25	0.69
1:o:166:SER:O	1:o:170:ARG:N	2.25	0.69
1:t:166:SER:O	1:t:170:ARG:N	2.25	0.69
1:X:166:SER:O	1:X:170:ARG:N	2.25	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:c:193:ASP:O	1:c:197:GLU:N	2.23	0.69
1:r:193:ASP:O	1:r:197:GLU:N	2.23	0.69
1:u:166:SER:O	1:u:170:ARG:N	2.25	0.69
1:4:166:SER:O	1:4:170:ARG:N	2.25	0.69
1:7:193:ASP:O	1:7:197:GLU:N	2.22	0.69
1:E:166:SER:O	1:E:170:ARG:N	2.25	0.69
1:Z:166:SER:O	1:Z:170:ARG:N	2.25	0.69
1:m:193:ASP:O	1:m:197:GLU:N	2.23	0.69
1:q:166:SER:O	1:q:170:ARG:N	2.25	0.69
1:1:193:ASP:O	1:1:197:GLU:N	2.22	0.69
1:W:166:SER:O	1:W:170:ARG:N	2.25	0.69
1:v:166:SER:O	1:v:170:ARG:N	2.25	0.69
1:S:166:SER:O	1:S:170:ARG:N	2.25	0.69
1:M:193:ASP:O	1:M:197:GLU:N	2.23	0.68
1:a:166:SER:O	1:a:170:ARG:N	2.25	0.68
1:g:193:ASP:O	1:g:197:GLU:N	2.23	0.68
1:F:213:LYS:O	1:F:217:SER:OG	2.07	0.68
1:w:193:ASP:O	1:w:197:GLU:N	2.23	0.68
1:I:212:MET:O	1:I:216:ARG:N	2.27	0.68
1:O:166:SER:O	1:O:170:ARG:N	2.25	0.68
1:V:166:SER:O	1:V:170:ARG:N	2.25	0.68
1:l:166:SER:O	1:l:170:ARG:N	2.25	0.68
1:4:212:MET:O	1:4:216:ARG:N	2.27	0.68
1:A:212:MET:O	1:A:216:ARG:N	2.27	0.68
1:W:193:ASP:O	1:W:197:GLU:N	2.22	0.68
1:c:212:MET:O	1:c:216:ARG:N	2.27	0.68
1:k:166:SER:O	1:k:170:ARG:N	2.25	0.68
1:o:212:MET:O	1:o:216:ARG:N	2.27	0.68
1:r:166:SER:O	1:r:170:ARG:N	2.25	0.68
1:w:166:SER:O	1:w:170:ARG:N	2.25	0.68
1:x:166:SER:O	1:x:170:ARG:N	2.25	0.68
1:0:212:MET:O	1:0:216:ARG:N	2.27	0.68
1:D:193:ASP:O	1:D:197:GLU:N	2.23	0.68
1:L:166:SER:O	1:L:170:ARG:N	2.25	0.68
1:T:212:MET:O	1:T:216:ARG:N	2.27	0.68
1:c:166:SER:O	1:c:170:ARG:N	2.25	0.68
1:f:212:MET:O	1:f:216:ARG:N	2.27	0.68
1:g:166:SER:O	1:g:170:ARG:N	2.25	0.68
1:h:166:SER:O	1:h:170:ARG:N	2.25	0.68
1:m:166:SER:O	1:m:170:ARG:N	2.25	0.68
1:w:212:MET:O	1:w:216:ARG:N	2.27	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:5:212:MET:O	1:5:216:ARG:N	2.27	0.68
1:D:212:MET:O	1:D:216:ARG:N	2.27	0.68
1:E:212:MET:O	1:E:216:ARG:N	2.27	0.68
1:K:166:SER:O	1:K:170:ARG:N	2.25	0.68
1:Q:212:MET:O	1:Q:216:ARG:N	2.27	0.68
1:U:212:MET:O	1:U:216:ARG:N	2.27	0.68
1:d:212:MET:O	1:d:216:ARG:N	2.27	0.68
1:j:166:SER:O	1:j:170:ARG:N	2.25	0.68
1:l:212:MET:O	1:l:216:ARG:N	2.27	0.68
1:s:212:MET:O	1:s:216:ARG:N	2.27	0.68
1:y:166:SER:O	1:y:170:ARG:N	2.25	0.68
1:z:212:MET:O	1:z:216:ARG:N	2.27	0.68
1:5:166:SER:O	1:5:170:ARG:N	2.25	0.68
1:7:166:SER:O	1:7:170:ARG:N	2.25	0.68
1:7:212:MET:O	1:7:216:ARG:N	2.27	0.68
1:A:166:SER:O	1:A:170:ARG:N	2.25	0.68
1:F:166:SER:O	1:F:170:ARG:N	2.25	0.68
1:H:212:MET:O	1:H:216:ARG:N	2.27	0.68
1:L:212:MET:O	1:L:216:ARG:N	2.27	0.68
1:M:166:SER:O	1:M:170:ARG:N	2.25	0.68
1:P:212:MET:O	1:P:216:ARG:N	2.27	0.68
1:Z:212:MET:O	1:Z:216:ARG:N	2.27	0.68
1:f:166:SER:O	1:f:170:ARG:N	2.25	0.68
1:N:166:SER:O	1:N:170:ARG:N	2.25	0.68
1:O:212:MET:O	1:O:216:ARG:N	2.27	0.68
1:b:166:SER:O	1:b:170:ARG:N	2.25	0.68
1:i:166:SER:O	1:i:170:ARG:N	2.25	0.68
1:i:212:MET:O	1:i:216:ARG:N	2.27	0.68
1:n:212:MET:O	1:n:216:ARG:N	2.27	0.68
1:x:212:MET:O	1:x:216:ARG:N	2.27	0.68
1:J:166:SER:O	1:J:170:ARG:N	2.25	0.68
1:W:212:MET:O	1:W:216:ARG:N	2.27	0.68
1:d:166:SER:O	1:d:170:ARG:N	2.25	0.68
1:p:212:MET:O	1:p:216:ARG:N	2.27	0.68
1:t:212:MET:O	1:t:216:ARG:N	2.27	0.68
1:1:212:MET:O	1:1:216:ARG:N	2.27	0.68
1:3:212:MET:O	1:3:216:ARG:N	2.27	0.68
1:F:212:MET:O	1:F:216:ARG:N	2.27	0.68
1:R:212:MET:O	1:R:216:ARG:N	2.27	0.68
1:k:212:MET:O	1:k:216:ARG:N	2.27	0.68
1:n:166:SER:O	1:n:170:ARG:N	2.25	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:U:166:SER:O	1:U:170:ARG:N	2.25	0.67
1:X:212:MET:O	1:X:216:ARG:N	2.27	0.67
1:a:212:MET:O	1:a:216:ARG:N	2.27	0.67
1:e:166:SER:O	1:e:170:ARG:N	2.25	0.67
1:B:212:MET:O	1:B:216:ARG:N	2.27	0.67
1:J:212:MET:O	1:J:216:ARG:N	2.27	0.67
1:u:212:MET:O	1:u:216:ARG:N	2.27	0.67
1:z:166:SER:O	1:z:170:ARG:N	2.25	0.67
1:B:166:SER:O	1:B:170:ARG:N	2.25	0.67
1:N:212:MET:O	1:N:216:ARG:N	2.27	0.67
1:6:166:SER:O	1:6:170:ARG:N	2.25	0.67
1:I:166:SER:O	1:I:170:ARG:N	2.25	0.67
1:T:166:SER:O	1:T:170:ARG:N	2.25	0.67
1:g:212:MET:O	1:g:216:ARG:N	2.27	0.67
1:s:166:SER:O	1:s:170:ARG:N	2.25	0.67
1:i:213:LYS:O	1:i:217:SER:OG	2.07	0.67
1:R:213:LYS:O	1:R:217:SER:OG	2.07	0.67
1:m:212:MET:O	1:m:216:ARG:N	2.27	0.67
1:t:213:LYS:O	1:t:217:SER:OG	2.07	0.67
1:M:212:MET:O	1:M:216:ARG:N	2.27	0.67
1:q:212:MET:O	1:q:216:ARG:N	2.27	0.67
1:2:212:MET:O	1:2:216:ARG:N	2.27	0.67
1:G:166:SER:O	1:G:170:ARG:N	2.25	0.67
1:G:212:MET:O	1:G:216:ARG:N	2.27	0.67
1:G:213:LYS:O	1:G:217:SER:OG	2.07	0.67
1:j:212:MET:O	1:j:216:ARG:N	2.27	0.67
1:6:212:MET:O	1:6:216:ARG:N	2.27	0.66
1:X:213:LYS:O	1:X:217:SER:OG	2.07	0.66
1:0:166:SER:O	1:0:170:ARG:N	2.25	0.66
1:4:213:LYS:O	1:4:217:SER:OG	2.07	0.66
1:L:213:LYS:O	1:L:217:SER:OG	2.07	0.66
1:S:212:MET:O	1:S:216:ARG:N	2.27	0.66
1:y:212:MET:O	1:y:216:ARG:N	2.27	0.66
1:V:212:MET:O	1:V:216:ARG:N	2.27	0.66
1:K:213:LYS:O	1:K:217:SER:OG	2.07	0.66
1:M:205:VAL:O	1:M:209:LEU:N	2.29	0.66
1:C:212:MET:O	1:C:216:ARG:N	2.27	0.66
1:0:205:VAL:O	1:0:209:LEU:N	2.29	0.66
1:E:205:VAL:O	1:E:209:LEU:N	2.29	0.66
1:I:213:LYS:O	1:I:217:SER:OG	2.07	0.66
1:b:212:MET:O	1:b:216:ARG:N	2.27	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:b:213:LYS:O	1:b:217:SER:OG	2.07	0.66
1:W:213:LYS:O	1:W:217:SER:OG	2.07	0.66
1:Y:213:LYS:O	1:Y:217:SER:OG	2.07	0.66
1:i:205:VAL:O	1:i:209:LEU:N	2.29	0.66
1:3:213:LYS:O	1:3:217:SER:OG	2.07	0.65
1:M:213:LYS:O	1:M:217:SER:OG	2.07	0.65
1:Q:205:VAL:O	1:Q:209:LEU:N	2.29	0.65
1:r:205:VAL:O	1:r:209:LEU:N	2.29	0.65
1:3:205:VAL:O	1:3:209:LEU:N	2.29	0.65
1:C:205:VAL:O	1:C:209:LEU:N	2.29	0.65
1:Y:212:MET:O	1:Y:216:ARG:N	2.27	0.65
1:r:212:MET:O	1:r:216:ARG:N	2.27	0.65
1:v:212:MET:O	1:v:216:ARG:N	2.27	0.65
1:T:213:LYS:O	1:T:217:SER:OG	2.07	0.65
1:e:212:MET:O	1:e:216:ARG:N	2.27	0.65
1:o:205:VAL:O	1:o:209:LEU:N	2.29	0.65
1:q:205:VAL:O	1:q:209:LEU:N	2.29	0.65
1:U:213:LYS:O	1:U:217:SER:OG	2.07	0.65
1:F:205:VAL:O	1:F:209:LEU:N	2.29	0.65
1:1:205:VAL:O	1:1:209:LEU:N	2.29	0.65
1:2:205:VAL:O	1:2:209:LEU:N	2.29	0.65
1:7:213:LYS:O	1:7:217:SER:OG	2.07	0.65
1:H:205:VAL:O	1:H:209:LEU:N	2.29	0.65
1:H:213:LYS:O	1:H:217:SER:OG	2.07	0.65
1:K:212:MET:O	1:K:216:ARG:N	2.27	0.65
1:N:205:VAL:O	1:N:209:LEU:N	2.29	0.65
1:h:213:LYS:O	1:h:217:SER:OG	2.07	0.65
1:R:205:VAL:O	1:R:209:LEU:N	2.29	0.65
1:B:205:VAL:O	1:B:209:LEU:N	2.29	0.65
1:p:205:VAL:O	1:p:209:LEU:N	2.29	0.65
1:y:205:VAL:O	1:y:209:LEU:N	2.29	0.65
1:G:205:VAL:O	1:G:209:LEU:N	2.29	0.65
1:c:205:VAL:O	1:c:209:LEU:N	2.29	0.65
1:5:205:VAL:O	1:5:209:LEU:N	2.29	0.64
1:J:213:LYS:O	1:J:217:SER:OG	2.07	0.64
1:g:213:LYS:O	1:g:217:SER:OG	2.07	0.64
1:h:212:MET:O	1:h:216:ARG:N	2.27	0.64
1:D:205:VAL:O	1:D:209:LEU:N	2.29	0.64
1:W:205:VAL:O	1:W:209:LEU:N	2.29	0.64
1:S:213:LYS:O	1:S:217:SER:OG	2.07	0.64
1:j:205:VAL:O	1:j:209:LEU:N	2.29	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:z:205:VAL:O	1:z:209:LEU:N	2.29	0.64
1:T:205:VAL:O	1:T:209:LEU:N	2.29	0.64
1:d:205:VAL:O	1:d:209:LEU:N	2.29	0.64
1:e:213:LYS:O	1:e:217:SER:OG	2.07	0.64
1:x:205:VAL:O	1:x:209:LEU:N	2.29	0.64
1:P:205:VAL:O	1:P:209:LEU:N	2.29	0.64
1:S:205:VAL:O	1:S:209:LEU:N	2.29	0.64
1:m:205:VAL:O	1:m:209:LEU:N	2.29	0.64
1:s:205:VAL:O	1:s:209:LEU:N	2.29	0.64
1:n:205:VAL:O	1:n:209:LEU:N	2.29	0.64
1:4:205:VAL:O	1:4:209:LEU:N	2.29	0.64
1:A:205:VAL:O	1:A:209:LEU:N	2.29	0.64
1:J:205:VAL:O	1:J:209:LEU:N	2.29	0.64
1:J:193:ASP:OD1	1:J:194:LYS:N	2.31	0.63
1:O:205:VAL:O	1:O:209:LEU:N	2.29	0.63
1:V:213:LYS:O	1:V:217:SER:OG	2.07	0.63
1:a:205:VAL:O	1:a:209:LEU:N	2.29	0.63
1:e:193:ASP:OD1	1:e:194:LYS:N	2.31	0.63
1:f:205:VAL:O	1:f:209:LEU:N	2.29	0.63
1:l:205:VAL:O	1:l:209:LEU:N	2.29	0.63
1:6:205:VAL:O	1:6:209:LEU:N	2.29	0.63
1:I:193:ASP:OD1	1:I:194:LYS:N	2.31	0.63
1:e:205:VAL:O	1:e:209:LEU:N	2.29	0.63
1:v:205:VAL:O	1:v:209:LEU:N	2.29	0.63
1:6:213:LYS:O	1:6:217:SER:OG	2.07	0.63
1:U:193:ASP:OD1	1:U:194:LYS:N	2.31	0.63
1:c:193:ASP:OD1	1:c:194:LYS:N	2.31	0.63
1:f:193:ASP:OD1	1:f:194:LYS:N	2.31	0.63
1:x:193:ASP:OD1	1:x:194:LYS:N	2.31	0.63
1:7:193:ASP:OD1	1:7:194:LYS:N	2.31	0.63
1:T:193:ASP:OD1	1:T:194:LYS:N	2.31	0.63
1:W:193:ASP:OD1	1:W:194:LYS:N	2.31	0.63
1:b:205:VAL:O	1:b:209:LEU:N	2.29	0.63
1:w:205:VAL:O	1:w:209:LEU:N	2.29	0.63
1:y:193:ASP:OD1	1:y:194:LYS:N	2.31	0.63
1:5:193:ASP:OD1	1:5:194:LYS:N	2.31	0.63
1:6:193:ASP:OD1	1:6:194:LYS:N	2.32	0.63
1:C:193:ASP:OD1	1:C:194:LYS:N	2.31	0.63
1:I:205:VAL:O	1:I:209:LEU:N	2.29	0.63
1:K:193:ASP:OD1	1:K:194:LYS:N	2.31	0.63
1:S:193:ASP:OD1	1:S:194:LYS:N	2.31	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:V:193:ASP:OD1	1:V:194:LYS:N	2.31	0.63
1:V:205:VAL:O	1:V:209:LEU:N	2.29	0.63
1:b:193:ASP:OD1	1:b:194:LYS:N	2.31	0.63
1:d:193:ASP:OD1	1:d:194:LYS:N	2.31	0.63
1:p:193:ASP:OD1	1:p:194:LYS:N	2.31	0.63
1:4:193:ASP:OD1	1:4:194:LYS:N	2.31	0.63
1:w:193:ASP:OD1	1:w:194:LYS:N	2.31	0.63
1:z:193:ASP:OD1	1:z:194:LYS:N	2.32	0.63
1:D:193:ASP:OD1	1:D:194:LYS:N	2.31	0.63
1:R:193:ASP:OD1	1:R:194:LYS:N	2.31	0.63
1:Z:205:VAL:O	1:Z:209:LEU:N	2.29	0.63
1:o:193:ASP:OD1	1:o:194:LYS:N	2.31	0.63
1:0:193:ASP:OD1	1:0:194:LYS:N	2.31	0.63
1:3:193:ASP:OD1	1:3:194:LYS:N	2.31	0.63
1:7:205:VAL:O	1:7:209:LEU:N	2.29	0.63
1:B:193:ASP:OD1	1:B:194:LYS:N	2.31	0.63
1:a:193:ASP:OD1	1:a:194:LYS:N	2.31	0.63
1:J:204:ALA:O	1:J:208:THR:OG1	2.14	0.62
1:W:204:ALA:O	1:W:208:THR:OG1	2.15	0.62
1:X:193:ASP:OD1	1:X:194:LYS:N	2.31	0.62
1:X:205:VAL:O	1:X:209:LEU:N	2.29	0.62
1:h:205:VAL:O	1:h:209:LEU:N	2.29	0.62
1:k:205:VAL:O	1:k:209:LEU:N	2.29	0.62
1:q:193:ASP:OD1	1:q:194:LYS:N	2.31	0.62
1:v:193:ASP:OD1	1:v:194:LYS:N	2.31	0.62
1:1:193:ASP:OD1	1:1:194:LYS:N	2.31	0.62
1:H:193:ASP:OD1	1:H:194:LYS:N	2.31	0.62
1:K:205:VAL:O	1:K:209:LEU:N	2.29	0.62
1:L:193:ASP:OD1	1:L:194:LYS:N	2.31	0.62
1:Q:193:ASP:OD1	1:Q:194:LYS:N	2.31	0.62
1:g:193:ASP:OD1	1:g:194:LYS:N	2.31	0.62
1:k:193:ASP:OD1	1:k:194:LYS:N	2.32	0.62
1:l:193:ASP:OD1	1:l:194:LYS:N	2.31	0.62
1:n:193:ASP:OD1	1:n:194:LYS:N	2.31	0.62
1:t:205:VAL:O	1:t:209:LEU:N	2.29	0.62
1:L:205:VAL:O	1:L:209:LEU:N	2.29	0.62
1:e:204:ALA:O	1:e:208:THR:OG1	2.15	0.62
1:t:193:ASP:OD1	1:t:194:LYS:N	2.31	0.62
1:A:193:ASP:OD1	1:A:194:LYS:N	2.31	0.62
1:M:193:ASP:OD1	1:M:194:LYS:N	2.31	0.62
1:O:193:ASP:OD1	1:O:194:LYS:N	2.31	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Z:193:ASP:OD1	1:Z:194:LYS:N	2.31	0.62
1:h:193:ASP:OD1	1:h:194:LYS:N	2.31	0.62
1:m:193:ASP:OD1	1:m:194:LYS:N	2.31	0.62
1:P:193:ASP:OD1	1:P:194:LYS:N	2.31	0.62
1:Y:193:ASP:OD1	1:Y:194:LYS:N	2.31	0.62
1:i:204:ALA:O	1:i:208:THR:OG1	2.15	0.62
1:u:193:ASP:OD1	1:u:194:LYS:N	2.31	0.62
1:2:193:ASP:OD1	1:2:194:LYS:N	2.31	0.62
1:E:193:ASP:OD1	1:E:194:LYS:N	2.31	0.62
1:G:193:ASP:OD1	1:G:194:LYS:N	2.32	0.62
1:N:193:ASP:OD1	1:N:194:LYS:N	2.31	0.62
1:U:205:VAL:O	1:U:209:LEU:N	2.29	0.62
1:Y:205:VAL:O	1:Y:209:LEU:N	2.29	0.62
1:i:193:ASP:OD1	1:i:194:LYS:N	2.31	0.62
1:j:193:ASP:OD1	1:j:194:LYS:N	2.31	0.62
1:s:193:ASP:OD1	1:s:194:LYS:N	2.31	0.62
1:w:163:SER:HG	1:w:164:ALA:H	1.48	0.62
1:u:205:VAL:O	1:u:209:LEU:N	2.29	0.62
1:F:193:ASP:OD1	1:F:194:LYS:N	2.31	0.62
1:g:205:VAL:O	1:g:209:LEU:N	2.29	0.61
1:r:193:ASP:OD1	1:r:194:LYS:N	2.31	0.61
1:H:204:ALA:O	1:H:208:THR:OG1	2.15	0.61
1:5:163:SER:HG	1:5:164:ALA:H	1.48	0.60
1:x:213:LYS:O	1:x:217:SER:OG	2.07	0.60
1:k:204:ALA:O	1:k:208:THR:OG1	2.14	0.60
1:U:204:ALA:O	1:U:208:THR:OG1	2.15	0.60
1:h:204:ALA:O	1:h:208:THR:OG1	2.14	0.60
1:Y:204:ALA:O	1:Y:208:THR:OG1	2.14	0.60
1:V:163:SER:HG	1:V:164:ALA:H	1.50	0.59
1:F:163:SER:HG	1:F:164:ALA:H	1.49	0.59
1:M:204:ALA:O	1:M:208:THR:OG1	2.14	0.59
1:p:213:LYS:O	1:p:217:SER:OG	2.07	0.58
1:O:213:LYS:O	1:O:217:SER:OG	2.07	0.58
1:l:204:ALA:O	1:l:208:THR:OG1	2.15	0.58
1:X:158:ASN:O	1:X:164:ALA:N	2.37	0.58
1:Y:158:ASN:O	1:Y:164:ALA:N	2.37	0.58
1:Z:158:ASN:O	1:Z:164:ALA:N	2.37	0.58
1:c:211:ALA:O	1:c:215:ARG:N	2.36	0.58
1:t:158:ASN:O	1:t:164:ALA:N	2.37	0.58
1:u:158:ASN:O	1:u:164:ALA:N	2.37	0.58
1:v:158:ASN:O	1:v:164:ALA:N	2.37	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:v:211:ALA:O	1:v:215:ARG:N	2.36	0.58
1:w:158:ASN:O	1:w:164:ALA:N	2.37	0.58
1:a:158:ASN:O	1:a:164:ALA:N	2.37	0.58
1:b:158:ASN:O	1:b:164:ALA:N	2.37	0.58
1:o:211:ALA:O	1:o:215:ARG:N	2.36	0.58
1:r:211:ALA:O	1:r:215:ARG:N	2.36	0.58
1:c:158:ASN:O	1:c:164:ALA:N	2.37	0.58
1:j:211:ALA:O	1:j:215:ARG:N	2.36	0.58
1:q:204:ALA:O	1:q:208:THR:OG1	2.14	0.58
1:x:158:ASN:O	1:x:164:ALA:N	2.37	0.58
1:V:204:ALA:O	1:V:208:THR:OG1	2.14	0.58
1:0:211:ALA:O	1:0:215:ARG:N	2.36	0.58
1:y:158:ASN:O	1:y:164:ALA:N	2.37	0.58
1:3:211:ALA:O	1:3:215:ARG:N	2.36	0.58
1:6:196:PHE:O	1:6:200:GLU:N	2.29	0.58
1:p:204:ALA:O	1:p:208:THR:OG1	2.14	0.58
1:E:211:ALA:O	1:E:215:ARG:N	2.36	0.58
1:l:211:ALA:O	1:l:215:ARG:N	2.36	0.58
1:C:204:ALA:O	1:C:208:THR:OG1	2.14	0.57
1:X:211:ALA:O	1:X:215:ARG:N	2.36	0.57
1:Z:204:ALA:O	1:Z:208:THR:OG1	2.15	0.57
1:Z:211:ALA:O	1:Z:215:ARG:N	2.36	0.57
1:d:158:ASN:O	1:d:164:ALA:N	2.37	0.57
1:o:204:ALA:O	1:o:208:THR:OG1	2.15	0.57
1:x:211:ALA:O	1:x:215:ARG:N	2.36	0.57
1:f:196:PHE:O	1:f:200:GLU:N	2.29	0.57
1:z:158:ASN:O	1:z:164:ALA:N	2.37	0.57
1:Q:211:ALA:O	1:Q:215:ARG:N	2.36	0.57
1:f:158:ASN:O	1:f:164:ALA:N	2.37	0.57
1:g:158:ASN:O	1:g:164:ALA:N	2.37	0.57
1:B:211:ALA:O	1:B:215:ARG:N	2.36	0.57
1:J:158:ASN:O	1:J:164:ALA:N	2.37	0.57
1:J:211:ALA:O	1:J:215:ARG:N	2.36	0.57
1:K:158:ASN:O	1:K:164:ALA:N	2.37	0.57
1:V:196:PHE:O	1:V:200:GLU:N	2.29	0.57
1:V:211:ALA:O	1:V:215:ARG:N	2.36	0.57
1:e:158:ASN:O	1:e:164:ALA:N	2.37	0.57
1:h:158:ASN:O	1:h:164:ALA:N	2.37	0.57
1:I:158:ASN:O	1:I:164:ALA:N	2.37	0.57
1:L:158:ASN:O	1:L:164:ALA:N	2.37	0.57
1:N:211:ALA:O	1:N:215:ARG:N	2.36	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:158:ASN:O	1:0:164:ALA:N	2.37	0.57
1:H:158:ASN:O	1:H:164:ALA:N	2.37	0.57
1:L:196:PHE:O	1:L:200:GLU:N	2.29	0.57
1:M:158:ASN:O	1:M:164:ALA:N	2.37	0.57
1:4:45:ARG:NH1	1:J:181:GLU:OE2	2.38	0.57
1:5:211:ALA:O	1:5:215:ARG:N	2.36	0.57
1:G:158:ASN:O	1:G:164:ALA:N	2.37	0.57
1:L:211:ALA:O	1:L:215:ARG:N	2.36	0.57
1:d:204:ALA:O	1:d:208:THR:OG1	2.14	0.57
1:i:158:ASN:O	1:i:164:ALA:N	2.37	0.57
1:7:45:ARG:NH1	1:M:181:GLU:OE2	2.38	0.57
1:J:45:ARG:NH1	1:W:181:GLU:OE2	2.38	0.57
1:N:158:ASN:O	1:N:164:ALA:N	2.37	0.57
1:T:45:ARG:NH1	1:g:181:GLU:OE2	2.38	0.57
1:e:211:ALA:O	1:e:215:ARG:N	2.36	0.57
1:g:45:ARG:NH1	1:t:181:GLU:OE2	2.38	0.57
1:j:158:ASN:O	1:j:164:ALA:N	2.37	0.57
1:q:211:ALA:O	1:q:215:ARG:N	2.36	0.57
1:I:45:ARG:NH1	1:V:181:GLU:OE2	2.38	0.57
1:R:196:PHE:O	1:R:200:GLU:N	2.29	0.57
1:W:211:ALA:O	1:W:215:ARG:N	2.36	0.57
1:0:204:ALA:O	1:0:208:THR:OG1	2.15	0.57
1:3:45:ARG:NH1	1:I:181:GLU:OE2	2.38	0.57
1:G:45:ARG:NH1	1:T:181:GLU:OE2	2.38	0.57
1:O:158:ASN:O	1:O:164:ALA:N	2.37	0.57
1:S:45:ARG:NH1	1:f:181:GLU:OE2	2.38	0.57
1:W:45:ARG:NH1	1:j:181:GLU:OE2	2.38	0.57
1:7:211:ALA:O	1:7:215:ARG:N	2.36	0.56
1:b:211:ALA:O	1:b:215:ARG:N	2.36	0.56
1:k:158:ASN:O	1:k:164:ALA:N	2.37	0.56
1:m:204:ALA:O	1:m:208:THR:OG1	2.15	0.56
1:1:158:ASN:O	1:1:164:ALA:N	2.37	0.56
1:2:211:ALA:O	1:2:215:ARG:N	2.36	0.56
1:4:181:GLU:OE2	1:r:45:ARG:NH1	2.38	0.56
1:F:45:ARG:NH1	1:S:181:GLU:OE2	2.38	0.56
1:F:158:ASN:O	1:F:164:ALA:N	2.37	0.56
1:I:211:ALA:O	1:I:215:ARG:N	2.36	0.56
1:s:158:ASN:O	1:s:164:ALA:N	2.37	0.56
1:v:204:ALA:O	1:v:208:THR:OG1	2.14	0.56
1:6:45:ARG:NH1	1:L:181:GLU:OE2	2.38	0.56
1:G:211:ALA:O	1:G:215:ARG:N	2.36	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:45:ARG:NH1	1:X:181:GLU:OE2	2.38	0.56
1:M:45:ARG:NH1	1:Z:181:GLU:OE2	2.38	0.56
1:P:158:ASN:O	1:P:164:ALA:N	2.37	0.56
1:R:211:ALA:O	1:R:215:ARG:N	2.36	0.56
1:U:45:ARG:NH1	1:h:181:GLU:OE2	2.38	0.56
1:l:158:ASN:O	1:l:164:ALA:N	2.37	0.56
1:r:158:ASN:O	1:r:164:ALA:N	2.37	0.56
1:u:211:ALA:O	1:u:215:ARG:N	2.36	0.56
1:1:45:ARG:NH1	1:G:181:GLU:OE2	2.38	0.56
1:2:158:ASN:O	1:2:164:ALA:N	2.37	0.56
1:Q:158:ASN:O	1:Q:164:ALA:N	2.37	0.56
1:R:158:ASN:O	1:R:164:ALA:N	2.37	0.56
1:h:45:ARG:NH1	1:u:181:GLU:OE2	2.38	0.56
1:n:211:ALA:O	1:n:215:ARG:N	2.36	0.56
1:3:158:ASN:O	1:3:164:ALA:N	2.37	0.56
1:3:181:GLU:OE2	1:q:45:ARG:NH1	2.38	0.56
1:5:45:ARG:NH1	1:K:181:GLU:OE2	2.38	0.56
1:A:158:ASN:O	1:A:164:ALA:N	2.37	0.56
1:C:213:LYS:O	1:C:217:SER:OG	2.07	0.56
1:H:163:SER:HG	1:H:164:ALA:H	1.52	0.56
1:P:204:ALA:O	1:P:208:THR:OG1	2.15	0.56
1:S:211:ALA:O	1:S:215:ARG:N	2.36	0.56
1:m:158:ASN:O	1:m:164:ALA:N	2.37	0.56
1:D:181:GLU:OE2	1:y:45:ARG:NH1	2.38	0.56
1:E:158:ASN:O	1:E:164:ALA:N	2.37	0.56
1:H:45:ARG:NH1	1:U:181:GLU:OE2	2.38	0.56
1:H:196:PHE:O	1:H:200:GLU:N	2.29	0.56
1:N:204:ALA:O	1:N:208:THR:OG1	2.14	0.56
1:a:45:ARG:NH1	1:n:181:GLU:OE2	2.38	0.56
1:b:45:ARG:NH1	1:o:181:GLU:OE2	2.38	0.56
1:q:158:ASN:O	1:q:164:ALA:N	2.37	0.56
1:4:158:ASN:O	1:4:164:ALA:N	2.37	0.56
1:D:45:ARG:NH1	1:Q:181:GLU:OE2	2.38	0.56
1:L:45:ARG:NH1	1:Y:181:GLU:OE2	2.38	0.56
1:X:45:ARG:NH1	1:k:181:GLU:OE2	2.38	0.56
1:g:211:ALA:O	1:g:215:ARG:N	2.36	0.56
1:k:196:PHE:O	1:k:200:GLU:N	2.29	0.56
1:C:181:GLU:OE2	1:x:45:ARG:NH1	2.38	0.56
1:E:181:GLU:OE2	1:z:45:ARG:NH1	2.38	0.56
1:Q:213:LYS:O	1:Q:217:SER:OG	2.07	0.56
1:S:158:ASN:O	1:S:164:ALA:N	2.37	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:i:211:ALA:O	1:i:215:ARG:N	2.36	0.56
1:j:45:ARG:NH1	1:w:181:GLU:OE2	2.38	0.56
1:z:211:ALA:O	1:z:215:ARG:N	2.36	0.56
1:0:45:ARG:NH1	1:F:181:GLU:OE2	2.38	0.56
1:D:158:ASN:O	1:D:164:ALA:N	2.37	0.56
1:H:162:ALA:N	1:H:166:SER:OG	2.39	0.56
1:R:45:ARG:NH1	1:e:181:GLU:OE2	2.38	0.56
1:Z:45:ARG:NH1	1:m:181:GLU:OE2	2.38	0.56
1:z:213:LYS:O	1:z:217:SER:OG	2.07	0.56
1:n:158:ASN:O	1:n:164:ALA:N	2.37	0.56
1:1:162:ALA:N	1:1:166:SER:OG	2.39	0.55
1:5:158:ASN:O	1:5:164:ALA:N	2.37	0.55
1:5:181:GLU:OE2	1:s:45:ARG:NH1	2.38	0.55
1:1:181:GLU:OE2	1:o:45:ARG:NH1	2.38	0.55
1:B:158:ASN:O	1:B:164:ALA:N	2.37	0.55
1:B:181:GLU:OE2	1:w:45:ARG:NH1	2.38	0.55
1:D:211:ALA:O	1:D:215:ARG:N	2.36	0.55
1:M:211:ALA:O	1:M:215:ARG:N	2.36	0.55
1:P:162:ALA:N	1:P:166:SER:OG	2.39	0.55
1:U:211:ALA:O	1:U:215:ARG:N	2.36	0.55
1:c:45:ARG:NH1	1:p:181:GLU:OE2	2.38	0.55
1:i:45:ARG:NH1	1:v:181:GLU:OE2	2.38	0.55
1:p:158:ASN:O	1:p:164:ALA:N	2.37	0.55
1:5:196:PHE:O	1:5:200:GLU:N	2.29	0.55
1:C:45:ARG:NH1	1:P:181:GLU:OE2	2.38	0.55
1:O:211:ALA:O	1:O:215:ARG:N	2.36	0.55
1:T:158:ASN:O	1:T:164:ALA:N	2.37	0.55
1:k:45:ARG:NH1	1:x:181:GLU:OE2	2.38	0.55
1:u:196:PHE:O	1:u:200:GLU:N	2.29	0.55
1:2:45:ARG:NH1	1:H:181:GLU:OE2	2.38	0.55
1:2:181:GLU:OE2	1:p:45:ARG:NH1	2.38	0.55
1:3:204:ALA:O	1:3:208:THR:OG1	2.14	0.55
1:Y:45:ARG:NH1	1:l:181:GLU:OE2	2.38	0.55
1:l:45:ARG:NH1	1:y:181:GLU:OE2	2.38	0.55
1:0:181:GLU:OE2	1:n:45:ARG:NH1	2.38	0.55
1:C:158:ASN:O	1:C:164:ALA:N	2.37	0.55
1:E:45:ARG:NH1	1:R:181:GLU:OE2	2.38	0.55
1:H:211:ALA:O	1:H:215:ARG:N	2.36	0.55
1:O:45:ARG:NH1	1:b:181:GLU:OE2	2.38	0.55
1:6:158:ASN:O	1:6:164:ALA:N	2.37	0.55
1:A:181:GLU:OE2	1:v:45:ARG:NH1	2.38	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:P:211:ALA:O	1:P:215:ARG:N	2.36	0.55
1:k:211:ALA:O	1:k:215:ARG:N	2.36	0.55
1:s:211:ALA:O	1:s:215:ARG:N	2.36	0.55
1:2:162:ALA:N	1:2:166:SER:OG	2.39	0.55
1:B:162:ALA:N	1:B:166:SER:OG	2.39	0.55
1:I:162:ALA:N	1:I:166:SER:OG	2.39	0.55
1:R:204:ALA:O	1:R:208:THR:OG1	2.14	0.55
1:d:45:ARG:NH1	1:q:181:GLU:OE2	2.38	0.55
1:e:196:PHE:O	1:e:200:GLU:N	2.29	0.55
1:h:211:ALA:O	1:h:215:ARG:N	2.36	0.55
1:n:204:ALA:O	1:n:208:THR:OG1	2.14	0.55
1:w:211:ALA:O	1:w:215:ARG:N	2.36	0.55
1:4:211:ALA:O	1:4:215:ARG:N	2.36	0.55
1:o:158:ASN:O	1:o:164:ALA:N	2.37	0.55
1:6:211:ALA:O	1:6:215:ARG:N	2.36	0.55
1:O:162:ALA:N	1:O:166:SER:OG	2.39	0.55
1:U:158:ASN:O	1:U:164:ALA:N	2.37	0.55
1:m:45:ARG:NH1	1:z:181:GLU:OE2	2.38	0.55
1:o:162:ALA:N	1:o:166:SER:OG	2.39	0.55
1:N:162:ALA:N	1:N:166:SER:OG	2.39	0.55
1:Q:162:ALA:N	1:Q:166:SER:OG	2.39	0.55
1:Y:211:ALA:O	1:Y:215:ARG:N	2.36	0.55
1:j:162:ALA:N	1:j:166:SER:OG	2.39	0.55
1:k:162:ALA:N	1:k:166:SER:OG	2.39	0.55
1:A:211:ALA:O	1:A:215:ARG:N	2.36	0.54
1:e:162:ALA:N	1:e:166:SER:OG	2.39	0.54
1:i:162:ALA:N	1:i:166:SER:OG	2.39	0.54
1:B:204:ALA:O	1:B:208:THR:OG1	2.14	0.54
1:N:196:PHE:O	1:N:200:GLU:N	2.29	0.54
1:d:211:ALA:O	1:d:215:ARG:N	2.36	0.54
1:y:213:LYS:O	1:y:217:SER:OG	2.07	0.54
1:7:158:ASN:O	1:7:164:ALA:N	2.37	0.54
1:C:162:ALA:N	1:C:166:SER:OG	2.39	0.54
1:M:162:ALA:N	1:M:166:SER:OG	2.39	0.54
1:U:196:PHE:O	1:U:200:GLU:N	2.29	0.54
1:K:211:ALA:O	1:K:215:ARG:N	2.36	0.54
1:N:45:ARG:NH1	1:a:181:GLU:OE2	2.38	0.54
1:l:162:ALA:N	1:l:166:SER:OG	2.39	0.54
1:3:162:ALA:N	1:3:166:SER:OG	2.39	0.54
1:V:45:ARG:NH1	1:i:181:GLU:OE2	2.40	0.54
1:V:158:ASN:O	1:V:164:ALA:N	2.37	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:X:196:PHE:O	1:X:200:GLU:N	2.29	0.54
1:w:204:ALA:O	1:w:208:THR:OG1	2.14	0.54
1:K:196:PHE:O	1:K:200:GLU:N	2.29	0.54
1:W:162:ALA:N	1:W:166:SER:OG	2.39	0.54
1:X:162:ALA:N	1:X:166:SER:OG	2.39	0.54
1:h:196:PHE:O	1:h:200:GLU:N	2.29	0.54
1:t:162:ALA:N	1:t:166:SER:OG	2.39	0.54
1:w:162:ALA:N	1:w:166:SER:OG	2.39	0.54
1:J:162:ALA:N	1:J:166:SER:OG	2.39	0.54
1:m:196:PHE:O	1:m:200:GLU:N	2.29	0.54
1:p:162:ALA:N	1:p:166:SER:OG	2.39	0.54
1:p:211:ALA:O	1:p:215:ARG:N	2.36	0.54
1:R:162:ALA:N	1:R:166:SER:OG	2.39	0.54
1:T:211:ALA:O	1:T:215:ARG:N	2.36	0.54
1:Y:162:ALA:N	1:Y:166:SER:OG	2.39	0.54
1:m:162:ALA:N	1:m:166:SER:OG	2.39	0.54
1:W:158:ASN:O	1:W:164:ALA:N	2.37	0.54
1:y:211:ALA:O	1:y:215:ARG:N	2.36	0.54
1:V:162:ALA:N	1:V:166:SER:OG	2.39	0.53
1:a:211:ALA:O	1:a:215:ARG:N	2.36	0.53
1:b:162:ALA:N	1:b:166:SER:OG	2.39	0.53
1:u:162:ALA:N	1:u:166:SER:OG	2.39	0.53
1:l:211:ALA:O	1:l:215:ARG:N	2.36	0.53
1:D:162:ALA:N	1:D:166:SER:OG	2.39	0.53
1:A:162:ALA:N	1:A:166:SER:OG	2.39	0.53
1:S:204:ALA:O	1:S:208:THR:OG1	2.14	0.53
1:7:162:ALA:N	1:7:166:SER:OG	2.39	0.53
1:Z:162:ALA:N	1:Z:166:SER:OG	2.39	0.53
1:Q:196:PHE:O	1:Q:200:GLU:N	2.29	0.53
1:c:162:ALA:N	1:c:166:SER:OG	2.39	0.53
1:f:162:ALA:N	1:f:166:SER:OG	2.39	0.53
1:x:162:ALA:N	1:x:166:SER:OG	2.39	0.53
1:1:196:PHE:O	1:1:200:GLU:N	2.29	0.53
1:4:162:ALA:N	1:4:166:SER:OG	2.39	0.53
1:6:202:GLN:O	1:6:206:GLU:N	2.36	0.53
1:F:211:ALA:O	1:F:215:ARG:N	2.36	0.53
1:O:204:ALA:O	1:O:208:THR:OG1	2.14	0.53
1:c:159:LEU:O	1:c:165:SER:OG	2.24	0.53
1:r:196:PHE:O	1:r:200:GLU:N	2.29	0.53
1:w:196:PHE:O	1:w:200:GLU:N	2.29	0.53
1:C:211:ALA:O	1:C:215:ARG:N	2.36	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:U:162:ALA:N	1:U:166:SER:OG	2.39	0.53
1:v:162:ALA:N	1:v:166:SER:OG	2.39	0.53
1:y:162:ALA:N	1:y:166:SER:OG	2.39	0.53
1:m:211:ALA:O	1:m:215:ARG:N	2.36	0.53
1:n:162:ALA:N	1:n:166:SER:OG	2.39	0.53
1:D:196:PHE:O	1:D:200:GLU:N	2.29	0.53
1:d:162:ALA:N	1:d:166:SER:OG	2.39	0.53
1:q:162:ALA:N	1:q:166:SER:OG	2.39	0.53
1:q:209:LEU:O	1:q:213:LYS:N	2.36	0.53
1:K:204:ALA:O	1:K:208:THR:OG1	2.15	0.53
1:S:162:ALA:N	1:S:166:SER:OG	2.39	0.53
1:b:204:ALA:O	1:b:208:THR:OG1	2.14	0.53
1:y:209:LEU:O	1:y:213:LYS:N	2.37	0.53
1:6:162:ALA:N	1:6:166:SER:OG	2.39	0.52
1:G:196:PHE:O	1:G:200:GLU:N	2.29	0.52
1:z:200:GLU:O	1:z:204:ALA:N	2.36	0.52
1:5:202:GLN:O	1:5:206:GLU:N	2.36	0.52
1:a:162:ALA:N	1:a:166:SER:OG	2.39	0.52
1:r:209:LEU:O	1:r:213:LYS:N	2.36	0.52
1:t:211:ALA:O	1:t:215:ARG:N	2.36	0.52
1:D:204:ALA:O	1:D:208:THR:OG1	2.14	0.52
1:E:162:ALA:N	1:E:166:SER:OG	2.39	0.52
1:T:162:ALA:N	1:T:166:SER:OG	2.39	0.52
1:V:159:LEU:O	1:V:165:SER:OG	2.24	0.52
1:l:202:GLN:O	1:l:206:GLU:N	2.36	0.52
1:s:202:GLN:O	1:s:206:GLU:N	2.36	0.52
1:w:159:LEU:O	1:w:165:SER:OG	2.24	0.52
1:z:162:ALA:N	1:z:166:SER:OG	2.39	0.52
1:A:200:GLU:O	1:A:204:ALA:N	2.36	0.52
1:M:202:GLN:O	1:M:206:GLU:N	2.36	0.52
1:X:204:ALA:O	1:X:208:THR:OG1	2.15	0.52
1:b:202:GLN:O	1:b:206:GLU:N	2.36	0.52
1:d:196:PHE:O	1:d:200:GLU:N	2.29	0.52
1:s:209:LEU:O	1:s:213:LYS:N	2.37	0.52
1:C:196:PHE:O	1:C:200:GLU:N	2.29	0.52
1:K:162:ALA:N	1:K:166:SER:OG	2.39	0.52
1:L:202:GLN:O	1:L:206:GLU:N	2.36	0.52
1:V:200:GLU:O	1:V:204:ALA:N	2.36	0.52
1:a:209:LEU:O	1:a:213:LYS:N	2.37	0.52
1:v:202:GLN:O	1:v:206:GLU:N	2.36	0.52
1:2:141:ALA:O	1:2:145:SER:N	2.43	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:3:141:ALA:O	1:3:145:SER:N	2.43	0.52
1:4:141:ALA:O	1:4:145:SER:N	2.42	0.52
1:N:202:GLN:O	1:N:206:GLU:N	2.36	0.52
1:P:141:ALA:O	1:P:145:SER:N	2.42	0.52
1:Q:141:ALA:O	1:Q:145:SER:N	2.42	0.52
1:R:141:ALA:O	1:R:145:SER:N	2.42	0.52
1:4:196:PHE:O	1:4:200:GLU:N	2.29	0.52
1:5:162:ALA:N	1:5:166:SER:OG	2.39	0.52
1:1:141:ALA:O	1:1:145:SER:N	2.42	0.52
1:5:141:ALA:O	1:5:145:SER:N	2.42	0.52
1:B:196:PHE:O	1:B:200:GLU:N	2.29	0.52
1:K:202:GLN:O	1:K:206:GLU:N	2.36	0.52
1:S:141:ALA:O	1:S:145:SER:N	2.42	0.52
1:f:211:ALA:O	1:f:215:ARG:N	2.36	0.52
1:0:162:ALA:N	1:0:166:SER:OG	2.39	0.52
1:4:202:GLN:O	1:4:206:GLU:N	2.36	0.52
1:A:202:GLN:O	1:A:206:GLU:N	2.36	0.52
1:c:196:PHE:O	1:c:200:GLU:N	2.29	0.52
1:f:200:GLU:O	1:f:204:ALA:N	2.36	0.52
1:r:200:GLU:O	1:r:204:ALA:N	2.36	0.52
1:F:141:ALA:O	1:F:145:SER:N	2.42	0.52
1:T:141:ALA:O	1:T:145:SER:N	2.43	0.52
1:r:202:GLN:O	1:r:206:GLU:N	2.36	0.52
1:x:204:ALA:O	1:x:208:THR:OG1	2.14	0.52
1:z:209:LEU:O	1:z:213:LYS:N	2.37	0.52
1:0:141:ALA:O	1:0:145:SER:N	2.43	0.51
1:A:209:LEU:O	1:A:213:LYS:N	2.37	0.51
1:G:141:ALA:O	1:G:145:SER:N	2.42	0.51
1:O:202:GLN:O	1:O:206:GLU:N	2.36	0.51
1:a:200:GLU:O	1:a:204:ALA:N	2.36	0.51
1:n:196:PHE:O	1:n:200:GLU:N	2.29	0.51
1:o:196:PHE:O	1:o:200:GLU:N	2.29	0.51
1:s:141:ALA:O	1:s:145:SER:N	2.42	0.51
1:6:141:ALA:O	1:6:145:SER:N	2.42	0.51
1:E:141:ALA:O	1:E:145:SER:N	2.43	0.51
1:E:204:ALA:O	1:E:208:THR:OG1	2.15	0.51
1:r:141:ALA:O	1:r:145:SER:N	2.42	0.51
1:r:162:ALA:N	1:r:166:SER:OG	2.39	0.51
1:3:202:GLN:O	1:3:206:GLU:N	2.36	0.51
1:6:200:GLU:O	1:6:204:ALA:N	2.36	0.51
1:U:141:ALA:O	1:U:145:SER:N	2.42	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:g:162:ALA:N	1:g:166:SER:OG	2.39	0.51
1:D:141:ALA:O	1:D:145:SER:N	2.42	0.51
1:J:202:GLN:O	1:J:206:GLU:N	2.36	0.51
1:L:162:ALA:N	1:L:166:SER:OG	2.39	0.51
1:g:204:ALA:O	1:g:208:THR:OG1	2.14	0.51
1:h:162:ALA:N	1:h:166:SER:OG	2.39	0.51
1:j:213:LYS:O	1:j:217:SER:OG	2.07	0.51
1:q:141:ALA:O	1:q:145:SER:N	2.43	0.51
1:z:141:ALA:O	1:z:145:SER:N	2.42	0.51
1:0:196:PHE:O	1:0:200:GLU:N	2.29	0.51
1:7:141:ALA:O	1:7:145:SER:N	2.42	0.51
1:H:141:ALA:O	1:H:145:SER:N	2.43	0.51
1:p:141:ALA:O	1:p:145:SER:N	2.42	0.51
1:0:200:GLU:O	1:0:204:ALA:N	2.36	0.51
1:c:202:GLN:O	1:c:206:GLU:N	2.36	0.51
1:k:209:LEU:O	1:k:213:LYS:N	2.37	0.51
1:m:202:GLN:O	1:m:206:GLU:N	2.36	0.51
1:w:202:GLN:O	1:w:206:GLU:N	2.36	0.51
1:B:202:GLN:O	1:B:206:GLU:N	2.36	0.51
1:C:141:ALA:O	1:C:145:SER:N	2.42	0.51
1:F:162:ALA:N	1:F:166:SER:OG	2.39	0.51
1:I:141:ALA:O	1:I:145:SER:N	2.42	0.51
1:Z:196:PHE:O	1:Z:200:GLU:N	2.29	0.51
1:d:141:ALA:O	1:d:145:SER:N	2.42	0.51
1:e:141:ALA:O	1:e:145:SER:N	2.42	0.51
1:z:196:PHE:O	1:z:200:GLU:N	2.29	0.51
1:0:209:LEU:O	1:0:213:LYS:N	2.36	0.51
1:2:202:GLN:O	1:2:206:GLU:N	2.36	0.51
1:b:209:LEU:O	1:b:213:LYS:N	2.36	0.51
1:p:196:PHE:O	1:p:200:GLU:N	2.29	0.51
1:z:202:GLN:O	1:z:206:GLU:N	2.36	0.51
1:A:196:PHE:O	1:A:200:GLU:N	2.29	0.51
1:E:196:PHE:O	1:E:200:GLU:N	2.29	0.51
1:I:200:GLU:O	1:I:204:ALA:N	2.36	0.51
1:T:196:PHE:O	1:T:200:GLU:N	2.29	0.51
1:r:204:ALA:O	1:r:208:THR:OG1	2.14	0.51
1:v:200:GLU:O	1:v:204:ALA:N	2.36	0.51
1:y:141:ALA:O	1:y:145:SER:N	2.42	0.51
1:B:200:GLU:O	1:B:204:ALA:N	2.36	0.51
1:I:202:GLN:O	1:I:206:GLU:N	2.36	0.51
1:V:141:ALA:O	1:V:145:SER:N	2.43	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:o:141:ALA:O	1:o:145:SER:N	2.43	0.51
1:s:200:GLU:O	1:s:204:ALA:N	2.36	0.51
1:0:202:GLN:O	1:0:206:GLU:N	2.36	0.50
1:1:202:GLN:O	1:1:206:GLU:N	2.36	0.50
1:B:141:ALA:O	1:B:145:SER:N	2.42	0.50
1:J:141:ALA:O	1:J:145:SER:N	2.43	0.50
1:b:196:PHE:O	1:b:200:GLU:N	2.29	0.50
1:b:200:GLU:O	1:b:204:ALA:N	2.36	0.50
1:c:141:ALA:O	1:c:145:SER:N	2.42	0.50
1:d:209:LEU:HD12	1:d:212:MET:HB2	1.94	0.50
1:B:209:LEU:O	1:B:213:LYS:N	2.36	0.50
1:D:209:LEU:HD12	1:D:212:MET:HB2	1.94	0.50
1:F:204:ALA:O	1:F:208:THR:OG1	2.15	0.50
1:L:209:LEU:HD12	1:L:212:MET:HB2	1.94	0.50
1:M:209:LEU:HD12	1:M:212:MET:HB2	1.94	0.50
1:c:204:ALA:O	1:c:208:THR:OG1	2.15	0.50
1:f:141:ALA:O	1:f:145:SER:N	2.42	0.50
1:j:196:PHE:O	1:j:200:GLU:N	2.29	0.50
1:1:209:LEU:O	1:1:213:LYS:N	2.37	0.50
1:6:209:LEU:O	1:6:213:LYS:N	2.36	0.50
1:B:209:LEU:HD12	1:B:212:MET:HB2	1.94	0.50
1:C:209:LEU:HD12	1:C:212:MET:HB2	1.94	0.50
1:E:209:LEU:HD12	1:E:212:MET:HB2	1.94	0.50
1:J:209:LEU:HD12	1:J:212:MET:HB2	1.94	0.50
1:K:209:LEU:HD12	1:K:212:MET:HB2	1.94	0.50
1:X:202:GLN:O	1:X:206:GLU:N	2.36	0.50
1:t:209:LEU:HD12	1:t:212:MET:HB2	1.94	0.50
1:u:209:LEU:HD12	1:u:212:MET:HB2	1.94	0.50
1:y:158:ASN:O	1:y:163:SER:OG	2.30	0.50
1:y:196:PHE:O	1:y:200:GLU:N	2.29	0.50
1:0:209:LEU:HD12	1:0:212:MET:HB2	1.94	0.50
1:1:209:LEU:HD12	1:1:212:MET:HB2	1.94	0.50
1:7:209:LEU:HD12	1:7:212:MET:HB2	1.94	0.50
1:A:158:ASN:O	1:A:163:SER:OG	2.30	0.50
1:C:202:GLN:O	1:C:206:GLU:N	2.36	0.50
1:H:202:GLN:O	1:H:206:GLU:N	2.36	0.50
1:Q:209:LEU:HD12	1:Q:212:MET:HB2	1.94	0.50
1:U:209:LEU:HD12	1:U:212:MET:HB2	1.94	0.50
1:V:209:LEU:HD12	1:V:212:MET:HB2	1.94	0.50
1:W:141:ALA:O	1:W:145:SER:N	2.43	0.50
1:W:209:LEU:HD12	1:W:212:MET:HB2	1.94	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:b:209:LEU:HD12	1:b:212:MET:HB2	1.94	0.50
1:c:209:LEU:HD12	1:c:212:MET:HB2	1.94	0.50
1:k:209:LEU:HD12	1:k:212:MET:HB2	1.94	0.50
1:n:141:ALA:O	1:n:145:SER:N	2.43	0.50
1:n:209:LEU:HD12	1:n:212:MET:HB2	1.94	0.50
1:t:209:LEU:O	1:t:213:LYS:N	2.36	0.50
1:x:141:ALA:O	1:x:145:SER:N	2.43	0.50
1:x:196:PHE:O	1:x:200:GLU:N	2.29	0.50
1:z:209:LEU:HD12	1:z:212:MET:HB2	1.94	0.50
1:3:209:LEU:HD12	1:3:212:MET:HB2	1.94	0.50
1:5:209:LEU:HD12	1:5:212:MET:HB2	1.94	0.50
1:A:141:ALA:O	1:A:145:SER:N	2.42	0.50
1:F:209:LEU:HD12	1:F:212:MET:HB2	1.94	0.50
1:G:209:LEU:HD12	1:G:212:MET:HB2	1.94	0.50
1:H:209:LEU:HD12	1:H:212:MET:HB2	1.94	0.50
1:I:209:LEU:HD12	1:I:212:MET:HB2	1.94	0.50
1:N:209:LEU:HD12	1:N:212:MET:HB2	1.94	0.50
1:P:209:LEU:HD12	1:P:212:MET:HB2	1.94	0.50
1:S:200:GLU:O	1:S:204:ALA:N	2.36	0.50
1:S:209:LEU:HD12	1:S:212:MET:HB2	1.94	0.50
1:W:202:GLN:O	1:W:206:GLU:N	2.36	0.50
1:l:209:LEU:HD12	1:l:212:MET:HB2	1.94	0.50
1:m:209:LEU:HD12	1:m:212:MET:HB2	1.94	0.50
1:o:209:LEU:HD12	1:o:212:MET:HB2	1.94	0.50
1:p:209:LEU:HD12	1:p:212:MET:HB2	1.94	0.50
1:x:158:ASN:O	1:x:163:SER:OG	2.30	0.50
1:z:158:ASN:O	1:z:163:SER:OG	2.30	0.50
1:2:209:LEU:HD12	1:2:212:MET:HB2	1.94	0.50
1:6:209:LEU:HD12	1:6:212:MET:HB2	1.94	0.50
1:A:209:LEU:HD12	1:A:212:MET:HB2	1.94	0.50
1:J:196:PHE:O	1:J:200:GLU:N	2.29	0.50
1:K:141:ALA:O	1:K:145:SER:N	2.43	0.50
1:O:209:LEU:HD12	1:O:212:MET:HB2	1.94	0.50
1:T:209:LEU:HD12	1:T:212:MET:HB2	1.94	0.50
1:Y:202:GLN:O	1:Y:206:GLU:N	2.36	0.50
1:a:209:LEU:HD12	1:a:212:MET:HB2	1.94	0.50
1:d:202:GLN:O	1:d:206:GLU:N	2.36	0.50
1:f:209:LEU:HD12	1:f:212:MET:HB2	1.94	0.50
1:g:209:LEU:HD12	1:g:212:MET:HB2	1.94	0.50
1:i:209:LEU:HD12	1:i:212:MET:HB2	1.94	0.50
1:j:209:LEU:HD12	1:j:212:MET:HB2	1.94	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:l:200:GLU:O	1:l:204:ALA:N	2.36	0.50
1:o:158:ASN:O	1:o:163:SER:OG	2.30	0.50
1:p:158:ASN:O	1:p:163:SER:OG	2.30	0.50
1:s:162:ALA:N	1:s:166:SER:OG	2.39	0.50
1:v:209:LEU:HD12	1:v:212:MET:HB2	1.94	0.50
1:w:209:LEU:HD12	1:w:212:MET:HB2	1.94	0.50
1:2:209:LEU:O	1:2:213:LYS:N	2.36	0.50
1:4:209:LEU:HD12	1:4:212:MET:HB2	1.94	0.50
1:B:158:ASN:O	1:B:163:SER:OG	2.30	0.50
1:G:162:ALA:N	1:G:166:SER:OG	2.39	0.50
1:R:209:LEU:HD12	1:R:212:MET:HB2	1.94	0.50
1:X:209:LEU:HD12	1:X:212:MET:HB2	1.94	0.50
1:a:158:ASN:O	1:a:163:SER:OG	2.30	0.50
1:b:141:ALA:O	1:b:145:SER:N	2.43	0.50
1:e:209:LEU:HD12	1:e:212:MET:HB2	1.94	0.50
1:g:141:ALA:O	1:g:145:SER:N	2.42	0.50
1:h:209:LEU:HD12	1:h:212:MET:HB2	1.94	0.50
1:q:209:LEU:HD12	1:q:212:MET:HB2	1.94	0.50
1:t:196:PHE:O	1:t:200:GLU:N	2.29	0.50
1:x:209:LEU:HD12	1:x:212:MET:HB2	1.94	0.50
1:0:158:ASN:O	1:0:163:SER:OG	2.30	0.50
1:5:209:LEU:O	1:5:213:LYS:N	2.37	0.50
1:G:202:GLN:O	1:G:206:GLU:N	2.36	0.50
1:V:202:GLN:O	1:V:206:GLU:N	2.36	0.50
1:Y:209:LEU:HD12	1:Y:212:MET:HB2	1.94	0.50
1:Z:158:ASN:O	1:Z:163:SER:OG	2.30	0.50
1:b:158:ASN:O	1:b:163:SER:OG	2.30	0.50
1:l:196:PHE:O	1:l:200:GLU:N	2.29	0.50
1:m:141:ALA:O	1:m:145:SER:N	2.42	0.50
1:n:202:GLN:O	1:n:206:GLU:N	2.36	0.50
1:q:158:ASN:O	1:q:163:SER:OG	2.30	0.50
1:r:209:LEU:HD12	1:r:212:MET:HB2	1.94	0.50
1:y:209:LEU:HD12	1:y:212:MET:HB2	1.94	0.50
1:2:196:PHE:O	1:2:200:GLU:N	2.29	0.49
1:3:209:LEU:O	1:3:213:LYS:N	2.36	0.49
1:4:209:LEU:O	1:4:213:LYS:N	2.37	0.49
1:D:202:GLN:O	1:D:206:GLU:N	2.36	0.49
1:P:196:PHE:O	1:P:200:GLU:N	2.29	0.49
1:X:141:ALA:O	1:X:145:SER:N	2.42	0.49
1:Z:209:LEU:HD12	1:Z:212:MET:HB2	1.94	0.49
1:n:158:ASN:O	1:n:163:SER:OG	2.30	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:w:141:ALA:O	1:w:145:SER:N	2.42	0.49
1:x:202:GLN:O	1:x:206:GLU:N	2.36	0.49
1:F:202:GLN:O	1:F:206:GLU:N	2.36	0.49
1:L:141:ALA:O	1:L:145:SER:N	2.42	0.49
1:M:200:GLU:O	1:M:204:ALA:N	2.36	0.49
1:O:158:ASN:O	1:O:163:SER:OG	2.30	0.49
1:a:141:ALA:O	1:a:145:SER:N	2.42	0.49
1:l:209:LEU:O	1:l:213:LYS:N	2.37	0.49
1:q:196:PHE:O	1:q:200:GLU:N	2.29	0.49
1:r:158:ASN:O	1:r:163:SER:OG	2.30	0.49
1:t:141:ALA:O	1:t:145:SER:N	2.42	0.49
1:y:204:ALA:O	1:y:208:THR:OG1	2.15	0.49
1:1:200:GLU:O	1:1:204:ALA:N	2.36	0.49
1:7:196:PHE:O	1:7:200:GLU:N	2.29	0.49
1:E:202:GLN:O	1:E:206:GLU:N	2.36	0.49
1:P:158:ASN:O	1:P:163:SER:OG	2.30	0.49
1:g:196:PHE:O	1:g:200:GLU:N	2.29	0.49
1:s:204:ALA:O	1:s:208:THR:OG1	2.14	0.49
1:s:209:LEU:HD12	1:s:212:MET:HB2	1.94	0.49
1:1:158:ASN:O	1:1:163:SER:OG	2.30	0.49
1:F:196:PHE:O	1:F:200:GLU:N	2.29	0.49
1:W:200:GLU:O	1:W:204:ALA:N	2.36	0.49
1:Y:141:ALA:O	1:Y:145:SER:N	2.43	0.49
1:Z:202:GLN:O	1:Z:206:GLU:N	2.36	0.49
1:c:158:ASN:O	1:c:163:SER:OG	2.30	0.49
1:t:200:GLU:O	1:t:204:ALA:N	2.36	0.49
1:v:141:ALA:O	1:v:145:SER:N	2.42	0.49
1:7:209:LEU:O	1:7:213:LYS:N	2.36	0.49
1:C:158:ASN:O	1:C:163:SER:OG	2.30	0.49
1:J:158:ASN:O	1:J:163:SER:OG	2.30	0.49
1:L:204:ALA:O	1:L:208:THR:OG1	2.14	0.49
1:M:196:PHE:O	1:M:200:GLU:N	2.29	0.49
1:T:158:ASN:O	1:T:163:SER:OG	2.30	0.49
1:Z:141:ALA:O	1:Z:145:SER:N	2.43	0.49
1:h:141:ALA:O	1:h:145:SER:N	2.42	0.49
1:i:200:GLU:O	1:i:204:ALA:N	2.36	0.49
1:j:200:GLU:O	1:j:204:ALA:N	2.36	0.49
1:j:205:VAL:HG23	1:j:206:GLU:N	2.28	0.49
1:u:141:ALA:O	1:u:145:SER:N	2.42	0.49
1:u:209:LEU:O	1:u:213:LYS:N	2.36	0.49
1:7:158:ASN:O	1:7:163:SER:OG	2.30	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:209:LEU:O	1:J:213:LYS:N	2.37	0.49
1:M:141:ALA:O	1:M:145:SER:N	2.42	0.49
1:Q:158:ASN:O	1:Q:163:SER:OG	2.30	0.49
1:U:202:GLN:O	1:U:206:GLU:N	2.36	0.49
1:W:196:PHE:O	1:W:200:GLU:N	2.29	0.49
1:X:158:ASN:O	1:X:163:SER:OG	2.30	0.49
1:a:205:VAL:HG23	1:a:206:GLU:N	2.28	0.49
1:c:209:LEU:O	1:c:213:LYS:N	2.37	0.49
1:d:200:GLU:O	1:d:204:ALA:N	2.36	0.49
1:e:158:ASN:O	1:e:163:SER:OG	2.30	0.49
1:f:158:ASN:O	1:f:163:SER:OG	2.30	0.49
1:g:205:VAL:HG23	1:g:206:GLU:N	2.28	0.49
1:j:158:ASN:O	1:j:163:SER:OG	2.30	0.49
1:k:158:ASN:O	1:k:163:SER:OG	2.30	0.49
1:3:200:GLU:O	1:3:204:ALA:N	2.36	0.49
1:6:158:ASN:O	1:6:163:SER:OG	2.30	0.49
1:C:209:LEU:O	1:C:213:LYS:N	2.37	0.49
1:N:205:VAL:HG23	1:N:206:GLU:N	2.28	0.49
1:O:205:VAL:HG23	1:O:206:GLU:N	2.28	0.49
1:P:205:VAL:HG23	1:P:206:GLU:N	2.28	0.49
1:U:158:ASN:O	1:U:163:SER:OG	2.30	0.49
1:X:205:VAL:HG23	1:X:206:GLU:N	2.28	0.49
1:a:196:PHE:O	1:a:200:GLU:N	2.29	0.49
1:c:200:GLU:O	1:c:204:ALA:N	2.36	0.49
1:h:200:GLU:O	1:h:204:ALA:N	2.36	0.49
1:l:141:ALA:O	1:l:145:SER:N	2.43	0.49
1:m:158:ASN:O	1:m:163:SER:OG	2.30	0.49
1:s:158:ASN:O	1:s:163:SER:OG	2.30	0.49
1:v:158:ASN:O	1:v:163:SER:OG	2.30	0.49
1:v:205:VAL:HG23	1:v:206:GLU:N	2.28	0.49
1:2:205:VAL:HG23	1:2:206:GLU:N	2.28	0.49
1:C:200:GLU:O	1:C:204:ALA:N	2.36	0.49
1:G:204:ALA:O	1:G:208:THR:OG1	2.14	0.49
1:I:158:ASN:O	1:I:163:SER:OG	2.30	0.49
1:K:205:VAL:HG23	1:K:206:GLU:N	2.28	0.49
1:L:205:VAL:HG23	1:L:206:GLU:N	2.28	0.49
1:N:158:ASN:O	1:N:163:SER:OG	2.30	0.49
1:W:205:VAL:HG23	1:W:206:GLU:N	2.28	0.49
1:f:204:ALA:O	1:f:208:THR:OG1	2.15	0.49
1:f:213:LYS:O	1:f:217:SER:OG	2.07	0.49
1:g:158:ASN:O	1:g:163:SER:OG	2.30	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:g:200:GLU:O	1:g:204:ALA:N	2.36	0.49
1:g:209:LEU:O	1:g:213:LYS:N	2.37	0.49
1:i:141:ALA:O	1:i:145:SER:N	2.43	0.49
1:i:158:ASN:O	1:i:163:SER:OG	2.30	0.49
1:t:205:VAL:HG23	1:t:206:GLU:N	2.28	0.49
1:l:205:VAL:HG23	1:l:206:GLU:N	2.28	0.49
1:4:158:ASN:O	1:4:163:SER:OG	2.30	0.49
1:A:205:VAL:HG23	1:A:206:GLU:N	2.28	0.49
1:K:158:ASN:O	1:K:163:SER:OG	2.30	0.49
1:L:158:ASN:O	1:L:163:SER:OG	2.30	0.49
1:Q:205:VAL:HG23	1:Q:206:GLU:N	2.28	0.49
1:T:209:LEU:O	1:T:213:LYS:N	2.37	0.49
1:d:158:ASN:O	1:d:163:SER:OG	2.30	0.49
1:j:141:ALA:O	1:j:145:SER:N	2.42	0.49
1:k:141:ALA:O	1:k:145:SER:N	2.42	0.49
1:k:200:GLU:O	1:k:204:ALA:N	2.36	0.49
1:l:158:ASN:O	1:l:163:SER:OG	2.30	0.49
1:w:205:VAL:HG23	1:w:206:GLU:N	2.28	0.49
1:2:158:ASN:O	1:2:163:SER:OG	2.30	0.49
1:6:205:VAL:HG23	1:6:206:GLU:N	2.28	0.49
1:N:141:ALA:O	1:N:145:SER:N	2.42	0.49
1:S:158:ASN:O	1:S:163:SER:OG	2.30	0.49
1:U:205:VAL:HG23	1:U:206:GLU:N	2.28	0.49
1:e:200:GLU:O	1:e:204:ALA:N	2.36	0.49
1:f:205:VAL:HG23	1:f:206:GLU:N	2.28	0.49
1:m:205:VAL:HG23	1:m:206:GLU:N	2.28	0.49
1:u:158:ASN:C	1:u:163:SER:HG	2.19	0.49
1:O:141:ALA:O	1:O:145:SER:N	2.42	0.48
1:R:158:ASN:O	1:R:163:SER:OG	2.30	0.48
1:T:205:VAL:HG23	1:T:206:GLU:N	2.28	0.48
1:W:158:ASN:O	1:W:163:SER:OG	2.30	0.48
1:Z:205:VAL:HG23	1:Z:206:GLU:N	2.28	0.48
1:h:158:ASN:O	1:h:163:SER:OG	2.30	0.48
1:i:205:VAL:HG23	1:i:206:GLU:N	2.28	0.48
1:k:205:VAL:HG23	1:k:206:GLU:N	2.28	0.48
1:0:205:VAL:HG23	1:0:206:GLU:N	2.28	0.48
1:3:205:VAL:HG23	1:3:206:GLU:N	2.28	0.48
1:7:200:GLU:O	1:7:204:ALA:N	2.36	0.48
1:D:158:ASN:O	1:D:163:SER:OG	2.30	0.48
1:E:158:ASN:O	1:E:163:SER:OG	2.30	0.48
1:F:200:GLU:O	1:F:204:ALA:N	2.36	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:158:ASN:O	1:G:163:SER:OG	2.30	0.48
1:T:202:GLN:O	1:T:206:GLU:N	2.36	0.48
1:W:209:LEU:O	1:W:213:LYS:N	2.36	0.48
1:Y:163:SER:HG	1:Y:164:ALA:H	1.61	0.48
1:Z:200:GLU:O	1:Z:204:ALA:N	2.36	0.48
1:C:205:VAL:HG23	1:C:206:GLU:N	2.28	0.48
1:M:158:ASN:O	1:M:163:SER:OG	2.30	0.48
1:n:205:VAL:HG23	1:n:206:GLU:N	2.28	0.48
1:o:202:GLN:O	1:o:206:GLU:N	2.36	0.48
1:p:205:VAL:HG23	1:p:206:GLU:N	2.28	0.48
1:3:196:PHE:O	1:3:200:GLU:N	2.29	0.48
1:B:205:VAL:HG23	1:B:206:GLU:N	2.28	0.48
1:H:205:VAL:HG23	1:H:206:GLU:N	2.28	0.48
1:a:202:GLN:O	1:a:206:GLU:N	2.36	0.48
1:o:205:VAL:HG23	1:o:206:GLU:N	2.28	0.48
1:q:159:LEU:O	1:q:165:SER:OG	2.24	0.48
1:q:200:GLU:O	1:q:204:ALA:N	2.36	0.48
1:s:159:LEU:O	1:s:165:SER:OG	2.24	0.48
1:3:158:ASN:O	1:3:163:SER:OG	2.30	0.48
1:G:205:VAL:HG23	1:G:206:GLU:N	2.28	0.48
1:P:200:GLU:O	1:P:204:ALA:N	2.36	0.48
1:R:205:VAL:HG23	1:R:206:GLU:N	2.28	0.48
1:V:205:VAL:HG23	1:V:206:GLU:N	2.28	0.48
1:Y:205:VAL:HG23	1:Y:206:GLU:N	2.28	0.48
1:b:205:VAL:HG23	1:b:206:GLU:N	2.28	0.48
1:d:205:VAL:HG23	1:d:206:GLU:N	2.28	0.48
1:h:205:VAL:HG23	1:h:206:GLU:N	2.28	0.48
1:o:159:LEU:O	1:o:165:SER:OG	2.24	0.48
1:r:159:LEU:O	1:r:165:SER:OG	2.25	0.48
1:v:196:PHE:O	1:v:200:GLU:N	2.29	0.48
1:y:205:VAL:HG23	1:y:206:GLU:N	2.28	0.48
1:z:205:VAL:HG23	1:z:206:GLU:N	2.28	0.48
1:5:205:VAL:HG23	1:5:206:GLU:N	2.28	0.48
1:6:204:ALA:O	1:6:208:THR:OG1	2.14	0.48
1:7:205:VAL:HG23	1:7:206:GLU:N	2.28	0.48
1:D:159:LEU:O	1:D:165:SER:OG	2.24	0.48
1:K:209:LEU:O	1:K:213:LYS:N	2.37	0.48
1:2:200:GLU:O	1:2:204:ALA:N	2.36	0.48
1:D:205:VAL:HG23	1:D:206:GLU:N	2.28	0.48
1:D:209:LEU:O	1:D:213:LYS:N	2.37	0.48
1:E:159:LEU:O	1:E:165:SER:OG	2.24	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:159:LEU:O	1:F:165:SER:OG	2.24	0.48
1:S:202:GLN:O	1:S:206:GLU:N	2.36	0.48
1:U:209:LEU:O	1:U:213:LYS:N	2.37	0.48
1:Y:203:GLN:O	1:Y:207:ALA:N	2.47	0.48
1:a:203:GLN:O	1:a:207:ALA:N	2.47	0.48
1:l:205:VAL:HG23	1:l:206:GLU:N	2.28	0.48
1:u:203:GLN:O	1:u:207:ALA:N	2.47	0.48
1:u:205:VAL:HG23	1:u:206:GLU:N	2.28	0.48
1:v:209:LEU:O	1:v:213:LYS:N	2.36	0.48
1:2:203:GLN:O	1:2:207:ALA:N	2.47	0.48
1:3:203:GLN:O	1:3:207:ALA:N	2.47	0.48
1:4:203:GLN:O	1:4:207:ALA:N	2.47	0.48
1:4:205:VAL:HG23	1:4:206:GLU:N	2.28	0.48
1:F:205:VAL:HG23	1:F:206:GLU:N	2.28	0.48
1:G:159:LEU:O	1:G:165:SER:OG	2.24	0.48
1:I:205:VAL:HG23	1:I:206:GLU:N	2.28	0.48
1:J:200:GLU:O	1:J:204:ALA:N	2.36	0.48
1:J:205:VAL:HG23	1:J:206:GLU:N	2.28	0.48
1:M:205:VAL:HG23	1:M:206:GLU:N	2.28	0.48
1:N:203:GLN:O	1:N:207:ALA:N	2.47	0.48
1:O:203:GLN:O	1:O:207:ALA:N	2.47	0.48
1:P:203:GLN:O	1:P:207:ALA:N	2.47	0.48
1:Q:203:GLN:O	1:Q:207:ALA:N	2.47	0.48
1:R:203:GLN:O	1:R:207:ALA:N	2.47	0.48
1:X:203:GLN:O	1:X:207:ALA:N	2.47	0.48
1:Z:203:GLN:O	1:Z:207:ALA:N	2.47	0.48
1:i:203:GLN:O	1:i:207:ALA:N	2.47	0.48
1:j:203:GLN:O	1:j:207:ALA:N	2.47	0.48
1:s:205:VAL:HG23	1:s:206:GLU:N	2.28	0.48
1:t:203:GLN:O	1:t:207:ALA:N	2.47	0.48
1:v:203:GLN:O	1:v:207:ALA:N	2.47	0.48
1:w:203:GLN:O	1:w:207:ALA:N	2.47	0.48
1:y:202:GLN:O	1:y:206:GLU:N	2.36	0.48
1:1:203:GLN:O	1:1:207:ALA:N	2.47	0.48
1:B:159:LEU:O	1:B:165:SER:OG	2.24	0.48
1:I:159:LEU:O	1:I:165:SER:OG	2.24	0.48
1:S:203:GLN:O	1:S:207:ALA:N	2.47	0.48
1:T:203:GLN:O	1:T:207:ALA:N	2.47	0.48
1:W:203:GLN:O	1:W:207:ALA:N	2.47	0.48
1:b:203:GLN:O	1:b:207:ALA:N	2.47	0.48
1:m:209:LEU:O	1:m:213:LYS:N	2.37	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:q:205:VAL:HG23	1:q:206:GLU:N	2.28	0.48
1:s:196:PHE:O	1:s:200:GLU:N	2.29	0.48
1:0:203:GLN:O	1:0:207:ALA:N	2.47	0.48
1:5:203:GLN:O	1:5:207:ALA:N	2.47	0.48
1:7:203:GLN:O	1:7:207:ALA:N	2.47	0.48
1:M:209:LEU:O	1:M:213:LYS:N	2.37	0.48
1:P:202:GLN:O	1:P:206:GLU:N	2.36	0.48
1:V:203:GLN:O	1:V:207:ALA:N	2.47	0.48
1:c:203:GLN:O	1:c:207:ALA:N	2.47	0.48
1:c:205:VAL:HG23	1:c:206:GLU:N	2.28	0.48
1:d:203:GLN:O	1:d:207:ALA:N	2.47	0.48
1:d:209:LEU:O	1:d:213:LYS:N	2.37	0.48
1:e:205:VAL:HG23	1:e:206:GLU:N	2.28	0.48
1:e:209:LEU:O	1:e:213:LYS:N	2.36	0.48
1:i:202:GLN:O	1:i:206:GLU:N	2.36	0.48
1:x:203:GLN:O	1:x:207:ALA:N	2.47	0.48
1:6:203:GLN:O	1:6:207:ALA:N	2.47	0.47
1:E:205:VAL:HG23	1:E:206:GLU:N	2.28	0.47
1:G:203:GLN:O	1:G:207:ALA:N	2.47	0.47
1:J:159:LEU:O	1:J:165:SER:OG	2.24	0.47
1:M:203:GLN:O	1:M:207:ALA:N	2.47	0.47
1:Q:202:GLN:O	1:Q:206:GLU:N	2.36	0.47
1:R:202:GLN:O	1:R:206:GLU:N	2.36	0.47
1:U:203:GLN:O	1:U:207:ALA:N	2.47	0.47
1:k:203:GLN:O	1:k:207:ALA:N	2.47	0.47
1:t:163:SER:HG	1:t:164:ALA:H	1.61	0.47
1:x:205:VAL:HG23	1:x:206:GLU:N	2.28	0.47
1:y:203:GLN:O	1:y:207:ALA:N	2.47	0.47
1:z:203:GLN:O	1:z:207:ALA:N	2.47	0.47
1:C:203:GLN:O	1:C:207:ALA:N	2.47	0.47
1:L:209:LEU:O	1:L:213:LYS:N	2.37	0.47
1:e:159:LEU:O	1:e:165:SER:OG	2.24	0.47
1:h:209:LEU:O	1:h:213:LYS:N	2.37	0.47
1:j:202:GLN:O	1:j:206:GLU:N	2.36	0.47
1:p:203:GLN:O	1:p:207:ALA:N	2.47	0.47
1:B:203:GLN:O	1:B:207:ALA:N	2.47	0.47
1:S:205:VAL:HG23	1:S:206:GLU:N	2.28	0.47
1:h:202:GLN:O	1:h:206:GLU:N	2.36	0.47
1:r:205:VAL:HG23	1:r:206:GLU:N	2.28	0.47
1:D:203:GLN:O	1:D:207:ALA:N	2.47	0.47
1:f:159:LEU:O	1:f:165:SER:OG	2.24	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:h:203:GLN:O	1:h:207:ALA:N	2.47	0.47
1:o:203:GLN:O	1:o:207:ALA:N	2.47	0.47
1:s:203:GLN:O	1:s:207:ALA:N	2.47	0.47
1:5:204:ALA:O	1:5:208:THR:OG1	2.14	0.47
1:E:203:GLN:O	1:E:207:ALA:N	2.47	0.47
1:E:209:LEU:O	1:E:213:LYS:N	2.36	0.47
1:T:200:GLU:O	1:T:204:ALA:N	2.36	0.47
1:l:159:LEU:O	1:l:165:SER:OG	2.24	0.47
1:p:202:GLN:O	1:p:206:GLU:N	2.36	0.47
1:q:203:GLN:O	1:q:207:ALA:N	2.47	0.47
1:D:200:GLU:O	1:D:204:ALA:N	2.36	0.47
1:F:203:GLN:O	1:F:207:ALA:N	2.47	0.47
1:L:159:LEU:O	1:L:165:SER:OG	2.24	0.47
1:A:203:GLN:O	1:A:207:ALA:N	2.47	0.47
1:G:209:LEU:O	1:G:213:LYS:N	2.36	0.47
1:H:203:GLN:O	1:H:207:ALA:N	2.47	0.47
1:I:203:GLN:O	1:I:207:ALA:N	2.47	0.47
1:L:203:GLN:O	1:L:207:ALA:N	2.47	0.47
1:M:159:LEU:O	1:M:165:SER:OG	2.24	0.47
1:N:209:LEU:O	1:N:213:LYS:N	2.37	0.47
1:O:159:LEU:O	1:O:165:SER:OG	2.24	0.47
1:e:203:GLN:O	1:e:207:ALA:N	2.47	0.47
1:g:159:LEU:O	1:g:165:SER:OG	2.24	0.47
1:g:202:GLN:O	1:g:206:GLU:N	2.36	0.47
1:g:203:GLN:O	1:g:207:ALA:N	2.47	0.47
1:k:159:LEU:O	1:k:165:SER:OG	2.24	0.47
1:l:203:GLN:O	1:l:207:ALA:N	2.47	0.47
1:n:203:GLN:O	1:n:207:ALA:N	2.47	0.47
1:r:203:GLN:O	1:r:207:ALA:N	2.47	0.47
1:u:204:ALA:O	1:u:208:THR:OG1	2.15	0.47
1:y:200:GLU:O	1:y:204:ALA:N	2.36	0.47
1:J:203:GLN:O	1:J:207:ALA:N	2.47	0.47
1:N:159:LEU:O	1:N:165:SER:OG	2.24	0.47
1:X:209:LEU:O	1:X:213:LYS:N	2.37	0.47
1:f:203:GLN:O	1:f:207:ALA:N	2.47	0.47
1:i:159:LEU:O	1:i:165:SER:OG	2.24	0.47
1:k:202:GLN:O	1:k:206:GLU:N	2.36	0.47
1:K:203:GLN:O	1:K:207:ALA:N	2.47	0.47
1:m:203:GLN:O	1:m:207:ALA:N	2.47	0.47
1:t:158:ASN:C	1:t:163:SER:HG	2.20	0.47
1:w:200:GLU:O	1:w:204:ALA:N	2.36	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:209:LEU:O	1:I:213:LYS:N	2.37	0.47
1:F:209:LEU:O	1:F:213:LYS:N	2.37	0.46
1:V:39:GLU:OE2	1:W:152:ILE:N	2.48	0.46
1:m:200:GLU:O	1:m:204:ALA:N	2.36	0.46
1:H:209:LEU:O	1:H:213:LYS:N	2.37	0.46
1:f:202:GLN:O	1:f:206:GLU:N	2.36	0.46
1:w:209:LEU:O	1:w:213:LYS:N	2.37	0.46
1:S:196:PHE:O	1:S:200:GLU:N	2.29	0.46
1:V:209:LEU:O	1:V:213:LYS:N	2.36	0.46
1:o:200:GLU:O	1:o:204:ALA:N	2.36	0.46
1:q:202:GLN:O	1:q:206:GLU:N	2.36	0.46
1:O:209:LEU:O	1:O:213:LYS:N	2.37	0.46
1:f:209:LEU:O	1:f:213:LYS:N	2.36	0.46
1:j:204:ALA:O	1:j:208:THR:OG1	2.15	0.46
1:4:200:GLU:O	1:4:204:ALA:N	2.36	0.46
1:O:196:PHE:O	1:O:200:GLU:N	2.29	0.46
1:n:209:LEU:O	1:n:213:LYS:N	2.37	0.46
1:e:202:GLN:O	1:e:206:GLU:N	2.36	0.46
1:Q:209:LEU:O	1:Q:213:LYS:N	2.37	0.46
1:j:209:LEU:O	1:j:213:LYS:N	2.37	0.46
1:N:200:GLU:O	1:N:204:ALA:N	2.36	0.46
1:X:200:GLU:O	1:X:204:ALA:N	2.36	0.46
1:G:200:GLU:O	1:G:204:ALA:N	2.36	0.45
1:P:209:LEU:O	1:P:213:LYS:N	2.36	0.45
1:R:209:LEU:O	1:R:213:LYS:N	2.36	0.45
1:I:196:PHE:O	1:I:200:GLU:N	2.29	0.45
1:I:204:ALA:O	1:I:208:THR:OG1	2.14	0.45
1:Y:196:PHE:O	1:Y:200:GLU:N	2.29	0.45
1:5:200:GLU:O	1:5:204:ALA:N	2.36	0.45
1:E:200:GLU:O	1:E:204:ALA:N	2.36	0.45
1:u:200:GLU:O	1:u:204:ALA:N	2.36	0.45
1:i:196:PHE:O	1:i:200:GLU:N	2.29	0.45
1:i:209:LEU:O	1:i:213:LYS:N	2.37	0.45
1:Q:200:GLU:O	1:Q:204:ALA:N	2.36	0.45
1:u:203:GLN:O	1:u:207:ALA:HB3	2.17	0.45
1:1:204:ALA:O	1:1:208:THR:OG1	2.15	0.45
1:X:203:GLN:O	1:X:207:ALA:HB3	2.17	0.45
1:Y:203:GLN:O	1:Y:207:ALA:HB3	2.17	0.45
1:V:203:GLN:O	1:V:207:ALA:HB3	2.17	0.45
1:W:203:GLN:O	1:W:207:ALA:HB3	2.17	0.45
1:Z:203:GLN:O	1:Z:207:ALA:HB3	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:n:200:GLU:O	1:n:204:ALA:N	2.36	0.45
1:t:202:GLN:O	1:t:206:GLU:N	2.36	0.45
1:t:203:GLN:O	1:t:207:ALA:HB3	2.17	0.45
1:7:203:GLN:O	1:7:207:ALA:HB3	2.17	0.45
1:Z:209:LEU:O	1:Z:213:LYS:N	2.37	0.45
1:v:203:GLN:O	1:v:207:ALA:HB3	2.17	0.45
1:a:203:GLN:O	1:a:207:ALA:HB3	2.17	0.45
1:u:202:GLN:O	1:u:206:GLU:N	2.36	0.45
1:w:203:GLN:O	1:w:207:ALA:HB3	2.17	0.45
1:A:204:ALA:O	1:A:208:THR:OG1	2.14	0.44
1:K:200:GLU:O	1:K:204:ALA:N	2.36	0.44
1:M:203:GLN:O	1:M:207:ALA:HB3	2.17	0.44
1:x:209:LEU:O	1:x:213:LYS:N	2.36	0.44
1:6:203:GLN:O	1:6:207:ALA:HB3	2.17	0.44
1:Y:209:LEU:O	1:Y:213:LYS:N	2.36	0.44
1:h:203:GLN:O	1:h:207:ALA:HB3	2.17	0.44
1:k:203:GLN:O	1:k:207:ALA:HB3	2.17	0.44
1:m:213:LYS:O	1:m:217:SER:OG	2.07	0.44
1:1:203:GLN:O	1:1:207:ALA:HB3	2.17	0.44
1:S:209:LEU:O	1:S:213:LYS:N	2.37	0.44
1:U:203:GLN:O	1:U:207:ALA:HB3	2.17	0.44
1:O:203:GLN:O	1:O:207:ALA:HB3	2.17	0.44
1:P:203:GLN:O	1:P:207:ALA:HB3	2.17	0.44
1:Q:93:GLN:O	1:Q:96:SER:OG	2.35	0.44
1:b:203:GLN:O	1:b:207:ALA:HB3	2.17	0.44
1:i:203:GLN:O	1:i:207:ALA:HB3	2.17	0.44
1:m:93:GLN:O	1:m:96:SER:OG	2.35	0.44
1:o:209:LEU:O	1:o:213:LYS:N	2.37	0.44
1:p:203:GLN:O	1:p:207:ALA:HB3	2.17	0.44
1:s:203:GLN:O	1:s:207:ALA:HB3	2.17	0.44
1:C:203:GLN:O	1:C:207:ALA:HB3	2.17	0.44
1:G:203:GLN:O	1:G:207:ALA:HB3	2.17	0.44
1:L:203:GLN:O	1:L:207:ALA:HB3	2.17	0.44
1:4:203:GLN:O	1:4:207:ALA:HB3	2.17	0.44
1:D:203:GLN:O	1:D:207:ALA:HB3	2.17	0.44
1:F:203:GLN:O	1:F:207:ALA:HB3	2.17	0.44
1:T:203:GLN:O	1:T:207:ALA:HB3	2.17	0.44
1:x:203:GLN:O	1:x:207:ALA:HB3	2.17	0.44
1:2:203:GLN:O	1:2:207:ALA:HB3	2.17	0.44
1:J:203:GLN:O	1:J:207:ALA:HB3	2.17	0.44
1:f:203:GLN:O	1:f:207:ALA:HB3	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:l:203:GLN:O	1:l:207:ALA:HB3	2.17	0.44
1:m:203:GLN:O	1:m:207:ALA:HB3	2.17	0.44
1:7:202:GLN:O	1:7:206:GLU:N	2.36	0.44
1:N:203:GLN:O	1:N:207:ALA:HB3	2.17	0.44
1:S:203:GLN:O	1:S:207:ALA:HB3	2.17	0.44
1:d:203:GLN:O	1:d:207:ALA:HB3	2.17	0.44
1:r:203:GLN:O	1:r:207:ALA:HB3	2.17	0.44
1:5:203:GLN:O	1:5:207:ALA:HB3	2.17	0.44
1:A:203:GLN:O	1:A:207:ALA:HB3	2.17	0.44
1:S:93:GLN:O	1:S:96:SER:OG	2.35	0.44
1:U:200:GLU:O	1:U:204:ALA:N	2.36	0.44
1:j:203:GLN:O	1:j:207:ALA:HB3	2.17	0.44
1:n:203:GLN:O	1:n:207:ALA:HB3	2.17	0.44
1:y:203:GLN:O	1:y:207:ALA:HB3	2.17	0.44
1:R:203:GLN:O	1:R:207:ALA:HB3	2.17	0.43
1:a:204:ALA:O	1:a:208:THR:OG1	2.14	0.43
1:c:203:GLN:O	1:c:207:ALA:HB3	2.17	0.43
1:g:203:GLN:O	1:g:207:ALA:HB3	2.17	0.43
1:o:203:GLN:O	1:o:207:ALA:HB3	2.17	0.43
1:v:93:GLN:O	1:v:96:SER:OG	2.35	0.43
1:B:203:GLN:O	1:B:207:ALA:HB3	2.17	0.43
1:E:203:GLN:O	1:E:207:ALA:HB3	2.17	0.43
1:K:203:GLN:O	1:K:207:ALA:HB3	2.17	0.43
1:e:203:GLN:O	1:e:207:ALA:HB3	2.17	0.43
1:g:93:GLN:O	1:g:96:SER:OG	2.35	0.43
1:q:203:GLN:O	1:q:207:ALA:HB3	2.17	0.43
1:0:203:GLN:O	1:0:207:ALA:HB3	2.17	0.43
1:I:203:GLN:O	1:I:207:ALA:HB3	2.17	0.43
1:Q:203:GLN:O	1:Q:207:ALA:HB3	2.17	0.43
1:o:93:GLN:O	1:o:96:SER:OG	2.35	0.43
1:z:203:GLN:O	1:z:207:ALA:HB3	2.17	0.43
1:0:152:ILE:N	1:z:39:GLU:OE2	2.52	0.43
1:2:204:ALA:O	1:2:208:THR:OG1	2.14	0.43
1:G:93:GLN:O	1:G:96:SER:OG	2.35	0.43
1:H:205:VAL:O	1:H:209:LEU:CB	2.67	0.43
1:I:205:VAL:O	1:I:209:LEU:CB	2.67	0.43
1:J:205:VAL:O	1:J:209:LEU:CB	2.67	0.43
1:Q:204:ALA:O	1:Q:208:THR:OG1	2.15	0.43
1:e:205:VAL:O	1:e:209:LEU:CB	2.67	0.43
1:1:39:GLU:OE2	1:2:152:ILE:N	2.52	0.43
1:A:205:VAL:O	1:A:209:LEU:CB	2.67	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:39:GLU:OE2	1:C:152:ILE:N	2.52	0.43
1:D:39:GLU:OE2	1:E:152:ILE:N	2.52	0.43
1:G:205:VAL:O	1:G:209:LEU:CB	2.67	0.43
1:H:203:GLN:O	1:H:207:ALA:HB3	2.17	0.43
1:K:205:VAL:O	1:K:209:LEU:CB	2.67	0.43
1:P:39:GLU:OE2	1:Q:152:ILE:N	2.52	0.43
1:R:39:GLU:OE2	1:S:152:ILE:N	2.52	0.43
1:U:93:GLN:O	1:U:96:SER:OG	2.35	0.43
1:d:205:VAL:O	1:d:209:LEU:CB	2.67	0.43
1:f:205:VAL:O	1:f:209:LEU:CB	2.67	0.43
1:n:39:GLU:OE2	1:o:152:ILE:N	2.52	0.43
1:p:159:LEU:O	1:p:165:SER:OG	2.24	0.43
1:t:39:GLU:OE2	1:u:152:ILE:N	2.52	0.43
1:w:205:VAL:O	1:w:209:LEU:CB	2.67	0.43
1:3:203:GLN:O	1:3:207:ALA:HB3	2.17	0.43
1:F:39:GLU:OE2	1:G:152:ILE:N	2.52	0.43
1:g:205:VAL:O	1:g:209:LEU:CB	2.67	0.43
1:h:39:GLU:OE2	1:i:152:ILE:N	2.52	0.43
1:p:39:GLU:OE2	1:q:152:ILE:N	2.52	0.43
1:x:39:GLU:OE2	1:y:152:ILE:N	2.52	0.43
1:D:205:VAL:O	1:D:209:LEU:CB	2.67	0.43
1:L:39:GLU:OE2	1:M:152:ILE:N	2.52	0.43
1:W:205:VAL:O	1:W:209:LEU:CB	2.67	0.43
1:Y:39:GLU:OE2	1:Z:152:ILE:N	2.52	0.43
1:Y:158:ASN:C	1:Y:163:SER:HG	2.20	0.43
1:l:39:GLU:OE2	1:m:152:ILE:N	2.52	0.43
1:m:205:VAL:O	1:m:209:LEU:CB	2.67	0.43
1:p:205:VAL:O	1:p:209:LEU:CB	2.67	0.43
1:y:205:VAL:O	1:y:209:LEU:CB	2.67	0.43
1:z:205:VAL:O	1:z:209:LEU:CB	2.67	0.43
1:2:93:GLN:O	1:2:96:SER:OG	2.35	0.43
1:3:39:GLU:OE2	1:4:152:ILE:N	2.52	0.43
1:4:39:GLU:OE2	1:5:152:ILE:N	2.52	0.43
1:C:205:VAL:O	1:C:209:LEU:CB	2.67	0.43
1:H:39:GLU:OE2	1:I:152:ILE:N	2.52	0.43
1:L:205:VAL:O	1:L:209:LEU:CB	2.67	0.43
1:M:39:GLU:OE2	1:N:152:ILE:N	2.52	0.43
1:V:205:VAL:O	1:V:209:LEU:CB	2.67	0.43
1:X:39:GLU:OE2	1:Y:152:ILE:N	2.52	0.43
1:X:205:VAL:O	1:X:209:LEU:CB	2.67	0.43
1:a:205:VAL:O	1:a:209:LEU:CB	2.67	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:b:39:GLU:OE2	1:c:152:ILE:N	2.52	0.43
1:b:205:VAL:O	1:b:209:LEU:CB	2.67	0.43
1:c:39:GLU:OE2	1:d:152:ILE:N	2.52	0.43
1:e:39:GLU:OE2	1:f:152:ILE:N	2.52	0.43
1:k:205:VAL:O	1:k:209:LEU:CB	2.67	0.43
1:n:205:VAL:O	1:n:209:LEU:CB	2.67	0.43
1:q:39:GLU:OE2	1:r:152:ILE:N	2.52	0.43
1:t:205:VAL:O	1:t:209:LEU:CB	2.67	0.43
1:1:205:VAL:O	1:1:209:LEU:CB	2.67	0.43
1:I:93:GLN:O	1:I:96:SER:OG	2.35	0.43
1:J:39:GLU:OE2	1:K:152:ILE:N	2.52	0.43
1:h:205:VAL:O	1:h:209:LEU:CB	2.67	0.43
1:r:39:GLU:OE2	1:s:152:ILE:N	2.52	0.43
1:s:205:VAL:O	1:s:209:LEU:CB	2.67	0.43
1:5:39:GLU:OE2	1:6:152:ILE:N	2.52	0.43
1:6:205:VAL:O	1:6:209:LEU:CB	2.67	0.43
1:F:205:VAL:O	1:F:209:LEU:CB	2.67	0.43
1:O:205:VAL:O	1:O:209:LEU:CB	2.67	0.43
1:T:205:VAL:O	1:T:209:LEU:CB	2.67	0.43
1:Y:205:VAL:O	1:Y:209:LEU:CB	2.67	0.43
1:Z:39:GLU:OE2	1:a:152:ILE:N	2.52	0.43
1:Z:93:GLN:O	1:Z:96:SER:OG	2.35	0.43
1:j:39:GLU:OE2	1:k:152:ILE:N	2.52	0.43
1:k:39:GLU:OE2	1:l:152:ILE:N	2.52	0.43
1:o:39:GLU:OE2	1:p:152:ILE:N	2.52	0.43
1:2:39:GLU:OE2	1:3:152:ILE:N	2.52	0.42
1:6:39:GLU:OE2	1:7:152:ILE:N	2.52	0.42
1:7:205:VAL:O	1:7:209:LEU:CB	2.67	0.42
1:I:39:GLU:OE2	1:J:152:ILE:N	2.52	0.42
1:M:205:VAL:O	1:M:209:LEU:CB	2.67	0.42
1:P:205:VAL:O	1:P:209:LEU:CB	2.67	0.42
1:U:205:VAL:O	1:U:209:LEU:CB	2.67	0.42
1:a:39:GLU:OE2	1:b:152:ILE:N	2.52	0.42
1:g:39:GLU:OE2	1:h:152:ILE:N	2.52	0.42
1:p:209:LEU:O	1:p:213:LYS:N	2.37	0.42
1:u:205:VAL:O	1:u:209:LEU:CB	2.67	0.42
1:v:39:GLU:OE2	1:w:152:ILE:N	2.52	0.42
1:0:39:GLU:OE2	1:1:152:ILE:N	2.52	0.42
1:3:205:VAL:O	1:3:209:LEU:CB	2.67	0.42
1:5:205:VAL:O	1:5:209:LEU:CB	2.67	0.42
1:A:39:GLU:OE2	1:B:152:ILE:N	2.52	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:205:VAL:O	1:B:209:LEU:CB	2.67	0.42
1:G:39:GLU:OE2	1:H:152:ILE:N	2.52	0.42
1:S:39:GLU:OE2	1:T:152:ILE:N	2.52	0.42
1:T:39:GLU:OE2	1:U:152:ILE:N	2.52	0.42
1:f:39:GLU:OE2	1:g:152:ILE:N	2.52	0.42
1:o:205:VAL:O	1:o:209:LEU:CB	2.67	0.42
1:q:93:GLN:O	1:q:96:SER:OG	2.35	0.42
1:2:205:VAL:O	1:2:209:LEU:CB	2.67	0.42
1:N:39:GLU:OE2	1:O:152:ILE:N	2.52	0.42
1:Q:205:VAL:O	1:Q:209:LEU:CB	2.67	0.42
1:c:205:VAL:O	1:c:209:LEU:CB	2.67	0.42
1:l:93:GLN:O	1:l:96:SER:OG	2.35	0.42
1:q:213:LYS:O	1:q:217:SER:OG	2.07	0.42
1:u:39:GLU:OE2	1:v:152:ILE:N	2.52	0.42
1:v:205:VAL:O	1:v:209:LEU:CB	2.67	0.42
1:K:39:GLU:OE2	1:L:152:ILE:N	2.52	0.42
1:R:205:VAL:O	1:R:209:LEU:CB	2.67	0.42
1:S:205:VAL:O	1:S:209:LEU:CB	2.67	0.42
1:Z:205:VAL:O	1:Z:209:LEU:CB	2.67	0.42
1:i:39:GLU:OE2	1:j:152:ILE:N	2.52	0.42
1:i:205:VAL:O	1:i:209:LEU:CB	2.67	0.42
1:q:205:VAL:O	1:q:209:LEU:CB	2.67	0.42
1:r:205:VAL:O	1:r:209:LEU:CB	2.67	0.42
1:w:39:GLU:OE2	1:x:152:ILE:N	2.52	0.42
1:x:205:VAL:O	1:x:209:LEU:CB	2.67	0.42
1:4:205:VAL:O	1:4:209:LEU:CB	2.67	0.42
1:C:39:GLU:OE2	1:D:152:ILE:N	2.52	0.42
1:u:163:SER:HG	1:u:164:ALA:H	1.68	0.42
1:Q:39:GLU:OE2	1:R:152:ILE:N	2.52	0.42
1:W:39:GLU:OE2	1:X:152:ILE:N	2.52	0.42
1:m:39:GLU:OE2	1:n:152:ILE:N	2.52	0.42
1:x:200:GLU:O	1:x:204:ALA:N	2.36	0.42
1:y:39:GLU:OE2	1:z:152:ILE:N	2.52	0.42
1:0:205:VAL:O	1:0:209:LEU:CB	2.67	0.42
1:E:39:GLU:OE2	1:F:152:ILE:N	2.52	0.42
1:E:205:VAL:O	1:E:209:LEU:CB	2.67	0.42
1:N:205:VAL:O	1:N:209:LEU:CB	2.67	0.42
1:j:205:VAL:O	1:j:209:LEU:CB	2.67	0.42
1:l:205:VAL:O	1:l:209:LEU:CB	2.67	0.42
1:B:93:GLN:O	1:B:96:SER:OG	2.35	0.42
1:1:201:GLU:HA	1:1:204:ALA:HB3	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:p:200:GLU:O	1:p:204:ALA:N	2.36	0.42
1:p:201:GLU:HA	1:p:204:ALA:HB3	2.02	0.42
1:E:201:GLU:HA	1:E:204:ALA:HB3	2.02	0.42
1:H:200:GLU:O	1:H:204:ALA:N	2.36	0.42
1:0:201:GLU:HA	1:0:204:ALA:HB3	2.02	0.41
1:F:201:GLU:HA	1:F:204:ALA:HB3	2.02	0.41
1:Q:201:GLU:HA	1:Q:204:ALA:HB3	2.02	0.41
1:d:201:GLU:HA	1:d:204:ALA:HB3	2.02	0.41
1:s:93:GLN:O	1:s:96:SER:OG	2.35	0.41
1:5:201:GLU:HA	1:5:204:ALA:HB3	2.02	0.41
1:B:201:GLU:HA	1:B:204:ALA:HB3	2.02	0.41
1:R:201:GLU:HA	1:R:204:ALA:HB3	2.02	0.41
1:x:201:GLU:HA	1:x:204:ALA:HB3	2.02	0.41
1:4:201:GLU:HA	1:4:204:ALA:HB3	2.02	0.41
1:A:201:GLU:HA	1:A:204:ALA:HB3	2.02	0.41
1:I:201:GLU:HA	1:I:204:ALA:HB3	2.02	0.41
1:q:201:GLU:HA	1:q:204:ALA:HB3	2.02	0.41
1:u:93:GLN:O	1:u:96:SER:OG	2.35	0.41
1:2:201:GLU:HA	1:2:204:ALA:HB3	2.02	0.41
1:O:200:GLU:O	1:O:204:ALA:N	2.36	0.41
1:a:213:LYS:O	1:a:217:SER:OG	2.07	0.41
1:o:201:GLU:HA	1:o:204:ALA:HB3	2.02	0.41
1:s:201:GLU:HA	1:s:204:ALA:HB3	2.02	0.41
1:L:200:GLU:O	1:L:204:ALA:N	2.36	0.41
1:P:201:GLU:HA	1:P:204:ALA:HB3	2.02	0.41
1:R:200:GLU:O	1:R:204:ALA:N	2.36	0.41
1:Y:200:GLU:O	1:Y:204:ALA:N	2.36	0.41
1:l:201:GLU:HA	1:l:204:ALA:HB3	2.02	0.41
1:y:162:ALA:H	1:y:166:SER:HG	1.67	0.41
1:G:201:GLU:HA	1:G:204:ALA:HB3	2.02	0.41
1:U:39:GLU:OE2	1:V:152:ILE:N	2.53	0.41
1:a:201:GLU:HA	1:a:204:ALA:HB3	2.02	0.41
1:m:201:GLU:HA	1:m:204:ALA:HB3	2.02	0.41
1:J:201:GLU:HA	1:J:204:ALA:HB3	2.02	0.41
1:P:93:GLN:O	1:P:96:SER:OG	2.35	0.41
1:U:201:GLU:HA	1:U:204:ALA:HB3	2.02	0.41
1:c:201:GLU:HA	1:c:204:ALA:HB3	2.02	0.41
1:d:159:LEU:O	1:d:165:SER:OG	2.24	0.41
1:u:213:LYS:O	1:u:217:SER:OG	2.07	0.41
1:0:159:LEU:O	1:0:165:SER:OG	2.24	0.41
1:1:59:ARG:O	1:1:63:ALA:N	2.53	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1:159:LEU:O	1:1:165:SER:OG	2.24	0.41
1:2:59:ARG:O	1:2:63:ALA:N	2.53	0.41
1:3:59:ARG:O	1:3:63:ALA:N	2.53	0.41
1:4:59:ARG:O	1:4:63:ALA:N	2.53	0.41
1:C:201:GLU:HA	1:C:204:ALA:HB3	2.02	0.41
1:D:201:GLU:HA	1:D:204:ALA:HB3	2.02	0.41
1:R:59:ARG:O	1:R:63:ALA:N	2.53	0.41
1:S:59:ARG:O	1:S:63:ALA:N	2.53	0.41
1:T:201:GLU:HA	1:T:204:ALA:HB3	2.02	0.41
1:a:159:LEU:O	1:a:165:SER:OG	2.24	0.41
1:f:201:GLU:HA	1:f:204:ALA:HB3	2.02	0.41
1:u:201:GLU:HA	1:u:204:ALA:HB3	2.02	0.41
1:y:201:GLU:HA	1:y:204:ALA:HB3	2.02	0.41
1:1:179:GLU:OE1	1:o:55:LYS:NZ	2.55	0.41
1:2:159:LEU:O	1:2:165:SER:OG	2.24	0.41
1:2:179:GLU:OE1	1:p:55:LYS:NZ	2.54	0.41
1:3:159:LEU:O	1:3:165:SER:OG	2.24	0.41
1:C:179:GLU:OE1	1:x:55:LYS:NZ	2.55	0.41
1:D:93:GLN:O	1:D:96:SER:OG	2.35	0.41
1:D:179:GLU:OE1	1:y:55:LYS:NZ	2.54	0.41
1:H:55:LYS:NZ	1:U:179:GLU:OE1	2.55	0.41
1:L:201:GLU:HA	1:L:204:ALA:HB3	2.02	0.41
1:M:55:LYS:NZ	1:Z:179:GLU:OE1	2.54	0.41
1:N:55:LYS:NZ	1:a:179:GLU:OE1	2.55	0.41
1:O:55:LYS:NZ	1:b:179:GLU:OE1	2.55	0.41
1:O:201:GLU:HA	1:O:204:ALA:HB3	2.02	0.41
1:P:59:ARG:O	1:P:63:ALA:N	2.53	0.41
1:w:201:GLU:HA	1:w:204:ALA:HB3	2.02	0.41
1:z:159:LEU:O	1:z:165:SER:OG	2.24	0.41
1:z:201:GLU:HA	1:z:204:ALA:HB3	2.02	0.41
1:A:213:LYS:O	1:A:217:SER:OG	2.07	0.40
1:C:3:LEU:HD23	1:C:3:LEU:HA	1.98	0.40
1:G:55:LYS:NZ	1:T:179:GLU:OE1	2.55	0.40
1:H:158:ASN:C	1:H:163:SER:HG	2.24	0.40
1:H:201:GLU:HA	1:H:204:ALA:HB3	2.02	0.40
1:I:55:LYS:NZ	1:V:179:GLU:OE1	2.54	0.40
1:J:55:LYS:NZ	1:W:179:GLU:OE1	2.55	0.40
1:K:55:LYS:NZ	1:X:179:GLU:OE1	2.55	0.40
1:L:55:LYS:NZ	1:Y:179:GLU:OE1	2.54	0.40
1:Q:59:ARG:O	1:Q:63:ALA:N	2.53	0.40
1:X:201:GLU:HA	1:X:204:ALA:HB3	2.02	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:4:159:LEU:O	1:4:165:SER:OG	2.25	0.40
1:5:159:LEU:O	1:5:165:SER:OG	2.24	0.40
1:B:179:GLU:OE1	1:w:55:LYS:NZ	2.55	0.40
1:F:55:LYS:NZ	1:S:179:GLU:OE1	2.54	0.40
1:S:201:GLU:HA	1:S:204:ALA:HB3	2.02	0.40
1:W:163:SER:OG	1:W:164:ALA:N	2.54	0.40
1:g:201:GLU:HA	1:g:204:ALA:HB3	2.02	0.40
1:i:201:GLU:HA	1:i:204:ALA:HB3	2.02	0.40
1:y:159:LEU:O	1:y:165:SER:OG	2.24	0.40
1:0:179:GLU:OE1	1:n:55:LYS:NZ	2.55	0.40
1:3:179:GLU:OE1	1:q:55:LYS:NZ	2.55	0.40
1:4:55:LYS:NZ	1:J:179:GLU:OE1	2.54	0.40
1:5:55:LYS:NZ	1:K:179:GLU:OE1	2.55	0.40
1:6:55:LYS:NZ	1:L:179:GLU:OE1	2.55	0.40
1:7:55:LYS:NZ	1:M:179:GLU:OE1	2.55	0.40
1:7:201:GLU:HA	1:7:204:ALA:HB3	2.02	0.40
1:A:179:GLU:OE1	1:v:55:LYS:NZ	2.55	0.40
1:E:55:LYS:NZ	1:R:179:GLU:OE1	2.54	0.40
1:E:179:GLU:OE1	1:z:55:LYS:NZ	2.55	0.40
1:N:163:SER:OG	1:N:164:ALA:N	2.54	0.40
1:O:163:SER:OG	1:O:164:ALA:N	2.54	0.40
1:P:159:LEU:O	1:P:165:SER:OG	2.24	0.40
1:Q:159:LEU:O	1:Q:165:SER:OG	2.24	0.40
1:b:201:GLU:HA	1:b:204:ALA:HB3	2.02	0.40
1:j:163:SER:OG	1:j:164:ALA:N	2.54	0.40
1:r:201:GLU:HA	1:r:204:ALA:HB3	2.02	0.40
1:z:93:GLN:O	1:z:96:SER:OG	2.35	0.40
1:6:159:LEU:O	1:6:165:SER:OG	2.25	0.40
1:6:201:GLU:HA	1:6:204:ALA:HB3	2.02	0.40
1:C:205:VAL:HG23	1:C:206:GLU:H	1.87	0.40
1:M:201:GLU:HA	1:M:204:ALA:HB3	2.02	0.40
1:R:55:LYS:NZ	1:e:179:GLU:OE1	2.55	0.40
1:R:159:LEU:O	1:R:165:SER:OG	2.24	0.40
1:S:55:LYS:NZ	1:f:179:GLU:OE1	2.55	0.40
1:X:55:LYS:NZ	1:k:179:GLU:OE1	2.55	0.40
1:X:163:SER:OG	1:X:164:ALA:N	2.54	0.40
1:Y:55:LYS:NZ	1:l:179:GLU:OE1	2.55	0.40
1:Z:201:GLU:HA	1:Z:204:ALA:HB3	2.02	0.40
1:3:201:GLU:HA	1:3:204:ALA:HB3	2.02	0.40
1:4:205:VAL:HG23	1:4:206:GLU:H	1.87	0.40
1:7:204:ALA:O	1:7:208:THR:OG1	2.14	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:205:VAL:HG23	1:F:206:GLU:H	1.87	0.40
1:S:159:LEU:O	1:S:165:SER:OG	2.24	0.40
1:T:55:LYS:NZ	1:g:179:GLU:OE1	2.54	0.40
1:U:55:LYS:NZ	1:h:179:GLU:OE1	2.54	0.40
1:V:55:LYS:NZ	1:i:179:GLU:OE1	2.54	0.40
1:W:55:LYS:NZ	1:j:179:GLU:OE1	2.55	0.40
1:Z:55:LYS:NZ	1:m:179:GLU:OE1	2.55	0.40
1:b:159:LEU:O	1:b:165:SER:OG	2.24	0.40
1:i:163:SER:OG	1:i:164:ALA:N	2.54	0.40
1:j:201:GLU:HA	1:j:204:ALA:HB3	2.02	0.40
1:k:163:SER:OG	1:k:164:ALA:N	2.54	0.40

There are no symmetry-related clashes.

## 5.3 Torsion angles [i](#)

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

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	0	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	1	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	2	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	3	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	4	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	5	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	6	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	7	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	A	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	B	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	C	215/246 (87%)	212 (99%)	3 (1%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	D	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	E	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	F	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	G	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	H	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	I	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	J	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	K	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	L	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	M	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	N	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	O	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	P	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	Q	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	R	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	S	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	T	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	U	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	V	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	W	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	X	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	Y	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	Z	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	a	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	b	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	c	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	d	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	e	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	f	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	g	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	h	215/246 (87%)	212 (99%)	3 (1%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	i	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	j	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	k	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	l	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	m	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	n	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	o	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	p	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	q	190/246 (77%)	187 (98%)	3 (2%)	0	100	100
1	r	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	s	52/246 (21%)	50 (96%)	2 (4%)	0	100	100
1	t	113/246 (46%)	112 (99%)	1 (1%)	0	100	100
1	u	164/246 (67%)	162 (99%)	2 (1%)	0	100	100
1	v	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	w	191/246 (78%)	189 (99%)	2 (1%)	0	100	100
1	x	66/246 (27%)	64 (97%)	2 (3%)	0	100	100
1	y	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
1	z	215/246 (87%)	212 (99%)	3 (1%)	0	100	100
All	All	12386/14760 (84%)	12212 (99%)	174 (1%)	0	100	100

There are no Ramachandran outliers to report.

### 5.3.2 Protein sidechains ⓘ

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	0	183/208 (88%)	183 (100%)	0	100	100
1	1	183/208 (88%)	183 (100%)	0	100	100
1	2	183/208 (88%)	183 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	3	183/208 (88%)	183 (100%)	0	100	100
1	4	183/208 (88%)	183 (100%)	0	100	100
1	5	183/208 (88%)	183 (100%)	0	100	100
1	6	183/208 (88%)	183 (100%)	0	100	100
1	7	183/208 (88%)	183 (100%)	0	100	100
1	A	183/208 (88%)	183 (100%)	0	100	100
1	B	183/208 (88%)	183 (100%)	0	100	100
1	C	183/208 (88%)	183 (100%)	0	100	100
1	D	183/208 (88%)	183 (100%)	0	100	100
1	E	183/208 (88%)	183 (100%)	0	100	100
1	F	183/208 (88%)	183 (100%)	0	100	100
1	G	183/208 (88%)	183 (100%)	0	100	100
1	H	183/208 (88%)	183 (100%)	0	100	100
1	I	183/208 (88%)	183 (100%)	0	100	100
1	J	183/208 (88%)	183 (100%)	0	100	100
1	K	183/208 (88%)	183 (100%)	0	100	100
1	L	183/208 (88%)	183 (100%)	0	100	100
1	M	183/208 (88%)	183 (100%)	0	100	100
1	N	183/208 (88%)	183 (100%)	0	100	100
1	O	183/208 (88%)	183 (100%)	0	100	100
1	P	183/208 (88%)	183 (100%)	0	100	100
1	Q	183/208 (88%)	183 (100%)	0	100	100
1	R	183/208 (88%)	183 (100%)	0	100	100
1	S	183/208 (88%)	183 (100%)	0	100	100
1	T	183/208 (88%)	183 (100%)	0	100	100
1	U	183/208 (88%)	183 (100%)	0	100	100
1	V	183/208 (88%)	183 (100%)	0	100	100
1	W	183/208 (88%)	183 (100%)	0	100	100
1	X	183/208 (88%)	183 (100%)	0	100	100
1	Y	183/208 (88%)	183 (100%)	0	100	100
1	Z	183/208 (88%)	183 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	a	183/208 (88%)	183 (100%)	0	100	100
1	b	183/208 (88%)	183 (100%)	0	100	100
1	c	183/208 (88%)	183 (100%)	0	100	100
1	d	183/208 (88%)	183 (100%)	0	100	100
1	e	183/208 (88%)	183 (100%)	0	100	100
1	f	183/208 (88%)	183 (100%)	0	100	100
1	g	183/208 (88%)	183 (100%)	0	100	100
1	h	183/208 (88%)	183 (100%)	0	100	100
1	i	183/208 (88%)	183 (100%)	0	100	100
1	j	183/208 (88%)	183 (100%)	0	100	100
1	k	183/208 (88%)	183 (100%)	0	100	100
1	l	183/208 (88%)	183 (100%)	0	100	100
1	m	183/208 (88%)	183 (100%)	0	100	100
1	n	183/208 (88%)	183 (100%)	0	100	100
1	o	183/208 (88%)	183 (100%)	0	100	100
1	p	183/208 (88%)	183 (100%)	0	100	100
1	q	183/208 (88%)	183 (100%)	0	100	100
1	r	183/208 (88%)	183 (100%)	0	100	100
1	s	183/208 (88%)	183 (100%)	0	100	100
1	t	183/208 (88%)	183 (100%)	0	100	100
1	u	183/208 (88%)	183 (100%)	0	100	100
1	v	183/208 (88%)	183 (100%)	0	100	100
1	w	183/208 (88%)	183 (100%)	0	100	100
1	x	183/208 (88%)	183 (100%)	0	100	100
1	y	183/208 (88%)	183 (100%)	0	100	100
1	z	183/208 (88%)	183 (100%)	0	100	100
All	All	10980/12480 (88%)	10980 (100%)	0	100	100

There are no protein residues with a non-rotameric sidechain to report.

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (76) such sidechains are listed below:

Mol	Chain	Res	Type
1	0	98	GLN
1	1	98	GLN
1	1	202	GLN
1	2	98	GLN
1	3	98	GLN
1	4	98	GLN
1	5	98	GLN
1	5	202	GLN
1	6	98	GLN
1	7	98	GLN
1	7	202	GLN
1	A	98	GLN
1	B	98	GLN
1	B	202	GLN
1	C	98	GLN
1	C	202	GLN
1	D	98	GLN
1	E	98	GLN
1	F	98	GLN
1	G	98	GLN
1	H	98	GLN
1	I	98	GLN
1	J	98	GLN
1	K	98	GLN
1	L	98	GLN
1	M	98	GLN
1	N	98	GLN
1	O	98	GLN
1	P	98	GLN
1	Q	98	GLN
1	R	98	GLN
1	S	98	GLN
1	T	98	GLN
1	U	98	GLN
1	V	98	GLN
1	W	98	GLN
1	X	98	GLN
1	Y	98	GLN
1	Y	202	GLN
1	Z	98	GLN
1	Z	202	GLN
1	a	98	GLN
1	b	98	GLN

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Mol	Chain	Res	Type
1	c	98	GLN
1	c	202	GLN
1	d	98	GLN
1	d	202	GLN
1	e	98	GLN
1	f	98	GLN
1	g	98	GLN
1	g	123	GLN
1	h	98	GLN
1	h	123	GLN
1	i	98	GLN
1	j	98	GLN
1	j	202	GLN
1	k	98	GLN
1	l	98	GLN
1	m	98	GLN
1	n	98	GLN
1	n	202	GLN
1	o	98	GLN
1	p	98	GLN
1	p	202	GLN
1	q	98	GLN
1	r	98	GLN
1	r	202	GLN
1	s	98	GLN
1	t	98	GLN
1	u	98	GLN
1	v	98	GLN
1	w	98	GLN
1	x	98	GLN
1	y	98	GLN
1	z	98	GLN
1	z	202	GLN

### 5.3.3 RNA ⓘ

There are no RNA molecules in this entry.

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

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

## 5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

## 5.6 Ligand geometry [i](#)

There are no ligands in this entry.

## 5.7 Other polymers [i](#)

There are no such residues in this entry.

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

There are no chain breaks in this entry.