



Full wwPDB EM Validation Report ⓘ

Aug 29, 2026 – 01:03 PM EDT

PDB ID : 11SR / pdb_000011sr
EMDB ID : EMD-76018
Title : EcMscM lacking the first periplasmic helical bundle and the cytoplasmic extension of TM7 in NaCl in a closed conformation
Authors : Hiotis, G.; Walz, T.
Deposited on : 2026-03-11
Resolution : 5.10 Å(reported)
Based on initial model : .

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

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

A user guide is available at

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

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

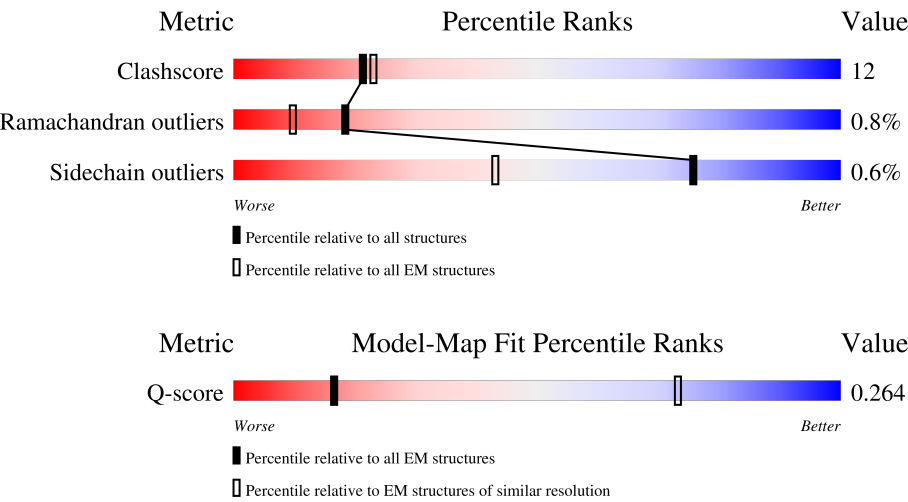
EMDB validation analysis : 0.0.1.dev133
MolProbity : 4-5-2 with Phenix2.0
Percentile statistics : 20250101.v01 (using entries in the PDB archive January 1st 2025)
EM percentile statistics : 202505.v01 (Using data in the EMDb archive up until May 2025)
MapQ : 1.9.13
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.50

1 Overall quality at a glance i

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

The reported resolution of this entry is 5.10 Å.

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



Metric	Whole archive (#Entries)	EM structures (#Entries)	Similar EM resolution (#Entries, resolution range(Å))
Clashscore	229148	23984	-
Ramachandran outliers	224038	23583	-
Sidechain outliers	223484	23102	-
Q-score	-	25397	844 (4.60 - 5.60)

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 ≥ 3 , 2, 1 and 0 types of geometric quality criteria respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions $\leq 5\%$. The upper red bar (where present) indicates the fraction of residues that have poor fit to the EM map (all-atom inclusion $< 40\%$). The numeric value is given above the bar.

Mol	Chain	Length	Quality of chain
1	A	873	8% 79% 11% 9%
1	B	873	9% 79% 11% 9%
1	C	873	8% 79% 11% 9%
1	D	873	8% 79% 11% 9%

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Mol	Chain	Length	Quality of chain
1	E	873	8% 79% 11% 9%
1	F	873	9% 79% 11% 9%
1	G	873	8% 79% 11% 9%

2 Entry composition

There is only 1 type of molecule in this entry. The entry contains 32277 atoms, of which 0 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a protein called Miniconductance mechanosensitive channel MscM.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	A	791	Total 4611	C 2857	N 863	O 884	S 7	0	0
1	B	791	Total 4611	C 2857	N 863	O 884	S 7	0	0
1	C	791	Total 4611	C 2857	N 863	O 884	S 7	0	0
1	D	791	Total 4611	C 2857	N 863	O 884	S 7	0	0
1	E	791	Total 4611	C 2857	N 863	O 884	S 7	0	0
1	F	791	Total 4611	C 2857	N 863	O 884	S 7	0	0
1	G	791	Total 4611	C 2857	N 863	O 884	S 7	0	0

There are 1729 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
A	?	-	ALA	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	PRO	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	VAL	deletion	UNP P39285
A	?	-	VAL	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
A	?	-	GLU	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	LYS	deletion	UNP P39285
A	?	-	GLY	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	ILE	deletion	UNP P39285
A	?	-	LYS	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	TYR	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	VAL	deletion	UNP P39285
A	?	-	ILE	deletion	UNP P39285
A	?	-	ASP	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	TYR	deletion	UNP P39285
A	?	-	PRO	deletion	UNP P39285
A	?	-	LYS	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	THR	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	MET	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	ASP	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	PRO	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	VAL	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
A	?	-	PRO	deletion	UNP P39285
A	?	-	GLY	deletion	UNP P39285
A	?	-	MET	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	THR	deletion	UNP P39285
A	?	-	ASP	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	ILE	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	VAL	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ASP	deletion	UNP P39285
A	?	-	LYS	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	ILE	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	ASP	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
A	?	-	LEU	deletion	UNP P39285
A	?	-	PRO	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	THR	deletion	UNP P39285
A	?	-	ASP	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	ILE	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	GLY	deletion	UNP P39285
A	?	-	THR	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	THR	deletion	UNP P39285
A	?	-	GLY	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	THR	deletion	UNP P39285
A	?	-	PRO	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	PHE	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	ASP	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
A	?	-	LEU	deletion	UNP P39285
A	?	-	LYS	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	VAL	deletion	UNP P39285
A	?	-	ASP	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	LYS	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ASP	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	TYR	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
A	?	-	LEU	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	THR	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	ASN	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	487	CYS	GLY	conflict	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	MET	deletion	UNP P39285
A	?	-	LEU	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	GLN	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	ARG	deletion	UNP P39285
A	?	-	GLY	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
A	?	-	HIS	deletion	UNP P39285
A	?	-	HIS	deletion	UNP P39285
A	?	-	HIS	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	?	-	PRO	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	GLY	deletion	UNP P39285
A	?	-	ALA	deletion	UNP P39285
A	?	-	ILE	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	VAL	deletion	UNP P39285
A	?	-	ASP	deletion	UNP P39285
A	?	-	GLU	deletion	UNP P39285
A	?	-	SER	deletion	UNP P39285
A	1108	LEU	-	expression tag	UNP P39285
A	1109	GLU	-	expression tag	UNP P39285
A	1110	VAL	-	expression tag	UNP P39285
A	1111	LEU	-	expression tag	UNP P39285
A	1112	PHE	-	expression tag	UNP P39285
A	1113	GLN	-	expression tag	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	PRO	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	VAL	deletion	UNP P39285
B	?	-	VAL	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ASN	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	LYS	deletion	UNP P39285
B	?	-	GLY	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
B	?	-	SER	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	ILE	deletion	UNP P39285
B	?	-	LYS	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	TYR	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	VAL	deletion	UNP P39285
B	?	-	ILE	deletion	UNP P39285
B	?	-	ASP	deletion	UNP P39285
B	?	-	ASN	deletion	UNP P39285
B	?	-	TYR	deletion	UNP P39285
B	?	-	PRO	deletion	UNP P39285
B	?	-	LYS	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	THR	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ASN	deletion	UNP P39285
B	?	-	ASN	deletion	UNP P39285
B	?	-	MET	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	ASP	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	PRO	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	VAL	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	PRO	deletion	UNP P39285
B	?	-	GLY	deletion	UNP P39285
B	?	-	MET	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	THR	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
B	?	-	ASP	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ASN	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	ILE	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	VAL	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ASP	deletion	UNP P39285
B	?	-	LYS	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	ILE	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	ASP	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ASN	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	PRO	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
B	?	-	THR	deletion	UNP P39285
B	?	-	ASP	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ASN	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	ILE	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	GLY	deletion	UNP P39285
B	?	-	THR	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	THR	deletion	UNP P39285
B	?	-	GLY	deletion	UNP P39285
B	?	-	ASN	deletion	UNP P39285
B	?	-	THR	deletion	UNP P39285
B	?	-	PRO	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ASN	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	ASN	deletion	UNP P39285
B	?	-	PHE	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	ASP	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	LYS	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	VAL	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
B	?	-	ASP	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	ASN	deletion	UNP P39285
B	?	-	ASN	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	LYS	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ASP	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	TYR	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	ASN	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
B	?	-	ASN	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	THR	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	ASN	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	487	CYS	GLY	conflict	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	MET	deletion	UNP P39285
B	?	-	LEU	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	GLN	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	ARG	deletion	UNP P39285
B	?	-	GLY	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	HIS	deletion	UNP P39285
B	?	-	HIS	deletion	UNP P39285
B	?	-	HIS	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
B	?	-	PRO	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	GLY	deletion	UNP P39285
B	?	-	ALA	deletion	UNP P39285
B	?	-	ILE	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	VAL	deletion	UNP P39285
B	?	-	ASP	deletion	UNP P39285
B	?	-	GLU	deletion	UNP P39285
B	?	-	SER	deletion	UNP P39285
B	1108	LEU	-	expression tag	UNP P39285
B	1109	GLU	-	expression tag	UNP P39285
B	1110	VAL	-	expression tag	UNP P39285
B	1111	LEU	-	expression tag	UNP P39285
B	1112	PHE	-	expression tag	UNP P39285
B	1113	GLN	-	expression tag	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	PRO	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	VAL	deletion	UNP P39285
C	?	-	VAL	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	LYS	deletion	UNP P39285
C	?	-	GLY	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	ILE	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
C	?	-	LYS	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	TYR	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	VAL	deletion	UNP P39285
C	?	-	ILE	deletion	UNP P39285
C	?	-	ASP	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	TYR	deletion	UNP P39285
C	?	-	PRO	deletion	UNP P39285
C	?	-	LYS	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	THR	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	MET	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	ASP	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	PRO	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	VAL	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	PRO	deletion	UNP P39285
C	?	-	GLY	deletion	UNP P39285
C	?	-	MET	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	THR	deletion	UNP P39285
C	?	-	ASP	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
C	?	-	GLU	deletion	UNP P39285
C	?	-	ILE	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	VAL	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ASP	deletion	UNP P39285
C	?	-	LYS	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	ILE	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	ASP	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	PRO	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	THR	deletion	UNP P39285
C	?	-	ASP	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
C	?	-	GLN	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	ILE	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	GLY	deletion	UNP P39285
C	?	-	THR	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	THR	deletion	UNP P39285
C	?	-	GLY	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	THR	deletion	UNP P39285
C	?	-	PRO	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	PHE	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	ASP	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	LYS	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	VAL	deletion	UNP P39285
C	?	-	ASP	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
C	?	-	ALA	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	LYS	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ASP	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	TYR	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
C	?	-	LEU	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	THR	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	ASN	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	487	CYS	GLY	conflict	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	MET	deletion	UNP P39285
C	?	-	LEU	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	GLN	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	ARG	deletion	UNP P39285
C	?	-	GLY	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	HIS	deletion	UNP P39285
C	?	-	HIS	deletion	UNP P39285
C	?	-	HIS	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	?	-	PRO	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	GLY	deletion	UNP P39285
C	?	-	ALA	deletion	UNP P39285
C	?	-	ILE	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
C	?	-	GLU	deletion	UNP P39285
C	?	-	VAL	deletion	UNP P39285
C	?	-	ASP	deletion	UNP P39285
C	?	-	GLU	deletion	UNP P39285
C	?	-	SER	deletion	UNP P39285
C	1108	LEU	-	expression tag	UNP P39285
C	1109	GLU	-	expression tag	UNP P39285
C	1110	VAL	-	expression tag	UNP P39285
C	1111	LEU	-	expression tag	UNP P39285
C	1112	PHE	-	expression tag	UNP P39285
C	1113	GLN	-	expression tag	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	PRO	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	VAL	deletion	UNP P39285
D	?	-	VAL	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ASN	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	LYS	deletion	UNP P39285
D	?	-	GLY	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	ILE	deletion	UNP P39285
D	?	-	LYS	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	TYR	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
D	?	-	VAL	deletion	UNP P39285
D	?	-	ILE	deletion	UNP P39285
D	?	-	ASP	deletion	UNP P39285
D	?	-	ASN	deletion	UNP P39285
D	?	-	TYR	deletion	UNP P39285
D	?	-	PRO	deletion	UNP P39285
D	?	-	LYS	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	THR	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ASN	deletion	UNP P39285
D	?	-	ASN	deletion	UNP P39285
D	?	-	MET	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	ASP	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	PRO	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	VAL	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	PRO	deletion	UNP P39285
D	?	-	GLY	deletion	UNP P39285
D	?	-	MET	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	THR	deletion	UNP P39285
D	?	-	ASP	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ASN	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	ILE	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	VAL	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
D	?	-	SER	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ASP	deletion	UNP P39285
D	?	-	LYS	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	ILE	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	ASP	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ASN	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	PRO	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	THR	deletion	UNP P39285
D	?	-	ASP	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ASN	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	ILE	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
D	?	-	GLU	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	GLY	deletion	UNP P39285
D	?	-	THR	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	THR	deletion	UNP P39285
D	?	-	GLY	deletion	UNP P39285
D	?	-	ASN	deletion	UNP P39285
D	?	-	THR	deletion	UNP P39285
D	?	-	PRO	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ASN	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	ASN	deletion	UNP P39285
D	?	-	PHE	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	ASP	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	LYS	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	VAL	deletion	UNP P39285
D	?	-	ASP	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
D	?	-	ASN	deletion	UNP P39285
D	?	-	ASN	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	LYS	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ASP	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	TYR	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	ASN	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ASN	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
D	?	-	ALA	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	THR	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	ASN	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	487	CYS	GLY	conflict	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	MET	deletion	UNP P39285
D	?	-	LEU	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	GLN	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	ARG	deletion	UNP P39285
D	?	-	GLY	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	HIS	deletion	UNP P39285
D	?	-	HIS	deletion	UNP P39285
D	?	-	HIS	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285
D	?	-	PRO	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	GLY	deletion	UNP P39285
D	?	-	ALA	deletion	UNP P39285
D	?	-	ILE	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	VAL	deletion	UNP P39285
D	?	-	ASP	deletion	UNP P39285
D	?	-	GLU	deletion	UNP P39285
D	?	-	SER	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
D	1108	LEU	-	expression tag	UNP P39285
D	1109	GLU	-	expression tag	UNP P39285
D	1110	VAL	-	expression tag	UNP P39285
D	1111	LEU	-	expression tag	UNP P39285
D	1112	PHE	-	expression tag	UNP P39285
D	1113	GLN	-	expression tag	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	PRO	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	VAL	deletion	UNP P39285
E	?	-	VAL	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	LYS	deletion	UNP P39285
E	?	-	GLY	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	ILE	deletion	UNP P39285
E	?	-	LYS	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	TYR	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	VAL	deletion	UNP P39285
E	?	-	ILE	deletion	UNP P39285
E	?	-	ASP	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	TYR	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
E	?	-	PRO	deletion	UNP P39285
E	?	-	LYS	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	THR	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	MET	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	ASP	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	PRO	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	VAL	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	PRO	deletion	UNP P39285
E	?	-	GLY	deletion	UNP P39285
E	?	-	MET	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	THR	deletion	UNP P39285
E	?	-	ASP	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	ILE	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	VAL	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
E	?	-	ASP	deletion	UNP P39285
E	?	-	LYS	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	ILE	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	ASP	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	PRO	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	THR	deletion	UNP P39285
E	?	-	ASP	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	ILE	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	GLY	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
E	?	-	THR	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	THR	deletion	UNP P39285
E	?	-	GLY	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	THR	deletion	UNP P39285
E	?	-	PRO	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	PHE	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	ASP	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	LYS	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	VAL	deletion	UNP P39285
E	?	-	ASP	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
E	?	-	LEU	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	LYS	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ASP	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	TYR	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	THR	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
E	?	-	GLU	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	ASN	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	487	CYS	GLY	conflict	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	MET	deletion	UNP P39285
E	?	-	LEU	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	GLN	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	ARG	deletion	UNP P39285
E	?	-	GLY	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	HIS	deletion	UNP P39285
E	?	-	HIS	deletion	UNP P39285
E	?	-	HIS	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	?	-	PRO	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	GLY	deletion	UNP P39285
E	?	-	ALA	deletion	UNP P39285
E	?	-	ILE	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	VAL	deletion	UNP P39285
E	?	-	ASP	deletion	UNP P39285
E	?	-	GLU	deletion	UNP P39285
E	?	-	SER	deletion	UNP P39285
E	1108	LEU	-	expression tag	UNP P39285
E	1109	GLU	-	expression tag	UNP P39285
E	1110	VAL	-	expression tag	UNP P39285
E	1111	LEU	-	expression tag	UNP P39285
E	1112	PHE	-	expression tag	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
E	1113	GLN	-	expression tag	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	PRO	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	VAL	deletion	UNP P39285
F	?	-	VAL	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ASN	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	LYS	deletion	UNP P39285
F	?	-	GLY	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	ILE	deletion	UNP P39285
F	?	-	LYS	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	TYR	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	VAL	deletion	UNP P39285
F	?	-	ILE	deletion	UNP P39285
F	?	-	ASP	deletion	UNP P39285
F	?	-	ASN	deletion	UNP P39285
F	?	-	TYR	deletion	UNP P39285
F	?	-	PRO	deletion	UNP P39285
F	?	-	LYS	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
F	?	-	THR	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ASN	deletion	UNP P39285
F	?	-	ASN	deletion	UNP P39285
F	?	-	MET	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	ASP	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	PRO	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	VAL	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	PRO	deletion	UNP P39285
F	?	-	GLY	deletion	UNP P39285
F	?	-	MET	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	THR	deletion	UNP P39285
F	?	-	ASP	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ASN	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	ILE	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	VAL	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ASP	deletion	UNP P39285
F	?	-	LYS	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
F	?	-	ALA	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	ILE	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	ASP	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ASN	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	PRO	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	THR	deletion	UNP P39285
F	?	-	ASP	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ASN	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	ILE	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	GLY	deletion	UNP P39285
F	?	-	THR	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	THR	deletion	UNP P39285
F	?	-	GLY	deletion	UNP P39285
F	?	-	ASN	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
F	?	-	THR	deletion	UNP P39285
F	?	-	PRO	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ASN	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	ASN	deletion	UNP P39285
F	?	-	PHE	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	ASP	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	LYS	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	VAL	deletion	UNP P39285
F	?	-	ASP	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	ASN	deletion	UNP P39285
F	?	-	ASN	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
F	?	-	SER	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	LYS	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ASP	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	TYR	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	ASN	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ASN	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	THR	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
F	?	-	ASN	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	487	CYS	GLY	conflict	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	MET	deletion	UNP P39285
F	?	-	LEU	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	GLN	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	ARG	deletion	UNP P39285
F	?	-	GLY	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	HIS	deletion	UNP P39285
F	?	-	HIS	deletion	UNP P39285
F	?	-	HIS	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	?	-	PRO	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	GLY	deletion	UNP P39285
F	?	-	ALA	deletion	UNP P39285
F	?	-	ILE	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	VAL	deletion	UNP P39285
F	?	-	ASP	deletion	UNP P39285
F	?	-	GLU	deletion	UNP P39285
F	?	-	SER	deletion	UNP P39285
F	1108	LEU	-	expression tag	UNP P39285
F	1109	GLU	-	expression tag	UNP P39285
F	1110	VAL	-	expression tag	UNP P39285
F	1111	LEU	-	expression tag	UNP P39285
F	1112	PHE	-	expression tag	UNP P39285
F	1113	GLN	-	expression tag	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	PRO	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
G	?	-	VAL	deletion	UNP P39285
G	?	-	VAL	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	LYS	deletion	UNP P39285
G	?	-	GLY	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	ILE	deletion	UNP P39285
G	?	-	LYS	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	TYR	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	VAL	deletion	UNP P39285
G	?	-	ILE	deletion	UNP P39285
G	?	-	ASP	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	TYR	deletion	UNP P39285
G	?	-	PRO	deletion	UNP P39285
G	?	-	LYS	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	THR	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
G	?	-	LEU	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	MET	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	ASP	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	PRO	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	VAL	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	PRO	deletion	UNP P39285
G	?	-	GLY	deletion	UNP P39285
G	?	-	MET	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	THR	deletion	UNP P39285
G	?	-	ASP	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	ILE	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	VAL	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ASP	deletion	UNP P39285
G	?	-	LYS	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
G	?	-	GLU	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	ILE	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	ASP	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	PRO	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	THR	deletion	UNP P39285
G	?	-	ASP	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	ILE	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	GLY	deletion	UNP P39285
G	?	-	THR	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	THR	deletion	UNP P39285
G	?	-	GLY	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	THR	deletion	UNP P39285
G	?	-	PRO	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
G	?	-	ALA	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	PHE	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	ASP	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	LYS	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	VAL	deletion	UNP P39285
G	?	-	ASP	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285

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Chain	Residue	Modelled	Actual	Comment	Reference
G	?	-	LYS	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ASP	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	TYR	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	THR	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	ASN	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	487	CYS	GLY	conflict	UNP P39285
G	?	-	GLU	deletion	UNP P39285

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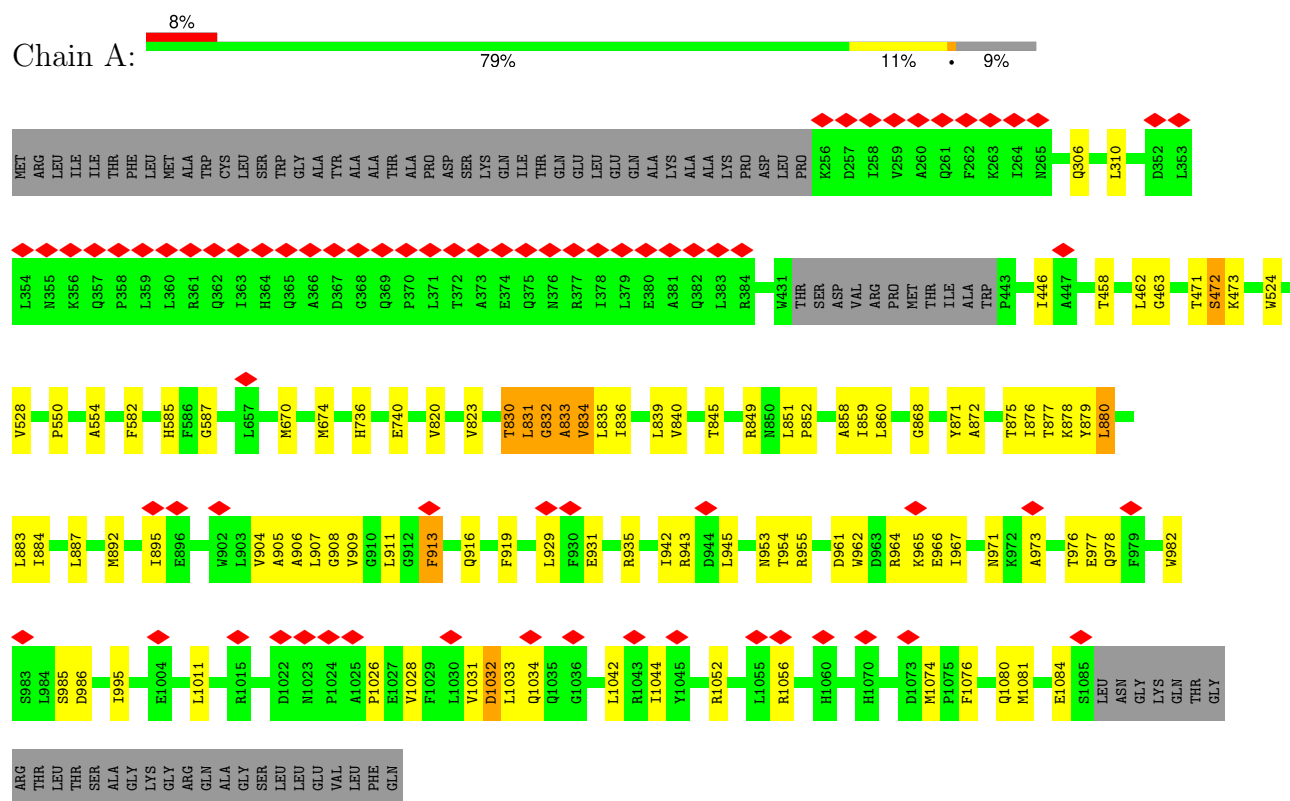
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Chain	Residue	Modelled	Actual	Comment	Reference
G	?	-	MET	deletion	UNP P39285
G	?	-	LEU	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	GLN	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	ARG	deletion	UNP P39285
G	?	-	GLY	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	HIS	deletion	UNP P39285
G	?	-	HIS	deletion	UNP P39285
G	?	-	HIS	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	?	-	PRO	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	GLY	deletion	UNP P39285
G	?	-	ALA	deletion	UNP P39285
G	?	-	ILE	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	VAL	deletion	UNP P39285
G	?	-	ASP	deletion	UNP P39285
G	?	-	GLU	deletion	UNP P39285
G	?	-	SER	deletion	UNP P39285
G	1108	LEU	-	expression tag	UNP P39285
G	1109	GLU	-	expression tag	UNP P39285
G	1110	VAL	-	expression tag	UNP P39285
G	1111	LEU	-	expression tag	UNP P39285
G	1112	PHE	-	expression tag	UNP P39285
G	1113	GLN	-	expression tag	UNP P39285

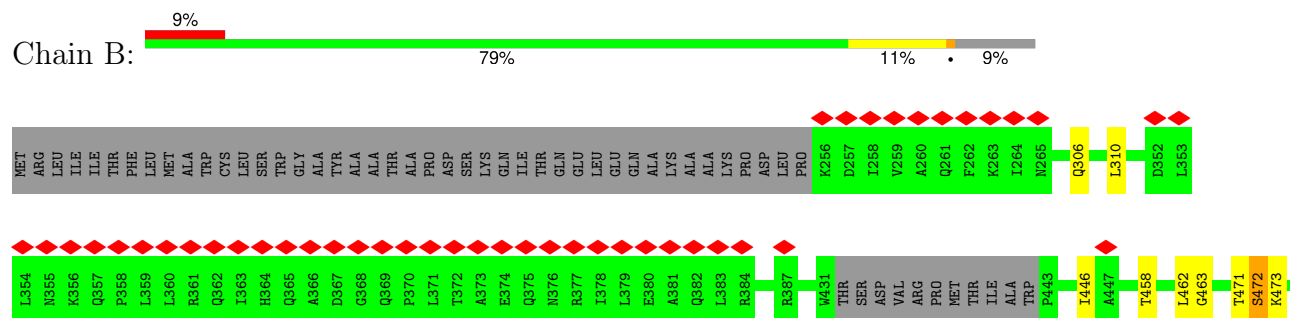
3 Residue-property plots

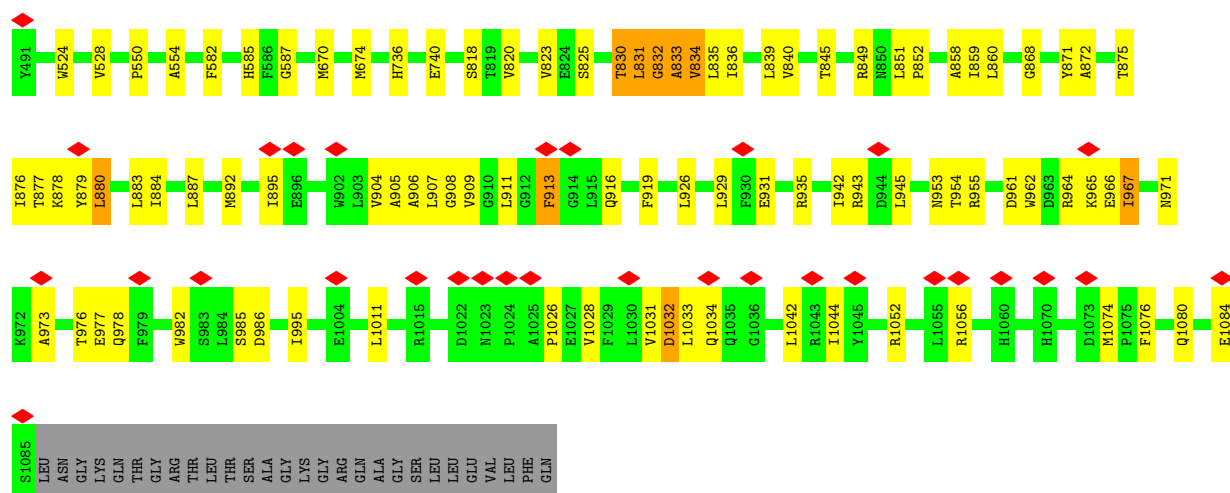
These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and atom inclusion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

• Molecule 1: Miniconductance mechanosensitive channel MscM

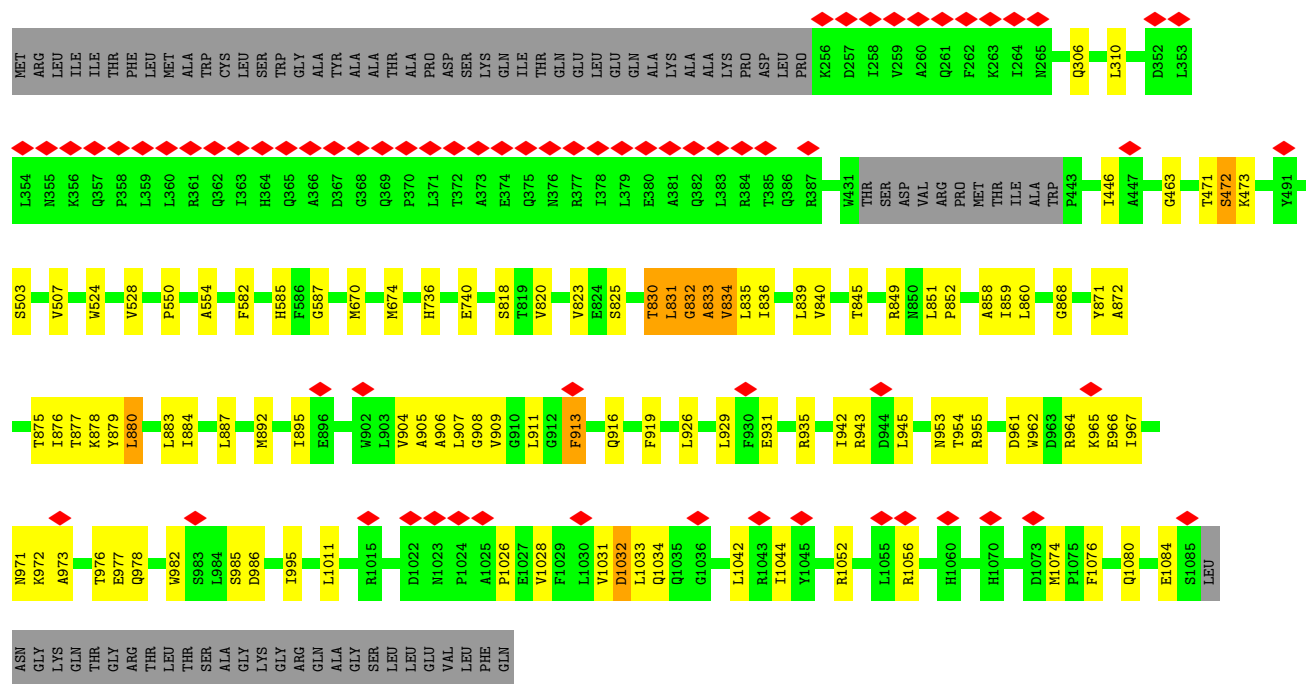
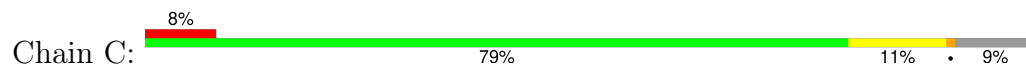


• Molecule 1: Miniconductance mechanosensitive channel MscM

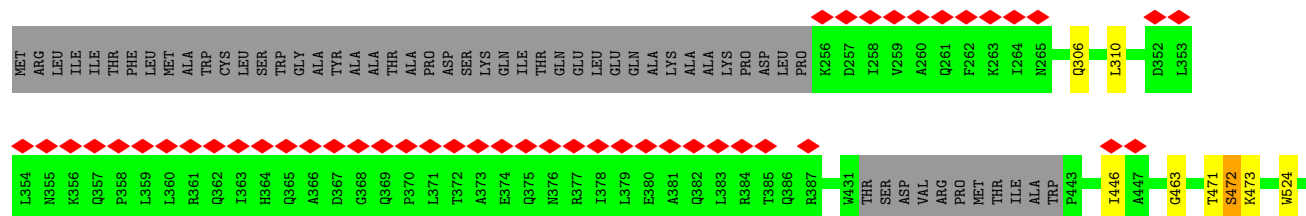
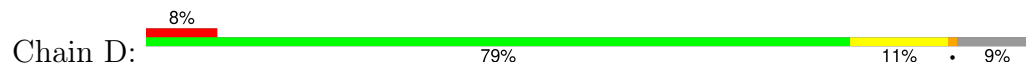


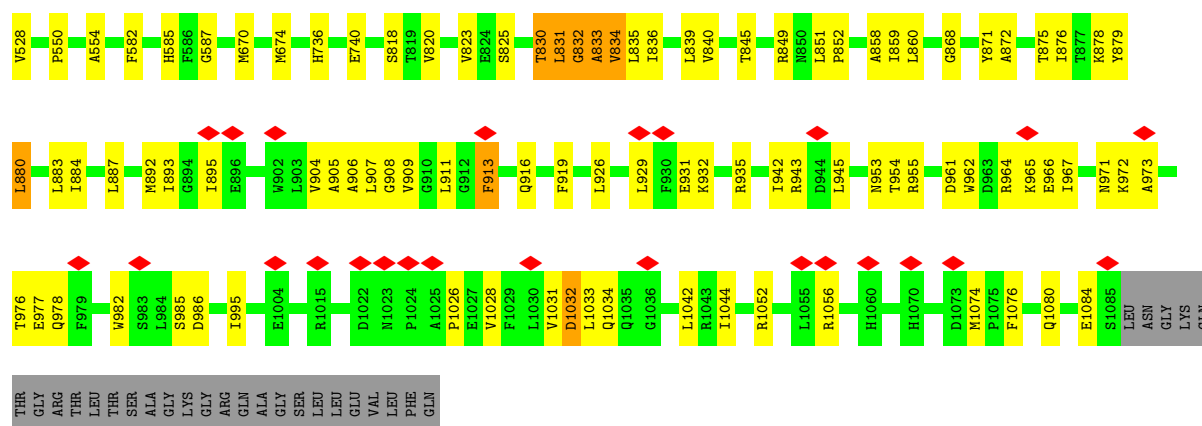


• Molecule 1: Miniconductance mechanosensitive channel MscM

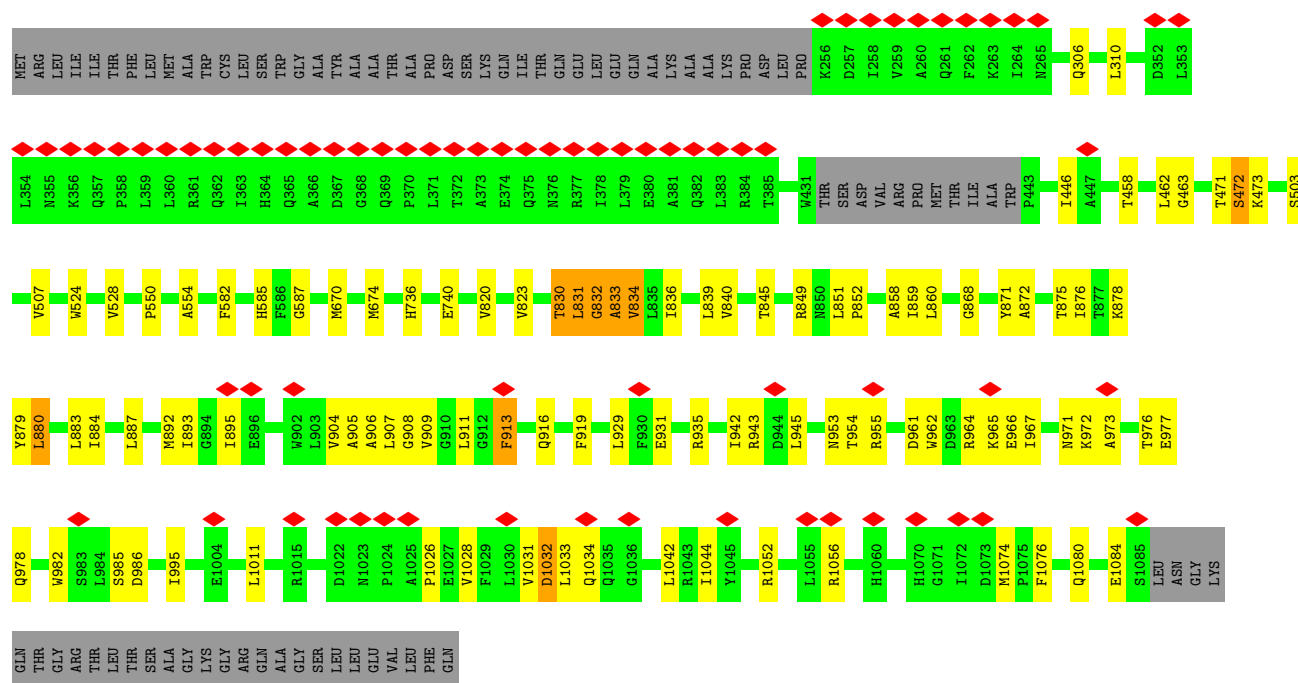
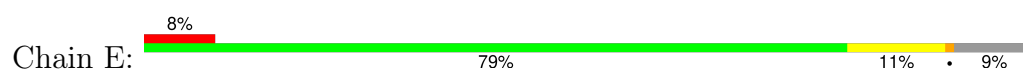


• Molecule 1: Miniconductance mechanosensitive channel MscM

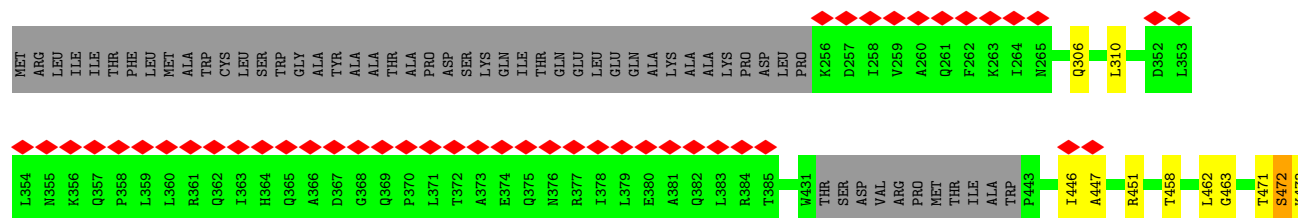
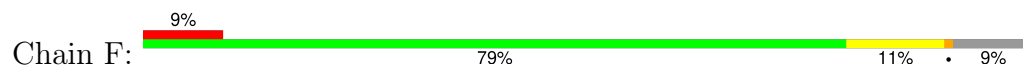




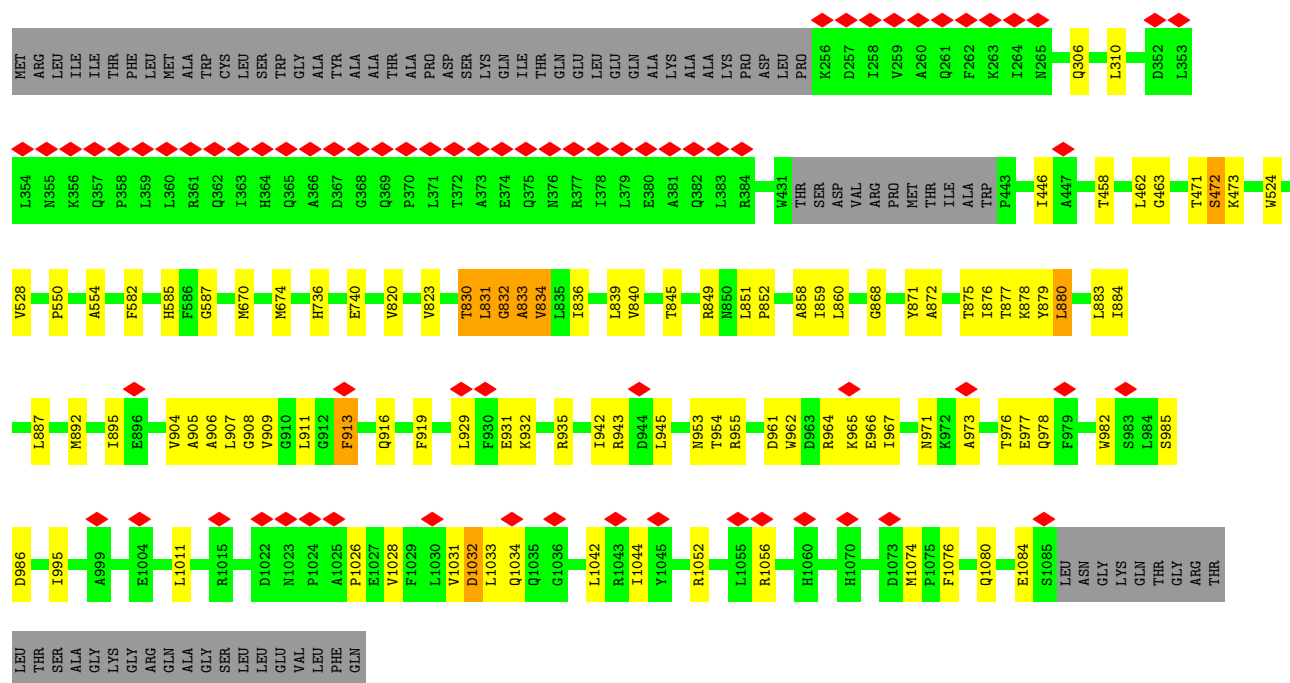
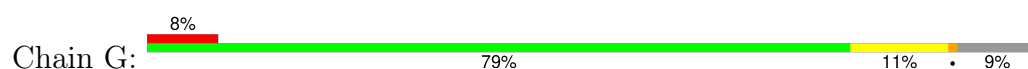
• Molecule 1: Miniconductance mechanosensitive channel MscM



• Molecule 1: Miniconductance mechanosensitive channel MscM



- Molecule 1: Miniconductance mechanosensitive channel MscM



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C7	Depositor
Number of particles used	48884	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	TFS KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	46.84	Depositor
Minimum defocus (nm)	800	Depositor
Maximum defocus (nm)	2000	Depositor
Magnification	105000	Depositor
Image detector	GATAN K3 (6k x 4k)	Depositor
Maximum map value	9.232	Depositor
Minimum map value	-0.125	Depositor
Average map value	0.030	Depositor
Map value standard deviation	0.491	Depositor
Recommended contour level	3.2	Depositor
Map size (Å)	381.15, 381.15, 381.15	wwPDB
Map dimensions	450, 450, 450	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	0.847, 0.847, 0.847	Depositor

5 Model quality

5.1 Standard geometry

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	A	0.30	1/4649 (0.0%)	0.55	7/6415 (0.1%)
1	B	0.30	0/4649	0.55	7/6415 (0.1%)
1	C	0.30	0/4649	0.55	7/6415 (0.1%)
1	D	0.30	1/4649 (0.0%)	0.55	7/6415 (0.1%)
1	E	0.30	0/4649	0.56	7/6415 (0.1%)
1	F	0.30	1/4649 (0.0%)	0.56	7/6415 (0.1%)
1	G	0.30	1/4649 (0.0%)	0.55	7/6415 (0.1%)
All	All	0.30	4/32543 (0.0%)	0.55	49/44905 (0.1%)

Chiral center outliers are detected by calculating the chiral volume of a chiral center and verifying if the center is modelled as a planar moiety or with the opposite hand. A planarity outlier is detected by checking planarity of atoms in a peptide group, atoms in a mainchain group or atoms of a sidechain that are expected to be planar.

Mol	Chain	#Chirality outliers	#Planarity outliers
1	A	0	1
1	B	0	1
1	C	0	1
1	D	0	1
1	E	0	1
1	F	0	1
1	G	0	1
All	All	0	7

All (4) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	G	831	LEU	CA-C	-5.08	1.45	1.52
1	A	831	LEU	CA-C	-5.01	1.45	1.52
1	F	831	LEU	CA-C	-5.01	1.45	1.52
1	D	831	LEU	CA-C	-5.00	1.45	1.52

All (49) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	B	832	GLY	N-CA-C	-10.12	89.20	113.18
1	C	832	GLY	N-CA-C	-10.11	89.21	113.18
1	A	832	GLY	N-CA-C	-10.11	89.22	113.18
1	D	832	GLY	N-CA-C	-10.11	89.21	113.18
1	F	832	GLY	N-CA-C	-10.11	89.22	113.18
1	G	832	GLY	N-CA-C	-10.11	89.22	113.18
1	E	832	GLY	N-CA-C	-10.11	89.23	113.18
1	F	446	ILE	N-CA-C	-8.15	102.31	110.62
1	D	446	ILE	N-CA-C	-8.13	102.33	110.62
1	A	446	ILE	N-CA-C	-8.12	102.33	110.62
1	G	446	ILE	N-CA-C	-8.12	102.34	110.62
1	C	446	ILE	N-CA-C	-8.11	102.35	110.62
1	B	446	ILE	N-CA-C	-8.11	102.35	110.62
1	E	446	ILE	N-CA-C	-8.09	102.36	110.62
1	F	831	LEU	N-CA-C	-7.31	104.09	113.16
1	D	831	LEU	N-CA-C	-7.31	104.10	113.16
1	C	831	LEU	N-CA-C	-7.30	104.10	113.16
1	E	831	LEU	N-CA-C	-7.30	104.10	113.16
1	B	831	LEU	N-CA-C	-7.30	104.10	113.16
1	A	831	LEU	N-CA-C	-7.29	104.12	113.16
1	G	831	LEU	N-CA-C	-7.26	104.16	113.16
1	G	463	GLY	N-CA-C	-5.58	106.03	112.73
1	C	463	GLY	N-CA-C	-5.57	106.04	112.73
1	B	463	GLY	N-CA-C	-5.57	106.04	112.73
1	D	463	GLY	N-CA-C	-5.57	106.05	112.73
1	F	463	GLY	N-CA-C	-5.57	106.05	112.73
1	A	463	GLY	N-CA-C	-5.55	106.06	112.73
1	E	463	GLY	N-CA-C	-5.54	106.08	112.73
1	G	1031	VAL	N-CA-C	5.40	116.21	111.56
1	B	1031	VAL	N-CA-C	5.40	116.20	111.56
1	E	1031	VAL	N-CA-C	5.39	116.20	111.56
1	A	1031	VAL	N-CA-C	5.39	116.19	111.56
1	F	1031	VAL	N-CA-C	5.38	116.19	111.56
1	D	1031	VAL	N-CA-C	5.38	116.18	111.56
1	C	1031	VAL	N-CA-C	5.34	116.16	111.56
1	D	1032	ASP	N-CA-C	5.17	117.32	108.90
1	A	880	LEU	CA-CB-CG	5.16	134.34	116.30
1	A	1032	ASP	N-CA-C	5.15	117.30	108.90
1	E	1032	ASP	N-CA-C	5.15	117.30	108.90
1	B	1032	ASP	N-CA-C	5.15	117.29	108.90
1	E	880	LEU	CA-CB-CG	5.15	134.32	116.30
1	G	880	LEU	CA-CB-CG	5.14	134.29	116.30
1	C	880	LEU	CA-CB-CG	5.14	134.28	116.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	F	1032	ASP	N-CA-C	5.14	117.28	108.90
1	B	880	LEU	CA-CB-CG	5.13	134.26	116.30
1	C	1032	ASP	N-CA-C	5.13	117.27	108.90
1	D	880	LEU	CA-CB-CG	5.13	134.26	116.30
1	F	880	LEU	CA-CB-CG	5.13	134.25	116.30
1	G	1032	ASP	N-CA-C	5.13	117.26	108.90

There are no chirality outliers.

All (7) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	A	830	THR	Mainchain
1	B	830	THR	Mainchain
1	C	830	THR	Mainchain
1	D	830	THR	Mainchain
1	E	830	THR	Mainchain
1	F	830	THR	Mainchain
1	G	830	THR	Mainchain

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	A	4611	0	3237	124	0
1	B	4611	0	3237	124	0
1	C	4611	0	3237	123	0
1	D	4611	0	3237	122	0
1	E	4611	0	3237	121	0
1	F	4611	0	3237	123	0
1	G	4611	0	3237	122	0
All	All	32277	0	22659	646	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 12.

All (646) 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:E:913:PHE:CD2	1:F:916:GLN:HB3	1.45	1.51
1:C:913:PHE:CD2	1:D:916:GLN:HB3	1.45	1.50
1:D:913:PHE:CD2	1:E:916:GLN:HB3	1.45	1.49
1:A:916:GLN:HB3	1:G:913:PHE:CD2	1.45	1.49
1:F:913:PHE:CD2	1:G:916:GLN:HB3	1.45	1.48
1:B:913:PHE:CD2	1:C:916:GLN:HB3	1.47	1.47
1:A:913:PHE:CD2	1:B:916:GLN:HB3	1.46	1.46
1:C:913:PHE:CE2	1:D:916:GLN:CD	2.00	1.40
1:A:916:GLN:CD	1:G:913:PHE:CE2	2.00	1.39
1:D:913:PHE:CE2	1:E:916:GLN:CD	2.01	1.39
1:F:913:PHE:CE2	1:G:916:GLN:CD	2.01	1.39
1:E:913:PHE:CE2	1:F:916:GLN:CD	2.01	1.38
1:A:913:PHE:CE2	1:B:916:GLN:CD	2.01	1.38
1:B:913:PHE:CE2	1:C:916:GLN:CD	2.01	1.37
1:A:916:GLN:CG	1:G:913:PHE:HE2	1.46	1.29
1:C:913:PHE:HE2	1:D:916:GLN:CG	1.46	1.28
1:E:913:PHE:HE2	1:F:916:GLN:CG	1.46	1.28
1:E:913:PHE:CE2	1:F:916:GLN:NE2	2.02	1.28
1:B:913:PHE:HE2	1:C:916:GLN:CG	1.47	1.28
1:C:913:PHE:HE2	1:D:916:GLN:CD	1.36	1.27
1:A:913:PHE:HE2	1:B:916:GLN:CG	1.46	1.27
1:D:913:PHE:HE2	1:E:916:GLN:CG	1.46	1.27
1:A:916:GLN:NE2	1:G:913:PHE:CE2	2.01	1.27
1:B:913:PHE:CE2	1:C:916:GLN:NE2	2.03	1.27
1:F:913:PHE:CE2	1:G:916:GLN:NE2	2.02	1.27
1:F:913:PHE:HE2	1:G:916:GLN:CG	1.46	1.27
1:A:913:PHE:CE2	1:B:916:GLN:NE2	2.03	1.26
1:C:913:PHE:CE2	1:D:916:GLN:NE2	2.02	1.26
1:F:913:PHE:HE2	1:G:916:GLN:CD	1.37	1.26
1:A:916:GLN:CD	1:G:913:PHE:HE2	1.36	1.26
1:A:913:PHE:HE2	1:B:916:GLN:CD	1.37	1.26
1:D:913:PHE:CE2	1:E:916:GLN:NE2	2.03	1.25
1:E:911:LEU:HD22	1:F:919:PHE:CE2	1.72	1.25
1:F:911:LEU:HD22	1:G:919:PHE:CE2	1.72	1.25
1:A:919:PHE:CE2	1:G:911:LEU:HD22	1.73	1.24
1:B:913:PHE:HE2	1:C:916:GLN:CD	1.37	1.24
1:D:911:LEU:HD22	1:E:919:PHE:CE2	1.72	1.23
1:A:911:LEU:HD22	1:B:919:PHE:CE2	1.73	1.23
1:C:911:LEU:HD22	1:D:919:PHE:CE2	1.72	1.22
1:D:913:PHE:HE2	1:E:916:GLN:CD	1.37	1.22
1:E:913:PHE:HE2	1:F:916:GLN:CD	1.37	1.22
1:B:911:LEU:HD22	1:C:919:PHE:CE2	1.73	1.21

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:913:PHE:CE2	1:B:916:GLN:HB3	1.83	1.13
1:D:913:PHE:CE2	1:E:916:GLN:CG	2.31	1.12
1:D:913:PHE:CE2	1:E:916:GLN:HB3	1.83	1.12
1:E:913:PHE:CE2	1:F:916:GLN:HB3	1.84	1.12
1:F:913:PHE:CE2	1:G:916:GLN:HB3	1.84	1.12
1:B:913:PHE:CE2	1:C:916:GLN:HB3	1.83	1.11
1:A:916:GLN:HB3	1:G:913:PHE:CE2	1.84	1.11
1:C:913:PHE:CE2	1:D:916:GLN:HB3	1.83	1.11
1:F:913:PHE:CE2	1:G:916:GLN:CG	2.32	1.10
1:A:916:GLN:CG	1:G:913:PHE:CE2	2.31	1.10
1:A:913:PHE:CE2	1:B:916:GLN:CB	2.37	1.08
1:B:913:PHE:CE2	1:C:916:GLN:CG	2.33	1.08
1:A:913:PHE:CE2	1:B:916:GLN:CG	2.32	1.08
1:C:913:PHE:CE2	1:D:916:GLN:CG	2.31	1.08
1:D:913:PHE:CE2	1:E:916:GLN:CB	2.36	1.08
1:E:913:PHE:CE2	1:F:916:GLN:CB	2.37	1.08
1:E:913:PHE:CE2	1:F:916:GLN:CG	2.31	1.08
1:B:913:PHE:CE2	1:C:916:GLN:CB	2.37	1.07
1:A:916:GLN:CB	1:G:913:PHE:CE2	2.37	1.07
1:F:913:PHE:CE2	1:G:916:GLN:CB	2.37	1.07
1:C:913:PHE:CE2	1:D:916:GLN:CB	2.36	1.07
1:D:913:PHE:CD2	1:E:916:GLN:CB	2.38	1.07
1:F:913:PHE:CD2	1:G:916:GLN:CB	2.38	1.06
1:A:916:GLN:CB	1:G:913:PHE:CD2	2.38	1.06
1:C:913:PHE:CD2	1:D:916:GLN:CB	2.38	1.06
1:E:913:PHE:CD2	1:F:916:GLN:CB	2.38	1.05
1:A:913:PHE:CD2	1:B:916:GLN:CB	2.39	1.04
1:B:913:PHE:CD2	1:C:916:GLN:CB	2.40	1.03
1:D:831:LEU:HA	1:D:834:VAL:HG13	1.44	1.00
1:C:911:LEU:HD22	1:D:919:PHE:CZ	1.97	0.99
1:C:831:LEU:HA	1:C:834:VAL:HG13	1.44	0.99
1:E:831:LEU:HA	1:E:834:VAL:HG13	1.44	0.99
1:E:911:LEU:HD22	1:F:919:PHE:CZ	1.97	0.99
1:F:831:LEU:HA	1:F:834:VAL:HG13	1.44	0.99
1:A:911:LEU:HD22	1:B:919:PHE:CZ	1.97	0.99
1:F:911:LEU:HD22	1:G:919:PHE:CZ	1.97	0.99
1:G:831:LEU:HA	1:G:834:VAL:HG13	1.44	0.99
1:A:919:PHE:CZ	1:G:911:LEU:HD22	1.97	0.99
1:D:911:LEU:HD22	1:E:919:PHE:CZ	1.96	0.98
1:B:911:LEU:HD22	1:C:919:PHE:CZ	1.97	0.98
1:A:831:LEU:HA	1:A:834:VAL:HG13	1.44	0.98

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:831:LEU:HA	1:B:834:VAL:HG13	1.44	0.98
1:F:913:PHE:HD2	1:G:916:GLN:HB3	1.14	0.96
1:D:913:PHE:HD2	1:E:916:GLN:HB3	1.14	0.94
1:C:913:PHE:HD2	1:D:916:GLN:HB3	1.13	0.92
1:E:913:PHE:HD2	1:F:916:GLN:HB3	1.13	0.92
1:A:913:PHE:HD2	1:B:916:GLN:HB3	1.15	0.90
1:B:913:PHE:HD2	1:C:916:GLN:HB3	1.15	0.90
1:A:916:GLN:HB3	1:G:913:PHE:HD2	1.13	0.89
1:A:916:GLN:NE2	1:G:913:PHE:CZ	2.45	0.85
1:F:913:PHE:CZ	1:G:916:GLN:NE2	2.46	0.84
1:A:913:PHE:CZ	1:B:916:GLN:NE2	2.46	0.83
1:C:913:PHE:CZ	1:D:916:GLN:NE2	2.45	0.83
1:D:913:PHE:CZ	1:E:916:GLN:NE2	2.45	0.83
1:E:913:PHE:CZ	1:F:916:GLN:NE2	2.46	0.83
1:A:964:ARG:NH2	1:B:985:SER:OG	2.12	0.82
1:B:913:PHE:CZ	1:C:916:GLN:NE2	2.46	0.82
1:E:911:LEU:CD2	1:F:919:PHE:CE2	2.61	0.82
1:F:911:LEU:CD2	1:G:919:PHE:CE2	2.62	0.82
1:F:964:ARG:NH2	1:G:985:SER:OG	2.13	0.82
1:C:964:ARG:NH2	1:D:985:SER:OG	2.13	0.82
1:B:964:ARG:NH2	1:C:985:SER:OG	2.13	0.81
1:E:964:ARG:NH2	1:F:985:SER:OG	2.14	0.81
1:D:964:ARG:NH2	1:E:985:SER:OG	2.13	0.81
1:A:985:SER:OG	1:G:964:ARG:NH2	2.13	0.81
1:B:911:LEU:CD2	1:C:919:PHE:CE2	2.62	0.80
1:D:911:LEU:CD2	1:E:919:PHE:CE2	2.62	0.79
1:A:919:PHE:CE2	1:G:911:LEU:CD2	2.62	0.79
1:C:911:LEU:CD2	1:D:919:PHE:CE2	2.62	0.79
1:A:911:LEU:CD2	1:B:919:PHE:CE2	2.62	0.78
1:A:858:ALA:O	1:A:860:LEU:N	2.18	0.77
1:G:858:ALA:O	1:G:860:LEU:N	2.18	0.77
1:F:858:ALA:O	1:F:860:LEU:N	2.18	0.76
1:C:913:PHE:CD2	1:D:916:GLN:CD	2.64	0.76
1:E:913:PHE:CD2	1:F:916:GLN:CD	2.64	0.76
1:D:858:ALA:O	1:D:860:LEU:N	2.18	0.76
1:D:913:PHE:CD2	1:E:916:GLN:CD	2.64	0.76
1:A:916:GLN:CD	1:G:913:PHE:CD2	2.63	0.75
1:C:858:ALA:O	1:C:860:LEU:N	2.18	0.75
1:E:858:ALA:O	1:E:860:LEU:N	2.18	0.75
1:B:913:PHE:CD2	1:C:916:GLN:CD	2.65	0.75
1:D:961:ASP:OD1	1:D:962:TRP:N	2.21	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:961:ASP:OD1	1:C:962:TRP:N	2.21	0.73
1:F:913:PHE:CD2	1:G:916:GLN:CD	2.64	0.73
1:G:961:ASP:OD1	1:G:962:TRP:N	2.21	0.73
1:F:961:ASP:OD1	1:F:962:TRP:N	2.21	0.73
1:E:961:ASP:OD1	1:E:962:TRP:N	2.21	0.73
1:A:961:ASP:OD1	1:A:962:TRP:N	2.21	0.73
1:B:858:ALA:O	1:B:860:LEU:N	2.18	0.73
1:B:961:ASP:OD1	1:B:962:TRP:N	2.21	0.73
1:A:913:PHE:CD2	1:B:916:GLN:CD	2.65	0.72
1:A:916:GLN:HG3	1:G:913:PHE:HE2	1.53	0.72
1:D:875:THR:HA	1:D:878:LYS:HZ2	1.53	0.72
1:A:953:ASN:OD1	1:A:954:THR:N	2.24	0.70
1:G:953:ASN:OD1	1:G:954:THR:N	2.24	0.70
1:F:953:ASN:OD1	1:F:954:THR:N	2.24	0.70
1:C:904:VAL:O	1:C:905:ALA:C	2.34	0.70
1:E:913:PHE:HE2	1:F:916:GLN:HG3	1.53	0.70
1:B:953:ASN:OD1	1:B:954:THR:N	2.24	0.70
1:E:953:ASN:OD1	1:E:954:THR:N	2.24	0.69
1:C:913:PHE:HE2	1:D:916:GLN:HG3	1.54	0.69
1:B:904:VAL:O	1:B:905:ALA:C	2.34	0.69
1:D:953:ASN:OD1	1:D:954:THR:N	2.24	0.69
1:G:904:VAL:O	1:G:905:ALA:C	2.34	0.69
1:F:913:PHE:HE2	1:G:916:GLN:NE2	1.63	0.69
1:C:953:ASN:OD1	1:C:954:THR:N	2.24	0.69
1:A:904:VAL:O	1:A:905:ALA:C	2.34	0.69
1:D:913:PHE:HE2	1:E:916:GLN:HG3	1.54	0.69
1:F:913:PHE:HE2	1:G:916:GLN:HG3	1.54	0.68
1:F:904:VAL:O	1:F:905:ALA:C	2.34	0.68
1:A:913:PHE:HE2	1:B:916:GLN:NE2	1.63	0.67
1:C:913:PHE:HE2	1:D:916:GLN:NE2	1.63	0.67
1:E:904:VAL:O	1:E:905:ALA:C	2.34	0.66
1:D:904:VAL:O	1:D:905:ALA:C	2.34	0.65
1:A:954:THR:O	1:A:971:ASN:ND2	2.30	0.65
1:D:954:THR:O	1:D:971:ASN:ND2	2.30	0.65
1:C:832:GLY:O	1:C:836:ILE:HD12	1.97	0.64
1:G:954:THR:O	1:G:971:ASN:ND2	2.30	0.64
1:B:832:GLY:O	1:B:836:ILE:HD12	1.97	0.64
1:B:954:THR:O	1:B:971:ASN:ND2	2.30	0.64
1:A:832:GLY:O	1:A:836:ILE:HD12	1.97	0.64
1:D:832:GLY:O	1:D:836:ILE:HD12	1.97	0.64
1:C:954:THR:O	1:C:971:ASN:ND2	2.30	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:832:GLY:O	1:E:836:ILE:HD12	1.97	0.64
1:D:306:GLN:O	1:D:310:LEU:N	2.32	0.63
1:E:954:THR:O	1:E:971:ASN:ND2	2.30	0.63
1:G:832:GLY:O	1:G:836:ILE:HD12	1.97	0.63
1:F:832:GLY:O	1:F:836:ILE:HD12	1.97	0.63
1:B:306:GLN:O	1:B:310:LEU:N	2.32	0.63
1:A:306:GLN:O	1:A:310:LEU:N	2.32	0.63
1:A:1080:GLN:HB2	1:G:1076:PHE:HB2	1.81	0.63
1:F:830:THR:O	1:F:833:ALA:HB3	1.99	0.63
1:C:306:GLN:O	1:C:310:LEU:N	2.32	0.63
1:E:830:THR:O	1:E:833:ALA:HB3	1.99	0.63
1:D:830:THR:O	1:D:833:ALA:HB3	1.99	0.62
1:D:913:PHE:HE2	1:E:916:GLN:NE2	1.63	0.62
1:E:306:GLN:O	1:E:310:LEU:N	2.32	0.62
1:F:954:THR:O	1:F:971:ASN:ND2	2.30	0.62
1:A:830:THR:O	1:A:833:ALA:HB3	1.99	0.62
1:F:306:GLN:O	1:F:310:LEU:N	2.32	0.62
1:G:306:GLN:O	1:G:310:LEU:N	2.32	0.62
1:G:830:THR:O	1:G:833:ALA:HB3	1.99	0.62
1:B:830:THR:O	1:B:833:ALA:HB3	1.99	0.62
1:C:830:THR:O	1:C:833:ALA:HB3	1.99	0.62
1:E:1076:PHE:HB2	1:F:1080:GLN:HB2	1.82	0.62
1:F:875:THR:HA	1:F:878:LYS:HZ2	1.65	0.62
1:F:1076:PHE:HB2	1:G:1080:GLN:HB2	1.82	0.62
1:B:1076:PHE:HB2	1:C:1080:GLN:HB2	1.82	0.62
1:C:977:GLU:OE2	1:C:977:GLU:N	2.33	0.61
1:A:832:GLY:O	1:A:833:ALA:C	2.43	0.61
1:C:1076:PHE:HB2	1:D:1080:GLN:HB2	1.82	0.61
1:D:1076:PHE:HB2	1:E:1080:GLN:HB2	1.82	0.61
1:B:832:GLY:O	1:B:833:ALA:C	2.43	0.61
1:D:977:GLU:N	1:D:977:GLU:OE2	2.33	0.61
1:F:832:GLY:O	1:F:833:ALA:C	2.43	0.61
1:G:977:GLU:N	1:G:977:GLU:OE2	2.33	0.61
1:A:1076:PHE:HB2	1:B:1080:GLN:HB2	1.83	0.61
1:A:977:GLU:N	1:A:977:GLU:OE2	2.33	0.61
1:B:977:GLU:N	1:B:977:GLU:OE2	2.33	0.61
1:E:875:THR:HA	1:E:878:LYS:HZ2	1.65	0.61
1:E:977:GLU:OE2	1:E:977:GLU:N	2.33	0.61
1:F:977:GLU:N	1:F:977:GLU:OE2	2.33	0.61
1:G:832:GLY:O	1:G:833:ALA:C	2.43	0.60
1:E:831:LEU:HA	1:E:834:VAL:CG1	2.27	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:966:GLU:N	1:B:966:GLU:OE2	2.35	0.60
1:A:966:GLU:OE2	1:A:966:GLU:N	2.35	0.60
1:C:832:GLY:O	1:C:833:ALA:C	2.43	0.60
1:C:966:GLU:OE2	1:C:966:GLU:N	2.35	0.60
1:F:831:LEU:HA	1:F:834:VAL:CG1	2.27	0.60
1:A:831:LEU:HA	1:A:834:VAL:CG1	2.27	0.59
1:D:966:GLU:N	1:D:966:GLU:OE2	2.35	0.59
1:E:832:GLY:O	1:E:833:ALA:C	2.43	0.59
1:E:966:GLU:N	1:E:966:GLU:OE2	2.35	0.59
1:G:966:GLU:N	1:G:966:GLU:OE2	2.35	0.59
1:F:966:GLU:OE2	1:F:966:GLU:N	2.35	0.59
1:B:871:TYR:HE1	1:C:929:LEU:HD21	1.67	0.58
1:D:831:LEU:HA	1:D:834:VAL:CG1	2.27	0.58
1:D:871:TYR:HE1	1:E:929:LEU:HD21	1.68	0.58
1:G:831:LEU:HA	1:G:834:VAL:CG1	2.27	0.58
1:E:1032:ASP:OD1	1:E:1034:GLN:HG3	2.04	0.58
1:F:1032:ASP:OD1	1:F:1034:GLN:HG3	2.04	0.58
1:B:1032:ASP:OD1	1:B:1034:GLN:HG3	2.04	0.58
1:B:907:LEU:O	1:B:911:LEU:HG	2.04	0.58
1:C:831:LEU:HA	1:C:834:VAL:CG1	2.27	0.58
1:G:524:TRP:O	1:G:528:VAL:N	2.35	0.58
1:G:1032:ASP:OD1	1:G:1034:GLN:HG3	2.04	0.58
1:D:832:GLY:O	1:D:833:ALA:C	2.43	0.58
1:D:1032:ASP:OD1	1:D:1034:GLN:HG3	2.04	0.58
1:D:907:LEU:O	1:D:911:LEU:HG	2.04	0.57
1:F:524:TRP:O	1:F:528:VAL:N	2.35	0.57
1:E:871:TYR:HE1	1:F:929:LEU:HD21	1.68	0.57
1:G:907:LEU:O	1:G:911:LEU:HG	2.04	0.57
1:A:1032:ASP:OD1	1:A:1034:GLN:HG3	2.04	0.57
1:B:524:TRP:O	1:B:528:VAL:N	2.35	0.57
1:C:871:TYR:HE1	1:D:929:LEU:HD21	1.68	0.57
1:C:907:LEU:O	1:C:911:LEU:HG	2.04	0.57
1:F:907:LEU:O	1:F:911:LEU:HG	2.04	0.57
1:C:1032:ASP:OD1	1:C:1034:GLN:HG3	2.04	0.57
1:C:875:THR:HA	1:C:878:LYS:HZ2	1.70	0.57
1:E:907:LEU:O	1:E:911:LEU:HG	2.04	0.56
1:F:871:TYR:HE1	1:G:929:LEU:HD21	1.69	0.56
1:E:524:TRP:O	1:E:528:VAL:N	2.35	0.56
1:A:871:TYR:HE1	1:B:929:LEU:HD21	1.68	0.56
1:A:875:THR:HA	1:A:878:LYS:HZ2	1.70	0.56
1:A:907:LEU:O	1:A:911:LEU:HG	2.04	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:929:LEU:HD21	1:G:871:TYR:HE1	1.69	0.56
1:A:913:PHE:HE2	1:B:916:GLN:HG3	1.55	0.56
1:E:832:GLY:C	1:E:836:ILE:HD12	2.31	0.55
1:F:832:GLY:C	1:F:836:ILE:HD12	2.31	0.55
1:D:832:GLY:C	1:D:836:ILE:HD12	2.31	0.55
1:D:524:TRP:O	1:D:528:VAL:N	2.35	0.55
1:C:832:GLY:C	1:C:836:ILE:HD12	2.31	0.55
1:B:832:GLY:C	1:B:836:ILE:HD12	2.31	0.55
1:A:851:LEU:HD23	1:A:852:PRO:HD3	1.89	0.55
1:D:832:GLY:C	1:D:836:ILE:CD1	2.80	0.55
1:A:832:GLY:C	1:A:836:ILE:HD12	2.31	0.55
1:B:851:LEU:HD23	1:B:852:PRO:HD3	1.89	0.55
1:D:670:MET:O	1:D:674:MET:N	2.38	0.55
1:E:832:GLY:C	1:E:836:ILE:CD1	2.80	0.55
1:A:916:GLN:NE2	1:G:913:PHE:HE2	1.62	0.55
1:B:670:MET:O	1:B:674:MET:N	2.38	0.55
1:B:832:GLY:C	1:B:836:ILE:CD1	2.80	0.55
1:A:1084:GLU:OE2	1:G:1080:GLN:NE2	2.36	0.55
1:E:913:PHE:HE2	1:F:916:GLN:NE2	1.63	0.55
1:B:875:THR:HA	1:B:878:LYS:HZ2	1.72	0.55
1:C:832:GLY:C	1:C:836:ILE:CD1	2.80	0.55
1:F:832:GLY:C	1:F:836:ILE:CD1	2.80	0.55
1:C:942:ILE:HG12	1:C:943:ARG:HG3	1.89	0.54
1:D:942:ILE:HG12	1:D:943:ARG:HG3	1.90	0.54
1:E:942:ILE:HG12	1:E:943:ARG:HG3	1.89	0.54
1:F:911:LEU:CD2	1:G:919:PHE:CZ	2.84	0.54
1:G:832:GLY:C	1:G:836:ILE:CD1	2.80	0.54
1:B:942:ILE:HG12	1:B:943:ARG:HG3	1.90	0.54
1:C:851:LEU:HD23	1:C:852:PRO:HD3	1.89	0.54
1:G:832:GLY:C	1:G:836:ILE:HD12	2.31	0.54
1:G:851:LEU:HD23	1:G:852:PRO:HD3	1.89	0.54
1:A:832:GLY:C	1:A:836:ILE:CD1	2.80	0.54
1:A:942:ILE:HG12	1:A:943:ARG:HG3	1.89	0.54
1:B:871:TYR:CE1	1:C:929:LEU:HD21	2.43	0.54
1:B:913:PHE:HE2	1:C:916:GLN:HG3	1.56	0.54
1:F:872:ALA:O	1:F:876:ILE:HG22	2.08	0.54
1:E:913:PHE:HD2	1:F:916:GLN:CB	2.01	0.54
1:F:942:ILE:HG12	1:F:943:ARG:HG3	1.90	0.54
1:C:872:ALA:O	1:C:876:ILE:HG22	2.08	0.54
1:G:942:ILE:HG12	1:G:943:ARG:HG3	1.89	0.54
1:A:736:HIS:O	1:A:740:GLU:N	2.34	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:911:LEU:CD2	1:D:919:PHE:CZ	2.84	0.54
1:E:736:HIS:O	1:E:740:GLU:N	2.34	0.54
1:F:851:LEU:HD23	1:F:852:PRO:HD3	1.89	0.54
1:D:851:LEU:HD23	1:D:852:PRO:HD3	1.89	0.54
1:A:986:ASP:OD1	1:A:986:ASP:N	2.39	0.53
1:G:872:ALA:O	1:G:876:ILE:HG22	2.08	0.53
1:A:871:TYR:CE1	1:B:929:LEU:HD21	2.44	0.53
1:B:831:LEU:HA	1:B:834:VAL:CG1	2.27	0.53
1:B:911:LEU:CD2	1:C:919:PHE:CZ	2.84	0.53
1:E:872:ALA:O	1:E:876:ILE:HG22	2.08	0.53
1:C:871:TYR:CE1	1:D:929:LEU:HD21	2.44	0.53
1:D:872:ALA:O	1:D:876:ILE:HG22	2.08	0.53
1:E:851:LEU:HD23	1:E:852:PRO:HD3	1.89	0.53
1:F:904:VAL:HA	1:F:907:LEU:HD23	1.89	0.53
1:A:524:TRP:O	1:A:528:VAL:N	2.35	0.53
1:B:872:ALA:O	1:B:876:ILE:HG22	2.08	0.53
1:B:904:VAL:HA	1:B:907:LEU:HD23	1.90	0.53
1:C:904:VAL:HA	1:C:907:LEU:HD23	1.90	0.53
1:A:872:ALA:O	1:A:876:ILE:HG22	2.08	0.53
1:B:875:THR:HA	1:B:878:LYS:NZ	2.24	0.53
1:F:471:THR:O	1:F:472:SER:C	2.52	0.53
1:G:904:VAL:HA	1:G:907:LEU:HD23	1.90	0.53
1:E:875:THR:HA	1:E:878:LYS:NZ	2.24	0.53
1:A:929:LEU:HD21	1:G:871:TYR:CE1	2.44	0.53
1:D:904:VAL:HA	1:D:907:LEU:HD23	1.89	0.53
1:D:913:PHE:HD2	1:E:916:GLN:CB	2.02	0.53
1:E:471:THR:O	1:E:472:SER:C	2.52	0.53
1:F:871:TYR:CE1	1:G:929:LEU:HD21	2.44	0.53
1:G:875:THR:HA	1:G:878:LYS:NZ	2.24	0.53
1:B:986:ASP:OD1	1:B:986:ASP:N	2.39	0.52
1:F:875:THR:HA	1:F:878:LYS:NZ	2.24	0.52
1:G:471:THR:O	1:G:472:SER:C	2.52	0.52
1:C:524:TRP:O	1:C:528:VAL:N	2.35	0.52
1:A:875:THR:HA	1:A:878:LYS:NZ	2.24	0.52
1:A:904:VAL:HA	1:A:907:LEU:HD23	1.90	0.52
1:C:875:THR:HA	1:C:878:LYS:NZ	2.24	0.52
1:D:875:THR:HA	1:D:878:LYS:NZ	2.24	0.52
1:E:871:TYR:CE1	1:F:929:LEU:HD21	2.44	0.52
1:D:871:TYR:CE1	1:E:929:LEU:HD21	2.44	0.52
1:E:670:MET:O	1:E:674:MET:N	2.38	0.52
1:G:875:THR:HA	1:G:878:LYS:HZ2	1.74	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:995:ILE:HG23	1:F:1074:MET:HE2	1.92	0.52
1:D:471:THR:O	1:D:472:SER:C	2.52	0.52
1:C:995:ILE:HG23	1:C:1074:MET:HE2	1.92	0.52
1:E:904:VAL:HA	1:E:907:LEU:HD23	1.90	0.52
1:B:995:ILE:HG23	1:B:1074:MET:HE2	1.92	0.52
1:C:1080:GLN:NE2	1:D:1084:GLU:OE2	2.37	0.52
1:E:995:ILE:HG23	1:E:1074:MET:HE2	1.92	0.52
1:G:995:ILE:HG23	1:G:1074:MET:HE2	1.92	0.52
1:B:471:THR:O	1:B:472:SER:C	2.52	0.52
1:D:995:ILE:HG23	1:D:1074:MET:HE2	1.92	0.52
1:A:995:ILE:HG23	1:A:1074:MET:HE2	1.92	0.51
1:C:471:THR:O	1:C:472:SER:C	2.52	0.51
1:D:736:HIS:O	1:D:740:GLU:N	2.34	0.51
1:C:670:MET:O	1:C:674:MET:N	2.38	0.51
1:C:913:PHE:HD2	1:D:916:GLN:CB	2.02	0.51
1:F:868:GLY:HA2	1:F:871:TYR:HB3	1.93	0.51
1:G:868:GLY:HA2	1:G:871:TYR:HB3	1.93	0.51
1:F:550:PRO:O	1:F:554:ALA:N	2.41	0.51
1:F:913:PHE:HD2	1:G:916:GLN:CB	2.02	0.51
1:A:471:THR:O	1:A:472:SER:C	2.52	0.51
1:E:1052:ARG:O	1:E:1056:ARG:HG3	2.11	0.51
1:F:1080:GLN:NE2	1:G:1084:GLU:OE2	2.38	0.51
1:G:1052:ARG:O	1:G:1056:ARG:HG3	2.11	0.51
1:B:868:GLY:HA2	1:B:871:TYR:HB3	1.93	0.51
1:C:1052:ARG:O	1:C:1056:ARG:HG3	2.11	0.51
1:E:550:PRO:O	1:E:554:ALA:N	2.41	0.51
1:F:736:HIS:O	1:F:740:GLU:N	2.34	0.51
1:A:868:GLY:HA2	1:A:871:TYR:HB3	1.93	0.50
1:B:1052:ARG:O	1:B:1056:ARG:HG3	2.11	0.50
1:D:868:GLY:HA2	1:D:871:TYR:HB3	1.93	0.50
1:E:868:GLY:HA2	1:E:871:TYR:HB3	1.93	0.50
1:C:736:HIS:O	1:C:740:GLU:N	2.34	0.50
1:F:670:MET:O	1:F:674:MET:N	2.38	0.50
1:G:670:MET:O	1:G:674:MET:N	2.38	0.50
1:C:986:ASP:OD1	1:C:986:ASP:N	2.39	0.50
1:C:868:GLY:HA2	1:C:871:TYR:HB3	1.93	0.50
1:A:1052:ARG:O	1:A:1056:ARG:HG3	2.11	0.50
1:D:911:LEU:CD2	1:E:919:PHE:CZ	2.84	0.50
1:D:1052:ARG:O	1:D:1056:ARG:HG3	2.11	0.50
1:F:892:MET:HA	1:F:892:MET:HE2	1.94	0.50
1:G:906:ALA:O	1:G:909:VAL:HG22	2.12	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:906:ALA:O	1:A:909:VAL:HG22	2.12	0.50
1:E:892:MET:HE2	1:E:892:MET:HA	1.94	0.50
1:E:911:LEU:CD2	1:F:919:PHE:CZ	2.84	0.50
1:E:976:THR:OG1	1:E:977:GLU:OE2	2.29	0.50
1:A:670:MET:O	1:A:674:MET:N	2.38	0.49
1:D:1080:GLN:NE2	1:E:1084:GLU:OE2	2.38	0.49
1:F:906:ALA:O	1:F:909:VAL:HG22	2.12	0.49
1:B:906:ALA:O	1:B:909:VAL:HG22	2.12	0.49
1:F:1052:ARG:O	1:F:1056:ARG:HG3	2.11	0.49
1:D:892:MET:HE2	1:D:892:MET:HA	1.94	0.49
1:B:913:PHE:HE2	1:C:916:GLN:NE2	1.63	0.49
1:C:906:ALA:O	1:C:909:VAL:HG22	2.12	0.49
1:E:911:LEU:HD22	1:F:919:PHE:CD2	2.41	0.49
1:G:976:THR:OG1	1:G:977:GLU:OE2	2.29	0.49
1:A:911:LEU:CD2	1:B:919:PHE:CZ	2.84	0.49
1:G:892:MET:HE2	1:G:892:MET:HA	1.94	0.49
1:D:906:ALA:O	1:D:909:VAL:HG22	2.12	0.49
1:E:906:ALA:O	1:E:909:VAL:HG22	2.12	0.49
1:B:820:VAL:N	1:B:823:VAL:O	2.46	0.48
1:D:976:THR:OG1	1:D:977:GLU:OE2	2.29	0.48
1:B:736:HIS:O	1:B:740:GLU:N	2.34	0.48
1:E:1080:GLN:NE2	1:F:1084:GLU:OE2	2.36	0.48
1:D:820:VAL:N	1:D:823:VAL:O	2.46	0.48
1:E:820:VAL:N	1:E:823:VAL:O	2.46	0.48
1:F:976:THR:OG1	1:F:977:GLU:OE2	2.29	0.48
1:A:916:GLN:CB	1:G:913:PHE:HD2	2.02	0.48
1:C:892:MET:HA	1:C:892:MET:HE2	1.94	0.48
1:F:820:VAL:N	1:F:823:VAL:O	2.46	0.48
1:A:820:VAL:N	1:A:823:VAL:O	2.46	0.48
1:G:820:VAL:N	1:G:823:VAL:O	2.46	0.48
1:A:892:MET:HA	1:A:892:MET:HE2	1.94	0.48
1:C:550:PRO:O	1:C:554:ALA:N	2.41	0.47
1:C:820:VAL:N	1:C:823:VAL:O	2.46	0.47
1:D:986:ASP:OD1	1:D:986:ASP:N	2.39	0.47
1:A:919:PHE:CZ	1:G:911:LEU:CD2	2.84	0.47
1:B:1080:GLN:NE2	1:C:1084:GLU:OE2	2.39	0.47
1:C:976:THR:OG1	1:C:977:GLU:OE2	2.29	0.47
1:B:892:MET:HE2	1:B:892:MET:HA	1.94	0.47
1:A:904:VAL:O	1:A:907:LEU:N	2.48	0.47
1:E:904:VAL:O	1:E:907:LEU:N	2.48	0.47
1:G:736:HIS:O	1:G:740:GLU:N	2.34	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:904:VAL:O	1:C:907:LEU:N	2.48	0.47
1:B:1028:VAL:HG22	1:B:1042:LEU:HD13	1.96	0.46
1:G:904:VAL:O	1:G:907:LEU:N	2.48	0.46
1:A:976:THR:OG1	1:A:977:GLU:OE2	2.29	0.46
1:B:976:THR:OG1	1:B:977:GLU:OE2	2.29	0.46
1:E:1028:VAL:HG22	1:E:1042:LEU:HD13	1.96	0.46
1:F:904:VAL:O	1:F:907:LEU:N	2.48	0.46
1:C:1028:VAL:HG22	1:C:1042:LEU:HD13	1.96	0.46
1:D:1028:VAL:HG22	1:D:1042:LEU:HD13	1.96	0.46
1:A:1028:VAL:HG22	1:A:1042:LEU:HD13	1.96	0.46
1:B:904:VAL:O	1:B:907:LEU:N	2.48	0.46
1:D:876:ILE:HA	1:D:879:TYR:CD2	2.51	0.46
1:D:904:VAL:O	1:D:907:LEU:N	2.48	0.46
1:E:876:ILE:HA	1:E:879:TYR:CD2	2.51	0.46
1:G:1028:VAL:HG22	1:G:1042:LEU:HD13	1.96	0.46
1:B:913:PHE:HD2	1:C:916:GLN:CB	2.04	0.46
1:E:986:ASP:OD1	1:E:986:ASP:N	2.39	0.46
1:F:986:ASP:OD1	1:F:986:ASP:N	2.39	0.46
1:D:1032:ASP:C	1:D:1033:LEU:HD12	2.41	0.46
1:G:986:ASP:OD1	1:G:986:ASP:N	2.39	0.46
1:A:876:ILE:HA	1:A:879:TYR:CD2	2.51	0.46
1:F:913:PHE:CE2	1:G:916:GLN:HG3	2.39	0.46
1:F:1028:VAL:HG22	1:F:1042:LEU:HD13	1.96	0.46
1:G:876:ILE:HA	1:G:879:TYR:CD2	2.51	0.46
1:D:1026:PRO:HB3	1:D:1044:ILE:HG22	1.98	0.46
1:E:1026:PRO:HB3	1:E:1044:ILE:HG22	1.98	0.46
1:C:1026:PRO:HB3	1:C:1044:ILE:HG22	1.98	0.46
1:D:550:PRO:O	1:D:554:ALA:N	2.41	0.46
1:F:876:ILE:HA	1:F:879:TYR:CD2	2.51	0.46
1:G:1026:PRO:HB3	1:G:1044:ILE:HG22	1.98	0.46
1:E:1032:ASP:C	1:E:1033:LEU:HD12	2.41	0.45
1:A:1026:PRO:HB3	1:A:1044:ILE:HG22	1.98	0.45
1:B:1026:PRO:HB3	1:B:1044:ILE:HG22	1.98	0.45
1:D:955:ARG:NH1	1:E:931:GLU:OE1	2.49	0.45
1:F:1026:PRO:HB3	1:F:1044:ILE:HG22	1.98	0.45
1:A:1080:GLN:NE2	1:B:1084:GLU:OE2	2.39	0.45
1:B:876:ILE:HA	1:B:879:TYR:CD2	2.51	0.45
1:A:955:ARG:NH1	1:B:931:GLU:CD	2.71	0.45
1:C:876:ILE:HA	1:C:879:TYR:CD2	2.51	0.45
1:C:911:LEU:CD2	1:D:919:PHE:CD2	3.00	0.45
1:C:1032:ASP:C	1:C:1033:LEU:HD12	2.41	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:883:LEU:O	1:F:887:LEU:HG	2.17	0.45
1:G:1032:ASP:C	1:G:1033:LEU:HD12	2.41	0.45
1:A:955:ARG:NH1	1:B:931:GLU:OE1	2.49	0.45
1:A:1032:ASP:C	1:A:1033:LEU:HD12	2.41	0.45
1:A:913:PHE:HD2	1:B:916:GLN:CB	2.03	0.45
1:A:931:GLU:CD	1:G:955:ARG:NH1	2.71	0.45
1:F:1032:ASP:C	1:F:1033:LEU:HD12	2.41	0.45
1:B:883:LEU:O	1:B:887:LEU:HG	2.17	0.45
1:D:911:LEU:CD2	1:E:919:PHE:CD2	3.00	0.45
1:E:883:LEU:O	1:E:887:LEU:HG	2.17	0.45
1:E:955:ARG:NH1	1:F:931:GLU:OE1	2.50	0.45
1:B:550:PRO:O	1:B:554:ALA:N	2.41	0.45
1:B:1032:ASP:C	1:B:1033:LEU:HD12	2.41	0.45
1:C:955:ARG:NH1	1:D:931:GLU:OE1	2.50	0.45
1:F:955:ARG:NH1	1:G:931:GLU:OE1	2.49	0.45
1:G:935:ARG:CZ	1:G:935:ARG:HA	2.47	0.45
1:C:955:ARG:NH1	1:D:931:GLU:CD	2.71	0.44
1:A:883:LEU:O	1:A:887:LEU:HG	2.17	0.44
1:A:935:ARG:CZ	1:A:935:ARG:HA	2.47	0.44
1:A:942:ILE:HD11	1:A:973:ALA:HB1	2.00	0.44
1:B:582:PHE:HA	1:B:587:GLY:H	1.82	0.44
1:B:942:ILE:HD11	1:B:973:ALA:HB1	2.00	0.44
1:C:935:ARG:HA	1:C:935:ARG:CZ	2.47	0.44
1:C:942:ILE:HD11	1:C:973:ALA:HB1	2.00	0.44
1:D:942:ILE:HD11	1:D:973:ALA:HB1	2.00	0.44
1:E:911:LEU:CD2	1:F:919:PHE:CD2	2.99	0.44
1:G:883:LEU:O	1:G:887:LEU:HG	2.17	0.44
1:D:832:GLY:O	1:D:834:VAL:N	2.51	0.44
1:D:935:ARG:CZ	1:D:935:ARG:HA	2.47	0.44
1:D:955:ARG:NH1	1:E:931:GLU:CD	2.71	0.44
1:E:1011:LEU:HD23	1:E:1011:LEU:HA	1.82	0.44
1:F:832:GLY:O	1:F:834:VAL:N	2.51	0.44
1:F:935:ARG:CZ	1:F:935:ARG:HA	2.47	0.44
1:B:935:ARG:CZ	1:B:935:ARG:HA	2.47	0.44
1:C:1011:LEU:HD23	1:C:1011:LEU:HA	1.83	0.44
1:D:883:LEU:O	1:D:887:LEU:HG	2.17	0.44
1:F:911:LEU:CD2	1:G:919:PHE:CD2	3.00	0.44
1:G:471:THR:O	1:G:473:LYS:N	2.51	0.44
1:A:964:ARG:NH2	1:B:986:ASP:OD1	2.51	0.44
1:G:832:GLY:O	1:G:834:VAL:N	2.51	0.44
1:G:942:ILE:HD11	1:G:973:ALA:HB1	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:550:PRO:O	1:A:554:ALA:N	2.41	0.44
1:A:832:GLY:O	1:A:834:VAL:N	2.51	0.44
1:B:955:ARG:NH1	1:C:931:GLU:OE1	2.50	0.44
1:C:471:THR:O	1:C:473:LYS:N	2.51	0.44
1:E:935:ARG:HA	1:E:935:ARG:CZ	2.47	0.44
1:E:942:ILE:HD11	1:E:973:ALA:HB1	2.00	0.44
1:A:911:LEU:CD2	1:B:919:PHE:CD2	3.01	0.44
1:A:919:PHE:CD2	1:G:911:LEU:CD2	3.00	0.44
1:D:839:LEU:HD12	1:D:840:VAL:N	2.33	0.44
1:A:471:THR:O	1:A:473:LYS:N	2.51	0.44
1:A:931:GLU:OE1	1:G:955:ARG:NH1	2.50	0.44
1:B:458:THR:O	1:B:462:LEU:N	2.48	0.44
1:D:582:PHE:HA	1:D:587:GLY:H	1.82	0.44
1:F:471:THR:O	1:F:473:LYS:N	2.51	0.44
1:F:972:LYS:HE3	1:F:972:LYS:HB3	1.89	0.44
1:B:471:THR:O	1:B:473:LYS:N	2.51	0.43
1:B:818:SER:O	1:B:825:SER:N	2.48	0.43
1:B:839:LEU:HD12	1:B:840:VAL:N	2.33	0.43
1:C:582:PHE:HA	1:C:587:GLY:H	1.82	0.43
1:B:955:ARG:NH1	1:C:931:GLU:CD	2.72	0.43
1:C:883:LEU:O	1:C:887:LEU:HG	2.17	0.43
1:D:471:THR:O	1:D:473:LYS:N	2.51	0.43
1:G:582:PHE:HA	1:G:587:GLY:H	1.82	0.43
1:B:832:GLY:O	1:B:834:VAL:N	2.51	0.43
1:D:818:SER:O	1:D:825:SER:N	2.48	0.43
1:F:839:LEU:HD12	1:F:840:VAL:N	2.33	0.43
1:B:911:LEU:CD2	1:C:919:PHE:CD2	3.01	0.43
1:C:964:ARG:NH2	1:D:986:ASP:OD1	2.51	0.43
1:E:582:PHE:HA	1:E:587:GLY:H	1.82	0.43
1:E:832:GLY:O	1:E:834:VAL:N	2.51	0.43
1:E:839:LEU:HD12	1:E:840:VAL:N	2.33	0.43
1:F:945:LEU:HD23	1:F:945:LEU:HA	1.91	0.43
1:G:839:LEU:HD12	1:G:840:VAL:N	2.33	0.43
1:A:582:PHE:HA	1:A:587:GLY:H	1.82	0.43
1:C:832:GLY:O	1:C:834:VAL:N	2.51	0.43
1:C:836:ILE:HA	1:C:839:LEU:HG	2.01	0.43
1:E:471:THR:O	1:E:473:LYS:N	2.51	0.43
1:F:942:ILE:HD11	1:F:973:ALA:HB1	2.00	0.43
1:B:945:LEU:HD23	1:B:945:LEU:HA	1.91	0.43
1:B:964:ARG:NH2	1:C:986:ASP:OD1	2.52	0.43
1:E:972:LYS:HE3	1:E:972:LYS:HB3	1.89	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:582:PHE:HA	1:F:587:GLY:H	1.82	0.43
1:B:836:ILE:HA	1:B:839:LEU:HG	2.01	0.43
1:C:908:GLY:HA2	1:C:911:LEU:HD12	2.00	0.43
1:E:884:ILE:O	1:E:887:LEU:HD12	2.19	0.43
1:A:839:LEU:HD12	1:A:840:VAL:N	2.33	0.43
1:B:908:GLY:HA2	1:B:911:LEU:HD12	2.00	0.43
1:C:503:SER:O	1:C:507:VAL:N	2.33	0.43
1:A:978:GLN:NE2	1:G:943:ARG:HH11	2.17	0.43
1:C:818:SER:O	1:C:825:SER:N	2.48	0.43
1:D:972:LYS:HE3	1:D:972:LYS:HB3	1.89	0.43
1:A:1011:LEU:HD23	1:A:1011:LEU:HA	1.83	0.43
1:D:908:GLY:HA2	1:D:911:LEU:HD12	2.00	0.43
1:A:884:ILE:O	1:A:887:LEU:HD12	2.18	0.42
1:B:884:ILE:O	1:B:887:LEU:HD12	2.19	0.42
1:B:1011:LEU:HD23	1:B:1011:LEU:HA	1.82	0.42
1:C:839:LEU:HD12	1:C:840:VAL:N	2.33	0.42
1:A:836:ILE:HA	1:A:839:LEU:HG	2.01	0.42
1:A:908:GLY:HA2	1:A:911:LEU:HD12	2.00	0.42
1:A:913:PHE:CE2	1:B:916:GLN:HG3	2.40	0.42
1:D:836:ILE:HA	1:D:839:LEU:HG	2.01	0.42
1:E:955:ARG:NH1	1:F:931:GLU:CD	2.71	0.42
1:F:836:ILE:HA	1:F:839:LEU:HG	2.01	0.42
1:G:908:GLY:HA2	1:G:911:LEU:HD12	2.00	0.42
1:E:836:ILE:HA	1:E:839:LEU:HG	2.01	0.42
1:G:884:ILE:O	1:G:887:LEU:HD12	2.19	0.42
1:A:982:TRP:HH2	1:G:965:LYS:HE2	1.85	0.42
1:D:884:ILE:O	1:D:887:LEU:HD12	2.19	0.42
1:D:964:ARG:NH2	1:E:986:ASP:OD1	2.51	0.42
1:F:943:ARG:HH11	1:G:978:GLN:NE2	2.17	0.42
1:C:884:ILE:O	1:C:887:LEU:HD12	2.19	0.42
1:D:926:LEU:HD23	1:D:926:LEU:HA	1.88	0.42
1:A:986:ASP:OD1	1:G:964:ARG:NH2	2.52	0.42
1:E:943:ARG:HH11	1:F:978:GLN:NE2	2.17	0.42
1:F:884:ILE:O	1:F:887:LEU:HD12	2.19	0.42
1:F:908:GLY:HA2	1:F:911:LEU:HD12	2.00	0.42
1:F:926:LEU:HD23	1:F:926:LEU:HA	1.88	0.42
1:C:943:ARG:HH11	1:D:978:GLN:NE2	2.18	0.42
1:E:876:ILE:O	1:E:880:LEU:HD23	2.20	0.42
1:E:908:GLY:HA2	1:E:911:LEU:HD12	2.00	0.42
1:F:845:THR:O	1:F:849:ARG:HG2	2.20	0.42
1:F:876:ILE:O	1:F:880:LEU:HD23	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:943:ARG:HH11	1:E:978:GLN:NE2	2.18	0.42
1:E:845:THR:O	1:E:849:ARG:HG2	2.20	0.42
1:G:845:THR:O	1:G:849:ARG:HG2	2.20	0.42
1:G:876:ILE:O	1:G:880:LEU:HD23	2.20	0.42
1:A:458:THR:O	1:A:462:LEU:N	2.48	0.41
1:B:845:THR:O	1:B:849:ARG:HG2	2.20	0.41
1:B:876:ILE:O	1:B:880:LEU:HD23	2.20	0.41
1:F:832:GLY:C	1:F:836:ILE:HD13	2.46	0.41
1:G:836:ILE:HA	1:G:839:LEU:HG	2.01	0.41
1:A:945:LEU:HD11	1:A:967:ILE:HG21	2.02	0.41
1:B:965:LYS:HE2	1:C:982:TRP:HH2	1.86	0.41
1:D:845:THR:O	1:D:849:ARG:HG2	2.20	0.41
1:G:945:LEU:HD11	1:G:967:ILE:HG21	2.02	0.41
1:A:832:GLY:C	1:A:836:ILE:HD13	2.46	0.41
1:C:972:LYS:HE3	1:C:972:LYS:HB3	1.89	0.41
1:C:834:VAL:O	1:C:835:LEU:C	2.62	0.41
1:D:876:ILE:O	1:D:880:LEU:HD23	2.20	0.41
1:E:832:GLY:C	1:E:836:ILE:HD13	2.45	0.41
1:A:834:VAL:O	1:A:835:LEU:C	2.62	0.41
1:C:876:ILE:O	1:C:880:LEU:HD23	2.20	0.41
1:F:834:VAL:O	1:F:835:LEU:C	2.62	0.41
1:A:845:THR:O	1:A:849:ARG:HG2	2.20	0.41
1:B:945:LEU:HD11	1:B:967:ILE:HG21	2.02	0.41
1:D:834:VAL:O	1:D:835:LEU:C	2.62	0.41
1:E:964:ARG:NH2	1:F:986:ASP:OD1	2.51	0.41
1:F:945:LEU:HD11	1:F:967:ILE:HG21	2.02	0.41
1:G:832:GLY:C	1:G:836:ILE:HD13	2.46	0.41
1:G:945:LEU:HD23	1:G:945:LEU:HA	1.91	0.41
1:A:943:ARG:HH11	1:B:978:GLN:NE2	2.19	0.41
1:A:945:LEU:HD23	1:A:945:LEU:HA	1.91	0.41
1:E:458:THR:O	1:E:462:LEU:N	2.48	0.41
1:F:447:ALA:O	1:F:451:ARG:N	2.50	0.41
1:F:813:LEU:N	1:F:829:ILE:O	2.51	0.41
1:G:1011:LEU:HD23	1:G:1011:LEU:HA	1.82	0.41
1:A:916:GLN:HG3	1:G:913:PHE:CE2	2.39	0.41
1:A:965:LYS:HE2	1:B:982:TRP:HH2	1.85	0.41
1:B:834:VAL:O	1:B:835:LEU:C	2.62	0.41
1:C:845:THR:O	1:C:849:ARG:HG2	2.20	0.41
1:C:945:LEU:HD11	1:C:967:ILE:HG21	2.02	0.41
1:D:932:LYS:HZ3	1:D:932:LYS:HB2	1.85	0.41
1:D:965:LYS:HE2	1:E:982:TRP:HH2	1.85	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:965:LYS:HE2	1:F:982:TRP:HH2	1.86	0.41
1:F:964:ARG:NH2	1:G:986:ASP:OD1	2.51	0.41
1:F:965:LYS:HE2	1:G:982:TRP:HH2	1.85	0.41
1:D:978:GLN:H	1:D:978:GLN:HG2	1.59	0.41
1:G:550:PRO:O	1:G:554:ALA:N	2.41	0.41
1:G:932:LYS:HZ3	1:G:932:LYS:HB2	1.85	0.41
1:A:876:ILE:O	1:A:880:LEU:HD23	2.20	0.40
1:A:1081:MET:HE3	1:A:1081:MET:HB2	1.98	0.40
1:E:945:LEU:HD11	1:E:967:ILE:HG21	2.02	0.40
1:G:880:LEU:O	1:G:884:ILE:HG22	2.21	0.40
1:A:877:THR:HA	1:A:880:LEU:HD23	2.04	0.40
1:B:943:ARG:HH11	1:C:978:GLN:NE2	2.20	0.40
1:D:880:LEU:O	1:D:884:ILE:HG22	2.21	0.40
1:B:880:LEU:O	1:B:884:ILE:HG22	2.21	0.40
1:D:945:LEU:HD11	1:D:967:ILE:HG21	2.02	0.40
1:G:877:THR:HA	1:G:880:LEU:HD23	2.03	0.40
1:A:953:ASN:OD1	1:A:953:ASN:C	2.65	0.40
1:B:877:THR:HA	1:B:880:LEU:HD23	2.03	0.40
1:C:832:GLY:C	1:C:836:ILE:HD13	2.46	0.40
1:C:877:THR:HA	1:C:880:LEU:HD23	2.03	0.40
1:C:965:LYS:HE2	1:D:982:TRP:HH2	1.85	0.40
1:D:833:ALA:HB1	1:D:893:ILE:HA	2.04	0.40
1:E:833:ALA:HB1	1:E:893:ILE:HA	2.04	0.40
1:G:458:THR:O	1:G:462:LEU:N	2.48	0.40
1:B:926:LEU:HD23	1:B:926:LEU:HA	1.88	0.40
1:B:953:ASN:OD1	1:B:953:ASN:C	2.65	0.40
1:C:880:LEU:O	1:C:884:ILE:HG22	2.21	0.40
1:C:926:LEU:HD23	1:C:926:LEU:HA	1.88	0.40
1:E:503:SER:O	1:E:507:VAL:N	2.33	0.40
1:F:458:THR:O	1:F:462:LEU:N	2.48	0.40

There are no symmetry-related clashes.

5.3 Torsion angles

5.3.1 Protein backbone

In the following table, the Percentiles column shows the percent Ramachandran outliers of the chain as a percentile score with respect to all 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	A	785/873 (90%)	742 (94%)	37 (5%)	6 (1%)	16	53
1	B	785/873 (90%)	742 (94%)	37 (5%)	6 (1%)	16	53
1	C	785/873 (90%)	742 (94%)	37 (5%)	6 (1%)	16	53
1	D	785/873 (90%)	742 (94%)	37 (5%)	6 (1%)	16	53
1	E	785/873 (90%)	742 (94%)	37 (5%)	6 (1%)	16	53
1	F	785/873 (90%)	742 (94%)	37 (5%)	6 (1%)	16	53
1	G	785/873 (90%)	742 (94%)	37 (5%)	6 (1%)	16	53
All	All	5495/6111 (90%)	5194 (94%)	259 (5%)	42 (1%)	18	53

All (42) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	A	472	SER
1	A	585	HIS
1	A	859	ILE
1	B	472	SER
1	B	585	HIS
1	B	859	ILE
1	C	472	SER
1	C	585	HIS
1	C	859	ILE
1	D	472	SER
1	D	585	HIS
1	D	859	ILE
1	E	472	SER
1	E	585	HIS
1	E	859	ILE
1	F	472	SER
1	F	585	HIS
1	F	859	ILE
1	G	472	SER
1	G	585	HIS
1	G	859	ILE
1	A	913	PHE
1	B	913	PHE
1	C	913	PHE
1	D	913	PHE
1	E	913	PHE

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Mol	Chain	Res	Type
1	F	913	PHE
1	G	913	PHE
1	A	833	ALA
1	A	895	ILE
1	B	833	ALA
1	B	895	ILE
1	C	833	ALA
1	C	895	ILE
1	D	833	ALA
1	D	895	ILE
1	E	833	ALA
1	E	895	ILE
1	F	833	ALA
1	F	895	ILE
1	G	833	ALA
1	G	895	ILE

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	A	208/741 (28%)	207 (100%)	1 (0%)	81	82
1	B	208/741 (28%)	206 (99%)	2 (1%)	68	77
1	C	208/741 (28%)	207 (100%)	1 (0%)	81	82
1	D	208/741 (28%)	207 (100%)	1 (0%)	81	82
1	E	208/741 (28%)	207 (100%)	1 (0%)	81	82
1	F	208/741 (28%)	206 (99%)	2 (1%)	68	77
1	G	208/741 (28%)	207 (100%)	1 (0%)	81	82
All	All	1456/5187 (28%)	1447 (99%)	9 (1%)	76	82

All (9) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	834	VAL

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Mol	Chain	Res	Type
1	B	834	VAL
1	B	967	ILE
1	C	834	VAL
1	D	834	VAL
1	E	834	VAL
1	F	834	VAL
1	F	967	ILE
1	G	834	VAL

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (7) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	971	ASN
1	B	971	ASN
1	C	971	ASN
1	D	971	ASN
1	E	971	ASN
1	F	971	ASN
1	G	971	ASN

5.3.3 RNA [i](#)

There are no RNA molecules in this entry.

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

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

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

There are no ligands in this entry.

5.7 Other polymers

There are no such residues in this entry.

5.8 Polymer linkage issues

The following chains have linkage breaks:

Mol	Chain	Number of breaks
1	A	1
1	B	1
1	C	1
1	D	1
1	E	1
1	F	1
1	G	1

All chain breaks are listed below:

Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	A	740:GLU	C	769:VAL	N	14.63
1	B	740:GLU	C	769:VAL	N	14.63
1	C	740:GLU	C	769:VAL	N	14.63
1	D	740:GLU	C	769:VAL	N	14.63
1	E	740:GLU	C	769:VAL	N	14.63
1	F	740:GLU	C	769:VAL	N	14.63
1	G	740:GLU	C	769:VAL	N	14.63

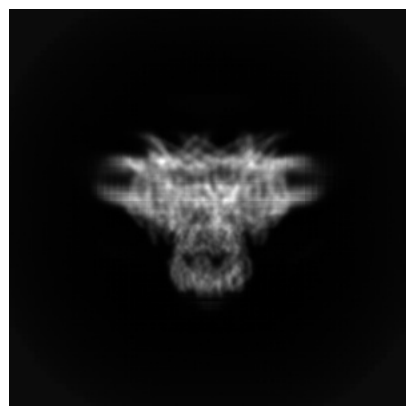
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-76018. These allow visual inspection of the internal detail of the map and identification of artifacts.

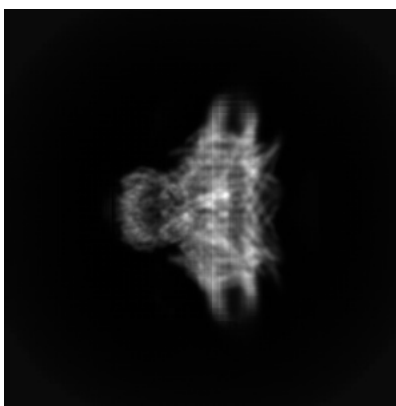
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections [i](#)

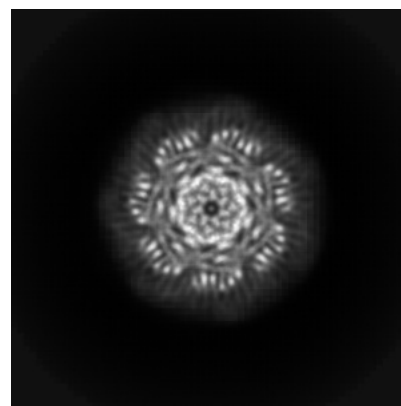
6.1.1 Primary map



X

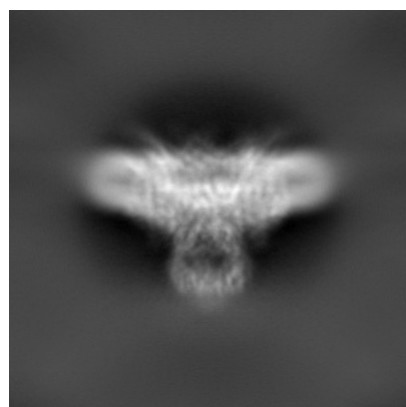


Y

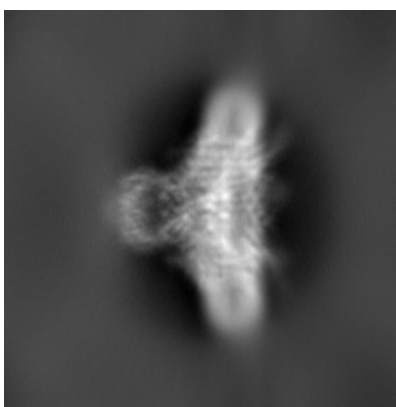


Z

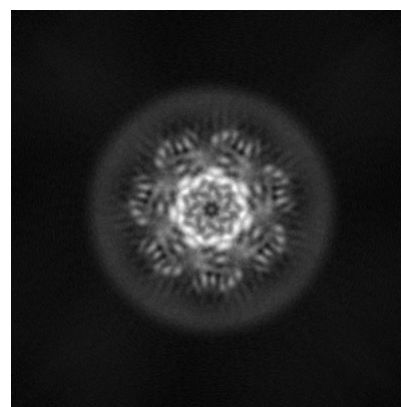
6.1.2 Raw map



X



Y



Z

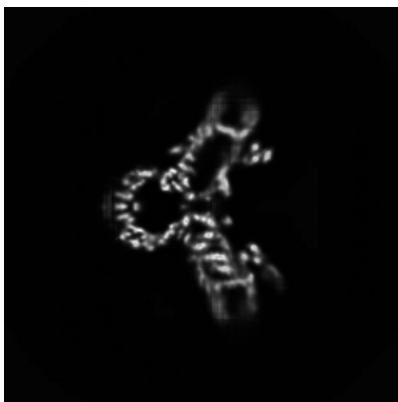
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

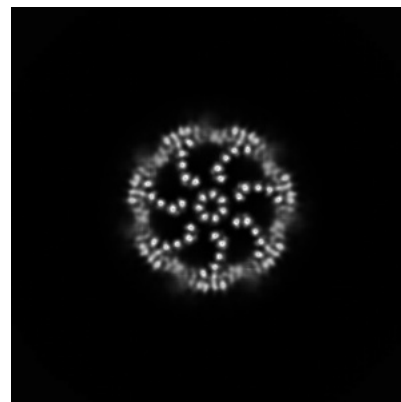
6.2.1 Primary map



X Index: 225

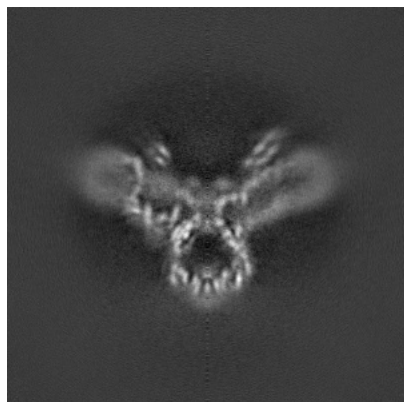


Y Index: 225

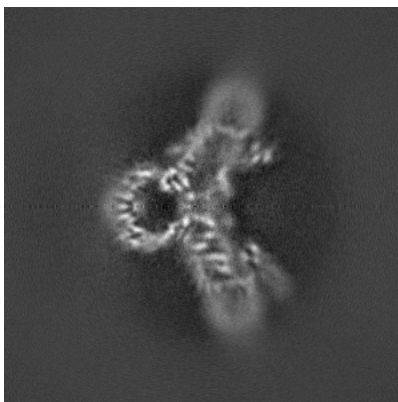


Z Index: 225

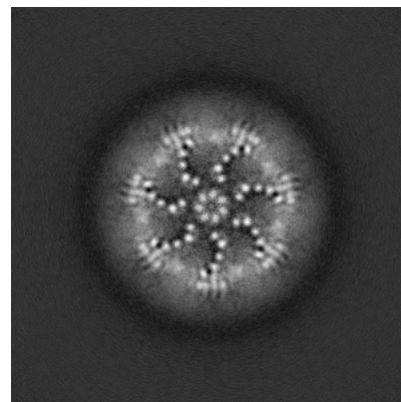
6.2.2 Raw map



X Index: 225



Y Index: 225



Z Index: 225

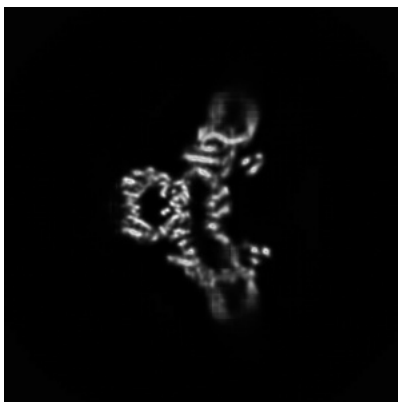
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

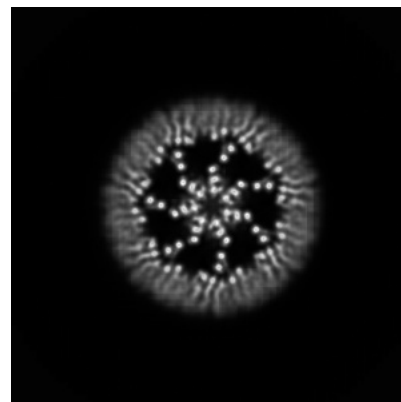
6.3.1 Primary map



X Index: 238

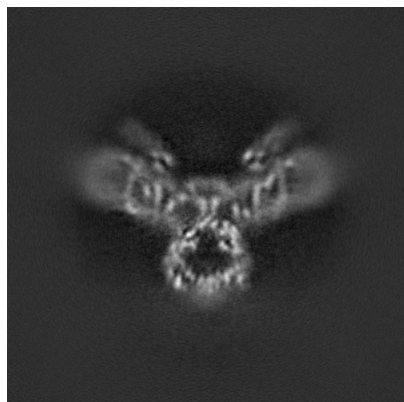


Y Index: 246

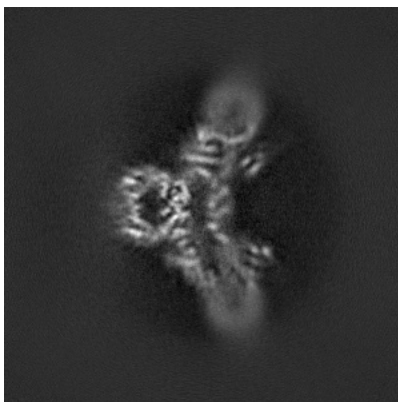


Z Index: 234

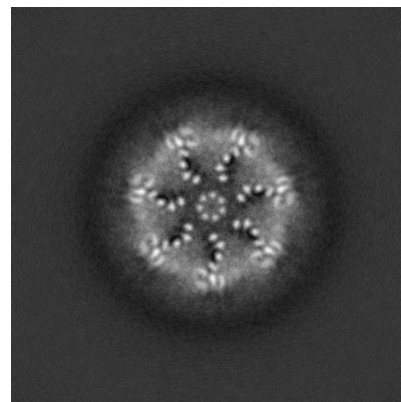
6.3.2 Raw map



X Index: 238



Y Index: 246



Z Index: 221

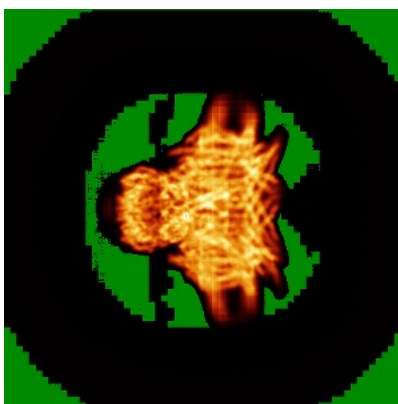
The images above show the largest variance slices of the map in three orthogonal directions.

6.4 Orthogonal standard-deviation projections (False-color) ⓘ

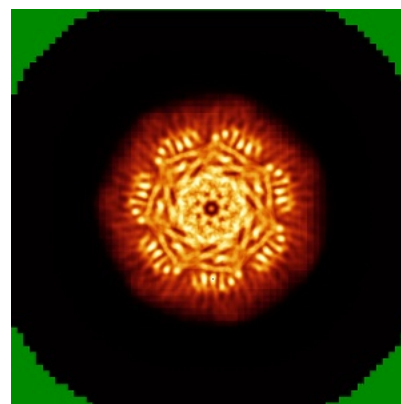
6.4.1 Primary map



X

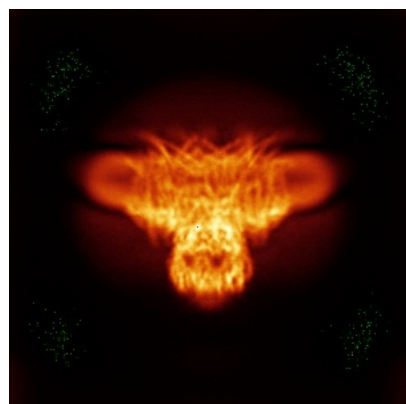


Y

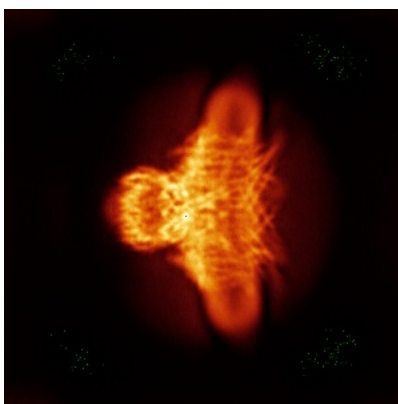


Z

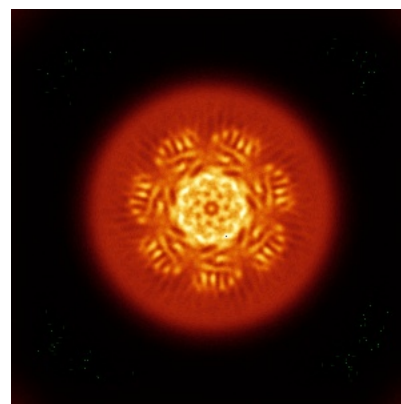
6.4.2 Raw map



X



Y

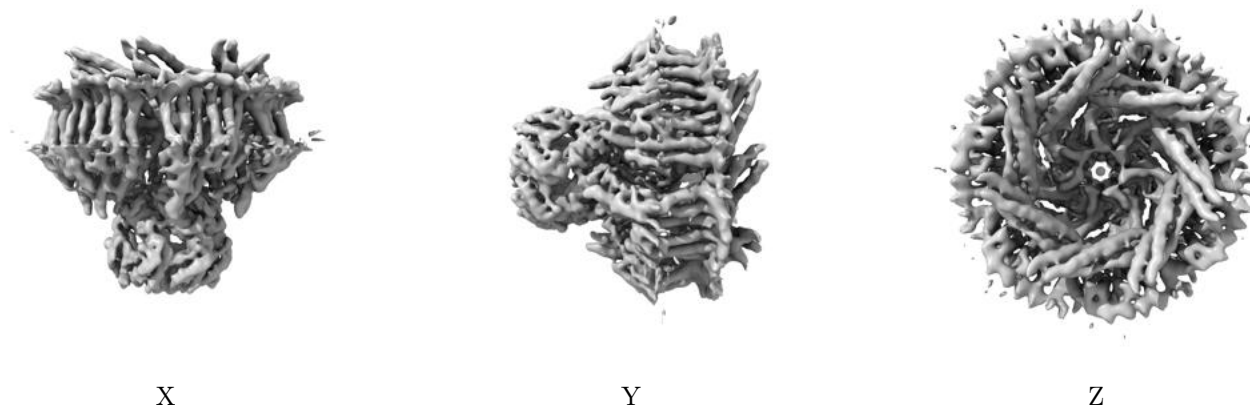


Z

The images above show the map standard deviation projections with false color in three orthogonal directions. Minimum values are shown in green, max in blue, and dark to light orange shades represent small to large values respectively.

6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 3.2. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

6.5.2 Raw map



These images show the 3D surface of the raw map. The raw map's contour level was selected so that its surface encloses the same volume as the primary map does at its recommended contour level.

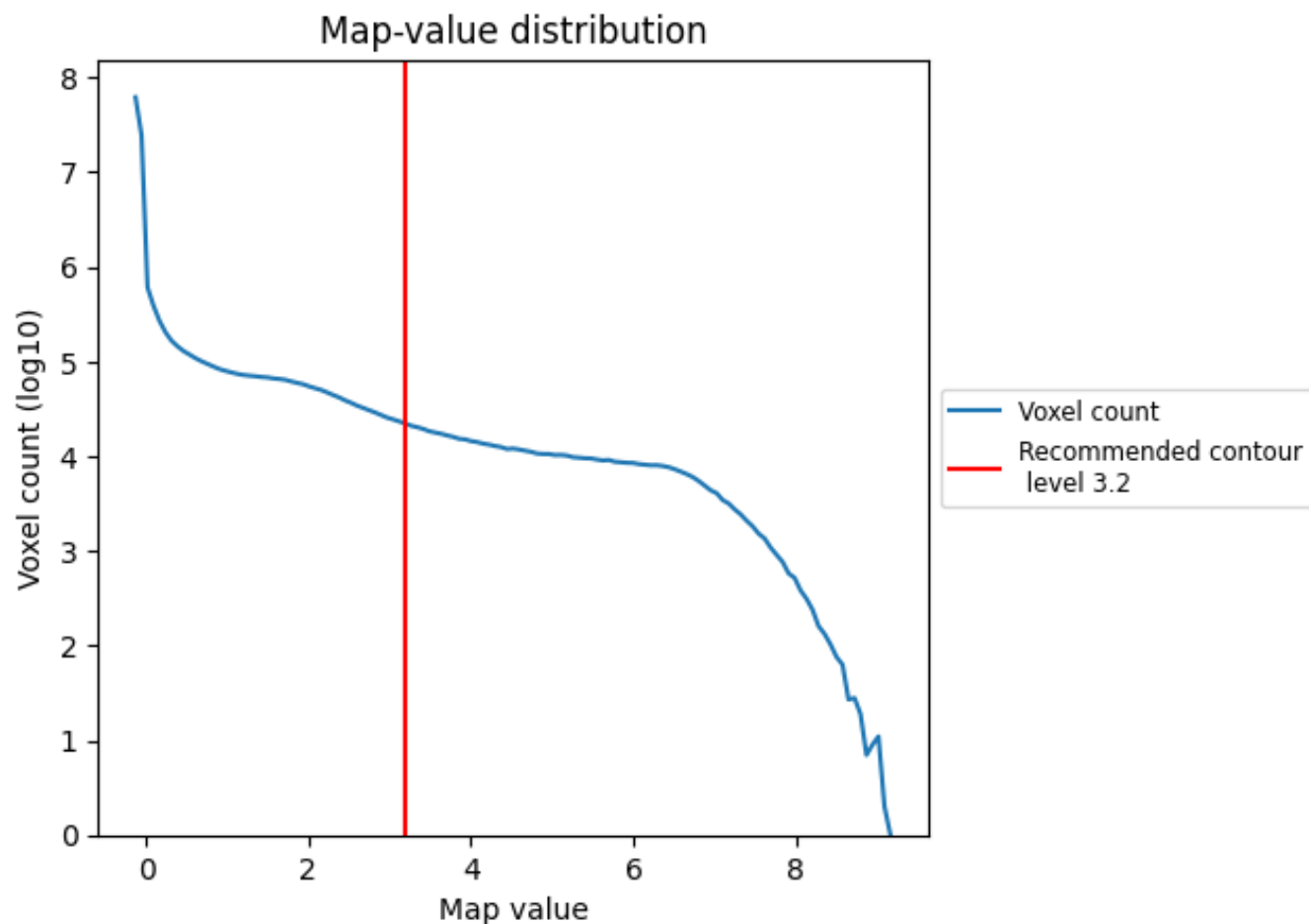
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

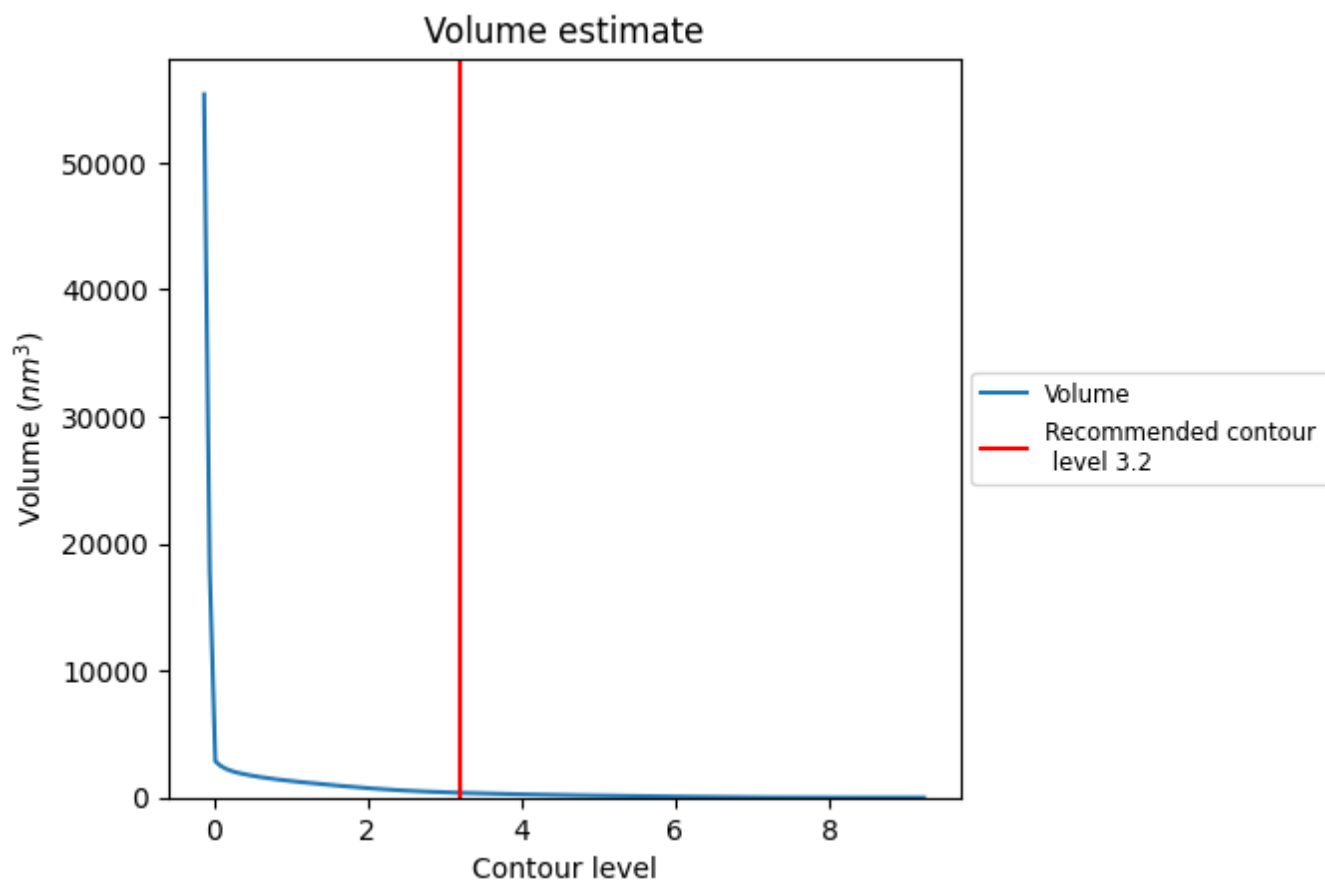
This section contains the results of statistical analysis of the map.

7.1 Map-value distribution [i](#)



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

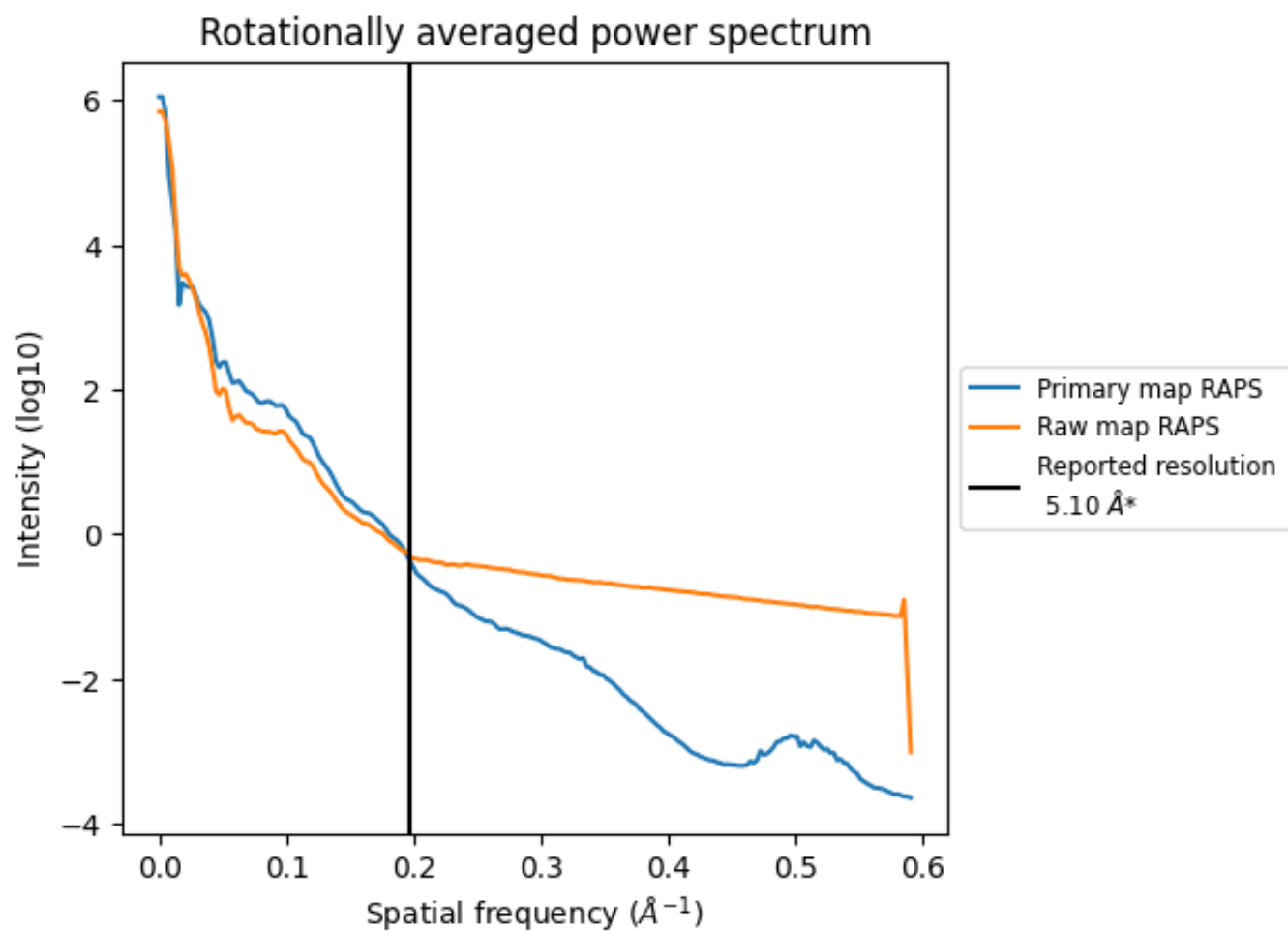
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 375 nm^3 ; this corresponds to an approximate mass of 339 kDa.

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

7.3 Rotationally averaged power spectrum ⓘ

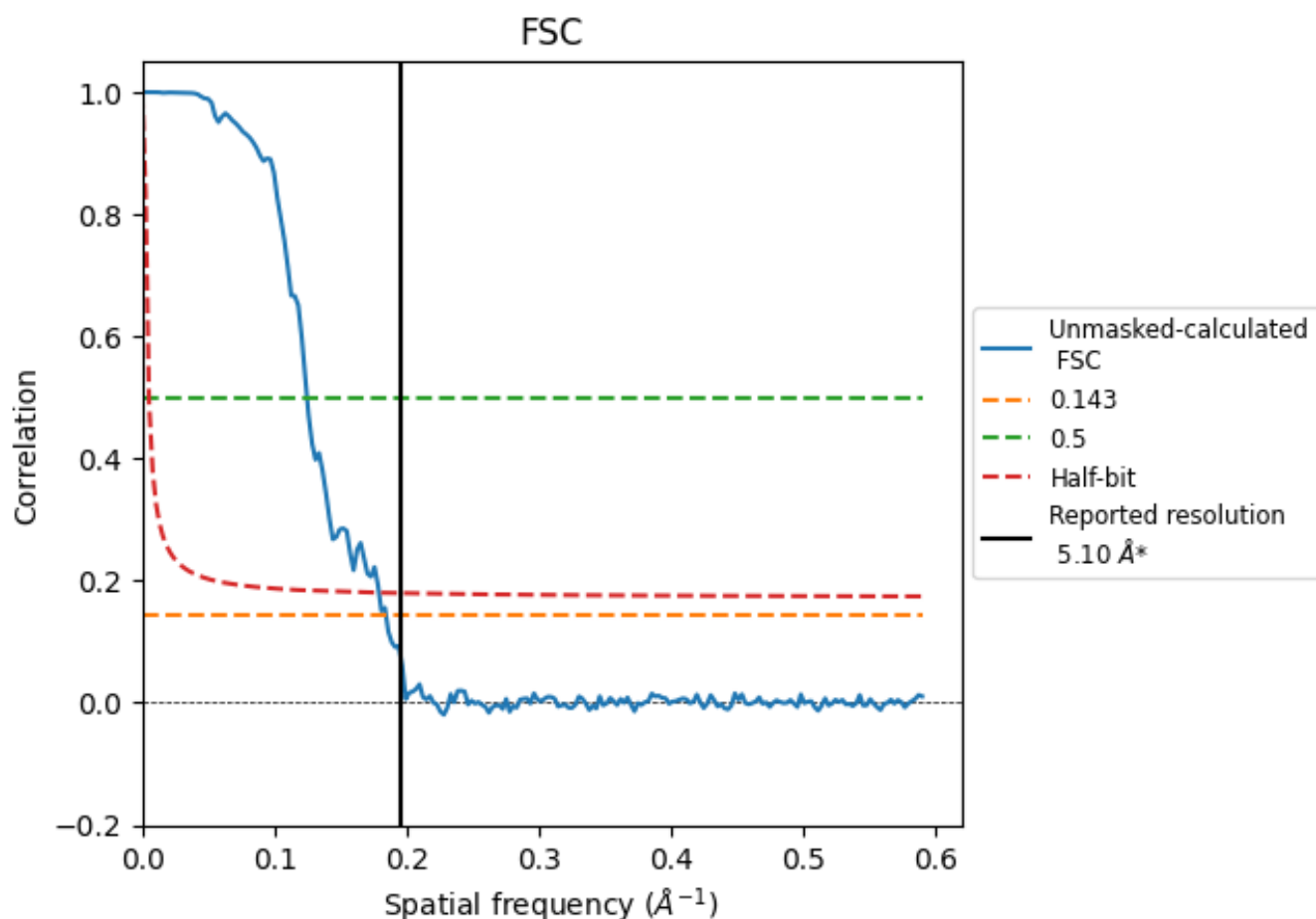


*Reported resolution corresponds to spatial frequency of 0.196 $\text{\AA}^{-1}$

8 Fourier-Shell correlation [i](#)

Fourier-Shell Correlation (FSC) is the most commonly used method to estimate the resolution of single-particle and subtomogram-averaged maps. The shape of the curve depends on the imposed symmetry, mask and whether or not the two 3D reconstructions used were processed from a common reference. The reported resolution is shown as a black line. A curve is displayed for the half-bit criterion in addition to lines showing the 0.143 gold standard cut-off and 0.5 cut-off.

8.1 FSC [i](#)



*Reported resolution corresponds to spatial frequency of 0.196 $\text{\AA}^{-1}$

8.2 Resolution estimates [i](#)

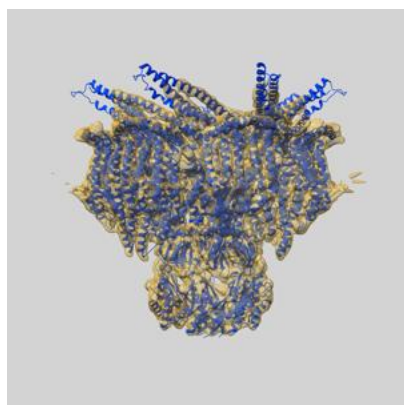
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	5.10	-	-
Author-provided FSC curve	-	-	-
Unmasked-calculated*	5.42	8.01	5.58

*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps.

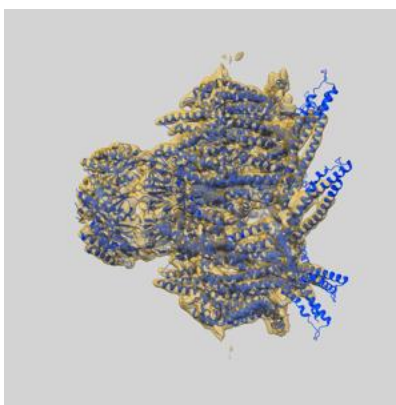
9 Map-model fit [i](#)

This section contains information regarding the fit between EMDB map EMD-76018 and PDB model 11SR. Per-residue inclusion information can be found in section [3](#) on page [46](#).

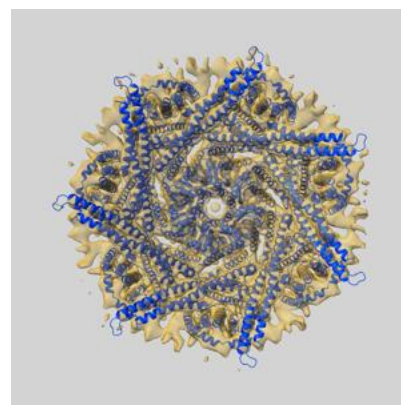
9.1 Map-model overlay [i](#)



X



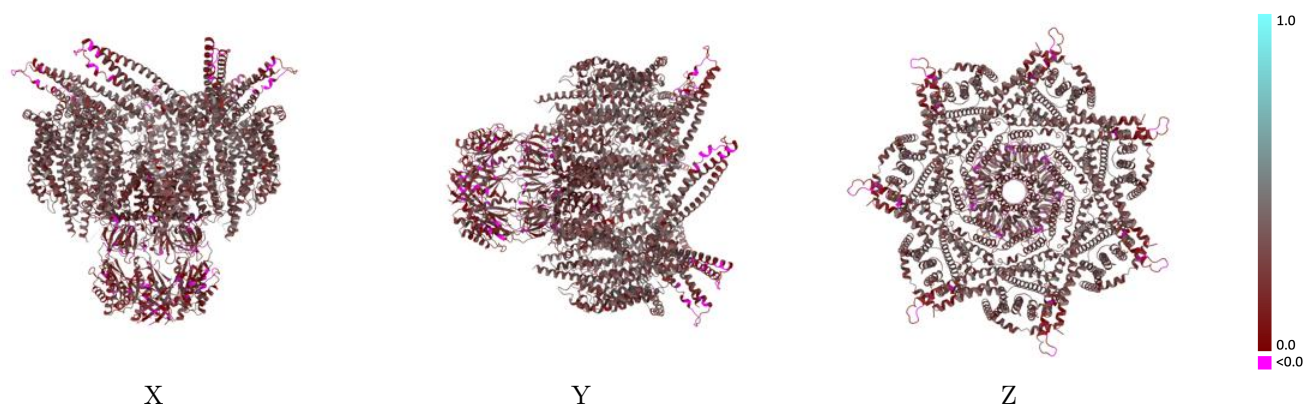
Y



Z

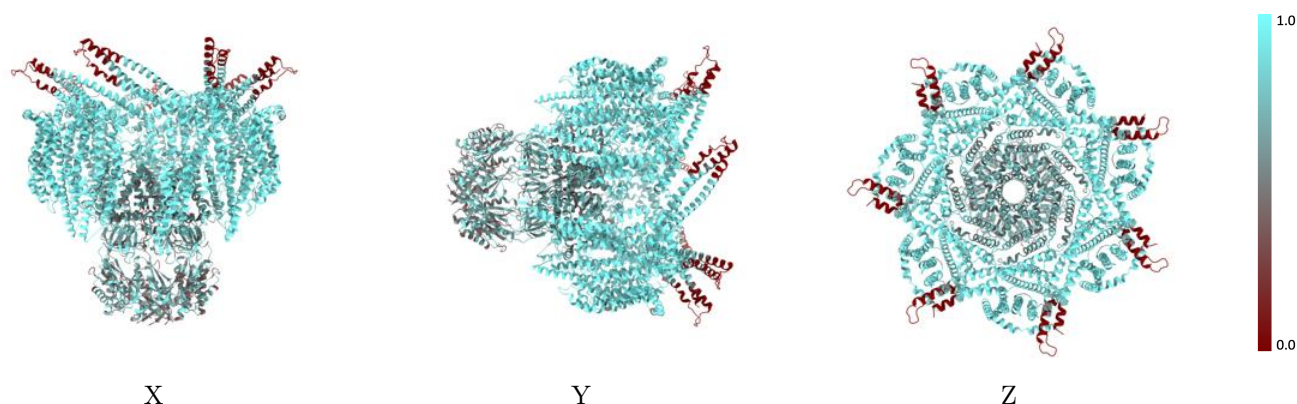
The images above show the 3D surface view of the map at the recommended contour level 3.2 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



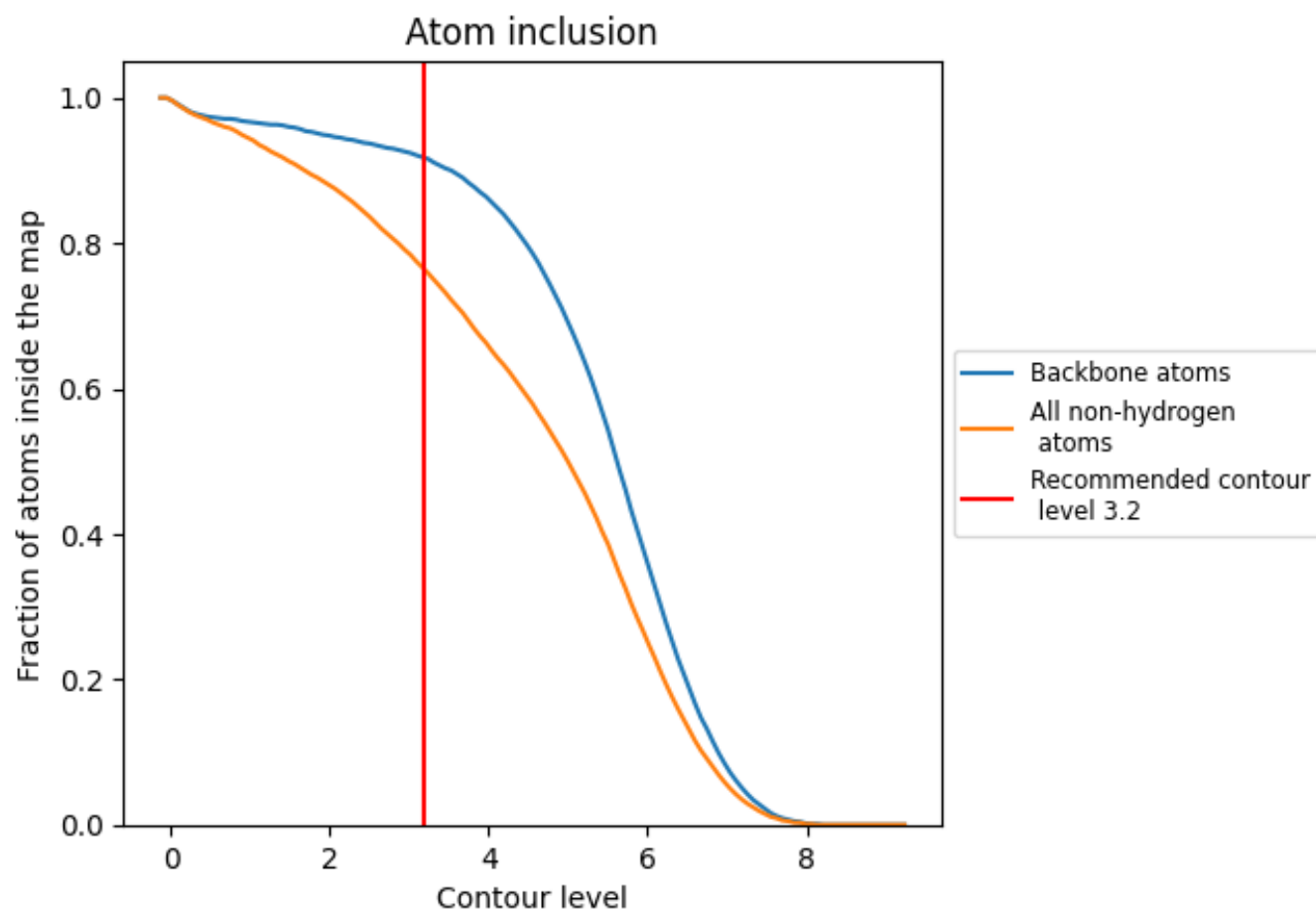
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions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

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



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (3.2).

9.4 Atom inclusion [i](#)



At the recommended contour level, 92% of all backbone atoms, 76% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary ⓘ

The table lists the average atom inclusion at the recommended contour level (3.2) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	0.7630	0.2640
A	0.7630	0.2640
B	0.7620	0.2650
C	0.7620	0.2630
D	0.7610	0.2640
E	0.7650	0.2640
F	0.7630	0.2630
G	0.7650	0.2650

1.0

0.0

<0.0