



## wwPDB EM Validation Summary Report ⓘ

Mar 11, 2025 – 06:09 pm GMT

PDB ID : 9FR3  
EMDB ID : EMD-50707  
Title : Structure of the SARS-CoV-2 spike glycoprotein in complex with nanobody 7F  
Authors : Debski-Antoniak, O.; Hurdiss, D.L.  
Deposited on : 2024-06-18  
Resolution : 3.30 Å(reported)  
Based on initial model : 7R40

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

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

A user guide is available at

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

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

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

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

EMDB validation analysis : **FAILED**  
Mogul : 1.8.4, CSD as541be (2020)  
MolProbity : 4.02b-467  
Percentile statistics : 20231227.v01 (using entries in the PDB archive December 27th 2023)  
MapQ : **FAILED**  
Ideal geometry (proteins) : Engh & Huber (2001)  
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)  
Validation Pipeline (wwPDB-VP) : 2.41

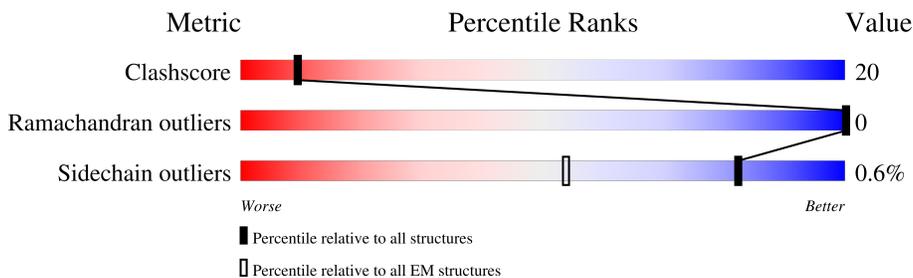
# 1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

*ELECTRON MICROSCOPY*

The reported resolution of this entry is 3.30 Å.

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



Metric	Whole archive (#Entries)	EM structures (#Entries)
Clashscore	210492	15764
Ramachandran outliers	207382	16835
Sidechain outliers	206894	16415

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

Mol	Chain	Length	Quality of chain
1	C	148	
1	E	148	
1	F	148	
1	I	148	
1	K	148	
1	L	148	
2	A	1275	
2	B	1275	
2	D	1275	

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Mol	Chain	Length	Quality of chain			
2	G	1275		57%	20%	23%
2	H	1275		56%	21%	23%
2	J	1275		57%	20%	23%

## 2 Entry composition [i](#)

There are 3 unique types of molecules in this entry. The entry contains 51750 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 Nanobody 7F.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
1	C	119	933	589	156	183	5	0	0
1	E	119	933	589	156	183	5	0	0
1	F	119	933	589	156	183	5	0	0
1	I	119	933	589	156	183	5	0	0
1	K	119	933	589	156	183	5	0	0
1	L	119	933	589	156	183	5	0	0

- Molecule 2 is a protein called Spike glycoprotein.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
2	A	986	7594	4864	1261	1435	34	0	0
2	B	986	7594	4864	1261	1435	34	0	0
2	D	986	7594	4864	1261	1435	34	0	0
2	G	986	7594	4864	1261	1435	34	0	0
2	H	986	7594	4864	1261	1435	34	0	0
2	J	986	7594	4864	1261	1435	34	0	0

There are 534 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
A	393	PHE	THR	conflict	UNP P0DTC2
A	607	GLU	GLN	conflict	UNP P0DTC2

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Chain	Residue	Modelled	Actual	Comment	Reference
A	682	ALA	ARG	conflict	UNP P0DTC2
A	683	ALA	ARG	conflict	UNP P0DTC2
A	892	PRO	ALA	conflict	UNP P0DTC2
A	899	PRO	ALA	conflict	UNP P0DTC2
A	942	PRO	ALA	conflict	UNP P0DTC2
A	986	PRO	LYS	conflict	UNP P0DTC2
A	987	PRO	VAL	conflict	UNP P0DTC2
A	1209	GLY	-	expression tag	UNP P0DTC2
A	1210	SER	-	expression tag	UNP P0DTC2
A	1211	GLY	-	expression tag	UNP P0DTC2
A	1212	TYR	-	expression tag	UNP P0DTC2
A	1213	ILE	-	expression tag	UNP P0DTC2
A	1214	PRO	-	expression tag	UNP P0DTC2
A	1215	GLU	-	expression tag	UNP P0DTC2
A	1216	ALA	-	expression tag	UNP P0DTC2
A	1217	PRO	-	expression tag	UNP P0DTC2
A	1218	ARG	-	expression tag	UNP P0DTC2
A	1219	ASP	-	expression tag	UNP P0DTC2
A	1220	GLY	-	expression tag	UNP P0DTC2
A	1221	GLN	-	expression tag	UNP P0DTC2
A	1222	ALA	-	expression tag	UNP P0DTC2
A	1223	TYR	-	expression tag	UNP P0DTC2
A	1224	VAL	-	expression tag	UNP P0DTC2
A	1225	ARG	-	expression tag	UNP P0DTC2
A	1226	LYS	-	expression tag	UNP P0DTC2
A	1227	ASP	-	expression tag	UNP P0DTC2
A	1228	GLY	-	expression tag	UNP P0DTC2
A	1229	GLU	-	expression tag	UNP P0DTC2
A	1230	TRP	-	expression tag	UNP P0DTC2
A	1231	VAL	-	expression tag	UNP P0DTC2
A	1232	LEU	-	expression tag	UNP P0DTC2
A	1233	LEU	-	expression tag	UNP P0DTC2
A	1234	SER	-	expression tag	UNP P0DTC2
A	1235	THR	-	expression tag	UNP P0DTC2
A	1236	PHE	-	expression tag	UNP P0DTC2
A	1237	LEU	-	expression tag	UNP P0DTC2
A	1238	GLY	-	expression tag	UNP P0DTC2
A	1239	ARG	-	expression tag	UNP P0DTC2
A	1240	SER	-	expression tag	UNP P0DTC2
A	1241	LEU	-	expression tag	UNP P0DTC2
A	1242	GLU	-	expression tag	UNP P0DTC2
A	1243	VAL	-	expression tag	UNP P0DTC2

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Chain	Residue	Modelled	Actual	Comment	Reference
A	1244	LEU	-	expression tag	UNP P0DTC2
A	1245	PHE	-	expression tag	UNP P0DTC2
A	1246	GLN	-	expression tag	UNP P0DTC2
A	1247	GLY	-	expression tag	UNP P0DTC2
A	1248	PRO	-	expression tag	UNP P0DTC2
A	1249	GLY	-	expression tag	UNP P0DTC2
A	1250	HIS	-	expression tag	UNP P0DTC2
A	1251	HIS	-	expression tag	UNP P0DTC2
A	1252	HIS	-	expression tag	UNP P0DTC2
A	1253	HIS	-	expression tag	UNP P0DTC2
A	1254	HIS	-	expression tag	UNP P0DTC2
A	1255	HIS	-	expression tag	UNP P0DTC2
A	1256	HIS	-	expression tag	UNP P0DTC2
A	1257	HIS	-	expression tag	UNP P0DTC2
A	1258	SER	-	expression tag	UNP P0DTC2
A	1259	ALA	-	expression tag	UNP P0DTC2
A	1260	TRP	-	expression tag	UNP P0DTC2
A	1261	SER	-	expression tag	UNP P0DTC2
A	1262	HIS	-	expression tag	UNP P0DTC2
A	1263	PRO	-	expression tag	UNP P0DTC2
A	1264	GLN	-	expression tag	UNP P0DTC2
A	1265	PHE	-	expression tag	UNP P0DTC2
A	1266	GLU	-	expression tag	UNP P0DTC2
A	1267	LYS	-	expression tag	UNP P0DTC2
A	1268	GLY	-	expression tag	UNP P0DTC2
A	1269	GLY	-	expression tag	UNP P0DTC2
A	1270	GLY	-	expression tag	UNP P0DTC2
A	1271	SER	-	expression tag	UNP P0DTC2
A	1272	GLY	-	expression tag	UNP P0DTC2
A	1273	GLY	-	expression tag	UNP P0DTC2
A	1274	GLY	-	expression tag	UNP P0DTC2
A	1275	GLY	-	expression tag	UNP P0DTC2
A	1276	SER	-	expression tag	UNP P0DTC2
A	1277	GLY	-	expression tag	UNP P0DTC2
A	1278	GLY	-	expression tag	UNP P0DTC2
A	1279	SER	-	expression tag	UNP P0DTC2
A	1280	ALA	-	expression tag	UNP P0DTC2
A	1281	TRP	-	expression tag	UNP P0DTC2
A	1282	SER	-	expression tag	UNP P0DTC2
A	1283	HIS	-	expression tag	UNP P0DTC2
A	1284	PRO	-	expression tag	UNP P0DTC2
A	1285	GLN	-	expression tag	UNP P0DTC2

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Chain	Residue	Modelled	Actual	Comment	Reference
A	1286	PHE	-	expression tag	UNP P0DTC2
A	1287	GLU	-	expression tag	UNP P0DTC2
A	1288	LYS	-	expression tag	UNP P0DTC2
B	393	PHE	THR	conflict	UNP P0DTC2
B	607	GLU	GLN	conflict	UNP P0DTC2
B	682	ALA	ARG	conflict	UNP P0DTC2
B	683	ALA	ARG	conflict	UNP P0DTC2
B	892	PRO	ALA	conflict	UNP P0DTC2
B	899	PRO	ALA	conflict	UNP P0DTC2
B	942	PRO	ALA	conflict	UNP P0DTC2
B	986	PRO	LYS	conflict	UNP P0DTC2
B	987	PRO	VAL	conflict	UNP P0DTC2
B	1209	GLY	-	expression tag	UNP P0DTC2
B	1210	SER	-	expression tag	UNP P0DTC2
B	1211	GLY	-	expression tag	UNP P0DTC2
B	1212	TYR	-	expression tag	UNP P0DTC2
B	1213	ILE	-	expression tag	UNP P0DTC2
B	1214	PRO	-	expression tag	UNP P0DTC2
B	1215	GLU	-	expression tag	UNP P0DTC2
B	1216	ALA	-	expression tag	UNP P0DTC2
B	1217	PRO	-	expression tag	UNP P0DTC2
B	1218	ARG	-	expression tag	UNP P0DTC2
B	1219	ASP	-	expression tag	UNP P0DTC2
B	1220	GLY	-	expression tag	UNP P0DTC2
B	1221	GLN	-	expression tag	UNP P0DTC2
B	1222	ALA	-	expression tag	UNP P0DTC2
B	1223	TYR	-	expression tag	UNP P0DTC2
B	1224	VAL	-	expression tag	UNP P0DTC2
B	1225	ARG	-	expression tag	UNP P0DTC2
B	1226	LYS	-	expression tag	UNP P0DTC2
B	1227	ASP	-	expression tag	UNP P0DTC2
B	1228	GLY	-	expression tag	UNP P0DTC2
B	1229	GLU	-	expression tag	UNP P0DTC2
B	1230	TRP	-	expression tag	UNP P0DTC2
B	1231	VAL	-	expression tag	UNP P0DTC2
B	1232	LEU	-	expression tag	UNP P0DTC2
B	1233	LEU	-	expression tag	UNP P0DTC2
B	1234	SER	-	expression tag	UNP P0DTC2
B	1235	THR	-	expression tag	UNP P0DTC2
B	1236	PHE	-	expression tag	UNP P0DTC2
B	1237	LEU	-	expression tag	UNP P0DTC2
B	1238	GLY	-	expression tag	UNP P0DTC2

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Chain	Residue	Modelled	Actual	Comment	Reference
B	1239	ARG	-	expression tag	UNP P0DTC2
B	1240	SER	-	expression tag	UNP P0DTC2
B	1241	LEU	-	expression tag	UNP P0DTC2
B	1242	GLU	-	expression tag	UNP P0DTC2
B	1243	VAL	-	expression tag	UNP P0DTC2
B	1244	LEU	-	expression tag	UNP P0DTC2
B	1245	PHE	-	expression tag	UNP P0DTC2
B	1246	GLN	-	expression tag	UNP P0DTC2
B	1247	GLY	-	expression tag	UNP P0DTC2
B	1248	PRO	-	expression tag	UNP P0DTC2
B	1249	GLY	-	expression tag	UNP P0DTC2
B	1250	HIS	-	expression tag	UNP P0DTC2
B	1251	HIS	-	expression tag	UNP P0DTC2
B	1252	HIS	-	expression tag	UNP P0DTC2
B	1253	HIS	-	expression tag	UNP P0DTC2
B	1254	HIS	-	expression tag	UNP P0DTC2
B	1255	HIS	-	expression tag	UNP P0DTC2
B	1256	HIS	-	expression tag	UNP P0DTC2
B	1257	HIS	-	expression tag	UNP P0DTC2
B	1258	SER	-	expression tag	UNP P0DTC2
B	1259	ALA	-	expression tag	UNP P0DTC2
B	1260	TRP	-	expression tag	UNP P0DTC2
B	1261	SER	-	expression tag	UNP P0DTC2
B	1262	HIS	-	expression tag	UNP P0DTC2
B	1263	PRO	-	expression tag	UNP P0DTC2
B	1264	GLN	-	expression tag	UNP P0DTC2
B	1265	PHE	-	expression tag	UNP P0DTC2
B	1266	GLU	-	expression tag	UNP P0DTC2
B	1267	LYS	-	expression tag	UNP P0DTC2
B	1268	GLY	-	expression tag	UNP P0DTC2
B	1269	GLY	-	expression tag	UNP P0DTC2
B	1270	GLY	-	expression tag	UNP P0DTC2
B	1271	SER	-	expression tag	UNP P0DTC2
B	1272	GLY	-	expression tag	UNP P0DTC2
B	1273	GLY	-	expression tag	UNP P0DTC2
B	1274	GLY	-	expression tag	UNP P0DTC2
B	1275	GLY	-	expression tag	UNP P0DTC2
B	1276	SER	-	expression tag	UNP P0DTC2
B	1277	GLY	-	expression tag	UNP P0DTC2
B	1278	GLY	-	expression tag	UNP P0DTC2
B	1279	SER	-	expression tag	UNP P0DTC2
B	1280	ALA	-	expression tag	UNP P0DTC2

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Chain	Residue	Modelled	Actual	Comment	Reference
B	1281	TRP	-	expression tag	UNP P0DTC2
B	1282	SER	-	expression tag	UNP P0DTC2
B	1283	HIS	-	expression tag	UNP P0DTC2
B	1284	PRO	-	expression tag	UNP P0DTC2
B	1285	GLN	-	expression tag	UNP P0DTC2
B	1286	PHE	-	expression tag	UNP P0DTC2
B	1287	GLU	-	expression tag	UNP P0DTC2
B	1288	LYS	-	expression tag	UNP P0DTC2
D	393	PHE	THR	conflict	UNP P0DTC2
D	607	GLU	GLN	conflict	UNP P0DTC2
D	682	ALA	ARG	conflict	UNP P0DTC2
D	683	ALA	ARG	conflict	UNP P0DTC2
D	892	PRO	ALA	conflict	UNP P0DTC2
D	899	PRO	ALA	conflict	UNP P0DTC2
D	942	PRO	ALA	conflict	UNP P0DTC2
D	986	PRO	LYS	conflict	UNP P0DTC2
D	987	PRO	VAL	conflict	UNP P0DTC2
D	1209	GLY	-	expression tag	UNP P0DTC2
D	1210	SER	-	expression tag	UNP P0DTC2
D	1211	GLY	-	expression tag	UNP P0DTC2
D	1212	TYR	-	expression tag	UNP P0DTC2
D	1213	ILE	-	expression tag	UNP P0DTC2
D	1214	PRO	-	expression tag	UNP P0DTC2
D	1215	GLU	-	expression tag	UNP P0DTC2
D	1216	ALA	-	expression tag	UNP P0DTC2
D	1217	PRO	-	expression tag	UNP P0DTC2
D	1218	ARG	-	expression tag	UNP P0DTC2
D	1219	ASP	-	expression tag	UNP P0DTC2
D	1220	GLY	-	expression tag	UNP P0DTC2
D	1221	GLN	-	expression tag	UNP P0DTC2
D	1222	ALA	-	expression tag	UNP P0DTC2
D	1223	TYR	-	expression tag	UNP P0DTC2
D	1224	VAL	-	expression tag	UNP P0DTC2
D	1225	ARG	-	expression tag	UNP P0DTC2
D	1226	LYS	-	expression tag	UNP P0DTC2
D	1227	ASP	-	expression tag	UNP P0DTC2
D	1228	GLY	-	expression tag	UNP P0DTC2
D	1229	GLU	-	expression tag	UNP P0DTC2
D	1230	TRP	-	expression tag	UNP P0DTC2
D	1231	VAL	-	expression tag	UNP P0DTC2
D	1232	LEU	-	expression tag	UNP P0DTC2
D	1233	LEU	-	expression tag	UNP P0DTC2

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Chain	Residue	Modelled	Actual	Comment	Reference
D	1234	SER	-	expression tag	UNP P0DTC2
D	1235	THR	-	expression tag	UNP P0DTC2
D	1236	PHE	-	expression tag	UNP P0DTC2
D	1237	LEU	-	expression tag	UNP P0DTC2
D	1238	GLY	-	expression tag	UNP P0DTC2
D	1239	ARG	-	expression tag	UNP P0DTC2
D	1240	SER	-	expression tag	UNP P0DTC2
D	1241	LEU	-	expression tag	UNP P0DTC2
D	1242	GLU	-	expression tag	UNP P0DTC2
D	1243	VAL	-	expression tag	UNP P0DTC2
D	1244	LEU	-	expression tag	UNP P0DTC2
D	1245	PHE	-	expression tag	UNP P0DTC2
D	1246	GLN	-	expression tag	UNP P0DTC2
D	1247	GLY	-	expression tag	UNP P0DTC2
D	1248	PRO	-	expression tag	UNP P0DTC2
D	1249	GLY	-	expression tag	UNP P0DTC2
D	1250	HIS	-	expression tag	UNP P0DTC2
D	1251	HIS	-	expression tag	UNP P0DTC2
D	1252	HIS	-	expression tag	UNP P0DTC2
D	1253	HIS	-	expression tag	UNP P0DTC2
D	1254	HIS	-	expression tag	UNP P0DTC2
D	1255	HIS	-	expression tag	UNP P0DTC2
D	1256	HIS	-	expression tag	UNP P0DTC2
D	1257	HIS	-	expression tag	UNP P0DTC2
D	1258	SER	-	expression tag	UNP P0DTC2
D	1259	ALA	-	expression tag	UNP P0DTC2
D	1260	TRP	-	expression tag	UNP P0DTC2
D	1261	SER	-	expression tag	UNP P0DTC2
D	1262	HIS	-	expression tag	UNP P0DTC2
D	1263	PRO	-	expression tag	UNP P0DTC2
D	1264	GLN	-	expression tag	UNP P0DTC2
D	1265	PHE	-	expression tag	UNP P0DTC2
D	1266	GLU	-	expression tag	UNP P0DTC2
D	1267	LYS	-	expression tag	UNP P0DTC2
D	1268	GLY	-	expression tag	UNP P0DTC2
D	1269	GLY	-	expression tag	UNP P0DTC2
D	1270	GLY	-	expression tag	UNP P0DTC2
D	1271	SER	-	expression tag	UNP P0DTC2
D	1272	GLY	-	expression tag	UNP P0DTC2
D	1273	GLY	-	expression tag	UNP P0DTC2
D	1274	GLY	-	expression tag	UNP P0DTC2
D	1275	GLY	-	expression tag	UNP P0DTC2

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Chain	Residue	Modelled	Actual	Comment	Reference
D	1276	SER	-	expression tag	UNP P0DTC2
D	1277	GLY	-	expression tag	UNP P0DTC2
D	1278	GLY	-	expression tag	UNP P0DTC2
D	1279	SER	-	expression tag	UNP P0DTC2
D	1280	ALA	-	expression tag	UNP P0DTC2
D	1281	TRP	-	expression tag	UNP P0DTC2
D	1282	SER	-	expression tag	UNP P0DTC2
D	1283	HIS	-	expression tag	UNP P0DTC2
D	1284	PRO	-	expression tag	UNP P0DTC2
D	1285	GLN	-	expression tag	UNP P0DTC2
D	1286	PHE	-	expression tag	UNP P0DTC2
D	1287	GLU	-	expression tag	UNP P0DTC2
D	1288	LYS	-	expression tag	UNP P0DTC2
G	393	PHE	THR	conflict	UNP P0DTC2
G	607	GLU	GLN	conflict	UNP P0DTC2
G	682	ALA	ARG	conflict	UNP P0DTC2
G	683	ALA	ARG	conflict	UNP P0DTC2
G	892	PRO	ALA	conflict	UNP P0DTC2
G	899	PRO	ALA	conflict	UNP P0DTC2
G	942	PRO	ALA	conflict	UNP P0DTC2
G	986	PRO	LYS	conflict	UNP P0DTC2
G	987	PRO	VAL	conflict	UNP P0DTC2
G	1209	GLY	-	expression tag	UNP P0DTC2
G	1210	SER	-	expression tag	UNP P0DTC2
G	1211	GLY	-	expression tag	UNP P0DTC2
G	1212	TYR	-	expression tag	UNP P0DTC2
G	1213	ILE	-	expression tag	UNP P0DTC2
G	1214	PRO	-	expression tag	UNP P0DTC2
G	1215	GLU	-	expression tag	UNP P0DTC2
G	1216	ALA	-	expression tag	UNP P0DTC2
G	1217	PRO	-	expression tag	UNP P0DTC2
G	1218	ARG	-	expression tag	UNP P0DTC2
G	1219	ASP	-	expression tag	UNP P0DTC2
G	1220	GLY	-	expression tag	UNP P0DTC2
G	1221	GLN	-	expression tag	UNP P0DTC2
G	1222	ALA	-	expression tag	UNP P0DTC2
G	1223	TYR	-	expression tag	UNP P0DTC2
G	1224	VAL	-	expression tag	UNP P0DTC2
G	1225	ARG	-	expression tag	UNP P0DTC2
G	1226	LYS	-	expression tag	UNP P0DTC2
G	1227	ASP	-	expression tag	UNP P0DTC2
G	1228	GLY	-	expression tag	UNP P0DTC2

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Chain	Residue	Modelled	Actual	Comment	Reference
G	1229	GLU	-	expression tag	UNP P0DTC2
G	1230	TRP	-	expression tag	UNP P0DTC2
G	1231	VAL	-	expression tag	UNP P0DTC2
G	1232	LEU	-	expression tag	UNP P0DTC2
G	1233	LEU	-	expression tag	UNP P0DTC2
G	1234	SER	-	expression tag	UNP P0DTC2
G	1235	THR	-	expression tag	UNP P0DTC2
G	1236	PHE	-	expression tag	UNP P0DTC2
G	1237	LEU	-	expression tag	UNP P0DTC2
G	1238	GLY	-	expression tag	UNP P0DTC2
G	1239	ARG	-	expression tag	UNP P0DTC2
G	1240	SER	-	expression tag	UNP P0DTC2
G	1241	LEU	-	expression tag	UNP P0DTC2
G	1242	GLU	-	expression tag	UNP P0DTC2
G	1243	VAL	-	expression tag	UNP P0DTC2
G	1244	LEU	-	expression tag	UNP P0DTC2
G	1245	PHE	-	expression tag	UNP P0DTC2
G	1246	GLN	-	expression tag	UNP P0DTC2
G	1247	GLY	-	expression tag	UNP P0DTC2
G	1248	PRO	-	expression tag	UNP P0DTC2
G	1249	GLY	-	expression tag	UNP P0DTC2
G	1250	HIS	-	expression tag	UNP P0DTC2
G	1251	HIS	-	expression tag	UNP P0DTC2
G	1252	HIS	-	expression tag	UNP P0DTC2
G	1253	HIS	-	expression tag	UNP P0DTC2
G	1254	HIS	-	expression tag	UNP P0DTC2
G	1255	HIS	-	expression tag	UNP P0DTC2
G	1256	HIS	-	expression tag	UNP P0DTC2
G	1257	HIS	-	expression tag	UNP P0DTC2
G	1258	SER	-	expression tag	UNP P0DTC2
G	1259	ALA	-	expression tag	UNP P0DTC2
G	1260	TRP	-	expression tag	UNP P0DTC2
G	1261	SER	-	expression tag	UNP P0DTC2
G	1262	HIS	-	expression tag	UNP P0DTC2
G	1263	PRO	-	expression tag	UNP P0DTC2
G	1264	GLN	-	expression tag	UNP P0DTC2
G	1265	PHE	-	expression tag	UNP P0DTC2
G	1266	GLU	-	expression tag	UNP P0DTC2
G	1267	LYS	-	expression tag	UNP P0DTC2
G	1268	GLY	-	expression tag	UNP P0DTC2
G	1269	GLY	-	expression tag	UNP P0DTC2
G	1270	GLY	-	expression tag	UNP P0DTC2

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Chain	Residue	Modelled	Actual	Comment	Reference
G	1271	SER	-	expression tag	UNP P0DTC2
G	1272	GLY	-	expression tag	UNP P0DTC2
G	1273	GLY	-	expression tag	UNP P0DTC2
G	1274	GLY	-	expression tag	UNP P0DTC2
G	1275	GLY	-	expression tag	UNP P0DTC2
G	1276	SER	-	expression tag	UNP P0DTC2
G	1277	GLY	-	expression tag	UNP P0DTC2
G	1278	GLY	-	expression tag	UNP P0DTC2
G	1279	SER	-	expression tag	UNP P0DTC2
G	1280	ALA	-	expression tag	UNP P0DTC2
G	1281	TRP	-	expression tag	UNP P0DTC2
G	1282	SER	-	expression tag	UNP P0DTC2
G	1283	HIS	-	expression tag	UNP P0DTC2
G	1284	PRO	-	expression tag	UNP P0DTC2
G	1285	GLN	-	expression tag	UNP P0DTC2
G	1286	PHE	-	expression tag	UNP P0DTC2
G	1287	GLU	-	expression tag	UNP P0DTC2
G	1288	LYS	-	expression tag	UNP P0DTC2
H	393	PHE	THR	conflict	UNP P0DTC2
H	607	GLU	GLN	conflict	UNP P0DTC2
H	682	ALA	ARG	conflict	UNP P0DTC2
H	683	ALA	ARG	conflict	UNP P0DTC2
H	892	PRO	ALA	conflict	UNP P0DTC2
H	899	PRO	ALA	conflict	UNP P0DTC2
H	942	PRO	ALA	conflict	UNP P0DTC2
H	986	PRO	LYS	conflict	UNP P0DTC2
H	987	PRO	VAL	conflict	UNP P0DTC2
H	1209	GLY	-	expression tag	UNP P0DTC2
H	1210	SER	-	expression tag	UNP P0DTC2
H	1211	GLY	-	expression tag	UNP P0DTC2
H	1212	TYR	-	expression tag	UNP P0DTC2
H	1213	ILE	-	expression tag	UNP P0DTC2
H	1214	PRO	-	expression tag	UNP P0DTC2
H	1215	GLU	-	expression tag	UNP P0DTC2
H	1216	ALA	-	expression tag	UNP P0DTC2
H	1217	PRO	-	expression tag	UNP P0DTC2
H	1218	ARG	-	expression tag	UNP P0DTC2
H	1219	ASP	-	expression tag	UNP P0DTC2
H	1220	GLY	-	expression tag	UNP P0DTC2
H	1221	GLN	-	expression tag	UNP P0DTC2
H	1222	ALA	-	expression tag	UNP P0DTC2
H	1223	TYR	-	expression tag	UNP P0DTC2

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Chain	Residue	Modelled	Actual	Comment	Reference
H	1224	VAL	-	expression tag	UNP P0DTC2
H	1225	ARG	-	expression tag	UNP P0DTC2
H	1226	LYS	-	expression tag	UNP P0DTC2
H	1227	ASP	-	expression tag	UNP P0DTC2
H	1228	GLY	-	expression tag	UNP P0DTC2
H	1229	GLU	-	expression tag	UNP P0DTC2
H	1230	TRP	-	expression tag	UNP P0DTC2
H	1231	VAL	-	expression tag	UNP P0DTC2
H	1232	LEU	-	expression tag	UNP P0DTC2
H	1233	LEU	-	expression tag	UNP P0DTC2
H	1234	SER	-	expression tag	UNP P0DTC2
H	1235	THR	-	expression tag	UNP P0DTC2
H	1236	PHE	-	expression tag	UNP P0DTC2
H	1237	LEU	-	expression tag	UNP P0DTC2
H	1238	GLY	-	expression tag	UNP P0DTC2
H	1239	ARG	-	expression tag	UNP P0DTC2
H	1240	SER	-	expression tag	UNP P0DTC2
H	1241	LEU	-	expression tag	UNP P0DTC2
H	1242	GLU	-	expression tag	UNP P0DTC2
H	1243	VAL	-	expression tag	UNP P0DTC2
H	1244	LEU	-	expression tag	UNP P0DTC2
H	1245	PHE	-	expression tag	UNP P0DTC2
H	1246	GLN	-	expression tag	UNP P0DTC2
H	1247	GLY	-	expression tag	UNP P0DTC2
H	1248	PRO	-	expression tag	UNP P0DTC2
H	1249	GLY	-	expression tag	UNP P0DTC2
H	1250	HIS	-	expression tag	UNP P0DTC2
H	1251	HIS	-	expression tag	UNP P0DTC2
H	1252	HIS	-	expression tag	UNP P0DTC2
H	1253	HIS	-	expression tag	UNP P0DTC2
H	1254	HIS	-	expression tag	UNP P0DTC2
H	1255	HIS	-	expression tag	UNP P0DTC2
H	1256	HIS	-	expression tag	UNP P0DTC2
H	1257	HIS	-	expression tag	UNP P0DTC2
H	1258	SER	-	expression tag	UNP P0DTC2
H	1259	ALA	-	expression tag	UNP P0DTC2
H	1260	TRP	-	expression tag	UNP P0DTC2
H	1261	SER	-	expression tag	UNP P0DTC2
H	1262	HIS	-	expression tag	UNP P0DTC2
H	1263	PRO	-	expression tag	UNP P0DTC2
H	1264	GLN	-	expression tag	UNP P0DTC2
H	1265	PHE	-	expression tag	UNP P0DTC2

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Chain	Residue	Modelled	Actual	Comment	Reference
H	1266	GLU	-	expression tag	UNP P0DTC2
H	1267	LYS	-	expression tag	UNP P0DTC2
H	1268	GLY	-	expression tag	UNP P0DTC2
H	1269	GLY	-	expression tag	UNP P0DTC2
H	1270	GLY	-	expression tag	UNP P0DTC2
H	1271	SER	-	expression tag	UNP P0DTC2
H	1272	GLY	-	expression tag	UNP P0DTC2
H	1273	GLY	-	expression tag	UNP P0DTC2
H	1274	GLY	-	expression tag	UNP P0DTC2
H	1275	GLY	-	expression tag	UNP P0DTC2
H	1276	SER	-	expression tag	UNP P0DTC2
H	1277	GLY	-	expression tag	UNP P0DTC2
H	1278	GLY	-	expression tag	UNP P0DTC2
H	1279	SER	-	expression tag	UNP P0DTC2
H	1280	ALA	-	expression tag	UNP P0DTC2
H	1281	TRP	-	expression tag	UNP P0DTC2
H	1282	SER	-	expression tag	UNP P0DTC2
H	1283	HIS	-	expression tag	UNP P0DTC2
H	1284	PRO	-	expression tag	UNP P0DTC2
H	1285	GLN	-	expression tag	UNP P0DTC2
H	1286	PHE	-	expression tag	UNP P0DTC2
H	1287	GLU	-	expression tag	UNP P0DTC2
H	1288	LYS	-	expression tag	UNP P0DTC2
J	393	PHE	THR	conflict	UNP P0DTC2
J	607	GLU	GLN	conflict	UNP P0DTC2
J	682	ALA	ARG	conflict	UNP P0DTC2
J	683	ALA	ARG	conflict	UNP P0DTC2
J	892	PRO	ALA	conflict	UNP P0DTC2
J	899	PRO	ALA	conflict	UNP P0DTC2
J	942	PRO	ALA	conflict	UNP P0DTC2
J	986	PRO	LYS	conflict	UNP P0DTC2
J	987	PRO	VAL	conflict	UNP P0DTC2
J	1209	GLY	-	expression tag	UNP P0DTC2
J	1210	SER	-	expression tag	UNP P0DTC2
J	1211	GLY	-	expression tag	UNP P0DTC2
J	1212	TYR	-	expression tag	UNP P0DTC2
J	1213	ILE	-	expression tag	UNP P0DTC2
J	1214	PRO	-	expression tag	UNP P0DTC2
J	1215	GLU	-	expression tag	UNP P0DTC2
J	1216	ALA	-	expression tag	UNP P0DTC2
J	1217	PRO	-	expression tag	UNP P0DTC2
J	1218	ARG	-	expression tag	UNP P0DTC2

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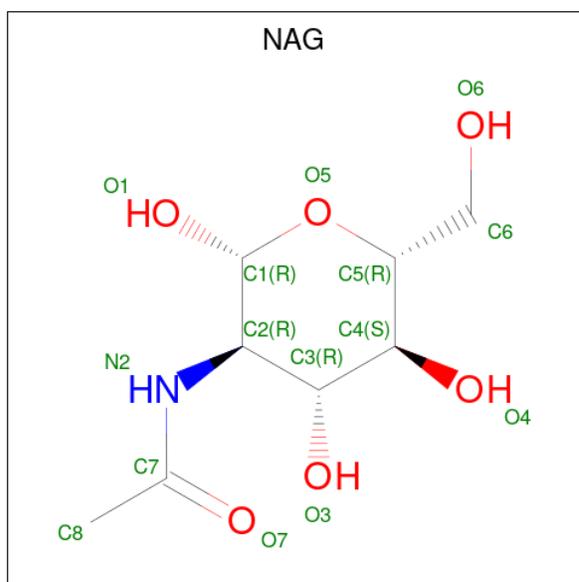
Chain	Residue	Modelled	Actual	Comment	Reference
J	1219	ASP	-	expression tag	UNP P0DTC2
J	1220	GLY	-	expression tag	UNP P0DTC2
J	1221	GLN	-	expression tag	UNP P0DTC2
J	1222	ALA	-	expression tag	UNP P0DTC2
J	1223	TYR	-	expression tag	UNP P0DTC2
J	1224	VAL	-	expression tag	UNP P0DTC2
J	1225	ARG	-	expression tag	UNP P0DTC2
J	1226	LYS	-	expression tag	UNP P0DTC2
J	1227	ASP	-	expression tag	UNP P0DTC2
J	1228	GLY	-	expression tag	UNP P0DTC2
J	1229	GLU	-	expression tag	UNP P0DTC2
J	1230	TRP	-	expression tag	UNP P0DTC2
J	1231	VAL	-	expression tag	UNP P0DTC2
J	1232	LEU	-	expression tag	UNP P0DTC2
J	1233	LEU	-	expression tag	UNP P0DTC2
J	1234	SER	-	expression tag	UNP P0DTC2
J	1235	THR	-	expression tag	UNP P0DTC2
J	1236	PHE	-	expression tag	UNP P0DTC2
J	1237	LEU	-	expression tag	UNP P0DTC2
J	1238	GLY	-	expression tag	UNP P0DTC2
J	1239	ARG	-	expression tag	UNP P0DTC2
J	1240	SER	-	expression tag	UNP P0DTC2
J	1241	LEU	-	expression tag	UNP P0DTC2
J	1242	GLU	-	expression tag	UNP P0DTC2
J	1243	VAL	-	expression tag	UNP P0DTC2
J	1244	LEU	-	expression tag	UNP P0DTC2
J	1245	PHE	-	expression tag	UNP P0DTC2
J	1246	GLN	-	expression tag	UNP P0DTC2
J	1247	GLY	-	expression tag	UNP P0DTC2
J	1248	PRO	-	expression tag	UNP P0DTC2
J	1249	GLY	-	expression tag	UNP P0DTC2
J	1250	HIS	-	expression tag	UNP P0DTC2
J	1251	HIS	-	expression tag	UNP P0DTC2
J	1252	HIS	-	expression tag	UNP P0DTC2
J	1253	HIS	-	expression tag	UNP P0DTC2
J	1254	HIS	-	expression tag	UNP P0DTC2
J	1255	HIS	-	expression tag	UNP P0DTC2
J	1256	HIS	-	expression tag	UNP P0DTC2
J	1257	HIS	-	expression tag	UNP P0DTC2
J	1258	SER	-	expression tag	UNP P0DTC2
J	1259	ALA	-	expression tag	UNP P0DTC2
J	1260	TRP	-	expression tag	UNP P0DTC2

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Chain	Residue	Modelled	Actual	Comment	Reference
J	1261	SER	-	expression tag	UNP P0DTC2
J	1262	HIS	-	expression tag	UNP P0DTC2
J	1263	PRO	-	expression tag	UNP P0DTC2
J	1264	GLN	-	expression tag	UNP P0DTC2
J	1265	PHE	-	expression tag	UNP P0DTC2
J	1266	GLU	-	expression tag	UNP P0DTC2
J	1267	LYS	-	expression tag	UNP P0DTC2
J	1268	GLY	-	expression tag	UNP P0DTC2
J	1269	GLY	-	expression tag	UNP P0DTC2
J	1270	GLY	-	expression tag	UNP P0DTC2
J	1271	SER	-	expression tag	UNP P0DTC2
J	1272	GLY	-	expression tag	UNP P0DTC2
J	1273	GLY	-	expression tag	UNP P0DTC2
J	1274	GLY	-	expression tag	UNP P0DTC2
J	1275	GLY	-	expression tag	UNP P0DTC2
J	1276	SER	-	expression tag	UNP P0DTC2
J	1277	GLY	-	expression tag	UNP P0DTC2
J	1278	GLY	-	expression tag	UNP P0DTC2
J	1279	SER	-	expression tag	UNP P0DTC2
J	1280	ALA	-	expression tag	UNP P0DTC2
J	1281	TRP	-	expression tag	UNP P0DTC2
J	1282	SER	-	expression tag	UNP P0DTC2
J	1283	HIS	-	expression tag	UNP P0DTC2
J	1284	PRO	-	expression tag	UNP P0DTC2
J	1285	GLN	-	expression tag	UNP P0DTC2
J	1286	PHE	-	expression tag	UNP P0DTC2
J	1287	GLU	-	expression tag	UNP P0DTC2
J	1288	LYS	-	expression tag	UNP P0DTC2

- Molecule 3 is 2-acetamido-2-deoxy-beta-D-glucofuranose (three-letter code: NAG) (formula: C<sub>8</sub>H<sub>15</sub>NO<sub>6</sub>).



Mol	Chain	Residues	Atoms				AltConf
			Total	C	N	O	
3	A	1	14	8	1	5	0
3	A	1	14	8	1	5	0
3	A	1	14	8	1	5	0
3	A	1	14	8	1	5	0
3	A	1	14	8	1	5	0
3	A	1	14	8	1	5	0
3	A	1	14	8	1	5	0
3	A	1	14	8	1	5	0
3	B	1	14	8	1	5	0
3	B	1	14	8	1	5	0
3	B	1	14	8	1	5	0
3	B	1	14	8	1	5	0
3	B	1	14	8	1	5	0
3	B	1	14	8	1	5	0
3	B	1	14	8	1	5	0

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Mol	Chain	Residues	Atoms				AltConf
			Total	C	N	O	
3	D	1	Total 14	C 8	N 1	O 5	0
3	D	1	Total 14	C 8	N 1	O 5	0
3	D	1	Total 14	C 8	N 1	O 5	0
3	D	1	Total 14	C 8	N 1	O 5	0
3	D	1	Total 14	C 8	N 1	O 5	0
3	D	1	Total 14	C 8	N 1	O 5	0
3	D	1	Total 14	C 8	N 1	O 5	0
3	D	1	Total 14	C 8	N 1	O 5	0
3	G	1	Total 14	C 8	N 1	O 5	0
3	G	1	Total 14	C 8	N 1	O 5	0
3	G	1	Total 14	C 8	N 1	O 5	0
3	G	1	Total 14	C 8	N 1	O 5	0
3	G	1	Total 14	C 8	N 1	O 5	0
3	G	1	Total 14	C 8	N 1	O 5	0
3	G	1	Total 14	C 8	N 1	O 5	0
3	G	1	Total 14	C 8	N 1	O 5	0
3	H	1	Total 14	C 8	N 1	O 5	0
3	H	1	Total 14	C 8	N 1	O 5	0
3	H	1	Total 14	C 8	N 1	O 5	0
3	H	1	Total 14	C 8	N 1	O 5	0
3	H	1	Total 14	C 8	N 1	O 5	0
3	H	1	Total 14	C 8	N 1	O 5	0
3	H	1	Total 14	C 8	N 1	O 5	0

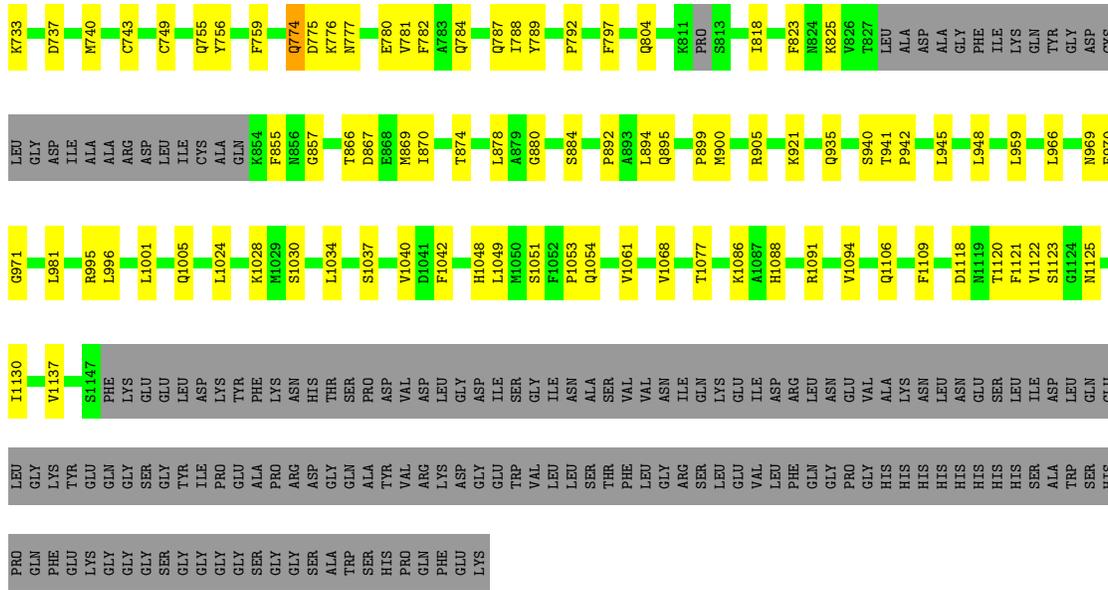
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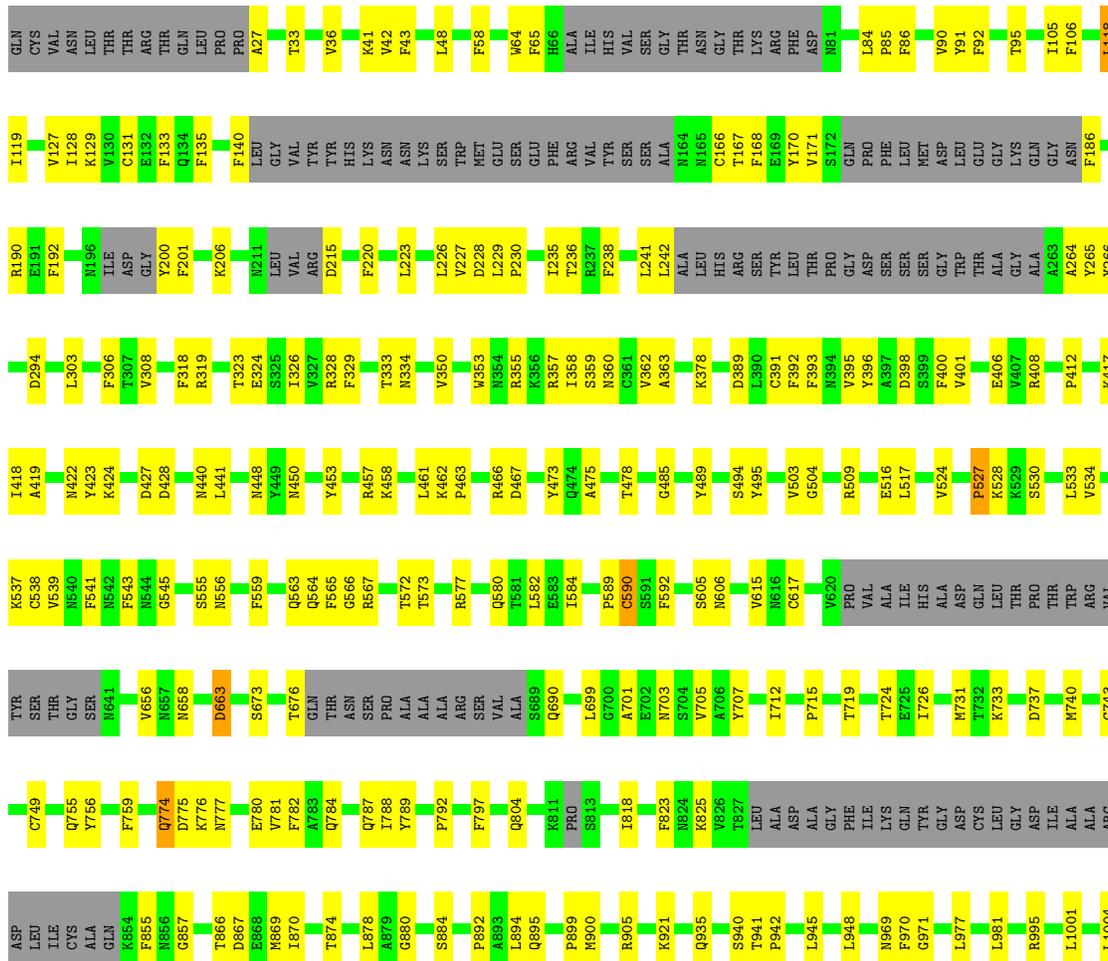
<b>Mol</b>	<b>Chain</b>	<b>Residues</b>	<b>Atoms</b>				<b>AltConf</b>
3	J	1	Total	C	N	O	0
			14	8	1	5	
3	J	1	Total	C	N	O	0
			14	8	1	5	
3	J	1	Total	C	N	O	0
			14	8	1	5	
3	J	1	Total	C	N	O	0
			14	8	1	5	
3	J	1	Total	C	N	O	0
			14	8	1	5	
3	J	1	Total	C	N	O	0
			14	8	1	5	







● Molecule 2: Spike glycoprotein











## 4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, Not provided	
Number of particles used	43791	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 ( $e^-/\text{\AA}^2$ )	50	Depositor
Minimum defocus (nm)	750	Depositor
Maximum defocus (nm)	1500	Depositor
Magnification	Not provided	
Image detector	FEI FALCON IV (4k x 4k)	Depositor

## 5 Model quality [i](#)

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

Bond lengths and bond angles in the following residue types are not validated in this section: NAG

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	C	0.27	0/956	0.51	0/1293
1	E	0.27	0/956	0.51	0/1293
1	F	0.26	0/956	0.51	0/1293
1	I	0.27	0/956	0.51	0/1293
1	K	0.27	0/956	0.51	0/1293
1	L	0.26	0/956	0.51	0/1293
2	A	0.38	0/7765	0.52	5/10575 (0.0%)
2	B	0.38	0/7765	0.52	5/10575 (0.0%)
2	D	0.38	0/7765	0.52	5/10575 (0.0%)
2	G	0.38	0/7765	0.52	5/10575 (0.0%)
2	H	0.38	0/7765	0.52	5/10575 (0.0%)
2	J	0.38	0/7765	0.52	5/10575 (0.0%)
All	All	0.37	0/52326	0.52	30/71208 (0.0%)

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
2	A	0	1
2	B	0	1
2	D	0	1
2	G	0	1
2	H	0	1
2	J	0	1
All	All	0	6

There are no bond length outliers.

The worst 5 of 30 bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	D	527	PRO	O-C-N	-9.06	108.20	122.70
2	J	527	PRO	O-C-N	-9.06	108.20	122.70
2	G	527	PRO	O-C-N	-9.06	108.21	122.70
2	H	527	PRO	O-C-N	-9.06	108.21	122.70
2	A	527	PRO	O-C-N	-9.05	108.22	122.70

There are no chirality outliers.

5 of 6 planarity outliers are listed below:

Mol	Chain	Res	Type	Group
2	A	527	PRO	Mainchain
2	B	527	PRO	Mainchain
2	D	527	PRO	Mainchain
2	G	527	PRO	Mainchain
2	H	527	PRO	Mainchain

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

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	C	933	0	865	52	0
1	E	933	0	865	51	0
1	F	933	0	865	53	0
1	I	933	0	865	51	0
1	K	933	0	865	52	0
1	L	933	0	865	54	0
2	A	7594	0	7294	390	0
2	B	7594	0	7294	385	0
2	D	7594	0	7294	391	0
2	G	7594	0	7294	387	0
2	H	7594	0	7294	397	0
2	J	7594	0	7294	389	0
3	A	98	0	91	1	0
3	B	98	0	91	1	0
3	D	98	0	91	1	0
3	G	98	0	91	1	0
3	H	98	0	91	1	0
3	J	98	0	91	1	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
All	All	51750	0	49500	2000	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 20.

The worst 5 of 2000 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)
2:J:565:PHE:CE2	2:J:567:ARG:HG3	1.01	1.54
2:B:565:PHE:CE2	2:B:567:ARG:HG3	1.01	1.54
2:A:565:PHE:CE2	2:A:567:ARG:HG3	1.01	1.52
2:G:565:PHE:HE2	2:G:567:ARG:CG	1.22	1.52
2:G:565:PHE:CE2	2:G:567:ARG:HG3	1.01	1.51

There are no symmetry-related clashes.

## 5.3 Torsion angles [i](#)

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

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	C	117/148 (79%)	108 (92%)	9 (8%)	0	100	100
1	E	117/148 (79%)	108 (92%)	9 (8%)	0	100	100
1	F	117/148 (79%)	108 (92%)	9 (8%)	0	100	100
1	I	117/148 (79%)	108 (92%)	9 (8%)	0	100	100
1	K	117/148 (79%)	108 (92%)	9 (8%)	0	100	100
1	L	117/148 (79%)	108 (92%)	9 (8%)	0	100	100
2	A	964/1275 (76%)	909 (94%)	55 (6%)	0	100	100
2	B	964/1275 (76%)	909 (94%)	55 (6%)	0	100	100
2	D	964/1275 (76%)	909 (94%)	55 (6%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
2	G	964/1275 (76%)	909 (94%)	55 (6%)	0	100	100
2	H	964/1275 (76%)	909 (94%)	55 (6%)	0	100	100
2	J	964/1275 (76%)	909 (94%)	55 (6%)	0	100	100
All	All	6486/8538 (76%)	6102 (94%)	384 (6%)	0	100	100

There are no Ramachandran outliers to report.

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

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	C	96/119 (81%)	95 (99%)	1 (1%)	73	84
1	E	96/119 (81%)	95 (99%)	1 (1%)	73	84
1	F	96/119 (81%)	95 (99%)	1 (1%)	73	84
1	I	96/119 (81%)	95 (99%)	1 (1%)	73	84
1	K	96/119 (81%)	95 (99%)	1 (1%)	73	84
1	L	96/119 (81%)	95 (99%)	1 (1%)	73	84
2	A	831/1102 (75%)	826 (99%)	5 (1%)	84	90
2	B	831/1102 (75%)	826 (99%)	5 (1%)	84	90
2	D	831/1102 (75%)	826 (99%)	5 (1%)	84	90
2	G	831/1102 (75%)	826 (99%)	5 (1%)	84	90
2	H	831/1102 (75%)	826 (99%)	5 (1%)	84	90
2	J	831/1102 (75%)	826 (99%)	5 (1%)	84	90
All	All	5562/7326 (76%)	5526 (99%)	36 (1%)	82	90

5 of 36 residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
2	H	774	GLN
2	J	1123	SER

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Mol	Chain	Res	Type
2	H	1123	SER
2	J	590	CYS
1	F	95	TYR

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. 5 of 32 such sidechains are listed below:

Mol	Chain	Res	Type
2	J	655	HIS
2	J	1048	HIS
2	D	360	ASN
2	B	1106	GLN
2	J	1071	GLN

### 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](#)

42 ligands are modelled in this entry.

In the following table, the Counts columns list the number of bonds (or angles) for which Mogul statistics could be retrieved, the number of bonds (or angles) that are observed in the model and the number of bonds (or angles) that are defined in the Chemical Component Dictionary. The Link column lists molecule types, if any, to which the group is linked. The Z score for a bond length (or angle) is the number of standard deviations the observed value is removed from the expected value. A bond length (or angle) with  $|Z| > 2$  is considered an outlier worth inspection. RMSZ is the root-mean-square of all Z scores of the bond lengths (or angles).

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z  > 2$	Counts	RMSZ	$\# Z  > 2$
3	NAG	D	1301	2	14,14,15	0.69	0	17,19,21	1.12	1 (5%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z  > 2	Counts	RMSZ	# Z  > 2
3	NAG	G	1302	2	14,14,15	0.70	0	17,19,21	1.01	1 (5%)
3	NAG	G	1303	2	14,14,15	0.75	0	17,19,21	0.83	0
3	NAG	J	1303	2	14,14,15	0.75	0	17,19,21	0.82	0
3	NAG	B	1305	2	14,14,15	0.70	0	17,19,21	1.24	2 (11%)
3	NAG	J	1302	2	14,14,15	0.68	0	17,19,21	1.01	1 (5%)
3	NAG	D	1303	2	14,14,15	0.75	0	17,19,21	0.82	0
3	NAG	B	1306	2	14,14,15	0.77	0	17,19,21	0.87	0
3	NAG	J	1307	2	14,14,15	0.81	0	17,19,21	0.88	0
3	NAG	D	1307	2	14,14,15	0.81	0	17,19,21	0.88	0
3	NAG	H	1307	2	14,14,15	0.80	0	17,19,21	0.89	0
3	NAG	B	1304	2	14,14,15	0.71	0	17,19,21	0.91	0
3	NAG	A	1307	2	14,14,15	0.80	0	17,19,21	0.88	0
3	NAG	D	1304	2	14,14,15	0.71	0	17,19,21	0.91	0
3	NAG	A	1302	2	14,14,15	0.70	0	17,19,21	1.01	1 (5%)
3	NAG	A	1304	2	14,14,15	0.72	0	17,19,21	0.92	0
3	NAG	H	1302	2	14,14,15	0.70	0	17,19,21	1.01	1 (5%)
3	NAG	G	1304	2	14,14,15	0.72	0	17,19,21	0.92	0
3	NAG	D	1306	2	14,14,15	0.77	0	17,19,21	0.87	0
3	NAG	J	1304	2	14,14,15	0.71	0	17,19,21	0.91	0
3	NAG	G	1306	2	14,14,15	0.77	0	17,19,21	0.87	0
3	NAG	H	1305	2	14,14,15	0.70	0	17,19,21	1.25	2 (11%)
3	NAG	H	1301	2	14,14,15	0.69	0	17,19,21	1.12	1 (5%)
3	NAG	J	1301	2	14,14,15	0.69	0	17,19,21	1.12	1 (5%)
3	NAG	B	1307	2	14,14,15	0.81	0	17,19,21	0.89	0
3	NAG	B	1302	2	14,14,15	0.70	0	17,19,21	1.01	1 (5%)
3	NAG	A	1305	2	14,14,15	0.71	0	17,19,21	1.24	2 (11%)
3	NAG	A	1306	2	14,14,15	0.77	0	17,19,21	0.87	0
3	NAG	G	1307	2	14,14,15	0.81	0	17,19,21	0.88	0
3	NAG	A	1301	2	14,14,15	0.69	0	17,19,21	1.13	1 (5%)
3	NAG	H	1304	2	14,14,15	0.72	0	17,19,21	0.92	0
3	NAG	D	1302	2	14,14,15	0.68	0	17,19,21	1.01	1 (5%)
3	NAG	A	1303	2	14,14,15	0.75	0	17,19,21	0.83	0
3	NAG	D	1305	2	14,14,15	0.70	0	17,19,21	1.25	2 (11%)
3	NAG	H	1306	2	14,14,15	0.77	0	17,19,21	0.87	0
3	NAG	G	1305	2	14,14,15	0.70	0	17,19,21	1.24	2 (11%)
3	NAG	J	1305	2	14,14,15	0.70	0	17,19,21	1.25	2 (11%)
3	NAG	J	1306	2	14,14,15	0.76	0	17,19,21	0.87	0
3	NAG	B	1303	2	14,14,15	0.75	0	17,19,21	0.82	0
3	NAG	G	1301	2	14,14,15	0.69	0	17,19,21	1.12	1 (5%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z  > 2	Counts	RMSZ	# Z  > 2
3	NAG	B	1301	2	14,14,15	0.69	0	17,19,21	1.12	1 (5%)
3	NAG	H	1303	2	14,14,15	0.74	0	17,19,21	0.82	0

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

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
3	NAG	D	1301	2	-	3/6/23/26	0/1/1/1
3	NAG	G	1302	2	-	0/6/23/26	0/1/1/1
3	NAG	G	1303	2	-	1/6/23/26	0/1/1/1
3	NAG	J	1303	2	-	1/6/23/26	0/1/1/1
3	NAG	B	1305	2	-	1/6/23/26	0/1/1/1
3	NAG	J	1302	2	-	0/6/23/26	0/1/1/1
3	NAG	D	1303	2	-	1/6/23/26	0/1/1/1
3	NAG	B	1306	2	-	2/6/23/26	0/1/1/1
3	NAG	J	1307	2	-	2/6/23/26	0/1/1/1
3	NAG	D	1307	2	-	2/6/23/26	0/1/1/1
3	NAG	H	1307	2	-	2/6/23/26	0/1/1/1
3	NAG	B	1304	2	-	2/6/23/26	0/1/1/1
3	NAG	A	1307	2	-	2/6/23/26	0/1/1/1
3	NAG	D	1304	2	-	2/6/23/26	0/1/1/1
3	NAG	A	1302	2	-	0/6/23/26	0/1/1/1
3	NAG	A	1304	2	-	2/6/23/26	0/1/1/1
3	NAG	H	1302	2	-	0/6/23/26	0/1/1/1
3	NAG	G	1304	2	-	2/6/23/26	0/1/1/1
3	NAG	D	1306	2	-	2/6/23/26	0/1/1/1
3	NAG	J	1304	2	-	2/6/23/26	0/1/1/1
3	NAG	G	1306	2	-	2/6/23/26	0/1/1/1
3	NAG	H	1305	2	-	1/6/23/26	0/1/1/1
3	NAG	H	1301	2	-	3/6/23/26	0/1/1/1
3	NAG	J	1301	2	-	3/6/23/26	0/1/1/1
3	NAG	B	1307	2	-	2/6/23/26	0/1/1/1
3	NAG	B	1302	2	-	0/6/23/26	0/1/1/1
3	NAG	A	1305	2	-	1/6/23/26	0/1/1/1
3	NAG	A	1306	2	-	2/6/23/26	0/1/1/1
3	NAG	G	1307	2	-	2/6/23/26	0/1/1/1

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
3	NAG	A	1301	2	-	3/6/23/26	0/1/1/1
3	NAG	H	1304	2	-	2/6/23/26	0/1/1/1
3	NAG	D	1302	2	-	0/6/23/26	0/1/1/1
3	NAG	A	1303	2	-	1/6/23/26	0/1/1/1
3	NAG	D	1305	2	-	1/6/23/26	0/1/1/1
3	NAG	H	1306	2	-	2/6/23/26	0/1/1/1
3	NAG	G	1305	2	-	1/6/23/26	0/1/1/1
3	NAG	J	1305	2	-	1/6/23/26	0/1/1/1
3	NAG	J	1306	2	-	2/6/23/26	0/1/1/1
3	NAG	B	1303	2	-	1/6/23/26	0/1/1/1
3	NAG	G	1301	2	-	3/6/23/26	0/1/1/1
3	NAG	B	1301	2	-	3/6/23/26	0/1/1/1
3	NAG	H	1303	2	-	1/6/23/26	0/1/1/1

There are no bond length outliers.

The worst 5 of 24 bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	A	1301	NAG	C2-N2-C7	3.01	127.19	122.90
3	G	1301	NAG	C2-N2-C7	3.01	127.19	122.90
3	D	1301	NAG	C2-N2-C7	3.00	127.17	122.90
3	J	1301	NAG	C2-N2-C7	3.00	127.17	122.90
3	B	1301	NAG	C2-N2-C7	2.99	127.15	122.90

There are no chirality outliers.

5 of 66 torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
3	A	1306	NAG	O5-C5-C6-O6
3	B	1306	NAG	O5-C5-C6-O6
3	D	1306	NAG	O5-C5-C6-O6
3	G	1306	NAG	O5-C5-C6-O6
3	H	1306	NAG	O5-C5-C6-O6

There are no ring outliers.

6 monomers are involved in 6 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
3	G	1302	NAG	1	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
3	J	1302	NAG	1	0
3	A	1302	NAG	1	0
3	H	1302	NAG	1	0
3	B	1302	NAG	1	0
3	D	1302	NAG	1	0

## 5.7 Other polymers [i](#)

There are no such residues in this entry.

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

There are no chain breaks in this entry.