


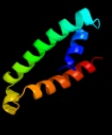



















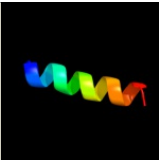


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d1l7va_</a>	 Alignment		92.7	16	<b>Fold:</b> ABC transporter involved in vitamin B12 uptake, BtuC <b>Superfamily:</b> ABC transporter involved in vitamin B12 uptake, BtuC <b>Family:</b> ABC transporter involved in vitamin B12 uptake, BtuC
2	<a href="#">c2nq2A_</a>	 Alignment		88.7	19	<b>PDB header:</b> metal transport <b>Chain:</b> A: <b>PDB Molecule:</b> hypothetical abc transporter permease protein <b>PDBTitle:</b> an inward-facing conformation of a putative metal-chelate2 type abc transporter.
3	<a href="#">c2ap8A_</a>	 Alignment		19.3	46	<b>PDB header:</b> antibiotic <b>Chain:</b> A: <b>PDB Molecule:</b> bombinin h4; <b>PDBTitle:</b> solution structure of bombinin h4 in dpc micelles
4	<a href="#">c2ap7A_</a>	 Alignment		19.1	46	<b>PDB header:</b> antibiotic <b>Chain:</b> A: <b>PDB Molecule:</b> bombinin h2; <b>PDBTitle:</b> solution structure of bombinin h2 in dpc micelles
5	<a href="#">c1xooA_</a>	 Alignment		12.0	45	<b>PDB header:</b> viral protein <b>Chain:</b> A: <b>PDB Molecule:</b> hemagglutinin; <b>PDBTitle:</b> nmr structure of g1s mutant of influenza hemagglutinin2 fusion peptide in dpc micelles at ph 5
6	<a href="#">c2l4gA_</a>	 Alignment		11.6	42	<b>PDB header:</b> viral protein <b>Chain:</b> A: <b>PDB Molecule:</b> haemagglutinin; <b>PDBTitle:</b> influenza haemagglutinin fusion peptide mutant g13a
7	<a href="#">c2kxaA_</a>	 Alignment		11.2	56	<b>PDB header:</b> viral protein, immune system <b>Chain:</b> A: <b>PDB Molecule:</b> haemagglutinin ha2 chain peptide; <b>PDBTitle:</b> the hemagglutinin fusion peptide (h1 subtype) at ph 7.4
8	<a href="#">c1iboA_</a>	 Alignment		11.0	45	<b>PDB header:</b> viral protein <b>Chain:</b> A: <b>PDB Molecule:</b> hemagglutinin ha2 chain peptide; <b>PDBTitle:</b> nmr structure of hemagglutinin fusion peptide in dpc2 micelles at ph 7.4
9	<a href="#">c1ibnA_</a>	 Alignment		11.0	45	<b>PDB header:</b> viral protein <b>Chain:</b> A: <b>PDB Molecule:</b> hemagglutinin ha2 chain peptide; <b>PDBTitle:</b> nmr structure of hemagglutinin fusion peptide in dpc2 micelles at ph 5
10	<a href="#">c1xopA_</a>	 Alignment		10.9	45	<b>PDB header:</b> viral protein <b>Chain:</b> A: <b>PDB Molecule:</b> hemagglutinin; <b>PDBTitle:</b> nmr structure of g1v mutant of influenza hemagglutinin2 fusion peptide in dpc micelles at ph 5
11	<a href="#">c2jrdA_</a>	 Alignment		7.0	45	<b>PDB header:</b> viral protein <b>Chain:</b> A: <b>PDB Molecule:</b> hemagglutinin; <b>PDBTitle:</b> influenza hemagglutinin fusion domain mutant f9a

12	<a href="#">c3cliA_</a>	Alignment		6.5	23	<b>PDB header:</b> transport protein <b>Chain:</b> A: <b>PDB Molecule:</b> ammonia channel; <b>PDBTitle:</b> substrate binding, deprotonation and selectivity at the2 periplasmic entrance of the e. coli ammonia channel amtb
13	<a href="#">c2kpeA_</a>	Alignment		6.3	33	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> glycophorin-a; <b>PDBTitle:</b> refined structure of glycophorin a transmembrane segment dimer in dpc2 micelles

14 [c2kpeB](#)

Alignment



6.3

33

**PDB header:**membrane protein  
**Chain:** B: **PDB Molecule:**glycophorin-a;  
**PDBTitle:** refined structure of glycophorin a transmembrane segment dimer in dpc2 micelles