Phyre^2

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Description	P0AAW9
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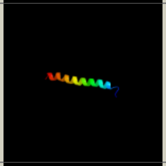
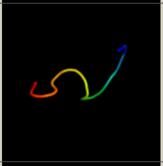
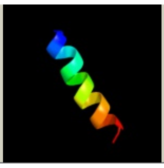
Unique Job ID	b9c036eb9d2f65e9

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2ajmA_	Alignment		23.7	46	**PDB header:**membrane protein **Chain:** A: **PDB Molecule:**nonstructural protein 5a; **PDBTitle:** nmr structure of the in-plane membrane anchor domain [1-28]2 of the monotopic nonstructural protein 5a (ns5a) from the3 bovine viral diarrhea virus (bvdv)
2	c3dinD_	Alignment		19.6	21	**PDB header:**membrane protein, protein transport **Chain:** D: **PDB Molecule:**preprotein translocase subunit sece; **PDBTitle:** crystal structure of the protein-translocation complex formed by the2 secy channel and the seca atpase
3	d1jb0i_	Alignment		7.9	33	**Fold:**Single transmembrane helix **Superfamily:**Subunit VIII of photosystem I reaction centre, PsaI **Family:**Subunit VIII of photosystem I reaction centre, PsaI
4	c3g79A_	Alignment		7.3	25	**PDB header:**oxidoreductase **Chain:** A: **PDB Molecule:**ndp-n-acetyl-d-galactosaminuronic acid dehydrogenase; **PDBTitle:** crystal structure of ndp-n-acetyl-d-galactosaminuronic acid2 dehydrogenase from methanosarcina mazei go1
5	c1afoB_	Alignment		6.4	27	**PDB header:**integral membrane protein **Chain:** B: **PDB Molecule:**glycophorin a; **PDBTitle:** dimeric transmembrane domain of human glycophorin a, nmr,2 20 structures

6	c2kv5A_	Alignment		6.0	50	**PDB header:**toxin **Chain:** A: **PDB Molecule:**putative uncharacterized protein rnai; **PDBTitle:** solution structure of the par toxin fst in dpc micelles