

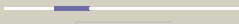








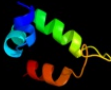










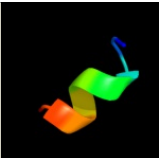
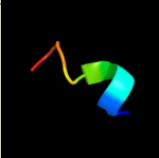





#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2i68B_	 Alignment		24.2	19	PDB header: transport protein Chain: B: PDB Molecule: protein emre; PDBTitle: cryo-em based theoretical model structure of transmembrane2 domain of the multidrug-resistance antiporter from e. coli3 emre
2	c1xooA_	 Alignment		11.9	70	PDB header: viral protein Chain: A: PDB Molecule: hemagglutinin; PDBTitle: nmr structure of g1s mutant of influenza hemagglutinin2 fusion peptide in dpc micelles at ph 5
3	c2y69W_	 Alignment		10.5	19	PDB header: electron transport Chain: W: PDB Molecule: cytochrome c oxidase polypeptide 7a1; PDBTitle: bovine heart cytochrome c oxidase re-refined with molecular2 oxygen
4	d1v54j_	 Alignment		10.3	19	Fold: Single transmembrane helix Superfamily: Mitochondrial cytochrome c oxidase subunit VIIa Family: Mitochondrial cytochrome c oxidase subunit VIIa
5	d1htjF_	 Alignment		10.2	22	Fold: Regulator of G-protein signaling, RGS Superfamily: Regulator of G-protein signaling, RGS Family: Regulator of G-protein signaling, RGS
6	c1htjF_	 Alignment		10.2	22	PDB header: signaling protein Chain: F: PDB Molecule: kiaa0380; PDBTitle: structure of the rgs-like domain from pdz-rhogef
7	c2kxaA_	 Alignment		7.5	71	PDB header: viral protein, immune system Chain: A: PDB Molecule: haemagglutinin ha2 chain peptide; PDBTitle: the hemagglutinin fusion peptide (h1 subtype) at ph 7.4
8	c1xopA_	 Alignment		6.7	71	PDB header: viral protein Chain: A: PDB Molecule: hemagglutinin; PDBTitle: nmr structure of g1v mutant of influenza hemagglutinin2 fusion peptide in dpc micelles at ph 5
9	c1iboA_	 Alignment		6.7	71	PDB header: viral protein Chain: A: PDB Molecule: hemagglutinin ha2 chain peptide; PDBTitle: nmr structure of hemagglutinin fusion peptide in dpc2 micelles at ph 7.4
10	c1ibnA_	 Alignment		6.7	71	PDB header: viral protein Chain: A: PDB Molecule: hemagglutinin ha2 chain peptide; PDBTitle: nmr structure of hemagglutinin fusion peptide in dpc2 micelles at ph 5
11	c2kn8A_	 Alignment		6.7	60	PDB header: protein binding, dna binding protein Chain: A: PDB Molecule: dna cleavage and packaging protein large subunit, ul89; PDBTitle: nmr structure of the c-terminal domain of pul89

12	c2l4gA_	Alignment		6.5	67	PDB header: viral protein Chain: A: PDB Molecule: haemagglutinin; PDBTitle: influenza haemagglutinin fusion peptide mutant g13a
13	c2dciA_	Alignment		6.2	67	PDB header: viral protein Chain: A: PDB Molecule: hemagglutinin; PDBTitle: nmr structure of influenza ha fusion peptide mutant w14a in2 dpc in ph5
14	c3a0hx_	Alignment		5.1	42	PDB header: electron transport Chain: X: PDB Molecule: photosystem ii reaction center protein x; PDBTitle: crystal structure of i-substituted photosystem ii complex
15	c3a0bX_	Alignment		5.1	42	PDB header: electron transport Chain: X: PDB Molecule: photosystem ii reaction center protein x; PDBTitle: crystal structure of br-substituted photosystem ii complex

16	c3a0bx_	Alignment		5.1	42	PDB header: electron transport Chain: X: PDB Molecule: photosystem ii reaction center protein x; PDBTitle: crystal structure of br-substituted photosystem ii complex
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