
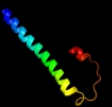

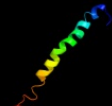

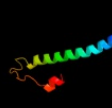





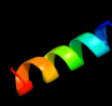





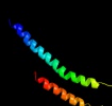







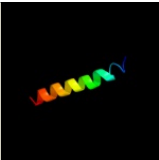


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d3ehbb2</a>	 Alignment		16.9	7	<b>Fold:</b> Transmembrane helix hairpin <b>Superfamily:</b> Cytochrome c oxidase subunit II-like, transmembrane region <b>Family:</b> Cytochrome c oxidase subunit II-like, transmembrane region
2	<a href="#">d1fftb2</a>	 Alignment		14.9	14	<b>Fold:</b> Transmembrane helix hairpin <b>Superfamily:</b> Cytochrome c oxidase subunit II-like, transmembrane region <b>Family:</b> Cytochrome c oxidase subunit II-like, transmembrane region
3	<a href="#">c1ar1B_</a>	 Alignment		14.3	7	<b>PDB header:</b> complex (oxidoreductase/antibody) <b>Chain:</b> B: <b>PDB Molecule:</b> cytochrome c oxidase; <b>PDBTitle:</b> structure at 2.7 angstrom resolution of the paracoccus2 denitrificans two-subunit cytochrome c oxidase complexed3 with an antibody fv fragment
4	<a href="#">c1qlcB_</a>	 Alignment		14.3	7	<b>PDB header:</b> oxidoreductase/immune system <b>Chain:</b> B: <b>PDB Molecule:</b> cytochrome c oxidase polypeptide ii; <b>PDBTitle:</b> cryo-structure of the paracoccus denitrificans four-subunit2 cytochrome c oxidase in the completely oxidized state3 complexed with an antibody fv fragment
5	<a href="#">d2osoa1</a>	 Alignment		12.4	16	<b>Fold:</b> Ligand-binding domain in the NO signalling and Golgi transport <b>Superfamily:</b> Ligand-binding domain in the NO signalling and Golgi transport <b>Family:</b> MJ1460-like
6	<a href="#">c2lbgA_</a>	 Alignment		10.9	29	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> major prion protein; <b>PDBTitle:</b> structure of the chr of the prion protein in dpc micelles
7	<a href="#">c1fftG_</a>	 Alignment		9.6	11	<b>PDB header:</b> oxidoreductase <b>Chain:</b> G: <b>PDB Molecule:</b> ubiquinol oxidase; <b>PDBTitle:</b> the structure of ubiquinol oxidase from escherichia coli
8	<a href="#">d3dtub2</a>	 Alignment		9.3	9	<b>Fold:</b> Transmembrane helix hairpin <b>Superfamily:</b> Cytochrome c oxidase subunit II-like, transmembrane region <b>Family:</b> Cytochrome c oxidase subunit II-like, transmembrane region
9	<a href="#">c2bbjB_</a>	 Alignment		8.8	16	<b>PDB header:</b> metal transport/membrane protein <b>Chain:</b> B: <b>PDB Molecule:</b> divalent cation transport-related protein; <b>PDBTitle:</b> crystal structure of the cora mg2+ transporter
10	<a href="#">c1p58F_</a>	 Alignment		8.4	24	<b>PDB header:</b> virus <b>Chain:</b> F: <b>PDB Molecule:</b> envelope protein m; <b>PDBTitle:</b> complex organization of dengue virus membrane proteins as revealed by2 9.5 angstrom cryo-em reconstruction
11	<a href="#">d2csba4</a>	 Alignment		7.7	14	<b>Fold:</b> SAM domain-like <b>Superfamily:</b> RuvA domain 2-like <b>Family:</b> Topoisomerase V repeat domain

12	<a href="#">c2l9uA_</a>	Alignment		6.3	19	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> receptor tyrosine-protein kinase erbb-3; <b>PDBTitle:</b> spatial structure of dimeric erbb3 transmembrane domain
13	<a href="#">d1w6ka2</a>	Alignment		6.1	9	<b>Fold:</b> alpha/alpha toroid <b>Superfamily:</b> Terpenoid cyclases/Protein prenyltransferases <b>Family:</b> Terpene synthases
14	<a href="#">d1y5ic1</a>	Alignment		5.9	14	<b>Fold:</b> Heme-binding four-helical bundle <b>Superfamily:</b> Respiratory nitrate reductase 1 gamma chain <b>Family:</b> Respiratory nitrate reductase 1 gamma chain

15 [c2hn8A](#)

Alignment



5.7

26

**PDB header:**viral protein  
**Chain:** A: **PDB Molecule:**protein pb1-f2;  
**PDBTitle:** structural characterization and oligomerization of pb1-f2,2 a pro-apoptotic influenza a virus protein