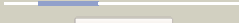


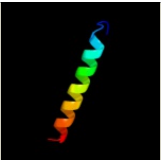
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1w5cL_	 Alignment		22.2	32	PDB header: photosynthesis Chain: L: PDB Molecule: cytochrome b559 beta subunit; PDBTitle: photosystem ii from thermosynechococcus elongatus
2	d2axtf1	 Alignment		21.5	32	Fold: Single transmembrane helix Superfamily: Cytochrome b559 subunits Family: Cytochrome b559 subunits
3	d1c8sa_	 Alignment		9.4	19	Fold: Family A G protein-coupled receptor-like Superfamily: Family A G protein-coupled receptor-like Family: Bacteriorhodopsin-like
4	d1v54e_	 Alignment		8.5	31	Fold: alpha-alpha superhelix Superfamily: Cytochrome c oxidase subunit E Family: Cytochrome c oxidase subunit E
5	c2y69R_	 Alignment		8.4	31	PDB header: electron transport Chain: R: PDB Molecule: cytochrome c oxidase subunit 5a; PDBTitle: bovine heart cytochrome c oxidase re-refined with molecular2 oxygen
6	c2k6tA_	 Alignment		7.3	75	PDB header: hormone Chain: A: PDB Molecule: insulin-like 3 a chain; PDBTitle: solution structure of the relaxin-like factor
7	c2h8bA_	 Alignment		7.3	75	PDB header: hormone/growth factor Chain: A: PDB Molecule: insulin-like 3; PDBTitle: solution structure of insl3
8	c2k6uA_	 Alignment		7.2	75	PDB header: hormone Chain: A: PDB Molecule: insulin-like 3 a chain; PDBTitle: the solution structure of a conformationally restricted2 fully active derivative of the human relaxin-like factor3 (rlf)

9

[c1m0kA](#)

Alignment



6.6

19

PDB header:ion transport
Chain: A: **PDB Molecule:**bacteriorhodopsin;
PDBTitle: bacteriorhodopsin k intermediate at 1.43 a resolution