
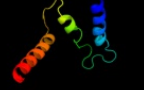
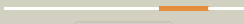


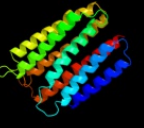





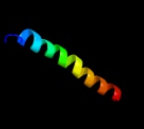






Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P31125
Date	Thu Jan 5 11:47:10 GMT 2012
Unique Job ID	566e556a278c732a

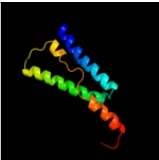
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1s7ba_	 Alignment		97.7	15	Fold: Multidrug resistance efflux transporter EmrE Superfamily: Multidrug resistance efflux transporter EmrE Family: Multidrug resistance efflux transporter EmrE
2	c2i68B_	 Alignment		89.2	24	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
3	c1xioA_	 Alignment		24.3	5	PDB header: signaling protein Chain: A: PDB Molecule: anabaena sensory rhodopsin; PDBTitle: anabaena sensory rhodopsin
4	d1xioa_	 Alignment		24.3	5	Fold: Family A G protein-coupled receptor-like Superfamily: Family A G protein-coupled receptor-like Family: Bacteriorhodopsin-like
5	c2jp3A_	 Alignment		12.5	18	PDB header: transcription Chain: A: PDB Molecule: fxyd domain-containing ion transport regulator 4; PDBTitle: solution structure of the human fxyd4 (chif) protein in sds2 micelles
6	c2rddB_	 Alignment		7.7	7	PDB header: membrane protein/transport protein Chain: B: PDB Molecule: upf0092 membrane protein yajc; PDBTitle: x-ray crystal structure of acrb in complex with a novel2 transmembrane helix.
7	c1w8xP_	 Alignment		7.4	25	PDB header: virus Chain: P: PDB Molecule: protein p16; PDBTitle: structural analysis of prd1
8	c2jwaA_	 Alignment		7.4	14	PDB header: transferase Chain: A: PDB Molecule: receptor tyrosine-protein kinase erbb-2; PDBTitle: erbb2 transmembrane segment dimer spatial structure

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[c3aqpB_](#)

Alignment



6.6

9

PDB header:membrane protein
Chain: B: **PDB Molecule:**probable secdf protein-export membrane protein;
PDBTitle: crystal structure of secdf, a translocon-associated membrane protein,2 from thermus thrmophilus