

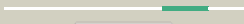
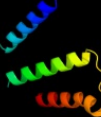















#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d1s7ba_</a>	 Alignment		97.3	10	<b>Fold:</b> Multidrug resistance efflux transporter EmrE <b>Superfamily:</b> Multidrug resistance efflux transporter EmrE <b>Family:</b> Multidrug resistance efflux transporter EmrE
2	<a href="#">c2i68B_</a>	 Alignment		42.8	16	<b>PDB header:</b> transport protein <b>Chain:</b> B: <b>PDB Molecule:</b> protein emre; <b>PDBTitle:</b> cryo-em based theoretical model structure of transmembrane2 domain of the multidrug-resistance antiporter from e. coli3 emre
3	<a href="#">c1lfpA_</a>	 Alignment		31.8	32	<b>PDB header:</b> virus <b>Chain:</b> A: <b>PDB Molecule:</b> major coat protein assembly; <b>PDBTitle:</b> inovirus (filamentous bacteriophage) strain pf3 major coat2 protein assembly
4	<a href="#">c2jx4A_</a>	 Alignment		9.2	38	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> vasopressin v2 receptor; <b>PDBTitle:</b> nmr structure of the intracellular loop (i3) of the2 vasopressin v2 receptor (gpcr)
5	<a href="#">c3aaqB_</a>	 Alignment		8.8	12	<b>PDB header:</b> membrane protein <b>Chain:</b> B: <b>PDB Molecule:</b> probable secdf protein-export membrane protein; <b>PDBTitle:</b> crystal structure of secdf, a translocon-associated membrane protein,2 from thermus thrmophilus
6	<a href="#">d1pw4a_</a>	 Alignment		8.5	11	<b>Fold:</b> MFS general substrate transporter <b>Superfamily:</b> MFS general substrate transporter <b>Family:</b> Glycerol-3-phosphate transporter
7	<a href="#">d1u7ga_</a>	 Alignment		8.0	12	<b>Fold:</b> Ammonium transporter <b>Superfamily:</b> Ammonium transporter <b>Family:</b> Ammonium transporter
8	<a href="#">d1iwga7</a>	 Alignment		7.6	14	<b>Fold:</b> Multidrug efflux transporter AcrB transmembrane domain <b>Superfamily:</b> Multidrug efflux transporter AcrB transmembrane domain <b>Family:</b> Multidrug efflux transporter AcrB transmembrane domain
9	<a href="#">c3rkoF_</a>	 Alignment		7.2	10	<b>PDB header:</b> oxidoreductase <b>Chain:</b> F: <b>PDB Molecule:</b> nadh-quinone oxidoreductase subunit j; <b>PDBTitle:</b> crystal structure of the membrane domain of respiratory complex i from2 e. coli at 3.0 angstrom resolution

10	<a href="#">c2kncA</a>	Alignment		5.8	18	<b>PDB header:</b> cell adhesion <b>Chain:</b> A: <b>PDB Molecule:</b> integrin alpha-iiB; <b>PDBTitle:</b> platelet integrin alphaIIb-beta3 transmembrane-cytoplasmic2 heterocomplex
----	------------------------	-----------	--	-----	----	---