





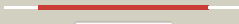













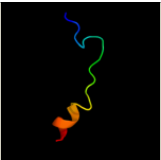


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c3giaA_</a>	 Alignment		100.0	31	<b>PDB header:</b> transport protein <b>Chain:</b> A: <b>PDB Molecule:</b> uncharacterized protein mj0609; <b>PDBTitle:</b> crystal structure of apct transporter
2	<a href="#">c3lrcC_</a>	 Alignment		100.0	18	<b>PDB header:</b> transport protein <b>Chain:</b> C: <b>PDB Molecule:</b> arginine/agmatine antiporter; <b>PDBTitle:</b> structure of e. coli adic (p1)
3	<a href="#">c2jlnA_</a>	 Alignment		100.0	11	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> mhp1; <b>PDBTitle:</b> structure of mhp1, a nucleobase-cation-symport-1 family2 transporter
4	<a href="#">c3dh4A_</a>	 Alignment		99.2	12	<b>PDB header:</b> transport protein <b>Chain:</b> A: <b>PDB Molecule:</b> sodium/glucose cotransporter; <b>PDBTitle:</b> crystal structure of sodium/sugar symporter with bound galactose from2 vibrio parahaemolyticus
5	<a href="#">c2xq2A_</a>	 Alignment		99.2	12	<b>PDB header:</b> transport protein <b>Chain:</b> A: <b>PDB Molecule:</b> sodium/glucose cotransporter; <b>PDBTitle:</b> structure of the k294a mutant of vsgl1
6	<a href="#">d2a65a1</a>	 Alignment		97.6	15	<b>Fold:</b> SNF-like <b>Superfamily:</b> SNF-like <b>Family:</b> SNF-like
7	<a href="#">c2w8aC_</a>	 Alignment		96.0	11	<b>PDB header:</b> membrane protein <b>Chain:</b> C: <b>PDB Molecule:</b> glycine betaine transporter betp; <b>PDBTitle:</b> crystal structure of the sodium-coupled glycine betaine2 symporter betp from corynebacterium glutamicum with bound3 substrate
8	<a href="#">c3hfxA_</a>	 Alignment		87.6	9	<b>PDB header:</b> transport protein <b>Chain:</b> A: <b>PDB Molecule:</b> l-carnitine/gamma-butyrobetaine antiporter; <b>PDBTitle:</b> crystal structure of carnitine transporter
9	<a href="#">d2iuba2</a>	 Alignment		15.5	11	<b>Fold:</b> Transmembrane helix hairpin <b>Superfamily:</b> Magnesium transport protein CorA, transmembrane region <b>Family:</b> Magnesium transport protein CorA, transmembrane region
10	<a href="#">c2wwbA_</a>	 Alignment		9.5	11	<b>PDB header:</b> ribosome <b>Chain:</b> A: <b>PDB Molecule:</b> protein transport protein sec61 subunit alpha isoform 1; <b>PDBTitle:</b> cryo-em structure of the mammalian sec61 complex bound to the2 actively translating wheat germ 80s ribosome

11	<a href="#">c1uj1A_</a>	Alignment		9.1	23	<p><b>PDB header:</b>membrane protein</p> <p><b>Chain:</b> A: <b>PDB Molecule:</b>potassium voltage-gated channel subfamily h</p> <p><b>PDBTitle:</b> solution structure of the herg k+ channel s5-p2 extracellular linker</p>
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