





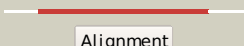

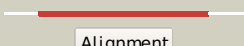

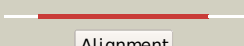

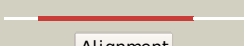





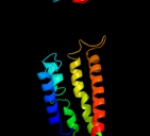

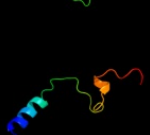
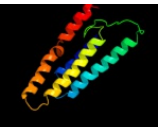


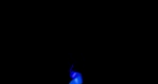
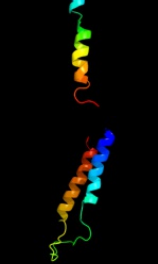


# Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	A0PK1
Date	Tue Apr 3 14:59:03 BST 2012
Unique Job ID	4119271ea0e6f686

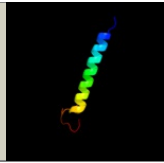
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c2xq2A_</a>	 Alignment		100.0	27	<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 vsglt
2	<a href="#">c3dh4A_</a>	 Alignment		100.0	26	<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
3	<a href="#">c2jlnA_</a>	 Alignment		99.0	12	<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="#">c3giaA_</a>	 Alignment		98.9	11	<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
5	<a href="#">c3lrcC_</a>	 Alignment		98.5	10	<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)
6	<a href="#">c4djiA_</a>	 Alignment		98.4	11	<b>PDB header:</b> transport protein <b>Chain:</b> A; <b>PDB Molecule:</b> probable glutamate/gamma-aminobutyrate antiporter; <b>PDBTitle:</b> structure of glutamate-gaba antiporter gadc
7	<a href="#">c2w8aC_</a>	 Alignment		96.1	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		95.2	11	<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="#">d2a65a1</a>	 Alignment		93.8	14	<b>Fold:</b> SNF-like <b>Superfamily:</b> SNF-like <b>Family:</b> SNF-like
10	<a href="#">c3rlbA_</a>	 Alignment		68.9	8	<b>PDB header:</b> thiamine-binding protein <b>Chain:</b> A; <b>PDB Molecule:</b> thit; <b>PDBTitle:</b> crystal structure at 2.0 a of the s-component for thiamin from an ecf-2 type abc transporter
11	<a href="#">c3hd6A_</a>	 Alignment		30.1	10	<b>PDB header:</b> membrane protein, transport protein <b>Chain:</b> A; <b>PDB Molecule:</b> ammonium transporter rh type c; <b>PDBTitle:</b> crystal structure of the human rhesus glycoprotein rhcg

12	<a href="#">c3riaC_</a>	Alignment		28.0	11	<p><b>PDB header:</b>transport protein/immune system</p> <p><b>Chain:</b> C: <b>PDB Molecule:</b> avermectin-sensitive glutamate-gated chloride channel glucl</p> <p><b>PDBTitle:</b> c. elegans glutamate-gated chloride channel (glucl) in complex with 2 fab, ivermectin and iodide.</p>
13	<a href="#">c3hd7A_</a>	Alignment		24.6	9	<p><b>PDB header:</b>exocytosis</p> <p><b>Chain:</b> A: <b>PDB Molecule:</b> vesicle-associated membrane protein 2;</p> <p><b>PDBTitle:</b> helical extension of the neuronal snares complex into the membrane, 2 spacegroup c 1 2 1</p>
14	<a href="#">c2k9yB_</a>	Alignment		6.7	24	<p><b>PDB header:</b>transferase</p> <p><b>Chain:</b> B: <b>PDB Molecule:</b> ephrin type-a receptor 2;</p> <p><b>PDBTitle:</b> epha2 dimeric structure in the lipidic bicelle at ph 5.0</p>
15	<a href="#">c2k9yA_</a>	Alignment		6.7	24	<p><b>PDB header:</b>transferase</p> <p><b>Chain:</b> A: <b>PDB Molecule:</b> ephrin type-a receptor 2;</p> <p><b>PDBTitle:</b> epha2 dimeric structure in the lipidic bicelle at ph 5.0</p>
16	<a href="#">c3mk7F_</a>	Alignment		5.8	10	<p><b>PDB header:</b>oxidoreductase</p> <p><b>Chain:</b> F: <b>PDB Molecule:</b> cytochrome c oxidase, cbb3-type, subunit p;</p> <p><b>PDBTitle:</b> the structure of cbb3 cytochrome oxidase</p>

17 [clzzaA](#)

Alignment



5.6

23

**PDB header:** membrane protein  
**Chain:** A: **PDB Molecule:** stannin;  
**PDBTitle:** solution nmr structure of the membrane protein stannin