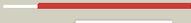
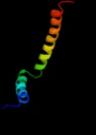
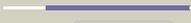
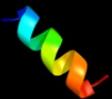
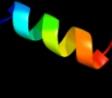
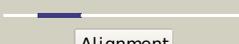
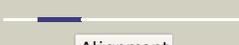


# Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P0A843
Date	Wed Jan 25 15:20:18 GMT 2012
Unique Job ID	34d777637fc236cf

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c2l16A_</a>	 Alignment		99.8	25	<b>PDB header:</b> protein transport <b>Chain:</b> A: <b>PDB Molecule:</b> sec-independent protein translocase protein tatad; <b>PDBTitle:</b> solution structure of bacillus subtilis tatad protein in dpc micelles
2	<a href="#">d2cqqa1</a>	 Alignment		18.5	12	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Myb/SANT domain
3	<a href="#">d2axti1</a>	 Alignment		15.0	23	<b>Fold:</b> Single transmembrane helix <b>Superfamily:</b> Photosystem II reaction center protein I, Psbl <b>Family:</b> Psbl-like
4	<a href="#">d2axti1</a>	 Alignment		9.2	29	<b>Fold:</b> Single transmembrane helix <b>Superfamily:</b> Photosystem II reaction center protein L, PsbL <b>Family:</b> PsbL-like
5	<a href="#">c3a0hl_</a>	 Alignment		9.2	29	<b>PDB header:</b> electron transport <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center protein I; <b>PDBTitle:</b> crystal structure of i-substituted photosystem ii complex
6	<a href="#">c3bz1L_</a>	 Alignment		9.2	29	<b>PDB header:</b> electron transport <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center protein I; <b>PDBTitle:</b> crystal structure of cyanobacterial photosystem ii (part 12 of 2). this file contains first monomer of psii dimer
7	<a href="#">c3prqL_</a>	 Alignment		9.2	29	<b>PDB header:</b> photosynthesis <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center protein I; <b>PDBTitle:</b> crystal structure of cyanobacterial photosystem ii in complex with 2 terbutryn (part 1 of 2). this file contains first monomer of psii3 dimer
8	<a href="#">c2axtl_</a>	 Alignment		9.2	29	<b>PDB header:</b> electron transport <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center I protein; <b>PDBTitle:</b> crystal structure of photosystem ii from thermosynechococcus elongatus
9	<a href="#">c3prrl_</a>	 Alignment		9.2	29	<b>PDB header:</b> photosynthesis <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center protein I; <b>PDBTitle:</b> crystal structure of cyanobacterial photosystem ii in complex with 2 terbutryn (part 2 of 2). this file contains second monomer of psii3 dimer
10	<a href="#">c3arcl_</a>	 Alignment		9.2	29	<b>PDB header:</b> electron transport, photosynthesis <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center protein I; <b>PDBTitle:</b> crystal structure of oxygen-evolving photosystem ii at 1.9 angstrom2 resolution
11	<a href="#">c3bz2L_</a>	 Alignment		9.2	29	<b>PDB header:</b> electron transport <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center protein I; <b>PDBTitle:</b> crystal structure of cyanobacterial photosystem ii (part 22 of 2). this file contains second monomer of psii dimer

12	<a href="#">c3a0bl_</a>	 Alignment		9.2	29	<b>PDB header:</b> electron transport <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center protein I; <b>PDBTitle:</b> crystal structure of br-substituted photosystem ii complex
13	<a href="#">c3kziL_</a>	 Alignment		9.2	29	<b>PDB header:</b> electron transport <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center protein I; <b>PDBTitle:</b> crystal structure of monomeric form of cyanobacterial photosystem ii
14	<a href="#">c1s5ll_</a>	 Alignment		9.2	29	<b>PDB header:</b> photosynthesis <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center I protein; <b>PDBTitle:</b> architecture of the photosynthetic oxygen evolving center
15	<a href="#">c3a0bl_</a>	 Alignment		9.2	29	<b>PDB header:</b> electron transport <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center protein I; <b>PDBTitle:</b> crystal structure of br-substituted photosystem ii complex
16	<a href="#">c1s5ll_</a>	 Alignment		9.2	29	<b>PDB header:</b> photosynthesis <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center I protein; <b>PDBTitle:</b> architecture of the photosynthetic oxygen evolving center
17	<a href="#">c2axtl_</a>	 Alignment		9.2	29	<b>PDB header:</b> electron transport <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center I protein; <b>PDBTitle:</b> crystal structure of photosystem ii from thermosynechococcus elongatus
18	<a href="#">c3a0hl_</a>	 Alignment		9.2	29	<b>PDB header:</b> electron transport <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center protein I; <b>PDBTitle:</b> crystal structure of i-substituted photosystem ii complex
19	<a href="#">d2axtj1</a>	 Alignment		8.3	14	<b>Fold:</b> Single transmembrane helix <b>Superfamily:</b> Photosystem II reaction center protein J, Psbj <b>Family:</b> Psbj-like
20	<a href="#">c3arcl_</a>	 Alignment		8.1	29	<b>PDB header:</b> electron transport, photosynthesis <b>Chain:</b> L: <b>PDB Molecule:</b> photosystem ii reaction center protein I; <b>PDBTitle:</b> crystal structure of oxygen-evolving photosystem ii at 1.9 angstrom2 resolution
21	<a href="#">c1by0A_</a>	 Alignment	not modelled	8.0	47	<b>PDB header:</b> rna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> protein (hepatitis delta antigen); <b>PDBTitle:</b> n-terminal leucine-repeat region of hepatitis delta antigen
22	<a href="#">c1a92B_</a>	 Alignment	not modelled	6.3	41	<b>PDB header:</b> leucine zipper <b>Chain:</b> B: <b>PDB Molecule:</b> delta antigen; <b>PDBTitle:</b> oligomerization domain of hepatitis delta antigen
23	<a href="#">c2kdcC_</a>	 Alignment	not modelled	5.9	9	<b>PDB header:</b> transferase <b>Chain:</b> C: <b>PDB Molecule:</b> diacylglycerol kinase; <b>PDBTitle:</b> nmr solution structure of e. coli diacylglycerol kinase2 (dagk) in dpc micelles