
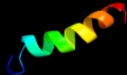
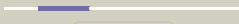

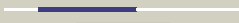






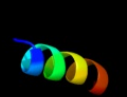










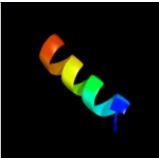



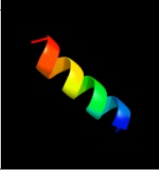


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c2k21A_</a>	 Alignment		11.7	18	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> potassium voltage-gated channel subfamily e <b>PDBTitle:</b> nmr structure of human kcne1 in Impg micelles at ph 6.0 and2 40 degree c
2	<a href="#">c1pd7B_</a>	 Alignment		11.1	56	<b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> mad1; <b>PDBTitle:</b> extended sid of mad1 bound to the pah2 domain of msin3b
3	<a href="#">c3a0ht_</a>	 Alignment		8.2	47	<b>PDB header:</b> electron transport <b>Chain:</b> T: <b>PDB Molecule:</b> photosystem ii reaction center protein t; <b>PDBTitle:</b> crystal structure of i-substituted photosystem ii complex
4	<a href="#">c2axtt_</a>	 Alignment		8.2	47	<b>PDB header:</b> electron transport <b>Chain:</b> T: <b>PDB Molecule:</b> photosystem ii reaction center t protein; <b>PDBTitle:</b> crystal structure of photosystem ii from thermosynechococcus elongatus
5	<a href="#">d2axttl</a>	 Alignment		8.2	47	<b>Fold:</b> Single transmembrane helix <b>Superfamily:</b> Photosystem II reaction center protein T, PsbT <b>Family:</b> PsbT-like
6	<a href="#">c2axtT_</a>	 Alignment		8.2	47	<b>PDB header:</b> electron transport <b>Chain:</b> T: <b>PDB Molecule:</b> photosystem ii reaction center t protein; <b>PDBTitle:</b> crystal structure of photosystem ii from thermosynechococcus elongatus
7	<a href="#">c3kziT_</a>	 Alignment		8.2	47	<b>PDB header:</b> electron transport <b>Chain:</b> T: <b>PDB Molecule:</b> photosystem ii reaction center protein t; <b>PDBTitle:</b> crystal structure of monomeric form of cyanobacterial photosystem ii
8	<a href="#">c3a0hT_</a>	 Alignment		8.2	47	<b>PDB header:</b> electron transport <b>Chain:</b> T: <b>PDB Molecule:</b> photosystem ii reaction center protein t; <b>PDBTitle:</b> crystal structure of i-substituted photosystem ii complex
9	<a href="#">c3arcT_</a>	 Alignment		8.2	47	<b>PDB header:</b> electron transport, photosynthesis <b>Chain:</b> T: <b>PDB Molecule:</b> photosystem ii reaction center protein t; <b>PDBTitle:</b> crystal structure of oxygen-evolving photosystem ii at 1.9 angstrom2 resolution
10	<a href="#">c3a0bt_</a>	 Alignment		8.2	47	<b>PDB header:</b> electron transport <b>Chain:</b> T: <b>PDB Molecule:</b> photosystem ii reaction center protein t; <b>PDBTitle:</b> crystal structure of br-substituted photosystem ii complex
11	<a href="#">c1s5lt_</a>	 Alignment		8.2	47	<b>PDB header:</b> photosynthesis <b>Chain:</b> T: <b>PDB Molecule:</b> photosystem ii psbt protein; <b>PDBTitle:</b> architecture of the photosynthetic oxygen evolving center

12	<a href="#">c1s5lT_</a>	Alignment		8.2	47	<b>PDB header:</b> photosynthesis <b>Chain:</b> T: <b>PDB Molecule:</b> photosystem ii psbt protein; <b>PDBTitle:</b> architecture of the photosynthetic oxygen evolving center
13	<a href="#">c3prpT_</a>	Alignment		8.2	47	<b>PDB header:</b> photosynthesis <b>Chain:</b> T: <b>PDB Molecule:</b> photosystem ii reaction center protein t; <b>PDBTitle:</b> crystal structure of cyanobacterial photosystem ii in complex with2 terbutryn (part 2 of 2). this file contains second monomer of psii3 dimer
14	<a href="#">c3prqT_</a>	Alignment		8.2	47	<b>PDB header:</b> photosynthesis <b>Chain:</b> T: <b>PDB Molecule:</b> photosystem ii reaction center protein t; <b>PDBTitle:</b> crystal structure of cyanobacterial photosystem ii in complex with2 terbutryn (part 1 of 2). this file contains first monomer of psii3 dimer
15	<a href="#">c3bz1T_</a>	Alignment		8.2	47	<b>PDB header:</b> electron transport <b>Chain:</b> T: <b>PDB Molecule:</b> photosystem ii reaction center protein t; <b>PDBTitle:</b> crystal structure of cyanobacterial photosystem ii (part 12 of 2). this file contains first monomer of psii dimer
16	<a href="#">c3bz2T_</a>	Alignment		8.2	47	<b>PDB header:</b> electron transport <b>Chain:</b> T: <b>PDB Molecule:</b> photosystem ii reaction center protein t; <b>PDBTitle:</b> crystal structure of cyanobacterial photosystem ii (part 22 of 2). this file contains second monomer of psii dimer

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[c1qgrB\\_](#)

Alignment



6.0

60

**PDB header:**transport receptor  
**Chain:** B: **PDB Molecule:**protein (importin alpha-2 subunit);  
**PDBTitle:** structure of importin beta bound to the ibb domain of2  
importin alpha (ii crystal form, grown at low ph)