
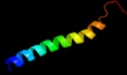





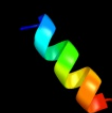
















#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2kncA_	 Alignment		23.9	30	PDB header: cell adhesion Chain: A: PDB Molecule: integrin alpha-iiB; PDBTitle: platelet integrin alpha-iiB-beta3 transmembrane-cytoplasmic2 heterocomplex
2	d2axtl1	 Alignment		18.6	14	Fold: Single transmembrane helix Superfamily: Photosystem II reaction center protein L, PsbL Family: PsbL-like
3	c3a0bl_	 Alignment		18.6	14	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of br-substituted photosystem ii complex
4	c3prrl_	 Alignment		18.6	14	PDB header: photosynthesis Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of cyanobacterial photosystem ii in complex with 2 terbutryn (part 2 of 2). this file contains second monomer of psii3 dimer
5	c1s5l_	 Alignment		18.6	14	PDB header: photosynthesis Chain: L: PDB Molecule: photosystem ii reaction center I protein; PDBTitle: architecture of the photosynthetic oxygen evolving center
6	c3a0bl_	 Alignment		18.6	14	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of br-substituted photosystem ii complex
7	c2axtl_	 Alignment		18.6	14	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center I protein; PDBTitle: crystal structure of photosystem ii from thermosynechococcus elongatus
8	c3bz2L_	 Alignment		18.6	14	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of cyanobacterial photosystem ii (part 22 of 2). this file contains second monomer of psii dimer
9	c1s5l_	 Alignment		18.6	14	PDB header: photosynthesis Chain: L: PDB Molecule: photosystem ii reaction center I protein; PDBTitle: architecture of the photosynthetic oxygen evolving center
10	c3a0hl_	 Alignment		18.6	14	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of i-substituted photosystem ii complex
11	c3kzil_	 Alignment		18.6	14	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of monomeric form of cyanobacterial photosystem ii

12	c3prqL_	Alignment		18.6	14	PDB header: photosynthesis Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of cyanobacterial photosystem ii in complex with2 terbutryn (part 1 of 2). this file contains first monomer of psii3 dimer
13	c3bz1L_	Alignment		18.6	14	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of cyanobacterial photosystem ii (part 12 of 2). this file contains first monomer of psii dimer
14	c3a0hL_	Alignment		18.6	14	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of i-substituted photosystem ii complex
15	c3arcL_	Alignment		18.6	14	PDB header: electron transport, photosynthesis Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of oxygen-evolving photosystem ii at 1.9 angstrom2 resolution
16	c2axtl_	Alignment		18.6	14	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center I protein; PDBTitle: crystal structure of photosystem ii from thermosynechococcus elongatus
17	c3aa6C_	Alignment		10.1	38	PDB header: protein binding Chain: C: PDB Molecule: 23mer peptide from cd2-associated protein; PDBTitle: crystal structure of actin capping protein in complex with the cp-2 binding motif derived from cd2ap
18	c2k1aA_	Alignment		8.9	35	PDB header: cell adhesion Chain: A: PDB Molecule: integrin alpha-iib; PDBTitle: bicelle-embedded integrin alpha(iib) transmembrane segment
19	d1fftb2	Alignment		8.3	10	Fold: Transmembrane helix hairpin Superfamily: Cytochrome c oxidase subunit II-like, transmembrane region Family: Cytochrome c oxidase subunit II-like, transmembrane region
20	d2j0sc1	Alignment		8.2	42	Fold: Mago nashi protein Superfamily: Mago nashi protein Family: Mago nashi protein
21	d1rk8b_	Alignment	not modelled	7.2	36	Fold: Mago nashi protein Superfamily: Mago nashi protein Family: Mago nashi protein
22	c2fd4A_	Alignment	not modelled	7.0	57	PDB header: ligase Chain: A: PDB Molecule: avirulence protein avrptob; PDBTitle: crystal structure of avrptob (436-553)
23	d2b9va2	Alignment	not modelled	7.0	32	Fold: alpha/beta-Hydrolases Superfamily: alpha/beta-Hydrolases Family: PepX catalytic domain-like
24	c2gidP_	Alignment	not modelled	6.4	42	PDB header: translation Chain: P: PDB Molecule: mitochondrial rna-binding protein 2; PDBTitle: crystal structures of trypanosoma brucei mrp1/mrp2
25	c1z65A_	Alignment	not modelled	6.2	26	PDB header: unknown function Chain: A: PDB Molecule: prion-like protein doppel; PDBTitle: mouse doppel 1-30 peptide