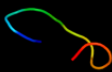
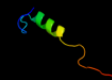
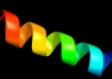
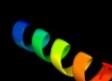
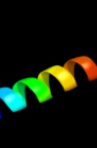


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2cuqa1	 Alignment		30.2	33	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
2	c4eazC_	 Alignment		8.1	46	PDB header: transcription Chain: C: PDB Molecule: transcription cofactor vestigial-like protein 1; PDBTitle: vgl11-tead4 structure
3	d1oe4a_	 Alignment		6.8	45	Fold: Uracil-DNA glycosylase-like Superfamily: Uracil-DNA glycosylase-like Family: Single-strand selective monofunctional uracil-DNA glycosylase SMUG1
4	c1s51L_	 Alignment		6.5	50	PDB header: photosynthesis Chain: L: PDB Molecule: photosystem ii reaction center I protein; PDBTitle: architecture of the photosynthetic oxygen evolving center
5	c3arcL_	 Alignment		6.5	50	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
6	c4fbyL_	 Alignment		6.5	50	PDB header: photosynthesis Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: fs x-ray diffraction of photosystem ii
7	c3bz1L_	 Alignment		6.5	50	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of cyanobacterial photosystem ii (part 1 of 2). this2 file contains first monomer of psii dimer
8	c2axtL_	 Alignment		6.5	50	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center I protein; PDBTitle: crystal structure of photosystem ii from thermosynechococcus elongatus
9	c3bz2L_	 Alignment		6.5	50	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of cyanobacterial photosystem ii (part 2 of 2). this2 file contains second monomer of psii dimer
10	c3prrl_	 Alignment		6.5	50	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 2 of 2). this file contains second monomer of psii3 dimer
11	c4fbyd_	 Alignment		6.5	50	PDB header: photosynthesis Chain: D: PDB Molecule: photosystem ii d2 protein; PDBTitle: fs x-ray diffraction of photosystem ii

12	c1s5lI_	Alignment		6.5	50	PDB header: photosynthesis Chain: L: PDB Molecule: photosystem ii reaction center I protein; PDBTitle: architecture of the photosynthetic oxygen evolving center
13	c2axtl_	Alignment		6.5	50	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center I protein; PDBTitle: crystal structure of photosystem ii from thermosynechococcus elongatus
14	c3prgL_	Alignment		6.5	50	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
15	c3a0hl_	Alignment		6.5	50	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of i-substituted photosystem ii complex
16	d2axtl1	Alignment		6.5	50	Fold: Single transmembrane helix Superfamily: Photosystem II reaction center protein L, PsbL Family: PsbL-like
17	c3a0bl_	Alignment		6.5	50	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of br-substituted photosystem ii complex
18	c3a0hL_	Alignment		6.5	50	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of i-substituted photosystem ii complex
19	c3kziL_	Alignment		6.5	50	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of monomeric form of cyanobacterial photosystem ii
20	c3a0bL_	Alignment		6.5	50	PDB header: electron transport Chain: L: PDB Molecule: photosystem ii reaction center protein I; PDBTitle: crystal structure of br-substituted photosystem ii complex
21	c2latA_	Alignment	not modelled	6.1	86	PDB header: membrane protein Chain: A: PDB Molecule: dolichyl-diphosphooligosaccharide--protein PDBTitle: solution structure of a human minimembrane protein ost4
22	c1h4uA_	Alignment	not modelled	6.0	35	PDB header: extracellular matrix protein Chain: A: PDB Molecule: nidogen-1; PDBTitle: domain g2 of mouse nidogen-1
23	c1dfwA_	Alignment	not modelled	5.8	100	PDB header: immune system Chain: A: PDB Molecule: lung surfactant protein b; PDBTitle: conformational mapping of the n-terminal segment of2 surfactant protein b in lipid using 13c-enhanced fourier3 transform infrared spectroscopy (ftir)
24	c4ei0B_	Alignment	not modelled	5.4	37	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: uncharacterized hypothetical protein; PDBTitle: crystal structure of a hypothetical protein (parmer_03218) from2 parabacteroides merdae atcc 43184 at 2.00 a resolution