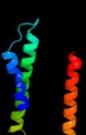
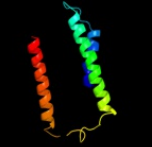
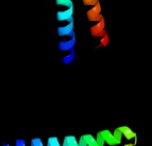
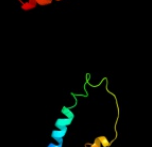
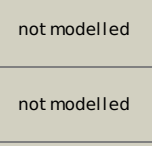


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1om2a_	 Alignment		32.8	30	Fold: Open three-helical up-and-down bundle Superfamily: Mitochondrial import receptor subunit Tom20 Family: Mitochondrial import receptor subunit Tom20
2	d1n1ca_	 Alignment		18.8	17	Fold: TorD-like Superfamily: TorD-like Family: TorD-like
3	c2d68A_	 Alignment		14.1	20	PDB header: cell cycle Chain: A: PDB Molecule: fop; PDBTitle: structure of the n-terminal domain of fop (fgfr1op) protein
4	c2lc2A_	 Alignment		12.4	33	PDB header: protein binding Chain: A: PDB Molecule: avr3a4; PDBTitle: solution structure of the rxlr effector p. capsici avr3a4
5	c3g43F_	 Alignment		11.8	50	PDB header: metal binding protein Chain: F: PDB Molecule: voltage-dependent l-type calcium channel subunit PDBTitle: crystal structure of the calmodulin-bound cav1.2 c-terminal2 regulatory domain dimer
6	d1p4ea1	 Alignment		10.9	50	Fold: SAM domain-like Superfamily: lambda integrase-like, N-terminal domain Family: lambda integrase-like, N-terminal domain
7	c2i0gA_	 Alignment		10.6	36	PDB header: protein binding Chain: A: PDB Molecule: dna polymerase iota; PDBTitle: solution nmr structure of ubiquitin-binding motif (ubm2) of human2 polymerase iota
8	c3b4qA_	 Alignment		10.6	46	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized protein; PDBTitle: crystal structure of a conserved protein domain (unknown2 function) from corynebacterium diphtheriae
9	c2khuA_	 Alignment		10.0	36	PDB header: transferase/protein binding Chain: A: PDB Molecule: immunoglobulin g-binding protein g, dna PDBTitle: solution structure of the ubiquitin-binding motif of human2 polymerase iota
10	d2ipqx1	 Alignment		10.0	40	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: STY4665 C-terminal domain-like
11	c2axtc_	 Alignment		9.4	21	PDB header: electron transport Chain: C: PDB Molecule: photosystem ii cp43 protein; PDBTitle: crystal structure of photosystem ii from thermosynechococcus elongatus

12	d2axtc1	Alignment		9.4	21	Fold: Photosystem II antenna protein-like Superfamily: Photosystem II antenna protein-like Family: Photosystem II antenna protein-like
13	c3mk7F_	Alignment		9.2	16	PDB header: oxidoreductase Chain: F: PDB Molecule: cytochrome c oxidase, cbb3-type, subunit p; PDBTitle: the structure of cbb3 cytochrome oxidase
14	d3proc1	Alignment		8.4	14	Fold: Alpha-lytic protease prodomain-like Superfamily: Alpha-lytic protease prodomain Family: Alpha-lytic protease prodomain
15	c2v1sD_	Alignment		7.7	30	PDB header: oxidoreductase Chain: D: PDB Molecule: mitochondrial import receptor subunit tom20 homolog; PDBTitle: crystal structure of rat tom20-albh presequence complex
16	c1h2sB_	Alignment		7.6	14	PDB header: membrane protein Chain: B: PDB Molecule: sensory rhodopsin ii transducer; PDBTitle: molecular basis of transmembrane signalling by sensory2 rhodopsin ii-transducer complex
17	d1h2sb_	Alignment		7.6	14	Fold: Transmembrane helix hairpin Superfamily: Htr2 transmembrane domain-like Family: Htr2 transmembrane domain-like
18	c1fftG_	Alignment		7.6	9	PDB header: oxidoreductase Chain: G: PDB Molecule: ubiquinol oxidase; PDBTitle: the structure of ubiquinol oxidase from escherichia coli
19	c2retE_	Alignment		6.8	25	PDB header: protein transport Chain: E: PDB Molecule: pseudopilin epsi; PDBTitle: the crystal structure of a binary complex of two pseudopilins: epsi2 and epsj from the type 2 secretion system of vibrio vulnificus
20	c2qm2B_	Alignment		6.7	19	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: putative hopj type iii effector protein; PDBTitle: putative hopj type iii effector protein from vibrio parahaemolyticus
21	d2reta1	Alignment	not modelled	6.3	25	Fold: Pili subunits Superfamily: Pili subunits Family: GSPII I/J protein-like
22	d1c99a_	Alignment	not modelled	6.3	21	Fold: Transmembrane helix hairpin Superfamily: F1F0 ATP synthase subunit C Family: F1F0 ATP synthase subunit C
23	d3b2ua2	Alignment	not modelled	5.8	40	Fold: Knottins (small inhibitors, toxins, lectins) Superfamily: Growth factor receptor domain Family: Growth factor receptor domain
24	d1dxa1	Alignment	not modelled	5.7	23	Fold: CH domain-like Superfamily: Calponin-homology domain, CH-domain Family: Calponin-homology domain, CH-domain
25	d2fj6a1	Alignment	not modelled	5.3	25	Fold: SAM domain-like Superfamily: YozE-like Family: YozE-like
26	c3m7kA_	Alignment	not modelled	5.2	100	PDB header: hydrolase/dna Chain: A: PDB Molecule: restriction endonuclease paci; PDBTitle: crystal structure of paci-dna enzyme product complex
27	d1bhda_	Alignment	not modelled	5.2	29	Fold: CH domain-like Superfamily: Calponin-homology domain, CH-domain Family: Calponin-homology domain, CH-domain