
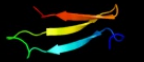







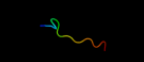

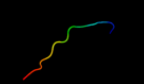

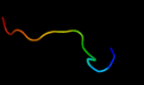









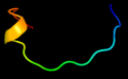




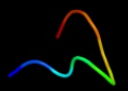
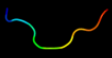



#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1fbvA_	 Alignment		28.2	31	PDB header: ligase Chain: A: PDB Molecule: signal transduction protein cbl; PDBTitle: structure of a cbl-ubch7 complex: ring domain function in2 ubiquitin-protein ligases
2	d3buxb3	 Alignment		27.1	31	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
3	c2cblA_	 Alignment		25.6	31	PDB header: complex (proto-oncogene/peptide) Chain: A: PDB Molecule: proto-oncogene cbl; PDBTitle: n-terminal domain of cbl in complex with its binding site2 on zap-70
4	c3bunB_	 Alignment		25.1	31	PDB header: ligase/signaling protein Chain: B: PDB Molecule: e3 ubiquitin-protein ligase cbl; PDBTitle: crystal structure of c-cbl-tyk domain complexed with its2 binding motif in sprouty4
5	d1bnla_	 Alignment		15.6	38	Fold: C-type lectin-like Superfamily: C-type lectin-like Family: Endostatin
6	c2gyrB_	 Alignment		11.5	38	PDB header: hormone/growth factor Chain: B: PDB Molecule: neurotrophic factor artemin, isoform 3; PDBTitle: crystal structure of human artemin
7	d1koea_	 Alignment		10.5	38	Fold: C-type lectin-like Superfamily: C-type lectin-like Family: Endostatin
8	c2qh7A_	 Alignment		8.6	44	PDB header: metal binding protein Chain: A: PDB Molecule: zinc finger cdgsh-type domain 1; PDBTitle: mitoneet is a uniquely folded 2fe-2s outer mitochondrial membrane2 protein stabilized by pioglitazone
9	c3fnvB_	 Alignment		8.5	44	PDB header: metal binding protein Chain: B: PDB Molecule: cdgsh iron sulfur domain-containing protein 2; PDBTitle: crystal structure of miner1: the redox-active 2fe-2s protein causative2 in wolfram syndrome 2
10	c2zkl_	 Alignment		8.0	60	PDB header: ribosomal protein/rna Chain: L: PDB Molecule: PDBTitle: structure of a mammalian ribosomal 40s subunit within an2 80s complex obtained by docking homology models of the rna3 and proteins into an 8.7 a cryo-em map
11	c3op0B_	 Alignment		7.9	34	PDB header: signaling protein/signaling protein regu Chain: B: PDB Molecule: signal transduction protein cbl-c; PDBTitle: crystal structure of cbl-c (cbl-3) tkb domain in complex with egfr2 py1069 peptide

12	c2pe4A_	Alignment		7.7	33	PDB header: hydrolase Chain: A: PDB Molecule: hyaluronidase-1; PDBTitle: structure of human hyaluronidase 1, a hyaluronan hydrolyzing enzyme2 involved in tumor growth and angiogenesis
13	d2uubl1	Alignment		7.4	47	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
14	c2xzml_	Alignment		7.3	60	PDB header: ribosome Chain: L: PDB Molecule: 40s ribosomal protein s12; PDBTitle: crystal structure of the eukaryotic 40s ribosomal2 subunit in complex with initiation factor 1. this file3 contains the 40s subunit and initiation factor for4 molecule 1
15	d2ilxa1	Alignment		6.7	32	Fold: LigT-like Superfamily: LigT-like Family: 2',3'-cyclic nucleotide 3'-phosphodiesterase, catalytic domain
16	c2zomC_	Alignment		6.5	31	PDB header: unknown function Chain: C: PDB Molecule: protein cuta, chloroplast, putative, expressed; PDBTitle: crystal structure of cuta1 from oryza sativa
17	d1agqa_	Alignment		6.5	54	Fold: Cystine-knot cytokines Superfamily: Cystine-knot cytokines Family: Transforming growth factor (TGF)-beta
18	c1s1hL_	Alignment		6.5	50	PDB header: ribosome Chain: L: PDB Molecule: 40s ribosomal protein s23; PDBTitle: structure of the ribosomal 80s-eef2-sordarin complex from2 yeast obtained by docking atomic models for rna and protein3 components into a 11.7 a cryo-em map. this file, 1s1h,4 contains 40s subunit. the 60s ribosomal subunit is in file5 1s1i.
19	d1ok8a1	Alignment		5.8	36	Fold: Immunoglobulin-like beta-sandwich Superfamily: E set domains Family: Class II viral fusion proteins C-terminal domain
20	c3egpA_	Alignment		5.7	27	PDB header: viral protein Chain: A: PDB Molecule: envelope protein; PDBTitle: crystal structure analysis of dengue-1 envelope protein2 domain iii
21	d2cb2a1	Alignment	not modelled	5.5	44	Fold: Ferredoxin-like Superfamily: Dimeric alpha+beta barrel Family: SOR-like
22	c2zfha_	Alignment	not modelled	5.5	24	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: cuta; PDBTitle: crystal structure of putative cuta1 from homo sapiens at 2.05a2 resolution
23	d1dy2a_	Alignment	not modelled	5.1	42	Fold: C-type lectin-like Superfamily: C-type lectin-like Family: Endostatin