























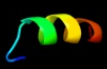
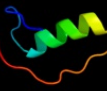




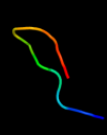


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2dhmA	 Alignment		100.0	100	PDB header: protein binding Chain: A: PDB Molecule: protein bola; PDBTitle: solution structure of the bola protein from escherichia coli
2	c1v60A	 Alignment		100.0	45	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: riken cdna 1810037g04; PDBTitle: solution structure of bola1 protein from mus musculus
3	c3o2eA	 Alignment		100.0	35	PDB header: unknown function Chain: A: PDB Molecule: bola-like protein; PDBTitle: crystal structure of a bol-like protein from babesia bovis
4	c2kdnA	 Alignment		100.0	39	PDB header: unknown function Chain: A: PDB Molecule: putative uncharacterized protein pfe0790c; PDBTitle: solution structure of pfe0790c, a putative bola-like2 protein from the protozoan parasite plasmodium falciparum.
5	d1v9ja	 Alignment		100.0	32	Fold: Alpha-lytic protease prodomain-like Superfamily: BolA-like Family: BolA-like
6	c3tr3A	 Alignment		100.0	30	PDB header: unknown function Chain: A: PDB Molecule: bola; PDBTitle: structure of a bola protein homologue from coxiella burnetii
7	d1ny8a	 Alignment		100.0	29	Fold: Alpha-lytic protease prodomain-like Superfamily: BolA-like Family: BolA-like
8	c1xs3A	 Alignment		100.0	35	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hypothetical protein xc975; PDBTitle: solution structure analysis of the xc975 protein
9	c2kz0A	 Alignment		99.9	31	PDB header: transcription Chain: A: PDB Molecule: bola family protein; PDBTitle: solution structure of a bola protein (ech_0303) from ehrlichia2 chaffeensis. seattle structural genomics center for infectious3 disease target ehcha.10365.a
10	d1ylqa1	 Alignment		96.8	20	Fold: Nucleotidyltransferase Superfamily: Nucleotidyltransferase Family: Catalytic subunit of bi-partite nucleotidyltransferase
11	c2rffA	 Alignment		44.5	15	PDB header: transferase Chain: A: PDB Molecule: putative nucleotidyltransferase; PDBTitle: crystal structure of a putative nucleotidyltransferase2 (np_343093.1) from sulfolobus solfataricus at 1.40 a3 resolution

12	c2hjjA_	Alignment		18.7	33	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hypothetical protein ykff; PDBTitle: solution nmr structure of protein ykff from escherichia coli.2 northeast structural genomics target er397.
13	d2hjjal	Alignment		18.7	33	Fold: dsRBD-like Superfamily: YcfA/nrd intein domain Family: Ykff-like
14	d1ubdc2	Alignment		16.3	27	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
15	c2hw4A_	Alignment		15.8	29	PDB header: structural genomics, hydrolase Chain: A: PDB Molecule: 14 kda phosphohistidine phosphatase; PDBTitle: crystal structure of human phosphohistidine phosphatase
16	d2hw4a1	Alignment		13.0	29	Fold: PHP14-like Superfamily: PHP14-like Family: Janus/Ocnus
17	c2wu8A_	Alignment		12.7	15	PDB header: isomerase Chain: A: PDB Molecule: glucose-6-phosphate isomerase; PDBTitle: structural studies of phosphoglucose isomerase from2 mycobacterium tuberculosis h37rv
18	d1xhja_	Alignment		10.3	19	Fold: Alpha-lytic protease prodomain-like Superfamily: Fe-S cluster assembly (FSCA) domain-like Family: NifU C-terminal domain-like
19	d1ug8a_	Alignment		9.7	31	Fold: IF3-like Superfamily: R3H domain Family: R3H domain
20	d1x63a1	Alignment		9.4	36	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
21	d1znfa_	Alignment	not modelled	9.0	25	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
22	c1znfA_	Alignment	not modelled	9.0	25	PDB header: zinc finger dna binding domain Chain: A: PDB Molecule: 31st zinc finger from xfin; PDBTitle: three-dimensional solution structure of a single zinc2 finger dna-binding domain
23	d1cf7b_	Alignment	not modelled	8.1	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Cell cycle transcription factor e2f-dp
24	d1pvza_	Alignment	not modelled	8.0	24	Fold: Knottins (small inhibitors, toxins, lectins) Superfamily: Scorpion toxin-like Family: Short-chain scorpion toxins
25	c2yv1A_	Alignment	not modelled	7.5	9	PDB header: ligase Chain: A: PDB Molecule: succinyl-coa ligase [adp-forming] subunit alpha; PDBTitle: crystal structure of succinyl-coa synthetase alpha chain from2 methanocaldococcus jannaschii dsm 2661
26	c3hjbA_	Alignment	not modelled	6.3	16	PDB header: isomerase Chain: A: PDB Molecule: glucose-6-phosphate isomerase; PDBTitle: 1.5 angstrom crystal structure of glucose-6-phosphate isomerase from2 vibrio cholerae.
27	c3htuE_	Alignment	not modelled	6.2	25	PDB header: protein transport Chain: E: PDB Molecule: vacuolar protein-sorting-associated protein 25; PDBTitle: crystal structure of the human vps25-vps20 subcomplex
28	c3ljka_	Alignment	not modelled	6.1	18	PDB header: isomerase Chain: A: PDB Molecule: glucose-6-phosphate isomerase; PDBTitle: glucose-6-phosphate isomerase from francisella tularensis.

29	c3ujhB_	Alignment	not modelled	6.1	20	PDB header: isomerase Chain: B: PDB Molecule: glucose-6-phosphate isomerase; PDBTitle: crystal structure of substrate-bound glucose-6-phosphate isomerase2 from toxoplasma gondii
30	d2g7oa1	Alignment	not modelled	6.0	16	Fold: TraM-like Superfamily: TraM-like Family: TraM-like
31	dliata_	Alignment	not modelled	5.9	16	Fold: SIS domain Superfamily: SIS domain Family: Phosphoglucose isomerase, PGI
32	d1pbya2	Alignment	not modelled	5.9	33	Fold: Cytochrome c Superfamily: Cytochrome c Family: Quinohemoprotein amine dehydrogenase A chain, domains 1 and 2
33	c2jsoA_	Alignment	not modelled	5.6	32	PDB header: signaling protein Chain: A: PDB Molecule: polymyxin resistance protein pmrd; PDBTitle: antimicrobial resistance protein
34	d1u0fa_	Alignment	not modelled	5.3	16	Fold: SIS domain Superfamily: SIS domain Family: Phosphoglucose isomerase, PGI
35	d1gzda_	Alignment	not modelled	5.1	16	Fold: SIS domain Superfamily: SIS domain Family: Phosphoglucose isomerase, PGI
36	c2btwA_	Alignment	not modelled	5.1	11	PDB header: transferase Chain: A: PDB Molecule: alr0975 protein; PDBTitle: crystal structure of alr0975