


























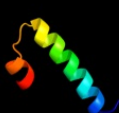





#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3utkA_	 Alignment		95.5	31	PDB header: protein transport Chain: A: PDB Molecule: lipoprotein outs; PDBTitle: structure of the pilotin of the type ii secretion system
2	c3solA_	 Alignment		93.1	25	PDB header: protein transport Chain: A: PDB Molecule: type ii secretion pathway related protein; PDBTitle: crystal structure of the type 2 secretion system pilotin gsp
3	c4a56A_	 Alignment		92.2	26	PDB header: protein transport Chain: A: PDB Molecule: pullulanase secretion protein puls; PDBTitle: crystal structure of the type 2 secretion system pilotin2 from klebsiella oxytoca
4	d2al6a2	 Alignment		42.5	43	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Third domain of FERM
5	c3pntD_	 Alignment		20.2	28	PDB header: hydrolase/hydrolase inhibitor Chain: D: PDB Molecule: immunity factor for spn; PDBTitle: crystal structure of the streptococcus pyogenes nad+ glycohydrolase2 spn in complex with ifs, the immunity factor for spn
6	d1q0qa1	 Alignment		15.3	19	Fold: Left-handed superhelix Superfamily: 1-deoxy-D-xylulose-5-phosphate reductoisomerase, C-terminal domain Family: 1-deoxy-D-xylulose-5-phosphate reductoisomerase, C-terminal domain
7	c1wn4A_	 Alignment		14.1	41	PDB header: plant protein Chain: A: PDB Molecule: vontr protein; PDBTitle: nmr structure of vontr
8	c3qb2A_	 Alignment		13.5	24	PDB header: hydrolase inhibitor Chain: A: PDB Molecule: immunity factor for spn; PDBTitle: the crystal structure of immunity factor for spn (ifs)
9	d1r0ka1	 Alignment		11.6	29	Fold: Left-handed superhelix Superfamily: 1-deoxy-D-xylulose-5-phosphate reductoisomerase, C-terminal domain Family: 1-deoxy-D-xylulose-5-phosphate reductoisomerase, C-terminal domain
10	c1mi1A_	 Alignment		11.5	28	PDB header: signaling protein Chain: A: PDB Molecule: neurobeachin; PDBTitle: crystal structure of the ph-beach domain of human2 neurobeachin
11	d1t77a1	 Alignment		10.9	30	Fold: BEACH domain Superfamily: BEACH domain Family: BEACH domain

12	c3dalA_	Alignment		9.7	21	PDB header: transferase Chain: A: PDB Molecule: pr domain zinc finger protein 1; PDBTitle: methyltransferase domain of human pr domain-containing2 protein 1
13	c2i2xO_	Alignment		8.5	83	PDB header: transferase Chain: O: PDB Molecule: methyltransferase 1; PDBTitle: crystal structure of methanol:cobalamin methyltransferase complex2 mtabc from methanosarcina barkeri
14	c3frwF_	Alignment		8.0	13	PDB header: structural genomics, unknown function Chain: F: PDB Molecule: putative trp repressor protein; PDBTitle: crystal structure of putative trpr protein from ruminococcus obeum
15	d1mi1a1	Alignment		7.9	23	Fold: BEACH domain Superfamily: BEACH domain Family: BEACH domain
16	c3a14B_	Alignment		7.8	33	PDB header: oxidoreductase Chain: B: PDB Molecule: 1-deoxy-d-xylulose 5-phosphate reductoisomerase; PDBTitle: crystal structure of dxr from thermotoga maritima, in complex with2 nadph
17	c3korD_	Alignment		7.5	13	PDB header: transcription Chain: D: PDB Molecule: possible trp repressor; PDBTitle: crystal structure of a putative trp repressor from staphylococcus2 aureus
18	c2wx3A_	Alignment		7.0	50	PDB header: structural protein Chain: A: PDB Molecule: mrna-decapping enzyme 1a; PDBTitle: asymmetric trimer of the human dcp1a c-terminal domain
19	c3mcrA_	Alignment		6.1	19	PDB header: oxidoreductase Chain: A: PDB Molecule: nadh dehydrogenase, subunit c; PDBTitle: crystal structure of nadh dehydrogenase subunit c (tfu_2693) from2 thermobifida fusca yx-er1 at 2.65 a resolution
20	c2jcyA_	Alignment		6.0	24	PDB header: oxidoreductase Chain: A: PDB Molecule: 1-deoxy-d-xylulose 5-phosphate reductoisomerase; PDBTitle: x-ray structure of mutant 1-deoxy-d-xylulose 5-phosphate2 reductoisomerase, dxr, rv2870c, from mycobacterium3 tuberculosis
21	d2p8ta1	Alignment	not modelled	6.0	26	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: PH0730 N-terminal domain-like
22	c3hk0B_	Alignment	not modelled	5.6	17	PDB header: signaling protein Chain: B: PDB Molecule: growth factor receptor-bound protein 10; PDBTitle: crystal structure of the ra and ph domains of grb10
23	c2hm2Q_	Alignment	not modelled	5.5	21	PDB header: apoptosis Chain: Q: PDB Molecule: pyrin-only protein 1; PDBTitle: solution structure of asc2