

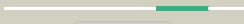

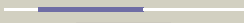


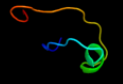

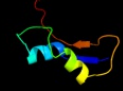












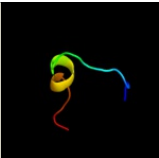
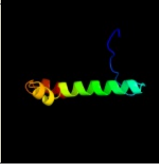


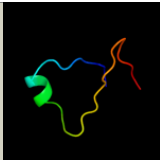
#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2kz6A_	 Alignment		98.3	20	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized protein; PDBTitle: solution structure of protein cv0426 from chromobacterium violaceum, 2 northeast structural genomics consortium (nesg) target cvt2
2	c3o0rC_	 Alignment		41.9	28	PDB header: immune system/oxidoreductase Chain: C: PDB Molecule: nitric oxide reductase subunit c; PDBTitle: crystal structure of nitric oxide reductase from pseudomonas2 aeruginosa in complex with antibody fragment
3	d1fcqa_	 Alignment		17.6	22	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Bee venom hyaluronidase
4	d1d4ba_	 Alignment		14.1	23	Fold: beta-Grasp (ubiquitin-like) Superfamily: CAD & PB1 domains Family: CAD domain
5	c2atmA_	 Alignment		10.9	27	PDB header: hydrolase Chain: A: PDB Molecule: hyaluronoglucosaminidase; PDBTitle: crystal structure of the recombinant allergen ves v 2
6	d1bgva2	 Alignment		8.9	8	Fold: Aminoacid dehydrogenase-like, N-terminal domain Superfamily: Aminoacid dehydrogenase-like, N-terminal domain Family: Aminoacid dehydrogenases
7	d1twfc1	 Alignment		8.7	24	Fold: DCoH-like Superfamily: RBP11-like subunits of RNA polymerase Family: RNA polymerase alpha subunit dimerisation domain
8	d2q22a1	 Alignment		8.3	16	Fold: Ava3019-like Superfamily: Ava3019-like Family: Ava3019-like
9	d1zaka2	 Alignment		7.7	18	Fold: Rubredoxin-like Superfamily: Microbial and mitochondrial ADK, insert "zinc finger" domain Family: Microbial and mitochondrial ADK, insert "zinc finger" domain
10	c2ahmG_	 Alignment		6.8	24	PDB header: viral protein, replication Chain: G: PDB Molecule: replicase polyprotein 1ab, heavy chain; PDBTitle: crystal structure of sars-cov super complex of non-structural 2 proteins: the hexadecamer
11	c1ibxB_	 Alignment		6.3	17	PDB header: hydrolase/hydrolase inhibitor Chain: B: PDB Molecule: chimera of igg binding protein g and dna PDBTitle: nmr structure of dff40 and dff45 n-terminal domain complex

12	dlibxb_	Alignment		6.3	17	Fold: beta-Grasp (ubiquitin-like) Superfamily: CAD & PB1 domains Family: CAD domain
13	dlx4ta1	Alignment		6.0	19	Fold: Long alpha-hairpin Superfamily: ISY1 domain-like Family: ISY1 N-terminal domain-like

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[c2eelA_](#)

Alignment



5.5

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PDB header:apoptosis
Chain: A: **PDB Molecule:**cell death activator cide-a;
PDBTitle: solution structure of the cide-n domain of human cell death2 activator cide-a