












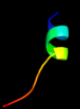















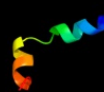



#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2wl8D_	 Alignment		25.0	12	PDB header: protein transport Chain: D: PDB Molecule: peroxisomal biogenesis factor 19; PDBTitle: x-ray crystal structure of pex19p
2	d1q2za_	 Alignment		18.6	11	Fold: alpha-alpha superhelix Superfamily: C-terminal domain of Ku80 Family: C-terminal domain of Ku80
3	c2jeeA_	 Alignment		15.0	36	PDB header: cell cycle Chain: A: PDB Molecule: yiiu; PDBTitle: xray structure of e. coli yiu
4	d1fafa_	 Alignment		14.6	7	Fold: Long alpha-hairpin Superfamily: Chaperone J-domain Family: Chaperone J-domain
5	d2pi8a1	 Alignment		13.5	19	Fold: Double psi beta-barrel Superfamily: Barwin-like endoglucanases Family: MLTA-like
6	c2cwoD_	 Alignment		12.4	19	PDB header: rna binding protein Chain: D: PDB Molecule: rna silencing suppressor; PDBTitle: crystal structure of rna silencing suppressor p21 from beet yellows2 virus
7	c3dveB_	 Alignment		11.0	23	PDB header: membrane protein Chain: B: PDB Molecule: voltage-dependent n-type calcium channel subunit alpha-1b; PDBTitle: crystal structure of ca2+/cam-cav2.2 iq domain complex
8	c2pjwV_	 Alignment		8.7	33	PDB header: endocytosis/exocytosis Chain: V: PDB Molecule: vacuolar protein sorting-associated protein 27; PDBTitle: the vps27/hse1 complex is a gat domain-based scaffold for2 ubiquitin-dependent sorting
9	c1k3vA_	 Alignment		8.3	19	PDB header: virus Chain: A: PDB Molecule: capsid protein vp2; PDBTitle: porcine parvovirus capsid
10	d1k3va_	 Alignment		8.3	19	Fold: Nucleoplasmin-like/VP (viral coat and capsid proteins) Superfamily: ssDNA viruses Family: Parvoviridae-like VP
11	d1c8da_	 Alignment		7.7	13	Fold: Nucleoplasmin-like/VP (viral coat and capsid proteins) Superfamily: ssDNA viruses Family: Parvoviridae-like VP

12	c2wx4C_	Alignment		7.4	50	PDB header: structural protein Chain: C: PDB Molecule: decapping protein 1; PDBTitle: asymmetric trimer of the drosophila melanogaster dcp1 c-2 terminal domain
13	d1kl1a_	Alignment		7.1	16	Fold: PLP-dependent transferase-like Superfamily: PLP-dependent transferases Family: GABA-aminotransferase-like
14	d2nw8a1	Alignment		7.1	20	Fold: Indolic compounds 2,3-dioxygenase-like Superfamily: Indolic compounds 2,3-dioxygenase-like Family: Bacterial tryptophan 2,3-dioxygenase
15	c2noxP_	Alignment		6.9	21	PDB header: oxidoreductase Chain: P: PDB Molecule: tryptophan 2,3-dioxygenase; PDBTitle: crystal structure of tryptophan 2,3-dioxygenase from ralstonia2 metallidurans
16	d2elba1	Alignment		6.8	7	Fold: BAR/IMD domain-like Superfamily: BAR/IMD domain-like Family: BAR domain
17	d1s58a_	Alignment		6.3	25	Fold: Nucleoplasmin-like/VP (viral coat and capsid proteins) Superfamily: ssDNA viruses Family: Parvoviridae-like VP
18	c3dvjB_	Alignment		5.8	30	PDB header: membrane protein Chain: B: PDB Molecule: voltage-dependent n-type calcium channel subunit alpha-1b; PDBTitle: crystal structure of ca2+/cam-cav2.2 iq domain (without cloning2 artifact, hm to tv) complex
19	c2nw7C_	Alignment		5.7	20	PDB header: oxidoreductase Chain: C: PDB Molecule: tryptophan 2,3-dioxygenase; PDBTitle: crystal structure of tryptophan 2,3-dioxygenase (tdo) from2 xanthomonas campestris in complex with ferric heme.3 northeast structural genomics target xcr13
20	c1fpvA_	Alignment		5.7	13	PDB header: virus Chain: A: PDB Molecule: feline panleukopenia virus (strain b) viral PDBTitle: structure determination of feline panleukopenia virus empty2 particles
21	d2ae0x1	Alignment	not modelled	5.4	19	Fold: Double psi beta-barrel Superfamily: Barwin-like endoglucanases Family: MLTA-like
22	c3nttA_	Alignment	not modelled	5.3	25	PDB header: virus Chain: A: PDB Molecule: capsid protein; PDBTitle: structural insights of adeno-associated virus 5. a gene therapy vector2 for cystic fibrosis
23	c3kieF_	Alignment	not modelled	5.2	19	PDB header: virus Chain: F: PDB Molecule: capsid protein vp1; PDBTitle: crystal structure of adeno-associated virus serotype 3b