


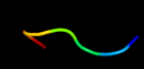











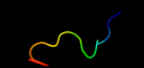





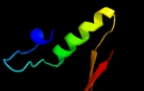
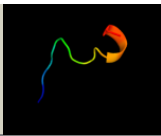


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	dlvqoq1	 Alignment		24.4	33	Fold: SH3-like barrel Superfamily: Translation proteins SH3-like domain Family: Ribosomal proteins L24p and L21e
2	dlag7a_	 Alignment		11.4	86	Fold: Knottins (small inhibitors, toxins, lectins) Superfamily: omega toxin-like Family: Conotoxin
3	c2qtdA_	 Alignment		9.8	29	PDB header: oxidoreductase Chain: A: PDB Molecule: uncharacterized protein mj0327; PDBTitle: crystal structure of a putative dinitrogenase (mj0327) from2 methanocaldococcus jannaschii dsm at 1.70 a resolution
4	c2khgA_	 Alignment		8.7	86	PDB header: antimicrobial protein Chain: A: PDB Molecule: plnj; PDBTitle: plantaricin j in tfe
5	c2pe4A_	 Alignment		8.3	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
6	c2khfA_	 Alignment		7.5	86	PDB header: antimicrobial protein Chain: A: PDB Molecule: plnj; PDBTitle: plantaricin j in dpc-micelles
7	c3m3nW_	 Alignment		6.9	100	PDB header: structural protein Chain: W: PDB Molecule: neural wiskott-aldrich syndrome protein; PDBTitle: structure of a longitudinal actin dimer assembled by tandem w domains
8	c3iz5U_	 Alignment		6.5	40	PDB header: ribosome Chain: U: PDB Molecule: 60s ribosomal protein l21 (l21e); PDBTitle: localization of the large subunit ribosomal proteins into a 5.5 a2 cryo-em map of triticum aestivum translating 80s ribosome
9	dluypa1	 Alignment		6.4	63	Fold: Concanavalin A-like lectins/glucanases Superfamily: Concanavalin A-like lectins/glucanases Family: Glycosyl hydrolases family 32 C-terminal domain
10	d2uuyb1	 Alignment		5.8	33	Fold: BPTI-like Superfamily: BPTI-like Family: Tick tryptase inhibitor-like
11	c2wb6A_	 Alignment		5.8	34	PDB header: viral protein Chain: A: PDB Molecule: afv1-102; PDBTitle: crystal structure of afv1-102, a protein from the acidianus2 filamentous virus 1

12 [c3izcU](#)

Alignment



5.4

47

PDB header: ribosome
Chain: U: **PDB Molecule:** 60s ribosomal protein rpl21 (l21e);
PDB Title: localization of the large subunit ribosomal proteins into a 6.1 a2 cryo-em map of saccharomyces cerevisiae translating 80s ribosome