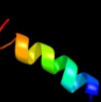
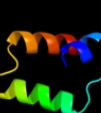


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3aqpB_	 Alignment		23.3	15	PDB header: membrane protein Chain: B: PDB Molecule: probable secdf protein-export membrane protein; PDBTitle: crystal structure of secdf, a translocon-associated membrane protein,2 from thermus thermophilus
2	c3k07A_	 Alignment		12.3	11	PDB header: transport protein Chain: A: PDB Molecule: cation efflux system protein cusa; PDBTitle: crystal structure of cusa
3	c3izcq_	 Alignment		9.5	33	PDB header: ribosome Chain: Q: PDB Molecule: 60s ribosomal protein rpl5 (l18p); PDBTitle: localization of the large subunit ribosomal proteins into a 6.1 a2 cryo-em map of saccharomyces cerevisiae translating 80s ribosome
4	c3u5in_	 Alignment		9.5	33	PDB header: ribosome Chain: N: PDB Molecule: 60s ribosomal protein l15-a; PDBTitle: the structure of the eukaryotic ribosome at 3.0 a resolution
5	c3u5en_	 Alignment		9.5	33	PDB header: ribosome Chain: N: PDB Molecule: 60s ribosomal protein l15-a; PDBTitle: the structure of the eukaryotic ribosome at 3.0 resolution
6	c3izsq_	 Alignment		9.5	33	PDB header: ribosome Chain: Q: PDB Molecule: 60s ribosomal protein rpl5 (l18p); PDBTitle: localization of the large subunit ribosomal proteins into a 6.1 a2 cryo-em map of saccharomyces cerevisiae translating 80s ribosome
7	c2kncA_	 Alignment		7.9	23	PDB header: cell adhesion Chain: A: PDB Molecule: integrin alpha-iiB; PDBTitle: platelet integrin alfaIIB-beta3 transmembrane-cytoplasmic2 heterocomplex
8	d1uoua1	 Alignment		6.1	12	Fold: Methionine synthase domain-like Superfamily: Nucleoside phosphorylase/phosphoribosyltransferase N-terminal domain Family: Nucleoside phosphorylase/phosphoribosyltransferase N-terminal domain
9	d2cpwa1	 Alignment		5.7	15	Fold: RuvA C-terminal domain-like Superfamily: UBA-like Family: UBA domain
10	c2rmzA_	 Alignment		5.2	24	PDB header: cell adhesion Chain: A: PDB Molecule: integrin beta-3; PDBTitle: bicelle-embedded integrin beta3 transmembrane segment

11 [c3iz5q](#)

Alignment



5.1

33

PDB header:ribosome
Chain: Q: **PDB Molecule:**60s ribosomal protein I5 (I18p);
PDBTitle: localization of the large subunit ribosomal proteins into a 5.5 a2 cryo-em map of triticum aestivum translating 80s ribosome