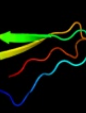
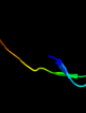
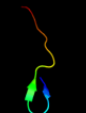
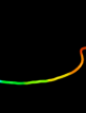
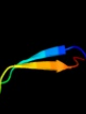
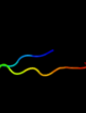
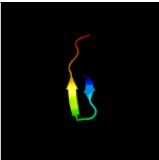


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3bbo6_	 Alignment		99.9	66	PDB header: ribosome Chain: 6: PDB Molecule: ribosomal protein l36; PDBTitle: homology model for the spinach chloroplast 50s subunit fitted to 9.4a2 cryo-em map of the 70s chlororibosome
2	c2qao4_	 Alignment		99.8	53	PDB header: ribosome Chain: 4: PDB Molecule: 50s ribosomal protein l36; PDBTitle: crystal structure of the bacterial ribosome from2 escherichia coli in complex with neomycin. this file3 contains the 50s subunit of the second 70s ribosome, with4 neomycin bound. the entire crystal structure contains two5 70s ribosomes and is described in remark 400.
3	c2qbk4_	 Alignment		99.8	53	PDB header: ribosome Chain: 4: PDB Molecule: 50s ribosomal protein l36; PDBTitle: crystal structure of the bacterial ribosome from escherichia2 coli in complex with gentamicin and ribosome recycling3 factor (rrf). this file contains the 50s subunit of the4 second 70s ribosome, with gentamicin and rrf bound. the5 entire crystal structure contains two 70s ribosomes and is6 described in remark 400.
4	d1dfea_	 Alignment		99.8	62	Fold: Ribosomal protein L36 Superfamily: Ribosomal protein L36 Family: Ribosomal protein L36
5	d2zjr41	 Alignment		99.8	68	Fold: Ribosomal protein L36 Superfamily: Ribosomal protein L36 Family: Ribosomal protein L36
6	d1j0ta_	 Alignment		50.3	23	Fold: Crustacean CHH/MIH/GIH neurohormone Superfamily: Crustacean CHH/MIH/GIH neurohormone Family: Crustacean CHH/MIH/GIH neurohormone
7	d3blza1	 Alignment		20.0	22	Fold: Cystatin-like Superfamily: NTF2-like Family: Sbal0622-like
8	c3dukD_	 Alignment		11.5	28	PDB header: unknown function Chain: D: PDB Molecule: ntf2-like protein of unknown function; PDBTitle: crystal structure of a ntf2-like protein of unknown function2 (mfla_0564) from methylobacillus flagellatus kt at 2.200 a resolution
9	c2zshB_	 Alignment		8.2	56	PDB header: hormone receptor Chain: B: PDB Molecule: della protein gai; PDBTitle: structural basis of gibberellin(ga3)-induced della2 recognition by the gibberellin receptor
10	c1cz6A_	 Alignment		7.8	29	PDB header: toxin Chain: A: PDB Molecule: protein (androctonin); PDBTitle: solution structure of androctonin
11	c3ccjU_	 Alignment		7.3	21	PDB header: ribosome Chain: U: PDB Molecule: 50s ribosomal protein l24e; PDBTitle: structure of anisomycin resistant 50s ribosomal subunit: 23s rrna2 mutation c2534u

12	c4a1dK_	Alignment		7.1	75	PDB header: ribosome Chain: K: PDB Molecule: ubiquitin-60s ribosomal protein l40; PDBTitle: t.thermophila 60s ribosomal subunit in complex with initiation2 factor 6. this file contains 26s rna and proteins of3 molecule 4.
13	c2zkru_	Alignment		6.8	21	PDB header: ribosomal protein/rna Chain: U: PDB Molecule: rna expansion segment es41; PDBTitle: structure of a mammalian ribosomal 60s subunit within an 80s complex2 obtained by docking homology models of the rna and proteins into an3 8.7 a cryo-em map
14	dlvqou1	Alignment		6.8	21	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Ribosomal protein L24e
15	c3hsbB_	Alignment		6.5	29	PDB header: rna binding protein/rna Chain: B: PDB Molecule: protein hfq; PDBTitle: crystal structure of ymah (hfq) from bacillus subtilis in complex with2 an rna aptamer
16	clqqlW_	Alignment		6.0	13	PDB header: translation Chain: W: PDB Molecule: host factor for q beta; PDBTitle: 1.55 a crystal structure of the pleiotropic translational2 regulator, hfq
17	c4a1eT_	Alignment		5.9	14	PDB header: ribosome Chain: T: PDB Molecule: rpl24; PDBTitle: t.thermophila 60s ribosomal subunit in complex with2 initiation factor 6. this file contains 5s rna, 5.8s rna3 and proteins of molecule 1
18	c3izcZ_	Alignment		5.9	14	PDB header: ribosome Chain: Z: PDB Molecule: 60s ribosomal protein rpl24 (l24e); PDBTitle: localization of the large subunit ribosomal proteins into a 6.1 a2 cryo-em map of saccharomyces cerevisiae translating 80s ribosome
19	dlkqla_	Alignment		5.8	13	Fold: Sm-like fold Superfamily: Sm-like ribonucleoproteins Family: Pleiotropic translational regulator Hfq

20	c3iz5Z	Alignment		5.2	14	PDB header: ribosome Chain: Z: PDB Molecule: 60s ribosomal protein l24 (l24e); PDBTitle: localization of the large subunit ribosomal proteins into a 5.5 a2 cryo-em map of triticum aestivum translating 80s ribosome
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