








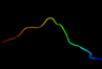

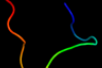





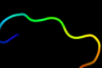




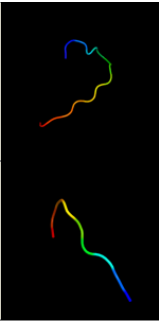
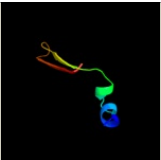


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2jtyA_	 Alignment		35.9	15	PDB header: structural protein Chain: A: PDB Molecule: type-1 fimbrial protein, a chain; PDBTitle: self-complemented variant of fima, the main subunit of type 1 pilus
2	c3jwnL_	 Alignment		30.3	33	PDB header: protein binding/cell adhesion Chain: L: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
3	c3jwnF_	 Alignment		28.7	33	PDB header: protein binding/cell adhesion Chain: F: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
4	c3jwnK_	 Alignment		27.1	33	PDB header: protein binding/cell adhesion Chain: K: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
5	c3jwnE_	 Alignment		27.1	33	PDB header: protein binding/cell adhesion Chain: E: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
6	c2jmrA_	 Alignment		22.3	33	PDB header: cell adhesion Chain: A: PDB Molecule: fimf; PDBTitle: nmr structure of the e. coli type 1 pilus subunit fimf
7	d2j2zb1	 Alignment		15.4	40	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
8	c3aq3A_	 Alignment		11.9	25	PDB header: toxin Chain: A: PDB Molecule: 6b protein; PDBTitle: molecular insights into plant cell proliferation disturbance by2 agrobacterium protein 6b
9	c2eouA_	 Alignment		10.3	25	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 473; PDBTitle: solution structure of the c2h2 type zinc finger (region 370-2 400) of human zinc finger protein 473
10	d2f09a1	 Alignment		8.4	28	Fold: Streptavidin-like Superfamily: YdhA-like Family: YdhA-like
11	d2pa2a1	 Alignment		8.2	18	Fold: alpha/beta-Hammerhead Superfamily: Ribosomal protein L16p/L10e Family: Ribosomal protein L10e

12	c2wkdA_	Alignment		8.0	31	PDB header: dna binding protein Chain: A: PDB Molecule: orf34p2; PDB Title: crystal structure of a double ile-to-met mutant of protein2 orf34 from lactococcus phage p2
13	d2ctda1	Alignment		7.7	56	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2

14 [c2ob9A](#)

Alignment



7.4

20

PDB header:chaperone
Chain: A: **PDB Molecule:**tail assembly chaperone;
PDBTitle: structure of bacteriophage hk97 tail assembly chaperone