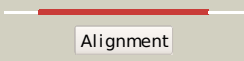
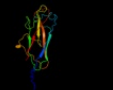
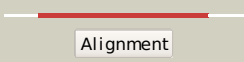
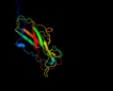
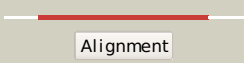
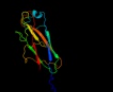
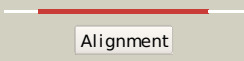
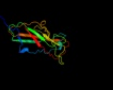
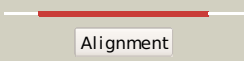
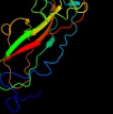
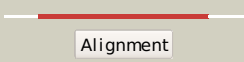
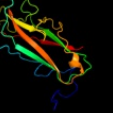
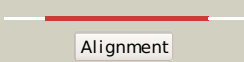
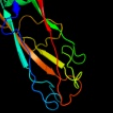
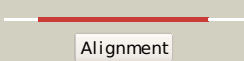
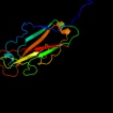
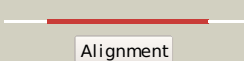
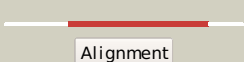
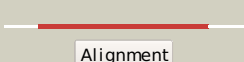
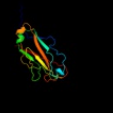
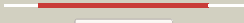
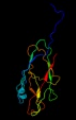
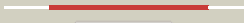
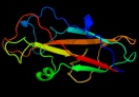
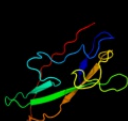


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3jwnK_	 Alignment		99.9	45	PDB header: protein binding/cell adhesion Chain: K: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
2	c3jwnL_	 Alignment		99.9	45	PDB header: protein binding/cell adhesion Chain: L: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
3	c3jwnE_	 Alignment		99.9	45	PDB header: protein binding/cell adhesion Chain: E: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
4	c3jwnF_	 Alignment		99.9	45	PDB header: protein binding/cell adhesion Chain: F: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
5	c2jmrA_	 Alignment		99.9	45	PDB header: cell adhesion Chain: A: PDB Molecule: fimf; PDBTitle: nmr structure of the e. coli type 1 pilus subunit fimf
6	c2jtyA_	 Alignment		99.9	26	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
7	d1pdkb_	 Alignment		99.9	30	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
8	d2j2zb1	 Alignment		99.8	18	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
9	c3bfwA_	 Alignment		99.8	23	PDB header: structural protein/structural protein Chain: A: PDB Molecule: protein fimg; PDBTitle: crystal structure of truncated fimg (fimgt) in complex with the donor2 strand peptide of fimf (dsf)
10	c3bwuF_	 Alignment		99.8	40	PDB header: chaperone, structural, membrane protein Chain: F: PDB Molecule: protein fimf; PDBTitle: crystal structure of the ternary complex of fimd (n-terminal domain,2 fimdn) with fimc and the n-terminally truncated pilus subunit fimf3 (fimft)
11	d2uy6b1	 Alignment		99.8	27	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits

12	c1klfP_	 Alignment		99.7	21	PDB header: chaperone/adhesin complex Chain: P: PDB Molecule: fimh protein; PDBTitle: fimh adhesin-fimc chaperone complex with d-mannose
13	d1ze3h1	 Alignment		99.7	25	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
14	c2w07B_	 Alignment		99.7	15	PDB header: cell adhesion Chain: B: PDB Molecule: minor pilin subunit papf; PDBTitle: structural determinants of polymerization reactivity of the2 p pilus adaptor subunit papf
15	d1n12a_	 Alignment		99.3	20	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
16	c2wmpB_	 Alignment		71.7	13	PDB header: chaperone Chain: B: PDB Molecule: papg protein; PDBTitle: structure of the e. coli chaperone papd in complex with the pilin2 domain of the papgii adhesin
17	d2jnaa1	 Alignment		41.9	35	Fold: Dodecin subunit-like Superfamily: YdgH-like Family: YdgH-like
18	dlp5vb_	 Alignment		39.3	17	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
19	dlz1za1	 Alignment		6.6	17	Fold: Phage tail protein-like Superfamily: Phage tail protein-like Family: Lambda phage gpU-like
20	c1dm9A_	 Alignment		5.2	6	PDB header: structural genomics Chain: A: PDB Molecule: hypothetical 15.5 kd protein in mrca-pcka PDBTitle: heat shock protein 15 kd
21	d1dm9a_	 Alignment	not modelled	5.2	6	Fold: Alpha-L RNA-binding motif Superfamily: Alpha-L RNA-binding motif Family: Heat shock protein 15 kD