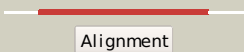
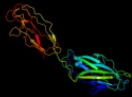
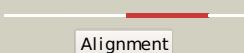
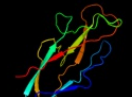
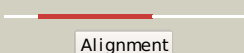

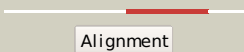
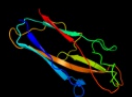
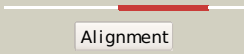




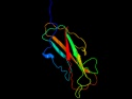
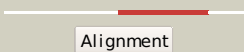

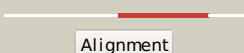
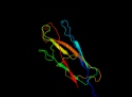


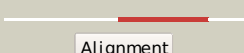

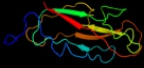


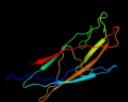
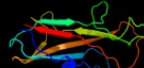

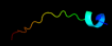




#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1klfP_	 Alignment		100.0	50	PDB header: chaperone/adhesin complex Chain: P: PDB Molecule: fimh protein; PDBTitle: fimh adhesin-fimc chaperone complex with d-mannose
2	dlze3h1	 Alignment		100.0	55	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
3	dluwfa1	 Alignment		99.9	46	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
4	c3bwuF_	 Alignment		99.9	16	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)
5	c3jwnK_	 Alignment		99.9	19	PDB header: protein binding/cell adhesion Chain: K: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
6	c3bfwA_	 Alignment		99.9	26	PDB header: structural protein/structural protein Chain: A: PDB Molecule: protein fimf; PDBTitle: crystal structure of truncated fimg (fimgt) in complex with the donor2 strand peptide of fimf (dsf)
7	c3jwnL_	 Alignment		99.9	19	PDB header: protein binding/cell adhesion Chain: L: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
8	c3jwnF_	 Alignment		99.9	19	PDB header: protein binding/cell adhesion Chain: F: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
9	c3jwnE_	 Alignment		99.9	17	PDB header: protein binding/cell adhesion Chain: E: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
10	d2j2zb1	 Alignment		99.9	17	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
11	c2jtyA_	 Alignment		99.8	24	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

12	c2jmrA_	Alignment		99.8	17	PDB header: cell adhesion Chain: A: PDB Molecule: fimf; PDBTitle: nmr structure of the e. coli type 1 pilus subunit fimf
13	d1pdkb_	Alignment		99.8	19	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
14	d2uy6b1	Alignment		99.8	14	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
15	c2w07B_	Alignment		99.7	17	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
16	d1n12a_	Alignment		99.4	19	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
17	c2wmpB_	Alignment		87.6	19	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
18	d2jnaa1	Alignment		25.3	23	Fold: Dodecin subunit-like Superfamily: YdgH-like Family: YdgH-like
19	d2oz4a1	Alignment		9.7	13	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C2 set domains
20	d1na8a_	Alignment		8.0	18	Fold: Immunoglobulin-like beta-sandwich Superfamily: Clathrin adaptor appendage domain Family: gamma-adaptin C-terminal appendage domain-like
21	d1p4ua_	Alignment	not modelled	7.6	17	Fold: Immunoglobulin-like beta-sandwich Superfamily: Clathrin adaptor appendage domain Family: gamma-adaptin C-terminal appendage domain-like
22	c1w3gA_	Alignment	not modelled	7.5	16	PDB header: toxin/lectin Chain: A: PDB Molecule: hemolytic lectin from laetiporus sulphureus; PDBTitle: hemolytic lectin from the mushroom laetiporus sulphureus2 complexed with two n-acetyl/lactosamine molecules.
23	d1kke2	Alignment	not modelled	7.4	17	Fold: Virus attachment protein globular domain Superfamily: Virus attachment protein globular domain Family: Reovirus attachment protein sigma 1 head domain
24	c1p58F_	Alignment	not modelled	6.0	33	PDB header: virus Chain: F: PDB Molecule: envelope protein m; PDBTitle: complex organization of dengue virus membrane proteins as revealed by2 9.5 angstrom cryo-em reconstruction