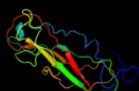
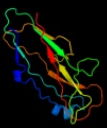
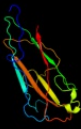
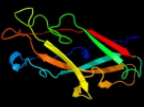
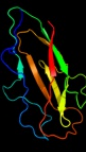
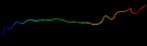
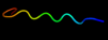
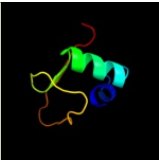


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3jwnL_	 Alignment		100.0	36	PDB header: protein binding/cell adhesion Chain: L: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
2	c3jwnK_	 Alignment		100.0	36	PDB header: protein binding/cell adhesion Chain: K: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
3	c3jwnE_	 Alignment		100.0	36	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	36	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	37	PDB header: cell adhesion Chain: A: PDB Molecule: fimf; PDBTitle: nmr structure of the e. coli type 1 pilus subunit fimf
6	d2j2zb1	 Alignment		99.9	20	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
7	c2jtyA_	 Alignment		99.9	27	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
8	d2uy6b1	 Alignment		99.9	24	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
9	c3bfwA_	 Alignment		99.9	18	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	d1pdkb_	 Alignment		99.9	22	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
11	c3bwuF_	 Alignment		99.8	37	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)

12	d1ze3h1	Alignment		99.8	16	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
13	c1klfp_	Alignment		99.8	14	PDB header: chaperone/adhesin complex Chain: P: PDB Molecule: fimh protein; PDBTitle: fimh adhesin-fimc chaperone complex with d-mannose
14	c2w07B_	Alignment		99.8	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		99.5	20	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
16	c2wmpB_	Alignment		61.8	17	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		7.5	31	Fold: Dodecin subunit-like Superfamily: YdgH-like Family: YdgH-like
18	d1z1za1	Alignment		5.2	33	Fold: Phage tail protein-like Superfamily: Phage tail protein-like Family: Lambda phage gpU-like

19 [c1dm9A](#)

Alignment



5.1

10

PDB header: structural genomics
Chain: A: **PDB Molecule:** hypothetical 15.5 kd protein in mrca-pcka
PDBTitle: heat shock protein 15 kd