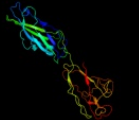
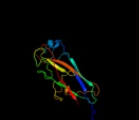
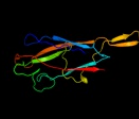
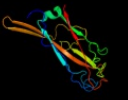
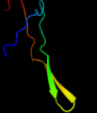
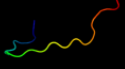
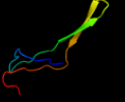
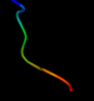


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1klfP_	 Alignment		99.9	16	PDB header: chaperone/adhesin complex Chain: P: PDB Molecule: fimh protein; PDBTitle: fimh adhesin-fimc chaperone complex with d-mannose
2	d1ze3h1	 Alignment		99.8	23	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
3	c2jtyA_	 Alignment		99.8	17	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
4	c3bfwA_	 Alignment		99.8	11	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)
5	c3jwnL_	 Alignment		99.8	17	PDB header: protein binding/cell adhesion Chain: L: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
6	c3jwnK_	 Alignment		99.8	16	PDB header: protein binding/cell adhesion Chain: K: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
7	c3jwnF_	 Alignment		99.8	17	PDB header: protein binding/cell adhesion Chain: F: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
8	c3jwnE_	 Alignment		99.8	16	PDB header: protein binding/cell adhesion Chain: E: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
9	c2jmrA_	 Alignment		99.7	17	PDB header: cell adhesion Chain: A: PDB Molecule: fimf; PDBTitle: nmr structure of the e. coli type 1 pilus subunit fimf
10	d2j2zb1	 Alignment		99.7	13	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
11	c3bwuF_	 Alignment		99.7	17	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	d1pdkb_	Alignment		99.7	16	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
13	d2uy6b1	Alignment		99.6	19	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
14	c2w07B_	Alignment		99.6	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
15	d1n12a_	Alignment		99.4	16	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
16	c2wmpB_	Alignment		42.1	16	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	d1ljma_	Alignment		9.0	14	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: p53-like transcription factors Family: RUNT domain
18	c2qxqB_	Alignment		8.2	27	PDB header: sugar binding protein Chain: B: PDB Molecule: endo-beta-n-acetylglucosaminidase d; PDBTitle: structure of the family 32 carbohydrate-binding module from2 streptococcus pneumoniae endod
19	d1eaqa_	Alignment		8.2	14	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: p53-like transcription factors Family: RUNT domain
20	d2qqia1	Alignment		7.2	0	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: Discoidin domain (FA58C, coagulation factor 5/8 C-terminal domain)
21	d1sddb3	Alignment	not modelled	7.2	23	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: Discoidin domain (FA58C, coagulation factor 5/8 C-terminal domain)
22	d1d7pm_	Alignment	not modelled	7.1	15	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: Discoidin domain (FA58C, coagulation factor 5/8 C-terminal domain)
23	c2qqjA_	Alignment	not modelled	6.7	44	PDB header: signaling protein Chain: A: PDB Molecule: neuropilin-2; PDBTitle: crystal structure of the b1b2 domains from human neuropilin-2
24	d1qbea_	Alignment	not modelled	5.6	27	Fold: RNA bacteriophage capsid protein Superfamily: RNA bacteriophage capsid protein Family: RNA bacteriophage capsid protein
25	c2orxA_	Alignment	not modelled	5.3	11	PDB header: signaling protein, membrane protein Chain: A: PDB Molecule: neuropilin-1; PDBTitle: structural basis for ligand binding and heparin mediated activation of2 neuropilin