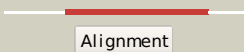

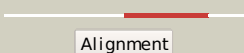
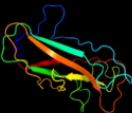


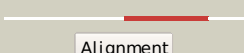





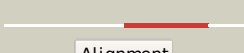
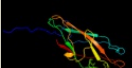



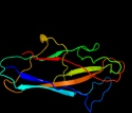

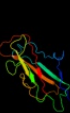







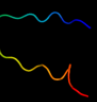
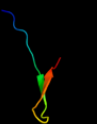




#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1klfP_	 Alignment		99.8	21	PDB header: chaperone/adhesin complex Chain: P: PDB Molecule: fimh protein; PDBTitle: fimh adhesin-fimc chaperone complex with d-mannose
2	c2jtyA_	 Alignment		99.8	25	PDB header: structural protein Chain: A: PDB Molecule: type-1 fimbrial protein, a chain; PDBTitle: self-complemented variant of fima, the main subunit of type 1 pilus
3	d1pdkb_	 Alignment		99.8	20	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
4	c3jwnK_	 Alignment		99.8	21	PDB header: protein binding/cell adhesion Chain: K: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
5	c3jwnL_	 Alignment		99.8	21	PDB header: protein binding/cell adhesion Chain: L: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
6	c3jwnE_	 Alignment		99.8	21	PDB header: protein binding/cell adhesion Chain: E: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
7	c3jwnF_	 Alignment		99.8	21	PDB header: protein binding/cell adhesion Chain: F: PDB Molecule: protein fimf; PDBTitle: complex of fimc, fimf, fimg and fimh
8	c3bfwA_	 Alignment		99.8	17	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)
9	c3bwuF_	 Alignment		99.8	25	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)
10	d2j2zb1	 Alignment		99.8	18	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
11	d1ze3h1	 Alignment		99.7	18	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits

12	c2jmrA_	Alignment		99.7	22	PDB header: cell adhesion Chain: A: PDB Molecule: fimf; PDBTitle: nmr structure of the e. coli type 1 pilus subunit fimf
13	d2uy6b1	Alignment		99.7	21	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
14	c2w07B_	Alignment		99.7	20	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	16	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: Bacterial adhesins Family: Pilus subunits
16	c2wmpB_	Alignment		51.6	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	c3cl5A_	Alignment		15.2	26	PDB header: hydrolase Chain: A: PDB Molecule: hemagglutinin-esterase; PDBTitle: structure of coronavirus hemagglutinin-esterase in complex with 4,9-o-2 diacetyl sialic acid
18	c3nqnB_	Alignment		7.6	13	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: uncharacterized protein; PDBTitle: crystal structure of a protein with unknown function. (dr_2006) from2 deinococcus radiodurans at 1.88 a resolution
19	d3c7bb3	Alignment		7.3	21	Fold: Nitrite and sulphite reductase 4Fe-4S domain-like Superfamily: Nitrite and sulphite reductase 4Fe-4S domain-like Family: Nitrite and sulphite reductase 4Fe-4S domain-like
20	d2akja4	Alignment		6.6	7	Fold: Nitrite and sulphite reductase 4Fe-4S domain-like Superfamily: Nitrite and sulphite reductase 4Fe-4S domain-like Family: Nitrite and sulphite reductase 4Fe-4S domain-like
21	c2i9lL_	Alignment	not modelled	6.5	21	PDB header: immune system/viral protein Chain: L: PDB Molecule: virion membrane protein m25; PDBTitle: structure of fab 7d11 from a neutralizing antibody against the2 poxvirus l1 protein
22	d2v4jb3	Alignment	not modelled	5.9	54	Fold: Nitrite and sulphite reductase 4Fe-4S domain-like Superfamily: Nitrite and sulphite reductase 4Fe-4S domain-like Family: Nitrite and sulphite reductase 4Fe-4S domain-like
23	d3c7ba3	Alignment	not modelled	5.7	31	Fold: Nitrite and sulphite reductase 4Fe-4S domain-like Superfamily: Nitrite and sulphite reductase 4Fe-4S domain-like Family: Nitrite and sulphite reductase 4Fe-4S domain-like
24	d1lvla1	Alignment	not modelled	5.3	22	Fold: FAD/NAD(P)-binding domain Superfamily: FAD/NAD(P)-binding domain Family: FAD/NAD-linked reductases, N-terminal and central domains