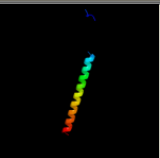
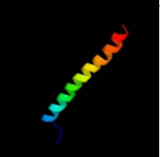
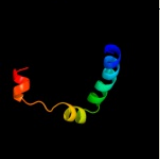
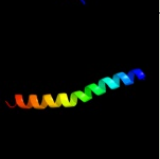
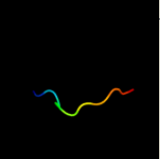
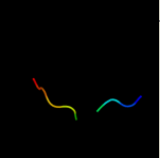
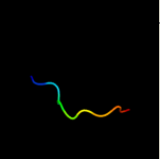
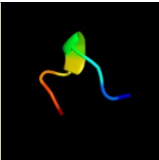


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3a69A_	 Alignment		98.7	13	PDB header: motor protein Chain: A: PDB Molecule: flagellar hook protein flge; PDBTitle: atomic model of the bacterial flagellar hook based on2 docking an x-ray derived structure and terminal two alpha-3 helices into an 7.1 angstrom resolution cryoem map
2	dlucua_	 Alignment		90.4	21	Fold: Phase 1 flagellin Superfamily: Phase 1 flagellin Family: Phase 1 flagellin
3	cloryB_	 Alignment		88.2	26	PDB header: chaperone Chain: B: PDB Molecule: flagellin; PDBTitle: flagellar export chaperone in complex with its cognate binding partner
4	c3k8vB_	 Alignment		44.7	14	PDB header: structural protein Chain: B: PDB Molecule: flagellin homolog; PDBTitle: crysatl structure of a bacterial cell-surface flagellin n20c20
5	dlr6fa_	 Alignment		39.3	23	Fold: Virulence-associated V antigen Superfamily: Virulence-associated V antigen Family: Virulence-associated V antigen
6	dl1voa_	 Alignment		21.7	36	Fold: Trypsin-like serine proteases Superfamily: Trypsin-like serine proteases Family: Viral cysteine protease of trypsin fold
7	c2q6fB_	 Alignment		19.5	55	PDB header: hydrolase Chain: B: PDB Molecule: infectious bronchitis virus (ibv) main protease; PDBTitle: crystal structure of infectious bronchitis virus (ibv) main2 protease in complex with a michael acceptor inhibitor n3
8	c3d23A_	 Alignment		16.0	36	PDB header: hydrolase Chain: A: PDB Molecule: 3c-like proteinase; PDBTitle: main protease of hcov-hku1
9	dlp9sa_	 Alignment		15.0	27	Fold: Trypsin-like serine proteases Superfamily: Trypsin-like serine proteases Family: Viral cysteine protease of trypsin fold
10	d2duca1	 Alignment		14.4	36	Fold: Trypsin-like serine proteases Superfamily: Trypsin-like serine proteases Family: Viral cysteine protease of trypsin fold

11

[dlnrja_](#)

Alignment



8.0

56

Fold: Profilin-like
Superfamily: SNARE-like
Family: SRP alpha N-terminal domain-like