






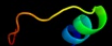

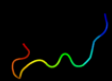










#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3ieeA_	 Alignment		13.3	22	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative exported protein; PDBTitle: crystal structure of hypothetical protein bf3319 from bacteroides2 fragilis (yp_212931.1) from bacteroides fragilis nctc 9343 at 1.70 a3 resolution
2	d2axpa1	 Alignment		8.7	18	Fold: P-loop containing nucleoside triphosphate hydrolases Superfamily: P-loop containing nucleoside triphosphate hydrolases Family: Nucleotide and nucleoside kinases
3	d2f1fa2	 Alignment		8.3	10	Fold: Ferredoxin-like Superfamily: ACT-like Family: IlvH-like
4	d2fgca1	 Alignment		7.5	15	Fold: Ferredoxin-like Superfamily: ACT-like Family: IlvH-like
5	c3m92B_	 Alignment		7.5	15	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: protein ycin; PDBTitle: the structure of ycin, an uncharacterized protein from shigella2 flexneri.
6	d2pc6a1	 Alignment		7.1	15	Fold: Ferredoxin-like Superfamily: ACT-like Family: IlvH-like
7	d1pjva_	 Alignment		7.1	33	Fold: Knottins (small inhibitors, toxins, lectins) Superfamily: Scorpion toxin-like Family: Short-chain scorpion toxins
8	d1pula_	 Alignment		5.7	27	Fold: Hypothetical protein MTH677 Superfamily: Hypothetical protein MTH677 Family: Hypothetical protein MTH677
9	d1zqlc1	 Alignment		5.5	57	Fold: GatB/YqeY motif Superfamily: GatB/YqeY motif Family: GatB/GatE C-terminal domain-like

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[c2fxpA](#)

Alignment



5.4

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PDB header:viral protein
Chain: A: **PDB Molecule:**spike glycoprotein;
PDBTitle: solution structure of the sars-coronavirus hr2 domain