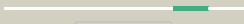
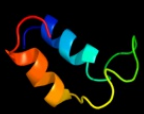
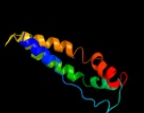
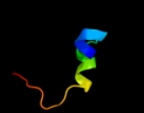
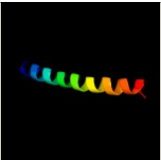


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2nwwa1	 Alignment		100.0	31	Fold: Proton glutamate symport protein Superfamily: Proton glutamate symport protein Family: Proton glutamate symport protein
2	d1a77a1	 Alignment		42.4	18	Fold: SAM domain-like Superfamily: 5' to 3' exonuclease, C-terminal subdomain Family: 5' to 3' exonuclease, C-terminal subdomain
3	d1mc8a1	 Alignment		23.3	18	Fold: SAM domain-like Superfamily: 5' to 3' exonuclease, C-terminal subdomain Family: 5' to 3' exonuclease, C-terminal subdomain
4	d1ul1x1	 Alignment		21.2	15	Fold: SAM domain-like Superfamily: 5' to 3' exonuclease, C-terminal subdomain Family: 5' to 3' exonuclease, C-terminal subdomain
5	d1b43a1	 Alignment		18.4	16	Fold: SAM domain-like Superfamily: 5' to 3' exonuclease, C-terminal subdomain Family: 5' to 3' exonuclease, C-terminal subdomain
6	d1iwga8	 Alignment		16.5	13	Fold: Multidrug efflux transporter AcrB transmembrane domain Superfamily: Multidrug efflux transporter AcrB transmembrane domain Family: Multidrug efflux transporter AcrB transmembrane domain
7	c3ah5E_	 Alignment		15.0	12	PDB header: transferase Chain: E: PDB Molecule: thymidylate synthase thyx; PDBTitle: crystal structure of flavin dependent thymidylate synthase thyx from2 helicobacter pylori complexed with fad and dump
8	d1rxwa1	 Alignment		13.8	19	Fold: SAM domain-like Superfamily: 5' to 3' exonuclease, C-terminal subdomain Family: 5' to 3' exonuclease, C-terminal subdomain
9	d1xo1a1	 Alignment		11.5	21	Fold: SAM domain-like Superfamily: 5' to 3' exonuclease, C-terminal subdomain Family: 5' to 3' exonuclease, C-terminal subdomain
10	d1o26a_	 Alignment		11.3	19	Fold: Thymidylate synthase-complementing protein Thy1 Superfamily: Thymidylate synthase-complementing protein Thy1 Family: Thymidylate synthase-complementing protein Thy1
11	c3aqpB_	 Alignment		10.0	11	PDB header: membrane protein Chain: B: PDB Molecule: probable secdf protein-export membrane protein; PDBTitle: crystal structure of secdf, a translocon-associated membrane protein,2 from thermus thermophilus

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[c3dl8D_](#)

Alignment



9.4

25

PDB header:protein transport
Chain: D: **PDB Molecule:**sece;
PDBTitle: structure of the complex of aquifex aeolicus secyeg and2
bacillus subtilis seca

13 [dlcmwa1](#)

Alignment



7.0

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Fold:SAM domain-like
Superfamily:5' to 3' exonuclease, C-terminal subdomain
Family:5' to 3' exonuclease, C-terminal subdomain