





















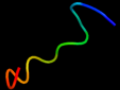




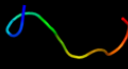


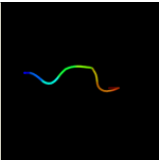
#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d1wila_</a>	 Alignment		19.4	29	<b>Fold:</b> FYVE/PHD zinc finger <b>Superfamily:</b> FYVE/PHD zinc finger <b>Family:</b> variant PHD-like domain
2	<a href="#">d1nekb1</a>	 Alignment		16.3	31	<b>Fold:</b> Globin-like <b>Superfamily:</b> alpha-helical ferredoxin <b>Family:</b> Fumarate reductase/Succinate dehydrogenase iron-sulfur protein, C-terminal domain
3	<a href="#">d1urja_</a>	 Alignment		15.4	24	<b>Fold:</b> Viral ssDNA binding protein <b>Superfamily:</b> Viral ssDNA binding protein <b>Family:</b> Viral ssDNA binding protein
4	<a href="#">d1kf6b1</a>	 Alignment		15.0	31	<b>Fold:</b> Globin-like <b>Superfamily:</b> alpha-helical ferredoxin <b>Family:</b> Fumarate reductase/Succinate dehydrogenase iron-sulfur protein, C-terminal domain
5	<a href="#">d2dlqa2</a>	 Alignment		12.6	44	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
6	<a href="#">d2c42a5</a>	 Alignment		9.6	55	<b>Fold:</b> Ferredoxin-like <b>Superfamily:</b> 4Fe-4S ferredoxins <b>Family:</b> Ferredoxin domains from multidomain proteins
7	<a href="#">c3gitA_</a>	 Alignment		9.2	23	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> carbon monoxide dehydrogenase/acetyl-coa synthase subunit <b>PDBTitle:</b> crystal structure of a truncated acetyl-coa synthase
8	<a href="#">c1v06A_</a>	 Alignment		9.0	8	<b>PDB header:</b> dna-binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> hmg box-containing protein 1; <b>PDBTitle:</b> axh domain of the transcription factor hbp1 from m.musculus
9	<a href="#">c2faoB_</a>	 Alignment		7.1	17	<b>PDB header:</b> hydrolase/transferase <b>Chain:</b> B: <b>PDB Molecule:</b> probable atp-dependent dna ligase; <b>PDBTitle:</b> crystal structure of pseudomonas aeruginosa ligd polymerase2 domain
10	<a href="#">d1xera_</a>	 Alignment		6.9	27	<b>Fold:</b> Ferredoxin-like <b>Superfamily:</b> 4Fe-4S ferredoxins <b>Family:</b> Archaeal ferredoxins
11	<a href="#">c1gx7A_</a>	 Alignment		6.7	36	<b>PDB header:</b> oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> periplasmic [fe] hydrogenase large subunit; <b>PDBTitle:</b> best model of the electron transfer complex between2 cytochrome c3 and [fe]-hydrogenase

12	<a href="#">c2b76N_</a>	Alignment		6.3	25	<b>PDB header:</b> oxidoreductase <b>Chain:</b> N: <b>PDB Molecule:</b> fumarate reductase iron-sulfur protein; <b>PDBTitle:</b> e. coli quinol fumarate reductase frda e49q mutation
13	<a href="#">c2iruA_</a>	Alignment		6.2	10	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> putative dna ligase-like protein rv0938/mt0965; <b>PDBTitle:</b> crystal structure of the polymerase domain from mycobacterium2 tuberculosis ligase d
14	<a href="#">d1jb0c_</a>	Alignment		6.2	27	<b>Fold:</b> Ferredoxin-like <b>Superfamily:</b> 4Fe-4S ferredoxins <b>Family:</b> 7-Fe ferredoxin
15	<a href="#">d1nnqa2</a>	Alignment		5.7	27	<b>Fold:</b> Rubredoxin-like <b>Superfamily:</b> Rubredoxin-like <b>Family:</b> Rubredoxin
16	<a href="#">c2xv7A_</a>	Alignment		5.5	44	<b>PDB header:</b> hormone <b>Chain:</b> A: <b>PDB Molecule:</b> vascular endothelial growth factor d; <b>PDBTitle:</b> crystal structure of vascular endothelial growth factor d
17	<a href="#">d1h98a_</a>	Alignment		5.2	45	<b>Fold:</b> Ferredoxin-like <b>Superfamily:</b> 4Fe-4S ferredoxins <b>Family:</b> 7-Fe ferredoxin

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[c3mjgB\\_](#)

Alignment



5.2

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**PDB header:**hormone/transferase  
**Chain:** B: **PDB Molecule:**platelet-derived growth factor subunit b;  
**PDBTitle:** the structure of a platelet derived growth factor receptor complex