








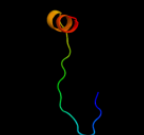



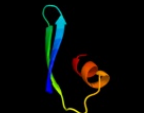




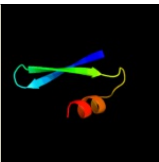
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d1xdpa3</a>	 Alignment		27.5	12	<b>Fold:</b> Phospholipase D/nuclease <b>Superfamily:</b> Phospholipase D/nuclease <b>Family:</b> Polyphosphate kinase C-terminal domain
2	<a href="#">d2p62a1</a>	 Alignment		22.9	18	<b>Fold:</b> PH0156-like <b>Superfamily:</b> PH0156-like <b>Family:</b> PH0156-like
3	<a href="#">c2k6lA_</a>	 Alignment		16.6	46	<b>PDB header:</b> unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> putative uncharacterized protein; <b>PDBTitle:</b> the solution structure of xacb0070 from xanthomonas2 axonopodis pv citri reveals this new protein is a member of3 the rhh family of transcriptional repressors
4	<a href="#">c1oxjA_</a>	 Alignment		15.5	28	<b>PDB header:</b> rna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> rna-binding protein smaug; <b>PDBTitle:</b> crystal structure of the smaug rna binding domain
5	<a href="#">d2o8ra3</a>	 Alignment		10.4	15	<b>Fold:</b> Phospholipase D/nuclease <b>Superfamily:</b> Phospholipase D/nuclease <b>Family:</b> Polyphosphate kinase C-terminal domain
6	<a href="#">c1favC_</a>	 Alignment		6.7	43	<b>PDB header:</b> viral protein <b>Chain:</b> C: <b>PDB Molecule:</b> protein (transmembrane glycoprotein); <b>PDBTitle:</b> the structure of an hiv-1 specific cell entry inhibitor in complex2 with the hiv-1 gp41 trimeric core
7	<a href="#">d1ef1a3</a>	 Alignment		5.4	20	<b>Fold:</b> beta-Grasp (ubiquitin-like) <b>Superfamily:</b> Ubiquitin-like <b>Family:</b> First domain of FERM
8	<a href="#">d1ny722</a>	 Alignment		5.4	33	<b>Fold:</b> Nucleoplasmin-like/VP (viral coat and capsid proteins) <b>Superfamily:</b> Positive stranded ssRNA viruses <b>Family:</b> Comoviridae-like VP

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[d1h4ra3](#)

Alignment



5.1

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**Fold:**beta-Grasp (ubiquitin-like)  
**Superfamily:**Ubiquitin-like  
**Family:**First domain of FERM