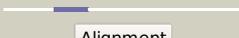
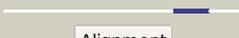
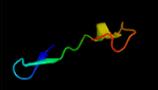
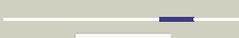
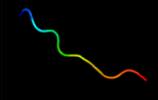


# Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD3424c_(-)_3841894_3842256
Date	Fri Aug 9 18:20:09 BST 2019
Unique Job ID	0f6e18c5ba3b8e1d

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d1u9da_</a>	 Alignment		23.3	27	<b>Fold:</b> Tautomerase/MIF <b>Superfamily:</b> Tautomerase/MIF <b>Family:</b> VC0714-like
2	<a href="#">d2al6a2</a>	 Alignment		13.3	17	<b>Fold:</b> PH domain-like barrel <b>Superfamily:</b> PH domain-like <b>Family:</b> Third domain of FERM
3	<a href="#">c2mx4A_</a>	 Alignment		11.9	75	<b>PDB header:</b> translation,protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> eukaryotic translation initiation factor 4e-binding protein <b>PDBTitle:</b> nmr structure of phosphorylated 4e-bp2
4	<a href="#">c5fceB_</a>	 Alignment		8.6	45	<b>PDB header:</b> cell adhesion <b>Chain:</b> B: <b>PDB Molecule:</b> lpxtg family cell surface protein fms2; <b>PDBTitle:</b> the crystal structure of the ligand binding region of serine-glutamate2 repeat protein a (sgra) of enterococcus faecium
5	<a href="#">c2vycA_</a>	 Alignment		7.9	13	<b>PDB header:</b> lyase <b>Chain:</b> A: <b>PDB Molecule:</b> biodegradative arginine decarboxylase; <b>PDBTitle:</b> crystal structure of acid induced arginine decarboxylase2 from e. coli
6	<a href="#">c2knpA_</a>	 Alignment		7.7	58	<b>PDB header:</b> unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> mcocc-1; <b>PDBTitle:</b> isolation and characterization of peptides from momordica2 cochinchinensis seeds.
7	<a href="#">d1e6vb2</a>	 Alignment		7.6	34	<b>Fold:</b> Ferredoxin-like <b>Superfamily:</b> Methyl-coenzyme M reductase subunits <b>Family:</b> Methyl-coenzyme M reductase alpha and beta chain N-terminal domain
8	<a href="#">c4ldnA_</a>	 Alignment		7.6	69	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> purine nucleoside phosphorylase deod-type; <b>PDBTitle:</b> crystal structure of a putative purine nucleoside phosphorylase from2 vibrio fischeri es114 (target nysgrc-029521)
9	<a href="#">c1z7IC_</a>	 Alignment		6.7	50	<b>PDB header:</b> ligase <b>Chain:</b> C: <b>PDB Molecule:</b> ubiquitin-activating enzyme e1 1; <b>PDBTitle:</b> crystal structure of fragment of mouse ubiquitin-activating2 enzyme

10

[c4lrvL\\_](#)

Alignment



5.9

63

**PDB header:** dna binding protein  
**Chain:** L: **PDB Molecule:** dna sulfur modification protein dnde;  
**PDB Title:** crystal structure of dnde from escherichia coli b7a  
involved in dna2 phosphorothioation modification