


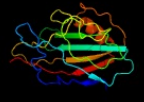











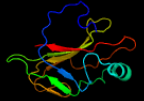

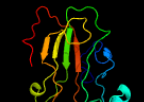




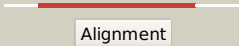

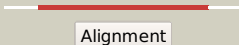

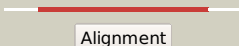
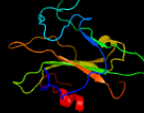

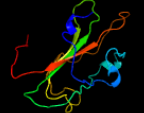
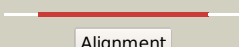

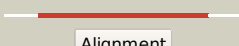

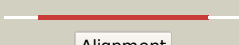



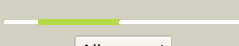
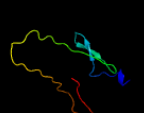



# Phyre2

Email [mdejesus@rockefeller.edu](mailto:mdejesus@rockefeller.edu)  
 Description RVBD2140c\_(TB18.6)\_2399806\_2400336  
 Date Mon Aug 5 13:25:26 BST 2019  
 Unique Job ID 4347753eacdf16e5

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c4begA_</a>	 Alignment		100.0	100	<b>PDB header:</b> lipid binding protein <b>Chain:</b> A; <b>PDB Molecule:</b> phosphatidylethanolamine binding protein; <b>PDBTitle:</b> structure of rv2140c, a phosphatidyl-ethanolamine binding protein from2 mycobacterium tuberculosis in complex with sulphate
2	<a href="#">d1fuxa_</a>	 Alignment		100.0	47	<b>Fold:</b> PEBP-like <b>Superfamily:</b> PEBP-like <b>Family:</b> Prokaryotic PEBP-like proteins
3	<a href="#">d1fjja_</a>	 Alignment		100.0	45	<b>Fold:</b> PEBP-like <b>Superfamily:</b> PEBP-like <b>Family:</b> Prokaryotic PEBP-like proteins
4	<a href="#">c3n08A_</a>	 Alignment		100.0	36	<b>PDB header:</b> phosphatidylethanolamine-binding protein <b>Chain:</b> A; <b>PDB Molecule:</b> putative phosphatidylethanolamine-binding protein (pebp); <b>PDBTitle:</b> crystal structure of a putative phosphatidylethanolamine-binding2 protein (pebp) homolog ct736 from chlamydia trachomatis d/uv-3/cx
5	<a href="#">c2evvD_</a>	 Alignment		100.0	27	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> D; <b>PDB Molecule:</b> hypothetical protein hp0218; <b>PDBTitle:</b> crystal structure of the pebp-like protein of unknown function hp02182 from helicobacter pylori
6	<a href="#">d1wpxb1</a>	 Alignment		99.9	23	<b>Fold:</b> PEBP-like <b>Superfamily:</b> PEBP-like <b>Family:</b> Phosphatidylethanolamine binding protein
7	<a href="#">c2gzaA_</a>	 Alignment		99.9	23	<b>PDB header:</b> lipid binding protein <b>Chain:</b> A; <b>PDB Molecule:</b> phosphatidylethanolamine-binding protein; <b>PDBTitle:</b> phosphatidylethanolamine-binding protein from plasmodium vivax
8	<a href="#">c2r77A_</a>	 Alignment		99.9	18	<b>PDB header:</b> lipid binding protein <b>Chain:</b> A; <b>PDB Molecule:</b> phosphatidylethanolamine-binding protein, putative; <b>PDBTitle:</b> crystal structure of phosphatidylethanolamine-binding protein,2 pfl0955c, from plasmodium falciparum
9	<a href="#">c2jyzA_</a>	 Alignment		99.9	28	<b>PDB header:</b> unknown function <b>Chain:</b> A; <b>PDB Molecule:</b> cg7054-pa; <b>PDBTitle:</b> cg7054 solution structure
10	<a href="#">c1wkpA_</a>	 Alignment		99.8	20	<b>PDB header:</b> signaling protein <b>Chain:</b> A; <b>PDB Molecule:</b> flowering locus t protein; <b>PDBTitle:</b> flowering locus t (ft) from arabidopsis thaliana
11	<a href="#">c1vw41_</a>	 Alignment		99.8	23	<b>PDB header:</b> ribosome <b>Chain:</b> 1; <b>PDB Molecule:</b> 54s ribosomal protein l35, mitochondrial; <b>PDBTitle:</b> structure of the yeast mitochondrial large ribosomal subunit

12	<a href="#">c5tvdA_</a>	 Alignment		99.8	34	<b>PDB header:</b> unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> tm16; <b>PDBTitle:</b> crystal structure of tm16
13	<a href="#">d1kn3a_</a>	 Alignment		99.8	32	<b>Fold:</b> PEBP-like <b>Superfamily:</b> PEBP-like <b>Family:</b> Phosphatidylethanolamine binding protein
14	<a href="#">d1qoua_</a>	 Alignment		99.8	19	<b>Fold:</b> PEBP-like <b>Superfamily:</b> PEBP-like <b>Family:</b> Phosphatidylethanolamine binding protein
15	<a href="#">c4ce4b_</a>	 Alignment		99.8	21	<b>PDB header:</b> ribosome <b>Chain:</b> B: <b>PDB Molecule:</b> unassigned rna; <b>PDBTitle:</b> 39s large subunit of the porcine mitochondrial ribosome
16	<a href="#">d2qyqa1</a>	 Alignment		99.8	29	<b>Fold:</b> PEBP-like <b>Superfamily:</b> PEBP-like <b>Family:</b> Phosphatidylethanolamine binding protein
17	<a href="#">d1a44a_</a>	 Alignment		99.8	26	<b>Fold:</b> PEBP-like <b>Superfamily:</b> PEBP-like <b>Family:</b> Phosphatidylethanolamine binding protein
18	<a href="#">c4v1ab_</a>	 Alignment		99.8	22	<b>PDB header:</b> ribosome <b>Chain:</b> B: <b>PDB Molecule:</b> <b>PDBTitle:</b> structure of the large subunit of the mammalian mitoribosome, part 22 of 2
19	<a href="#">c3j7y6_</a>	 Alignment		99.7	23	<b>PDB header:</b> ribosome <b>Chain:</b> 6: <b>PDB Molecule:</b> ml38; <b>PDBTitle:</b> structure of the large ribosomal subunit from human mitochondria
20	<a href="#">c3ks7D_</a>	 Alignment		63.2	21	<b>PDB header:</b> hydrolase <b>Chain:</b> D: <b>PDB Molecule:</b> putative putative pngase f; <b>PDBTitle:</b> crystal structure of putative peptide:n-glycosidase f (pngase f)2 (yp_210507.1) from bacteroides fragilis nctc 9343 at 2.30 a3 resolution
21	<a href="#">c3kmlB_</a>	 Alignment	not modelled	9.6	31	<b>PDB header:</b> viral protein <b>Chain:</b> B: <b>PDB Molecule:</b> coat protein; <b>PDBTitle:</b> circular permutant of the tobacco mosaic virus