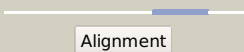
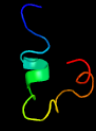
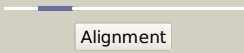

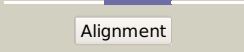



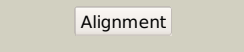

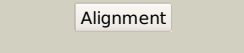

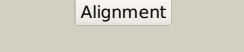





Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD2481c_(-)_2786585_2786908
Date	Wed Aug 7 12:50:11 BST 2019
Unique Job ID	922f901029253405

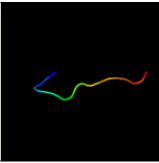
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1pyvA_	 Alignment		21.7	46	PDB header: hydrolase Chain: A: PDB Molecule: atp synthase beta chain, mitochondrial precursor; PDBTitle: nmr solution structure of the mitochondrial f1b presequence2 peptide from nicotiana plumbaginifolia
2	c5xzsA_	 Alignment		16.1	45	PDB header: transcription Chain: A: PDB Molecule: histone acetyltransferase p300; PDBTitle: cryo-em structure of p300
3	c2vldA_	 Alignment		13.0	37	PDB header: hydrolase Chain: A: PDB Molecule: endonuclease nucs; PDBTitle: crystal structure of a repair endonuclease from pyrococcus abyssi
4	c5u7gB_	 Alignment		12.2	45	PDB header: transferase Chain: B: PDB Molecule: creb-binding protein; PDBTitle: crystal structure of the catalytic core of cbp
5	c4bhwB_	 Alignment		11.2	45	PDB header: transferase Chain: B: PDB Molecule: histone acetyltransferase p300; PDBTitle: structural basis for autoinhibition of the acetyltransferase activity2 of p300
6	d2pu9b1	 Alignment		11.0	27	Fold: SH3-like barrel Superfamily: Electron transport accessory proteins Family: Ferredoxin thioredoxin reductase (FTR), alpha (variable) chain
7	c3u5rG_	 Alignment		9.4	21	PDB header: structural genomics, unknown function Chain: G: PDB Molecule: uncharacterized protein; PDBTitle: crystal structure of a hypothetical protein smc02350 from2 sinorhizobium meliloti 1021
8	c5cfdD_	 Alignment		5.6	64	PDB header: virus Chain: D: PDB Molecule: vp4; PDBTitle: crystal structure of dtt treated human coronavirus safv-3

9

[c5cfcD_](#)

Alignment



5.6

64

PDB header:virus
Chain: D: **PDB Molecule:**vp4;
PDBTitle: crystal structure of human coronavirus safv-3