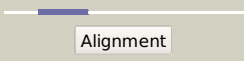
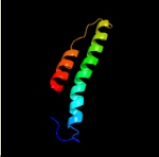
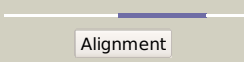
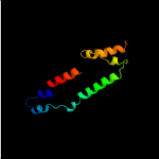
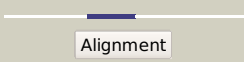

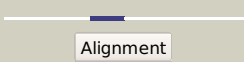
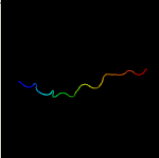


Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD2743c_(-)_3056430_3057242
Date	Wed Aug 7 12:50:40 BST 2019
Unique Job ID	7e850efe1a08ee71

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c6hwhB_	 Alignment		19.0	22	PDB header: electron transport Chain: B: PDB Molecule: ubiquinol-cytochrome c reductase iron-sulfur subunit; PDBTitle: structure of a functional obligate respiratory supercomplex from2 mycobacterium smegmatis
2	c2p7nA_	 Alignment		10.3	20	PDB header: cell invasion Chain: A: PDB Molecule: pathogenicity island 1 effector protein; PDBTitle: crystal structure of the pathogenicity island 1 effector protein from2 chromobacterium violaceum. northeast structural genomics consortium3 (nesgc) target cvr69.
3	c4yo3G_	 Alignment		9.9	25	PDB header: transport protein Chain: G: PDB Molecule: tssa; PDBTitle: enteroaggregative escherichia coli tssa n-terminal fragment
4	c5sxpF_	 Alignment		8.3	32	PDB header: signaling protein/ligase Chain: F: PDB Molecule: e3 ubiquitin-protein ligase itchy homolog; PDBTitle: structural basis for the interaction between itch prr and beta-pix