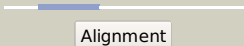

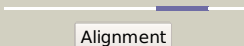

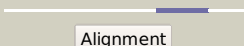
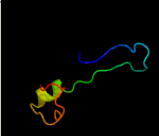
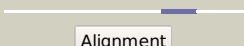
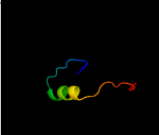
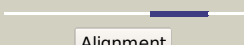
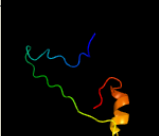


Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD2784c_(ppU)_3092961_3093476
Date	Wed Aug 7 12:50:44 BST 2019
Unique Job ID	2d456e6cd7f9f75f

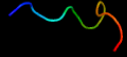
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2m5sA_	 Alignment		24.6	28	PDB header: viral protein Chain: A: PDB Molecule: coat protein; PDBTitle: high-resolution nmr structure and cryo-em imaging support multiple2 functional roles for the accessory i-domain of phage p22 coat protein
2	c3heif_	 Alignment		13.2	16	PDB header: transferase/signaling protein Chain: F: PDB Molecule: ephrin-a1; PDBTitle: ligand recognition by a-class eph receptors: crystal structures of the2 epha2 ligand-binding domain and the epha2/ephrin-a1 complex
3	c3czuB_	 Alignment		11.5	16	PDB header: transferase/signaling protein Chain: B: PDB Molecule: ephrin-a1; PDBTitle: crystal structure of the human ephrin a2- ephrin a1 complex
4	c3g74B_	 Alignment		11.3	18	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: protein of unknown function; PDBTitle: crystal structure of a functionally unknown protein from eubacterium2 ventriosum atcc 27560
5	c2wo3B_	 Alignment		7.8	11	PDB header: transferase/signaling protein Chain: B: PDB Molecule: ephrin-a2; PDBTitle: crystal structure of the epha4-ephrina2 complex

6

[c5zz9E_](#)

Alignment



5.3

8

PDB header:protein binding

Chain: E: **PDB Molecule:**peptide from drebrin;

PDBTitle: crystal structure of homer2 evh1/drebrin ppxxf complex