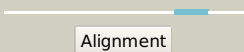

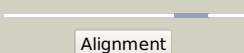

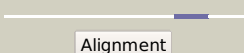


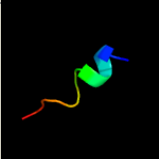
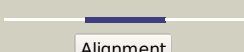
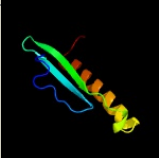
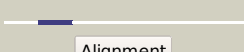



Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD1486c_(-)_1676023_1676889
Date	Fri Aug 2 13:30:07 BST 2019
Unique Job ID	1cda5dfa1919e6c0

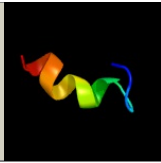
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2n3xA_	 Alignment		30.1	47	PDB header: dna binding protein Chain: A; PDB Molecule: tar dna-binding protein 43; PDBTitle: solution structure of tdp-43 amyloidogenic core region
2	c1wnkA_	 Alignment		24.8	54	PDB header: transcription Chain: A; PDB Molecule: fibroin-modulator-binding-protein-1; PDBTitle: nmr structure of fmbp-1 tandem repeat 3 in 30%(v/v) tfe2 solution
3	c2n2cA_	 Alignment		19.4	47	PDB header: dna binding protein Chain: A; PDB Molecule: tar dna-binding protein 43; PDBTitle: nmr structure of tdp-43 prion-like hydrophobic helix in dpc
4	c1vd9A_	 Alignment		12.6	54	PDB header: transcription Chain: A; PDB Molecule: fibroin-modulator-binding-protein-1; PDBTitle: solution structure of fmbp-1 tandem repeat 3
5	c4guaB_	 Alignment		9.5	21	PDB header: hydrolase Chain: B; PDB Molecule: non-structural polyprotein; PDBTitle: alphavirus p23pro-zbd
6	c3v24V_	 Alignment		8.5	31	PDB header: ribosome Chain: V; PDB Molecule: ribosome modulation factor; PDBTitle: crystal structure of rmf bound to the 70s ribosome. this pdb entry2 contains coordinates for the 30s subunit with bound rmf of the 2nd3 ribosome in the asu

7

[c2jrmA_](#)

Alignment



5.7

25

PDB header: structural genomics, unknown function
Chain: A: **PDB Molecule:** ribosome modulation factor;
PDBTitle: solution nmr structure of ribosome modulation factor
vp1593 from2 vibrio parahaemolyticus. northeast structural
genomics target vpr55