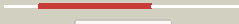













Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD1952_(-) _2200733_2200948
Date	Mon Aug 5 13:25:05 BST 2019
Unique Job ID	7320c5fa0dea40b7

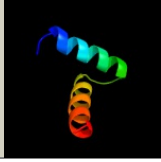
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2h1oH_	 Alignment		98.4	43	PDB header: gene regulation/dna complex Chain: H: PDB Molecule: trafficking protein a; PDBTitle: structure of fitab bound to ir36 dna fragment
2	d2bsqe1	 Alignment		98.2	43	Fold: Ribbon-helix-helix Superfamily: Ribbon-helix-helix Family: Trafficking protein A-like
3	c4p5nB_	 Alignment		60.2	22	PDB header: structural genomics Chain: B: PDB Molecule: hypothetical protein cnag_02591; PDBTitle: structure of cnag_02591 from cryptococcus neoformans
4	c3qoqC_	 Alignment		45.1	40	PDB header: transcription/dna Chain: C: PDB Molecule: alginate and motility regulator z; PDBTitle: crystal structure of the transcription factor amrz in complex with the2 18 base pair amrz1 binding site
5	c4ki0A_	 Alignment		38.5	29	PDB header: calcium binding protein Chain: A: PDB Molecule: putative uncharacterized protein; PDBTitle: crystal structure of the effector protein xoo4466
6	d1mnta_	 Alignment		16.9	19	Fold: Ribbon-helix-helix Superfamily: Ribbon-helix-helix Family: Arc/Mnt-like phage repressors

7

[c1u9pA](#)

Alignment



15.2

20

PDB header: unknown function
Chain: A: **PDB Molecule:** parc;
PDBTitle: permuted single-chain arc