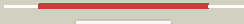









Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD0804 (-) _898084_898713
Date	Fri Jul 26 01:50:39 BST 2019
Unique Job ID	9ba18badb431d0dd

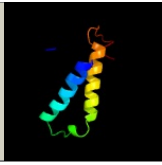
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c4cadC_	 Alignment		100.0	12	PDB header: protein binding Chain: C; PDB Molecule: ras and a-factor converting enzyme 1, rce1; PDBTitle: mechanism of farnesylated caax protein processing by the integral2 membrane protease rce1
2	c4cadF_	 Alignment		99.9	11	PDB header: protein binding Chain: F; PDB Molecule: ras and a-factor converting enzyme 1, rce1; PDBTitle: mechanism of farnesylated caax protein processing by the integral2 membrane protease rce1
3	c5wwoB_	 Alignment		57.0	17	PDB header: rna binding protein Chain: B; PDB Molecule: essential nuclear protein 1; PDBTitle: crystal structure of enp1
4	c6g4ww_	 Alignment		26.4	17	PDB header: ribosome Chain: W; PDB Molecule: 40s ribosomal protein s15a; PDBTitle: cryo-em structure of a late human pre-40s ribosomal subunit - state a

5

[c4z7fD_](#)

Alignment



19.4

18

PDB header:transport protein
Chain: D: **PDB Molecule:**folate ecf transporter;
PDBTitle: crystal structure of folt bound with folic acid