

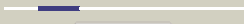


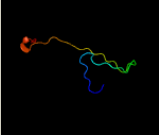

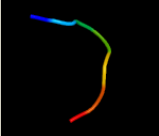


Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD2077A (-)_2334303_2334602
Date	Mon Aug 5 13:25:19 BST 2019
Unique Job ID	4aef478c9eadb574

Detailed template
information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2fpoa1	 Alignment		17.8	24	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: YhhF-like
2	c4wsfA_	 Alignment		6.5	28	PDB header: signaling protein Chain: A: PDB Molecule: serine/threonine-protein phosphatase 4 regulatory subunit PDBTitle: falafel evh1 domain bound to cenp-c fim
3	c5jvkB_	 Alignment		5.8	18	PDB header: hydrolase Chain: B: PDB Molecule: uncharacterized protein; PDBTitle: structural insights into a family 39 glycoside hydrolase from the gut2 symbiont bacteroides cellulosilyticus wh2.
4	c3pg6D_	 Alignment		5.1	29	PDB header: ligase Chain: D: PDB Molecule: e3 ubiquitin-protein ligase dtx3l; PDBTitle: the carboxyl terminal domain of human deltex 3-like