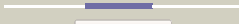

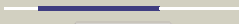

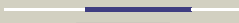
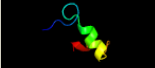




# Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD3202A_(RVBD3202A)_3580283_3580492
Date	Thu Aug 8 16:20:39 BST 2019
Unique Job ID	ab6b86f7cb020668

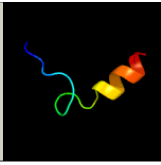
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c2zp2B_</a>	 Alignment		11.7	19	<b>PDB header:</b> transferase inhibitor <b>Chain:</b> B: <b>PDB Molecule:</b> kinase a inhibitor; <b>PDBTitle:</b> c-terminal domain of kipi from bacillus subtilis
2	<a href="#">c2jy8A_</a>	 Alignment		9.4	29	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> ubiquitin-binding protein p62; <b>PDBTitle:</b> nmr structure of the ubiquitin associated (uba) domain of2 p62 (sqstm1) in complex with ubiquitin. rdc refined
3	<a href="#">d2phcb1</a>	 Alignment		9.3	28	<b>Fold:</b> Cyclophilin-like <b>Superfamily:</b> Cyclophilin-like <b>Family:</b> PH0987 C-terminal domain-like
4	<a href="#">d2k0bx1</a>	 Alignment		6.4	29	<b>Fold:</b> RuvA C-terminal domain-like <b>Superfamily:</b> UBA-like <b>Family:</b> UBA domain

5

[c5dudB\\_](#)

Alignment



5.8

24

**PDB header:** unknown function  
**Chain:** B: **PDB Molecule:** ybgj;  
**PDB title:** crystal structure of e. coli ybgjk