
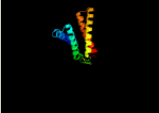
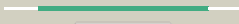




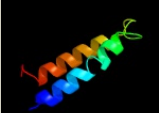




# Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD3789_(- )_4235552_4235917
Date	Fri Aug 9 18:20:49 BST 2019
Unique Job ID	98bbe00f32de0ed8

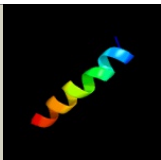
Detailed template  
information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c5mm1A_</a>	 Alignment		99.2	14	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> dolichol monophosphate mannose synthase; <b>PDBTitle:</b> dolichyl phosphate mannose synthase in complex with gdp and dolichyl2 phosphate mannose
2	<a href="#">c6cc4A_</a>	 Alignment		45.3	10	<b>PDB header:</b> transport protein <b>Chain:</b> A: <b>PDB Molecule:</b> soluble cytochrome b562, lipid ii flippase murj chimera; <b>PDBTitle:</b> structure of murj from escherichia coli
3	<a href="#">c4z7fD_</a>	 Alignment		20.3	16	<b>PDB header:</b> transport protein <b>Chain:</b> D: <b>PDB Molecule:</b> folate ecf transporter; <b>PDBTitle:</b> crystal structure of folt bound with folic acid
4	<a href="#">d2rc6gf1</a>	 Alignment		15.2	15	<b>Fold:</b> MalF N-terminal region-like <b>Superfamily:</b> MalF N-terminal region-like <b>Family:</b> MalF N-terminal region-like
5	<a href="#">c4huqS_</a>	 Alignment		14.4	15	<b>PDB header:</b> hydrolase <b>Chain:</b> S: <b>PDB Molecule:</b> uncharacterized protein; <b>PDBTitle:</b> crystal structure of a transporter

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[c4b19A](#)

Alignment



12.0

20

**PDB header:** toxin  
**Chain:** A: **PDB Molecule:** pepa1;  
**PDBTitle:** s. aureus pepa1 nmr structure