
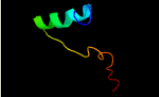








# Phyre2

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Description	RVBD0664 (-) _758535_758807
Date	Fri Jul 26 01:50:23 BST 2019
Unique Job ID	e2564b7b97a3afd5

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d1saza1</a>	 Alignment		52.2	26	<b>Fold:</b> Ribonuclease H-like motif <b>Superfamily:</b> Actin-like ATPase domain <b>Family:</b> Acetokinase-like
2	<a href="#">c3dboA</a>	 Alignment		9.3	56	<b>PDB header:</b> toxin/antitoxin <b>Chain:</b> A: <b>PDB Molecule:</b> uncharacterized protein; <b>PDBTitle:</b> crystal structure of a member of the vapbc family of toxin-antitoxin2 systems, vapbc-5, from mycobacterium tuberculosis
3	<a href="#">c2o98P</a>	 Alignment		8.5	48	<b>PDB header:</b> protein binding <b>Chain:</b> P: <b>PDB Molecule:</b> plasma membrane h+ atpase; <b>PDBTitle:</b> structure of the 14-3-3 / h+-atpase plant complex
4	<a href="#">c1onvB</a>	 Alignment		7.3	79	<b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> serine phosphatase fcp1a; <b>PDBTitle:</b> nmr structure of a complex containing the tfiif subunit2 rap74 and the rnap ii ctd phosphatase fcp1