
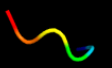









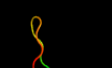

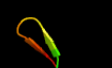


# Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD1046c_(-)_1168640_1169233
Date	Wed Jul 31 22:05:12 BST 2019
Unique Job ID	549190cfb4369bc4

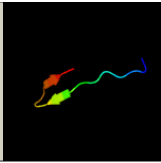
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c5wknD_</a>	 Alignment		17.2	100	<b>PDB header:</b> viral protein <b>Chain:</b> D: <b>PDB Molecule:</b> phosphoprotein; <b>PDBTitle:</b> crystal structure of the parainfluenza virus 5 nucleoprotein-2 phosphoprotein complex
2	<a href="#">c5wknC_</a>	 Alignment		16.6	100	<b>PDB header:</b> viral protein <b>Chain:</b> C: <b>PDB Molecule:</b> phosphoprotein; <b>PDBTitle:</b> crystal structure of the parainfluenza virus 5 nucleoprotein-2 phosphoprotein complex
3	<a href="#">c1qcrl_</a>	 Alignment		10.7	48	<b>PDB header:</b> oxidoreductase <b>Chain:</b> I: <b>PDB Molecule:</b> ubiquinol cytochrome c oxidoreductase; <b>PDBTitle:</b> crystal structure of bovine mitochondrial cytochrome bc12 complex, alpha carbon atoms only
4	<a href="#">c4wlpB_</a>	 Alignment		9.0	40	<b>PDB header:</b> protein binding <b>Chain:</b> B: <b>PDB Molecule:</b> nuclear factor related to kappa-b-binding protein; <b>PDBTitle:</b> crystal structure of uch37-nfrkb inhibited deubiquitylating complex
5	<a href="#">c5ejoA_</a>	 Alignment		7.4	33	<b>PDB header:</b> nuclear protein <b>Chain:</b> A: <b>PDB Molecule:</b> chromatin assembly factor 1 subunit p90; <b>PDBTitle:</b> crystal structure of the winged helix domain in chromatin assembly2 factor 1 subunit p90
6	<a href="#">c3dkbA_</a>	 Alignment		7.3	38	<b>PDB header:</b> hydrolase <b>Chain:</b> A: <b>PDB Molecule:</b> tumor necrosis factor, alpha-induced protein 3; <b>PDBTitle:</b> crystal structure of a20, 2.5 angstrom
7	<a href="#">c5lrvA_</a>	 Alignment		7.2	44	<b>PDB header:</b> hydrolase <b>Chain:</b> A: <b>PDB Molecule:</b> otu domain-containing protein 7b; <b>PDBTitle:</b> structure of cezanne/otud7b otu domain bound to lys11-linked2 diubiquitin

8

[c2vfjA\\_](#)

Alignment



5.7

38

**PDB header:**hydrolase

**Chain:** A: **PDB Molecule:**tumor necrosis factor;

**PDBTitle:** structure of the a20 ovarian tumour (otu) domain