

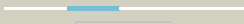











Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	A5A613
Date	Thu Jan 5 10:55:18 GMT 2012
Unique Job ID	2bd7531caeb3a158

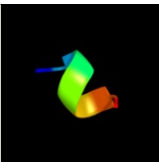
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1etfB_	 Alignment		30.7	64	PDB header: viral protein/rna Chain: B: PDB Molecule: rev peptide; PDBTitle: rev response element (rre) rna complexed with rev peptide,2 nmr, minimized average structure
2	c1etgB_	 Alignment		30.7	64	PDB header: viral protein/rna Chain: B: PDB Molecule: rev peptide; PDBTitle: rev response element (rre) rna complexed with rev peptide,2 nmr, 19 structures
3	d3etja2	 Alignment		7.4	83	Fold: PreATP-grasp domain Superfamily: PreATP-grasp domain Family: BC N-terminal domain-like
4	c2kzkA_	 Alignment		7.2	46	PDB header: protein transport Chain: A: PDB Molecule: uncharacterized protein yol083w; PDBTitle: solution structure of alpha-mannosidase binding domain of atg34
5	c2kzbA_	 Alignment		7.1	46	PDB header: protein transport Chain: A: PDB Molecule: autophagy-related protein 19; PDBTitle: solution structure of alpha-mannosidase binding domain of atg19
6	c2plxB_	 Alignment		7.0	43	PDB header: hydrolase Chain: B: PDB Molecule: peptide inhibitor; PDBTitle: trypsin complexed to a synthetic peptide from veronica hederifolia

7

[d2cmyb1](#)

Alignment



6.8

43

Fold:Toxic hairpin
Superfamily:VhTI-like
Family:VhTI-like