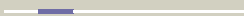
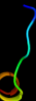


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P36675
Date	Thu Jan 5 11:53:40 GMT 2012
Unique Job ID	27742b53322ac220

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1vf6a_	 Alignment		24.1	36	Fold: L27 domain Superfamily: L27 domain Family: L27 domain
2	c1vf6B_	 Alignment		23.6	36	PDB header: protein binding/protein transport Chain: B: PDB Molecule: pals1-associated tight junction protein; PDBTitle: 2.1 angstrom crystal structure of the pals-1-l27n and patj2 l27 heterodimer complex
3	d1y76a1	 Alignment		16.7	43	Fold: L27 domain Superfamily: L27 domain Family: L27 domain
4	d2pa2a1	 Alignment		10.9	33	Fold: alpha/beta-Hammerhead Superfamily: Ribosomal protein L16p/L10e Family: Ribosomal protein L10e