





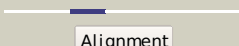
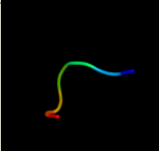


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	Q9JMR4
Date	Thu Jan 5 12:37:48 GMT 2012
Unique Job ID	66e1530f0870e12b

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2l61A_	 Alignment		20.9	67	PDB header: metal binding protein Chain: A: PDB Molecule: ec protein i/ii; PDBTitle: protein and metal cluster structure of the wheat metallothionein2 domain g-ec-1. the second part of the puzzle.
2	c3ushB_	 Alignment		8.1	28	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: uncharacterized protein; PDBTitle: crystal structure of the q2s0r5 protein from salinibacter ruber,2 northeast structural genomics consortium target srr207
3	c2l53B_	 Alignment		8.1	41	PDB header: ca-binding protein/proton transport Chain: B: PDB Molecule: voltage-gated sodium channel type v alpha isoform b PDBTitle: solution nmr structure of apo-calmodulin in complex with the iq motif2 of human cardiac sodium channel nav1.5
4	d1ld3b_	 Alignment		6.5	57	Fold: Histone-fold Superfamily: Histone-fold Family: Nucleosome core histones