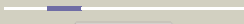


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P45570
Date	Thu Jan 5 12:03:21 GMT 2012
Unique Job ID	6d4426a7872d47a7

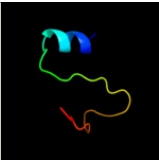
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1mn3a_	 Alignment		16.6	23	Fold: RuvA C-terminal domain-like Superfamily: UBA-like Family: CUE domain
2	d1p3qq_	 Alignment		11.9	33	Fold: RuvA C-terminal domain-like Superfamily: UBA-like Family: CUE domain
3	d1p3qr_	 Alignment		8.8	43	Fold: RuvA C-terminal domain-like Superfamily: UBA-like Family: CUE domain
4	c2jvyA_	 Alignment		8.2	44	PDB header: metal binding protein Chain: A: PDB Molecule: nf-kappa-b essential modulator; PDBTitle: solution structure of the eda-id-related c417f mutant of2 human nemo zinc finger

5

[d2gjva1](#)

Alignment



5.9

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Fold: Phage tail protein-like
Superfamily: Phage tail protein-like
Family: STM4215-like