



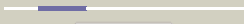

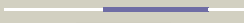
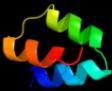




Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	P0AFV8
Date	Thu Jan 5 11:27:23 GMT 2012
Unique Job ID	cc1f351f49928cb7

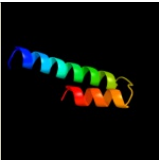
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2b6ca1	 Alignment		27.6	18	Fold: alpha-alpha superhelix Superfamily: ARM repeat Family: BC3264-like
2	c2zshB_	 Alignment		15.0	18	PDB header: hormone receptor Chain: B: PDB Molecule: della protein gai; PDBTitle: structural basis of gibberellin(ga3)-induced della2 recognition by the gibberellin receptor
3	c2ketA_	 Alignment		10.9	29	PDB header: antibiotic Chain: A: PDB Molecule: cathelicidin-6; PDBTitle: solution structure of bmap-27
4	c3bvsA_	 Alignment		10.0	13	PDB header: hydrolase Chain: A: PDB Molecule: alkylpurine dna glycosylase alkd; PDBTitle: crystal structure of bacillus cereus alkylpurine dna glycosylase alkd
5	c2ekgB_	 Alignment		8.1	29	PDB header: oxidoreductase Chain: B: PDB Molecule: proline dehydrogenase/delta-1-pyrroline-5-carboxylate PDBTitle: structure of thermus thermophilus proline dehydrogenase inactivated by2 n-propargylglycine

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[d1rtyb_](#)

Alignment



7.5

15

Fold: Ferritin-like
Superfamily: Cobalamin adenosyltransferase-like
Family: Cobalamin adenosyltransferase