




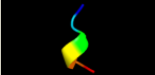






Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P27243
Date	Thu Jan 5 11:43:23 GMT 2012
Unique Job ID	7d4d6de964902b7a

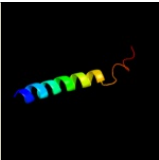
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2d07A_	 Alignment		9.3	15	PDB header: hydrolase Chain: A: PDB Molecule: g/t mismatch-specific thymine dna glycosylase; PDBTitle: crystal structure of sumo-3-modified thymine-dna glycosylase
2	d1muga_	 Alignment		8.9	12	Fold: Uracil-DNA glycosylase-like Superfamily: Uracil-DNA glycosylase-like Family: Mug-like
3	c1a92B_	 Alignment		8.4	17	PDB header: leucine zipper Chain: B: PDB Molecule: delta antigen; PDBTitle: oligomerization domain of hepatitis delta antigen
4	c2rbaB_	 Alignment		7.6	12	PDB header: hydrolase/dna Chain: B: PDB Molecule: g/t mismatch-specific thymine dna glycosylase; PDBTitle: structure of human thymine dna glycosylase bound to abasic and2 undamaged dna
5	d1wraa1	 Alignment		6.8	7	Fold: Metallo-hydrolase/oxidoreductase Superfamily: Metallo-hydrolase/oxidoreductase Family: Pce catalytic domain-like

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[c2kncA](#)

Alignment



6.4

17

PDB header:cell adhesion
Chain: A: **PDB Molecule:**integrin alpha-iiB;
PDBTitle: platelet integrin alphaIIb-beta3 transmembrane-cytoplasmic2 heterocomplex