








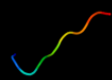


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P0A710
Date	Thu Jan 5 11:04:25 GMT 2012
Unique Job ID	7a2344abde334231

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2a5yb2	 Alignment		11.8	31	Fold: DEATH domain Superfamily: DEATH domain Family: Caspase recruitment domain, CARD
2	c2x11A_	 Alignment		10.0	40	PDB header: translation Chain: A: PDB Molecule: arginine attenuator peptide; PDBTitle: structural basis of translational stalling by human cytomegalovirus2 (hcmv) and fungal arginine attenuator peptide (aap)
3	c2kygC_	 Alignment		9.0	38	PDB header: protein binding Chain: C: PDB Molecule: protein cbfa2t1; PDBTitle: structure of the aml1-eto nervy domain - pka(riia) complex and its2 contribution to aml1-eto activity
4	d1ckma2	 Alignment		9.0	33	Fold: ATP-grasp Superfamily: DNA ligase/mRNA capping enzyme, catalytic domain Family: mRNA capping enzyme
5	d1p16a2	 Alignment		7.7	33	Fold: ATP-grasp Superfamily: DNA ligase/mRNA capping enzyme, catalytic domain Family: mRNA capping enzyme

6

[c3s0xB](#)

Alignment



7.1

14

PDB header:hydrolase
Chain: B: **PDB Molecule:**peptidase a24b, flak domain protein;
PDBTitle: the crystal structure of gxgd membrane protease flak