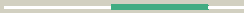







Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	Q9JMR8
Date	Thu Jan 5 12:37:52 GMT 2012
Unique Job ID	08bbab574202cc2a

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3nicA_	 Alignment		40.5	47	PDB header: hydrolase/dna Chain: A: PDB Molecule: eco29kir; PDBTitle: dna binding and cleavage by the giy-yig endonuclease r.eco29ki2 inactive variant y49f
2	dlxcra1	 Alignment		13.8	71	Fold: AF0104/ALDC/Ptd012-like Superfamily: AF0104/ALDC/Ptd012-like Family: PTD012-like
3	dlq1ra3	 Alignment		9.1	20	Fold: CO dehydrogenase flavoprotein C-domain-like Superfamily: FAD/NAD-linked reductases, dimerisation (C-terminal) domain Family: FAD/NAD-linked reductases, dimerisation (C-terminal) domain