Phyre^2

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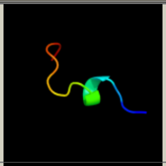
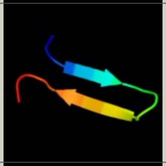
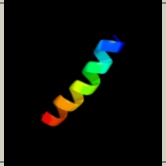
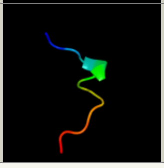
Description	Q47272
Date	Thu Jan 5 12:36:43 GMT 2012
Unique Job ID	41ce00c30883272e

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3b8fB_	Alignment		6.9	21	**PDB header:**hydrolase **Chain:** B: **PDB Molecule:**putative blasticidin s deaminase; **PDBTitle:** crystal structure of the cytidine deaminase from bacillus anthracis
2	d1rq0a_	Alignment		6.1	47	**Fold:**Release factor **Superfamily:**Release factor **Family:**Release factor
3	d1tafa_	Alignment		5.8	32	**Fold:**Histone-fold **Superfamily:**Histone-fold **Family:**TBP-associated factors, TAFs
4	c3eudE_	Alignment		5.6	58	**PDB header:**nuclear protein **Chain:** E: **PDB Molecule:**protein shq1; **PDBTitle:** structure of the cs domain of the essential h/aca rnp2 assembly protein shq1p