

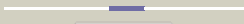
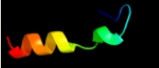



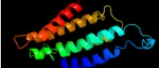


Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	P0ABM1
Date	Thu Jan 5 11:15:49 GMT 2012
Unique Job ID	0d1b4f54c4d48fd1

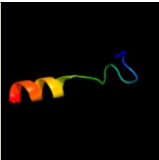
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2oqma1	 Alignment		15.4	16	Fold: DinB/YfiT-like putative metalloenzymes Superfamily: DinB/YfiT-like putative metalloenzymes Family: Sden0562-like
2	d1n7oa1	 Alignment		13.0	27	Fold: alpha/alpha toroid Superfamily: Chondroitin AC/alginate lyase Family: Hyaluronate lyase-like catalytic, N-terminal domain
3	c3qthA_	 Alignment		12.5	11	PDB header: unknown function Chain: A: PDB Molecule: uncharacterized protein; PDBTitle: crystal structure of a hypothetical dinb-like protein (cps_3021) from <i>Colwellia psychrerythraea</i> 34h at 2.20 Å resolution
4	c3mk7K_	 Alignment		9.1	15	PDB header: oxidoreductase Chain: K: PDB Molecule: cytochrome c oxidase, cbb3-type, subunit n; PDBTitle: the structure of cbb3 cytochrome oxidase

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[dlrwha1](#)

Alignment



7.5

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Fold:alpha/alpha toroid
Superfamily:Chondroitin AC/alginate lyase
Family:Hyaluronate lyase-like catalytic, N-terminal domain