

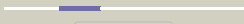


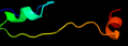

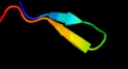

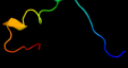

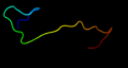




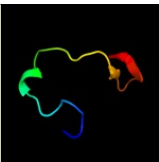
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1y6uA_	 Alignment		50.0	42	PDB header: dna binding protein Chain: A; PDB Molecule: excisionase from transposon tn916; PDBTitle: the structure of the excisionase (xis) protein from 2 conjugative transposon tn916 provides insights into the 3 regulation of heterobivalent tyrosine recombinases
2	c2lcyA_	 Alignment		10.9	33	PDB header: viral protein Chain: A; PDB Molecule: virion spike glycoprotein; PDBTitle: nmr structure of the complete internal fusion loop from ebolavirus gp22 at ph 5.5
3	d2a5yb3	 Alignment		9.1	37	Fold: P-loop containing nucleoside triphosphate hydrolases Superfamily: P-loop containing nucleoside triphosphate hydrolases Family: Extended AAA-ATPase domain
4	c3l51A_	 Alignment		9.0	29	PDB header: cell cycle Chain: A; PDB Molecule: structural maintenance of chromosomes protein 2; PDBTitle: crystal structure of the mouse condensin hinge domain
5	d1f3ud_	 Alignment		7.9	38	Fold: triple barrel Superfamily: Rap30/74 interaction domains Family: Rap30/74 interaction domains
6	c2j8qB_	 Alignment		7.1	23	PDB header: nuclear protein Chain: B; PDB Molecule: cleavage and polyadenylation specificity factor 5; PDBTitle: crystal structure of human cleavage and polyadenylation 2 specificity factor 5 (cpsf5) in complex with a sulphate 3 ion.
7	c3s88j_	 Alignment		6.8	44	PDB header: immune system/viral protein Chain: J; PDB Molecule: envelope glycoprotein; PDBTitle: crystal structure of sudan ebolavirus glycoprotein (strain gulu) bound 2 to 16f6

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[c2w27A_](#)

Alignment



6.2

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PDB header:signaling protein
Chain: A: **PDB Molecule:**ykui protein;
PDBTitle: crystal structure of the bacillus subtilis ykui protein,2
with an eal domain, in complex with substrate c-di-gmp and3
calcium