

















Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2kp6A_	 Alignment		25.6	30	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized protein; PDBTitle: solution nmr structure of protein cv0237 from2 chromobacterium violaceum. northeast structural genomics3 consortium (nesg) target cvt1
2	c2hcnA_	 Alignment		15.8	30	PDB header: transferase Chain: A: PDB Molecule: rna-directed rna polymerase (ns5); PDBTitle: crystal structure of rna dependent rna polymerase domain from west2 nile virus
3	c2j7uA_	 Alignment		14.2	30	PDB header: viral protein Chain: A: PDB Molecule: rna dependent rna polymerase; PDBTitle: dengue virus ns5 rna dependent rna polymerase domain
4	c3lygA_	 Alignment		10.6	33	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: ntf2-like protein of unknown function; PDBTitle: crystal structure of ntf2-like protein of unknown function2 (yp_270605.1) from colwellia psychrerythraea 34h at 1.61 a resolution
5	d1t6la2	 Alignment		9.4	29	Fold: DNA clamp Superfamily: DNA clamp Family: DNA polymerase processivity factor
6	c3o59X_	 Alignment		8.2	17	PDB header: transferase Chain: X: PDB Molecule: dna polymerase ii large subunit; PDBTitle: dna polymerase d large subunit dp2(1-300) from pyrococcus horikoshii
7	d1j5wa_	 Alignment		7.6	23	Fold: Class II aaRS and biotin synthetases Superfamily: Class II aaRS and biotin synthetases Family: Class II aminoacyl-tRNA synthetase (aaRS)-like, catalytic domain
8	c2hfaA_	 Alignment		5.6	30	PDB header: transferase Chain: A: PDB Molecule: rna-directed rna polymerase(ns5); PDBTitle: crystal structure of rna dependent rna polymerase domain2 from west nile virus

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[c3rf1B_](#)

Alignment



5.3

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PDB header:ligase
Chain: B: **PDB Molecule:**glycyl-trna synthetase alpha subunit;
PDBTitle: the crystal structure of glycyl-trna synthetase subunit
alpha from2 campylobacter jejuni subsp. jejuni nctc 11168