




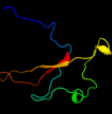

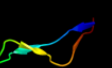












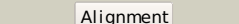







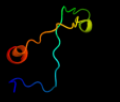


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2i06a1	 Alignment		100.0	100	Fold: Replication terminator protein (Tus) Superfamily: Replication terminator protein (Tus) Family: Replication terminator protein (Tus)
2	d2a1ja1	 Alignment		51.5	20	Fold: SAM domain-like Superfamily: RuvA domain 2-like Family: Hef domain-like
3	d2geca1	 Alignment		33.7	18	Fold: Coronavirus RNA-binding domain Superfamily: Coronavirus RNA-binding domain Family: Coronavirus RNA-binding domain
4	d2bxxa1	 Alignment		21.9	22	Fold: Coronavirus RNA-binding domain Superfamily: Coronavirus RNA-binding domain Family: Coronavirus RNA-binding domain
5	c2d3jA_	 Alignment		18.7	13	PDB header: signaling protein inhibitor Chain: A: PDB Molecule: wnt inhibitory factor-1; PDBTitle: nmr structure of the wif domain from human wif-1
6	c3kx6C_	 Alignment		14.8	22	PDB header: lyase Chain: C: PDB Molecule: fructose-bisphosphate aldolase; PDBTitle: crystal structure of fructose-1,6-bisphosphate aldolase from babesia2 bovis at 2.1a resolution
7	d2aq0a1	 Alignment		14.8	18	Fold: SAM domain-like Superfamily: RuvA domain 2-like Family: Hef domain-like
8	d1vdda_	 Alignment		14.4	16	Fold: Recombination protein RecR Superfamily: Recombination protein RecR Family: Recombination protein RecR
9	c1vddC_	 Alignment		13.1	16	PDB header: recombination Chain: C: PDB Molecule: recombination protein recr; PDBTitle: crystal structure of recombinational repair protein recr
10	c2km6A_	 Alignment		12.5	24	PDB header: signaling protein, protein binding Chain: A: PDB Molecule: nacht, lrr and pyd domains-containing protein 7; PDBTitle: nmr structure of the nlrp7 pyrin domain
11	c1ci6B_	 Alignment		11.4	25	PDB header: transcription Chain: B: PDB Molecule: transcription factor c/ebp beta; PDBTitle: transcription factor atf4-c/ebp beta bzip heterodimer

12	c3qf2B_	Alignment		11.3	19	PDB header: apoptosis Chain: B: PDB Molecule: nacht, Irr and pyd domains-containing protein 3; PDBTitle: crystal structure of nalp3 pyd
13	c2pnvA_	Alignment		10.5	25	PDB header: membrane protein Chain: A: PDB Molecule: small conductance calcium-activated potassium PDBTitle: crystal structure of the leucine zipper domain of small-2 conductance ca2+-activated k+ (skca) channel from rattus3 norvegicus
14	c1s1cY_	Alignment		9.4	38	PDB header: signaling protein Chain: Y: PDB Molecule: rho-associated, coiled-coil containing protein PDBTitle: crystal structure of the complex between the human rhoa and2 rho-binding domain of human rocki
15	d1pn5a1	Alignment		8.7	17	Fold: DEATH domain Superfamily: DEATH domain Family: Pyrin domain, PYD
16	c1pn5A_	Alignment		8.7	17	PDB header: apoptosis Chain: A: PDB Molecule: nacht-, Irr- and pyd-containing protein 2; PDBTitle: nmr structure of the nalp1 pyrin domain (pyd)
17	d1a9xa3	Alignment		8.2	24	Fold: PreATP-grasp domain Superfamily: PreATP-grasp domain Family: BC N-terminal domain-like
18	c2gefA_	Alignment		7.3	17	PDB header: hydrolase Chain: A: PDB Molecule: protease vp4; PDBTitle: crystal structure of a novel viral protease with a2 serine/lysine catalytic dyad mechanism

19 [c2do9A](#)

Alignment



7.2

16

PDB header:signaling protein
Chain: A: **PDB Molecule:**nacht-, Irr- and pyd-containing protein 10;
PDBTitle: solution structure of the pyrin/paad-dapin domain in mouse2 nalp10 (nacht, leucine rich repeat and pyd containing 10)