








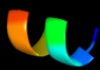




Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1wiga2	 Alignment		21.8	67	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
2	c2kafA	 Alignment		7.7	50	PDB header: viral protein, rna binding protein Chain: A: PDB Molecule: non-structural protein 3; PDBTitle: solution structure of the sars-unique domain-c from the2 nonstructural protein 3 (nsp3) of the severe acute3 respiratory syndrome coronavirus
3	d1vpsa	 Alignment		6.6	42	Fold: Nucleoplasmin-like/VP (viral coat and capsid proteins) Superfamily: Group I dsDNA viruses Family: Papovaviridae-like VP
4	c2rmzA	 Alignment		6.2	33	PDB header: cell adhesion Chain: A: PDB Molecule: integrin beta-3; PDBTitle: bicelle-embedded integrin beta3 transmembrane segment
5	c1nauA	 Alignment		6.0	56	PDB header: hormone/growth factor Chain: A: PDB Molecule: glucagon; PDBTitle: nmr solution structure of the glucagon antagonist [deshis1,2 desphe6, glu9]glucagon amide in the presence of3 perdeuterated dodecylphosphocholine micelles
6	d1x7da	 Alignment		5.8	21	Fold: NAD(P)-binding Rossmann-fold domains Superfamily: NAD(P)-binding Rossmann-fold domains Family: Ornithine cyclodeaminase-like

7

[c3ostA](#)

Alignment



5.5

63

PDB header:lipid binding protein
Chain: A: **PDB Molecule:**serine/threonine-protein kinase kcc4;
PDBTitle: structure of the kinase associated-1 (ka1) from kcc4p