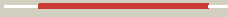





















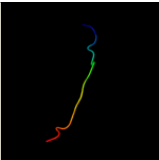


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2kvvA_	 Alignment		100.0	73	PDB header: hydrolase Chain: A: PDB Molecule: putative excisionase; PDBTitle: solution nmr of putative excisionase from klebsiella pneumoniae,2 northeast structural genomics consortium target kpr49
2	d2foka4	 Alignment		14.0	38	Fold: Restriction endonuclease-like Superfamily: Restriction endonuclease-like Family: Restriction endonuclease FokI, C-terminal (catalytic) domain
3	c3pc3A_	 Alignment		13.0	11	PDB header: lyase Chain: A: PDB Molecule: cg1753, isoform a; PDBTitle: full length structure of cystathionine beta-synthase from drosophila2 in complex with aminoacrylate
4	d1ffkf_	 Alignment		9.3	29	Fold: alpha/beta-Hammerhead Superfamily: Ribosomal protein L16p/L10e Family: Ribosomal protein L10e
5	d1efea_	 Alignment		9.1	32	Fold: Insulin-like Superfamily: Insulin-like Family: Insulin-like
6	d2ih2a2	 Alignment		8.6	21	Fold: DNA methylase specificity domain Superfamily: DNA methylase specificity domain Family: TaqI C-terminal domain-like
7	c1hp9A_	 Alignment		8.5	60	PDB header: toxin Chain: A: PDB Molecule: kappa-hefutoxin 1; PDBTitle: kappa-hefutoxins: a novel class of potassium channel toxins2 from scorpion venom
8	d2o2aa1	 Alignment		6.6	20	Fold: SecB-like Superfamily: SecB-like Family: SP1558-like
9	d1liua3	 Alignment		6.2	29	Fold: Pyruvate kinase C-terminal domain-like Superfamily: PK C-terminal domain-like Family: Pyruvate kinase, C-terminal domain
10	c2lk2A_	 Alignment		5.8	23	PDB header: transcription Chain: A: PDB Molecule: homeobox protein tgif1; PDBTitle: solution nmr structure of homeobox domain (171-248) of human homeobox2 protein tgif1, northeast structural genomics consortium target3 hr4411b
11	d1a3xa3	 Alignment		5.7	29	Fold: Pyruvate kinase C-terminal domain-like Superfamily: PK C-terminal domain-like Family: Pyruvate kinase, C-terminal domain

12 [d2hnga1](#)

Alignment



5.6

25

Fold:SecB-like
Superfamily:SecB-like
Family:SP1558-like