

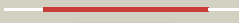






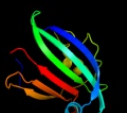


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1tt8a_	 Alignment		100.0	99	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: Chorismate lyase
2	d2nwia1	 Alignment		100.0	21	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: AF1396-like
3	c3ft8lC_	 Alignment		98.4	16	PDB header: transcription Chain: C: PDB Molecule: hth-type transcriptional repressor phnf; PDBTitle: crystal structure of the effector domain of phnf from mycobacterium2 smegmatis
4	c3hfiA_	 Alignment		97.8	12	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative regulator; PDBTitle: the crystal structure of the putative regulator from escherichia coli2 cft073
5	d3cnva1	 Alignment		97.8	11	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
6	d2pkha1	 Alignment		97.6	12	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
7	d2fa1a1	 Alignment		97.6	13	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
8	d2p19a1	 Alignment		97.4	17	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
9	d2ikka1	 Alignment		97.4	12	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
10	d2ogga1	 Alignment		97.2	15	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
11	d2ra5a1	 Alignment		97.0	18	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain

12	c2ra5A_	Alignment		97.0	18	PDB header: transcription Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of the putative transcriptional regulator2 from streptomyces coelicolor
13	d3bwga2	Alignment		96.9	8	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
14	d2ooia1	Alignment		96.9	11	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
15	d3ddva1	Alignment		96.3	19	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
16	c3lhea_	Alignment		95.1	21	PDB header: transcription regulator Chain: A: PDB Molecule: gntr family transcriptional regulator; PDBTitle: the crystal structure of the c-terminal domain of a gntr2 family transcriptional regulator from bacillus anthracis3 str. sterne
17	c3f8mA_	Alignment		84.3	16	PDB header: transcription Chain: A: PDB Molecule: gntr-family protein transcriptional regulator; PDBTitle: crystal structure of phnf from mycobacterium smegmatis
18	c3eetA_	Alignment		80.9	11	PDB header: transcription regulator Chain: A: PDB Molecule: putative gntr-family transcriptional regulator; PDBTitle: crystal structure of putative gntr-family transcriptional2 regulator
19	c3edpB_	Alignment		57.7	9	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: lin2111 protein; PDBTitle: the crystal structure of the protein lin2111 (functionally unknown)2 from listeria innocua clip11262
20	c2wv0H_	Alignment		35.7	10	PDB header: transcription Chain: H: PDB Molecule: hth-type transcriptional repressor yvoa; PDBTitle: crystal structure of the gntr-hutc family member yvoa from2 bacillus subtilis
21	c3bwgA_	Alignment	not modelled	10.8	8	PDB header: transcription regulator Chain: A: PDB Molecule: uncharacterized hth-type transcriptional regulator yydk; PDBTitle: the crystal structure of possible transcriptional regulator yydk from2 bacillus subtilis subsp. subtilis str. 168
22	d1dsva_	Alignment	not modelled	9.1	25	Fold: Retrovirus zinc finger-like domains Superfamily: Retrovirus zinc finger-like domains Family: Retrovirus zinc finger-like domains