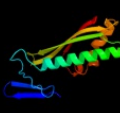


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1ml8a_	 Alignment		100.0	99	Fold: OsmC-like Superfamily: OsmC-like Family: YhfA-like
2	d1vlaa_	 Alignment		100.0	36	Fold: OsmC-like Superfamily: OsmC-like Family: YhfA-like
3	c2egtA_	 Alignment		100.0	36	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hypothetical protein aq_1549; PDBTitle: crystal structure of hypothetical protein (aq1549) from aquifex2 aeolicus
4	d2onfa1	 Alignment		100.0	19	Fold: OsmC-like Superfamily: OsmC-like Family: Ohr/OsmC resistance proteins
5	d2pn2a1	 Alignment		99.9	15	Fold: OsmC-like Superfamily: OsmC-like Family: Ohr/OsmC resistance proteins
6	d1qwia_	 Alignment		99.9	14	Fold: OsmC-like Superfamily: OsmC-like Family: Ohr/OsmC resistance proteins
7	c1lqlE_	 Alignment		99.9	16	PDB header: unknown function Chain: E: PDB Molecule: osmotic inducible protein c like family; PDBTitle: crystal structure of osmC like protein from mycoplasma2 pneumoniae
8	d1lqla_	 Alignment		99.9	16	Fold: OsmC-like Superfamily: OsmC-like Family: Ohr/OsmC resistance proteins
9	c3eerA_	 Alignment		99.9	11	PDB header: oxidoreductase Chain: A: PDB Molecule: organic hydroperoxide resistance protein, putative; PDBTitle: high resolution structure of putative organic hydroperoxide resistance2 protein from vibrio cholerae o1 biovar eltor str. n16961
10	d2opla1	 Alignment		99.9	22	Fold: OsmC-like Superfamily: OsmC-like Family: Ohr/OsmC resistance proteins
11	c2bjoa_	 Alignment		99.9	13	PDB header: oxidoreductase Chain: A: PDB Molecule: organic hydroperoxide resistance protein ohrb; PDBTitle: crystal structure of the organic hydroperoxide resistance2 protein ohrb of bacillus subtilis

12	dlukka_	Alignment		99.9	16	Fold: OsmC-like Superfamily: OsmC-like Family: Ohr/OsmC resistance proteins
13	dlnyed_	Alignment		99.9	15	Fold: OsmC-like Superfamily: OsmC-like Family: Ohr/OsmC resistance proteins
14	dln2fa_	Alignment		99.9	12	Fold: OsmC-like Superfamily: OsmC-like Family: Ohr/OsmC resistance proteins
15	c2ql8A_	Alignment		99.9	10	PDB header: oxidoreductase Chain: A: PDB Molecule: putative redox protein; PDBTitle: crystal structure of a putative redox protein (Isei_0423) from2 lactobacillus casei atcc 334 at 1.50 a resolution
16	dluspa_	Alignment		99.9	10	Fold: OsmC-like Superfamily: OsmC-like Family: Ohr/OsmC resistance proteins
17	c3cjeA_	Alignment		99.9	16	PDB header: oxidoreductase Chain: A: PDB Molecule: osmc-like protein; PDBTitle: crystal structure of an osmc-like hydroperoxide resistance protein2 (jann_2040) from jannaschia sp. ccs1 at 1.70 a resolution
18	d2d7va1	Alignment		99.9	20	Fold: OsmC-like Superfamily: OsmC-like Family: Ohr/OsmC resistance proteins
19	c1zb8B_	Alignment		99.9	11	PDB header: oxidoreductase Chain: B: PDB Molecule: organic hydroperoxide resistance protein; PDBTitle: crystal structure of xylella fastidiosa organic peroxide resistance2 protein
20	dljma_	Alignment		26.5	28	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: p53-like transcription factors Family: RUNT domain
21	dleaga_	Alignment	not modelled	25.7	24	Fold: Common fold of diphtheria toxin/transcription factors/cytochrome f Superfamily: p53-like transcription factors Family: RUNT domain
22	d1xo3a_	Alignment	not modelled	21.8	13	Fold: beta-Grasp (ubiquitin-like) Superfamily: MoaD/ThiS Family: C9orf74 homolog
23	c2qj1A_	Alignment	not modelled	16.6	22	PDB header: signaling protein Chain: A: PDB Molecule: ubiquitin-related modifier 1; PDBTitle: crystal structure of urm1
24	d2qgra2	Alignment	not modelled	10.9	22	Fold: SH3-like barrel Superfamily: Tudor/PWWP/MBT Family: Tudor domain
25	d1wgka_	Alignment	not modelled	10.3	13	Fold: beta-Grasp (ubiquitin-like) Superfamily: MoaD/ThiS Family: C9orf74 homolog
26	d2qqa2	Alignment	not modelled	10.3	26	Fold: SH3-like barrel Superfamily: Tudor/PWWP/MBT Family: Tudor domain
27	c3kl7A_	Alignment	not modelled	7.9	12	PDB header: hydrolase Chain: A: PDB Molecule: putative metal-dependent hydrolase; PDBTitle: crystal structure of putative metal-dependent hydrolase2 (yp_001302908.1) from parabacteroides distasonis atcc 8503 at 2.30 a3 resolution
28	d1v5ta_	Alignment	not modelled	5.9	13	Fold: beta-Grasp (ubiquitin-like) Superfamily: Ubiquitin-like Family: Ubiquitin-related