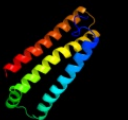
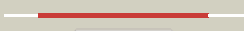
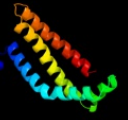
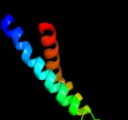
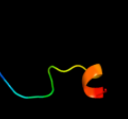
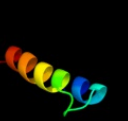
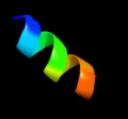
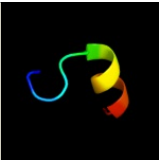


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1nekd_	 Alignment		100.0	100	Fold: Heme-binding four-helical bundle Superfamily: Fumarate reductase respiratory complex transmembrane subunits Family: Succinate dehydrogenase/Fumarate reductase transmembrane subunits (SdhC/FrdC and SdhD/FrdD)
2	d2bs2c1	 Alignment		99.3	11	Fold: Heme-binding four-helical bundle Superfamily: Fumarate reductase respiratory complex transmembrane subunits Family: Fumarate reductase respiratory complex cytochrome b subunit, FrdC
3	d1nekc_	 Alignment		99.0	17	Fold: Heme-binding four-helical bundle Superfamily: Fumarate reductase respiratory complex transmembrane subunits Family: Succinate dehydrogenase/Fumarate reductase transmembrane subunits (SdhC/FrdC and SdhD/FrdD)
4	c1zp0D_	 Alignment		99.0	14	PDB header: oxidoreductase Chain: D: PDB Molecule: small cytochrome binding protein; PDBTitle: crystal structure of mitochondrial respiratory complex ii2 bound with 3-nitropropionate and 2-thenoyltrifluoroacetone
5	c1yq3C_	 Alignment		98.5	16	PDB header: oxidoreductase Chain: C: PDB Molecule: succinate dehydrogenase cytochrome b, large subunit; PDBTitle: avian respiratory complex ii with oxaloacetate and ubiquinone
6	d1kf6c_	 Alignment		96.4	17	Fold: Heme-binding four-helical bundle Superfamily: Fumarate reductase respiratory complex transmembrane subunits Family: Succinate dehydrogenase/Fumarate reductase transmembrane subunits (SdhC/FrdC and SdhD/FrdD)
7	d1kf6d_	 Alignment		95.6	20	Fold: Heme-binding four-helical bundle Superfamily: Fumarate reductase respiratory complex transmembrane subunits Family: Succinate dehydrogenase/Fumarate reductase transmembrane subunits (SdhC/FrdC and SdhD/FrdD)
8	d1kcfa1	 Alignment		10.6	33	Fold: LEM/SAP HeH motif Superfamily: SAP domain Family: SAP domain
9	c2do5A_	 Alignment		10.4	18	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: splicing factor 3b subunit 2; PDBTitle: solution structure of the sap domain of human splicing2 factor 3b subunit 2
10	c3da4B_	 Alignment		10.2	30	PDB header: antibiotic Chain: B: PDB Molecule: colicin-m; PDBTitle: crystal structure of colicin m, a novel phosphatase2 specifically imported by escherichia coli
11	c2gzdC_	 Alignment		8.1	38	PDB header: protein transport Chain: C: PDB Molecule: rab11 family-interacting protein 2; PDBTitle: crystal structure of rab11 in complex with rab11-fip2

12	dly8oa1	Alignment		6.8	5	Fold: Bromodomain-like Superfamily: alpha-ketoacid dehydrogenase kinase, N-terminal domain Family: alpha-ketoacid dehydrogenase kinase, N-terminal domain
13	c3mvnA_	Alignment		6.3	22	PDB header: ligase Chain: A: PDB Molecule: udp-n-acetylmuramate:l-alanyl-gamma-d-glutamyl-medo- PDBTitle: crystal structure of a domain from a putative udp-n-acetylmuramate:l-2 alanyl-gamma-d-glutamyl-medo-diaminopimelate ligase from haemophilus3 ducreyi 35000hp

14 [d2eyqa6](#)

Alignment



5.4

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Fold:TRCF domain-like
Superfamily:TRCF domain-like
Family:TRCF domain