



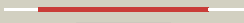








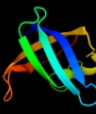










#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1h95a_	 Alignment		99.9	37	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
2	d1mjca_	 Alignment		99.9	47	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
3	c3camB_	 Alignment		99.9	38	PDB header: gene regulation Chain: B: PDB Molecule: cold-shock domain family protein; PDBTitle: crystal structure of the cold shock domain protein from neisseria2 meningitidis
4	d2es2a1	 Alignment		99.9	41	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
5	c2kcmA_	 Alignment		99.9	28	PDB header: nucleic acid binding protein Chain: A: PDB Molecule: cold shock domain family protein; PDBTitle: solution nmr structure of the n-terminal ob-domain of so_1732 from2 shewanella oneidensis. northeast structural genomics consortium3 target sor210a.
6	c3a0jB_	 Alignment		99.9	35	PDB header: transcription Chain: B: PDB Molecule: cold shock protein; PDBTitle: crystal structure of cold shock protein 1 from thermus2 thermophilus hb8
7	c2k5nA_	 Alignment		99.9	22	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative cold-shock protein; PDBTitle: solution nmr structure of the n-terminal domain of protein2 eca1580 from erwinia carotovora, northeast structural3 genomics consortium target ewr156a
8	d1g6pa_	 Alignment		99.9	44	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
9	d1c9oa_	 Alignment		99.9	41	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
10	c3aqqD_	 Alignment		99.8	29	PDB header: dna binding protein Chain: D: PDB Molecule: calcium-regulated heat stable protein 1; PDBTitle: crystal structure of human crhsp-24
11	c3trzE_	 Alignment		99.8	29	PDB header: rna binding protein/rna Chain: E: PDB Molecule: protein lin-28 homolog a; PDBTitle: mouse lin28a in complex with let-7d microRNA pre-element

12	c2ytyA_	Alignment		99.7	30	PDB header: rna binding protein Chain: A: PDB Molecule: cold shock domain-containing protein e1; PDBTitle: solution structure of the fourth cold-shock domain of the human2 kiaa0885 protein (unr protein)
13	d1wfga_	Alignment		99.7	25	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
14	c2ytxA_	Alignment		99.7	35	PDB header: rna binding protein Chain: A: PDB Molecule: cold shock domain-containing protein e1; PDBTitle: solution structure of the second cold-shock domain of the human2 kiaa0885 protein (unr protein)
15	c1x65A_	Alignment		99.6	25	PDB header: rna binding protein Chain: A: PDB Molecule: unr protein; PDBTitle: solution structure of the third cold-shock domain of the human2 kiaa0885 protein (unr protein)
16	c2bh8B_	Alignment		99.6	46	PDB header: transcription Chain: B: PDB Molecule: 1b11; PDBTitle: combinatorial protein 1b11
17	c2yvtA_	Alignment		99.4	30	PDB header: rna binding protein Chain: A: PDB Molecule: cold shock domain-containing protein e1; PDBTitle: solution structure of the fifth cold-shock domain of the human2 kiaa0885 protein (unr protein)
18	d2ix0a2	Alignment		97.5	16	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
19	c2ix1A_	Alignment		96.9	20	PDB header: hydrolase Chain: A: PDB Molecule: exoribonuclease 2; PDBTitle: rnase ii d209n mutant
20	d1a62a2	Alignment		95.6	17	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
21	c2a8vA_	Alignment	not modelled	95.6	20	PDB header: protein/rna Chain: A: PDB Molecule: rna binding domain of rho transcription PDBTitle: rho transcription termination factor/rna complex
22	d1smxa_	Alignment	not modelled	94.9	14	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
23	d2ix0a1	Alignment	not modelled	92.2	16	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
24	d1hh2p1	Alignment	not modelled	91.3	18	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
25	c1hh2P_	Alignment	not modelled	91.0	18	PDB header: transcription regulation Chain: P: PDB Molecule: n utilization substance protein a; PDBTitle: crystal structure of nusa from thermotoga maritima
26	c3go5A_	Alignment	not modelled	90.1	14	PDB header: gene regulation Chain: A: PDB Molecule: multidomain protein with s1 rna-binding domains; PDBTitle: crystal structure of a multidomain protein with nucleic acid binding2 domains (sp_0946) from streptococcus pneumoniae tigr4 at 1.40 a3 resolution
27	c2vnuD_	Alignment	not modelled	89.6	21	PDB header: hydrolase/rna Chain: D: PDB Molecule: exosome complex exonuclease rrp44; PDBTitle: crystal structure of sc rrp44
28	c2k52A_	Alignment	not modelled	89.0	13	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized protein mj1198; PDBTitle: structure of uncharacterized protein mj1198 from2 methanocaldococcus jannaschii. northeast structural3 genomics

					target mjr117b
29	c2cqoA_	Alignment	not modelled	88.8	22 PDB header: ribosome Chain: A: PDB Molecule: nucleolar protein of 40 kda; PDBTitle: solution structure of the s1 rna binding domain of human2 hypothetical protein flj11067
30	c1l2fA_	Alignment	not modelled	88.3	18 PDB header: transcription Chain: A: PDB Molecule: n utilization substance protein a; PDBTitle: crystal structure of nusa from thermotoga maritima: a2 structure-based role of the n-terminal domain
31	d1u0la1	Alignment	not modelled	88.2	16 Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
32	d1kl9a2	Alignment	not modelled	82.0	13 Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
33	c1xpuB_	Alignment	not modelled	79.9	20 PDB header: transcription/rna Chain: B: PDB Molecule: rho transcription termination factor; PDBTitle: structural mechanism of inhibition of the rho transcription2 termination factor by the antibiotic 5a-(3-formylphenylsulfanyl)-3 dihydrobicyclomycin (fpdb)
34	c1u0lB_	Alignment	not modelled	79.7	16 PDB header: hydrolase Chain: B: PDB Molecule: probable gtpase engc; PDBTitle: crystal structure of yjeq from thermotoga maritima
35	c1zeqX_	Alignment	not modelled	78.4	21 PDB header: metal binding protein Chain: X: PDB Molecule: cation efflux system protein cusf; PDBTitle: 1.5 a structure of apo-cusf residues 6-88 from escherichia2 coli
36	c2c4rL_	Alignment	not modelled	69.4	12 PDB header: hydrolase Chain: L: PDB Molecule: ribonuclease e; PDBTitle: catalytic domain of e. coli rnase e
37	d1y14b1	Alignment	not modelled	68.4	18 Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
38	c2l55A_	Alignment	not modelled	67.1	26 PDB header: metal binding protein Chain: A: PDB Molecule: silb,silver efflux protein, mfp component of the three PDBTitle: solution structure of the c-terminal domain of silb from cupriavidus2 metallidurans
39	d2nn6h1	Alignment	not modelled	66.9	14 Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
40	d2c35b1	Alignment	not modelled	66.7	18 Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
41	d2z0sa1	Alignment	not modelled	64.6	14 Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
42	c2wp8j_	Alignment	not modelled	63.3	15 PDB header: hydrolase Chain: J: PDB Molecule: exosome complex exonuclease dis3; PDBTitle: yeast rrp44 nuclease
43	d2ba0a1	Alignment	not modelled	58.3	13 Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
44	c2oceA_	Alignment	not modelled	57.0	21 PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hypothetical protein pa5201; PDBTitle: crystal structure of tex family protein pa5201 from2 pseudomonas aeruginosa
45	c3l0oB_	Alignment	not modelled	55.7	17 PDB header: hydrolase Chain: B: PDB Molecule: transcription termination factor rho; PDBTitle: structure of rna-free rho transcription termination factor from2 thermotoga maritima
46	d1wi5a_	Alignment	not modelled	54.9	9 Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
47	d3bzka4	Alignment	not modelled	53.8	20 Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
48	c2khiA_	Alignment	not modelled	48.8	25 PDB header: ribosomal protein Chain: A: PDB Molecule: 30s ribosomal protein s1; PDBTitle: nmr structure of the domain 4 of the e. coli ribosomal2 protein s1
49	c1j6qa_	Alignment	not modelled	47.5	19 PDB header: chaperone Chain: A: PDB Molecule: cytochrome c maturation protein e; PDBTitle: solution structure and characterization of the heme2 chaperone ccme
50	d1j6qa_	Alignment	not modelled	47.5	19 Fold: OB-fold Superfamily: Heme chaperone CcmE Family: Heme chaperone CcmE
51	c2k4kA_	Alignment	not modelled	41.8	25 PDB header: rna binding protein Chain: A: PDB Molecule: general stress protein 13; PDBTitle: solution structure of gsp13 from bacillus subtilis
52	d2asba1	Alignment	not modelled	41.8	20 Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
53	c2khjA_	Alignment	not modelled	39.0	25 PDB header: ribosomal protein Chain: A: PDB Molecule: 30s ribosomal protein s1; PDBTitle: nmr structure of the domain 6 of the e. coli ribosomal2 protein s1
54	d1t9ha1	Alignment	not modelled	37.6	18 Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like

55	d2je6i1	Alignment	not modelled	36.2	17	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
56	c2asbA	Alignment	not modelled	34.9	20	PDB header: transcription/rna Chain: A: PDB Molecule: transcription elongation protein nusa; PDBTitle: structure of a mycobacterium tuberculosis nusa-rna complex
57	c2z0sA	Alignment	not modelled	32.1	14	PDB header: rna binding protein Chain: A: PDB Molecule: probable exosome complex rna-binding protein 1; PDBTitle: crystal structure of putative exosome complex rna-binding2 protein
58	c2xngA	Alignment	not modelled	31.7	31	PDB header: rna binding protein Chain: A: PDB Molecule: nuclear polyadenylated rna-binding protein 3; PDBTitle: structural insights into cis element recognition of non-2 polyadenylated rnas by the nab3-rrm
59	c2eqsA	Alignment	not modelled	28.2	25	PDB header: hydrolase Chain: A: PDB Molecule: atp-dependent rna helicase dhx8; PDBTitle: solution structure of the s1 rna binding domain of human2 atp-dependent rna helicase dhx8
60	d1ueqa	Alignment	not modelled	27.9	16	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
61	d1q46a2	Alignment	not modelled	26.8	11	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
62	d1h9ma2	Alignment	not modelled	25.7	19	Fold: OB-fold Superfamily: MOP-like Family: BiMOP, duplicated molybdate-binding domain
63	c2b8kG	Alignment	not modelled	25.5	16	PDB header: transferase Chain: G: PDB Molecule: dna-directed rna polymerase ii 19 kda PDBTitle: 12-subunit rna polymerase ii
64	c2dc2A	Alignment	not modelled	24.4	23	PDB header: structural protein Chain: A: PDB Molecule: golgi associated pdz and coiled-coil motif PDBTitle: solution structure of pdz domain
65	c2opgB	Alignment	not modelled	24.1	18	PDB header: structural protein Chain: B: PDB Molecule: multiple pdz domain protein; PDBTitle: the crystal structure of the 10th pdz domain of mpdz
66	c3h0gS	Alignment	not modelled	23.4	29	PDB header: transcription Chain: S: PDB Molecule: dna-directed rna polymerase ii subunit rpb7; PDBTitle: rna polymerase ii from schizosaccharomyces pombe
67	c2gu1A	Alignment	not modelled	23.4	16	PDB header: hydrolase Chain: A: PDB Molecule: zinc peptidase; PDBTitle: crystal structure of a zinc containing peptidase from2 vibrio cholerae
68	c2je6l	Alignment	not modelled	22.6	20	PDB header: hydrolase Chain: I: PDB Molecule: exosome complex rna-binding protein 1; PDBTitle: structure of a 9-subunit archaeal exosome
69	c1q46A	Alignment	not modelled	20.9	9	PDB header: translation Chain: A: PDB Molecule: translation initiation factor 2 alpha subunit; PDBTitle: crystal structure of the eif2 alpha subunit from2 saccharomyces cerevisia
70	d1rgwa	Alignment	not modelled	19.8	27	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
71	c2dhxA	Alignment	not modelled	19.6	21	PDB header: rna binding protein Chain: A: PDB Molecule: poly (adp-ribose) polymerase family, member 10 PDBTitle: solution structure of the rrm domain in the human poly (adp-2 ribose) polymerase family, member 10 variant
72	c2iwnA	Alignment	not modelled	19.3	16	PDB header: signaling protein Chain: A: PDB Molecule: multiple pdz domain protein; PDBTitle: 3rd pdz domain of multiple pdz domain protein mpdz (casp2 target)
73	d2bs2b2	Alignment	not modelled	19.1	33	Fold: beta-Grasp (ubiquitin-like) Superfamily: 2Fe-2S ferredoxin-like Family: 2Fe-2S ferredoxin domains from multidomain proteins
74	c2c35F	Alignment	not modelled	18.8	14	PDB header: polymerase Chain: F: PDB Molecule: dna-directed rna polymerase ii 19 kda PDBTitle: subunits rpb4 and rpb7 of human rna polymerase ii
75	d1n7ea	Alignment	not modelled	18.3	18	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
76	c2i04B	Alignment	not modelled	18.0	18	PDB header: peptide binding protein Chain: B: PDB Molecule: membrane-associated guanylate kinase, ww and pdz PDBTitle: x-ray crystal structure of magi-1 pdz1 bound to the c-2 terminal peptide of hpv18 e6
77	c1nt9G	Alignment	not modelled	17.9	16	PDB header: transcription, transferase Chain: G: PDB Molecule: dna-directed rna polymerase ii 19 kd polypeptide; PDBTitle: complete 12-subunit rna polymerase ii
78	c2o2tB	Alignment	not modelled	17.8	14	PDB header: structural protein Chain: B: PDB Molecule: multiple pdz domain protein; PDBTitle: the crystal structure of the 1st pdz domain of mpdz
79	c3bpuA	Alignment	not modelled	17.7	19	PDB header: transferase Chain: A: PDB Molecule: membrane-associated guanylate kinase, ww and pdz domain- PDBTitle: crystal structure of the 3rd pdz domain of human membrane associated2 guanylate kinase, c677s and c709s double mutant

80	c2iwoA	Alignment	not modelled	17.6	28	PDB header: signaling protein Chain: A: PDB Molecule: multiple pdz domain protein; PDBTitle: 12th pdz domain of multiple pdz domain protein mpdz (casp2 target)
81	c1yz6A	Alignment	not modelled	17.0	22	PDB header: translation Chain: A: PDB Molecule: probable translation initiation factor 2 alpha PDBTitle: crystal structure of intact alpha subunit of aif2 from2 pyrococcus abyssi
82	d1kq1a	Alignment	not modelled	16.9	27	Fold: Sm-like fold Superfamily: Sm-like ribonucleoproteins Family: Pleiotropic translational regulator Hfq
83	d1wi2a	Alignment	not modelled	16.6	26	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
84	c3psiA	Alignment	not modelled	16.5	14	PDB header: transcription Chain: A: PDB Molecule: transcription elongation factor spt6; PDBTitle: crystal structure of the spt6 core domain from saccharomyces2 cerevisiae, form spt6(239-1451)
85	c1go3E	Alignment	not modelled	16.5	25	PDB header: transferase Chain: E: PDB Molecule: dna-directed rna polymerase subunit e; PDBTitle: structure of an archeal homolog of the eukaryotic rna2 polymerase ii rpb4/rpb7 complex
86	c2pkua	Alignment	not modelled	16.2	13	PDB header: protein binding Chain: A: PDB Molecule: prkca-binding protein; PDBTitle: solution structure of pick1 pdz in complex with the2 carboxyl tail peptide of glur2
87	d1ujda	Alignment	not modelled	16.2	24	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
88	d1hk9a	Alignment	not modelled	16.1	36	Fold: Sm-like fold Superfamily: Sm-like ribonucleoproteins Family: Pleiotropic translational regulator Hfq
89	c3qhsD	Alignment	not modelled	16.1	36	PDB header: rna binding protein Chain: D: PDB Molecule: protein hfq; PDBTitle: crystal structure of full-length hfq from escherichia coli
90	c2qt7B	Alignment	not modelled	16.0	23	PDB header: hydrolase Chain: B: PDB Molecule: receptor-type tyrosine-protein phosphatase-like PDBTitle: crystallographic structure of the mature ectodomain of the2 human receptor-type protein-tyrosine phosphatase ia-2 at3 1.30 angstroms
91	d1u1sa1	Alignment	not modelled	15.9	36	Fold: Sm-like fold Superfamily: Sm-like ribonucleoproteins Family: Pleiotropic translational regulator Hfq
92	d2vnud3	Alignment	not modelled	15.7	15	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
93	c3hsbB	Alignment	not modelled	15.6	45	PDB header: rna binding protein/rna Chain: B: PDB Molecule: protein hfq; PDBTitle: crystal structure of ymah (hfq) from bacillus subtilis in complex with2 an rna aptamer
94	c2kviA	Alignment	not modelled	15.2	31	PDB header: rna binding protein Chain: A: PDB Molecule: nuclear polyadenylated rna-binding protein 3; PDBTitle: structure of nab3 rrm
95	d1whxa	Alignment	not modelled	14.6	6	Fold: Ferredoxin-like Superfamily: RNA-binding domain, RBD Family: Canonical RBD
96	c2ba0A	Alignment	not modelled	14.5	15	PDB header: rna binding protein Chain: A: PDB Molecule: archaeal exosome rna binding protein rrp4; PDBTitle: archaeal exosome core
97	d1qaua	Alignment	not modelled	14.5	13	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
98	c3ggeA	Alignment	not modelled	14.5	11	PDB header: protein binding Chain: A: PDB Molecule: pdz domain-containing protein gipc2; PDBTitle: crystal structure of the pdz domain of pdz domain-containing protein2 gipc2
99	c1kq1W	Alignment	not modelled	14.3	27	PDB header: translation Chain: W: PDB Molecule: host factor for q beta; PDBTitle: 1.55 a crystal structure of the pleiotropic translational2 regulator, hfq