
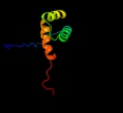



















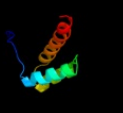

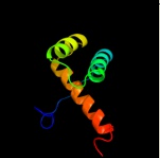
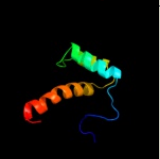
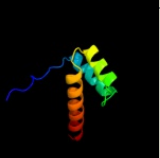
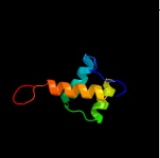

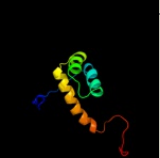

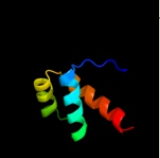




#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2dmtA_	 Alignment		99.9	40	PDB header: dna binding protein Chain: A; PDB Molecule: homeobox protein barh-like 1; PDBTitle: solution structure of the homeobox domain of homeobox2 protein barh-like 1
2	d1pufa_	 Alignment		99.9	33	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
3	c2dmsA_	 Alignment		99.9	51	PDB header: dna binding protein Chain: A; PDB Molecule: homeobox protein otx2; PDBTitle: solution structure of the homeobox domain of homeobox2 protein otx2
4	c2djnA_	 Alignment		99.9	37	PDB header: transcription Chain: A; PDB Molecule: homeobox protein dlx-5; PDBTitle: the solution structure of the homeobox domain of human2 homeobox protein dlx-5
5	c2kt0A_	 Alignment		99.9	29	PDB header: transcription regulation Chain: A; PDB Molecule: homeobox protein nanog; PDBTitle: solution structure of human stem cell transcription factor nanog2 homeodomain fragment
6	d1vnDA_	 Alignment		99.9	32	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
7	c3a01A_	 Alignment		99.9	33	PDB header: gene regulation/dna Chain: A; PDB Molecule: homeodomain-containing protein; PDBTitle: crystal structure of aristaless and clawless homeodomains bound to dna
8	d1ftza_	 Alignment		99.9	29	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
9	d1b72a_	 Alignment		99.8	38	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
10	d1zq3p1	 Alignment		99.8	31	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
11	d1nk3p_	 Alignment		99.8	35	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain

12	d1ahdp_	Alignment		99.8	30	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
13	c2dmuA_	Alignment		99.8	56	PDB header: dna binding protein Chain: A: PDB Molecule: homeobox protein gooseoid; PDBTitle: solution structure of the homeobox domain of homeobox2 protein gooseoid
14	d2lfba_	Alignment		99.8	27	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
15	d1fjla_	Alignment		99.8	62	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
16	c2l9rA_	Alignment		99.8	31	PDB header: transcription Chain: A: PDB Molecule: homeobox protein nkx-3.1; PDBTitle: solution nmr structure of homeobox domain of homeobox protein nkx-3.12 from homo sapiens, northeast structural genomics consortium target3 hr6470a
17	d1ftta_	Alignment		99.8	30	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
18	c2dmqA_	Alignment		99.8	33	PDB header: dna binding protein Chain: A: PDB Molecule: lim/homeobox protein lhx9; PDBTitle: solution structure of the homeobox domain of lim/homeobox2 protein lhx9
19	c3d1nK_	Alignment		99.8	36	PDB header: transcription regulator/dna Chain: K: PDB Molecule: pou domain, class 6, transcription factor 1; PDBTitle: structure of human brn-5 transcription factor in complex2 with corticotrophin-releasing hormone gene promoter
20	d2e1oa1	Alignment		99.8	39	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
21	d2craa1	Alignment	not modelled	99.8	30	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
22	d1b8ia_	Alignment	not modelled	99.8	34	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
23	d1uhsa_	Alignment	not modelled	99.8	27	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
24	d1ig7a_	Alignment	not modelled	99.8	40	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
25	c2da3A_	Alignment	not modelled	99.8	34	PDB header: transcription Chain: A: PDB Molecule: alpha-fetoprotein enhancer binding protein; PDBTitle: solution structure of the third homeobox domain of at-2 binding transcription factor 1 (atbf1)
26	d2cufa1	Alignment	not modelled	99.8	32	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
27	d2cuea1	Alignment		99.8	63	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
						PDB header: dna binding protein

28	c2k40A_	Alignment	not modelled	99.8	47	Chain: A: PDB Molecule: homeobox expressed in es cells 1; PDBTitle: nmr structure of hesx-1 homeodomain double mutant r31l/e42l
29	d1fjlb_	Alignment		99.8	64	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
30	d1jgga_	Alignment	not modelled	99.8	49	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
31	d1ztra1	Alignment	not modelled	99.8	36	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
32	c2da2A_	Alignment	not modelled	99.8	40	PDB header: transcription Chain: A: PDB Molecule: alpha-fetoprotein enhancer binding protein; PDBTitle: solution structure of the second homeobox domain of at-2 binding transcription factor 1 (atbf1)
33	d1bw5a_	Alignment	not modelled	99.8	32	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
34	c2da1A_	Alignment	not modelled	99.8	29	PDB header: transcription Chain: A: PDB Molecule: alpha-fetoprotein enhancer binding protein; PDBTitle: solution structure of the first homeobox domain of at-2 binding transcription factor 1 (atbf1)
35	d2hddb_	Alignment	not modelled	99.8	39	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
36	c2vi6F_	Alignment	not modelled	99.8	36	PDB header: transcription Chain: F: PDB Molecule: homeobox protein nanog; PDBTitle: crystal structure of the nanog homeodomain
37	d1sana_	Alignment	not modelled	99.8	30	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
38	d9anta_	Alignment	not modelled	99.8	35	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
39	d1ocpa_	Alignment	not modelled	99.8	30	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
40	d1hdpa_	Alignment	not modelled	99.8	27	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
41	d1wh5a_	Alignment	not modelled	99.8	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
42	c3l1pA_	Alignment	not modelled	99.8	28	PDB header: transcription/dna Chain: A: PDB Molecule: pou domain, class 5, transcription factor 1; PDBTitle: pou protein:dna complex
43	d1octc1	Alignment	not modelled	99.8	31	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
44	d1yz8p1	Alignment	not modelled	99.8	65	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
45	d1gt0c1	Alignment	not modelled	99.8	31	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
46	c2dn0A_	Alignment	not modelled	99.8	25	PDB header: transcription Chain: A: PDB Molecule: zinc fingers and homeoboxes protein 3; PDBTitle: solution structure of the second homeobox domain of human2 zinc fingers and homeoboxes protein 3
47	c3a03A_	Alignment	not modelled	99.8	35	PDB header: gene regulation Chain: A: PDB Molecule: t-cell leukemia homeobox protein 2; PDBTitle: crystal structure of hox11l1 homeodomain
48	d1f43a_	Alignment	not modelled	99.8	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
49	d1mh3a1	Alignment	not modelled	99.8	24	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
50	d1du0a_	Alignment	not modelled	99.8	37	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
51	d1p7ia_	Alignment	not modelled	99.8	40	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
52	c1hf0A_	Alignment	not modelled	99.8	31	PDB header: transcription factor Chain: A: PDB Molecule: octamer-binding transcription factor 1; PDBTitle: crystal structure of the dna-binding domain of oct-1 bound2 to dna as a dimer
53	d1le8a_	Alignment	not modelled	99.8	23	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain

54	d2hi3a1	Alignment	not modelled	99.8	30	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
55	clic8B_	Alignment	not modelled	99.8	29	PDB header: transcription/dna Chain: B: PDB Molecule: hepatocyte nuclear factor 1-alpha; PDBTitle: hepatocyte nuclear factor 1a bound to dna : mody3 gene2 product
56	d1au7a1	Alignment	not modelled	99.8	31	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
57	c2h8rA_	Alignment	not modelled	99.8	30	PDB header: transcription activator/dna Chain: A: PDB Molecule: hepatocyte nuclear factor 1-beta; PDBTitle: hepatocyte nuclear factor 1b bound to dna: mody5 gene2 product
58	d1e3oc1	Alignment	not modelled	99.8	31	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
59	c2da4A_	Alignment	not modelled	99.8	23	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hypothetical protein dkfzp686k21156; PDBTitle: solution structure of the homeobox domain of the2 hypothetical protein, dkfzp686k21156
60	d2ecca1	Alignment	not modelled	99.7	19	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
61	c2dmnA_	Alignment	not modelled	99.7	19	PDB header: transcription Chain: A: PDB Molecule: homeobox protein tgif2lx; PDBTitle: the solution structure of the homeobox domain of human2 homeobox protein tgif2lx
62	c2da5A_	Alignment	not modelled	99.7	24	PDB header: dna binding protein Chain: A: PDB Molecule: zinc fingers and homeoboxes protein 3; PDBTitle: solution structure of the second homeobox domain of zinc2 fingers and homeoboxes protein 3 (triple homeobox 13 protein)
63	c1au7B_	Alignment	not modelled	99.7	30	PDB header: transcription/dna Chain: B: PDB Molecule: protein pit-1; PDBTitle: pit-1 mutant/dna complex
64	d1lfup_	Alignment	not modelled	99.7	27	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
65	c2dmpA_	Alignment	not modelled	99.7	19	PDB header: dna binding protein Chain: A: PDB Molecule: zinc fingers and homeoboxes protein 2; PDBTitle: solution structure of the third homeobox domain of zinc2 fingers and homeoboxes protein 2
66	d1p7jb_	Alignment	not modelled	99.7	35	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
67	d2ecba1	Alignment	not modelled	99.7	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
68	cls7eA_	Alignment	not modelled	99.7	20	PDB header: transcription Chain: A: PDB Molecule: hepatocyte nuclear factor 6; PDBTitle: solution structure of hnf-6
69	c3narA_	Alignment	not modelled	99.7	22	PDB header: transcription Chain: A: PDB Molecule: zinc fingers and homeoboxes protein 1; PDBTitle: crystal structure of zhx1 hd4 (zinc-fingers and homeoboxes protein 1,2 homeodomain 4)
70	c2da7A_	Alignment	not modelled	99.7	16	PDB header: dna binding protein Chain: A: PDB Molecule: zinc finger homeobox protein 1b; PDBTitle: solution structure of the homeobox domain of zinc finger2 homeobox protein 1b (smad interacting protein 1)
71	d1s7ea1	Alignment	not modelled	99.7	24	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
72	d1pufb_	Alignment	not modelled	99.7	31	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
73	c2xsdC_	Alignment	not modelled	99.7	30	PDB header: transcription/dna Chain: C: PDB Molecule: pou domain, class 3, transcription factor 1; PDBTitle: crystal structure of the dimeric oct-6 (pou3f1) pou domain2 bound to palindromic more dna
74	d1lfba_	Alignment	not modelled	99.7	29	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
75	d1mnmc_	Alignment	not modelled	99.7	29	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
76	c2e19A_	Alignment	not modelled	99.7	22	PDB header: transcription Chain: A: PDB Molecule: transcription factor 8; PDBTitle: solution structure of the homeobox domain from human nil-2-2 a zinc finger protein, transcription factor 8
77	c3nauA_	Alignment	not modelled	99.7	25	PDB header: transcription Chain: A: PDB Molecule: zinc fingers and homeoboxes protein 2; PDBTitle: crystal structure of zhx2 hd2 (zinc-fingers and homeoboxes protein 2,2 homeodomain 2)
78	d1le8b_	Alignment	not modelled	99.7	28	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
79	d1o4xa1	Alignment	not modelled	99.7	29	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
						Fold: DNA/RNA-binding 3-helical bundle

80	d1du6a_	Alignment	not modelled	99.7	26	Superfamily: Homeodomain-like Family: Homeodomain
81	d1yrnb_	Alignment	not modelled	99.7	28	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
82	d1akha_	Alignment	not modelled	99.7	25	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
83	d2cqa1_	Alignment	not modelled	99.6	25	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
84	d1x2na1_	Alignment	not modelled	99.6	22	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
85	c2lk2A_	Alignment	not modelled	99.6	14	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
86	d1x2ma1_	Alignment	not modelled	99.6	34	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
87	d2r5yb1_	Alignment	not modelled	99.6	31	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
88	d1k61a_	Alignment	not modelled	99.6	29	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
89	d2p81a1_	Alignment	not modelled	99.5	33	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
90	c2nzzA_	Alignment	not modelled	98.0	41	PDB header: membrane protein Chain: A: PDB Molecule: penetratin conjugated gas (374-394) peptide; PDBTitle: nmr structure analysis of the penetratin conjugated gas2 (374-394) peptide
91	c1mh3A_	Alignment	not modelled	97.5	25	PDB header: sugar binding, dna binding protein Chain: A: PDB Molecule: maltose binding-a1 homeodomain protein chimera; PDBTitle: maltose binding-a1 homeodomain protein chimera, crystal2 form i
92	d1wi3a_	Alignment	not modelled	97.0	26	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
93	d1wh7a_	Alignment	not modelled	96.5	27	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Homeodomain
94	c2elha_	Alignment	not modelled	91.8	19	PDB header: dna binding protein Chain: A: PDB Molecule: cg11849-pa; PDBTitle: solution structure of the cenp-b n-terminal dna-binding2 domain of fruit fly distal antenna cg11849-pa
95	d1hlva1_	Alignment	not modelled	90.8	14	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Centromere-binding
96	c1hlva_	Alignment	not modelled	87.1	14	PDB header: dna binding protein/dna Chain: A: PDB Molecule: major centromere autoantigen b; PDBTitle: crystal structure of cenp-b(1-129) complexed with the cenp-2 b box dna
97	c3bdnB_	Alignment	not modelled	86.0	22	PDB header: transcription/dna Chain: B: PDB Molecule: lambda repressor; PDBTitle: crystal structure of the lambda repressor
98	d1smvf2_	Alignment	not modelled	85.6	9	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Sigma3 and sigma4 domains of RNA polymerase sigma factors Family: Sigma4 domain
99	c2ys9A_	Alignment	not modelled	83.4	20	PDB header: transcription Chain: A: PDB Molecule: homeobox and leucine zipper protein homez; PDBTitle: structure of the third homeodomain from the human homeobox2 and leucine zipper protein, homez
100	c2ktga_	Alignment	not modelled	79.1	15	PDB header: ca-binding protein Chain: A: PDB Molecule: calmodulin, putative; PDBTitle: calmodulin like protein from entamoeba histolytica: solution structure2 and calcium binding properties of a partially folded protein
101	d1rp3a2_	Alignment	not modelled	78.6	10	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Sigma3 and sigma4 domains of RNA polymerase sigma factors Family: Sigma4 domain
102	c1or7A_	Alignment	not modelled	78.5	20	PDB header: transcription Chain: A: PDB Molecule: rna polymerase sigma-e factor; PDBTitle: crystal structure of escherichia coli sigmae with the cytoplasmic2 domain of its anti-sigma rsea
103	d1avsa_	Alignment	not modelled	76.5	8	Fold: EF Hand-like Superfamily: EF-hand Family: Calmodulin-like
104	c3mzyA_	Alignment	not modelled	74.2	9	PDB header: rna binding protein Chain: A: PDB Molecule: rna polymerase sigma-h factor; PDBTitle: the crystal structure of the rna polymerase sigma-h factor from2 fusobacterium nucleatum to 2.5a
105	c1rp3G_	Alignment	not modelled	73.0	10	PDB header: transcription Chain: G: PDB Molecule: rna polymerase sigma factor sigma-28 (fliA); PDBTitle: cocrystal structure of the flagellar sigma/anti-sigma2

						complex, sigma-28/flgm
106	d1rioa_	Alignment	not modelled	71.5	22	Fold: lambda repressor-like DNA-binding domains Superfamily: lambda repressor-like DNA-binding domains Family: Phage repressors
107	c3hugA_	Alignment	not modelled	71.4	11	PDB header: transcription/membrane protein Chain: A: PDB Molecule: rna polymerase sigma factor; PDBTitle: crystal structure of mycobacterium tuberculosis anti-sigma factor rslA2 in complex with -35 promoter binding domain of sigI
108	d1bw6a_	Alignment	not modelled	70.1	15	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Centromere-binding
109	d1or7a1	Alignment	not modelled	69.9	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Sigma3 and sigma4 domains of RNA polymerase sigma factors Family: Sigma4 domain
110	c2lfwA_	Alignment	not modelled	67.9	20	PDB header: signaling protein Chain: A: PDB Molecule: phyr sigma-like domain; PDBTitle: nmr structure of the phyrsl-nepr complex from sphingomonas sp. fr1
111	c3fmyA_	Alignment	not modelled	66.0	8	PDB header: dna binding protein Chain: A: PDB Molecule: hth-type transcriptional regulator mqsa PDBTitle: structure of the c-terminal domain of the e. coli protein2 mqsa (ygit/b3021)
112	d1iufa1	Alignment	not modelled	63.2	29	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Centromere-binding
113	d1l3la1	Alignment	not modelled	61.2	14	Fold: DNA/RNA-binding 3-helical bundle Superfamily: C-terminal effector domain of the bipartite response regulators Family: GerE-like (LuxR/UhpA family of transcriptional regulators)
114	c2o8xA_	Alignment	not modelled	59.9	12	PDB header: transcription Chain: A: PDB Molecule: probable rna polymerase sigma-c factor; PDBTitle: crystal structure of the "-35 element" promoter recognition domain of2 mycobacterium tuberculosis sigc
115	c2fjrB_	Alignment	not modelled	58.9	6	PDB header: transcription regulator Chain: B: PDB Molecule: repressor protein ci; PDBTitle: crystal structure of bacteriophage 186
116	c3sztB_	Alignment	not modelled	58.8	18	PDB header: transcription Chain: B: PDB Molecule: quorum-sensing control repressor; PDBTitle: quorum sensing control repressor, qscr, bound to n-3-oxo-dodecanoyl-l-2 homoserine lactone
117	d1sw8a_	Alignment	not modelled	58.1	19	Fold: EF Hand-like Superfamily: EF-hand Family: Calmodulin-like
118	d1g8ia_	Alignment	not modelled	55.7	29	Fold: EF Hand-like Superfamily: EF-hand Family: Calmodulin-like
119	c2q1zA_	Alignment	not modelled	54.9	19	PDB header: transcription Chain: A: PDB Molecule: rpoe, ecf sigE; PDBTitle: crystal structure of rhodobacter sphaeroides sigE in complex with the2 anti-sigma chrr
120	d1fsea_	Alignment	not modelled	54.1	19	Fold: DNA/RNA-binding 3-helical bundle Superfamily: C-terminal effector domain of the bipartite response regulators Family: GerE-like (LuxR/UhpA family of transcriptional regulators)