

Detailed template information

Into	<u>information</u>							
#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information		
1	d1pufa_	Alignment		99.8	39	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain		
2	d1nk3p_	Alignment	and a	99.8	34	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain		
3	<u>c3a01A</u> _	Alignment	Zean	99.8	37	PDB header:gene regulation/dna Chain: A: PDB Molecule:homeodomain-containing protein; PDBTitle: crystal structure of aristaless and clawless homeodomains bound to dna		
4	<u>c2m34A</u> _	Alignment	EE	99.8	39	PDB header:transcription Chain: A: PDB Molecule:homeobox protein gbx-1; PDBTitle: nmr structure of the homeodomain transcription factor gbx1 from homo2 sapiens		
5	c5luxK_	Alignment	EE	99.8	40	PDB header:transcription Chain: K: PDB Molecule:homeobox protein cdx-1; PDBTitle: homeobox transcription factor cdx1 bound to methylated dna		
6	d1zq3p1	Alignment	Eum)	99.8	32	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain		
7	<u>d2cuea1</u>	Alignment	E. E.	99.8	31	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain		
8	<u>d1jgga</u>	Alignment	E.E.	99.8	40	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain		
9	d1fjla_	Alignment	The state of the s	99.8	29	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain		
10	<u>d1b72a</u> _	Alignment	Elem	99.8	38	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain		
11	<u>dlahdp</u>	Alignment		99.8	38	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain		

12	 d2e1oa1	Alignment		99.8	46	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
13	c2dmtA_	Alignment	and the same of th	99.8	41	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
14	d9anta_	Alignment		99.8	39	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
15	dlig7a_	Alignment	E	99.8	35	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
16	c2mgqA_	Alignment		99.8	24	PDB header:dna binding protein Chain: A: PDB Molecule:homeobox protein ceh-37; PDBTitle: structure of ceh37 homeodomain
17	<u>c4uutA</u>	Alignment	Euro	99.8	34	PDB header:transcription Chain: A: PDB Molecule:homeotic protein ultrabithorax; PDBTitle: crystal structure of the ultrabithorax protein
18	<u>d1b8ia</u> _	Alignment		99.8	36	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
19	dlle8a_	Alignment	1003	99.8	34	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
20	d2craa1	Alignment	Since.	99.8	24	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
21	<u>c3a03A_</u>	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
22	d1fjlb_	Alignment	not modelled	99.8	31	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
23	c2dmuA_	Alignment	not modelled	99.8	35	PDB header:dna binding protein Chain: A: PDB Molecule:homeobox protein goosecoid; PDBTitle: solution structure of the homeobox domain of homeobox2 protein goosecoid
24	<u>c2l9rA_</u>	Alignment	not modelled	99.8	36	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
25	c2dmsA_	Alignment	not modelled	99.8	28	PDB header:dna binding protein Chain: A: PDB Molecule:homeobox protein otx2; PDBTitle: solution structure of the homeobox domain of homeobox2 protein otx2
26	<u>c2k40A_</u>	Alignment	not modelled	99.8	29	PDB header:dna binding protein Chain: A: PDB Molecule:homeobox expressed in es cells 1; PDBTitle: nmr structure of hesx-1 homeodomain double mutant r31l/e42l
27	d2hddb_	Alignment	not modelled	99.8	34	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
28	d1ftza_	Alignment	not modelled	99.8	35	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain

29	d1ftta_	Alignment	not modelled	99.8	33	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
30	dluhsa_	Alignment	not modelled	99.8	34	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
31	d2cufa1	Alignment	not modelled	99.8	19	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
32	d1mh3a1	Alignment	not modelled	99.8	32	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
33	dloctcl	Alignment	not modelled	99.8	30	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
34	c2vi6F_	Alignment	not modelled	99.8	28	PDB header:transcription Chain: F: PDB Molecule:homeobox protein nanog; PDBTitle: crystal structure of the nanog homeodomain
35	d1p7ia_	Alignment	not modelled	99.8	35	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
36	d1yz8p1	Alignment	not modelled	99.7	29	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
37	d1sana_	Alignment	not modelled	99.7	40	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
38	c2djnA_	Alignment	not modelled	99.7	31	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
39	c2kt0A_	Alignment	not modelled	99.7	28	PDB header:transcription regulation Chain: A: PDB Molecule:homeobox protein nanog; PDBTitle: solution structure of human stem cell transcription factor nanog2 homeodomain fragment
40	d1du0a_	Alignment	not modelled	99.7	35	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
41	c2dmqA_	Alignment	not modelled	99.7	27	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
42	c2m0cA_	Alignment	not modelled	99.7	37	PDB header:gene regulation Chain: A: PDB Molecule:homeobox protein aristaless-like 4; PDBTitle: solution nmr structure of homeobox domain of human alx4, northeast2 structural genomics consortium (nesg) target hr4490c
43	c5hodD_	Alignment	not modelled	99.7	36	PDB header:transcription Chain: D: PDB Molecule:lim/homeobox protein lhx4; PDBTitle: structure of lhx4 transcription factor complexed with dna
44	dle3oc1	Alignment	not modelled	99.7	30	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
45	c2da2A_	Alignment	not modelled	99.7	30	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)
46	d1bw5a_	Alignment	not modelled	99.7	31	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
47	dlau7al	Alignment	not modelled	99.7	31	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
48	d1gt0c1	Alignment	not modelled	99.7	30	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
49	d2lfba_	Alignment	not modelled	99.7	21	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
50	dlvnda_	Alignment	not modelled	99.7	33	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain DDB to a day the properties.
51	c2ly9A_	Alignment	not modelled	99.7	17	PDB header:transcription Chain: A: PDB Molecule:zinc fingers and homeoboxes protein 1; PDBTitle: solution nmr structure of homeobox 2 domain from human zhx1 repressor,2 northeast structural genomics consortium (nesg) target hr7907f
52	d1ztra1	Alignment	not modelled	99.7	31	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
53	c2da3A_	Alignment	not modelled	99.7	33	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)
54	d2ecca1	Alignment	not modelled	99.7	24	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
						Fold:DNA/RNA-binding 3-helical bundle

55	<u>d1p7jb_</u>	Alignment	not modelled	99.7	38	Superfamily:Homeodomain-like Family:Homeodomain
56	d1hdpa_	Alignment	not modelled	99.7	31	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
57	<u>c2da1A_</u>	Alignment	not modelled	99.7	26	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)
58	c2dn0A_	Alignment	not modelled	99.7	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
59	d1f43a_	Alignment	not modelled	99.7	34	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
60	d2hi3a1	Alignment	not modelled	99.7	37	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
61	d2ecba1	Alignment	not modelled	99.7	21	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
62	dlocpa_	Alignment	not modelled	99.7	31	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
63	c2da5A_	Alignment	not modelled	99.7	22	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)
64	c3narA_	Alignment	not modelled	99.7	30	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)
65	c1hf0A_	Alignment	not modelled	99.7	30	PDB header:transcription 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
66	<u>c3d1nK_</u>	Alignment	not modelled	99.7	28	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
67	c2dmpA_	Alignment	not modelled	99.7	28	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
68	c3l1pA_	Alignment	not modelled	99.7	31	PDB header:transcription/dna Chain: A: PDB Molecule:pou domain, class 5, transcription factor 1; PDBTitle: pou protein:dna complex
69	d1mnmc_	Alignment	not modelled	99.7	30	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
70	d1lfup_	Alignment	not modelled	99.7	27	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
71	c2mg4B_	Alignment	not modelled	99.7	26	PDB header:de novo protein Chain: B: PDB Molecule:computational designed homodimer; PDBTitle: computational design and experimental verification of a symmetric2 protein homodimer
72	c3nauA_	Alignment	not modelled	99.7	22	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)
73	d2r5yb1	Alignment	not modelled	99.7	31	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
74	d1pufb_	Alignment	not modelled	99.7	27	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
75	d1le8b_	Alignment	not modelled	99.7	28	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
76	dlyrnb_	Alignment	not modelled	99.7	28	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
77	<u>c2da7A_</u>	Alignment	not modelled	99.7	17	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)
78	dls7eal	Alignment	not modelled	99.7	24	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
79	d1k61a_	Alignment	not modelled	99.7	28	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
80	c2lk2A_	Alignment	not modelled	99.7	27	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
Ω1	c2e10A	Allenance	not modelled	99.7	17	PDB header:transcription Chain: A: PDB Molecule:transcription factor 8;

01	CZETAW_	Alignment	not modelled	99.1	1/	PDBTitle: solution structure of the homeobox domain from human nil-2-2 a zinc finger protein, transcription factor 8
82	dlakha_	Alignment	not modelled	99.7	33	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
83	dlo4xal	Alignment	not modelled	99.7	31	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
84	c2dmnA_	Alignment	not modelled	99.7	27	PDB header:transcription Chain: A: PDB Molecule:homeobox protein tgif2lx; PDBTitle: the solution structure of the homeobox domain of human2 homeobox protein tgif2lx
85	c1x2mA_	Alignment	not modelled	99.7	26	PDB header:transcription Chain: A: PDB Molecule:lag1 longevity assurance homolog 6; PDBTitle: solution structure of the homeobox domain of mouse lag12 longevity assurance homolog 6
86	c1au7B_	Alignment	not modelled	99.7	31	PDB header:transcription/dna Chain: B: PDB Molecule:protein pit-1; PDBTitle: pit-1 mutant/dna complex
87	d1lfba_	Alignment	not modelled	99.6	25	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
88	c2xsdC_	Alignment	not modelled	99.6	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
89	d1x2na1	Alignment	not modelled	99.6	22	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
90	c2da4A_	Alignment	not modelled	99.6	20	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
91	c1ic8B_	Alignment	not modelled	99.6	26	PDB header:transcription/dna Chain: B: PDB Molecule:hepatocyte nuclear factor 1-alpha; PDBTitle: hepatocyte nuclear factor 1a bound to dna: mody3 gene product
92	d1du6a_	Alignment	not modelled	99.6	31	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
93	d2cqxa1	Alignment	not modelled	99.6	25	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
94	d1wh5a_	Alignment	not modelled	99.6	18	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
95	c2h8rA_	Alignment	not modelled	99.6	27	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
96	c1s7eA_	Alignment	not modelled	99.6	24	PDB header:transcription Chain: A: PDB Molecule:hepatocyte nuclear factor 6; PDBTitle: solution structure of hnf-6
97	d1x2ma1	Alignment	not modelled	99.6	29	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
98	<u>c5flvA_</u>	Alignment	not modelled	99.5	37	PDB header:transcription Chain: A: PDB Molecule:homeobox protein nkx-2.5, t-box transcription factor tbx5; PDBTitle: crystal structure of nkx2-5 and tbx5 bound to the nppa2 promoter region
99	<u>d2p81a1</u>	Alignment	not modelled	99.4	40	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
100	c4egcA_	Alignment	not modelled	99.4	31	PDB header:transcription/hydrolase Chain: A: PDB Molecule:maltose-binding periplasmic protein, homeobox protein six1 PDBTitle: crystal structure of mbp-fused human six1 bound to human eya2 eya2 domain
101	c1mh3A_	Alignment	not modelled	97.4	34	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
102	d1wi3a_	Alignment	not modelled	96.7	18	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
103	c4ndlC_	Alignment	not modelled	96.4	33	PDB header:de novo protein Chain: C: PDB Molecule:enh-c2b, computational designed homodimer; PDBTitle: computational design and experimental verification of a symmetric2 homodimer
104	c2nzzA_	Alignment	not modelled	96.1	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
105	d1wh7a_	Alignment	not modelled	95.1	21	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Homeodomain-like Family:Homeodomain
106	c4ndkA_	Alignment	not modelled	91.7	27	PDB header:fluorescent protein, de novo protein Chain: A: PDB Molecule:e23p-yfp, chimeric construct; PDBTitle: crystal structure of a computational designed engrailed homeodomain2 variant fused with yfp

107	c1hlvA_	Alignment	not modelled	89.2	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
108	d1hlva1	Alignment	not modelled	88.0	Fold:DNA/RNA-binding 3-helical bundle 24 Superfamily:Homeodomain-like Family:Centromere-binding
109	<u>c2ys9A_</u>	Alignment	not modelled	85.1	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
110	d1smyf2	Alignment	not modelled	82.0	Fold:DNA/RNA-binding 3-helical bundle Superfamily:Sigma3 and sigma4 domains of RNA polymerase sigma factors Family:Sigma4 domain
111	d1bw6a_	Alignment	not modelled	80.6	Fold:DNA/RNA-binding 3-helical bundle 20 Superfamily:Homeodomain-like Family:Centromere-binding
112	c2rn7A_	Alignment	not modelled	79.4	PDB header:unknown function Chain: A: PDB Molecule:is629 orfa; PDBTitle: nmr solution structure of tnpe protein from shigella2 flexneri. northeast structural genomics target sfr125
113	d2jn6a1	Alignment	not modelled	75.7	Fold:DNA/RNA-binding 3-helical bundle 14 Superfamily:Homeodomain-like Family:Cgl2762-like
114	c4cxfA_	Alignment	not modelled	75.6	PDB header:transcription Chain: A: PDB Molecule:rna polymerase sigma factor cnrh; PDBTitle: structure of cnrh in complex with the cytosolic domain of cnry
115	c3vdoA_	Alignment	not modelled	74.6	PDB header:dna binding protein/protein binding Chain: A: PDB Molecule:rna polymerase sigma factor sigk; PDBTitle: structure of extra-cytoplasmic function(ecf) sigma factor sigk in 2 complex with its negative regulator rska from mycobacterium3 tuberculosis
116	c3mzyA_	Alignment	not modelled	73.0	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
117	clor7A_	Alignment	not modelled	72.8	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
118	d1 3 a1	Alignment	not modelled	72.7	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)
119	c2pmyB_	Alignment	not modelled	71.9	PDB header:structural genomics, unknown function Chain: B: PDB Molecule:ras and ef-hand domain-containing protein; PDBTitle: ef-hand domain of human rasef
120	c4nn1A_	Alignment	not modelled	70.5	PDB header:transcription Chain: A: PDB Molecule:transcriptional regulator; PDBTitle: crystal structure of transcriptional regulator rv1219c of2 mycobacterium tuberculosis