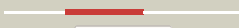
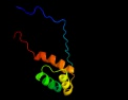



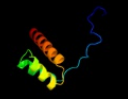

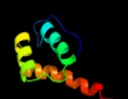














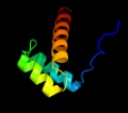



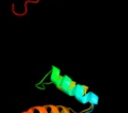






#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c2dmtA_</a>	 Alignment		99.9	35	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> homeobox protein barh-like 1; <b>PDBTitle:</b> solution structure of the homeobox domain of homeobox2 protein barh-like 1
2	<a href="#">c2kt0A_</a>	 Alignment		99.9	26	<b>PDB header:</b> transcription regulation <b>Chain:</b> A: <b>PDB Molecule:</b> homeobox protein nanog; <b>PDBTitle:</b> solution structure of the homeobox domain of nanog2 homeodomain fragment
3	<a href="#">d1pufa_</a>	 Alignment		99.9	37	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
4	<a href="#">c2m34A_</a>	 Alignment		99.9	46	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> homeobox protein gbx-1; <b>PDBTitle:</b> nmr structure of the homeodomain transcription factor gbx1 from homo2 sapiens
5	<a href="#">c2dmsA_</a>	 Alignment		99.9	47	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> homeobox protein otx2; <b>PDBTitle:</b> solution structure of the homeobox domain of homeobox2 protein otx2
6	<a href="#">d1zq3p1</a>	 Alignment		99.9	34	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
7	<a href="#">d1nk3p_</a>	 Alignment		99.8	40	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
8	<a href="#">c3a01A_</a>	 Alignment		99.8	36	<b>PDB header:</b> gene regulation/dna <b>Chain:</b> A: <b>PDB Molecule:</b> homeodomain-containing protein; <b>PDBTitle:</b> crystal structure of aristaless and clawless homeodomains bound to dna
9	<a href="#">d1b72a_</a>	 Alignment		99.8	46	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
10	<a href="#">d1ahdp_</a>	 Alignment		99.8	40	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
11	<a href="#">c2dmuA_</a>	 Alignment		99.8	48	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> homeobox protein gooseoid; <b>PDBTitle:</b> solution structure of the homeobox domain of homeobox2 protein gooseoid

12	<a href="#">d1fjla_</a>	Alignment		99.8	59	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
13	<a href="#">c2djna_</a>	Alignment		99.8	33	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> homeobox protein dlx-5; <b>PDBTitle:</b> the solution structure of the homeobox domain of human2 homeobox protein dlx-5
14	<a href="#">c2dmqa_</a>	Alignment		99.8	30	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> lim/homeobox protein lhx9; <b>PDBTitle:</b> solution structure of the homeobox domain of lim/homeobox2 protein lhx9
15	<a href="#">d1ftza_</a>	Alignment		99.8	38	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
16	<a href="#">d2lfba_</a>	Alignment		99.8	28	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
17	<a href="#">d1b8ia_</a>	Alignment		99.8	41	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
18	<a href="#">d1ftta_</a>	Alignment		99.8	38	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
19	<a href="#">c2da3A_</a>	Alignment		99.8	32	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> alpha-fetoprotein enhancer binding protein; <b>PDBTitle:</b> solution structure of the third homeobox domain of at-2 binding transcription factor 1 (atbf1)
20	<a href="#">c2m0cA_</a>	Alignment		99.8	62	<b>PDB header:</b> gene regulation <b>Chain:</b> A: <b>PDB Molecule:</b> homeobox protein aristaless-like 4; <b>PDBTitle:</b> solution nmr structure of homeobox domain of human alx4, northeast2 structural genomics consortium (nesg) target hr4490c
21	<a href="#">d2cufa1</a>	Alignment	not modelled	99.8	29	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
22	<a href="#">c2l9ra_</a>	Alignment	not modelled	99.8	31	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> homeobox protein nkx-3.1; <b>PDBTitle:</b> solution nmr structure of homeobox domain of homeobox protein nkx-3.12 from homo sapiens, northeast structural genomics consortium target3 hr6470a
23	<a href="#">d1ig7a_</a>	Alignment	not modelled	99.8	46	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
24	<a href="#">d1jgga_</a>	Alignment	not modelled	99.8	46	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
25	<a href="#">d1uhsa_</a>	Alignment	not modelled	99.8	30	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
26	<a href="#">d2e1oa1</a>	Alignment	not modelled	99.8	40	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
27	<a href="#">d2cuea1</a>	Alignment	not modelled	99.8	59	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
28	<a href="#">d1vnda_</a>	Alignment	not modelled	99.8	33	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
						<b>Fold:</b> DNA/RNA-binding 3-helical bundle

29	<a href="#">d2craa1</a>	Alignment	not modelled	99.8	32	<b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
30	<a href="#">c2k40A</a>	Alignment	not modelled	99.8	41	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> homeobox expressed in es cells 1; <b>PDBTitle:</b> nmr structure of hex-1 homeodomain double mutant r31l/e42l
31	<a href="#">d1fjlb</a>	Alignment	not modelled	99.8	60	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
32	<a href="#">d2hddb</a>	Alignment	not modelled	99.8	43	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
33	<a href="#">c2da2A</a>	Alignment	not modelled	99.8	36	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> alpha-fetoprotein enhancer binding protein; <b>PDBTitle:</b> solution structure of the second homeobox domain of at-2 binding transcription factor 1 (atbf1)
34	<a href="#">c3d1nK</a>	Alignment	not modelled	99.8	27	<b>PDB header:</b> transcription regulator/dna <b>Chain:</b> K: <b>PDB Molecule:</b> pou domain, class 6, transcription factor 1; <b>PDBTitle:</b> structure of human brn-5 transcription factor in complex2 with corticotrophin-releasing hormone gene promoter
35	<a href="#">d1bw5a</a>	Alignment	not modelled	99.8	30	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
36	<a href="#">c2da1A</a>	Alignment	not modelled	99.8	26	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> alpha-fetoprotein enhancer binding protein; <b>PDBTitle:</b> solution structure of the first homeobox domain of at-2 binding transcription factor 1 (atbf1)
37	<a href="#">d9anta</a>	Alignment	not modelled	99.8	45	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
38	<a href="#">d1octc1</a>	Alignment	not modelled	99.8	27	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
39	<a href="#">d1yz8p1</a>	Alignment	not modelled	99.8	55	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
40	<a href="#">c2vi6F</a>	Alignment	not modelled	99.8	35	<b>PDB header:</b> transcription <b>Chain:</b> F: <b>PDB Molecule:</b> homeobox protein nanog; <b>PDBTitle:</b> crystal structure of the nanog homeodomain
41	<a href="#">d1ztra1</a>	Alignment	not modelled	99.8	42	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
42	<a href="#">d1gt0c1</a>	Alignment	not modelled	99.8	26	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
43	<a href="#">d1sana</a>	Alignment	not modelled	99.8	41	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
44	<a href="#">d1wh5a</a>	Alignment	not modelled	99.8	19	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
45	<a href="#">d1hdpa</a>	Alignment	not modelled	99.8	24	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
46	<a href="#">c3l1pA</a>	Alignment	not modelled	99.8	23	<b>PDB header:</b> transcription/dna <b>Chain:</b> A: <b>PDB Molecule:</b> pou domain, class 5, transcription factor 1; <b>PDBTitle:</b> pou protein:dna complex
47	<a href="#">c3a03A</a>	Alignment	not modelled	99.8	40	<b>PDB header:</b> gene regulation <b>Chain:</b> A: <b>PDB Molecule:</b> it-cell leukemia homeobox protein 2; <b>PDBTitle:</b> crystal structure of hox11l1 homeodomain
48	<a href="#">d1ocpa</a>	Alignment	not modelled	99.8	25	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
49	<a href="#">c2ly9A</a>	Alignment	not modelled	99.8	21	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> zinc fingers and homeoboxes protein 1; <b>PDBTitle:</b> solution nmr structure of homeobox 2 domain from human zhx1 repressor,2 northeast structural genomics consortium (nesg) target hr7907f
50	<a href="#">c2dn0A</a>	Alignment	not modelled	99.8	20	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> zinc fingers and homeoboxes protein 3; <b>PDBTitle:</b> solution structure of the second homeobox domain of human2 zinc fingers and homeoboxes protein 3
51	<a href="#">d1p7ia</a>	Alignment	not modelled	99.8	47	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
52	<a href="#">d1du0a</a>	Alignment	not modelled	99.8	42	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
53	<a href="#">d1e3oc1</a>	Alignment	not modelled	99.8	26	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
54	<a href="#">d2hi3a1</a>	Alignment	not modelled	99.8	30	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
55	<a href="#">c1hf0A</a>	Alignment	not modelled	99.8	28	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> octamer-binding transcription factor 1; <b>PDBTitle:</b> crystal structure of the dna-binding domain of oct-1

						bound2 to dna as a dimer
56	<a href="#">d1au7a1</a>	Alignment	not modelled	99.8	27	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
57	<a href="#">c2da4A</a>	Alignment	not modelled	99.8	21	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> hypothetical protein dkfzp686k21156; <b>PDBTitle:</b> solution structure of the homeobox domain of the2 hypothetical protein, dkfzp686k21156
58	<a href="#">d1le8a</a>	Alignment	not modelled	99.8	17	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
59	<a href="#">d1mh3a1</a>	Alignment	not modelled	99.8	17	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
60	<a href="#">d1f43a</a>	Alignment	not modelled	99.8	17	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
61	<a href="#">c2dmnA</a>	Alignment	not modelled	99.8	18	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> homeobox protein tgif2lx; <b>PDBTitle:</b> the solution structure of the homeobox domain of human2 homeobox protein tgif2lx
62	<a href="#">d2ecca1</a>	Alignment	not modelled	99.7	18	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
63	<a href="#">c1ic8B</a>	Alignment	not modelled	99.7	26	<b>PDB header:</b> transcription/dna <b>Chain:</b> B: <b>PDB Molecule:</b> hepatocyte nuclear factor 1-alpha; <b>PDBTitle:</b> hepatocyte nuclear factor 1a bound to dna : mody3 gene2 product
64	<a href="#">c2h8rA</a>	Alignment	not modelled	99.7	30	<b>PDB header:</b> transcription activator/dna <b>Chain:</b> A: <b>PDB Molecule:</b> hepatocyte nuclear factor 1-beta; <b>PDBTitle:</b> hepatocyte nuclear factor 1b bound to dna: mody5 gene2 product
65	<a href="#">c1au7B</a>	Alignment	not modelled	99.7	25	<b>PDB header:</b> transcription/dna <b>Chain:</b> B: <b>PDB Molecule:</b> protein pit-1; <b>PDBTitle:</b> pit-1 mutant/dna complex
66	<a href="#">c2da5A</a>	Alignment	not modelled	99.7	19	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> zinc fingers and homeoboxes protein 3; <b>PDBTitle:</b> solution structure of the second homeobox domain of zinc2 fingers and homeoboxes protein 3 (triple homeobox 13 protein)
67	<a href="#">c2dmpA</a>	Alignment	not modelled	99.7	15	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> zinc fingers and homeoboxes protein 2; <b>PDBTitle:</b> solution structure of the third homeobox domain of zinc2 fingers and homeoboxes protein 2
68	<a href="#">d2ecba1</a>	Alignment	not modelled	99.7	15	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
69	<a href="#">c2da7A</a>	Alignment	not modelled	99.7	14	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> zinc finger homeobox protein 1b; <b>PDBTitle:</b> solution structure of the homeobox domain of zinc finger2 homeobox protein 1b (smad interacting protein 1)
70	<a href="#">d1lfup</a>	Alignment	not modelled	99.7	28	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
71	<a href="#">d1p7jb</a>	Alignment	not modelled	99.7	43	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
72	<a href="#">c1s7eA</a>	Alignment	not modelled	99.7	20	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> hepatocyte nuclear factor 6; <b>PDBTitle:</b> solution structure of hnf-6
73	<a href="#">d1pufb</a>	Alignment	not modelled	99.7	26	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
74	<a href="#">c1x2mA</a>	Alignment	not modelled	99.7	24	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> lag1 longevity assurance homolog 6; <b>PDBTitle:</b> solution structure of the homeobox domain of mouse lag12 longevity assurance homolog 6
75	<a href="#">c3narA</a>	Alignment	not modelled	99.7	27	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> zinc fingers and homeoboxes protein 1; <b>PDBTitle:</b> crystal structure of zhx1 hd4 (zinc-fingers and homeoboxes protein 1,2 homeodomain 4)
76	<a href="#">d1mnmc</a>	Alignment	not modelled	99.7	26	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
77	<a href="#">d1s7ea1</a>	Alignment	not modelled	99.7	24	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
78	<a href="#">c2xsdC</a>	Alignment	not modelled	99.7	25	<b>PDB header:</b> transcription/dna <b>Chain:</b> C: <b>PDB Molecule:</b> pou domain, class 3, transcription factor 1; <b>PDBTitle:</b> crystal structure of the dimeric oct-6 (pou3f1) pou domain2 bound to palindromic more dna
79	<a href="#">c3nauA</a>	Alignment	not modelled	99.7	24	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> zinc fingers and homeoboxes protein 2; <b>PDBTitle:</b> crystal structure of zhx2 hd2 (zinc-fingers and homeoboxes protein 2,2 homeodomain 2)
80	<a href="#">d1le8b</a>	Alignment	not modelled	99.7	29	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
81	<a href="#">d1lfba</a>	Alignment	not modelled	99.7	27	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
						<b>Fold:</b> DNA/RNA-binding 3-helical bundle

82	<a href="#">d1yrnb_</a>	Alignment	not modelled	99.7	24	<b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
83	<a href="#">d1o4xa1</a>	Alignment	not modelled	99.7	24	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
84	<a href="#">c2e19A_</a>	Alignment	not modelled	99.7	22	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcription factor 8; <b>PDBTitle:</b> solution structure of the homeobox domain from human nil-2-2 a zinc finger protein, transcription factor 8
85	<a href="#">c2lk2A_</a>	Alignment	not modelled	99.7	18	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> homeobox protein tgif1; <b>PDBTitle:</b> solution nmr structure of homeobox domain (171-248) of human homeobox2 protein tgif1, northeast structural genomics consortium target3 hr4411b
86	<a href="#">d1akha_</a>	Alignment	not modelled	99.7	19	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
87	<a href="#">d1x2na1</a>	Alignment	not modelled	99.7	22	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
88	<a href="#">d1du6a_</a>	Alignment	not modelled	99.6	20	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
89	<a href="#">d2cqxa1</a>	Alignment	not modelled	99.6	25	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
90	<a href="#">d2r5yb1</a>	Alignment	not modelled	99.6	28	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
91	<a href="#">d1x2ma1</a>	Alignment	not modelled	99.6	26	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
92	<a href="#">d1k61a_</a>	Alignment	not modelled	99.6	25	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
93	<a href="#">d2p81a1</a>	Alignment	not modelled	99.5	43	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
94	<a href="#">c4egcA_</a>	Alignment	not modelled	99.2	24	<b>PDB header:</b> transcription/hydrolase <b>Chain:</b> A: <b>PDB Molecule:</b> maltose-binding periplasmic protein, homeobox protein six1 <b>PDBTitle:</b> crystal structure of mbp-fused human six1 bound to human eya2 eya2 domain
95	<a href="#">c2nzza_</a>	Alignment	not modelled	97.9	53	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> penetratin conjugated gas (374-394) peptide; <b>PDBTitle:</b> nmr structure analysis of the penetratin conjugated gas2 (374-394) peptide
96	<a href="#">d1wh7a_</a>	Alignment	not modelled	97.5	19	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
97	<a href="#">c1mh3A_</a>	Alignment	not modelled	97.3	18	<b>PDB header:</b> sugar binding, dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> maltose binding-a1 homeodomain protein chimera; <b>PDBTitle:</b> maltose binding-a1 homeodomain protein chimera, crystal2 form i
98	<a href="#">d1wi3a_</a>	Alignment	not modelled	97.3	31	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
99	<a href="#">c2elhA_</a>	Alignment	not modelled	91.1	19	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> cg11849-pa; <b>PDBTitle:</b> solution structure of the cenp-b n-terminal dna-binding2 domain of fruit fly distal antenna cg11849-pa
100	<a href="#">c2ys9A_</a>	Alignment	not modelled	87.5	23	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> homeobox and leucine zipper protein homez; <b>PDBTitle:</b> structure of the third homeodomain from the human homeobox2 and leucine zipper protein, homez
101	<a href="#">d1hlva1</a>	Alignment	not modelled	86.8	19	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Centromere-binding
102	<a href="#">c3bdnB_</a>	Alignment	not modelled	86.8	16	<b>PDB header:</b> transcription/dna <b>Chain:</b> B: <b>PDB Molecule:</b> lambda repressor; <b>PDBTitle:</b> crystal structure of the lambda repressor
103	<a href="#">c1hlvA_</a>	Alignment	not modelled	85.9	15	<b>PDB header:</b> dna binding protein/dna <b>Chain:</b> A: <b>PDB Molecule:</b> major centromere autoantigen b; <b>PDBTitle:</b> crystal structure of cenp-b(1-129) complexed with the cenp-2 b box dna
104	<a href="#">d1smvf2</a>	Alignment	not modelled	85.8	8	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Sigma3 and sigma4 domains of RNA polymerase sigma factors <b>Family:</b> Sigma4 domain
105	<a href="#">d1rp3a2</a>	Alignment	not modelled	85.0	11	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Sigma3 and sigma4 domains of RNA polymerase sigma factors <b>Family:</b> Sigma4 domain
106	<a href="#">c2ktgA_</a>	Alignment	not modelled	84.2	15	<b>PDB header:</b> ca-binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> calmodulin, putative; <b>PDBTitle:</b> calmodulin like protein from entamoeba histolytica: solution structure2 and calcium binding properties of a partially folded protein
107	<a href="#">d1avsa_</a>	Alianment	not modelled	83.4	12	<b>Fold:</b> EF Hand-like <b>Superfamily:</b> EF-hand

						<b>Family:</b> Calmodulin-like
108	<a href="#">c3vdoA_</a>	Alignment	not modelled	81.3	13	<b>PDB header:</b> dna binding protein/protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> rna polymerase sigma factor sigk; <b>PDBTitle:</b> structure of extra-cytoplasmic function(ecf) sigma factor sigk in2 complex with its negative regulator rska from mycobacterium3 tuberculosis
109	<a href="#">c1or7A_</a>	Alignment	not modelled	80.2	14	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> rna polymerase sigma-e factor; <b>PDBTitle:</b> crystal structure of escherichia coli sigmae with the cytoplasmic2 domain of its anti-sigma rsea
110	<a href="#">c2pmyB_</a>	Alignment	not modelled	78.2	19	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> B: <b>PDB Molecule:</b> ras and ef-hand domain-containing protein; <b>PDBTitle:</b> ef-hand domain of human rasef
111	<a href="#">d1rioA_</a>	Alignment	not modelled	77.5	16	<b>Fold:</b> lambda repressor-like DNA-binding domains <b>Superfamily:</b> lambda repressor-like DNA-binding domains <b>Family:</b> Phage repressors
112	<a href="#">d1sw8a_</a>	Alignment	not modelled	77.2	21	<b>Fold:</b> EF Hand-like <b>Superfamily:</b> EF-hand <b>Family:</b> Calmodulin-like
113	<a href="#">c3hugA_</a>	Alignment	not modelled	74.0	13	<b>PDB header:</b> transcription/membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> rna polymerase sigma factor; <b>PDBTitle:</b> crystal structure of mycobacterium tuberculosis anti-sigma factor rsla2 in complex with -35 promoter binding domain of sigl
114	<a href="#">c3mzyA_</a>	Alignment	not modelled	73.6	15	<b>PDB header:</b> rna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> rna polymerase sigma-h factor; <b>PDBTitle:</b> the crystal structure of the rna polymerase sigma-h factor from2 fusobacterium nucleatum to 2.5a
115	<a href="#">d1or7a1</a>	Alignment	not modelled	72.5	9	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Sigma3 and sigma4 domains of RNA polymerase sigma factors <b>Family:</b> Sigma4 domain
116	<a href="#">c2kgrA_</a>	Alignment	not modelled	72.2	11	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> intersectin-1; <b>PDBTitle:</b> solution structure of protein itsn1 from homo sapiens.2 northeast structural genomics consortium target hr5524a
117	<a href="#">d1l3la1</a>	Alignment	not modelled	70.3	7	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> C-terminal effector domain of the bipartite response regulators <b>Family:</b> GerE-like (LuxR/UhpA family of transcriptional regulators)
118	<a href="#">c4if4A_</a>	Alignment	not modelled	70.0	14	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> response regulator protein vvar; <b>PDBTitle:</b> crystal structure of the magnesium and berylliofluoride-activated vvar2 from staphylococcus aureus
119	<a href="#">c2lfwA_</a>	Alignment	not modelled	69.1	17	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> phyr sigma-like domain; <b>PDBTitle:</b> nmr structure of the phyrsl-nepr complex from sphingomonas sp. fr1
120	<a href="#">c2o8xA_</a>	Alignment	not modelled	67.1	12	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> probable rna polymerase sigma-c factor; <b>PDBTitle:</b> crystal structure of the "-35 element" promoter recognition domain of2 mycobacterium tuberculosis sigc