
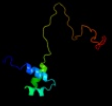


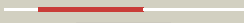








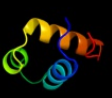
















#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c2rn7A_</a>	 Alignment		99.8	19	<b>PDB header:</b> unknown function <b>Chain:</b> A; <b>PDB Molecule:</b> is629 orfa; <b>PDBTitle:</b> nmr solution structure of tnpe protein from shigella2 flexneri. northeast structural genomics target sfr125
2	<a href="#">d2jn6a1</a>	 Alignment		99.8	21	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Cgl2762-like
3	<a href="#">c2elha_</a>	 Alignment		98.3	20	<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
4	<a href="#">c1hlvA_</a>	 Alignment		98.2	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
5	<a href="#">d1pdnc_</a>	 Alignment		98.1	12	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Paired domain
6	<a href="#">c6paxA_</a>	 Alignment		98.0	12	<b>PDB header:</b> gene regulation/dna <b>Chain:</b> A; <b>PDB Molecule:</b> homeobox protein pax-6; <b>PDBTitle:</b> crystal structure of the human pax-6 paired domain-dna2 complex reveals a general model for pax protein-dna3 interactions
7	<a href="#">d6paxa1</a>	 Alignment		98.0	12	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Paired domain
8	<a href="#">d1k78a1</a>	 Alignment		98.0	10	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Paired domain
9	<a href="#">c1u78A_</a>	 Alignment		97.9	11	<b>PDB header:</b> dna binding protein/dna <b>Chain:</b> A; <b>PDB Molecule:</b> transposable element tc3 transposase; <b>PDBTitle:</b> structure of the bipartite dna-binding domain of tc32 transposase bound to transposon dna
10	<a href="#">d1hlva1</a>	 Alignment		97.9	18	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Centromere-binding
11	<a href="#">d1bw6a_</a>	 Alignment		97.8	18	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Centromere-binding

12	<a href="#">c3hefB_</a>	Alignment		97.4	13	<b>PDB header:</b> viral protein <b>Chain:</b> B: <b>PDB Molecule:</b> gene 1 protein; <b>PDBTitle:</b> crystal structure of the bacteriophage sf6 terminase small2 subunit
13	<a href="#">c2gloA_</a>	Alignment		97.3	22	<b>PDB header:</b> transcription/dna <b>Chain:</b> A: <b>PDB Molecule:</b> brinker cg9653-pa; <b>PDBTitle:</b> solution structure of the brinker dna binding domain in2 complex with the omb enhancer
14	<a href="#">d2coba1</a>	Alignment		97.1	14	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Psq domain
15	<a href="#">c2k27A_</a>	Alignment		96.6	10	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> paired box protein pax-8; <b>PDBTitle:</b> solution structure of human pax8 paired box domain
16	<a href="#">c2w48D_</a>	Alignment		96.2	17	<b>PDB header:</b> transcription <b>Chain:</b> D: <b>PDB Molecule:</b> sorbitol operon regulator; <b>PDBTitle:</b> crystal structure of the full-length sorbitol operon2 regulator sorc from klebsiella pneumoniae
17	<a href="#">d2ao9a1</a>	Alignment		96.1	31	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Nanomeric phage protein-like
18	<a href="#">c2r0qF_</a>	Alignment		95.4	11	<b>PDB header:</b> recombination/dna <b>Chain:</b> F: <b>PDB Molecule:</b> putative transposon tn552 dna-invertase bin3; <b>PDBTitle:</b> crystal structure of a serine recombinase- dna regulatory2 complex
19	<a href="#">d1ntca_</a>	Alignment		95.3	24	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> FIS-like
20	<a href="#">d1fipa_</a>	Alignment		95.2	21	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> FIS-like
21	<a href="#">cliufA_</a>	Alignment	not modelled	95.1	8	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> centromere abp1 protein; <b>PDBTitle:</b> low resolution solution structure of the two dna-binding2 domains in schizosaccharomyces pombe abp1 protein
22	<a href="#">c1umqA_</a>	Alignment	not modelled	94.8	24	<b>PDB header:</b> dna-binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> photosynthetic apparatus regulatory protein; <b>PDBTitle:</b> solution structure and dna binding of the effector domain2 from the global regulator prra(rega) from r. sphaeroides:3 insights into dna binding specificity
23	<a href="#">d1umqa_</a>	Alignment	not modelled	94.8	24	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> FIS-like
24	<a href="#">d1etxa_</a>	Alignment	not modelled	94.8	21	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> FIS-like
25	<a href="#">c2ao9H_</a>	Alignment	not modelled	94.7	26	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> H: <b>PDB Molecule:</b> phage protein; <b>PDBTitle:</b> structural genomics, the crystal structure of a phage protein2 (phbc6a51) from bacillus cereus atcc 14579
26	<a href="#">d1ocpa_</a>	Alignment	not modelled	94.7	20	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
27	<a href="#">c2l9rA_</a>	Alignment	not modelled	94.5	18	<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
28	<a href="#">d1gt0c1</a>	Alignment	not modelled	94.2	20	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like

					<b>Family:</b> Homeodomain
29	<a href="#">c3e7ID_</a>	Alignment	not modelled	94.2	23 <b>PDB header:</b> transcription regulator <b>Chain:</b> D: <b>PDB Molecule:</b> transcriptional regulator (ntrc family); <b>PDBTitle:</b> crystal structure of sigma54 activator ntrc4's dna binding2 domain
30	<a href="#">d2ezla_</a>	Alignment	not modelled	94.2	22 <b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Recombinase DNA-binding domain
31	<a href="#">c2gm4B_</a>	Alignment	not modelled	94.0	13 <b>PDB header:</b> recombination, dna <b>Chain:</b> B: <b>PDB Molecule:</b> transposon gamma-delta resolvase; <b>PDBTitle:</b> an activated, tetrameric gamma-delta resolvase: hin chimaera bound to2 cleaved dna
32	<a href="#">c2da1A_</a>	Alignment	not modelled	93.7	16 <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)
33	<a href="#">d2cuea1</a>	Alignment	not modelled	93.7	21 <b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
34	<a href="#">d1rzsa_</a>	Alignment	not modelled	93.7	14 <b>Fold:</b> lambda repressor-like DNA-binding domains <b>Superfamily:</b> lambda repressor-like DNA-binding domains <b>Family:</b> Phage repressors
35	<a href="#">c2da2A_</a>	Alignment	not modelled	93.7	11 <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)
36	<a href="#">c2dmuA_</a>	Alignment	not modelled	93.6	23 <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
37	<a href="#">c2dmsA_</a>	Alignment	not modelled	93.6	18 <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
38	<a href="#">d2eloal</a>	Alignment	not modelled	93.5	27 <b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
39	<a href="#">d2craa1</a>	Alignment	not modelled	93.1	21 <b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
40	<a href="#">c2pbxB_</a>	Alignment	not modelled	93.1	35 <b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> hemagglutinin/protease regulatory protein; <b>PDBTitle:</b> vibrio cholerae hapr
41	<a href="#">c1or7A_</a>	Alignment	not modelled	93.0	5 <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
42	<a href="#">d1b8ia_</a>	Alignment	not modelled	93.0	24 <b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
43	<a href="#">d1ahdp_</a>	Alignment	not modelled	93.0	27 <b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
44	<a href="#">d1or7a1</a>	Alignment	not modelled	92.8	5 <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
45	<a href="#">d1bw5a_</a>	Alignment	not modelled	92.8	7 <b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
46	<a href="#">c3a01A_</a>	Alignment	not modelled	92.8	27 <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
47	<a href="#">c3mzyA_</a>	Alignment	not modelled	92.7	20 <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
48	<a href="#">d1wh7a_</a>	Alignment	not modelled	92.7	21 <b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
49	<a href="#">c2djnA_</a>	Alignment	not modelled	92.7	23 <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
50	<a href="#">d1ig7a_</a>	Alignment	not modelled	92.7	22 <b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
51	<a href="#">d1zq3p1</a>	Alignment	not modelled	92.7	14 <b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
52	<a href="#">d1p7ia_</a>	Alignment	not modelled	92.6	25 <b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
53	<a href="#">c2k40A_</a>	Alignment	not modelled	92.6	17 <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 hexs-1 homeodomain double mutant r311/e42l
54	<a href="#">d2hddb_</a>	Alignment	not modelled	92.5	27 <b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain

55	<a href="#">c2o8xA</a>	Alignment	not modelled	92.5	10	<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
56	<a href="#">d1fjla</a>	Alignment	not modelled	92.5	21	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
57	<a href="#">d1wh5a</a>	Alignment	not modelled	92.4	27	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
58	<a href="#">c3bd1B</a>	Alignment	not modelled	92.3	30	<b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> cro protein; <b>PDBTitle:</b> structure of the cro protein from putative prophage element xfaso 1 in2 xylella fastidiosa strain ann-1
59	<a href="#">d1vnda</a>	Alignment	not modelled	92.3	16	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
60	<a href="#">d1octc1</a>	Alignment	not modelled	92.3	20	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
61	<a href="#">d1f43a</a>	Alignment	not modelled	92.2	20	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
62	<a href="#">d1rp3a2</a>	Alignment	not modelled	92.1	10	<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
63	<a href="#">d1nk3p</a>	Alignment	not modelled	92.1	16	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
64	<a href="#">d1fjlb</a>	Alignment	not modelled	92.1	22	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
65	<a href="#">d1etob</a>	Alignment	not modelled	92.0	18	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> FIS-like
66	<a href="#">d1ftza</a>	Alignment	not modelled	92.0	28	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
67	<a href="#">c1s7eA</a>	Alignment	not modelled	91.9	23	<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
68	<a href="#">d1au7a1</a>	Alignment	not modelled	91.8	20	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
69	<a href="#">d1ztra1</a>	Alignment	not modelled	91.7	29	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
70	<a href="#">d1xsva</a>	Alignment	not modelled	91.6	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> YlxM/p13-like
71	<a href="#">d1ftta</a>	Alignment	not modelled	91.6	20	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
72	<a href="#">d1yz8p1</a>	Alignment	not modelled	91.6	15	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
73	<a href="#">d1pufa</a>	Alignment	not modelled	91.5	25	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
74	<a href="#">c3d1nK</a>	Alignment	not modelled	91.5	23	<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
75	<a href="#">d1jgga</a>	Alignment	not modelled	91.4	20	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
76	<a href="#">c2dmtA</a>	Alignment	not modelled	91.4	19	<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
77	<a href="#">c2vz4A</a>	Alignment	not modelled	91.4	13	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> hth-type transcriptional activator tipa; <b>PDBTitle:</b> the n-terminal domain of merr-like protein tipal bound to2 promoter dna
78	<a href="#">d1zs4a1</a>	Alignment	not modelled	91.3	24	<b>Fold:</b> lambda repressor-like DNA-binding domains <b>Superfamily:</b> lambda repressor-like DNA-binding domains <b>Family:</b> Bacteriophage CII protein
79	<a href="#">d1ijwc</a>	Alignment	not modelled	91.1	16	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Recombinase DNA-binding domain
80	<a href="#">c2e7xA</a>	Alignment	not modelled	90.7	23	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> 150aa long hypothetical transcriptional regulator; <b>PDBTitle:</b> structure of the Irp/asnc like transcriptional regulator from2 sulfobolus tokodaii 7 complexed with its cognate ligand
81	<a href="#">c2ds4A</a>	Alignment	not modelled	90.7	20	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> hypothetical protein dkfzp686k21156;

81	<a href="#">c2da4A</a>	Alignment	not modelled	90.7	20	<b>PDBTitle:</b> solution structure of the homeobox domain of the2 hypothetical protein, dkfzp686k21156 <b>PDB header:</b> transcription/membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> rna polymerase sigma factor;
82	<a href="#">c3hugA</a>	Alignment	not modelled	90.6	18	<b>PDBTitle:</b> crystal structure of mycobacterium tuberculosis anti-sigma factor rslA2 in complex with -35 promoter binding domain of sigl <b>PDB header:</b> transcription regulation <b>Chain:</b> A: <b>PDB Molecule:</b> homeobox protein nanog;
83	<a href="#">c2kt0A</a>	Alignment	not modelled	90.4	18	<b>PDBTitle:</b> solution structure of human stem cell transcription factor nanog2 homeodomain fragment <b>PDB header:</b> transcription <b>Chain:</b> H: <b>PDB Molecule:</b> transcriptional regulator, Irp/asnc family;
84	<a href="#">c2p6tH</a>	Alignment	not modelled	90.4	10	<b>PDBTitle:</b> crystal structure of transcriptional regulator nmb0573 and l-leucine2 complex from neisseria meningitidis <b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> regulatory protein asnc;
85	<a href="#">c2cg4B</a>	Alignment	not modelled	90.4	8	<b>PDBTitle:</b> structure of e.coli asnc <b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> zinc fingers and homeoboxes protein 3;
86	<a href="#">c2da5A</a>	Alignment	not modelled	90.3	14	<b>PDBTitle:</b> solution structure of the second homeobox domain of zinc2 fingers and homeoboxes protein 3 (triple homeobox 13 protein) <b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Recombinase DNA-binding domain
87	<a href="#">d1hcra</a>	Alignment	not modelled	90.3	16	<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	90.1	14	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
89	<a href="#">c3i4pA</a>	Alignment	not modelled	90.0	12	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, asnc family; <b>PDBTitle:</b> crystal structure of asnc family transcriptional regulator from2 agrobacterium tumefaciens
90	<a href="#">d2ecba1</a>	Alignment	not modelled	90.0	9	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
91	<a href="#">c2cfxD</a>	Alignment	not modelled	89.9	5	<b>PDB header:</b> transcription <b>Chain:</b> D: <b>PDB Molecule:</b> hth-type transcriptional regulator lrpc; <b>PDBTitle:</b> structure of b.subtilis lrpc
92	<a href="#">c2ia0A</a>	Alignment	not modelled	89.9	18	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> putative hth-type transcriptional regulator pf0864; <b>PDBTitle:</b> transcriptional regulatory protein pf0864 from pyrococcus furiosus a2 member of the asnc family (pf0864)
93	<a href="#">d1hdpa</a>	Alignment	not modelled	89.9	17	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
94	<a href="#">c2dmqA</a>	Alignment	not modelled	89.8	16	<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
95	<a href="#">d2ecca1</a>	Alignment	not modelled	89.7	11	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
96	<a href="#">c3l1pA</a>	Alignment	not modelled	89.6	20	<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 <b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> alpha-fetoprotein enhancer binding protein;
97	<a href="#">c2da3A</a>	Alignment	not modelled	89.5	18	<b>PDBTitle:</b> solution structure of the third homeobox domain of at-2 binding transcription factor 1 (atbf1) <b>PDB header:</b> dna-binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulatory protein;
98	<a href="#">c2vbzA</a>	Alignment	not modelled	89.5	10	<b>PDBTitle:</b> feast or famine regulatory protein (rv3291c)from m.2 tuberculosis complexed with l-tryptophan <b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
99	<a href="#">d1b72a</a>	Alignment	not modelled	89.4	24	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
100	<a href="#">c2kpiA</a>	Alignment	not modelled	89.4	9	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> sos-response transcriptional repressor, lexa; <b>PDBTitle:</b> solution structure of protein sos-response transcriptional2 repressor, lexa from eubacterium rectale. northeast3 structural genomics consortium target err9a
101	<a href="#">d1q06a</a>	Alignment	not modelled	88.9	13	<b>Fold:</b> Putative DNA-binding domain <b>Superfamily:</b> Putative DNA-binding domain <b>Family:</b> DNA-binding N-terminal domain of transcription activators
102	<a href="#">c2krfB</a>	Alignment	not modelled	88.6	13	<b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> transcriptional regulatory protein coma; <b>PDBTitle:</b> nmr solution structure of the dna binding domain of competence protein2 a
103	<a href="#">c1hf0A</a>	Alignment	not modelled	88.5	20	<b>PDB header:</b> transcription factor <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
104	<a href="#">d1ulya</a>	Alignment	not modelled	88.4	17	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> "Winged helix" DNA-binding domain <b>Family:</b> Hypothetical protein PH1932
105	<a href="#">d1pufb</a>	Alignment	not modelled	88.3	13	<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

106	<a href="#">d1biaa1</a>	Alignment	not modelled	88.2	5	<b>Superfamily:</b> "Winged helix" DNA-binding domain <b>Family:</b> Biotin repressor-like
107	<a href="#">d2esna1</a>	Alignment	not modelled	88.1	22	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> "Winged helix" DNA-binding domain <b>Family:</b> LysR-like transcriptional regulators
108	<a href="#">d1e3oc1</a>	Alignment	not modelled	88.1	18	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
109	<a href="#">c3op9A_</a>	Alignment	not modelled	88.0	16	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> pli0006 protein; <b>PDBTitle:</b> crystal structure of transcriptional regulator from listeria innocua
110	<a href="#">c1rp3G_</a>	Alignment	not modelled	87.7	10	<b>PDB header:</b> transcription <b>Chain:</b> G: <b>PDB Molecule:</b> rna polymerase sigma factor sigma-28 (fliA); <b>PDBTitle:</b> cocrystal structure of the flagellar sigma/anti-sigma2 complex, sigma-28/flgm
111	<a href="#">c3e7qB_</a>	Alignment	not modelled	87.4	4	<b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> transcriptional regulator; <b>PDBTitle:</b> the crystal structure of the putative transcriptional regulator from2 pseudomonas aeruginosa pao1
112	<a href="#">c2l4aA_</a>	Alignment	not modelled	87.0	13	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> leucine responsive regulatory protein; <b>PDBTitle:</b> nmr structure of the dna-binding domain of e.coli lrp
113	<a href="#">d1adra_</a>	Alignment	not modelled	87.0	17	<b>Fold:</b> lambda repressor-like DNA-binding domains <b>Superfamily:</b> lambda repressor-like DNA-binding domains <b>Family:</b> Phage repressors
114	<a href="#">d1x2na1</a>	Alignment	not modelled	86.6	15	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
115	<a href="#">c1ilgA_</a>	Alignment	not modelled	86.6	18	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator lrpA; <b>PDBTitle:</b> crystal structure of the lrp-like transcriptional regulator from the2 archaeon pyrococcus furiosus
116	<a href="#">d2cg4a1</a>	Alignment	not modelled	86.6	8	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> "Winged helix" DNA-binding domain <b>Family:</b> Lrp/AsnC-like transcriptional regulator N-terminal domain
117	<a href="#">d1s7ea1</a>	Alignment	not modelled	86.5	18	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> Homeodomain-like <b>Family:</b> Homeodomain
118	<a href="#">c2dn0A_</a>	Alignment	not modelled	86.2	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
119	<a href="#">d2oa4a1</a>	Alignment	not modelled	86.2	8	<b>Fold:</b> DNA/RNA-binding 3-helical bundle <b>Superfamily:</b> TrpR-like <b>Family:</b> SPO1678-like
120	<a href="#">c2jmlA_</a>	Alignment	not modelled	86.1	15	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> dna binding domain/transcriptional regulator; <b>PDBTitle:</b> solution structure of the n-terminal domain of cara repressor