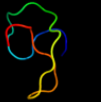
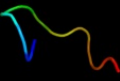
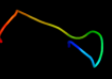
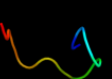


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1lv3a_	 Alignment		54.7	41	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Hypothetical zinc finger protein YacG
2	d2jw6a1	 Alignment		52.6	38	Fold: HIT/MYND zinc finger-like Superfamily: HIT/MYND zinc finger-like Family: MYND zinc finger
3	c2odda_	 Alignment		47.8	33	PDB header: metal binding protein Chain: A: PDB Molecule: protein cbfa2t1; PDBTitle: solution structure of the mynd domain from aml1-eto2 complexed with smrt, a corepressor
4	d2d8va1	 Alignment		43.5	53	Fold: B-box zinc-binding domain Superfamily: B-box zinc-binding domain Family: B-box zinc-binding domain
5	c2jw6A_	 Alignment		42.6	36	PDB header: transcription Chain: A: PDB Molecule: deformed epidermal autoregulatory factor 1 homolog; PDBTitle: solution structure of the deaf1 mynd domain
6	d2dj8a1	 Alignment		40.4	31	Fold: HIT/MYND zinc finger-like Superfamily: HIT/MYND zinc finger-like Family: MYND zinc finger
7	c2yqqa_	 Alignment		39.9	30	PDB header: protein binding Chain: A: PDB Molecule: zinc finger hit domain-containing protein 3; PDBTitle: solution structure of the zf-hit domain in zinc finger hit2 domain-containing protein 3 (trip-3)
8	c2dj8A_	 Alignment		39.6	31	PDB header: metal binding protein Chain: A: PDB Molecule: protein cbfa2t1; PDBTitle: solution structure of zf-mynd domain of protein cbfa2t2 (protein mtg8)
9	d2dana1	 Alignment		38.2	32	Fold: HIT/MYND zinc finger-like Superfamily: HIT/MYND zinc finger-like Family: MYND zinc finger
10	d2d8qa1	 Alignment		32.3	29	Fold: HIT/MYND zinc finger-like Superfamily: HIT/MYND zinc finger-like Family: MYND zinc finger
11	c2danA_	 Alignment		31.8	32	PDB header: metal binding protein Chain: A: PDB Molecule: zinc finger mynd domain containing protein 10; PDBTitle: the solution structure of the mynd domain (leu384-cys430)2 of human zinc finger mynd domain containing protein 10

12	d1u5ka2	Alignment		29.4	21	Fold: ArfGap/RecO-like zinc finger Superfamily: ArfGap/RecO-like zinc finger Family: RecO C-terminal domain-like
13	c3o37D_	Alignment		27.4	33	PDB header: transcription/protein binding Chain: D: PDB Molecule: transcription intermediary factor 1-alpha; PDBTitle: crystal structure of trim24 phd-bromo complexed with h3(1-10)k42 peptide
14	c3q8dB_	Alignment		25.6	29	PDB header: dna binding protein Chain: B: PDB Molecule: dna repair protein reco; PDBTitle: e. coli reco complex with ssb c-terminus
15	c1u5kA_	Alignment		25.3	13	PDB header: recombination,replication Chain: A: PDB Molecule: hypothetical protein; PDBTitle: recombinational repair protein reco
16	d1hcqa_	Alignment		22.9	29	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
17	d1libia2	Alignment		18.2	71	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
18	d2nlla_	Alignment		17.0	46	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
19	d1hraa_	Alignment		14.0	31	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
20	c2a66A_	Alignment		13.0	36	PDB header: transcription/dna Chain: A: PDB Molecule: orphan nuclear receptor nr5a2; PDBTitle: human liver receptor homologue dna-binding domain (hlrh-12 dbd) in complex with dsdna from the hcyp7a1 promoter
21	d1dsza_	Alignment	not modelled	12.4	31	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
22	d1cita_	Alignment	not modelled	12.4	31	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
23	c2gb5B_	Alignment	not modelled	12.2	32	PDB header: hydrolase Chain: B: PDB Molecule: nadh pyrophosphatase; PDBTitle: crystal structure of nadh pyrophosphatase (ec 3.6.1.22) (1790429) from2 escherichia coli k12 at 2.30 a resolution
24	d2hana1	Alignment	not modelled	12.2	46	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
25	c3u5pF_	Alignment	not modelled	11.8	33	PDB header: transcription Chain: F: PDB Molecule: e3 ubiquitin-protein ligase trim33; PDBTitle: crystal structure of the complex of trim33 phd-bromo and h3(1-28)2 k9me3k14ack18ack23ac histone peptide
26	d2cona1	Alignment	not modelled	11.7	38	Fold: Rubredoxin-like Superfamily: NOB1 zinc finger-like Family: NOB1 zinc finger-like
27	c3cbbA_	Alignment	not modelled	11.3	54	PDB header: transcription/dna Chain: A: PDB Molecule: hepatocyte nuclear factor 4-alpha, dna binding PDBTitle: crystal structure of hepatocyte nuclear factor 4alpha in2 complex with dna: diabetes gene product
28	d1r4ra_	Alignment	not modelled	11.1	31	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
						PDB header: glycoprotein

29	c1brvA_	Alignment	not modelled	11.0	43	Chain: A: PDB Molecule: protein g; PDBTitle: solution nmr structure of the immunodominant region of2 protein g of bovine respiratory syncytial virus, 483 structures
30	d1glua_	Alignment	not modelled	10.9	31	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
31	c1r4iA_	Alignment	not modelled	10.8	46	PDB header: transcription/dna Chain: A: PDB Molecule: androgen receptor; PDBTitle: crystal structure of androgen receptor dna-binding domain2 bound to a direct repeat response element
32	d1r4ia_	Alignment	not modelled	10.8	46	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
33	d1lo1a_	Alignment	not modelled	10.6	36	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
34	d1p91a_	Alignment	not modelled	10.5	27	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: rRNA methyltransferase RImA
35	d1r0na_	Alignment	not modelled	10.5	46	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
36	d2hanb1	Alignment	not modelled	10.0	33	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
37	c1r0nB_	Alignment	not modelled	9.8	33	PDB header: transcription/dna Chain: B: PDB Molecule: ecdysone receptor; PDBTitle: crystal structure of heterodimeric ecdysone receptor dna2 binding complex
38	c4acrB_	Alignment	not modelled	9.6	31	PDB header: membrane protein Chain: B: PDB Molecule: glypican-1; PDBTitle: crystal structure of n-glycosylated, c-terminally truncated2 human glypican-1
39	d1a6ya_	Alignment	not modelled	9.6	36	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
40	d1lata_	Alignment	not modelled	9.5	31	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
41	d1r0oa_	Alignment	not modelled	9.4	46	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
42	d1kb2a_	Alignment	not modelled	9.4	36	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
43	d2jz6a1	Alignment	not modelled	9.0	55	Fold: L28p-like Superfamily: L28p-like Family: Ribosomal protein L28
44	d2nllb_	Alignment	not modelled	8.9	46	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
45	c1zr9A_	Alignment	not modelled	8.9	36	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 593; PDBTitle: solution structure of a human c2h2-type zinc finger protein
46	d1zr9a1	Alignment	not modelled	8.9	36	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: HkH motif-containing C2H2 finger
47	d1dszb_	Alignment	not modelled	8.8	46	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
48	c1t60R_	Alignment	not modelled	8.6	25	PDB header: structural protein Chain: R: PDB Molecule: type iv collagen; PDBTitle: crystal structure of type iv collagen nc1 domain from2 bovine lens capsule
49	d1kb6b_	Alignment	not modelled	8.6	38	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
50	c2hepA_	Alignment	not modelled	8.5	40	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: upf0291 protein ynzC; PDBTitle: solution nmr structure of the upf0291 protein ynzC from2 bacillus subtilis. northeast structural genomics target3 sr384.
51	d2hepa1	Alignment	not modelled	8.5	40	Fold: Long alpha-hairpin Superfamily: YnzC-like Family: YznC-like
52	c2jl81_	Alignment	not modelled	8.4	45	PDB header: ribosome Chain: 1: PDB Molecule: 50s ribosomal protein l28; PDBTitle: insights into translational termination from the structure2 of rf2 bound to the ribosome (part 4 of 4).3 this file contains the 50s subunit.
53	c3lqiA_	Alignment	not modelled	8.2	18	PDB header: transferase Chain: A: PDB Molecule: ml11 phd3-bromo; PDBTitle: crystal structure of ml11 phd3-bromo complexed with h3(1-9)k4me22 peptide
54	d1ynwa1	Alignment	not modelled	8.1	38	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
55	c2eb1A_	Alignment	not modelled	7.6	38	PDB header: transcription Chain: A: PDB Molecule: coup transcription factor 1; PDBTitle: solution structure of the zinc finger, c4-type domain of2 human coup transcription factor 1

56	c2wg7B_	Alignment	not modelled	7.4	55	PDB header: hydrolase Chain: B: PDB Molecule: putative phospholipase a2; PDBTitle: structure of oryza sativa (rice) pla2
57	d1dx5i3	Alignment	not modelled	7.4	57	Fold: Knottins (small inhibitors, toxins, lectins) Superfamily: EGF/Laminin Family: EGF-type module
58	c2ybfB_	Alignment	not modelled	7.4	22	PDB header: ligase Chain: B: PDB Molecule: e3 ubiquitin-protein ligase rad18; PDBTitle: complex of rad18 (rad6 binding domain) with rad6b
59	d2dara1	Alignment	not modelled	7.3	57	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
60	c2q8aA_	Alignment	not modelled	7.1	29	PDB header: immune system Chain: A: PDB Molecule: apical membrane antigen 1; PDBTitle: structure of the malaria antigen ama1 in complex with a growth-2 inhibitory antibody
61	c3bboY_	Alignment	not modelled	7.1	36	PDB header: ribosome Chain: Y: PDB Molecule: ribosomal protein l28; PDBTitle: homology model for the spinach chloroplast 50s subunit fitted to 9.4a2 cryo-em map of the 70s chlororibosome
62	d2yt9a2	Alignment	not modelled	7.0	71	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
63	d2qamz1	Alignment	not modelled	7.0	55	Fold: L28p-like Superfamily: L28p-like Family: Ribosomal protein L28
64	d2cora1	Alignment	not modelled	6.9	64	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
65	c1t60W_	Alignment	not modelled	6.7	25	PDB header: structural protein Chain: W: PDB Molecule: type iv collagen; PDBTitle: crystal structure of type iv collagen nc1 domain from2 bovine lens capsule
66	c2k0aA_	Alignment	not modelled	6.7	32	PDB header: metal binding protein Chain: A: PDB Molecule: pre-mrna-splicing factor rds3; PDBTitle: 1h, 15n and 13c chemical shift assignments for rds3 protein
67	c3cngC_	Alignment	not modelled	6.7	21	PDB header: hydrolase Chain: C: PDB Molecule: nudix hydrolase; PDBTitle: crystal structure of nudix hydrolase from nitrosomonas europaea
68	c2x2zE_	Alignment	not modelled	6.7	35	PDB header: membrane protein Chain: E: PDB Molecule: apical membrane antigen 1, putative; PDBTitle: crystal structure ama1 from toxoplasma gondii
69	d1xx6a2	Alignment	not modelled	6.5	56	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Type II thymidine kinase zinc finger
70	c2envA_	Alignment	not modelled	6.5	38	PDB header: transcription Chain: A: PDB Molecule: peroxisome proliferator-activated receptor delta; PDBTitle: solution sturcture of the c4-type zinc finger domain from2 human peroxisome proliferator-activated receptor delta
71	d1wila_	Alignment	not modelled	6.4	33	Fold: FYVE/PHD zinc finger Superfamily: FYVE/PHD zinc finger Family: variant PHD-like domain
72	c3qvvA_	Alignment	not modelled	5.7	25	PDB header: transferase Chain: A: PDB Molecule: set and mynd domain-containing protein 2; PDBTitle: crystal structure of histone lysine methyltransferase smyd2 in complex2 with the cofactor product adohcy
73	d1oj8a_	Alignment	not modelled	5.4	23	Fold: RNase A-like Superfamily: RNase A-like Family: Ribonuclease A-like
74	d1t0za_	Alignment	not modelled	5.3	50	Fold: Knottins (small inhibitors, toxins, lectins) Superfamily: Scorpion toxin-like Family: Long-chain scorpion toxins
75	c2jmiA_	Alignment	not modelled	5.1	29	PDB header: protein binding Chain: A: PDB Molecule: protein yng1; PDBTitle: nmr solution structure of phd finger fragment of yeast yng12 protein in free state
76	c2iybG_	Alignment	not modelled	5.1	50	PDB header: metal-binding Chain: G: PDB Molecule: testin; PDBTitle: structure of complex between the 3rd lim domain of tes and2 the evh1 domain of mena
77	d1ryqa_	Alignment	not modelled	5.1	35	Fold: Rubredoxin-like Superfamily: RNA polymerase subunits Family: RpoE2-like
78	d1hc7a3	Alignment	not modelled	5.0	57	Fold: IF3-like Superfamily: C-terminal domain of ProRS Family: C-terminal domain of ProRS