
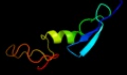
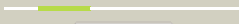

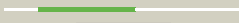
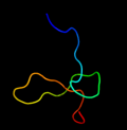

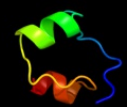
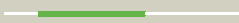






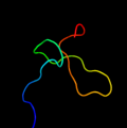



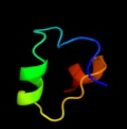

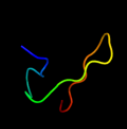
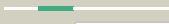


















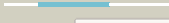




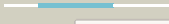



#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1lv3a_	 Alignment		100.0	100	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Hypothetical zinc finger protein YacG
2	d2cu8a1	 Alignment		68.0	44	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
3	d2dasa1	 Alignment		57.5	26	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: TRASH domain
4	d2d8qa1	 Alignment		55.1	25	Fold: HIT/MYND zinc finger-like Superfamily: HIT/MYND zinc finger-like Family: MYND zinc finger
5	c2yqqa_	 Alignment		54.7	24	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)
6	c3n71A_	 Alignment		49.9	24	PDB header: transcription Chain: A: PDB Molecule: histone lysine methyltransferase smyd1; PDBTitle: crystal structure of cardiac specific histone methyltransferase smyd1
7	c1yuzB_	 Alignment		49.8	27	PDB header: oxidoreductase Chain: B: PDB Molecule: nigerythrin; PDBTitle: partially reduced state of nigerythrin
8	c2dasA_	 Alignment		48.9	26	PDB header: metal transport Chain: A: PDB Molecule: zinc finger mym-type protein 5; PDBTitle: solution structure of trash domain of zinc finger mym-type2 protein 5
9	c3mekA_	 Alignment		47.8	30	PDB header: transferase Chain: A: PDB Molecule: set and mynd domain-containing protein 3; PDBTitle: crystal structure of human histone-lysine n-2 methyltransferase smyd3 in complex with s-adenosyl-l-3 methionine
10	c2jw6A_	 Alignment		45.1	33	PDB header: transcription Chain: A: PDB Molecule: deformed epidermal autoregulatory factor 1 homolog; PDBTitle: solution structure of the deaf1 mynd domain
11	c2jrpA_	 Alignment		43.3	17	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative cytoplasmic protein; PDBTitle: solution nmr structure of yfgj from salmonella typhimurium2 modeled with two zn+2 bound, northeast structural genomics3 consortium target str86

12	c2k5cA_	 Alignment		41.5	45	PDB header: metal binding protein Chain: A: PDB Molecule: uncharacterized protein pf0385; PDBTitle: nmr structure for pf0385
13	c2dj8A_	 Alignment		40.4	29	PDB header: metal binding protein Chain: A: PDB Molecule: protein cbfa2t1; PDBTitle: solution structure of zf-mynd domain of protein cbfa2t12 (protein mtg8)
14	d2jw6a1	 Alignment		39.5	33	Fold: HIT/MYND zinc finger-like Superfamily: HIT/MYND zinc finger-like Family: MYND zinc finger
15	c2odda_	 Alignment		38.9	29	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
16	d1imla1	 Alignment		38.7	46	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
17	d2jneal	 Alignment		35.6	17	Fold: Rubredoxin-like Superfamily: Yfgj-like Family: Yfgj-like
18	c2jneA_	 Alignment		35.6	17	PDB header: metal binding protein Chain: A: PDB Molecule: hypothetical protein yfgj; PDBTitle: nmr structure of e.coli yfgj modelled with two zn+2 bound.2 northeast structural genomics consortium target er317.
19	d2dlqa4	 Alignment		35.5	60	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
20	c2jr6A_	 Alignment		35.1	14	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: upf0434 protein nma0874; PDBTitle: solution structure of upf0434 protein nma0874. northeast structural2 genomics target mr32
21	d2dj8a1	 Alignment	not modelled	33.8	28	Fold: HIT/MYND zinc finger-like Superfamily: HIT/MYND zinc finger-like Family: MYND zinc finger
22	c2js4A_	 Alignment	not modelled	32.1	17	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: upf0434 protein bb2007; PDBTitle: solution nmr structure of bordetella bronchiseptica protein2 bb2007. northeast structural genomics consortium target3 bor54
23	c2danA_	 Alignment	not modelled	32.1	25	PDB header: metal binding protein Chain: A: PDB Molecule: zinc finger mynd domain containing protein 10; PDBTitle: the solution structure of the mynd domain (Ileu384-cys430)2 of human zinc finger mynd domain containing protein 10
24	d2dana1	 Alignment	not modelled	31.9	25	Fold: HIT/MYND zinc finger-like Superfamily: HIT/MYND zinc finger-like Family: MYND zinc finger
25	d1fbva4	 Alignment	not modelled	31.5	30	Fold: RING/U-box Superfamily: RING/U-box Family: RING finger domain, C3HC4
26	d2jnva1	 Alignment	not modelled	31.1	13	Fold: Trm112p-like Superfamily: Trm112p-like Family: Trm112p-like
27	c2ecga_	 Alignment	not modelled	30.0	28	PDB header: apoptosis Chain: A: PDB Molecule: baculoviral iap repeat-containing protein 4; PDBTitle: solution structure of the ring domain of the baculoviral2 iap repeat-containing protein 4 from homo sapiens
28	c2ecta_	 Alignment	not modelled	29.9	25	PDB header: metal binding protein Chain: A: PDB Molecule: ring finger protein 126; PDBTitle: solution structure of the zinc finger, c3hc4 type (ring2

						finger) domain of ring finger protein 126
29	c3qvvA	Alignment	not modelled	28.8	35	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
30	c3u50C	Alignment	not modelled	28.7	32	PDB header: dna binding protein Chain: C: PDB Molecule: telomerase-associated protein 82; PDBTitle: crystal structure of the tetrahymena telomerase processivity factor2 teb1 ob-c
31	c3nw0A	Alignment	not modelled	28.4	19	PDB header: metal binding protein Chain: A: PDB Molecule: non-structural maintenance of chromosomes element 1 PDBTitle: crystal structure of mageg1 and nse1 complex
32	c2l8eA	Alignment	not modelled	27.4	26	PDB header: dna binding protein Chain: A: PDB Molecule: polyhomeotic-like protein 1; PDBTitle: solution nmr structure of fcs domain of human polyhomeotic homolog 12 (hph1)
33	c2hl7A	Alignment	not modelled	26.9	22	PDB header: oxidoreductase Chain: A: PDB Molecule: cytochrome c-type biogenesis protein ccmh; PDBTitle: crystal structure of the periplasmic domain of ccmh from pseudomonas2 aeruginosa
34	d2pw4a1	Alignment	not modelled	26.2	26	Fold: Jann2411-like Superfamily: Jann2411-like Family: Jann2411-like
35	d1p91a	Alignment	not modelled	25.6	17	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: rRNA methyltransferase RImA
36	d1imla2	Alignment	not modelled	24.0	35	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
37	d2hf1a1	Alignment	not modelled	23.7	24	Fold: Trm112p-like Superfamily: Trm112p-like Family: Trm112p-like
38	c3oxgA	Alignment	not modelled	23.5	30	PDB header: transferase Chain: A: PDB Molecule: set and mynd domain-containing protein 3; PDBTitle: human lysine methyltransferase smyd3 in complex with adohcy (form iii)
39	d1bora	Alignment	not modelled	23.5	27	Fold: RING/U-box Superfamily: RING/U-box Family: RING finger domain, C3HC4
40	d2pk7a1	Alignment	not modelled	23.3	18	Fold: Trm112p-like Superfamily: Trm112p-like Family: Trm112p-like
41	d1yuza2	Alignment	not modelled	22.9	27	Fold: Rubredoxin-like Superfamily: Rubredoxin-like Family: Rubredoxin
42	c2kw0A	Alignment	not modelled	22.9	22	PDB header: oxidoreductase Chain: A: PDB Molecule: ccmh protein; PDBTitle: solution structure of n-terminal domain of ccmh from escherichia.coli
43	d1cta1	Alignment	not modelled	22.3	40	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
44	c2pptA	Alignment	not modelled	21.2	36	PDB header: oxidoreductase Chain: A: PDB Molecule: thioredoxin-2; PDBTitle: crystal structure of thioredoxin-2
45	d1libia1	Alignment	not modelled	19.9	36	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
46	c3iz6X	Alignment	not modelled	19.8	21	PDB header: ribosome Chain: X: PDB Molecule: 40s ribosomal protein s27 (s27e); PDBTitle: localization of the small subunit ribosomal proteins into a 5.5 a2 cryo-em map of triticum aestivum translating 80s ribosome
47	c2hr5B	Alignment	not modelled	19.3	36	PDB header: metal binding protein Chain: B: PDB Molecule: rubrerythrin; PDBTitle: pf1283- rubrerythrin from pyrococcus furiosus iron bound form
48	c3eb6A	Alignment	not modelled	18.8	39	PDB header: apoptosis, ligase Chain: A: PDB Molecule: baculoviral iap repeat-containing protein 3; PDBTitle: structure of the ciap2 ring domain bound to ubch5b
49	c2w0tA	Alignment	not modelled	18.6	37	PDB header: transcription Chain: A: PDB Molecule: lethal(3)malignant brain tumor-like 2 protein; PDBTitle: solution structure of the fcs zinc finger domain of human2 lmb12
50	c2ct2A	Alignment	not modelled	18.4	26	PDB header: ligase Chain: A: PDB Molecule: tripartite motif protein 32; PDBTitle: solution structure of the ring domain of the tripartite2 motif protein 32
51	d1ur6b	Alignment	not modelled	18.1	25	Fold: RING/U-box Superfamily: RING/U-box Family: RING finger domain, C3HC4
52	c2kpiA	Alignment	not modelled	17.9	20	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized protein sco3027; PDBTitle: solution nmr structure of streptomyces coelicolor sco30272 modeled with zn+2 bound, northeast structural genomics3 consortium target rr58
53	c1dvbA	Alignment	not modelled	17.6	27	PDB header: electron transport Chain: A: PDB Molecule: rubrerythrin; PDBTitle: rubrerythrin

54	d1pfta_	Alignment	not modelled	17.4	27	Fold: Rubredoxin-like Superfamily: Zinc beta-ribbon Family: Transcriptional factor domain
55	d1g25a_	Alignment	not modelled	16.9	47	Fold: RING/U-box Superfamily: RING/U-box Family: RING finger domain, C3HC4
56	c2ecwA_	Alignment	not modelled	16.7	26	PDB header: apoptosis Chain: A: PDB Molecule: tripartite motif-containing protein 30; PDBTitle: solution structure of the zinc finger, c3hc4 type (ring2 finger) domain tripartite motif protein 30
57	c3mhsC_	Alignment	not modelled	16.5	29	PDB header: hydrolase/transcription regulator/protei Chain: C: PDB Molecule: saga-associated factor 11; PDBTitle: structure of the saga ubp8/sgf11/sus1/sgf73 dub module bound to2 ubiquitin aldehyde
58	c3izbX_	Alignment	not modelled	15.9	31	PDB header: ribosome Chain: X: PDB Molecule: 40s ribosomal protein rps27 (s27e); PDBTitle: localization of the small subunit ribosomal proteins into a 6.1 a2 cryo-em map of saccharomyces cerevisiae translating 80s ribosome
59	d1v87a_	Alignment	not modelled	15.8	21	Fold: RING/U-box Superfamily: RING/U-box Family: RING finger domain, C3HC4
60	c3floD_	Alignment	not modelled	15.7	31	PDB header: transferase Chain: D: PDB Molecule: dna polymerase alpha catalytic subunit a; PDBTitle: crystal structure of the carboxyl-terminal domain of yeast2 dna polymerase alpha in complex with its b subunit
61	c2djbA_	Alignment	not modelled	15.6	40	PDB header: gene regulation Chain: A: PDB Molecule: polycomb group ring finger protein 6; PDBTitle: solution structure of the ring domain of the human polycomb2 group ring finger protein 6
62	c2csyA_	Alignment	not modelled	15.6	22	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: zinc finger protein 183-like 1; PDBTitle: solution structure of the ring domain of the zinc finger2 protein 183-like 1
63	c2gb5B_	Alignment	not modelled	15.1	24	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
64	c3t6pA_	Alignment	not modelled	15.1	30	PDB header: apoptosis Chain: A: PDB Molecule: baculoviral iap repeat-containing protein 2; PDBTitle: iap antagonist-induced conformational change in ciap1 promotes e32 ligase activation via dimerization
65	c2yslA_	Alignment	not modelled	14.9	21	PDB header: protein binding Chain: A: PDB Molecule: tripartite motif-containing protein 31; PDBTitle: solution structure of the ring domain (1-66) from2 tripartite motif-containing protein 31
66	c2ea5A_	Alignment	not modelled	14.7	40	PDB header: cell cycle Chain: A: PDB Molecule: cell growth regulator with ring finger domain PDBTitle: solution structure of the ring domain of the human cell2 growth regulator with ring finger domain 1 protein
67	c3i2dA_	Alignment	not modelled	14.7	20	PDB header: ligase Chain: A: PDB Molecule: e3 sumo-protein ligase siz1; PDBTitle: crystal structure of s. cerevisiae sumo e3 ligase siz1
68	d2apob1	Alignment	not modelled	14.6	30	Fold: Rubredoxin-like Superfamily: Nop10-like SnoRNP Family: Nucleolar RNA-binding protein Nop10-like
69	d1twfj_	Alignment	not modelled	14.6	41	Fold: DNA/RNA-binding 3-helical bundle Superfamily: RNA polymerase subunit RPB10 Family: RNA polymerase subunit RPB10
70	c2ec1A_	Alignment	not modelled	14.4	36	PDB header: metal binding protein Chain: A: PDB Molecule: ring-box protein 2; PDBTitle: solution structure of the ring domain of the human ring-box2 protein 2
71	d1nnqa2	Alignment	not modelled	14.4	50	Fold: Rubredoxin-like Superfamily: Rubredoxin-like Family: Rubredoxin
72	c2y0fD_	Alignment	not modelled	14.2	29	PDB header: oxidoreductase Chain: D: PDB Molecule: 4-hydroxy-3-methylbut-2-en-1-yl diphosphate synthase; PDBTitle: structure of gcpe (ispg) from thermus thermophilus hb27
73	d1dgsa1	Alignment	not modelled	14.2	32	Fold: SAM domain-like Superfamily: RuvA domain 2-like Family: NAD+-dependent DNA ligase, domain 3
74	d2ey4e1	Alignment	not modelled	13.9	30	Fold: Rubredoxin-like Superfamily: Nop10-like SnoRNP Family: Nucleolar RNA-binding protein Nop10-like
75	c2ak1A_	Alignment	not modelled	13.8	22	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: phna-like protein pa0128; PDBTitle: solution structure for phn-a like protein pa0128 from2 pseudomonas aeruginosa
76	c1z6uA_	Alignment	not modelled	13.7	33	PDB header: ligase Chain: A: PDB Molecule: np95-like ring finger protein isoform b; PDBTitle: np95-like ring finger protein isoform b [homo sapiens]
77	c2yurA_	Alignment	not modelled	13.5	36	PDB header: protein binding Chain: A: PDB Molecule: retinoblastoma-binding protein 6; PDBTitle: solution structure of the ring finger of human2 retinoblastoma-binding protein 6
78	d2baya1	Alignment	not modelled	13.5	26	Fold: RING/U-box Superfamily: RING/U-box Family: U-box
79	d1lkoa2	Alignment	not modelled	13.4	38	Fold: Rubredoxin-like Superfamily: Rubredoxin-like

						Family: Rubredoxin
80	c2vjfB_	Alignment	not modelled	13.4	44	PDB header: ligase Chain: B: PDB Molecule: mdm4 protein; PDBTitle: crystal structure of the mdm2-mdmx ring domain heterodimer
81	d2j0151	Alignment	not modelled	13.4	24	Fold: Rubredoxin-like Superfamily: Zn-binding ribosomal proteins Family: Ribosomal protein L32p
82	d2qam01	Alignment	not modelled	13.4	25	Fold: Rubredoxin-like Superfamily: Zn-binding ribosomal proteins Family: Ribosomal protein L32p
83	c2yhoE_	Alignment	not modelled	13.2	40	PDB header: ligase Chain: E: PDB Molecule: e3 ubiquitin-protein ligase mylip; PDBTitle: the idol-ube2d complex mediates sterol-dependent degradation of the2 Idl receptor
84	d1b8ta3	Alignment	not modelled	12.7	56	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
85	d1bia2	Alignment	not modelled	12.7	31	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
86	c2emwA_	Alignment	not modelled	12.6	17	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 268; PDBTitle: solution structure of the c2h2 type zinc finger (region 301-2 331) of human zinc finger protein 268
87	c3b08H_	Alignment	not modelled	12.6	29	PDB header: signaling protein/metal binding protein Chain: H: PDB Molecule: ranbp-type and c3hc4-type zinc finger-containing protein 1; PDBTitle: crystal structure of the mouse hoil1-l-nzf in complex with linear di-2 ubiquitin
88	d1e7la2	Alignment	not modelled	12.5	25	Fold: His-Me finger endonucleases Superfamily: His-Me finger endonucleases Family: Recombination endonuclease VII, N-terminal domain
89	c2ecyA_	Alignment	not modelled	12.4	23	PDB header: apoptosis Chain: A: PDB Molecule: tnf receptor-associated factor 3; PDBTitle: solution structure of the zinc finger, c3hc4 type (ring2 finger)" domain of tnf receptor-associated factor 3
90	c2hdpB_	Alignment	not modelled	12.3	22	PDB header: ligase Chain: B: PDB Molecule: ubiquitin-protein ligase e3 mdm2; PDBTitle: solution structure of hdm2 ring finger domain
91	d1e4ua_	Alignment	not modelled	12.2	21	Fold: RING/U-box Superfamily: RING/U-box Family: RING finger domain, C3HC4
92	c3fl2A_	Alignment	not modelled	12.2	30	PDB header: ligase Chain: A: PDB Molecule: e3 ubiquitin-protein ligase uhrf1; PDBTitle: crystal structure of the ring domain of the e3 ubiquitin-2 protein ligase uhrf1
93	c2xeuA_	Alignment	not modelled	12.1	33	PDB header: transcription Chain: A: PDB Molecule: ring finger protein 4; PDBTitle: ring domain
94	d1jm7a_	Alignment	not modelled	12.0	15	Fold: RING/U-box Superfamily: RING/U-box Family: RING finger domain, C3HC4
95	d1qxfa_	Alignment	not modelled	11.7	26	Fold: Rubredoxin-like Superfamily: Zn-binding ribosomal proteins Family: Ribosomal protein S27e
96	d1qcva_	Alignment	not modelled	11.7	27	Fold: Rubredoxin-like Superfamily: Rubredoxin-like Family: Rubredoxin
97	c2ep4A_	Alignment	not modelled	11.5	40	PDB header: protein binding Chain: A: PDB Molecule: ring finger protein 24; PDBTitle: solution structure of ring finger from human ring finger2 protein 24
98	c2d8tA_	Alignment	not modelled	11.4	22	PDB header: metal binding protein Chain: A: PDB Molecule: ring finger protein 146; PDBTitle: solution structure of the ring domain of the human ring2 finger protein 146
99	d1x6ma_	Alignment	not modelled	11.4	26	Fold: Mss4-like Superfamily: Mss4-like Family: Glutathione-dependent formaldehyde-activating enzyme, Gfa