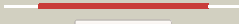



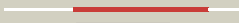




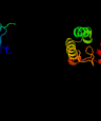



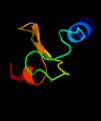



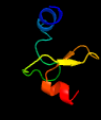

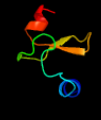


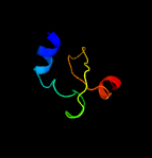

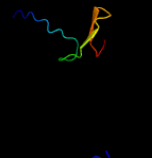
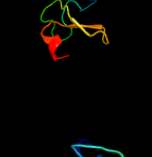




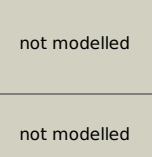


Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD2362c_(recO)_2643471_2644268
Date	Mon Aug 5 13:25:51 BST 2019
Unique Job ID	012a79b58871ce07

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1u5kA_	 Alignment		100.0	22	PDB header: recombination,replication Chain: A: PDB Molecule: hypothetical protein; PDBTitle: recombinational repair protein reco
2	c3q8dB_	 Alignment		100.0	20	PDB header: dna binding protein Chain: B: PDB Molecule: dna repair protein reco; PDBTitle: e. coli reco complex with ssb c-terminus
3	d1u5ka2	 Alignment		100.0	20	Fold: ArfGap/RecO-like zinc finger Superfamily: ArfGap/RecO-like zinc finger Family: RecO C-terminal domain-like
4	d1u5ka1	 Alignment		98.3	34	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: RecO N-terminal domain-like
5	c5fz5M_	 Alignment		93.2	16	PDB header: transcription Chain: M: PDB Molecule: transcription initiation factor iib; PDBTitle: transcription initiation complex structures elucidate dna opening (cc)
6	c2crwA_	 Alignment		89.4	17	PDB header: signaling protein Chain: A: PDB Molecule: adp-ribosylation factor gtpase-activating PDBTitle: solution structure of the arfgap domain of adp-ribosylation2 factor gtpaseactivating protein 3 (arfgap 3)
7	c2d9lA_	 Alignment		88.3	19	PDB header: gene regulation Chain: A: PDB Molecule: nucleoporin-like protein rip; PDBTitle: solution structure of the arfgap domain of human rip
8	c2p57A_	 Alignment		88.0	17	PDB header: metal binding protein Chain: A: PDB Molecule: gtpase-activating protein znf289; PDBTitle: gap domain of znf289, an id1-regulated zinc finger protein
9	c2owaB_	 Alignment		87.9	14	PDB header: protein transport Chain: B: PDB Molecule: arfgap-like finger domain containing protein; PDBTitle: crystal structure of putative gtpase activating protein for2 adp ribosylation factor from cryptosporidium parvum3 (cgd5_1040)
10	c3dwdB_	 Alignment		87.6	12	PDB header: transport protein Chain: B: PDB Molecule: adp-ribosylation factor gtpase-activating protein 1; PDBTitle: crystal structure of the arfgap domain of human arfgap1
11	d1dcqa2	 Alignment		87.4	14	Fold: ArfGap/RecO-like zinc finger Superfamily: ArfGap/RecO-like zinc finger Family: Pyk2-associated protein beta ARF-GAP domain

12	c3o47A_	Alignment		87.4	12	PDB header: hydrolase, hydrolase activator Chain: A: PDB Molecule: adp-ribosylation factor gtpase-activating protein 1, adp- PDBTitle: crystal structure of arfgap1-arf1 fusion protein
13	c2lqjB_	Alignment		87.2	17	PDB header: protein transport Chain: B: PDB Molecule: stromal membrane-associated protein 1-like; PDBTitle: crystal structure of the gap domain of smap1l (loc64744) stromal2 membrane-associated protein 1-like
14	c3k1fM_	Alignment		87.1	31	PDB header: transcription Chain: M: PDB Molecule: transcription initiation factor iib; PDBTitle: crystal structure of rna polymerase ii in complex with tfiib
15	c3lvrE_	Alignment		84.9	14	PDB header: protein transport Chain: E: PDB Molecule: arf-gap with sh3 domain, ank repeat and ph domain- PDBTitle: the crystal structure of asap3 in complex with arf6 in transition2 state soaked with calcium
16	c3fehA_	Alignment		84.2	24	PDB header: hydrolase activator Chain: A: PDB Molecule: centaurin-alpha-1; PDBTitle: crystal structure of full length centaurin alpha-1
17	c2crrA_	Alignment		83.6	17	PDB header: protein transport Chain: A: PDB Molecule: stromal membrane-associated protein smap1b; PDBTitle: solution structure of arfgap domain from human smap1
18	d1libia2	Alignment		81.8	23	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
19	c1dcqA_	Alignment		81.6	18	PDB header: metal binding protein Chain: A: PDB Molecule: pyk2-associated protein beta; PDBTitle: crystal structure of the arf-gap domain and ankyrin repeats of2 papbeta.
20	c2b0oF_	Alignment		81.4	18	PDB header: metal binding protein Chain: F: PDB Molecule: uplc1; PDBTitle: crystal structure of uplc1 gap domain
21	c3jueA_	Alignment	not modelled	80.1	21	PDB header: protein transport/endocytosis Chain: A: PDB Molecule: arfgap with coiled-coil, ank repeat and ph domain- PDBTitle: crystal structure of arfgap and ank repeat domain of acap1
22	c4v1oM_	Alignment	not modelled	78.6	30	PDB header: transcription Chain: M: PDB Molecule: transcription initiation factor iib; PDBTitle: architecture of the rna polymerase ii-mediator core transcription2 initiation complex
23	c3t9kA_	Alignment	not modelled	77.9	17	PDB header: protein transport Chain: A: PDB Molecule: arf-gap with coiled-coil, ank repeat and ph domain- PDBTitle: crystal structure of acap1 c-portion mutant s554d fused with integrin2 beta1 peptide
24	c6cnbR_	Alignment	not modelled	76.5	16	PDB header: transcription/dna Chain: R: PDB Molecule: transcription factor iib 70 kda subunit,tata-box-binding PDBTitle: yeast rna polymerase iii initial transcribing complex
25	c3k7aM_	Alignment	not modelled	70.9	30	PDB header: transcription Chain: M: PDB Molecule: transcription initiation factor iib; PDBTitle: crystal structure of an rna polymerase ii-tfiib complex
26	c5iy9Q_	Alignment	not modelled	69.7	24	PDB header: transcription, transferase/dna/rna Chain: Q: PDB Molecule: general transcription factor iie subunit 1; PDBTitle: human holo-pic in the initial transcribing state (no iis)
27	c1x68A_	Alignment	not modelled	69.4	17	PDB header: protein binding Chain: A: PDB Molecule: fhl5 protein; PDBTitle: solution structures of the c-terminal lim domain of human2 fh15 protein
28	c5iy9M_	Alignment	not modelled	68.1	16	PDB header: transcription, transferase/dna/rna Chain: M: PDB Molecule: transcription initiation factor iib; PDBTitle: human holo-pic in the initial transcribing state (no iis)

29	d1x4la2	Alignment	not modelled	61.8	23	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
30	c2gb5B_	Alignment	not modelled	61.8	16	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
31	d1hk8a_	Alignment	not modelled	61.6	18	Fold: PFL-like glycy radical enzymes Superfamily: PFL-like glycy radical enzymes Family: Class III anaerobic ribonucleotide reductase NRDD subunit
32	c1hk8A_	Alignment	not modelled	61.6	18	PDB header: oxidoreductase Chain: A: PDB Molecule: anaerobic ribonucleotide-triphosphate reductase; PDBTitle: structural basis for allosteric substrate specificity regulation in2 class iii ribonucleotide reductases: nrdd in complex with dgtp
33	c1x64A_	Alignment	not modelled	61.2	20	PDB header: contractile protein Chain: A: PDB Molecule: alpha-actinin-2 associated lim protein; PDBTitle: solution structure of the lim domain of alpha-actinin-2 associated lim protein
34	c5fmfR_	Alignment	not modelled	61.1	18	PDB header: transcription Chain: R: PDB Molecule: transcription initiation factor iie subunit alpha, tfa1; PDBTitle: the p-lobe of rna polymerase ii pre-initiation complex
35	d2dara1	Alignment	not modelled	60.5	27	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
36	c6f42V_	Alignment	not modelled	58.9	27	PDB header: transcription Chain: V: PDB Molecule: transcription factor iiii 70 kda subunit; PDBTitle: rna polymerase iii closed complex cc1.
37	c6o9lQ_	Alignment	not modelled	58.8	25	PDB header: transcription/dna Chain: Q: PDB Molecule: general transcription factor iie subunit 1; PDBTitle: human holo-pic in the closed state
38	c5fywW_	Alignment	not modelled	57.8	18	PDB header: transcription Chain: W: PDB Molecule: transcription initiation factor iie subunit alpha; PDBTitle: transcription initiation complex structures elucidate dna opening (oc)
39	c1x63A_	Alignment	not modelled	55.7	19	PDB header: contractile protein Chain: A: PDB Molecule: skeletal muscle lim-protein 1; PDBTitle: solution structure of the second lim domain of skeletal2 muscle lim protein 1
40	c1wyhA_	Alignment	not modelled	55.6	31	PDB header: metal binding protein Chain: A: PDB Molecule: skeletal muscle lim-protein 2; PDBTitle: solution structure of the lim domain from human skeletal2 muscle lim-protein 2
41	c3f6qB_	Alignment	not modelled	55.4	31	PDB header: signaling protein/signaling protein Chain: B: PDB Molecule: lim and senescent cell antigen-like-containing PDBTitle: crystal structure of integrin-linked kinase ankyrin repeat2 domain in complex with pinch1 lim1 domain
42	c1x4lA_	Alignment	not modelled	55.1	20	PDB header: metal binding protein Chain: A: PDB Molecule: skeletal muscle lim-protein 3; PDBTitle: solution structure of lim domain in four and a half lim2 domains protein 2
43	d1x68a2	Alignment	not modelled	54.9	18	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
44	c3vrhA_	Alignment	not modelled	54.4	20	PDB header: rna binding protein Chain: A: PDB Molecule: putative uncharacterized protein ph0300; PDBTitle: crystal structure of ph0300
45	c5oqmW_	Alignment	not modelled	54.2	18	PDB header: transcription Chain: W: PDB Molecule: transcription factor tfiie subunit; PDBTitle: structure of yeast transcription pre-initiation complex with tfiih and2 core mediator
46	c6gymW_	Alignment	not modelled	53.8	18	PDB header: transcription Chain: W: PDB Molecule: transcription initiation factor iie subunit alpha, PDBTitle: structure of a yeast closed complex with distorted dna (ccd1st)
47	d1pfta_	Alignment	not modelled	53.3	29	Fold: Rubredoxin-like Superfamily: Zinc beta-ribbon Family: Transcriptional factor domain
48	c3d00A_	Alignment	not modelled	53.1	24	PDB header: metal binding protein Chain: A: PDB Molecule: tungsten formylmethanofuran dehydrogenase subunit e; PDBTitle: crystal structure of a tungsten formylmethanofuran dehydrogenase2 subunit e (fmde)-like protein (syn_00638) from syntrophus3 aciditrophicus at 1.90 a resolution
49	c3dzuD_	Alignment	not modelled	52.7	23	PDB header: transcription/dna Chain: D: PDB Molecule: peroxisome proliferator-activated receptor gamma; PDBTitle: intact ppar gamma - rxr alpha nuclear receptor complex on dna bound2 with bvt.13, 9-cis retinoic acid and ncoa2 peptide
50	c2gviA_	Alignment	not modelled	52.5	20	PDB header: oxidoreductase Chain: A: PDB Molecule: conserved hypothetical protein; PDBTitle: crystal structure of a putative formylmethanofuran dehydrogenase2 subunit e (ta1109) from thermoplasma acidophilum at 1.87 a resolution
51	c5oqiW_	Alignment	not modelled	52.4	18	PDB header: transcription Chain: W: PDB Molecule: transcription initiation factor iie subunit alpha; PDBTitle: structure of yeast transcription pre-initiation complex with tfiih
52	d1xpa2	Alignment	not modelled	52.0	17	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: DNA repair factor XPA DNA- and RPA-binding domain, N-

						terminal subdomain
53	d2cupa2	Alignment	not modelled	51.3	27	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
54	c3izci	Alignment	not modelled	51.0	37	PDB header: ribosome Chain: I: PDB Molecule: 60s ribosomal protein rp10 (I10e); PDBTitle: localization of the large subunit ribosomal proteins into a 6.1 a2 cryo-em map of saccharomyces cerevisiae translating 80s ribosome
55	d1wfla	Alignment	not modelled	50.5	19	Fold: AN1-like Zinc finger Superfamily: AN1-like Zinc finger Family: AN1-like Zinc finger
56	c1d4uA	Alignment	not modelled	50.4	15	PDB header: dna binding protein Chain: A: PDB Molecule: nucleotide excision repair protein xpa (xpa-mbd); PDBTitle: interactions of human nucleotide excision repair protein2 xpa with rpa70 and dna: chemical shift mapping and 15n nmr3 relaxation studies
57	c2eheA	Alignment	not modelled	49.5	17	PDB header: structural protein Chain: A: PDB Molecule: four and a half lim domains 3; PDBTitle: solution structure of the first lim domain from human four2 and a half lim domains protein 3
58	c2darA	Alignment	not modelled	49.3	28	PDB header: metal binding protein Chain: A: PDB Molecule: pdz and lim domain protein 5; PDBTitle: solution structure of first lim domain of enigma-like pdz2 and lim domains protein
59	c4nqaL	Alignment	not modelled	49.2	14	PDB header: transcription regulator/dna Chain: I: PDB Molecule: liver x nuclear receptor beta; PDBTitle: crystal structure of liganded hrxr-alpha/hlrx-beta heterodimer on dna
60	c5uanA	Alignment	not modelled	49.1	27	PDB header: transcription/dna Chain: A: PDB Molecule: retinoic acid receptor rxr-alpha; PDBTitle: crystal structure of multi-domain rar-beta-rxr-alpha heterodimer on2 dna
61	d1dl6a	Alignment	not modelled	48.9	17	Fold: Rubredoxin-like Superfamily: Zinc beta-ribbon Family: Transcriptional factor domain
62	c2o10A	Alignment	not modelled	48.9	15	PDB header: metal binding protein Chain: A: PDB Molecule: muscle lim protein; PDBTitle: solution structure of the n-terminal lim domain of mlp/crp3
63	c1libA	Alignment	not modelled	48.6	23	PDB header: metal binding protein Chain: A: PDB Molecule: cysteine-rich protein 2; PDBTitle: quail cysteine and glycine-rich protein, nmr, 15 minimized2 model structures
64	c2d8xA	Alignment	not modelled	48.2	25	PDB header: structural protein, cell cycle Chain: A: PDB Molecule: protein pinch; PDBTitle: solution structure of the second lim domain of particularly2 interesting new cys-his protein (pinch)
65	c2cuqA	Alignment	not modelled	47.8	25	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: four and a half lim domains 3; PDBTitle: solution structure of second lim domain from human skeletal2 muscle lim-protein 2
66	c2egqA	Alignment	not modelled	47.4	19	PDB header: structural protein Chain: A: PDB Molecule: fhl1 protein; PDBTitle: solution structure of the fourth lim domain from human four2 and a half lim domains 1
67	c1x3hA	Alignment	not modelled	47.2	25	PDB header: metal binding protein Chain: A: PDB Molecule: leupaxin; PDBTitle: solution structure of the lim domain of human leupaxin
68	c3iz5i	Alignment	not modelled	46.5	10	PDB header: ribosome Chain: I: PDB Molecule: 60s ribosomal protein I10 (I10e); PDBTitle: localization of the large subunit ribosomal proteins into a 5.5 a2 cryo-em map of triticum aestivum translating 80s ribosome
69	c2curA	Alignment	not modelled	46.4	19	PDB header: metal binding protein Chain: A: PDB Molecule: skeletal muscle lim-protein 1; PDBTitle: solution structure of skeletal muscle lim-protein 1
70	d1wfpA	Alignment	not modelled	46.4	22	Fold: AN1-like Zinc finger Superfamily: AN1-like Zinc finger Family: AN1-like Zinc finger
71	c2mbvA	Alignment	not modelled	46.4	34	PDB header: transcription Chain: A: PDB Molecule: fusion protein of lim domain transcription factor lmo4 (77- PDBTitle: lmo4-lim2 in complex with deaf1 (404-418)
72	c3j39g	Alignment	not modelled	46.3	23	PDB header: ribosome Chain: G: PDB Molecule: 60s ribosomal protein I7a; PDBTitle: structure of the d. melanogaster 60s ribosomal proteins
73	d3gata	Alignment	not modelled	45.1	27	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Erythroid transcription factor GATA-1
74	c5uanB	Alignment	not modelled	44.8	27	PDB header: transcription/dna Chain: B: PDB Molecule: retinoic acid receptor beta; PDBTitle: crystal structure of multi-domain rar-beta-rxr-alpha heterodimer on2 dna
75	d1wffa	Alignment	not modelled	44.7	22	Fold: AN1-like Zinc finger Superfamily: AN1-like Zinc finger Family: AN1-like Zinc finger
76	c1x4kA	Alignment	not modelled	44.7	25	PDB header: metal binding protein Chain: A: PDB Molecule: skeletal muscle lim-protein 3; PDBTitle: solution structure of lim domain in lim-protein 3
77	c3dzyA	Alignment	not modelled	44.6	27	PDB header: transcription/dna Chain: A: PDB Molecule: retinoic acid receptor rxr-alpha; PDBTitle: intact ppar gamma - rxr alpha nuclear receptor complex on dna bound2 with rosiglitazone, 9-cis retinoic acid and ncoa2 peptide PDB header: transcription/dna

78	c4iqrE_	Alignment	not modelled	43.9	32	Chain: E: PDB Molecule: hepatocyte nuclear factor 4-alpha; PDBTitle: multi-domain organization of the hnf4alpha nuclear receptor complex on2 dna
79	c3j3bg_	Alignment	not modelled	43.6	40	PDB header: ribosome Chain: G: PDB Molecule: 60s ribosomal protein l7a; PDBTitle: structure of the human 60s ribosomal proteins
80	d1wg2a_	Alignment	not modelled	43.5	17	Fold: AN1-like Zinc finger Superfamily: AN1-like Zinc finger Family: AN1-like Zinc finger
81	d2vuti1	Alignment	not modelled	43.4	26	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Erythroid transcription factor GATA-1
82	c2d8zA_	Alignment	not modelled	42.2	19	PDB header: signaling protein, structural protein Chain: A: PDB Molecule: four and a half lim domains 2; PDBTitle: solution structure of the third lim domain of four and a2 half lim domains protein 2 (fhl-2)
83	c1a7iA_	Alignment	not modelled	42.1	16	PDB header: lim domain containing proteins Chain: A: PDB Molecule: qcrp2 (lim1); PDBTitle: amino-terminal lim domain from quail cysteine and glycine-2 rich protein, nmr, minimized average structure
84	d1y0ja1	Alignment	not modelled	42.1	24	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Erythroid transcription factor GATA-1
85	d5gata_	Alignment	not modelled	42.1	23	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Erythroid transcription factor GATA-1
86	c1x61A_	Alignment	not modelled	41.1	16	PDB header: cell adhesion Chain: A: PDB Molecule: thyroid receptor interacting protein 6; PDBTitle: solution structure of the first lim domain of thyroid2 receptor interacting protein 6 (trip6)
87	d1wfhA_	Alignment	not modelled	40.8	17	Fold: AN1-like Zinc finger Superfamily: AN1-like Zinc finger Family: AN1-like Zinc finger
88	d2ds5a1	Alignment	not modelled	40.4	18	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: ClpX chaperone zinc binding domain
89	c1nypA_	Alignment	not modelled	40.1	22	PDB header: cell adhesion Chain: A: PDB Molecule: pinch protein; PDBTitle: 4th lim domain of pinch protein
90	d1b8ta2	Alignment	not modelled	40.0	16	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
91	c1ovxB_	Alignment	not modelled	40.0	18	PDB header: metal binding protein Chain: B: PDB Molecule: atp-dependent clp protease atp-binding subunit clpx; PDBTitle: nmr structure of the e. coli clpx chaperone zinc binding domain dimer
92	d1wyha1	Alignment	not modelled	38.7	32	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
93	d1b8ta4	Alignment	not modelled	38.5	23	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
94	c2ds8A_	Alignment	not modelled	38.0	15	PDB header: metal binding protein, protein binding Chain: A: PDB Molecule: atp-dependent clp protease atp-binding subunit PDBTitle: structure of the zbd-xb complex
95	d1gnfa_	Alignment	not modelled	37.4	23	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Erythroid transcription factor GATA-1
96	c2l3kA_	Alignment	not modelled	37.1	16	PDB header: oncoprotein Chain: A: PDB Molecule: rhombotin-2, linker, lim domain-binding protein 1; PDBTitle: solution structure of lmo2(lim2)-ldb1(lid)
97	c1x6aA_	Alignment	not modelled	36.5	16	PDB header: protein binding Chain: A: PDB Molecule: lim domain kinase 2; PDBTitle: solution structures of the second lim domain of human lim-2 kinase 2 (limk2)
98	c2o13A_	Alignment	not modelled	36.5	22	PDB header: metal binding protein Chain: A: PDB Molecule: muscle lim protein; PDBTitle: solution structure of the c-terminal lim domain of mlp/crp3
99	d1vd4a_	Alignment	not modelled	36.4	25	Fold: Rubredoxin-like Superfamily: Zinc beta-ribbon Family: Transcriptional factor domain
100	d1hraa_	Alignment	not modelled	35.9	23	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
101	d1hcqa_	Alignment	not modelled	35.9	24	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
102	c4u3eA_	Alignment	not modelled	35.6	28	PDB header: oxidoreductase Chain: A: PDB Molecule: ribonucleoside triphosphate reductase; PDBTitle: anaerobic ribonucleotide reductase
103	c2co8A_	Alignment	not modelled	35.4	17	PDB header: metal binding protein Chain: A: PDB Molecule: nedd9 interacting protein with calponin homology PDBTitle: solution structures of the lim domain of human nedd92 interacting protein with calponin homology and lim domains
104	c2miuA_	Alignment	not modelled	34.8	13	PDB header: protein binding Chain: A: PDB Molecule: four and a half lim domains protein 2; PDBTitle: structure of fhl2 lim adaptor and its interaction with ski

105	d1kb6b_	Alignment	not modelled	34.7	24	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
106	c2jtnA_	Alignment	not modelled	34.5	13	PDB header: protein binding/transcription Chain: A: PDB Molecule: lim domain-binding protein 1, lim/homeobox PDBTitle: nmr solution structure of a ldb1-lid:lhx3-lim complex
107	d2fiya1	Alignment	not modelled	34.4	18	Fold: FdhE-like Superfamily: FdhE-like Family: FdhE-like
108	c1ee8A_	Alignment	not modelled	34.4	26	PDB header: dna binding protein Chain: A: PDB Molecule: mutm (fpg) protein; PDBTitle: crystal structure of mutm (fpg) protein from thermus thermophilus hb8
109	c2cu8A_	Alignment	not modelled	34.3	23	PDB header: metal binding protein Chain: A: PDB Molecule: cysteine-rich protein 2; PDBTitle: solution structure of the lim domain of human cysteine-rich2 protein 2
110	d2cura1	Alignment	not modelled	34.3	20	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
111	c2d8yA_	Alignment	not modelled	34.2	13	PDB header: structural protein Chain: A: PDB Molecule: eplin protein; PDBTitle: solution structure of the lim domain of epithelial protein2 lost in neoplasm
112	c3j21d_	Alignment	not modelled	34.0	24	PDB header: ribosome Chain: D: PDB Molecule: 50s ribosomal protein l4p; PDBTitle: promiscuous behavior of proteins in archaeal ribosomes revealed by2 cryo-em: implications for evolution of eukaryotic ribosomes (50s3 ribosomal proteins)
113	c2corA_	Alignment	not modelled	33.6	26	PDB header: structural protein Chain: A: PDB Molecule: pinch protein; PDBTitle: solution structure of the third lim domain of particularly2 interesting new cys-his protein
114	c1x62A_	Alignment	not modelled	33.0	16	PDB header: structural protein Chain: A: PDB Molecule: c-terminal lim domain protein 1; PDBTitle: solution structure of the lim domain of carboxyl terminal2 lim domain protein 1
115	c5ghaC_	Alignment	not modelled	33.0	17	PDB header: transferase/transport protein Chain: C: PDB Molecule: sulfur transferase ttua; PDBTitle: sulfur transferase ttua in complex with sulfur carrier ttub
116	c4v195_	Alignment	not modelled	32.8	18	PDB header: ribosome Chain: 5: PDB Molecule: mitoribosomal protein bl32m, mrpl32; PDBTitle: structure of the large subunit of the mammalian mitoribosome, part 12 of 2
117	d2nllb_	Alignment	not modelled	32.7	19	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
118	d1kb2a_	Alignment	not modelled	32.5	24	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor
119	c2lzuA_	Alignment	not modelled	32.4	16	PDB header: metal binding protein Chain: A: PDB Molecule: lim domain-containing protein 2; PDBTitle: solution structure of limd2
120	d1glua_	Alignment	not modelled	31.8	19	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: Nuclear receptor