
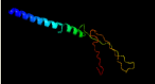
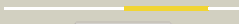
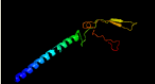

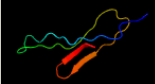

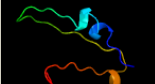



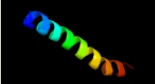

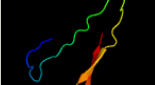

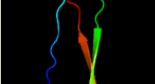






Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD2091c_(-)_2348566_2349300
Date	Mon Aug 5 13:25:20 BST 2019
Unique Job ID	10141c7bffe6ee59

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c5o4uK_	 Alignment		81.5	14	PDB header: cell adhesion Chain: K; PDB Molecule: flagellin; PDBTitle: the flagellin of pyrococcus furiosus
2	c5tfyJ_	 Alignment		77.1	15	PDB header: cell adhesion Chain: J; PDB Molecule: flagellin; PDBTitle: the archaeal flagellum of methanospirillum hungatei strain jf1.
3	d1qfha2	 Alignment		72.2	25	Fold: Immunoglobulin-like beta-sandwich Superfamily: E set domains Family: Filamin repeat (rod domain)
4	d2oz4a3	 Alignment		70.9	7	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
5	c2k1aA_	 Alignment		69.0	27	PDB header: cell adhesion Chain: A; PDB Molecule: integrin alpha-iiB; PDBTitle: bicelle-embedded integrin alpha(iiB) transmembrane segment
6	c2kncA_	 Alignment		65.9	27	PDB header: cell adhesion Chain: A; PDB Molecule: integrin alpha-iiB; PDBTitle: platelet integrin alphaIIb-beta3 transmembrane-cytoplasmic2 heterocomplex
7	d2nxyb2	 Alignment		63.7	21	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C2 set domains
8	c3ginB_	 Alignment		57.5	25	PDB header: metal binding protein Chain: B; PDB Molecule: sodium/calcium exchanger 1; PDBTitle: crystal structure of e454k-cbd1
9	c3s37X_	 Alignment		55.8	15	PDB header: immune system/transferase Chain: X; PDB Molecule: vascular endothelial growth factor receptor 2; PDBTitle: structural basis for the function of two anti-vegF receptor antibodies
10	d1o75a3	 Alignment		55.6	19	Fold: Tp47 lipoprotein, N-terminal domain Superfamily: Tp47 lipoprotein, N-terminal domain Family: Tp47 lipoprotein, N-terminal domain
11	c3osvC_	 Alignment		54.0	27	PDB header: structural protein Chain: C; PDB Molecule: flagellar basal-body rod modification protein flgd; PDBTitle: the crytsal structure of flgd from p. aeruginosa

12	c2bk8A_	Alignment		54.0	13	PDB header: ig domain Chain: A; PDB Molecule: titin heart isoform n2-b; PDBTitle: m1 domain from titin
13	c2mkIC_	Alignment		52.5	15	PDB header: immune system Chain: C; PDB Molecule: novel antigen receptor; PDBTitle: solution structure of the fourth constant immunoglobulin domain of f2 nurse shark ignar
14	d1ql4b_	Alignment		51.7	15	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
15	d1bjha1	Alignment		51.7	11	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
16	c2rrfA_	Alignment		50.7	29	PDB header: unknown function Chain: A; PDB Molecule: zinc finger fyve domain-containing protein 21; PDBTitle: the solution structure of the c-terminal region of zinc finger fyve2 domain-containing protein 21
17	c5lxkA_	Alignment		50.6	18	PDB header: viral protein Chain: A; PDB Molecule: decoration protein; PDBTitle: nmr structure of the c-terminal domain of the bacteriophage t52 decoration protein pb10.
18	c5z1lL_	Alignment		50.5	15	PDB header: protein fibril Chain: L; PDB Molecule: flagellin; PDBTitle: cryo-em structure of methanococcus maripaludis archaellum
19	d2avga1	Alignment		49.7	26	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
20	c6nhwC_	Alignment		48.0	33	PDB header: immune system Chain: C; PDB Molecule: tumor necrosis factor receptor superfamily member 10b; PDBTitle: structure of the transmembrane domain of the death receptor 5 - dimer2 of trimer
21	c2qvka_	Alignment	not modelled	47.4	22	PDB header: metal binding protein Chain: A; PDB Molecule: sodium/calcium exchanger 1; PDBTitle: the second ca2+-binding domain of the na+-ca2+ exchanger is2 essential for regulation: crystal structures and3 mutational analysis
22	c2l7JA_	Alignment	not modelled	47.4	9	PDB header: membrane protein Chain: A; PDB Molecule: poliovirus receptor-related 1; PDBTitle: solution structure of the third immunoglobulin-like domain of nectin-1
23	c6nhwB_	Alignment	not modelled	47.1	33	PDB header: immune system Chain: B; PDB Molecule: tumor necrosis factor receptor superfamily member 10b; PDBTitle: structure of the transmembrane domain of the death receptor 5 - dimer2 of trimer
24	c6nhwA_	Alignment	not modelled	47.0	33	PDB header: immune system Chain: A; PDB Molecule: tumor necrosis factor receptor superfamily member 10b; PDBTitle: structure of the transmembrane domain of the death receptor 5 - dimer2 of trimer
25	c1o75A_	Alignment	not modelled	47.0	19	PDB header: lipoprotein nullIntigen Chain: A; PDB Molecule: 47 kda membrane antigen; PDBTitle: tp47, the 47-kilodalton lipoprotein of treponema pallidum
26	d1ulva2	Alignment	not modelled	46.7	31	Fold: Immunoglobulin-like beta-sandwich Superfamily: E set domains Family: E-set domains of sugar-utilizing enzymes
27	d1olza1	Alignment	not modelled	46.4	20	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
28	c3rnhA_	Alignment	not modelled	45.7	26	PDB header: cell adhesion Chain: A; PDB Molecule: filamin-a;

28	c0rgnA_	Alignment	not modelled	43.7	20	PDBTitle: structure of filamin a immunoglobulin-like repeat 10 from homo sapiens PDB header: structural protein Chain: A; PDB Molecule: bk158_1; PDBTitle: solution structure of the filamin domain from human bk158_12 protein
29	c2di7A_	Alignment	not modelled	44.8	18	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
30	d2dava1	Alignment	not modelled	44.7	18	PDB header: cell adhesion Chain: X; PDB Molecule: cox sackievirus and adenovirus receptor; PDBTitle: nmr structure of card d2 domain
31	d1rhfa2	Alignment	not modelled	44.0	21	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
32	c2nplX_	Alignment	not modelled	43.7	13	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
33	d1epfa1	Alignment	not modelled	42.4	13	PDB header: immune system Chain: E; PDB Molecule: tumor necrosis factor receptor superfamily member 10b; PDBTitle: structure of the transmembrane domain of the death receptor 5 - dimer2 of trimer
34	c6nhwE_	Alignment	not modelled	41.4	33	PDB header: immune system Chain: F; PDB Molecule: tumor necrosis factor receptor superfamily member 10b; PDBTitle: structure of the transmembrane domain of the death receptor 5 - dimer2 of trimer
35	c6nhwF_	Alignment	not modelled	41.4	33	PDB header: hydrolase Chain: C; PDB Molecule: mucosa-associated lymphoid tissue lymphoma PDBTitle: crystal structure of ig-like c2-type 2 domain of the human2 mucosa-associated lymphoid tissue lymphoma translocation3 protein 1
36	c3bfoC_	Alignment	not modelled	41.3	12	PDB header: immune system Chain: D; PDB Molecule: tumor necrosis factor receptor superfamily member 10b; PDBTitle: structure of the transmembrane domain of the death receptor 5 - dimer2 of trimer
37	c6nhwD_	Alignment	not modelled	41.3	33	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
38	d1adqa1	Alignment	not modelled	40.8	31	PDB header: cell adhesion Chain: B; PDB Molecule: down syndrome cell adhesion molecule (dscam) isoform PDBTitle: crystal structure of down syndrome cell adhesion molecule (dscam)2 isoform 1.30.30, n-terminal eight ig domains
39	c3dmkB_	Alignment	not modelled	40.7	9	PDB header: cell adhesion Chain: A; PDB Molecule: integrin alpha-1; PDBTitle: solution nmr structure of transmembrane and cytosolic regions of2 integrin alpha1 in detergent micelles
40	c2l8sA_	Alignment	not modelled	40.2	30	PDB header: metal binding protein Chain: B; PDB Molecule: putative periplasmic or exported protein; PDBTitle: crystal structure of s.cuep
41	c4gqzB_	Alignment	not modelled	39.3	25	PDB header: transferase Chain: A; PDB Molecule: fibroblast growth factor receptor 2; PDBTitle: solution structure of the d2 domain of the fibroblast2 growth factor
42	c1wvzA_	Alignment	not modelled	37.8	13	PDB header: virus/immune system Chain: 4; PDB Molecule: protein (immunoglobulin light chain); PDBTitle: structure of the complex of a fab fragment of a neutralizing antibody2 with foot and mouth disease virus
43	c1qgc4_	Alignment	not modelled	37.7	16	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
44	d1l6xa1	Alignment	not modelled	37.6	26	PDB header: cell adhesion Chain: D; PDB Molecule: down syndrome cell adhesion molecule, isoform ak; PDBTitle: crystal structure of dscam1 ig7 domain, isoform 5
45	c4wvrD_	Alignment	not modelled	37.6	16	Fold: WSSV envelope protein-like Superfamily: WSSV envelope protein-like Family: WSSV envelope protein-like
46	d2edma1	Alignment	not modelled	36.9	41	PDB header: transport protein/contractile protein Chain: A; PDB Molecule: maltose-binding periplasmic protein,myosin-binding protein PDBTitle: mbp-fusion protein of myosin-binding protein c residues 149-269
47	c4edqA_	Alignment	not modelled	36.3	15	PDB header: immune system Chain: F; PDB Molecule: beta-2-microglobulin; PDBTitle: structure of the beta2_microglobulin involved in amyloidogenesis
48	c2x89F_	Alignment	not modelled	36.1	13	PDB header: immune system Chain: B; PDB Molecule: beta-2-microglobulin; PDBTitle: crystal structure of 6218 tcr in complex with the h2db-pa224
49	c3pqyB_	Alignment	not modelled	36.0	23	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
50	d2fcbal	Alignment	not modelled	35.8	10	PDB header: immune system Chain: G; PDB Molecule: beta-2-microglobulin; PDBTitle: crystal structure of 6218 tcr in complex with the h2db-pa224
51	c3pqyG_	Alignment	not modelled	35.0	23	PDB header: immune system Chain: A; PDB Molecule: ig gamma-1 chain c region; PDBTitle: crystal structure of an isolated, unglycosylated antibody ch2 domain
52	c3dj9A_	Alignment	not modelled	34.7	26	

53	c2k9yB_	Alignment	not modelled	34.0	21	PDB header: transferase Chain: B: PDB Molecule: ephrin type-a receptor 2; PDBTitle: epha2 dimeric structure in the lipidic bicelle at ph 5.0
54	d1fp5a1	Alignment	not modelled	33.8	19	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
55	d1f97a1	Alignment	not modelled	33.8	23	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: V set domains (antibody variable domain-like)
56	d1gxea_	Alignment	not modelled	33.7	18	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
57	d1biha2	Alignment	not modelled	33.6	16	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
58	d2dpka1	Alignment	not modelled	33.3	28	Fold: Immunoglobulin-like beta-sandwich Superfamily: CalX-like Family: CalX-beta domain
59	c6f0ka_	Alignment	not modelled	33.2	14	PDB header: membrane protein Chain: A: PDB Molecule: cytochrome c family protein; PDBTitle: alternative complex iii
60	c2k9yA_	Alignment	not modelled	33.1	21	PDB header: transferase Chain: A: PDB Molecule: ephrin type-a receptor 2; PDBTitle: epha2 dimeric structure in the lipidic bicelle at ph 5.0
61	d1v8ha1	Alignment	not modelled	33.0	10	Fold: Immunoglobulin-like beta-sandwich Superfamily: E set domains Family: SoxZ-like
62	d1epfa2	Alignment	not modelled	32.9	8	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
63	c4kmgA_	Alignment	not modelled	32.7	12	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: lmo2446 protein; PDBTitle: 1.9 angstrom resolution crystal structure of uncharacterized protein2 lmo2446 from listeria monocytogenes egg-e
64	c4x5lB_	Alignment	not modelled	32.7	8	PDB header: cell adhesion Chain: B: PDB Molecule: down syndrome cell adhesion molecule, isoform am; PDBTitle: crystal structure of dscam1 ig7 domain, isoform 9
65	d1fcga1	Alignment	not modelled	31.9	8	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
66	d1uvqb1	Alignment	not modelled	31.2	20	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
67	d1ry7b1	Alignment	not modelled	30.9	13	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
68	c2cknA_	Alignment	not modelled	30.5	6	PDB header: transferase Chain: A: PDB Molecule: basic fibroblast growth factor receptor 1; PDBTitle: nmr structure of the first ig module of mouse fgfr1
69	c3ilqD_	Alignment	not modelled	30.3	19	PDB header: immune system Chain: D: PDB Molecule: beta-2 microglobulin; PDBTitle: structure of mcd1d with bound glycolipid bbgl-2c from2 borrelia burgdorferi
70	d1h3ua1	Alignment	not modelled	30.3	26	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
71	c6fhjA_	Alignment	not modelled	30.0	19	PDB header: hydrolase Chain: A: PDB Molecule: protein,protein; PDBTitle: structural dynamics and catalytic properties of a multi-modular2 xanthanase, native.
72	c3cu1A_	Alignment	not modelled	29.8	16	PDB header: transferase/hormone Chain: A: PDB Molecule: fibroblast growth factor receptor 2; PDBTitle: crystal structure of 2:2:2 fgfr2d2:fgf1:sos complex
73	c3euuA_	Alignment	not modelled	29.8	16	PDB header: transferase Chain: A: PDB Molecule: fibroblast growth factor receptor 2; PDBTitle: crystal structure of the fgfr2 d2 domain
74	c3cafA_	Alignment	not modelled	29.8	16	PDB header: transferase Chain: A: PDB Molecule: fibroblast growth factor receptor 2; PDBTitle: crystal structure of hfgfr2 d2 domain
75	c3euuB_	Alignment	not modelled	29.8	16	PDB header: transferase Chain: B: PDB Molecule: fibroblast growth factor receptor 2; PDBTitle: crystal structure of the fgfr2 d2 domain
76	c3cu1C_	Alignment	not modelled	29.8	16	PDB header: transferase/hormone Chain: C: PDB Molecule: fibroblast growth factor receptor 2; PDBTitle: crystal structure of 2:2:2 fgfr2d2:fgf1:sos complex
77	d2ifga1	Alignment	not modelled	29.6	10	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
78	d1qz1a3	Alignment	not modelled	29.4	16	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
79	c1igtB_	Alignment	not modelled	29.2	14	PDB header: immunoglobulin Chain: B: PDB Molecule: igg2a intact antibody - mab231; PDBTitle: structure of immunoglobulin
						Fold: Immunoglobulin-like beta-sandwich

80	d1zs8a1	Alignment	not modelled	29.0	17	Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
81	d1iray3	Alignment	not modelled	29.0	12	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
82	d1igyb4	Alignment	not modelled	28.9	17	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
83	c3u6gB_	Alignment	not modelled	28.9	38	PDB header: structural genomics, unknown function Chain: B; PDB Molecule: uncharacterized protein duf4425; PDBTitle: crystal structure of a domain of unknown function, duf4425 (bvu_3708)2 from bacteroides vulgatus atcc 8482 at 1.35 a resolution
84	d1hdma1	Alignment	not modelled	28.8	25	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
85	d1hwxa2	Alignment	not modelled	28.5	20	Fold: Aminoacid dehydrogenase-like, N-terminal domain Superfamily: Aminoacid dehydrogenase-like, N-terminal domain Family: Aminoacid dehydrogenases
86	d1iakb1	Alignment	not modelled	28.4	21	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
87	d1klub1	Alignment	not modelled	28.2	21	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
88	c2e7cA_	Alignment	not modelled	28.2	16	PDB header: structural protein Chain: A; PDB Molecule: myosin-binding protein c, fast-type; PDBTitle: solution structure of the 6th ig-like domain from human2 myosin-binding protein c, fast-type
89	c4jvwD_	Alignment	not modelled	27.8	17	PDB header: protein binding Chain: D; PDB Molecule: ig mu chain c region secreted form; PDBTitle: igm c4-domain from mouse
90	d1dn0b2	Alignment	not modelled	27.7	13	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
91	c3o9wB_	Alignment	not modelled	27.6	19	PDB header: immune system Chain: B; PDB Molecule: beta-2-microglobulin; PDBTitle: recognition of a glycolipid antigen by the inkt cell tcr
92	d3b5ha1	Alignment	not modelled	27.6	12	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
93	d2fb4I2	Alignment	not modelled	27.4	32	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
94	c3rbgB_	Alignment	not modelled	27.2	15	PDB header: immune system Chain: B; PDB Molecule: cytotoxic and regulatory t-cell molecule; PDBTitle: crystal structure analysis of class-i mhc restricted t-cell associated2 molecule
95	c2n56A_	Alignment	not modelled	27.2	15	PDB header: protein binding Chain: A; PDB Molecule: obscurin; PDBTitle: ig59 domain of human obscurin a
96	d1rhfa1	Alignment	not modelled	27.2	18	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: V set domains (antibody variable domain-like)
97	c2vajA_	Alignment	not modelled	26.8	13	PDB header: cell adhesion Chain: A; PDB Molecule: neural cell adhesion molecule 2; PDBTitle: crystal structure of ncam2 ig1 (i4122 cell unit)
98	c5xviA_	Alignment	not modelled	26.7	29	PDB header: oxidoreductase Chain: A; PDB Molecule: glutamate dehydrogenase; PDBTitle: crystal structure of aspergillus niger apo- glutamate dehydrogenase
99	d1o0va1	Alignment	not modelled	26.4	15	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
100	d1iray1	Alignment	not modelled	26.3	8	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
101	c6nhyb_	Alignment	not modelled	26.2	33	PDB header: immune system Chain: B; PDB Molecule: tumor necrosis factor receptor superfamily member 10b; PDBTitle: structure of the transmembrane domain of the death receptor 5 mutant2 (g217y) - trimer only
102	c2brqA_	Alignment	not modelled	26.1	15	PDB header: structural protein Chain: A; PDB Molecule: filamin a; PDBTitle: crystal structure of the filamin a repeat 21 complexed with2 the integrin beta7 cytoplasmic tail peptide
103	c2w0pB_	Alignment	not modelled	26.1	15	PDB header: cell adhesion Chain: B; PDB Molecule: filamin-a; PDBTitle: crystal structure of the filamin a repeat 21 complexed with2 the migfilin peptide
104	c3iswb_	Alignment	not modelled	26.1	15	PDB header: structural protein Chain: B; PDB Molecule: filamin-a; PDBTitle: crystal structure of filamin-a immunoglobulin-like repeat 21 bound to2 an n-terminal peptide of cfr
105	c3cnkB_	Alignment	not modelled	25.8	17	PDB header: structural protein Chain: B; PDB Molecule: filamin-a; PDBTitle: crystal structure of the dimerization domain of human filamin a
106	c1cqkA_	Alignment	not modelled	25.6	27	PDB header: immune system Chain: A; PDB Molecule: ch3 domain of mak33 antibody; PDBTitle: crystal structure of the ch3 domain from the mak33

						antibody
107	c1cqkB_	Alignment	not modelled	25.6	27	PDB header: immune system Chain: B: PDB Molecule: ch3 domain of mak33 antibody; PDBTitle: crystal structure of the ch3 domain from the mak33 antibody
108	d1cqka_	Alignment	not modelled	25.6	27	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
109	d1l6za2	Alignment	not modelled	25.2	17	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
110	c2id5D_	Alignment	not modelled	25.1	14	PDB header: ligand binding protein,membrane protein Chain: D: PDB Molecule: leucine rich repeat neuronal 6a; PDBTitle: crystal structure of the lingo-1 ectodomain
111	d1g1ca_	Alignment	not modelled	25.1	18	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: I set domains
112	c1ntIA_	Alignment	not modelled	25.0	15	PDB header: immune system Chain: A: PDB Molecule: complement component receptor 1-like protein,ig gamma-1 PDBTitle: model of mouse crry-ig determined by solution scattering, curve2 fitting and homology modelling
113	c6nhyC_	Alignment	not modelled	24.6	33	PDB header: immune system Chain: C: PDB Molecule: tumor necrosis factor receptor superfamily member 10b; PDBTitle: structure of the transmembrane domain of the death receptor 5 mutant2 (g217y) - trimer only
114	c6nhyA_	Alignment	not modelled	24.6	33	PDB header: immune system Chain: A: PDB Molecule: tumor necrosis factor receptor superfamily member 10b; PDBTitle: structure of the transmembrane domain of the death receptor 5 mutant2 (g217y) - trimer only
115	c3i0uA_	Alignment	not modelled	24.6	25	PDB header: lyase Chain: A: PDB Molecule: phosphothreonine lyase ospf; PDBTitle: structure of the type iii effector/phosphothreonine lyase ospf from2 shigella flexneri
116	d1es0b1	Alignment	not modelled	24.4	21	Fold: Immunoglobulin-like beta-sandwich Superfamily: Immunoglobulin Family: C1 set domains (antibody constant domain-like)
117	c2yuzA_	Alignment	not modelled	24.3	21	PDB header: immune system Chain: A: PDB Molecule: myosin-binding protein c, slow-type; PDBTitle: solution structure of 4th immunoglobulin domain of slow2 type myosin-binding protein c
118	d1bgva2	Alignment	not modelled	24.3	27	Fold: Aminoacid dehydrogenase-like, N-terminal domain Superfamily: Aminoacid dehydrogenase-like, N-terminal domain Family: Aminoacid dehydrogenases
119	c3sgjA_	Alignment	not modelled	24.1	12	PDB header: immune system Chain: A: PDB Molecule: human fc fragment; PDBTitle: unique carbohydrate-carbohydrate interactions are required for high2 affinity binding between fcgiii and antibodies lacking core fucose
120	c1xmeB_	Alignment	not modelled	24.0	6	PDB header: oxidoreductase Chain: B: PDB Molecule: cytochrome c oxidase polypeptide ii; PDBTitle: structure of recombinant cytochrome ba3 oxidase from thermus2 thermophilus