
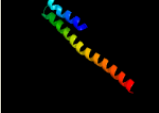
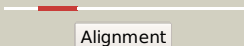
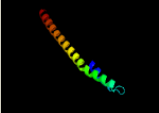
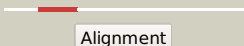
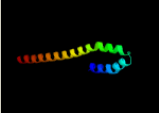
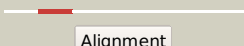

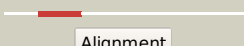
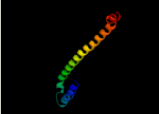
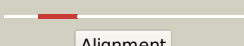
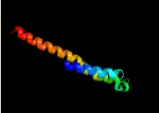
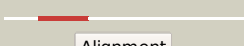
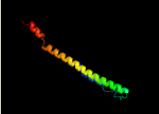

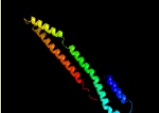

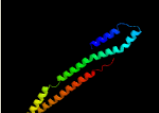




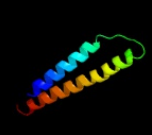
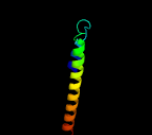
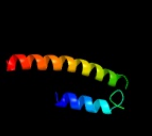

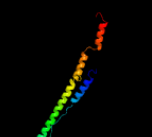
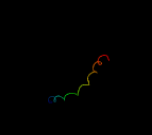
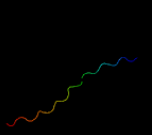

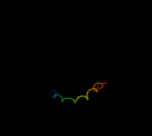


Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD3616c_(-)_4055376_4056554
Date	Fri Aug 9 18:20:30 BST 2019
Unique Job ID	12a67a633c580f50

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c4iogD_	 Alignment		96.8	12	PDB header: unknown function Chain: D: PDB Molecule: secreted protein esxb; PDBTitle: the crystal structure of a secreted protein esxb (wild-type, in p212 space group) from bacillus anthracis str. Sterne
2	c3zbhC_	 Alignment		96.7	13	PDB header: unknown function Chain: C: PDB Molecule: esxa; PDBTitle: geobacillus thermodenitrificans esxa crystal form i
3	c3gvmA_	 Alignment		96.6	18	PDB header: viral protein Chain: A: PDB Molecule: putative uncharacterized protein sag1039; PDBTitle: structure of the homodimeric wxg-100 family protein from streptococcus2 agalactiae
4	c2vs0B_	 Alignment		96.3	15	PDB header: cell invasion Chain: B: PDB Molecule: virulence factor esxa; PDBTitle: structural analysis of homodimeric staphylococcal aureus2 virulence factor esxa
5	c4lwsA_	 Alignment		96.2	5	PDB header: unknown function Chain: A: PDB Molecule: uncharacterized protein; PDBTitle: esxa : esxb (semet) hetero-dimer from thermomonospora curvata
6	c5xfsB_	 Alignment		95.9	15	PDB header: protein transport Chain: B: PDB Molecule: ppe family protein ppe15; PDBTitle: crystal structure of pe8-ppe15 in complex with espg5 from m.2 tuberculosis
7	c4wj2A_	 Alignment		95.7	15	PDB header: unknown function Chain: A: PDB Molecule: antigen mtb48; PDBTitle: mycobacterial protein
8	d2g38b1	 Alignment		95.7	7	Fold: Ferritin-like Superfamily: PE/PPE dimer-like Family: PPE
9	c2g38B_	 Alignment		95.7	7	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: ppe family protein; PDBTitle: a pe/ppe protein complex from mycobacterium tuberculosis
10	d1wa8b1	 Alignment		95.2	18	Fold: Ferritin-like Superfamily: EsxAB dimer-like Family: ESAT-6 like
11	c4lwsB_	 Alignment		95.1	16	PDB header: unknown function Chain: B: PDB Molecule: uncharacterized protein; PDBTitle: esxa : esxb (semet) hetero-dimer from thermomonospora curvata

12	d1wa8a1	Alignment		93.8	6	Fold: Ferritin-like Superfamily: EsxAB dimer-like Family: ESAT-6 like
13	c4i0xA	Alignment		93.0	20	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: esat-6-like protein mab_3112; PDBTitle: crystal structure of the mycobacterium abscessus esxf (mab_3112-2 mab_3113) complex
14	c4i0xJ	Alignment		92.9	19	PDB header: structural genomics, unknown function Chain: J: PDB Molecule: esat-6-like protein mab_3113; PDBTitle: crystal structure of the mycobacterium abscessus esxf (mab_3112-2 mab_3113) complex
15	c2kg7B	Alignment		89.0	13	PDB header: unknown function Chain: B: PDB Molecule: esat-6-like protein esxh; PDBTitle: structure and features of the complex formed by the tuberculosis2 virulence factors rv0287 and rv0288
16	c4xy3A	Alignment		76.1	18	PDB header: protein transport Chain: A: PDB Molecule: esx-1 secretion-associated protein espb; PDBTitle: structure of esx-1 secreted protein espb
17	c2cuoC	Alignment		66.0	19	PDB header: structural protein Chain: C: PDB Molecule: collagen model peptide (pro-pro-gly)9; PDBTitle: collagen model peptide (pro-pro-gly)9
18	c2cuoF	Alignment		66.0	19	PDB header: structural protein Chain: F: PDB Molecule: collagen model peptide (pro-pro-gly)9; PDBTitle: collagen model peptide (pro-pro-gly)9
19	c2cuoB	Alignment		58.7	15	PDB header: structural protein Chain: B: PDB Molecule: collagen model peptide (pro-pro-gly)9; PDBTitle: collagen model peptide (pro-pro-gly)9
20	c2cuoA	Alignment		58.7	15	PDB header: structural protein Chain: A: PDB Molecule: collagen model peptide (pro-pro-gly)9; PDBTitle: collagen model peptide (pro-pro-gly)9
21	c2cuoE	Alignment	not modelled	58.7	15	PDB header: structural protein Chain: E: PDB Molecule: collagen model peptide (pro-pro-gly)9; PDBTitle: collagen model peptide (pro-pro-gly)9
22	c2cuoD	Alignment	not modelled	58.7	15	PDB header: structural protein Chain: D: PDB Molecule: collagen model peptide (pro-pro-gly)9; PDBTitle: collagen model peptide (pro-pro-gly)9
23	c3ah9A	Alignment	not modelled	51.0	17	PDB header: structural protein Chain: A: PDB Molecule: collagen-like peptide; PDBTitle: crystal structure of (pro-pro-gly)9 at 1.1 a resolution
24	c3ah9F	Alignment	not modelled	50.8	17	PDB header: structural protein Chain: F: PDB Molecule: collagen-like peptide; PDBTitle: crystal structure of (pro-pro-gly)9 at 1.1 a resolution
25	c3ah9E	Alignment	not modelled	49.2	16	PDB header: structural protein Chain: E: PDB Molecule: collagen-like peptide; PDBTitle: crystal structure of (pro-pro-gly)9 at 1.1 a resolution
26	c3ah9B	Alignment	not modelled	49.2	16	PDB header: structural protein Chain: B: PDB Molecule: collagen-like peptide; PDBTitle: crystal structure of (pro-pro-gly)9 at 1.1 a resolution
27	c3ah9C	Alignment	not modelled	49.2	16	PDB header: structural protein Chain: C: PDB Molecule: collagen-like peptide; PDBTitle: crystal structure of (pro-pro-gly)9 at 1.1 a resolution
28	c3a19B	Alignment	not modelled	45.7	23	PDB header: structural protein Chain: B: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-oog-(ppg)4_h monoclinic, twinned crystal
						PDB header: structural protein Chain: A: PDB Molecule: collagen model peptides (pro-pro-gly)4-

29	c2d3hA_	Alignment	not modelled	45.7	23	hyp-hyp- PDBTitle: crystal structures of collagen model peptides (pro-pro-gly)2 4-hyp-hyp-gly-(pro-pro-gly)4 PDB header: structural protein Chain: B: PDB Molecule: collagen model peptides (pro-pro-gly)4-hyp-hyp- PDBTitle: crystal structures of collagen model peptides (pro-pro-gly)2 4-hyp-hyp-gly-(pro-pro-gly)4
30	c2d3hB_	Alignment	not modelled	45.7	23	hyp-hyp- PDBTitle: crystal structures of collagen model peptides (pro-pro-gly)2 4-hyp-hyp-gly-(pro-pro-gly)4 PDB header: structural protein Chain: D: PDB Molecule: collagen-like peptide; PDBTitle: crystal structure of (pro-pro-gly)9 at 1.1 a resolution
31	c3ah9D_	Alignment	not modelled	44.1	14	PDB header: structural protein Chain: E: PDB Molecule: collagen model peptides (pro-pro-gly)4-hyp-hyp- PDBTitle: crystal structures of collagen model peptides (pro-pro-gly)2 4-hyp-hyp-gly-(pro-pro-gly)4 PDB header: structural protein Chain: D: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-oog-(ppg)4_h monoclinic, twinned crystal
32	c2d3hE_	Alignment	not modelled	43.3	22	PDB header: structural protein Chain: A: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-oog-(ppg)4_h monoclinic, twinned crystal
33	c3a19D_	Alignment	not modelled	40.2	21	PDB header: structural protein Chain: C: PDB Molecule: collagen model peptides (pro-pro-gly)4-hyp-hyp- PDBTitle: crystal structures of collagen model peptides (pro-pro-gly)2 4-hyp-hyp-gly-(pro-pro-gly)4 PDB header: viral protein Chain: A: PDB Molecule: polymerase cofactor vp35; PDBTitle: crystal structure of the oligomerization domain of vp35 from reston2 virus, mercury derivative
34	c3a19A_	Alignment	not modelled	39.8	22	PDB header: structural protein Chain: B: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-oog-(ppg)4, monoclinic, twinned crystal
35	c2d3hC_	Alignment	not modelled	39.8	22	PDB header: structural protein Chain: A: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-oog-(ppg)4, monoclinic, twinned crystal
36	c6gbrA_	Alignment	not modelled	39.8	28	PDB header: structural protein Chain: C: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-oog-(ppg)4_h monoclinic, twinned crystal
37	c3a08B_	Alignment	not modelled	38.3	19	PDB header: structural protein Chain: E: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-oog-(ppg)4_h monoclinic, twinned crystal
38	c3a08A_	Alignment	not modelled	38.3	19	PDB header: structural protein Chain: C: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-oog-(ppg)4, monoclinic, twinned crystal
39	c3a19C_	Alignment	not modelled	38.1	21	PDB header: structural protein Chain: E: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-oog-(ppg)4_h monoclinic, twinned crystal
40	c3a19E_	Alignment	not modelled	38.1	21	PDB header: structural protein Chain: C: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-oog-(ppg)4, monoclinic, twinned crystal
41	c3a08C_	Alignment	not modelled	27.5	24	PDB header: structural protein Chain: F: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-ovg-(ppg)4, monoclinic, twinned crystal
42	c3a08E_	Alignment	not modelled	26.7	23	PDB header: structural protein Chain: F: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-ovg-(ppg)4, monoclinic, twinned crystal
43	c3a0mF_	Alignment	not modelled	26.6	13	PDB header: structural protein Chain: F: PDB Molecule: host-guest peptide (pro-pro-gly)4-(pro-allohyp- PDBTitle: host-guest peptide (pro-pro-gly)4-(pro-allohyp-gly)-(pro-2 pro-gly)4
44	c1x1kF_	Alignment	not modelled	26.4	26	PDB header: structural protein Chain: F: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-oog-(ppg)4_h monoclinic, twinned crystal
45	c3a19F_	Alignment	not modelled	24.1	22	PDB header: structural protein Chain: D: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-oog-(ppg)4, monoclinic, twinned crystal
46	c3a08D_	Alignment	not modelled	24.1	22	PDB header: structural protein Chain: D: PDB Molecule: collagen model peptides (pro-pro-gly)4-hyp-hyp- PDBTitle: crystal structures of collagen model peptides (pro-pro-gly)2 4-hyp-hyp-gly-(pro-pro-gly)4
47	c2d3hD_	Alignment	not modelled	24.1	22	Fold: Orange carotenoid protein, N-terminal domain Superfamily: Orange carotenoid protein, N-terminal domain Family: Orange carotenoid protein, N-terminal domain
48	d1m98a1	Alignment	not modelled	21.2	15	PDB header: structural protein Chain: C: PDB Molecule: collagen alpha-1(i) chain, collagen alpha-3(ix) chain; PDBTitle: crystal structure of the type ix collagen nc2 heterotrimerization2 domain with a guest fragment a2a1a1 of type i collagen (native form)
49	c5ctiC_	Alignment	not modelled	19.1	24	PDB header: structural protein Chain: F: PDB Molecule: collagen model peptides (pro-pro-gly)4-hyp-hyp- PDBTitle: crystal structures of collagen model peptides (pro-pro-gly)2 4-hyp-hyp-gly-(pro-pro-gly)4
50	c2d3hF_	Alignment	not modelled	18.8	23	PDB header: plant protein Chain: A: PDB Molecule: wa352; PDBTitle: crystal structure of wa352 from oryza sativa
51	c5zt3A_	Alignment	not modelled	17.0	23	PDB header: motor protein/metal-bindng protein Chain: A: PDB Molecule: unconventional myosin-vi, green fluorescent protein; PDBTitle: myosin vi (mdinsert2-gfp fusion) pre-powerstroke state (mg.adp.alf4)
52	c4anjA_	Alignment	not modelled	16.6	16	

53	c3a08F_	Alignment	not modelled	16.4	24	PDB header: structural protein Chain: F: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-oog-(ppg)4, monoclinic, twinned crystal
54	c2k48A_	Alignment	not modelled	16.1	26	PDB header: viral protein Chain: A: PDB Molecule: nucleoprotein; PDBTitle: nmr structure of the n-terminal coiled coil domain of the2 andes hantavirus nucleocapsid protein
55	c1m98A_	Alignment	not modelled	16.0	15	PDB header: unknown function Chain: A: PDB Molecule: orange carotenoid protein; PDBTitle: crystal structure of orange carotenoid protein
56	c3admF_	Alignment	not modelled	14.0	23	PDB header: structural protein Chain: F: PDB Molecule: collagen-like peptide; PDBTitle: crystal structure of (pro-pro-gly)4-hyp-ser-gly-(pro-pro-gly)4
57	c3admE_	Alignment	not modelled	14.0	23	PDB header: structural protein Chain: E: PDB Molecule: collagen-like peptide; PDBTitle: crystal structure of (pro-pro-gly)4-hyp-ser-gly-(pro-pro-gly)4
58	c3admB_	Alignment	not modelled	14.0	23	PDB header: structural protein Chain: B: PDB Molecule: collagen-like peptide; PDBTitle: crystal structure of (pro-pro-gly)4-hyp-ser-gly-(pro-pro-gly)4
59	c3admA_	Alignment	not modelled	14.0	23	PDB header: structural protein Chain: A: PDB Molecule: collagen-like peptide; PDBTitle: crystal structure of (pro-pro-gly)4-hyp-ser-gly-(pro-pro-gly)4
60	c2i9nA_	Alignment	not modelled	13.9	46	PDB header: de novo protein Chain: A: PDB Molecule: mhb4a peptide; PDBTitle: design of bivalent miniprotein consisting of two2 independent elements, a b-hairpin peptide and a-helix3 peptide, tethered by four glycines
61	c6mcjA_	Alignment	not modelled	12.7	15	PDB header: protein binding Chain: A: PDB Molecule: orange carotenoid-binding protein; PDBTitle: structure of helical carotenoid protein 2 from fremyella diplosiphon
62	c2d3fE_	Alignment	not modelled	12.3	19	PDB header: structural protein Chain: E: PDB Molecule: collagen model peptides (pro-pro-gly)4-pro-hyp- PDBTitle: crystal structures of collagen model peptides (pro-pro-gly)2 4-pro-hyp-gly-(pro-pro-gly)4
63	c2d3fD_	Alignment	not modelled	12.3	19	PDB header: structural protein Chain: D: PDB Molecule: collagen model peptides (pro-pro-gly)4-pro-hyp- PDBTitle: crystal structures of collagen model peptides (pro-pro-gly)2 4-pro-hyp-gly-(pro-pro-gly)4
64	c2d3fF_	Alignment	not modelled	12.3	19	PDB header: structural protein Chain: F: PDB Molecule: collagen model peptides (pro-pro-gly)4-pro-hyp- PDBTitle: crystal structures of collagen model peptides (pro-pro-gly)2 4-pro-hyp-gly-(pro-pro-gly)4
65	c1x1kA_	Alignment	not modelled	12.3	19	PDB header: structural protein Chain: A: PDB Molecule: host-guest peptide (pro-pro-gly)4-(pro-allohyp- PDBTitle: host-guest peptide (pro-pro-gly)4-(pro-allohyp-gly)-(pro-2 pro-gly)4
66	c1x1kD_	Alignment	not modelled	12.3	19	PDB header: structural protein Chain: D: PDB Molecule: host-guest peptide (pro-pro-gly)4-(pro-allohyp- PDBTitle: host-guest peptide (pro-pro-gly)4-(pro-allohyp-gly)-(pro-2 pro-gly)4
67	c1x1kC_	Alignment	not modelled	12.3	19	PDB header: structural protein Chain: C: PDB Molecule: host-guest peptide (pro-pro-gly)4-(pro-allohyp- PDBTitle: host-guest peptide (pro-pro-gly)4-(pro-allohyp-gly)-(pro-2 pro-gly)4
68	c1x1kB_	Alignment	not modelled	12.3	19	PDB header: structural protein Chain: B: PDB Molecule: host-guest peptide (pro-pro-gly)4-(pro-allohyp- PDBTitle: host-guest peptide (pro-pro-gly)4-(pro-allohyp-gly)-(pro-2 pro-gly)4
69	c2vqkA_	Alignment	not modelled	12.2	19	PDB header: membrane protein Chain: A: PDB Molecule: uncharacterized protein cgl0972; PDBTitle: crystal structure of porb from corynebacterium glutamicum (2 crystal form iv)
70	c2vqkD_	Alignment	not modelled	11.1	19	PDB header: membrane protein Chain: D: PDB Molecule: uncharacterized protein cgl0972; PDBTitle: crystal structure of porb from corynebacterium glutamicum (2 crystal form iii)
71	c4rvqA_	Alignment	not modelled	10.8	17	PDB header: protein binding Chain: A: PDB Molecule: pre-mrna splicing helicase-like protein; PDBTitle: pwi-like domain of chaetomium thermophilum brr2
72	c3admC_	Alignment	not modelled	10.3	24	PDB header: structural protein Chain: C: PDB Molecule: collagen-like peptide; PDBTitle: crystal structure of (pro-pro-gly)4-hyp-ser-gly-(pro-pro-gly)4
73	c5ctdA_	Alignment	not modelled	10.2	24	PDB header: structural protein Chain: A: PDB Molecule: collagen alpha-1(i) chain,collagen alpha-1(ix) chain; PDBTitle: crystal structure of the type ix collagen nc2 heterotrimerization2 domain with a guest fragment a2a1a1 of type i collagen
74	d2ffga1	Alignment	not modelled	9.5	29	Fold: Ykuj-like Superfamily: Ykuj-like Family: Ykuj-like
75	c6j9hA_	Alignment	not modelled	9.5	25	PDB header: peptide binding protein/hydrolase Chain: A: PDB Molecule: small vasohibin-binding protein; PDBTitle: crystal structure of svbp-vash1 complex
76	c3n42D_	Alignment	not modelled	9.1	15	PDB header: unknown function Chain: D: PDB Molecule: predicted protein;

76	c3p4zD_	Alignment	not modelled	9.1	15	PDBTitle: structure of gfcc (ymcb), protein encoded by the e. coli group 42 capsule operon
77	d1a9xa1	Alignment	not modelled	9.1	17	Fold: Carbamoyl phosphate synthetase, large subunit connection domain Superfamily: Carbamoyl phosphate synthetase, large subunit connection domain Family: Carbamoyl phosphate synthetase, large subunit connection domain
78	c6gboG_	Alignment	not modelled	9.0	21	PDB header: viral protein Chain: G: PDB Molecule: polymerase cofactor vp35; PDBTitle: crystal structure of the oligomerization domain of vp35 from ebola2 virus
79	c3a0aB_	Alignment	not modelled	8.8	20	PDB header: structural protein Chain: B: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-opg-(ppg)4, monoclinic, twinned crystal
80	c2ic6B_	Alignment	not modelled	8.7	26	PDB header: viral protein Chain: B: PDB Molecule: nucleocapsid protein; PDBTitle: the coiled-coil domain (residues 1-75) structure of the sin nombre2 virus nucleocapsid protein
81	c3abnA_	Alignment	not modelled	8.4	12	PDB header: structural protein Chain: A: PDB Molecule: collagen-like peptide; PDBTitle: crystal structure of (pro-pro-gly)4-hyp-asp-gly-(pro-pro-gly)4 at 1.022 a
82	c2i9oA_	Alignment	not modelled	7.6	44	PDB header: de novo protein Chain: A: PDB Molecule: mhb8a peptide; PDBTitle: design of bivalent miniprotein consisting of two2 independent elements, a b-hairpin peptide and a-helix3 peptide, tethered by eight glycines
83	c3admD_	Alignment	not modelled	7.4	20	PDB header: structural protein Chain: D: PDB Molecule: collagen-like peptide; PDBTitle: crystal structure of (pro-pro-gly)4-hyp-ser-gly-(pro-pro-gly)4
84	c2d3fA_	Alignment	not modelled	7.3	21	PDB header: structural protein Chain: A: PDB Molecule: collagen model peptides (pro-pro-gly)4-pro-hyp- PDBTitle: crystal structures of collagen model peptides (pro-pro-gly)2 4-pro-hyp-gly-(pro-pro-gly)4
85	c2d3fB_	Alignment	not modelled	7.3	21	PDB header: structural protein Chain: B: PDB Molecule: collagen model peptides (pro-pro-gly)4-pro-hyp- PDBTitle: crystal structures of collagen model peptides (pro-pro-gly)2 4-pro-hyp-gly-(pro-pro-gly)4
86	c1x1kE_	Alignment	not modelled	7.3	21	PDB header: structural protein Chain: E: PDB Molecule: host-guest peptide (pro-pro-gly)4-(pro-allohyp- PDBTitle: host-guest peptide (pro-pro-gly)4-(pro-allohyp-gly)-(pro-2 pro-gly)4
87	c2d3fC_	Alignment	not modelled	7.3	21	PDB header: structural protein Chain: C: PDB Molecule: collagen model peptides (pro-pro-gly)4-pro-hyp- PDBTitle: crystal structures of collagen model peptides (pro-pro-gly)2 4-pro-hyp-gly-(pro-pro-gly)4
88	c1mofA_	Alignment	not modelled	7.0	14	PDB header: viral protein Chain: A: PDB Molecule: moloney murine leukemia virus p15; PDBTitle: coat protein
89	c3j38T_	Alignment	not modelled	6.7	29	PDB header: ribosome Chain: T: PDB Molecule: 40s ribosomal protein s19a; PDBTitle: structure of the d. melanogaster 40s ribosomal proteins
90	c2mc7A_	Alignment	not modelled	6.7	17	PDB header: membrane protein Chain: A: PDB Molecule: regulatory peptide; PDBTitle: structure of salmonella mgtr
91	c6j7bB_	Alignment	not modelled	6.6	25	PDB header: hydrolase Chain: B: PDB Molecule: small vasohibin-binding protein; PDBTitle: crystal structure of vash1-svbp in complex with epoxy
92	c3ggzC_	Alignment	not modelled	6.6	11	PDB header: protein transport, endocytosis Chain: C: PDB Molecule: increased sodium tolerance protein 1; PDBTitle: crystal structure of s.cerevisiae ist1 n-terminal domain in complex2 with did2 mim motif
93	c4jf3A_	Alignment	not modelled	6.1	11	PDB header: viral protein Chain: A: PDB Molecule: envelope glycoprotein; PDBTitle: crystal structure of the mpvm tm retroviral fusion core
94	c5tszA_	Alignment	not modelled	6.0	25	PDB header: cell invasion Chain: A: PDB Molecule: pv cell-traversal protein; PDBTitle: crystal structure of plasmodium vivax celtsos
95	c5ctdB_	Alignment	not modelled	6.0	21	PDB header: structural protein Chain: B: PDB Molecule: collagen alpha-2(i) chain,collagen alpha-2(ix) chain; PDBTitle: crystal structure of the type ix collagen nc2 heterotrimerization2 domain with a guest fragment a2a1a1 of type i collagen
96	c3abnC_	Alignment	not modelled	6.0	13	PDB header: structural protein Chain: C: PDB Molecule: collagen-like peptide; PDBTitle: crystal structure of (pro-pro-gly)4-hyp-asp-gly-(pro-pro-gly)4 at 1.022 a
97	c3a0mA_	Alignment	not modelled	6.0	24	PDB header: structural protein Chain: A: PDB Molecule: collagen-like peptide; PDBTitle: structure of (ppg)4-ovg-(ppg)4, monoclinic, twinned crystal
98	c6b87C_	Alignment	not modelled	5.8	14	PDB header: membrane protein Chain: C: PDB Molecule: tmhc2_e; PDBTitle: crystal structure of transmembrane protein tmhc2_e
99	d1emya_	Alignment	not modelled	5.8	15	Fold: Globin-like Superfamily: Globin-like Family: Globins