Phyre2

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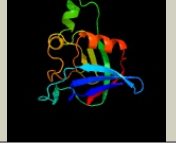
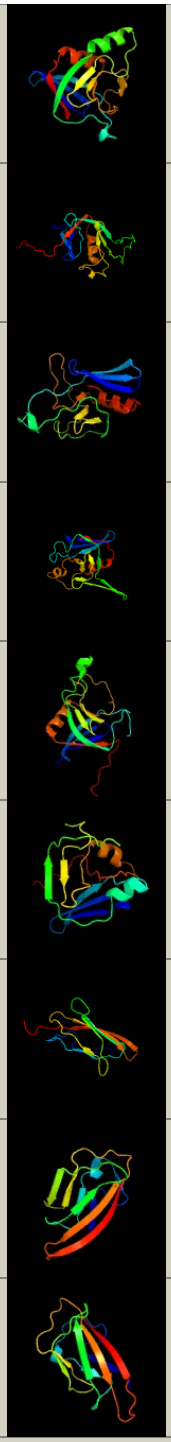
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c4z7aA_	Alignment		100.0	100	**PDB header:**transferase **Chain:** A: **PDB Molecule:**mycobacterium tuberculosis (3,3)l,d-transpeptidase type 5; **PDBTitle:** structural and biochemical characterization of a non-functionally2 redundant m. tuberculosis (3,3) l,d-transpeptidase, ldtmt5.
2	c3vynB_	Alignment		100.0	35	**PDB header:**transferase **Chain:** B: **PDB Molecule:**probable conserved lipoprotein lpps; **PDBTitle:** crystal structure of mycobacterium tuberculosis l,d-transpeptidase2 ldtmt2 n55 truncation mutant (resideus 55-408)
3	c3u1qB_	Alignment		100.0	35	**PDB header:**peptidoglycan binding protein **Chain:** B: **PDB Molecule:**mycobacteria tuberculosis ld-transpeptidase type 2; **PDBTitle:** crystal structure of m. tuberculosis ld-transpeptidase type 2 with 2-2 mercaptoethanol
4	c4xvoB_	Alignment		100.0	31	**PDB header:**transferase **Chain:** B: **PDB Molecule:**l,d-transpeptidase; **PDBTitle:** l,d-transpeptidase from mycobacterium smegmatis
5	c4jmxA_	Alignment		100.0	33	**PDB header:**transferase/transferase inhibitor **Chain:** A: **PDB Molecule:**probable l,d-transpeptidase ldta; **PDBTitle:** structure of ld transpeptidase ldtmt1 in complex with imipenem
6	c4k73A_	Alignment		100.0	35	**PDB header:**transferase **Chain:** A: **PDB Molecule:**l,d-transpeptidase; **PDBTitle:** x-ray crystal structure of an l,d-transpeptidase from mycobacterium2 tuberculosis h37rv
7	c4hu2A_	Alignment		100.0	39	**PDB header:**unknown function **Chain:** A: **PDB Molecule:**probable conserved lipoprotein lpps; **PDBTitle:** crystal structure of ldtmt2, a l,d-transpeptidase from mycobacterium2 tuberculosis: domain a and b
8	c2hklB_	Alignment		100.0	16	**PDB header:**transferase **Chain:** B: **PDB Molecule:**l,d-transpeptidase; **PDBTitle:** crystal structure of enterococcus faecium l,d-2 transpeptidase c442s mutant
9	c5bmqA_	Alignment		100.0	22	**PDB header:**hydrolase **Chain:** A: **PDB Molecule:**erfk/ybis/ycfs/ynhg family protein; **PDBTitle:** crystal structure of l,d-transpeptidase (yku) from stackebrandtia2 nassauensis
10	d1zata1	Alignment		100.0	18	**Fold:**L,D-transpeptidase catalytic domain-like **Superfamily:**L,D-transpeptidase catalytic domain-like **Family:**L,D-transpeptidase catalytic domain-like
11	d1y7ma1	Alignment		100.0	15	**Fold:**L,D-transpeptidase catalytic domain-like **Superfamily:**L,D-transpeptidase catalytic domain-like **Family:**L,D-transpeptidase catalytic domain-like

12	c1y7mB_	Alignment		100.0	15	**PDB header:**structural genomics, unknown function **Chain:** B: **PDB Molecule:**hypothetical protein bsu14040; **PDBTitle:** crystal structure of the b. subtilis ykud protein at 2 a2 resolution
13	c4lzhA_	Alignment		99.9	13	**PDB header:**transferase **Chain:** A: **PDB Molecule:**l,d-transpeptidase; **PDBTitle:** l,d-transpeptidase from klebsiella pneumoniae
14	c4lpqA_	Alignment		99.9	15	**PDB header:**transferase **Chain:** A: **PDB Molecule:**erfk/ybis/ycfs/ynhg family protein; **PDBTitle:** crystal structure of the l,d-transpeptidase (residues 123-326) from2 xylanimonas cellulosilytica dsm 15894
15	c4xxtA_	Alignment		99.9	19	**PDB header:**hydrolase **Chain:** A: **PDB Molecule:**fusion of predicted zn-dependent amidase/peptidase (cell **PDBTitle:** crystal structure of fused zn-dependent2 amidase/peptidase/peptodoglycan-binding domain-containing protein3 from clostridium acetobutylicum atcc 824
16	c4y4vB_	Alignment		99.8	16	**PDB header:**hydrolase **Chain:** B: **PDB Molecule:**conserved hypothetical secreted protein; **PDBTitle:** structure of helicobacter pylori csd6 in the d-ala-bound state
17	c6ntwA_	Alignment		99.3	15	**PDB header:**transferase/transferase inhibitor **Chain:** A: **PDB Molecule:**probable l,d-transpeptidase ycbb; **PDBTitle:** crystal structure of e. coli ycbb
18	c5icuA_	Alignment		98.0	16	**PDB header:**chaperone **Chain:** A: **PDB Molecule:**copc; **PDBTitle:** the crystal structure of copc from methylosinus trichosporium ob3b
19	c6nfqC_	Alignment		97.7	22	**PDB header:**metal binding protein **Chain:** C: **PDB Molecule:**copc; **PDBTitle:** copc from pseudomonas fluorescens
20	d2c9qa1	Alignment		97.3	24	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**E set domains **Family:**Copper resistance protein C (CopC, PcoC)
21	d1ix2a_	Alignment	not modelled	97.0	25	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**E set domains **Family:**Copper resistance protein C (CopC, PcoC)
22	c5n1tM_	Alignment	not modelled	96.6	14	**PDB header:**oxidoreductase **Chain:** M: **PDB Molecule:**copc; **PDBTitle:** crystal structure of complex between flavocytochrome c and copper2 chaperone copc from t. paradoxus
23	d2cuha1	Alignment	not modelled	93.6	18	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
24	d2mfna1	Alignment	not modelled	93.2	25	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
25	c3t1wA_	Alignment	not modelled	92.5	16	**PDB header:**protein binding **Chain:** A: **PDB Molecule:**four-domain fibronectin fragment; **PDBTitle:** structure of the four-domain fragment fn7b89 of oncofetal fibronectin
26	d1tena_	Alignment	not modelled	92.4	26	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
27	c3tesD_	Alignment	not modelled	92.0	21	**PDB header:**de novo protein **Chain:** D: **PDB Molecule:**tencon; **PDBTitle:** crystal structure of tencon
28	d2vkwa2	Alignment	not modelled	92.0	17	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
						PDB header:unknown function **Chain:** B: **PDB Molecule:**ten-d3;

29	c3b83B_	Alignment	not modelled	91.0	22	**PDBTitle:** computer-based redesign of a beta sandwich protein suggests that2 extensive negative design is not required for de novo beta sheet3 design.
30	c3teuA_	Alignment	not modelled	91.0	23	**PDB header:**de novo protein **Chain:** A: **PDB Molecule:**fibcon; **PDBTitle:** crystal structure of fibcon
31	d1j8ka_	Alignment	not modelled	90.9	17	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
32	d1wfna1	Alignment	not modelled	90.5	20	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
33	d1tdqa1	Alignment	not modelled	90.4	18	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
34	d1fnha1	Alignment	not modelled	90.3	22	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
35	d2fnba_	Alignment	not modelled	89.8	18	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
36	d2a9da1	Alignment	not modelled	89.8	25	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**E set domains **Family:**Molybdenum-containing oxidoreductases-like dimerisation domain
37	d1owwa_	Alignment	not modelled	89.6	22	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
38	c2n1kA_	Alignment	not modelled	89.6	18	**PDB header:**structural protein **Chain:** A: **PDB Molecule:**fibronectin; **PDBTitle:** structure of the third type iii domain from human fibronectin
39	c3l5jB_	Alignment	not modelled	89.1	25	**PDB header:**immune system **Chain:** B: **PDB Molecule:**interleukin-6 receptor subunit beta; **PDBTitle:** crystal structure of fniii domains of human gp130 (domains 4-6)
40	d1fnfa1	Alignment	not modelled	89.0	17	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
41	c3rzwA_	Alignment	not modelled	88.5	18	**PDB header:**protein binding **Chain:** A: **PDB Molecule:**monobody ysmb-9; **PDBTitle:** crystal structure of the monobody ysmb-9 bound to human sumo1
42	c3pe9B_	Alignment	not modelled	88.2	14	**PDB header:**unknown function **Chain:** B: **PDB Molecule:**fibronectin(iii)-like module; **PDBTitle:** structures of clostridium thermocellum cbha fibronectin(iii)-like2 modules
43	d1qr4a1	Alignment	not modelled	88.2	28	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
44	d1k85a_	Alignment	not modelled	88.1	18	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
45	d2cuia1	Alignment	not modelled	87.9	22	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
46	c1qr4A_	Alignment	not modelled	87.9	21	**PDB header:**structural protein **Chain:** A: **PDB Molecule:**protein (tenascin); **PDBTitle:** two fibronectin type-iii domain segment from chicken tenascin
47	d1fnfa3	Alignment	not modelled	87.8	19	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
48	c3pe9D_	Alignment	not modelled	87.0	11	**PDB header:**unknown function **Chain:** D: **PDB Molecule:**fibronectin(iii)-like module; **PDBTitle:** structures of clostridium thermocellum cbha fibronectin(iii)-like2 modules
49	c5fm8B_	Alignment	not modelled	86.6	16	**PDB header:**structural protein **Chain:** B: **PDB Molecule:**myomesin-1; **PDBTitle:** structure of the c-terminally extended domain my4 of human myomesin2 (space group p65)
50	c2ee3A_	Alignment	not modelled	86.6	32	**PDB header:**signaling protein **Chain:** A: **PDB Molecule:**collagen alpha-1(xx) chain; **PDBTitle:** solution structures of the fn3 domain of human collagen2 alpha-1(xx) chain
51	c2ha1A_	Alignment	not modelled	86.6	16	**PDB header:**cell adhesion, structural protein **Chain:** A: **PDB Molecule:**fibronectin; **PDBTitle:** complex of the first and second type iii domains of human2 fibronectin in solution
52	c5fmvB_	Alignment	not modelled	86.5	23	**PDB header:**hydrolase **Chain:** B: **PDB Molecule:**receptor-type tyrosine-protein phosphatase c; **PDBTitle:** crystal structure of human cd45 extracellular region, domains d1-d4
53	c1fnhA_	Alignment	not modelled	86.2	26	**PDB header:**heparin and integrin binding **Chain:** A: **PDB Molecule:**protein (fibronectin); **PDBTitle:** crystal structure of heparin and integrin binding segment2 of human fibronectin
54	c5m0aA_	Alignment	not modelled	85.9	13	**PDB header:**cell adhesion **Chain:** A: **PDB Molecule:**fibronectin; **PDBTitle:** solution structure of isolated 15th fibronectin iii domain from human2 fibronectin
55	d1qr4a2	Alignment	not modelled	85.7	13	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III

						Family: Fibronectin type III
56	d2cuma1	Alignment	not modelled	85.2	17	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
57	c3l5hA_	Alignment	not modelled	84.9	24	**PDB header:**immune system **Chain:** A: **PDB Molecule:**interleukin-6 receptor subunit beta; **PDBTitle:** crystal structure of the full ectodomain of human gp130: new insights2 into the molecular assembly of receptor complexes
58	d1fnfa2	Alignment	not modelled	84.1	22	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
59	c3pe9C_	Alignment	not modelled	83.8	18	**PDB header:**unknown function **Chain:** C: **PDB Molecule:**fibronectin(iii)-like module; **PDBTitle:** structures of clostridium thermocellum cbha fibronectin(iii)-like2 modules
60	c3pe9A_	Alignment	not modelled	83.8	18	**PDB header:**unknown function **Chain:** A: **PDB Molecule:**fibronectin(iii)-like module; **PDBTitle:** structures of clostridium thermocellum cbha fibronectin(iii)-like2 modules
61	c5dftC_	Alignment	not modelled	83.6	25	**PDB header:**cell adhesion **Chain:** C: **PDB Molecule:**fibronectin; **PDBTitle:** structure of the eleventh type iii domain from human fibronectin
62	c4yg8A_	Alignment	not modelled	83.5	13	**PDB header:**transport protein **Chain:** A: **PDB Molecule:**chitin biosynthesis protein chs5; **PDBTitle:** crystal structure of the chs5-chs6 exomer cargo adaptor complex
63	c4gnsA_	Alignment	not modelled	83.5	13	**PDB header:**transport protein **Chain:** A: **PDB Molecule:**chitin biosynthesis protein chs5; **PDBTitle:** crystal structure of the chs5-chs6 exomer cargo adaptor complex
64	d1q38a_	Alignment	not modelled	83.4	20	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
65	d1fnha2	Alignment	not modelled	83.1	19	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
66	c2vkxE_	Alignment	not modelled	82.5	26	**PDB header:**cell adhesion **Chain:** E: **PDB Molecule:**neural cell adhesion molecule; **PDBTitle:** human ncam, fn3 domains 1 and 2, m610r mutant
67	c2geeA_	Alignment	not modelled	82.4	29	**PDB header:**protein binding, oncoprotein **Chain:** A: **PDB Molecule:**hypothetical protein; **PDBTitle:** crystal structure of human type iii fibronectin extradomain2 b and domain 8
68	c2kbgA_	Alignment	not modelled	82.3	23	**PDB header:**cell adhesion **Chain:** A: **PDB Molecule:**neural cell adhesion molecule 2; **PDBTitle:** solution structure of the second fibronectin type-iii module2 of ncam2
69	c4u3hA_	Alignment	not modelled	81.8	24	**PDB header:**de novo protein **Chain:** A: **PDB Molecule:**fn3con; **PDBTitle:** crystal structure of fn3con
70	c3asiA_	Alignment	not modelled	81.7	12	**PDB header:**cell adhesion **Chain:** A: **PDB Molecule:**neurexin-1-alpha; **PDBTitle:** alpha-neurexin-1 ectodomain fragment; lns5-egf3-lns6
71	d1fnha3	Alignment	not modelled	81.5	31	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
72	c2dleA_	Alignment	not modelled	81.0	24	**PDB header:**hydrolase **Chain:** A: **PDB Molecule:**receptor-type tyrosine-protein phosphatase eta; **PDBTitle:** solution structure of the fourth fn3 domain of human2 receptor-type tyrosine-protein phosphatase eta
73	d1x5la1	Alignment	not modelled	81.0	12	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
74	c5kf4A_	Alignment	not modelled	80.8	22	**PDB header:**contractile protein **Chain:** A: **PDB Molecule:**collagen alpha-1(xx) chain; **PDBTitle:** crystal structure of fn3 domain (residues p368-p466) of human collagen2 xx
75	c2h45A_	Alignment	not modelled	80.1	15	**PDB header:**cell adhesion, structural protein **Chain:** A: **PDB Molecule:**fibronectin; **PDBTitle:** solution structure of the second type iii domain of human2 fibronectin: ensemble of 25 structures
76	c2r0fA_	Alignment	not modelled	77.7	16	**PDB header:**sugar binding protein **Chain:** A: **PDB Molecule:**cgl3 lectin; **PDBTitle:** ligand free structure of fungal lectin cgl3
77	c2dm4A_	Alignment	not modelled	76.9	18	**PDB header:**lipid transport **Chain:** A: **PDB Molecule:**sortilin-related receptor; **PDBTitle:** solution structure of the second fn3 domain of human2 sorla/lr11
78	c2yrzA_	Alignment	not modelled	76.9	21	**PDB header:**cell adhesion **Chain:** A: **PDB Molecule:**integrin beta-4; **PDBTitle:** solution structure of the fibronectin type iii domain of2 human integrin beta-4
79	d1x5xa1	Alignment	not modelled	76.3	31	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
80	c2w1nA_	Alignment	not modelled	75.9	17	**PDB header:**hydrolase **Chain:** A: **PDB Molecule:**o-glcnacase nagj; **PDBTitle:** cohesin and fibronectin type-iii double module construct2 from the clostridium perfringens glycoside hydrolase gh84c
81	c3utoA_	Alignment	not modelled	74.9	18	**PDB header:**transferase **Chain:** A: **PDB Molecule:**twitchin; **PDBTitle:** twitchin kinase region from c.elegans (fn31-nl-kin-crd-ig26)

82	d2dn7a1	Alignment	not modelled	74.6	23	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
83	c4lsdF_	Alignment	not modelled	74.6	19	**PDB header:**hormone **Chain:** F: **PDB Molecule:**fibronectin type iii domain-containing protein 5; **PDBTitle:** myokine structure
84	c4agrB_	Alignment	not modelled	73.7	18	**PDB header:**sugar binding protein **Chain:** B: **PDB Molecule:**galectin; **PDBTitle:** structure of a tetrameric galectin from cinachyrella sp. (ball2 sponge)
85	c2ibgC_	Alignment	not modelled	73.6	15	**PDB header:**protein binding **Chain:** C: **PDB Molecule:**cg9211-pa; **PDBTitle:** crystal structure of hedgehog bound to the fniii domains of ihog
86	c1w0pA_	Alignment	not modelled	72.8	21	**PDB header:**hydrolase **Chain:** A: **PDB Molecule:**sialidase; **PDBTitle:** vibrio cholerae sialidase with alpha-2,6-sialyllactose
87	c4kdwA_	Alignment	not modelled	72.6	16	**PDB header:**cell adhesion **Chain:** A: **PDB Molecule:**antifreeze protein; **PDBTitle:** crystal structure of a bacterial immunoglobulin-like domain from the2 m. primoryensis ice-binding adhesin
88	d2bvya1	Alignment	not modelled	72.6	21	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**E set domains **Family:**E-set domains of sugar-utilizing enzymes
89	c2dmkA_	Alignment	not modelled	71.9	15	**PDB header:**protein binding **Chain:** A: **PDB Molecule:**midline 2 isoform 2; **PDBTitle:** the solution structure of the fn3 domain of human midline 22 protein
90	c3pdgA_	Alignment	not modelled	71.8	13	**PDB header:**unknown function **Chain:** A: **PDB Molecule:**fibronectin(iii)-like module; **PDBTitle:** structures of clostridium thermocellum cbha fibronectin(iii)-like2 modules
91	c2mfnA_	Alignment	not modelled	71.6	24	**PDB header:**cell adhesion protein **Chain:** A: **PDB Molecule:**fibronectin; **PDBTitle:** solution nmr structure of linked cell attachment modules of2 mouse fibronectin containing the rgd and synergy regions,3 10 structures
92	c2ekjA_	Alignment	not modelled	71.6	20	**PDB header:**signaling protein **Chain:** A: **PDB Molecule:**collagen alpha-1(xx) chain; **PDBTitle:** solution structures of the fn3 domain of human collagen2 alpha-1(xx) chain
93	d1lkta_	Alignment	not modelled	70.6	35	**Fold:**Head-binding domain of phage P22 tailspike protein **Superfamily:**Head-binding domain of phage P22 tailspike protein **Family:**Head-binding domain of phage P22 tailspike protein
94	c4qt6A_	Alignment	not modelled	69.4	13	**PDB header:**transport protein **Chain:** A: **PDB Molecule:**probable e3 ubiquitin-protein ligase herc1; **PDBTitle:** crystal structure of the spry domain of human herc1
95	c2a9dB_	Alignment	not modelled	69.4	26	**PDB header:**oxidoreductase **Chain:** B: **PDB Molecule:**sulfite oxidase; **PDBTitle:** crystal structure of recombinant chicken sulfite oxidase with arg at2 residue 161
96	d2haza1	Alignment	not modelled	69.1	28	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
97	c3fl7A_	Alignment	not modelled	68.5	15	**PDB header:**transferase, signaling protein **Chain:** A: **PDB Molecule:**ephrin receptor; **PDBTitle:** crystal structure of the human ephrin a2 ectodomain
98	d1x5ka1	Alignment	not modelled	68.3	28	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
99	c2dkmA_	Alignment	not modelled	68.3	11	**PDB header:**signaling protein **Chain:** A: **PDB Molecule:**collagen alpha-1(xx) chain; **PDBTitle:** solution structures of the fn3 domain of human collagen2 alpha-1(xx) chain
100	c4yfeA_	Alignment	not modelled	68.0	19	**PDB header:**hydrolase **Chain:** A: **PDB Molecule:**receptor-type tyrosine-protein phosphatase delta; **PDBTitle:** crystal structure of ptp delta fn1-fn2
101	d2djsa1	Alignment	not modelled	67.4	19	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
102	d1uena_	Alignment	not modelled	66.8	24	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
103	c2qbwA_	Alignment	not modelled	66.7	18	**PDB header:**unknown function **Chain:** A: **PDB Molecule:**pdz-fibronectin fusion protein; **PDBTitle:** the crystal structure of pdz-fibronectin fusion protein
104	c2vkyB_	Alignment	not modelled	66.3	35	**PDB header:**viral protein **Chain:** B: **PDB Molecule:**tail protein, piigcn4; **PDBTitle:** headbinding domain of phage p22 tailspike c-terminally fused to2 isoleucine zipper piigcn4 (chimera i)
105	d2gala_	Alignment	not modelled	65.9	21	**Fold:**Concanavalin A-like lectins/glucanases **Superfamily:**Concanavalin A-like lectins/glucanases **Family:**Galectin (animal S-lectin)
106	c4urtB_	Alignment	not modelled	65.7	21	**PDB header:**protein binding **Chain:** B: **PDB Molecule:**netrin receptor dcc; **PDBTitle:** the crystal structure of a fragment of netrin-1 in complex with fn5-2 fn6 of dcc
107	c2ed9A_	Alignment	not modelled	65.7	19	**PDB header:**apoptosis **Chain:** A: **PDB Molecule:**netrin receptor dcc; **PDBTitle:** solution structure of the third fibronectin type iii domain2 of human netrin receptor dcc
108	c1zlgA_	Alignment	not modelled	65.3	19	**PDB header:**hormone/growth factor **Chain:** A: **PDB Molecule:**anosmin 1;

108	c1zlgA_	Alignment	not modelled	65.5	19	**PDBTitle:** solution structure of the extracellular matrix protein anosmin-1
109	d3csba1	Alignment	not modelled	65.0	19	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
110	c5e7lA_	Alignment	not modelled	65.0	12	**PDB header:**cell adhesion **Chain:** A: **PDB Molecule:**contactin-2; **PDBTitle:** crystal structure of mouse cntn2 fn1-fn3 domains
111	d1wfua_	Alignment	not modelled	64.8	16	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
112	c2yroA_	Alignment	not modelled	64.6	9	**PDB header:**sugar binding protein **Chain:** A: **PDB Molecule:**galectin-8; **PDBTitle:** solution structure of the c-terminal gal-bind lectin2 protein from human galectin-8
113	d1tdqa2	Alignment	not modelled	64.5	21	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
114	c5fn8A_	Alignment	not modelled	64.4	16	**PDB header:**hydrolase **Chain:** A: **PDB Molecule:**receptor-type tyrosine-protein phosphatase c; **PDBTitle:** crystal structure of rat cd45 extracellular region, domains d3-d4
115	d1bqua1	Alignment	not modelled	64.2	17	**Fold:**Immunoglobulin-like beta-sandwich **Superfamily:**Fibronectin type III **Family:**Fibronectin type III
116	c5e53B_	Alignment	not modelled	64.0	16	**PDB header:**cell adhesion **Chain:** B: **PDB Molecule:**contactin-1; **PDBTitle:** crystal structure of chicken cntn1 fn1-fn3 domains
117	c2zgtB_	Alignment	not modelled	63.5	16	**PDB header:**hydrolase **Chain:** B: **PDB Molecule:**anti-tumor lectin; **PDBTitle:** crystal structure of agrocybe aegerita lectin aal mutant2 f93g
118	c4plnD_	Alignment	not modelled	63.1	15	**PDB header:**protein binding **Chain:** D: **PDB Molecule:**neogenin; **PDBTitle:** crystal structure of chicken netrin-1 (ln-le3) complexed with mouse2 neogenin (fn4-5)
119	c3qhtC_	Alignment	not modelled	62.8	16	**PDB header:**de novo protein **Chain:** C: **PDB Molecule:**monobody ysmb-1; **PDBTitle:** crystal structure of the monobody ysmb-1 bound to yeast sumo
120	c5e4sA_	Alignment	not modelled	62.6	23	**PDB header:**cell adhesion **Chain:** A: **PDB Molecule:**contactin-4; **PDBTitle:** crystal structure of mouse cntn4 fn1-fn3 domains