




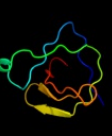



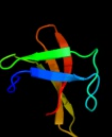

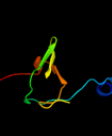





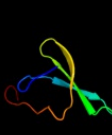



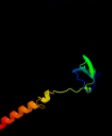









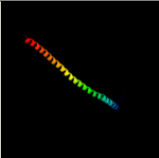
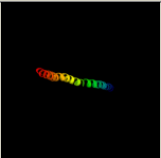



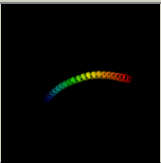
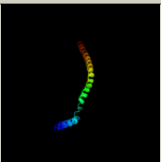
#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2kt8A_	 Alignment		99.0	18	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: probable surface protein; PDBTitle: solution nmr structure of the cpe1231(468-535) protein from2 clostridium perfringens, northeast structural genomics3 consortium target cpr82b
2	c2krsA_	 Alignment		98.9	18	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: probable enterotoxin; PDBTitle: solution nmr structure of sh3 domain from cpf_0587 (fragment2 415-479) from clostridium perfringens. northeast structural3 genomics consortium (nesg) target cpr74a.
3	c2kybA_	 Alignment		98.6	15	PDB header: toxin Chain: A: PDB Molecule: mannosyl-glycoprotein endo-beta-n-acetylglucosamidase PDBTitle: solution structure of cpr82g from clostridium perfringens. north east2 structural genomics consortium target cpr82g
4	c2kq8A_	 Alignment		98.6	28	PDB header: hydrolase Chain: A: PDB Molecule: cell wall hydrolase; PDBTitle: solution nmr structure of a domain from bt9727_4915 from2 bacillus thuringiensis, northeast structural genomics3 consortium target bur95a
5	c3npfB_	 Alignment		98.4	16	PDB header: hydrolase Chain: B: PDB Molecule: putative dipeptidyl-peptidase vi; PDBTitle: crystal structure of a putative dipeptidyl-peptidase vi (bacova_00612)2 from bacteroides ovatus at 1.72 a resolution
6	c3h41A_	 Alignment		97.4	24	PDB header: hydrolase Chain: A: PDB Molecule: nlp/p60 family protein; PDBTitle: crystal structure of a nlp/p60 family protein (bce_2878) from2 bacillus cereus atcc 10987 at 1.79 a resolution
7	c1xovA_	 Alignment		96.5	10	PDB header: hydrolase Chain: A: PDB Molecule: ply protein; PDBTitle: the crystal structure of the listeria monocytogenes bacteriophage psa2 endolysin plypsa
8	c3m1uB_	 Alignment		96.5	18	PDB header: hydrolase Chain: B: PDB Molecule: putative gamma-d-glutamyl-l-diamino acid endopeptidase; PDBTitle: crystal structure of a putative gamma-d-glutamyl-l-diamino acid2 endopeptidase (dnu_0896) from desulfovibrio vulgaris hildenborough at3 1.75 a resolution
9	d1wfwA_	 Alignment		96.3	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
10	d1cska_	 Alignment		96.0	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
11	c3pe0B_	 Alignment		95.9	26	PDB header: structural protein Chain: B: PDB Molecule: plectin; PDBTitle: structure of the central region of the plakin domain of plectin

12	c2dx1A_	Alignment		95.8	9	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
13	c3qwyA_	Alignment		95.7	18	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
14	c1k9aB_	Alignment		95.7	20	PDB header: transferase Chain: B: PDB Molecule: carboxyl-terminal src kinase; PDBTitle: crystal structure analysis of full-length carboxyl-terminal2 src kinase at 2.5 a resolution
15	d1gl5a_	Alignment		95.7	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
16	c3qwxX_	Alignment		95.6	18	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
17	c2x3wD_	Alignment		95.6	21	PDB header: endocytosis Chain: D: PDB Molecule: protein kinase c and casein kinase substrate in neurons PDBTitle: structure of mouse syndapin i (crystal form 2)
18	d1k9aa1	Alignment		95.6	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
19	d1ckaa_	Alignment		95.5	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
20	c2eyyA_	Alignment		95.5	17	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
21	c2ct4A_	Alignment	not modelled	95.4	20	PDB header: signaling protein Chain: A: PDB Molecule: cdc42-interacting protein 4; PDBTitle: solution struture of the sh3 domain of the cdc42-2 interacting protein 4
22	d1ng2a2	Alignment	not modelled	95.4	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
23	d1k1za_	Alignment	not modelled	95.3	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
24	c1s1nA_	Alignment	not modelled	95.2	21	PDB header: cell adhesion Chain: A: PDB Molecule: nephrocystin 1; PDBTitle: sh3 domain of human nephrocystin
25	c1griA_	Alignment	not modelled	95.1	14	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
26	d1qlya_	Alignment	not modelled	95.1	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
27	c2kbtA_	Alignment	not modelled	95.1	17	PDB header: signaling protein Chain: A: PDB Molecule: chimera of proto-oncogene vav, linker, tag2 (inset) using a sortase-mediated protein ligation method PDB header: transferase Chain: D: PDB Molecule: mitogen-activated protein kinase kinase kinase 10; PDBTitle: crystal structure of human mixed lineage kinase map3k10 sh3 domain
28	c2rf0D_	Alignment	not modelled	95.1	23	PDB header: structural genomics, unknown function

29	c2dl5A_	Alignment	not modelled	95.1	14	Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the first sh3 domain of human2 kiaa0769 protein
30	c1zlmA_	Alignment	not modelled	95.1	24	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
31	dl0pka1	Alignment	not modelled	95.1	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
32	dl0eba_	Alignment	not modelled	95.0	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
33	c2egcA_	Alignment	not modelled	95.0	15	PDB header: signaling protein Chain: A: PDB Molecule: sh3 and px domain-containing protein 2a; PDBTitle: solution structure of the fifth sh3 domain from human2 kiaa0418 protein
34	c1z9zA_	Alignment	not modelled	95.0	18	PDB header: structural protein Chain: A: PDB Molecule: cytoskeleton assembly control protein sla1; PDBTitle: crystal structure of yeast sla1 sh3 domain 3
35	c1ov3A_	Alignment	not modelled	95.0	16	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
36	dlnega_	Alignment	not modelled	95.0	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
37	c1negA_	Alignment	not modelled	95.0	22	PDB header: structural protein Chain: A: PDB Molecule: spectrin alpha chain, brain; PDBTitle: crystal structure analysis of n-and c-terminal labeled sh3-2 domain of alpha-chicken spectrin
38	dluj0a_	Alignment	not modelled	94.9	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
39	dlpwta_	Alignment	not modelled	94.9	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
40	dlwlpb2	Alignment	not modelled	94.9	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
41	c2xmFA_	Alignment	not modelled	94.9	14	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
42	c1ei3E_	Alignment		94.8	14	PDB header: PDB COMPND:
43	c1m9sA_	Alignment	not modelled	94.8	13	PDB header: signaling protein Chain: A: PDB Molecule: internalin b; PDBTitle: crystal structure of internalin b (inlb), a listeria2 monocytogenes virulence protein containing sh3-like3 domains.
44	dlutia_	Alignment	not modelled	94.8	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
45	c2azsA_	Alignment	not modelled	94.8	12	PDB header: signaling protein Chain: A: PDB Molecule: sh2-sh3 adapter protein drk; PDBTitle: nmr structure of the n-terminal sh3 domain of drk2 (calculated without noe restraints)
46	c2fg0B_	Alignment	not modelled	94.8	13	PDB header: hydrolase Chain: B: PDB Molecule: cog0791: cell wall-associated hydrolases (invasion- PDBTitle: crystal structure of a putative gamma-d-glutamyl-l-diamino acid2 endopeptidase (npun_r0659) from nostoc punctiforme pcc 73102 at 1.793 a resolution
47	c1x6gA_	Alignment	not modelled	94.8	25	PDB header: signaling protein Chain: A: PDB Molecule: megakaryocyte-associated tyrosine-protein kinase; PDBTitle: solution structures of the sh3 domain of human2 megakaryocyte-associated tyrosine-protein kinase.
48	c2df6A_	Alignment	not modelled	94.8	16	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 7; PDBTitle: crystal structure of the sh3 domain of betapix in complex2 with a high affinity peptide from pak2
49	c2j6kE_	Alignment	not modelled	94.8	16	PDB header: protein binding Chain: E: PDB Molecule: cd2-associated protein; PDBTitle: n-terminal sh3 domain of cms (cd2ap human homolog)
50	dlfmka1	Alignment	not modelled	94.8	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
51	dlu06a1	Alignment	not modelled	94.8	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
52	c2ekhA_	Alignment	not modelled	94.8	23	PDB header: signaling protein Chain: A: PDB Molecule: sh3 and px domain-containing protein 2a; PDBTitle: solution structures of the sh3 domain of human kiaa0418
53	c2epdA_	Alignment	not modelled	94.8	18	PDB header: protein binding Chain: A: PDB Molecule: rho gtpase-activating protein 4; PDBTitle: solution structure of sh3 domain in rho-gtpase-

						activating2 protein 4
54	d1ov3a2	Alignment	not modelled	94.8	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
55	c1zx6A	Alignment	not modelled	94.8	22	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
56	c2fo0A	Alignment	not modelled	94.7	18	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase abl1 (1b isoform); PDBTitle: organization of the sh3-sh2 unit in active and inactive forms of the2 c-abl tyrosine kinase
57	c1x2pA	Alignment	not modelled	94.7	17	PDB header: transferase Chain: A: PDB Molecule: protein arginine n-methyltransferase 2; PDBTitle: solution structure of the sh3 domain of the protein2 arginine n-methyltransferase 2
58	c2cucA	Alignment	not modelled	94.7	18	PDB header: signaling protein Chain: A: PDB Molecule: sh3 domain containing ring finger 2; PDBTitle: solution structure of the sh3 domain of the mouse2 hypothetical protein sh3rf2
59	c2enmA	Alignment	not modelled	94.7	21	PDB header: endocytosis Chain: A: PDB Molecule: sorting nexin-9; PDBTitle: solution structure of the sh3 domain from mouse sorting2 nexin-9
60	c3cqtA	Alignment	not modelled	94.7	21	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase fyn; PDBTitle: n53i v55l mutant of fyn sh3 domain
61	c2dmoA	Alignment	not modelled	94.7	28	PDB header: signaling protein Chain: A: PDB Molecule: neutrophil cytosol factor 2; PDBTitle: refined solution structure of the 1st sh3 domain from human2 neutrophil cytosol factor 2 (ncf-2)
62	d1oota	Alignment	not modelled	94.6	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
63	d1gcqc	Alignment	not modelled	94.6	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
64	c1w70A	Alignment	not modelled	94.6	25	PDB header: sh3 domain Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: sh3 domain of p40phox complexed with c-terminal polyproline2 region of p47phox
65	d1uuea	Alignment	not modelled	94.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	c1degO	Alignment		94.6	13	PDB header: PDB COMPND:
67	d1k4us	Alignment	not modelled	94.6	15	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
68	c3ojaB	Alignment		94.6	11	PDB header: protein binding Chain: B: PDB Molecule: anopheles plasmodium-responsive leucine-rich repeat protein PDBTitle: crystal structure of lrim1/apl1c complex
69	d1gria2	Alignment	not modelled	94.6	14	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
70	c2ed0A	Alignment	not modelled	94.5	10	PDB header: signaling protein Chain: A: PDB Molecule: abl interactor 2; PDBTitle: solution structure of the sh3 domain of abl interactor 22 (abelson interactor 2)
71	d2rn8a1	Alignment	not modelled	94.5	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
72	d2fo0a1	Alignment	not modelled	94.5	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
73	d1e6ga	Alignment	not modelled	94.5	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	d1u5sa1	Alignment	not modelled	94.5	14	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
75	c1oplA	Alignment	not modelled	94.5	18	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase; PDBTitle: structural basis for the auto-inhibition of c-abl tyrosine2 kinase
76	c2jxbA	Alignment	not modelled	94.5	21	PDB header: signaling protein complex Chain: A: PDB Molecule: t-cell surface glycoprotein cd3 epsilon chain, PDBTitle: structure of cd3epsilon-nck2 first sh3 domain complex
77	c2csqA	Alignment	not modelled	94.5	11	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: rim binding protein 2; PDBTitle: solution structure of the second sh3 domain of human rim-2 binding protein 2

78	c2d1xD_	Alignment	not modelled	94.5	11	PDB header: cell invasion Chain: D: PDB Molecule: cortactin isoform a; PDBTitle: the crystal structure of the cortactin-sh3 domain and amap1-2 peptide complex
79	d1tuca_	Alignment	not modelled	94.5	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
80	d1sema_	Alignment	not modelled	94.5	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
81	c1ng2A_	Alignment	not modelled	94.4	15	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
82	c1zuyB_	Alignment	not modelled	94.4	8	PDB header: contractile protein Chain: B: PDB Molecule: myosin-5 isoform; PDBTitle: high-resolution structure of yeast myo5 sh3 domain
83	c2nwmA_	Alignment	not modelled	94.4	20	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the first sh3 domain of human vinexin2 and its interaction with the peptides from vinculin
84	d1gcqa_	Alignment	not modelled	94.3	14	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
85	d1e6ha_	Alignment	not modelled	94.3	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
86	c2ablA_	Alignment	not modelled	94.3	18	PDB header: transferase Chain: A: PDB Molecule: abl tyrosine kinase; PDBTitle: sh3-sh2 domain fragment of human bcr-abl tyrosine kinase
87	c1wxtA_	Alignment	not modelled	94.3	32	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hypothetical protein flj21522; PDBTitle: solution structure of the sh3 domain of human hypothetical2 protein flj21522
88	c2l0aA_	Alignment	not modelled	94.3	21	PDB header: signaling protein Chain: A: PDB Molecule: signal transducing adapter molecule 1; PDBTitle: solution nmr structure of signal transducing adapter molecule 1 stam-12 from homo sapiens, northeast structural genomics consortium target3 hr4479e
89	c2pqhA_	Alignment	not modelled	94.3	23	PDB header: structural protein Chain: A: PDB Molecule: spectrin alpha chain, brain; PDBTitle: structure of sh3 chimera with a type ii ligand linked to the chain c-2 terminal
90	c2js0A_	Alignment	not modelled	94.3	15	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
91	d1shfa_	Alignment	not modelled	94.3	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
92	c2bz8B_	Alignment	not modelled	94.2	19	PDB header: sh3 domain Chain: B: PDB Molecule: sh3-domain kinase binding protein 1; PDBTitle: n-terminal sh3 domain of cin85 bound to cbl-b peptide
93	d1ycsb2	Alignment	not modelled	94.2	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	c2k2mA_	Alignment	not modelled	94.2	23	PDB header: signaling protein Chain: A: PDB Molecule: eps8-like protein 1; PDBTitle: structural basis of pxxdy motif recognition in sh3 binding
95	d1jo8a_	Alignment	not modelled	94.2	11	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
96	c1y57A_	Alignment	not modelled	94.2	18	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase src; PDBTitle: structure of unphosphorylated c-src in complex with an inhibitor
97	c2h8hA_	Alignment	not modelled	94.2	18	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase src; PDBTitle: src kinase in complex with a quinazoline inhibitor
98	c2ct3A_	Alignment	not modelled	94.2	11	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
99	c2egaA_	Alignment	not modelled	94.1	24	PDB header: signaling protein Chain: A: PDB Molecule: sh3 and px domain-containing protein 2a; PDBTitle: solution structure of the first sh3 domain from human2 kiaa0418 protein
100	d1h8ka_	Alignment	not modelled	94.1	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	c2c0iA_	Alignment	not modelled	94.1	14	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck; PDBTitle: src family kinase hck with bound inhibitor a-420983
102	c2yuoA_	Alignment	not modelled	94.0	13	PDB header: signaling protein Chain: A: PDB Molecule: run and tbc1 domain containing 3; PDBTitle: solution structure of the sh3 domain of mouse run and tbc12 domain containing 3
103	c2yunA_	Alignment	not modelled	94.0	15	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin

104	c2drmB_	Alignment	not modelled	94.0	19	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
105	c1wxbA_	Alignment	not modelled	94.0	19	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: epidermal growth factor receptor pathway PDBTitle: solution structure of the sh3 domain from human epidermal2 growth factor receptor pathway substrate 8-like protein
106	c1u3oA_	Alignment	not modelled	94.0	21	PDB header: signaling protein Chain: A: PDB Molecule: huntingtin-associated protein-interacting PDBTitle: solution structure of rat kalirin n-terminal sh3 domain
107	c2dnuA_	Alignment	not modelled	94.0	14	PDB header: structural genomics, structural protein Chain: A: PDB Molecule: sh3 multiple domains 1; PDBTitle: solution structure of rsgi ruh-061, a sh3 domain from human
108	c2jw4A_	Alignment	not modelled	93.9	23	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: nmr solution structure of the n-terminal sh3 domain of2 human nckalpha
109	d2v1ra1	Alignment	not modelled	93.9	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
110	c2dila_	Alignment	not modelled	93.9	16	PDB header: cell adhesion Chain: A: PDB Molecule: proline-serine-threonine phosphatase-interacting PDBTitle: solution structure of the sh3 domain of the human proline-2 serine-threonine phosphatase-interacting protein 1
111	d1lcka1	Alignment	not modelled	93.9	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	c2a28D_	Alignment	not modelled	93.7	14	PDB header: signaling protein Chain: D: PDB Molecule: bzz1 protein; PDBTitle: atomic-resolution crystal structure of the second sh32 domain of yeast bzz1 determined from a pseudomerohedrally3 twinned crystal
113	c1x27F_	Alignment	not modelled	93.6	18	PDB header: signaling protein Chain: F: PDB Molecule: proto-oncogene tyrosine-protein kinase lck; PDBTitle: crystal structure of lck sh2-sh3 with sh2 binding site of2 p130cas
114	c1x2qA_	Alignment	not modelled	93.6	14	PDB header: signaling protein Chain: A: PDB Molecule: signal transducing adapter molecule 2; PDBTitle: solution structure of the sh3 domain of the signal2 transducing adaptor molecule 2
115	d1gbra_	Alignment	not modelled	93.6	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	c2wpqA_	Alignment		93.6	8	PDB header: membrane protein Chain: A: PDB Molecule: trimeric autotransporter adhesin fragment; PDBTitle: salmonella enterica sada 479-519 fused to gcn4 adaptors (2 sadak3, in-register fusion)
117	c2djqa_	Alignment	not modelled	93.6	19	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: sh3 domain containing ring finger 2; PDBTitle: the solution structure of the first sh3 domain of mouse sh32 domain containing ring finger 2
118	d1ov3a1	Alignment	not modelled	93.6	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	c1degF_	Alignment		93.6	6	PDB header: PDB COMPND:
120	c2eqia_	Alignment	not modelled	93.6	8	PDB header: immune system, hydrolase Chain: A: PDB Molecule: phospholipase c, gamma 2; PDBTitle: solution structure of the sh3 domain from phospholipase c,2 gamma 2