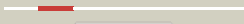
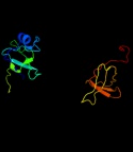
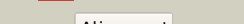
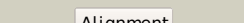
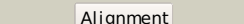
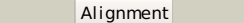
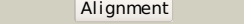
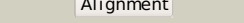
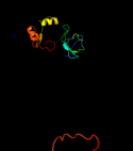
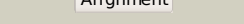
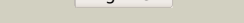
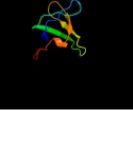
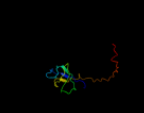
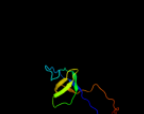
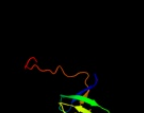


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1griA_	 Alignment		100.0	25	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
2	c1ng2A_	 Alignment		100.0	15	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
3	c2lqnA_	 Alignment		100.0	20	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
4	c2l3sA_	 Alignment		100.0	19	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
5	c1ov3A_	 Alignment		100.0	19	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
6	c2eyzA_	 Alignment		100.0	15	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
7	d1udla_	 Alignment		99.8	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
8	c1mv3A_	 Alignment		99.7	26	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: myc box dependent interacting protein 1; PDBTitle: nmr structure of the tumor suppressor bin1: alternative2 splicing in melanoma and interaction with c-myc
9	c2dybA_	 Alignment		99.7	17	PDB header: oxidoreductase Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: the crystal structure of human p40(phox)
10	d1ng2a2	 Alignment		99.7	11	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
11	c2dx1A_	 Alignment		99.6	21	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef

12	c2jmcA_	Alignment		99.6	28	PDB header: signaling protein Chain: A: PDB Molecule: spectrin alpha chain, brain and p41 peptide PDBTitle: chimera between spc-sh3 and p41
13	c3nmzD_	Alignment		99.6	17	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of apc complexed with asef
14	c1x2qA_	Alignment		99.6	23	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
15	d1ug1a_	Alignment		99.6	10	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
16	c1x43A_	Alignment		99.6	27	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: sh3 domain grb2-like protein b1; PDBTitle: solution structure of the sh3 domain of endophilin b12 (sh3g1b1)
17	d1uffa_	Alignment		99.6	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
18	c2dlpA_	Alignment		99.6	18	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
19	c2yunA_	Alignment		99.6	19	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
20	d1wfwA_	Alignment		99.6	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
21	c2cubA_	Alignment	not modelled	99.6	17	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
22	c2nwmA_	Alignment	not modelled	99.6	22	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
23	c2ysqA_	Alignment	not modelled	99.6	15	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 9; PDBTitle: solution structure of the sh3 domain from rho guanine2 nucleotide exchange factor 9
24	c2d8jA_	Alignment	not modelled	99.6	19	PDB header: transferase Chain: A: PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh3 domain of fyn-related kinase
25	c2ekhA_	Alignment	not modelled	99.6	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
26	d1wiea_	Alignment	not modelled	99.6	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
27	c2yuoA_	Alignment	not modelled	99.6	23	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
28	c2eyyA_	Alignment	not modelled	99.6	21	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i

29	c2ed1A_	Alignment	not modelled	99.6	19	PDB header: signaling protein Chain: A: PDB Molecule: 130 kda phosphatidylinositol 4,5-biphosphate- PDBTitle: solution structure of the sh3 domain of 130 kda2 phosphatidylinositol 4,5-biphosphate-dependent arf1 gtpase-3 activating protein
30	d1ov3a2	Alignment	not modelled	99.6	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
31	c2yt6A_	Alignment	not modelled	99.6	28	PDB header: cell cycle Chain: A: PDB Molecule: adult male urinary bladder cdna, riken full- PDBTitle: solution structure of the sh3_1 domain of yamaguchi sarcoma2 viral (v-yes) oncogene homolog 1
32	c2egcA_	Alignment	not modelled	99.6	23	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
33	c2egeA_	Alignment	not modelled	99.6	14	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized protein kiaa1666; PDBTitle: solution structure of the third sh3 domain from human2 kiaa1666 protein
34	c2rqrA_	Alignment	not modelled	99.6	15	PDB header: protein binding Chain: A: PDB Molecule: engulfment and cell motility protein 1, linker, dedicator PDBTitle: the solution structure of human dock2 sh3 domain - elmo1 peptide2 chimera complex
35	c2dmoA_	Alignment	not modelled	99.6	20	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)
36	c2csqA_	Alignment	not modelled	99.6	29	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
37	c2df6A_	Alignment	not modelled	99.6	22	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
38	c2yupA_	Alignment	not modelled	99.6	15	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
39	c2ed0A_	Alignment	not modelled	99.6	34	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)
40	c2epdA_	Alignment	not modelled	99.6	23	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
41	d1ue9a_	Alignment	not modelled	99.6	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
42	c2dila_	Alignment	not modelled	99.6	29	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
43	c3qwyA_	Alignment	not modelled	99.5	19	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
44	c2dl4A_	Alignment	not modelled	99.5	18	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
45	c2kymA_	Alignment	not modelled	99.5	21	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of the bem1p sh3-ci domain from l.elongisporus in2 complex with ste20p peptide
46	c2e5kA_	Alignment	not modelled	99.5	24	PDB header: protein binding Chain: A: PDB Molecule: suppressor of t-cell receptor signaling 1; PDBTitle: solution structure of sh3 domain in suppressor of t-cell2 receptor signaling 1
47	c2dvjA_	Alignment	not modelled	99.5	17	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii
48	c2l0aA_	Alignment	not modelled	99.5	24	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
49	c2drmB_	Alignment	not modelled	99.5	33	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
50	c2dl3A_	Alignment	not modelled	99.5	23	PDB header: cell adhesion, signaling protein Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: solution structure of the first sh3 domain of human sorbin2 and sh3 domain-containing protein 1
51	c2xmfA_	Alignment	not modelled	99.5	36	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
52	c1wx6A_	Alignment	not modelled	99.5	20	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: cytoplasmic protein nck2; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck2

53	c1zlmA	Alignment	not modelled	99.5	25	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
54	c2djgA	Alignment	not modelled	99.5	20	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
55	c2dnuA	Alignment	not modelled	99.5	24	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
56	c2dbmA	Alignment	not modelled	99.5	22	PDB header: transferase, signaling protein Chain: A: PDB Molecule: sh3-containing grb2-like protein 2; PDBTitle: solution structures of the sh3 domain of human sh3-2 containing grb2-like protein 2
57	c2kbtA	Alignment	not modelled	99.5	22	PDB header: signaling protein Chain: A: PDB Molecule: chimera of proto-oncogene vav, linker, PDBTitle: attachment of an nmr-invisible solubility enhancement tag2 (inset) using a sortase-mediated protein ligation method
58	c1wi7A	Alignment	not modelled	99.5	17	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase binding protein 1; PDBTitle: solution structure of the sh3 domain of sh3-domain kinase2 binding protein 1
59	d1utia	Alignment	not modelled	99.5	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
60	c2fpeB	Alignment	not modelled	99.5	16	PDB header: signaling protein Chain: B: PDB Molecule: c-jun-amino-terminal kinase interacting protein PDBTitle: conserved dimerization of the ib1 src-homology 3 domain
61	d1k4us	Alignment	not modelled	99.5	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
62	d1oeba	Alignment	not modelled	99.5	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
63	d1gria2	Alignment	not modelled	99.5	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
64	c2ebpA	Alignment	not modelled	99.5	13	PDB header: cell adhesion Chain: A: PDB Molecule: sam and sh3 domain-containing protein 1; PDBTitle: solution structure of the sh3 domain from human sam and sh32 domain containing protein 1
65	d1wlpb2	Alignment	not modelled	99.5	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	d1ujya	Alignment	not modelled	99.5	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
67	d1gcqa	Alignment	not modelled	99.5	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
68	d1uj0a	Alignment	not modelled	99.5	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
69	c2eqiA	Alignment	not modelled	99.5	24	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
70	c2d1xD	Alignment	not modelled	99.5	36	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
71	c2bz8B	Alignment	not modelled	99.5	23	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
72	d1kjaw1	Alignment	not modelled	99.5	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
73	d1sema	Alignment	not modelled	99.5	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	c1x6gA	Alignment	not modelled	99.5	16	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.
75	d1phta	Alignment	not modelled	99.5	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
76	c4esrB	Alignment	not modelled	99.5	22	PDB header: protein binding Chain: B: PDB Molecule: jouberin; PDBTitle: molecular and structural characterization of the sh3 domain of ahi-12 in regulation of cellular resistance of bcr-abl + chronic myeloid3 leukemia cells to tyrosine kinase inhibitors
77	c1zukA	Alignment	not modelled	99.5	24	PDB header: contractile protein Chain: A: PDB Molecule: myosin tail region-interacting protein mti 1; PDBTitle: yeast bbc1 sh3 domain complexed with a peptide from las17
						PDB header: signaling protein complex

78	c2jxbA_	Alignment	not modelled	99.5	28	Chain: A: PDB Molecule: t-cell surface glycoprotein cd3 epsilon chain; PDBTitle: structure of cd3epsilon-nck2 first sh3 domain complex
79	dlugva_	Alignment	not modelled	99.5	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
80	c1zx6A_	Alignment	not modelled	99.5	21	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
81	c2cudA_	Alignment	not modelled	99.5	18	PDB header: signaling protein Chain: A: PDB Molecule: src-like-adaptor; PDBTitle: solution structure of the sh3 domain of the human src-like2 adpater protein (slap)
82	c3qwxX_	Alignment	not modelled	99.5	21	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
83	dlfmka1	Alignment	not modelled	99.5	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	c1x69A_	Alignment	not modelled	99.5	32	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
85	dluhca_	Alignment	not modelled	99.5	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
86	dlfna_	Alignment	not modelled	99.5	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
87	c2ct3A_	Alignment	not modelled	99.5	26	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
88	c2pqhA_	Alignment	not modelled	99.5	32	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
89	c2eywA_	Alignment	not modelled	99.5	23	PDB header: signaling protein Chain: A: PDB Molecule: v-crK sarcoma virus ct10 oncogene homolog PDBTitle: n-terminal sh3 domain of ct10-regulated kinase
90	c1wyxA_	Alignment	not modelled	99.5	25	PDB header: cell adhesion Chain: A: PDB Molecule: crk-associated substrate; PDBTitle: the crystal structure of the p130cas sh3 domain at 1.1 a2 resolution
91	c2js0A_	Alignment	not modelled	99.5	20	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
92	dlloota_	Alignment	not modelled	99.5	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
93	c1w70A_	Alignment	not modelled	99.5	24	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
94	c2egaA_	Alignment	not modelled	99.5	22	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
95	c2dl8A_	Alignment	not modelled	99.5	27	PDB header: signaling protein Chain: A: PDB Molecule: slit-robo rho gtpase-activating protein 2; PDBTitle: solution structure of the sh3 domain of human slit-robo rho2 gtpase-activating protein 2
96	c1x6bA_	Alignment	not modelled	99.5	13	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine exchange factor (gef) 16; PDBTitle: solution structures of the sh3 domain of human rho guanine2 exchange factor (gef) 16
97	dlnega_	Alignment	not modelled	99.5	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
98	c1negA_	Alignment	not modelled	99.5	29	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
99	dluuea_	Alignment	not modelled	99.5	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	dlshfa_	Alignment	not modelled	99.5	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	c2o2oA_	Alignment	not modelled	99.5	17	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
102	dlpwta_	Alignment	not modelled	99.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	dlspka_	Alignment	not modelled	99.5	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
104	c2oi3A_	Alignment	not modelled	99.5	24	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck; PDBTitle: nmr structure analysis of the hematopoetic cell kinase

						sh32 domain complexed with an artificial high affinity ligand3 (pd1)
105	c2j06B_	Alignment	not modelled	99.5	25	PDB header: signal transduction Chain: B: PDB Molecule: ras gtpase-activating protein 1; PDBTitle: crystal structure of the rasgap sh3 domain at 1.8 angstrom2 resolution
106	c3rnjA_	Alignment	not modelled	99.5	16	PDB header: protein binding Chain: A: PDB Molecule: brain-specific angiogenesis inhibitor 1-associated protein PDBTitle: crystal structure of the sh3 domain from irsp53 (baiap2)
107	c2dm1A_	Alignment	not modelled	99.5	24	PDB header: signaling protein Chain: A: PDB Molecule: protein vav-2; PDBTitle: solution structure of the second sh3 domain of human2 protein vav-2
108	d1g2ba_	Alignment	not modelled	99.5	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
109	c2krnA_	Alignment	not modelled	99.5	27	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
110	c1x2pA_	Alignment	not modelled	99.5	21	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
111	d1ng2a1	Alignment	not modelled	99.5	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	c2d8hA_	Alignment	not modelled	99.5	21	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
113	c2csiA_	Alignment	not modelled	99.5	29	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: rim binding protein 2; PDBTitle: solution structure of the third sh3 domain of human rim-2 binding protein 2
114	d1j3ta_	Alignment	not modelled	99.5	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	d1u06a1	Alignment	not modelled	99.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	c2i0nA_	Alignment	not modelled	99.5	17	PDB header: structural protein Chain: A: PDB Molecule: class vii unconventional myosin; PDBTitle: structure of dictyostelium discoideum myosin vii sh3 domain2 with adjacent proline rich region
117	d1bb9a_	Alignment	not modelled	99.5	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
118	c1bb9A_	Alignment	not modelled	99.5	25	PDB header: transferase Chain: A: PDB Molecule: amphiphysin 2; PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
119	d1uhfa_	Alignment	not modelled	99.5	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	c2lcsA_	Alignment	not modelled	99.5	18	PDB header: transferase, signaling protein Chain: A: PDB Molecule: nap1-binding protein 2; PDBTitle: yeast nbp2p sh3 domain in complex with a peptide from ste20p