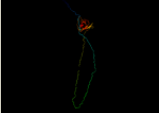
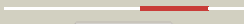
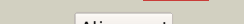
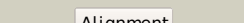
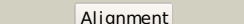
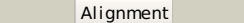
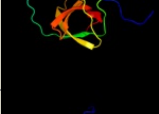
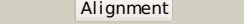
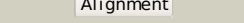
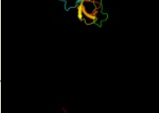
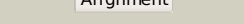
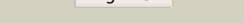
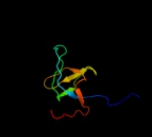
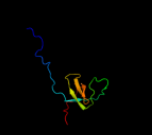
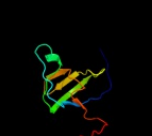


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1mv3A_	 Alignment		99.9	10	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: myc box dependent interacting protein 1; PDBTitle: nmr structure of the tumor suppressor bin1: alternative2 splicing in melanoma and interaction with c-myc
2	c2rqrA_	 Alignment		99.8	10	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
3	c1griA_	 Alignment		99.8	26	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
4	c2lqnA_	 Alignment		99.7	14	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
5	c2ebpA_	 Alignment		99.7	15	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
6	c2vknA_	 Alignment		99.7	100	PDB header: membrane protein Chain: A: PDB Molecule: protein ssu81; PDBTitle: yeast sho1 sh3 domain complexed with a peptide from pbs2
7	c2jxbA_	 Alignment		99.7	25	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
8	c2dbkA_	 Alignment		99.7	21	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
9	c2oi3A_	 Alignment		99.7	29	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck; PDBTitle: nmr structure analysis of the hematopoietic cell kinase sh32 domain complexed with an artificial high affinity ligand3 (pd1)
10	c2dx1A_	 Alignment		99.7	27	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
11	c2ed0A_	 Alignment		99.7	32	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)

12	d1ng2a2	Alignment		99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
13	c2l3sA	Alignment		99.7	16	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
14	c2eyxA	Alignment		99.7	24	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
15	c1x6bA	Alignment		99.7	22	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
16	c2i0nA	Alignment		99.7	22	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
17	c2eyyA	Alignment		99.7	23	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
18	c2ysqA	Alignment		99.7	23	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
19	c2cudA	Alignment		99.7	26	PDB header: signaling protein Chain: A: PDB Molecule: src-like-adapter; PDBTitle: solution structure of the sh3 domain of the human src-like2 adppter protein (slap)
20	d2hsaA	Alignment		99.7	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
21	c3nmzD	Alignment	not modelled	99.7	23	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with ascf
22	c2eyzA	Alignment	not modelled	99.7	15	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
23	c2yupA	Alignment	not modelled	99.7	23	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
24	d1spka	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
25	c2kymA	Alignment	not modelled	99.7	30	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
26	c1wxtA	Alignment	not modelled	99.7	35	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
27	c3qwyA	Alignment	not modelled	99.7	32	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
28	c2yt6A	Alignment	not modelled	99.7	33	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

29	d1jqqa_	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
30	c2ed1A_	Alignment	not modelled	99.7	25	PDB header: signaling protein Chain: A: PDB Molecule: 130 kda phosphatidylinositol 4,5-bisphosphate- PDBTitle: solution structure of the sh3 domain of 130 kda2 phosphatidylinositol 4,5-bisphosphate-dependent arf1 gtpase-3 activating protein
31	c3qwxX_	Alignment	not modelled	99.6	29	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
32	c2enmA_	Alignment	not modelled	99.6	24	PDB header: endocytosis Chain: A: PDB Molecule: sorting nexin-9; PDBTitle: solution structure of the sh3 domain from mouse sorting2 nexin-9
33	c2dl3A_	Alignment	not modelled	99.6	26	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
34	c2epdA_	Alignment	not modelled	99.6	28	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
35	c2cubA_	Alignment	not modelled	99.6	25	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
36	c3rniA_	Alignment	not modelled	99.6	17	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)
37	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)
38	c1x2qA_	Alignment	not modelled	99.6	28	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
39	c1x43A_	Alignment	not modelled	99.6	25	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)
40	d2v1ra1	Alignment	not modelled	99.6	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
41	c2djgA_	Alignment	not modelled	99.6	28	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
42	c1wx6A_	Alignment	not modelled	99.6	27	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
43	c1zukA_	Alignment	not modelled	99.6	24	PDB header: contractile protein Chain: A: PDB Molecule: myosin tail region-interacting protein mtl1; PDBTitle: yeast bbc1 sh3 domain complexed with a peptide from las17
44	c2e5kA_	Alignment	not modelled	99.6	20	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
45	d1udla_	Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
46	c2yuoA_	Alignment	not modelled	99.6	31	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
47	c2ke9A_	Alignment	not modelled	99.6	21	PDB header: protein binding Chain: A: PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
48	c2yunA_	Alignment	not modelled	99.6	30	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
49	c2dl4A_	Alignment	not modelled	99.6	24	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
50	c2eqiA_	Alignment	not modelled	99.6	36	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
51	c2dlpA_	Alignment	not modelled	99.6	22	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
52	d1wlpb2	Alignment	not modelled	99.6	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
53	d1phta_	Alignment	not modelled	99.6	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
						PDB header: signaling protein Chain: A: PDB Molecule: megakaryocyte-associated tyrosine-

54	c1x6gA_	Alignment	not modelled	99.6	10	protein kinase; PDBTitle: solution structures of the sh3 domain of human2 megakaryocyte-associated tyrosine-protein kinase.
55	c2bzxA_	Alignment	not modelled	99.6	24	PDB header: sh3 domain Chain: A: PDB Molecule: crk-like protein; PDBTitle: atomic model of crkl-sh3c monomer
56	c2nwmA_	Alignment	not modelled	99.6	21	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
57	d1wiea_	Alignment	not modelled	99.6	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
58	c2df6A_	Alignment	not modelled	99.6	35	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
59	c2dl5A_	Alignment	not modelled	99.6	30	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the first sh3 domain of human2 kiaa0769 protein
60	c2csiA_	Alignment	not modelled	99.6	31	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
61	d1j3ta_	Alignment	not modelled	99.6	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
62	c2dl8A_	Alignment	not modelled	99.6	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
63	c4esrB_	Alignment	not modelled	99.6	28	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
64	d1gria2	Alignment	not modelled	99.6	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
65	c2pqhA_	Alignment	not modelled	99.6	28	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
66	d1uhfa_	Alignment	not modelled	99.6	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
67	d1utia_	Alignment	not modelled	99.6	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
68	c1wi7A_	Alignment	not modelled	99.6	31	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
69	c1bb9A_	Alignment	not modelled	99.6	20	PDB header: transferase Chain: A: PDB Molecule: amphiphysin 2; PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
70	d1bb9a_	Alignment	not modelled	99.6	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
71	c2ct3A_	Alignment	not modelled	99.6	25	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
72	d1u5sa1	Alignment	not modelled	99.6	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
73	c2d1xD_	Alignment	not modelled	99.6	43	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
74	c2ct4A_	Alignment	not modelled	99.6	25	PDB header: signaling protein Chain: A: PDB Molecule: cdc42-interacting protein 4; PDBTitle: solution strcture of the sh3 domain of the cdc42-2 interacting protein 4
75	d1i07a_	Alignment	not modelled	99.6	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
76	d1jo8a_	Alignment	not modelled	99.6	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
77	c1w70A_	Alignment	not modelled	99.6	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
78	d1ov3a2	Alignment	not modelled	99.6	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
79	c2k2mA_	Alignment	not modelled	99.6	24	PDB header: signaling protein Chain: A: PDB Molecule: eps8-like protein 1; PDBTitle: structural basis of pxxdy motif recognition in sh3 binding
						Fold: SH3-like barrel

80	d1ugva_	Alignment	not modelled	99.6	35	Superfamily: SH3-domain Family: SH3-domain
81	c2csqA_	Alignment	not modelled	99.6	22	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
82	c2dilA_	Alignment	not modelled	99.6	31	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
83	c2l0aA_	Alignment	not modelled	99.6	32	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
84	d1ue9a_	Alignment	not modelled	99.6	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
85	c2dl7A_	Alignment	not modelled	99.6	27	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
86	c2dnuA_	Alignment	not modelled	99.6	25	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
87	c2bz8B_	Alignment	not modelled	99.6	33	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
88	c1x69A_	Alignment	not modelled	99.6	33	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
89	c1yn8E_	Alignment	not modelled	99.6	26	PDB header: unknown function Chain: E: PDB Molecule: nap1-binding protein 2; PDBTitle: sh3 domain of yeast nbp2
90	d1sema_	Alignment	not modelled	99.6	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
91	c2rqvA_	Alignment	not modelled	99.6	25	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
92	c2j06B_	Alignment	not modelled	99.6	29	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
93	d1uj0a_	Alignment	not modelled	99.6	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	d1pwta_	Alignment	not modelled	99.6	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
95	d1uuea_	Alignment	not modelled	99.6	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
96	c2fpeB_	Alignment	not modelled	99.6	23	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
97	c2ekhA_	Alignment	not modelled	99.6	29	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
98	c2creA_	Alignment	not modelled	99.6	34	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hef-like protein; PDBTitle: solution structure of rsgi ruh-036, an sh3 domain from2 human cdna
99	d1k4us_	Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	d1gcqa_	Alignment	not modelled	99.6	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	c2rf0D_	Alignment	not modelled	99.6	30	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
102	c2kt1A_	Alignment	not modelled	99.6	23	PDB header: protein binding Chain: A: PDB Molecule: phosphatidylinositol 3-kinase regulatory subunit PDBTitle: solution nmr structure of the sh3 domain from the p85beta2 subunit of phosphatidylinositol 3-kinase from h.sapiens,3 northeast structural genomics consortium target hr5531e
103	c2d8jA_	Alignment	not modelled	99.6	35	PDB header: transferase Chain: A: PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh3 domain of fyn-related kinase
104	c2drnB_	Alignment	not modelled	99.6	40	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
						PDB header: motor protein

105	c2xmfA_	Alignment	not modelled	99.6	41	Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
106	c2egaA_	Alignment	not modelled	99.6	18	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
107	c2o2oA_	Alignment	not modelled	99.6	28	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
108	c1wyxA_	Alignment	not modelled	99.6	33	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
109	c2dbmA_	Alignment	not modelled	99.6	24	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
110	c1x2pA_	Alignment	not modelled	99.6	31	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	d1oota_	Alignment	not modelled	99.6	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	d1ov3a1	Alignment	not modelled	99.6	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	c2krnA_	Alignment	not modelled	99.6	28	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
114	d1ujya_	Alignment	not modelled	99.6	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	d1gbra_	Alignment	not modelled	99.6	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	d1e6ga_	Alignment	not modelled	99.6	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	c2cucA_	Alignment	not modelled	99.6	32	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
118	c2js0A_	Alignment	not modelled	99.6	31	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
119	c2egeA_	Alignment	not modelled	99.6	15	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
120	c1s1nA_	Alignment	not modelled	99.6	26	PDB header: cell adhesion Chain: A: PDB Molecule: nephrocystin 1; PDBTitle: sh3 domain of human nephrocystin