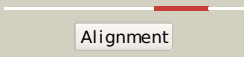
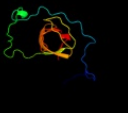
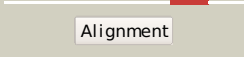
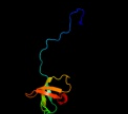
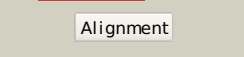
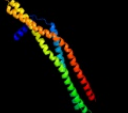
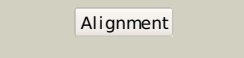
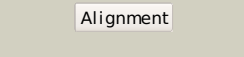
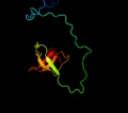
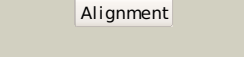
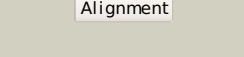
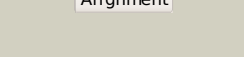
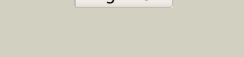
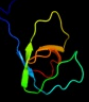


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1mv3A_	 Alignment		99.9	13	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: myc box dependent interacting protein 1; PDBTitle: nmr structure of the tumor suppressor bin1: alternative2 splicing in melanoma and interaction with c-myc
2	c2rqra_	 Alignment		99.7	17	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		99.7	31	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
4	c4avmA_	 Alignment		99.7	26	PDB header: protein binding Chain: A: PDB Molecule: bridging integrator 2; PDBTitle: crystal structure of the n-bar domain of human bridging2 integrator 2.
5	d1ng2a2	 Alignment		99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
6	c2lqnA_	 Alignment		99.7	20	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
7	c2eyzA_	 Alignment		99.6	21	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
8	c2l3sA_	 Alignment		99.6	19	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
9	c3nmzD_	 Alignment		99.6	29	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of apc complexed with ascf
10	c2jxbA_	 Alignment		99.6	23	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
11	c2dx1A_	 Alignment		99.6	40	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein ascf

12	c2ed0A_	Alignment		99.6	38	PDB header: signaling protein Chain: A: PDB Molecule: abl interactor 2; PDBTitle: solution structure of the sh3 domain of abl interactor 2 (abelson interactor 2)
13	c2dlpA_	Alignment		99.6	30	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
14	c2oi3A_	Alignment		99.6	35	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)
15	c3qwyA_	Alignment		99.6	35	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
16	d1ov3a2	Alignment		99.6	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
17	c3qwxX_	Alignment		99.6	34	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
18	c1x43A_	Alignment		99.6	23	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)
19	c1x2qA_	Alignment		99.6	30	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
20	c2eyyA_	Alignment		99.6	22	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
21	c1zukA_	Alignment	not modelled	99.6	31	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
22	d1oota_	Alignment	not modelled	99.6	50	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
23	c1x6gA_	Alignment	not modelled	99.6	13	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.
24	c2yt6A_	Alignment	not modelled	99.6	31	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
25	c2ysqA_	Alignment	not modelled	99.6	30	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
26	d1wiea_	Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
27	c2epdA_	Alignment	not modelled	99.6	31	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
						PDB header: unknown function

28	c2d8hA	Alignment	not modelled	99.6	39	Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
29	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
30	c2csqA	Alignment	not modelled	99.6	28	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
31	d1uj0a	Alignment	not modelled	99.6	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
32	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
33	c2dl7A	Alignment	not modelled	99.6	32	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
34	c2djgA	Alignment	not modelled	99.6	32	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
35	c2df6A	Alignment	not modelled	99.6	36	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
36	c2kbtA	Alignment	not modelled	99.6	33	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
37	c2yuoA	Alignment	not modelled	99.6	30	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
38	d1udla	Alignment	not modelled	99.6	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
39	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
40	c2d1xD	Alignment	not modelled	99.6	41	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
41	c2xmFA	Alignment	not modelled	99.6	42	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
42	c2nwmA	Alignment	not modelled	99.5	29	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
43	c2cubA	Alignment	not modelled	99.5	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
44	c2dbkA	Alignment	not modelled	99.5	24	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
45	c2ebpA	Alignment	not modelled	99.5	23	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
46	d1jqqa	Alignment	not modelled	99.5	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
47	c1zlmA	Alignment	not modelled	99.5	31	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
48	c2kymA	Alignment	not modelled	99.5	29	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
49	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
50	c1wyxA	Alignment	not modelled	99.5	36	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
51	c2yupA	Alignment	not modelled	99.5	30	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
52	c2dmoA	Alignment	not modelled	99.5	22	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)
53	d2v1ra1	Alignment	not modelled	99.5	30	Fold: SH3-like barrel Superfamily: SH3-domain

						Family: SH3-domain
54	c2dl8A_	Alignment	not modelled	99.5	29	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
55	dlu5sa1	Alignment	not modelled	99.5	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
56	dlycsb2	Alignment	not modelled	99.5	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
57	c1wx6A_	Alignment	not modelled	99.5	25	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
58	c3rniA_	Alignment	not modelled	99.5	26	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)
59	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
60	dl oe ba_	Alignment	not modelled	99.5	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
61	c2dl3A_	Alignment	not modelled	99.5	25	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
62	dlwlpb2	Alignment	not modelled	99.5	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
63	c2dl4A_	Alignment	not modelled	99.5	33	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
64	c2eyxA_	Alignment	not modelled	99.5	22	PDB header: signaling protein Chain: A: PDB Molecule: v-cr k sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
65	dluffa_	Alignment	not modelled	99.5	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	dluhca_	Alignment	not modelled	99.5	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
67	c2dbmA_	Alignment	not modelled	99.5	35	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
68	dlutia_	Alignment	not modelled	99.5	40	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
69	c2egeA_	Alignment	not modelled	99.5	25	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
70	c2cudA_	Alignment	not modelled	99.5	20	PDB header: signaling protein Chain: A: PDB Molecule: src-like-adaptor; PDBTitle: solution structure of the sh3 domain of the human src-like2 adppter protein (slap)
71	c2k6dA_	Alignment	not modelled	99.5	31	PDB header: sh3 domain/ubiquitin Chain: A: PDB Molecule: sh3 domain-containing kinase-binding protein 1; PDBTitle: cin85 sh3-c domain in complex with ubiquitin
72	dlpwta_	Alignment	not modelled	99.5	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
73	dlue9a_	Alignment	not modelled	99.5	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	dlgria2	Alignment	not modelled	99.5	41	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
75	dlilja_	Alignment	not modelled	99.5	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
76	clov3A_	Alignment	not modelled	99.5	28	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
77	c2azsA_	Alignment	not modelled	99.5	28	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)
78	c1w70A_	Alignment	not modelled	99.5	38	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
79	c2ed1A_	Alignment	not modelled	99.5	30	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
80	c1x6bA_	Alignment	not modelled	99.5	17	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
81	d1uuea_	Alignment	not modelled	99.5	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
82	c2pqhA_	Alignment	not modelled	99.5	31	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
83	d1gcqa_	Alignment	not modelled	99.5	43	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	c2kt1A_	Alignment	not modelled	99.5	30	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
85	d1k4us_	Alignment	not modelled	99.5	39	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
86	c4esrB_	Alignment	not modelled	99.5	44	PDB header: protein binding Chain: B: PDB Molecule: joubertin; 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
87	c2bz8B_	Alignment	not modelled	99.5	34	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	c1wi7A_	Alignment	not modelled	99.5	32	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
89	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
90	d1ujya_	Alignment	not modelled	99.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
91	c2dila_	Alignment	not modelled	99.5	36	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
92	c2ct4A_	Alignment	not modelled	99.5	21	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
93	c2dm1A_	Alignment	not modelled	99.5	34	PDB header: signaling protein Chain: A: PDB Molecule: protein vav-2; PDBTitle: solution structure of the second sh3 domain of human2 protein vav-2
94	d1phta_	Alignment	not modelled	99.5	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
95	c2jteA_	Alignment	not modelled	99.5	32	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: third sh3 domain of cd2ap
96	c1zx6A_	Alignment	not modelled	99.5	42	PDB header: protein binding Chain: A: PDB Molecule: yp154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
97	c2krnA_	Alignment	not modelled	99.5	33	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
98	c1x2pA_	Alignment	not modelled	99.5	35	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
99	c2a28D_	Alignment	not modelled	99.5	33	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
100	c2ct3A_	Alignment	not modelled	99.5	32	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
101	c2drmB_	Alignment	not modelled	99.5	46	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
102	d1sema_	Alignment	not modelled	99.5	43	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	c2csiA_	Alignment	not modelled	99.5	33	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
104	c2egaA_	Alignment	not modelled	99.5	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
105	dlspka_	Alignment	not modelled	99.5	25 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
106	c2o2oA_	Alignment	not modelled	99.5	30 PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
107	dlu06a1	Alignment	not modelled	99.5	35 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	c2rf0D_	Alignment	not modelled	99.5	38 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
109	dlugva_	Alignment	not modelled	99.5	31 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
110	dle6ga_	Alignment	not modelled	99.5	32 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
111	dlnega_	Alignment	not modelled	99.5	35 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	c1negA_	Alignment	not modelled	99.5	35 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
113	c2creA_	Alignment	not modelled	99.5	27 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
114	c2i0nA_	Alignment	not modelled	99.5	30 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
115	dlgbra_	Alignment	not modelled	99.5	33 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	dlbb9a_	Alignment	not modelled	99.5	25 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	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
118	dljo8a_	Alignment	not modelled	99.5	41 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	dlgria1	Alignment	not modelled	99.5	41 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	c2fpeB_	Alignment	not modelled	99.5	20 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