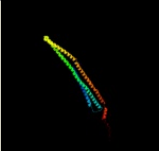
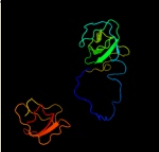
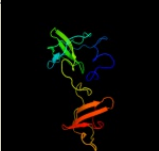
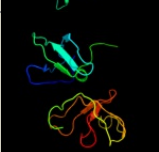
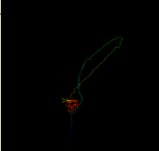
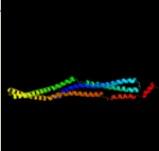


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1griA_	 Alignment		100.0	37	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
2	d2efla1	 Alignment		99.9	21	Fold: BAR/IMD domain-like Superfamily: BAR/IMD domain-like Family: FCH domain
3	c2lqnA_	 Alignment		99.9	26	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
4	c2eyzA_	 Alignment		99.9	20	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
5	d2efka1	 Alignment		99.9	20	Fold: BAR/IMD domain-like Superfamily: BAR/IMD domain-like Family: FCH domain
6	c1ng2A_	 Alignment		99.9	25	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
7	c1ov3A_	 Alignment		99.9	25	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
8	c2l3sA_	 Alignment		99.9	25	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
9	c1mv3A_	 Alignment		99.9	18	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
10	c3m3wA_	 Alignment		99.9	19	PDB header: endocytosis Chain: A: PDB Molecule: protein kinase c and casein kinase ii substrate protein 3; PDBTitle: crystal strcuture of mouse pacsin3 bar domain mutant
11	c3i2wB_	 Alignment		99.9	24	PDB header: endocytosis Chain: B: PDB Molecule: syndapin; PDBTitle: crystal structure of efc/f-bar domain of drosophila2 syndapin/pacsin

12	c2v0oB_	Alignment		99.9	17	PDB header: lipid-binding protein Chain: B: PDB Molecule: fch domain only protein 2; PDBTitle: fcho2 f-bar domain
13	c3hajA_	Alignment		99.9	20	PDB header: endocytosis Chain: A: PDB Molecule: human pacsin2 f-bar; PDBTitle: crystal structure of human pacsin2 f-bar domain (p2121212 lattice)
14	c3lllA_	Alignment		99.9	21	PDB header: endocytosis Chain: A: PDB Molecule: protein kinase c and casein kinase substrate in neurons PDBTitle: crystal structure of mouse pacsin2 f-bar domain
15	c3acoB_	Alignment		99.8	21	PDB header: endocytosis Chain: B: PDB Molecule: protein kinase c and casein kinase substrate in PDBTitle: crystal structure of the efc/f-bar domain of human2 pacsin2/syndapin ii (2.7 a)
16	c3hahA_	Alignment		99.8	17	PDB header: endocytosis Chain: A: PDB Molecule: human pacsin1 f-bar; PDBTitle: crystal structure of human pacsin1 f-bar domain (c2 lattice)
17	c3syvB_	Alignment		99.8	14	PDB header: endocytosis Chain: B: PDB Molecule: protein kinase c and casein kinase ii substrate protein 3; PDBTitle: crystal structure of mpacsin 3 f-bar domain mutant
18	c3qweA_	Alignment		99.8	14	PDB header: protein binding Chain: A: PDB Molecule: gem-interacting protein; PDBTitle: crystal structure of the n-terminal domain of the gem interacting2 protein
19	d1udla_	Alignment		99.8	39	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
20	d1ng2a2	Alignment		99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
21	c2csqA_	Alignment	not modelled	99.7	24	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
22	c2dx1A_	Alignment	not modelled	99.7	32	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
23	c2dlpA_	Alignment	not modelled	99.7	32	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
24	c2egeA_	Alignment	not modelled	99.7	21	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
25	c2eyyA_	Alignment	not modelled	99.7	21	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
26	d1ov3a2	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
27	c3g9gA_	Alignment	not modelled	99.7	13	PDB header: endocytosis Chain: A: PDB Molecule: suppressor of yeast profilin deletion; PDBTitle: crystal structure of the n-terminal efc/f-bar domain of syp1
28	c3rnjA_	Alignment	not modelled	99.7	30	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) PDB header: signaling protein

29	c1x6gA_	Alignment	not modelled	99.7	13	Chain: A; PDB Molecule: megakaryocyte-associated tyrosine-protein kinase; PDBTitle: solution structures of the sh3 domain of human2 megakaryocyte-associated tyrosine-protein kinase.
30	c3nmzD_	Alignment	not modelled	99.7	27	PDB header: cell adhesion/cell cycle Chain: D; PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of apc complexed with asf
31	c2df6A_	Alignment	not modelled	99.7	41	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
32	c3qwyA_	Alignment	not modelled	99.7	20	PDB header: signaling protein Chain: A; PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
33	c2rqrA_	Alignment	not modelled	99.7	19	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
34	c1x43A_	Alignment	not modelled	99.7	20	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)
35	c2kbtA_	Alignment	not modelled	99.7	39	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
36	c2xmfA_	Alignment	not modelled	99.7	45	PDB header: motor protein Chain: A; PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
37	c2e5kA_	Alignment	not modelled	99.7	29	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
38	c2csiA_	Alignment	not modelled	99.7	32	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
39	d1wiea_	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
40	c2cubA_	Alignment	not modelled	99.7	29	PDB header: signaling protein Chain: A; PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
41	d1gcqa_	Alignment	not modelled	99.7	46	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
42	c2ed0A_	Alignment	not modelled	99.6	42	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)
43	d1sema_	Alignment	not modelled	99.6	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
44	c2nwmA_	Alignment	not modelled	99.6	36	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
45	d1bb9a_	Alignment	not modelled	99.6	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
46	c1bb9A_	Alignment	not modelled	99.6	31	PDB header: transferase Chain: A; PDB Molecule: amphiphysin 2; PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
47	c2jxbA_	Alignment	not modelled	99.6	26	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
48	c2dbkA_	Alignment	not modelled	99.6	28	PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
49	c2yuoA_	Alignment	not modelled	99.6	29	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
50	d1k4us_	Alignment	not modelled	99.6	40	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
51	c2dl7A_	Alignment	not modelled	99.6	41	PDB header: structural genomics, unknown function Chain: A; PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
52	c2djgA_	Alignment	not modelled	99.6	29	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
53	d1utiA_	Alignment	not modelled	99.6	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
54	c2dmoA_	Alignment	not modelled	99.6	23	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)
55	c1x2qA_	Alignment	not modelled	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
56	c2l0aA_	Alignment	not modelled	99.6	30	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
57	d1wlpb2	Alignment	not modelled	99.6	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
58	d1oota_	Alignment	not modelled	99.6	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
59	c2yunA_	Alignment	not modelled	99.6	37	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
60	d1gria2	Alignment	not modelled	99.6	45	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
61	c1zlmA_	Alignment	not modelled	99.6	39	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
62	c3qwxX_	Alignment	not modelled	99.6	22	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
63	d1uuea_	Alignment	not modelled	99.6	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
64	c2eyxA_	Alignment	not modelled	99.6	26	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
65	c2ebpA_	Alignment	not modelled	99.6	24	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
66	d1ugva_	Alignment	not modelled	99.6	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
67	c1zx6A_	Alignment	not modelled	99.6	43	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
68	d1spka_	Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
69	d1uhca_	Alignment	not modelled	99.6	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
70	c2bz8B_	Alignment	not modelled	99.6	39	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
71	c2fpeB_	Alignment	not modelled	99.6	28	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
72	c2j6kE_	Alignment	not modelled	99.6	40	PDB header: protein binding Chain: E: PDB Molecule: cd2-associated protein; PDBTitle: n-terminal sh3 domain of cms (cd2ap human homolog)
73	c1w70A_	Alignment	not modelled	99.6	35	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
74	d1pwta_	Alignment	not modelled	99.6	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
75	c2dnuA_	Alignment	not modelled	99.6	22	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
76	d1ujya_	Alignment	not modelled	99.6	42	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
77	d1u06a1	Alignment	not modelled	99.6	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
78	c2ysqA_	Alignment	not modelled	99.6	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
79	c1x69A_	Alignment	not modelled	99.6	32	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
						PDB header: signaling protein Chain: D: PDB Molecule: bzz1 protein;

80	c2a28D_	Alignment	not modelled	99.6	45	PDBTitle: atomic-resolution crystal structure of the second sh32 domain of yeast bzz1 determined from a pseudomerohedrally3 twinned crystal
81	dluj0a_	Alignment	not modelled	99.6	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
82	dlphta_	Alignment	not modelled	99.6	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
83	dlilja_	Alignment	not modelled	99.6	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	c1wyxA_	Alignment	not modelled	99.6	38	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
85	c2i0nA_	Alignment	not modelled	99.6	29	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
86	dl0eba_	Alignment	not modelled	99.6	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
87	clzuka_	Alignment	not modelled	99.6	22	PDB header: contractile protein Chain: A: PDB Molecule: myosin tail region-interacting protein mti1; PDBTitle: yeast bbc1 sh3 domain complexed with a peptide from las17
88	c2dlxD_	Alignment	not modelled	99.6	40	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
89	c2pqhA_	Alignment	not modelled	99.6	33	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	c2drnB_	Alignment	not modelled	99.6	52	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
91	dljqqa_	Alignment	not modelled	99.6	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
92	d2v1ra1	Alignment	not modelled	99.6	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
93	dlnega_	Alignment	not modelled	99.6	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	clnegA_	Alignment	not modelled	99.6	33	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
95	c2cudA_	Alignment	not modelled	99.6	21	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)
96	c2rf0D_	Alignment	not modelled	99.6	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
97	dlu5sa1	Alignment	not modelled	99.6	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
98	c2dl4A_	Alignment	not modelled	99.6	27	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
99	c2oi3A_	Alignment	not modelled	99.6	27	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)
100	dlgbra_	Alignment	not modelled	99.6	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	dle6ga_	Alignment	not modelled	99.6	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
102	c2kt1A_	Alignment	not modelled	99.6	29	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	clx6bA_	Alignment	not modelled	99.6	26	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
104	c2yt6A_	Alignment	not modelled	99.6	30	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
						PDB header: protein binding

105	c2ke9A_	Alignment	not modelled	99.6	28	Chain: A: PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
106	c2kymA_	Alignment	not modelled	99.6	23	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of the bem1p sh3-ci domain from <i>I.elongisporus</i> in2 complex with ste20p peptide
107	c2epdA_	Alignment	not modelled	99.6	30	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
108	c2krnA_	Alignment	not modelled	99.6	34	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
109	c2ed1A_	Alignment	not modelled	99.6	29	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
110	c2azsA_	Alignment	not modelled	99.6	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)
111	c2jteA_	Alignment	not modelled	99.6	36	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: third sh3 domain of cd2ap
112	d1uffa_	Alignment	not modelled	99.6	41	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	d1cska_	Alignment	not modelled	99.6	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	c2dbmA_	Alignment	not modelled	99.6	39	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
115	c2ct3A_	Alignment	not modelled	99.6	32	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
116	d1e6ha_	Alignment	not modelled	99.6	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	d1ycsb2	Alignment	not modelled	99.6	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
118	c2k6dA_	Alignment	not modelled	99.6	45	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
119	c2dilA_	Alignment	not modelled	99.6	32	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
120	c4esrB_	Alignment	not modelled	99.6	44	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