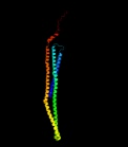
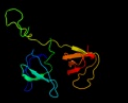
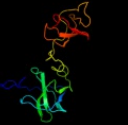
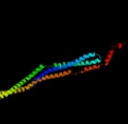
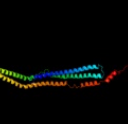


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2efla1	 Alignment		100.0	18	Fold: BAR/IMD domain-like Superfamily: BAR/IMD domain-like Family: FCH domain
2	d2efka1	 Alignment		100.0	19	Fold: BAR/IMD domain-like Superfamily: BAR/IMD domain-like Family: FCH domain
3	c1griA_	 Alignment		100.0	29	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
4	c2lqnA_	 Alignment		100.0	19	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
5	c2eyzA_	 Alignment		100.0	18	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
6	c3m3wA_	 Alignment		100.0	15	PDB header: endocytosis Chain: A: PDB Molecule: protein kinase c and casein kinase ii substrate protein 3; PDBTitle: crystal structure of mouse pacsin3 bar domain mutant
7	c2v0oB_	 Alignment		100.0	14	PDB header: lipid-binding protein Chain: B: PDB Molecule: fch domain only protein 2; PDBTitle: fcho2 f-bar domain
8	c2l3sA_	 Alignment		99.9	22	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
9	c3i2wB_	 Alignment		99.9	18	PDB header: endocytosis Chain: B: PDB Molecule: syndapin; PDBTitle: crystal structure of efc/f-bar domain of drosophila2 syndapin/pacsin
10	c3l1IA_	 Alignment		99.9	16	PDB header: endocytosis Chain: A: PDB Molecule: protein kinase c and casein kinase ii substrate in neurons PDBTitle: crystal structure of mouse pacsin2 f-bar domain
11	c3hajA_	 Alignment		99.9	15	PDB header: endocytosis Chain: A: PDB Molecule: human pacsin2 f-bar; PDBTitle: crystal structure of human pacsin2 f-bar domain (p2121212 lattice)

12	c1ng2A_	Alignment		99.9	23	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
13	c1ov3A_	Alignment		99.9	22	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of the p22phox-p47phox complex
14	c3acoB_	Alignment		99.9	16	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)
15	c3hahA_	Alignment		99.9	16	PDB header: endocytosis Chain: A: PDB Molecule: human pacsin1 f-bar; PDBTitle: crystal structure of human pacsin1 f-bar domain (c2 lattice)
16	c3syvB_	Alignment		99.9	17	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
17	c1mv3A_	Alignment		99.9	16	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
18	c3qweA_	Alignment		99.9	9	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	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
20	c3g9gA_	Alignment		99.8	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
21	d1ng2a2	Alignment	not modelled	99.8	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
22	c2csqA_	Alignment	not modelled	99.8	26	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
23	c2rqrA_	Alignment	not modelled	99.8	24	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
24	c2jxbA_	Alignment	not modelled	99.8	13	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
25	c2dx1A_	Alignment	not modelled	99.7	28	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asecf
26	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 asecf
27	c2eyyA_	Alignment	not modelled	99.7	14	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
28	d1ov3a2	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

29	c2dlpA	Alignment	not modelled	99.7	23	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
30	c2egeA	Alignment	not modelled	99.7	33	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
31	c2l0aA	Alignment	not modelled	99.7	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
32	c1x6gA	Alignment	not modelled	99.7	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.
33	c2ed0A	Alignment	not modelled	99.7	35	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)
34	c2kbtA	Alignment	not modelled	99.7	32	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
35	c1x2qA	Alignment	not modelled	99.7	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
36	c2csiA	Alignment	not modelled	99.7	40	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
37	d1wiea	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
38	c2cubA	Alignment	not modelled	99.7	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
39	c2dl7A	Alignment	not modelled	99.7	34	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
40	c2yuoA	Alignment	not modelled	99.7	34	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
41	c2a28D	Alignment	not modelled	99.7	96	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
42	c1zlmA	Alignment	not modelled	99.7	32	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
43	c2ysqA	Alignment	not modelled	99.7	25	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
44	c2nwmA	Alignment	not modelled	99.7	28	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	d1k4us	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
46	c1x43A	Alignment	not modelled	99.7	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)
47	c2e5kA	Alignment	not modelled	99.7	31	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
48	c2epdA	Alignment	not modelled	99.7	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
49	d1utia	Alignment	not modelled	99.7	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
50	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)
51	d1oeba	Alignment	not modelled	99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
52	c2df6A	Alignment	not modelled	99.7	30	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
53	c3qwxX	Alignment	not modelled	99.7	10	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174

54	d1sema_	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
55	c3qwyA_	Alignment	not modelled	99.7	10	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
56	c2oi3A_	Alignment	not modelled	99.7	18	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)
57	c2yunA_	Alignment	not modelled	99.7	34	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
58	d1gcqa_	Alignment	not modelled	99.7	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
59	d1uj0a_	Alignment	not modelled	99.7	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
60	c2xmfa_	Alignment	not modelled	99.7	43	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
61	c2kymA_	Alignment	not modelled	99.7	29	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure analysis of the bem1p sh3-ci domain from l.elongisporus in2 complex with ste20p peptide
62	c2eyxA_	Alignment	not modelled	99.7	23	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
63	c2yt6A_	Alignment	not modelled	99.7	23	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
64	c2ebpA_	Alignment	not modelled	99.7	20	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	c2djqa_	Alignment	not modelled	99.7	27	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
66	d1i1ja_	Alignment	not modelled	99.7	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
67	d1gbra_	Alignment	not modelled	99.7	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
68	c1zukA_	Alignment	not modelled	99.7	25	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
69	c2dl8A_	Alignment	not modelled	99.7	21	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
70	c2ekhA_	Alignment	not modelled	99.7	27	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
71	c2dmoA_	Alignment	not modelled	99.7	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)
72	c2dbkA_	Alignment	not modelled	99.7	24	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
73	c1zx6A_	Alignment	not modelled	99.7	32	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
74	d1ugva_	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
75	c2dila_	Alignment	not modelled	99.7	33	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
76	d1gria2	Alignment	not modelled	99.7	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
77	d1loota_	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
78	d1wlpb2	Alignment	not modelled	99.7	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
						PDB header: cell invasion

79	c2d1xD	Alignment	not modelled	99.7	53	Chain: D: PDB Molecule: cortactin isoform a; PDBTitle: the crystal structure of the cortactin-sh3 domain and amap1-2 peptide complex
80	d1uuea	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
81	c1wyxA	Alignment	not modelled	99.7	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
82	c2i0nA	Alignment	not modelled	99.7	24	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
83	c1bb9A	Alignment	not modelled	99.7	23	PDB header: transferase Chain: A: PDB Molecule: amphiphysin 2; PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
84	d1bb9a	Alignment	not modelled	99.7	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
85	c2dl3A	Alignment	not modelled	99.7	31	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
86	c2drmB	Alignment	not modelled	99.7	38	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
87	c2dl4A	Alignment	not modelled	99.7	27	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
88	c2o2oA	Alignment	not modelled	99.7	32	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
89	c2dnuA	Alignment	not modelled	99.7	30	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	c2bz8B	Alignment	not modelled	99.7	42	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
91	d1u5sa1	Alignment	not modelled	99.7	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
92	c2j6kE	Alignment	not modelled	99.7	37	PDB header: protein binding Chain: E: PDB Molecule: cd2-associated protein; PDBTitle: n-terminal sh3 domain of cms (cd2ap human homolog)
93	c2azsA	Alignment	not modelled	99.7	32	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)
94	c1w70A	Alignment	not modelled	99.7	27	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
95	c1x69A	Alignment	not modelled	99.7	42	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
96	c2pqhA	Alignment	not modelled	99.7	26	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
97	d1pwta	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
98	c2dbmA	Alignment	not modelled	99.7	31	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
99	c2krnA	Alignment	not modelled	99.7	32	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
100	d2v1ra1	Alignment	not modelled	99.7	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	c2ed1A	Alignment	not modelled	99.7	32	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
102	d1ycsb2	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	d1ujya	Alignment	not modelled	99.7	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
104	d1spka	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
						PDB header: transferase

105	c1x2pA_	Alignment	not modelled	99.7	25	Chain: A: PDB Molecule: protein arginine n-methyltransferase 2; PDBTitle: solution structure of the sh3 domain of the protein2 arginine n-methyltransferase 2
106	d1u06a1	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
107	d1uhca_	Alignment	not modelled	99.7	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	c1wi7A_	Alignment	not modelled	99.7	34	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
109	c2d8hA_	Alignment	not modelled	99.7	26	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
110	d1nega_	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
111	c1negA_	Alignment	not modelled	99.7	26	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
112	d1ov3a1	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	d1ue9a_	Alignment	not modelled	99.7	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	d1jqqa_	Alignment	not modelled	99.7	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	d1e6ga_	Alignment	not modelled	99.6	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	c2yupA_	Alignment	not modelled	99.6	28	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
117	c2egaA_	Alignment	not modelled	99.6	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
118	d1e6ha_	Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	c2js0A_	Alignment	not modelled	99.6	29	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
120	c2x3wD_	Alignment	not modelled	99.6	38	PDB header: endocytosis Chain: D: PDB Molecule: protein kinase c and casein kinase substrate in neurons PDBTitle: structure of mouse syndapin i (crystal form 2)