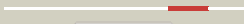
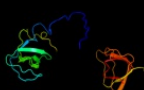


Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	dlhqz1_	 Alignment		100.0	32	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
2	c2lqnA_	 Alignment		100.0	21	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
3	c1griA_	 Alignment		100.0	38	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
4	c2eyzA_	 Alignment		100.0	22	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
5	c1x67A_	 Alignment		100.0	34	PDB header: protein binding Chain: A: PDB Molecule: drebrin-like protein; PDBTitle: solution structure of the cofilin homology domain of hip-552 (drebrin-like protein)
6	dludma_	 Alignment		100.0	21	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
7	c2l3sA_	 Alignment		100.0	23	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
8	c1ng2A_	 Alignment		100.0	25	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
9	dlwnja_	 Alignment		100.0	19	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
10	c1ov3A_	 Alignment		100.0	25	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
11	dl13ya1	 Alignment		100.0	19	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like

12	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
13	d1udla_	Alignment		99.8	43	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
14	d1cfya_	Alignment		99.8	20	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
15	d1cnua_	Alignment		99.8	13	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
16	d1ng2a2	Alignment		99.8	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
17	c3nmzD_	Alignment		99.7	27	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
18	c2kymA_	Alignment		99.7	26	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
19	c2xfaA_	Alignment		99.7	8	PDB header: protein binding Chain: A: PDB Molecule: actin depolymerization factor 2; PDBTitle: crystal structure of plasmodium berghei actin2 depolymerization factor 2
20	c1x2qA_	Alignment		99.7	41	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
21	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 asef
22	d1f7sa_	Alignment	not modelled	99.7	12	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
23	c2ebpA_	Alignment	not modelled	99.7	19	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
24	c2kd5A_	Alignment	not modelled	99.7	16	PDB header: hormone Chain: A: PDB Molecule: actin severing and dynamics regulatory protein; PDBTitle: solution structure of adf/cofilin (ldcof) from leishmania donovani
25	c2djqa_	Alignment	not modelled	99.7	33	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
26	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
27	c2rqrA_	Alignment	not modelled	99.7	18	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

28	c2dybA	Alignment	not modelled	99.7	19	PDB header: oxidoreductase Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: the crystal structure of human p40(phox)
29	c2ysqA	Alignment	not modelled	99.7	27	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
30	c2xmfA	Alignment	not modelled	99.7	40	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
31	c2cubA	Alignment	not modelled	99.7	31	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
32	c2drmB	Alignment	not modelled	99.7	50	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
33	c3qwyA	Alignment	not modelled	99.7	20	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
34	d1ov3a2	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
35	c2d1xD	Alignment	not modelled	99.7	51	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
36	c2dbmA	Alignment	not modelled	99.7	43	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
37	c2ed0A	Alignment	not modelled	99.7	38	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)
38	c2dl3A	Alignment	not modelled	99.7	36	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
39	c2egeA	Alignment	not modelled	99.7	27	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
40	d1v6fa	Alignment	not modelled	99.7	12	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
41	c2yunA	Alignment	not modelled	99.7	36	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
42	c2l72A	Alignment	not modelled	99.7	21	PDB header: protein binding Chain: A: PDB Molecule: actin depolymerizing factor, putative; PDBTitle: solution structure and dynamics of adf from toxoplasma gondii (tgadf)
43	c2nwmA	Alignment	not modelled	99.7	44	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
44	c2i2qA	Alignment	not modelled	99.7	18	PDB header: actin-binding protein Chain: A: PDB Molecule: cofilin; PDBTitle: fission yeast cofilin
45	c2yuoA	Alignment	not modelled	99.7	36	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
46	c2eyyA	Alignment	not modelled	99.7	24	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
47	d1k4us	Alignment	not modelled	99.7	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
48	d1gcqa	Alignment	not modelled	99.7	48	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
49	c3qwxX	Alignment	not modelled	99.7	20	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
50	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
51	c2yupA	Alignment	not modelled	99.7	27	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
52	c2csqA	Alignment	not modelled	99.7	29	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: rim binding protein 2; PDBTitle: solution structure of the second sh3 domain of human rim-2 binding protein 2
53	c2dl4A	Alignment	not modelled	99.7	28	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
						PDB header: signaling protein

54	c2krnA_	Alignment	not modelled	99.7	36	Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
55	c2rqvA_	Alignment	not modelled	99.7	22	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
56	d1uj0a_	Alignment	not modelled	99.7	51	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
57	c1w70A_	Alignment	not modelled	99.7	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
58	c2dmoA_	Alignment	not modelled	99.7	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)
59	c1wi7A_	Alignment	not modelled	99.7	36	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
60	c2dlpA_	Alignment	not modelled	99.7	25	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
61	c2ekbA_	Alignment	not modelled	99.7	21	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
62	c2eyxA_	Alignment	not modelled	99.7	27	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	d1utia_	Alignment	not modelled	99.7	40	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
64	c2fpeB_	Alignment	not modelled	99.7	26	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
65	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
66	c1x43A_	Alignment	not modelled	99.7	27	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: sh3 domain grb2-like protein b1; PDBTitle: solution structure of the sh3 domain of endophilin b12 (sh3g1b1)
67	c1x69A_	Alignment	not modelled	99.7	43	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
68	c2js0A_	Alignment	not modelled	99.7	31	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
69	c2oi3A_	Alignment	not modelled	99.7	31	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)
70	d1wiea_	Alignment	not modelled	99.7	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
71	c2j06B_	Alignment	not modelled	99.7	25	PDB header: signal transduction Chain: B: PDB Molecule: ras gtpase-activating protein 1; PDBTitle: crystal structure of the rasgap sh3 domain at 1.8 angstrom2 resolution
72	c1x6gA_	Alignment	not modelled	99.7	14	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.
73	d1sema_	Alignment	not modelled	99.7	45	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	c2dl8A_	Alignment	not modelled	99.7	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
75	d1ujya_	Alignment	not modelled	99.7	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
76	d1uuea_	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
77	c2bz8B_	Alignment	not modelled	99.7	50	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
78	c2ed1A_	Alignment	not modelled	99.7	38	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
						PDB header: oxidoreductase activator

79	c1z9qA_	Alignment	not modelled	99.7	23	Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: solution structure of sh3 domain of p40phox
80	c2l0aA_	Alignment	not modelled	99.7	56	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
81	c1wyxA_	Alignment	not modelled	99.7	30	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	d1pwta_	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
83	c1zlmA_	Alignment	not modelled	99.7	44	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
84	c2dnuA_	Alignment	not modelled	99.7	16	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
85	d1j3ta_	Alignment	not modelled	99.7	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
86	d1gria2_	Alignment	not modelled	99.7	48	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
87	c2dila_	Alignment	not modelled	99.7	29	PDB header: cell adhesion Chain: A: PDB Molecule: proline-serine-threonine phosphatase-interacting PDBTitle: solution structure of the sh3 domain of the human proline-2 serine-threonine phosphatase-interacting protein 1
88	c2eqiA_	Alignment	not modelled	99.7	43	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
89	c1zuka_	Alignment	not modelled	99.7	27	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
90	d1loota_	Alignment	not modelled	99.7	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
91	d1vkka_	Alignment	not modelled	99.7	11	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
92	d1oeba_	Alignment	not modelled	99.7	40	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
93	c2epdA_	Alignment	not modelled	99.7	27	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
94	c2bzxA_	Alignment	not modelled	99.7	30	PDB header: sh3 domain Chain: A: PDB Molecule: crk-like protein; PDBTitle: atomic model of crkl-sh3c monomer
95	c2cudA_	Alignment	not modelled	99.7	17	PDB header: signaling protein Chain: A: PDB Molecule: src-like-adapter; PDBTitle: solution structure of the sh3 domain of the human src-like2 adpoter protein (slap)
96	c2pqhA_	Alignment	not modelled	99.7	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
97	d1l1ja_	Alignment	not modelled	99.7	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
98	d1ue9a_	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
99	c1x2pA_	Alignment	not modelled	99.7	34	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
100	d1u06a1_	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	c4esrB_	Alignment	not modelled	99.7	35	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
102	d1nega_	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	c1negaA_	Alignment	not modelled	99.7	27	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
104	d1spka_	Alignment	not modelled	99.7	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

105	d1gbra_	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
106	c2ct3A_	Alignment	not modelled	99.7	30	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
107	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
108	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
109	c3rnjA_	Alignment	not modelled	99.6	33	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)
110	d1e6ga_	Alignment	not modelled	99.6	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
111	d1phta_	Alignment	not modelled	99.6	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	c1zx6A_	Alignment	not modelled	99.6	31	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
113	c2i0nA_	Alignment	not modelled	99.6	25	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
114	c1yn8E_	Alignment	not modelled	99.6	30	PDB header: unknown function Chain: E: PDB Molecule: nap1-binding protein 2; PDBTitle: sh3 domain of yeast nbp2
115	d1uhfa_	Alignment	not modelled	99.6	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	c2ke9A_	Alignment	not modelled	99.6	24	PDB header: protein binding Chain: A: PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
117	c2d8hA_	Alignment	not modelled	99.6	24	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypotheticala2 protein sh3yl1
118	c2e5kA_	Alignment	not modelled	99.6	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
119	c2jxbA_	Alignment	not modelled	99.6	28	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
120	d1uhca_	Alignment	not modelled	99.6	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain