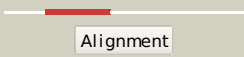
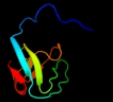
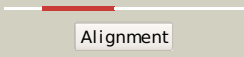
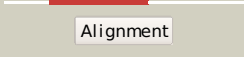
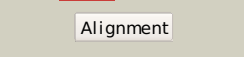
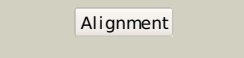
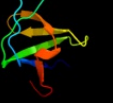
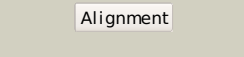
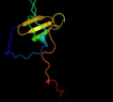
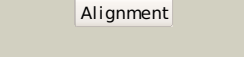
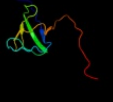
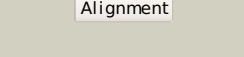
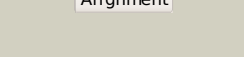
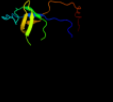
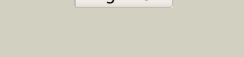
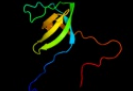
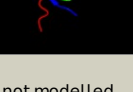


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1mv3A_	 Alignment		99.9	29	PDB header: endocytosis/exocytosis Chain: A; PDB Molecule: myc box dependent interacting protein 1; PDBTitle: nmr structure of the tumor suppressor bin1: alternative2 splicing in melanoma and interaction with c-myc
2	c1griA_	 Alignment		99.9	34	PDB header: signal transduction adaptor Chain: A; PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
3	d1ng2a2	 Alignment		99.8	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
4	c2dx1A_	 Alignment		99.8	31	PDB header: signaling protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asecf
5	c3nmzD_	 Alignment		99.8	29	PDB header: cell adhesion/cell cycle Chain: D; PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asecf
6	c1x2qA_	 Alignment		99.8	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
7	c2ysqA_	 Alignment		99.8	28	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
8	c2yunA_	 Alignment		99.8	32	PDB header: protein transport Chain: A; PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
9	c2yuoA_	 Alignment		99.8	28	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
10	c2dlpA_	 Alignment		99.8	28	PDB header: structural protein Chain: A; PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
11	c2rqrA_	 Alignment		99.8	30	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

12	c2eyyA_	Alignment		99.8	28	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
13	c2cubA_	Alignment		99.8	27	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
14	c2ed0A_	Alignment		99.8	29	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)
15	c2kymA_	Alignment		99.8	22	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
16	c2ekhA_	Alignment		99.8	22	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
17	c2djgA_	Alignment		99.8	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
18	c2epdA_	Alignment		99.8	23	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
19	c2dbmA_	Alignment		99.8	34	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
20	c2nwmA_	Alignment		99.8	31	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
21	c2lqnA_	Alignment	not modelled	99.8	21	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
22	c2ebpA_	Alignment	not modelled	99.8	28	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
23	d1udla_	Alignment	not modelled	99.8	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
24	c2pqhA_	Alignment	not modelled	99.8	32	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
25	c2yupA_	Alignment	not modelled	99.8	23	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
26	c2dl4A_	Alignment	not modelled	99.8	36	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
27	c2xmFA_	Alignment	not modelled	99.8	31	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
28	c2dmoA_	Alignment	not modelled	99.8	24	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)

29	c2df6A_	Alignment	not modelled	99.8	31	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
30	c2yt6A_	Alignment	not modelled	99.8	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
31	d1utia_	Alignment	not modelled	99.8	43	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
32	d1sema_	Alignment	not modelled	99.8	46	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
33	c1zx6A_	Alignment	not modelled	99.8	73	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
34	d1oeba_	Alignment	not modelled	99.8	41	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
35	c2dbkA_	Alignment	not modelled	99.8	20	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
36	c2dl3A_	Alignment	not modelled	99.8	28	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
37	d1ov3a2	Alignment	not modelled	99.8	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
38	c2csqA_	Alignment	not modelled	99.8	21	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
39	c2ed1A_	Alignment	not modelled	99.8	24	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
40	d1gria2	Alignment	not modelled	99.8	42	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
41	d1gcqa_	Alignment	not modelled	99.8	44	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
42	c2dl8A_	Alignment	not modelled	99.8	24	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
43	c1wi7A_	Alignment	not modelled	99.8	28	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
44	d1nega_	Alignment	not modelled	99.8	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
45	c1negA_	Alignment	not modelled	99.8	31	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
46	c2oi3A_	Alignment	not modelled	99.8	30	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)
47	c2eyxA_	Alignment	not modelled	99.8	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
48	d1uuea_	Alignment	not modelled	99.8	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
49	c1x43A_	Alignment	not modelled	99.8	22	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)
50	c2d1xD_	Alignment	not modelled	99.8	37	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
51	d1pwta_	Alignment	not modelled	99.8	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
52	c2l0aA_	Alignment	not modelled	99.8	37	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
53	d1k4us_	Alignment	not modelled	99.8	42	Fold: SH3-like barrel Superfamily: SH3-domain

						Family: SH3-domain
54	c2i0nA_	Alignment	not modelled	99.8	21	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
55	d1ue9a_	Alignment	not modelled	99.8	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
56	c1x69A_	Alignment	not modelled	99.8	30	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
57	c2o2oA_	Alignment	not modelled	99.8	25	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
58	c1w70A_	Alignment	not modelled	99.8	37	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
59	c2cudA_	Alignment	not modelled	99.8	14	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)
60	c2dila_	Alignment	not modelled	99.8	25	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
61	d1ujya_	Alignment	not modelled	99.8	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
62	c2l3sA_	Alignment	not modelled	99.8	21	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
63	c2eyzA_	Alignment	not modelled	99.8	20	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
64	d1j3ta_	Alignment	not modelled	99.8	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
65	d1uj0a_	Alignment	not modelled	99.8	41	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	c1x2pA_	Alignment	not modelled	99.8	28	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
67	c1wx6A_	Alignment	not modelled	99.8	24	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
68	d1u06a1	Alignment	not modelled	99.8	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
69	c2dnuA_	Alignment	not modelled	99.8	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
70	c3qwxX_	Alignment	not modelled	99.8	26	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
71	c2drnB_	Alignment	not modelled	99.8	44	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
72	c2bz8B_	Alignment	not modelled	99.8	29	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
73	c3qwyA_	Alignment	not modelled	99.8	26	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
74	c1x6gA_	Alignment	not modelled	99.8	17	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.
75	c1zlmA_	Alignment	not modelled	99.8	30	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
76	d1wiea_	Alignment	not modelled	99.8	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
77	d1wlpb2	Alignment	not modelled	99.8	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
78	c2kbtA_	Alignment	not modelled	99.8	37	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
79	d1uhfa_	Alignment	not modelled	99.8	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

80	d1e6ga_	Alignment	not modelled	99.8	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
81	c2jxbA_	Alignment	not modelled	99.7	27	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
82	c2fpeB_	Alignment	not modelled	99.7	24	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
83	d1uffa_	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	c1wyxA_	Alignment	not modelled	99.7	29	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	c2krnA_	Alignment	not modelled	99.7	36	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
86	c2egaA_	Alignment	not modelled	99.7	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
87	c2e5kA_	Alignment	not modelled	99.7	27	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
88	c1z9qA_	Alignment	not modelled	99.7	33	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: solution structure of sh3 domain of p40phox
89	c2eqiA_	Alignment	not modelled	99.7	27	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
90	c2dvjA_	Alignment	not modelled	99.7	25	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii
91	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
92	c1zukA_	Alignment	not modelled	99.7	26	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
93	c2dl7A_	Alignment	not modelled	99.7	37	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
94	c4esrB_	Alignment	not modelled	99.7	36	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
95	d1uhca_	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
96	d1h8ka_	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
97	d1e6ha_	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
98	c2js0A_	Alignment	not modelled	99.7	36	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
99	d1oota_	Alignment	not modelled	99.7	46	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	c2cucA_	Alignment	not modelled	99.7	31	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
101	d1ug1a_	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
102	c2egeA_	Alignment	not modelled	99.7	23	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
103	c1x6bA_	Alignment	not modelled	99.7	20	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	c2k2mA_	Alignment	not modelled	99.7	26	PDB header: signaling protein Chain: A: PDB Molecule: eps8-like protein 1; PDBTitle: structural basis of pxxdy motif recognition in sh3 binding
105	c2i06B_	Alignment	not modelled	99.7	18	PDB header: signal transduction Chain: B: PDB Molecule: ras gtpase-activating protein 1;

105	c2jv0b	Alignment	not modelled	99.7	18	PDBTitle: crystal structure of the rasgap sh3 domain at 1.8 angstrom2 resolution
106	c1ng2A	Alignment	not modelled	99.7	24	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
107	d1spka	Alignment	not modelled	99.7	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	c3rnjA	Alignment	not modelled	99.7	27	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)
109	c2d8jA	Alignment	not modelled	99.7	35	PDB header: transferase Chain: A: PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh3 domain of fyn-related kinase
110	c2bzxA	Alignment	not modelled	99.7	25	PDB header: sh3 domain Chain: A: PDB Molecule: crkl-like protein; PDBTitle: atomic model of crkl-sh3c monomer
111	c2d8hA	Alignment	not modelled	99.7	36	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypotheticala2 protein sh3yl1
112	d1jo8a	Alignment	not modelled	99.7	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	c2csiA	Alignment	not modelled	99.7	27	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
114	d1ugva	Alignment	not modelled	99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	d2v1ra1	Alignment	not modelled	99.7	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	c2enmA	Alignment	not modelled	99.7	21	PDB header: endocytosis Chain: A: PDB Molecule: sorting nexin-9; PDBTitle: solution structure of the sh3 domain from mouse sorting2 nexin-9
117	d1phta	Alignment	not modelled	99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
118	c2rqvA	Alignment	not modelled	99.7	19	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
119	c2eczA	Alignment	not modelled	99.7	24	PDB header: signaling protein Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: solution structure of the sh3 domain of sorbin and sh32 domain-containing protein 1
120	c2lcsA	Alignment	not modelled	99.7	30	PDB header: transferase, signaling protein Chain: A: PDB Molecule: nap1-binding protein 2; PDBTitle: yeast nbp2p sh3 domain in complex with a peptide from ste20p