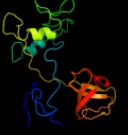
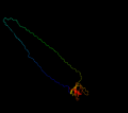
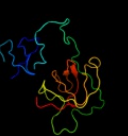
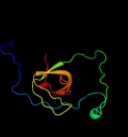
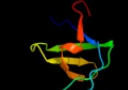
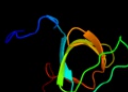
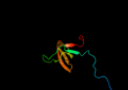


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1griA_	 Alignment		99.9	20	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
2	c1mv3A_	 Alignment		99.9	16	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: myc box dependent interacting protein 1; PDBTitle: nmr structure of the tumor suppressor bin1: alternative2 splicing in melanoma and interaction with c-myc
3	c2lqnA_	 Alignment		99.9	17	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
4	c2eyzA_	 Alignment		99.8	22	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
5	c2l3sA_	 Alignment		99.8	22	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
6	d1ng2a2	 Alignment		99.8	15	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
7	c1ng2A_	 Alignment		99.8	21	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
8	c2rqrA_	 Alignment		99.8	20	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
9	c2dx1A_	 Alignment		99.8	37	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
10	c1ov3A_	 Alignment		99.8	22	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
11	c1x2qA_	 Alignment		99.8	32	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

12	c2ysqA_	Alignment		99.8	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
13	d1udla_	Alignment		99.8	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
14	c3nmzD_	Alignment		99.8	37	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
15	c2yunA_	Alignment		99.8	29	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
16	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)
17	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
18	c2yupA_	Alignment		99.7	27	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
19	c2oi3A_	Alignment		99.7	29	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck; PDBTitle: nmr structure analysis of the hematopoetic cell kinase sh32 domain complexed with an artificial high affinity ligand3 (pd1)
20	c2cubA_	Alignment		99.7	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
21	c2dlpA_	Alignment	not modelled	99.7	19	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
22	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
23	c2eyyA_	Alignment	not modelled	99.7	23	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
24	c2djgA_	Alignment	not modelled	99.7	30	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
25	c2kymA_	Alignment	not modelled	99.7	19	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
26	c2i0nA_	Alignment	not modelled	99.7	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
27	c2nwmA_	Alignment	not modelled	99.7	30	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
28	c2dl3A_	Alignment	not modelled	99.7	24	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
						PDB header: cell adhesion

29	c2ebpA	Alignment	not modelled	99.7	28	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
30	d1ov3a2	Alignment	not modelled	99.7	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
31	d1ue9a	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
32	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
33	c2dbmA	Alignment	not modelled	99.7	30	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
34	c2csqA	Alignment	not modelled	99.7	22	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
35	c2dbkA	Alignment	not modelled	99.7	18	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
36	c2df6A	Alignment	not modelled	99.7	38	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
37	c2xmFA	Alignment	not modelled	99.7	32	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
38	c2ed1A	Alignment	not modelled	99.7	33	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
39	c2dl4A	Alignment	not modelled	99.7	30	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
40	c2jxbA	Alignment	not modelled	99.7	18	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
41	c2bz8B	Alignment	not modelled	99.7	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
42	c2eyxA	Alignment	not modelled	99.7	30	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
43	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)
44	c2d1xD	Alignment	not modelled	99.7	31	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
45	c2dmoA	Alignment	not modelled	99.7	21	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)
46	c2pqhA	Alignment	not modelled	99.7	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
47	d1uffa	Alignment	not modelled	99.7	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
48	c1w70A	Alignment	not modelled	99.7	42	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
49	c2dnuA	Alignment	not modelled	99.7	21	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
50	d1gria2	Alignment	not modelled	99.7	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
51	d1utia	Alignment	not modelled	99.7	45	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
52	c1wi7A	Alignment	not modelled	99.7	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
53	c1x69A	Alignment	not modelled	99.7	27	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
54	c3qwxX	Alignment	not modelled	99.7	27	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2;

						PDBTitle: ced-2 1-174
55	c1wx6A	Alignment	not modelled	99.7	28	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
56	c3qwyA	Alignment	not modelled	99.7	33	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
57	d1wlpb2	Alignment	not modelled	99.7	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
58	c2e5kA	Alignment	not modelled	99.7	20	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
59	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
60	c2yt6A	Alignment	not modelled	99.7	29	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
61	c2l0aA	Alignment	not modelled	99.7	35	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
62	c1x6gA	Alignment	not modelled	99.7	23	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.
63	d1gcqa	Alignment	not modelled	99.7	39	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
64	d1k4us	Alignment	not modelled	99.7	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
65	d1pwta	Alignment	not modelled	99.7	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	c1x2pA	Alignment	not modelled	99.7	31	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	c2dvjA	Alignment	not modelled	99.7	22	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii
68	c2dila	Alignment	not modelled	99.7	23	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
69	c2drmB	Alignment	not modelled	99.7	43	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
70	c1zukA	Alignment	not modelled	99.7	29	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
71	c2dl7A	Alignment	not modelled	99.7	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
72	d1uuea	Alignment	not modelled	99.7	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
73	c2egaA	Alignment	not modelled	99.7	19	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
74	d1nega	Alignment	not modelled	99.7	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
75	c1negA	Alignment	not modelled	99.7	32	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
76	d1ujya	Alignment	not modelled	99.7	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
77	d1loota	Alignment	not modelled	99.7	51	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
78	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
79	c1x6bA	Alignment	not modelled	99.7	22	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
80	dlsema_	Alignment	not modelled	99.7	42	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
81	dlwiea_	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
82	dluj0a_	Alignment	not modelled	99.7	41	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
83	dl oe ba_	Alignment	not modelled	99.7	43	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	clwyxA_	Alignment	not modelled	99.7	31	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	clz9qA_	Alignment	not modelled	99.7	39	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: solution structure of sh3 domain of p40phox
86	dlu06a1	Alignment	not modelled	99.7	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
87	dle6ga_	Alignment	not modelled	99.7	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
88	dlphta_	Alignment	not modelled	99.7	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
89	dlj3ta_	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
90	c2ct3A_	Alignment	not modelled	99.7	22	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
91	c2cudA_	Alignment	not modelled	99.7	24	PDB header: signaling protein Chain: A: PDB Molecule: src-like-adaptor; PDBTitle: solution structure of the sh3 domain of the human src-like2 adopter protein (slap)
92	c2o2oA_	Alignment	not modelled	99.7	26	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
93	dlug1a_	Alignment	not modelled	99.7	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	dluhca_	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
95	c2krnA_	Alignment	not modelled	99.7	37	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
96	c2fpeB_	Alignment	not modelled	99.7	17	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
97	clzx6A_	Alignment	not modelled	99.7	42	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
98	c4esrB_	Alignment	not modelled	99.7	38	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
99	dluhfa_	Alignment	not modelled	99.7	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	c2js0A_	Alignment	not modelled	99.7	34	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
101	c3rnjA_	Alignment	not modelled	99.7	29	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)
102	c2lcsA_	Alignment	not modelled	99.7	20	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
103	dlugva_	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
104	c2bzxA_	Alignment	not modelled	99.7	29	PDB header: sh3 domain Chain: A: PDB Molecule: crk-like protein; PDBTitle: atomic model of crkl-sh3c monomer
105	dlspka_	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
						Fold: SH3-like barrel

106	d2v1ra1	Alignment	not modelled	99.7	27	Superfamily: SH3-domain Family: SH3-domain
107	d1jqga	Alignment	not modelled	99.7	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	c2rqvA	Alignment	not modelled	99.7	21	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
109	c2kbtA	Alignment	not modelled	99.7	34	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
110	c2eqiA	Alignment	not modelled	99.7	25	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
111	c2ke9A	Alignment	not modelled	99.7	34	PDB header: protein binding Chain: A: PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
112	d1e6ha	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	d1gbra	Alignment	not modelled	99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	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
115	d1h8ka	Alignment	not modelled	99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	d1i1ja	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	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
118	c2k2mA	Alignment	not modelled	99.6	24	PDB header: signaling protein Chain: A: PDB Molecule: eps8-like protein 1; PDBTitle: structural basis of pxxdy motif recognition in sh3 binding
119	d1ycsb2	Alignment	not modelled	99.6	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	d1jo8a	Alignment	not modelled	99.6	44	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain