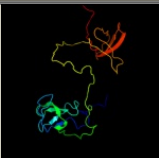
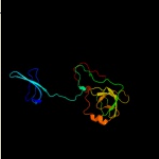
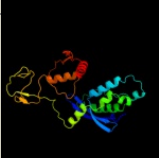
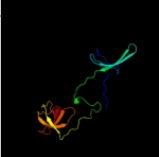
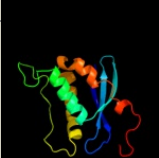
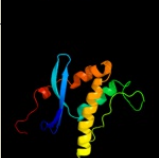
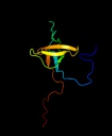
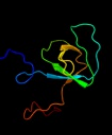
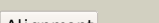
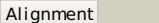
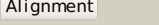
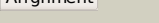
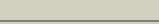
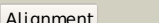
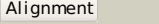
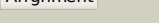
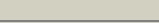
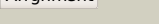
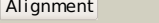
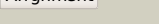
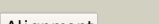
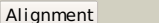
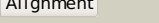


Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2lqnA_	 Alignment		100.0	18	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
2	c2l3sA_	 Alignment		100.0	20	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
3	c1griA_	 Alignment		100.0	22	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
4	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
5	c1ng2A_	 Alignment		99.9	20	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
6	c2dybA_	 Alignment		99.9	18	PDB header: oxidoreductase Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: the crystal structure of human p40(phox)
7	c1ov3A_	 Alignment		99.9	23	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
8	c2czoA_	 Alignment		99.9	38	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of the px domain of bem1p
9	c2l73A_	 Alignment		99.9	20	PDB header: oxidoreductase regulator Chain: A: PDB Molecule: nadph oxidase organizer 1; PDBTitle: structure of the noxo1b px domain
10	d1h6ha_	 Alignment		99.8	22	Fold: PX domain Superfamily: PX domain Family: PX domain
11	d1kq6a_	 Alignment		99.8	23	Fold: PX domain Superfamily: PX domain Family: PX domain

12	c2kymA_	Alignment		99.8	41	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
13	c2rqvA_	Alignment		99.8	42	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
14	d1o7ka_	Alignment		99.8	23	Fold: PX domain Superfamily: PX domain Family: PX domain
15	c1mv3A_	Alignment		99.8	25	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
16	c2csqA_	Alignment		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
17	c2rqrA_	Alignment		99.7	16	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
18	d1wiea_	Alignment		99.7	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
19	c2e5kA_	Alignment		99.7	13	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
20	c1x6gA_	Alignment		99.7	22	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.
21	c3rnjA_	Alignment	not modelled	99.7	22	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)
22	c2yt6A_	Alignment	not modelled	99.6	26	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
23	c3qwxX_	Alignment	not modelled	99.6	21	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
24	d1udla_	Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
25	c2dx1A_	Alignment	not modelled	99.6	23	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
26	d1ng2a2	Alignment	not modelled	99.6	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
27	c2jmcA_	Alignment	not modelled	99.6	23	PDB header: signaling protein Chain: A: PDB Molecule: spectrin alpha chain, brain and p41 peptide PDBTitle: chimera between spc-sh3 and p41
28	d1uffa_	Alignment	not modelled	99.6	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

29	c2cubA	Alignment	not modelled	99.6	17	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
30	c2yupA	Alignment	not modelled	99.6	23	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
31	c2oi3A	Alignment	not modelled	99.6	21	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)
32	c2cudA	Alignment	not modelled	99.6	16	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)
33	c2eyyA	Alignment	not modelled	99.6	20	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
34	c2wweA	Alignment	not modelled	99.6	16	PDB header: transferase Chain: A: PDB Molecule: phosphoinositide-3-kinase, class 2, gamma PDBTitle: crystal structure of the phox homology domain of human2 phosphoinositide-3-kinase-c2-gamma
35	c2yunA	Alignment	not modelled	99.6	21	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
36	c2dlpA	Alignment	not modelled	99.6	19	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
37	c3qwyA	Alignment	not modelled	99.6	21	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
38	d1spka	Alignment	not modelled	99.6	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
39	c1x2qA	Alignment	not modelled	99.6	20	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
40	d1phta	Alignment	not modelled	99.6	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
41	c1wx6A	Alignment	not modelled	99.6	22	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
42	c2pqhA	Alignment	not modelled	99.6	27	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
43	c2reaA	Alignment	not modelled	99.6	22	PDB header: transferase Chain: A: PDB Molecule: phosphatidylinositol-4-phosphate 3-kinase c2 PDBTitle: crystal structures of c2alpha-pi3 kinase px-domain domain2 indicate conformational change associated with ligand3 binding.
44	c2egeA	Alignment	not modelled	99.6	25	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
45	c2jxbA	Alignment	not modelled	99.6	23	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
46	c2ysqA	Alignment	not modelled	99.6	18	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
47	c2fpeB	Alignment	not modelled	99.6	27	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
48	c2enmA	Alignment	not modelled	99.6	24	PDB header: endocytosis Chain: A: PDB Molecule: sorting nexin-9; PDBTitle: solution structure of the sh3 domain from mouse sorting2 nexin-9
49	c2kt1A	Alignment	not modelled	99.6	24	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
50	c1x43A	Alignment	not modelled	99.6	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)
51	c3nmzD	Alignment	not modelled	99.6	19	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
52	d1u5sa1	Alignment	not modelled	99.6	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

53	c2iwlX	 Alignment	not modelled	99.6	25	PDB header: transferase Chain: X: PDB Molecule: phosphatidylinositol-4-phosphate 3-kinase c2 PDBTitle: structure of the px domain of phosphoinositide 3-kinase-2 c2alpha
54	d1ue9a	 Alignment	not modelled	99.6	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
55	c2yuoA	 Alignment	not modelled	99.6	23	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
56	c2ed0A	 Alignment	not modelled	99.6	31	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)
57	c2j06B	 Alignment	not modelled	99.6	34	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
58	d1jqga	 Alignment	not modelled	99.6	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
59	c1z9zA	 Alignment	not modelled	99.6	22	PDB header: structural protein Chain: A: PDB Molecule: cytoskeleton assembly control protein sla1; PDBTitle: crystal structure of yeast sla1 sh3 domain 3
60	d1ug1a	 Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
61	c2ct3A	 Alignment	not modelled	99.6	19	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
62	c2i0nA	 Alignment	not modelled	99.6	23	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
63	d1ugva	 Alignment	not modelled	99.6	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
64	c2cucA	 Alignment	not modelled	99.6	19	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
65	d1uuea	 Alignment	not modelled	99.6	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	c2dbmA	 Alignment	not modelled	99.6	29	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
67	c2ed1A	 Alignment	not modelled	99.6	23	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
68	c1negA	 Alignment	not modelled	99.6	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
69	d1nega	 Alignment	not modelled	99.6	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
70	d1jo8a	 Alignment	not modelled	99.6	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
71	c2epdA	 Alignment	not modelled	99.6	24	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
72	c1zukA	 Alignment	not modelled	99.6	19	PDB header: contractile protein Chain: A: PDB Molecule: myosin tail region-interacting protein mtl1; PDBTitle: yeast bbc1 sh3 domain complexed with a peptide from las17
73	d1k1za	 Alignment	not modelled	99.6	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	d1pwta	 Alignment	not modelled	99.6	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
75	d2v1ra1	 Alignment	not modelled	99.6	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
76	c1x6bA	 Alignment	not modelled	99.6	17	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
77	d1uhca	 Alignment	not modelled	99.6	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
78	d1ujya	Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain

					Family:SH3-domain
79	c2xmfa_	Alignment	not modelled	99.6	28 PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
80	c2d8hA_	Alignment	not modelled	99.6	16 PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
81	c2dl5A_	Alignment	not modelled	99.6	20 PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the first sh3 domain of human2 kiaa0769 protein
82	c2ct4A_	Alignment	not modelled	99.5	19 PDB header: signaling protein Chain: A: PDB Molecule: cdc42-interacting protein 4; PDBTitle: solution strcture of the sh3 domain of the cdc42-2 interacting protein 4
83	dlarka_	Alignment	not modelled	99.5	35 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	c1k9aB_	Alignment	not modelled	99.5	21 PDB header: transferase Chain: B: PDB Molecule: carboxyl-terminal src kinase; PDBTitle: crystal structure analysis of full-length carboxyl-terminal2 src kinase at 2.5 a resolution
85	dlzuua1	Alignment	not modelled	99.5	24 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
86	c2dvjA_	Alignment	not modelled	99.5	17 PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii
87	c2dbkA_	Alignment	not modelled	99.5	21 PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
88	c2lcsA_	Alignment	not modelled	99.5	23 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
89	c2df6A_	Alignment	not modelled	99.5	34 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
90	dlgbra_	Alignment	not modelled	99.5	29 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
91	c2eqiA_	Alignment	not modelled	99.5	23 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
92	c2nwmA_	Alignment	not modelled	99.5	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
93	c2dl3A_	Alignment	not modelled	99.5	26 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
94	c2dnuA_	Alignment	not modelled	99.5	24 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
95	c1x69A_	Alignment	not modelled	99.5	26 PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
96	dlsema_	Alignment	not modelled	99.5	26 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
97	c2ekhA_	Alignment	not modelled	99.5	13 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
98	dlgria2	Alignment	not modelled	99.5	28 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
99	dle6ga_	Alignment	not modelled	99.5	24 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	dlk9aa1	Alignment	not modelled	99.5	20 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	c2dl4A_	Alignment	not modelled	99.5	15 PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
102	dlu06a1	Alignment	not modelled	99.5	31 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	c2jteA_	Alignment	not modelled	99.5	21 PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: thir d sh3 domain of cd2ap
104	c2creA_	Alignment	not modelled	99.5	19 PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hef-like protein; PDBTitle: solution structure of rsgi ruh-036, an sh3 domain from2

					human cdna
105	c2dmoA_	Alignment	not modelled	99.5	19 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)
106	c2kbtA_	Alignment	not modelled	99.5	25 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
107	d1ycsb2	Alignment	not modelled	99.5	28 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	c1yn8E_	Alignment	not modelled	99.5	25 PDB header: unknown function Chain: E: PDB Molecule: nap1-binding protein 2; PDBTitle: sh3 domain of yeast nbp2
109	c2csiA_	Alignment	not modelled	99.5	30 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
110	c2dl8A_	Alignment	not modelled	99.5	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
111	c2ebpA_	Alignment	not modelled	99.5	14 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
112	c2l0aA_	Alignment	not modelled	99.5	21 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
113	d1gcqa_	Alignment	not modelled	99.5	27 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	d1l1ja_	Alignment	not modelled	99.5	19 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	c2dl7A_	Alignment	not modelled	99.5	29 PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
116	d1fmka1	Alignment	not modelled	99.5	24 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	d1gl5a_	Alignment	not modelled	99.5	19 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
118	c1wi7A_	Alignment	not modelled	99.5	20 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
119	d1efna_	Alignment	not modelled	99.5	29 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	c2drmb_	Alignment	not modelled	99.5	34 PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125