




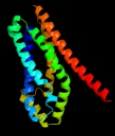

























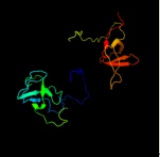
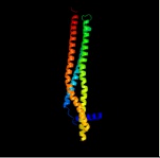
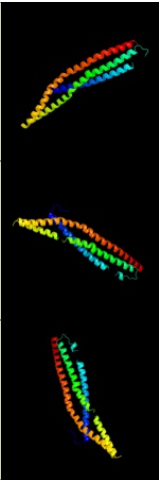
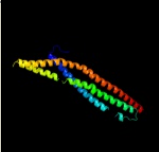
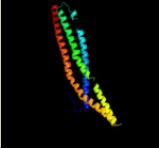



#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2dx1A_	 Alignment		100.0	26	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
2	c3ky9B_	 Alignment		100.0	28	PDB header: apoptosis Chain: B: PDB Molecule: proto-oncogene vav; PDBTitle: autoinhibited vav1
3	c2vrwB_	 Alignment		100.0	27	PDB header: signaling protein Chain: B: PDB Molecule: proto-oncogene vav; PDBTitle: critical structural role for the ph and c1 domains of the2 vav1 exchange factor
4	c3odoA_	 Alignment		100.0	23	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 1; PDBTitle: crystal structure of the dh/ph domains of p115-rhogef
5	c1xcgA_	 Alignment		100.0	24	PDB header: signaling protein activator/signaling pr Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 11; PDBTitle: crystal structure of human rhoa in complex with dh/ph2 fragment of pdzrhogef
6	c4gouA_	 Alignment		100.0	20	PDB header: signaling protein Chain: A: PDB Molecule: ehrgs-rhogef; PDBTitle: crystal structure of an rgs-rhogef from entamoeba histolytica
7	d1xcga1	 Alignment		100.0	24	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
8	c1x86A_	 Alignment		100.0	22	PDB header: signaling protein/membrane protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 12; PDBTitle: crystal structure of the dh/ph domains of leukemia-associated rhogef2 in complex with rhoa
9	c3ksyA_	 Alignment		100.0	21	PDB header: signaling protein Chain: A: PDB Molecule: son of sevenless homolog 1; PDBTitle: crystal structure of the histone domain, dh-ph unit, and2 catalytic unit of the ras activator son of sevenless (sos)
10	c3jv3A_	 Alignment		100.0	24	PDB header: protein binding Chain: A: PDB Molecule: intersectin-1; PDBTitle: structure of sh3e-dh unit of murine intersectin-1l
11	c1foeA_	 Alignment		100.0	23	PDB header: signaling protein, immune system/signali Chain: A: PDB Molecule: t-lymphoma invasion and metastasis inducing PDBTitle: crystal structure of rac1 in complex with the guanine2 nucleotide exchange region of tiam1

12	c3t06E_	Alignment		100.0	24	PDB header: signaling protein Chain: E: PDB Molecule: rho guanine nucleotide exchange factor 11; PDBTitle: crystal structure of the dh/ph fragment of pdzrhogef with n-terminal2 regulatory elements in complex with human rhoa
13	c2dfkA_	Alignment		100.0	26	PDB header: cell cycle Chain: A: PDB Molecule: collybistin ii; PDBTitle: crystal structure of the cdc42-collybistin ii complex
14	d1txda1	Alignment		100.0	21	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
15	c3bjjA_	Alignment		100.0	27	PDB header: signaling protein Chain: A: PDB Molecule: proto-oncogene vav; PDBTitle: structural basis of promiscuous guanine nucleotide exchange2 by the t-cell essential vav1
16	d2dfka1	Alignment		100.0	27	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
17	c2z0qA_	Alignment		100.0	23	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 3; PDBTitle: crystal structure of dh-ph domain of rhogef3(xpln)
18	c1xdvB_	Alignment		100.0	20	PDB header: signaling protein Chain: B: PDB Molecule: son of sevenless protein homolog 1; PDBTitle: experimentally phased structure of human the son of2 sevenless protein at 4.1 ang.
19	c4gyvA_	Alignment		100.0	23	PDB header: signaling protein Chain: A: PDB Molecule: ferm, rhogef and pleckstrin domain-containing protein 2; PDBTitle: crystal strucure of the dh domain of farp2
20	c1ki1B_	Alignment		100.0	25	PDB header: signaling protein Chain: B: PDB Molecule: intersectin long form; PDBTitle: guanine nucleotide exchange region of intersectin in2 complex with cdc42
21	d1dbha1	Alignment	not modelled	100.0	21	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
22	d1foea1	Alignment	not modelled	100.0	26	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
23	c2rgnB_	Alignment	not modelled	100.0	21	PDB header: signaling protein complex Chain: B: PDB Molecule: rhoa/rac/cdc42 exchange factor; PDBTitle: crystal structure of p63rhogef complex with galpha-q and rhoa
24	c4gzvB_	Alignment	not modelled	100.0	24	PDB header: signaling protein Chain: B: PDB Molecule: ferm, rhogef and pleckstrin domain-containing protein 2; PDBTitle: crystal strucure of the dh-ph-ph domain of farp2
25	c3eo2A_	Alignment	not modelled	100.0	26	PDB header: splicing Chain: A: PDB Molecule: neuroepithelial cell-transforming gene 1 protein; PDBTitle: crystal structure of the rhogef domain of human neuroepithelial cell-2 transforming gene 1 protein
26	c1lb1C_	Alignment	not modelled	100.0	28	PDB header: signaling protein Chain: C: PDB Molecule: guanine nucleotide exchange factor dbs; PDBTitle: crystal structure of the dbl and pleckstrin homology2 domains of dbs in complex with rhoa
27	d1ki1b1	Alignment	not modelled	100.0	23	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
28	c4h6yA_	Alignment	not modelled	100.0	18	PDB header: signaling protein Chain: A: PDB Molecule: ferm, rhogef and pleckstrin domain-containing protein 1; PDBTitle: crystal strucure of the dh-ph-ph domain of farp1
						Fold: DBL homology domain (DH-domain)

29	d1by1a_	Alignment	not modelled	100.0	23	Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
30	c3mpxA_	Alignment	not modelled	100.0	21	PDB header: lipid binding protein Chain: A: PDB Molecule: fyve, rhogef and ph domain-containing protein 5; PDBTitle: crystal structure of the dh and ph-1 domains of human fgd5
31	d1f5xa_	Alignment	not modelled	100.0	25	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
32	c3odwB_	Alignment	not modelled	100.0	23	PDB header: signaling protein Chain: B: PDB Molecule: rho guanine nucleotide exchange factor 1; PDBTitle: crystal structure of the linker-dh/ph domains of p115-rhogef
33	c1dbbA_	Alignment	not modelled	100.0	18	PDB header: gene regulation Chain: A: PDB Molecule: protein (human sos 1); PDBTitle: dbl and pleckstrin homology domains from hsos1
34	c3odxA_	Alignment	not modelled	100.0	23	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 1; PDBTitle: crystal structure of an n-terminally truncated linker-dh/ph domains of f2 p115-rhogef
35	d1kz7a1	Alignment	not modelled	100.0	28	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
36	c1ntyA_	Alignment	not modelled	100.0	22	PDB header: signaling protein Chain: A: PDB Molecule: triple functional domain protein; PDBTitle: crystal structure of the first dh/ph domain of trio to 1.7 a
37	d1ntyA1	Alignment	not modelled	100.0	22	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
38	c3gf9A_	Alignment	not modelled	100.0	28	PDB header: endocytosis Chain: A: PDB Molecule: intersectin 2; PDBTitle: crystal structure of human intersectin 2 rhogef domain
39	c2lqnA_	Alignment		99.9	17	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
40	c1griA_	Alignment		99.9	18	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
41	c2eyzA_	Alignment		99.9	21	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
42	c2l3sA_	Alignment		99.9	21	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
43	c1ng2A_	Alignment		99.9	23	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
44	c1ov3A_	Alignment		99.9	22	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
45	c1mv3A_	Alignment	not modelled	99.9	15	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
46	c4avmA_	Alignment		99.9	15	PDB header: protein binding Chain: A: PDB Molecule: bridging integrator 2; PDBTitle: crystal structure of the n-bar domain of human bridging2 integrator 2.

47	c2ficA_	Alignment		99.8	13	PDB header: endocytosis/exocytosis, membrane protein Chain: A: PDB Molecule: bridging integrator 1; PDBTitle: the crystal structure of the bar domain from human bin1/amphiphysin ii2 and its implications for molecular recognition
48	c1uruA_	Alignment		99.8	20	PDB header: endocytosis Chain: A: PDB Molecule: amphiphysin; PDBTitle: amphiphysin bar domain from drosophila
49	d1urua_	Alignment		99.8	20	Fold: BAR/IMD domain-like Superfamily: BAR/IMD domain-like Family: BAR domain
50	c2e5kA_	Alignment	not modelled	99.8	14	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
51	d1ng2a2	Alignment	not modelled	99.8	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
52	d1udla_	Alignment	not modelled	99.8	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
53	c3sogA_	Alignment		99.7	15	PDB header: structural protein Chain: A: PDB Molecule: amphiphysin; PDBTitle: crystal structure of the bar domain of human amphiphysin, isoform 1
54	c2ed0A_	Alignment	not modelled	99.7	22	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)
55	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
56	c2rqrA_	Alignment	not modelled	99.7	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
57	c2eyyA_	Alignment	not modelled	99.7	12	PDB header: signaling protein Chain: A: PDB Molecule: v-crkl sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
58	c2csqA_	Alignment	not modelled	99.7	16	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
59	d1uhca_	Alignment	not modelled	99.7	42	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
60	c2jxbA_	Alignment	not modelled	99.7	20	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
61	c2dl7A_	Alignment	not modelled	99.7	24	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
62	c2cudA_	Alignment	not modelled	99.7	14	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)
63	c3nmzD_	Alignment	not modelled	99.7	17	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
64	d1ug1a_	Alignment	not modelled	99.7	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
65	c1x6gA_	Alignment	not modelled	99.7	24	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.
66	c4glmD_	Alignment	not modelled	99.7	28	PDB header: signaling protein Chain: D: PDB Molecule: dynammin-binding protein; PDBTitle: crystal structure of the sh3 domain of dnmbp protein [homo sapiens]
67	c2oi3A_	Alignment	not modelled	99.7	28	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)
68	d1ov3a2	Alignment	not modelled	99.7	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

69	c2dnuA_	Alignment	not modelled	99.7	19	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	c2nwmA_	Alignment	not modelled	99.7	32	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
71	c2egeA_	Alignment	not modelled	99.7	22	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
72	d1ycsb2	Alignment	not modelled	99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
73	d1l1ja_	Alignment	not modelled	99.7	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	c2yt6A_	Alignment	not modelled	99.7	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
75	c2ekha_	Alignment	not modelled	99.7	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
76	c2kbtA_	Alignment	not modelled	99.7	32	PDB header: signaling protein Chain: A; PDB Molecule: chimera of proto-oncogene vav, linker, PDBTitle: attachment of an nmr-invisible solubility enhancement tag2 (inset) using a sortase-mediated protein ligation method
77	c2cubA_	Alignment	not modelled	99.7	21	PDB header: signaling protein Chain: A; PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
78	c2df6A_	Alignment	not modelled	99.7	28	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
79	c2dilA_	Alignment	not modelled	99.7	30	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
80	c2yuoa_	Alignment	not modelled	99.7	21	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
81	c1zlmA_	Alignment	not modelled	99.7	28	PDB header: signaling protein Chain: A; PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
82	c2djqa_	Alignment	not modelled	99.7	17	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
83	c1x2qa_	Alignment	not modelled	99.7	22	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
84	c2dmoA_	Alignment	not modelled	99.7	29	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)
85	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
86	c2kymA_	Alignment	not modelled	99.7	24	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
87	c2epdA_	Alignment	not modelled	99.7	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
88	d2v1ra1	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
89	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
90	d1u5sa1	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
91	c2dbkA_	Alignment	not modelled	99.7	17	PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
92	c2csiA_	Alignment	not modelled	99.7	32	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
93	c2yupA_	Alignment	not modelled	99.6	20	PDB header: cell adhesion Chain: A; PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
94	c2egaA_	Alignment	not modelled	99.6	29	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
95	d1gcqa_	Alignment	not modelled	99.6	26 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
96	d1gbra_	Alignment	not modelled	99.6	28 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
97	c1x2pA_	Alignment	not modelled	99.6	25 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
98	c3qwxX_	Alignment	not modelled	99.6	25 PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
99	c2ebpA_	Alignment	not modelled	99.6	16 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
100	c2l0aA_	Alignment	not modelled	99.6	32 PDB header: signaling protein Chain: A: PDB Molecule: signal transducing adapter molecule 1; PDBTitle: solution nmr structure of signal transducing adapter molecule 1 stam-12 from homo sapiens, northeast structural genomics consortium target3 hr4479e
101	c2ed1A_	Alignment	not modelled	99.6	20 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
102	c1x43A_	Alignment	not modelled	99.6	21 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)
103	c2xmfA_	Alignment	not modelled	99.6	25 PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
104	d1wiea_	Alignment	not modelled	99.6	25 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
105	c2yunA_	Alignment	not modelled	99.6	29 PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
106	c1wyxA_	Alignment	not modelled	99.6	36 PDB header: cell adhesion Chain: A: PDB Molecule: crk-associated substrate; PDBTitle: the crystal structure of the p130cas sh3 domain at 1.1 a2 resolution
107	c2ysqA_	Alignment	not modelled	99.6	21 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
108	d1oeba_	Alignment	not modelled	99.6	28 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
109	c1bb9A_	Alignment	not modelled	99.6	28 PDB header: transferase Chain: A: PDB Molecule: amphiphysin 2; PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
110	d1bb9a_	Alignment	not modelled	99.6	28 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
111	c3qwyA_	Alignment	not modelled	99.6	27 PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
112	c2dvjA_	Alignment	not modelled	99.6	14 PDB header: signaling protein Chain: A: PDB Molecule: iv-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii
113	d1lutia_	Alignment	not modelled	99.6	29 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	c2dl8A_	Alignment	not modelled	99.6	20 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
115	d1loota_	Alignment	not modelled	99.6	29 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	d1wlpb2	Alignment	not modelled	99.6	20 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	d1nega_	Alignment	not modelled	99.6	27 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
118	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
119	d1uuea_	Alignment	not modelled	99.6	29 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	d1u06a1	Alignment	not modelled	99.6	29 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

