


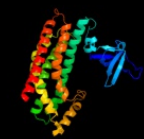

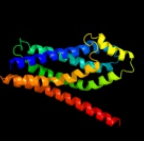













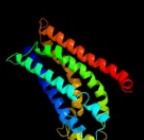




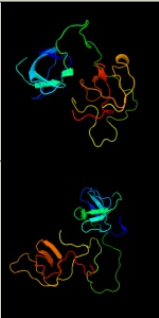


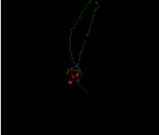

Phyre2

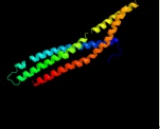
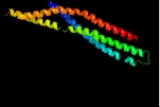






Email	l.a.kelley@imperial.ac.uk
Description	Q6TXD4
Date	Tue Jul 30 13:06:16 BST 2013
Unique Job ID	3e4e7cfa0db1a39a

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2dx1A_	 Alignment		100.0	41	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein a5ef
2	c3jv3A_	 Alignment		100.0	38	PDB header: protein binding Chain: A: PDB Molecule: intersectin-1; PDBTitle: structure of sh3e-dh unit of murine intersectin-1l
3	c3ky9B_	 Alignment		100.0	24	PDB header: apoptosis Chain: B: PDB Molecule: proto-oncogene vav; PDBTitle: autoinhibited vav1
4	c4gouA_	 Alignment		100.0	15	PDB header: signaling protein Chain: A: PDB Molecule: ehrgs-rhogef; PDBTitle: crystal structure of an rgs-rhogef from entamoeba histolytica
5	c3ksyA_	 Alignment		100.0	20	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)
6	c2vrwB_	 Alignment		100.0	25	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
7	c3odoA_	 Alignment		100.0	21	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
8	c1xcgA_	 Alignment		100.0	19	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
9	c1x86A_	 Alignment		100.0	18	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
10	c1foeA_	 Alignment		100.0	27	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
11	c1griA_	 Alignment		100.0	24	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2

12	c1xdvB_	Alignment		100.0	19	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.
13	c3t06E_	Alignment		100.0	19	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
14	d1xcqa1	Alignment		100.0	19	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
15	c2dfkA_	Alignment		100.0	29	PDB header: cell cycle Chain: A: PDB Molecule: collybistin ii; PDBTitle: crystal structure of the cdc42-collybistin ii complex
16	c3bjjA_	Alignment		100.0	24	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
17	c1ki1B_	Alignment		100.0	24	PDB header: signaling protein Chain: B: PDB Molecule: intersectin long form; PDBTitle: guanine nucleotide exchange region of intersectin in2 complex with cdc42
18	d1txda1	Alignment		100.0	17	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
19	d1dbha1	Alignment		100.0	19	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
20	c2z0qA_	Alignment		100.0	27	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 3; PDBTitle: crystal structure of dh-ph domain of rhogef3(xpln)
21	d2dfka1	Alignment	not modelled	100.0	30	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
22	c30dwB_	Alignment	not modelled	100.0	20	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
23	c2lqnA_	Alignment		100.0	21	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
24	c30dxA_	Alignment	not modelled	99.9	20	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 of2 p115-rhogef
25	d1ki1b1	Alignment	not modelled	99.9	23	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
26	c1dbhA_	Alignment	not modelled	99.9	18	PDB header: gene regulation Chain: A: PDB Molecule: protein (human sos 1); PDBTitle: dbl and pleckstrin homology domains from hsos1
27	c4gyvA_	Alignment	not modelled	99.9	26	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
						Fold: DBL homology domain (DH-domain)

28	d1foea1	Alignment	not modelled	99.9	29	Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
29	c1lb1C	Alignment	not modelled	99.9	27	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
30	c4gzub	Alignment	not modelled	99.9	28	PDB header: signaling protein Chain: B: PDB Molecule: ferm, rhogef and pleckstrin domain-containing protein 2; PDBTitle: crystal structure of the dh-ph-ph domain of farp2
31	c2rgnB	Alignment	not modelled	99.9	27	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
32	c3eo2A	Alignment	not modelled	99.9	27	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
33	c4h6yA	Alignment	not modelled	99.9	22	PDB header: signaling protein Chain: A: PDB Molecule: ferm, rhogef and pleckstrin domain-containing protein 1; PDBTitle: crystal structure of the dh-ph-ph domain of farp1
34	d1f5xa	Alignment	not modelled	99.9	26	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
35	c3mpxA	Alignment	not modelled	99.9	14	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
36	c2l3sa	Alignment		99.9	26	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
37	c2eyzA	Alignment		99.9	24	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
38	c1ng2A	Alignment	not modelled	99.9	21	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
39	d1by1a	Alignment	not modelled	99.9	23	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
40	d1kz7a1	Alignment	not modelled	99.9	27	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
41	c3gf9A	Alignment	not modelled	99.9	28	PDB header: endocytosis Chain: A: PDB Molecule: intersectin 2; PDBTitle: crystal structure of human intersectin 2 rhogef domain
42	c1ntyA	Alignment	not modelled	99.9	24	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
43	d1nty1	Alignment	not modelled	99.9	24	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
44	c1ov3A	Alignment		99.9	27	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
45	c1mv3A	Alignment		99.8	10	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		99.8	23	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	21	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	d1urua_	Alignment		99.7	23	Fold: BAR/IMD domain-like Superfamily: BAR/IMD domain-like Family: BAR domain
49	c1uruA_	Alignment		99.7	23	PDB header: endocytosis Chain: A: PDB Molecule: amphiphysin; PDBTitle: amphiphysin bar domain from drosophila
50	c3sogA_	Alignment		99.7	23	PDB header: structural protein Chain: A: PDB Molecule: amphiphysin; PDBTitle: crystal structure of the bar domain of human amphiphysin, isoform 1
51	c2rqra_	Alignment		99.7	17	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
52	c2e5kA_	Alignment		99.6	18	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
53	d1ng2a2	Alignment		99.6	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
54	c1bb9A_	Alignment		99.6	17	PDB header: transferase Chain: A: PDB Molecule: amphiphysin 2; PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
55	d1bb9a_	Alignment		99.6	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
56	d1uhca_	Alignment	not modelled	99.6	83	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
57	d1udla_	Alignment	not modelled	99.6	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
58	c2jxbA_	Alignment	not modelled	99.6	21	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
59	c2dlpA_	Alignment	not modelled	99.6	20	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
60	c2dybA_	Alignment	not modelled	99.6	21	PDB header: oxidoreductase Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: the crystal structure of human p40(phox)
61	d1ug1a_	Alignment	not modelled	99.6	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
62	c3nmzD_	Alignment	not modelled	99.6	15	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
63	d1ov3a2	Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
64	d1i1ja_	Alignment	not modelled	99.6	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
65	d1i4da_	Alignment	not modelled	99.6	13	Fold: BAR/IMD domain-like Superfamily: BAR/IMD domain-like Family: Arfaptin, Rac-binding fragment
66	c2eyyA_	Alignment	not modelled	99.5	20	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog

						PDBTitle: ct10-regulated kinase isoform i
67	c2dl7A_	Alignment	not modelled	99.5	27	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
68	c2ed0A_	Alignment	not modelled	99.5	20	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)
69	c1zlmA_	Alignment	not modelled	99.5	37	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
70	d1wlpb2	Alignment	not modelled	99.5	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
71	c2dmoA_	Alignment	not modelled	99.5	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)
72	c1x2qA_	Alignment	not modelled	99.5	21	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
73	c2d1xD_	Alignment	not modelled	99.5	24	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
74	c1x43A_	Alignment	not modelled	99.5	25	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)
75	c2cubA_	Alignment	not modelled	99.5	23	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
76	c2ysqA_	Alignment	not modelled	99.5	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
77	c2df6A_	Alignment	not modelled	99.5	40	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
78	d1k4us_	Alignment	not modelled	99.5	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
79	d1jqqa_	Alignment	not modelled	99.5	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
80	c2ebpA_	Alignment	not modelled	99.5	13	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
81	c2egeA_	Alignment	not modelled	99.5	16	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
82	c2djqa_	Alignment	not modelled	99.5	33	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: sh3 domain containing ring finger 2; PDBTitle: the solution structure of the first sh3 domain of mouse sh32 domain containing ring finger 2
83	d1uuea_	Alignment	not modelled	99.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	c2yunA_	Alignment	not modelled	99.5	24	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
85	d2v1ra1	Alignment	not modelled	99.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
86	d1oeba_	Alignment	not modelled	99.5	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
87	c2epdA_	Alignment	not modelled	99.5	18	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	c1x6gA_	Alignment	not modelled	99.5	14	PDB header: signaling protein Chain: A: PDB Molecule: megakaryocyte-associated tyrosine-protein kinase; PDBTitle: solution structures of the sh3 domain of human2 megakaryocyte-associated tyrosine-protein kinase.
89	c2yt6A_	Alignment	not modelled	99.5	27	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
90	c2dilA_	Alignment	not modelled	99.5	27	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
91	c4glmD_	Alignment	not modelled	99.5	27	PDB header: signaling protein Chain: D: PDB Molecule: dynammin-binding protein; PDBTitle: crystal structure of the sh3 domain of dnmbp protein [homo sapiens]
92	c2numA_	Alignment	not modelled	99.5	22	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin;

92	c2iwmA	Alignment	not modelled	99.5	30	PDBTitle: solution structure of the first sh3 domain of human vinxin2 and its interaction with the peptides from vinculin PDB header: signaling protein Chain: A; PDB Molecule: 130 kda phosphatidylinositol 4,5-biphosphate-
93	c2ed1A	Alignment	not modelled	99.5	31	PDBTitle: solution structure of the sh3 domain of 130 kda2 phosphatidylinositol 4,5-biphosphate-dependent arf1 gtpase-3 activating protein PDB header: transferase Chain: A; PDB Molecule: tyrosine-protein kinase hck;
94	c2oi3A	Alignment	not modelled	99.5	18	PDBTitle: nmr structure analysis of the hematopoietic cell kinase sh32 domain complexed with an artificial high affinity ligand3 (pd1)
95	c2dl4A	Alignment	not modelled	99.5	34	PDB header: signaling protein Chain: A; PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
96	c2dl3A	Alignment	not modelled	99.5	33	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
97	c1wx6A	Alignment	not modelled	99.5	21	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
98	c2pqhA	Alignment	not modelled	99.5	21	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
99	c2kbtA	Alignment	not modelled	99.5	36	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
100	d1utia	Alignment	not modelled	99.5	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	c2jmcA	Alignment	not modelled	99.5	19	PDB header: signaling protein Chain: A; PDB Molecule: spectrin alpha chain, brain and p41 peptide PDBTitle: chimera between spc-sh3 and p41
102	c1wyxA	Alignment	not modelled	99.5	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
103	c2kymA	Alignment	not modelled	99.5	20	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
104	c3qwyA	Alignment	not modelled	99.5	28	PDB header: signaling protein Chain: A; PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
105	d1ujya	Alignment	not modelled	99.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
106	d1ycsb2	Alignment	not modelled	99.5	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
107	c2xmfA	Alignment	not modelled	99.5	19	PDB header: motor protein Chain: A; PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
108	c2dbmA	Alignment	not modelled	99.5	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
109	c2bz8B	Alignment	not modelled	99.5	24	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
110	d1uj0a	Alignment	not modelled	99.5	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
111	d1pwta	Alignment	not modelled	99.5	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	c2dbkA	Alignment	not modelled	99.5	15	PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
113	c3qwxX	Alignment	not modelled	99.5	22	PDB header: signaling protein Chain: X; PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
114	c2yuoA	Alignment	not modelled	99.5	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
115	c2eqiA	Alignment	not modelled	99.5	29	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
116	c2drmB	Alignment	not modelled	99.5	28	PDB header: contractile protein Chain: B; PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
117	c1negA	Alignment	not modelled	99.5	20	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 Fold: SH3-like barrel

118	d1nega_	Alignment	not modelled	99.5	20	Superfamily: SH3-domain Family: SH3-domain
119	d1u06a1	Alignment	not modelled	99.5	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	c2csqA_	Alignment	not modelled	99.5	25	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