






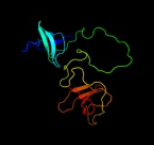













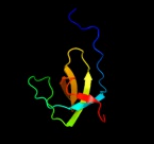




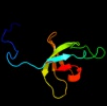





Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	Q18204
Date	Sat Apr 13 12:59:27 BST 2013
Unique Job ID	8d9382dba279366a

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1griA_	 Alignment		100.0	17	PDB header: signal transduction adaptor Chain: A; PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
2	c2lqnA_	 Alignment		100.0	20	PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
3	c2l3sA_	 Alignment		100.0	16	PDB header: structural protein Chain: A; PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
4	c2eyzA_	 Alignment		100.0	14	PDB header: signaling protein Chain: A; PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
5	c1ng2A_	 Alignment		100.0	13	PDB header: oxidoreductase activator Chain: A; PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
6	c1ov3A_	 Alignment		100.0	13	PDB header: oxidoreductase activator Chain: A; PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
7	c1mv3A_	 Alignment		99.9	11	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
8	d1udla_	 Alignment		99.8	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
9	c2rqrA_	 Alignment		99.8	12	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
10	c2oi3A_	 Alignment		99.8	18	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)
11	d1ng2a2	 Alignment		99.8	11	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

12	c2yt6A_	Alignment		99.8	20	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
13	c2yupA_	Alignment		99.8	19	PDB header: cell adhesion Chain: A; PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
14	c2kymA_	Alignment		99.8	19	PDB header: signaling protein Chain: A; PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of the bem1p sh3-ci domain from I.elongisporus in2 complex with ste20p peptide
15	c3nmzD_	Alignment		99.8	16	PDB header: cell adhesion/cell cycle Chain: D; PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
16	c1x43A_	Alignment		99.8	15	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)
17	c2cudA_	Alignment		99.8	15	PDB header: signaling protein Chain: A; PDB Molecule: src-like-adapter; PDBTitle: solution structure of the sh3 domain of the human src-like2 adppter protein (slap)
18	c1x6gA_	Alignment		99.8	9	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.
19	c2fpeB_	Alignment		99.8	16	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
20	c2d8hA_	Alignment		99.8	16	PDB header: unknown function Chain: A; PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
21	d1wiea_	Alignment	not modelled	99.8	15	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
22	c2lcsA_	Alignment	not modelled	99.8	17	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
23	d1k9aa1	Alignment	not modelled	99.8	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
24	c2dybA_	Alignment	not modelled	99.8	14	PDB header: oxidoreductase Chain: A; PDB Molecule: neutrophil cytosol factor 4; PDBTitle: the crystal structure of human p40(phox)
25	c2ct4A_	Alignment	not modelled	99.8	16	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
26	c1x6bA_	Alignment	not modelled	99.8	13	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
27	c1wx6A_	Alignment	not modelled	99.8	20	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
28	c1x2qA_	Alignment	not modelled	99.8	16	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
						Fold: SH3-like barrel

29	d1ue9a_	Alignment	not modelled	99.8	18	Superfamily: SH3-domain Family: SH3-domain
30	c2enmA_	Alignment	not modelled	99.8	21	PDB header: endocytosis Chain: A: PDB Molecule: sorting nexin-9; PDBTitle: solution structure of the sh3 domain from mouse sorting2 nexin-9
31	d1spka_	Alignment	not modelled	99.8	15	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
32	c2dlpA_	Alignment	not modelled	99.7	16	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
33	c2dx1A_	Alignment	not modelled	99.7	16	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
34	c2ct3A_	Alignment	not modelled	99.7	16	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
35	c2jteA_	Alignment	not modelled	99.7	21	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: third sh3 domain of cd2ap
36	c2e5kA_	Alignment	not modelled	99.7	16	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
37	c2dl5A_	Alignment	not modelled	99.7	17	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the first sh3 domain of human2 kiaa0769 protein
38	c2j06B_	Alignment	not modelled	99.7	16	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
39	d1uffa_	Alignment	not modelled	99.7	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
40	c1yn8E_	Alignment	not modelled	99.7	17	PDB header: unknown function Chain: E: PDB Molecule: nap1-binding protein 2; PDBTitle: sh3 domain of yeast nbp2
41	c3rnjA_	Alignment	not modelled	99.7	13	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)
42	c1zukA_	Alignment	not modelled	99.7	17	PDB header: contractile protein Chain: A: PDB Molecule: myosin tail region-interacting protein mt1; PDBTitle: yeast bbc1 sh3 domain complexed with a peptide from las17
43	c2ed0A_	Alignment	not modelled	99.7	16	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)
44	c4afzC_	Alignment	not modelled	99.7	18	PDB header: hydrolase/de novo protein Chain: C: PDB Molecule: fynomer; PDBTitle: human chymase - fynomer complex
45	d1phta_	Alignment	not modelled	99.7	14	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
46	c2kt1A_	Alignment	not modelled	99.7	22	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
47	c2eczA_	Alignment	not modelled	99.7	17	PDB header: signaling protein Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: solution structure of the sh3 domain of sorbin and sh32 domain-containing protein 1
48	c2jxbA_	Alignment	not modelled	99.7	13	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
49	d1jo8a_	Alignment	not modelled	99.7	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
50	c2yunA_	Alignment	not modelled	99.7	14	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
51	c2cubA_	Alignment	not modelled	99.7	15	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
52	c2m0yA_	Alignment	not modelled	99.7	11	PDB header: apoptosis Chain: A: PDB Molecule: dedicator of cytokinesis protein 1; PDBTitle: solution structure of the sh3 domain of dock180
53	c2cucA_	Alignment	not modelled	99.7	22	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
54	d1arka_	Alignment	not modelled	99.7	10	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
						PDB header: signaling protein

55	c2ysqA_	Alignment	not modelled	99.7	21	Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 9; PDBTitle: solution structure of the sh3 domain from rho guanine2 nucleotide exchange factor 9
56	c2lj0A_	Alignment	not modelled	99.7	13	PDB header: signaling protein Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: the third sh3 domain of r85fl
57	c2ed1A_	Alignment	not modelled	99.7	19	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
58	c3jv3A_	Alignment	not modelled	99.7	20	PDB header: protein binding Chain: A: PDB Molecule: intersectin-1; PDBTitle: structure of sh3e-dh unit of murine intersectin-1l
59	c2yu0A_	Alignment	not modelled	99.7	13	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
60	d1ov3a2	Alignment	not modelled	99.7	12	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
61	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
62	c2i0nA_	Alignment	not modelled	99.7	11	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	c1zuyB_	Alignment	not modelled	99.7	20	PDB header: contractile protein Chain: B: PDB Molecule: myosin-5 isoform; PDBTitle: high-resolution structure of yeast myo5 sh3 domain
64	c2ekhA_	Alignment	not modelled	99.7	9	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
65	c2dmoA_	Alignment	not modelled	99.7	18	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)
66	c2djgA_	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
67	d1fmka1	Alignment	not modelled	99.7	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
68	c2epdA_	Alignment	not modelled	99.7	21	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
69	c2rqvA_	Alignment	not modelled	99.7	19	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
70	c1x69A_	Alignment	not modelled	99.7	14	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
71	c2dbmA_	Alignment	not modelled	99.7	17	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
72	d1cska_	Alignment	not modelled	99.7	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
73	c2azsA_	Alignment	not modelled	99.7	19	PDB header: signaling protein Chain: A: PDB Molecule: sh2-sh3 adapter protein drk; PDBTitle: nmr structure of the n-terminal sh3 domain of drk2 (calculated without noe restraints)
74	c3qwyA_	Alignment	not modelled	99.7	14	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
75	c2dl4A_	Alignment	not modelled	99.7	22	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
76	c2df6A_	Alignment	not modelled	99.7	16	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
77	c1z9zA_	Alignment	not modelled	99.7	21	PDB header: structural protein Chain: A: PDB Molecule: cytoskeleton assembly control protein sla1; PDBTitle: crystal structure of yeast sla1 sh3 domain 3
78	c2egeA_	Alignment	not modelled	99.7	11	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
79	c2eyxA_	Alignment	not modelled	99.7	19	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
80	c2ebpA_	Alignment	not modelled	99.7	15	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	c3qwxX	Alignment	not modelled	99.7	19	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
82	c2lx7A	Alignment	not modelled	99.7	26	PDB header: protein binding Chain: A: PDB Molecule: growth arrest-specific protein 7; PDBTitle: solution nmr structure of sh3 domain of growth arrest-specific protein2 7 (gas7) (fragment 1-60) from homo sapiens, northeast structural3 genomics consortium (nesg) target hr8574a
83	c2eqiA	Alignment	not modelled	99.7	13	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
84	c2nwmA	Alignment	not modelled	99.7	20	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
85	c4glmD	Alignment	not modelled	99.7	22	PDB header: signaling protein Chain: D: PDB Molecule: dynammin-binding protein; PDBTitle: crystal structure of the sh3 domain of dnmbp protein [homo sapiens]
86	c2dl8A	Alignment	not modelled	99.7	18	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
87	c4esrB	Alignment	not modelled	99.7	19	PDB header: protein binding Chain: B: PDB Molecule: jouberein; 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
88	d1k4us	Alignment	not modelled	99.7	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
89	d1uj0a	Alignment	not modelled	99.7	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
90	c2l0aA	Alignment	not modelled	99.7	18	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
91	d2iima1	Alignment	not modelled	99.7	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
92	c2dl3A	Alignment	not modelled	99.7	11	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
93	c2xmfA	Alignment	not modelled	99.7	15	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
94	d1jqqa	Alignment	not modelled	99.7	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
95	c2d1xD	Alignment	not modelled	99.7	16	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
96	c2pqhA	Alignment	not modelled	99.7	22	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
97	c1s1nA	Alignment	not modelled	99.7	23	PDB header: cell adhesion Chain: A: PDB Molecule: nephrocystin 1; PDBTitle: sh3 domain of human nephrocystin
98	c2bz8B	Alignment	not modelled	99.7	20	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
99	d1gria2	Alignment	not modelled	99.7	15	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	d1qcfa1	Alignment	not modelled	99.7	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	c2eyyA	Alignment	not modelled	99.7	20	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
102	d1wlpb2	Alignment	not modelled	99.7	11	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	c2dm1A	Alignment	not modelled	99.7	17	PDB header: signaling protein Chain: A: PDB Molecule: protein vav-2; PDBTitle: solution structure of the second sh3 domain of human2 protein vav-2
104	c2creA	Alignment	not modelled	99.7	9	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	d1ugva	Alignment	not modelled	99.7	10	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
106	c3cqtA	Alignment	not modelled	99.7	20	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase fyn;

				PDBTitle: n53i v55l mutant of fyn sh3 domain	
107	d1utia_	Alignment	not modelled	99.7	18 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	d1zuua1	Alignment	not modelled	99.7	16 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
109	d1pwta_	Alignment	not modelled	99.7	22 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
110	d1y57a1	Alignment	not modelled	99.7	22 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
111	c2dila_	Alignment	not modelled	99.7	14 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
112	c1w70A_	Alignment	not modelled	99.7	25 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
113	c1wi7A_	Alignment	not modelled	99.7	16 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
114	c2egaA_	Alignment	not modelled	99.7	17 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
115	d1uhca_	Alignment	not modelled	99.7	17 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	c1x27F_	Alignment	not modelled	99.7	22 PDB header: signaling protein Chain: F; PDB Molecule: proto-oncogene tyrosine-protein kinase lck; PDBTitle: crystal structure of lck sh2-sh3 with sh2 binding site of2 p130cas
117	d1uuea_	Alignment	not modelled	99.7	24 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
118	d1u5sa1	Alignment	not modelled	99.7	22 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	c2k6dA_	Alignment	not modelled	99.7	20 PDB header: sh3 domain/ubiquitin Chain: A; PDB Molecule: sh3 domain-containing kinase-binding protein 1; PDBTitle: cin85 sh3-c domain in complex with ubiquitin
120	c2dnuA_	Alignment	not modelled	99.7	17 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