


















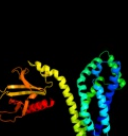













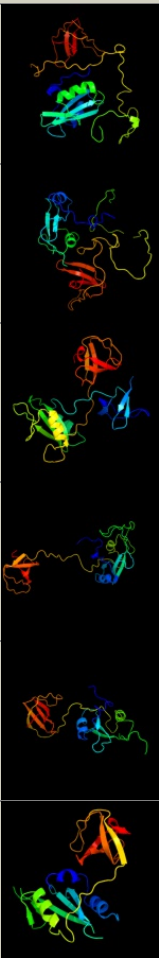
Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	P52735
Date	Wed Jun 6 09:50:28 BST 2012
Unique Job ID	3adc006cc7b0d8b4

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3ky9B_	 Alignment		100.0	56	PDB header: apoptosis Chain: B; PDB Molecule: proto-oncogene vav; PDBTitle: autoinhibited vav1
2	c2vrwB_	 Alignment		100.0	54	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
3	c3bjjA_	 Alignment		100.0	54	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
4	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)
5	c2dx1A_	 Alignment		100.0	27	PDB header: signaling protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
6	c3odoA_	 Alignment		100.0	19	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
7	c1xdvB_	 Alignment		100.0	21	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.
8	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
9	c1xcgA_	 Alignment		100.0	21	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
10	c2dfkA_	 Alignment		100.0	24	PDB header: cell cycle Chain: A; PDB Molecule: collybistin ii; PDBTitle: crystal structure of the cdc42-collybistin ii complex
11	c3t06E_	 Alignment		100.0	20	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

12	c3odxA	Alignment		100.0	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 of p115-rhogef
13	c3odwB	Alignment		100.0	22	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
14	c2rgnB	Alignment		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
15	c1foeA	Alignment		100.0	19	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
16	c3mpxA	Alignment		100.0	16	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
17	c2z0qA	Alignment		100.0	22	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	c1ki1B	Alignment		100.0	20	PDB header: signaling protein Chain: B: PDB Molecule: intersectin long form; PDBTitle: guanine nucleotide exchange region of intersectin in2 complex with cdc42
19	c1dbhA	Alignment		100.0	20	PDB header: gene regulation Chain: A: PDB Molecule: protein (human sos 1); PDBTitle: dbl and pleckstrin homology domains from hso1
20	c1lb1C	Alignment		100.0	22	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
21	c1ntyA	Alignment	not modelled	100.0	17	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
22	d1xcga1	Alignment	not modelled	100.0	22	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
23	d1bda1	Alignment	not modelled	100.0	19	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
24	d1dbha1	Alignment	not modelled	100.0	20	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
25	d1f5xa	Alignment	not modelled	100.0	49	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
26	d1ki1b1	Alignment	not modelled	100.0	25	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
27	d2dfka1	Alignment	not modelled	100.0	27	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
28	c2d86A	Alignment	not modelled	100.0	71	PDB header: signaling protein, protein binding Chain: A: PDB Molecule: vav-3 protein; PDBTitle: solution structure of the ch domain from human vav-3 protein
29	c3eo2A	Alignment	not modelled	100.0	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	
30	d1kz7a1	Alignment	not modelled	100.0	26 Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)	
31	d1foea1	Alignment	not modelled	100.0	28 Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)	
32	d1ntya1	Alignment	not modelled	100.0	21 Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)	
33	c3gf9A_	Alignment	not modelled	100.0	24 PDB header: endocytosis Chain: A; PDB Molecule: intersectin 2; PDBTitle: crystal structure of human intersectin 2 rhogef domain	
34	d1by1a	Alignment	not modelled	100.0	27 Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)	
35	c21qnA_	Alignment		100.0	25 PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl	
36	c2eyzA_	Alignment		100.0	25 PDB header: signaling protein Chain: A; PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii	
37	c1griA_	Alignment		100.0	28 PDB header: signal transduction adaptor Chain: A; PDB Molecule: growth factor bound protein 2; PDBTitle: grb2	
38	c2eyyA_	Alignment		100.0	20 PDB header: signaling protein Chain: A; PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i	
39	c2dviA_	Alignment		99.9	24 PDB header: signaling protein Chain: A; PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii	
40	c3qwxX_	Alignment		99.9	21 PDB header: signaling protein Chain: X; PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174	
41	c3qwyA_	Alignment		not modelled	99.9	21 PDB header: signaling protein Chain: A; PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
42	c213gA_	Alignment		not modelled	99.9	37 PDB header: signaling protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 7; PDBTitle: solution nmr structure of ch domain of rho guanine nucleotide exchange2 factor 7 from homo sapiens, northeast structural genomics consortium3 target hr4495e
43	d1ujoa_	Alignment		not modelled	99.9	23 Fold: CH domain-like Superfamily: Calponin-homology domain, CH-domain Family: Calponin-homology domain, CH-domain
44	c1wymA_	Alignment		not modelled	99.9	18 PDB header: structural protein Chain: A; PDB Molecule: transgelin-2; PDBTitle: solution structure of the ch domain of human transgelin-2
45	c1wyrA_	Alignment	not modelled	99.9	32 PDB header: structural protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 6; PDBTitle: solution structure of the ch domain of human rho guanine2 nucleotide exchange factor 6	
46	c1mv3A_	Alignment	not modelled	99.9	18 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	
47	c213sA_	Alignment	not modelled	99.9	26 PDB header: structural protein Chain: A; PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk	
48	c1wypA_	Alignment	not modelled	99.8	23 PDB header: structural protein Chain: A; PDB Molecule: calponin 1; PDBTitle: solution structure of the ch domain of human calponin 1	
49	c21g1A_	Alignment	not modelled	99.8	10 PDB header: metal binding protein Chain: A; PDB Molecule: a-kinase anchor protein 13; PDBTitle: solution structure of the human akap13 ph domain and	

						stabilizing dh2 helix
50	c1wynA	Alignment	not modelled	99.8	23	PDB header: structural protein Chain: A: PDB Molecule: calponin-2; PDBTitle: solution structure of the ch domain of human calponin-2
51	c1ov3A	Alignment	not modelled	99.8	28	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
52	c1ng2A	Alignment	not modelled	99.8	31	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
53	d1p2xa	Alignment	not modelled	99.8	21	Fold: CH domain-like Superfamily: Calponin-homology domain, CH-domain Family: Calponin-homology domain, CH-domain
54	c1p5sA	Alignment	not modelled	99.8	21	PDB header: cytokine Chain: A: PDB Molecule: ras gtpase-activating-like protein rng2; PDBTitle: structure and function of the calponin-homology domain of2 an iqgap protein from schizosaccharomyces pombe
55	d1h67a	Alignment	not modelled	99.8	26	Fold: CH domain-like Superfamily: Calponin-homology domain, CH-domain Family: Calponin-homology domain, CH-domain
56	c2rqrA	Alignment	not modelled	99.8	22	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	c3i6xC	Alignment	not modelled	99.8	23	PDB header: calmodulin-binding, membrane protein Chain: C: PDB Molecule: ras gtpase-activating-like protein iqgap1; PDBTitle: crystal structure of the calponin homology domain of iqgap1
58	c2jxbA	Alignment	not modelled	99.8	16	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	d1ng2a2	Alignment	not modelled	99.8	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
60	c3nmzD	Alignment	not modelled	99.7	23	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
61	c1x43A	Alignment	not modelled	99.7	28	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)
62	c2dlpA	Alignment	not modelled	99.7	23	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
63	c1zlmA	Alignment	not modelled	99.7	40	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
64	c1x2qA	Alignment	not modelled	99.7	30	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
65	c2ed0A	Alignment	not modelled	99.7	30	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)
66	d1ov3a2	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
67	d1uj0a	Alignment	not modelled	99.7	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
68	c1k9aB	Alignment	not modelled	99.7	23	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
69	c2csqA	Alignment	not modelled	99.7	26	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
70	c2l0aA	Alignment	not modelled	99.7	31	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
71	c2e5kA	Alignment	not modelled	99.7	21	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
72	c2cubA	Alignment	not modelled	99.7	34	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
73	d1k4us	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	d2dfka2	Alignment	not modelled	99.7	17	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
75	c2oi3A	Alignment	not modelled	99.7	24	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)
76	c2xmfA_	Alignment	not modelled	99.7	34	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
77	d1l1ja_	Alignment	not modelled	99.7	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
78	c2nwmA_	Alignment	not modelled	99.7	41	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
79	d1loota_	Alignment	not modelled	99.7	44	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
80	c2ysqA_	Alignment	not modelled	99.7	30	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 9; PDBTitle: solution structure of the sh3 domain from rho guanine2 nucleotide exchange factor 9
81	c2yt6A_	Alignment	not modelled	99.7	24	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
82	c1wx6A_	Alignment	not modelled	99.7	24	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
83	d1u5sa1	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	c2df6A_	Alignment	not modelled	99.7	42	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
85	d1gcqa_	Alignment	not modelled	99.7	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
86	c1x6gA_	Alignment	not modelled	99.7	19	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.
87	c2kbtA_	Alignment	not modelled	99.7	61	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
88	d1sema_	Alignment	not modelled	99.7	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
89	c2d8hA_	Alignment	not modelled	99.7	25	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
90	c2cudA_	Alignment	not modelled	99.7	20	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)
91	d1utia_	Alignment	not modelled	99.7	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
92	d1wlpb2	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
93	c2yupA_	Alignment	not modelled	99.7	31	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
94	c1x6bA_	Alignment	not modelled	99.7	24	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
95	c2k6dA_	Alignment	not modelled	99.7	34	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
96	c2yunA_	Alignment	not modelled	99.7	42	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
97	c2dbkA_	Alignment	not modelled	99.7	37	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
98	c2dmoA_	Alignment	not modelled	99.7	23	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)
99	d1udla_	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	c2js0A_	Alignment	not modelled	99.7	42	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck

101	d1uuea_	Alignment	not modelled	99.7	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
102	d1wiea_	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	c2yuoA_	Alignment	not modelled	99.7	37	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
104	d1oeba_	Alignment	not modelled	99.7	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
105	d1uhca_	Alignment	not modelled	99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
106	c2epdA_	Alignment	not modelled	99.7	38	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
107	c1wyxA_	Alignment	not modelled	99.7	33	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
108	d1u06a1	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
109	d1ov3a1	Alignment	not modelled	99.7	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
110	c2dl4A_	Alignment	not modelled	99.7	32	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
111	d1ycsb2	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	c1zx6A_	Alignment	not modelled	99.7	35	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
113	d1gria2	Alignment	not modelled	99.7	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	c2d1xD_	Alignment	not modelled	99.7	40	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
115	c2fpeB_	Alignment	not modelled	99.7	26	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
116	c2dm1A_	Alignment	not modelled	99.7	91	PDB header: signaling protein Chain: A: PDB Molecule: protein vav-2; PDBTitle: solution structure of the second sh3 domain of human2 protein vav-2
117	c3rnjA_	Alignment	not modelled	99.7	30	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)
118	c2ekhA_	Alignment	not modelled	99.7	20	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
119	c2ebpA_	Alignment	not modelled	99.7	31	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
120	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