





















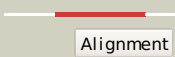

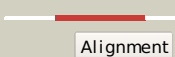

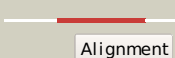

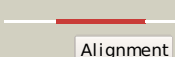

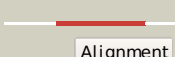



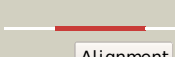

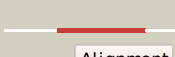


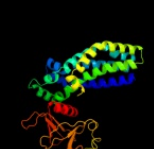
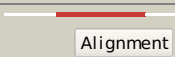

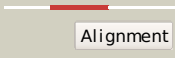
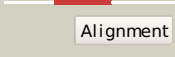
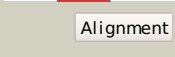

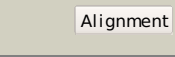
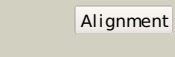




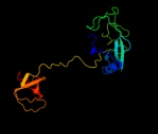




Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	Q9UKW4
Date	Thu Apr 26 09:47:10 BST 2012
Unique Job ID	9a0d39b8b1617d36

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3ky9B_	 Alignment		100.0	63	PDB header: apoptosis Chain: B; PDB Molecule: proto-oncogene vav; PDBTitle: autoinhibited vav1
2	c2vrwB_	 Alignment		100.0	60	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	59	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	19	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	23	PDB header: signaling protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
6	c1xdvB_	 Alignment		100.0	18	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.
7	c2dfkA_	 Alignment		100.0	23	PDB header: cell cycle Chain: A; PDB Molecule: collybistin ii; PDBTitle: crystal structure of the cdc42-collybistin ii complex
8	c3odoA_	 Alignment		100.0	20	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
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	c1x86A_	 Alignment		100.0	17	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
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	c1foeA	 Alignment		100.0	24	PDB header: signaling protein, immune system/signal 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
13	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
14	c3mpxA	 Alignment		100.0	17	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
15	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 of2 p115-rhogef
16	c3odwB	 Alignment		100.0	19	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
17	c1ki1B	 Alignment		100.0	19	PDB header: signaling protein Chain: B: PDB Molecule: intersectin long form; PDBTitle: guanine nucleotide exchange region of intersectin in2 complex with cdc42
18	c2z0qA	 Alignment		100.0	20	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 3; PDBTitle: crystal structure of dh-ph domain of rhogef3(xpln)
19	c1lb1C	 Alignment		100.0	20	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
20	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
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	d1f5xa	 Alignment	not modelled	100.0	62	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
24	d1txda1	 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	d1dbha1	 Alignment	not modelled	100.0	17	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	22	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	23	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	96	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	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)

30	c3eo2A	Alignment	not modelled	100.0	24	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
31	d1kz7a1	Alignment	not modelled	100.0	22	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
32	c3gf9A	Alignment	not modelled	100.0	25	PDB header: endocytosis Chain: A: PDB Molecule: intersectin 2; PDBTitle: crystal structure of human intersectin 2 rhogef domain
33	d1by1a	Alignment	not modelled	100.0	24	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
34	d1ntya1	Alignment	not modelled	100.0	19	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
35	c1griA	Alignment		100.0	23	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
36	c2eyyA	Alignment		100.0	21	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
37	c2dvjA	Alignment		99.9	29	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii
38	c2eyzA	Alignment		99.9	26	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
39	c3qwxX	Alignment		99.9	23	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
40	c3qwyA	Alignment		99.9	23	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
41	c213gA	Alignment	not modelled	99.9	34	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
42	d1ujoa	Alignment	not modelled	99.9	23	Fold: CH domain-like Superfamily: Calponin-homology domain, CH-domain Family: Calponin-homology domain, CH-domain
43	c1wymA	Alignment	not modelled	99.9	21	PDB header: structural protein Chain: A: PDB Molecule: transgelin-2; PDBTitle: solution structure of the ch domain of human transgelin-2
44	c1mv3A	Alignment	not modelled	99.9	17	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
45	c1wyrA	Alignment	not modelled	99.9	31	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	c213sA	Alignment	not modelled	99.9	21	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
47	c1wypA	Alignment	not modelled	99.8	24	PDB header: structural protein Chain: A: PDB Molecule: calponin 1; PDBTitle: solution structure of the ch domain of human calponin 1
48	c21g1A	Alignment	not modelled	99.8	11	PDB header: metal binding protein Chain: A: PDB Molecule: α-kinase anchor protein 13; PDBTitle: solution structure of the human akap13 ph domain and stabilizing dh2 helix
49	c1wynA	Alignment	not modelled	99.8	27	PDB header: structural protein Chain: A: PDB Molecule: calponin-2;

						PDBTitle: solution structure of the ch domain of human calponin-2
50	c1ov3A_	Alignment	not modelled	99.8	26	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
51	d1h67a_	Alignment	not modelled	99.8	29	Fold: CH domain-like Superfamily: Calponin-homology domain, CH-domain Family: Calponin-homology domain, CH-domain
52	c1ng2A_	Alignment	not modelled	99.8	25	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
53	d1p2xa_	Alignment	not modelled	99.8	25	Fold: CH domain-like Superfamily: Calponin-homology domain, CH-domain Family: Calponin-homology domain, CH-domain
54	c1p5sA_	Alignment	not modelled	99.8	25	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	d1ng2a2_	Alignment	not modelled	99.8	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
56	c2rqrA_	Alignment	not modelled	99.8	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	c3i6xC_	Alignment	not modelled	99.8	24	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	c3nmzD_	Alignment	not modelled	99.8	31	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
59	d1kz7a2_	Alignment	not modelled	99.7	21	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
60	c2csgA_	Alignment	not modelled	99.7	28	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
61	c1k9aB_	Alignment	not modelled	99.7	24	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
62	c1x6gA_	Alignment	not modelled	99.7	22	PDB header: signaling protein Chain: A: PDB Molecule: megakaryocyte-associated tyrosine-protein kinase; PDBTitle: solution structures of the sh3 domain of human2 megakaryocyte-associated tyrosine-protein kinase.
63	d2dfka2_	Alignment	not modelled	99.7	22	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
64	d1ov3a2_	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
65	c2kbtA_	Alignment	not modelled	99.7	64	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
66	c1x2qA_	Alignment	not modelled	99.7	32	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
67	c1x43A_	Alignment	not modelled	99.7	30	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)
68	c3rniA_	Alignment	not modelled	99.7	24	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)
69	c2dlpA_	Alignment	not modelled	99.7	27	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
70	d1wlpb2_	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
71	c2e5kA_	Alignment	not modelled	99.7	19	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	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)
73	c2egeA_	Alignment	not modelled	99.7	33	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
74	c1zlmA_	Alignment	not modelled	99.7	34	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
						PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor

75	c2ysqA	Alignment	not modelled	99.7	33	9; PDBTitle: solution structure of the sh3 domain from rho guanine2 nucleotide exchange factor 9
76	c2cubA	Alignment	not modelled	99.7	38	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
77	c2jxbA	Alignment	not modelled	99.7	23	PDB header: signaling protein complex Chain: A: PDB Molecule: t-cell surface glycoprotein cd3 epsilon chain, PDBTitle: structure of cd3epsilon-non-nck2 first sh3 domain complex
78	c2dbkA	Alignment	not modelled	99.7	38	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
79	d1uj0a	Alignment	not modelled	99.7	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
80	c2nwmA	Alignment	not modelled	99.7	42	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
81	d1wiea	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
82	c2ebpA	Alignment	not modelled	99.7	36	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
83	c2d1xD	Alignment	not modelled	99.7	36	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
84	d1rj2a2	Alignment	not modelled	99.7	22	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
85	c2xmfA	Alignment	not modelled	99.7	34	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
86	d1uhca	Alignment	not modelled	99.7	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
87	c2epdA	Alignment	not modelled	99.7	35	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	c2l0aA	Alignment	not modelled	99.7	27	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
89	d1uuea	Alignment	not modelled	99.7	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
90	c2dl7A	Alignment	not modelled	99.7	36	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
91	d1k4us	Alignment	not modelled	99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
92	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
93	d1pwta	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	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
95	c2oi3A	Alignment	not modelled	99.7	21	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck; PDBTitle: nmr structure analysis of the hematopoietic cell kinase sh32 domain complexed with an artificial high affinity ligand3 (pd1)
96	c2dl8A	Alignment	not modelled	99.7	35	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
97	c2j6kE	Alignment	not modelled	99.7	36	PDB header: protein binding Chain: E: PDB Molecule: cd2-associated protein; PDBTitle: n-terminal sh3 domain of cms (cd2ap human homolog)
98	c2df6A	Alignment	not modelled	99.7	41	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
99	c2ekhA	Alignment	not modelled	99.7	24	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
100	c1hb9A	Alignment	not modelled	99.7	30	PDB header: transferase Chain: A: PDB Molecule: amphiphysin 2;

100	c1b9xA	Alignment	not modelled	99.7	30	PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
101	d1bb9a	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
102	c2yua	Alignment	not modelled	99.7	46	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
103	d1gcqa	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
104	c1wyxA	Alignment	not modelled	99.7	32	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
105	c2yuoA	Alignment	not modelled	99.7	36	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
106	c1zukA	Alignment	not modelled	99.7	24	PDB header: contractile protein Chain: A: PDB Molecule: myosin tail region-interacting protein mtl1; PDBTitle: yeast bbc1 sh3 domain complexed with a peptide from las17
107	d1ugva	Alignment	not modelled	99.7	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	d1udla	Alignment	not modelled	99.7	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
109	c2eyxA	Alignment	not modelled	99.7	31	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
110	d1i1ja	Alignment	not modelled	99.7	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
111	d1loota	Alignment	not modelled	99.7	46	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	c2dl4A	Alignment	not modelled	99.7	26	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
113	d1ov3a1	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	c1w70A	Alignment	not modelled	99.7	31	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
115	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)
116	d1utia	Alignment	not modelled	99.7	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	c2bz8B	Alignment	not modelled	99.7	37	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
118	d1gria2	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	d1sema	Alignment	not modelled	99.7	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	d1e6ga	Alignment	not modelled	99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain