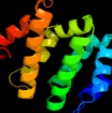
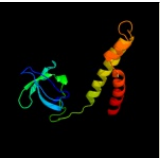
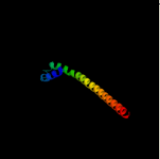
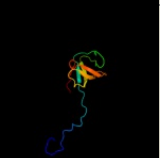
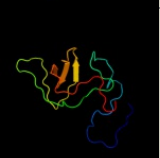
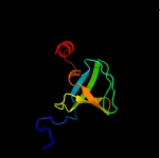
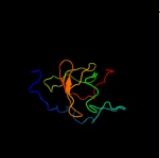
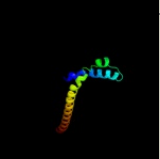
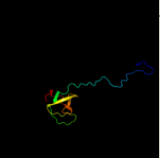
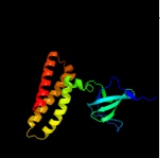
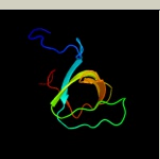


Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	O74749
Date	Sun Jul 8 11:51:36 BST 2012
Unique Job ID	1192323390d339dd

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1dvpA_	 Alignment		100.0	20	PDB header: transferase Chain: A: PDB Molecule: hepatocyte growth factor-regulated tyrosine PDBTitle: crystal structure of the vhs and fyve tandem domains of hrs,2 a protein involved in membrane trafficking and signal3 transduction
2	c1x5bA_	 Alignment		100.0	30	PDB header: protein binding Chain: A: PDB Molecule: signal transducing adaptor molecule 2; PDBTitle: the solution structure of the vhs domain of human signal2 transducing adaptor molecule 2
3	c3zyqA_	 Alignment		100.0	19	PDB header: signaling Chain: A: PDB Molecule: hepatocyte growth factor-regulated tyrosine kinase PDBTitle: crystal structure of the tandem vhs and fyve domains of hepatocyte2 growth factor-regulated tyrosine kinase substrate (hgs-hrs) at 1.483 a resolution
4	d1dvpA1	 Alignment		100.0	20	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
5	d1elka_	 Alignment		100.0	25	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
6	d1juqa_	 Alignment		100.0	23	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
7	d1ujka_	 Alignment		100.0	23	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
8	d1mhqa_	 Alignment		100.0	20	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
9	c3rruA_	 Alignment		100.0	24	PDB header: signaling protein Chain: A: PDB Molecule: tom1l1 protein; PDBTitle: x-ray crystal structure of the vhs domain of human tom1-like protein,2 northeast structural genomics consortium target hr3050e
10	c2dcpA_	 Alignment		99.6	19	PDB header: unknown function Chain: A: PDB Molecule: hypothetical protein (rafl09-17-b18); PDBTitle: fully automated nmr structure determination of the enth-vhs2 domain at3g16270 from arabidopsis thaliana
11	c1mv3A_	 Alignment		99.6	24	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

12	c2dx1A_	Alignment		99.6	18	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asaf
13	c2pjwV_	Alignment		99.6	18	PDB header: endocytosis/exocytosis Chain: V: PDB Molecule: vacuolar protein sorting-associated protein 27; PDBTitle: the vps27/hse1 complex is a gat domain-based scaffold for2 ubiquitin-dependent sorting
14	c1griA_	Alignment		99.5	36	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
15	c2eyzA_	Alignment		99.5	16	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
16	d1ng2a2	Alignment		99.4	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
17	c2l3sA_	Alignment		99.4	16	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
18	c2pjwH_	Alignment		99.4	31	PDB header: endocytosis/exocytosis Chain: H: PDB Molecule: uncharacterized protein yhl002w; PDBTitle: the vps27/hse1 complex is a gat domain-based scaffold for2 ubiquitin-dependent sorting
19	c2eyyA_	Alignment		99.4	22	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
20	c3a98A_	Alignment		99.4	21	PDB header: signaling protein Chain: A: PDB Molecule: dedicator of cytokinesis protein 2; PDBTitle: crystal structure of the complex of the interacting regions of dock22 and elmo1
21	c2lqnA_	Alignment	not modelled	99.4	14	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
22	c3nmzD_	Alignment	not modelled	99.4	25	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of apc complexed with asaf
23	c2jxbA_	Alignment	not modelled	99.3	20	PDB header: signaling protein complex Chain: A: PDB Molecule: t-cell surface glycoprotein cd3 epsilon chain, PDBTitle: structure of cd3epsilon-nck2 first sh3 domain complex
24	c2ed0A_	Alignment		99.3	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)
25	c2dbkA_	Alignment	not modelled	99.3	16	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
26	c2o2oA_	Alignment	not modelled	99.3	24	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
27	c1wi7A_	Alignment		99.3	31	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
28	d1udla_	Alignment	not modelled	99.3	36 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
29	c2dlpA_	Alignment	not modelled	99.3	32 PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
30	c2ed1A_	Alignment	not modelled	99.3	30 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
31	c2nwmA_	Alignment	not modelled	99.3	36 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
32	c2djqA_	Alignment	not modelled	99.3	32 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
33	d1ov3a2	Alignment	not modelled	99.3	22 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
34	c2dmoA_	Alignment	not modelled	99.3	27 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)
35	c2rqrA_	Alignment	not modelled	99.3	18 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
36	c1x2qA_	Alignment	not modelled	99.3	36 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
37	c2dl3A_	Alignment	not modelled	99.3	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
38	c2yunA_	Alignment	not modelled	99.3	34 PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
39	c2l0aA_	Alignment	not modelled	99.3	38 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
40	c2oi3A_	Alignment	not modelled	99.3	29 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)
41	d1k4us_	Alignment	not modelled	99.3	42 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
42	d1awja_	Alignment	not modelled	99.3	22 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
43	c2yuoA_	Alignment	not modelled	99.3	33 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
44	c2dl4A_	Alignment	not modelled	99.3	42 PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
45	c1ng2A_	Alignment	not modelled	99.3	25 PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
46	c2cubA_	Alignment	not modelled	99.3	24 PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
47	c2dila_	Alignment	not modelled	99.3	30 PDB header: cell adhesion Chain: A: PDB Molecule: proline-serine-threonine phosphatase-interacting PDBTitle: solution structure of the sh3 domain of the human proline-2 serine-threonine phosphatase-interacting protein 1
48	d1oeba_	Alignment	not modelled	99.3	46 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
49	c2csqA_	Alignment	not modelled	99.3	24 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
50	c2ebpA_	Alignment	not modelled	99.3	28 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
51	d1utia_	Alignment	not modelled	99.3	46 Fold: SH3-like barrel Superfamily: SH3-domain

						Family: SH3-domain
52	d2hspa_	Alignment	not modelled	99.3	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
53	c2yt6A_	Alignment	not modelled	99.3	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
54	c2egaA_	Alignment	not modelled	99.3	23	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
55	c2dnuA_	Alignment	not modelled	99.2	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
56	c2ysqA_	Alignment	not modelled	99.2	33	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
57	c2xmfA_	Alignment	not modelled	99.2	33	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
58	c2ekhA_	Alignment	not modelled	99.2	19	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
59	dlsema_	Alignment	not modelled	99.2	45	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
60	c2eyxA_	Alignment	not modelled	99.2	18	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
61	dlujya_	Alignment	not modelled	99.2	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
62	c2df6A_	Alignment	not modelled	99.2	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
63	c2bz8B_	Alignment	not modelled	99.2	27	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
64	dluj0a_	Alignment	not modelled	99.2	46	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
65	c1x43A_	Alignment	not modelled	99.2	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)
66	c1z9qA_	Alignment	not modelled	99.2	38	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: solution structure of sh3 domain of p40phox
67	c3qwxX_	Alignment	not modelled	99.2	25	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
68	dlawwa_	Alignment	not modelled	99.2	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
69	c1x2pA_	Alignment	not modelled	99.2	35	PDB header: transferase Chain: A: PDB Molecule: protein arginine n-methyltransferase 2; PDBTitle: solution structure of the sh3 domain of the protein2 arginine n-methyltransferase 2
70	c2csiA_	Alignment	not modelled	99.2	35	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: rim binding protein 2; PDBTitle: solution structure of the third sh3 domain of human rim-2 binding protein 2
71	c2eqiA_	Alignment	not modelled	99.2	35	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
72	c2dbmA_	Alignment	not modelled	99.2	39	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
73	c2dl8A_	Alignment	not modelled	99.2	25	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
74	c2epdA_	Alignment	not modelled	99.2	25	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
75	dlgria2	Alignment	not modelled	99.2	43	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
76	c2dvjA_	Alignment	not modelled	99.2	24	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii PDB header: signaling protein

77	c2kymA	Alignment	not modelled	99.2	23	Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of the bem1p sh3-ci domain from <i>I.elongisporus</i> in2 complex with ste20p peptide
78	c2yupA	Alignment	not modelled	99.2	31	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
79	c2drmB	Alignment	not modelled	99.2	40	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
80	c3qwyA	Alignment	not modelled	99.2	27	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
81	c1zx6A	Alignment	not modelled	99.2	45	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
82	d1uhfa	Alignment	not modelled	99.2	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
83	c2cudA	Alignment	not modelled	99.2	16	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)
84	c2pqhA	Alignment	not modelled	99.2	37	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
85	c1w70A	Alignment	not modelled	99.2	42	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
86	c2k2mA	Alignment	not modelled	99.2	26	PDB header: signaling protein Chain: A: PDB Molecule: eps8-like protein 1; PDBTitle: structural basis of pxxdy motif recognition in sh3 binding
87	d1ng2a1	Alignment	not modelled	99.2	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
88	d1wlpb2	Alignment	not modelled	99.2	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
89	d1gcqa	Alignment	not modelled	99.2	49	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
90	d1ugva	Alignment	not modelled	99.2	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
91	d1ov3a1	Alignment	not modelled	99.2	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
92	c2egeA	Alignment	not modelled	99.2	19	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
93	c2e5kA	Alignment	not modelled	99.2	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
94	c2eywA	Alignment	not modelled	99.2	31	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: n-terminal sh3 domain of ct10-regulated kinase
95	c2ct3A	Alignment	not modelled	99.2	31	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
96	c2d1xD	Alignment	not modelled	99.2	43	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
97	c2bzxA	Alignment	not modelled	99.2	20	PDB header: sh3 domain Chain: A: PDB Molecule: crkl-like protein; PDBTitle: atomic model of crkl-sh3c monomer
98	c2rqvA	Alignment	not modelled	99.2	18	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
99	d1uuea	Alignment	not modelled	99.2	42	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	c1zlmA	Alignment	not modelled	99.2	42	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
101	c1x69A	Alignment	not modelled	99.2	31	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
102	d1gbra	Alignment	not modelled	99.2	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	c2dl7A	Alignment	not modelled	99.2	37	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein

104	c2fpeB	Alignment	not modelled	99.2	27	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
105	c2ke9A	Alignment	not modelled	99.2	37	PDB header: protein binding Chain: A: PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
106	d1wiea	Alignment	not modelled	99.2	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
107	c1wxtA	Alignment	not modelled	99.2	24	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hypothetical protein flj21522; PDBTitle: solution structure of the sh3 domain of human hypothetical2 protein flj21522
108	c2krnA	Alignment	not modelled	99.2	35	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
109	c2d8hA	Alignment	not modelled	99.2	28	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
110	c1wxba	Alignment	not modelled	99.2	32	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: epidermal growth factor receptor pathway PDBTitle: solution structure of the sh3 domain from human epidermal2 growth factor receptor pathway substrate 8-like protein
111	c1negA	Alignment	not modelled	99.2	35	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
112	d1nega	Alignment	not modelled	99.2	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	d1pwta	Alignment	not modelled	99.2	40	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	d1qlya	Alignment	not modelled	99.2	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	d1spka	Alignment	not modelled	99.1	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	d1i07a	Alignment	not modelled	99.1	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	c1wyxA	Alignment	not modelled	99.1	43	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
118	d1u06a1	Alignment	not modelled	99.1	42	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	c2creA	Alignment	not modelled	99.1	35	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
120	d1h8ka	Alignment	not modelled	99.1	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain