

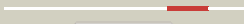






















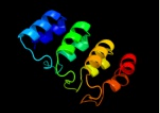







#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	clycsB_	 Alignment		100.0	100	PDB header: complex (anti-oncogene/ankyrin repeats) Chain: B: PDB Molecule: 53bp2; PDBTitle: p53-53bp2 complex
2	c2vgeA_	 Alignment		100.0	52	PDB header: transcription Chain: A: PDB Molecule: rela-associated inhibitor; PDBTitle: crystal structure of the c-terminal region of human iaspp
3	dlycsb1	 Alignment		99.8	100	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
4	dln11a_	 Alignment		99.8	33	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
5	c1n11A_	 Alignment		99.8	33	PDB header: structural protein Chain: A: PDB Molecule: ankyrin; PDBTitle: d34 region of human ankyrin-r and linker
6	c3jueA_	 Alignment		99.7	23	PDB header: protein transport/endocytosis Chain: A: PDB Molecule: arfgap with coiled-coil, ank repeat and ph domain- PDBTitle: crystal structure of arfgap and ank repeat domain of acap1
7	c2b0oF_	 Alignment		99.7	26	PDB header: metal binding protein Chain: F: PDB Molecule: uplc1; PDBTitle: crystal structure of uplc1 gap domain
8	c3lvrE_	 Alignment		99.7	22	PDB header: protein transport Chain: E: PDB Molecule: arf-gap with sh3 domain, ank repeat and ph domain- PDBTitle: the crystal structure of asap3 in complex with arf6 in transition2 state soaked with calcium
9	d1s70b_	 Alignment		99.7	45	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
10	c3ui2A_	 Alignment		99.7	20	PDB header: transport protein Chain: A: PDB Molecule: signal recognition particle 43 kda protein, chloroplastic; PDBTitle: crystal structure of the cpsrp54 tail bound to cpsrp43
11	c1dcqA_	 Alignment		99.6	21	PDB header: metal binding protein Chain: A: PDB Molecule: pyk2-associated protein beta; PDBTitle: crystal structure of the arf-gap domain and ankyrin repeats2 of papbeta.

12	c3keaB_	Alignment		99.6	19	PDB header: viral protein Chain: B: PDB Molecule: k1l; PDBTitle: structure function studies of vaccinia virus host-range protein k12 reveal a novel ankyrin repeat interaction surface for k1s function
13	c3utmA_	Alignment		99.6	24	PDB header: transferase/signaling protein Chain: A: PDB Molecule: tankyrase-1; PDBTitle: crystal structure of a mouse tankyrase-axin complex
14	d2ajaa1	Alignment		99.6	16	Fold: alpha-alpha superhelix Superfamily: Pseudo ankyrin repeat-like Family: Pseudo ankyrin repeat
15	d1dcqa1	Alignment		99.6	25	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
16	c2fo1E_	Alignment		99.6	21	PDB header: gene regulation/signalling protein/dna Chain: E: PDB Molecule: lin-12 protein; PDBTitle: crystal structure of the csl-notch-mastermind ternary2 complex bound to dna
17	c2ajaa_	Alignment		99.6	15	PDB header: protein binding Chain: A: PDB Molecule: ankyrin repeat family protein; PDBTitle: x-ray structure of an ankyrin repeat family protein q5zsv02 from legionella pneumophila. northeast structural genomics3 consortium target lgr21.
18	d2fo1e1	Alignment		99.6	20	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
19	c3ljnA_	Alignment		99.5	24	PDB header: unknown function Chain: A: PDB Molecule: hypothetical protein; PDBTitle: ankyrin repeat protein from leishmania major
20	c2rfaA_	Alignment		99.5	36	PDB header: membrane protein Chain: A: PDB Molecule: transient receptor potential cation channel subfamily v PDBTitle: crystal structure of the mouse trpv6 ankyrin repeat domain
21	c2zgdA_	Alignment	not modelled	99.5	30	PDB header: de novo protein Chain: A: PDB Molecule: 3 repeat synthetic ankyrin; PDBTitle: asn-hydroxylation stabilises the ankyrin repeat domain fold
22	d1ycsb2	Alignment	not modelled	99.5	100	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
23	c1sw6A_	Alignment	not modelled	99.5	18	PDB header: transcription regulation Chain: A: PDB Molecule: regulatory protein swi6; PDBTitle: s. cerevisiae swi6 ankyrin-repeat fragment
24	d1sw6a_	Alignment	not modelled	99.5	18	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
25	d1uoha_	Alignment	not modelled	99.5	47	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
26	c2xenA_	Alignment	not modelled	99.5	29	PDB header: de novo protein Chain: A: PDB Molecule: nilc mut4; PDBTitle: structural determinants for improved thermal stability of2 designed ankyrin repeat proteins with a redesigned c-3 capping module.
27	c3hraA_	Alignment	not modelled	99.5	34	PDB header: structural protein Chain: A: PDB Molecule: ankyrin repeat family protein; PDBTitle: crystal structure of ef0377 an ankyrin repeat protein
28	c3deoA_	Alignment	not modelled	99.5	25	PDB header: protein transport, membrane protein Chain: A: PDB Molecule: signal recognition particle 43 kda protein; PDBTitle: structural basis for specific substrate recognition by the2 chloroplast signal recognition particle protein cpsrp43
						Fold: beta-hairpin-alpha-hairpin repeat

29	d1wdya_	Alignment	not modelled	99.4	24	Superfamily: Ankyrin repeat Family: Ankyrin repeat
30	c3ehrB_	Alignment	not modelled	99.4	24	PDB header: signaling protein Chain: B: PDB Molecule: osteoclast-stimulating factor 1; PDBTitle: crystal structure of human osteoclast stimulating factor
31	c2f8xK_	Alignment	not modelled	99.4	18	PDB header: transcription/dna Chain: K: PDB Molecule: neurogenic locus notch homolog protein 1; PDBTitle: crystal structure of activated notch, csf and mam1 on hes-12 promoter dna sequence
32	c2kbxA_	Alignment	not modelled	99.4	23	PDB header: cell adhesion Chain: A: PDB Molecule: integrin-linked protein kinase; PDBTitle: solution structure of ilk-pinch complex
33	c1n0qB_	Alignment	not modelled	99.4	41	PDB header: structural protein Chain: B: PDB Molecule: 3 ankyrin repeats; PDBTitle: 3ank: a designed ankyrin repeat protein with three identical consensus2 repeats
34	c2xaiD_	Alignment	not modelled	99.4	27	PDB header: transcription Chain: D: PDB Molecule: ankyrin repeat and socs box protein 9; PDBTitle: crystal structure of ankyrin repeat and socs box-containing2 protein 9 (asb9) in complex with elonginb and elonginc
35	d1iknd_	Alignment	not modelled	99.4	21	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
36	c3eu9B_	Alignment	not modelled	99.4	25	PDB header: protein binding, methyl-lysine-binding p Chain: B: PDB Molecule: huntingtin-interacting protein 14; PDBTitle: the ankyrin repeat domain of huntingtin interacting protein 14
37	d1ot8a_	Alignment	not modelled	99.4	18	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
38	c3b7bB_	Alignment	not modelled	99.3	36	PDB header: transferase Chain: B: PDB Molecule: euchromatic histone-lysine n-methyltransferase 1; PDBTitle: ehmt1 (glp) ankyrin repeat domain (structure 1)
39	c3d9hA_	Alignment	not modelled	99.3	27	PDB header: structural protein, protein binding Chain: A: PDB Molecule: cdna flj77766, highly similar to homo sapiens PDBTitle: crystal structure of the splice variant of human asb92 (hasb9-2), an ankyrin repeat protein
40	c3t8kB_	Alignment	not modelled	99.3	13	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: uncharacterized protein; PDBTitle: the crystal structure of a functionally unknown protein lebu_0176 from2 leptotrichia buccalis c-1013-b
41	c3jxiA_	Alignment	not modelled	99.3	22	PDB header: membrane protein Chain: A: PDB Molecule: vanilloid receptor-related osmotically activated channel PDBTitle: crystal structure of the chicken trpv4 ankyrin repeat domain
42	d1ixva_	Alignment	not modelled	99.3	35	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
43	d1blxb_	Alignment	not modelled	99.3	30	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
44	c3v30A_	Alignment	not modelled	99.3	28	PDB header: protein binding Chain: A: PDB Molecule: dna-binding protein rfxank; PDBTitle: crystal structure of the peptide bound complex of the ankyrin repeat2 domains of human rfxank
45	c2f37B_	Alignment	not modelled	99.3	24	PDB header: membrane protein Chain: B: PDB Molecule: transient receptor potential cation channel PDBTitle: crystal structure of the ankyrin repeat domain of human2 trpv2
46	c1oy3D_	Alignment	not modelled	99.3	24	PDB header: dna binding protein Chain: D: PDB Molecule: transcription factor inhibitor i-kappa-b-beta; PDBTitle: crystal structure of an ikbbeta/nf-kb p65 homodimer complex
47	d1oy3d_	Alignment	not modelled	99.3	24	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
48	c2rfmB_	Alignment	not modelled	99.3	33	PDB header: protein binding Chain: B: PDB Molecule: putative ankyrin repeat protein tv1425; PDBTitle: structure of a thermophilic ankyrin repeat protein
49	d1k1aa_	Alignment	not modelled	99.3	22	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
50	d1bi7b_	Alignment	not modelled	99.3	24	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
51	c1n0rA_	Alignment	not modelled	99.3	66	PDB header: structural protein Chain: A: PDB Molecule: 4 ankyrin repeats; PDBTitle: 4ank: a designed ankyrin repeat protein with four identical2 consensus repeats
52	c2v5qC_	Alignment	not modelled	99.3	41	PDB header: transferase Chain: C: PDB Molecule: design ankyrin repeat protein; PDBTitle: crystal structure of wild-type plk-1 kinase domain in2 complex with a selective darpin
53	d2a5ea_	Alignment	not modelled	99.2	21	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
54	c2jabC_	Alignment	not modelled	99.2	30	PDB header: de novo protein Chain: C: PDB Molecule: h10-2-g3; PDBTitle: a designed ankyrin repeat protein evolved to picomolar2

					affinity to her2
55	d1ihba_	Alignment	not modelled	99.2	32 Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
56	c2l6bA_	Alignment	not modelled	99.2	32 PDB header: de novo protein Chain: A: PDB Molecule: nr1c; PDBTitle: nrc consensus ankyrin repeat protein solution structure
57	d1awcb_	Alignment	not modelled	99.2	24 Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
58	c1ympB_	Alignment	not modelled	99.2	22 PDB header: transcription Chain: B: PDB Molecule: notch 1 protein; PDBTitle: the crystal structure of a partial mouse notch-1 ankyrin2 domain: repeats 4 through 7 preserve an ankyrin fold
59	c3twtd_	Alignment	not modelled	99.2	28 PDB header: signaling protein/peptide Chain: D: PDB Molecule: tankyrase-2; PDBTitle: crystal structure of arc4 from human tankyrase 2 in complex with2 peptide from human mcl1 (chimeric peptide)
60	c3nmzD_	Alignment	not modelled	99.2	36 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	d1myoa_	Alignment	not modelled	99.2	28 Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
62	c1ympA_	Alignment	not modelled	99.2	22 PDB header: transcription Chain: A: PDB Molecule: notch 1 protein; PDBTitle: the crystal structure of a partial mouse notch-1 ankyrin2 domain: repeats 4 through 7 preserve an ankyrin fold
63	d1bd8a_	Alignment	not modelled	99.1	32 Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
64	c2nyiA_	Alignment	not modelled	99.1	17 PDB header: transport protein Chain: A: PDB Molecule: transient receptor potential cation channel PDBTitle: crystal structure of the ankyrin repeat domain of trpv1
65	c1wyxA_	Alignment	not modelled	99.1	29 PDB header: cell adhesion Chain: A: PDB Molecule: crk-associated substrate; PDBTitle: the crystal structure of the p130cas sh3 domain at 1.1 a2 resolution
66	c2kymA_	Alignment	not modelled	99.1	27 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
67	d1ug1a_	Alignment	not modelled	99.1	19 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
68	c2dnuA_	Alignment	not modelled	99.1	29 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
69	c1griA_	Alignment	not modelled	99.1	34 PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
70	c2yupA_	Alignment	not modelled	99.1	21 PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
71	c2jxbA_	Alignment	not modelled	99.1	35 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
72	c2egeA_	Alignment	not modelled	99.1	32 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
73	d1ng2a2	Alignment	not modelled	99.1	33 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	c2ysqA_	Alignment	not modelled	99.1	28 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
75	d1ilja_	Alignment	not modelled	99.1	20 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
76	c1u3oA_	Alignment	not modelled	99.1	28 PDB header: signaling protein Chain: A: PDB Molecule: huntingtin-associated protein-interacting PDBTitle: solution structure of rat kalirin n-terminal sh3 domain
77	c2i0nA_	Alignment	not modelled	99.1	32 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
78	d1k9aa1	Alignment	not modelled	99.1	22 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
79	c2d8hA_	Alignment	not modelled	99.1	28 PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
80	c2yunA_	Alignment	not modelled	99.1	31 PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
					PDB header: structural genomics, unknown function

81	c2creA	Alignment	not modelled	99.0	31	Chain: A: PDB Molecule: hef-like protein; PDBTitle: solution structure of rsgi ruh-036, an sh3 domain from2 human cdna
82	c2k2mA	Alignment	not modelled	99.0	31	PDB header: signaling protein Chain: A: PDB Molecule: eps8-like protein 1; PDBTitle: structural basis of ppxdy motif recognition in sh3 binding
83	c1w70A	Alignment	not modelled	99.0	21	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
84	c2azsA	Alignment	not modelled	99.0	30	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)
85	c2ct4A	Alignment	not modelled	99.0	26	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
86	c1x2pA	Alignment	not modelled	99.0	24	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
87	c2a28D	Alignment	not modelled	99.0	31	PDB header: signaling protein Chain: D: PDB Molecule: bzz1 protein; PDBTitle: atomic-resolution crystal structure of the second sh32 domain of yeast bzz1 determined from a pseudomerohedrally3 twinned crystal
88	c2ekhA	Alignment	not modelled	99.0	23	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
89	d1uffa	Alignment	not modelled	99.0	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
90	c1wx6A	Alignment	not modelled	99.0	28	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
91	c1x2qA	Alignment	not modelled	99.0	25	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
92	c2eywA	Alignment	not modelled	99.0	34	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: n-terminal sh3 domain of ct10-regulated kinase
93	d1ov3a2	Alignment	not modelled	99.0	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	c2dl5A	Alignment	not modelled	99.0	29	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the first sh3 domain of human2 kiaa0769 protein
95	d1gbra	Alignment	not modelled	99.0	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
96	c2xmfA	Alignment	not modelled	99.0	33	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
97	c3c5rB	Alignment	not modelled	99.0	28	PDB header: protein binding Chain: B: PDB Molecule: brca1-associated ring domain protein 1; PDBTitle: crystal structure of the bard1 ankyrin repeat domain and its2 functional consequences
98	c2csqA	Alignment	not modelled	99.0	31	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
99	c1x6gA	Alignment	not modelled	99.0	20	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.
100	c2dl7A	Alignment	not modelled	99.0	33	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
101	c1mv3A	Alignment	not modelled	99.0	30	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
102	d1udla	Alignment	not modelled	99.0	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	c2dl8A	Alignment	not modelled	99.0	27	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
104	c2yuoA	Alignment	not modelled	99.0	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
105	c2l0aA	Alignment	not modelled	99.0	30	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
106	d1uhca	Alianment	not modelled	99.0	28	Fold: SH3-like barrel Superfamily: SH3-domain

					Family: SH3-domain
107	d1uuea_	Alignment	not modelled	99.0	31 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	c2cubA_	Alignment	not modelled	99.0	22 PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
109	c2cudA_	Alignment	not modelled	99.0	24 PDB header: signaling protein Chain: A: PDB Molecule: src-like-adaptor; PDBTitle: solution structure of the sh3 domain of the human src-like2 adpoter protein (slap)
110	d1e6ga_	Alignment	not modelled	99.0	31 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
111	c2dmoA_	Alignment	not modelled	99.0	26 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)
112	d1k4us_	Alignment	not modelled	99.0	31 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	d1uj0a_	Alignment	not modelled	99.0	32 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	c2qrqA_	Alignment	not modelled	99.0	17 PDB header: protein binding Chain: A: PDB Molecule: engulfment and cell motility protein 1, linker, dedicator PDBTitle: the solution structure of human dock2 sh3 domain - elmo1 peptide2 chimera complex
115	d1j3ta_	Alignment	not modelled	99.0	23 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	c2yt6A_	Alignment	not modelled	99.0	33 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
117	c1x43A_	Alignment	not modelled	99.0	24 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)
118	c1zlmA_	Alignment	not modelled	99.0	35 PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
119	d1pwta_	Alignment	not modelled	99.0	35 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	c3nhnA_	Alignment	not modelled	99.0	39 PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck; PDBTitle: crystal structure of the src-family kinase hck sh3-sh2-linker2 regulatory region