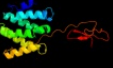
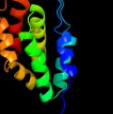


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P38753
Date	Mon Jul 2 19:20:20 BST 2012
Unique Job ID	574dcf2a2f45462e

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1dvpA_	 Alignment		100.0	16	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	c3zyqA_	 Alignment		100.0	17	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
3	c1x5bA_	 Alignment		100.0	21	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
4	d1dvpal	 Alignment		100.0	18	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
5	d1elka_	 Alignment		100.0	20	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
6	d1juqa_	 Alignment		100.0	22	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
7	d1mhqa_	 Alignment		100.0	20	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
8	d1ujka_	 Alignment		100.0	15	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
9	c3rruA_	 Alignment		100.0	21	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		99.8	17	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	d1lnza_	 Alignment		99.5	15	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: ENTH domain

12	c1mv3A_	Alignment		99.4	25	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
13	d1hf8a2	Alignment		99.3	16	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: Phosphoinositide-binding clathrin adaptor, N-terminal domain
14	d1hx8a2	Alignment		99.3	19	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: Phosphoinositide-binding clathrin adaptor, N-terminal domain
15	d1eyha_	Alignment		99.2	14	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: ENTH domain
16	c1griA_	Alignment		99.1	35	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
17	c2dx1A_	Alignment		99.1	36	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
18	c2qy7A_	Alignment		99.0	17	PDB header: protein binding Chain: A: PDB Molecule: clathrin interactor 1; PDBTitle: crystal structure of human epsinr enth domain
19	d1ng2a2	Alignment		99.0	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
20	c2eyzA_	Alignment		99.0	18	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
21	c2l3sA_	Alignment	not modelled	99.0	16	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
22	c3onlB_	Alignment	not modelled	99.0	15	PDB header: protein transport Chain: B: PDB Molecule: epsin-3; PDBTitle: yeast ent3_enth-vti1p_habc complex structure
23	c3nmzD_	Alignment		99.0	29	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
24	c2yuaA_	Alignment		99.0	29	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
25	d1udla_	Alignment	not modelled	98.9	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
26	c2eyyA_	Alignment	not modelled	98.9	23	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i PDB header: signaling protein

27	c2ed0A	Alignment	not modelled	98.9	33	Chain: A: PDB Molecule: abl interactor 2; PDBTitle: solution structure of the sh3 domain of abl interactor 22 (abelson interactor 2)
28	c1wi7A	Alignment	not modelled	98.9	32	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
29	c2o2oA	Alignment	not modelled	98.9	33	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
30	c1xgwA	Alignment	not modelled	98.9	16	PDB header: endocytosis Chain: A: PDB Molecule: epsin 4; PDBTitle: the crystal structure of human enthoprotin n-terminal domain
31	c1x2qA	Alignment	not modelled	98.9	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
32	c2l0aA	Alignment	not modelled	98.9	41	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
33	c2nwmA	Alignment	not modelled	98.9	34	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
34	c2ysqA	Alignment	not modelled	98.9	34	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
35	c2rqrA	Alignment	not modelled	98.9	15	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	c2yuoA	Alignment	not modelled	98.9	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
37	c1ng2A	Alignment	not modelled	98.9	24	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
38	d1k4us	Alignment	not modelled	98.9	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
39	c2dl3A	Alignment	not modelled	98.9	30	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
40	c2ed1A	Alignment	not modelled	98.9	31	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
41	c2dlpA	Alignment	not modelled	98.9	28	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
42	c2djgA	Alignment	not modelled	98.9	35	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
43	c2jxbA	Alignment	not modelled	98.9	21	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
44	c2lqnA	Alignment	not modelled	98.9	14	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
45	c2dmoA	Alignment	not modelled	98.9	28	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)
46	c2dilA	Alignment	not modelled	98.9	31	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
47	c2dl4A	Alignment	not modelled	98.8	36	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
48	c2cubA	Alignment	not modelled	98.8	32	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
49	c2ebpA	Alignment	not modelled	98.8	27	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
50	d2hsaA	Alignment	not modelled	98.8	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
						PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck;

51	c2oi3A_	Alignment	not modelled	98.8	27	PDBTitle: nmr structure analysis of the hematopoietic cell kinase sh32 domain complexed with an artificial high affinity ligand3 (pd1)
52	d1uj0a_	Alignment	not modelled	98.8	47	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
53	c2xmfA_	Alignment	not modelled	98.8	41	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
54	c1w70A_	Alignment	not modelled	98.8	45	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
55	c2ekhA_	Alignment	not modelled	98.8	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
56	c2dbkA_	Alignment	not modelled	98.8	18	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
57	c2dnuA_	Alignment	not modelled	98.8	20	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
58	c1z9gA_	Alignment	not modelled	98.8	42	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: solution structure of sh3 domain of p40phox
59	d1sema_	Alignment	not modelled	98.8	44	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
60	d1gria2	Alignment	not modelled	98.8	48	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
61	c2dbmA_	Alignment	not modelled	98.8	38	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
62	c2kymA_	Alignment	not modelled	98.8	19	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
63	c2pqhA_	Alignment	not modelled	98.8	33	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
64	c2bz8B_	Alignment	not modelled	98.8	30	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
65	d1ujya_	Alignment	not modelled	98.8	43	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	c1x2pA_	Alignment	not modelled	98.8	31	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
67	c2bzxA_	Alignment	not modelled	98.8	22	PDB header: sh3 domain Chain: A: PDB Molecule: crk-like protein; PDBTitle: atomic model of crkl-sh3c monomer
68	d1gcqa_	Alignment	not modelled	98.8	51	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
69	d1ov3a2	Alignment	not modelled	98.8	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
70	c2yt6A_	Alignment	not modelled	98.8	25	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
71	c2df6A_	Alignment	not modelled	98.8	45	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
72	c2drmB_	Alignment	not modelled	98.8	44	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
73	d1awwa_	Alignment	not modelled	98.8	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	c2eyxA_	Alignment	not modelled	98.8	19	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
75	c2e5kA_	Alignment	not modelled	98.8	23	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
76	c2rqvA_	Alignment	not modelled	98.8	16	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
						PDB header: signaling protein Chain: A: PDB Molecule: sh3 and px domain-containing protein

77	c2egaA	Alignment	not modelled	98.8	19	2a; PDBTitle: solution structure of the first sh3 domain from human2 kiaa0418 protein
78	c2dlxD	Alignment	not modelled	98.8	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
79	d1uhfa	Alignment	not modelled	98.8	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
80	c2yupA	Alignment	not modelled	98.8	27	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
81	d1utia	Alignment	not modelled	98.8	49	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
82	c2dl8A	Alignment	not modelled	98.8	26	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
83	c2epdA	Alignment	not modelled	98.8	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
84	d1oeba	Alignment	not modelled	98.8	48	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
85	c2eqiA	Alignment	not modelled	98.8	36	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
86	c1zx6A	Alignment	not modelled	98.8	40	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
87	d1ng2a1	Alignment	not modelled	98.8	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
88	c1zlmA	Alignment	not modelled	98.8	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
89	c2csqA	Alignment	not modelled	98.8	25	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
90	d1awja	Alignment	not modelled	98.8	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
91	d1nega	Alignment	not modelled	98.8	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
92	c1negA	Alignment	not modelled	98.8	33	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
93	d1ov3a1	Alignment	not modelled	98.8	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	c3qwxX	Alignment	not modelled	98.8	22	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
95	d1uuea	Alignment	not modelled	98.7	40	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
96	c2js0A	Alignment	not modelled	98.7	34	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
97	c1x43A	Alignment	not modelled	98.7	27	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)
98	d1ugva	Alignment	not modelled	98.7	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
99	c2fpeB	Alignment	not modelled	98.7	25	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
100	d1j3ta	Alignment	not modelled	98.7	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	d1phta	Alignment	not modelled	98.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
102	c1wyxA	Alignment	not modelled	98.7	46	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
103	d1uffa	Alignment	not modelled	98.7	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

104	c1x69A_	Alignment	not modelled	98.7	36	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
105	c2krnA_	Alignment	not modelled	98.7	39	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
106	c2i0nA_	Alignment	not modelled	98.7	23	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
107	c3qwyA_	Alignment	not modelled	98.7	25	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
108	c2dvjA_	Alignment	not modelled	98.7	21	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii
109	c2ke9A_	Alignment	not modelled	98.7	43	PDB header: protein binding Chain: A: PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
110	dlpwta_	Alignment	not modelled	98.7	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
111	dlu06a1	Alignment	not modelled	98.7	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	c2ct3A_	Alignment	not modelled	98.7	33	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
113	dlspka_	Alignment	not modelled	98.7	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	dlwiea_	Alignment	not modelled	98.7	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	c3rnjA_	Alignment	not modelled	98.7	33	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)
116	dlue9a_	Alignment	not modelled	98.7	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	c2csiA_	Alignment	not modelled	98.7	38	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
118	dlwlpb2	Alignment	not modelled	98.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	dlgbra_	Alignment	not modelled	98.7	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	dlqlya_	Alignment	not modelled	98.7	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain