




















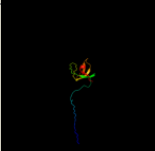







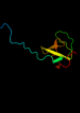



Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	O01498
Date	Fri May 3 21:45:00 BST 2013
Unique Job ID	461520a87ceaff19

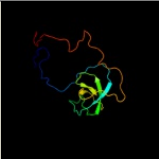
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1dvpA_	 Alignment		100.0	18	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	29	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	26	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	d1dvpal	 Alignment		100.0	24	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
5	d1juqa_	 Alignment		100.0	27	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
6	d1ujka_	 Alignment		100.0	24	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
7	d1elka_	 Alignment		100.0	29	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
8	d1mhqa_	 Alignment		100.0	24	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: VHS domain
9	c3rruA_	 Alignment		100.0	23	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		99.5	17	PDB header: unknown function Chain: A: PDB Molecule: hypothetical protein (raf109-17-b18); PDBTitle: fully automated nmr structure determination of the enth-vhs2 domain at3g16270 from arabidopsis thaliana
11	c1mv3A_	 Alignment		99.1	18	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: myc box dependent interacting protein 1; PDBTitle: nmr structure of the tumor suppressor bin1: alternative2 splicing in melanoma and interaction with c-myc

12	d1hf8a2	Alignment		99.0	20	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: Phosphoinositide-binding clathrin adaptor, N-terminal domain
13	d1inza	Alignment		99.0	16	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: ENTH domain
14	d1hx8a2	Alignment		98.9	16	Fold: alpha-alpha superhelix Superfamily: ENTH/VHS domain Family: Phosphoinositide-binding clathrin adaptor, N-terminal domain
15	c1griA	Alignment		98.8	32	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
16	c2dx1A	Alignment		98.7	35	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
17	c3jv3A	Alignment		98.7	37	PDB header: protein binding Chain: A: PDB Molecule: intersectin-1; PDBTitle: structure of sh3e-dh unit of murine intersectin-1l
18	d1ng2a2	Alignment		98.7	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
19	c3nmzD	Alignment		98.6	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
20	c2eyyA	Alignment		98.6	22	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
21	c2l3sA	Alignment	not modelled	98.6	15	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
22	c2yunA	Alignment		98.6	28	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
23	d1udla	Alignment	not modelled	98.6	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
24	c2nwmA	Alignment	not modelled	98.6	31	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
25	c2eyzA	Alignment	not modelled	98.6	13	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
26	c1wi7A	Alignment	not modelled	98.6	24	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
27	c1ng2A	Alignment	not modelled	98.6	20	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
						PDB header: signaling protein

28	c2l0aA	Alignment	not modelled	98.6	51	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
29	c3zymA	Alignment	not modelled	98.5	20	PDB header: endocytosis Chain: A: PDB Molecule: phosphatidylinositol-binding clathrin assembly protein, PDBTitle: structure of calm (picalm) in complex with vamp8
30	c1x2qA	Alignment	not modelled	98.5	40	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
31	c2o2oA	Alignment	not modelled	98.5	22	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
32	c2ed0A	Alignment	not modelled	98.5	26	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)
33	c2cubA	Alignment	not modelled	98.5	23	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
34	c2yu0A	Alignment	not modelled	98.5	32	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
35	c2jxbA	Alignment	not modelled	98.5	18	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
36	c2dbmA	Alignment	not modelled	98.5	34	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
37	c1z9qA	Alignment	not modelled	98.5	32	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: solution structure of sh3 domain of p40phox
38	c2dl3A	Alignment	not modelled	98.5	25	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
39	c2ysqA	Alignment	not modelled	98.5	36	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
40	c2ed1A	Alignment	not modelled	98.5	37	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	d1k4us	Alignment	not modelled	98.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
42	c2ebpA	Alignment	not modelled	98.5	25	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
43	c2lqnA	Alignment	not modelled	98.5	16	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
44	c2oi3A	Alignment	not modelled	98.5	25	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)
45	d1uj0a	Alignment	not modelled	98.5	54	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
46	c2djqa	Alignment	not modelled	98.5	24	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
47	c1x43A	Alignment	not modelled	98.5	28	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: sh3 domain grb2-like protein b1; PDBTitle: solution structure of the sh3 domain of endophilin b12 (sh3g1b1)
48	c2xmfA	Alignment	not modelled	98.5	44	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
49	d1ov3a2	Alignment	not modelled	98.5	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
50	c4iqzA	Alignment	not modelled	98.5	33	PDB header: protein binding Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 2; PDBTitle: crystal structure of the sh3 domain of human sorbin and sh3 domain-2 containing protein 2
51	c2dilA	Alignment	not modelled	98.5	29	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
52	c2rqrA	Alignment	not modelled	98.5	21	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 PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
53	c2dlpA_	Alignment	not modelled	98.4	29	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
54	c2dl4A_	Alignment	not modelled	98.4	35	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)
55	c2dmoA_	Alignment	not modelled	98.4	23	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
56	c1w70A_	Alignment	not modelled	98.4	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
57	d2hsa_	Alignment	not modelled	98.4	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
58	d1gria2	Alignment	not modelled	98.4	40	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
59	c2bzxA_	Alignment	not modelled	98.4	24	PDB header: sh3 domain Chain: A: PDB Molecule: crk-like protein; PDBTitle: atomic model of crkl-sh3c monomer
60	c2kymA_	Alignment	not modelled	98.4	21	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
61	d1ujya_	Alignment	not modelled	98.4	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
62	c2pqhA_	Alignment	not modelled	98.4	27	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
63	d1uhfa_	Alignment	not modelled	98.4	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
64	c2yt6A_	Alignment	not modelled	98.4	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
65	c2dnuA_	Alignment	not modelled	98.4	18	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
66	c2csgA_	Alignment	not modelled	98.4	22	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
67	c2eyxA_	Alignment	not modelled	98.4	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
68	c2dbkA_	Alignment	not modelled	98.4	22	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
69	d1sema_	Alignment	not modelled	98.4	42	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
70	c2ekhA_	Alignment	not modelled	98.4	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
71	c1x69A_	Alignment	not modelled	98.4	33	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
72	c2drmB_	Alignment	not modelled	98.4	33	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
73	c2df6A_	Alignment	not modelled	98.4	36	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
74	c1x2pA_	Alignment	not modelled	98.4	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
75	c2d1xD_	Alignment	not modelled	98.4	37	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
76	c2eqiA_	Alignment	not modelled	98.4	34	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
77	c2bz8B_	Alignment	not modelled	98.4	32	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
78	c2dl8A_	Alignment	not modelled	98.4	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
						Fold: alpha-alpha superhelix

79	d1saa_	Alignment	not modelled	98.4	14	Superfamily: ENTH/VHS domain Family: RPR domain (SMART 00582)
80	c2egaA_	Alignment	not modelled	98.4	20	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
81	c2yupA_	Alignment	not modelled	98.4	29	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
82	c4glmD_	Alignment	not modelled	98.4	39	PDB header: signaling protein Chain: D: PDB Molecule: dynamin-binding protein; PDBTitle: crystal structure of the sh3 domain of dnmbp protein [homo sapiens]
83	c1wyxA_	Alignment	not modelled	98.4	31	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
84	d1oeba_	Alignment	not modelled	98.4	47	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
85	c2e5kA_	Alignment	not modelled	98.4	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
86	d1wiea_	Alignment	not modelled	98.4	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
87	d1gcqa_	Alignment	not modelled	98.4	45	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
88	d1utia_	Alignment	not modelled	98.4	48	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
89	c1zx6A_	Alignment	not modelled	98.4	28	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
90	d1ue9a_	Alignment	not modelled	98.3	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
91	d1awja_	Alignment	not modelled	98.3	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
92	c2eywA_	Alignment	not modelled	98.3	26	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	c2epdA_	Alignment	not modelled	98.3	24	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
94	c2dviA_	Alignment		98.3	18	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii
95	d1gbra_	Alignment	not modelled	98.3	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
96	c1zlmA_	Alignment	not modelled	98.3	46	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
97	c2krnA_	Alignment	not modelled	98.3	32	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
98	d1j3ta_	Alignment	not modelled	98.3	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
99	d1uuea_	Alignment	not modelled	98.3	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	c2ke9A_	Alignment	not modelled	98.3	37	PDB header: protein binding Chain: A: PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
101	d1ng2a1	Alignment	not modelled	98.3	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
102	d1ov3a1	Alignment	not modelled	98.3	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	c1negA_	Alignment	not modelled	98.3	27	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
104	d1nega_	Alignment	not modelled	98.3	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

105	c3qwyA	Alignment	not modelled	98.3	27	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
106	c3qwxX	Alignment	not modelled	98.3	26	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
107	c2fpeB	Alignment	not modelled	98.3	19	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
108	c2d8jA	Alignment	not modelled	98.3	35	PDB header: transferase Chain: A: PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh3 domain of fyn-related kinase
109	d1phta	Alignment	not modelled	98.3	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
110	c2cudA	Alignment	not modelled	98.3	12	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)
111	c2rqvA	Alignment	not modelled	98.3	25	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
112	d1pwta	Alignment	not modelled	98.3	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	c2ct3A	Alignment	not modelled	98.3	20	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
114	d1u06a1	Alignment	not modelled	98.3	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	c2d8hA	Alignment	not modelled	98.3	29	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
116	d1uffa	Alignment	not modelled	98.3	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	c2lx7A	Alignment	not modelled	98.3	31	PDB header: protein binding Chain: A: PDB Molecule: growth arrest-specific protein 7; PDBTitle: solution nmr structure of sh3 domain of growth arrest-specific protein2 7 (gas7) (fragment 1-60) from homo sapiens, northeast structural3 genomics consortium (nesg) target hr8574a
118	d1awwa	Alignment	not modelled	98.3	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	d1ugva	Alignment	not modelled	98.3	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	d1ycsb2	Alignment	not modelled	98.3	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain