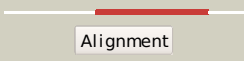

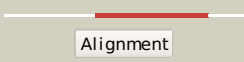

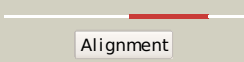
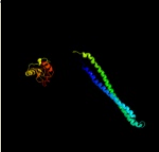




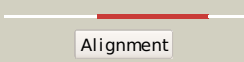

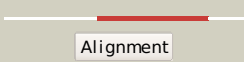

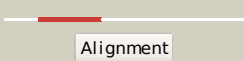
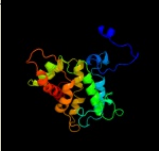
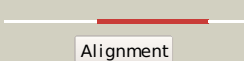

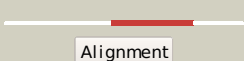
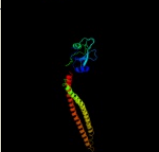
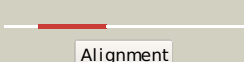



Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	O00459
Date	Fri May 25 09:44:30 BST 2012
Unique Job ID	0ae9a79684a9af6d

Detailed template information


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2oq1A_	 Alignment		100.0	29	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase zap-70; PDBTitle: tandem sh2 domains of zap-70 with 19-mer zeta1 peptide
2	c2ozoA_	 Alignment		100.0	27	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase zap-70; PDBTitle: autoinhibited intact human zap-70
3	c2y3aB_	 Alignment		100.0	100	PDB header: transferase Chain: B: PDB Molecule: phosphatidylinositol 3-kinase regulatory subunit beta; PDBTitle: crystal structure of p110beta in complex with icsh2 of p85beta and2 the drug gdc-0941
4	c2b3oA_	 Alignment		100.0	35	PDB header: hydrolase Chain: A: PDB Molecule: tyrosine-protein phosphatase, non-receptor type PDBTitle: crystal structure of human tyrosine phosphatase shp-1
5	c2shpA_	 Alignment		100.0	30	PDB header: tyrosine phosphatase Chain: A: PDB Molecule: shp-2; PDBTitle: tyrosine phosphatase shp-2
6	c1a81E_	 Alignment		100.0	32	PDB header: complex (transferase/peptide) Chain: E: PDB Molecule: syk kinase; PDBTitle: crystal structure of the tandem sh2 domain of the syk kinase bound to2 a dually tyrosine-phosphorylated itam
7	c3ps5A_	 Alignment		100.0	29	PDB header: hydrolase, signaling protein Chain: A: PDB Molecule: tyrosine-protein phosphatase non-receptor type 6; PDBTitle: crystal structure of the full-length human protein tyrosine2 phosphatase shp-1
8	c1xa6A_	 Alignment		100.0	20	PDB header: signaling protein Chain: A: PDB Molecule: beta2-chimaerin; PDBTitle: crystal structure of the human beta2-chimaerin
9	c3gqiB_	 Alignment		100.0	30	PDB header: transferase/transferase inhibitor Chain: B: PDB Molecule: phospholipase c-gamma-1; PDBTitle: crystal structure of activated receptor tyrosine kinase in complex2 with substrates
10	c3hizB_	 Alignment		100.0	77	PDB header: transferase/oncoprotein Chain: B: PDB Molecule: phosphatidylinositol 3-kinase regulatory subunit PDBTitle: crystal structure of p110alpha h1047r mutant in complex with2 nish2 of p85alpha
11	c2ee5A_	 Alignment		100.0	20	PDB header: hydrolase Chain: A: PDB Molecule: rho gtpase activating protein 5 variant; PDBTitle: solution structure of the n-terminus extended rhogap2 domain from human rho gtpase activating protein 5 variant

12	c2xs6A_	Alignment		100.0	93	PDB header: transferase Chain: A: PDB Molecule: phosphatidylinositol 3-kinase regulatory subunit beta; PDBTitle: crystal structure of the rhogap domain of human pik3r2
13	c3byiA_	Alignment		100.0	21	PDB header: signaling protein Chain: A: PDB Molecule: rho gtpase activating protein 15; PDBTitle: crystal structure of human rho gtpase activating protein 15 (arhgap15)
14	c2ngrB_	Alignment		100.0	21	PDB header: hydrolase Chain: B: PDB Molecule: protein (gtpase activating protein (rhg)); PDBTitle: transition state complex for gtp hydrolysis by cdc42:2 comparisons of the high resolution structures for cdc423 bound to the active and catalytically compromised forms of4 the cdc42-gap.
15	c3qisA_	Alignment		100.0	20	PDB header: hydrolase/protein binding Chain: A: PDB Molecule: inositol polyphosphate 5-phosphatase ocl1-1; PDBTitle: recognition of the f&h motif by the lowe syndrome protein ocl1
16	d1xa6a1	Alignment		100.0	21	Fold: GTPase activation domain, GAP Superfamily: GTPase activation domain, GAP Family: BCR-homology GTPase activation domain (BH-domain)
17	d1bx4a_	Alignment		100.0	21	Fold: GTPase activation domain, GAP Superfamily: GTPase activation domain, GAP Family: BCR-homology GTPase activation domain (BH-domain)
18	c3msxB_	Alignment		100.0	15	PDB header: protein binding Chain: B: PDB Molecule: rho gtpase-activating protein 20; PDBTitle: crystal structure of rhoa.gdp.mgf3 in complex with gap domain of2 arhgap20
19	c3kugA_	Alignment		100.0	15	PDB header: hydrolase activator Chain: A: PDB Molecule: rho gtpase-activating protein 7; PDBTitle: crystal structure of the dlc1 rhogap domain
20	c3iugA_	Alignment		100.0	22	PDB header: splicing Chain: A: PDB Molecule: rho/cdc42/rac gtpase-activating protein rics; PDBTitle: crystal structure of the rhogap domain of rics
21	d1pbwa_	Alignment	not modelled	100.0	35	Fold: GTPase activation domain, GAP Superfamily: GTPase activation domain, GAP Family: BCR-homology GTPase activation domain (BH-domain)
22	c3fk2B_	Alignment	not modelled	100.0	19	PDB header: signaling protein, hydrolase activator Chain: B: PDB Molecule: glucocorticoid receptor dna-binding factor 1; PDBTitle: crystal structure of the rhogap domain of human2 glucocorticoid receptor dna-binding factor 1
23	c1pbwB_	Alignment	not modelled	100.0	35	PDB header: phosphotransferase Chain: B: PDB Molecule: phosphatidylinositol 3-kinase; PDBTitle: structure of bcr-homology (bh) domain
24	d1f7ca_	Alignment	not modelled	100.0	23	Fold: GTPase activation domain, GAP Superfamily: GTPase activation domain, GAP Family: BCR-homology GTPase activation domain (BH-domain)
25	c1f7cA_	Alignment	not modelled	100.0	23	PDB header: signaling protein Chain: A: PDB Molecule: rhogap protein; PDBTitle: crystal structure of the bh domain from graf, the gtpase2 regulator associated with focal adhesion kinase
26	c2oviA_	Alignment	not modelled	100.0	27	PDB header: signaling protein Chain: A: PDB Molecule: rac gtpase-activating protein 1; PDBTitle: the crystal structure of the human rac gtpase activating protein 12 (racgap1) mgcracgap.
27	c2qv2A_	Alignment	not modelled	100.0	22	PDB header: hydrolase Chain: A: PDB Molecule: inositol polyphosphate 5-phosphatase ocl1-1; PDBTitle: a role of the lowe syndrome protein ocl1 in early steps of2 the endocytic pathway
28	c3eapD_	Alignment	not modelled	99.9	20	PDB header: hydrolase activator Chain: D: PDB Molecule: rho gtpase-activating protein 11a; PDBTitle: crystal structure of the rhogap domain of arhgap11a

29	d2oqla1	Alignment	not modelled	99.9	29	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
30	dlpica	Alignment	not modelled	99.9	73	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
31	dlqada	Alignment	not modelled	99.9	79	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
32	dla81a1	Alignment	not modelled	99.9	28	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
33	c2dlyA	Alignment	not modelled	99.9	35	PDB header: transferase Chain: A: PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh2 domain of murine fyn-related2 kinase
34	d2oqla2	Alignment	not modelled	99.9	28	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
35	c2crhA	Alignment	not modelled	99.9	28	PDB header: signaling protein Chain: A: PDB Molecule: vav proto-oncogene; PDBTitle: solution structure of the sh2 domain of human proto-2 oncogene protein vav1
36	dla81e1	Alignment	not modelled	99.9	28	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
37	dla81a2	Alignment	not modelled	99.9	34	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
38	c2dlzA	Alignment	not modelled	99.9	26	PDB header: signaling protein Chain: A: PDB Molecule: protein vav-2; PDBTitle: solution structure of the sh2 domain of human protein vav-2
39	dla81e2	Alignment	not modelled	99.9	33	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
40	c2eo6A	Alignment	not modelled	99.9	29	PDB header: signaling protein Chain: A: PDB Molecule: b-cell linker protein; PDBTitle: solution structure of the sh2 domain from mouse b-cell2 linker protein blk
41	c2fo0A	Alignment	not modelled	99.9	29	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase abl1 (1b isoform); PDBTitle: organization of the sh3-sh2 unit in active and inactive forms of the2 c-abl tyrosine kinase
42	c2ablA	Alignment	not modelled	99.9	34	PDB header: transferase Chain: A: PDB Molecule: abl tyrosine kinase; PDBTitle: sh3-sh2 domain fragment of human bcr-abl tyrosine kinase
43	c2lctA	Alignment	not modelled	99.9	29	PDB header: signaling protein Chain: A: PDB Molecule: proto-oncogene vav; PDBTitle: solution structure of the vav1 sh2 domain complexed with a syk-derived2 doubly phosphorylated peptide
44	c2ge9A	Alignment	not modelled	99.9	29	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase btk; PDBTitle: solution structures of the sh2 domain of bruton's tyrosine2 kinase
45	c2eobA	Alignment	not modelled	99.9	31	PDB header: hydrolase Chain: A: PDB Molecule: 1-phosphatidylinositol-4,5-bisphosphate PDBTitle: solution structure of the second sh2 domain from rat plc2 gamma-2
46	cloplA	Alignment	not modelled	99.9	28	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase; PDBTitle: structural basis for the auto-inhibition of c-abl tyrosine2 kinase
47	dlcsya	Alignment	not modelled	99.9	31	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
48	c2gsbA	Alignment	not modelled	99.9	23	PDB header: signaling protein Chain: A: PDB Molecule: ras gtpase-activating protein 1; PDBTitle: solution structure of the second sh2 domain of human ras2 gtpase-activating protein 1
49	d1opka2	Alignment	not modelled	99.9	32	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
50	d1lkka	Alignment	not modelled	99.9	37	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
51	d1qcfa2	Alignment	not modelled	99.9	37	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
52	d2shpa2	Alignment	not modelled	99.9	27	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
53	d1jwoa	Alignment	not modelled	99.8	27	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
54	d1nrva	Alignment	not modelled	99.8	24	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
55	d1k9aa2	Alignment	not modelled	99.8	33	Fold: SH2-like Superfamily: SH2 domain

						Family: SH2 domain
56	c2ysxA	Alignment	not modelled	99.8	29	PDB header: signaling protein Chain: A; PDB Molecule: signaling inositol polyphosphate phosphatase PDBTitle: solution structure of the human ship sh2 domain
57	c1y57A	Alignment	not modelled	99.8	31	PDB header: transferase Chain: A; PDB Molecule: proto-oncogene tyrosine-protein kinase src; PDBTitle: structure of unphosphorylated c-src in complex with an inhibitor
58	c2knoA	Alignment	not modelled	99.8	32	PDB header: hydrolase Chain: A; PDB Molecule: tensin-like c1 domain-containing phosphatase; PDBTitle: nmr solution structure of sh2 domain of the human tensin like c12 domain containing phosphatase (tenc1)
59	d1o48a	Alignment	not modelled	99.8	34	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
60	d1rjaa	Alignment	not modelled	99.8	33	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
61	c2lqnA	Alignment	not modelled	99.8	26	PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
62	c1k9aB	Alignment	not modelled	99.8	29	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
63	d1bkla	Alignment	not modelled	99.8	34	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
64	c1ka6A	Alignment	not modelled	99.8	29	PDB header: immune system Chain: A; PDB Molecule: sh2 domain protein 1a; PDBTitle: sap/sh2d1a bound to peptide n-py
65	d1luia	Alignment	not modelled	99.8	28	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
66	d1blja	Alignment	not modelled	99.8	32	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
67	d2qmsa1	Alignment	not modelled	99.8	27	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
68	c3nhnA	Alignment	not modelled	99.8	36	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
69	d1fhsa	Alignment	not modelled	99.8	33	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
70	d1g83a2	Alignment	not modelled	99.8	34	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
71	c2eo3A	Alignment	not modelled	99.8	30	PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein; PDBTitle: solution structure of the sh2 domain from human crk-like2 protein
72	d1f2fa	Alignment	not modelled	99.8	34	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
73	d1ayaa	Alignment	not modelled	99.8	28	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
74	c1x27F	Alignment	not modelled	99.8	34	PDB header: signaling protein Chain: F; PDB Molecule: proto-oncogene tyrosine-protein kinase lck; PDBTitle: crystal structure of lck sh2-sh3 with sh2 binding site of f2 p130cas
75	d1i3za	Alignment	not modelled	99.8	32	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
76	c2dm0A	Alignment	not modelled	99.8	29	PDB header: transferase Chain: A; PDB Molecule: tyrosine-protein kinase btk; PDBTitle: solution structure of the sh2 domain of human tyrosine-2 protein kinase btk
77	d1d4ta	Alignment	not modelled	99.8	29	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
78	c1x6cA	Alignment	not modelled	99.8	28	PDB header: signaling protein Chain: A; PDB Molecule: tyrosine-protein phosphatase, non-receptor type PDBTitle: solution structures of the sh2 domain of human protein-2 tyrosine phosphatase shp-1
79	c2vifA	Alignment	not modelled	99.8	35	PDB header: signaling protein Chain: A; PDB Molecule: suppressor of cytokine signalling 6; PDBTitle: crystal structure of socs6 sh2 domain in complex with a c-kit2 phosphopeptide
80	d2izva2	Alignment	not modelled	99.8	26	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
81	d3c7ia1	Alignment	not modelled	99.8	35	Fold: SH2-like Superfamily: SH2 domain

					Family:SH2 domain
82	c2ekxA	Alignment	not modelled	99.8	32 PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic tyrosine-protein kinase bmx; PDBTitle: solution structure of the human bmx sh2 domain
83	d2cs0a1	Alignment	not modelled	99.8	27 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
84	c2h8hA	Alignment	not modelled	99.8	30 PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase src; PDBTitle: src kinase in complex with a quinazoline inhibitor
85	c2c0iA	Alignment	not modelled	99.8	34 PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck; PDBTitle: src family kinase hck with bound inhibitor a-420983
86	c2hmhA	Alignment	not modelled	99.8	26 PDB header: cytokine regulator Chain: A: PDB Molecule: suppressor of cytokine signaling 3; PDBTitle: crystal structure of socs3 in complex with gp130(pyr757)2 phosphopeptide.
87	d1cwea	Alignment	not modelled	99.8	40 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
88	d1fu6a	Alignment	not modelled	99.8	32 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
89	d2eyva1	Alignment	not modelled	99.8	33 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
90	d2shpa3	Alignment	not modelled	99.8	29 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
91	d2fcia1	Alignment	not modelled	99.8	28 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
92	c2dcrA	Alignment	not modelled	99.8	26 PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase fes/fps; PDBTitle: fully automated solution structure determination of the fes2 sh2 domain
93	c3mazA	Alignment	not modelled	99.8	17 PDB header: signaling protein Chain: A: PDB Molecule: signal-transducing adaptor protein 1; PDBTitle: crystal structure of the human brdg1/stap-1 sh2 domain in complex with2 the ntal ptyr136 peptide
94	d1r1qa	Alignment	not modelled	99.8	33 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
95	c1g83A	Alignment	not modelled	99.8	35 PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase fyn; PDBTitle: crystal structure of fyn sh3-sh2
96	c2c9wA	Alignment	not modelled	99.8	27 PDB header: transcription regulation Chain: A: PDB Molecule: suppressor of cytokine signaling 2; PDBTitle: crystal structure of socs-2 in complex with elongin-b and2 elongin-c at 1.9a resolution
97	c2ci8A	Alignment	not modelled	99.8	29 PDB header: translation Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: sh2 domain of human nck1 adaptor protein - uncomplexed
98	d1jyra	Alignment	not modelled	99.8	37 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
99	c3or8A	Alignment	not modelled	99.8	29 PDB header: transcription Chain: A: PDB Molecule: transcription elongation factor spt6; PDBTitle: a tandem sh2 domain in transcription elongation factor spt6
100	c2izvA	Alignment	not modelled	99.8	24 PDB header: transcription Chain: A: PDB Molecule: suppressor of cytokine signaling 4; PDBTitle: crystal structure of socs-4 in complex with elongin-b and2 elongin-c at 2.55a resolution
101	d1mila	Alignment	not modelled	99.7	25 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
102	c2el8A	Alignment	not modelled	99.7	18 PDB header: signaling protein Chain: A: PDB Molecule: signal-transducing adaptor protein 2; PDBTitle: solution structure of the human stap2 sh2 domain
103	c2kk6A	Alignment	not modelled	99.7	30 PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase fer; PDBTitle: solution structure of sh2 domain of proto-oncogene tyrosine-2 protein kinase fer from homo sapiens, northeast structural3 genomics consortium (nesg) target hr3461d
104	d2c9wa2	Alignment	not modelled	99.7	25 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
105	c2cr4A	Alignment	not modelled	99.7	24 PDB header: signaling protein Chain: A: PDB Molecule: sh3 domain-binding protein 2; PDBTitle: solution structure of the sh2 domain of human sh3bp2 protein
106	d1xa6a2	Alignment	not modelled	99.7	20 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
107	c2dx0B	Alignment	not modelled	99.7	36 PDB header: hydrolase Chain: B: PDB Molecule: phospholipase c, gamma 2; PDBTitle: crystal structure of the n-terminal sh2 domain of mouse2

						phospholipase c-gamma 2 PDB header: transcription Chain: B; PDB Molecule: transcription elongation factor spt6; PDBTitle: structure of the sh2 domain of the candida glabrata2 transcription elongation factor spt6, crystal form b
108	c3gxxB_	Alignment	not modelled	99.7	25	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
109	d1rpya_	Alignment	not modelled	99.7	20	PDB header: transcription Chain: A; PDB Molecule: spt6; PDBTitle: structure of the tandem sh2 domains from antonospora locustae2 transcription elongation factor spt6
110	c2xp1A_	Alignment	not modelled	99.7	18	PDB header: transferase/signaling protein Chain: C; PDB Molecule: adaptor protein aps; PDBTitle: crystal structure of the insulin receptor kinase in complex2 with the sh2 domain of aps
111	c1rqcC_	Alignment	not modelled	99.7	22	PDB header: signaling protein Chain: A; PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
112	c2eyzA_	Alignment	not modelled	99.7	30	PDB header: signaling protein Chain: X; PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
113	c3qwxX_	Alignment	not modelled	99.7	25	PDB header: signaling protein Chain: A; PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii
114	c2dviA_	Alignment	not modelled	99.7	36	PDB header: signaling protein Chain: A; PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
115	c2eyyA_	Alignment	not modelled	99.7	29	PDB header: signaling protein Chain: A; PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
116	c3qwyA_	Alignment	not modelled	99.6	25	PDB header: signaling protein Chain: A; PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
117	c2v1yB_	Alignment	not modelled	99.5	73	PDB header: transferase Chain: B; PDB Molecule: phosphatidylinositol 3-kinase regulatory subunit alpha; PDBTitle: structure of a phosphoinositide 3-kinase alpha adaptor-2 binding domain (abd) in a complex with the ish2 domain3 from p85 alpha
118	d1uura3	Alignment	not modelled	99.4	18	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
119	c2kt1A_	Alignment		99.4	100	PDB header: protein binding Chain: A; PDB Molecule: phosphatidylinositol 3-kinase regulatory subunit PDBTitle: solution nmr structure of the sh3 domain from the p85beta2 subunit of phosphatidylinositol 3-kinase from h.sapiens,3 northeast structural genomics consortium target hr5531e
120	d1phta_	Alignment	not modelled	99.3	51	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain