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Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3ps5A_	 Alignment		100.0	59	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
2	c2b3oA_	 Alignment		100.0	60	PDB header: hydrolase Chain: A; PDB Molecule: tyrosine-protein phosphatase, non-receptor type PDBTitle: crystal structure of human tyrosine phosphatase shp-1
3	c2shpA_	 Alignment		100.0	100	PDB header: tyrosine phosphatase Chain: A; PDB Molecule: shp-2; PDBTitle: tyrosine phosphatase shp-2
4	c1larB_	 Alignment		100.0	35	PDB header: hydrolase Chain: B; PDB Molecule: protein (lar); PDBTitle: crystal structure of the tandem phosphatase domains of rtp2 lar
5	c2nlkA_	 Alignment		100.0	42	PDB header: hydrolase Chain: A; PDB Molecule: protein tyrosine phosphatase, receptor type, g variant PDBTitle: crystal structure of d1 and d2 catalytic domains of human protein2 tyrosine phosphatase gamma (d1+d2 ptprg)
6	c1yguA_	 Alignment		100.0	31	PDB header: hydrolase Chain: A; PDB Molecule: leukocyte common antigen; PDBTitle: crystal structure of the tandem phosphatase domains of rtp2 cd45 with a ptyr peptide
7	d1wcha_	 Alignment		100.0	39	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Higher-molecular-weight phosphotyrosine protein phosphatases
8	d1jlna_	 Alignment		100.0	40	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Higher-molecular-weight phosphotyrosine protein phosphatases
9	d2shpa1	 Alignment		100.0	100	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Higher-molecular-weight phosphotyrosine protein phosphatases
10	d1fpra_	 Alignment		100.0	60	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Higher-molecular-weight phosphotyrosine protein phosphatases
11	c2nz6A_	 Alignment		100.0	37	PDB header: hydrolase Chain: A; PDB Molecule: receptor-type tyrosine-protein phosphatase eta; PDBTitle: crystal structure of the ptpri inactivating mutant c1239s

12	c3s3fA	Alignment		100.0	37	PDB header: hydrolase/hydrolase inhibitor Chain: A: PDB Molecule: tyrosine-protein phosphatase 10d; PDBTitle: crystal structure of the catalytic domain of ptp10d from drosophila2 melanogaster with a small molecule inhibitor vanadate
13	c3i36A	Alignment		100.0	40	PDB header: hydrolase Chain: A: PDB Molecule: vascular protein tyrosine phosphatase 1; PDBTitle: crystal structure of rat protein tyrosine phosphatase eta catalytic2 domain
14	c2pa5A	Alignment		100.0	37	PDB header: hydrolase Chain: A: PDB Molecule: tyrosine-protein phosphatase non-receptor type 9; PDBTitle: crystal structure of human protein tyrosine phosphatase ptpn9
15	d1lara1	Alignment		100.0	38	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Higher-molecular-weight phosphotyrosine protein phosphatases
16	c2jjdA	Alignment		100.0	32	PDB header: hydrolase Chain: A: PDB Molecule: receptor-type tyrosine-protein phosphatase epsilon; PDBTitle: protein tyrosine phosphatase, receptor type, e isoform
17	d2f71a1	Alignment		100.0	35	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Higher-molecular-weight phosphotyrosine protein phosphatases
18	c2vexA	Alignment		100.0	35	PDB header: hydrolase Chain: A: PDB Molecule: tyrosine-protein phosphatase non-receptor type 1; PDBTitle: crystal structure of protein tyrosine phosphatase 1b in2 complex with an isothiazolidinone-containing inhibitor
19	c2bzlA	Alignment		100.0	36	PDB header: hydrolase Chain: A: PDB Molecule: tyrosine-protein phosphatase, non-receptor type PDBTitle: crystal structure of the human protein tyrosine phosphatase2 n14 at 1.65 a resolution
20	c2c7sA	Alignment		100.0	42	PDB header: hydrolase Chain: A: PDB Molecule: receptor-type tyrosine-protein phosphatase kappa; PDBTitle: crystal structure of human protein tyrosine phosphatase2 kappa at 1.95a resolution
21	c2hy3A	Alignment	not modelled	100.0	43	PDB header: hydrolase Chain: A: PDB Molecule: receptor-type tyrosine-protein phosphatase gamma; PDBTitle: crystal structure of the human tyrosine receptor phosphate gamma in2 complex with vanadate
22	c2qdmA	Alignment	not modelled	100.0	38	PDB header: hydrolase Chain: A: PDB Molecule: tyrosine-protein phosphatase non-receptor type 7; PDBTitle: crystal structure of the hept catalytic domain c270s/d236a/q314a2 mutant
23	d1rpma	Alignment	not modelled	100.0	41	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Higher-molecular-weight phosphotyrosine protein phosphatases
24	c2oc3A	Alignment	not modelled	100.0	37	PDB header: hydrolase Chain: A: PDB Molecule: tyrosine-protein phosphatase non-receptor type PDBTitle: crystal structure of the catalytic domain of human protein2 tyrosine phosphatase non-receptor type 18
25	c2h04A	Alignment	not modelled	100.0	38	PDB header: hydrolase Chain: A: PDB Molecule: protein tyrosine phosphatase, receptor type, b,; PDBTitle: structural studies of protein tyrosine phosphatase beta2 catalytic domain in complex with inhibitors
26	d1yfoa	Alignment	not modelled	100.0	43	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Higher-molecular-weight phosphotyrosine protein phosphatases
27	c2qcjB	Alignment	not modelled	100.0	38	PDB header: hydrolase Chain: B: PDB Molecule: tyrosine-protein phosphatase non-receptor type PDBTitle: native structure of lyp

28	c3jrlA	Alignment	not modelled	100.0	100	PDB header: hydrolase Chain: A: PDB Molecule: oncogenic tyrosine phosphatase shp2; PDBTitle: crystal structure of the oncogenic tyrosine phosphatase shp2 complexed2 with a salicylic acid-based small molecule inhibitor
29	c1l8kA	Alignment	not modelled	100.0	37	PDB header: hydrolase Chain: A: PDB Molecule: t-cell protein-tyrosine phosphatase; PDBTitle: t cell protein-tyrosine phosphatase structure
30	d1l8ka	Alignment	not modelled	100.0	37	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Higher-molecular-weight phosphotyrosine protein phosphatases
31	c2i75A	Alignment	not modelled	100.0	36	PDB header: hydrolase Chain: A: PDB Molecule: tyrosine-protein phosphatase non-receptor type 4; PDBTitle: crystal structure of human protein tyrosine phosphatase n4 (ptpn4)
32	c2bijA	Alignment	not modelled	100.0	40	PDB header: hydrolase Chain: A: PDB Molecule: tyrosine-protein phosphatase, non-receptor type 5; PDBTitle: crystal structure of the human protein tyrosine phosphatase2 ptpn5 (step, striatum enriched enriched phosphatase)
33	d1larb1	Alignment	not modelled	100.0	42	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Higher-molecular-weight phosphotyrosine protein phosphatases
34	c2b49A	Alignment	not modelled	100.0	37	PDB header: hydrolase Chain: A: PDB Molecule: protein tyrosine phosphatase, non-receptor type 3; PDBTitle: crystal structure of the catalytic domain of protein tyrosine2 phosphatase, non-receptor type 3
35	c2g59B	Alignment	not modelled	100.0	36	PDB header: hydrolase Chain: B: PDB Molecule: receptor-type tyrosine-protein phosphatase o; PDBTitle: crystal structure of the catalytic domain of protein2 tyrosine phosphatase from homo sapiens
36	c2qepA	Alignment	not modelled	100.0	38	PDB header: hydrolase Chain: A: PDB Molecule: receptor-type tyrosine-protein phosphatase n2; PDBTitle: crystal structure of the d1 domain of ptpn2 (ia2beta)
37	c1g4wR	Alignment	not modelled	100.0	15	PDB header: signaling protein Chain: R: PDB Molecule: protein tyrosine phosphatase sptp; PDBTitle: crystal structure of the salmonella tyrosine phosphatase2 and gtpase activating protein sptp
38	d1lara2	Alignment	not modelled	100.0	41	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Higher-molecular-weight phosphotyrosine protein phosphatases
39	c3m4uB	Alignment	not modelled	100.0	32	PDB header: hydrolase Chain: B: PDB Molecule: tyrosine specific protein phosphatase, putative; PDBTitle: crystal structure of trypanosoma brucei protein tyrosine phosphatase2 tbtp1
40	d1p15a	Alignment	not modelled	100.0	40	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Higher-molecular-weight phosphotyrosine protein phosphatases
41	d1lyva	Alignment	not modelled	100.0	20	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Higher-molecular-weight phosphotyrosine protein phosphatases
42	d1g4us2	Alignment	not modelled	100.0	19	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Higher-molecular-weight phosphotyrosine protein phosphatases
43	c3gqiB	Alignment	not modelled	100.0	33	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
44	c2oq1A	Alignment	not modelled	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
45	c2ozoA	Alignment	not modelled	100.0	30	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase zap-70; PDBTitle: autoinhibited intact human zap-70
46	c1a81E	Alignment	not modelled	100.0	31	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
47	c3or8A	Alignment	not modelled	100.0	17	PDB header: transcription Chain: A: PDB Molecule: transcription elongation factor spt6; PDBTitle: a tandem sh2 domain in transcription elongation factor spt6
48	c2xp1A	Alignment	not modelled	100.0	23	PDB header: transcription Chain: A: PDB Molecule: spt6; PDBTitle: structure of the tandem sh2 domains from antonospora locustae2 transcription elongation factor spt6
49	d1v3aa	Alignment	not modelled	100.0	24	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Dual specificity phosphatase-like
50	d1rxda	Alignment	not modelled	100.0	28	Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Dual specificity phosphatase-like
51	c3rz2B	Alignment	not modelled	99.9	25	PDB header: hydrolase Chain: B: PDB Molecule: protein tyrosine phosphatase type iva 1; PDBTitle: crystal of pri-1 complexed with peptide
52	d2shpa2	Alignment	not modelled	99.9	99	Fold: SH2-like Superfamily: SH2 domain

						Family: SH2 domain
53	c1x6cA	Alignment	not modelled	99.9	50	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
54	c3s4oB	Alignment	not modelled	99.9	18	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: protein tyrosine phosphatase-like protein; PDBTitle: protein tyrosine phosphatase (putative) from Leishmania major
55	d2oq1a1	Alignment	not modelled	99.9	25	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
56	c2ci8A	Alignment	not modelled	99.9	36	PDB header: translation Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: sh2 domain of human nck1 adaptor protein - uncomplexed
57	d3c7ia1	Alignment	not modelled	99.9	28	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
58	d2shpa3	Alignment	not modelled	99.9	100	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
59	d1ayaa	Alignment	not modelled	99.9	100	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
60	c3hizB	Alignment	not modelled	99.9	27	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
61	c2eobA	Alignment	not modelled	99.9	28	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
62	c2y3aB	Alignment	not modelled	99.9	26	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
63	c2lqnA	Alignment	not modelled	99.9	23	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
64	d1a81a1	Alignment	not modelled	99.9	29	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
65	d1nrva	Alignment	not modelled	99.9	30	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
66	d2oq1a2	Alignment	not modelled	99.9	32	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
67	d1fhsa	Alignment	not modelled	99.9	31	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
68	c2ablA	Alignment	not modelled	99.9	30	PDB header: transferase Chain: A: PDB Molecule: abl tyrosine kinase; PDBTitle: sh3-sh2 domain fragment of human bcr-abl tyrosine kinase
69	c2fo0A	Alignment	not modelled	99.9	28	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
70	d1rjaa	Alignment	not modelled	99.9	32	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
71	d1a81e1	Alignment	not modelled	99.9	28	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
72	c2gsbA	Alignment	not modelled	99.9	29	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
73	c2crhA	Alignment	not modelled	99.9	27	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
74	c2eo6A	Alignment	not modelled	99.9	26	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
75	d1a81a2	Alignment	not modelled	99.9	28	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
76	c2dlzA	Alignment	not modelled	99.9	29	PDB header: signaling protein Chain: A: PDB Molecule: protein vav-2; PDBTitle: solution structure of the sh2 domain of human protein vav-2
77	c2h8hA	Alignment	not modelled	99.9	28	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase src; PDBTitle: src kinase in complex with a quinazoline inhibitor

78	c3nhnA	Alignment	not modelled	99.9	26	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
79	c1k9aB	Alignment	not modelled	99.9	28	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
80	c2dlyA	Alignment	not modelled	99.9	31	PDB header: transferase Chain: A; PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh2 domain of murine fyn-related2 kinase
81	d1lkka	Alignment	not modelled	99.9	34	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
82	c1y57A	Alignment	not modelled	99.9	29	PDB header: transferase Chain: A; PDB Molecule: proto-oncogene tyrosine-protein kinase src; PDBTitle: structure of unphosphorylated c-src in complex with an inhibitor
83	c2ysxA	Alignment	not modelled	99.9	31	PDB header: signaling protein Chain: A; PDB Molecule: signaling inositol polyphosphate phosphatase PDBTitle: solution structure of the human ship sh2 domain
84	d1a81e2	Alignment	not modelled	99.9	28	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
85	d1pica	Alignment	not modelled	99.9	29	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
86	c2ge9A	Alignment	not modelled	99.9	27	PDB header: transferase Chain: A; PDB Molecule: tyrosine-protein kinase btk; PDBTitle: solution structures of the sh2 domain of bruton's tyrosine2 kinase
87	c2lctA	Alignment	not modelled	99.9	27	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
88	c1g83A	Alignment	not modelled	99.9	29	PDB header: transferase Chain: A; PDB Molecule: proto-oncogene tyrosine-protein kinase fyn; PDBTitle: crystal structure of fyn sh3-sh2
89	c2eo3A	Alignment	not modelled	99.8	28	PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein; PDBTitle: solution structure of the sh2 domain from human crk-like2 protein
90	c1x27F	Alignment	not modelled	99.8	31	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 p130cas
91	c1oplA	Alignment	not modelled	99.8	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
92	c2el8A	Alignment	not modelled	99.8	22	PDB header: signaling protein Chain: A; PDB Molecule: signal-transducing adaptor protein 2; PDBTitle: solution structure of the human stap2 sh2 domain
93	d1d4ta	Alignment	not modelled	99.8	26	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
94	d1jwoa	Alignment	not modelled	99.8	28	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
95	c2c0iA	Alignment	not modelled	99.8	28	PDB header: transferase Chain: A; PDB Molecule: tyrosine-protein kinase hck; PDBTitle: src family kinase hck with bound inhibitor a-420983
96	d1blja	Alignment	not modelled	99.8	35	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
97	c2kk6A	Alignment	not modelled	99.8	29	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
98	c1ka6A	Alignment	not modelled	99.8	26	PDB header: immune system Chain: A; PDB Molecule: sh2 domain protein 1a; PDBTitle: sap/sh2d1a bound to peptide n-py
99	d1jyra	Alignment	not modelled	99.8	36	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
100	c2vifA	Alignment	not modelled	99.8	25	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
101	d1qada	Alignment	not modelled	99.8	30	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
102	d2izva2	Alignment	not modelled	99.8	18	Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
103	d1bkla	Alignment	not modelled	99.8	31	Fold: SH2-like Superfamily: SH2 domain

					Family: SH2 domain
104	d1k9aa2	Alignment	not modelled	99.8	30 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
105	d2qmsa1	Alignment	not modelled	99.8	32 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
106	d1qcfa2	Alignment	not modelled	99.8	33 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
107	d1opka2	Alignment	not modelled	99.8	31 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
108	d1o48a_	Alignment	not modelled	99.8	31 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
109	d1r1qa_	Alignment	not modelled	99.8	36 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
110	d1f2fa_	Alignment	not modelled	99.8	32 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
111	d1csya_	Alignment	not modelled	99.8	27 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
112	c2knoA	Alignment	not modelled	99.8	24 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)
113	c2dcrA	Alignment	not modelled	99.8	29 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
114	d1g83a2	Alignment	not modelled	99.8	31 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
115	d1ohea2	Alignment	not modelled	99.8	28 Fold: (Phosphotyrosine protein) phosphatases II Superfamily: (Phosphotyrosine protein) phosphatases II Family: Dual specificity phosphatase-like
116	d1xa6a2	Alignment	not modelled	99.8	24 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
117	d1luia_	Alignment	not modelled	99.8	29 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
118	d1mila_	Alignment	not modelled	99.8	26 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
119	d1i3za_	Alignment	not modelled	99.8	33 Fold: SH2-like Superfamily: SH2 domain Family: SH2 domain
120	c2c9wA	Alignment	not modelled	99.8	26 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