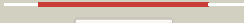
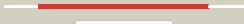
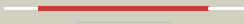


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3jr1A_	 Alignment		100.0	40	PDB header: transferase Chain: A: PDB Molecule: putative fructosamine-3-kinase; PDBTitle: crystal structure of putative fructosamine-3-kinase2 (yp_719053.1) from haemophilus somnus 129pt at 2.32 a3 resolution
2	c3f7wA_	 Alignment		100.0	31	PDB header: transferase Chain: A: PDB Molecule: putative fructosamine-3-kinase; PDBTitle: crystal structure of putative fructosamine-3-kinase (yp_290396.1) from2 thermobifida fusca yx-er1 at 1.85 a resolution
3	c3ovcA_	 Alignment		100.0	13	PDB header: transferase/antibiotic Chain: A: PDB Molecule: hygromycin-b 4-o-kinase; PDBTitle: crystal structure of aminoglycoside phosphotransferase aph(4)-ia
4	c3lzhA_	 Alignment		100.0	14	PDB header: transferase Chain: A: PDB Molecule: aph(2'')-id/aph(2'')-iva; PDBTitle: crystal structure of aminoglycoside phosphotransferase aph(2'')-2 id/aph(2'')-iva
5	c3dxaA_	 Alignment		100.0	15	PDB header: transferase Chain: A: PDB Molecule: putative acyl-coa dehydrogenase; PDBTitle: crystal structure of a putative aminoglycoside phosphotransferase2 (reut_a1007) from ralstonia eutropha jmp134 at 2.32 a resolution
6	c3hamA_	 Alignment		100.0	12	PDB header: transferase Chain: A: PDB Molecule: aminoglycoside phosphotransferase; PDBTitle: structure of the gentamicin-aph(2'')-iia complex
7	d2ppqa1	 Alignment		100.0	14	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: APH phosphotransferases
8	dlzylal	 Alignment		100.0	14	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: APH phosphotransferases
9	c3attA_	 Alignment		100.0	19	PDB header: transferase Chain: A: PDB Molecule: putative uncharacterized protein; PDBTitle: crystal structure of rv3168 with atp
10	c3i0oA_	 Alignment		100.0	13	PDB header: transferase Chain: A: PDB Molecule: spectinomycin phosphotransferase; PDBTitle: crystal structure of spectinomycin phosphotransferase,2 aph(9)-ia, in complex with adp and spectinomycin
11	dlj7la_	 Alignment		100.0	14	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: APH phosphotransferases

12	d1nd4a_	Alignment		100.0	17	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: APH phosphotransferases
13	d2pula1	Alignment		99.9	15	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: APH phosphotransferases
14	c3mesB_	Alignment		99.9	11	PDB header: transferase Chain: B: PDB Molecule: choline kinase; PDBTitle: crystal structure of choline kinase from cryptosporidium2 parvum iowa ii, cgd3_2030
15	c2pywA_	Alignment		99.9	15	PDB header: transferase Chain: A: PDB Molecule: uncharacterized protein; PDBTitle: structure of a. thaliana 5-methylthioribose kinase in complex with adp2 and mtr
16	c3r78B_	Alignment		99.9	13	PDB header: transferase Chain: B: PDB Molecule: aminoglycoside 3'-phosphotransferase aphA1-iab; PDBTitle: crystal structure of the aminoglycoside phosphotransferase aph(3')-ia,2 atp-bound
17	c3dxqB_	Alignment		99.9	15	PDB header: transferase Chain: B: PDB Molecule: choline/ethanolamine kinase family protein; PDBTitle: crystal structure of choline/ethanolamine kinase family protein2 (np_106042.1) from mesorhizobium loti at 2.55 a resolution
18	c3c5iD_	Alignment		99.9	11	PDB header: transferase Chain: D: PDB Molecule: choline kinase; PDBTitle: crystal structure of plasmodium knowlesi choline kinase, pkh_134520
19	c2q83A_	Alignment		99.9	15	PDB header: transferase Chain: A: PDB Molecule: ytAA protein; PDBTitle: crystal structure of ytAA (2635576) from bacillus subtilis at 2.50 a2 resolution
20	c2ig7A_	Alignment		99.9	16	PDB header: transferase Chain: A: PDB Molecule: choline/ethanolamine kinase; PDBTitle: crystal structure of human choline kinase b
21	c1nw1A_	Alignment	not modelled	99.9	15	PDB header: transferase Chain: A: PDB Molecule: choline kinase (49.2 kd); PDBTitle: crystal structure of choline kinase
22	d1nw1a_	Alignment	not modelled	99.9	15	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Choline kinase
23	c3f2rA_	Alignment	not modelled	99.9	13	PDB header: transferase Chain: A: PDB Molecule: choline kinase alpha; PDBTitle: crystal structure of human choline kinase alpha in complex2 with hemicholinium-3
24	c3csvA_	Alignment	not modelled	99.9	16	PDB header: transferase Chain: A: PDB Molecule: aminoglycoside phosphotransferase; PDBTitle: crystal structure of a putative aminoglycoside phosphotransferase2 (yp_614837.1) from silicibacter sp. tm1040 at 2.15 a resolution
25	c2qg7A_	Alignment	not modelled	99.8	10	PDB header: transferase Chain: A: PDB Molecule: ethanolamine kinase pv091845; PDBTitle: plasmodium vivax ethanolamine kinase pv091845
26	c3en9B_	Alignment	not modelled	99.3	22	PDB header: hydrolase Chain: B: PDB Molecule: o-sialoglycoprotein endopeptidase/protein kinase; PDBTitle: structure of the methanococcus jannaschii kae1-bud32 fusion2 protein
27	c2pziA_	Alignment	not modelled	99.1	17	PDB header: transferase Chain: A: PDB Molecule: probable serine/threonine-protein kinase pkng; PDBTitle: crystal structure of protein kinase pkng from mycobacterium2 tuberculosis in complex with tetrahydrobenzothiophene ax20017
						PDB header: ribosome

28	c1tqmA	Alignment	not modelled	98.9	21	Chain: A: PDB Molecule: conserved hypothetical protein; PDBTitle: crystal structure of a. fulgidus rio2 serine protein kinase bound to2 amppnp
29	dlzara2	Alignment	not modelled	98.9	23	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: RIO1-like kinases
30	c3e7eA	Alignment	not modelled	98.8	20	PDB header: transferase Chain: A: PDB Molecule: mitotic checkpoint serine/threonine-protein kinase bub1; PDBTitle: structure and substrate recruitment of the human spindle checkpoint2 kinase bub
31	c2vuwA	Alignment	not modelled	98.7	18	PDB header: transferase Chain: A: PDB Molecule: serine/threonine-protein kinase haspin; PDBTitle: structure of human haspin kinase domain
32	c3qa8A	Alignment	not modelled	98.6	20	PDB header: immune system, signaling protein Chain: A: PDB Molecule: mgc80376 protein; PDBTitle: crystal structure of inhibitor of kappa b kinase beta
33	c1ym7C	Alignment	not modelled	98.5	16	PDB header: transferase Chain: C: PDB Molecule: beta-adrenergic receptor kinase 1; PDBTitle: g protein-coupled receptor kinase 2 (grk2)
34	c1zp9A	Alignment	not modelled	98.5	19	PDB header: transferase Chain: A: PDB Molecule: rio1 kinase; PDBTitle: crystal structure of full-length a.fulgidus rio1 serine kinase bound2 to atp and mn2+ ions.
35	c3c4yA	Alignment	not modelled	98.4	19	PDB header: transferase Chain: A: PDB Molecule: rhodopsin kinase; PDBTitle: crystal structure of apo form of g protein coupled receptor kinase 12 at 7.51a
36	c3ckxA	Alignment	not modelled	98.4	20	PDB header: transferase Chain: A: PDB Molecule: serine/threonine-protein kinase 24; PDBTitle: crystal structure of sterile 20-like kinase 3 (mst3, stk24)2 in complex with staurosporine
37	c2acxB	Alignment	not modelled	98.4	18	PDB header: transferase Chain: B: PDB Molecule: g protein-coupled receptor kinase 6; PDBTitle: crystal structure of g protein coupled receptor kinase 6 bound to2 amppnp
38	c2fo0A	Alignment	not modelled	98.4	15	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
39	c3pfqA	Alignment	not modelled	98.3	18	PDB header: transferase Chain: A: PDB Molecule: protein kinase c beta type; PDBTitle: crystal structure and allosteric activation of protein kinase c beta2 ii
40	c2vwiC	Alignment	not modelled	98.3	16	PDB header: transferase Chain: C: PDB Molecule: serine/threonine-protein kinase osr1; PDBTitle: structure of the osr1 kinase, a hypertension drug target
41	c1oplA	Alignment	not modelled	98.2	15	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase; PDBTitle: structural basis for the auto-inhibition of c-abl tyrosine2 kinase
42	c1k9aB	Alignment	not modelled	98.2	21	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
43	c3fbvL	Alignment	not modelled	98.2	15	PDB header: transferase, hydrolase Chain: L: PDB Molecule: serine/threonine-protein kinase/endoribonuclease ire1; PDBTitle: crystal structure of the oligomer formed by the kinase-ribonuclease2 domain of ire1
44	c3q60A	Alignment	not modelled	98.2	15	PDB header: transferase Chain: A: PDB Molecule: rop5b; PDBTitle: crystal structure of virulent allele rop5b pseudokinase domain bound2 to atp
45	c3p23B	Alignment	not modelled	98.2	19	PDB header: hydrolase, transferase Chain: B: PDB Molecule: serine/threonine-protein kinase/endoribonuclease ire1; PDBTitle: crystal structure of the human kinase and rnase domains in complex2 with adp
46	c2c30A	Alignment	not modelled	98.1	19	PDB header: transferase Chain: A: PDB Molecule: serine/threonine-protein kinase pak 6; PDBTitle: crystal structure of the human p21-activated kinase 6
47	c3d9vA	Alignment	not modelled	98.1	19	PDB header: transferase Chain: A: PDB Molecule: rho-associated protein kinase 1; PDBTitle: crystal structure of rock i bound to h-1152p a di-2 methylated variant of fasudil
48	c3cokB	Alignment	not modelled	98.1	17	PDB header: transferase Chain: B: PDB Molecule: serine/threonine-protein kinase plk4; PDBTitle: crystal structure of plk4 kinase
49	dlqcfa3	Alignment	not modelled	98.0	19	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
50	c2j0kB	Alignment	not modelled	98.0	12	PDB header: transferase Chain: B: PDB Molecule: focal adhesion kinase 1; PDBTitle: crystal structure of a fragment of focal adhesion kinase2 containing the ferm and kinase domains.
51	c3dzoA	Alignment	not modelled	98.0	20	PDB header: transferase Chain: A: PDB Molecule: rhoptry kinase domain; PDBTitle: crystal structure of a rhoptry kinase from toxoplasma gondii
52	dlolja	Alignment	not modelled	98.0	16	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
53	c2clqA	Alignment	not modelled	98.0	17	PDB header: transferase Chain: A: PDB Molecule: mitogen-activated protein kinase kinase kinase 5;

					PDBTitle: structure of mitogen-activated protein kinase kinase kinase2 5
54	dliasa_	Alignment	not modelled	97.9	16 Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
55	c2wu6A_	Alignment	not modelled	97.9	13 PDB header: transferase Chain: A: PDB Molecule: dual specificity protein kinase clk3; PDBTitle: crystal structure of the human clk3 in complex with dki
56	c2wtkC_	Alignment	not modelled	97.9	17 PDB header: transferase/metal-binding protein Chain: C: PDB Molecule: serine/threonine-protein kinase 11; PDBTitle: structure of the heterotrimeric lkb1-stradalpha-mo25alpha2 complex
57	c2j51A_	Alignment	not modelled	97.9	17 PDB header: transferase Chain: A: PDB Molecule: ste20-like serine/threonine-protein kinase; PDBTitle: crystal structure of human ste20-like kinase bound to 5-2 amino-3-((4-(aminosulfonyl)phenyl)amino)-n-(2,6-3 difluorophenyl)-1h-1,2,4-triazole-1-carbothioamide
58	c3soaA_	Alignment	not modelled	97.9	21 PDB header: transferase/transferase inhibitor Chain: A: PDB Molecule: calcium/calmodulin-dependent protein kinase type ii subunit PDBTitle: full-length human camkii
59	c2bujA_	Alignment	not modelled	97.9	18 PDB header: transferase Chain: A: PDB Molecule: serine/threonine-protein kinase 16; PDBTitle: crystal structure of the human serine-threonine kinase 162 in complex with staurosporine
60	d2j4za1	Alignment	not modelled	97.9	20 Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
61	c2c0iA_	Alignment	not modelled	97.9	17 PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck; PDBTitle: src family kinase hck with bound inhibitor a-420983
62	c2h8hA_	Alignment	not modelled	97.9	20 PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase src; PDBTitle: src kinase in complex with a quinazoline inhibitor
63	d1yhwa1	Alignment	not modelled	97.8	21 Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
64	c2jiiA_	Alignment	not modelled	97.8	14 PDB header: transferase Chain: A: PDB Molecule: serine/threonine-protein kinase vrk3 molecule: vaccinia PDBTitle: structure of vaccinia related kinase 3
65	c2y7jB_	Alignment	not modelled	97.8	20 PDB header: transferase Chain: B: PDB Molecule: phosphorylase b kinase gamma catalytic chain, PDBTitle: structure of human phosphorylase kinase, gamma 2
66	c2ozoA_	Alignment	not modelled	97.8	22 PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase zap-70; PDBTitle: autoinhibited intact human zap-70
67	c3lItA_	Alignment	not modelled	97.8	16 PDB header: transferase Chain: A: PDB Molecule: serine/threonine kinase-1, pflammer; PDBTitle: crystal structure of pf14_0431, kinase domain.
68	c3dfaA_	Alignment	not modelled	97.8	19 PDB header: transferase Chain: A: PDB Molecule: calcium-dependent protein kinase cgd3_920; PDBTitle: crystal structure of kinase domain of calcium-dependent protein kinase2 cgd3_920 from cryptosporidium parvum
69	c3q5iA_	Alignment	not modelled	97.8	18 PDB header: transferase Chain: A: PDB Molecule: protein kinase; PDBTitle: crystal structure of pbanka_031420
70	c2cgvA_	Alignment	not modelled	97.7	15 PDB header: transferase Chain: A: PDB Molecule: serine/threonine-protein kinase chk1; PDBTitle: identification of chemically diverse chk1 inhibitors by2 receptor-based virtual screening
71	c2a1aB_	Alignment	not modelled	97.7	17 PDB header: protein synthesis/transferase Chain: B: PDB Molecule: interferon-induced, double-stranded rna-activated protein PDBTitle: pkr kinase domain-eif2alpha complex
72	c1y57A_	Alignment	not modelled	97.7	21 PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase src; PDBTitle: structure of unphosphorylated c-src in complex with an inhibitor
73	c3kvwA_	Alignment	not modelled	97.7	21 PDB header: transferase Chain: A: PDB Molecule: dual specificity tyrosine-phosphorylation-regulated kinase PDBTitle: crystal structure of dual-specificity tyrosine phosphorylation2 regulated kinase 2 (dyrk2) in complex with an indirubin ligand
74	c2wtkB_	Alignment	not modelled	97.7	18 PDB header: transferase/metal-binding protein Chain: B: PDB Molecule: ste20-related kinase adapter protein alpha; PDBTitle: structure of the heterotrimeric lkb1-stradalpha-mo25alpha2 complex
75	c2qg5D_	Alignment	not modelled	97.7	18 PDB header: transferase Chain: D: PDB Molecule: calcium/calmodulin-dependent protein kinase; PDBTitle: cryptosporidium parvum calcium dependent protein kinase cgd7_1840
76	d1o6ya_	Alignment	not modelled	97.7	20 Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
77	c2bmcD_	Alignment	not modelled	97.7	22 PDB header: transferase Chain: D: PDB Molecule: serine threonine-protein kinase 6; PDBTitle: aurora-2 t287d t288d complexed with pha-680632

78	c2wqmA_	Alignment	not modelled	97.6	17	PDB header: transferase Chain: A: PDB Molecule: serine/threonine-protein kinase nek7; PDBTitle: structure of apo human nek7
79	c2vd5A_	Alignment	not modelled	97.6	20	PDB header: transferase Chain: A: PDB Molecule: dmpk protein; PDBTitle: structure of human myotonic dystrophy protein kinase in2 complex with the bisindolmaleide inhibitor bim viii
80	c3byvA_	Alignment	not modelled	97.6	19	PDB header: transferase Chain: A: PDB Molecule: rhoprtry kinase; PDBTitle: crystal structure of toxoplasma gondii specific rhoprtry2 antigen kinase domain
81	d1qpca_	Alignment	not modelled	97.6	19	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
82	c2zv2A_	Alignment	not modelled	97.6	17	PDB header: transferase/transferase inhibitor Chain: A: PDB Molecule: calcium/calmodulin-dependent protein kinase kinase 2; PDBTitle: crystal structure of human calcium/calmodulin-dependent protein kinase2 kinase 2, beta, camkk2 kinase domain in complex with sto-609
83	d1phka_	Alignment	not modelled	97.6	18	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
84	c2chlA_	Alignment	not modelled	97.6	20	PDB header: transferase Chain: A: PDB Molecule: casein kinase i isoform gamma-3; PDBTitle: structure of casein kinase 1 gamma 3
85	d1fmka3	Alignment	not modelled	97.6	17	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
86	c1mrub_	Alignment	not modelled	97.5	19	PDB header: transferase Chain: B: PDB Molecule: probable serine/threonine-protein kinase pknb; PDBTitle: intracellular ser/thr protein kinase domain of2 mycobacterium tuberculosis pknb.
87	c2hakH_	Alignment	not modelled	97.5	20	PDB header: signaling protein,transferase Chain: H: PDB Molecule: serine/threonine-protein kinase mark1; PDBTitle: catalytic and ubiquitin-associated domains of mark1/par-1
88	c2qluA_	Alignment	not modelled	97.5	14	PDB header: transferase Chain: A: PDB Molecule: activin receptor type iib; PDBTitle: crystal structure of activin receptor type ii kinase domain2 from human
89	d1omwa3	Alignment	not modelled	97.5	17	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
90	d1nvra_	Alignment	not modelled	97.5	16	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
91	c3is5E_	Alignment	not modelled	97.5	20	PDB header: transferase Chain: E: PDB Molecule: calcium-dependent protein kinase; PDBTitle: crystal structure of cdpk kinase domain from toxoplasma gondii, 2 tgme49_018720
92	c3zzwA_	Alignment	not modelled	97.4	15	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase transmembrane receptor ror2; PDBTitle: crystal structure of the kinase domain of ror2
93	c2ivsA_	Alignment	not modelled	97.4	16	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase receptor PDBTitle: crystal structure of non-phosphorylated ret tyrosine kinase2 domain
94	c1koaA_	Alignment	not modelled	97.4	14	PDB header: kinase Chain: A: PDB Molecule: twitchin; PDBTitle: twitchin kinase fragment (c.elegans), autoregulated protein2 kinase and immunoglobulin domains
95	c2r7bA_	Alignment	not modelled	97.3	18	PDB header: signaling protein, transferase Chain: A: PDB Molecule: phosphoinositide-dependent protein kinase 1; PDBTitle: crystal structure of the phosphoinositide-dependent kinase-2 1 (pdk-1)catalytic domain bound to a dibenzonaphthyridine3 inhibitor
96	c1zrzA_	Alignment	not modelled	97.3	17	PDB header: transferase Chain: A: PDB Molecule: protein kinase c, iota; PDBTitle: crystal structure of the catalytic domain of atypical2 protein kinase c-iota
97	c3g0eA_	Alignment	not modelled	97.3	14	PDB header: transferase Chain: A: PDB Molecule: mast/stem cell growth factor receptor; PDBTitle: kit kinase domain in complex with sunitinib
98	c3lijA_	Alignment	not modelled	97.3	21	PDB header: transferase Chain: A: PDB Molecule: calcium/calmodulin dependent protein kinase with PDBTitle: crystal structure of full length cpcdpk3 (cgd5_820) in2 complex with ca2+ and amppnp
99	d1lufa_	Alignment	not modelled	97.3	20	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
100	c1lufA_	Alignment	not modelled	97.3	20	PDB header: transferase Chain: A: PDB Molecule: muscle-specific tyrosine kinase receptor musk; PDBTitle: crystal structure of the musk tyrosine kinase: insights2 into receptor autoregulation
101	c3hztA_	Alignment	not modelled	97.3	17	PDB header: transferase Chain: A: PDB Molecule: calcium-dependent protein kinase 3; PDBTitle: crystal structure of toxoplasma gondii cdpk3, tgme49_105860
102	d2jfla1	Alignment	not modelled	97.3	19	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like)

						Family: Protein kinases, catalytic subunit
103	c2pvfA_	Alignment	not modelled	97.3	12	PDB header: transferase Chain: A: PDB Molecule: fibroblast growth factor receptor 2; PDBTitle: crystal structure of tyrosine phosphorylated activated fgf2 receptor 2 (fgfr2) kinase domain in complex with atp3 analog and substrate peptide
104	c2a27B_	Alignment	not modelled	97.2	18	PDB header: transferase Chain: B: PDB Molecule: death-associated protein kinase 2; PDBTitle: human drp-1 kinase, w305s s308a d40 mutant, crystal form with 82 monomers in the asymmetric unit
105	c3dpkA_	Alignment	not modelled	97.2	15	PDB header: transferase Chain: A: PDB Molecule: macrophage colony-stimulating factor 1 receptor; PDBTitle: cfms tyrosine kinase in complex with a pyridopyrimidinone2 inhibitor
106	c2x7fA_	Alignment	not modelled	97.2	16	PDB header: transferase Chain: A: PDB Molecule: traf2 and nck-interacting protein kinase; PDBTitle: crystal structure of the kinase domain of human traf2-and2 nck-interacting kinase with wee1chk1 inhibitor
107	c3begA_	Alignment	not modelled	97.1	15	PDB header: transferase/splicing Chain: A: PDB Molecule: serine/threonine-protein kinase srp1; PDBTitle: crystal structure of sr protein kinase 1 complexed to its substrate2 asf/sf2
108	d2psqa1	Alignment	not modelled	97.1	16	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
109	c2psqA_	Alignment	not modelled	97.1	16	PDB header: transferase Chain: A: PDB Molecule: fibroblast growth factor receptor 2; PDBTitle: crystal structure of unphosphorylated unactivated wild type2 fgf receptor 2 (fgfr2) kinase domain
110	c3h9rA_	Alignment	not modelled	97.1	16	PDB header: isomerase/protein kinase Chain: A: PDB Molecule: activin receptor type-1; PDBTitle: crystal structure of the kinase domain of type i activin receptor2 (acvr1) in complex with fkbp12 and dorsomorphin
111	d1csna_	Alignment	not modelled	97.0	22	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
112	c3eygA_	Alignment	not modelled	97.0	18	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase; PDBTitle: crystal structures of jak1 and jak2 inhibitor complexes
113	d1ckia_	Alignment	not modelled	97.0	22	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
114	d1uu3a_	Alignment	not modelled	97.0	19	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
115	c2i0eB_	Alignment	not modelled	97.0	19	PDB header: transferase Chain: B: PDB Molecule: protein kinase c-beta ii; PDBTitle: structure of catalytic domain of human protein kinase c2 beta ii complexed with a bisindolylmaleimide inhibitor
116	c2zmcA_	Alignment	not modelled	97.0	16	PDB header: transferase Chain: A: PDB Molecule: dual specificity protein kinase ttk; PDBTitle: crystal structure of human mitotic checkpoint kinase mps1 catalytic2 domain apo form
117	d1sm2a_	Alignment	not modelled	97.0	17	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
118	d1fgka_	Alignment	not modelled	97.0	14	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
119	d1u59a_	Alignment	not modelled	96.9	23	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit
120	d1k2pa_	Alignment	not modelled	96.9	15	Fold: Protein kinase-like (PK-like) Superfamily: Protein kinase-like (PK-like) Family: Protein kinases, catalytic subunit