

























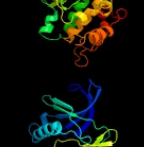



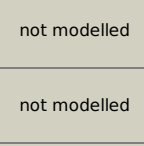


# Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD1266c_(pknH)_1413966_1415846
Date	Wed Jul 31 22:05:36 BST 2019
Unique Job ID	b339e5a7b7453c68

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c5ebzF_</a>	 Alignment		100.0	26	<b>PDB header:</b> transferase/transferase inhibitor <b>Chain:</b> F: <b>PDB Molecule:</b> inhibitor of nuclear factor kappa-b kinase subunit alpha; <b>PDBTitle:</b> crystal structure of human ikk1
2	<a href="#">c2pziA_</a>	 Alignment		100.0	25	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> probable serine/threonine-protein kinase pkng; <b>PDBTitle:</b> crystal structure of protein kinase pkng from mycobacterium2 tuberculosis in complex with tetrahydrobenzothiothiophene ax20017
3	<a href="#">c3qa8H_</a>	 Alignment		100.0	25	<b>PDB header:</b> immune system, signaling protein <b>Chain:</b> H: <b>PDB Molecule:</b> mgc80376 protein; <b>PDBTitle:</b> crystal structure of inhibitor of kappa b kinase beta
4	<a href="#">c3nyoB_</a>	 Alignment		100.0	25	<b>PDB header:</b> transferase <b>Chain:</b> B: <b>PDB Molecule:</b> g protein-coupled receptor kinase 6; <b>PDBTitle:</b> crystal structure of g protein-coupled receptor kinase 6 in complex2 with amp
5	<a href="#">c3qa8A_</a>	 Alignment		100.0	26	<b>PDB header:</b> immune system, signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> mgc80376 protein; <b>PDBTitle:</b> crystal structure of inhibitor of kappa b kinase beta
6	<a href="#">c4wnkA_</a>	 Alignment		100.0	22	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> g protein-coupled receptor kinase 5; <b>PDBTitle:</b> crystal structure of bovine g protein coupled-receptor kinase 5 in2 complex with ccg215022
7	<a href="#">c4im2A_</a>	 Alignment		100.0	22	<b>PDB header:</b> transferase/transferase inhibitor <b>Chain:</b> A: <b>PDB Molecule:</b> serine/threonine-protein kinase tbk1; <b>PDBTitle:</b> structure of tank-binding kinase 1
8	<a href="#">c2acxB_</a>	 Alignment		100.0	24	<b>PDB header:</b> transferase <b>Chain:</b> B: <b>PDB Molecule:</b> g protein-coupled receptor kinase 6; <b>PDBTitle:</b> crystal structure of g protein coupled receptor kinase 6 bound to2 amppnp
9	<a href="#">c3pfqA_</a>	 Alignment		100.0	22	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> protein kinase c beta type; <b>PDBTitle:</b> crystal structure and allosteric activation of protein kinase c beta2 ii
10	<a href="#">c4yhjA_</a>	 Alignment		100.0	23	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> g protein-coupled receptor kinase 4; <b>PDBTitle:</b> structure and function of the hypertension variant a486v of g protein-2 coupled receptor kinase 4 (grk4)
11	<a href="#">c1ym7C_</a>	 Alignment		100.0	22	<b>PDB header:</b> transferase <b>Chain:</b> C: <b>PDB Molecule:</b> beta-adrenergic receptor kinase 1; <b>PDBTitle:</b> g protein-coupled receptor kinase 2 (grk2)

12	<a href="#">c3c4wB_</a>	Alignment		100.0	24	<b>PDB header:</b> transferase <b>Chain:</b> B: <b>PDB Molecule:</b> rhodopsin kinase; <b>PDBTitle:</b> crystal structure of g protein coupled receptor kinase 1 bound to atp2 and magnesium chloride at 2.7a
13	<a href="#">c1koaA_</a>	Alignment		100.0	21	<b>PDB header:</b> kinase <b>Chain:</b> A: <b>PDB Molecule:</b> twitchin; <b>PDBTitle:</b> twitchin kinase fragment (c.elegans), autoregulated protein2 kinase and immunoglobulin domains
14	<a href="#">c3c4yA_</a>	Alignment		100.0	26	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> rhodopsin kinase; <b>PDBTitle:</b> crystal structure of apo form of g protein coupled receptor kinase 12 at 7.51a
15	<a href="#">c4i6hA_</a>	Alignment		100.0	25	<b>PDB header:</b> transferase/transferase inhibitor <b>Chain:</b> A: <b>PDB Molecule:</b> serine/threonine-protein kinase plk2; <b>PDBTitle:</b> selective & brain-permeable polo-like kinase-2 (plk-2) inhibitors that2 reduce alpha-synuclein phosphorylation in rat brain
16	<a href="#">c4czuC_</a>	Alignment		100.0	23	<b>PDB header:</b> transferase <b>Chain:</b> C: <b>PDB Molecule:</b> cbl-interacting serine/threonine-protein kinase 23; <b>PDBTitle:</b> crystal structure of the kinase domain of cipk23 t190d mutant
17	<a href="#">c6c9dB_</a>	Alignment		100.0	25	<b>PDB header:</b> transferase <b>Chain:</b> B: <b>PDB Molecule:</b> serine/threonine-protein kinase mark1,serine/threonine- <b>PDBTitle:</b> crystal structure of ka1-autoinhibited mark1 kinase
18	<a href="#">c5nclA_</a>	Alignment		100.0	23	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> serine/threonine-protein kinase cbk1; <b>PDBTitle:</b> crystal structure of the cbk1-mob2 kinase-coactivator complex with an2 ssd1 peptide
19	<a href="#">c4wb7B_</a>	Alignment		100.0	23	<b>PDB header:</b> transferase/transferase inhibitor <b>Chain:</b> B: <b>PDB Molecule:</b> dnaj homolog subfamily b member 1,camp-dependent protein <b>PDBTitle:</b> crystal structure of a chimeric fusion of human dnaj (hsp40) and camp-2 dependent protein kinase a (catalytic alpha subunit)
20	<a href="#">c2fo0A_</a>	Alignment		100.0	22	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> proto-oncogene tyrosine-protein kinase abl1 (1b isoform); <b>PDBTitle:</b> organization of the sh3-sh2 unit in active and inactive forms of the2 c-abl tyrosine kinase
21	<a href="#">c4myiA_</a>	Alignment	not modelled	100.0	25	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> cgmp-dependent protein kinase, putative; <b>PDBTitle:</b> crystal structure of pvx_084705
22	<a href="#">c4c0tA_</a>	Alignment	not modelled	100.0	21	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> likely protein kinase; <b>PDBTitle:</b> candida albicans pkh kinase domain
23	<a href="#">c3tkuB_</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase/transferase inhibitor <b>Chain:</b> B: <b>PDB Molecule:</b> serine/threonine-protein kinase mrck beta; <b>PDBTitle:</b> mrck beta in complex with fasudil
24	<a href="#">c5uuuA_</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase/transferase inhibitor <b>Chain:</b> A: <b>PDB Molecule:</b> beta-adrenergic receptor kinase 1; <b>PDBTitle:</b> design, synthesis, and evaluation of the first selective and potent g-2 protein-coupled receptor kinase 2 (grk2) inhibitor for the potential3 treatment of heart failure
25	<a href="#">c3q5iA_</a>	Alignment	not modelled	100.0	22	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> protein kinase; <b>PDBTitle:</b> crystal structure of pbanka_031420
26	<a href="#">c3soaA_</a>	Alignment	not modelled	100.0	25	<b>PDB header:</b> transferase/transferase inhibitor <b>Chain:</b> A: <b>PDB Molecule:</b> calcium/calmodulin-dependent protein kinase type ii subunit <b>PDBTitle:</b> full-length human camkii
27	<a href="#">c4cfhA_</a>	Alignment	not modelled	100.0	22	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> 5'-amp-activated protein kinase catalytic subunit alpha-1; <b>PDBTitle:</b> structure of an active form of mammalian ampk
28	<a href="#">c4b6lA_</a>	Alignment	not modelled	100.0	26	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> serine/threonine-protein kinase plk3; <b>PDBTitle:</b> discovery of oral polo-like kinase (plk) inhibitors with enhanced2 selectivity profile using residue targeted drug design

29	<a href="#">c1oplA</a>	Alignment	not modelled	100.0	22	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> proto-oncogene tyrosine-protein kinase; <b>PDBTitle:</b> structural basis for the auto-inhibition of c-abl tyrosine2 kinase
30	<a href="#">c4eqmE</a>	Alignment	not modelled	100.0	30	<b>PDB header:</b> transferase <b>Chain:</b> E: <b>PDB Molecule:</b> protein kinase; <b>PDBTitle:</b> structural analysis of staphylococcus aureus serine/threonine kinase2 pknb
31	<a href="#">c5dfzB</a>	Alignment	not modelled	100.0	18	<b>PDB header:</b> transferase <b>Chain:</b> B: <b>PDB Molecule:</b> serine/threonine-protein kinase vps15; <b>PDBTitle:</b> structure of vps34 complex ii from s. cerevisiae.
32	<a href="#">d2j4za1</a>	Alignment	not modelled	100.0	22	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
33	<a href="#">c3ckxA</a>	Alignment	not modelled	100.0	24	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> serine/threonine-protein kinase 24; <b>PDBTitle:</b> crystal structure of sterile 20-like kinase 3 (mst3, stk24) in complex2 with staurosporine
34	<a href="#">c2vwiC</a>	Alignment	not modelled	100.0	26	<b>PDB header:</b> transferase <b>Chain:</b> C: <b>PDB Molecule:</b> serine/threonine-protein kinase osr1; <b>PDBTitle:</b> structure of the osr1 kinase, a hypertension drug target
35	<a href="#">c4xi2A</a>	Alignment	not modelled	100.0	16	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> tyrosine-protein kinase btk; <b>PDBTitle:</b> crystal structure of an auto-inhibited form of bruton's tryrosine2 kinase
36	<a href="#">c4y93A</a>	Alignment	not modelled	100.0	16	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> non-specific protein-tyrosine kinase,non-specific protein- <b>PDBTitle:</b> crystal structure of the ph-th-kinase construct of bruton's tyrosine2 kinase (btk)
37	<a href="#">c2c30A</a>	Alignment	not modelled	100.0	25	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> serine/threonine-protein kinase pak 6; <b>PDBTitle:</b> crystal structure of the human p21-activated kinase 6
38	<a href="#">c1mruB</a>	Alignment	not modelled	100.0	36	<b>PDB header:</b> transferase <b>Chain:</b> B: <b>PDB Molecule:</b> probable serine/threonine-protein kinase pknb; <b>PDBTitle:</b> intracellular ser/thr protein kinase domain of mycobacterium2 tuberculosis pknb.
39	<a href="#">c6ccfB</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase/transferase inhibitor <b>Chain:</b> B: <b>PDB Molecule:</b> calcium/calmodulin-dependent protein kinase kinase 1; <b>PDBTitle:</b> crystal structure of the human camkk1a in complex with hesperadin
40	<a href="#">c5hmvA</a>	Alignment	not modelled	100.0	26	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> ppka n terminal; <b>PDBTitle:</b> crystal structure of ppka
41	<a href="#">c4x3fA</a>	Alignment	not modelled	100.0	38	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> serine/threonine-protein kinase pkna; <b>PDBTitle:</b> crystal structure of the intracellular domain of the m. tuberculosis2 ser/thr kinase pkna
42	<a href="#">c2jamB</a>	Alignment	not modelled	100.0	24	<b>PDB header:</b> transferase <b>Chain:</b> B: <b>PDB Molecule:</b> calcium/calmodulin-dependent protein kinase type 1g; <b>PDBTitle:</b> crystal structure of human calmodulin-dependent protein kinase i g
43	<a href="#">d1o6ya</a>	Alignment	not modelled	100.0	37	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
44	<a href="#">c2j0kB</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase <b>Chain:</b> B: <b>PDB Molecule:</b> focal adhesion kinase 1; <b>PDBTitle:</b> crystal structure of a fragment of focal adhesion kinase containing2 the ferm and kinase domains.
45	<a href="#">c4fieB</a>	Alignment	not modelled	100.0	24	<b>PDB header:</b> transferase <b>Chain:</b> B: <b>PDB Molecule:</b> serine/threonine-protein kinase pak 4; <b>PDBTitle:</b> full-length human pak4
46	<a href="#">c4fl2A</a>	Alignment	not modelled	100.0	21	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> tyrosine-protein kinase syk; <b>PDBTitle:</b> structural and biophysical characterization of the syk activation2 switch
47	<a href="#">c4fijA</a>	Alignment	not modelled	100.0	24	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> serine/threonine-protein kinase pak 4; <b>PDBTitle:</b> catalytic domain of human pak4
48	<a href="#">c2y7jB</a>	Alignment	not modelled	100.0	26	<b>PDB header:</b> transferase <b>Chain:</b> B: <b>PDB Molecule:</b> phosphorylase b kinase gamma catalytic chain, testis/liver <b>PDBTitle:</b> structure of human phosphorylase kinase, gamma 2
49	<a href="#">d1phka</a>	Alignment	not modelled	100.0	24	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
50	<a href="#">c2ozoA</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> tyrosine-protein kinase zap-70; <b>PDBTitle:</b> autoinhibited intact human zap-70
51	<a href="#">c1y57A</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> proto-oncogene tyrosine-protein kinase src; <b>PDBTitle:</b> structure of unphosphorylated c-src in complex with an inhibitor
52	<a href="#">c4wnoA</a>	Alignment	not modelled	100.0	25	<b>PDB header:</b> transferase/transferase inhibitor <b>Chain:</b> A: <b>PDB Molecule:</b> serine/threonine-protein kinase ulk1; <b>PDBTitle:</b> structure of ulk1 bound to an inhibitor
53	<a href="#">c4eutA</a>	Alignment	not modelled	100.0	20	<b>PDB header:</b> transferase/transferase inhibitor <b>Chain:</b> A: <b>PDB Molecule:</b> serine/threonine-protein kinase tbk1; <b>PDBTitle:</b> structure of bx-795 complexed with unphosphorylated human tbk1 kinase-2 uld domain
						<b>PDB header:</b> transferase

54	<a href="#">c3d9vA</a>	Alignment	not modelled	100.0	25	<b>Chain:</b> A; <b>PDB Molecule:</b> rho-associated protein kinase 1; <b>PDBTitle:</b> crystal structure of rock i bound to h-1152p a di-2 methylated variant of fasudil
55	<a href="#">c4btfA</a>	Alignment	not modelled	100.0	18	<b>PDB header:</b> transferase <b>Chain:</b> A; <b>PDB Molecule:</b> mixed lineage kinase domain-like protein; <b>PDBTitle:</b> structure of mlkl
56	<a href="#">c6fdzU</a>	Alignment	not modelled	100.0	25	<b>PDB header:</b> transferase <b>Chain:</b> U; <b>PDB Molecule:</b> serine/threonine-protein kinase ulk3; <b>PDBTitle:</b> unc-51-like kinase 3 (ulk3) in complex with momelotinib
57	<a href="#">c2h8hA</a>	Alignment	not modelled	100.0	22	<b>PDB header:</b> transferase <b>Chain:</b> A; <b>PDB Molecule:</b> proto-oncogene tyrosine-protein kinase src; <b>PDBTitle:</b> src kinase in complex with a quinazoline inhibitor
58	<a href="#">c2a27B</a>	Alignment	not modelled	100.0	22	<b>PDB header:</b> transferase <b>Chain:</b> B; <b>PDB Molecule:</b> death-associated protein kinase 2; <b>PDBTitle:</b> human drp-1 kinase, w305s s308a d40 mutant, crystal form with 82 monomers in the asymmetric unit
59	<a href="#">c3lijA</a>	Alignment	not modelled	100.0	22	<b>PDB header:</b> transferase <b>Chain:</b> A; <b>PDB Molecule:</b> calcium/calmodulin dependent protein kinase with <b>PDBTitle:</b> crystal structure of full length cpdpk3 (cgd5_820) in2 complex with ca2+ and amppnp
60	<a href="#">c2qg5D</a>	Alignment	not modelled	100.0	26	<b>PDB header:</b> transferase <b>Chain:</b> D; <b>PDB Molecule:</b> calcium/calmodulin-dependent protein kinase; <b>PDBTitle:</b> cryptosporidium parvum calcium dependent protein kinase cgd7_1840
61	<a href="#">c4xhgA</a>	Alignment	not modelled	100.0	20	<b>PDB header:</b> transferase <b>Chain:</b> A; <b>PDB Molecule:</b> similar to uniprot p29295 saccharomyces cerevisiae ypl204w
62	<a href="#">c6eqiC</a>	Alignment	not modelled	100.0	21	<b>PDB header:</b> transferase <b>Chain:</b> C; <b>PDB Molecule:</b> serine/threonine-protein kinase pink1, putative; <b>PDBTitle:</b> structure of pink1 bound to ubiquitin
63	<a href="#">c3dfaA</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase <b>Chain:</b> A; <b>PDB Molecule:</b> calcium-dependent protein kinase cgd3_920; <b>PDBTitle:</b> crystal structure of kinase domain of calcium-dependent protein kinase2 cgd3_920 from cryptosporidium parvum
64	<a href="#">c4y83B</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase <b>Chain:</b> B; <b>PDB Molecule:</b> mitogen-activated protein kinase kinase kinase 8; <b>PDBTitle:</b> crystal structure of cot kinase domain in complex with 5-(2-amino-5-(2-(quinolin-3-yl)pyridin-3-yl)-1,3,4-oxadiazole-2(3h)-thione
65	<a href="#">d1yhwa1</a>	Alignment	not modelled	100.0	24	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
66	<a href="#">c2wtkC</a>	Alignment	not modelled	100.0	19	<b>PDB header:</b> transferase/metal-binding protein <b>Chain:</b> C; <b>PDB Molecule:</b> serine/threonine-protein kinase 11; <b>PDBTitle:</b> structure of the heterotrimeric lkb1-stradalphamo25alpha2 complex
67	<a href="#">c2bmcD</a>	Alignment	not modelled	100.0	22	<b>PDB header:</b> transferase <b>Chain:</b> D; <b>PDB Molecule:</b> serine threonine-protein kinase 6; <b>PDBTitle:</b> aurora-2 t287d t288d complexed with pha-680632
68	<a href="#">c4dn5A</a>	Alignment	not modelled	100.0	28	<b>PDB header:</b> transferase <b>Chain:</b> A; <b>PDB Molecule:</b> mitogen-activated protein kinase kinase kinase 14; <b>PDBTitle:</b> crystal structure of nf-kb-inducing kinase (nik)
69	<a href="#">d1ksa</a>	Alignment	not modelled	100.0	22	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
70	<a href="#">c3zzwA</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase <b>Chain:</b> A; <b>PDB Molecule:</b> tyrosine-protein kinase transmembrane receptor ror2; <b>PDBTitle:</b> crystal structure of the kinase domain of ror2
71	<a href="#">c1k9aB</a>	Alignment	not modelled	100.0	20	<b>PDB header:</b> transferase <b>Chain:</b> B; <b>PDB Molecule:</b> carboxyl-terminal src kinase; <b>PDBTitle:</b> crystal structure analysis of full-length carboxyl-terminal2 src kinase at 2.5 a resolution
72	<a href="#">c5zjbB</a>	Alignment	not modelled	100.0	24	<b>PDB header:</b> transferase <b>Chain:</b> B; <b>PDB Molecule:</b> serine/threonine-protein kinase dclk1; <b>PDBTitle:</b> crystal structure of dclk1-kd in complex with amppn
73	<a href="#">d1qcfa3</a>	Alignment	not modelled	100.0	24	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
74	<a href="#">c5x3fB</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase <b>Chain:</b> B; <b>PDB Molecule:</b> zpa963,camp-dependent protein kinase catalytic subunit <b>PDBTitle:</b> crystal structure of the ygig-protein a-zpa963-pka catalytic domain
75	<a href="#">c1xjdA</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase <b>Chain:</b> A; <b>PDB Molecule:</b> protein kinase c, theta type; <b>PDBTitle:</b> crystal structure of pkc-theta complexed with staurosporine2 at 2a resolution
76	<a href="#">d1xjda</a>	Alignment	not modelled	100.0	23	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
77	<a href="#">c3hztA</a>	Alignment	not modelled	100.0	22	<b>PDB header:</b> transferase <b>Chain:</b> A; <b>PDB Molecule:</b> calcium-dependent protein kinase 3; <b>PDBTitle:</b> crystal structure of toxoplasma gondii cpdpk3, tgme49_105860
78	<a href="#">c5oatF</a>	Alignment	not modelled	100.0	22	<b>PDB header:</b> kinase <b>Chain:</b> F; <b>PDB Molecule:</b> serine/threonine-protein kinase pink1, mitochondrial-like <b>PDBTitle:</b> pink1 structure
						<b>PDB header:</b> transferase

79	<a href="#">c5d7vD_</a>	Alignment	not modelled	100.0	21	<b>Chain:</b> D: <b>PDB Molecule:</b> protein-tyrosine kinase 6; <b>PDBTitle:</b> crystal structure of ptk6 kinase domain
80	<a href="#">c5ig1A_</a>	Alignment	not modelled	100.0	22	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> camk/camk2 protein kinase; <b>PDBTitle:</b> crystal structure of s. rosetta camkii kinase domain
81	<a href="#">c4lqsA_</a>	Alignment	not modelled	100.0	24	<b>PDB header:</b> transferase/transferase activator <b>Chain:</b> A: <b>PDB Molecule:</b> serine/threonine-protein kinase cbk1; <b>PDBTitle:</b> crystal structure of the cbk1-mob2 kinase-coactivator complex
82	<a href="#">c4ci6B_</a>	Alignment	not modelled	100.0	15	<b>PDB header:</b> transferase/structural protein <b>Chain:</b> B: <b>PDB Molecule:</b> protein kinase yopo; <b>PDBTitle:</b> mechanisms of crippling actin-dependent phagocytosis by yopo
83	<a href="#">d1lopja_</a>	Alignment	not modelled	100.0	22	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
84	<a href="#">c2ya9A_</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> death-associated protein kinase 2; <b>PDBTitle:</b> crystal structure of the autoinhibited form of mouse dapk2
85	<a href="#">c2bdwB_</a>	Alignment	not modelled	100.0	25	<b>PDB header:</b> transferase <b>Chain:</b> B: <b>PDB Molecule:</b> hypothetical protein k11e8.1d; <b>PDBTitle:</b> crystal structure of the auto-inhibited kinase domain of2 calcium/calmodulin activated kinase ii
86	<a href="#">d1koaa2</a>	Alignment	not modelled	100.0	21	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
87	<a href="#">c2vd5A_</a>	Alignment	not modelled	100.0	21	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> dmpk protein; <b>PDBTitle:</b> structure of human myotonic dystrophy protein kinase in2 complex with the bisindoylmaleide inhibitor bim viii
88	<a href="#">d1o6la_</a>	Alignment	not modelled	100.0	22	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
89	<a href="#">c4cdsA_</a>	Alignment	not modelled	100.0	24	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> tyrosine kinase as - a common ancestor of src and abl; <b>PDBTitle:</b> tyrosine kinase as - a common ancestor of src and abl
90	<a href="#">c6cqdB_</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase <b>Chain:</b> B: <b>PDB Molecule:</b> mitogen-activated protein kinase kinase kinase kinase 1; <b>PDBTitle:</b> crystal structure of hpk1 in complex with atp analogue (amppnp)
91	<a href="#">c2zv2A_</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase/transferase inhibitor <b>Chain:</b> A: <b>PDB Molecule:</b> calcium/calmodulin-dependent protein kinase kinase 2; <b>PDBTitle:</b> crystal structure of human calcium/calmodulin-dependent protein kinase2 kinase 2, beta, camkk2 kinase domain in complex with sto-609
92	<a href="#">c1kobB_</a>	Alignment	not modelled	100.0	19	<b>PDB header:</b> kinase <b>Chain:</b> B: <b>PDB Molecule:</b> twitchin; <b>PDBTitle:</b> twitchin kinase fragment (aplysia), autoregulated protein2 kinase domain
93	<a href="#">c4qfrA_</a>	Alignment	not modelled	100.0	22	<b>PDB header:</b> signaling protein/inhibitor/activator <b>Chain:</b> A: <b>PDB Molecule:</b> 5'-amp-activated protein kinase catalytic subunit alpha-1; <b>PDBTitle:</b> structure of ampk in complex with ci-a769662 activator and2 staurosporine inhibitor
94	<a href="#">c2j51A_</a>	Alignment	not modelled	100.0	23	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> ste20-like serine/threonine-protein kinase; <b>PDBTitle:</b> 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
95	<a href="#">c2a1aB_</a>	Alignment	not modelled	100.0	24	<b>PDB header:</b> protein synthesis/transferase <b>Chain:</b> B: <b>PDB Molecule:</b> interferon-induced, double-stranded rna-activated protein <b>PDBTitle:</b> pkr kinase domain-eif2alpha complex
96	<a href="#">c2wtkB_</a>	Alignment	not modelled	100.0	18	<b>PDB header:</b> transferase/metal-binding protein <b>Chain:</b> B: <b>PDB Molecule:</b> ste20-related kinase adapter protein alpha; <b>PDBTitle:</b> structure of the heterotrimeric lkb1-stradalpha-mo25alpha2 complex
97	<a href="#">c2c0iA_</a>	Alignment	not modelled	100.0	24	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> tyrosine-protein kinase hck; <b>PDBTitle:</b> src family kinase hck with bound inhibitor a-420983
98	<a href="#">d1koba_</a>	Alignment	not modelled	100.0	20	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
99	<a href="#">c2rsvA_</a>	Alignment	not modelled	100.0	15	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> serine/threonine-protein kinase vrk1; <b>PDBTitle:</b> solution structure of human full-length vaccinia related kinase 12 (vrk1)
100	<a href="#">c2vz6A_</a>	Alignment	not modelled	100.0	25	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> calcium calmodulin dependent protein kinase type ii alpha <b>PDBTitle:</b> structure of human calcium calmodulin dependent protein kinase type ii2 alpha (camk2a) in complex with indirubin e804
101	<a href="#">c2r7bA_</a>	Alignment	not modelled	100.0	20	<b>PDB header:</b> signaling protein, transferase <b>Chain:</b> A: <b>PDB Molecule:</b> phosphoinositide-dependent protein kinase 1; <b>PDBTitle:</b> crystal structure of the phosphoinositide-dependent kinase-2 1 (pdk-1)catalytic domain bound to a dibenzonaphthyridine3 inhibitor
102	<a href="#">d2jfla1</a>	Alignment	not modelled	100.0	24	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit <b>PDB header:</b> signaling protein

103	<a href="#">c4lqdB_</a>	Alignment	not modelled	100.0	24	<b>Chain:</b> B: <b>PDB Molecule:</b> serine/threonine-protein kinase 3; <b>PDBTitle:</b> structural basis for autoactivation of human mst2 kinase and its2 regulation by rassf5
104	<a href="#">c6jkmA_</a>	Alignment	not modelled	100.0	18	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> mitotic checkpoint control protein kinase bub1; <b>PDBTitle:</b> crystal structure of bubr1 kinase domain
105	<a href="#">c2x4fA_</a>	Alignment	not modelled	100.0	20	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> myosin light chain kinase family member 4; <b>PDBTitle:</b> the crystal structure of the human myosin light chain kinase2 loc340156.
106	<a href="#">c5j5tA_</a>	Alignment	not modelled	100.0	22	<b>PDB header:</b> transferase/transferase inhibitor <b>Chain:</b> A: <b>PDB Molecule:</b> mitogen-activated protein kinase kinase kinase 3; <b>PDBTitle:</b> glk co-crystal structure with aminopyrrolopyrimidine inhibitor
107	<a href="#">d1qpca_</a>	Alignment	not modelled	100.0	23	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
108	<a href="#">c2cgvA_</a>	Alignment	not modelled	100.0	26	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> serine/threonine-protein kinase chk1; <b>PDBTitle:</b> identification of chemically diverse chk1 inhibitors by2 receptor-based virtual screening
109	<a href="#">d1tkia_</a>	Alignment	not modelled	100.0	20	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
110	<a href="#">c3cblA_</a>	Alignment	not modelled	100.0	22	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> proto-oncogene tyrosine-protein kinase fes/fps; <b>PDBTitle:</b> crystal structure of human feline sarcoma viral oncogene homologue (v-2 fes) in complex with staurosporine and a consensus peptide
111	<a href="#">c2ivsA_</a>	Alignment	not modelled	100.0	21	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> proto-oncogene tyrosine-protein kinase receptor ret; <b>PDBTitle:</b> crystal structure of non-phosphorylated ret tyrosine kinase domain
112	<a href="#">c5dvtA_</a>	Alignment	not modelled	100.0	20	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> calmodulin-domain protein kinase 1; <b>PDBTitle:</b> crystal structure of tgcdpk1 from toxoplasma gondii complexed with 5gb
113	<a href="#">c3g0eA_</a>	Alignment	not modelled	100.0	21	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> mast/stem cell growth factor receptor; <b>PDBTitle:</b> kit kinase domain in complex with sunitinib
114	<a href="#">c5o0yA_</a>	Alignment	not modelled	100.0	26	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> serine/threonine-protein kinase tousled-like 2; <b>PDBTitle:</b> tlk2 kinase domain from human
115	<a href="#">d1nvra_</a>	Alignment	not modelled	100.0	24	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
116	<a href="#">d1uu3a_</a>	Alignment	not modelled	100.0	21	<b>Fold:</b> Protein kinase-like (PK-like) <b>Superfamily:</b> Protein kinase-like (PK-like) <b>Family:</b> Protein kinases, catalytic subunit
117	<a href="#">c1zrzA_</a>	Alignment	not modelled	100.0	22	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> protein kinase c, iota; <b>PDBTitle:</b> crystal structure of the catalytic domain of atypical2 protein kinase c-iota
118	<a href="#">c3c0hB_</a>	Alignment	not modelled	100.0	24	<b>PDB header:</b> transferase <b>Chain:</b> B: <b>PDB Molecule:</b> peripheral plasma membrane protein cask; <b>PDBTitle:</b> cask cam-kinase domain- amppnp complex, p1 form
119	<a href="#">c5u6bD_</a>	Alignment	not modelled	100.0	21	<b>PDB header:</b> transferase/transferase inhibitor <b>Chain:</b> D: <b>PDB Molecule:</b> tyrosine-protein kinase receptor ufo; <b>PDBTitle:</b> structure of the axl kinase domain in complex with a macrocyclic2 inhibitor
120	<a href="#">c3fbvL_</a>	Alignment	not modelled	100.0	21	<b>PDB header:</b> transferase, hydrolase <b>Chain:</b> L: <b>PDB Molecule:</b> serine/threonine-protein kinase/endoribonuclease ire1; <b>PDBTitle:</b> crystal structure of the oligomer formed by the kinase-ribonuclease2 domain of ire1