	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q6PCN3	TTBK1_MOUSE	Mus musculus	Tau-tubulin kinase 1	25.810548	NaN	S441;S541	29187734	MQCLAAALKDETNMSGGGEQADIL PANYVVKDRWKVLKKIGGGGFGEIY EAMDLLTRENVALKVESAQQPKQVL KMEVAVLKKLQGKDHVCRFIGCGR NEKFNYVVMQLQGRNLADLRRSQP RGTFTLSTTLRLGKQILESIEAIHSVG FLHRDIKPSNFAMGRLPSTYRKCYM LDFGLARQYTNTTGDVRPPRNVAGF RGTVRYASVNAHKNREMGRHDDL WSLFYMLVEFAVGQLPWRKIKDKEQ VGMIKEKYEHRMLLKHMPSEFHLF LDHIASLDYFTKPDYQLIMSVFENS MKERGIAENEAFDWEKAGTDALLST STSTPPQQNTRQTAAMFGVVNVTPV PGDLLRENTEDVLQGEHLSDQENAP PILPGRPPEGLGPGPHLVPHPGGPEA EVWEETDVNRNKLRINIGKTPCVEE EQSRGVGVPSSPVRAPPDSPTTPVRS LCYRRVNSPESERLSTAADGRVELQ ERRSRMDLPGSPSRQACSSQPAQM LSVDTGHADRQASGRMDVSASVEQ EALSNAFRSVPLAEEEDFDSKEWVII DKETELKDFPPGAEPSTSGTTDEEPE ELRPLPEEGEERRRLGTEPTVRPRG RGMHTLTEEDPRQMLPQPAPPQLS QADGRSETSQPPTPGSPSHSPLHSG PRPRRRESDPTGPQRQVFSVAPPFE VNGLPRAVPLALPYQDFKRDLSDYR ERARLLNRVRRVGFSHMLLTAPQVP LAPFQPQANGKEGEEEEEEEEEEEE EEEEEEEEEEEEEEEEEEEEAGAL GEVLGPRSGSSSEGSERSTERSQEG APSTLLADDQKEARGRASMADGDLE PEEGSKTLVLVSPGDMKKSPVTAEL APDPDLGTLAALTPQHERPQPTGSQ LDVSEPGTLSSILKSEPKPSGPGAGG GVGLVAPGAGVTAVTSPFTKVERTFV HIAEKSHLNVMSSGGQASRPEELST GGELGLEVLSEGGIAEEGAPAPLEN GMALAGLDGTEMESCALSGPPGETP SEVVTDSLPNGPALADGPAPASQQE PVTKKGTTISPSRHAMPGSRPRSRIP VLLSEEDTGSEPSGSLSAKERWSKR ARPQQDLARLVMEKRQGRLLLRLAS GASSSSSEEQRRASETLSGTGSEED TPASEPTTALPRKAVRAATTRSRIPRP ISVSMPVEGQQLPGRPHGAASATDL AITSRLQLQKPSGLAPAADLRPKQSA SRGPGPGRAQVSKPAAPRSPGLPAST ARHPSGSPRSQSLSRKESSSPSHQA RPGVPPSRGVLQVRSQPEASPVAPK KGPKGKQLQTQRAATKGRAVVSEGR PGAR	False	False	3.882	3.514	1.228	0.93	0.808	1.059	1.162