

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q3UYK3	TBCD9_MOUSE	Mus musculus	TBC1 domain family member 9	34.74265	S346;T984	NaN	22645316;34418053	MWVNPEEVLLANALWITERANPYFI LQRRKGHGDDGGGGGLAGLLVGT LDVVLDSSARVAPYRILYQTPDSL VYWTIACGGSRKEVTEHWEWLEQ NLLQTL SIFENENDVTTFVRGKI QGIIAEY NKINDVKEDEDETE KFKEAIVKFHRL FGMPEEEL LVNYYSCSYWKG RVP R QG WMYLSINHLCFSSFLMGREAK L VIRWVDITQLEKNATLLLPD MIKVST RSSEHFFSVFLNIN ETFKLMEQLANI AMRQLLDN EGFEQDRSLPKLKKKSP K KVSALKRDL DARAKSERYR ALFRLP KDEKLDGHTDCTL WTPFNKMHLG QMFVSTNY ICFTSKEENLCSLIPLR E VTIVEKADSCSVLP SPLS ISTRNRM TFLFANLKDRDF LVQRISDFLQQTTS RIYSD KEFGSGSCNSSDEVYSRP SSLVSSSPQRSTSSDADG GERPFNLNGNS VPTATQ TLMTMYRRRSPEEFNP KLA KEFLKEQAWKIHFAE YGGIGCMYRT EKTREL VLKGIPESMRGE LWLL SG AINEKATHPGY YGLVEKSMGKYNL ATEEIERDLHRSL PEHPAFQNE MGI AALRRVLTAYAF RNPNI GYQAMNI VTSVLLLYAKEE EAFWLLVALCERM LPDYNTTRVVGAL VDQGVFEELARD YVPQLYDCMQD LGVISTISLSW FLTL FLSVMPFES AVVVVDCFFYEG IKVIF QLALAVLDAN VDKLNCKDDGEAM TVLGRYLDSTN KDSTLPPIPHLH SL LSDDVGPYP AVDIFRLIGTSY EKFGTI RADLIEQ MRFKQRLKVIQ TLEDTTKR NVVRTI VTETSTIDELE ELYALFKAE HLTSCYWGGS SNALDRHDP SLPYLE QYRIDFE QFKGMFVLL FPWACGTHS DVLASRLFQ LLENGDSLINF REFV SGLSAACH GDLTEKLLY KMHVLP EPSCDQ DEPDSAFEAT QYFFEDITPE CTHVVG LDSRGKQSA DDGFVTVSLK QDRGKRAN SQENRNYL KLWTAENK SKSKTAKD LPKLNQGO FIELCKTMY NMFSEDP NEQELYHATA AAVTSLLLE IGEVGKFF ITQPAKEDAV PGPPCGQA IPGMLFP PKGSSQSY VVESTEPL TAS LAVDSEE HSLGGQME DIKLEDS SPR DNGACSS MLISDDDT KDDSSM SSYS VLSAGS HEEDKLH CEDIGED TVLVR SSQGRAT LPRSSSL DRDWAIT FEQFL ASLLTEP ALVRYFDK PVCMMAR VTS AKNIRMM GKPLTSAS DYEISALS G	None	None	None	None	None	None	None		

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