	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus
0	P33981	TTK_HUMAN	Homo sapiens	Dual specificity protein kinase TTK	23.627257	S699;S700;S701	S7;T33;S37;S80;S281;S317;S321;S393;S436;S455;S821	34019948;31492838;23301498	MESEDLSGRELTIDSIMNKVRDIKN KFKNEDLTDELSLNKISADTTDNSG TVNQIMMMANNPEDWLSLLLKLEK NSVPLSDALLNKLIGRYSQAIEALPP DKYGQNESFARIQVRFAELKAIQEPD DARDYFQMARANCKKFAFVHISFAQ FELSQGNVKKSKQLLQKAVERGAVP LEMLEIALRNLNLQKKQLLSEEEKK NLSASTVLTAQESFSGSLGHLQNRN NSCDSRGQTTKARFLYGENMPPQD AEIGYRNSLRQTNKTKQSCPFGRVP VNLLNSPDCDVKTDDSVVPCFMKR QTSRSECRDLVVPGSKPSGNDSCEL RNLKSVQNSHFKEPLVSDEKSSELII TDSITLKNKTESSLLAKLEETKEYQE PEVPESNQKQWQSKRKSECINQNP AASSNHWQIPELARKVNTEQKHTTF EQPVFSVSKQSPPISTSKWFDPKSIC KTPSSNTLDDYMSCFRTPVVKNDFP PACQLSTPYGQPACFQQQQHQILAT PLQNLQVLASSSANECISVKGRIYSIL KQIGSGGSSKVFQVLNEKKQIYAIKY VNLEEADNQTLDSYRNEIAYLNKLQ QHSDKIIRLYDYEITDQYIYMVMECG NIDLNSWLKKKKSIDPWERKSYWK NMLEAVHTIHQHGIVHSDLKPANFL IVDGMLKLIDFGIANQMQPDTTSVV KDSQVGTVNYMPPEAIKDMSSSREN GKSKSKISPKSDVWSLGCILYYMTYG KTPFQQIINQISKLHAIIDPNHEIEFP DIPEKDLQDVLKCCLKRDPKQRISIP ELLAHPYVQIQTHPVNQMAKGTTEE MKYVLGQLVGLNSPNSILKAAKTLY EHYSGGESHNSSSSKTFEKKRGKK	None	None	None	None	None	None	None