

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	extracell region
Q9ERD7	TBB3_MOUSE	Mus musculus	Tubulin beta-3 chain	43.237047	S339	S172;S444	40684658;20665223;34418053;3330544;22645316	MREIVHIQAGQCNGNQIGAKFWEVIS DEHGIDPSGNYVGDSDLQLERISVYY NEASSHKYVPRAILVDLEPGTMDSV RSGAFGHLFRPDNFIFGQSGAGNN WAKGHYTEGAELVDSVLDVVRKECE NCDCLQGFQLTHSLGGGTGSGMGT LLISKVREEYPDRIMNTFSVVPSPKV SDTVVEPYNATLSIHQLVENTDETYC IDNEALYDICFRTLKLATPTYGDLNH LVSATMSGVTTSLRFPQQLNADLRK LAVNMVPPRLHFFMPGFAPLTARG SQQYRALTVPELTQQMFDAKNMMA ACDPRHGRLTVATVFRGRMSMKE VDEQMLAIOSKNSSYFVEWIPNNVK VAVCDIPPRGLKMSSTFIGNSTAIQE LFRKRISEQFTAMFRRKAFLHWYTGE GMDEMEFTEAESNMNDLVSEYQQY QDATAEEEGEMYEDDDEESEAQGP K	False	False	2.757	1.312	False	False	False	False	2.0