	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	P08841	TBB3_DROME	Drosophila melanogaster	Tubulin beta-3 chain	27.146871	NaN	NaN	15952742	MREIVNLQAGQCGNQIGAKFWEIIS EEHGIDSNGIYVGDSDLQLERVSVYY NEASAVTRSSGGKYVPRAILLDLEPG TMESVRSGPYGQLFRPDNFVYGQSG AGNNWAKGHYTEGAELVDNVLDVV RKECENCDCLQGFQLTHSLGGGTG SGMGTLLISKIREEYPDRIMNTYSVV PSPKVSDTVVEPYNATLSIHQLVENT DETYCIDNEALYDICFRTLKVSNPSY GDLNHLVSLTMSGVTTCLRFPGQLN ADLRKLAVNMVPFPRLHFFMPGFAP LTSRGSQQYRALTVPELTQQMFDAK NMMAACDPRHGRYLTVAAVFRGRM SMKEVDEQMLAVQNKNSSYFVEWI PNNVKTAVCDIPPKGLKMSSTFIGN TTAIQELFKRISEQFSAMFRRKAFLH WYTGEGMDEMEFTEAESNMNDLV SEYQQYQEATADDEFDPEVNQEEVE GDCI	None	None	None	None	None	None	None	None	None