

| 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 | 37.222262 | S339 | S172;S444 | 22645316;20665223;33300544;34418053 | MREIVHIQAGQCGNQIGAKFWEVIS DEHGIDPSGNYVGDSDLQLERISVYY NEASSHKYVPRAILVDLEPGTMSV RSGAFGHLFRPDNFIFGQSGAGNN WAKGHYTEGAELVDSVLDVVRKECE NCDCLQGFQLTHSLGGGTGSGMGT LLISKVREEYPDRIMNTFSVVPSPKV SDTVVEPYNATLSIHQLVENTDETYC IDNEALYDICFRTLKLATPTYGDLNH LVSATMSGVTTSLRFPGQLNADLRK LAVNMVPPRLHFFMPGFAPLTARG SQQYRALTVPELTQQMFDAKNMMA ACDPRHGRLTVATVFRGRMSMKE VDEQMLAIQSKNSSYFVEWIPNNVK VAVCDIPPRGLKMSSTFIGNSTAIQE LFKRISEQFTAMFRRKAFLHWYTGE GMDEMEFTEAESNMNDLVSEYQQY QDATAEEEGEMYEDDDEESEAQGP K | False | False | 3.809 | 3.69 | 2.615 | 2.1 | 1.775 | 2.709 | 3.609 |