

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochondrion | endoplasmic reticulum | golgi appara |
|--------------|------------|--------------|------------------------------------|--------------|--------------------------|--|-------------------|--|---------------|---------------|---------|---------|---------------|-----------------------|--------------|
| Q08122 | TLE3_MOUSE | Mus musculus | Transducin-like enhancer protein 3 | 37.551664 | T285;S289;T328;S332;T333 | S203;S207;S211;S217;S240;S245;T259;S263;S267;S286;T312;S317;T319;T321;T328;T334;S413 | 40885482;36852467 | MYPQGRHPAPHQPGQPGFKFTVAE SCDRIKDEFQFLQAOYHSLKVEYDK LANEKTEMQRHYVMMYYEMSYGLNI EMHKQTEIAKRLNTILAQIMPFLSQE HQQQVAQAVERAKQVTMTELNAIIG QQQLQAQHLSHATHGPPVQLPPHP SGLQPPGIPPVTGSSSGLLALGALGS QAHLAVKDEKNHHELDHRRERESST NNSVSPSESLRASEKHRGSADYSME AKKRKAEKDSLRSRYDSGDGKSDDL VVDVSNEDPATPRVSPAHSPPENGL DKARGLKKDAPTSVASVSSSTPSS KTKDLGHNDKSSTPGLKSNTPTPRN DAPTPTGTSTTPGLRSMPPGKPPGMDP IGIMASALRTPITLTSSYPAPFAMMS HHEMNGSLTSPSAYAGLHNIPSQM SAAAAAAAAAYGRSPMVSGAVGFD PHPPMRATGLPSSLASIPGGKPAYSF HVSADGQMOPVPPHDLAAGPGIPR HARQINTLSHGEVVCVAVTISNPTRH VYTGKGKCVKIWDISQPGSKSPISQL DCLNRDNYIRSKLLPDGRTLIVGGE ASTLTIWDLASPTPRIKAELTSSAPAC YALAI SPDAKVCFSCCSDGNIAVWDL HNQTLVRQFQGH TDGASCIDISHDG TKLWTGGLDNTVRSWDLREGRQLQ QHDFTSQIFSLGYCPTGEWLAVGME SSNVEVLHHTKPKYQLHLHESCVL SLKFAYCGKWFVSTGKDNLLNAWR TPYGASIFQSKSSSVLSCDISADDK YIVTGS GDKKATVYEVY | True | False | 2.309 | 4.319 | 1.657 | 1.058 | False |