

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q04726	TLE3_HUMAN	Homo sapiens	Transducin-like enhancer protein 3	37.329985	S253;S286;S289;T328;T331;S332;T333;T334;T352 (Q04726-2);T359;S362;S413;S417	S203;S207;S211;S217;S240;S245;T259;S263;S267;S286;T312;S317;T319;T321;T328;T334;S413	40307207	MYPQGRHPAPHQPGQPGKFTVAE SCDRKDEFQFLQAQYHSLKVEYDK LANEKTEMQRHYVMYEYMSYGLNI EMHKQTEIAKRLNTILAQIMPFLSQE HQQQVAQAVERAKQVTMTLNAIIG QQQLQAQHLSHATHGPPVQLPPHP SGLQPPGIPPVTGSSSGLLALGALGS QAHLTVKDEKNHHELDHRERESSA NNSVSPSESLRASEKHRGSADYSME AKKRKAEEKDLSRYSDGDKSDDL VVDVSNEDPATPRVSPAHSPENGL DKARSLKKDAPTSPASVASSSTPSS KTKDLGHNDKSSTPGLKSNTPTRN DAPTPTGTTTPGLRSMGKPPGMDP IGIMASALRTPISITSSYAAPFAMMS HHEMNGSLTSPGAYAGLHNIPPQM SAAAAAAAAAYGRSPMVSGAVGDF PHPPMRATGLPSSLASIPGGKPAYSF HVSADGQMOPVPPHDLALAGPGIPR HARQINTLSHGEVVCVAVTISNPTRH VYTGKGCVKIWDISQPGSKSPISQL DCLNRDNYIRSKLLPDGRTLIVGGE ASTLTIWDLASPTPRIKAELTSSAPAC YALAIKPAKVCFCSSDGNIAVWDL HNQTLVRQFQGHDTGASCIDISHDG TKLWTGGLDNTVRSWDLREGRLQ QHDFTSQIFSLGYCPTGEWLAVGME SSNVEVLHHTKPKDYQLHLHESCVL SLKFAYCGKWFVSTGKDNLLNAWR TPYGASIFQSKESSVLSCDISADDK YIVTGSGDKKATVYEVY	True	False	1.409	5.0	1.673	1.099	0.612