

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	extracellular region
Q3U7R1	ESYT1_MOUSE	Mus musculus	Extended synaptotagmin-1	25.685855	NaN	S314;S807;S937;S951;T997	39627609	MEHSPEEGASPEPSGPPATDSTRD GGSGVPPAGPGAASEALAVLTSFGR RLLVLPVYLAGAAGLSVGFVLFGLA LYLGWRRVRDGGKERSLRAARQLLDD EERITAETLYMSHRELPAWVSFPDV EKAEWLNKIVAQVWPFLGQYMEKL LAETVAVAVRGANPHLQTFTRVE LGEKPLRIIGVKVHPSQRKDQILLDL NVSIVGDVQIDVEVKYFCKAGVKG MQLHGVLRVILEPLTGDLPVGAVS MFFIKRPTLDINWTGMTNLLDIPGL SSLSDTMIMDSIAAFLVLPNRLLVPL VPDLQDVAQLRSPLPRGIIRIHLAAR GLSSKDKYVKGKIEGKSDPYALVRVG TQTFCSRVIDEELNPHWGETYEVIV HEVPGQEIEVEVFDKDPKDDFLGR MKLDVGVKVLQAGVLDNWPYPLQGGQ GOVHLRLEWLSLLPDAEKLDQVLQ WNRGITSRPEPPSAAILVVYLDRAQD LPLKGNKEPNPMVQLSVQDVTRES KATYSTNSPVWEEAFRFFLQDPRSQ ELDVQVKDDSRALTLGALTLPLARLL TASELTLDQWFQLSSSGPNSRLYMK LVMRILYLDYSEIRFPTVPGAQDWDR ESLETGSSVDAPPRPYHTTPNSHFG TENVLRIHVLEAQDLIAKDRFLGGLV KGKSDPYVVKLVAGKSFRTHVRED LNPRWNEVFEVIVTSIPGQELEIEVF DKDLKDDDFLGRYKVSLLTVLNSGF LDEWLTLEDVPSGRLHLRLRLTPR PTAAELEEVLVQVNSLIQTQKSSSELA ALLSVFLERAEDLPLRKGTKPPSPYA TITVGETSHKTKTVSQSSAPVWEESA SFLIRKPHAESLELQVRGEGTGLGS VSLPSELLQEDQLCLDHWFALSGQ GOVLMRAQLGILVSOHSGVEAHSHS YSHSHSSSLNDEPEALGGPHTPAS PVLEVRHRLTHGDSPEAPVGPLGQ VKLTVWYHSDEQKLSIIHSCRALRQ NGRDLPDPYVSVLLPDKNRSTKRK TPQKKRTLNPEFNERFEWDLPLDGT LRRKLDVSVKSNSSFMSRERELLGK VQLDLAEIDLQGAQWYDLMDDR DKGGS	False	True	3.54	2.062	2.11	4.405	1.484	3.827	1.584