

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
Q8C0D5	EFL1_MOUSE	Mus musculus	Elongation factor-like GTPase 1	25.685855	NaN	NaN	39627609	MVLSGVDKMIRLQKNTANIRNICVL AHVDHGKTTLADCLISSNGISSLRA GKLRYMDSREDEQVRGITMKSSAIS LHYAEGHEEYLINLIDSPGHVDFSS VSTAVRICDGCIIIVDAVEGVCPTQ AVLRQAWLENIRPVLVINKIDRLIVE LKFTTPQEAYSHLKNILEQINALTGTL FTSKVLEERAERETESQAKPHSEQG EQYVDWSAGLEDVDDSQLYFSPEQG NVVFTSAIDGWGFGIEHFARIYSQKI GIKKEVLLKTLWGDYYINMKAKKIM KVDQAKGKKPLFVQLILENIWSLYDA VLKDKKEKIDKIVTSLGLKIGAREAR HSDPKVQINAICSQWLPISHAVLAM VCHKLPSPLDMTSEKLLCTGSQ TFESLPPETQALKAAMKCGSEDTA PVIIIFVSKMFAVDVKALPQNKPRPLT QEEMAQRREARORHAEKLAQAQ QTSQGPTQDGALETSPHEDEPRGD EPDVASVSRQPVSQEESQEAFAFA RVFSGIARRGKIFVLGPKYSPVDFL QRVPLGFSAPLEDLPPVPHMACCTL ENLYLLMGRELEDLEEVPPGNVLI GGLQDFVLKATLCSLPSCPPFIPLN FEATPIVRVAVEPKHPSEMPQLVKG MKLLNQADPCVQVLIQETGEHVLT AGEVHLQRCLDDLRFERFAKIHISVSE PIIPFRETTIKPKVDMVNEEIGRQQ KVAVIHQTKEEQSKIPEGIHVSDGL ITIPTPNKLATLSVRAIPLPEEVTRILE ENSDLIRSMELTSSLNEGRNTQAI HQKTQEKIWEFKGKLEKHLTGRKW RNTVDQIWSFGPRKCGPNILVSRSE DFQNSVWSGPAGRESKEASRFRDF GNSIVSGFQLATLSGPMCEEPLMGV CFVLEKWELNKCAEQGASDKQHQQ QCDLAGEGQGGKTC HVGDENQEQ QDVCSEPFEEETSQKGDSVIDCYGPF SGQLIATMKEACRYALQVKPQLMA AMYTCDIMATSDVLGRVYAVLSKRE GRVLQEEMKEGTD MFIIKAVLPVAE SFGFADEIRKRTSGLASPQLVFSHW EVIPSDPFVWPTTEEEYLHFGEKADS ENQARKYMNAVRKRKGLYVEEKIVE HAEKQRTLSKNK	False	False	2.507	2.024	1.067	0.83	False	False	False