

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
Q3TH57	Q3TH57_MOUSE	Mus musculus	Transcription elongation regulator 1	26.09834	NaN	NaN	36064721	MAERGGDGGEGERFNPGLRMAQ QQALRFRGPAPPNAVMRGPPPLM RPPPPFGMMRGPppppRPPFRPPF DPNMPPMPPPGGIPPPMGPPHLQRP PFMPPPMGAMPPPGMMFPPGMPP GTAPGAPALPPTTEIIVENKTPDGK VYYNARTRESAWTKPDGVKVIQOS ELTPMLAAQAQVQAQAQAQAQAQA QAQAQAQAQAQAQAQAQAQAQAQA QAQAQAQAQAQAQAQAQAQAQAQA QAQAQAQAQVQAQAVGAPTPTTSSP APAVSTSTPTSTPSSTTATTTTATSV QTVSTPTTQDQTPSSAVSATPTVSV SAPAPTATPVQTVQPHQTLPPAVP HSVPQAAAIPAFPPVMVPPFRVPLP GMPIPLPGVAMMQIVSPPYKTVAT TKTGVLPGMAPPVPMIHPQVAIAAS PATLAGATAVSEWTEYKTADGKTYYY NNRTLESTWEKPQELKEKELDEKI KEPIKEASEEPLPMETEEEDPKEEPV KEIKEEPKEEEMTEEEKAAQKAPV ATTPIGTPWCVVWTGDERVFFYNP TTRLMSWDRPDDLIGRADVDKIIQEP PHKKGLEDKMLRHPAPTMLSIQK WQFSMSAIKEEQELMEEMNEDEPI KAKKRKRDDNKDIDSEKAAAMEAEI KAARERAIVPLEARMKQFKDMLLER GVSAFSTWEKELHKIVFDPYLLLN KERKQVFDQYVKTRAEERREKKNK IMQAKEDFKKMMEEAKFNPRATFS EFAAKHAKDSRFKAIEKMKDREALF NEFVAAARKKEKEDSKTRGEKIKSD FFELLSNHLLDSQSRWSKVVDKVE SDPRYKAVDSSSMREDLFKQYIEKIA KNLDSEKEKELERQARIEASLRERE REVQKARSEQTKRIDREREQHKREE AIQNFKALLSDMVRSSDVSWSDTRR TLRKDHRWESGSLLEEEKEKLFNE HIEALTKKREHFRQLLDETSAILT STWKEVKKIIEKEDPRCIKFSSDRKK QREFEYIRDKYITAKADFRLLKET KFITYRSKLIQESDQHLKDVEKILQ NDKRYLVLDVPEERRKLIVAYVDDL DRRGPPPTASEPTRRSTK	None	None	None	None	None	None	None	None	

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