	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q2TBA4	Q2TBA4_MOUSE	Mus musculus	NaN	27.829469	S437	NaN	21606357	MQFGTRTTAAEPGFMGTWQNADTN LLFRMSQQAIRCTLVNCTCECFQPG KINLRTCDQCKHGWVAHALDKLST QHLYHPTQVEIVQSNVVFDISSLMLY GTQAVPVRLKILLDRLFSVLKQEEVL HILHGLGWTLRDYVRGYILQDAAGK VLDRWAIMSREEEIITLQQFLRFGET KSIVELMAIQEKEGQAVAVPSSKTDS DIRTFIESNNRTRSPSLLAHLENSNP SSIHHFENIPNSLAFLLPFQYINPVS APLLGLPPNGLLLEQPGLRLREPSIS TQNEYNESSESEVSPTPYKSDQTPN RNALTSITNVEPKTEPACVSPIQNSA PVSDLSKTEHPKSSFRIHRMRRMGS ASRKGRVFCNACGKTFYDKGTLKIH YNAVHLKIKHRCTIEGCNMVFSSLR SRNRHSANPNPRLHMPMLRNNRD KDLIRATSGAATPVIASTKSNLTLTSP GRPPMGFTTPPLDPVLQNPLPSQLV FSGLKTVQPVPPFYRSLLTPGEMVSP PTSLPTSPIIPTSGTIEQHPPPPSEPIV PAVMMGTHEPSADLAPKKKPRKSS MPVKIEKEIIDTADEFDDEDDDPND GGTVVNDMSHDNHCHSQDEMSPG MSVKDFSKHNRTRCISRTEIRRADS MTSEDQEPERDYENESESSEPKLGE ESMEGDEHLHSEVSEKVLMNSERP DENHSEPSHQDVIKVKEEFTDPTYD MFYMSQYGLYNGGGASMAALHESF TSSLNYGSPQKFSPEGDLCSSPDPKI CYVCKKSFKSSYSVKLHYRNVHLKE MHVCTVAGCNAAFPSRRSRDRHSA NINLHRKLLTKELDDMSLDSSQPSL SKDLRDEFLMKIYGAQHPLGLDGRE DASSPAGTEDSHLNGYGRGMAEDY MVLDLSTTSSLQSSSSVHSSRESDA GSDEGILLDDIDGASDSGESTHKAEA PTLPGSLGAEVSGSLMFSSLSGSNG GIMCNICHKMYSNKGTLRVHYKTV HLREMHKCKVPGCNMMFSSVRSR NRHSQNPNLHKNIPFTSID	None	None	None	None	None	None	None	None	None