

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
E0CYT1	E0CYT1_MOUSE	Mus musculus	non-specific serine/threonine protein kinase	29.185005	NaN	NaN	22645316	MDGDGRRDAPGALMEAGRGTGSA GMAEPRARAARLGPQRFLRRSVVES DQEEPPGLEAAETPSAQPPLQRR VLLCKTRRLIAERARGRPAAPAPAA PAAPPGSPSPSDPGPERAGTOEPS DPTTASAAAQVPDGGPRQEEAFAP TQEDAGTTEAKPEPGRARKDEPEEE EDDEDDLKAVATSLDGRFLKFDIEL GRGSFKTVYKGLDTETWVEVAWCEL QDRKLTCLERQRFKEEAEMKGLQ HPNIVRFYDFWESSAKGKRCIVLVE LMTSGTLKTYLKRFKVMKPKVLRWS CRQILKGLFLHTRTPPIHRDLKCD NIFITGPTGSVKIGDLGLATLKRSFA KSVIGTPEFMAPEMYEEHYDESVDV YAFGMCMLEMATSEYPYSECQNA QIYRKVTCGKIPASFEKVHDEPEKEI GECICKNEEREYEIKDLLSHAFFAED TGVVVELAEEDHGRKSTIALRLWVE DPKLLKGGPKDNGAIEFTFDLEKETP DEVAQEMIDSGFFHESDVKIVAKSIR DRVALIQWRREIWPALQSQEPKDS GSPDKARGLPAPLQVQVYHAQSGQ PGQPEPEPEADQHLLPPTLPASVTS LASDSTFDSGGSTVYSDSQSSQSS MVLSSLVDTAPTASCVCSPVSEGP GLTHSLPTLGAFAQPATVPGLSVGPV PPPARPPLQQHFPSSMSFTPVLP PSTPVPTGPSQAPPVQVLAPQPM GTVPVPSHLPPYLAPTSQVVAQAQL KPLQMPQPLQPLAQVPPQMPQMP VVPITPLTGLDGLPQTLTDLPAANV APVPPPQYFSPAVILPSLTTPLTSPA LPMQAVKLPHPGTP LAVPCQTIVP NAPAAIPLAVAPQGVAAALSIHPAVA QIPAAQVYPAAFPMVPGDIPSPHH TVQSLRATPPQLASPVPPQVQPSVI HLPQAAPTAASGTQVLLGHPPSYT ADVAAPVSAVSLPPAVLSPPLDPTLL PTVPDLLPKVPSSLAPTVVAASQSA AQTSSLLPTNPPLPTGPAVAGPCPA VQLMVEVAQEEQVSQDKPPGPPQSS ESFGGSDVTSGRDLSDSCEGTFGGG RLEGRTARKHHRRSTRARSRQERAS RPRLTILNVCNTGDKMVECCQLETHN HKMVTFKFDLDGDAPDEIATYMVEH DFILPAERETFIEQMKDVMDKAEDM LSEDTDADHGS DTGTSPPHLGTCGL ATGEENRQSQANAPVYQNVLHTG KRWFIIICPVAEHPATDTSSESPPLPL SSLQPEASQDPAPYPDQLSLTDKPSF PAAQQLLSQAGSSNPPGGASAPLAP SSPPVTVIPAAPATSTVPESAAGTA MQAGGPGTHQGPASVHETLQPLAE TRSAQTAQPLSTGQGPCPTALEAS RCSTGLGEPSTREVSTQGEPLPASV PEPSPPTGATQSVPGQPPPLPITVG AISLAAPQLPSPPLGPTAPPPPSALE SDGEGPPPRVGFVDNTIKSLDEKLRT LLYQEHVPTSSASAGTPMEASDRDF TLEPLRGDLPSALS DKTPSLTQQTQP SLEKSETAPAGWALAQREQGASSPM TAESSSNTLGCSDAGQVADSST APSVQDASGSSVPTHMDPKDQNSS VPREALAAPMQSGPGSFTVGSQAQL RGARDSGSPHKRPGQDQNSPAKTV GRFVSVSTQDEWTLASPHSLRYSAP PDVYLDEIPSSPEVKLAVRRVQTASSI EVGVEEPASSDSGDERPRRRSQVQK QSSLPGTGGVASDFVKKATAFLHRS SRAGSLGPETPSRAGVKVPTISITSFH SQSSYSSDNDSEFEDADIKKELRSL REKHLKEISELQSQKQIEALYRRL GKPLPPNVGFFHTAPP MGRRRKTSK SKLKAGKLLNPLVQQLKVVASSTGH LSDSSRGPPTKDPAHASTPPGTKAV QTQPCSVRASLSTDICGLASDGG	None	None	None	None	None	None	None		

