	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasm reticulum
0	Q61194	P3C2A_MOUSE	Mus musculus	Phosphatidylinositol 4-phosphate 3-kinase C2 domain-containing subunit alpha	29.800457	S251;S1098	S60;S261;S328;S339;S630;S1281;S1553	30059200;36852467;33300544;35822049	MAQISNNSEFKQCSSSHPEPIRTKD VNKAEALQMEAEALAKLQKDRQMT DSPRGFELSSSTRQRTQGFNKQDYD LMVFPELDSQKRAVDIDVEKLTQAE LEKILLDDNFETRKPPALPVTPVLSP SFSTQLYLRPSGQRGQWPPGLCGPS TYTLPSTYPSAYSKQATFQNGFSPRM PTFPSTESVYLRLPGQSPYFSYPLTPA TPFHPQGSLPVYRPLVSPDMAKLFE KIASTSEFLKNGKARTDLEIANSKAS VCNLQISPKSEDINKFDWLDLDPLS KPKVDYVEVLEHEEEKKDPVLLAED PWDAVLLEERSPSCHLERKVNGKSL SGATVTRSQSLIIRTAQFTKAQGQVS QKDPNGTSSLPTGSSLLQEFEVQND EVAAFCQSIMKLKTKFPYTDHCTNP GYLLSPVTVQRNMCGENASVKVSIEI EGLQLPVTFTCDVSSTVEIIIMQALC WVHDDLNQVDVGSYILKVCGQEEVL QNNHCLGSHEHIQNCRKWDTEIKL QLLTLSAMCQNLARTAEDDEAPVDL NKYLYQIEKPYKEVMTRHPVEELLD SYHYQVELALQTENQHRAVDQVIKA VRKICSALDGVETPSVTEAVKKLKRA VNLPRNKSADVTSLSGSDTRKNSTK GSLNPENPVQVSMDHLTTAIYDLLR LHANSSRCSTACPRGSRNIKEAWTA TEQLQFTVYAAHGISSNWVSNYEKY YLICSLSHNGKDLFKPIQSKKVGTYK NFFYLIKWDELIIFPIQISQLPLESVL HLTLFGVLNQSSGSSPDSNKQRKGP EALGKVSLTLFDFKRFLTCGTKLLYL WTSSHTNSIPGAIPKKSYVMERIVLQ VDFPSPAFDIIYTSPQIDRNIIQQDKL ETLESDIKGKLLDIIHRDSSFGLSKE DKVFLWENRYYCLKHPNCLPKILAS APNWKWANLAKTYSLLHQWPPLCP LAALELLDAKFADQEVRSLAVSWME AISDDELADLLPQFVQALKYEIYLNS SLVRFLLSRALGNIQIAHSLYWLLKD ALHDTHFGSRYEHVLGALLSVGGKG LREELSKQMKLVQLLGGVAEKVRQA SGSTRQVVLQKSMERVQSFFLRNKC RLPLKPSLVAKELNIKSCSFFSSNAM PLKVTMVNADPLGEEINVMFKVGED LRQDMLALQMIKIMDKIWLKEGLDL RMVIFRCLSTGRDRGMVELVPASDT LRKIQVEYGVTGSFKDKPLAEWLRK YNPSEEEYEKASENFIYSCAGCCVAT YVLGICDRHNDNIMLRSTGHMFHID FGKFLGHAQMFGSFKRDRAPFVLTS DMAYVINGGEKPTIRFQLFVDLCCQ AYNLIRKQTNLFLNLLSLMIPSGLPE LTSIQDLKYVRDALQPQTTDAEATIF FTRLIESSLGSIATKFNFFIHNLAQLR FSGLPSNDEPILSFSPKTYSFRQDGR IKEVSVFTYHKKYNPDKHYIYVVRILR EGHLEPSFVFRTFDEFQELHNKLSII FPLWKLPGFPNRMVLGRTHIKDVAA KRKIELNSYLQSLMNASTDVAECDL VCTFFHPLLRDEKAEGIARSAGAVPF SPTLGQIGGAVKLSVSYRNGTLFIMV MHIKDLVTEDGADPNPYVKTYLLPD THKTSKRKTKISRKTRNPTFNEMLV YSGYSKETLRQRELQLSVLSAESLRE NFFLGGITLPLKDFNLSKETVKWYQ LTAATYL	None	None	None	None	None	None