

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q61194	P3C2A_MOUSE	Mus musculus	Phosphatidylinositol 4-phosphate 3-kinase C2 domain-containing subunit alpha	19.638248	S251;S1098	S60;S261;S328;S339;S630;S1281;S1553	35822049;30059200;36852467;33300544	MAQISNNSEFKQCSSSHPEPIRTKD VNKAERALQMEAEALAKLQKDRQMT DSPRGFELSSSTRQRTQGFNKQDYD LMVFPELTSQKRAVDIDVEKLTQAE LEKILLDDNFETRKPALPVPVLS SFSTQLYLRPSGQRGQWPPGLCGPS TYTLPSTYPSAYSQATFQNGFSPRM PTFPSTESVYLRLPGQSPYFSYPLTPA TPFHPPQGSPLVYRPLVSPDMAKLFE KIASTSEFLKNGKARTDLEIANSKAS VCNLQISPKSEDINKFDWLDLPLS KPKVDYVEVLEHEEEKDPVLLAED PWDVAVLLEERSPSCHLERKVNKSL SGATVTRSQSLIIRTAQFTKAQGQVS QKDPNGTSSLPTGSSLLQEFEVQND EVA AFCQSIMKLT KFPYTDHCTNP GYLLSPVTVQRNMCGENASVKVSI EGLQLPVTFTCDVSSTVEIIMQALC WVHDDL NQVDVGSYILKVCQEEVL QNNHCLGSHEHIQNCRKWDTEIKL QLLTL SAMCQNLARTAEDDEAPVDL NKYLYQIEKPYKEVMTRHPVEELLD SYHYQVELALQTENQHRAVDQVIKA VRKICSALDGVETPSVTEAVKCLKRA VNLPRNKSADVTSLSGSDTRKNSTK GSLNPENPVQVSMHLLTAYDLLR LHANSSRCSTACPRGSRNIKEAWTA TEQLQFTVYAAHGISSNWVSNEYKY YLICSLSHNGKDLFKPIQSKKVGTYK NFFYLIKWDELIFPIQISQLPLESVL HLTLFGVLNQSSGSSPDSNKQRKGP EALGKVSLTLDFKRF LTCGTKLLYL WTSSTHTNSIPGAIPKKS YVMERIVLQ VDFPSPAFDIYTSPQIDRNIIQQDKL ETLES DIKGLLDIIHRDSSFGLSKE DKVFLWENRYCLKHPNCLPKILAS APNWKWANLAKTYSLLHQWPP LCP LAALELLDAKFADQEVRS LAVSWME AISDDELADLLPQFVQALKYEIY LNS SLVRFLLSRALGNIQIAHSLYWLLKD ALHDTHFGSRYEHVLGALLSVGGKG LREELSKQMKLVQLLGGVAEKVRQA SGSTRQVVLQKSMERVQSFFLRNKC RLPLKPSLVAKELNIKSCSFFSSNAM PLKVTMVNADPLGEEINVMFKVGED LRQDMLALQMIKIMDKIWLKEGLDL RMVIFRCLSTGRDRGMVELVPASDT LRKIQVEYGVTSF KDKPLAEWLRK YNPSEEEYEKASENFIYSCAGCCVAT YVLGICDRHNDNIMLRSTGHMFHID FGKFLGHAQMFGSFKRDRAPFV LTS DMAYVINGGEKPTIRFQLFVDLCCQ AYNLIRKQTNLFLNLLSLMIPSGLPE LTSIQDLKYVRDALQPQTDAEATIF FTRLIESSLGSIATKFNFFIHNLAQLR FSGLPSNDEPILSFSPKTYSF RQDGR IKEVSVFTYHKKYNPDKHYIYVVRILR EGHLEPSFVFRTFDEFQELHNKLSII

