

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
Q9R0K7	AT2B2_MOUSE	Mus musculus	Plasma membrane calcium-transporting ATPase 2	24.474581	NaN	S18;T1094;S1133;T1143;S1156;S1166	33300544	MGDMTNSDFYSKNQRNESHGGE FGCTMEELRSLMELRGTEAVVKKKE TYGDTEAICRRLKTSPEVGLPGTAPD LEKRKQIFGQNFIPPKPKTFLQLVW EALQDVTLIILEIAAIIISLGLSFYHPPG ESNEG CATAQGGAEDEGEAEAGWIE GAAILLSVICVVLVTA FN DWSKEKQF RGLQSRIEQE QKFTVVRAGQVVQIPV AEIVVGDIAQIKYGDLLPADGLFIQG NDLKIDESSLTGESDQVRKSVDKDP MLLSGTHVMEGSGRMVVAVGVNS QTGIIFTLLGAGGEEEEKDKKAKQQ DGAAMEMQPLKSAEGGDADDKCK ANMHKKEKSVLQGLTKLAVOIGKA GLVMSAITVHILVLYFTVDTFVFNKPP WLTECTPVVYQYFVKFFIIGVTVLV AVPEGLPLAVTISLAYSVKMMKDN NLVRHLDACETMGNATAICSDKTGT LTTNRMTVVQAYVDVHYKEIPDPS SINAKTLELLVNAIAINSAYTTKILPP EKEGALPROVGNKTECGLLGFVLDL RQDYEPVRSQMPEEKLYKYTFNSV RKSMSVVIKMPDESFRMYSKGASEI VLKCKKILSGAGEARVFRPRDRDE MVKKVIEPMACDGLRTICVAYRDFP SSPEPDWDNENDILNELTICVVGIE DPVRPEVPEAIRKCORAGITVRMVTG DNINTARAIKCGIHPGEDFLCLEG KEFNRRIRNEKEIEQERIDKIWPKL RVLARSSPTDKHTLVKGIIDSTHTEQ RQVAVTGDGTNDGPALKKADVGFA MGIAGTDVAKEASDIILTDNFSSIV KAVMWGRNVYDSISKFLQFQLTVNV VAVIVAF TGACITQDSPLKAVQMLW VNLIMDTFASLALATEPPTETLLLRK PYGRNKPLISRTMMKNILGHAVYQL TLIFTLLFVGEKMFQIDSGRNAPLHS PPSEHYTHFNTFVMMQLFNEINARK IHGERNVFDGIFRNPIFCTIVLGTFAI QIVIVQFGGKPFSCSPLQLDOWMWC IFIGLGELVWGQVIATIPSTRKFLKE AGRLTQKEIPEEELNEDVEIDHAE RELRRGOILWFRGLNRIQTQIRVKA FRSSLYEGLEKPESTRSIHNFMAHP EFRIEDSQPHIPLIDDTLEEDAALK QNSSPPSSLNKNNSAIDSGINLTTDT SKSATSSSPGSPHISLETSL	None	None	None	None	None	None	None		