	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plas men
0	Q923T9	KCC2G_MOUSE	Mus musculus	Calcium/calmodulin-dependent protein kinase type II subunit gamma	27.053486	NaN	T287;T306;T307;S311;S334;S349;S352;S455	28528544;33300544	MATTATCTRFTDDYQLFEELGKGAF SVVRRCVKKTSTQEYAAKIINTKKLS ARDHQKLEREARICRLLKHPNIVRL HDSISEEGFHYLVFDLVTGGELFEDI VAREYYSEADASHCIHQILESVNHIH QHDIVHRDLKPENLLLASKCKGAAV KLADFGLAIEVQGEQQAWFGFAGTP GYLSPEVLRKDPYGKPVDIWACGVIL YILLVGYPPFWDEDQHKLYQQIKAGA YDFPSPEWDTVTPEAKNLINQMLTI NPAKRITADQALKHPWVCQRSTVAS MMHRQETVECLRKFNARRKLKGAI LTTMLVSRNFSAAKSLLNKKSDGGV KKRKSSSSVHLMPQSNNKNSLVSPA QEPAPLQTAMEPQTTVVHNATDGIK GSTESCNTTTEDEDLKVRKQEIIKIT EQLIEAINNGDFEAYTKICDPGLTSF EPEALGNLVEGMDFHKFYFENLLSK NSKPIHTTILNPHVHVIGEDAACIAYI RLTQYIDGQGRPRTSQSEETRVWHR RDGKWLNVHYHCSGAPAAPLQ	False	False	3.742	2.957	2.821	3.927	1.407	2.52