

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
Q9Z2W9	GRIA3_MOUSE	Mus musculus	Glutamate receptor 3	26.328835	NaN	T871;T881	33300544;34678516	MGQSVLRAVFFLVLLGSHGGFP NTISIGGLFMRNTVQEHSAFRFAVQ LYNTNQNTTEKPFHLNHYVDHLDS SNSFSVTNAFCSQFSRGVYAIFFGYD QMSMNTLTSFCGALHTSFVTPSFPT DADVQFVIQMRPALKGAILSLGYYK WEKVVLYDTERGFSILQAIMAAV QNNWQVTARSVGNIKDIQEFRRIIEE MDRRQEKRYLIDCEVERINTILEQVV ILGKHSRGYHYMLANLGFDTIVLER VMHGGANITGFQIVNENPMVQQF IQRWVRLDEREFPEAKNAPLKYTSA LTHDAILVIAEAFRYLRRQRVDVSRR GSAGDCLANPAVPWSQGIDIERALK MVVQVQGMTGNIQFDTYGRRTNYTID VYEMKVSRSRKAQYWNEYERFVPPFS DQQISNDSSSENRTIVVTTILESPPV MYKKNHEQLEGNERYEGYCVDLAY EIAKHVRIKYLIVGDKYKARDPE TKIWNMVGELVYGRADIAVAPLTTT LVREEVIDFSKPFMSLGISIMIKKPQ KSKPGVFSFLDPLAYEIVMCIWFAYI GVSVVLFVSRFSPYEWHLNNEE PRDPQSPDPNNEFGIFNSLWFSLG AFMQGCDISPRSLSGRIVGGVWWF FTLIHSSYTANLAAFLTVERMVSPIES AEDLAKQTEIAYGTLDSGSTKEFFRR SKIAVYEKMWSYMKSAEPSVFTKTT ADGVARVRKSKGKFAFLESTMNEY IEQRKPCDTMKVGGNLDKGYGVAT PKGSALRTPVNLAVLKLSEQGILDKL KNKWWYDKGECGAKDSGSKDKTSA LLSNVAGVFYILVGGGLGLAMMVALI EFCYKSRAESKRMKLTKNQNFKPA PATNTQNYATYREGYNNVGTESVKI	False	True	1.601	1.87	1.566	1.635	1.279	5.0	4.429