

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
Q61627	GRID1_MOUSE	Mus musculus	Glutamate receptor ionotropic, delta-1	46.009546	T521;T523;T942;S944	NaN	33300544;34418053;40885482;22645316	MEALTLWLLPWICQCVTVRADSIHI GAIFEENAAKDDRVFQLAVSDLSLN DDILOSEKITYSIKVIENNPFAVQVE ACDLMTQGILALVSTGCASANALQ SLTDAMHHPHLFVQRNPPGSPRTAC HLNPSPDGEAYTLASRPPVRLNDVM LRLVTELRWQKFMFYDSEYDIRGL QSFLDQASRLGLDVSLQKVDKNISH VFTSLFTTMKTEELNRYRDTLRRAIL LLSPQGAHSFINEAVETNLASKDSH WVFNVEEISDPEILDVHLSALGRMT VVRQIFPSAKDNQKCMRNNHRISSL LCDPOEGYLQMLQISNLYLYDSVLM LANAFHRKLEDRKWHSMASLNCIR KSTKPWNGGRSMLDTIKKGHITGLT GVMEFREDSSNPYVQFEILGTTYSET FGKDMRKLATWDSEKGLNGSLQER PMGSRLQGLTLKVVTVLEEPFVMVA ENILGQPKRYKGFSDVLDALAKALG FKYEIYQAPDGRYGHQLHNTSWNG MIGELISKRADLAISAITITPERESVV DFSKRYMDYSVILIKKPEEKISIFSL FAPDFAVWACIAAAIPVVGVLIFVL NRIQAVRSQSATQPRPSASATLHSAI WIVYGAFVQGGESSVNSVAMRIVM GSWWLFTLIVCSSYTANLAFLTVSR MDNPIRTFQDLKQLEMSYGTVRDS AVYEYFRAKGTNPLEQDSTFAELWR TISKNGGADNCVSNPSEGIRKAKKG NYAFLWDVAVVEYAALTDDDCSVTV IGNSISSKGYGIALQHGSFYRDLFSQ RILELQDTGDLVLLKQKWWPHTGR CDLTSHSSTQTEGKSLKLSFAGVF CILAIGLLLACLVALELWVNSNRC HOETPKEDKEVNLEQVHRRINSLM DEDIAHKQISPASIELSALEMGGGLAP SQALEPTREYQNTQLSVSTFLPEQSS HGTSRTLSSGPPSNLPLPLSSSATMP SIQCKHRSPNGGLFRQSPVKTPIPMS FQVPVGGVLPALDTSHGTSI	False	True	0.916	1.484	1.172	1.174	False	4.70