

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
0 Q61626	GRIK5_MOUSE	Mus musculus	Glutamate receptor ionotropic, kainate 5	24.474581	NaN	NaN	33300544	MPAELLLLLIVAFANPSCQVLSLRM AAILDDQTVCGRGERLALALAREQIN GIIEVPAKARVEVDIFELQRDSQYETT DTMCQILPKGVVSVLGPSSPASAST VSHICGEKEIPHIKVGPEETPRLOYL RFASVSLYPSNEDVSLAVSRILKSFN YPSASLICKAKAECLLRLEELVVRGFLIS KETLSVRMLDDSRDPTPLLKEIRDD KVSTIIDANASISHLVLRKASELGMT SAFYKYILTTMDFPILHLDGIVEDSS NILGFSMFNTSHPPYFPEFVRSLNMS WRENCEASTYPGPALSAAALMFDVAV HVVVSAVRELNRSQEIGVKPLACTSA NIWPHGTSMLNMYLRMVEYDGLTGR VEFNKSGQRTNYTLRILEKSRQGHR EIGVWYSNRTLAMNATTLNINLSQT LANKTLVVTTILENPYVMRRPNFQA LSGNEREFEGFCVDMLELAELLRFR YRLRLVEDGLYGAPENGSWTGMV GELINRKADLAVAAFTITAEREKVIDF SKPFMTLGISILYRVHMGRKPGYFSF LDPFSPAVWLFMLLAYLAVSCVLF AARLSPYEWYNPHPCLRARPHILEN QYTLGNLWFPVGGFMQQGSEIMP RALSTRCVSGVWWAFTLIHSSYTAN LAAFLTIVQRMEVPVESADDLADQTN IEYGTIHAGSTMFFQNSRYQTYQR MWNMYQSKQPSVFKSTEEGIARV LNSRYAFLESTMNEYHRRNLNCLT QIGGLDTKGYGIGMPLGSPFRDEIT LAILQLQENNRLEILKRKWEGGRC PKEEDHRAKGLGMENIGGIFVVIC GLIIAVFVAVMEFIWSTRRSAESEEV SVCQEMLQELRHAVSCRKTSRSTRR RRPGGPRALLSLRAVREMRSLNGK LYSAGAGDAGAHGGPQRLLDDPGP PGGPRPQAPTCTHVRVCQECRRIQ ALRASGAGAPPRGLGTPAEATSPPRP RPGPTGPRELTEHE	False	True	1.194	2.893	1.246	5.0	0.955	5.0	1.712