

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
P42264	GRIK3_RAT	Rattus norvegicus	Glutamate receptor ionotropic, kainate 3	23.439218	NaN	S869	38843836	MTAPWRRRLRSLVWEYWAGFLVCAF WIPDSRGMPhVIRIGGIFEYADGPNA QVMNAEEHAFRFSANIINRRNTLLP NTTLTYDIQRIHFHDSFEATKKACDQ LALGVVAIFGPSQGSCTNAVQSIENA LEVPHIQLRWKHHPLDNKDTFYVNL YPDYASLSHAIDLVLQSLKWRSATVV YDDSTGLIRLQELIMAPSRYNIRLKIR QLPIDSDDSRPLLKEMKRGREFRIIF DCSHTMAAQILKQAMAMGMMTEY YHFIFTTLDLYALDLEPYRYSGVNLT GFRILNVDNAHVSAIVEKWSMERLQ AAPRAESGLLDGVMMTDAALLYDAV HIVSVCYQRASQMTVNSLQCHRHK PWRFGGRFMNFIKEAQWELTGRIV FNKTSGLRTDFDLIISLKEDGLEKV GVWSPADGLNITEVAKGRGPNVTD LTNRSLIVTTLLEPFVFRKSDRTL YGNDRFEGYCIDLLKELAHILGFSYE IRLVEDGKYGAQDDKGQWNGMVKE LIDHKADLAVAPLTITHVREKAIDFS KPFMTLGVSILYRKPNGTNPSVFSFL NPLSPDIWMYVLLAYLGVSCVLFVIA RFSPEYWDYDAHPCNPGSEVENNF TLLNSFWFGMGSMLMQGSELMPKA LSTRIGGIWVFFTLIISSYTANLAAF LTVERMESPIDSAADLAKQTKIEYGA VKDGATMTFFKKSISTFEKMWAF MSSKPSALVKNNEEGIQRTLTADYA LLMESTTIEYITQRNCNLQIGGLIDS KGYGIGTPMGSPYRDKITAILQLEE DKLHIMKEKWWRGSGCPEEENKEA SALGIQKIGGIFIVLAAGLVLSVLVAV GEFIYKLRKTAEREQRSFCSTVADEI RFSLTQRRRLKHKPQPPMMVKTD VINMHTFNDRRLPGKDSMSCSTSLA PVFP	False	True	0.986	1.373	1.128	0.942	0.699	5.0	1.723