	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q925T6	GRIP1_MOUSE	Mus musculus	Glutamate receptor-interacting protein 1	25.775171	NaN	S43	34678516	MIAVSFKCRCQILRRLTKDESPYTKS ASQTKPPDGALAVRRQSIPEEFKGST VVELMKKEGTTLGLTVSGGIDKDGK PRVSNLRQGGIAARSDQLDVGDYIKA VNGINLAKFRHDEIISLLKNVGERVV LEVEYELPPVSVQGSSVMFRTVEVTL HKEGNTFGFVIRGGAHDDRNKSRPV VITCVRPGGPADREGTIKPGDRLLSV DGIRLLGTTHAEAMSILKQCGQEAT LLIEYDVSVMDSVATASGPLLVEVAK TPGASLGVALTTSVCCNKQVIVIDKIK SASIADRCGALHVGDHILSIDGTSME YCTLAEATQFLANTTDQVKLEILPHH QTRLALKGPDHVKIQRSDRQHPWD AWASNQCGVHTNHHHNTYHPDHC RVPALTFPKALPPNSPPAMVPSSSPT SMSAYSLSSLNMGTLPRSLYSTSPR GTMMRRRLKKKDFKSSLSLASSTVG LAGQVVHTETTEVVLTADPVTGFGIQ LQGSVFATETLSSPPLISYIEADSPAE RCGVLQIGDRVMAINGIPTEDSTFEE ANQLLRDSSITSKVTLEIEFDVAESVI PSSGTFHVKLPKKHSVELGITISSPSS RKPGDPLVISDIKKGSVAHRTGTLEL GDKLLAIDNIRLDNCSMEDAVQILQ QCEDLVKLKIRKDEDNSDEQESSGA IIYTVELKRYGGPLGITISGTEEPFDPI IISSLTKGGLAERTGAIHIGDRILAINS SSLKGKPLSEAIHLLQMAGETVTLKI KKQTDAQSASSPKKFPIPGHSGDLG DGEEDPSPIQKPGKLSDAYPSTVPSV DSAVDSWDGSGIDASYGSQGSTFQT SGYNYNTYDWRSPKQRTSLSPVPKP RSQTYPDVGLSNEDWDRSTASGFVG ASDSADAEQEENFWSQALEDLETC GQSGILRELEEKADRRVSLRNMTLL ATIMSGSTMSLNHEAPMARSQLGR QASFQERSSSRPHYSQTTRSNTLPS DVGRKSVTLRKMKQEIKEIMSPTPV ELHKVTLYKDSGMEDFGFSVADGLL EKGVYVKNIRPAGPGDVGGLKPYDR LLQVNHVRTRDFDCCLVVPLIAESG NKLDLVISRNPLASQKSIEQPALPSD WSEQNSAFFQQPSHGGNLETREPT NTL	None	None	None	None	None	None	None	None	None