

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q3UVX5	GRM5_MOUSE	Mus musculus	Metabotropic glutamate receptor 5	21.665133	NaN	S860;S1014;S1016	22645316;33300544;34678516;35822049	MVLLLLSVLLLKEDVRGSAQSSERR VVAHMPPGDIIIGALFSVHHQPTVDKV HERKCGAVREQYGIQRVEAMLHTLE RINSDPTLLPNITLGCIRDSCWWSA VALEQSIEFIRDLSISSEEEGLVRCV DGSSSFRSKKPIVGVIGPGSSSVAIQV QNLLQLFNIPQIAYSATSMDLSDKTL FKYFMRVVPDAQQARAMVDIVKRY NWTYVSAVHTEGNYGESGMEAFKD MSAKEGICIAHSYKIYSNAGEQSFDK LLKLRSHLPKARVVACFCEGMTVR GLLMAMRRLGLAGEFLLLGS DGWA DRYDVT DGYQREAVGGITIKLQSPDV KWFDDY LKLRPETNLRNPWFQEF WQHRFQCRLEGFAQENSKYNKTCN SSLTLRTHHVQDSKMGFVINAIYSM AYGLHNMQMSLCPGYAGLCDAMKP IDGRKLLDSLMTNFTGVSGDMILF DENGDS PGRYEIMNFKEMGKDYFD YINVGSWDNGELKMDDDDEVWSKKN NIIRSV CSEPCEKQIKVIRKGEVSC CWTCTPCKENEYVFDEYTCACQLG SWPTDDL TGC DLIPVQYLRWGDPEP IAAVVFAC LGLLATLFVTVIFIIYRDTP VVKSSSREL CYIILAGICLGYLCTFCLI AKPKQIYCYLQRIGIGLSPAMSYSALV TKTNRIARILAGSKKKICTKKPRFMS ACAQLVIAFILICIQLGIIVALFIMEPP DIMHDYPSIREVYLICNTTNLGVVTP LGYNGLLILSCTFYAFKTRNVPANFN EAKYIAFTMYTTCHWLAFVPIYFGSN YKIITMCF SVLSATVALGCMFVPKV YIILAKPERNVRS AFTTSTVVRMHVG DGKSSSAASRSSLVNLWKRRGSSG ETLRYKDRRLAQHKSEIECFTPKGS MGNGGRATMSSSNGKSVTWAQNE KSTRGQHLWQRLSVHINKKENPNQ TAVIKPFPKSTESRGQAGAGGGSG PGAAGAGSAGCTATGGPEPPDAGPK ALYDVAEAEERFPAAARPRSPSPIST LSHLAGSAGRTDDDAPSLHSETAAR SSSSQGS LMEQISSVTRFTANITEL NSMMLSTAAAPGPPGTPICSSYLIPK EIQLPTTMTTFAEIQLPAIEVTGGAQ PATGPPAQETPAGAEAAPGKPDLEE LVALTPSPFRDSVDSGSTTPNSPVS ESALCIPSSPKYDTLIIRDY TQSSSSL