

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
Q9WV18	GABR1_MOUSE	Mus musculus	Gamma-aminobutyric acid type B receptor subunit 1	24.474581	NaN	T872;T929	33300544	MLLLLLVPLFLRPLGAGGAQTPNVT SEGCGIHPWEGGIRYRGLTRDQVK AINFLPVDYIEIYVCRGEREVVGPVK RKCLANGSWTDMDTPSRCVTRICKS YLTLENGKVFLTGGLPALD GARVD FRCDPDFHLVGSRRSICSGQWSTP KPHCQVNRTPHSERRAVYIGALFPM SGGWPGGQACQPAVEMALEDVNSR RDILPDYELKLIHHD SKCDPGQATKY LYELLYNDPIKILMPCGSSVSTLVAE AARMWNLIVLSYGSSPALS NRQRF PTFFRTHPSATLHNPTRVKLFKFWG WKKIATIQQTTEVFTSTLDDLEERVK EAGIEITFRQSFSDPAVPVKNLKRQ DARIIVGLFYETEARKVFCVYKERL FGKKYVWFLIGWYADNWFKTYDPSI NCTVEEMTEAVEGHITTEIVMLNPA NTRSISNMTSQEFVEKLTKRKRHP EETGGFQEAPLAYDAIWALALANK TSGGGGRSGVRLEDFNYYNNQITTDQ IYRAMNSSFEVSGHVVFDA SCSR MAWTLIEQLQGGSYKKIGYDSTKD DLSWSKTDKWIGGSPADQTLVIKT FRFLSQKLFISVSVLSSLGIVLAVVCL SFNIYN SHVRYIQNSQP NLNLTAV GC SLALAAVPLGLDGYHIGRSQPPF VCQARLWLLGLGFLGYSMFTKIW WVHTVFTKKEKKEWRKLEPWKL YATVGLLVGMDILTAIWQIVDPLHR TIETFAKEPKEDIDVSILPQLEHCSS KKMNTWLGIFYGKGLLLLLGIFLAY ETKSVSTEKINDHRAVGMAYNVAV LCLITAPVTMILSSQDAAFASLAI VFSSYITLVVLFVPMRRLITRGEWQ SEAQDTMKTGSSTNNNEEKSRLL EKENRELEKIIAEKEERVSELRHQLQ SRQQIRSRRHPTPPDPSGGLPRGPS EPPDRLSCDGRVHLLYK	True	True	1.716	2.039	4.319	4.381	1.398	5.0	4.409