	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	Q9VR07	INE_DROME	Drosophila melanogaster	Sodium- and chloride-dependent GABA transporter ine	25.496522	NaN	NaN	33925313	MAENKDVSQVLNTTPDQVVINRAPP TGLAPLRHSQLSDSGAESCYEGDEQ ARLIRSSSSRAHKFIIIPATPGTPPGS GAVAGPLPFRQTSSTTSLAAMAAAL HKQSPLRHTSVRTRPSSEVLQPGQV LANQPTAAVSTSTVTFTIDDCNEEGD AIPAAASVPAPVTMPAISPTPATLTCT TTTTMAGDSVAVSPLSMELKSRLGS HHSSMRSVVSGYLPGLSDSSGNLVG GVSMATSGLQAPNPLYMQPQASLSG SSYHFHELAGNQIYSDVTSVRSLASI GIGSTDGRKLVIRRVPTTANELFDMV NPQTPPPLGVDDDDSYLDMSDETAQ LKPRQQHWANKMQFVLACIGYSVG LGNVWRFPYMCYKSGGGVFLVPYCI ILFICSIPLLFMELSVGQYTGRGPIGA LGQLCPLFKGAGLASVVVSFLMSTY YSVIIGYSIYYFFTSFKTEMPWIDCNN RWNTPDCWVPQRKGINASAPDTSRT PSEEFFENKVLQISGGLEYPGMMR WELFACLICAWLMVYFATWKSIKSS AKVRYFTATFPFVLIIILMVRAVTLDG AAEGLRFFFRPKWSELKNANVWINA ASQNFNSLGITFGSMISFASYNKYN NNILRDTVAVSAVNMITSLLVGIFAF STLGNLALEQNTNVRDVIGDGPGMI FVVYPQAMAKMPYAQLWAVMFFFM LLCLGLNSQFAIVEVVVTSIQDGFPR WIKRHLGYHEIVVLFVCVISCLFGMP NIIQGGIYYFQLMDHYAASVTIMFLA FCQMIAIAWFYGTGRLSKNVKQMTG KAPSFYLRSCWLVLGPCLLFAIWVLS LINYHEPTYHNGRYTYPDWAYGIGW MFASFSLICIPGYAVINFLRSSGDTF WERIRNTLRPNIYECKICGEHHCEH DYPEQEQFMLAQEMATVYKPTNPH LLNLGQKCGYNAMQASPSHAEAGG PCGQ	None	None	None	None	None	None	None	None	None