

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
P32037	GTR3_MOUSE	Mus musculus	Solute carrier family 2, facilitated glucose transporter member 3	27.478975	NaN	T232;S471;S482;T489	29187734;33300544	MGTTKVTPLVFAVTVATIGSFQFGY NTGVINAPETILKDFLNYTLERLED LPSEGLLTALWSLCVAIFSVGGMIGS FSVGLFVNRFGRRNSMLLVNLLAIIA GCLMGFAKIAESVEMLILGRLLIGIF CGLCTGFVPMYIGEVSPTALRGAFGT LNQLGIVVGILVAQIFGLDFILGSEEL WPGLLGLTIIPAILQSAALPFCPEspr FLLINKKEEDQATEILQRLWGTSDVV QEIQEMKDESVRMSQEKQVTVLELF RSPNYVQPLLSIVLQLSQQLSGINAV FYYSTGIFKDAGVQEPYATIGAGVVN TIFTVVSLFLVERAGRRTLHMIGLGG MAVCSVFM TISLLKDDYEAMSFVC IVAILIYVAFFEIGPGPIPWFIVAE LFS QGPRPAIAVAGCCNWTSNFLVGML FPSAAAYLGAYVFIIFAAFLIFLIFTF FKVPETKGRTFEDIARAFEGQAHSg KGPAGVELNSMQPVKETPGNA	False	True	2.504	2.569	2.859	2.32	1.477	5.0	2.621