

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0 P15626	GSTM2_MOUSE	Mus musculus	Glutathione S-transferase Mu 2	25.629171	NaN	S27;S44;S117	36288343	MPMTLGYWDIRGLAHAIRLLLEYTD TSYEDKKYTMGDAPDYDRSQWLSE KFKLGLDFPNLPYLIDGSHKITQSNA ILRYLARKHNLCGETEEERIRVDILE NQAMDTRIQLAMVCYSPDFEKKKPE YLEGLPEKMKLYSEFLGKQPWFAGN KVITYVDFLVYDVLQHRIFEPKCLDA FPNLKDFMGRFELGKKISDYMKSSR FLSKPIFAKMAFWNPK	True	False	4.713	2.695	1.84	1.433	0.567	1.499	1.58