

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
0 P48774	GSTM5_MOUSE	Mus musculus	Glutathione S-transferase Mu 5	27.312347	NaN	S5	28135057;33300544	MSSKSMVLGYWDIRGLAHAIRMLLE FTDTSYEEKRYICGEAPDYDRSQWL DVKFKLDLDFPNLPYLMGKKNITQ SNAILRYIARKHNMC GDTEEEKIRV DIMENQIMDFRMQLVRLCYNSNHE NLKPQYLEQLPAQLKQFSLFLGKFT WFAGEKLT FVDFLTYDVLQNRIFE PKCLDEFPNLKAFMCRFEALEKIAA FLQSDRFFKMPINNKMAKWGNKCL C	True	False	4.146	2.547	2.172	1.465	0.595	1.717	3.191