

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
Q9Z2V4	PCKGC_MOUSE	Mus musculus	Phosphoenolpyruvate carboxykinase, cytosolic [GTP]	22.380232	NaN	S19;S90;S118;S286	37507081	MPPQLHNGLDLFSKVIQGLDLSLPQ AVRKFEVGNALCQPEYIHCIDGSE EEYGQLLAHQEVEGIRKLLKDYDNC WLALDPRDVARIESKTVIITQEQRD TVPIPKTGLSQLGRWMSEEDFEKAF NARFPGCMKGRTMYVIPFSMGPLG SPLAKIGIELTDSPPYVVASMRIMTRM GISVLEALGDGEFIKCLHSGCPLPL KKPLVNNWACNPELTLIAHLPDRRE IISFGSGYGGNSLLGKKCFALRIASRL AKEEGWLAEHMLLIGITNPEGKKKY LAAAFPSACGKTNLMMNPSLPGW KVECVGDDIAWMKFDAQGNLRIN PENGFFGVAPGTSVKTNPNAIKTIQK NTIFTNVAETSDGGVYWEIDEPLAP GVTITSWKNKEWRPQDAEPCAHPN SRFCTPASQCPIDPAWESPEGVPIE GIIFGRRRPEGVPLVYEALSWQHGV FVGAAMRSEATAAAEHKGIIMHDP FAMRPFPGYNFGKYLAHWLSMAHR PAAKLPKIFHVNWFRKDKDGKFLWP GFGENSRVLEWFMFGRIEGEDSAKLT PIGYIPKENALNLKGLGGVNVVEELFG ISKEFWEKEVEIDRYLEDQVNTDLP YEIERELRALKQRISQM	True	True	5.0	3.326	2.714	4.435	1.315	2.14	2.393