

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P35558	PCKGC_HUMAN	Homo sapiens	Phosphoenolpyruvate carboxykinase, cytosolic [GTP]	27.26403	T102	S19;S90;S118;T178;S286	31492838	MPPQLQNLNLSAKVVQGSLSLDP QAVREFLENNAEELCQPDHIIICDGS EEENGRLLGQMEEEGILRRLKKYDN CWLALTDPRDVARIESTVIVTQEQR DTVPIPKTGLSQLGRWMSEEDFEKA FNARFPGCMKGRMYVIFPSMGPL GSPLSKIGIELTDSFYVVASMRIMTR MGTPVLEAVGDGEFVKCLHLSVGCPL PLQKPLVNNWPCNPELTLIAHLDPDR REIISFGSGYGGNSLLGKKCFALRMA SRLAKEEGWLAEHMLLIGITNPEGE KKYLAAPFACGKTNLAMMNPSLP GWKVECVGDDIAWMKFDAQGHLRA INPENGFFGVAPGTSVKTNPNAIKTI QKNTIFTNVAETSDDGGVYWEIDEP LASGVTTTSWKNKEWSSSEDEGEPCAH PNSRFCTPASQCPIIDAAWESPEGVP IEGIIFGGRRPAGVPLVYEALSQWQHG VFVGAAMRSEATAAAEHKGIIMHD PFAMRPFYFGYNFGKYLAWLSMAQ HPAAKLPKIFHVNWFRKDKGKFL WPGFGENSRVLEWFMFNRIDGKAST KLTPIGYIPKEDALNLKGLGHINMM ELFSISKEFWKEVEDIEKYLEQVN ADLPCIEIEREILALKQRISQM	True	True	5.0	3.295	2.775	5.0	1.534	2.113	4.468