

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	pl m
Q16822	PCKGM_HUMAN	Homo sapiens	Phosphoenolpyruvate carboxykinase [GTP], mitochondrial	12.241267	S34	T42;S115;T196;S304	34229054;30379171;30397120;28510447;34725712;33465208;38665916	MAALYRPGRLRNWHGLSPLGWPS RSIQLRLVLSGDLGQLPTGIRDFVEH SARLCQPEGIHCDCGTEAENTATLTL LEQQGLIRKLPKYNNCWLARTDPKD VARVESKTVIVTPSQRDTVPLPPGGA RGQLGNWMSPADFORAVDERFPGC MQGRTMYVLPFSMGPVGSPLSRIGV QLTDSAYVVASMRIMTRLGTFVLQA LGDGDFVKCLHSVGQPLTGGQEPVS QWPCNPEKTLIGHVPDQREIISFGSG YGGNSLLGKKCFALRIASRLARDEG WLAEHMLILGITSAGKRYVAAAFF SACGKTNLAMMRPALPGWKVECVG DDIAWMRFDSEGLRAINPENGFFG VAPGTSATTNPNAMATIQSNTIFTNV AETSDGGVYWEQIDQPLPPGVTVTS WLGKPPWPGDKEPCAHPNSRFCAP ARQCPIMDPAWEAPEGVPIDAIIFGG RRPKGVPLVYEFNWRHGVVFGSA MRSESTAAAEHKGKIIIMHDPFAMRP FFGYNFGHYLEHWLSMEGRKGAQL PRIFHVNWFRRDEAGHFLWPGFGE NARVLDWICRRLEGEDSARETPIGL VPKEGALDLSGLRAIDTTLFSLPKD FWEQEVDRIRSYLTEQVNQDLPKEV LAELEALERRVHKM	True	False	3.765	2.888	4.978	2.079	1.196	2.1