

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	extr regi
P11498	PYC_HUMAN	Homo sapiens	Pyruvate carboxylase, mitochondrial	15.337785	NaN	T1003	38665916;35083852;34105348;33214551;30397120;32119511;34725712;37217939;28510447	MLKFRVTVHGGLRLLGIRRTSTAPAAS PNVRRLEYKPIKKVMVANRGEIAIRV FRACTELGIRTVAIYSEQDTGQMHR QKADEAYLIGRGLAPVQAYLHIPDIK VAKENNVDAVHPGYGLSERADFAQ ACQDAGVRFIGPSPEVVRKMGDKVE ARAIAlAAGVPVPGTDAPIITSLHEAH EFSNTYGFPIIFKAAYGGGGGRGMRV VHSYEELEENYTRAYSEALAAFNGG ALFVEKFIKPRHIEVQILGDQYGNIL HLYERDCSIQRRHQKVVIEIAPAAHL DPQLRTRLTSDSVKLAKQVGYENAG TVEFLVDRHGKHYFIEVNSRLQVEH TVTEEITDVLVHAQIHVAEGRSLPD LGLRQENIRINGCAIQCRVTTEDPAR SFQPDGTGRIEVFRSGEGMGIRLDNA SAFQGAIVSPHYDSLVLVAVIAGKDH PTAATKMSRALAEFRVRGVTNIAFL QNVLNNQQFLAGTVDTFIDENPEL FQLRPAQNRAQKLLHLYLGHVMVNG PTTPIPVKASPSPTDPVVPVAVIGPPP AGFRDILLREGPEGFARAVRNHPGL LLMDTTFRDAHQSLLATRVRTHDLK KIAPYVAHNFSKLFSMENWGGATFD VAMRFLYECPWRRRLQELRELIPNIPF QMLLRGANAVGYTNYPDNVVFKFC EVAKENGMDVFRVFDLNYLPNML LGMEAAGSAGGVVEAASITGDVAD PSRTKYSLOYMGLAEELVRAGTHIL CIKDMAGLLKPTACTMLVSSLRDRF PDLPLHIHTHDTSGAGVAAMLACAQ AGADVVDVAADSMGMSQPSMGA LVACTRGTPLDTEVPMERVFYDSEY WEGARGLYAAFDTATMKSNGSDV YENEIPGGQYTNLHFQAHSMGLGSK FKEVKKAYVEANQMLGDLIKVTPSS KIVGDLAQFMVQNGLSRAEAEQAQAE ELSFPRSVVEFLQGYIGVPHGGFPEP FRSKVLKDLPRVEGRPGASLPLDLQ ALEKELVDRHGEEVTPEDVLSAAMY PDVFAHFKDFATFGPLDSLNTRLF LQGPKIAEEFEVELERGKTLHIKALA VSDLNRAGQRQVFFELNGQLRSILV KDTQAMKEMHFHFKALKDVKQGIG APMPGKVIDIKVVAGAKVAKGQPLC VLSAMKMETVVTSPMEGTVRKVHV TKDMTLEGDDLILEIE	None	None	None	None	None	None	None	None	

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