

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P11498	PYC_HUMAN	Homo sapiens	Pyruvate carboxylase, mitochondrial	12.757578	NaN	T1003	30397120;34105348;28510447;33214551;35083852;34725712;32119511	MLKFRTVHGGLRLLGIRRTSTAPAAS PNVRRLEYKPIKKVMVANRGEIAIRV FRACTELGIRTVAIYSEQDTGQMHR QKADEAYLIGRGLAPVQAYLHIPDIK VAKENNVDAVHPGYGFLSERADFAQ ACQDAGVRFIGPSPEVVRKMGDKVE ARAI AIAAGVPVVPGT DAPITSLHEAH EFSNTYGFPIIFKAAYGGGGRGMRV VHSYEELEENYTRAYSEALAAFNG ALFVEKFIEKPRHIEVQILGDQYGNIL HLYERDCSIQRRHQKVVEIAPAAHL DPQLRTRLTSDSVKLAKQVGYENAG TVEFLVDRHGKHYFIEVNSRLQVEH TVTEEITDVDLVHAQIHVAEGRSLPD LGLRQENIRINGCAIQCRVTTEDPAR SFQPD TGRIEVFRSGEGMGIRLDNA SAFQGAVISPHYDSL LVKVIAHGKDH PTAATKMSRALAEFRVRGVKTNIAFL QNVLNNQQFLAGTVD TQFIDENPEL FQLRPAQNRAQKLLHYLGHV MVNG PTTPIPVKASPSPTDPV VPAVPIGPPP AGFRDILLREGPEGFARAVRNHPGL LLMDTTFRDAHQSLLATRV RTHDLK KIAPYVAHNFSKLFSMENWGGATFD VAMRFLYECPWRRLQELRELIPNIPF QMLLRGANAVGYTNYPDNVVFKFC EVAKENGMDVFRVFD SLNYLPNML LGMEAAGSAGGVVEAAISYTG DVAD PSRTKYSLQYYMGLAEELVRAGTHIL CIKDMAGLLKPTACTMLVSSLRDRF PDLPLHIHTHDTSGAGVAAMLACAQ AGADVVDVAADSMG MTSQPSMGA LVACTRGTPLDTEVP MERVFDYSEY WEGARGLYAAFDC TATMKSGNSDV YENEIPGGQYTNLHFQAHSMGLGSK FKEVKKAYVEANQMLGD LKIVTPSS KIVGDLAQFMVQNGLSRAEAEAEAE ELSFPRSVEFLQGYIGVPHGGFPEP FRSKVLKDLPRVEGRPGASLPPLDLQ ALEKELVDRHGEEVTPEDVLSAAMY PDVFAHFKDFTATFGPLDSL NTRLF LQGPKIAEEFEVELER GKTLHIKALA VSDLNRAGQRQVFFELNGQLRSILV KDTQAMKEMHFHPKALKDVKGQIG APMPGKVIDIKVVAGAKVAKGQPLC VLSAMKMETVVTSPMEGT VRKVHV TKDMTLEGDDLILEIE