

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q05920	PYC_MOUSE	Mus musculus	Pyruvate carboxylase, mitochondrial	26.836479	NaN	S21;T1003	33300544;37453647;22645316	MLKFQTVRGGRLRLGVRSSSAPVA SPNVRRLLEYKPIKKVMVANRGEIAIR VFRACTELGIRTVAVYSEQDTGQMH RQKADEAYLIGRGLAPVQAYLHIPDII KVAKENGVDVHVPYGYGFLSERADFA QACQDAGVRFI GPSPEVVRKMGDKV EARAIAAAGVPVVPGTDSPISSLHEA HEFSNTFGFPPIIFKAAYGGGGRGMR VVHSYEELEENYTRAYSEALAAFNG GALFVEKFIEKPRHIEVQILGDQYGN ILHLYERDCSIQRRHQKVVEIAPATH LDPQLRSRLTSDSVKLAQVGYENA GTVEFLVDKHKHYFIEVNSRLQVE HTVTEEITDVDLVHAQIHVSEGRSLP DLGLRQENIRINGCAIQCRVTTEDPA RSFQPD TGRIE VFRSGEGMGIRLDN ASAFQGAVISPHYDSSLVKVIAHGKD HPTAATKMSRALAEFRVRGVKTNIP FLQNVLNNQQFLAGTVDTQFIDENP ELFQLRPAQNRAQKLLHYLGHVMV NGPTTPIPVNVSPSPVDPVAVPVVPIGP PPAGFRDILLREGPEGFARAVRNHQ GLLLMDTTFRDAHQSLLATRVRTHD LKKIAPYVAHNFNKLFSMENWGGA TFDVAMRFLYECPWRRQLQELRELIP NIPFQMLLRGANAVGYTNYPDNVVF KFCEVAKENGMDVFRVFDLSLNYLP NMLLGMEAAGSAGGVVEAAISYTG VADPSRTKYSLEYMGLAEELVRAG THILCIKDMAGLLKPAACTMLVSSLR DRFPDLPLHIHHTDTSAGVAAMLA CAQAGADVVDVAVDSMSGMTSQPS MGALVACTKGTPLDTEVPLERVFDY SEYWEGARGLYAAFDCTATMKSGN SDVYENEIPGGQYTNLHFQAHSMGL GSKFKEVKKAYVEANQMLGDLIKVT PSSKIVGDLAQFMVQNGLSRAEAEA QAEELSFPRSVVEFLQGYIGIPHGGF PEPFRSKVLKDLPRIEGRPGASLPPL NLKELEKDLIDRHGEEVTPEDVLSA AMYPDVFAQFKDFTATFGPLDSLNT RLFLQGPKIAEEFEVELERGKTLHIK ALAVSDLNRAGQRQVFFELNGQLRS ILVKDTQAMKEMHFHPKALKDVKG QIGAPMPGKVIDIKVAAGDKVAKGQP LCVLSAMKMETVVTSPMEGTIRKVH VTKDMTLEGDDLILEIE