	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	P52873	PYC_RAT	Rattus norvegicus	Pyruvate carboxylase, mitochondrial	26.220974	NaN	T1003	21933451	MLKFQTVRGGLRLLGVRRSSTAPVA SPNVRRLEYKPIKKVMVANRGEIAIR VFRACTELGIRTVAVYSEQDTGQMH RQKADEAYLIGRGLAPVQAYLHIPDII KVAKENGVDAVHPGYGFLSERADFA QACQDAGVRFIGPSPEVVRKMGDKV EARAIAIAAGVPVVPGTNSPINSLHE AHEFSNTYGFPIIFKAAYGGGGRGM RVVHSYEELEENYTRAYSEALAAFG NGALFVEKFIEKPRHIEVQILGDQYG NILHLYERDCSIQRRHQKVVEIAPAT HLDPQLRSRLTSDSVKLAKQVGYEN AGTVEFLVDKHGKHYFIEVNSRLQV EHTVTEEITDVDLVHAQIHVSEGRSL PDLGLRQENIRINGCAIQCRVTTEDP ARSFQPDTGRIEVFRSGEGMGIRLD NASAFQGAVISPHYDSLLVKVIAHGK DHPTAATKMSRALAEFRVRGVKTNI PFLQNVLNNQQFLAGIVDTQFIDEN PELFQLRPAQNRAQKLLHYLGHVM VNGPTTPIPVKVSPSPVDPIVPVVPIG PPPAGFRDILLREGPEGFARAVRNH QGLLLMDTTFRDAHQSLLATRVRTH DLKKIAPYVAHNFNNLFSIENWGGA TFDVAMRFLYECPWRRLQELRELIP NIPFQMLLRGANAVGYTNYPDNVVF KFCEVAKENGMDVFRIFDSLNYLPN MLLGMEAAGSAGGVVEAAISYTGDV ADPSRTKYSLEYYMGLAEELVRAGT HILCIKDMAGLLKPAACTMLVSSLR DRFPDLPLHIHTHDTSGSGVAAMLA CAQAGADVVDVAVDSMSGMTSQPS MGALVACTKGTPLDTEVPLERVFDY SEYWEGARGLYAAFDCTATMKSGN SDVYENEIPGGQYTNLHFQAHSMGL GSKFKEVKKAYVEANQMLGDLIKVT PSSKIVGDLAQFMVQNGLSRAEAEA QAEELSFPRSVVEFLQGYIGIPHGGF PEPFRSKVLKDLPRIEGRPGASLPPL NLKELEKDLIDRHGEEVTPEDVLSA AMYPDVFAQFKDFTATFGPLDSLNT RLFLQGPKIAEEFEVELERGKTLHIK ALAVSDLNRAGQRQVFFELNGQLRS ILVKDTQAMKEMHFHPKALKDVKG QIGAPMPGKVIDVKVAAGAKVVKGQ PLCVLSAMKMETVVTSPMEGTIRKV HVTKDMTLEGDDLILEIE	None	None	None	None	None	None	None	None	None