

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	extracellular region
E9QPD7	E9QPD7_MOUSE	Mus musculus	Pyruvate carboxylase	22.380232	NaN	NaN	37507081	MMLKFQTVRGGRLRLGVRSSAPV ASPNNRRLEYKPIKVMVANRGEIAI RVFRACTELGIRTVAVYSEQDTGQM HRQKADEAYLIGRGLAPVOAYLHIPD IIVAKENGVDAVHPGYGFLSERADF AQACQDAGVRFIGPSPVVRKMGDK VEARAIAAAGVPVVPGTDSPISSLHE AHEFSNTYGFPIIFKAAAYGGGGRGM RVVHSYEELEENYTRAYSEALAAF NGALFVEKFIKPRHIEVQILGDQYG NILHLYERDCSIQRRHQKVEIAPAT HLDPQLRSRLTSDSVKLAKQVGYEN AGTVEFLVDKHKHYFIEVNSRLQV EHTVTEITDVDLVHAAQIHVSEGRSL PDLGLRQENIRINGCAIQCRVTTEDP ARSFQPDGTGRIEVFRSGEGMGIRLD NASAFQGAVISPHYDSLKLVIAHGK DHPTAATKMSRALAEFRVRGKVTNI PFLQNVLNNQQFLAGTVDTFIDEN PELFQLRPAQNRAQKLLHYLGHVM VNGPTTPIPNNVSPSPVDPAPVVP GPPPAGFRDILLREGPEGFARAVRN HQGLLLMDTTFRDAHQSLLATRVRT HDLKKIAPYVAHNFNKLFMENWGW GATFDVAMRFLYECPWRRLOELREL IPNIPFQMLLRGANAVGYTNYPDNV VFKFCEVAKENGMDFRVFDSLNYL PNMLLGMEAAGSAGGVVEAISYTG DVADPSRTKYSLEYMGLAEELVRA GTHILCIKDMAGLLKPAACTMLVSSL RDRFPDLPLHIHTHDTSGAGVAAML ACAQAGADVVDVAVDSMSGMTSQP SMGALVACTKGTPLDTEVPLERVFD YSEYWEGARGLYAAFDTATMKSG NSDVYENEIPGGQYTNLHFQAHSM GLGSKFKEVKAYVEANQMLGDLIK VTPSSKIVGDLAQFMVQNGLSRAEA EAQAEELSFPRSVVEFLQGYIGIPHG GFPEPFRSKVLKDLPRIEGRPGASLP PLNLKELEKDLIDRHGEVTPEDVLS AAMYPDVFAQFKDFTATFGPLDSL TRLFLQGPKIAEEFEVELERGKTLHI KALAVSDLNRAGQRQVFFELNGQLR SILVKDTQAMKEMHFFPKALKDVK GQIGAPMPGKVIDIKVAAGDKVAKG QPLCVLSAMKMETVVTSPMEGTIRK VHVTKDMTLEGDDLILEIE	True	False	3.354	2.344	5.0	2.073	1.406	2.206	2.32