

UniprotKB ID	Entry name	organism	full name	oglcnaScore	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
G5E8R3	G5E8R3_MOUSE	Mus musculus	Pyruvate carboxylase	26.09834	NaN	NaN	36064721	MLKFQTVRGGLLRLLGVRSSAPVA SPNVRRLLEYKPIKKVMVANRGEIAIR VFRACTELGIQRTVAVYSEQDTGQMH RQKADEAYLIGRGLAPVQAYLHIPDII KVAKENGVDVHHPGYGFLSERADFA QACQDAGVRFVIGPSPEVVRKMGDKV EARAIAAAGVPVVPVGTDSPISSLHEA HEFSNTYGFPIIFKAAAYGGGRGMR VVHSYEELEENYTRAYSEALAAFGN GALFVEKFIKPRHIEVQILGDQYGN ILHLYERDCSIQRRHQKVEIAPATH LDPQLRSRLTSDSVKLAQVGYENA GTVEFLVDKHKHYFIEVNSRLQVE HTVTEEITDIDLVAQIHVSEGRSLP DLGLRQENIRINGCAIQCRVTEDPA RSFQPDTRGRIEVRFRSEGGMGIRLDN ASAFQGAVISPHYDSLIVKVIHAKGD HPTAATKMSRALAEFRVRGKTNIP FLQNVLNNQQFLAGTVDVDTQFIDENP ELFQLRPAQNRAQKLLHVLGHVMV NGPTTPIPVNVSPSPVDPVAVVPIGP PPAGFRDILLREGPEGFARAVRNHQ GLLLMDTTFRDAHQSLLATRVTRTHD LKKIAPYVAHNFNKLFSMENWGGGA TFDVAMRFLYECPWRRQLRELIP NIPFQMLLRGANAVGYTNYPDNVVF KFCEVAKENGMDVFRVFDLSNLYLP NMLLGMEAAGSAGGVVEAAISYTG VADPSRTKYSLEYMGLAEELVRAG THILCIKDMAGLLKPAACTMLVSSLR DRFPDLPLHIHTHDTSGAGVAAMLA CAQAGADVDDVAVDSMSGMTSOPS MGALVACTKGTPLDTEVPLERVFDY SEYWEGARGLYAAFDCATMKSGN SDVYENEIPGGQYTNLHFQAHSMGL GSKFKEVKAYVEANQMLGDLIKVT PSSKIVGDLAQFMVQNGLSRAEAEA QAEELSFRPSVVEFLQYIGIPHGGF PEPFRSKVLKDLPRIEGRPGASLPPL NLKELEKDLIDRHGEEVTPEDVLSA AMYPDVFAQFKDFTATFGPLDSLNT RLFLQGPKIAEEFEVELERKTLHIK ALAVSDLNRAGORQVFFELNGQLRS ILVKDTQAMKEMHHPKALKDKVKG QIGAPMPGKVIDIKVAAGDKVAKGQP LCVLSAMKMETVVTSPMEGTIRKVVH VTKDMTLEGDDLILEIE	None	None	None	None	None	None	None	None	