

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q91V92	ACLY_MOUSE	Mus musculus	ATP-citrate synthase	22.336416	NaN	T131;S263;T447;S451;S455;S459;T629;S653;T672;S829;S1090	22645316;25153642;33300544	MSAKAISEQTGKELLYKYICTTSAIQ NRFKYARVTPD TDWAHLLQDHPWL LSQSLVVKPDQLIKRRGKLGVLGVN LSLDGVKSWLKPRLGHEATV GKAKG FLKNFLIEPFVPHSQAEEFYVCIYAT REGDYVLFHHEGGVDVGDVDAKAQ KLLVGVDEKLNTE DIKRHLLVHAPE DKKEVLASFISGLFNFYEDLYFTYLEI NPLVVTKDG VYILD LAAKVDATADYI CKVKWGDIEFPPPF GREAYPEEAYIA DLDAKSGASLKL TLLNPKGRIW TMV AGGGASVVYSD TICDLGGVNELANY GEYSGAPSEQQTYDYAKTILSLMTRE KHPEGKIL IIGGSIANFTNVAATFKGI VRAIRDYQG PLKEHEVTIFVRRGGPN YQEGLRVMGEV GKT TGPIHVFGTET HMTAIVGMALGHRPIPNQPPTAAHT ANFLLNASGSTSTPAPSRTASFSESR ADEVAPAKKAKPAMPQ GK SATLFSR HTKAI VWGMQTRAVQGM LDFDYVC SRDEPSVAAMVYPFTGDHKQKFYW GHKEILIPVFKNMADAMKKHPEVDV LINFASLRSAYDSTMETMNYAQIRTI AIIAEGIPEALTRKLIK KADQKGVTIIG PATVGGIKPGCFKIGN TGGMLDNILA SKLYRPGSVAYVSRSGGMSNELNNII SRTTDGVYEGVAIGGDRYPGSTFMD HVLRYQDTPGVKMIVVLGEIGGTEEY KICRGIKEGRLTKPVVCW CIGTCATM FSSEVQFGHAGACANQAS ETAVAKN QALKEAGVFVPRSFDELGEI IQSVYE DLVAKGAI VPAQEVPPPTVPM DYSW ARELGLIRKPASFMTSICDERGQELI YAGMPITEVFKEEMGIGGVLG LLWF QRRLPKYSCQFIEMCLMVTADHGPA VSGAHNTIICARAGKDLVSSLT SGLL TIGDRFGGALDAAAKMFSKAFDSGII PMEFVNKMKKEGKLIMGIGHRVKSI NNPDMRVQILKDFVKQHFPATPLLD YALEVEKITT SKKPNLILNVDGFIGVA FVDMLRNCGSFTREEADEYVDIGAL NGIFVLGRSMGFIGHYLDQKRLKQG LYRHPWDDISYVLP EHMMSM