

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
Q99NB1	ACS2L_MOUSE	Mus musculus	Acetyl-coenzyme A synthetase 2-like, mitochondrial	24.088486	NaN	NaN	37507081;36288343	MAARSLGSGVGRLLRGLQGRSGQS GWSLSVSRSTATRLPGCVPAQAQPG SYPALSAQAQAEPAAFWGPLARDTL VWDTPYHTVWDCDFRTGKIGWFLG GQLNVSVNCLDQHVQKSPETIALIW ERDEPGTEVRITYRELLETTCLRANT LKRHGVRGRDRAIYMPVSPPLAVAA MLACARIGAIHTVVFAGFSAESLAGR INDAKCKAVITFNQGLRGGRRVELK KIVDEAVKSCPTVQHVLVAHRTDTK VPMGSLDIPLEQEMAKEAPVCTPES MSSEDMFLMLYTSGSTGTPKGLVH TQAGYLLYAAMTHKLVFDYQPGDVF GCVADIGWITGHSYVYVYGPLCNGAT TVLFESTPVYYPDAGRYWETVQRLKIN QFYGAPTAVRLLKYGDAWVKKYDR SSLRTLGSVGEPIHEAWEWLHKVV GDGRCTLVDTWWTETGGICIAPRP SEDGAEILPGMAMRPFVGVPLMD EKGNVLEGGDVSGALCISQAWPGM ARTIYGDHQRFDAYFRAYPGYYFTG DGAHRTEGGYYQITGRMDDVINISG HRLGTAEIEDAMADHPAVPETAVIG YPHDIKGEAAFAFVLDNISDENMV VNELKLSVATKIAYAVPDQILVVKR LPKTRSGKVMRRLRKIITSRGQDLG DTTTLEDPSVITEILSAFQKYEEQRAA TN	True	False	2.681	2.579	5.0	1.774	1.121	1.823	1.94