

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
Q99LB7	SARDH_MOUSE	Mus musculus	Sarcosine dehydrogenase, mitochondrial	23.627151	NaN	T778	37453647	MASLSRVLRLVAATCPRGRAAWNGLGL QPLATEARPTTEKSVPYQRTLKEEAQ GASVVPQGPSPQLPSTANVVVIGGGS LGCQTLYHLAKLGVGGAVLLERERL TSGTTWHTAGLLWQLRPSDVEVELL AHTRQVVSRLDEEETGLHTGWIQN GGLFIASNQQRRLNEYKRLMSLGKAY GIESHVLSPAETKSLYPLMNVDLTY GTLYVPQDGTMDPAGTCTTLTRAAV ARGAQVIENCAVTGIRVRTDDFGVR RVAAVETEHSIQTPCVVNCAGVWA SKVGRMAGVKVPLVAMHHAYVTE RIEIQNMPNVRDHDASVYLRLLQGD ALSVGGYEANPIFWEEVSDKFAFGL FDLDWDVFTQHIEGAINRVPVLEKT GIKSTVCGPESFTPDHKPLMGGAPE LRGFFLGCGFNSAGMMLGGGCGQE LAHWIVHGRPEKDMYSYDIRRFHHS LTDHTRWIRERSHESYAKNYSVFFP HDEPLAGRNMRRDPLHEELLGQGC VFQERQGWERPWFNPQETAQVLD YDYYGAYGNQAHKDYTYSRLLGDEY TFDFPPHHMIQKECLACRGA AAVF NMSYFGKFYLLGVDARKAADWLFSA DVNRPPGSTVYTCMLNQRGGTESDL TVSRLAPGTQASPLVPAFEGDCYLLA VGGAVAQHNWSHINTVLQDQEFRC QLMDSSEDLGMLSIQGPASRDILQD VLDADLSNEAFPSTHQLVRAAGHL VRAIRLSFVGGELGWELHVPRASCLP VYRAVMAAGARHGLV NAGYRAIDSL SIEKGYRHHHADLRPDDSPLEAGLA FTCKLKTSPFLGREALEKQRATGLR RRLICLTVEEEVPMFGLA IWRNGQ VVGHVRRADFGFTVNKTIAYGYIRDP SGGPVSLDFVKNGEYALERMGVTYA AQVHLKSPFDPDNKRKVG IY	None	None	None	None	None	None	None	None	

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