

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
Q91W43	GCSP_MOUSE	Mus musculus	Glycine dehydrogenase (decarboxylating), mitochondrial	23.627151	NaN	NaN	37453647	MQLCARAWGLRLGRGAGGGHRLAR GTGLSWAQRSDSSGGGGGGGGD RGAAGASRLLERLPRHDDFSRRHI GPGDKDRREMLQALGLASIDELIEKT VPASIRLKRPLKMEDPICENEILETL HAISKQIWRSYIGMGYYNCSVPO TILRNLENSGWVTQYTPYQPEVSQ GRLESLLNYQTMVSDITGLDMANAS LLDEATAAAEAMQLCHRRHNRKKKF FVDPRCHPQTIAVVQTRAKYRGVLV ELKLPHEMDFSGKDVCVLFQYPDT EGKVEDFTELVDRAHQTGSLTCCAT DLLALCILRPPGEFGVDIALGNSQRF GVPLGYGGPHAAFFAVKENLVRMM PGRMVGVTRDATGKEVYRLALQTRE QHRRDKATSNICTAQALLANMAAM FAIHGSGQLKHIKRVHNATLILSE GLKRAGHLQHDLFFDTLKVQCGC SVKEVLGRAAQQRQINFRLFDDGTLGI SLDETVTEKDLLLWIFGCESSAEL VAEGMGEERRGLLGSSFKRTSPFLT HQVFNSYHSETNLVRYMKLENKDI SLVHSMIPLGSCTMKLNSSELAPIT WREFANIHFPVPLDQAQGYQLFQG LEKDLCEITGYDRVSFQPNSSGAQGE YAGLATIRAYLDQKGERHRTVCLIPK SAHGTPASAHMAGMKIQPVEVDR YGNIDVAHLKAMVDQHKNLAAIMI TYPSTNGVFEENIGDVCALIHQHGG QVYLDGANMNAQVGCIRPGDFGSD VSHLNLHKTFCIPHGGGGPDMGPIG VKKHLSPFPSHPVISIKPTEGTWPV GTVSAPWGSSSILPISWAYIKMMG GKGLKEATEIAILNANYMAKRLEKH YRVLFRGARGYVAHEFILDTRPFKKS ANVEAVDVAKRLQDYGFHAPTMSW PVAGTLMIEPTESEDKAELDRFCDA MISIRQEIADIEEGRIDPRVNPLKMS PHSLTCVTSSCWDPRPYSREVAAPPL PFVKPENKFWPTIARIDDIYGDQHLV CTCPPMEVYESPFSEQKRASS	True	False	3.262	3.102	4.718	0.923	0.528	2.755	1.466

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