

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
0 Q9HCL2	GPAT1_HUMAN	Homo sapiens	Glycerol-3-phosphate acyltransferase 1, mitochondrial	18.379595	NaN	S380;S688;S695	30379171	MDESALTLGTTIDVSYLPHSSEYSVGR CKHTSEEWGECGFRPTIFRSATLKW KESLMSRKRPFVGRCCYSCTPQSWD KFFNPSIPSLGLRNVIYINETHTRHR GWLARRLSYVLFIQERDVHKGMFAT NVTENVLNSRVQEAIAEVAELNP DGSAQQQSKAVNKVKKAKRILQE MVATVSPAMIRLTGWVLLKLFNSFF WNIQIHKGQLEMVKAATEINLPLLF LPVHRSHIDYLLLTFFILFCHNIKAPYI ASGNNLNIPFSTLIHKLGGFFIRRRLL DETPDGRKDVLYRALLHGHIPELLR QQQFLEIFLEGTRSRSGKTSARAGL LSVVVDTLSTNVIPDILIPVGISYDRII EGHYNGEQLGKPKKNESLWSVARG VIRMLRKNYGCVRVDFAPQFSLKEY LESQSQKPVSAALLSLEQALLPAILPS RPSDAADEGRDTSINESRNATDESLL RRRLIANLAEHILFTASKSCAIMSTHI VACLLYRHRQGDIDLSTLVEDFFVM KEEVLARDFDLGFSGNSEDVVMHAI QLLGNCVTIHTSRNDEFFITPSTTV PSVFELNFYSNGVLHVFMIAIACS LYAVLNKRGLGGPTSTPPNLISQEQL VRKAASLCYLLSNEGTSILPCQTFYQ VCHETVKGFIQYGILTVAEHDDQEDI SPSLAEQQWDKKLPEPLSWRSDEED EDSDFGEEQRDCYLKVSQSKEHQQ FITFLQRLGPLEAYSSAIFVHNFS GPVPEPEYLQKLHKYLITRTERNVAV YAESATYCLVKNVAVKMFKDIGVFKE TKQKRVSVLELSSTFLPQCNRQKLL EYILSFVVL	True	False	2.675	2.534	4.559	2.481	1.151	3.055	2.096