

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
E9PUQ8	DGKD_MOUSE	Mus musculus	Diacylglycerol kinase delta	29.185005	NaN	NaN	22645316	MAAAAGAPPPGPPQPPPPPEESS DSEPEAEPGSPQKLRKVSTSGQIRQ KTLKKEGMLTKQNNSFQRSKRRYFK LRGRTLYYAKTAKSIIFDEVLDTASV AESSTKNVNNSFVTITPCRKLILCAD NRKEMEDWIAALKTVQNEHFPT QYSMDHFGSMHNWYACSHARPTY CNVCREVLGSGVTSGLSCEVCKFKA HKRCAVRATSNCKWTTLASIGKDIIE DEDGIAMPHQWLEGNLPVSAKIVC DKTCGSVLRQDWRCLWCKAMVHT SCKESLVMKCPGLGLCKVSVIPTALN SIDSDFGWKATCPPSCTSPLLVFNVS KSGDNQGVKFLRRFKQLLPAQVF DLMNGGPHLGLRFLQKFDTRILVC GGDGSVGVVLEIDSLNLHKQCQL GVLPLGTGNDLARVLGWGSACDDD TQLPQILAKLERASTKMLDRWSVMA YETKLPQASSTVTEDFSESEVQ QILFYEDSVAHLSKILTSQHSVVIS SAKVLCETVKDFVARVKGAYEKTTE SQESEVMAKKCSVLKEKLDLLKTL DDESQASSLSNPPPTIAEEAEDGDG SGNICSTGDHLVGSACPSRQIFRP REQMLRANSLKKAIRQIIEHTEKAV DEQNAQTQEQGFVGLSESEKKDL KTDNRVCTSSVHSESCVIAKGRSQR KASRAPCEKLVSKGLSLGSSASLPPG TGSRDSPALNTKILYPSVRAGMSG LPGGSVISRLINADPFNAEPENLEY YTEKVMNMYFGIGLDAKISLDFNN KRDEHPEKCRSRTKNMMWYGVLT KELLHRTYRNLEQKVLECDGRPIPL PSLQGIAVLNIPSYAGGTNFWGGTK EDDTFAAPSFDDKILEVVAVFGSMQ MAVSRVIKQHHRIAQCRTVKISILG DEGVPVQVDGEAWIQPPGYIRIVHK NRAQTLTRDRAFENTLKSWEKQK CELSRPPSFLHPEILSEEEATQMDQ FGQAAGGLIHSIREIAQSHRAMEQE LAHAVNASSKAMERVYKPRTAEGL NCSFVLEMVNNIRALRSETELLG KMALQDPPQKERLGAALIMDQQL RKLTDTPWLCQPLEPGEESLQQNV MLDLTKRSRSGKFRLVTKFKKEKNN KNKEVHNSLGGPVHLWGTEEVAAW LEHLSLCEYKDFTRHDIRGSELLHL ERRDLKDLGVTKVGHMKRILCGIKE LSRSSPAAEA	True	False	4.672	2.376	1.114	1.256	1.341	4.288	0.796