

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
D3YXJ0	DGKH_MOUSE	Mus musculus	Diacylglycerol kinase eta	29.185005	NaN	NaN	22645316	MAGAGSQHHPQGVAGGAVAGASAV SPTAAGPGEDSSDSEAEQEGPQLIR KVSTSGQMRTKTSIKEGQLLKQTSS FQRWKKRYFKLRGRTLYYAKDSKSLI FDEVLDSDASVAEASTKNANNSFTII TPFRRLMLCAENRKEMEVWISSLKS VQSREPYEVAQFNVEHFSGMHNWY ACSHARPTFCNVCRESLGSVTSHGL SCEVCKFKAHKRCAVRATNNCKWT TLASIGKDIIIEDEGVMHPHQWLEG NLPVSAKCAVCDKTCGSLRLQDVK CLWCKTMVHTACKDVYHPVCLGQ CKVSIIPPIALNSTSDGFCRATFSFC VSPLLVFVNSKSGDNQGVKFLRRFK QLLNPAQVFDLMNGGPYLGLRFLQK FDNFRILVCGGDGSGVWVLEIDKL NLHKQCQLGVLPLGTGNDLARVLG WGGSYDDDTQLPQILEKLERASTKM LDRWSIMTYELKLPKSSLLPEPVAA TEEFYMTYEDSVANHLTKIVNSDE HAVVSSAKILCETVKDFVAKVEKAQ DRTLENTVVAEAVASKCSVLNEKLE QLLQALHADSQASRVPPGIGPAIPEE DTVESASDES LGESKDQLVNDIGKP SSQKAVKPREIMLRANSLKKAQRQVI EEAEKVMDEPAVQPEPVPSPCDYD TETDEAKEDDAKESLSAKTTSQSPD AQASCGRPOTDSVAGPAMATTKEN LPVLNTRIIICPLRAGLAASIAGSSIIN KMLLANIDPFGATPFIDPDLDSLQGY SEKCVMMNYFGIGLDAKISLEFNK REEHPEKCRSRTKNLMWYGVLGTR ELLQRSYKNLEQRVQLECDGQYIPLP SLOGIAVLNIPSYAGGTNFVWGCTKE DDIFAAPSFDDKILEVVAVFDSVQMA VSRVIKLQHHRIAQCRTVKITIFGDE GVPVQVDGEAWVQPPGIKIVHKNR AQMLTRDRAFESTLKSWEKQKCD SGKPVLRNLYIHPAPDLATEEVSQ MRLCSQAAEELITRICDAATHCCLLE QELAHAVNACSHALNKANPRFPESL TRDTATEIAINVKALYNTEALLVGR VPLHLESPHEERVSSALHSVEMELQ KLT EIPWLYILRPSEDEEPLDCTK RNNKSTVFRIVPKFKKEKAQKQKTS SQPVQNWGT EEEVAAWLDDLNLGEY KEIFIRHDVVRGAE LLHLERRDLKDLG IPKVGHMKRILQGIKELERNPPNLV	False	True	3.432	2.497	1.31	1.415	1.104	5.0	1.701