

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 Q6NS52	DGKB_MOUSE	Mus musculus	Diacylglycerol kinase beta	24.474581	NaN	T117;S418;S791	33300544	MTNQEKWAHLSPSEFSQLQKYAEY STKLLKDVLEEFHGNVGLAKYNPEG KQDILNQITDFEGFKLFMKTFLAEEL PDDFTAHLFMSFSNKFPHSSPNVKS KPALLSGGLRMNKGAITPPRSSPAN TCSPEVIHLKDIVCYLSLLERGRPED KLEFMFRLYDTDGNGLDSSSELENII GQMMHVAEYLEWDVTELNPIIHEM MEEIDYDRDGTVSLEEWIQQGMMTI PLLVLLGLENNVKDDGQHVWRLLKH FNKPAYCNLCLNMLIGVKGQLCCS FCKYTVHERCVARAPPSCIKTYVKSK KNTDVMHHYVWEGNCPKCDKCH KTVKCYQGLTGLHCVWCQTTLHNK CASHLKPECDGPKLDHILPPTTICP VVLTMPSAGASVPEERQSTAKKEKS SSQPNKATDRNKMQRANSVTMD GGGLQITPVPGTHPLLVFVNPKSGG KQGERIYRKFQYLLNPRQVYSLSGN GPMPGLHFFRDVPDFRVLACGGDG TVGWILDICIEKANVVKHPPVAILPLG TGNDLARCLRWGGGYEGENLMKIL KDIESSTEIMLDRWKFEVTPNDKDE KGDVPVYSIINNYFSIGVDASIAHRF HIMREKHPEKFNSRMKNKFWYFEF GTSETFSATCKKLHESVEIEDGVQI DLINISLEGIAILNIPSMHGGSNLWG ESKKRSHRRIEKKGSDKRPTLDA KELKFASQDLSDQLLEVVGLEGAME MGQIYTGKLSAGRRLAQCSSVIRTS KSLPMQIDGEPWMQTPCTIKITHKN QAPMLMGPPPKTGLFCSLIKTRRNR SKE	False	True	3.372	2.465	1.98	1.247	0.941	5.0	1.21