	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q13574	DGKZ_HUMAN	Homo sapiens	Diacylglycerol kinase zeta	19.994096	NaN	S705;S781	37217939	MEPRDGSPEARSSDSESASASSSGS ERDAGPEPDKAPRRLNKRRFPGLRL FGHRKAITKSGLQHLAPPPPTPGAPC SESERQIRSTVDWSESATYGEHIWF ETNVSGDFCYVGEQYCVARMLKSVS RRKCAACKIVVHTPCIEQLEKINFRC KPSFRESGSRNVREPTFVRHHWVH RRRQDGKCRHCGKGFQQKFTFHSK EIVAISCSWCKQAYHSKVSCFMLQQI EEPCSLGVHAAVVIPPTWILRARRPQ NTLKASKKKKRASFKRKSSKKGPEE GRWRPFIIRPTPSPLMKPLLVFVNPK SGGNQGAKIIQSFLWYLNPRQVFDL SQGGPKEALEMYRKVHNLRILACGG DGTVGWILSTLDQLRLKPPPPVAILP LGTGNDLARTLNWGGGYTDEPVSKI LSHVEEGNVVQLDRWDLHAEPNPE AGPEDRDEGATDRLPLDVFNNYFSL GFDAHVTLEFHESREANPEKFNSRF RNKMFYAGTAFSDFLMGSSKDLAK HIRVVCDGMDLTPKIQDLKPQCVVF LNIPRYCAGTMPWGHPGEHHDFEP QRHDDGYLEVIGFTMTSLAALQVGG HGERLTQCREVVLTTSKAIPVQVDG EPCKLAASRIRIALRNQATMVQKAK RRSAAPLHSDQQPVPEQLRIQVSRV SMHDYEALHYDKEQLKEASVPLGTV VVPGDSDLELCRAHIERLQQEPDGA GAKSPTCQKLSPKWCFLDATTASRF YRIDRAQEHLNYVTEIAQDEIYILDPE LLGASARPDLPTPTSPLPTSPCSPTP RSLQGDAAPPQGEELIEAAKRNDFC KLQELHRAGGDLMHRDEQSRTLLH HAVSTGSKDVVRYLLDHAPPEILDAV EENGETCLHQAAALGQRTICHYIVE AGASLMKTDQQGDTPRQRAEKAQD TELAAYLENRQHYQMIQREDQETAV	None	None	None	None	None	None	None	None	None