

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
P61460	DEPD5_MOUSE	Mus musculus	GATOR complex protein DEPDC5	25.019509	T1000;S1032	S505	34418053	MRTTKVYKLVIIHKKGFGGSDDELVV NPKVFPHKLGDIIVEIAHPNDEYSPL LLQVKSLKEDLQKETISVDQTVTQVF RLRPYQDVYVNVVDPKDVTLDLVEL TFKDQYIGRGMWRLKKSIVSTCAY ITQKVEFAGIRAQAGELWVKNEKVM CGYISEETRVVFRSTSAMVYIFIQMS CEMWDFDIYGDLYFEKAVNGFLADL FTKWKEKNCSHEVTVVLFSRTFYDA KSIDFPEINRASIQEDHKGRFYEDF YKVVVQNERREEWTSLLVTIKKLFIQ YPVLVRLAQAGGFPGDNSTSAQGN YLEAINLSFNVFDKHYINRNFRDTG QMSVVITPGVGVFEVDRLMLTKQ RMIDNGIGVDLVCMEQPLHAVPLF KLHNRVPRDSRLGDDYNIPHWINH SFYTSKSQLFCNSFTPRIKLAGKKA SEKTKNGRDTSLGTPKESENTLPQV DYDAYDAQVFRPLPGPSRAQRLATCR SVREQENHSRKSASSCDVSSPSLP SRALPTEEVRSQASDDSSLGKSTNII MIPNPHLHQYEVSSSLGYTSTRDVL ENMIEPPORDSSAPGRFHVGSAESM LHVRPGGYTPQRALINPFAPSRMPM KLTSNRRRWMHTFPVGPGEAIQIH HQTRQNMAELQGSQRDPHSSAE LLELAYHEAAGRHSRQPGDSMSL NFSGTEELSVLLSNSSTGVNPRTO NKDSLEDSVSTSPDPMGFCCTVGV DWKSLTTPACLPLTDDYFPDRQGLQ NDYTEGCYDLLPEADMRRDEEGV QMTAQQVFEFICQRLMQGYQIIVQ PKTQKPNITVPPPLSSSPLYSRGLVS RNRPEEEGQYWL SMGRTFHKVTLK DKMITVTRYLPKYPYESAQIHITYSL CPSHSDSEFVSCWVDFCHERLEEYK WNYLDQYICSAAGSEDFSLIESLKF RTRFLLPACVTATKRITEGEVHCDI YGDKPRADEDEWQLLDGFIRFVEGL NRIRRRHRSRDMIRKGTAMKGLQM TGPISAHSLAAGPPVGGKGTSA LLEMEASQKSLGEQTTVHGKSS PAENSSVAMTPTYVDSPRKDGAF EFVRSPTASSAFYQASVDQTAPLV LDSTSLGVSTGQPMDRGNNQTFGN SQNIEQAFPSANSQYSSQHVASS LTSSTLVEILEAMKHPSTGVQLLSE QKGLSPCCFISAEVVHWMNNVEG VQTQAMGIDIMQKMLEEQLITHASG EAWRTFIYGFYFYKIVMDKEPERVA MQQPSAPWYTAGADDFASFQRKWF EVAFVAEELVHSEIPAFLLPWLPSRP ASYASRHSSFSRSGGRSQAALLA ATVPEQRTVTLDVDVNNRDRLEW CSCYYHGNFSLNAAFEIKLHWMMAVT ATVLFEMVQGWHRKATSCGFLLPV LEGPFALPSYLYGDPLRAQLFIPLNL SCLLKEGSEHLFDSFEPETYWDRM HLFQEAIAHRFGFVQDKYSVSAFNF PAENKPYIHTGTVFLQLPYSKRKF SGQQRNRNSTSSTNQNMFCCEERV GYNWAYNTMLTKTWRSSATGDEKF ADRLKDFTFDCINRDNRLVTFWTN CLEKMHASAP	True	False	5.0	2.84	1.497	1.079	0.875	1.306	1.167