	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	O75140	DEPD5_HUMAN	Homo sapiens	GATOR complex protein DEPDC5	27.086674	T513	S505	28657654	MRTTKVYKLVIHKKGFGGSDDELVV NPKVFPHIKLGDIVEIAHPNDEYSPL LLQVKSLKEDLQKETISVDQTVTQVF RLRPYQDVYVNVVDPKDVTLDLVEL TFKDQYIGRGDMWRLKKSLVSTCAY ITQKVEFAGIRAQAGELWVKNEKVM CGYISEDTRVVFRSTSAMVYIFIQMS CEMWDFDIYGDLYFEKAVNGFLADL FTKWKEKNCSHEVTVVLFSRTFYDA KSVDEFPEINRASIRQDHKGRFYEDF YKVVVQNERREEWTSLLVTIKKLFIQ YPVLVRLEQAEGFPQGDNSTSAQGN YLEAINLSFNVFDKHYINRNFDRTG QMSVVITPGVGVFEVDRLLMILTKQ RMIDNGIGVDLVCMGEQPLHAVPLF KLHNRSAPRDSRLGDDYNIPHWINH SFYTSKSQLFCNSFTPRIKLAGKKPA SEKAKNGRDTSLGSPKESENALPIQ VDYDAYDAQVFRLPGPSRAQCLTTC RSVRERESHSRKSASSCDVSSSPSLP SRTLPTEEVRSQASDDSSLGKSANIL MIPHPHLHQYEVSSSLGYTSTRDVL ENMMEPPQRDSSAPGRFHVGSAES MLHVRPGGYTPQRALINPFAPSRMP MKLTSNRRRWMHTFPVGPSGEAIQI HHQTRQNMAELQGSGQRDPTHSSA ELLELAYHEAAGRHSNSRQPGDGM SFLNFSGTEELSVGLLSNSGAGMNP RTQNKDSLEDSVSTSPDPILTLSAPP VVPGFCCTVGVDWKSLTTPACLPLT TDYFPDRQGLQNDYTEGCYDLLPEA DIDRRDEDGVQMTAQQVFEEFICQR LMQGYQIIVQPKTQKPNPAVPPPLSS SPLYSRGLVSRNRPEEEDQYWLSMG RTFHKVTLKDKMITVTRYLPKYPYES AQIHYTYSLCPSHSDSEFVSCWVEF SHERLEEYKWNYLDQYICSAGSEDF SLIESLKFWRTRFLLLPACVTATKRIT EGEAHCDIYGDRPRADEDEWQLLD GFVRFVEGLNRIRRRHRSDRMMRK GTAMKGLQMTGPISTHSLESTAPPV GKKGTSALSALLEMEASQKCLGEQQ AAVHGGKSSAQSAESSSVAMTPTYM DSPRKDGAFFMEFVRSPRTASSAFY PQVSVDQTATPMLDGTSLGICTGQS MDRGNSQTFGNSQNIGEQGYSSTN SSDSSSQQLVASSLTSSSTLTEILEA MKHPSTGVQLLSEQKGLSPYCFISA EVVHWLVNHVEGIQTQAMAIDIMQ KMLEEQLITHASGEAWRTFIYGFYFY KIVTDKEPDRVAMQQPATTWHTAGV DDFASFQRKWFEVAFVAEELVHSEI PAFLLPWLPSRPASYASRHSSFSRSF GGRSQAAALLAATVPEQRTVTLDVD VNNRTDRLEWCSCYYHGNFSLNAA FEIKLHWMAVTAAVLFEMVQGWHR KATSCGFLLVPVLEGPFALPSYLYGD PLRAQLFIPLNISCLLKEGSEHLFDS FEPETYWDRMHLFQEAIAHRFGFVQ DKYSASAFNFPAENKPQYIHVTGTVF LQLPYSKRKFSGQQRRRRNSTSSTN QNMFCEERVGYNWAYNTMLTKTW RSSATGDEKFADRLLKDFTDFCINR DNRLVTFWTSCLEKMHASAP	None	None	None	None	None	None	None	None	None