	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma memb
0	O75153	CLU_HUMAN	Homo sapiens	Clustered mitochondria protein homolog	20.754878	T1282	S279;S281;S654;S664;S723	30379171;35138101;23301498;41350704;28510447;33214551;34019948	MLLNGDCPESLKKEAAAAEPPRENG LDEAGPGDETTGQEVIVIQDTGFSVK ILAPGIEPFSLQVSPQEMVQEIHQVL MDREDTCHRTCFSLHLDGNVLDHF SELRSVEGLQEGSVLRVVEEPYTVRE ARIHVRHVRDLLKSLDPSDAFNGVD CNSLSFLSVFTDGDLGDSGKRKKGL EMDPIDCTPPEYILPGSRERPLCPLQ PQNRDWKPLQCLKVLTMSGWNPPP GNRKMHGDLMYLFVITAEDRQVSIT ASTRGFYLNQSTAYHFNPKPASPRF LSHSLVELLNQISPTFKKNFAVLQKK RVQRHPFERIATPFQVYSWTAPQAE HAMDCVRAEDAYTSRLGYEEHIPGQ TRDWNEELQTTRELPRKNLPERLLR ERAIFKVHSDFTAAATRGAMAVIDG NVMAINPSEETKMQMFIWNNIFFSL GFDVRDHYKDFGGDVAAYVAPTNDL NGVRTYNAVDVEGLYTLGTVVVDYR GYRVTAQSIIPGILERDQEQSVIYGSI DFGKTVVSHPRYLELLERTSRPLKIL RHQVLNDRDEEVELCSSVECKGIIG NDGRHYILDLLRTFPPDLNFLPVPGE ELPEECARAGFPRAHRHKLCCLRQE LVDAFVEHRYLLFMKLAALQLMQQ NASQLETPSSLENGGPSSLESKSED PPGQEAGSEEEGSSASGLAKVKELA ETIAADDGTDPRSREVIRNACKAVGS ISSTAFDIRFNPDIFSPGVRFPESCQD EVRDQKQLLKDAAAFLLSCQIPGLVK DCMEHAVLPVDGATLAEVMRQRGI NMRYLGKVLELVLRSPARHQLDHVF KIGIGELITRSAKHIFKTYLQGVELSG LSAAISHFLNCFLSSYPNPVAHLPAD ELVSKKRNKRRKNRPPGAADNTAW AVMTPQELWKNICQEAKNYFDFDL ECETVDQAVETYGLQKITLLREISLK TGIQVLLKEYSFDSRHKPAFTEEDVL NIFPVVKHVNPKASDAFHFFQSGQA KVQQGFLKEGCELINEALNLFNNVY GAMHVETCACLRLLARLHYIMGDYA EALSNQQKAVLMSERVMGTEHPNT IQEYMHLALYCFASSQLSTALSLLYR ARYLMLLVFGEDHPEMALLDNNIGL VLHGVMEYDLSLRFLENALAVSTKY HGPKALKVALSHHLVARVYESKAEF RSALQHEKEGYTIYKTQLGEDHEKT KESSEYLKCLTQQAVALQRTMNEIY RNGSSANIPPLKFTAPSMASVLEQL NVINGILFIPLSQKDLENLKAEVARR HQLQEASRNRDRAEEPMATEPAPA GAPGDLGSQPPAAKDPSPSVQG	False	False	2.35	2.797	3.841	1.124	0.188	0.851