	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	extracellul region
0	P58854	GCP3_MOUSE	Mus musculus	Gamma-tubulin complex component 3	30.023367	NaN	S113	34678516;35822049;28528544	MATPDQKSPNVLLQNLCCRILGRSE ADVAQQFQFAVRVIGSNFAPTVERD EFLVAEKIKKELIRQRREADAALFSE LHRKLHSQGVLKNKWSILYLLLNLS EDPRKQASKVTSYASLFAQALPRDA HSTPYYYARPQTLPLNYQDRSTQAQ SSGSLGSSGISSIGMCGLSGPTPVQP FLPGQSHQAPGVGDGLRQQLGPRLA WTLTGNQPSSQTPTSKGFPNALSRN LTRSRREGDPGGTLEVTEAALVRDIL YVFQGIDGKNIKMSSTENCYKVEAK ANLNKSLRDTAVRLAELGWLHNKIR KYADQRSLDRSFGLVGQSFCAALHQ ELKEYYRLLSVLHSQLQLEDDQGVN LGLESSLTLRRLLVWTYDPKIRLKTL AALVDHCQGRKGGELASAVHAYTKT GDPYMKSLVQHILSLVSHPVLSFLYR WIYDGELEDTYHEFFVASDPTVKTD RLWHDKYTLRKSMIPSFITMDQSRK VLLIGKSINFLHQVCHDQTPTTKMIA VTKSAESPRDAADLFTDLENAFQGKI DAAYFETSKYLLDVLNKKYSLLEHM QAMRRYLLLGQGDFIRHLMDLLKPE LVRPATTLYQHNLTGILETAVRATNA QFDSPEILKRLDVRLLEVSPGDTGW DVFSLDYHVDGPIATVFTRECMSHY LRVFNFLWRAKRMEYILTDIRKGHM CNAKLLRNMPEFSGVLHQCHILASE MVHFIHQMQYYITFEVLECSWDEL WNRVQQAQDLDHIIAAHEAFLNTIT SRCLLDSNSRVLLNQLRAVFDQIIEL QNAQDVMYRAALEELQRRLQFEEK KKQREIEGQWGVTAAEEEEENKRIR EFQDSIPKMCSQLRILTHFYQGVVQ QFLVLLTTSSDESLQFLSFRLDFNE HYKAREPRLRVSLGSRGRRSSHT	None	None	None	None	None	None	None	None	None