	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	Q62638	GSLG1_RAT	Rattus norvegicus	Golgi apparatus protein 1	23.439218	NaN	S953	38843836	MAVCGRVRRMFRLSAALQLLLLVAA GVQNSHGQGQGLGVNFGPFAGQAG GGNPVGQQPPQLPQLSQQQQQQP PPQQQQPFPAGGLPARRGGAGPGGT GGGWKLAEEESCREDVTRVCPKHT WSNNLAVLECLQDVREPENEISSDC NHLLWNYKLNLTTDPKFESVAREVC KSTISEIKECAEEPVGKGYMVSCLVD HRGNITEYQCHQYITKMTAIIFSDYR LICGFMDDCKNDINLLKCGSIRLGE KDAHSQGEVVSCLEKGLVKEAEEKE PKIQVSELCKKAILRVAELSSDDFHL DRHLYFACRDDRERFCENTQAGEG RVYKCLFNHKFEESMSEKCREALTT RQKLIAQDYKVSYSLAKSCKSDLKKY RCNVENLPRSREARLSYLLMCLESA VHRGRQVSSECQGEMLDYRRMLME DFSLSPEIILSCRGEIEHHCSGLHRK GRTLHCLMKVVRGEKGSLGMNCQQ ALQTLIQETDPGADYRIDRALNEACE SVIQTACKHIRSGDPMILSCLMEHLY TEKMVEDCEHRLLELQYFISRDWKL DPVLYRKCQGDASRLCHTHGWNET SELMPPGAVFSCLYRHAYRTEEQGR RLSRECRAEVQRILHQRAMDVKLDP ALQDKCLIDLGKWCSEKTETGQELE CLQDHLDDLAVECRDIVGNLTELES EDIQIEALLMRACEPIIHNFCHDVAD NQIDSGDLMECLIQNKHQKDMNEK CAIGVTHFQLVQMKDFRFSYKFKMA CKEDVLKLCPNIKKKVDVVICLSTTV RNDTLQEAKEHRVSLKCRKQLRVEE LEMTEDIRLEPDLYEACKSDIKNYCS TVQYGNAQIIECLKENKKQLSTRCH QRVFKLQETEMMDPELDYTLMRVC KQMIKRFCPEADSKTMLQCLKQNK NSELMDPKCKQMITKRQITQNTDYR LNPVLRKACKADIPKFCHGILTKAKD DSELEGQVISCLKLRYADQRLSSDCE DQIRIITQESALDYRLDPQLQLHCSD EIANLCAEEAAAQEQTGQVEECLKV NLLKIRTELCKKEVLNMLKESKADIF VDPVLHTACALDIKHHCAAITPGRGR QMSCLMEALEDKRVRLQPECKKRL NDRIEMWSYAAKVAPADGFSDLAM QVMTSPSKNYILSVISGSICILFLIGL MCGRITKRVTRELKDR	False	True	1.091	1.381	1.608	1.606	5.0	2.523	1.633