	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	extra regio
0	Q711U4	Q711U4_DROME	Drosophila melanogaster	Alpha-1,6-mannosyl-glycoprotein 2-beta-N-acetylglucosaminyltransferase	28.06426	S388;S392	NaN	40245499	MGRKRNNFYMRSLFLLALGIFGLLQ YNNFNYLDSRDNVLGDAVTNDSDD AILAMVPATLHKYLTPHSRNHSASG AGVLNGAALLLNASSPGAATASTISF DVYHPPNITEIKRQIVRYNDMQMVL NEDVFGPLQNDSVIIVVQVHTRITYL RHLIVSLAQARDISKVLLVFSHDYYD DDINDLVQQIDFCKVMQIFYPYSIQT HPNEYPGVDPNDCPRNIKKEQALIT NCNNAMYPDLYGHYREAKFTQTKH HWIWKANRVFNELEVTRYHTGLVLF LEEDHYVAEDFLYLLAMMQQRTKD LCPQCNVLSLGTYLKTFNYYTYHSKT NKKSYASSLISTNSLLGYNRNNNRN SVQLAVSSSYSSSSSAQSPPPSKVYV GDTSDNQRASDSSSRRGNLINTVTA TATATAATSDTSNNEQSSHSSSKNG AQTWNYHVLPSLYSVYQKVEVMPW VSSKHNMGFAFNRTTWSNIRKCAR HFCTYDDYNWDWSLQHVSQQCLRR KLHAMIVKGPRVFHIGECGVHHKNK NCESNQVISKVQHVLRIARNSHQLF PRSLTLTVPSLMKKSKLRKGNGGWG DMRDHELCLNMTLATR	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None