	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	O09159	MA2B1_MOUSE	Mus musculus	Lysosomal alpha-mannosidase	25.810548	NaN	NaN	29187734	MGTGPLTSGVRAGGGNTGWLWMS SCNLGSPVLPISFLFWLLLAAPGARA AGYKTCPPTKPGMLNVHLLPHTHD DVGWLKTVDQYYYGILSDVQHASVQ YILDSVVSSLLEKPTRRFIYVEMAFFS RWWKQQTSATQDAVRNLVRQGRLE FVNGGWVMNDEAATHYGAIVDQMT LGLRFLQDTFGSDGLPRVAWHIDPF GHSREQASLFAQMGFDGFFLGRIDY QDKLNRKKKLRMEELWRASDSLEP PAADLFTGVLPNNYNPPKYLCWDVL CTDPPVVDNPRSPEFNAKTLVNYFL KLASSQKGFYRTNHTVMTMGSDFH YENANMWFKNMDKLIRLVNAQQV NGSLVHVLYSTPTCYLWELNKANLT WTVKEDDFFPYADGPHMFWTGYFS SRPALKRYERLSYNFLQVCNQLEAL VGPEANVGPYGSGDSAPLQEAMAVL QHHDAVSGTARQNVVNDYARQLAA GWGPCEVLVSNALARLSHYKQNFSF CRELNISICPVSQTSERFQVTLYNPL GRKVDQMVRLPVYEGNFIVKDPHDK NISSNVVMVPSYYSETYQWELLFPA SVPALGFSTYSVAKMSDLNHQAHNL LSRPRKHKSHHVLVIENKYMRATFD SGTGLLMKIENLEQNLSLPVSQGFF WYNASVGDEESSQASGAYIFRPNVG KPIPVSRWAQISLVKTALVQEVHQNF SAWCSQVIRLYKGQRHLELEWTVGP IPVRDDWGKEVISRFDTPMKTKGQF FTDSNGREILKRRDDYRPTWTLNQT EPVAGNYYPVNTRIYITDGQMQLTVL TDRSQGGSSLQDGSLELMVHRRLLV DDDRGVSEPLLETDTGDKVRGRHLV LLSSVSDAAARHRLLAEQEVLAPQV VLSLGGSSPYHSRATPKTQFSGLRQ ELPPQVHLLTLARWGPKMLLLRLEH QFALKEDSDRNLSSPVTLNVQNLFQ TFTINYLQETTLAANQPLSRASRLKW MTNTGPTSYPEPSKLDPTSVTLKPM EIRTFLASVQWQEHRPA	False	False	1.506	3.611	1.336	1.198	2.755	1.243	2.833