

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q80UM7	MOGS_MOUSE	Mus musculus	Mannosyl-oligosaccharide glucosidase	28.30369	NaN	NaN	29187734;37453647	MARGERRRRAAAEGARPLERARAA GRRDGRAGGARGSASGAALAVVLA LAFGLSGRWVLAWLVRRAATLHPA PSALPPDSSSPAVAPELFWGTYRPHV YFGMKTRSPKPLLTGLMWAQQGAT PGTTPKLRHTCEQGDGVGYPGWEF HDGRTFGRQHIHDGALRLITTEFVKR PGGQHGDDWSWRVTVEPQASGTPS FPLVSLFFYVVDGQEVLLPEIGAKG QLKSISGHTSELGDFRLTLLPPTSPG DTVPKHGSYNVFWSSNPGLPQLTD MVKSRLLNSWFQHRPPGASPDRLG LPGSLKWEERGPGSGQGQFLIQVTL KAPFSVEFVFESGSAATGGNQASGR LVGSOLTQALESHAAAFKERFEKTF QLKEKGLSPEEQALGQVALSGLLGGI GYFYQGGLVLPDTSMEGSEQKMDP ALFPPVPLFSGVPSRSFFPRGFLWDE GFHQLVVQRWDPHLTREALGHWLG LLNADGWIGREQILGDEARARVPPE FLVQRAAHANPPTLLPVVHMLEGH DPDDLAFLRKAFPRLHAWFSLHQ SQAGPVPLSYRWRGRDLALPTLLNP KTLPSGLDDYPRASHPSTAERHLDL RCWVALGARVLSQLAEQLGETEAAA ELGPLAASLEEPGSLDELHWAPELG VFADFGNHTKAVQLKSRPPQGLVRV VGRPPPRLQYVDALGYVSLFLLLQL LDPSSPRLGPLLDVLAISRHLWSPF GLRSLSASSLFYKQRNTEHDPYWR GAVWLNINYLALGALHHYGHVEGP HKVQAAKLYHELRLANVVRNVQGY QATGFLWEQYSDQDGRMGCRPFQ GWTSVLVLLIMAEY	False	False	1.869	1.786	1.66	3.83	1.654	2.005	1.333