	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	Q8TET4	GANC_HUMAN	Homo sapiens	Neutral alpha-glucosidase C	25.829053	NaN	NaN	34105348	MEAAVKEEISLEDEAVDKNIFRDCN KIAFYRRQKQWLSKKSTYQALLDSVT TDEDSTRFQIINEASKVPLLAEIYGIE GNIFRLKINEETPLKPRFEVPDVLTS KPSTVRLISCSGDTGSLILADGKGDL KCHITANPFKVDLVSEEEVVISINSL GQLYFEHLQILHKQRAAKENEEETS VDTSQENQEDLGLWEEKFGKFVDIK ANGPSSIGLDFSLHGFEHLYGIPQHA ESHQLKNTGDGDAYRLYNLDVYGYQ IYDKMGIYGSVPYLLAHKLGRTIGIF WLNASETLVEINTEPAVEYTLTQMG PVAAKQKVRSRTHVHWMSESGIIDV FLLTGPTPSDVFKQYSHLTGTQAMP PLFSLGYHQCRWNYEDEQDVKAVD AGFDEHDIPYDAMWLDIEHTEGKRY FTWDKNRFPNPKRMQELLRSKKRK LVVISDPHIKIDPDYSVYVKAKDQGF FVKNQEGEDFEGVCWPGLSSYLDFT NPKVREWYSSLFAFPVYQGSTDILFL WNDMNEPSVFRGPEQTMQKNAIH HGNWEHRELHNIYGFYHQMATAEG LIKRSKGKERPFVLTRSFFAGSQKYG AVWTGDNTAEWSNLKISIPMLLTLSI TGISFCGADIGGFIGNPETELLVRWY QAGAYQPFFRGHATMNTKRREPWL FGEEHTRLIREAIRERYGLLPYWYSL FYHAHVASQPVMRPLWVEFPDELKT FDMEDEYMLGSALLVHPVTEPKATT VDVFLPGSNEVWYDYKTFAHWEGG CTVKIPVALDTIPVFQRGGSVIPIKTT VGKSTGWMTESSYGLRVALSTKGSS VGELYLDDGHSFQYLHQKQFLHRKF SFCSSVLINSFADQRGHYPSKCVVE KILVLGFRKEPSSVTTHSSDGKDQPV AFTYCAKTSILSLEKLSLNIATDWEV RII	None	None	None	None	None	None	None	None	None