	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	extracel region
0	P70699	LYAG_MOUSE	Mus musculus	Lysosomal alpha-glucosidase	37.855485	S142;T469	NaN	33300544;40885482;40021952	MNIRKPLCSNSVVGACTLISLTTAVI LGHLMLRELMLLPQDLHESSSGLW KTYRPHHQEGYKPGPLHIQEQTEQP KEAPTQCDVPPSSRFDCAPDKGISQE QCEARGCCYVPAGQVLKEPQIGQPW CFFPPSYPSYRLENLSSTESGYTATL TRTSPTFFPKDVLTLQLEVLMETDSR LHFKIKDPASKRYEVPLETPRVLSQA PSPLYSVEFSEEPFGVIVRRKLGGRV LLNTTVAPLFFADQFLQLSTSLPSQH ITGLGEHLSPLMLSTDWARITLWNR DTPPSQGTNLYGSHPFYLALEDGGL AHGVFLLNSNAMDVILQPSPALTWR STGGILDVYVFLGPEPKSVVQQYLDV VGYPFMPPYWGLGFHLCRWGYSST AIVRQVVENMTRTHFPLDVQWNDL DYMDARRDFTFNQDSFADFPDMVR ELHQDGRRYMMIVDPAISSAGPAGS YRPYDEGLRRGVFITNETGQPLIGKV WPGTTAFPDFTNPETLDWWQDMVS EFHAQVPFDGMWLDMNEPSNFVR GSQQGCPNNELENPPYVPGVVGGIL QAATICASSHQFLSTHYNLHNLYGL TEAIASSRALVKTRGTRPFVISRSTFS GHGRYAGHWTGDVRSSWEHLAYSV PDILQFNLLGVPLVGADICGFIGDTS EELCVRWTQLGAFYPFMRNHNDLN SVPQEPYRFSETAQQAMRKAFALRY ALLPYLYTLFHRAHVRGDTVARPLFL EFPEDPSTWSVDRQLLWGPALLITP VLEPGKTEVTGYFPKGTWYNMQMV SVDSLGTLPSPSSASSFRSAVQSKGQ WLTLEAPLDTINVHLREGYIIPLQGP SLTTTESRKQPMALAVALTASGEAD GELFWDDGESLAVLERGAYTLVTFS AKNNTIVNKLVRVTKEGAELQLREV TVLGVATAPTQVLSNGIPVSNFTYSP DNKSLAIPVSLLMGELFQISWS	False	False	1.868	1.153	1.941	1.421	1.07	1.681	1.976