	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	extracellula region
0	Q9DBE8	ALG2_MOUSE	Mus musculus	Alpha-1,3/1,6-mannosyltransferase ALG2	26.986372	S291	NaN	33300544;30059200	MAENLYRARSRVYSPSVLFLHPDMG IGGAERLVLDAALALQEYGCDVKIWT AHYDPNHCFIETRELSVQCAGDWLP RSLGWGGRGAAICSYVRMVFLALYV LFLSGEEFDVVVCDQVSACIPVFKLA RRRKRVLFYCHFPDLLLTQRNSALK KFYRAPIDWIEEYTTGMADRILVNSQ YTASVFKETFKTLSHRNPDVLYPSLN IGSFDLAIPEKIDDLVPKGKQFLFLSI NRYERKKNLPLALRSLVQLRNRLPS QEWDKVHLFMAGGYDDRIPENVEH YKELKKMVQESDLERHVTFLRSFSD RQKISLLHGCLCVLYTPSNEHFGIVP LEAMYMQCPVIAVNNGGPLESIVHK VTGFLCEPDPVHFSEAMEKFIHKPS LKATMGLAGKARVAEKFSADAFADQ LYQYVTKLV	None	None	None	None	None	None	None	None	None