

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
0 Q9Y673	ALG5_HUMAN	Homo sapiens	Dolichyl-phosphate beta-glucosyltransferase	21.366884	NaN	NaN	28510447	MAPLLQLAVLGAALAAAALVLISIV AFTTATKMPALHRHEEEKFFLNAKG QKETLPSIWDSPTKQLSVVPSYNEE KRLPVMMDEALSYLEKRQKRDPAFT YEVIVDDGSKDQTSKVAFKYCQKY GSDKVRVITLVKNRGKGGAIRMGIFS SRGEKILMADADGATKFPDVEKLEK GLNDLQPWPNQMAIACGSRHLEK ESIAQRSYFRITLLMYGFHFLVWFLC VKGIRDTQCGFKLFTREASRTFSSL HVERWAFDVELLYIAQFFKPIAEIAV NWTEIEGSKLVPFWSWLQMGKDLL FIRLRYLTGAWRLEQTRKMN	False	False	2.893	1.27	1.159	4.077	1.494	1.348	1.086