	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	Q9NP73	ALG13_HUMAN	Homo sapiens	Putative bifunctional UDP-N-acetylglucosamine transferase and deubiquitinase ALG13	25.835862	NaN	NaN	26374642	MKCVFVTVGTTSFDDLIACVSAPDSL QKIESLGYNRLILQIGRGTVVPEPFST ESFTLDVYRYKDSLKEDIQKADLVIS HAGAGSCLETLEKGKPLVVVINEKL MNNHQLELAKQLHKEGHLFYCTCR VLTCPGQAKSIASAPGKCQDSAALTS TAFSGLDFGLLSGYLHKQALVTATH PTCTLLFPSCHAFFPLPLTPTLYKMH KGWKNYCSQKSLNEASMDEYLGSL GLFRKLTAKDASCLFRAISEQLFCSQ VHHLEIRKACVSYMRENQQTFESYV EGSFEKYLERLGDPKESAGQLEIRAL SLIYNRDFILYRFPGKPPTYVTDNGY EDKILLCYSSSGHYDSVYSKQFQSSA AVCQAVLYEILYKDVFVVDEEELKTA IKLFRSGSKKNRNNAVTGSEDAHTD YKSSNQNRMEEWGACYNAENIPEG YNKGTEETKSPENPSKMPFPYKVLK ALDPEIYRNVEFDVWLDSRKELQKS DYMEYAGRQYYLGDKCQVCLESEGR YYNAHIQEVGNENNSVTVFIEELAE KHVVPLANLKPVTQVMSVPAWNAM PSRKGRGYQKMPGGYVPEIVISEMDI KQQKKMFKKIRGKEVYMTMAYGKG DPLLPPRLQHSMHYGHDPPMHYSQ TAGNVMSNEHFHPQHPSPRQGRGY GMPRNSSRFINRHNMPGPKVDFYP GPGKRCCQSYDNFSYRSRSFRRSHR QMSCVNKESQYGFTPGNGQMPRGL EETITFYEVEEGDETAYPTLPNHGGP STMVPATSGYCVGRRGHSSGKQTLN LEEGNGQSENGRYHEEYLYRAEPDY ETSGVYSTTASTANLSLQDRKSCSM SPQDTVTSYNYPQKMMGNIAAVAAS CANNVPAPVLSNGAAANQAISTTSV SSQNAIQPLFVSPPTHGRPVIASPSY PCHSAIPHAGASLPPPPPPPPPPPPP PPPPPPPPPPPPPALDVGETSNLQPP PPLPPPYSCDPSGSDLPQDTKVLQY YFNLGLQCYYHSYWHSMVYVPQMQ QQLHVENYPVYTEPPLVDQTVPQCY SEVRREDGIQAEASANDTFPNADSS SVPHGAVYYPVMSDPYGQPPLPGFD SCLPVVPDYSCVPPWHPVGTAYGGS SQIHGAINPGPIGCIAPSPPASHYVPQ GM	None	None	None	None	None	None	None	None	None