

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q08379	GOGA2_HUMAN	Homo sapiens	Golgin subfamily A member 2	11.996054	S607	S37;S66;S273;S438;S690;S937;S953;S981	34019948;34725712;37217939;33465208;23301498	MWPQPRLPPRPAMSEETRQSKLAA AKKKLREYQQRNSPGVPTGAKKKKK IKNGSNPETTTTSGGCHSPEDTPKDN AATLQPSDDTVLPGGVSPGASLTS MAASQNHADNVPNLMDETKTFSS TESLRQLSQNLGLVCESATCVNGE GPASSANLKDLESRYQLAVALDSS YVTNKQLNITIEKCLKQONQEITDQLE EEKKECHQKOGALREQLQVHIQTIGI LVSEKAEQLTALAHTQHAARQKEGE SEDLASRLQYSRRRVGELERALSAVS TQQKKADRYNKELTKERDALRLELY KNTQSNEDLKQEKSELEEKLRVLVT EKAGMQLNLEELQKKLEMTLELLQ QFSSRCEAPDANQQQLQAMEERAQ LEAHLGQVMESVRQLQMERDKYAE NLKGESAMWRQRMQQMSEQVHTL REEKECSMSRVQELETSLAELRNQ MAEPPPPPEPPAGPSEVEQQLQAEAE HLRKELEGLAGQLQAVQDNEGLS RLNREQEERLLELERAELWGEQAE ARRQILETMQNDRTTISRALSQNRE LKEQLAELQSGFVKLTNENMEITSA LQSEQHVKRELGKKLGELQEKLSEL KETVELKSQEAQSLQQQRDQYLGHL QQYVAAYQQLTSEKEVLHNQLLQ QLVDQLQQQEAQGKAVAEMARQEL QETQERLEAATQONQQLRAQLSLM AHPGEGDGLDREEEDEEEEEEEAV AVPQPMPSIPEDLESREAMVAFFNS AVASAEQQARLRGQLKEQRVRCRR LAHLLASAQKEPEAAAPAPGTGGDS VCGETHRALQGAMEKLQSRFMELM QEKADLKERVEELEHRCIQLSGETD TIGEYIALYQSQRAVLKERHREKEYI SRLAQDKEEMKVKLLELQELVLRV GDRNEWHGRFLAAAQNPADPTSG APAPQELGAANQQGDLCEVSLAGSV EPAQGEAREGSPRDNPTAQQIMQLL REMQNPRERPGLGSNPCIPFFYRAD ENDEVKITVI