

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q08379	GOGA2_HUMAN	Homo sapiens	Golgin subfamily A member 2	10.721651	S607	S37;S66;S273;S438;S690;S937;S953;S981	23301498;33465208;34019948	MWPQPRLPPRPAMSEETRQSKLAA AKKKLREYQQRNSPGVPTGAKKKKK IKNGSNPETTTTSGGCHSPEDTPKDN AATLQPSDDTVLPGGVPSPGASLTS MAASQNHADNVPNLMDETKTFSS TESLRQLSQQLNGLVCESATCVNGE GPASSANLKDLESRYQQLAVALDSS YVTNKQLNITIEKLLKQONQEITDQLE EEKKECHQKQGALREQLQVHIQTIGI LVSEKAELQTALAHTQHAARQKEGE SEDLASRLQYSRRRVGELERALSAVS TQKKADRYNKELTKERDALRLELY KNTQSNEDLKQEKSELEEKLRVLVT EKAGMQLNLEELQKKLEMTELLLQ QFSSRCEAPDANQQLQQAMEERAQ LEAHLGQVMESVRQLQMERDKYAE NLKGESAMWRQRMQQMSEQVHTL REEKECSMSRVQELETSLAELRNQ MAEPPPEPPAGPSEVEQQQQAQAEAE HLRKELEGLAGQLQAQVQDNEGLS RLNREQEERLLELERAELWGEQAE ARRQILETMQNDRTTISRALSQNRE LKEQLAELQSGFVKLTNENMEITSA LQSEQHVKRELGKKLGELQEKLSSEL KETVELKSQEAQSLQQQRDQYLGHL QQYVAAYQQLTSEKEVLHNQLLLQT QLVDQLQQQEAQGKAVAMARQEL QETQERLEAATQQNQQLRAQLSLM AHPGEGDGLDREEEEDDEEEEEEAV AVPQPMPSIPEDLESREAMVAFFNS AVASAEQARLRGQLKEQVRRCRR LAHLLASQKEPEAAAPAPGTGGDS VCGETHRALQGAMEKLQSRFMELM QEKADLKERVEELEHRCIQLSGETD TIGEYIALYQSQRVLRKERHREKEYI SRLAQDKEEMKVKLELQELVLRVL GDRNEWHGRFLAAAQNPADPTSG APAPQELGAANQQGDLCEVSLAGSV EPAQGEAREGSPRDNPTAQQIMQLL REMQNPRERPGLGSNPCIPFFYRAD ENDEVKITVI