

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q921M4	GOGA2_MOUSE	Mus musculus	Golgin subfamily A member 2	9.739272	S385;S686	S37;S66;S273;S438;S697;S934;S978	34418053	MWPPRFPPPRPGMSEETRQSKLAA AKKKLREYQQKNSPGVPAGAKKKKK IKNGHSPERPTASDCQSPENVPTDH IAPAPPTAATDTMFLGVTPSPDADLT QSHDAGNCSNLMEETKTFSSSTESLR QLSQQLNGLVSESTSYINGEGLTSS NMKELENRYQELAVALDSSYVTNK QLSSTIEELKQONQDTLDQLEKEKK DYQQKLAKEQGSREQLQVHIQTIGI LVSEKAELQATALAHTQQAARQKAGE SEDLASRLQSSRQRVGELERTLSTVS TQKKQADRYNKDLTKERDALKLELY KNSKSNEDLRQONSELEEKLRLVVA EKAAAQLGVEELQKKLEMSELLLQQ FSSQSSAAGGNEQLQHAMEERAQL ETHVSQLMESLKQLQVERDQYAE LKGESAMWQQRVQQMAEQVHTLK EEKEHRERQVQELETSLAALRSQME EPPPEPPAGPSEAEELQGEVEQL HKELERLTGQLRAQVQDNESLSHL NREQEGRLLELERAQRWSEQAE RKQILESMQSDRTTISRALSQRELK EQLAELQNGFVRLTNNMEITSALQ SEQHVKKELARKLQERLQERLQELKE TVELKSQEAQGLQEQRDQCLSHLQ QYAAAYQQHLAAYEQLTSEKEAIHK QLLLQTQLMDQLQHHEEVQGMMAE LARQELQEAQERLQKATSQENQQLQA QLSLLVLPGEQDQVDEEEDVEVQPS SLAIPEDLDSREAMVAFFNAAIARAE EEQARLRVQLKEQKARCRSLSHLAA PVQSKLEKEAVVPRNVDDSAEESN QALHVAMEKLQSRFLEVMQEKVEL KERVEELEHCCIQLSGETDTIGEYIA LYQNQRAVLKARHLEKEEYISRLAQ DKEEMKVKLELQELVLRVNERNE WQKFLAVSQNPGDVLTPVPTGSQE FGAADQDDLREVS LADDIEPAQGE AGVPAPHENPTAQIMQLLREIQNP RERPGLGSNPCIPFFYRADENDEVKI MVV