

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	g
P57740	NU107_HUMAN	Homo sapiens	Nuclear pore complex protein Nup107	23.321817	T64;S67;S652;S893	S4;S10;S11;S37;T46;T55;S57;S58;T64;S69;S86	32574038;30379171;31492838;35132862;33214551;34019948;30397120;35254053;35289036;32119511;30059200;34725712;22121020;33465208;38665916	MDRSGFGEISSPVIREAEVTRTARKQ SAQKRVLLOASQDENFGNTTPRNO VIPRTIPSSFRQPFPTPSRLLRQPDIS CILGTGGKSPRLTQSSGFFGNLSMV TNLDDSNWAAAFSSQRSGLFTNTEP HSITEDVTISAVMLREDDPGEAASM SMFSDFLQSFLLKHSSTVFDLVEEY ENICGSQVNLISKIVSRATPGLQKFS KTASMLWLLQEMVTWRLLASLYR DRIQSALEEEVSFAVTAVNASEKTVV EALFQRDSLVRQSQLVVDWLESIAK DEIGEFSDNIEFYAKSVYWENTLHTL KQRQLTSYVGSVRPLVTELDPAPIR QKMPLDDLDREDEVRLKYLFTLIR AGMTEEAQRLCKRCGQAWRAATLE GWKLYHDPNVNGGTELEPVEGNPY RRIWKISCWRMAEDELFNRYERAIY AALSGNLKQLLPVCDTWEDTVWAYF RVMVDSLVEQEIQTSVATLDETEELP REYLGANWTLEKVFEELQATDKKRV LEENQEHYHIVQKFLILGDIDGLMD EFSKWLKSRNNLPGHLLRFMTHLI LFFRTLGLQTKEEVSIEVLKTYIQLLI REKHTNLIAYFTCHLPQDLAVAQYAL FLESVTEFEQRHHCLELAKEADLDV ATITKTVVENIRKKNDFESHHDLP ALDTGTTEEDRLKIDVIDWLVFDPAQ RAEALKQGNAMRKFSLAKKHEAAK EVFVKIPQDSIAEINYQCEEQGMESP LPAEDDNAIREHLICIRAYLEAHETFN EWFKHMNSVPQKALIPQPTFTTEKV AHEHKEKYE MDFGIWKGHLDAL ADVKEKMYNVLVFDGGWMDVRE DAKEDHERTHQMVLLRKLCLPMLC FLLHTILHSTGQYQECLQLADMVSS ERHKLYLVFSKEELRKLQKLRSSL MLLDQGLDPLGYEIQ	True	False	4.591	5.0	1.474	1.523	1