

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrar
075694	NU155_HUMAN	Homo sapiens	Nuclear pore complex protein Nup155	22.294592	T587;S992;T1258	S1057	30379171;35083852;33214551;33465208;34019948;30397120;38253038;34725712;25367160;22121020;35008409;28510447	MPSSLLGAAMPASTSAAALQEALEN AGRLIDRQLQEDRMYPDLSELLMVS APNNPTVSGMSMDMDYPLQGPGLLS VPNLPEISSIRRVLPPELVEQFGHM QCNCMMGVFPPIISRAWLTIDSDIFM WNYEDGGDLAYFDGLSETILAVGLV KPKAGIFQPHVRHLLVLATPVDIVIL GLSYANLQTGSGVLNDSLGGMQLL PDPYLSLPTDNTYLLTTSTDNGRIFL AGKDGCLYEVAYQAEAGWFSQRCP KINHSSLSFLVPSLLQFTFSEDDP ILQIAIDNSRNILYTRSEKGVIVYDL GODGQGMSRVASVSQNAIVSAAGNI ARTIDRSVFKPIVQIAVIENSELDCQ LLAVTHAGVRLYFSTCFRQPLARP TLTLVHVRLPPGFSASSTVEKPSKVH RALYSKGILLMAASENEDNDILWCV NHDTFFQKPMMETQMTAGVDGH SWALSAIDELKVDKIITPLNKDHPIT DSPVVVQQHMLPPKKFVLLSAQGS MFHKLRPVDQLRHLLVSNVGGDGE EIERFFKLHQEDQACATCLILACSTA ACDREVSAWATRAFFRYGGEAQM FPTTLPPSNVGPILGSPVYSSSPVPS GSPYPNPSFLGTPSHGIQPPAMSTPV CALGNPATQATNMSCVTPGEIVVSG KHNGICIYFSRIMGNIWDASLVVERI FKSGNREITAIESSVPCOLLESVLQE LKGLEFLDRNSQFAGGPLGNPNTT AKVQQRLLIGMRPENGNPQQMQQE LQRKFHEAQLSEKISLQAIQQLVRKS YQALALWKLCEHQFTIIVAELOKEL QEQLKITTFKDLVIRKELTGALIASL INCYIRDNAAVDGISLHLQDICPLLYS TDDAICSKANELQRSROVQNKTEK ERMLRESLKEYQKISNOVDLSNVCA QYRQVRFYEGVVELSLTAAEKDPQ GLGLHFYKHGPEEDIVGLQAFQER LNSYKCITDTLQELVNQSKAAPQSPS VPKKPGPPVLSDDPNMLSNEEAGH HFEQMLKLSQRSKDELFSIALYNWL IQVDLADKLLQVASFLEPHLVRMA KVDQNRVRYMDLLWRYYEKNRSFS NAARVLSRLADMHSTEISLQQRLEYI ARAILSAKSSTAISSIAADGEFLHELE EKMEVARIQLQIQETLQRQYSHHSS VQDAVSQDSELMITKLYGEFADP FKLAECKLAIHCAGYSDPILVQTLW QDIIEKELSDSVTLSSDRMHASLKL IVLLGKIYAGTPRFFPLDFIVQFLEQQ VCTLNWDVGFVIQTMNEIGVPLPRL LEVYDQLFKSRDPFWRNRMKPLHL LDCIHVLLIRYVENPSQVLNCERRRF TNLCLDAVCGYLVELQSMSSSVAVQ AITGNFKSLQAKLERLH	True	False	4.425	5.0	1.123	1.572	0.997	2.354