

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P37199	NU155_RAT	Rattus norvegicus	Nuclear pore complex protein Nup155	29.853773	S525	S1056	12438562	MPSMLGSMVASTSAPSLQEALEN AGRLIDRQLQEDRMYPDLSELLMVS APNSPTVSGMSDMDYPLQGPGLLSV PSLPEISTIRRVPLRLSWLNSLDTCS VTAMMGVFPPIRAWLTISSDIFMW NYEDGGDLAYFDGLSETILAVGLVKP KAGIFQPHVRHLLVLATPVDIVILGLS YANVQTGSGILNDSVCGGLQLLPDP LYSLPTDNTYLLTITSTDNGRIFLAGK DGCLYEYVAYQAEAGWFSQRCKINH SKSSLSFLVPSLLQFTFSEDDPIVQIE IDNSRNILYTRSEKGVIOVYDLGHDG QGMSRVASVSQNAIVCAAGNIARTI DRSVFKPIVQIAVIENSESLDCQLLA VTHAGVRLYFSTCFRQPLARPNTLT LVHVRLPPGFASSTVEKPSKVHKAL YSKGILLMTASENEDNDILWCVNHD TFPFQKPMMETQMTTRVDGHSWAL SAIDELKVDKIITPLNKDHIPTDSPV VVQOHMLPPKFFVLLSAQGSMLFH KLRPVDQLRHLLVSNVGGDGEIEER FFKLHQEDQACATCLILACSTAACD REVSAAWATRAFFRYGGEAQMRFPAT LPTPSNVGPILGSPMYSSSPVPTGSP YPNPSSLGTPSHGAQPPTMSTPMSA VGNPAMQAASLSGLTGPEIVYSGKH NGICIFYSRIMGNIWDASLVVERVFK SSNREITAIESSVPIQLLESVLQELKG LQEFLDNRNSQFSGGPLGNPNTTAKV QQRLLGVMRPENGNTQQMQQELQ RKFHEAQLSEKISLQAIQQLVRKSYQ ALALWKLCEHQFTVIVGELQKEFQ EQLKITTFKDLVIREKEVTGALIASLI NCYIRDNAAVDGLSLHLQDTCPLLYS TDDAVCSKANELLQRSRQVQSKSER ERMLRESLKEYQKISNQVDLPSVCA QYRQVRFYEGVVLSLTAEEKDPQ GLGLHFYKHGEPEEDVVGLQTFQER LNSYKCIDTLQELVNQSKAAPQSPS VPKPGPPVLSDDPNMLSNEEAGH HFEQMLKLAQRSKDELFSIALYNWL IQADLADKLLQIASPFLEPHLVRMAK VDQNRVRYMDLLWRYYEKNRSFSS AARVLSKLADMHSTEISLQORLEYIA RAILSAKSSTAISIAADGEFLHELEE KMEVARIQLQIQETLQRQYSHHSSV QDAISQLDSELMDITKLYGEFADPFK LAECKLAIHCAGYSDPILVHTLWQD IIEKELSDSVTLSSDRMHALSCLKV LLGKIYAGTPRFFPLDFIVQFLEQQV CTLNWDVGFVIQTMNEIGVPLPRL EVDQLFKSRDPFVNRVKSPLHLLD CIHVLLTRYVENPSLVLCERRRFT NLCLDAVCGYLVELQSMSSSVAVQA ITGNFKSLQAKLERLH	None	None	None	None	None	None	None		

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