

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9W1X4	NU214_DROME	Drosophila melanogaster	Nuclear pore complex protein Nup214	10.212237	NaN	NaN	33925313	MAQNAPDCEDTQDFQFKLHHKIVA FKKCGEPRSNVNLLAVSSSRGLLFA GSPTQPELKVIVKDLVNAKSTAQQP QARLVPLSPIPNYIACSSDGNLLAVN HTQNGTSLLSIYAVLSFMTDPDVRPVY NIRLAAEDHVHGVQLLWNPVLPNSL AVVLSNGALAMYALKEGGNFEMHS LDKNQQVKCGCWSPKGGKQIVLGFP GGTVKQFKPDLTLAKTLLCPPNIHDA PFDTIAIQWLSTFQFAVIFLQHGEDC SPSLYILNAPKAGAPSYINYYDICYSM NGPRNHQFVFSHVPQWNLLLVSVA NGVEVGIMRSTEAGDTPAWQQLTLL DEARIEMPLSEDKDETFFPLGFAFDTS TTHQLTINEKKLQTMPMVHVLSSDG ELLSFNFLNVLPTAVSVCSPPPPVAD TSGQFKPLNMLLASEEEEEQPAAWAAS PSKAPAATPAASSDISFAFTPNTVTST PAPSKDKQPSLFSGFGAAAAKAPAP QLSFGTAPTSSPVSGAPTTNAAKPT TPFGGFGTQATTTAMGSMFASGA NAFGGMALNKPAIASVTPRTAAPGS TVPATPASAPANPLYTVPLTFTPVD TKPATSAPPQIADESLKPDDEPIIKD MIALQIEAFSKDIQKQKEQTKELLKG IAAPSALRAYAKRLDDLQELNEQAK DVEFELDVQGLRQGLNEAYIVAEC RGKLEIYRKPEITRLMNSSSCDPSGR RMLARLQSYVAANEACLRLAQQHV DLQWEQFQDVVRRNSKSRMHMPC LEGIYQRLTRLQNLTSNQRIVQNNIK SKLKERGLLQAALLDQEKSRTRTNE AVDTLTDLSILTMSLSQVVDNSNAAKLT RERLQKIRNIVQLQKINVICPQRPDR VGLKSEVILETKRRAEQIKRAAAKPA TANKYTQAAVAPPSPPDVAPTPAVAP MPQATVTVAPPLPKPMPSIPSVVEKP GVPTHPTTPVATPFSFSQSIPFVKTST VTPTNTVTPGAAKPGLSIFGGSTIS SFSFGGGAAKSALSFGGTGSPAVAAPT PKPNPLSAVEKPTPEPTKPKKEQKAAE SKEFKAVQPETEESKVPQKPKAETE NKSFGFGGFTGTGGTVGNTSSSPFS FGGLGSSLGFGGTAAAVPKSEPSSTA TTSVATSASTAPFGIFSAALAKPSNS EPITTVTSNTTTTITSKPTNVIASSSVT DAPSVTTTNAVTSSTDPIGGLFSSVTI CKPNTPADTTK PANIFGGGLSAGSFS FGGTIDASKGLFGSTKPVATAPTSVT

							EANNKTDPISTTPSAISTTTATTTVSS PAVVPAAVTAAVPATSSTTVTSSTAV PGSAFSSNAFSTLSAGGAAAPTSA SAPLAAKSPTATSTGNNSSNSVFGG GFAVATSTAAPVASPFQSAAKSPVSS ANIFGSIPKAETSVFGGATTAPSNTT AAATPDAPPAGLFAAAISNPSPFGS PTTRAPASGGNIFGQAVKPSVFGQPA QAGDSGGSIFGGGSASTPFGSSSIFG GGNTQGAVGAPAAGSTSIFGQKVFG QSSAAAPAAGGNIFSNPVGSPQASPF GGGNSIFGSPATAPPASGGSIFGGG SSGGFGSFTQTTPAQGGFGGGFGQ GGGGSVAQTGFGSPQAPQQQTTTPG GFGAKPVFGGSPAFGASPTFGGGAT FGSPKGFGGFGGASPVASPPPFGAA AKPAQGNIFETLGGQESGLSFGNLA QTGNSNAQKPAFGGSSFMNYR
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