

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q17602	NU214_CAEL	Caenorhabditis elegans	Nuclear pore complex protein 14	12.013438	T885	NaN	31682414	MSNEDVAEDVSQVTDFHFHTCRKF RLFSSKSDGYSQNEINIRNRVATSSQ LGVTFTVNSNQLSCFHTKSLLGYKI TRENMNVEVTDLPKIRTIRLHGVLIN DMGVNSDGTVLGVLHTKNNDVSVD VFDIKKICTSSSIEPFKPLCTTRVGTE QINQGSCLWNPAPFDTFASSTDR SILVAKINVQSPANQKLVGIGKFGAV TTAISWSPKGGKLTIGDSLKIVQLK PELEVRSQHGPENKPNYGRITGLC WLATTEWLVSLENGTDQDAYLMRC KKDKPTEWIQFHELSSSSKWPLPP QLFPATQLLDVNWVIVGNSKTSEIS TVGKRDDWQTVVPEGESIYLPPTS SGKDTVPIGVAVDRSMTDEVLLNPD GSQRHRPSPLVCLTNDGILTAHHII STFAAHIPCQMSSQNLAINDLKKLQ FDSQKPISAPSDQTPVTKPSTVFGQ KPEAETLKSSLVGSPPSVQTPKSSS LFNPKSIASNIETSQLTESKPSTPAAP SSQPKIASPKSEAIPKISDKTLEHKK AELIATKKQVLIERMDKINDSMAGA KDATMKLSFAVGKVTTIMECADVV RASLGDSKEVMDELKNLILSIERMS DRTQHTVKEMDFEIDEKMELVAGV EDGNQVLEKLRNMSETEKLMRFNK LETAADLLNGKYECSDLIKLRMS LSEKESLRKQAILSPLRLSSNLNQLR SGSETELALKVMRNVSKIIMDTREIQI QRTELEFVRFQRDVVKFQNFKKGKE NLNFTQPLEMSSLDGDAPQGKSLTD AESIKVRQALVNQIQKRGIVKTRNVI VESYKKESENSAAMKNDLLDTSNLSN AILKLSMTPRRVMPSSSLFSASPSTP STKSDAATQADEPPIVKTVVVTVESE AKPIASAPAVSSPLIKLNTTTATTTMT TPKVTVPKEEANKTQDQKPIISTPAS SSIFSSGSLFGTKTQPLVSKEESTLT TGVPSLINSSLSISPQIEKASKVET LNKTEEVKDEKSENEVTPDLKSEEP KSLETKVKEEPKPAVQTPVKEEETGS NIQKTPSFSFNSTTTPKSTSSTSSIFG GGLKTQTPSSSNSTNIFGARTTTTAT PTPASNTSSIFGGGSKAASSPFGSFG QAGCQPAKTSNPATSTASVTFNFNT GATSASAKPAGFGSFGAGASAKPSS VFGGSVTAPTVPNVDDGMEDDSMA NGGGSGGFMSGLGNARTSNTSGGN NPFAPKTSTGTSASSSSWLFGGGGN

								QQQQQQKPSFSFNTAGSSAQQAS APATGTSSVFGGAPKFGSQPAFGAK PFGGANAGLSKNASIFGGATSSTT NNPATGGFAQFASGQKTSSLFGGGA TPQTNTSIFGGANTTPAPTSSVFGG GASANANKPTSFTSWR
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