

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q17602	NU214_CAEL	Caenorhabditis elegans	Nuclear pore complex protein 14	10.335743	T885	NaN	31682414	MSNEDVAEDVSQVTDHFHHTCRKF RLFSSKSDGYSQNEINIRNRVATSSQ LGVTFTVNSNQLSCFHTKSLLGYKI TRENMNVEVTDLPKIRTIRLHGVLIN DMGVNSDGTVLGVLHTKNNDVSVD VFDIKKICTSSSIEPFKPLCTTRVGTE QINQGSLEWNPAPFDTFASSTDR SILVAKINVQSPANQKLVGIGKFGAV TTAISWSPKGGKLTIGDSLKIVQLK PELEVRSQHGPENKPNYGRITGLC WLATTEWLVSLENGTDQDAYLMRC KKDKPTEWIOFHESYSSSKWPLPP QLFPATQLLDVNVVIVGNSKTSEIS TVGKRDDWQTVVPEGESIYLPPTS SGKDTVPIGVAVDRSMTDEVLLNPD GSQRHRPSPLVCLTNDGILTAHHII STFAAHIPCQMSSQNLAINDLKKLQ FDSQKPISAPSDQTPVTKPSTVFGQ KPEAETLKSSLVGSPPSVQTPKSSS LFNPKSIASNIETSQLTESKPSTPAAP SSQPKIASPKSEAIPKISDKTLEHKK AELIATKKQVLIERMDKINDSMAGA KDATMKLSFAVGKVKTTIMECADVV RASLGDSKEVMDELKNLILSIERMS DRTQHTVKEMDFEIDEKMELVAGV EDGNQVLEKLRNMSETEKLMRFNK LETAADLLNGKYECSDLIKLRMS LSEKESLRKQAILSPLRLSSNLNQLR SGSETELALKVMRNVSKIIMDTREIQI QRTELEFVRFQRDVVKFQNFKKGKE NLNFTQPLEMSSLDGDAPQGKSLTD AESIKVRQALVNQIQKRGIVKTRNVI VESYKKESENSAAMKNDLLDTSNLSN AILKLSMTPRRVMPSSSLFSASPSTP STKSDAATQADEPPIVKTVVVTVESE AKPIASAPAVSSPLIKLNTTTATTTMT TPKVTVPKEEANKTQDQKPIISTPAS SSIFSSGSLFGTKTQPLVSKEESTLT TGVPSLINSSLSISPQEIEKASKVET LNKTEEVKDEKSENEVTPDLKSEEP KSLETKVKEEPKPAVQTPVKEEETGS NIQKTPSFSFNSTTTPKSTSSTSSIFG GGLKTQTPSSSNSTNIFGARTTTTAT PTPASNTSSIFGGGSKAASSPFGSFG QAGCQPAKTSNPATSTASVTFNFNT GATSASAKPAGFGSFGAGASAKPSS VFGGSVTAPTVPNVDDGMEDDSMA NGGGSGGFMSGLGNARTSNTSGGN NPFAPKTSTGTSASSSSWLFGGGGN

								QQQQQQKPSFSFNTAGSSAQQAS APATGTSSVFGGAPKFGSQPAFGAK PFGGANAGLSKNASIFGGATSSTT NNPATGGFAQFASGQKTSSLFGGGA TPQTNTSIFGGANTTPAPTSSVFGG GASANANKPTSFTSWR
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