

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
Q2M1P5	KIF7_HUMAN	Homo sapiens	Kinesin-like protein KIF7	19.994096	NaN	S898	37217939	MGLEAQRLPGAEEAPVRVALRVRPL LPKELLHGHSCLQVEPGLGRVTLG RDRHFQGFHVLAEDAGQEAQVQAC VQPLLEAFFEFGFNATVFAYGQTGSG KTYTMGEASVASLLEDEQGVPRAM AEAFKLIDENLLDCLVHVSYLEVYK EEFRDLLEVGTASRDIQLREDERGN VVLGCVKEVDVEGLDEVLSLEMGN AARHTGATHLNHLSSRSHTVFTVTL EQRGRAPSRLPRPAPGQLLVSKFHF VDLAGSERVLTGSGTERLKEIQIN SLLALGNVISALGDPQRRGSHIPYR DSKITRILKDSLGGNAKTVMIACVSP SSSDFDETLNLTNYASRAQNIENRA TVNWRPEAERPPEETASGARGPPRH RSETRIIHRGRRAPGPATASAAAAMR LGAECARYRACTDAAYSLLRELQAEP GLPGAAARKVRDWCALVEGERSALS SASGPDSGIESASVEDQAAQAGGGR KEDEGAQQLLTQNOVARLEENR DFLALEDAMEQYKLOSDRLREQQE EMVELRLRLELVRPVGWGGPRLNNG LPPGSFVPRPHTAPLGGAAHVLGM VPPACLPGDEVGSEQRGEQVTNGRE AGAEELLTEVNRLGSGSSAASEEEEE EEPPRRTLHLRRNRISNCSQRAGA RPGSLPERKGPCLCEELDAAIPGSR AVGGSKARVQARQVPPATASEWRLA QAQKIRELAINIRMKEELIGELVRT GKAAQALNRQHSQRIRELEQEAQV RAELSEGQRQLRELECKELQDAGER SRLQEFRRRVAAAQSQVQLKEKKQ ATERLVLSAQSEKRLQELERNVQL MRQQQGQLQRRLREETEQRRLLEA EMSKRQHRVKELELKHEQQQKILKI KTEEIAAFQKRRRSGSNGSVVLEQ QQKIEQKKWLDQEMEKVLQORRA LEELGELHKKREAILAKKEALMQEK TGLESKRLRSSQALNEDIVRVSSRLE HLEKELSEKSGQLRQGSAAQSQQIR GEIDSLRQEKDSLKQRLIDGKLRQ GSLLSPEEERTLFQLDEAIEALDAAIE YKNEAITCRQVLRASASLLSQCEM NLMAKLSYLSSSETRALLCKYFDKV VTLREEHQHQQIAFSELEMQLEEQQ RLVYWLEVALERQRLEMDRQLTLQ QKEHEQNMQLLLQQRDLHLEGLA DSRRQYEAIRIQALEKELGRYMWINQ ELKQKLGGVNAVGHSRGGEKRSLS EGRQAPGNEDELHLAPELLWLSPLT EGAPRTRETRDLVHAPLPLTWKRS SLCGEEQGSPEELRQREAAEPLVGR VLPVGEAGLPWNFGPLSKPRRELRR ASPGMIDVRKNPL	False	False	3.46	3.029	1.014	0.992	1.129	1.694	1.343