

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
Q8MSU4	SYMPK_DROME	Drosophila melanogaster	Symplekin	28.06426	T1000;T1001	NaN	40245499	MDSIIGRSQFVSETANLFTDEKTATA RAKVVWDWCNELVIASPTKCELLAK VQETVLGSCAEAEFLESVLSLAH DSNMEVRKQVVAFVEQVCKVKVEL LPHVINVVSMLLRDNASQVIKRVQA CGSIYKNGLYLCSLMEPGDSEAOA WNILSLIKAQILDIDNENDGIRTNA IKFLEGVVVLQSFADSDSLKRDGDFS LADVDPDHCTLFREKLQEEGNNILD ILLQFHGTTHISSVNLIACTSSSLCTIA KMRPIFMGAVVEAFKQLNANLPPTL TDSQVSSVRKSLKMQLOTLLKNRGA FEFASTIRGMLVDLGSSTNEIQKLIP KMDKQEMARRQKRILENAOQSLAK RARLACEQQDQQQREMELDTEELE RQKQKSTRVNEKFLAEHFRNPETVV TLVLEFLPSLPTVEVPQKFLQEYTPIRE MSIQQVVTNISRFQGEQLSEKRLGP GAATFSREPPMRVKVQAIESTLTA MEVDEDAVQKLSSEEFQRKEEATKK LRETMERAKGEQTVIEKMKERAKTL KLQEITKPLPRNLKEKFLTDVRRIL NSERQCICKGGVSSKRRKLVTVIAATF PDNVRYGIMEFILEDIKQRIDLAFSW LFEEYSLQGFTRHTYVKTENRPDH AYNELLNKLFIGIGERCDHKDKILIR RVYLEAPILPEVSIHGLVQLSLDDEF SQHGLELIKDLAVLRPPRKNRFVRVL LNFSVHERLDLRDLAQAHVLSLYHV HKILPARIDEFALEWLKFIEQESPPA AVFSQDFGRPTEEPDWREDITTKVCF GLAFTLLPYKPEVYLQICQVVFVSTS AELKRTILRSLDIPKMGVESPTLLQ LIEDCPKGMETLVIRIYILTERVPSP HEELVRRVRDLYQNKVKDVRVMIPV LSGLTRSELISVLPKLIKLNPAVVKEV FNRLLGIGAEFAHQTMAMTPTDILV ALHTIDTSVCDIKAIKATSLCLAERD LYTQEVLMVAVLQQLVEVTPPLTMM RTTIQSLTLYPRLANFVMNLLQRLIK QVWRQKVIWEGFLKTVQRLKQPSM PILLHLPPAQLVDALQCPDLRPLALS EYAESMQDEPMNGSGITQQVLDIIS GKSVDFVTDSEGGYISAEHIKKEAP DPSEISVISTVPVLTSLVPLPVPPPIGS DLNQPLPPGED	True	False	2.366	5.0	0.281	False	False	0.967	0.723