

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
P30665	MCM4_YEAST	Saccharomyces cerevisiae	DNA replication licensing factor MCM4	24.475439	NaN	S52;S56;S69	33229814	MSQQSSPTKEDNNSSSPVVNPDS VPPQLSSPALFYSSSSSQGDIYGRNN SQNLSSQGEGNIRAAIGSSPLNFPSSS QRQNSDVVFQSQGROGRIRSSASASG RSRYHSDLRSDRALPTSSSSLGRNG QNRVHMRRNDIHTSDLSSPRRIVDF DTRSGVNTLDTSSSSAPPSEASEPLR IIWGTNVSIECTTFRNFLMSFKYK FRKILDEREEFINNTTDEELYIKQL NEMRELGTSNLNDARNLLAYKOT EDLYHQLLNYPQEVISIMDQTIKDC MVSLIVDNNLDYDLDEIETKFKYKVRP YNVGSCKGMRELNPNDIDKLINLKG LVLIRSTPVIPDMKVAFKCNVCDHT MAVEIDRGVIQEPARCERIDCNEPN SMSLIHNRCSFADKQVIKQETPDF VPDGGTPHSISLCVYDELVDSCRAG DRIEVTGTFRSIPIRANSRQRLKSLY KTYVDVVHVKKVSDKRLDVTSTIE QELMQNKVDHNEVEEVROITDQDL AKIREVAAREDLYSLLARSIAPSIYEL EDVKKGILLQLFGGTNKTFTKGGRY RGDINILLCGDPSTSKSQILQYVHKIT PRGVYTSKGSSAVGLTAYITRDVDT KQLVLESGALVLSDGGVCCIDEFDK MSDSTRSVLHEVMEQQTISIAGKAGII TTLNARSSILASANPIGSRYPNLPV TENIDLPPPLLSRFDLVYLVLDKQVDE KNDRELAKHLTNLYLEDKPEHISQD DVLVPEFLTMYISYAKEHIHPITEAA KTELVRAYVGMKMGDDSRSEKRI TATTRQLESMIRLAEAHAKMKLKNV VELEDVQEAURLIRSAIKDYATDPKT GKIDMNLVQTGKSVIQRKLOEDLSR EIMNVLKQASDSMSFNELIKQINE HSQDRVSSDIQEALSRLQQEDKVIV LGEVRRSVRLNRRV	None	None	None	None	None	None	None		

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