

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q01846	MDM1_YEAST	Saccharomyces cerevisiae	Structural protein MDM1	9.365641	NaN	S670;S673;S692	33229814	MPKFPQFRLILVLFYLISMIQWSVITF SLGFFLNVCIFAYFVFFKSLPDLPKP QPRFVDIVPESSNTVDVDKELKSVE GLIQDGNAQIGKELESIVNLIKDFVQ PWFTKIDKNSDAEFLKVIKWRLLQT LLVVKDKLMKNDSASLIVLKLLPIFN KHFSTFCDAREAVLSDLTLERHKSA NIDLQIAVEFNKNYKIHKSLSLKPNA LQKEIEKSIRKTVIGLLPHLFDNDEL DSSLVFTLMTEVLTTCIISPLIFKFTD PDSWNLRIVSLSQNYFEEKHKVHKI RRMLSKELODHRKVMNDVANKDV GEPSSKLELNAEYTGKQFEHYLNQ LDSLLDLSDIKYVAYSLALKIYQLKEN EHLTKENLKYKRLLLSLNLIESKLS FPGSEIDTASKKLAREANYPDNLMD NGIVLKEMASFLTSITLKDIVDDSEF LPPFESFLGSPETQGSTFLEYSQTI ESFKNPLEDATSEDIISGYSGISTMQ LQEISSKFFHNNNLQNMKLLDEGL VKNILFRNSFQINNDEDTFILARKS VLLLQTEAIKYLDDRFLPLFKKTPSFL KMLSTSHIISTDIYAHFLSRIGGVNN PEQNKIKDNVKTDFMNPVRFANP GITDALDNIVNGSGSKPHKSRISSNP RYSQLFGSENDNIFKDKLFDENDN TSEISVVEDQLDHPRNMEKVSVSSG NSGLNPSQFYGSNNFRDNIASTISI DQIEKELELLRHLILKADLTNNQMQ LKILKKSQRTLLKELEMKELLKQQY MVQENGNSLFRKTKIYIRSYFSENS NGLKEITYYIINIHHFNNGQVSSWD MARRYNEFFELNTYLKKNFRDLMR QLQDLFPSKVKMSLKYHVTKTLLYE ERKQKLEKYLRELLSISEICEDNIFRR FLTDPTPFKLNKEYMHDDILEPLH EPIGSSNSTSNSSSVVDLQSSDGG ELNFYEDERHFFTDSGYPFYSONKS FVKQICDLFISLFALNKANAGWLRG RAIITVLQQLLGSTIEKYIKVSIQKLS EDQVFEAIVTFKNMLWGDNGLFR KRNETAEATRSEGERLRTEQLALTS LQRLFADTCGRVVGLRDSHEAAGR HAMLQNPYLNASLLLEALDAILLDII CND