

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P25846	MSH1_YEAST	Saccharomyces cerevisiae	DNA mismatch repair protein MSH1, mitochondrial	9.365641	NaN	NaN	33229814	MKHFFRLPTAFRPISRVSRLRYSSTDT AQPKISKLKISFNKISESNSEKKDNL GSIDTRNCLSTQDDKLSSTEPKAS LPPSLQYVRDLMDLYKDHVVLTMQ GSFYELYFEQAIRYAPELNISLTNRAY SHGKVPFAGFPVHQLSRHLKMLVN NCGYSVTIAEQFKKDVADNEANKF YRRVTRIVTPGTFIDEAFENLRENTY LLNIEFPENCMSQVADTSLKVGICW CDVSTGEIFVQVYLRDLVSAITRIQP KEILLDERLLEFHIESGTWYPELVEL KKFFIKYQKMPSQHRTIESFYGLFNL GGKEATERQLKIQFQFTQKELAALR NTLIYVSNHLPDFSINFQIPQRQLAT AIMQIDSRTSTALELHSTVRDNNKK GSLSSIRRTVTPSGTRLLSQWLSGP SLDLKEIKKRQKIVAFFKDNRDITET LRTMLKKVNDLSRILQKFSFGRGEA LELIQMARSLEVSREIRKYLLNNTSL MKATLKSQITQLTESLNFENLIDDI LKFLNEEELAKSQDAKQNAVTRM LDIDVKDKKESNKDEIFELRDFIVNP SFNTKLRKLHDTYQGVWQKTEYN ALLKGFFVGDGAKTFTLKERQNGE YALHVTGTASSLKKIDELISKSTEYH GSCFHILQSSQTRWLSHKIWTDLG HELELLNLKIRNEEANIIDLFKRKF DRSNEVRQVATTLGYLDTLSSFAVLA NERNLVCVKVDESNKLEVNGRHL MVEEGLSARSLETFTANNCELAKDN LWVITGPNMGGKSTFLRQNAIIVILA QIGCFVPCSKARVGIVDKLFSRVGSA DDLYNEMSTFMVEMIETSFILQGAT ERSLAILDEIGRGTSGKEGISIAYATL KYLLENQCRTLFATHFGQELKQIID NKCSKGMSEKVKFYQSGITDLGGN NFCYNHKLKPGICTKSDAIRVAELAG FPMEALKEAREILG