

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	extracell region
P25588	MRC1_YEAST	Saccharomyces cerevisiae	Mediator of replication checkpoint protein 1	24.475439	NaN	S144;S409;S411;S434;S605;S607;T609;S801;S807;S911	33229814	MDDALHALSSLTAKKRTTTYKKVAV PILDENDNTNGNPNDIDNPPELTG NGFLFANATLNRVKNRLEGKKAPEQ NHNNGKDRSENLSPTQLISNLYDGG EELEKSEVKDINSYSEKNSVSSFTQT QRIPVSIQQDKVFNVPVHNSVNDGKPT QLIKEDGLVNETSQALKTPLTTGRPG ATQRIDSSGATSQTOPIKSIIEPQSQH TTSSNHSNALS PKIPIITELIGTSPLF QSIQNRGPDQMDVPPQTAHDEDK TQAIGIPQATHQEQTQIDTVAQTLQ DEVPHTLKIREIQSELASEDSKREKA RNVEYKPKPKPIPTKFFSKESFLAD FDDSSSNEDDDIKLENAHPKPVOND DELHENKSVELNLTDETRINEKRVP LLSSYANNLKREIDSSKCITLDLSD SDEYGGDDMDSIKLSKDES VLPISQL SKATILNLKARLSKQNKLSQRPNK SKDPKVDHNVLLNLTLRKASRQILD HQKEVIETKGLKLEDMAKEKEIVEN LLEQEI LRNRIRQKEKRREKLEEN DFQLNAHDSGSDSGSESSGFALSGN EIADYESSGSENDNRRES DSEKEDD EILKQKSHHVKHIIINESDSTEVE AKPKEKADESLPKRIAINLGHYGDNI GEDTDKFQETNVLDTONIEEVMAER NTIENEVKDDVYVNEEAEAIRROLI DKEKLQKQKEKEHEAKIKELKRRG VTNFFEMEAESEDEWHGIGGADG EGSDDYDSDLEKMIDDYSKNNFNP HEIREMLAAENKEMDIKINKILYDI KNGGFRNKRAKNSLELESDDDDED DVLQYQYRLKRELMRKRRLIEIGDDA KLVKNPKSSAFFESMVEDIIEYKNPF GAEEYNLDITSTATDLDTQDINSINV GDNTGNNEQKPVQKNNKVIISED VQKSLSFLKSNYEDFETDKELSRIQ HGNDEAIEDLYTLKQNSSIKSFTNS QTDSTTSKTVNTIIDLEKRPEDDEV ENGDTSLVGVFKHPSIISFASRTDI NDKFKEGNKTVKILKSYKTVGSSKA SITYMGKTRKLIAPKRKTEGSHRYH HDHNNKMKMKTTKSNKLFESG QDSFDN	None	None	None	None	None	None	None	None	

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