	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q02866	MUK1_YEAST	Saccharomyces cerevisiae	Protein MUK1	24.475439	NaN	S67;S163;S185;S245	33229814	MARQLFTPPITNPRFDPNQSIRESYK NTTGGMQFQQNLHEDQNDNERSS CDGDENSTTGERLENNKSPILTKQEI DEALNTVTNLPPELSKLIDIFIDDLKQ PKYVRPLSVLQLSSLFQSFYIKFDKA SFQHVSSANNNGYYFSGGGSSSFLA AKETLSSGLSGIFGRSRSSSGNSLM RPRRSSSLFSNESISNSTNATQMLSP EEIKKQLKINELNNMKIEKYMELCE RDVFKKILIVGTSVSSPNKMKTFKPH QLQTFKVGNLFRNSVEFTEYNKLLN EKILCLSKLSTMNKINLIKFLSLNNG IDPEPKFEEIKDILYEFTYHSISPCEKI KALLKLHEIMTYSQEMSNDDYLSLLI YYIITIVPRDIFLNAEFIRLFRYKKKLV ETESFALTNLEAALVFVEGLTKNDFS NELQDKLTVNESKILENSISSRVSLP SKTAIMHKNNGNNGSNLGDIVTPTI QRPDVTRSNSYDGFRTVFDSSLKNII GKIRSYTPPHPNNTSNNNLHSSNNL NIPRSSSQLSMELSNRDTTEMSRDG SRSTSSSSRSSASLEHGNREFTGDL TVTASINGADKKEFQKSWKKYKGYK FEDLTICELRDLFEIYQKMMQ	None	None	None	None	None	None	None	None	None