

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				
P40002	MIT1_YEAST	Saccharomyces cerevisiae	Transcriptional regulator MIT1	24.475439	NaN	S152	33229814	MDIEPTFKGYIEDEDDALLILQATLD GCLKHIPRRPYEIERPYLIVSGSIFVFI EEISGIKRWTDGVSWSPPSRISGKFLI YKELDKENAGSNANATSSGSTDSAV ITDGTSGARNNPSSSKIKLPPLKNHQ FDLPPTMGHSSFESQDTSISPSNR SNLPLKYTGLVKKTISVKLRPPFNSI ENLHIVSYYSVKDIKQNCVTPKASP FLKDVRPSQELIVAMGNTTLGNVKN NSTTTGNGPNNINNKSNSSTPLNTV ISTNNNSANINAAGSNQFTSANKNY YYKNDESSGYPITQFAPALPSTILMY TANPPYITQSPDNTNATGMNTHVN NNNNNSNNSSNNSNNNNNNNNNN NNNNNNNNNNININNVNTNAGN GNNPNRFHNASFAYNTTGDFINPO QQGQISYPFYTTIPINPNYTTQPP NPVTNASTNENQGYSTSSTQHPYYG HPTESQSASAAAAGATGTPGTAENVL PVSSMQLLHQANNSASSATSTAP YPVYSMNVNVPYYNSASAYKRAQE NTTSNTNAEPGATSTNSGTMLSNP AYANSQYTPSQVYYQGFPOYAMASA QNPSMYQHQQHPLPTVYPIATPQQ NIMSSGHTLSTIGSDPQH HHYQQEP NDHKNFAMGHANNILNITNNDT MNNLNTNTSTTTQ	None	None	None	None	None	None	None	None	None	None	None	None	None

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