

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
P37304	PAM1_YEAST	Saccharomyces cerevisiae	Protein PAM1	24.475439	NaN	S659;S732;S767	33229814	<p>MTSALRVLVCGDHPNLILYTSRFQH  AKNIEFYLVNNSKNANYEVSSLFYG  TERFQIQNHFFQSLLDLVDLNNENG  GLVFDLIIMSASSLQEIPQVLRDIKP  MMNKTTKILFESSGFYILEPFIKASV  DLSLSNIFSIPTDYDIRRLDNGSYKQ  FTTANAKSFSVSIQQTTSVHENSYS  DIIPILNTFQKLFQKLFPRDVVTLYDH  SPSAFLAKEWELALPQICFDPLLILE  EKNPSTLDDHVLAKPLISGLLGESLL  IIKKMGIAMNPNFQNEQTIKHWK  NKCEDLPDGPALLYNFIHKASSLNID  LLLLQPILLADDFGVKTPYLECLFTM  MTQYQLLNKGDSEWFIRKIDENTALT  RVDDLQNSIALKDGKIMQLQNSEST  LKNEIKELQSQVLSLKQEVSSSKAN  NGQEILEILKKKVQMGDNSLFDPRPN  SNTNGISPSDNIVDVLNYESDQG  NNSSGNSRRQSFNNTSDTTLSTRD  ETSLKERELEVRMKELELQERELEL  QRKALQQQQYQQRPPKQVYSGPS  GTPTSGNNNKSYNPNRKSSYSQP  QHVAMMTSRGLHGPSAASSSPVISA  NNFVDPVSSGTPYSSNSSRFSQQIPS  QQYMHTVKPTSRRKNRSSVMPNIGYV  PGLTNNEYGRKFNNGMNGTQSRL  NSLSNQSTFRSQGPPITQOKSFQN  NGGSMRTNRIPSANYNISNQSGFV  NSISSPNLSNLENRNTVQNSRNADS  APCVNQLNSDSPPQLQSLSONGTSK  VPQINITQPSPIQTNFATSDNPAAVIK  LGTPESEDVSAATANNISTMGDES  RKEDVKEKKKKKFSFFGKRKK</p>	None	None	None	None	None	None	None	None	

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