

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
Q06673	ECM30_YEAST	Saccharomyces cerevisiae	Protein ECM30	24.475439	NaN	S635;S1065	33229814	MGNTDSKSSSILLNHCIALVRPEDA DASSPSRTSSPSPSLVDADPLSLNL SIFKLDSPDVEALFSDKPNVPLDTV FNDFYLDLDFISVDVQDFSNSSFKKIL HIHSSLNPPNFNNLIVFLSLYIILSAN SLPASRTGLHSSRLINAIKTLILPIY FDRVKSSTQDHYDVFWATQHEIEGL PLQNIPLGERLLLAIKLAFQDNFTT AVTAHPSELWEIGILTNSNKYRSLLN MHHQWHLFANRLLRLRLAALFSS DLYTSGGKQDINMFLVYWCTQMPK DKSIQFTSLLNCTMRFILNKKDF QSLKANFFSSDATASWOTLYFQFV QSCLVHVLNLSMSYKAQDNVITIFLT QLQREYDLKLLSSFIKIFKYPIDLAIE QESNIFNFTNNKHIDASRRRAVSTS SHDNSSSSHASLPSSSAAYHTKPQ TKPQLPEIHPLIPMTILMTNLIDCN KCFQNYFADKFASRFIIFSIYLYKYD YSSLSSSSTTRSNSSTTSNGTSNDT SNERSIVELNENSVSQILLPLLNHLL LILTSKKLVLFKMLQTFNLNYTNNL PNFYKLSNINGDINNLTFRDFTVIQL SNLILDNIKFNLPNPIFYELIYNLLP INDEILTSSHKNDDSHDDLILLSAKK KSASPSAATSSHTSSKLSYNAAMSL LYVLSKSSNKVYLTTYATPVFKTDIP YMISPGFKMDLLALLRSITIFFTYLF DDAENLLFAMVRHQSTHQINDSIN SISKALDMNPNLNSHIMTLKQMGF NRKVQWKDFYQFEEITDLPQVNLYS SANQQHQNQOQQGQNDNRGQONQ EDPGQENESPTPYLLFPASLENET PGTVKHFSSTNHDKNYQVIAFDKFS DSNLNLQHLEYWPHRPQWPTPLT FTHKCKNPKYENFNEVWSGTVYLOI LLRVIKQILSKVPEIPRIKSVQYFETLS KLSALRSDILTTIHPRLPLDVRRLTTF QPLSMHTNDKLLMWFHATWANIF TQTSFKYEETFSHELRQFESLLDISI DECEGNTISKPTDRLGYIRRSRGQS SVSLERTISAGSGVSTPTMALNRTKS NGSGNLMNYFFQNTAQNHFFQHLR SSSSSSSITLKTTSNSSIRTRPNH HVAPETNNNNSTNGNSNNSNGG FSFFKWKWGGNNSNGGSDDTKASQ RDPNVSTSIITDNLNSYMFEEIEISPG VVNNHIENNIWWGTDIRLFKIANFRK ESFSFLEMTSSFFKFKFINSNDNDN YNNNEFDDNTQLRYTSRGLYR	None	None	None	None	None	None	None	None	

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