

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P13586	ATC1_YEAST	Saccharomyces cerevisiae	Calcium-transporting ATPase 1	20.556082	NaN	S227	38179648	MSDNPFNASLLDEDSNREREILDAT AEALSKPSPSLEYCTLSVDEALEKLD TDKNGGLRSSNEANNRRSLYGPNEI TVEDDESLFKKFLSNFIEDRMILLI GSAVVSFLFMGNIDDAVSITLAIFIVVT VGFVQEYRSEKSLEALNKLVPAECH LMRCGQESHVLASTLVPGDLVHFRI GDRIPADIRIIEAIDLSIDESNLTGEN EPVHKTSQTIEKSSFNDQPNSIVPIS ERSCIAYMGTLVKEGHGKGIVVGTG TNTSFGAVFEMMNNIEKPKTPLQLT MDKLGKDSLVSFIVIGMICLVGIIQ GRSWLEMFQISVSLAVAAIPEGLPIIV TVTLALGVLRMAKRKAIVRRLPSVET LGSVNVICSDKTGTLTSNHMTVSKL WCLDSMSNKLNVLSLDKNKKTNS NGNLKNYLTEDVRETLTIGNLCNNA SFSQEHAIFLGNPTDVALLEQLANF EMPDIRNTVQKVQELPFNSKRKLMA TKILNPVDNKCTVYVKGAFERILEYS TSYLKSKGKKTEKLEAQKATINECA NSMASEGLRVFGFAKLTLSDSSTPL TEDLIKDLTFTGLIGMNDPPRPNVKF AIEQLLQGGVHIIMITGDSSENTAVNI AKQIGIPVIDPKLSVLSGDKLDEMSD DQLANVIDHVNIFARATPEHKLNIVR ALRKRGDVVAMTGDGVNDAPALKL SDIGVSMGRIGTDVAKEASDMVLTD DDFSTILTAIEEGKGIFNNIQNFLT QLSTVAALSLVALSTAFKLPNPLNA MQILWINILMDGPPAQSLGVEPVDH EVMKKPPRKRTDKILTHDVMKRLLT TAACIIVGTVYIFVKEMAEDGKVTAR DTTMTFTCFVFFDMFNALACRHNT KSIFEIGFFTnkMFNYAVGLSLLGQ MCAIYIPFFQSIFKTEKLGISDILLLLL ISSVFIVDELRLKWTRKKNEEDSTY FSNV