

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q12276	HER1_YEAST	Saccharomyces cerevisiae	HMG2-induced ER-remodeling protein 1	9.365641	NaN	S102;T128;S277;S1013;T1130;S1200;S1204;S1207	33229814	MSSKLYTDIDVPLDWLYKGKRRNR TKSAASTRTSEATTTSVKKTATLPST AAVPTKTIASPQRPLSGQNVNNE NSKPAVSAEKVSQOGQVPTRRTRSH SVSYGLLQKKNNDTDDSPKISRIR TAQDQPVKETKSSTLAEPVSKKGRS RSSSISTSLNERSKKSFLGSLFGRRP STTPSHVVERPLSSQNDHKKSTELP PIDTRQSKISTPTSTPTTASSKPSSSG GNRHSDGSLTSKLLSIPHNILETSST NFNAHHHIQSHHSSGREQDSPHSE SSDLPPILEKETTQKQLQKVSQVNLK RVTIAVQEFNSDPPQQLPSRKPGRG NVLIPEDMISAPPLISLGITNSSDQSS FQSNISPSYSKDSKEYKLALENFKKA AKEAEKHQKDAYVAERMAQEVAN YKARQLKTSPLTGATNSAADSATDQ ESSSLDARASKLHIDKPINVGHPFE THQDDNIKYSHLEQTLDVAYTRCC HLREILPIPSTLRQVKGKTAPLQTLKF LNPKPTLVDILSFCDFIAITPIHNIIFD NVSLTHDMFKIVICSLVTSPPVEKLG LRNVVINEQSWKLLCKFLLQNKTLI KLDISQTKARTDLNDSNYRDQMDW ELFCEVLRNREGRPLEELLNGLRF DKMSFSHFKNILLTFAQMNPKNPIR LGMANVEFSTECDFDFLNMSEYN VQGVDLAYNNLESLAKRMIKKLARL PYKHLEYFTLNSTNITSVDDMSYILK YLSRLPSIKFLDLSNLPQLFPGILTSG YKYFPQFPQLKRIHFDLDDLSIKETT MLVSILAKCETLSHVSLIGQSPMPDA SKISDSTDEPKSKDEKKEQIVFMR NTLWASLYAFVRDSHNLVSLDLDVYD QVPDEIQSRIALCLMHNMKRIMDSS FKLDELTVQDDLIFDGLITETAEEV LKRLNDKSLQNDVGGKYLKYYFE KMEKVHHNVQNTIDSMFEKRKSGE LPLQEKENLLRLLLEKNLSNILDIF ASMPNIADVVPFSKADNSFPNIGDS TVSANYNDGIRPSLKHLDSDRLIND VSIPENDSSIRPHLMATDSGRIIDVTT GKALLFKSSNTSLAGKRQEEEEGE LHKWGVFVQHQSSRHNSGLPSSAN SSRISGSLTPDSSVAGGKKGESSRTS GTRPKILPKIPTGAELRDAIHKAGIDS VDDLKINVTSEKVGLESYLGDELNS RSPNSDSLQESQKAPLQRPLVEDE TVTKKYDKLLNDSLNRHRSKT

