

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P06701	SIR3_YEAST	Saccharomyces cerevisiae	Regulatory protein SIR3	9.365641	NaN	NaN	33229814	MAKTLKDLDGWQVIITDDQGRVIDD NNRRRSRKRGGENVFLKRISDGLSF GKGESVIFNDNVTETYSVYLIHEIRL NTLNNVVEIWVFSYLRWFELKPKLY YEQFRPDLIKEDHPLEFYKDKFFNE VNKSELYLTAELSEIWLKDFIavgQI LPESQWNDSSIDKIEDRDFLVRYAC EPTAEKfVpidIFQIIRrvKEMEPKQS DEYLKRVSVPVSGQKTNRQVMHKM GVERSSKRLAKKPSMKKIKIEPSADD DVNNGNIPSQRGTSTTHGSISPQEE SVSPNISSASPSALTSPTDSSKILQKR SISKELIVSEEIPINSSEQESDYEPNN ETSVLSSKPGSKPEKTSTELVDGREN FVYANNPEVSDDGGLEETDEVSSE SSDEAIIPVnkRRGAHGSELSSKIRKI HIQETQEFskNYTTETDNEMNGNG KPGIPRGNTKIHSMNENPTPEKGNA KMIDFATLSKLKKYQIILDRFAPDN QVTDSSQLNKLTDQSSLDVAGLED KFRKACSSSGRETILSNFNADINLEE SIRESLQRELLKSQVEDFTRIFLPIY DSLMSSQNklFYITNADDSTKFQLV NDVMDELITSSARKELPIFDYIHIDAL ELAGMDALYEKIWFaSKENLCGDIS LEALNFYITNVPKAKKRKTLILIQNPE NLLSEKILQYFEKwISSKNSKLSIICV GGHNVTIREQINIMPSLKAHFTEIKL NKVDKNELQqMIITRLKSLKPFHV KVNDKKEMTIYNNIREGQNQKIPDN VIVINHKINNKITQLIAKNVANVSGS TEKAFKICEAAVEISKKDFVRKGGGLQ KGKLVVSQEMVPRYFSEAINGFKDE TISKKIIGMSLLMRTFLYTLAQETEG TNRHTLALETVLIKMVKMLRDNPGY KASKEIKVICGAWEPaITIEKLKQFS WISVVNDLVGEKLVVVVLEEPSASIM VELKLPLEINYAFSMDEEFKNMDCI