

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P11978	SIR4_YEAST	Saccharomyces cerevisiae	Regulatory protein SIR4	9.365641	NaN	S692	33229814	MPNDNKTPNRSSTPKFTKKPVT PNDKIPEREKSNVKT PKIPLFTFAKSKNYSR PSTAIHTSPHQPSDVKPT SHKQLQQPKSSPLKKNYNS FPHSNLEKISNSKLLS LLRSKTSAGRIESNNPSH DASRLASFEQTAFSRHAQ QQTSTFNSKPVRTIVPIST SQTNNNSFLSGVKSLL SEEKIRDYSKEILGINLANE QPVLEKPLKKSADIGASV ISLTKDKSIRKDTVEEK KEEKLNIGKNFAHSDSL SVPKVSAGDSGISPEESKAR SPGIKPNAIQTEVYGINEE STNERLEINQEKPVKLD ENSANSTVASALDTNGT SATTETLTSKKIVPSPK KVAIDQDKITLHDEKTL APSKHQIPITSEQMKED ADLKRMEILKSPHLSKSP ADRPQGRNRSRNFSTR DEETTKLAFVVEYEGQEN NYNSTSRTEKKNDMNTS AKNKNGENKKIGKRPPE IMSTEAHVNKVTEETTK QIQSVRIDGRKVLQKVQ GESHIDSRNNTLNVT PSKRPQLGEIPNPMKKH KPNEGRTPNISNGTINI QKKLEPKEIVRDILHT KESSNEAKKTIQNPLN KSQNTALPSTHKVTQK KDIKIGTNDLQVESAP KISSEIDRENVKSKDE PVS KAVESKSLNLN LFSNVLKAPFIKSES KPFSSDALSKKANFLE TIASTEKPENKTDK VSLSQPVSASKHEYS DNFPVLSLQPSKKS FANHTEDEQIEKKK ICRGRMNTIITH PGKMELVYVSDSD SSDNDSDLTDL ESLSSGESNEIKV TNDLDTSAEKDQI QAGKWFDPVLDWR KSDRELTKNIL WRIADKTTYDKET ITDLIEQGIPKHSY LSGNPLTSVTNDI CSVENYETSSAFF YQQVHKKDRLO YLPYAVSTFENTN NTEKNDVTNKNIN IGKHSQEQNSSS AKPSQIPTVSSPL GFEETKLSTTPTKS NRRVSHSDTNSS KPKNTKENLSKSS WRQEWLANLKLIS VSLVDEFPS ELSDSDRQIINEK MQLLKDIFANNL KSAISNNFRESDI IILKGEIEDYPM SSEIKIYYNELQNK PDAKKARFWSF MKTQRFVSNMGFDI QKSCEPVSIST SVKPHVVEPEHMAD AKIMPKDILQIT KKPLMVKNVKPSS PPDVKSLVQLST METKTLPEKKQFD SIFNSNKAKIIP GNGKHA SENISLSFSRPA SYGYFSVGKRV PIVEDRRVKQLDD ITDSNTTEILTS VDVLGTHSQTGTQ QSNMYTSTQKTE LEIDN

