

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P25302	SWI4_YEAST	Saccharomyces cerevisiae	Regulatory protein SWI4	9.365641	NaN	S255;S806	33229814	MPFDVLISNQKDNTNHQNITPISKS VLLAPHSNHPVIEIATYSETDVYECYI RGFETKIVMRRTKDDWINITQVFKIA QFSKTKRTKILEKESNDMQHEKVQG GYGRFQGTWIPLDSAKFLVNKYEIID PVVNSILTFQFDPNNPPPKRSKNSIL RKTSPGTKITSPSSYNKTPRKNSSS STSATTTAANKKGKKNASINQPNPS PLQNLVFQTPQQFQVNSSMNIMNN NDNHTTMNFNNDTRHNLINNISNN SNQSTIIQQQKSIHENSFNNNYSAT QKPLQFFPIPTNLQKNVALNPNPN NDSNSYSHNIDNVINSSNNNNNGN NNNLIIVPDGPMQSQQQQHHHEY LTNMFNHSMMDSITNGNSKRRRKK LNQSNEQQFYNQEQEKIQRHFKLMK QPLLWQSFQNPNDHHECYCDSNGS NNNNNTVASNGSSIEVFSSNENDN SMNMSSRSMTPFSAGNTSSQNKLE NKMTDQEYKQTILTILSSERSSDVDQ ALLATLYPAPKNFNINFEIDDQGHTP LHWATAMANIPLIKMLITLNANALQ CNKLGFNCTKISIFYNNCYKENAFD EIIKILKICLITPDVNGRLPFHYLIELS VNKSKNPMIISYMSIISLGLGQDY NLLKICLNYQDNIGNTPLHLSALNL NFEVYNRLVYLGASTDILNLDNESP ASIMNKFNTPAGGSNSRNNNTKAD RKLARNLPQKNYYQQQQQQQPQN NVKIPKIIKTQHPDKEDSTADVNIK TDSEVNESQYLHSNQPSTNMNTI MEDLSNINSFVTSSVIKDIKSTPSKIL ENSPILYRRRSQISDEKEKAKDNEN QVEKKKDPNLSVKTAMPSLESPESS LPIQMSPLGKYSKPLSQINKLNTKV SSLQRIMGEEIKNLDNEVVETESSIS NNKKRLITIAHQIEDAFDSVSNKTPI NSISDLQSRIKETSSKLNSEKQNFQI SLEKSQALKLATIVQDEESKVDMMNT NSSSHPEKQDEEPIPKSTSETSSPK NTKADAKFSNTVQESYDVNETLRLA TELTILQFKRRMTTLKISEAKSKINSS VKLDKYRNLIIGITENIDSKLDDIEKD LRANA