

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P09547	SWI1_YEAST	Saccharomyces cerevisiae	SWI/SNF chromatin-remodeling complex subunit SWI1	9.365641	NaN	NaN	33229814	MDFFFLNNNNNNNNTTTTTTTN NNNTNNNTNNNNNPANNTNNN NSTGHSSTNNNTNNNTNTGAS GVDDFQNFDPKPFQNLDSNNNN SNSNNNDNNNSNTVASSTNFTSPT AVVNNAAPANVTGGKAANFIQNQSP QFNSPYDSNNSNTNLNSLSPQAILA KNSIIDSSNLPLQAQQQLYGGNNNN NSTGIANDNVITPHFITNVQISQNS SSSTPNTNSNSTPNANQQFLPFNNS ASNNGNLTSNQLISNYAASNSMDR SSSASNEFVPNTSDNNNNNSNNHN MRNNSNNKTSNNNNVTAVPAATPA NTNNSTSNANTVFSERAAMFAALQ QKQQRFQALQQQQQQQNNQQQQ NQPQQQQQQQNPKFLQSQRQQ QORSILQSLNPALQEKISTELNNKQY ELFMKSLIENCKKRNMPLQSIPEIG NRKINLFYLYMLVQKFGGADQVTRT QQWSMVAQRLQISDYQQLESYFRIL LPYERHMISQEGIKETQAKRIFLQF LQELLKKVQQQQAAAALANANNNI NSASSAPTPAAPGASVPATAAPGTEA GIVPVSANTPKSLNSNININVNNNI GQQQVKKPRKQRVKKKTKKELELER KEREDFQKRQKLEDQQRQKLL LETCLRQYIEIKKLPKVYKRSIVR NYKPLINRLKHYNGYDINYISKIGEKI DSNKPIFLFAPELGAINLHALMSLQ SKNLGEINTALNTLLVTSADSNLKIS LVKYPELLDSLAILGMNLLSNLSQN VVPYHRNTSDYYYEDAGSNQYYVTQ HDKMVDKIFEKVN NATLTPNDSN DEKVTILVDSL TGNQLPTPTPEMEP DLDECFISMQSTSPAVKQWDLLEPE PIRFLPNQFPLKIHRTPYLTSKKIKD EIDDPFTKINTRGAEDPKVLINDQLS TISMILRNISFSDNNSRIMSRNFYK RFISDLLWLVLIHPENFTCNRKILNF KKDLVIVLSNISHLLEIASSIDCLLILI LVISFGQPKLNPMASSSSFGSESLTF NEFQLQWGKYQTFGVDILAKLFSLE KPNLNYFKSILLNKNTGNNLYDRNS NNNHKDKKLLRLLNLYNDNNKN NNNRHNLLNDVVSFLFSAIPLQQVL SQSADPSLLIDQFSPVISQSLTSILVIV QKILPLSNEVFEISENNSDSNSNNN GNKDSSFNFNKNL PFVWLSSEENIG SGLLKLSEILNINNSTSKNTLLQQQ

								NYSKVLLPSINISCVQLIKCLVEKSIC FENCLNNDPEILKKIASIPNLFPTDL EIFQLFTNPSVDIQIINQYQLLYNLKN DILTNLE
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