

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P18480	SNF5_YEAST	Saccharomyces cerevisiae	SWI/SNF chromatin-remodeling complex subunit SNF5	9.365641	NaN	S818	33229814	MNNQPQGTNSVPNSIGNIFSNIGTP SFNMAQIPQQLYQSLTPQQLQMIQQ RHQQLLRSLRQQQQQQQQTSPPP QTHQSPPPPPQQSQPIANQSATSTPP PPPAPHNLHPQIGQVPLAPAPINLPP QIAQLPLATQQQVLNKLROQAIKKN NPQVVNAITVAQQQVQRQIEQQKGGQ QTAQTQLEQQRQLLVQQQQQQQLR NQIQRQQQQQFRHHVQIQQQQQKQ QQQQQQHQQQQQQQQQQQQQQQQ QQQQQQQQQQQQQQQQQQQQQQQQ GQIPQSQQVPQVRMSGQPPTNVQP TIGQLPQLPKLNLPKYQTIQYDPPET KLPYPTYWSDKKADTDLLYEQIIQR DKINKYSLIRETNGYDPFSIYGFSNK EYISRLWHTLKYQDLKNTRMKSTS TSQKIPSASIWGNGYSGYGNGITNTT TRVIPQVEVGNRKHYLEDKLKVKQ AMNETSEQLVPIRLEFDQDRDRFFL RDTLLWNKNDKLIKIEDFVDDMLRD YRFEDATREQHIDTICQSIQEQIQEF QGNPYIELNQDRLGGDDLRIKLDI VVGQNQLIDQFEWDISNSDNCPEEF AESMCQELELPGEFVTAIAHSIREQV HMYHKSALLGYNFDGSAIEDDDIR SRMLPTITLDDVYRPAAESKIFTPNL LQISAAELERLDKDKDRDTRRKRQ GRSNRRGMLALSGETSASNTSMNGV HNTVAAGNASSLPPGEILLPDIADIP RTFRTPVPSTLMPGGVDVGPSVESY ELRNTTTYKSRPDRPKPVSPPCYIID HIPGHSLLSIKLPGKVNTKEEFAAA PNDTSSGTNAMLPSPELTKLNSN IRAGVTIPSIPNPIANHTVTNSPNPTL QPVIPGGAASKSVPTPSLPIAPPVAPH DSEATLLTNSNNGSSNNNTQNT

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