

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P22082	SNF2_YEAST	Saccharomyces cerevisiae	Transcription regulatory protein SNF2	9.365641	NaN	S358;T383;S716;S1340	33229814	MNIPQRQFSNEEVNRCYLRWQHLLR NEHGMNAPSVEFIYLTQVLFQFAAK QRQELQMQRQQGGISGSQQNIVPN SSDQAEPLPNNASSHISASASPHLAP NMQLNGNETFSTSAHQSPIMQTQM PLNSNGGNNMLPQRQSSVGSLSNAT NFSPTPANNGENAAEKPDNSNHNN LNLNNSLQPNRSLQEHNIQDSN VMPGSQINSPMPQQAQMQQAQFQA QQAQQAQQAQQAQQAQARLQQGRR LPMTMFTAQSELLKAQITSLKCLV NRKPIPFQAVIQKSINHPPDFKRM LLSLSEFARRRQPTDQNNQSNLNG GNNTQQPGTNSHYNNTNTDNVSGL TRNAPLDSKDEFASVSPAGPSSVH NAKNGTLDKNSQTVSGTPITQTESK KEENETISNVAKTAPNSNKTHTEQN NPPKPQKPVPLNVLQDQYKEGKVV DIDDPDMMVDSFTMPNISHSNIDYQ TLLANS DHAKFTIEPGVLPVGDTHT ATDIYQTLIALNLDTTVNDCLDKLLN DECTESTRENALYDYALQLLPLQKA VRGHVLQFEWHQNSLLTNTHPNFL SKIRNINVQDALLTNQLYKNHELLK LERKKTEAVARLKS MNKSAINQYNR RQDKKNRRLKFGHRLIATHTNLERD EQKRAEKKAKERLQALKANDEEAYI KLLDQTKDTRITHLLRQTNAFLDSL TAVKDDQKYTKEMIDSHIKEASEEV DDL SMVPKMKDEEYDDDDNSNV DYINVAHRIKEDIKKQPSILVGGTLK DYQIKGLQW MVSLFNNHLNGILAD EMGLGKTIQTISLLTYLYEMKNIRGP YLVIVPLSTLSNWSSEFAKWAPTLRT ISFKGSPNERKAKQAKIRAGEFDVVL TTFEYI IKERALLSKVKVWHMIIDEG HRMKN AQSKLSLTLNTHYHADYRLI LTGTPLQNNLPELWALLNFVLPKIF NSVKS FDEWFNTPFANTGGQDKIEL SEETLLVIRRLHKVLRPFLRLRLKK DVEKELPDKVEKVVCKMSALQQIM YQQMLKYRRLFIGDQNNK KMVGLR GFNNQIMQLKKICNHPVFEEVEDQ INPTRETNDDIWRVAGKFELDRILP KLKATGHRVLIFFQMTQIMDIMEDF LRYINIKYLRDLGHTKSDERSELLRL FNAPDSEYLCFILSTRAGGLGLNLQT ADTVIIFD TDWNP HQDLQAQDRAHR IGQKNEVRILRLITNSVEEVILERAY

<p>KKLDIDGKVIQAGKFDNKSTSEEQ ALLRSLDAEEERRKKRESGVVEEE ELKDSEINEILARNDEEMAVLTRMD EDRSKKEEELGVKSRLLEKSELPDIY SRDIGAELKREESESAAVYNRGRGAR ERKTATYNDNMSEEQWLRQFEVSD DEKNDKQARKQRTKKEDKSEIDG NGEIKGENIDADNDGPRINNISAED RADTDLAMNDDDFLSKKRKAGRPR GRPKKVLEGESENSEPPALESSPVT GDNPSSEDFMDIPKPRTAGKTSVKS ARTSTRGRGRGRGRGRGRGRGRGR PPKARNGLDYVRTPAAATSPIDIREK VAKQALDLYHFALNYENEAGRKLS IFLSKPSKALYDPYYMIKYPVAFDNI NTHIETLAYNSLKETLQDFHLIFSNA RIYNTEGSVVYEDSLELEKVVTKKYC EIMGDNSQLDFTEFDEQYGTRPLVL PPVVTSSVAESFTDEADSSMTEASV</p>
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