

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q06683	IRC3_YEAST	Saccharomyces cerevisiae	Putative ATP-dependent helicase IRC3	9.365641	NaN	NaN	33229814	MTSTLATRLSTYSISLILQRIKIIRKRCY SAPVLRDYQQDAIDACVNSIRQGTGR IGVSLATGGGKTVIFSNLINQLRQNY FKERQGNFKSLILVHRRELALQATAT LKKIFPDLKVHIEMGKYDCDIEDSDV IVASVQTLIRRLHKYDTNSVNLIIIDE AHHSVANSYRSILDHFKASTAETKIP VIGFSATFERADKRALSMVMDKIVY HRGILEMIDDKWLCEAKFTSVKIEA DLSDVKSTADDFQLAPLSSLMNTKE INEVILKTYLHKKQEKSLKSTLLFGV DKAHVQSLHKLFKDNGINTDYVTS TKQIERDNIIQFKNGETEVLNCGI FTEGTDMPNIDCILLCRPTKSRSLLI QMIGRGLRLHHSKDHCHIIDFIGASS VGVVSAPTLGIRSDDIEFDDATVED LKAIQGEIIAKQQKIDERLRALFQTDE AAMENVTERNSVADWIHSANSVDL TLCSEFDSFRNFTQSNNNSYPSGKEFD EASEAVKEMELLMNSQYPWVKFAS NAWGLPLKGKNHLRIYKEKSEDKLS MVYHLKMYRQLPCFITNKYADYVVK SIKDANLWNVMSKVEKIINTLNSDL EGQTMQYQAISSKYSKWRQTVPTSK QRDFVFRKLKKVYGESSKDFIRLSL DVTTYVNTKMTKGDASNLI FASSLAP VYPLKSLLRILEYQKRRSFIK

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