

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region				
P32831	NGR1_YEAST	Saccharomyces cerevisiae	Negative growth regulatory protein NGR1	24.475439	NaN	S524	33229814	MMSNVANASQRQENPYIPLPPSST VETSTEPRTLWMGDLDPSPFDEATI EEIWSKLDKKVIVKLIRAKKNLLIPCS STSSSNNTSEENAENQOSASNST DQLDNSQMININGISFIDPSTTQLHH AGYCFVEFETQKDAKFALSLNATPLP NFYSPTTNSQTNPTFKRTFRLNWAS GATLQSSIPSTPEFSLFVGDLSPTATE ADLLSLFQTRFKSVKTVRVMTDPLT GSSRCFGFVRFQDEDEERRRALIEMS GKWFQGRALRVAYATPRNNMMLQL QEQQQQQQQLQQQHQLDQEDNN GPLLIKANNLIQNNSNMLPLNALH NAPPMHLNEGGISNMRVNDLPSN TYNTDPTNTTVFVGGLVPKTTEFQL RSLFKPFGPILNVRIPNGKNCGFVKF EKRIDAEASIQGLQGFI VGGSPIRLS WGRPSSNAKTNSTIMGASQYMSS NGLRAPSAASSVDNSKQILEQYAED KRRLFLHQQQQQQQQDGNFS MEQMAHNNYNNYDYHRNKNG SHSDLVNLQRSNVPMQEDGALYP HQYSSPSYSLHPTGNQFSNATNLP QFGNAMSISMLPNGNSNKTASSM NTNPNTNMIMNSNMNMNVNP VPYGMGNGANMYDVSRMMTPPLNI APNSNNSKSSIMNKHPNRNNVPPI HPSLH	None	None	None	None	None	None	None	None	None	None	None	None	None

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