	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	P18494	GLN3_YEAST	Saccharomyces cerevisiae	Nitrogen regulatory protein GLN3	24.475439	NaN	S251;S267;S285;S469;S552;S562	33229814	MQDDPENSKLYDLLNSHLDVHGRS NEEPRQTGDSRSQSSGNTGENEEDI AFASGLNGGTFDSMLEALPDDLYFT DFVSPFTAAATTSVTTKTVKDTTPAT NHMDDDIAMFDSLATTQPIDIAASN QQNGEIAQLWDFNVDQFNMTPSNS SGSATISAPNSFTSDIPQYNHGSLGN SVSKSSLFPYNSSTSNSNINQPSINN NSNTNAQSHHSFNIYKLQNNNSSS SAMNITNNNNSNNSNIQHPFLKKS DSIGLSSSNTTNSVRKNSLIKPMSST SLANFKRAASVSSSISNMEPSGQNK KPLIQCFNCKTFKTPLWRRSPEGNT LCNACGLFQKLHGTMRPLSLKSDVI KKRISKKRAKQTDPNIAQNTPSAPAT ASTSVTTTNAKPIRSRKKSLQQNSLS RVIPEEIIRDNIGNTNNILNVNRGGY NFNSVPSPVLMNSQSYNSSNANFN GASNANLNSNNLMRHNSNTVTPNF RRSSRRSSTSSNTSSSSKSSSRSVVP ILPKPSPNSANSQQFNMNMNLMNT TNNVSAGNSVASSPRIISSANFNSNS PLQQNLLSNSFQRQGMNIPRRKMS RNASYSSSFMAASLQQLHEQQQVD VNSNTNTNSNRQNWNSSNSVSTNS RSSNFVSQKPNFDIFNTPVDSPSVSR PSSRKSHTSLLSQQLQNSESNSFIS NHKFNNRLSSDSTSPIKYEADVSAG GKISEDNSTKGSSKESSAIADELDWL KFGI	None	None	None	None	None	None	None	None	None