

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P20676	NUP1_YEAST	Saccharomyces cerevisiae	Nucleoporin NUP1	24.475439	NaN	S54;S161;T381;S383;S637	33229814	MSSNTSSVMSSPRVEKRSFSSTLKS FFTNPKNKRPPSSKVKVSSNLSYANH LEESDVEDTLHVNKRKRVSQHS DSLQNNNNAPIIHYGTENTERPPLL PILPIQRLRLREKQVRNMRELGLI QSTEFPSITSSVILGSQSKSDEGGSYL CTSSTPSPIKNGSCTRQLAGSGEDT NVGLPILKSLKNRSNRKRFRHSQSKG TVWSANFEYDLSEYDAIQKKNKDK EGNAGGDQKTSNRNNKSSISNGN LATGPNLTSEIEDLRADINSNRLSNP QKNLLKGPASTVAKTAPIQESFVFN SERSGTPTLKKNIEPKKDKESIVLPT VGFDFIKDNETPSKKTSPKATSSAGA VFKSSVEMGKTDKSTKTAEAPTLSF NFSQKANKTKAVDNTVPSTTLFNFG GKSDTVTSASQPFKFGKTSKSENH TESDAPPKSTAPIFSGKQEEENGDEG DDENEPKRRRLPVSEDTNTKPLFD FGKTGDQKETKKGSEKDSAGKPSF VFGASDKQAEGTPLFTFGKKADVTS NIDSSAQFTFGKAATAKETHTKPSET PATIVKKPTFTFGQSTSENKISEGSA KPTFSFSKSEERKSSPISNEAAKPS FSFPGKPVVQAPTDDKTLKPTFSFT EPAQKDSVVSEPKKPSFTFASKTS QPKPLFSFGKSDAAKEPPGSNTSFSF TKPPANETDKRTPPSFTFGGSTTN NTTTTSTKPSFSGAPESMKSTASTA AANTEKLSNGFSFTKFNHNKEKSN SPTSFDFGSASSTPIPVLGKPTDATG NTTSKSAFSFGTANTNGTNASANST SFSFNAPATGNGTTTTNTSGTNIAG TFNVGKPDQSIASGNTNGAGSAFGF SSSGTAATGAASNQSSFNFGNNGAG GLNPFTSATSSNANAGLNFNKPST NAQNVNVPFAFNFTGNNSTPGGGS VENMNGNTNANTVFAGSNNQPHQ SQTPSFNTNSSFTPSTVPNINFGSLN GGITNTATNALRPSDIFGANAASGSN SNVTNPSSIFGGAGGVPITTSFGQPQ SAPNQMGMTNNGMSMGGGVMA NRKIARMRHSKR	None	None	None	None	None	None	None		

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