	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	extrac region
0	P40477	NU159_YEAST	Saccharomyces cerevisiae	Nucleoporin NUP159	24.475439	NaN	S404;S657;S724;S735;S745;T803;S805;S819;S889;S940	33229814	MSSLKDEVPTETSEDFGFKFLGQKQ ILPSFNEKLPFASLQNLDISNSKSLF VAASGSKAVVGELQLLRDHITSDSTP LTFKWEKEIPDVIFVCFHGDQVLVST RNALYSLDLEELSEFRTVTSFEKPVF QLKNVNNTLVILNSVNDLSALDLRT KSTKQLAQNVTSFDVTNSQLAVLLK DRSFQSFAWRNGEMEKQFEFSLPS ELEELPVEEYSPLSVTILSPQDFLAVF GNVISETDDEVSYDQKMYIIKHIDGS ASFQETFDITPPFGQIVRFPYMYKVT LSGLIEPDANVNVLASSCSSEVSIWD SKQVIEPSQDSERAVLPISEETDKDT NPIGVAVDVVTSGTILEPCSGVDTIER LPLVYILNNEGSLQIVGLFHVAAIKS GHYSINLESLEHEKSLSPTSEKIPIAG QEQEEKKKNNESSKALSENPFTSAN TSGFTFLKTQPAAANSLQSQSSSTFG APSFGSSAFKIDLPSVSSTSTGVASS EQDATDPASAKPVFGKPAFGAIAKEP STSEYAFGKPSFGAPSFGSGKSSVES PASGSAFGKPSFGTPSFGSGNSSVEP PASGSAFGKPSFGTPSFGSGNSSAEP PASGSAFGKPSFGTSAFGTASSNET NSGSIFGKAAFGSSSFAPANNELFG SNFTISKPTVDSPKEVDSTSPFPSSG DQSEDESKSDVDSSSTPFGTKPNTS TKPKTNAFDFGSSSFGSGFSKALES VGSDTTFKFGTQASPFSSQLGNKSP FSSFTKDDTENGSLSKGSTSEINDD NEEHESNGPNVSGNDLTDSTVEQT SSTRLPETPSDEDGEVVEEEAQKSPI GKLTETIKKSANIDMAGLKNPVFGN HVKAKSESPFSAFATNITKPSSTTPA FSFGNSTMNKSNTSTVSPMEEADT KETSEKGPITLKSVENPFLPAKEERT GESSKKDHNDDPKDGYVSGSEISVR TSESAFDTTANEEIPKSQDVNNHEK SETDPKYSQHAVVDHDNKSKEMNE TSKNNERSGQPNHGVQGDGIALKK DNEKENFDSNMAIKQFEDHQSSEE DASEKDSRQSSEVKESDDNMSLNS DRDESISESYDKLEDINTDELPHGGE AFKAREVSASADFDVQTSLEDNYAE SGIQTDLSESSKENEVQTDAIPVKHN STQTVKKEAVDNGLQTEPVETCNFS VQTFEGDENYLAEQCKPKQLKEYYT SAKVSNIPFVSQNSTLRLIESTFQTV EAEFTVLMENIRNMDTFFTDQSSIP LVKRTVRSINNLYTWRIPEAEILLNIQ NNIKCEQMQITNANIQDLKEKVTDY VRKDIAQITEDVANAKEEYLFLMHF DDASSGYVKDLSTHQFRMQKTLRQ KLFDVSAKINHTEELLNILKLFTVKN KRLDDNPLVAKLAKESLARDGLLKEI KLLREQVSRLQLEEKGKKASSFDAS SSITKDMKGFKVVEVGLAMNTKKQI GDFFKNLNMAK	None	None	None	None	None	None	None	None	None