

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q9CAF4	NUP1_ARATH	Arabidopsis thaliana	Nuclear pore complex protein NUP1	27.144759	NaN	NaN	28154133	MASAARGESSNPYGGGLGTGGKFR KPTARRSQKTPYDRPTTSVRNAGLG GGDVRGGGWLSKLVDPQAORLITYSA QRLFGSLSRKRLGSGETPLQSPQQ KQLPERGVNQETKVGHKEDVSNLS MKNGLIRMEDTNASVDPKDGFTD LEKILQKKTFRSEVDRLTLLRSKA ADSSTMNEEQRNEVGMVVRHPPSH ERDRTHPDNGSMNTLVSTPPGSLRT LDECIA SPAQLAKAYMGSRPSEVTPS MLGLRQGAGREDSVFLNRTFPFQKS PTMSLVTKPSGQRPLENGFVTPRSR GRSAVYSMARTPYSRQSSVKIGSLF QASPSKWEESLPSGSRQGFQSGLKR RSSVLDNDIGSVGPVRRIRQKSNLSS RSLALPVSESPLSVRANGGEKTTHTS KDSAEDIPGSSFNLVPTKSSEMASKI LQQLDKLVSTREKSPSKLSPMLRG PALKSLQNV EAPKFLGNLPEKKANS PDSSYQKQEISRESVSREVLAQSEKT GDAVDGTSKTGSSKDQDMRGKGVY MPLTNSLEEHPKKRSFRMSAHED FLELDDDLGAASTPCEVAEKQNAFE VEKSHISMPIGEKLTPSEAMPSTSY ISNGDASQGT SNGSLETERNKFVAF PIEAVQQSNMASEPTS KFIQTEKSS ISSGKPTSEKRIPL EPKKPAAVFPN ISFSPPATGLLNQNSGASADIKLEKT SSTAFGVSEAWAKPTESKKTFSNSA SGAESSTSAAPTLNGSIF SAGANAVT PPPSNGSLTSSPFPSPSINIPSDNSV GDMPSTVQSFAATHNSSIFGKLP SNDSNSQSTSASPLSSTSPFKFGQPA APFSAPAVSESSGQISKETE VKNATF GNTSTFKFGGMASADQSTGIVFGAK SAENKSRPGFVFGSSSVVGGSTLNP STAAASAPESSGSLIFGVTSSSTPGTE TSKISASSAATNTGNSVFGTSSFAFT SSGSSMVGGVSASTGSSVFGFNAV SASATSSSQASNLFGAGNAQTGNT GSGTTTSTQSI PFQFGSSPAPS FGL SGNSSLASNSSPFGFSKSEPAVFTSV STPQLSSTNSSASSSTMSPLFGTS WQAPNSSPN SGPVFSSSFTTSTPTT FSGGSSAATVSTTTPIFGASTNNT PSPSPIFGFGSTPPTTPQQPVFGNSG TPSQSLFGNSTPGFAFGAPNNGNGI NNNQVSMEDSMAEDTDQANRAS MVAPMFGQAAVSMQPQNFAGGGA ATQPPSMANPFQGGQPMAS TLQN ASPFQASQSLFQGGGSFLGSTGG GDKSGRRIFKAKKSTRKK	None	None	None	None	None	None	None		