	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	P39923	NPR2_YEAST	Saccharomyces cerevisiae	Nitrogen permease regulator 2	24.475439	NaN	S362	33229814	MLSYFQGFVPIHTIFYSVFHPTEGSK IKYEFPPNNLKNHGINFNTFKNYIIP KPILCHKLITFKYGTYRIVCYPVTINS PIYARNFFSFNFVFVFPYDCETSPYE PAITRLGKMFKVLEEQNQLLSKSER DPVFFDLKVLENSTTTPSTAGPSSTP NPSSNTTPTHPTSEKDTKDMRSSRY SDLIKDLGLPQSAFSIQDLLMRIFQD LNNYSECLIPIDEGNAVDIKIFPLLRP PTTCVSLEDVPLSSVNLKKIIDVNWD PTMMSIVPYIDGLNSIAKISKLSNSD PGLVIECIRHLIYYKCVTLSDIFQFSNI YAPSSLIRNFLTDPLMASDCQSYVTF PEVSKISNLPLNKSLGSGDQDSPSFS VRRKSKSSSIPSNPDSRTTSFSSTSR VSQNSSLNSSFSSIYKDWRQSQTSC SSSNIHVINNRNRFLPTRSCLFDLYR SLSQGQTLKTWYESKYMILKENNIDI RRFITFGLEKRIIYRCYSFPVMINAGS REPKEMTPIITKDLVNNDKLLEKRN HNHLLSATGSRNTAQSGNLKPERPS KVSFEMQRVSSLATGKSTMPKLSDE EEGILEESIRNAETFDKICVLLSKPKL EVESYLNELGEFKVINS	None	None	None	None	None	None	None	None	None