	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	P33420	NIP80_YEAST	Saccharomyces cerevisiae	Protein NIP100	24.475439	NaN	NaN	33229814	MRNAGVQVDTNMQKISLQDTVLVN EMKGRVKFIGETQFAKGIWYGIELD KPLGKNDGSANGIRYFDIDLKKANS NGGYYGLFCKKDTLQFYKPDDDEHS LLNGNAAQETIKNLQVKCESLASKL NKIKIENHELKTSVEKLSTNETVLLS KISRLDKLVKELKVENGNMKTHLDN FNHLLDASDSVMAPDLDKGTLLERS HLLQGLLDQTKLSYDKAMKVQEDLL EENTQLLEENAVLSKKISDLGLQLQ QTNNTIGDLALQIEAQSKSSNIVDKL TNDNILLTSNIKALNNELEELQAKEK LDENLRITYEQLEQELRLQLSNLQSA LENEKEIAGTYIEENSRLKATLESIEA KTSHKFQSLELKVNTLQEELYQNKL LKKFYQIYEPFAQPHLAALSSQLQYL AEVIESENFGKLENIEIHIILKVLSSIS YALHIYTIKNTPDHLETTLQCFKVNI APISMWLSEFLQRKFSSKQETAFSIC QFLEDNKFLDKDVTLILKILHPILETT VPKLLAFLRTNSNFNDNDTLCLIGS LYERSLSLIARIDKLIGKEEISKQDNR LFLYPSCDITLSSILTILFSDALFLRQ DYKRISSLKKLEVFFQGIESLLENITI FPEQPSQQTSDSESQCNIKEGNFSN SLLSDRLNEENIRLKEVLVQKENML TELETKIKIIIGRDLERKTLEENIKTLK VELNNKNEENCGKTEILNKLKEENF NLVNRLKNMELKLYQIKDNNTLNKI YLDREKVDRVNLVSEIMELRETIRRQ IKEQKRVSIDFSWLDELPAVENKQPF KEHINHSLDTLGIEMFNFVSTSRILD LKLDQPLAEDELWHERDHSYISYLK RKRKNIRLKSQNVVTYYK	None	None	None	None	None	None	None	None	None