	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	P40473	POG1_YEAST	Saccharomyces cerevisiae	Transcriptional activator POG1	24.475439	NaN	S152;S168;S314	33229814	MKQEPHRQSEEKEKPKGPMAVERE QHTSLSSGTTVTASTGDESTNSRPV ESSQTEKSLSLRIRILKQLGFDDIQEL NACDTGLVEQFLNVRLINDTKELEKI RESNLAKLNQIIDKCMESDKISDSTL NKILDMSMNRDTNNDNNNHLTIPS PITTKKRKINASELASPRGHRRYRSD IPTVSEVETGVGYPQIHQQPGAYTLP MPANQWMSNPYMQPPQPQVQQIM PQYLYPPGMGPQAQLPTMSSNSESQ TPVMSSQFLSLNQHGLYQQNIGAHP VMSMGPQANIYGQQHQLQPGQERD QSRKSFSHRRSQSANISMANFRSP MRNPQPASSQRPVNFLIHTPKHPPP T	None	None	None	None	None	None	None	None	None