	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	Q03674	PLB2_YEAST	Saccharomyces cerevisiae	Lysophospholipase 2	24.475439	NaN	NaN	33229814	MQLRNILQASSLISGLSLAADSSSTT GDGYAPSIIPCPSDDTSLVRNASGLS TAETDWLKKRDAYTKEALHSFLSRA TSNFSDTSLLSTLFSSNSSNVPKIGIA CSGGGYRAMLGGAGMIAAMDNRTD GANEHGLGGLLQSSTYLSGLSGGN WLTGTLAWNNWTSVQEIVDHMSES DSIWNITKSIVNPGGSNLTYTIERWE SIVQEVQAKSDAGFNISLSDLWARAL SYNFFPSLPDAGSALTWSSLRDVDV FKNGEMPLPITVADGRYPGTTVINLN ATLFEFTPFEMGSWDPSLNAFTDVK YLGTNVTNGKPVNKDQCVSGYDNA GFVIATSASLFNEFSLEASTSTYYKMI NSFANKYVNNLSQDDDDIAIYAANP FKDTEFVDRNYTSSIVDADDLFLVD GGEDGQNLPLVPLIKKERDLDVVFA LDISDNTDESWPSGVCMTNTYERQY SKQGKGMAFPYVPDVNTFLNLGLTN KPTFFGCDAKNLTDLEYIPPLVVYIP NTKHSFNGNQSTLKMNYNVTERLG MIRNGFEAATMGNFTDDSNFLGCIG CAIIRRKQESLNATLPPECTKCFADY CWNGTLSTSANPELSGNSTYQSGAI ASAISEATDGIPITALLGSSTSGNTTS NSTTSTSSNVTSNSNSSSNTTLNSN SSSSSISSSTARSSSSTANKANAAAIS YANTNTLMSLLGAITALFGLI	None	None	None	None	None	None	None	None	None