	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	P27514	PHO91_YEAST	Saccharomyces cerevisiae	Low-affinity phosphate transporter PHO91	24.475439	NaN	S295;S311;S312	33229814	MKFSHSLQFNSVPEWSTKYLAYSQL KKLIYSLQKDKLYSNNKHHVVEPHD ANDENLPLLADASPDDQFYISKFVAA LNQELKKIDKFYISQETGLIANYNEL KDDVMELENTNKATQLFNQQQQH QLQSVARNRKSKSQQRQRRFSSVSS TDSNPSLTDMSIDSAPVIHTQVSNTT NNGNSMQNLASASVSLSNSNPVYL SPFTQHRLSLKKRLISIYTQLSELKDF IELNQTGFSKICKKFDKSLNTNLKQ NYLNYIKFHSHVFNPATINRIQHHIT ETILTYASLNKGTRRPSNTFNLDADR INNDENSSGNEEDEDGNRQEVLDF QDAERELSSHLRDHVVWERNTVWK DMMNLERKYQSAKTDNKKFSKLSS SQLRPNANITESMAMSSGGAGIIAPS TDSLTFRELMHLPPKQWLQFIMGQT SLLKFLLITSCFIALLTFNLTPFTQDS LQKNCFAILIYASLLWATETIPLFVTS LMIPLLIVVFPVIKDPITSQPMSPRDS SQFILSTMWSSVIMLLLGGFTLAAAL SKYNIAKVLSTHILASAGTNPHFILLT NMFVALFVSMWVSNVAAPVLCYSIV QPLLRTLPRNCSYAKALILGIALASNI GGMSSPIASPQNIFSIGIMDPSPSWA EWFMIALPVCFICVMAIWVLLIITFPP EPNVKILQLHPSRDPFTLKQWFVTL VCIITIVLWCLSNQISGIFGEMGIISII PIVVFFGTGLLTSDDFNNFMWTIVV LAMGGTTLGKAVSSSGLLSTMAQLI KAQVEHEPIFIIVLIFGLVILVMATFV SHTVAAMIIVPLMSEIGSNLPSGDHS RLLIVIAALLCSSAMGLPTSGFPNVT AISMIDEVGDRYLTVGTFITRGVPAS LLSYAAIVTVGYGILKVMGF	None	None	None	None	None	None	None	None	None