	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	Q12051	GGPPS_YEAST	Saccharomyces cerevisiae	Geranylgeranyl pyrophosphate synthase BTS1	24.475439	NaN	NaN	33229814	MEAKIDELINNDPVWSSQNESLISK PYNHILLKPGKNFRLNLIVQINRVM NLPKDQLAIVSQIVELLHNSSLLIDDI EDNAPLRRGQTTSHLIFGVPSTINTA NYMYFRAMQLVSQLTTKEPLYHNLI TIFNEELINLHRGQGLDIYWRDFLPE IIPTQEMYLNMVMNKTGGLFRLTLR LMEALSPSSHHGHSLVPFINLLGIIY QIRDDYLNLKDFQMSSEKGFAEDIT EGKLSFPIVHALNFTKTKGQTEQHN EILRILLLRTSDKDIKLKLIQILEFDTN SLAYTKNFINQLVNMIKNDNENKYL PDLASHSDTATNLHDELLYIIDHLSE L	None	None	None	None	None	None	None	None	None