	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	P08468	PT111_YEAST	Saccharomyces cerevisiae	Protein PET111, mitochondrial	24.475439	NaN	NaN	33229814	MLQRRFISSSGIKRLLHRESNKVMH TVFFKVRYYSTELIKKKHKEDIEDWV KAQLKDSSTISGVYESRNKLDWMDS ITKSPSSLDILKNQYNIVKDKDFGIL WKQKFESADPDILMTIISLSTNQKVL FSIQQLLILINSLHFLKRDYDIGQIYT TYEQFTPLLASHTDKGTYGQFIEIML VVQHNLHHFDVCETLFAEYIKYCKV KPQMISLGLNSFIRSNNTQLAVEFYT QAITNPDTFPITEKQLFEFLRCMERY LDMSSMKHIFYLWLKVKCGDEQSS STNLPSFKTLAIIHRMLLRFSNTDEL NDFLTNPVVLSTGYTSSVQFELIEFC HSLYCIKGDRTKSIDDSILMERVDKF ITRLNNNISTRKELYMSVVQAYVSTN NFENLKVILEKIQRDNDISIDGSFHL CISRYFVNTNQFEGLFKYYRSVVKTT DGKTRLRPAFIQQLWSCAVNVYPML AKEITNDLLVTLKRSQYSKCLTWVYT FLQENAHIHTRKINGGEDSSLSGFN AVDFERFEEFKKKVSHNDVYGAELV ISNSLKEGIAPQFSFLYSVLALCLRN SLTGLARVVDVILRTRFRYIPLKVDIL WLKWEIISNYRSFEKLSAEHLKELEF KLKEFERVHQKELSVQNYLQLTQIC FHTRDFKYACYLISQARKNLDTSNN KQWMMYYMTSLKLASRMHESERFS RILKDWNCNHRASLITPGCIRQIKGF MKYFEKRPAYISTAASIDNKEIKDRID ELVLRYVDYKYQGLENMRKLTLFLK EWFDEEISLLKLEQNERKMKLFEEN KKEEE	None	None	None	None	None	None	None	None	None