	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	Q28851	ATPK_BOVIN	Bos taurus	ATP synthase subunit f, mitochondrial	27.604395	NaN	S3	28100784	MASVVPLKEKKLLEVKLGELPSWIL MRDFTPSGIAGAFQRGYYRYYNKYV NVKKGSIAGLSMVLAAYVFLNYCRS YKELKHERLRKYH	None	None	None	None	None	None	None	None	None