	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	extrace region
0	P56134	ATPK_HUMAN	Homo sapiens	ATP synthase subunit f, mitochondrial	23.324282	NaN	S3	35083852;34725712;35138101	MASVGECPAPVPVKDKKLLEVKLGE LPSWILMRDFSPSGIFGAFQRGYYRY YNKYINVKKGSISGITMVLACYVLFS YSFSYKHLKHERLRKYH	True	False	2.375	4.347	5.0	1.074	0.281	1.057	1.326