	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	D3ZAF6	ATPK_RAT	Rattus norvegicus	ATP synthase subunit f, mitochondrial	23.439218	NaN	S3	38843836	MASIVPLKEKKLMEVKLRELPSWIL MRDFTPSGIAGAFRRGYDRYYNKYI NVRKGSISGINMVLAAYVVFSYCISY KELKHERRRKYH	True	False	2.345	1.785	5.0	1.074	0.281	0.738	0.597