	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	P13620	ATP5H_BOVIN	Bos taurus	ATP synthase subunit d, mitochondrial	27.604395	NaN	NaN	28100784	MAGRKLALKTIDWVAFGEIIPRNQK AVANSLKSWNETLTSRLATLPEKPP AIDWAYYKANVAKAGLVDDFEKKFN ALKVPIPEDKYTAQVDAEEKEDVKSC AEFLTQSKTRIQEYEKELEKMRNIIP FDQMTIEDLNEVFPETKLDKKKYPY WPHRPIETL	None	None	None	None	None	None	None	None	None