	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	P05631	ATPG_BOVIN	Bos taurus	ATP synthase subunit gamma, mitochondrial	27.604395	NaN	S146	28100784	MFSRAGVAGLSAWTVQPQWIQVRN MATLKDITRRLKSIKNIQKITKSMKM VAAAKYARAERELKPARVYGVGSLAL YEKADIKTPEDKKKHLIIGVSSDRGL CGAIHSSVAKQMKSEAANLAAAGKE VKIIGVGDKIRSILHRTHSDQFLVTFK EVGRRPPTFGDASVIALELLNSGYEF DEGSIIFNRFRSVISYKTEEKPIFSLD TISSAESMSIYDDIDADVLRNYQEYS LANIIYYSLKESTTSEQSARMTAMDN ASKNASEMIDKLTLTFNRTRQAVITK ELIEIISGAAALD	None	None	None	None	None	None	None	None	None