	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	Q4R5B0	ATPG_MACFA	Macaca fascicularis	ATP synthase F(1) complex subunit gamma, mitochondrial	39.03173	NaN	S146	41018975	MFSRAGVAGLSAWTLQPQWIQVRN MATLKDITRRLKSIKNIQKITKSMKM VAAAKYARAERELKPARIYGLGSLAL YEKADIKVPEDKKKHLLIGVSSDRGL CGAIHSSIAKQMKSEVATLTAAGKEV MLVGIGDKIRGILYRTHSDQFLVAFK EVGRKPPTFGDASVIALELLNSGYEF DEGSVIFNRFRSVISYKTEEKPIFSLN TIASAESMSIYDDIDADVLQNYQEYN LANIIYYSLKESSTSEQSARMTAMDN ASKNASEMIDKLTLTFNRTRQAVITK ELIEIISGAAALD	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None