

	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
0	Q91VR2	ATPG_MOUSE	Mus musculus	ATP synthase subunit gamma, mitochondrial	19.496418	NaN	S146	22645316;33300544	MFSRASVVGLSACAVQPQWIQVRN MATLKDITRRLKSIKNIQKITKSMKM VAAAKYARAERELKPARVYGTGSLAL YEKADIKAPEDKKKHLIIGVSSDRGL CGAIHSSVAKQMKNEVAALTAAGKE VMIVGVGEKIKGILYRTHSDQFLVSF KDVGRKPPTFGDASVIALELLNSGYE FDEGSIIFNQFKSVISYKTEEKPIFSL NTIATAETMSIYDDIDADVLQNYQEY NLANLIYYSLKESTTSEQSARMTAM DNASKNASDMIDKLTLTFNRTRQAV ITKELIEIISGAAALD