

	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	26.056242	NaN	S146	33300544;37453647;36288343;22645316	MFSRASVVGLSACAVQPQWIVRN MATLKDITRRLKSIKNIQKITKSMKM VAAAKYARAERELKPARVYGTGSLAL YEKADIKAPEDKKKHLIIGVSSDRGL CGAIHSSVAKQMKNEVAALTAAGKE VMIVGVGEKIKGILYRTHSDQFLVSF KDVGRKPPTFGDASVIALELLNSGYE FDEGSIIFNQFKSVISYKTEEKPIFSL NTIATAETMSIYDDIDADVLQNYQEY NLANLIYYSLKESTTSEQSARMTAM DNASKNASDMIDKLTTLTFNRTRQAV ITKELIEIISGAAALD