	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	P00829	ATPB_BOVIN	Bos taurus	ATP synthase subunit beta, mitochondrial	27.604395	NaN	T312;S415;S433	28100784	MLGLVGRVVAASASGALRGLSPSAP LPQAQLLLRAAPAALQPARDYAAQA SPSPKAGATTGRIVAVIGAVVDVQFD EGLPPILNALEVQGRETRLVLEVAQH LGESTVRTIAMDGTEGLVRGQKVLD SGAPIRIPVGPETLGRIMNVIGEPIDE RGPIKTKQFAAIHAEAPEFVEMSVEQ EILVTGIKVVDLLAPYAKGGKIGLFGG AGVGKTVLIMELINNVAKAHGGYSV FAGVGERTREGNDLYHEMIESGVIN LKDATSKVALVYGQMNEPPGARARV ALTGLTVAEYFRDQEGQDVLLFIDNI FRFTQAGSEVSALLGRIPSAVGYQPT LATDMGTMQERITTTKKGSITSVQAI YVPADDLTDPAPATTFAHLDATTVLS RAIAELGIYPAVDPLDSTSRIMDPNIV GSEHYDVARGVQKILQDYKSLQDIIA ILGMDELSEEDKLTVSRARKIQRFLS QPFQVAEVFTGHLGKLVPLKETIKGF QQILAGEYDHLPEQAFYMVGPIEEA VAKADKLAEEHS	None	None	None	None	None	None	None	None	None