	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	P13619	AT5F1_BOVIN	Bos taurus	ATP synthase F(0) complex subunit B1, mitochondrial	27.604395	NaN	NaN	28100784	MLSRVVLSAAAAAAPSLKNAALLGP GVLQATRIFHTGQPSLAPVPPLPEHG GKVRFGLIPEEFFQFLYPKTGVTGPY VLGTGLILYLLSKEIYVITPETFSAISTI GFLVYIVKKYGASVGEFADKLNEQKI AQLEEVKQASIKQIQDAIDMEKSQQ ALVQKRHYLFDVQRNNIAMALEVTY RERLHRVYREVKNRLDYHISVQNM MRQKEQEHMINWVEKRVVQSISAQ QEKETIAKCIADLKLLSKKAQAQPVM	None	None	None	None	None	None	None	None	None