

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	ex re
0 Q9CQQ7	AT5F1_MOUSE	Mus musculus	ATP synthase F(0) complex subunit B1, mitochondrial	24.348384	NaN	NaN	37453647;33300544;37507081;36288343	MLSRVLSAAATAAPCLKNAAALGP GVLQATRAFHTGQPR LAPLPPEYG GKVR LGLIPEEFFQFLYPKTGVTG PY VLGTGLSLYFLSKEIYVITPETFSTISV VGLIVYVIKYGASFGFIDKLNEEKI AQL EEVKQSSMKIQDAIDMEKAQQ ALVQKRHYLFDVQRN NIALALEVTY RERLHKAYKEVKNRLDYHISVQNM MRK EEEHMIDWVEKHVVK SISVQ QEKETIAKCIEDLKLLAKKAQAQ PIM	True	False	1.581	1.541	5.0	1.217	0.829	1.302	1..