

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	extrac region
P50516	VATA_MOUSE	Mus musculus	V-type proton ATPase catalytic subunit A	35.850493	T257;S262	S384	33300544;40885482;37507081;35822049;37453647;28135057;39627609	MDFSKLPKIRDEDKSTFGYVHGVS GPVVTACDMAGAAMYELVRVGHSE LVGEIIRLEGDMATIQQVYEETSGVSV GDPVLRRTGKPLSVELGPGIMGAIFDG IQRPLSDISSQTQSIYIPRGVNVSALS RDIKWEFIPSKNLRVGSHTGGDIYGI VNENSLIKHKIMLPPRNRGVSVTYIAP PGNYDASDVVLELEFEGVKEKFSMV QVWPVROVRPVTEKLPANHPLLTGO RVLDALFPCVQGGTTAIPGAFGCGKT VISQSLSKYSNSDVIIYVCGGERGNE MSEVLRDFPELTMEVDGKVESIMKR TALVANTSMPVAAREASIYTGITLS EYFRDMGYHVSMMADSTRWAEAL REISGRLAEMPADSGYPAYLGARLAS FYERAGRVKCLGNPEREGSVIVGA VSPGGDFSDPVTSATLGIVQVFWG LDKKLAQRKHFPVSNWLISYSKYMR ALDEYYDKHFEFVPLRTRKAKEILQE EEDLAEIVQLVGKASLAETDKITLEV AKLIKDDFLQQNGYTPYDRFCPFYKT VGMLSNMISFYDMARRAVETTAQS DNKITWSIIREHMGEILYKLSMKFK DPVKDGEAKIKADYAQLLEDMQLNAF RSLED	True	True	5.0	3.392	4.359	1.452	1.523	5.0	1.646