	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasm memb
0	P13637	AT1A3_HUMAN	Homo sapiens	Sodium/potassium-transporting ATPase subunit alpha-3	22.133582	NaN	S37;S56;S218;S265;S442;T548;S933	34019948;30397120	MGDKKDDKDSPKKNKGKERRDLDD LKKEVAMTEHKMSVEEVCRKYNTD CVQGLTHSKAQEILARDGPNALTPP PTTPEWVKFCRQLFGGFSILLWIGAI LCFLAYGIQAGTEDDPSGDNLYLGIV LAAVVIITGCFSYYQEAKSSKIMESFK NMVPQQALVIREGEKMQVNAEEVV VGDLVEIKGGDRVPADLRIISAHGCK VDNSSLTGESEPQTRSPDCTHDNPL ETRNITFFSTNCVEGTARGVVVATG DRTVMGRIATLASGLEVGKTPIAIEIE HFIQLITGVAVFLGVSFFILSLILGYT WLEAVIFLIGIIVANVPEGLLATVTVC LTLTAKRMARKNCLVKNLEAVETLG STSTICSDKTGTLTQNRMTVAHMW FDNQIHEADTTEDQSGTSFDKSSHT WVALSHIAGLCNRAVFKGGQDNIPV LKRDVAGDASESALLKCIELSSGSVK LMRERNKKVAEIPFNSTNKYQLSIH ETEDPNDNRYLLVMKGAPERILDRC STILLQGKEQPLDEEMKEAFQNAYL ELGGLGERVLGFCHYYLPEEQFPKG FAFDCDDVNFTTDNLCFVGLMSMI DPPRAAVPDAVGKCRSAGIKVIMVTG DHPITAKAIAKGVGIISEGNETVEDIA ARLNIPVSQVNPRDAKACVIHGTDL KDFTSEQIDEILQNHTEIVFARTSPQ QKLIIVEGCQRQGAIVAVTGDGVNDS PALKKADIGVAMGIAGSDVSKQAAD MILLDDNFASIVTGVEEGRLIFDNLK KSIAYTLTSNIPEITPFLLFIMANIPLP LGTITILCIDLGTDMVPAISLAYEAAE SDIMKRQPRNPRTDKLVNERLISMA YGQIGMIQALGGFFSYFVILAENGFL PGNLVGIRLNWDDRTVNDLEDSYG QQWTYEQRKVVEFTCHTAFFVSIVV VQWADLIICKTRRNSVFQQGMKNKI LIFGLFEETALAAFLSYCPGMDVALR MYPLKPSWWFCAFPYSFLIFVYDEIR KLILRRNPGGWVEKETYY	None	None	None	None	None	None	None	None