

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P05023	AT1A1_HUMAN	Homo sapiens	Sodium/potassium-transporting ATPase subunit alpha-1	19.017347	S65	T10;S16;S40;S47;S228;T260;S452;S484;T542;S668;S675;S943	35083852;32119511;34725712;37217939;30397120;26853435;33214551;29351928	MGKGVGRDKYEPAAVSEQGDKKGGK KGKKDRDMDELKKEVSMDDHKLSL DELHRKYGTDLSRGLTSARAAEILAR DGNALTPPPPTTPEWIKFCRQLFGG FSMLLWIGAILCFLAYSIAATEEPEP QNDNLYLGVVLSAVVIITGCFSYQE AKSSKIMESFKNMVPQQALVIRNGE KMSINAEVVVGDLEVKGGDRIPA DLRIISANGCKVDNSSLTGESEPOTR SPDFTNENPLETRNIAFFSTNCVEG TARGIVVYTGDRVTMGRIATLASGLE GGQTPIAAEIEHFIHIITGVAVFLGVS FFILSLILEYTWLEAVIFLIGIIVANVP EGLLATVTVCLTLTAKRMARKNCLV KNLEAVETLGSTSTICSDKTGTLTQN RMTVAHMWFDNQHIEADTTENQS GVSFDKTSATWLALSRIAGLCNRAV FQANQENLPILKRAVAGDASESALL KCIELCCGSVKEMRERYAKIVEIPFN STNKYQLSIHKNPNTSEPOHLLVMK GAPERILDRCSSILLHGKEQPLDEEL KDAFQNAVLELGGGERVLGFCHLF LPDEQFPEGFQFDTDDVNFIDNLC FVGLISMIDPPRAAVPDAVGKCRSAG IKVIMVTGDHPITAKAIKGVGIISEG NETVEDIAARLNIPVSQVNPRAKAC VVHGSDLKDMTSEQLDDILKYHTEI VFARTSPQQKLIIVEGCQRQGAIVAV TGDGVNDSPALKKADIGVAMGIAGS DVSKQAADMILLDDNFASIVTGVEE GRLIFDNLKKSIAYTLSNIPEITPFLI FIANIPLPLGTVTILCIDLGTDMVPAI SLAYEQAESDIMKRQPRNPKTDKLV NERLISMAYGQIGMIQALGGFFTYFV ILAENGFPIHLLGLRVDWDDRWIN DVEDSYGQWQWYEQRKIVEFTCHTA FFVSIVVVQWADLVICKTRRNSVFQ QGMKNKILIFGLFEETALAAFLSYCP GMGVALRMYPLKPTWWFCAPYSL LIFVYDEVKLIIRRRPGGWVEKETY Y