

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence
Q8VDN2	AT1A1_MOUSE	Mus musculus	Sodium/potassium-transporting ATPase subunit alpha-1	25.365296	NaN	T10;S16;S40;S47;S228;T260;S452;S484;T542;S668;S675;S943	24788674;22645316;37507081;33300544;36288343;37453647	MGKGVGRDKYEPAAVSEHGDKKGGK KAKKERDMDELKKEVSMDDHKLSL DELHRKYGTDLRSGLTPARAAEILAR DGNALTPPPTTPEWVKFCRQLFGG FSMLLWIGAILCFLAYGIRSATEEEP PNDDLVLGVLSAVVIITGCFSYQEQE AKSSKIMESFKNMVPPQALVIRNGE KMSINAEDVVVGDLEVKGGDRIPA DLRIISANGCKVDNSSLTGESEPPQTR SPDFTNENPLETRNIAFFSTNCVEG TARGIVVYTGDRVTVMGRIATLASGLE GGQTPIAEEIEHFIHLITGVAVFLGVS FFILSLILEYTWLEAVIFLIGIIVANVP EGLLATVTVCLTLTAKRMARKNCLV KNLEAVETLGSTSTICSDKTGTLTQN RMTVAHMWFDNQNIEHEADTTENQS GVSFDKTSATWFALSRIAGLCNRAV FQANQENLPILKRAVAGDASESALL KCIEVCCGSMEMREKYSKIVEIPFN STNKYQLSIHKPNASEPKHLLVMK GAPERILDRCSILLHGKEQPLDEEL KDAFQDAYLELGGGLGERVLGFCHLL LPDEQFPEGFQFDTDDVNFVVDNLC FVGLISMIDPPRAAVPDAVGKCRSAG IKVIMVTGDHPITAKAIKGVGIISEG NETVEDIAARLNIPVNQVNPRDAKA CVVHGSCLKDMTSEELDDILRYHTE IVFARTSPQQKLIVEGCQRQGAIVAV TGDGVNDSPALKKADIGVAMGIVGS DVSQQAADMILLDDNFASIVTGVEE GRLIFDNLKKSIAYTLLTSNIPEITPFLI FIANIPLPLGTVTILCIDLGTDMVPAI SLAYEQAESDIMKQPRNPKTDKLV NERLISMAYGQIGMIQALGGFFTYFV ILAENGFLPFHLLGIRETWDDRWWVN DVEDSYGQQWYEQRKIVEFTCHTA FFVSIVVVQWADLVICKTRRNSVFQ QGMKNKILIFGLFEETALAAFLSYCP GMGAALRMYPLKPTWWFCAPYSL LIFVYDEVKRLIIRRRPGGWVEKETY Y