

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence
Q6PIE5	AT1A2_MOUSE	Mus musculus	Sodium/potassium-transporting ATPase subunit alpha-2	20.458298	NaN	S10;S439;S450;S496;S559;T570;S587;S672;S826;S940	22645316;33300544;36288343;29187734	MGRGAGREYSPAATTAENGGGKKK QKEKELDELKKEVAMDDHKLSLDE LGRKYQVDLSKGLTNQRAQDILARD GPNALTPPPTTPEWVKFCRQLFGGF SILLWIGALLCFLAYGILAAMEDEPS NDNLYLGIVLAADVIVTGCFSYQEA KSSKIMDSFKNMVPPQALVIREGEK MQINAEVVVVDLVEVKGGDRVPAD LRISSHGCKVDNSSLTGESEPOTRS PEFTHENPLETRNICFFSTNCVEGT ARGIVIATGDRTVMGRIATLASGLEV GQTPIAMEIEHFQILITGVAVFLGVSF FVLSLILGYSWLEAVIFLIGIIVANVPE GLLATVTVCLTLTAKRMARKNCLVK NLEAVETLSTSTICSDKTGTLTQNR MTVAHMWFDNQIHEADTTEDQSGA TFDKRSPTWTALSRIAGLCNRAVFK AGQENISVSKRDTAGDASESALLKCI ELSCGSRKMRDRNPKVAEIPFNST NKYQLSIHEREDSPQSHVLMKGGAP ERILDRCASTILVQGKEIPLDKEMQDA FQNAVMEGLGGERVLGFCQLNLP SGKFPRGFKFDDELNFPTEKLCFV GLMSMIDPPRAAVPDVAVGKCRSAGI KVIMVTGDHPITAKAIKGVGIISEGN ETVEDIAARLNIPVSQVNPREAKACV VHGSDLKDMTSEQLDEILRDHTEIV FARTSPQQLIIVEGCQRQGAIVAVT GDGVNDSPALKKADIGIAMGISGSD VSKQAADMILLDDNFASIVTGVEEG RLIFDNLKKSIAAYTLTSNIPEITPFLF IIANIPLPLGTVTILCIDLGTDMVPAIS LAYEAAESDIMKRQPRNSQTDKLVN ERLISMAYGQIGMIQALGGFFTYFVI LAENGFPLSRLGIRLDWDDRTTND LEDSYGQEWTYEQRKVVEFTCHTAF FASIVVVQWADLIICKTRRNSVFQQG MKNKILIFGLLEETALAAFLSYCPGM GVALRMYPLKVTWWFCAFPYLLIFI YDEVRLKLLRRYPGGWVEKETYY