



Ammonium Accumulation Caused by Reduced Tonoplast V-ATPase Activity in *Arabidopsis thaliana*

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Table S1 Primers of the genes targeted in the qRT-PCR assays.

Gene ID	GenBank accession	Primer	Sequence(5'-3')	Method
AT2G38290	NM129385	AMT2;1-F	CGGGAAAGATAGAATAACAAAATGG	Real-time qPCR
		AMT2;1-R	ATTGCTCCGATGACAGAAAGG	
AT1G32450	NM102980	NRT1.5-F	TGTCATTGGACTTTCATCGC	Real-time qPCR
		NRT1.5-R	CCCACAACCTCTGGTCTAAC	
AT4G21680	NM118288	NRT1.8-F	GGCTTCAGATTCTGGATAG	Real-time qPCR
		NRT1.8-R	AACCACAGAGTAGAGGATGG	
AT5G13170	NM121320	SAG29-F	GCCACCAGGGAGAAAAGG	Real-time qPCR
		SAG29-R	CCACGAAATGTGTTACCATTAGAA	
AT3G18780	NM112764	Actin2-F	CCACGAAATGTGTTACCATTAGAA	Real-time qPCR
		Actin2-R	TTTCCCGCTTGCTGTTGT	

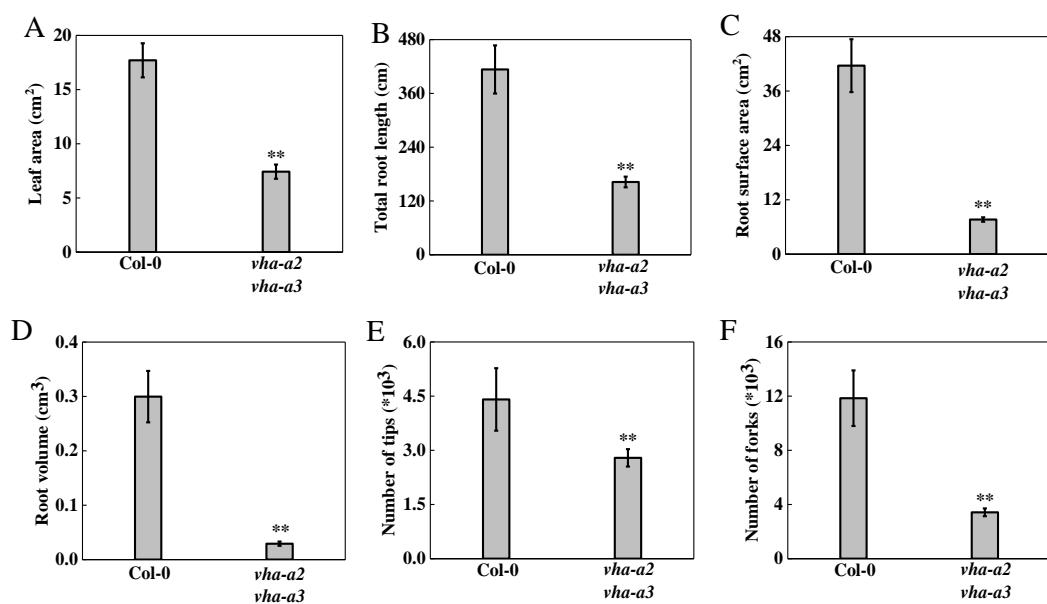


Figure S1 The leaf area and root configuration of wild type and the *vha-a2 vha-a3* double mutant. The rosette area (A), total root length (B), root surface area (C), root volume (D), the number of tips (E) and forks (F) were determined for Col-0 and *vha-a2 vha-a3*. Error bars represent S.D. of $n = 3$ biological replicates. Asterisks (**) indicate significant differences at $p < 0.01$.

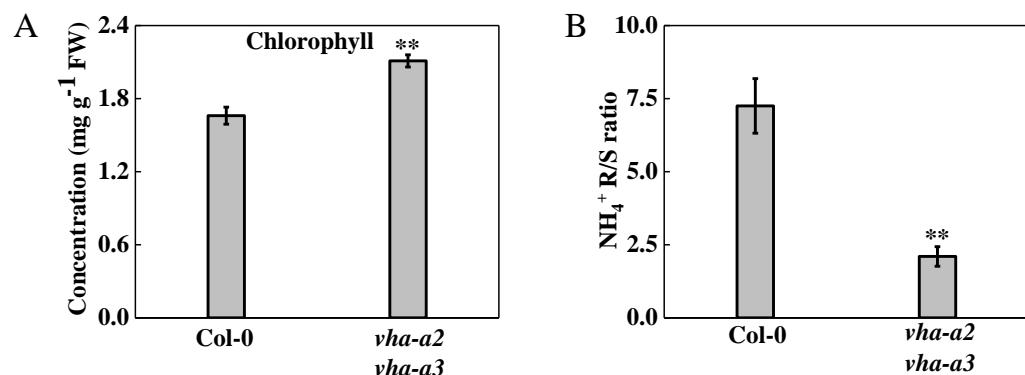


Figure S2 Chlorophyll concentration and NH_4^+ root/shoot ratio (R/S) of wild type and *vha-a2 vha-a3* mutant. The seedlings were grown for four-week-old under normal condition. Error bars were defined as S.D. of $n = 3$ biological replicates. Asterisks (**) indicate significant differences at $p < 0.01$. FW, Fresh weight.

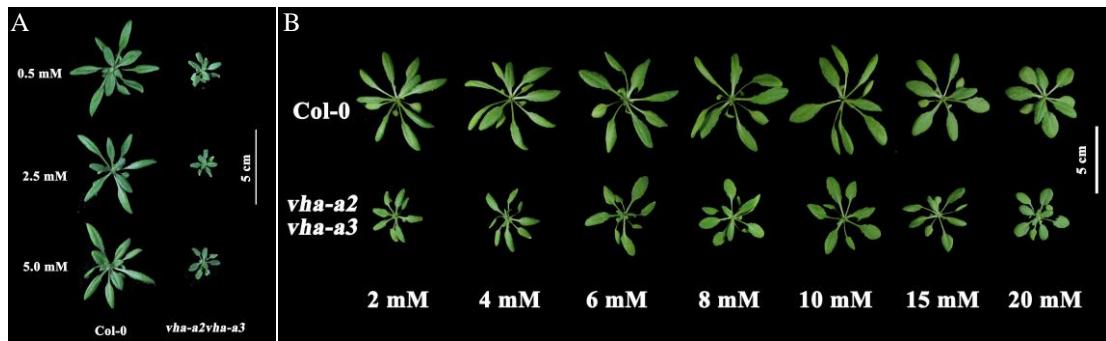


Figure S3 The phenotype of wild type and *vha-a2 vha-a3* mutant with different Ca²⁺ and K⁺ supply levels. (A) The phenotype of wild type and *vha-a2 vha-a3* mutant with additional Ca²⁺ supplied. (B) The phenotype of wild type and *vha-a2 vha-a3* mutant with additional K⁺ supplied. Pictures show four-week-old plants with different Ca²⁺ and K⁺ supply levels.

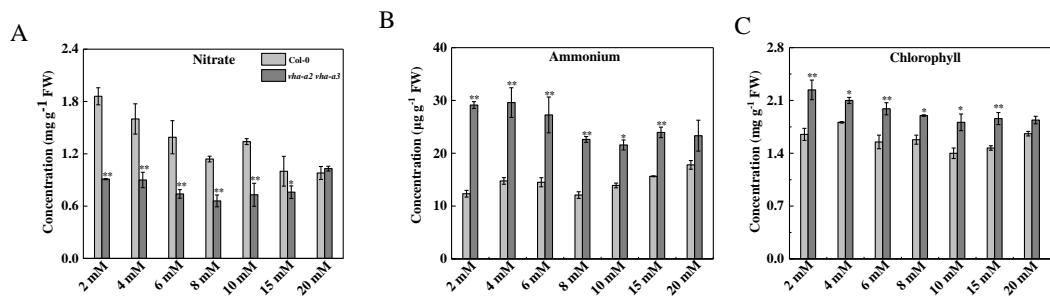


Figure S4 Nitrogen metabolites in wild type and *vha-a2 vha-a3* mutant with different potassium concentrations applied. The concentration of nitrate (A), ammonium (B), and chlorophyll (C) were determined with different K⁺ supply levels from shoots. Error bars were defined as S.D. of $n = 3$ technical replicates. Asterisks (*) and (**) indicate significant differences at $p < 0.05$ and $p < 0.01$, respectively.

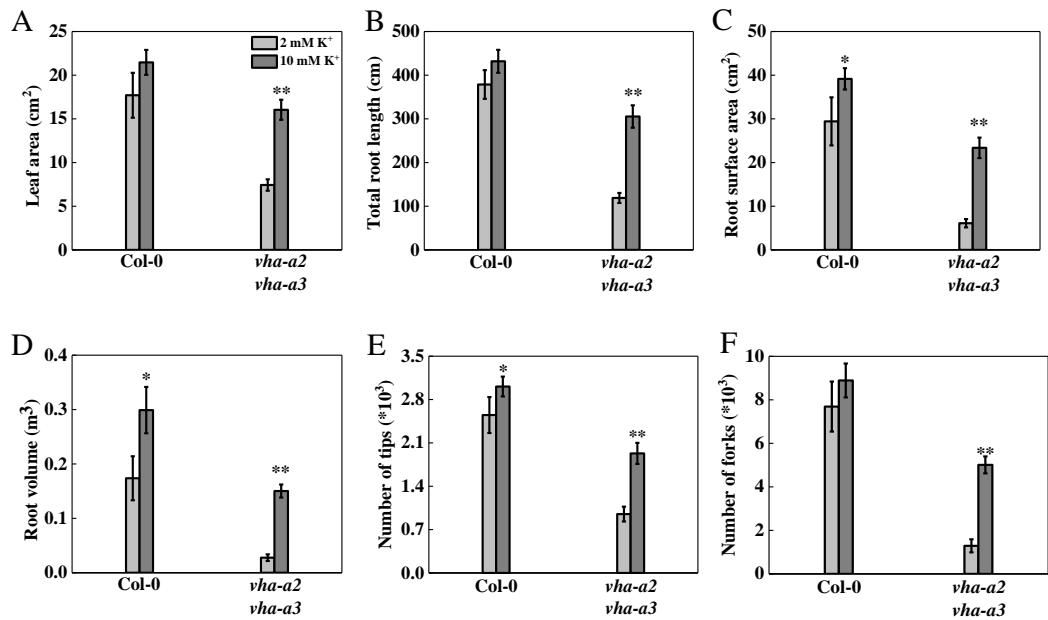


Figure S5 The leaf area and root configuration of wild type and the *vha-a2 vha-a3* mutant with different K^+ doses applied. The rosette area (A), total root length (B), root surface area (C), root volume (D), the number of tips (E) and forks (F) were determined for Col-0 and *vha-a2 vha-a3* mutant with different K^+ doses. Error bars represent S.D. of $n = 3$ biological replicates. Asterisks (*) and (**) indicate significant differences at $p < 0.05$ and $p < 0.01$, respectively.