

Bundle-sheath cells are leaf "water valves" controlled by an H⁺-ATPase via xylem acidification

One-sentence summary:

Cells enwrapping the leaf veins control the leaf hydraulic conductance by xylem sap pH regulated by a proton-pump

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Abstract

Bundle-sheath cells (BSCs), parenchymatous cells tightly enwrapping the leaf veins, constitute a selective and dynamic barrier to solutes and water. The over three-fold-higher abundance of the AHA2 (Arabidopsis H⁺-ATPase2) transcript we recently reported in Arabidopsis BSCs (compared to mesophyll cells) suggests AHA2 participation in this barrier function. Manipulating AHA2 activity in BSCs pharmacologically and genetically, while monitoring the xylem sap pH by fluorescence imaging of FITC-dextran fed into veins of detached leaves, demonstrated AHA2 indispensability for low sap pH, resolving the decades-lasting controversy about the molecular *origin* of xylem sap acidification. Furthermore, both AHA2 activity and sap acidification by added buffers enhanced the hydraulic leaf conductance, while alkaline sap pH – buffered or in AHA2 knockouts – reduced it. Similarly, the osmotic water permeability of isolated BSC protoplasts was reduced by alkaline and enhanced by acidic pH, thereby revealing the cellular mechanism underlying the above causative link between AHA2 activity and leaf water influx. This positions the BSCs as a transpiration-controlling valve in series with the stomata.

Introduction

The majority (95%) of nutrients and water which enter the plant through the roots move upward as bulk via xylem vessels (Taiz and Zeiger, 2014) all the way into the leaf veins. On its way out of the leaf xylem and into the leaf photosynthetic tissue, the mesophyll, this transpiration stream crosses a layer of parenchymatous cells (bundle sheath cells, BSCs) which tightly enwrap the entire vasculature (Kinsman and Pyke, 1998). BSCs have been shown to act as a selective barrier between the vein and the mesophyll, capable of impeding, for example, the transport of sodium (in the banana leaf (Shapira et al., 2009)) and of boron (in the leaves of *Thelungiella* (Lamdan et al., 2012), and of Arabidopsis (Shatil-Cohen and Moshelion, 2012), and of banana (Shapira et al., 2013)), as well as the passage of water (Shatil-Cohen et al., 2011; Pantin et al., 2013). However, it is still not clear (Geilfus, 2017) whether and how the apoplastic (xylem) milieu

modulates the “barrier behavior” of the bundle sheath, and in particular, whether this behavior depends on the xylem sap pH.

While under normal growth conditions the leaf apoplast pH is around 5.5-6 (reviewed by Grignon and Sentenac, 1991; Geilfus, 2017). The rare direct determinations of leaf xylem pH placed it at 6.3-6.7 in cotton leaves (Hartung et al., 1988), or at 5.3 in sunflower leaves (Jia and Davies, 2007). The leaf xylem sap pH can change in response to external conditions. For example, the leaf xylem sap alkalinized when tomato roots were exposed to drying soil or supplied with nitrate (Jia and Davies, 2007), or when detached sunflower leaves were supplied with HCO_3^- (Mengel, 1994). In turn, changes in the leaf xylem sap pH regulate physiological processes. For example, feeding high-pH solutions to detached leaves reduced stomatal conductance and transpiration both in *Commelina communis* and in tomato (Wilkinson and Davies, 1997; Wilkinson et al., 1998; Jia and Davies, 2007). As changes in pH define the dissociation state of weak acids, among them the major phytohormones abscisic acid (ABA) and indol acetic acid (IAA, auxin), they would thereby affect both their distribution between the apoplast and the cellular compartments of the plant, and hence, their biological activities. This has been shown, for example, for ABA accumulation in the alkalinizing xylem sap of a gradually dehydrated detached cotton leaf, which could be prevented by a pretreatment with an H^+ -ATPase-activating fungal toxin, fusicoccin (Hartung et al., 1988).

Moreover, xylem sap pH is likely a crucial component of the proton-motive force governing transmembrane transport between the xylem and the surrounding BSCs. Yet, in spite of its importance to the plant basic life processes, molecular evidence linking a specific H^+ -ATPase to the pH regulation of the leaf xylem sap and to the ensuing physiological changes in the leaf is missing.

H^+ -ATPases constitute a family of proton pumps driven by hydrolysis of ATP and are found in the plasma membrane of plants and fungi (Axelsen and Palmgren, 2001). 11 H^+ -ATPases isoforms have been reported in Arabidopsis; among them, the AHA1 and AHA2 are by far the most abundantly expressed members of this family throughout plant

life and tissues (Haruta et al., 2010). The expression of a GUS reporter gene in Arabidopsis, under the *AHA2* promoter, revealed abundant expression specifically in roots and leaves, and especially in the vascular tissue (Fuglsang et al., 2007). Our transcriptome analysis of protoplasts isolated from Arabidopsis BSCs and mesophyll cells (MCs) showed that the BSCs express the *AHA2* gene at a threefold higher level than the MCs, while the *AHA1* gene expression, though abundant, was not different between these two cell types (Wigoda et al., 2017). Here we show that BSCs act as an active barrier between xylem and mesophyll: (a) it is mainly the activity of *AHA2* of the BSCs which acidifies the sap, (b) the xylem sap pH regulates the radial hydraulic conductance of the leaf and (c) pH regulates the osmotic water permeability of the individual isolated BSCs protoplasts. We thus demonstrate a causative link between the activity of *AHA2* and the hydraulic conductance of the leaf and reveal that its underlying mechanism is the pH-controlled osmotic water permeability of the BSCs.

Results

The effect of pharmacological agents – a pump stimulator and an inhibitor – on xylem pH

We used the fluorescence of FITC-dextran (10 kDa) perfused via petioles into detached WT Arabidopsis leaves to monitor the pH within the leaf minor veins (Materials and methods, Suppl. Figs S1-S2). When 10 μ M fusicoccin (a fungal stimulator of P-type proton pumps, Serrano, 1988) was fed into the xylem, we observed a sharp decrease in xylem sap pH of about two pH units compared with leaves in control conditions (Figs. 1Ai, 1Aii, 1B) approximately 30 minutes after addition of fusicoccin. ETOH, the fusicoccin solvent, by itself, had no impact on xylem pH (Suppl. Fig. S3).

Further, WT leaves treated with 1 mM of vanadate (a commonly used P-type H^+ -pump inhibitor (reviewed by Palmgren, 2001), in a high- K^+ solution (10 mM KNO_3), resulted in xylem sap alkalization of about one pH unit within 30-40 min (Figs. 1Aiii, 1Aiv, 1C).

Interestingly, neither a similar exposure to vanadate in the low-K⁺ solution (XPS without added KNO₃, suppl. Fig.S4A), nor the high-K⁺ solution by itself (Suppl. Fig. S4B) had any effect on the xylem sap pH.

AHA2 acidifies the xylem pH

In an attempt to resolve between the relative contributions of the two abundant H⁺-ATPases of the BSCs, AHA1 and AHA2 (Wigoda et al., 2017), to the acidification of the xylem pH, we compared the xylem sap pH of WT plants to that in T-DNA-insertion mutants of either pump. We used two independent mutant lines with a homozygous loss of function of the *AHA2* gene (*aha2-4* and *aha2-5*) and three such lines with mutated *AHA1* (*aha1-6*, *aha1-7*, *aha1-8*). The xylem sap pH of both *AHA2* mutants, *aha2-4* and *aha2-5*, was consistently higher, by 0.5-1 pH units, compared to the WT plants (Fig. 2A). In contrast, there was no significant difference between the xylem sap pH in WT vs. the three lines of *AHA1* mutants (Suppl. Fig. S5).

To further test the ability of *AHA2* to acidify the xylem sap pH, we complemented the *AHA2* deficient plants (the *aha2-4* mutant), with the *AHA2* gene directed specifically to the BSCs (under the BSCs-specific promotor Scarecrow, SCR; Wysocka-Diller et al., 2000). Using qRT-PCR on whole leaves of mature transgenic plants, we confirmed *AHA2* absence in the *aha2-4* mutant (Fig. 3A, as in Haruta et al., 2010), and demonstrated successful complementation in two lines (T55 and T56). The transcript level of *AHA2* in these lines was even higher than in the WT (Fig. 3A). While the xylem sap pH in the *AHA2*-knockout mutant, *aha2-4*, was significantly more alkaline than in WT plants (repeating the results of Fig. 2A above), upon the complementation of the *aha2-4* mutant with the BSC-directed *AHA2*, the xylem sap pH in both lines became no higher than WT (Fig 3B). The level of *AHA1* transcript in the whole leaf was not affected by the genetic manipulations of *AHA2*, neither by *AHA2* mutation (as in Haruta et al., 2010), nor by the BSC-specific mutant complementation with *AHA2* (Fig. S6).

Basic pH lowers the leaf hydraulic conductance, K_{leaf}

To examine the impact of AHA2 on the water economy of the whole leaf we determined the leaf water potential (Ψ_{leaf}) and its transpiration (E), and from these two we calculated the leaf hydraulic conductance (K_{leaf} , Eq. 1 in Materials and methods). We compared the K_{leaf} in detached leaves of WT and of the *aha2-4* and *aha2-5* knockouts fed with unbuffered XPS. Notably, the K_{leaf} of the mutants was appreciably lower than that of WT leaves, about 50 % in *aha2-4* and about 30 % in *aha2-5* (Fig. 4A) although all three leaf types transpired at a similar rate (Suppl. Figs. S7A and S7E). Furthermore, WT leaves perfused with XPS^b buffered at pH 7.5 had K_{leaf} lower by over 50% compared to WT leaves perfused with XPS^{db} buffered at pH 6 (Fig. 4B), again, without a difference in their transpiration rate (Suppl. Figs. S7B and S7F).

Basic pH lowers the osmotic water permeability coefficient of BSCs protoplast membrane, P_f

In order to test the hypothesis that the pH-dependent reduction in K_{leaf} is due to a reduction in the water permeability of the BSCs membranes, we measured the osmotic water permeability coefficient, P_f , of BSC under two pH treatments, using GFP labeled BSCs, from SCR:GFP plants. The mean initial P_f (see Materials and methods) of BSCs treated with pH 6 was ~8.5 fold higher than that of BSCs treated with pH 7.5. ($5.39 \pm 1 \mu\text{m sec}^{-1}$ and $0.63 \pm 0.2 \mu\text{m sec}^{-1}$, respectively, Fig. 5).

Discussion

The AHA2 of the BSCs is indispensable for the leaf xylem sap acidification

H^+ -ATPases are well known regulators of the apoplastic pH which has a key role in cell ion homeostasis (Taiz and Zeiger, 2014). Here we demonstrate, for the first time, that AHA2 which resides in the BSCs plasma membrane, exerts a dominant role in the acidification of the xylem sap in leaves. This conclusion is based on four different complementing approaches used in our study.

(1) *In-vivo measurement of xylem pH.* The ratio of fluorescence intensity of FITC-dextran excited at two different wavelengths is independent of the absolute dye concentration and is related only to pH. By this dual excitation technique, any variations in dye concentration caused by dye loading, compartmentation, leakage and photo-bleaching can be compensated for, since with this dye these parameters have a similar effect on fluorescence intensities at both wavelengths. Hoffmann and Kosengarten (Hoffmann et al., 1992; Hoffmann and Kosegarten, 1995) successfully measured the apoplastic pH, pH gradient between various cell types in the apoplast of sunflower leaves, and changes of leaf apoplastic pH in response to the uptake of NH_4^+ and NO_3^- via petioles. In maize, measurements using pH-sensitive microelectrodes validated FITC-dextran as a faithful reporter of apoplastic pH-conditions (Pitann et al., 2009). In using the membrane-impermeable FITC-Dextran fed to the Arabidopsis leaf via a petiole, we relied on earlier demonstrations of the tight isolation of the vascular system due to the bundle sheath (Kinsman and Pyke, 1998; Shatil-Cohen et al., 2011; Shatil-Cohen and Moshelion, 2012; Sade et al., 2014;). We were thus satisfied that the FITC probe remained confined and reported reliably on the pH from within the BSC-lined xylem apoplast of the minor veins (Fig. 1A).

(2) *The use of classical pharmacological agents.* An H^+ -ATPase specific enhancer (fusicoocin) and inhibitor (vanadate), fed directly to the leaf xylem, established that pH acidification of the leaf xylem sap involves P-type proton pump(s) (AHAs), which extrude protons – most likely, from the BSCs – into the xylem lumen. Notably, in our experiments, vanadate increased the xylem sap pH only when it was administered in high- K^+ XPS, while vanadate in low- K^+ XPS did not cause any detectable pH change. We explain this requirement for 10 mM KNO_3 to show vanadate inhibition of the H^+ -pump, as resulting from an increased concentration of substrate (the added K^+ and NO_3^- ions) which can participate in secondary H^+ -co-transport into the BSCs, thereby dissipating the protons from the xylem lumen. This conforms with the aforementioned general notion that the pH of an apoplastic compartment reflects, among others, a balance in bi-directional proton movements (Serrano, 1988). That proton-coupled K^+

transmembrane co-transport plays a particularly important role in the BSCs is suggested by the 10% higher expression in BSCs (relative to mesophyll cells) of two K^+ uptake permeases, AtKT2 and AtKUP11 (likely to be H^+ -coupled (Wigoda et al., 2017) and references therein).

(3) *Genetic manipulation – knockouts.* We have established here, for the first time, the dominant role of AHA2 in regulating the xylem sap pH in Arabidopsis leaf. While the pH of xylem sap in both AHA2 mutant lines, *aha2-4* and *aha2-5* was consistently higher than WT, xylem sap in the *aha1* mutant lines was no different than in WT.

(4) *Genetic manipulation – knockout complementation.* The xylem sap pH was restored to WT-like levels in two lines resulting from *aha2-4* complementation with the AHA2 gene directed specifically into the BSCs, using the specific promotor SCR. Somewhat surprisingly, although the complementation was directed to BSCs, the AHA2 transcript levels in the whole leaf of the complemented plants were considerably higher than in the WT leaf. Nevertheless, the level of AHA2 activity – as reflected in the xylem sap pH (especially in the complemented line T55, where pH was intermediate between the mutant *aha2-4* and the WT) – was somewhat lower than could be expected from the transcript level. This may suggest a post-transcriptional regulation of the AHA2 in BSCs.

Moreover, we noticed a lack of a change in the transcript level of AHA1, not only in the mutant *aha2-4*, as already noted by Haruta et al., (2010), but also in the BSC-directed complementation lines T55 and T56 (Fig. S6). This lack of *inverted* correlation between the transcript levels of AHA1 and AHA2 in the above examples, contrary to what might be expected had they shared responsibility for the same function and, therefore would have been linked by a mutual feedback, is another indication for the already inferred relative specificity of AHA2 for its BSCs related function – the maintenance of xylem sap pH.

The physiological relevance of extracellular pH

External alkalization reduces the osmotic water permeability (P_f) of the BSCs. It had been established that low cytosolic pH inhibits aquaporin gating and in doing so reduces P_f (Tournaire-Roux et al., 2003; Alleva et al., 2006). This phenomenon has been studied in roots under stress conditions such as flooding or drought, where a change in the water hydraulic permeability of the roots is considered to be part of the plant stress defense mechanism. External medium acidification also reduced P_f in “right-side-out” red beet root PM vesicles (i.e., vesicles with aquaporin membrane topology the same as in the cell), which, because of a reduction of the activation energy for water transport was attributed to aquaporins (Alleva et al., 2006). In contrast, the fungal aquaporin RdAQP1, when expressed in *Arabidopsis* protoplasts, *elevated* the cell membrane water permeability at acidic external pH, but increasing the external pH to 7 abolished this elevation (Turgeman et al., 2016). The reduction in the fungal aquaporin activity with rising external pH was mediated by histidines facing the outside of the cell (*ibid.*). Here we demonstrate, for the first time in any plant cell, a reduction in P_f driven by external alkalization in a leaf bundle sheath cell, the BSC. Future work is required to clarify whether this P_f reduction in BSCs is mediated by aquaporins and their external histidines.

AHA2 role in the regulation of the whole leaf water balance. The notion that AHA pumps, in general, power secondary H^+ -cotransport across cell membranes facing the apoplast is decades old (Serrano, 1988; Sze et al., 1999; Palmgren, 2001). The plant plasma membrane H^+ -ATPase has since been recognized as essential for plant growth (reviewed by Falhof et al., 2016). The practically sole “celebrated product” of its action to date is the proton motive force (PMF) – a gradient of protons concentration without- or in combination with an electrical gradient – which drives the movement of solutes across cellular membranes. In only a few cases has a *specific* physiological role been assigned to the plasma membrane H^+ -ATPase – mainly in stomatal physiology and in roots (reviewed by Falhof et al., 2016). Here we show for the first time another, novel aspect of transport activity *regulated* (rather than *driven*) by AHA2 – that of water fluxes

from the xylem into the leaf and across the bundle sheath layer – evident as the leaf hydraulic conductance, K_{leaf} . We demonstrate a causative inverse correlation between the resulting K_{leaf} and the xylem sap pH, be it pH manipulated directly by buffers, or altered genetically by abolishing AHA2 activity (in *AHA2* knockouts).

In an earlier work, Shatil-Cohen et al. (Shatil-Cohen et al., 2011) localized K_{leaf} to the BSCs layer by showing that K_{leaf} decreased when the BSCs were brought into direct contact with the plant stress hormone, abscisic acid (ABA) fed to the detached leaf, and not when ABA was smeared on the leaf surface. Our current work outlines a mechanism for this phenomenon: stress-induced xylem alkalization (already reported by others (Jia and Davies, 2007; Wang et al., 2012; Korovetska et al., 2014), likely due to ABA inhibition of the BSCs AHA2, similar to the H^+ -ATPase inhibition seen in guard cells (Goh et al., 1996; Schroeder et al., 2001; Zhang et al., 2004), reduces the water permeability of the BSCs membranes. Consequently, the water permeability of the entire bundle sheath layer declines, demonstrating that under stress conditions it is an active hydraulic barrier. Here we show directly, using individual isolated BSCs, that P_f , the measure of the rate of water passage via the BSCs membranes, declines with external medium alkalization. Such a pH rise could mediate the effect of ABA on BSCs in the detached leaf.

The results of this study broaden our basic understanding of how a leaf controls its water influx. Our results also support the notion that xylem sap alkalization mediates the decline in plant shoot nutrient uptake due to abiotic stress. Here we focus specifically on the stress-induced cessation of activity of AHA2 in the BSCs, where we found the AHA2 instrumental in generating the low pH in the leaf xylem sap. The consequence of this decline of AHA2 activity and xylem sap alkalization, is a decline in the PMF. Since AHA2 – via the PMF – powers the secondary, proton-coupled export and import of solutes across the BSCs membranes, not only does it regulate plant nutrition, but, very likely also drives the extrusion of toxic compounds from the xylem-lining cells into the

xylem, hence governing also the plant toxicity tolerance. These hypotheses await future experimentation.

In conclusion, rapid growth rate and high yields of crop plants are positively correlated with enhanced transpiration and K_{leaf} . Our finding that AHA2 regulates K_{leaf} via xylem sap pH provides a molecular basis for understanding a novel aspect, other than just PMF, of the control that the xylem sap pH can exert in the leaf. This control, in combination with effects of PMF, is likely to underlie the plant's key physiological activities. These results provide a new focus for exploration and understanding the role of the involvement of BSCs in determining the xylem sap composition, and, in particular, their role as a transpiration-controlling valve in series with the stomata. BSCs are likely to become a key target tissue for the development of a new generation of manipulations for plant adaptation to environmental challenges.

Materials and Methods

Plant material

Plant types. We used WT (wild type) *Arabidopsis thaliana* plants ecotype Columbia, Col-0 (aka Col) and T-DNA insertion *AHA* mutants (Col) , *AHA1* mutants: *aha1-6* (SALK_ 016325), *aha1-7* (SALK_ 065288) and *aha1-8* (SALK_ 118350), and *AHA2* mutants; *aha2-4* (SALK_ 082786) and *aha2-5* (SALK_022010) obtained from the Arabidopsis Biological Resource Center (Ohio State University). The plants' genotypes were determined by PCR with allele-specific primers and then the single-gene knockout were confirmed for the absence of *AHA2* (or *AHA1*) RNA using real time PCR (RT-PCR). The sequences of the PCR primers are provided in the supplemental Table S1. For the experiments of Fig. 5, we used SCR:GFP (Col) Arabidopsis plants expressing GFP specifically in the BSCs ER generated in our lab and described in detail by Attia et al., (2018)

Plant Growth Conditions. All plants were grown in soil as described by (Shatil-Cohen et al., 2011). Plants were kept in a growth chamber under short-day conditions (10-h light) and a controlled temperature of 20–22°C, with 70% humidity. Light intensity at the plant level of 100-150 $\mu\text{mol m}^{-2} \text{sec}^{-1}$, using either fluorescent lights (Osram cool white, T5 FH 35W/840 HE, Germany) or LED light (EnerLED 24V-5630, 24W/m, 3000K (50%)/6000K(50%)). The plants were irrigated twice a week.

Determination of xylem sap pH in detached leaves

Leaf perfusion. Leaves from 6-7 week old plants, approximately 2.5 cm long and 1 cm wide, were excised at the petiole base using a sharp blade and immediately dipped in “xylem perfusion solution” (XPS, see Solutions below), in 0.5 ml Eppendorf tubes for 30 minutes. Perfusion experiments using Safranin O (Sigma cat. #: S2255; 1% w/v in XPS) demonstrated that 30 minutes incubation sufficed for the whole leaf perfusion via the petiole by means of the transpiration stream (Fig. S1). All experiments were conducted between 1-4 hours from lights on.

The leaf xylem pH was determined using the dual-excitation membrane-impermeant fluorescent dye fluorescein isothiocyanate conjugated to 10 kD dextran (FITC-D; (Hoffmann and Kosegarten, 1995; Mühlhng et al., 1995)) dissolved in the solution perfused into the detached leaf. An *in-vivo* pH calibration curve was constructed using XPS buffered to predefined pH values (5, 5.5, 6, 6.5, 7 and 7.5; see Solutions below). Effects of vanadate in high-K⁺ solution or of fusicoccin in low-K⁺ solution (i.e., in XPS) was tested in the absence of pH buffers (see Solutions below).

Sample preparation on the microscope stage. Immediately after the leaf xylem perfusion, the leaf was washed with that leaf's intended perfusate (the specific XPS to be tested) without the dye, laid on a microscope slide abaxial side up, and a drop of XPS (without the dye) was placed on top. Then, a thin layer of silicone high vacuum grease (Merck cat. #: 1.07921.0100) was applied around the leaf edge and a large coverslip (50 by 20 mm, larger than the leaf) was placed on the leaf and affixed to the slide beneath. The minor veins on the abaxial side were imaged via the coverslip.

Fluorescence microscopy was performed using an inverted microscope (Olympus-IX8 integrated within the Cell-R system, <http://www.olympus-global.com>), via an UPlanSApo 10X/0.40 (∞ /0.17/ FN26.5) objective. Pairs of images were recorded at a quick succession (at an approx. 200 ms interval) at two excitation wavelengths of 488 nm and 450 nm, and fluorescence at a single emission wavelength of 520 nm (Hoffmann and Kosegarten, 1995; Mühlhng et al., 1995) was recorded with a 12-bit CCD camera, Orca-AG (Hamamatsu, <http://www.hamamatsu.com>) and saved in a linear 16 bit tiff format.

In all fluorescence microscopy experiments, each treatment was performed on at least five leaves, each from a different plant, on three days (all together, five biological repetitions in three independent experiments per treatment), alternating randomly among different treatments. Four to six paired images of a minor vein were obtained from different areas of each leaf (four-six technical repetitions per a biological repeat). For background values, leaves were perfused with experimental solutions without the dye; one leaf was sampled per treatment or plant type on each day of an experiment (all

together, at least three biological repetitions per treatment or plant type, a total of 20 leaves, with three technical repetitions per each biological repeat).

Image analysis. The acquired pairs of images were exported in a linear 16-bit tiff format for further processing with ImageJ (ver. 1.49V; <http://rsbweb.nih.gov/ij/>). The pixel by pixel ratio of the two images in each pair was calculated using the ImageJ ‘Ratio-plus’ plugin (Paulo J. Magalhães, 2004). The ratio calculation took into account the average background fluorescence at the corresponding excitation wavelength, which was subtracted from each pixel’s fluorescence. Further analysis was performed on selected pixels, based on the image resulting from excitation at 488 nm. Only pixels with fluorescence values below the camera saturation level (<4095) and, at the same time, of at least 3 fold higher than the mean background value were selected (areas enclosed within yellow lines in the supplemental Fig. S2C). For each leaf (biological repeat), the mean fluorescence ratio obtained from a vein segment was converted to a pH value, using an *in-vivo* calibration curve, which was obtained, as described above, from paired images of WT leaves fed with pH-calibration solutions (see Solutions below). We occasionally verified the system stability using an *in-vitro* calibration curve, derived from imaging drops of the calibration solutions placed on microscope slides rather than fed into leaf veins. Also, changing the composition of the buffers (MES and HEPES) in the XPS had no effect on the calibration curve.

Physiological characterization of the leaf (gas exchange and hydraulic conductance, K_{leaf})

Sample preparation. For Fig. 4A (and Suppl Figs. S7A, 7C, 7E), the leaves were excised in the evening preceding the measurements, in the light, shortly before the “lights OFF” transition, their petioles were dipped in the unbuffered High- K^+ XPS and placed in gas-sealed transparent 25 x 25 x 15 cm plastic boxes with water-soaked tissue paper on the bottom to provide ~100% humidity. Each leaves-containing box was then placed in another, lightproof box for the duration of the night. Measurements were conducted in the morning, between 10:00 AM and 12:00 PM, after 20 min illumination in the growth chamber.

For Fig. 4B (and Suppl. Figs. S7B, S7D, S7F), leaves were excised on the morning of the experiment, in the dark, about 10 min before the regular “lights ON” transition and their petioles were dipped in vials with buffered XPSdb (at pH, 6 and 7.5; see Solutions below). The vials were placed in gas-sealed transparent plastic boxes as described above. The boxes were then kept for two hours in the growth room under the regular light and temperature conditions. Prior to the measurements, the boxes were opened for 3 minutes to equilibrate with the ambient vapor pressure deficit (VPD) of 1.3-1.5 kPa. The measurements were conducted between 10:00AM and 1:00 PM (1-4 hours after lights ON).

Gas-exchange assays, i.e., g_s (stomatal conductance) and E (transpiration) were performed (as in Sade et al., 2014) using a Li-Cor 6400 portable gas-exchange system (LI-COR, USA <https://www.licor.com/>) equipped with a standard leaf cuvette with an aperture of 2 x 3 cm). The transfer time from box to Licor was at most 5 s. The measuring conditions were set to be as similar as possible to the growth chamber conditions (illumination: $150 \mu\text{mol m}^{-2} \text{s}^{-1}$; the amount of blue light was set to 10% of the photosynthetically active photon flux density to optimize stomatal aperture, the leaf temperature: approximately 24 °C, the VPD: approximately 1.3 kPa, and $[\text{CO}_2]$ surrounding the leaf: $400 \mu\text{mol mol}^{-1}$).

Measuring the leaf water potential, Ψ_{leaf} . Immediately following the gas exchange measurement which took about 5 min, the leaf was transferred to a pressure chamber (ARIMAD-3000; MRC Israel) equipped with a home-made silicon adaptor especially designed to fit Arabidopsis petioles into the chamber’s O-ring. Ψ_{leaf} was determined as described in (Sade et al., 2014).

Determination of the leaf hydraulic conductance, K_{leaf} . K_{leaf} was calculated for each individual leaf as follows (Martre et al., 2000):

Eq. 1:
$$K_{\text{leaf}} = E / (\Psi_{\text{Leaf}} - \Psi_{\text{XPS}}) \approx E / \Psi_{\text{Leaf}},$$

where E is the whole-leaf’s transpiration, i.e., the water flux, and Ψ_{Leaf} is the leaf water potential and Ψ_{XPS} is the water potential of XPS (or of the XPS^{db}); as Ψ_{XPS} is nearly null, in this calculator we used solely the leaf water potential value.

Protoplast isolation

For protoplast isolation, Protoplasts were isolated from 6- to 8-week-old plants using the rapid method (Shatil-Cohen et al., 2014) with our pH 6 isotonic solution in the extraction process (600 mOsmol, see solutions below).

Osmotic water permeability coefficient (P_f) measurements

P_f was determined as described by Shatil-Cohen et al. (2014), except here we used an inverted epifluorescent microscope (Nikon eclipse TS100) with a 20x/NA 0.40 objective (Nikon) and a CCD 12 bit camera Manta G-235B (<https://www.alliedvision.com>), and an image-processing software AcquireControl® v5.0.0 (<https://www.alliedvision.com>). We recorded the BSCs swelling in response to hypo-osmotic challenge (of 0.37 MPa) generated by changing the bath solutions from an isotonic one (600 mOsm) to a hypotonic one (450 mOsm, *ibid.*). The challenges were performed at pH 6 and at pH 7.5. The osmolarity of these solutions (variants of the XPS^{db}; see Solutions) was adjusted with the appropriate amounts of D-sorbitol and was verified within 1% of the target value using a vapour pressure osmometer (Wescor). P_f was determined from the initial rate of the cell volume increase using a numerical approach, in an offline curve-fitting procedure of the P_f Fit program, as described in Shatil-Cohen et al., (2014) and detailed in Moshelion et al., (2004). We present here the values of the initial P_f (P_{fi}) obtained by fitting model 5 (*ibid.*).

Solutions

FITC-D dye (Sigma cat. #: FD10S) was added from a 10 mM stock in water (kept aliquoted, protected from light, at -20 °C) to all the XPS solutions (except in the autofluorescence assays) to a final conc. of 100 μ M.

XPS, basal Xylem Perfusion Solution: 1 mM KCl, 0.3 mM CaCl₂ and 20 mM D-sorbitol (Sigma cat# 8143). Upon preparation, when unbuffered, the pH of this solution was 5.6 - 5.8 and its osmolarity was approx. 23 mOsm/L.

Low-K⁺ XPS: the same as XPS above.

High-K⁺ XPS: XPS + 10 mM KNO₃ and D-sorbitol adjustment to approx. 23 mOsm/L. Upon preparation, when unbuffered, the pH of this solution was 5.6 -5.8.

XPS^b (pH calibration solutions): XPS buffered with 20 mM MES and HCl or NMG to pH 5, 5.5, or 6), or with 20 mM HEPES and NMG to pH 6.5, 7, or 7.5. Osmolarity of these solutions was adjusted with D-sorbitol to 23 mOsm/L.

XPS^{db}: High-K⁺ XPS buffered with 10 mM MES and 10 mM HEPES and adjusted to pH 6 or 7.5 with N-Methyl D-glucamine (NMG). These solutions were adjusted with D-sorbitol to a final osmolarity of 40 mOsm/L.

Solutions for leaf physiological characterization: Non-buffered High-K⁺ XPS (as above) was used for Figure 4A (and Suppl. Figs. S7A,S7C, S7E). XPS^{db} was used for Fig. 4B (and Suppl. Figs. S7B, S7D, S7F).

Solutions for P_f determination: pH 6 solutions : XPS^{db} adjusted to pH 6 by NMG, and adjusted with D-sorbitol to either 450 mOsmol (hypotonic) or 600 mOsmol (isotonic); pH 7.5 solutions: XPS^{db} adjusted to pH 7.5 by NMG, and adjusted with D-sorbitol to either 450 mOsmol (hypotonic) or 600 mOsmol (isotonic).

Solution for protoplasts isolation: pH 6 isotonic solution as for P_f determination.

Generation of SCR:AHA2-complemented plants

Vector construction: *AHA2* gene was cloned into pDONR™ 221 (Invitrogene) vector and the SCR promoter into pDONRP4P1r using Gateway® compatible by BP reactions, and later cloned into the pB7M24GW (Invitrogene) two fragment binary vector by LR reaction according to the manufacturer's instructions. The binary *SCR:AHA2* vector was transformed into *Agrobacterium* by electroporation; transformants were selected on LB plates containing 25 µg/mL gentamycin and 50 µg/mL spectinomycin.

aha2-4 and aha2-5 mutant lines transformation with SCR:AHA2: was performed using the floral dip method (Clough and Bent, 1998). Transformants were selected based on their BASTA resistance, grown on plates with MS (Murashige and Skoog, Duchefa cat# M222.0050) Basal medium + 1 % sucrose and 20 µg/ml BASTA (Glufosinate Ammonium, Sigma cat # 45520). DNA insertion was verified in selected lines by PCR targeting the junction of *AHA2* and the 35S terminator with forward primer about 1000bp

from the 3' end of AHA2 and reverse primer on the 35S terminator (see primer list in supplemental Table S1).

AHA2 gene expression in the whole leaf by qRT-PCR

RNA extraction and quantitative real-time (qRT-) PCR. Total RNA was extracted from leaves using Tri-Reagent (Molecular Research Center, cat. #: TR 118) and treated with RNase-free DNase (Thermo Scientific™, cat. #: EN0525). Complementary DNA (cDNA) was prepared using the EZ-First Strand cDNA synthesis kit (Biological Industries cat. #: 2080050) according to the manufacturer's instructions. qRT-PCR was performed using C1000 Thermal Cycler (Bio-Rad), in the presence of EvaGreen (BIO-RAD cat.# 172-5204) and PCR primers to amplify specific regions of the genome (Haruta et al., 2010; suppl. Table S1). The results were analyzed using Bio Rad CFX manager™ software. Dissociation curve analysis was performed at the end of each qRT-PCR reaction to validate the presence of a single reaction product and lack of primer dimerization. Expression levels of examined genes were normalized using two normalizing genes (AT5G12240 and AT2G07734, Wigoda et al., 2017).

Statistics

The Student's unpaired two-tailed t-test was used for comparison of two means, which were considered to be significantly different at $P < 0.05$. For comparisons of three or more population means we used ANOVA, all-pairs Tuckey HSD (JMP® Pro 13), wherein different letters represent statistical differences at $P < 0.05$. Images which yielded extreme ratio values which were more than 2.5 SD above the mean were discarded.

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Author Contributions

Author contributions:

N. Wigoda and Y. Grunwald planned performed and analyzed the experiments, and wrote the paper. A. Yaaran and T. Torne participated in generating the complemented plant lines, and A. Yaaran and S. Gosa participated in the leaf hydraulics determinations. N. Sade participated in the genotyping of the mutant AHA lines. M. Moshelion and N. Moran conceived the experiments, guided the students and wrote the paper.

Competing interests

All authors declare they have no competing interests.

Supplementary Materials

Figure S1. Perfusion of detached leaves via petioles

Figure S2. Image analysis details

Figure S3. Fusicoccin (and not its solvent EtOH by itself) lowered the leaf xylem sap pH

Figure S4. Neither vanadate in the *low-K⁺* XPS nor the high-*K⁺* XPS *without* vanadate affect the xylem sap pH in WT Arabidopsis

Figure S5. Knockout of AHA1 does not increase xylem sap pH in minor leaf veins of Arabidopsis leaf.

Figure S6. Expression levels of *AHA1* in leaves of WT, *aha2-4* and of two independent bundle-sheath-specific *AHA2*-complimented lines (T55, T56)

Figure S7. Knockout of *AHA2* or alkaline xylem sap pH decrease K_{leaf} in detached leaves.

Table S1. List of primers used for genotyping (PCR) and expression quantification (RT-PCR) of *AHA1* and *AHA2* in mutants and transformed plants.

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Main Figure Legends

FIGURE 1. Leaf xylem sap pH reduced by petiole-fed fusicoccin (an AHA stimulator), and increased by vanadate (an AHA inhibitor) in minor veins in detached leaves of WT (wild type) Arabidopsis. A. Representative images of treatment effects with color-coded pH values calculated ratiometrically for all pixels which passed the selection criteria (Supplementary Materials and methods), with black masking all other pixels. i, low- K^+ control (i.e., XPS, xylem perfusion solution). ii, XPS + fusicoccin (10 μ M). iii, high- K^+ control solution (XPS +10 mM KNO_3). iv, XPS +10 mM KNO_3 + vanadate (1 mM Na_3VO_4). **B.** The mean (\pm SE) xylem sap pH without and with fusicoccin in the indicated number of leaves (biological repeats) from at least three independent experiments. Asterisks denote significant differences from the respective control, using Student's two-tailed unpaired t-test (*: $P < 0.05$, **: $P < 0.001$). Where error bar is invisible, SE= 0.009. Other details as in A. **C.** The mean (\pm SE) calculated values of the xylem sap pH without and with vanadate. Other details as in B.

FIGURE 2. Knockout of *AHA2* increases xylem sap pH in minor leaf veins of Arabidopsis leaf. *aha2* knockout lines and WT plants. The mean (\pm SE) xylem sap pH in the indicated number of leaves, from three independent

experiments. Different letters denote significantly different pH values ($P < 0.05$; ANOVA).

FIGURE 3. Expression levels of *AHA2* and *AHA1* and xylem sap pH in leaves of WT, *aha2-4*, and in two independent bundle-sheath-specific *AHA2*-complimented lines (T55, T56). A. Mean normalized (\pm SE) *AHA2* expression levels obtained by qRT-PCR on whole leaf RNA ($n=5$ biological repetitions, leaves; see Suppl. Materials and methods). **B.** Mean (\pm SE) xylem sap pH in the indicated number of leaves from three independent experiments. Different letters indicate significantly different means ($P < 0.05$; ANOVA). **C.** Mean normalized (\pm SE) *AHA1* expression levels; other details as in A.

FIGURE 4. The leaf hydraulic conductance (K_{leaf}) depends on the activity of *AHA2* and the pH of the xylem perfusion solution (XPS). Mean (\pm SE) K_{leaf} (calculated as in Suppl. Fig. S5A) in detached Arabidopsis leaves. **A.** leaves of WT and *AHA2* knockout plants perfused with non-buffered XPS. Numbers are those of assayed leaves. Different letters indicate significantly different means ($P < 0.05$; ANOVA). Note that K_{leaf} is lower in the *AHA2* knockouts relative to WT. **B.** WT Arabidopsis leaves fed with XPS^b pH-buffered as indicated, from three independent experiments. Asterisk indicates significantly different means ($P < 0.05$; Student's non-paired two-tailed t-test). Note that K_{leaf} is lower at alkaline pH relative to acidic pH. A and B results are from experiments done at different time and with different solutions (see Materials and Methods).

Figure 5: The effect of pH treatment on the membrane osmotic water permeability coefficient (P_f) of BSCs from SCR:GFP plants. A. Time course (60 sec) of bundle sheath protoplasts swelling upon exposure to a hypotonic XPS^{db} solution at pH 6 or 7.5. The arrow indicates onset of bath flush. **B.** Time course of the osmotic concentration change in the bath (C_{out}) during the hypotonic challenge (calculated as in Moshelion et al., 2004). **C.** Mean (\pm SE) initial P_f values of the

indicated number of bundle sheath protoplasts under the different pHs from three independent experiments. The asterisk denotes a significant difference between the treatments using Student's two-tailed unpaired t-test ($P < 0.01$)

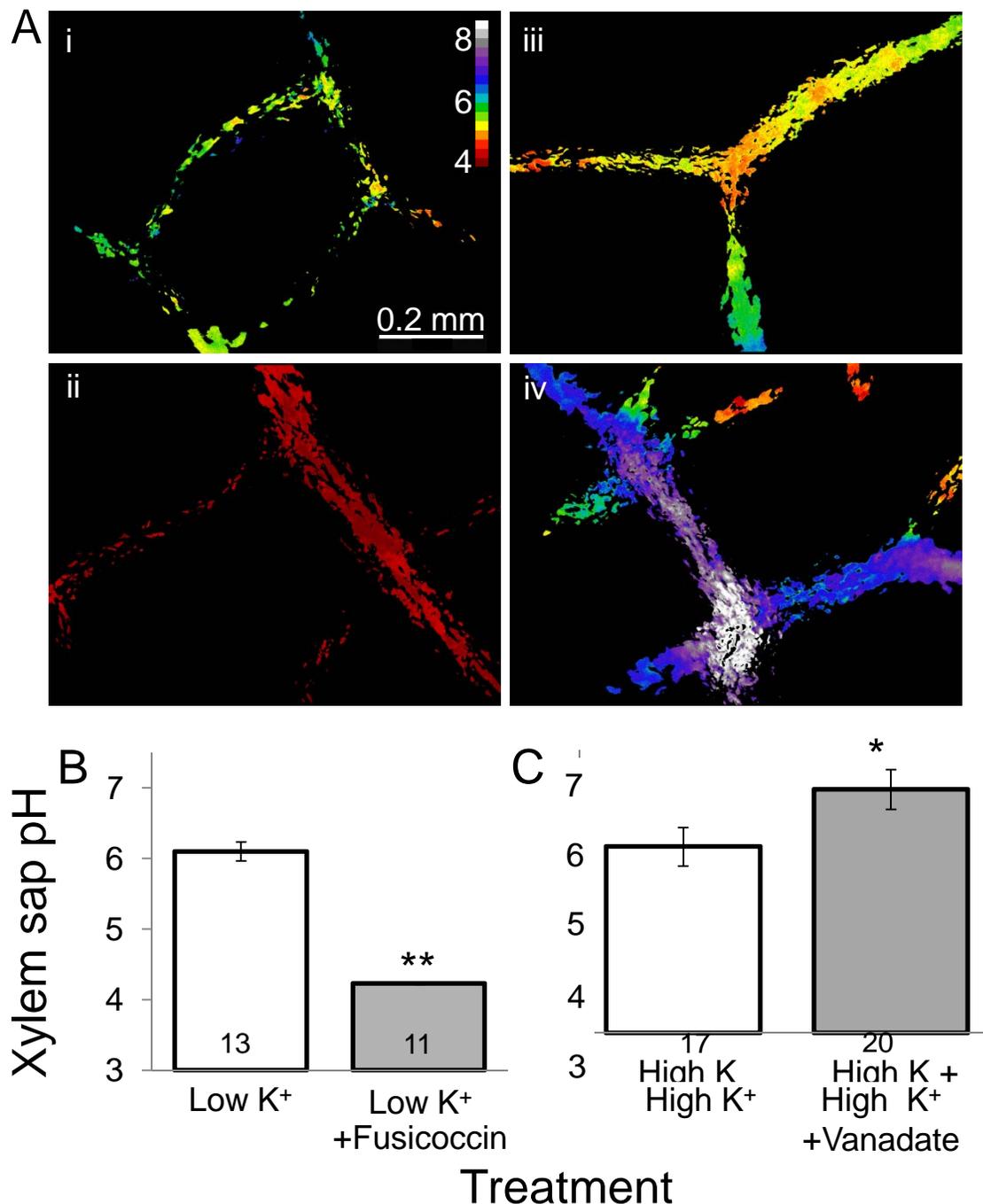


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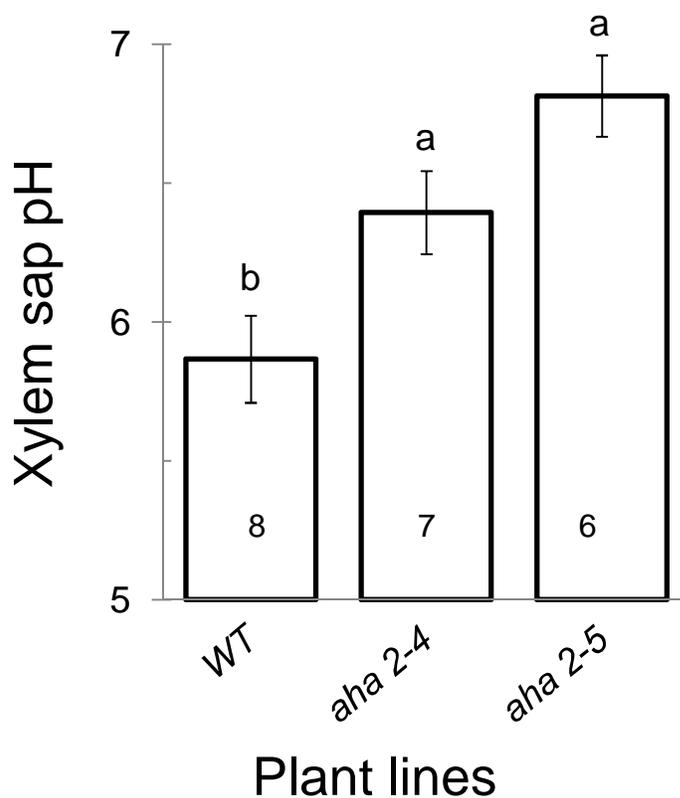


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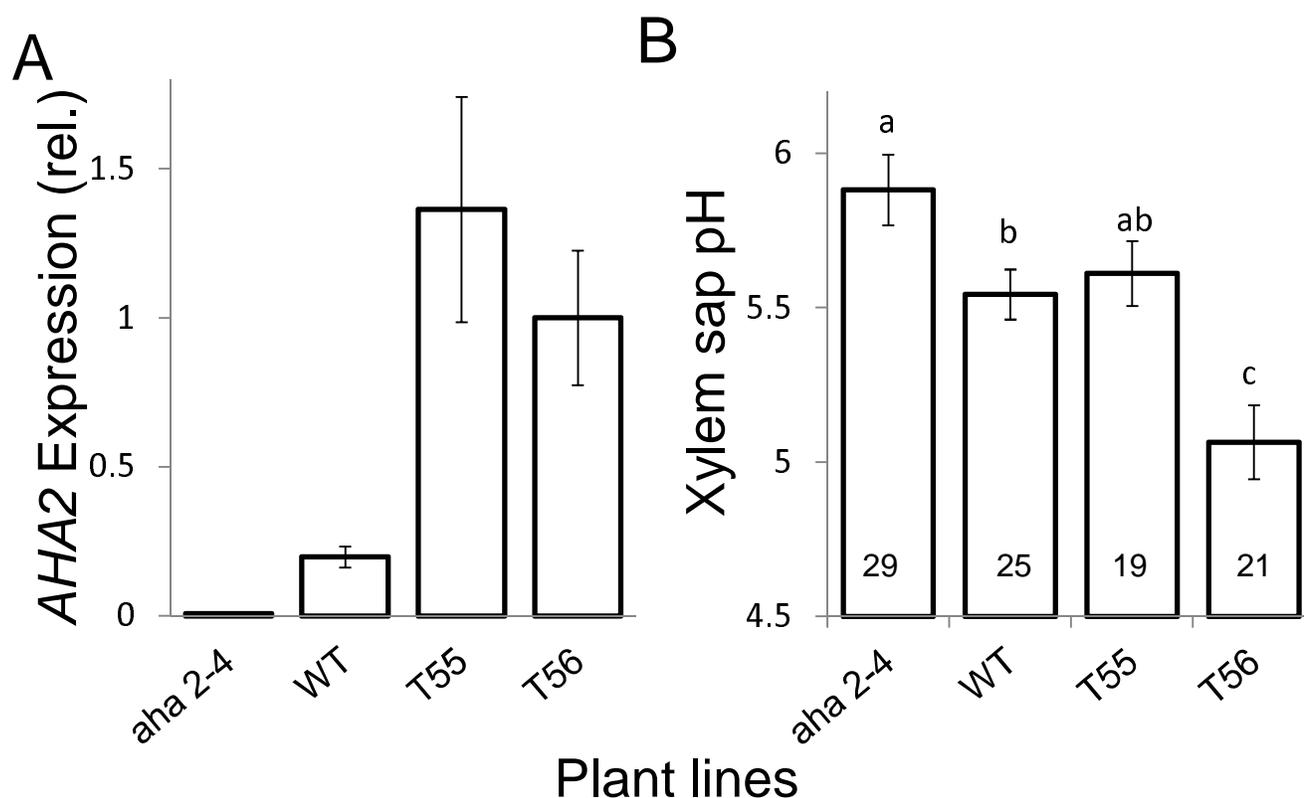


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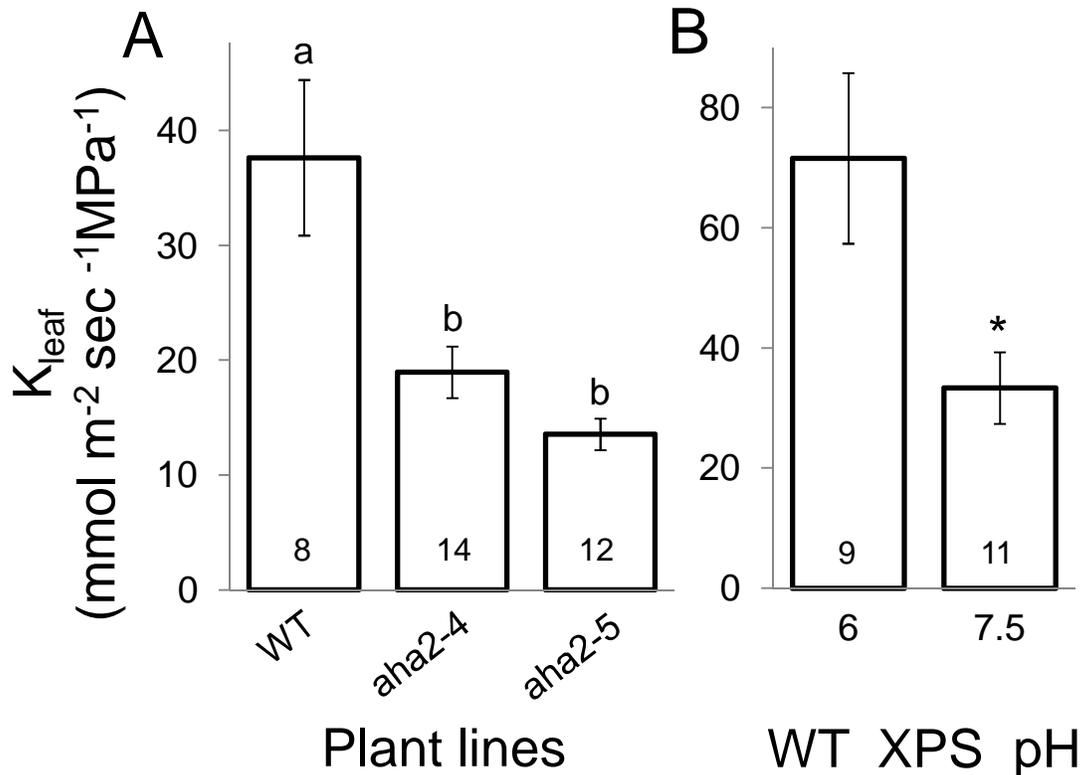


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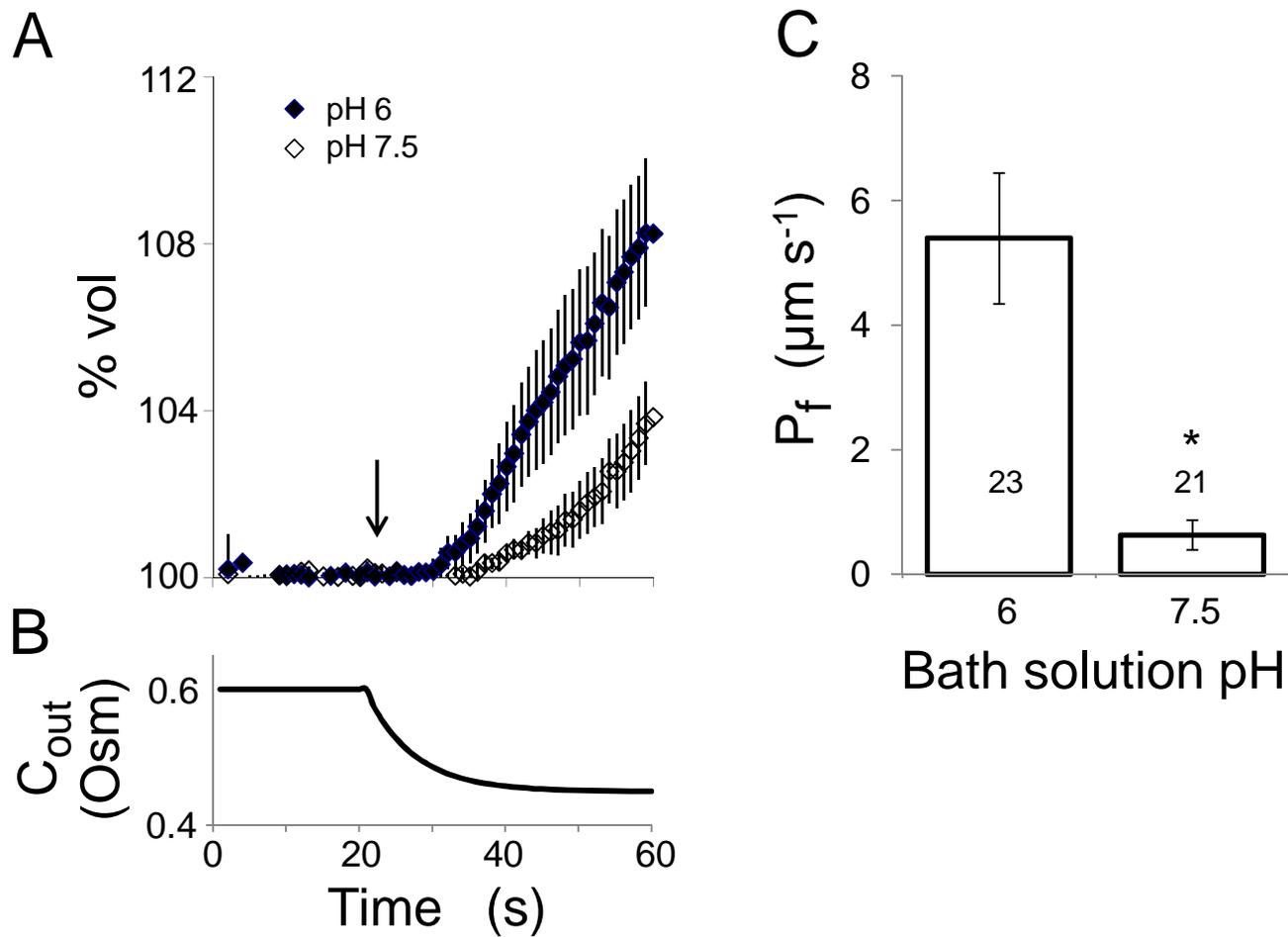


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