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Leaf hydraulic conductance varies with vein anatomy  
across *Arabidopsis thaliana* wild-type and leaf vein mutants

A thesis submitted in partial satisfaction  
of the requirements for the degree Master of Science  
in Biology

by

Marissa Anne Caringella

2015

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## ABSTRACT OF THE THESIS

Leaf hydraulic conductance varies with vein anatomy  
across *Arabidopsis thaliana* wild-type and leaf vein mutants

by

Marissa Anne Caringella

Master of Science in Biology

University of California, Los Angeles, 2016

Professor Lawren Sack, Chair

Leaf venation is diverse across plant species and has practical applications from paleobotany to modern agriculture. However, the impact of vein traits on plant performance has not been tested in a model system such as *Arabidopsis thaliana*. Previous studies analyzed cotyledons of *A. thaliana* vein mutants, and identified visible differences in their vein systems from the wild type (WT). We measured leaf hydraulic conductance ( $K_{\text{leaf}}$ ), vein traits, and xylem and mesophyll anatomy for *A. thaliana* WT and four vein mutants. Mutant true leaves did not possess the venation anomalies previously shown in cotyledons, but varied quantitatively in vein traits and leaf anatomy across genotypes. The WT had significantly higher mean  $K_{\text{leaf}}$ . Across all genotypes there was strong correlation of  $K_{\text{leaf}}$  with traits related to hydraulic conductance across the bundle sheath, as influenced by the number and radial diameter of bundle sheath cells and

vein length per area. These findings support the hypothesis that vein traits influence  $K_{\text{leaf}}$ , indicate the usefulness of this mutant system for testing theory primarily established comparatively across species, and support a strong role for the bundle sheath in influencing  $K_{\text{leaf}}$ .

The thesis of Marissa Anne Caringella is approved.

Philip Rundel

Thomas Gillespie

Lawren Sack, Committee Chair

University of California, Los Angeles

2016

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## PREFACE

This thesis is a modified version of Caringella MA, Bongers FJ, Sack L (2015) Leaf hydraulic conductance varies with vein anatomy across *Arabidopsis thaliana* wild-type and leaf vein mutants. *Plant, Cell & Environment*, doi:10.1111/pce.12584.

Co-author information: Franca J Bongers assisted with data collection and provided feedback on the manuscript, Lawren Sack mentored at all stages of the project.

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# THESIS

Leaf hydraulic conductance varies with vein anatomy  
across *Arabidopsis thaliana* wild-type and leaf vein mutants

## ABSTRACT

Leaf venation is diverse across plant species and has practical applications from paleobotany to modern agriculture. However, the impact of vein traits on plant performance has not yet been tested in a model system such as *Arabidopsis thaliana*. Previous studies analyzed cotyledons of *A. thaliana* vein mutants, and identified visible differences in their vein systems from the wild type (WT). We measured leaf hydraulic conductance ( $K_{\text{leaf}}$ ), vein traits, and xylem and mesophyll anatomy for *A. thaliana* WT (Col-0) and four vein mutants (*dot3-111* and *dot3-134*, and *cvp1-3* and *cvp2-1*). Mutant true leaves did not possess the venation anomalies previously shown in the cotyledons, but varied quantitatively in vein traits and leaf anatomy across genotypes. The WT had significantly higher mean  $K_{\text{leaf}}$ . Across all genotypes there was a strong correlation of  $K_{\text{leaf}}$  with traits related to hydraulic conductance across the bundle sheath, as influenced by the number and radial diameter of bundle sheath cells and vein length per area. These findings support the hypothesis that vein traits influence  $K_{\text{leaf}}$ , indicating the usefulness of this mutant system for testing theory that was primarily established comparatively across species, and supports a strong role for the bundle sheath in influencing  $K_{\text{leaf}}$ .

## INTRODUCTION

Across terrestrial ecosystems leaves are diverse in size, structure, and function. In particular, the leaf vein network is extremely variable across species, and angiosperms display the most diverse set of vein structures and systems (Ellis *et al.* 2009, Sack and Scoffoni 2013). Much of this diversity is linked to hydraulic design, as the leaf is a key component of the plant hydraulic system, which plays an important role in determining the maximum rate of photosynthetic gas exchange and growth (Tyree and Zimmerman 2002, Sack and Holbrook 2006, Brodribb *et al.* 2007). Leaf hydraulic conductance ( $K_{\text{leaf}}$ ) provides a measure of how efficiently water is transported through the leaf, but important questions remain about the complex pathways that water follows to sites of evaporation (Rockwell *et al.* 2014, Buckley 2014). This is especially true regarding dynamic pathways outside the xylem, where water moves across the bundle sheath (BS) and mesophyll tissue to stomata. Therefore, in addition to the xylem conduits, leaf venation architecture and mesophyll structure and cellular anatomy are integral components in the leaf hydraulic pathway. We studied leaf vein mutants of the model species *Arabidopsis thaliana* to test hypotheses for the anatomical determinants of leaf hydraulic conductance.

A number of studies conducted across a broad range of terrestrial plants have shown the importance of leaf hydraulic conductance ( $K_{\text{leaf}}$ ), i.e., the efficiency of water transport through the leaf, and indicated possible determinants of  $K_{\text{leaf}}$ . Thus, across diverse species, the maximum photosynthetic rate ( $A_{\text{max}}$ ) and stomatal conductance ( $g_s$ ) was positively correlated with  $K_{\text{leaf}}$  (Brodribb *et al.* 2007). Further, across diverse species, both  $A_{\text{max}}$ ,  $g_s$  and  $K_{\text{leaf}}$  are often positively correlated with vein traits, including vein length per leaf area (VLA or “vein density”; Sack and Frole 2006, Brodribb *et al.* 2007, Boyce *et al.* 2008, Brodribb *et al.* 2010, Sack and Scoffoni

2013). Additionally, greater xylem conduit numbers and diameters, high major VLA, more free-ending veins (FEVs) per unit area, BS anatomical traits and potentially the total area of mesophyll cells per leaf area ( $A_{mes}/A$ ) may be positively correlated with higher  $K_{leaf}$  (Brodrribb *et al.* 2007, McKown *et al.* 2010, Griffiths *et al.* 2013, Sack and Scoffoni 2013, Chatelet *et al.* 2013). However, while these correlations observed among  $K_{leaf}$  and vein traits that have been established across diverse species have also been supported by computer or physical simulations (Noblin *et al.* 2008, McKown *et al.* 2010), they have just begun to be tested across genotypes within given species. Such tests have potential to establish a stronger causality, given the assumption of a similar background of other traits. Two very recent studies focused on 4 genotypes of *Coffea arabica* (Nardini *et al.* 2014) and 11 cultivars of *Oryza* (Xiong *et al.* 2014), and both confirmed the relationship of  $K_{leaf}$  to  $g_s$  and  $A_{max}$ , but did not find a positive correlation of  $K_{leaf}$  with VLA. To our knowledge there have been no previous studies of the hydraulic properties of wild-type (WT) and leaf vein mutant genotypes of *Arabidopsis thaliana*, which provide a premier platform for testing hypotheses for trait linkages.

Knowledge about the molecular and developmental basis of leaf vein development has increased greatly by focusing on model species *A. thaliana* (Kang and Dengler 2004, Sieburth and Deyholos 2006, Carland and Nelson 2004, Petricka *et al.* 2008). As for typical angiosperm leaves, in *A. thaliana* the venation system is constructed in a hierarchy, with the first three orders known as “major veins”. In general, one or more first-order ( $1^\circ$ ) veins enter the lamina from the petiole, multiple secondary ( $2^\circ$ ) veins branch off along the length of the  $1^\circ$  vein(s), and third ( $3^\circ$ ) and higher order veins (known as “minor veins”) form a mesh throughout the lamina (Ellis *et al.* 2009). Wild-type (WT) *A. thaliana* leaves have a central mid-vein with pairs of smaller high-arching secondary veins, forming closed loops nearly reaching the margins (known as

“brochidodromous” venation; Figure 1A), enclosing a network of smaller tertiary and higher order veins (Kang and Dengler 2004). The development of leaf vein mutants with extreme phenotypes compared to the WT (Candela *et al.* 1999, Carland *et al.* 1999, Carland *et al.* 2002, Turner and Sieburth 2003, Carland and Nelson 2004, Clay and Nelson 2005, Sieburth and Deyholos 2006, Petricka *et al.* 2008, Robles *et al.* 2010) presents exciting possibilities for *A. thaliana* as a model system for leaf hydraulics. A recent study found variation among *A. thaliana* genotypes in stem hydraulic conductance and anatomy (Tixier *et al.* 2013). Here we link  $K_{\text{leaf}}$  to vein traits and comprehensive mesophyll and xylem cellular anatomy in true leaves of *A. thaliana* WT and four vein mutants.

Among vein mutants, six classes have been described (Petricka *et al.* 2008); here we focus on two mutants from the parallel class (*defectively organized tributaries, dot3*) and two mutants from the open network class (*cotyledon vascular pattern, cvp*; Table 1). Cotyledons and juvenile leaves of parallel class mutants have more monocot-like parallel venation (Petricka *et al.* 2008), while open network class mutants have unclosed 2° veins and vascular islands (Carland *et al.* 1999; see Table 1 for previously documented cotyledon/juvenile leaf vein traits). We focused on higher node (adult) leaves to determine if mutant phenotypes of cotyledons and juvenile leaves were retained, and to analyze linkage with hydraulic function.

We tested hypotheses for the relationships among venation architecture, hydraulic performance, and anatomical structure among *A. thaliana* WT and mutants. We expected that (1) adult mutant leaves would display significant quantitative differences in venation traits from the WT, and the two mutant types (*dot* and *cvp*) and venation classes (parallel and open network) would differ. (2) Given that the WT has a vein system optimized during evolution, whereas mutant cotyledons/juvenile leaves have abnormal phenotypes, the WT would have higher  $K_{\text{leaf}}$

than the mutants. Indeed, previous studies of *A. thaliana* stomatal density and patterning mutants found reduced carbon assimilation and other morphological abnormalities due to direct and/or pleiotropic effects that cause measureable changes outside the targeted trait (Dow *et al.* 2014; Lawson *et al.* 2014). Most importantly, (3) we hypothesized based on previous studies conducted across species that shifts in specific vein and anatomical traits would be associated with shifts in  $K_{\text{leaf}}$  across *A. thaliana* genotypes (hypotheses listed in Table 2).

## METHODS

### *Plant material and growth conditions*

*Arabidopsis thaliana* wild-type (WT) Col-0 (ecotype Columbia), and four venation mutants were studied. The mutants were selected based on previously documented variation in venation traits, including two mutants from two types and venation classes (Table 1); *defectively organized tributaries* mutants of the parallel class (*dot3-111* and *dot3-134*), and *cotyledon vascular pattern* mutants of the open network class (*cvp1-3* and *cvp2-1*). All mutants were originally generated from mutagenized M<sub>2</sub> seeds of the ecotype Columbia: *dot3-111* and *dot3-134* using diepoxybutane (Petricka *et al.* 2008), and *cvp1-3* and *cvp2-1* using methanesulfonate (Carland *et al.* 1999).

Seeds were sown on a 3:3:1:1 (peat moss, sandy loam, perlite, vermiculite) soil mix in a germination tray, kept in a dark chamber at 4°C for 4 days to synchronize and optimize germination, and then grown in a greenhouse at 24°C for three weeks. All plants were then transplanted to pots (d 3.8cm, h 14 cm; Cone-tainer 610-09645, Li-Cor, Lincoln, NE, USA), and placed in a growth chamber at 22°C and 70% relative humidity under a short-day light regime (9h light, 15h dark) with 200  $\mu\text{mol m}^{-2} \text{s}^{-1}$  photosynthetically active radiation (after Wagner *et al.* 2011). Plants were watered daily and were six to eight weeks old with  $\geq 20$  leaves before anatomical and hydraulic traits were assessed. We sampled fully mature, higher node (adult) leaves not yet showing signs of degradation (See Tables 2 and S1 for measured traits).

### *Leaf hydraulic conductance*

Leaf hydraulic conductance ( $K_{\text{leaf}}$ ) was determined for 10 leaves collected from 4 individuals per mutant genotype, and 11 leaves for the WT. The evaporative flux method (EFM; Sack *et al.* 2002, Sack and Scoffoni 2012) was refined for small, delicate leaves. Pots were watered to full hydration the night before and morning of measurements. Leaves were cut from shoots with a fresh razor blade under ultrapure water (0.22  $\mu\text{m}$  Thornton 200 CR, Millipore, Billerica, MA, USA), petioles were wrapped in Parafilm to ensure a seal, and then < 1 mm was cut from the end underwater to ensure a fresh surface. The petiole was then rapidly connected under water to silicone tubing which ran to a cylinder on a balance ( $\pm 10 \mu\text{g}$ , models XS205 and AB265, Mettler Toledo, Columbus, OH, USA) connected to a computer which logged data in 30 second intervals and calculated steady-state transpirational flow rate through the leaf ( $E$ ,  $\text{mmol m}^{-2} \text{s}^{-1}$ ). The flow solution used was ultrapure water (0.22  $\mu\text{m}$  Thornton 200 CR, Millipore, Billerica, MA, USA) previously degassed under vacuum for >8 h using a vacuum pump (Model DOA-P704AA, Gast, Benton Harbor, MI, USA) and then refiltered (0.2  $\mu\text{m}$ , Syringe filter, Cole-Parmer, Vernon Hills, IL, USA).

The leaf attached to the tubing was placed abaxial side down in a wooden frame strung with fishing line (for support) above a box fan (Lakewood Engineering & Manufacturing Co., Chicago, IL, USA). Photosynthetically active radiation of >1000  $\mu\text{mol m}^{-2} \text{s}^{-1}$  (LI-250 light meter, Li-Cor, Lincoln, NE, USA) was provided by floodlights (model 73828, 1000 W UV filter, Sears Roebuck, Hoffman Estates, IL, USA) suspended above the leaf surface, with a water-filled Pyrex container (Corning Inc., Corning, NY, USA) between leaf and light to absorb the heat. Leaf temperature was measured with a thermocouple (Cole-Parmer) and maintained at 23-28° C (mean temperature was  $24.6 \pm 0.083$  °C). Relative humidity was monitored with a weather



station (HOBO Micro Station with Smart Sensors, Onset, Bourne, MA, USA); mean %RH was  $40.7 \pm 0.49$ . To achieve a stable flow rate and allow the leaves to acclimate to high irradiance, leaves were allowed to transpire for at least 30 minutes or longer as necessary to achieve a coefficient of variation  $<0.05$  for at least five measurements (Scoffoni *et al.* 2012). For all leaves the mean time on the system was  $45 \pm 1.3$  minutes, with a minimum of 30 minutes and a maximum of 75 minutes. Leaves were discarded if there was any sudden change in flow rate. When flow rate stabilized, the average of the final ten flow rate measurements was recorded, and leaf temperature was measured with the thermocouple. The leaf was then immediately removed from the tubing, any excess water removed from the petiole, and placed in a Whirl-Pak bag (Whirl-Pak, Nasco, Fort Atkinson, WI, USA) that had been exhaled in for humidity. A final water potential measurement was then made for each leaf.

Given that the fragile petioles of *A. thaliana* mutants were not well suited to the pressure chamber, we used an osmometer (Vapro, model 5520, Wescor, Logan, UT, USA) to measure final water potential ( $\Psi_{\text{leaf}}$ ). The sensitive thermocouple and electronic design of the osmometer enable a relatively short equilibration time of  $\sim 1$  hour (Ball and Oosterhuis, 2005). Published comparisons of pressure chamber and psychrometer water potential measurements found that with proper insulation and equilibration (both built-in to the Vapro osmometer) the two methods yield values that are tightly correlated, often with a nearly 1:1 relationship (Oosterhuis *et al.* 1983); although psychrometer values may be more negative at higher water potentials (Barigah *et al.* 2013). A leaf disc was removed with a cork-borer, and placed abaxial side up in the osmometer chamber within 10 s. The lamina disc equilibrated for 40 min in the chamber, and 2-10 additional measurements were taken at 5-10 min intervals in *Auto Repeat* mode until the difference in values was  $<5\%$ , for a minimum of 60 min total in the osmometer. We tested the

seal of the osmometer using a solution of known osmolality that was left to equilibrate for 1 hr in the same manner as the leaf discs for water potential measurements. In three tests of each of the two Vapro osmometers (six tests total) using a 1000 mmol/kg solution, the mean measured value was  $1004 \pm 1.6$  mmol/kg (this falls within the range of acceptable error for calibration of the Vapro osmometers,  $\pm 5$  mmol/kg). The rest of the leaf was then rehydrated with the petiole under water, and leaf area was determined by scanning (flatbed scanner, Epson Perfection 4490) and analyzing the image (ImageJ).  $K_{\text{leaf}}$  was calculated as  $E/\Delta\Psi_{\text{leaf}}$ , where  $\Delta\Psi_{\text{leaf}} = -\Psi_{\text{leaf}}$  (because the water potential at the leaf petiole was 0), normalized by leaf area, and standardized to 25° C to correct for changes in  $K_{\text{leaf}}$  caused by the temperature dependence of water viscosity (Weast 1974, Yang and Tyree 1993). We note that the leaves measured for  $K_{\text{leaf}}$  by the evaporative flux method are transpiring and thus partially dehydrated when steady state flow is achieved; final leaf water potentials were observed in the range previously published for *Arabidopsis* leaves transpiring under high evaporative demand (Levin *et al.* 2007; Caldeira *et al.* 2014).

#### *Leaf clearing and vein system analysis*

We determined venation traits from one leaf from each of four individuals per genotype ( $n = 4$  for each of the five genotypes). Leaves were cleared, stained, imaged, and analyzed according to standard protocols (Berlyn and Miksche 1976, Sack *et al.* 2012). Fully hydrated leaves were fixed in FAA (70% formalin–acetic acid–alcohol, 48% ethanol:10% formalin:5% glacial acetic acid:37% water), cleared in 5% sodium hydroxide and water, and stained with safranin and fast green. Cleared leaves were mounted with water in transparency film (AF4300, 3M, St. Paul, MN, USA) and scanned (1200 dpi; flatbed scanner, Epson Perfection 4490, Long Beach, CA, USA) for quantification of leaf area and major vein lengths and diameters. Measurements of

minor veins were made from images obtained using a light microscope (DMRB, Leica Microsystems, Buffalo Grove, IL, USA) with a 5× objective and digital camera (14. 2 Color Mosaic, Diagnostic Instruments, MI, USA) utilizing SPOT advanced imaging software (SPOT Imaging Solutions, Diagnostic Instruments, Sterling Heights, MI, USA). All images were manually analyzed with Image J software (Version 1. 46r, NIH). We used sufficient magnification to ensure that all major veins were observed in the whole leaf scans (1.00 magnification, 47pixels/mm) and multiple areoles were observed in the microscopic images of the minor veins (287× magnification, 813 pixels/mm in scale, 2.84 pixels/mm in resolution; Sack *et al.* 2014).

Major vein (1°, 2°, and 3°) measurements were made for the entire leaf. We measured midrib (1°) projected area and length, and averaged three diameter measurements made in the middle of the top, center, and bottom thirds of the midrib. We measured the branching angle of the 2° veins from the midrib for the two 2° veins on either side of the midrib closest to the center of the leaf. We also measured the number of vascular strands entering the lamina from the petiole (becoming 2° veins), the number of large 2° veins, the 2° vein length, the 3° vein length, and the total major (1° + 2° + 3°) vein length per area (major VLA). From the microscope images we measured the diameters of randomly selected 2° and 3° veins, the lengths of minor veins (4° and 5°), and the number of free-ending veins (FEVs). Minor VLA and number of free ending veins per leaf area (FEV/A) were calculated by dividing the minor vein length and the number of FEVs by the microscope image size corrected for 2° vein area (area image – area 2° veins), and total VLA was calculated as the sum of major and minor vein VLA. Given that minor VLA values might be affected by excluding major veins from the images, we confirmed that results

were robust by calculating leaf-scale values using a novel calculation that corrected for the area of the leaf taken up by major veins using the formula:

$$\text{Total VLA} = 1^\circ\text{VLA} + 2^\circ\text{VLA} + 3^\circ\text{VLA} + \text{minor VLA} \times [1 - (1^\circ\text{VPAA} + 2^\circ\text{VPAA} + 3^\circ\text{VPAA})]$$

(Eqn. 1)

where 1°, 2° and 3° VPAA are respectively the vein projected areas per leaf area of the first, second, and third-order veins, determined as the product of their VLA and diameters. Values were very similar (for total VLA within 2-3%) using the two calculation methods; we present analyses in this paper based on values determined with the first method, the most commonly used in the literature (results of both methods are compared in Table S2).

#### *Anatomical measurements*

One leaf fixed in FAA from each of four individuals per genotype was sampled for cross-sectioning to determine leaf tissue, cell, and cell wall dimensions (Oguchi *et al.* 2005, Tosens *et al.* 2012). A 1 × 0.5 cm rectangle was cut from the center of each leaf, with the midrib centered lengthwise. Samples were slowly infiltrated with a mixture of ethanol and low viscosity acrylic resin (L. R. White, London Resin Co., London, UK) under vacuum, until completely infiltrated. Samples were placed in resin-filled gelatin capsules and allowed to set overnight in an oven at 55° C. Embedded samples were then sectioned in the transverse plane with glass knives (LKB 7800 KnifeMaker, LKB Produkter, Bromma, Sweden) at 1 μm thickness using a rotary microtome (Leica Ultracut E, Reichert-Jung, Arcadia, CA, USA). Cross-sections were stained on slides with 0.01% toluidine blue in 1% sodium borate and were imaged using the 20× and 40× objectives of a light microscope and camera (Leica DMRB; 14.2 Color Mosaic with SPOT advanced imaging software), and manually analyzed with ImageJ.

To comprehensively phenotype the leaf cross-sectional anatomy (Fig. 2), we measured tissue thicknesses (whole lamina, spongy and palisade mesophyll, epidermis, and cuticle), the cross-sectional areas, perimeters, and diameters of cells (spongy, palisade, BS, epidermal pavement cells, and guard cells), the cross-sectional areas and perimeters (outer and inner) of minor veins, the interveinal distance (IVD), and the adaxial and abaxial minor vein-to-epidermal distances (from BS to epidermis; VED). For measurement of tissue and cell dimensions, the lamina image was divided into three sections and each trait was averaged from a measurement made near the center of each of the three sections. Minor vein traits were averaged from two minor veins per individual; the radial diameter of the smallest, medium, and largest cell per BS were measured and the means were averaged.

We calculated traits that would contribute to the conductance of water through the BS (see Fig. 2 for images of minor veins with BS). We expected that for a given BS cell membrane conductivity, the BS conductance would increase with the number of cells arranged around the perimeter of the BS (BSC), and decline with the radial diameter of the BS cells ( $d_{bsc}$ ) given a greater distance for symplastic and apoplastic flow across the BS. We thus derived an anatomical index of minor vein BS conductance per area ( $K_{bs}'$ ;  $\text{mm}^{-2}$ ):

$$K_{bs}' = \text{BSC}/d_{bsc} \times \text{Minor VLA} \quad (\text{Eqn. 2})$$

For the midrib and minor veins, all xylem conduits were measured; conduits were considered as ellipses, and lumen area was calculated using the major and minor axes. The theoretical conductivities ( $K_x'$ ;  $\text{mmols}^{-1} \text{ m MPa}^{-1}$ ) of the major and minor veins were calculated using the Hagen-Poiseuille equation modified forellipses (Lewis and Boose 1995, Cochard *et al.* 2004, Dettmann *et al.* 2013):

$$K_x' = \frac{J}{\Delta p} = \frac{V}{t \times \Delta p} = \frac{\pi (a^3 \times b^3)}{4\eta L (a^2 + b^2)} \quad (\text{Eqn. 3})$$

where  $J$  is the rate of volume flow,  $\Delta p$  is the pressure gradient along a shoot of length  $L$ ,  $V$  is volume water,  $t$  is time,  $\eta$  is viscosity of water,  $a$  is the major axis, and  $b$  is the minor axis. Conduit conductivities were calculated individually and summed for each vein. Resulting conductivities were normalized by leaf area to generate theoretical leaf specific conductivity ( $K_x'$ ,  $\text{mmols}^{-1}\text{m}^{-1}\text{MPa}^{-1}$ ; Choat 2011).

Maximum vein-to-stomatal distance ( $D_m$ ) has been described as an anatomical trait closely correlated with VLA, and which may correlate with or determine conductance outside of the xylem (Brodribb *et al.* 2007). The  $D_m$  was measured as the hypotenuse of a right triangle bound by the interveinal distance and vein-to-epidermal distance:

$$D_m = \sqrt{(\text{IVD}/2)^2 + \text{VED}^2} \quad (\text{Eqn. 4})$$

All genotypes were amphistomatous, so  $D_{m, \text{upper}}$  and  $D_{m, \text{lower}}$  were measured separately, and because the two epidermises represent flow pathways in parallel, the overall  $D_m$  was determined as the harmonic mean of  $D_{m, \text{upper}}$  and  $D_{m, \text{lower}}$  (i. e., the inverse of the mean of their inverses).

The surface area of mesophyll cells per leaf area ( $A_{\text{mes}}/A$ ) was calculated as a measure of the cell surface available for  $\text{CO}_2$  absorption or water evaporation within the leaf (Nobel *et al.* 1975, Tosens *et al.* 2012). The  $A_{\text{mes}}/A$  was determined separately for spongy and palisade mesophyll tissue layers (spongy and palisade cells modeled as spheres and capsules respectively; Chatelet *et al.* 2013), and additionally we included a novel correction to account for minor vein vascular bundles, considering half the bundle to occur in spongy and half in palisade tissue:

$$\frac{A_{\text{mes}}}{A} = \text{SA}_{\text{cell}} \times \frac{(T_{\text{tissue}} - (\text{ASF}_{\text{tissue}} \times T_{\text{tissue}})) - (0.5 \text{CSA}_{\text{bundle}} \times \text{VLA}_{\text{minor}})}{V_{\text{cell}}} \quad (\text{Eqn. 5})$$

Where SA is surface area,  $V$  is volume, ASF is airspace fraction,  $T$  is thickness,  $\text{CSA}_{\text{bundle}}$  is cross-sectional area of the vascular bundle, and  $\text{VLA}_{\text{minor}}$  is minor vein length per area.  $A_{\text{mes}}/A$  was then calculated for the BS:

$$\frac{A_{\text{mes,bs}}}{A} = P_{\text{bs}} \times VLA_{\text{minor}} \quad (\text{Eqn. 6})$$

Where  $P_{\text{bs}}$  is the outer perimeter of the BS. The three tissues (spongy, palisade, and BS) were then summed to achieve total  $A_{\text{mes}}/A$ .

### *Statistical analysis*

We used two ANOVAs to test for differences in measured traits among genotypes. First, we tested for differences across all five genotypes in each trait with a one-way ANOVA and, when significant, we applied Tukey's post-hoc tests to resolve differences among given genotypes. Second, to test for putative differences among mutant classes we used an ANOVA with mutant genotype nested within mutant class (see Table S5 for n, F, and P-values for the two ANOVAs). For all comparisons, mean  $K_{\text{leaf}}$  was also analyzed including steady-state leaf water potential ( $\Psi_{\text{leaf}}$ ) as a cofactor, to account for  $K_{\text{leaf}}$  being dynamic with leaf water status (Blackman *et al.* 2009, Scoffoni *et al.* 2012).

Correlations were performed on raw and logged data for measured traits across all genotypes (see Table S3 for correlation matrix, Pearson's r shown).

We focused on testing only previously hypothesized relationships and thus did not generally apply a correction for multiple correlation tests, as that would have reduced the power to test *a priori* hypotheses. We thus discuss as significant (1) those differences resolved by ANOVAs for traits hypothesized *a priori* to differ with venation architecture among genotypes, including vein traits and palisade and spongy mesophyll cell layers (Wylie 1939), and (2) the significant correlations of  $K_{\text{leaf}}$  with traits expected to be influential based on previous studies (see hypotheses in Table 2). However, we provided data on a wider set of phenotypic traits to fully characterize these mutants, and for readers interested in applying a multiple test correction

for “mining” for unhyposthesized significant phenotypic differences from the ANOVA results (Table 2) we provided the significance level required by the false detection rate method to avoid the risk of inflated type I error (Benjamini and Hochberg 1995).

All analyses were performed in R version 3. 0.1 ([http://www. r-project. org/](http://www.r-project.org/)).



## RESULTS

### *Variation in venation of true leaves among mutants and WT*

The true leaves of the mutants did not show the qualitative phenotypes previously described for cotyledons. The true leaves of the parallel class mutants (*dot3-111* and *dot3-134*) did not have parallel venation, and we found no significant differences in 2° branching angle, number of 2° veins exiting the petiole, or diameter of the midrib ( $d_{\text{midrib}}$ ) between the WT and *dot* mutants (Tables 2 and S1). Similarly, the true leaves of the open network class mutants (*cvp1-3* and *cvp2-1*) did not differ from the WT in 3° or minor vein length per area (Figs. 1 and 3; Table 2). The open network class mutants did not generally have distinctively higher FEV/A; the mutant *cvp2-1* had the highest mean value, 1.59 mm<sup>2</sup>, while the other mutant of this class, *cvp1-3* had the lowest value, 0.648 mm<sup>2</sup> (Fig. 4C; Table 2).

However, we found statistically significant quantitative variation in vein traits among the genotypes. There were 1.5-fold differences across genotypes in 2° VLA and major VLA, with lowest values for the WT ( $P < 0.05$ ; Fig. 3, Table 2),  $0.711 \pm 0.0391$  mm mm<sup>2</sup> and  $1.25 \pm 0.0548$  mm mm<sup>2</sup> respectively, and highest values for *dot3-111*,  $1.09 \pm 0.0752$  mm mm<sup>2</sup> and  $1.88 \pm 0.136$  mm mm<sup>2</sup>. Mutants varied by 1.6-fold in 3° VLA ( $P < 0.05$ ; Fig 3, Table 2), with *dot3-111* again having the highest value,  $0.667 \pm 0.0794$  mm mm<sup>2</sup>. There was a 2.5-fold difference across genotypes in FEV/A ( $P < 0.05$ ; Fig. 4C, Table 2). We found no significant differences in 1°, minor, or total VLA.

### *Variation among A. thaliana vein mutants and WT in leaf hydraulic efficiency*

Mean  $K_{\text{leaf}}$  varied 3-fold across all *A. thaliana* genotypes. The mean  $K_{\text{leaf}} \pm \text{SE}$  of the WT was  $4.36 \pm 1.31 \text{ mmol m}^{-2} \text{ s}^{-1} \text{ MPa}^{-1}$ , a value close to that reported in a previous study for the *A. thaliana* WT,  $4 \text{ mmol m}^{-2} \text{ s}^{-1} \text{ MPa}^{-1}$ , using the same Columbia ecotype and EFM method, though with a pressure chamber rather than the osmometer used here (Sade *et al.* 2014). The WT  $K_{\text{leaf}}$  values arose from a mean transpiration rate  $\pm \text{SE}$  of  $3.50 \pm 0.739 \text{ mmol m}^{-2} \text{ s}^{-1}$ , and a mean water potential driving force of  $-1.02 \pm 0.0809 \text{ MPa}$ . The mean  $K_{\text{leaf}}$  of the WT was significantly greater than that of the mutants, which varied from  $1.43 \pm 0.229$  to  $2.80 \pm 0.579 \text{ mmol m}^{-2} \text{ s}^{-1} \text{ MPa}^{-1}$  for *dot3-134* and *dot3-111* respectively ( $P < 0.001$ ; Tables 2 and 3). Final water potential values were similar across genotypes, therefore differences in  $K_{\text{leaf}}$  followed the same pattern as differences in transpiration (Table S4). Among the mutants, differences were significant among genotypes but not between venation classes, indicating more variation within *dot* and *cvp* mutant types than between the types ( $P < 0.05$ ; Tables 2 and 3).

### *Differences in leaf gross structure, and mesophyll and vein anatomy*

*A. thaliana* genotypes varied significantly in aspects of leaf structure and tissue organization, though they were very similar in many mesophyll anatomy traits. Individual leaf area differed 2-fold among all genotypes ( $P < 0.001$ ), with significant differences between *cvp* and *dot* mutant types ( $P < 0.05$ ), and among all mutants ( $P < 0.01$ ; Tables 2 and 3). There was significant variation in the number of layers of spongy and palisade mesophyll cells (SCL and PCL), and in the ratio of PCL:SCL among all genotypes ( $P < 0.05$ ; Fig 4D, E, F, Table 2). Spongy mesophyll surface area per leaf area ( $A_{\text{mes, sp}}/A$ ) differed among all genotypes ( $P < 0.05$ ; Fig. 4A, Table 2). The number of conduits in the midrib ( $\text{CN}_{\text{midrib}}$ ) significantly differed by 2.4-fold among all

genotypes, and also among mutants ( $P < 0.05$ ; Fig. 4B, Table 2); the number of conduits in the minor veins ( $CN_{\text{minor}}$ ) varied by 1.6-fold among mutants ( $P < 0.05$ ; Table 2). Across all genotypes there were significant differences in the number of minor vein BS cells (BSC) and minor vein BS conductance per area ( $K_{\text{bs}'}$ ;  $P < 0.05$ ; Fig. 4G, H, Table 2).

*Correlation of mesophyll and bundle sheath traits with leaf hydraulic conductance*

The higher  $K_{\text{leaf}}$  of the WT was not related to several vein traits that had been hypothesized to be influential based on comparisons made across diverse species, including VLA,  $d_{\text{bsc}}$  and BSC (see *Introduction*; Table S3; Fig. 5A-C). However, across all genotypes we found a strong correlation of  $K_{\text{leaf}}$  with the anatomical index of minor vein BS conductance per area ( $K_{\text{bs}'} = \text{BSC}/d_{\text{bsc}} \times \text{Minor VLA}$ ;  $r = 0.92$ ,  $P < 0.05$ ; Fig. 5D).

## DISCUSSION

The true leaves of the *A. thaliana* vein mutants studied did not reflect the differences previously documented for cotyledon or juvenile leaves. However, we found strong, novel variation in leaf hydraulic function, and quantitative differences in anatomy and venation. In addition, strong correlations between  $K_{\text{leaf}}$  and the anatomy of the path of water through the bundle sheath (BS) point to this tissue as an important locus for the determination of leaf hydraulic efficiency.

### *True leaves of mutants did not retain distinct phenotypes associated with venation class*

The true leaves of *A. thaliana* did not retain the extreme phenotypes previously documented in cotyledons and juvenile leaves of the vein mutants. Contrary to hypothesis (1), the class distinctions documented for these mutants did not apply to mature leaves, and in general the variation in hydraulic performance and xylem and mesophyll anatomy was not greater between than within mutant classes. Studies for over a decade into *A. thaliana* leaf and vein network development have shown that vein pattern depends on gene expression and signal transduction pathways that can be altered (Scarpella *et al.* 2004). Thus, differential vein patterning, including the formation of free vein endings or interconnected veins, arises through alteration in the initiation of preprocambial branches, and the timing of their development into veins before mesophyll differentiation. Most models attribute mutational defects to interruptions in patterns of auxin transport during organ development (Sachs 1975, Aloni 2001). Our findings of strong differences in venation between cotyledons (as previously described) and true leaves suggest either differential gene expression affecting vein development in leaves of different ontogenetic stages, or that plasticity of the developmental system could cover up defects in the true leaves.

*Extra-xylary traits contribute to significantly higher leaf hydraulic conductance in WT*

The WT had substantially higher  $K_{\text{leaf}}$  than the vein mutants (Table 3, Fig. 5), which supported hypothesis (2). The higher  $K_{\text{leaf}}$  of the WT was found despite lack of significantly different values for traits such as total VLA or larger xylem conduit sizes and/or numbers, which we had expected to correlate positively with  $K_{\text{leaf}}$  (McKown *et al.* 2010, Sack and Scoffoni 2013). However, we found strong correlation across all genotypes between  $K_{\text{leaf}}$  and the anatomical index of conductance of water across the BS (Fig. 5D, Table S3), which supports the expectation that hydraulic function is highly dependent on BS conductance. The BS has been suggested a bottleneck to hydraulic transport in the leaf based on studies of turgor pressure and aquaporin activity in *A. thaliana* (Ache *et al.* 2010, Prado *et al.* 2013). Indeed, our findings support the proposal by Griffiths *et al.* (2013) that the importance of BS anatomy for overall hydraulic conductance of water to the mesophyll would be equal to or stronger than that of VLA.

We found that  $K_{\text{leaf}}$  was associated with the length and structure of the outside xylem path through BS cells (BS radial diameter, number of BS cells, and the relationship to minor VLA;  $K_{\text{bs}}$ , Figs. 4G, H and 5A-D), as expected if significant resistance is found in the apoplastic or symplastic flow through the BS (Cochard *et al.* 2007). The flow across membranes within the vein and across the BS would also be expected to play a role, as aquaporin-mediated water transport strongly influences  $K_{\text{leaf}}$  (Postaire *et al.* 2010). Recently, Sade *et al.* (2014) found that  $K_{\text{leaf}}$  was significantly reduced in mutants with BS aquaporin expression silenced as compared to the WT. Further, aquaporins are known to play a general role in regulation of transmembrane water transport during plant growth and stress responses (Maurel 1997), and in particular have been found to decrease BS permeability during drought (Shatil-Cohen *et al.* 2011). In addition, the degree of suberization of the BS walls may have an effect. Future work is needed to further

clarify the influence of BS anatomy on hydraulic flow pathways in *A. thaliana* and across diverse species. Indeed, our findings suggest that anatomy, along with permeability and aquaporin activation/expression per area can overshadow the differences in vein length or numbers of xylem conduits in determining  $K_{\text{leaf}}$  differences among genotypes of a species. Such a finding is consistent with, and provides a potential explanation for the findings of recent studies of genotypes of *Coffea arabica* (Nardini *et al.* 2014) and cultivars of *Oryza* (Xiong *et al.* 2014), which did not find a positive correlation of  $K_{\text{leaf}}$  with VLA.

#### *Importance of vein traits and biochemistry in determining differences in hydraulic performance*

Our study supported the influence of vein traits, and in particular the BS on  $K_{\text{leaf}}$ . We generally expect traits that vary most to have the greatest influence on  $K_{\text{leaf}}$  against a background of other traits that remain similar, and this depends on the set of plants examined (Sack & Scoffoni 2013). In *A. thaliana*, overall leaf morphology was relatively similar across the tested genotypes, yielding different results than those across diverse species. For example, in most angiosperm species minor veins account for >85% of total VLA (Sack *et al.* 2012), but in *A. thaliana* the ratio of minor:total VLA is much lower, and major veins are more important (in this study minor:total VLA ranged from 0.51 to 0.59). Thus, while across diverse species VLA is often a strong driver of  $K_{\text{leaf}}$  (Sack & Frole 2006; Brodribb *et al.* 2007), among genotypes the anatomical differences in BS traits may be a stronger influence. A general importance of the BS in determining  $K_{\text{leaf}}$  differences within a species requires confirmation, especially for species with a larger proportion of high order veins.

Further research is needed to uncover the relative roles of vein and outside xylem traits in determining  $K_{\text{leaf}}$  across a broader range of *A. thaliana* genotypes. Our work is novel in showing

strong variation in  $K_{\text{leaf}}$  and a first support for the hypothesis (3) of vein and anatomical traits in influencing hydraulic conductance across leaves for a mutant system. Our study demonstrates that structural variation in these genotypes scales up to functional consequences for the hydraulic system. Future studies can therefore clarify on one hand the genetic basis for these traits, including aquaporin genes and expression, and on the other hand, the consequences for plant performance under different resource supplies. Such detailed knowledge for a model system will confirm and extend a unified understanding of leaf venation and its influence on plant growth and adaptation. Further work on the hydraulic properties of *A. thaliana* can also clarify its role in the dynamics of growth. For example,  $K_{\text{leaf}}$  is sensitive to external factors such as environmental conditions, canopy position and time of day (Õunapuu and Sellin 2013), and the substantial variation in  $K_{\text{leaf}}$  between WT and mutants may be echoed in whole leaf performance based on evidence of the positive relationship between  $K_{\text{leaf}}$ , photosynthesis and growth (Sack and Frole 2006, Brodribb *et al.* 2007, Maherali *et al.* 2008). Thus, future studies are needed to determine to what degree variation in hydraulic traits and leaf anatomy, and in particular, BS traits, might scale up to influencing gas exchange and growth across *A. thaliana* genotypes.

#### ACKNOWLEDGEMENTS

We thank F. Carland and T. Nelson for providing seeds and advice for mutant selection and plant cultivation; W. Deng, G. John, C. Scoffoni, M. Bartlett, R. Mendez-Alonzo, A. Rowat, M. Ng, and J. Smith for logistical assistance; K. F. Hein Foundation for travel funds to F. Bongers. This work was supported by NSF grant IOS-1147292 and NSF GRFP grant DGE-1144087.

## TABLES

**Table 1.** *Arabidopsis thaliana* genotypes tested, mutant type, mutant cotyledon venation class, associated class traits (not found in mature leaves), and previous studies documenting cotyledon and juvenile leaf venation traits.

<b>Genotype</b>	<b>Mutant type</b>	<b>Venation class (cotyledon)</b>	<b>Venation class traits (cotyledon/juvenile leaf)</b>	<b>Reference</b>
Col-0	wild-type(WT) ecotype Columbia	WT	WT	Carland <i>et al.</i> 1999 Carland <i>et al.</i> 2002
<i>dot3-111</i> <i>dot3-134</i>	<i>defectively organized tributaries (dot)</i>	parallel	Monocot-like parallel venation More 2° veins exiting petiole Acute midrib to 2° branch angle Midrib same size as 2° veins	Petricka <i>et al.</i> 2008
<i>cvp1-3</i> <i>cvp2-1</i>	<i>cotyledon vascular pattern (cvp)</i>	open network	Unclosed 2° veins Disconnected 3° veins More FEV and vascular islands Fewer high order veins	Carland <i>et al.</i> 1999 Turner & Sieburth 2003 Carland & Nelson 2004



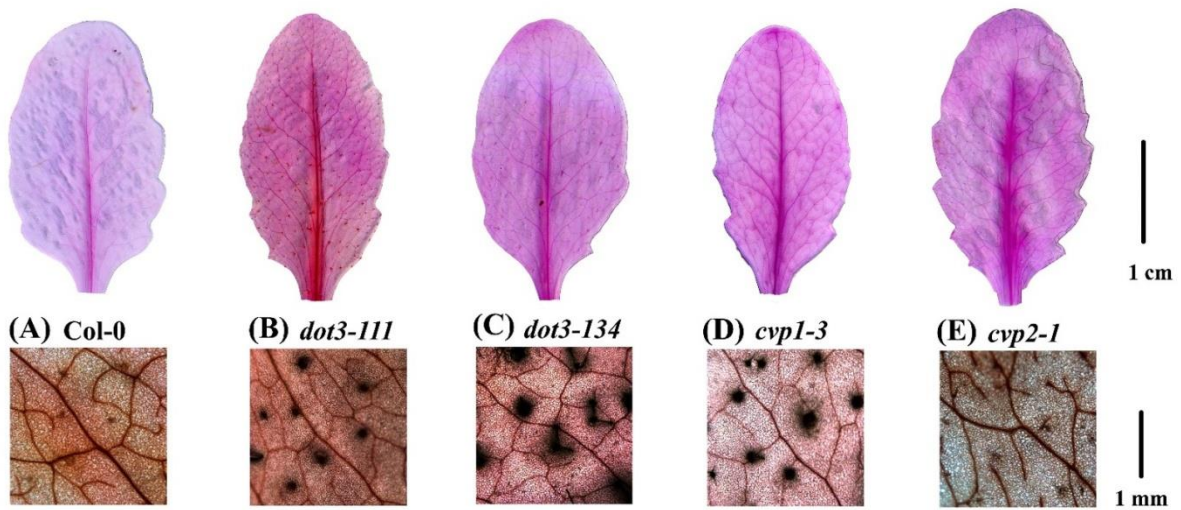
**Table 2.** Traits tested across Arabidopsis genotypes with symbols and units, followed by *a priori* hypotheses (H) for how each trait would affect  $K_{leaf}$  (i.e. as the trait increases in value will  $K_{leaf}$  increase  $\uparrow$  or decrease  $\downarrow$ ; based on previous studies comparing diverse species: Aasamaa *et al.* 2001, Aasamaa *et al.* 2005, Brodribb *et al.* 2007, Sack & Scoffoni 2013, Krober *et al.* 2014), and significance in analyses of variance. ANOVA 1 tested trait variation across all 5 genotypes; ANOVA 2 tested trait variation across mutant genotypes nested within class. See Table S5 for complete ANOVA results (including n, F, and P-values). \*  $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ . <sup>x</sup> Rendered not significant by false detection rate method (Benjamini and Hochberg 1995) applied to traits not hypothesized *a priori* to vary across genotypes differing in venation architecture (see *Methods*).  $\dagger$  indicates effect expected for amphistomatous leaves, as all 5 genotypes were.

Trait	Symbol	Units	H	ANOVA 1	ANOVA 2	
				all genotypes	Differences among mutants	Class
<b>Leaf hydraulics</b>						
Leaf hydraulic conductance (leaf area basis)	$K_{leaf}$	$\text{mmol m}^{-2} \text{s}^{-1} \text{MPa}^{-1}$		*		*
Measurement leaf water potential (covariate)	$\Psi_{leaf}$	MPa		***	***	
<b>Leaf venation architecture</b>						
Total vein length per area	VLA	$\text{mm mm}^{-2}$	$\uparrow$			
Major (1°, 2°, 3°) vein length per area	Major VLA	$\text{mm mm}^{-2}$	$\uparrow$	*		
Primary vein length per area	1° VLA	$\text{mm mm}^{-2}$	$\uparrow$			
Secondary vein length per area	2° VLA	$\text{mm mm}^{-2}$	$\uparrow$	*		
Tertiary vein length per area	3° VLA	$\text{mm mm}^{-2}$	$\uparrow$			*
Minor (4° and 5°) vein length per area	Minor VLA	$\text{mm mm}^{-2}$	$\uparrow$			
Free-ending veins per area	FEV/A	$\# \text{mm}^{-2}$	$\uparrow$	**		
Midrib diameter	$d_{midrib}$	mm	$\uparrow$			
Secondary vein diameter	$d_2^*$	mm	$\uparrow$			
Angle at which secondary branches off midrib			NA			
Total number of secondaries exiting the petiole		#	NA			
<b>Gross leaf anatomy</b>						
Leaf area	LA	$\text{cm}^2$	NA	***	* x	**
Leaf (lamina) thickness	LT	$\mu\text{m}$	$\uparrow$			
<b>Epidermal and mesophyll anatomy</b>						
Upper cuticle thickness	$T_{cut, upper}$	$\mu\text{m}$	NA	* x	**	
Lower cuticle thickness	$T_{cut, lower}$	$\mu\text{m}$	NA	* x	**	
Upper epidermal thickness	$T_{epi, upper}$	$\mu\text{m}$	NA			
Lower epidermal thickness	$T_{epi, lower}$	$\mu\text{m}$	NA	* x	* x	
Spongy mesophyll tissue thickness	$T_{spongy}$	$\mu\text{m}$	$\downarrow$			
Spongy mesophyll cell layers	SCL	#	$\downarrow$	*		
Palisade mesophyll thickness	$T_{palisade}$	$\mu\text{m}$	$\downarrow \dagger$			
Palisade mesophyll cell layers	PCL	#	$\downarrow \dagger$	*		
Ratio palisade to spongy cell layers	PCL/SCL	units cancel	$\uparrow$	*		
Upper epidermal cell cross-sectional area	$CA_{epi, upper}$	$\mu\text{m}^2$	NA			
Lower epidermal cell cross-sectional area	$CA_{epi, lower}$	$\mu\text{m}^2$	NA			
Spongy mesophyll cell cross-sectional area	$CA_{spongy}$	$\mu\text{m}^2$	NA			
Spongy mesophyll tissue percent airspace	$Air_{spongy}$	%	NA			
Palisade mesophyll cell cross-sectional area	$CA_{palisade}$	$\mu\text{m}^2$	NA			
Palisade mesophyll tissue percent airspace	$Air_{palisade}$	%	NA			
Total leaf lamina percent airspace	$Air_{leaf}$	%	NA			
Stomatal guard cell depth	GCD	$\mu\text{m}$	NA			
Bundle sheath extensions	BSE	not present	NA			
Mean maximum vein-to-stomatal distance	$D_m$	$\mu\text{m}$	$\downarrow$			
Spongy mesophyll surface area per leaf area	$A_{mes, sp} / A$	$\mu\text{m}^2 \mu\text{m}^{-2}$	$\uparrow$	*		
Palisade mesophyll surface area per leaf area	$A_{mes, pal} / A$	$\mu\text{m}^2 \mu\text{m}^{-2}$	$\uparrow$			
Bundle sheath cell surface area per leaf area	$A_{mes, bs} / A$	$\mu\text{m}^2 \mu\text{m}^{-2}$	$\uparrow$			
Total mesophyll surface area per leaf area	$A_{mes} / A$	$\mu\text{m}^2 \mu\text{m}^{-2}$	$\uparrow$			
<b>Leaf vein cross-sectional anatomy</b>						
Vascular bundle cross-sectional area	$AC_{bundle}$	$\mu\text{m}^2$	$\uparrow$			
Number of bundle sheath cells (minor veins)	BSC	#	$\uparrow$	*		**
Mean bundle sheath cell radial diameter (minor veins)	$d_{bsc}$	$\mu\text{m}$	$\downarrow$			
Anatomical index of BS conductance (minor veins)	$K_{bs}'$	$\text{mm}^{-2}$	$\uparrow$	*		*
Interveinal distance (minor veins)	IVD	$\mu\text{m}$	$\downarrow$			
Vein-to-epidermal distance, upper	$VED_{upper}$	$\mu\text{m}$	$\downarrow \dagger$			
Vein-to-epidermal distance, lower	$VED_{lower}$	$\mu\text{m}$	$\downarrow$			
Midrib number of xylem conduits	$CN_{midrib}$	#	$\uparrow$	*		*
Midrib mean maximum conduit diameter	$MCD_{midrib}$	$\mu\text{m}$	$\uparrow$			
Midrib mean conduit lumen area	$LUA_{midrib}$	$\mu\text{m}^2$	$\uparrow$			
Minor vein number of xylem conduits	$CN_{minor}$	#	$\uparrow$			*
Minor vein mean maximum conduit diameter	$MCD_{minor}$	$\mu\text{m}$	$\uparrow$			
Minor vein mean conduit lumen area	$LUA_{minor}$	$\mu\text{m}^2$	$\uparrow$			
Midrib theoretical leaf specific conductivity	$K_{x', midrib}$	$\text{mmol m}^{-1} \text{s}^{-1} \text{MPa}^{-1}$	$\uparrow$			
Minor vein theoretical leaf specific conductivity	$K_{x', minor}$	$\text{mmol m}^{-1} \text{s}^{-1} \text{MPa}^{-1}$	$\uparrow$			

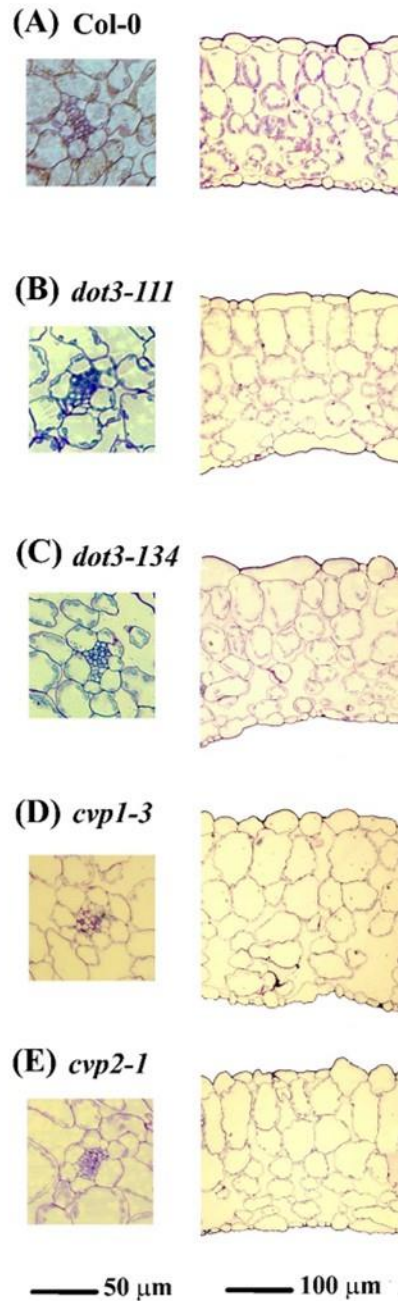
**Table 3.** Leaf morphological traits (mean  $\pm$  SE) and hydraulic conductance for true leaves of the WT (Col-0) and four venation mutants: leaf area (LA), lamina thickness (LT), total surface area of mesophyll cells per leaf area ( $A_{\text{mes}}/A$ ), total vein length per area (VLA), and mean leaf hydraulic conductance ( $K_{\text{leaf}}$ ).

Genotype	LA (cm <sup>2</sup> )	LT ( $\mu\text{m}$ )	$A_{\text{mes}}/A$ (mm <sup>2</sup> mm <sup>-2</sup> )	VLA (mm mm <sup>-2</sup> )	$K_{\text{leaf}}$ (mmol m <sup>-2</sup> s <sup>-1</sup> MPa <sup>-1</sup> )
Col-0	2.48 $\pm$ 0.271	216 $\pm$ 13.5	15.8 $\pm$ 0.366	3.04 $\pm$ 0.0800	4.36 $\pm$ 1.31
<i>dot3-111</i>	1.91 $\pm$ 0.202	233 $\pm$ 12.9	18.6 $\pm$ 1.66	3.87 $\pm$ 0.266	2.80 $\pm$ 0.579
<i>dot3-134</i>	3.52 $\pm$ 0.270	219 $\pm$ 8.92	18.6 $\pm$ 0.723	3.18 $\pm$ 0.222	1.43 $\pm$ 0.229
<i>cvp1-3</i>	3.00 $\pm$ 0.252	234 $\pm$ 15.8	18.3 $\pm$ 0.496	3.32 $\pm$ 0.343	2.60 $\pm$ 0.587
<i>cvp2-1</i>	3.57 $\pm$ 0.303	233 $\pm$ 6.80	16.7 $\pm$ 0.426	3.00 $\pm$ 0.184	1.62 $\pm$ 0.244

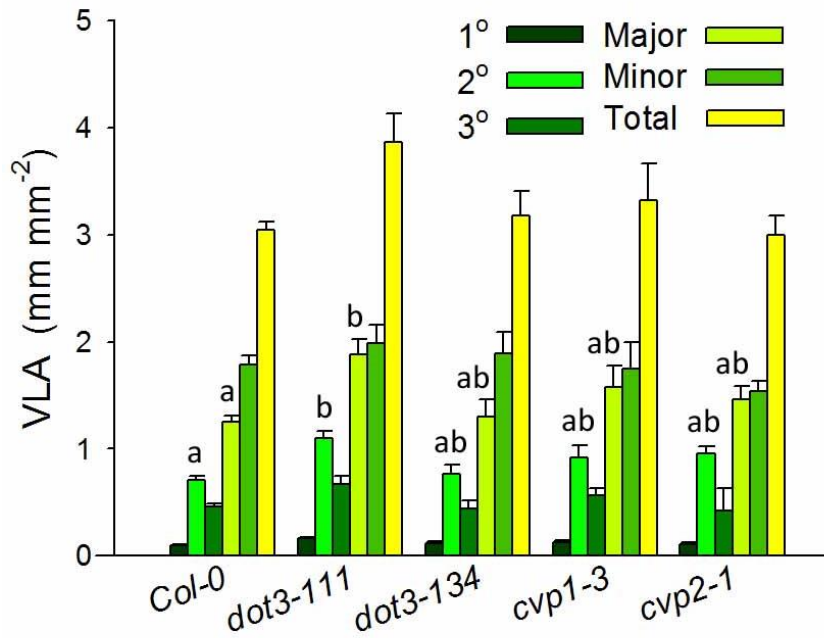
## FIGURES



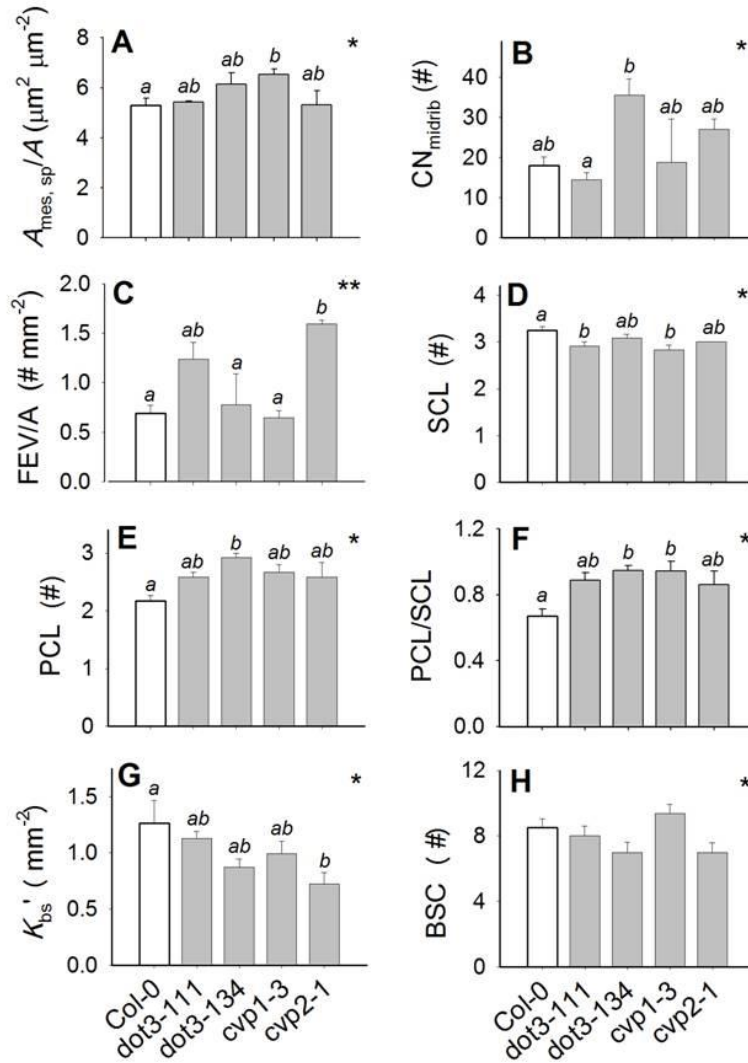
**Figure 1.** Whole cleared leaves and microscope images of cleared leaves (5×, dark spots are trichomes) for *A. thaliana* genotypes (A) WT Col-0 (B) *dot3-111* (C) *dot3-134* (D) *cvp1-3* (E) *cvp2-1*. Whole cleared leaves show that mature leaves of mutant genotypes did not retain traits seen in cotyledon classes. Microscope images display the network of 3° and minor veins branching from a central 2° vein.



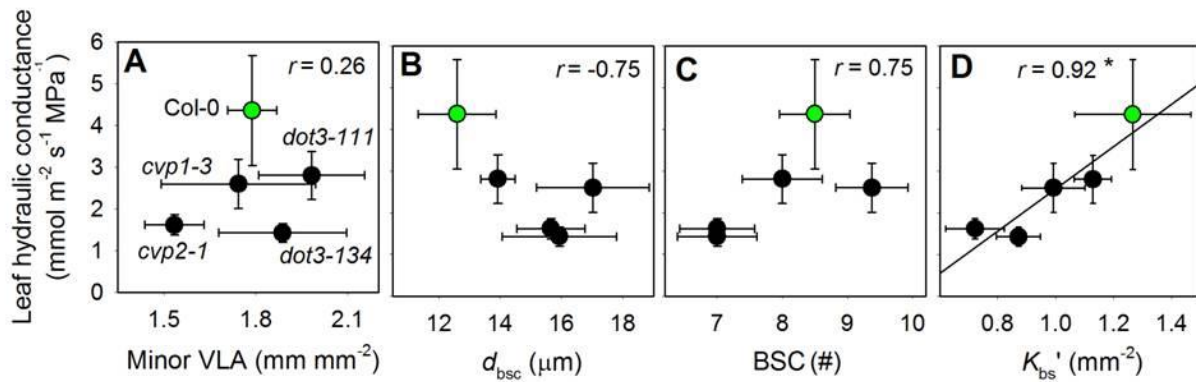
**Figure 2.** Microscope images of leaf cross-sections showing a minor vein with bundle sheath and entire lamina for *A. thaliana* genotypes (A) WT Col-0 (B) *dot3-111* (C) *dot3-134* (D) *cvp1-3* (E) *cvp2-1*. Images were all 20×, but minor veins were zoomed in to 200%. See Table 2 for significant differences in epidermal and mesophyll anatomy and leaf vein cross-sectional anatomy.



**Figure3.** Mean vein length per area (VLA)  $\pm$  SE for Col-0 (WT), *dot3-111*, *dot3-134*, *cvp1-3*, and *cvp2-1*. There were significant differences in major VLA and 2° VLA among all 5 genotypes ( $P < 0.05$ ; ANOVA 1). Differences (significant Tukey's test) indicated by different letters (a, b).



**Figure 4.** Barplots of structural and anatomical traits ( $\pm$  SE) for Col-0 (WT), *dot3-111*, *dot3-134*, *cvp1-3*, and *cvp2-1*. (A) Area spongy mesophyll per leaf area,  $A_{mes, sp}/A$  (B) number of xylem conduits in the midrib,  $CN_{midrib}$  (C) Number of free-ending veins per area, FEV/A (D) Number of spongy mesophyll cell layers, SCL (E) Number of palisade mesophyll cell layers, PCL (F) Ratio of palisade to spongy cell layers, PCL/SCL (G) Anatomical index of minor vein BS conductance per area,  $K_{bs}' = BSC/d_{bsc} \times$  Minor VLA (H) Number of BS cells (minor veins), BSC. Significant differences are shown for ANOVA 1 only (comparing all 5 genotypes) in the upper right hand corner of plots; \*  $P < 0.05$ , \*\*  $P < 0.01$ . Differences between genotypes (significant Tukey's test) are indicated by different letters (a, b).



**Figure 5.** Correlations between  $K_{leaf}$  and (A) Minor VLA (B)  $d_{bsc}$  (C) BSC, to illustrate how each component contributes to the anatomical index of minor vein bundle sheath conductance per area,  $K_{bs}'$  (D). Pearson's  $r$  shown (Table S3); \*  $P < 0.05$ .

## SUPPLEMENTARY TABLE DESCRIPTIONS

All supplementary tables can be found in the thesis supplementary data excel spreadsheet file.

**Table S1.** Mean and SE for measured traits

**Table S2.** Comparison of minor VLA, total VLA, and FEV/A means  $\pm$  SE using method 1 and method 2 (corrected for VPAA 1°, 2°, 3°).

**Table S3.** Correlation matrix for the 5 genotypes (Col-O, dot3-111, dot3-134, cvp1-3, cvp2-1); correlations for untransformed and log-transformed data shown with significance.

**Table S4.** Mean  $\pm$  SE Kleaf and corresponding E (transpiration) and final  $\Psi_{\text{leaf}}$  (leaf water potential) values measured for the WT and four mutants.

**Table S5.** ANOVA results comparing among all 5 genotypes, and among 4 mutant genotypes nested in class/type (dot mutants are parallel class and cvp mutants are open network class).



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