

# Scaling between stomatal size and density in forest plants

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The size and density of stomatal pores limit the maximum rate of leaf carbon gain and water loss ( $g_{\max}$ ) in land plants. Stomatal size and density are negatively correlated at broad phylogenetic scales, such that species with small stomata tend to have greater stomatal density, but the consequences of this relationship for leaf function has been controversial. The prevailing hypothesis posits that the negative scaling of size and density arises because species that evolved higher  $g_{\max}$  also achieved reduced allocation of epidermal area to stomata (stomatal-area minimization). Alternatively, the negative scaling of size and density might reflect the maintenance of a stable mean and variance in  $g_{\max}$  despite variation in stomatal size and density, which would result in a higher allocation of epidermal area to achieve high  $g_{\max}$  (stomatal-area increase). Here, we tested these hypotheses by comparing their predictions for the structure of the covariance of stomatal size and density across species, applying macroevolutionary models and phylogenetic regression to data for 2408 species of angiosperms, gymnosperms, and ferns from forests worldwide. The observed stomatal size-density scaling and covariance supported the stomatal area increase hypothesis for high  $g_{\max}$ . Thus, contrary to the prevailing view, higher  $g_{\max}$  is not achieved while minimizing stomatal area allocation but requires increasing epidermal area allocated to stomata. Understanding of optimal stomatal conductance, photosynthesis, and plant water-use efficiency used in Earth System and crop productivity models will thus be improved by including the cost of higher  $g_{\max}$  both in construction cost of stomata and opportunity cost in epidermal space.

Stomatal pores are critical determinants of the function of plants and the composition of the atmosphere<sup>1</sup>. The stomatal conductance to diffusion of water vapor and CO<sub>2</sub> ( $g_s$ ) influences a broad spectrum of ecological processes at leaf, community, and ecosystem scales, including photosynthesis, net primary production, and water use efficiency<sup>2,3</sup>. Theoretically, stomata can regulate  $g_s$  through evolutionary or plastic shifts in stomatal size or numbers<sup>4</sup> or through short-term stomatal aperture changes<sup>5</sup>. The  $g_s$ , and its typical operational value ( $g_{op}$ ), can thus vary from near zero with stomata fully closed and  $g_{max}$  with stomata fully open. However,  $g_{op}$  and  $g_{max}$  are correlated across plants measured under controlled conditions<sup>6,7</sup>, possibly to maintain the ratio  $g_{op}:g_{max}$  in which stomatal pore aperture is most responsive to guard cell turgor pressure<sup>8</sup>. There has been substantial debate about whether to increase  $g_{max}$ , plants must allocate a greater fraction of limited epidermal area to stomata, which might be costly. Stomatal optimization used to predict water use, productivity, and drought responses in critical global vegetation and crop models could be sensitive to costs of stomata, but they are rarely included<sup>9</sup>. Because of their importance in controlling leaf water and CO<sub>2</sub> fluxes, stomatal anatomy could be integrated with other plant traits to inform global vegetation and crop models<sup>10-13</sup> if we better understood what drives variation in these traits. Indeed, the anatomical traits stomatal density ( $D_s$ , number of pores per unit epidermal area) and size ( $A_s$ , area of guard cells surrounding each pore) have been widely used to study the adaptation and competition of plants because they are reliable indicators of  $g_{max}$  to water vapor and CO<sub>2</sub><sup>14-20</sup>. Yet, while the covariation of stomatal size and density within and across species has emerged as a critical constraint on  $g_{max}$  in controlled experiments and specific ecosystems<sup>21,22</sup>, these traits have engendered controversy when considering the diversity of species in natural vegetation at regional or global scales.

Variation in stomatal size and density may be constrained to minimize leaf surface area (stomatal-area minimization) or leaves may allocate more area to increase stomatal conductance (stomatal-area increase). We adjudicate between these competing hypotheses by considering how they affect stomatal trait covariance differently and test competing predictions with a global data set of stomatal anatomy in forest plants. A leaf's  $g_{\max}$  is determined by stomatal anatomy:

$$g_{\max} = bmD_sA_s^{0.5}, \quad (1)$$

where  $b$  and  $m$  are biophysical and morphological constants, respectively<sup>23</sup> (see Methods for equations to calculate these constants). The fraction of epidermal area allocated to stomata ( $f_s$ ) is a fundamental physical constraint on  $g_{\max}$  because  $f_s$  cannot exceed unity, and

$$f_s = D_sA_s, \quad (2)$$

Therefore,  $D_s$  and  $A_s$  define both  $g_{\max}$  and  $f_s$ , two related traits that affect gas exchange, productivity, and fitness.

Stomatal size and density covary negatively across species, reducing the variation in  $g_{\max}$  and  $f_s$  compared to what it would be if these anatomical traits were uncorrelated<sup>4,22–27</sup>. Despite that the inverse relationship between stomatal size and density has been recognized since 1865<sup>24</sup>, its evolutionary origin and its consequences have not been understood. The leading view, established by Franks and Beerling<sup>25</sup>, is that the size-density relationship arises due to the minimization of the area of epidermis allocated to stomata, such that the evolution of more stomata involves the reduction of stomatal size. By Eq. 2, a higher  $g_{\max}$  can be achieved with a smaller total stomatal area by increasing stomatal number and reducing stomatal size, because smaller stomata also have a shorter channel for diffusion. Thus, selection for higher  $g_{\max}$  would result in more numerous, smaller stomata, to minimize epidermal allocation to stomata. However, this hypothesis, and its implications that epidermal allocation to stomata is minimized,

and can be strongly reduced, have not been tested. Such reduction of allocation would greatly reduce the construction and maintenance cost of guard cells, and also the opportunity cost in other epidermal leaf structures, and in light penetration to the mesophyll. A counter-hypothesis to the ‘stomatal-area minimization’ is the ‘stomatal-area increase’ hypothesis in which selection for higher  $g_{\max}$  would lead to greater surface allocation, and thus incur substantial cost (Fig. 1). It is critical to determine which of these hypotheses is consistent with the observed stomatal size-density relationship. An implication of stomatal-area minimization is that extremely high  $f_s$  is rare because of costs associated with allocating too much epidermal area to stomata<sup>27–32</sup>, and  $g_{\max}$  is ultimately constrained by the costs of high  $f_s$ . By contrast, an implication of the stomatal-area increase hypothesis is that on average high  $g_{\max}$  incur major costs in stomatal volume construction cost and in epidermal space.

Both the stomatal-area minimization and stomatal increase hypotheses for high  $g_{\max}$  would be consistent with an inverse stomatal size-density relationship. We developed theory showing that the scaling exponents of the relationship would differ between the hypotheses using a new, additional criterion based on models of macroevolutionary landscapes<sup>26–29</sup> originally derived from quantitative genetics. Over macroevolutionary time, variation in phenotypic traits is usually constrained by some combination of physical, developmental, or ecological factors. Theoretically, this process can be described by an Ornstein-Uhlenbeck (OU) model in which trait variation expands through time until it reaches a stationary distribution around a long-term average<sup>27</sup>. In macroevolutionary models, the OU process describes the movement of the adaptive optima itself. Species should be close to their current adaptive optimum if there is sufficient genetic variation, so phenotypic constraint is caused by limits on movement in the adaptive optima, not limits on response to selection.

We propose that scaling between stomatal size and density does not reflect constraints on either trait alone, but rather the constraint on a composite trait associated with stomatal size and density. That is, if the composite traits  $f_S$  and/or  $g_{\max}$  were constrained following an OU process then constituent traits (size and density) should *appear to evolve* as if they are mutually constrained. This prediction is based on quantitative genetic theory which shows that a constraint on composite traits imposed by stabilizing selection limits variation in constituent traits<sup>30</sup>. We tested whether the observed size-density scaling across 2408 species was consistent with a stronger constraint on  $f_S$  or  $g_{\max}$ . Note that both  $f_S$  and  $g_{\max}$  equations have the form:

$$Z_S = \lambda D_S A_S^\beta, \quad (3)$$

where  $Z_S$  is a composite stomatal trait that is proportional to the product of constituent traits, stomatal density and stomatal size with scaling exponent  $\beta$  multiplied by a scalar  $\lambda$ . For  $g_{\max}$ ,  $\lambda = bm$  and  $\beta = 0.5$  (Eq. 1); for  $f_S$ ,  $\lambda = 1$  and  $\beta = 1$  (Eq. 2). Next, we linearize Eq. 3 by log-transformation:

$$z_S = \log(\lambda) + d_S + \beta a_S. \quad (4)$$

Lowercase variables are the log-transformed versions of their uppercase counterparts. Log-transformation is also desirable here because  $D_S$  and  $A_S$  are log-normally distributed<sup>27</sup>. Using random variable algebra, the variance in  $z_S$  is defined as:

$$\text{Var}(z_S) = \text{Var}(d_S) + \beta^2 \text{Var}(a_S) + 2\beta \text{Cov}(d_S, a_S). \quad (5)$$

The scaling exponent  $\beta$  minimizes  $\text{Var}(z_S)$ , the most constrained composite trait, is:

$$\beta = -\frac{\text{Cov}(d_S, a_S)}{\text{Var}(d_S)} \quad (6)$$

The right-hand side of this equation is the negative of the ordinary linear regression slope of log-stomatal size against log-density, which means that  $\beta$  can be estimated using regression methods, but a negative relationship will result in positive value of  $\beta$ . The stomatal-area

minimization hypothesis predicts that  $\hat{\beta} = 1$  whereas the stomatal-area increase hypothesis predicts that  $\hat{\beta} = 0.5$ . Forward-time, individual based, macroevolutionary quantitative genetic simulations confirm that these predictions hold true regardless of underlying assumptions about mutational and genetic variance (Supplementary Information).

We estimated stomatal size-density scaling in 2408 forest plant species from new field-collected samples over 28 sites in China and global synthesis of data from the literature (Fig. 2) and estimated the scaling exponent  $\beta$  using OU phylogenetic multiple regression with group (Angiosperm, Pteridophyte, Gymnosperm) and growth form (tree, shrub, herb) as covariates (see Methods)

Our analysis shows that stomatal size-density scaling among forest plant species is consistent with selection for higher  $g_{\max}$  increasing stomatal area allocation, and not minimizing area allocation (Fig. 3). Given the variance in stomatal density, the covariance between size and density among forest species minimizes the variance in  $g_{\max}$ . There is no evidence that scaling differs between major groups, Angiosperms, Gymnosperms, and Pteridophytes (Fig. 3a; Table S1), but  $g_{\max}$  is 49% (17-88% 95% CI;  $P = 0.001$ ) and 14% (1-30% 95% CI;  $P = 0.04$ ) higher in Angiosperms than Gymnosperms and Pteridophytes, respectively (Table S2). Trees also have 18% (8-28% 95% CI;  $P < 0.0001$ ) and 48% (39-59% 95% CI;  $P < 0.0001$ ) greater  $g_{\max}$  than shrubs and herbs, respectively (Table S2). The mean and variance in  $\log(g_{\max})$  are nearly invariant across latitude, temperature, and precipitation gradients, indicating that most of the variation in  $g_{\max}$  occurs within rather than between forest sites (Fig. 4).

Our results overturn the prevailing view that size-density scaling results in minimized stomatal area allocation. Instead, we find that negative covariance between stomatal size and density is consistent with stomatal area allocation increasing with  $g_{\max}$ , and, furthermore that

limits on the fraction of epidermis allocated to stomatal ( $f_s$ ) are a secondary consequence of limits on  $g_{\max}$ . Our results differ from previous studies because we adopted a new theoretical framework based on quantitative genetic and macroevolutionary theory. Previous analyses<sup>31</sup> estimated scaling exponents close to 1 for hypostomatous leaves using standardized major axis (SMA) regression, which finds the scaling exponent which minimizes residual variance in both  $d_s$  and  $a_s$ . SMA regression, adopted from allometry, does not enable analysis of how stomatal size-density scaling shapes covariance between constituent traits (see Supplementary Information). Although estimated scaling using standard phylogenetic regression approaches (see Methods), it is more appropriate to interpret the results not as minimizing residual variance, but rather estimating the  $\beta$  consistent with the covariance structure of stomatal size and density (Fig. 1).

Our results have at least three important implications for understanding the evolutionary consequences of the stomatal size-density scaling relationship. First, size-density scaling implies that extreme values of  $g_{\max}$  do not provide competitive advantage across environments. Low  $g_{\max}$  limits plants productivity by reducing the  $\text{CO}_2$  supply for photosynthesis under favorable conditions. Excessively high  $g_{\max}$  may be rare because selection on  $g_{\max}$  most likely ensures a proper  $g_{\text{op}}:g_{\max}$  ratio that situates leaf  $g_{\text{op}}$  in a region of maximal control to respond rapidly to changing environments<sup>8</sup>. High anatomical  $g_{\max}$  may also increase the risk of hydraulic failure<sup>32</sup> especially under transient conditions, thereby setting a physical upper limit to leaf gas exchange. Other possible costs include interference of stomatal movements and diffusion, as well as energetic costs<sup>31,33</sup>. The scaling between size and density is consistent with excessive  $g_{\max}$  incurring a major cost that cannot be obviated by stomatal area-minimization. Future work should prioritize identifying the fitness costs and evolutionary constraints that prevent higher



$g_{\max}$  from evolving. Second, if  $g_{\max}$  is the primary constraint, this implies that plants could allocate a greater fraction of their epidermal area to stomata than they currently do without incurring a major cost. For example, we predict that if stomatal size and density could be manipulated independently, anatomies with the same  $g_{\max}$ , but different  $f_s$ , would have similar fitness. Another important corollary is that smaller stomata were not directly required to increase  $g_{\max}$  in angiosperms. All three major land plant lineages have similar variance in  $g_{\max}$  (Fig. 3b); gymnosperms and pteridophytes have lower average  $g_{\max}$  because they have on average a lower  $d_s$  for a given  $a_s$ , not because of differences in the scaling relationship. Increased  $g_{\text{op}}$  may be an indirect consequence of higher stomatal density being linked to increases in leaf water transport capacity, for example, by decreasing the distance between vein and stomata, allowing stomata to stay open<sup>40</sup>. The primary constraint on maximum stomatal conductance appears to be selection itself, implying that vegetation and crop models should incorporate costs of extreme trait values in their predictions.

# Methods

## *Stomatal trait data from global forests*

The stomatal dataset of global forests involved the information of a total of 2408 plant species from natural forests. The data consists of novel field data collected from Chinese forest communities and a compilation of published literature values.

Our field data were collected from 28 typical forest communities occurring between 18.7 °N and 53.3 °N latitude in China. The field sites were selected to cover most of the forest types in the northern hemisphere, including cold-temperate coniferous forest, temperate deciduous forest, subtropical evergreen forest, and tropical rain forest (Fig. 2). In total, we sampled 28 forest sites. We used the Worldclim database <sup>34</sup> to extract additional data on mean annual temperature (MAT) and precipitation (MAP) over the period 1960-1990 using latitude and longitude. Among these forests, mean annual temperature (MAT) ranged from -5.5-23.2 °C, and mean annual precipitation (MAP) varied from 320 to 2266 mm. The field investigation was conducted in July-August, during the peak period of growth for forests. Sampling plots were located within well-protected national nature reserves or long-term monitoring plots of field ecological stations, with relatively continuous vegetation. Four experimental plots (30 × 40 m) were established in each forest.

Leaves from trees, shrubs, and herbs were collected within and around each plot. For trees, mature leaves were collected from the top of the canopy in four healthy trees and mixed as a composite sample. Eight to 10 leaves from the pooled samples were cut into roughly 1.0 × 0.5 cm pieces along the main vein, and were fixed in formalin-aceto-alcohol (FAA) solution (5 ml 38 % formalin, 90 ml 75 % ethanol, 5 ml 100 % glacial acetic acid, and 5 ml 37 % methanol) <sup>35</sup>.

In the laboratory, three small pieces were randomly sampled, and each replicate was photographed twice using a scanning electron microscopy (Hitachi SN-3400, Hitachi, Tokyo, Japan) on the lower surface at different positions. We focused on the lower epidermis<sup>22</sup>, because a previous study has demonstrated that most of leaf upper epidermis has no stomata for forest plants<sup>36</sup>.

In each photograph, the number of stomata was recorded, and  $D_s$  was calculated as the number of stomata per unit leaf area. Simultaneously, five typical stomata were selected to measure stomatal size using an electronic image analysis equipment (MIPS software, Optical Instrument Co. Ltd., Chongqing, China).

Peer-reviewed papers on leaf stomata were collected using an all-databases search of Web of Science ([www.webofknowledge.com](http://www.webofknowledge.com)) from 1900 to 2018 using “forest” and “stomata” as a topic, in line with the principle of “nature forest, non-intervention, species name”. There were a total of 90 papers (see Supporting Table S3) which met our requirements, yielding  $D_s$  and  $L$  measurements from 413 plant species (Fig. 2) from which we calculated  $g_{\max}$  and  $f_s$ .  $f_s$  is proportional to the stomatal pore area index (SPI), which defined as the product of  $D_s$  and stomatal length ( $L$ ) squared<sup>37</sup>, because  $A_s = mL^2$ <sup>23</sup>.

We calculated  $g_{\max}$  (Equation 1) to water vapor at a reference leaf temperature ( $T_{\text{leaf}} = 25^\circ$  C) following Sack and Buckley<sup>23</sup>. They defined a biophysical and morphological constant as:

$$b = \frac{D_{\text{wv}}}{v}$$

$$m = \frac{\pi c^2}{j^{0.5}(4hj + \pi c)}$$

$b$  is the diffusion coefficient of water vapor in air ( $D_{wv}$ ) divided by the kinematic viscosity of dry air ( $\nu$ ).  $D_{wv} = 2.49 \times 10^{-5} \text{ m}^2 \text{ s}^{-1}$  and  $\nu = 2.24 \times 10^{-2} \text{ m}^3 \text{ mol}^{-1}$  at  $25^\circ\text{C}$ <sup>38</sup>. For kidney-shaped guard cells,  $c = h = j = 0.5$ ; for dumbbell-shaped guard cells in the Poaceae,  $c = h = 0.5$  and  $j = 0.125$ . We used the species average  $g_{\max}$  and  $f_s$  for all analyses.

### *Phylogenetic regression*

By positing that the least variable composite of stomatal size and density indicates the trait with the most constraint (Fig. 1), we identify a new way to estimate the scaling exponent  $\beta$  (Eq. 6) using linear regression estimates, but these estimates should still account for phylogenetic nonindependence. We used the Plant List (<http://www.theplantlist.org>) to confirm species names, then we assembled a synthetic phylogeny using S.PhyloMaker<sup>39</sup>. We fit phylogenetic regression models using the **phylolm** version 2.6 package in R<sup>40</sup>. As we derived in the main text, the scaling exponent  $\beta$  can be estimated from the slope of the regression of  $a_s$  on  $d_s$ , where  $\hat{\beta} = -\text{slope}$ . We estimated separate scaling exponents for major groups, Angiosperms, Pteridophytes, and Gymnosperms. We also estimated different intercepts, corresponding with different average  $g_{\max}$  values, for functional types (herbs, shrubs, and trees) and grasses, because of their unique stomatal anatomy. We used the “OUrandomRoot” model of trait evolution. 95% confidence intervals for all parameters were estimated from 1000 parametric bootstrap samples generated by simulating from the best-fit model and re-fitting.  $P$ -values for coefficients are based on  $t$ -tests. We used the same methods to test whether  $g_{\max}$  (log-transformed for homoskedasticity) was affected by  $|\text{latitude}|$ , MAP, MAT, group (Angiosperms, Pteridophytes, Gymnosperms), and/or functional type (herb, shrub, tree). One gymnosperm species, *Torreya fargesii*, had substantially lower stomatal size than would be predicted from its density (Fig. 3a). These results of the paper did not change if this outlier was excluded because the confidence intervals for stomatal-density

scaling are very wide for Gymnosperms regardless. Therefore, we excluded this species from statistical analyses but show it in the figure for completeness. All data were analyzed in R

<sup>41</sup>version 4.0.5

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## Acknowledgements

Financial support was supported by the National Natural Science Foundation of China (31988102, 42071303), the National Key R&D Program of China (2017YFA0604803), National Science and Technology Basic Resources Survey Program of China (2019FY101300) , US National Science Foundation 1929167 (to CDM), and the Project funded by China Postdoctoral Science Foundation (2020M680663). We thank “Functional Trait Database of Terrestrial Ecosystems in China (China\_Trait)” for sharing data, further information for other materials should contact to N.P. He ([henp@igsnr.ac.cn](mailto:henp@igsnr.ac.cn)). There are no conflicts of interest to declare.

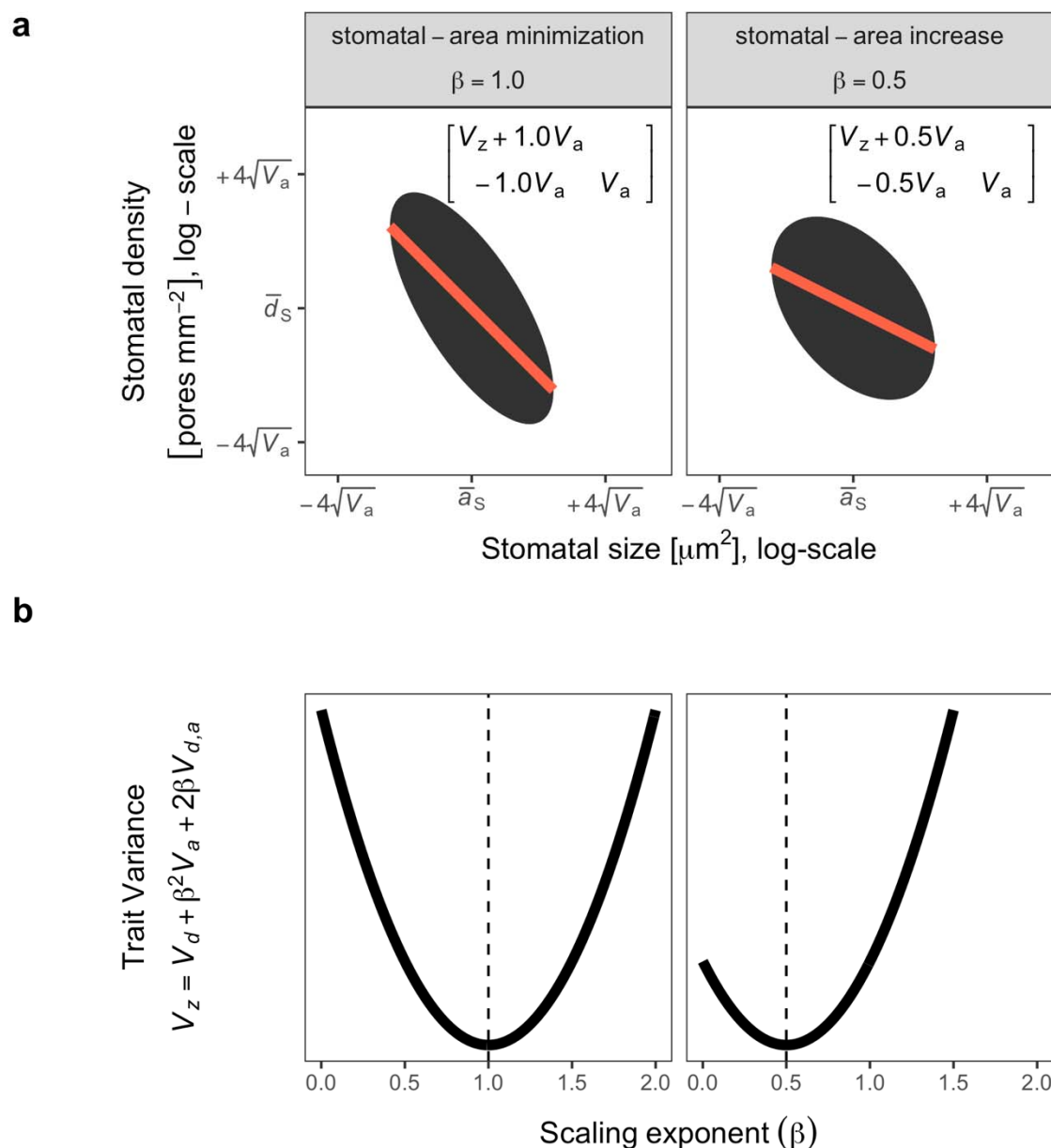
## Author contributions

N.H. and G.Y. designed field sampling; N.H., C.D.M, and C.L. conceived the initial ideas; C.L., N.H., Y.L. J.Z., M.L. and L.X collected the data; C.L. writed the first draft, and C.D.M. made major revision on it, and C.D.M. contributed the final mathematical derivations, data analysis, and wrote the final manuscript; L.S., H.J.B., C.L., N.H., G.Y., and X.H. revised the manuscript. All authors gave final approval for publication.



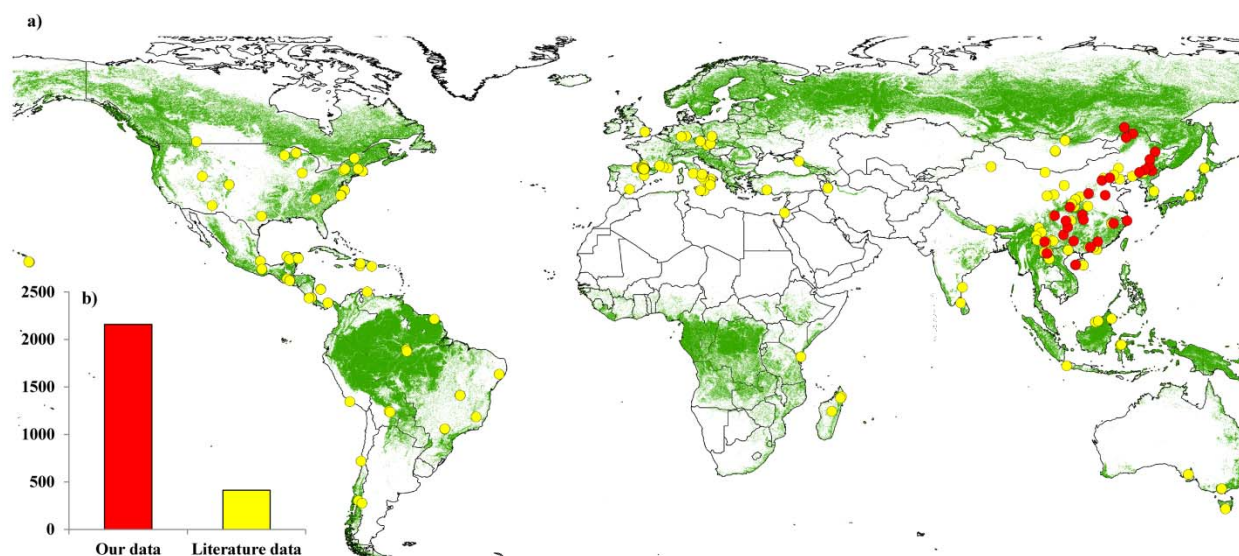
# Supplementary information

## Figures

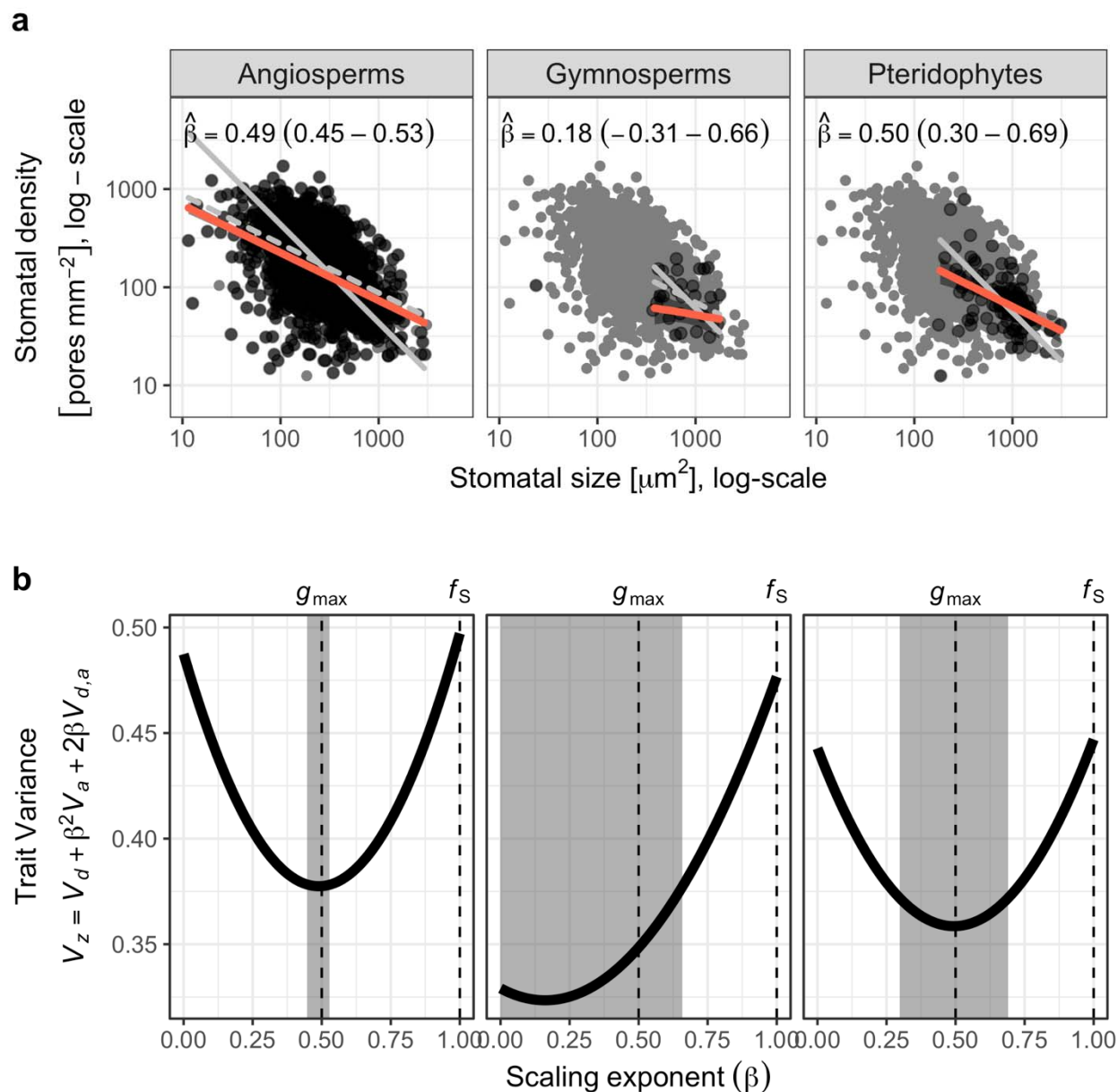


**Fig. 1 | Competing hypotheses for the origin of negative stomatal size-density scaling make different predictions about the trait covariance structure.** Maximum stomatal conductance ( $g_{\text{max}}$ ) and the fraction of epidermal area allocated to stomata ( $f_s$ ) are composite traits determined by stomatal density and size. On a log-scale, they are the sum of log-stomatal density ( $d_s$ ) and

399 log-stomatal size ( $a_s$ ) times a scaling exponent ( $\beta$ ), 0.5 for  $g_{\max}$  and 1.0 for  $f_s$  (see Methods).  
400 There are infinite number of other composite traits, which we denote  $z_s$ , associated with different  
401 values of  $\beta$  (Eq. 4). **a.** A unique stomatal size-density covariance structure is associated with the  
402  $\beta$  that minimizes the variance in the resulting composite trait. If the means ( $\bar{d}_s, \bar{a}_s$ ) and variances  
403 ( $V_d, V_a$ ) of stomatal density and size, respectively, can be measured, the covariance between  
404 them ( $V_{d,a}$ ) is equal to  $-\beta V_a$ . Under the stomatal-area increase (left panel) and stomatal-area  
405 minimization (right panel) hypotheses,  $\beta$  should be 0.5 and 1, respectively. The ellipse is the  
406 0.95 quantile of covariance ellipse associated with the covariance matrix (upper right corner of  
407 the plot); the orange line is the scaling exponent fit through the constituent trait means. **b.** The  
408 covariance structure depicted in **a** is that which minimizes the variance in a composite trait ( $V_z$ )  
409 given a hypothesized scaling exponent.

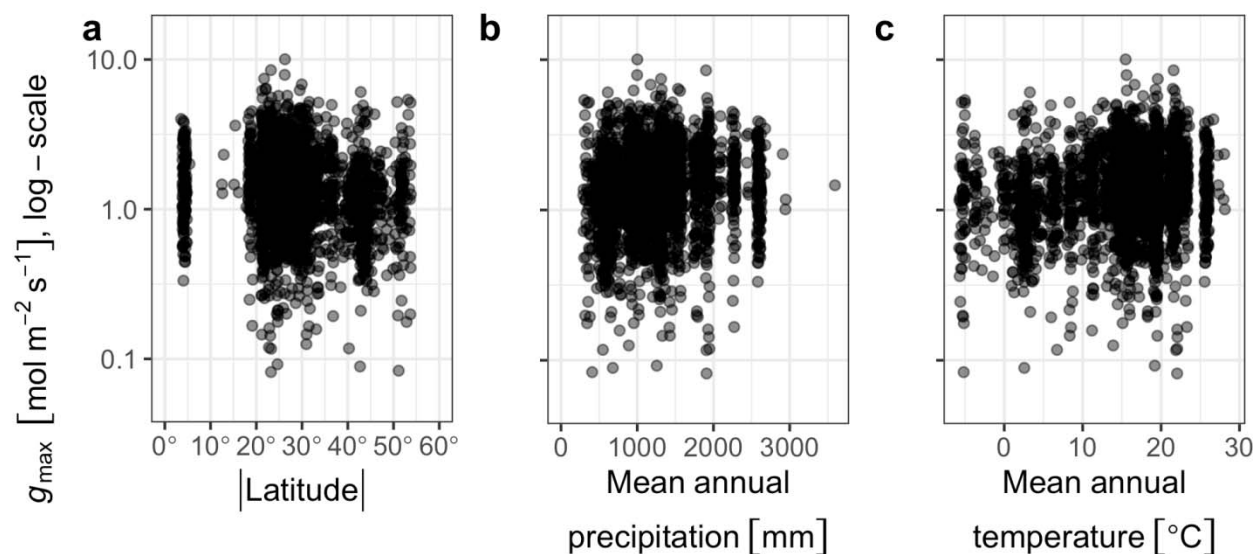


**Fig. 2 | Geographic distribution of sampling sites (a) and the number of plant species (b) in this study.**



**Fig. 3 | Stomatal size-density scaling is consistent with stomata-area increase but not area-minimization.** **a.** In both angiosperms (left panel) and pteridophytes (right panel), the scaling exponent ( $\beta$ ) estimated as the phylogenetic linear regression slope of stomatal size against density (Methods) is close to 0.5 as predicted by the stomatal-area increase hypothesis, but much less than 1.0, as predicted by the stomatal-area minimization hypothesis. For comparison, thin gray lines in the background show predicted slopes for each group when  $\beta = 1.0$  (solid line) and

421  $\beta = 0.5$  (dashed line). The bootstrap 95% confidence intervals are in parentheses and shown  
 422 graphically by the width of the grey rectangle in **b**. Dark points represent species mean trait  
 423 values from the focal group; grey background points are from all groups for comparison. Orange  
 424 line and ribbon are the estimated phylogenetic regression line and the 95% bootstrap confidence  
 425 intervals. Scaling in gymnosperms (middle panel) is not significantly different from 0 or 0.5, but  
 426 the confidence intervals do not include 1.0. **b**. The variance of the composite trait ( $V_z$ ) is  
 427 minimized near  $\beta = 0.5$ , as predicted under the stomatal-area increase hypothesis (dashed-line  
 428 under  $g_{\max}$ ) but not where  $\beta = 1.0$  as predicted by the stomatal-area minimization hypothesis  
 429 (dashed-line under  $f_s$ ).



**Fig. 4 | Anatomical maximum stomatal conductance varies little with latitude, mean annual precipitation, or mean annual temperature.** Each point is the species' mean |latitude| (a.), mean annual precipitation (b.), or mean annual temperature (c.) on the  $x$ -axis and the maximum stomatal conductance ( $g_{\max}$ ) on the  $y$ -axis (log-scale).