Appendix 1: Supplemental Methods and Spatially Implicit Model

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SUPPLEMENTAL METHODS

$g_{s,max}$ calculation

I calculated $g_{s,max}$ (Equation 1) to water vapor at a reference leaf temperature ($T_{leaf} = 25^{\circ}$ C) following Sack and Buckley (2016). They defined a biophysical and morphological constant as:

$$b = D_{wv}/v$$
$$m = \frac{\pi c^2}{j^{0.5}(4hj + \pi c)}$$

b is the diffusion coefficient of water vapor in air (D_{wv}) divided by the kinematic viscosity of dry air (v). $D_{wv} = 2.49 \times 10^{-5} \text{ m}^2 \text{ s}^{-1}$ and $v = 2.24 \times 10^{-2} \text{ m}^3 \text{ mol}^{-1}$ at 25° (Monteith and Unsworth, 2013). For kidney-shaped guard cells, c = h = j = 0.5.

$f_{\rm S}$ is proportional to the stomatal pore area index

The stomatal pore area index (SPI; Sack et al., 2003) is calculated as the product of the stomatal density and guard cell length (GCL) squared:

$$SPI = D \times GCL^2$$

Assuming that the stomatal radius R is half the GCL, then stomatal size S is equivalent to:

$$S = \pi R^2 = \frac{\pi \times \text{GCL}^2}{4}$$

Based on equation 2, it follows that f_{S} and SPI are proportional:

$$f_{\mathbf{S}} = \frac{\pi \times \mathrm{SPI}}{4}$$

SPATIALLY IMPLICIT MODEL

A limitation of the spatially explicit model is that a pathogen could only infect stomata in the focal triangle where it landed. Here I analyze an alternative, spatially implicit, model that relaxes this assumptions. Instead, I assume that a pathogen can potentially infect any stomate on the leaf. It searches through a random walk and has a continuous, constant probability of encountering a stomate that is determined by stomatal cover (f_S). If $f_S \ll 1$, this can be modeled as a homogeneous Poisson process and the distance x a pathogen must travel before reaching a stomate follows an exponential distribution:

$$f(x) = f_{\mathbf{S}}e^{-f_{\mathbf{S}}x}$$

Given the constant death rate per unit distance H, the probability of surviving to distance x is e^{-Hx} . The probability of locating a stomate the probability of surviving to distance x multiplied by f(x) and integrated over all x from 0 to ∞ :

$$p_{\text{locate}} = \int_0^\infty f(x)e^{-Hx}dx$$
$$= \int_0^\infty f_{\text{S}}e^{-(f_{\text{S}}+H)x}dx$$
$$= \frac{f_{\text{S}}}{H+f_{\text{S}}}$$

Substituting p_{locate} above into Equation 5:

$$p_{\text{colonize}} = f_{\text{S}} + (1 - f_{\text{S}})p_{\text{locate}}$$
$$= f_{\text{S}} + (1 - f_{\text{S}})\frac{f_{\text{S}}}{H + f_{\text{S}}}$$
$$= f_{\text{S}}(1 + \frac{1 - f_{\text{S}}}{H + f_{\text{S}}})$$

With the spatially implicit model, because pathogens can potentially reach any stomate on the leaf, p_{colonize} is greater than that in the spatially explicit model for the same value of H. For example, if the pathogen can search forever (H = 0), then it will always colonize ($p_{\text{colonize}} = 1$; Figure S3a). But even when H > 0, p_{colonize} is significantly higher than in the spatially implicit than spatially explicit model for the same f_{S} because pathogens can potentially colonize any stomate on the leaf.

Whereas the spatially explicit model probably underestimates p_{locate} for pathogens that can search over long distances, the implicit model overestimates because it assumes that the probability of encountering a stomate is constant (i.e. homogeneous Poisson process). This is not true because stomata are discrete areas on the leaf. If a pathogen is searching far away from a stomate, its probability of encountering a stomate in the near future is lower than that for a pathogen searching near a stomate. This should be modeled as a *nonhomogenous* Poisson process. Future work should derive p_{locate} for the stomatal anatomies presented here under a nonhomogeneous process. Despite the quantitative differences in the the spatially explicit and implicit models, they have similar qualitative properties when H > 0, which is reasonable since the leaf surface is a relatively hostile environment for most pathogens (see [Introduction]). In both models, p_{colonize} increases with, but is higher than $f_{\rm S}$. In the spatially explicit model, size-density scaling that preserves p_{colonize} is 1 when H = 0 and slightly less than 1 otherwise (Figure 4). In the spatially implicit model, the scaling coefficient is always 1. Rearranging the equation for p_{colonize} above and substituting $f_{\rm S} = DS$, the following relationship holds:

$$SD = \frac{p_{\text{colonize}}H}{1 - p_{\text{colonize}} + H}$$

Since H is a constant the right-hand side of the equation above is constant for a given value of p_{colonize} . Hence the β_p that would preserve the relationship above is simply 1 (Figure S4).



Figure S2. The probability of colonization increases with both stomatal size (S) and density D I simulated the probability of colonization ($p_{colonize}$, y-axis) over a range of S, D, and H (see [Materials and Methods]) **a.** Each line shows how $p_{colonize}$ increases with S (x-axis, log-scale) for selected values of $D \in \{10, 100, 1000\} \text{ mm}^{-2}$. **b.** Each line shows how $p_{colonize}$ increases with D (x-axis, log-scale) for selected values of $S \in \{10, 100, 1000\} \mu \text{m}^2$. The facets show results for different values of H

Figures



Figure S3. The probability of colonization increases with both stomatal cover and conductance in the spatially implicit model. As in Figure 3, I simulated the probability of colonization ($p_{colonize}$, y-axis) over a range of stomatal densities and sizes (see [Materials and Methods]), but a subset of results are shown here. Stomatal size and density determine stomatal cover (f_S ; Equation 2) and theoretical maximum stomatal conductance ($g_{s,max}$; Equation 1). **a.** $p_{colonize}$ initially increases rapidly with f_S (x-axis), then slows down to a linear relationship. Overall, $p_{colonize}$ is lower when pathogens can die on the leaf surface (H > 0). The relationship between f_S and $p_{colonize}$ increases sigmoidally with $g_{s,max}$ at all stomatal densities, but $p_{colonize}$ is lower at higher densities for a given $g_{s,max}$. The relationship between $g_{s,max}$ and $p_{colonize}$ is similar for all values of H > 0.



Figure S4. Log-log scaling relationships between stomatal density (*D*, *x*-axis) and size (*S*, *y*-axis) that preserve the probability of colonization ($p_{colonize}$) in the spatially implicit model. As in Figure 4, in each panel, solid lines indicate values of *D* and *S* where $p_{colonize}$ is 0.25 (lowest line), 0.5, or 0.75 (highest line). For reference, dashed grey lines show scaling relationships that preserve f_S ($\beta = 1$, slope $= -1/\beta = -1$) and $g_{s,max}$ ($\beta = 0.5$, slope $= -1/\beta = -2$) drawn through the centroid of the plotting region. The scaling exponent is unity $\beta = 1$ when H > 0.

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