## ARTICLE ADDENDUM

# Stomate-based defense and environmental cues

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# ABSTRACT

Environmental conditions play crucial roles in modulating immunity and disease in plants. For instance, many bacterial disease outbreaks occur after periods of high humidity and rain. A critical step in bacterial infection is entry into the plant interior through wounds or natural openings, such as stomata. Bacterium-triggered stomatal closure is an integral part of the plant immune response to reduce pathogen invasion. Recently, we found that high humidity compromises stomatal defense, which is accompanied by regulation of the salicylic acid and jasmonic acid pathways in guard cells. Periods of darkness, when most stomata are closed, are effective in decreasing pathogen penetration into leaves. However, coronatine produced by *Pseudomonas syringae* pv. *tomato (Pst)* DC3000 cells can open dark-closed stomata facilitating infection. Thus, a well-known disease-promoting environmental condition (high humidity) acts in part by suppressing stomatal defense, whereas an anti-stomatal defense factor such as coronatine, may provide epidemiological advantages to ensure bacterial infection when environmental conditions (darkness and insufficient humidity) favor stomatal defense.

Plant disease is a successful culmination of three important factors viz. high pathogen virulence, ineffective plant immunity, and favorable environmental conditions. This central dogma of plant pathology is a 50-year-old concept of the disease triangle<sup>1</sup> and is relevant in all aspects of plant-pathogen interactions<sup>2</sup>. Environmental abiotic factors such as relative humidity (RH) and light conditions have a drastic effect on prevalence of disease in different geographical regions. Plants need to adapt to simultaneous exposure to variable biotic and abiotic stresses, sometimes with opposing effects, for maintenance of healthy whole plant physiology. For instance, high disease incidence can be explained by the occurrence of climatic conditions that favor pathogen growth and weaken the plant immune system<sup>3</sup>. It is well known that the outbreak of late blight of potato caused by Phytophthora infestans that lead to the unfortunate Irish potato famine of 1845 was initiated and spread rapidly mainly because of the unusually wet and cool climatic conditions chronicled for that year<sup>2</sup>. Still, current knowledge on the molecular basis of environment-mediated regulation of plant responses to pathogens is still in its infancy. Moreover, we have gathered evidence that different cell types (e.g., guard cell and mesophyll cell) may have variable molecular responses to the same environmental condition<sup>3</sup> adding additional levels of complexity in plant immune responses.

Plant immune system consists of a complex network of signals tuned to respond to specific types of biotic stresses. One of the first outputs of pattern-triggered immunity (PTI) consists of stomatal defense<sup>4</sup>. The microscopic stomatal pores in the leaves are important not only for transpiration and exchange of gases, but also as entry points for some pathogenic microbes, which otherwise could not transit from the phylloplane to the leaf apoplast. However, recognition of microbe-associated molecular patterns (MAMPs) by plant pattern-recognition receptors (PRRs) is a signal to close stomata that serve as guarding gates against microbe invasion<sup>5</sup>. A rapid (< 2h) bacterium-triggered stomatal closure is also observed when the plant perceives non-pathogens such as *Escherichia coli*, *Salmonella enterica*, and *Bacillus subtilis*<sup>4,6-8</sup>.

Molecular mechanisms underlying stomatal defense have been studied mostly in the Arabidopsis-*Pst* pathosystem. This well-studied system has been very useful to decipher both stomatal defense and counter-defense mainly due to the initial PTI response and subsequent induction of coronatine production in the bacterium that overrides PTI<sup>9,10</sup>. This temporal response in the Arabidopsis guard cell is mediated by phytohormones<sup>5</sup>. For instance, abscisic acid (ABA), salicylic acid (SA), and jasmonic acid (JA) play important roles in guard cell signaling during Arabidopsis/*P. syringae* interaction.

Endogenous ABA and SA are important for stomatal closure in response to bacteria or purified MAMPs<sup>4,11-17</sup>. By contrast, strong evidence suggests that, similar to its structural and functional mimic coronatine, jasmonoyl-L-isoleucine (JA-Ile) mediates stomatal opening<sup>3,18</sup>. Intriguingly, control of stomatal movement by air RH also seems to operate through hormone signaling. As an example, low RH induced-stomatal closure is associated with ABA biosynthesis<sup>19</sup>, whereas activation of

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stomatal opening by high RH is associated with ABA catabolism<sup>20</sup>. However, we have found that exogenous treatment of ABA does not close stomata to the full extent under high RH as compared with plants at moderate RH<sup>3</sup>. This finding indicates that while ABA has a prominent role in RH-mediated stomatal movement, it does not seem to be the only target of high RH in guard cells.

Previously, SA-dependent phenotypes have also been shown to be suppressed under high RH<sup>21</sup>, including the suppression of SA-dependent activation of PR genes in Arabidopsis leaves at 24 h after shifting plants to high RH<sup>22</sup>. As SA signaling is required for stomatal closure<sup>4,13</sup>, we performed guard cell-specific analysis and determined that high RH also repressed the expression of PR1 gene in this cell type<sup>3</sup> (Fig. 1). On the other hand, JA-responsive genes are upregulated in guard cells within 1h of plant exposure to high RH<sup>3</sup>. However, this regulation is independent of the JA-Ile receptor, COI1. COI1-independent and JA-dependent signaling pathway has been previously proposed and induction of some JAZ genes in coil plants has been reported when Arabidopsis leaves are infected with Sclerotinia sclerotiorum<sup>23</sup>. In addition, P. syringae pv. maculicola ES4326 infection in coi1-1 plants also leads to induction of JA-regulated genes, indicating that JA response can be activated downstream or independent of COI1<sup>24</sup>. Moreover, an effector from Pst DC3000, HopX1 triggers degradation of JAZ proteins in a COI1-independent manner and promotes stomatal opening<sup>25</sup>. Consistent with this, we observed that the JA biosynthesis genes, LOX3 and OPR3 are repressed within 1 h of exposure to high RH<sup>3</sup>. This finding suggests that JA-Ile replenishment may not be required as the signaling occurs independent of COI1 in guard cells. Specific branches of the SA and JA signaling pathways regulated by RH are yet to be determined.

In several circumstances, JA and SA act antagonistically and some key regulators in this crosstalk have been identified. SA inhibits JA signaling through the regulatory protein, NONEX-PRESSOR OF PR GENES 1 (NPR1)<sup>26</sup>. By contrast, JA and coronatine inhibit SA biosynthesis genes (isochorismate synthase, *ICS1*) and activate SA degradation genes (benzoic acid/SA carboxyl methyltransferase 1, *BSMT1*) through 3 NAC transcription factors, ANAC019, ANAC055, and ANAC072<sup>27</sup>. However, we observed that both activation of JA and suppression of SA occur simultaneously in guard cells of plants exposed to high RH<sup>3</sup> and hence these pathways are likely to be regulated independently by RH. Guard cell response to RH is much quicker (< 1h) than that of whole leaves (> 8h) suggesting the existence of an independent regulation of guard cell signaling by RH. However, it is possible that JA/SA antagonism exist in guard cell under high RH at a step downstream of the signaling components tested so far, which still needs further investigation. Based on current evidence, we propose that the shift of balance between SA and JA signaling leads to repression of bacterium-triggered stomatal closure and consequently bacteria that are otherwise unable to overcome PTI can still penetrate leaf tissue under high RH (Fig. 1).

High humidity also promotes rapid proliferation of bacteria in the epiphytic phase<sup>28</sup>. However, in general, phyllosphere is a water-limiting environment<sup>29</sup> that imposes a challenge for epiphytic survival of pathogens in this niche. To counter this challenge, bacteria produce extracellular polymeric substances (EPS) to maintain hydration and form aggregates on the leaf surface<sup>30,31</sup>. High humidity positively affects such aggregate formation of P. syringae pv syringae B728a on bean leaf surface and aids in rapid proliferation of the bacteria and subsequent entry into the endophytic phase<sup>30</sup>. To maintain epiphytic fitness, virulent bacteria can physically alter the wettability of the leaf surface by producing biosurfactants<sup>32,33</sup>. Furthermore, bacterial-dependency on high RH to establish apoplastic infection while suppressing host immunity has also been demonstrated recently<sup>34</sup>. These observations emphasize that RH participates in multiple steps of molecular plant-pathogen interaction and influences its outcome.

In contrast to high RH that aids plant susceptibility and counteracts stomatal defense, several other abiotic factors may favor a robust stomatal defense. In particular, absence of light may lead to stomatal closure; indeed, most stomata of C3 and C4 plants are closed at night. This suggests that bacterial penetration of leaves through stomata would be minimal at night. Interestingly, the clock proteins CIRCADIAN CLOCK ASSO-CIATED 1 (CCA1) and LATE ELONGATED HYPOCOTYL (LHY) not only control the circadian stomatal movement, but they are also required for flagellin-mediated immune response<sup>35</sup>. Disruption of the clock activity through CCA1 and LHY resulted in stomata that are less responsive to dark and P. syringae pv. maculicola, thus rendering Arabidopsis plants more susceptible to infection at night. Furthermore, surfaceinoculated plants, but not syringe-infiltrated plants, are more resistant to bacterium infection at dusk than at dawn<sup>35</sup>. These findings mechanistically link stomatal defense and the circadian clock.

Interestingly, the levels of the two most well-known hormones associated with biotic stress, JA and SA, naturally oscillate throughout a 24 h cycle. While the JA level peaks in the



Figure 1. A simplified model depicting distinct modulation of JA and SA signaling pathways by different external factors in guard cells. Left: stomatal defense is reduced under high air relative humidity (RH) or in the presence of the phytotoxin coronatine by repressing SA signaling and activation of JA signaling. Right: stomatal defense is enhanced in natural conditions (darkness) or induced by MAMPs, which is accompanied by upregulation of SA signaling and downregulation of JA signaling.



Figure 2. Stomata can be closed in the dark, a condition that may prevent bacterial internalization of leaves (A). However, virulent pathogens, such as the COR-producing bacterium *Pst* DC3000, can open dark-closed stomata and gain entry into the leaf apoplast (B). Colored bacterial cells in (B) indicate induction of coronatine biosynthesis.

daytime, the SA level is highest during the night in whole leaves<sup>36,37</sup>. These oscillations are under the control of the clock and several clock-associated proteins<sup>37</sup>. If the JA/SA hormone balance determines the opening and closing of stomata (Fig. 1), then one would assume that inducing JA signaling at night could promote stomatal opening. Previously, others and we have determined that coronatine, a molecular mimic of JA-Ile, overcomes bacterium-triggered stomatal closure by upregulating JA signaling and repressing SA signaling<sup>4,38</sup>. Consistently, Pst DC3000 senses the leaf surface, produces coronatine, and opens dark-closed stomata<sup>39</sup>. It remains to be determined whether coronatine disrupts the natural guard cell circadian movement by actively suppressing CCA and LHY1 mediated signaling. Nonetheless, it is evident that a stomatal defensefavoring environmental condition such as darkness can be overcome by a virulent pathogen that shifts the hormone balance in guard cell toward JA<sup>3,39</sup> (Fig. 2).

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