Agroinfiltration Reduces ABA Levels and Suppresses Pseudomonas syringae-Elicited Salicylic Acid Production in Nicotiana tabacum

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Abstract

Background: Agrobacterium tumefaciens strain GV3101 (pMP90) is widely used in transient gene expression assays, including assays to study pathogen effectors and plant disease resistance mechanisms. However, inoculation of A. tumefaciens GV3101 into Nicotiana tabacum (tobacco) leaves prior to infiltration with pathogenic and non-host strains of Pseudomonas syringae results in suppression of macroscopic symptoms when compared with leaves pre-treated with a buffer control.

Methodology/Findings: To gain further insight into the mechanistic basis of symptom suppression by A. tumefaciens we examined the effect of pre-treatment with A. tumefaciens on the growth of P. syringae, the production of the plant signalling molecules salicylic acid (SA) and abscisic acid (ABA), and the presence of callose deposits. Pre-treatment with A. tumefaciens reduced ABA levels, P. syringae multiplication and P. syringae-elicited SA and ABA production, but promoted increased callose deposition. However, pre-treatment with A. tumefaciens did not suppress necrosis or SA production in leaves inoculated with the elicitor HrpZ.

Conclusions/Significance: Collectively, these results show that inoculation of N. tabacum leaves with A. tumefaciens alters plant hormone levels and plant defence responses to P. syringae, and demonstrate that researchers should consider the impact of A. tumefaciens on plant signal transduction when using A. tumefaciens-mediated transient expression assays to investigate ABA-regulated processes or pathogenicity and plant defence mechanisms.

Introduction

Agrobacterium tumefaciens is a soil-borne pathogen that causes crown gall disease in dicotyledoneous plants. The bacterium transfers the T-DNA region from its tumor-inducing (Ti) plasmid into plant cells, where it integrates into the plant genome. Once integrated in the plant chromosome, expression of T-DNA encoded genes elicits a hyperplastic response that results in gall formation [1,2]. The ability of A. tumefaciens to transfer DNA across kingdoms has been exploited for biotechnological applications and A. tumefaciens-mediated transformation is routinely used to engineer plants and to express, disrupt or silence genes in functional genomic studies [3,4]. A. tumefaciens can also be used in transient expression assays, where high densities of A. tumefaciens are infiltrated into plant tissues and transgene properties can be assayed within a few days of infiltration. Transient expression is of particular relevance in studies of gene function, protein localization and protein production [5–7].

Agrobacterium-mediated transient expression (agroinfiltration) assays are commonly carried out using Nicotiana tabacum (tobacco), because its leaves are easy to infiltrate and manipulate, and because A. tumefaciens does not have any visible effect on plant health, in contrast to the necrosis elicited by A. tumefaciens when it is introduced into plants such as tomato and Arabidopsis thaliana [8,9]. Although N. tabacum does not exhibit macroscopic responses to A. tumefaciens during transient expression assays, numerous studies have shown that asymptomatic challenge of plants with microorganisms or with microbe-derived molecules can prime or modify plant defences [10–12]. It is therefore logical to hypothesise that infiltration of high densities of A. tumefaciens into bacterial leaves could have a significant impact on the behaviour of the plant immune system. In this work we have investigated how inoculation of A. tumefaciens into N. tabacum leaves at concentrations typically used for transient expression assays affects the production of plant hormones and plant defence responses to bacterial pathogens.

Early studies of A. tumefaciens-plant interactions suggested that A. tumefaciens did not produce elicitors that elicit plant immune responses, as flagellin and flagellin-derived peptides from A. tumefaciens failed to act as pathogen-associated molecular patterns (PAMPs, also known as microbial associated molecular patterns...
(MAMPs)), in Arabidopsis thaliana [13,14]. Flagellins from other plant pathogens, such as Pseudomonas syringae, are recognised by Arabidopsis cells, resulting in ROS production and callose deposition [15,16]. However, recent studies have provided increasing evidence that A. tumefaciens does elicit PAMP-triggered immunity (PTI), although the nature of the elicitors it produces is not fully understood [3,9,17,18]. For example, although Arabidopsis does not recognise A. tumefaciens flagellin, A. tumefaciens can elicit PTI in Arabidopsis via recognition of elongation factor Tu (EF-Tu) by the EF-Tu receptor kinase (EFR) [9]; and via recognition of peptidoglycan (PGN) [19]. In addition, Diet et al. [3] have proposed that a diffusible elicitor from A. tumefaciens activates basal defence responses in Agrostis conyzoides.

There is species-specific variation in PAMP recognition by plants, and neither flagellin nor EF-Tu account for N. tabacum responses to A. tumefaciens [9,13]. However, N. tabacum has been shown to respond to the conserved RNA-binding motif RNP-1 of bacterial cold shock protein (CSP) [19,20], which is present in A. tumefaciens.

The mounting evidence that A. tumefaciens can trigger PTI in many, and possibly all of its plant hosts supports the hypothesis that one function of T-DNA encoded proteins and other A. tumefaciens virulence factors is to modify and suppress plant defence responses [18]. For example, A. tumefaciens may use auxins not only to promote host cell division and tumor proliferation but also to suppress host defences by an antagonistic interaction with salicylate (SA)-mediated defence signalling [21]. Pruss and collaborators [17] have demonstrated that infiltration of N. tabacum with the oncogenic A. tumefaciens strain C58 elicits accumulation of miR393, a microRNA that represses auxin signalling and promotes antibacterial resistance. However, miR393 expression was repressed in A. tumefaciens induced tumors, which led them to speculate that miR393 is down-regulated in later stages of A. tumefaciens infection due to increased auxin synthesis resulting from transfer and expression of T-DNA genes.

We have observed that infiltration of A. tumefaciens into N. tabacum leaves reduces their susceptibility to bacterial pathogens such as P. syringae pv. tabaci (Pta) and attenuates or eliminates the hypersensitive response (HR) to the avirulent pathogen P. syringae pv. tomato (Pto). A similar phenomenon was reported by Pruss and collaborators [17], who observed that infiltration with a disarmed A. tumefaciens strain protected tobacco leaves from infection with tobacco mosaic virus (TMV). The observation that pre-treatment with a non-pathogenic bacterium elicits PTI and suppresses disease or the HR arising from subsequent inoculation of a pathogenic bacterium is not new, as previous studies have described this phenomenon with non-pathogenic strains of Pseudomonas [22-24]. However, this is a relatively uncharacterised phenomenon in the context of A. tumefaciens-plant interactions, despite the popularity of A. tumefaciens as a delivery mechanism for transient expression.

In this report we demonstrate that infiltration of A. tumefaciens GV3101 into N. tabacum leaves not only reduces necrosis associated with disease or the HR, but also suppresses P. syringae growth. We examine the effect of A. tumefaciens on plant defence responses and plant hormones and show that infiltration of A. tumefaciens into N. tabacum leaves elicits a low level of callose deposition and results in reduced abscisic acid (ABA) levels. We also show that when A. tumefaciens-treated leaves are challenged with P. syringae they display reduced salicylic acid (SA) production and increased callose deposition. This clearly demonstrates that agroinfiltration can have effects on plant signal transduction and plant defence responses that are unrelated to transient expression of specific genes. A preliminary report outlining some of these findings was presented at the 7th International Conference on Pseudomonas syringae pathovars and related pathogens [25].

**Results**

**Infiltration of A. tumefaciens into N. tabacum Leaves Prior to Inoculation with P. syringae Results in Reduced Macroscopic Symptoms and Reduced P. syringae Population Levels**

While performing transient expression assays using agroinfiltration to monitor protein expression and localisation in N. tabacum leaves infected with the pathogen P. syringae pv. tabaci (Pta) we observed reduced or complete elimination of necrosis in leaves infected with Pta, compared to leaves treated with the buffer control (10 mM MgCl$_2$ supplemented with 0.5 mM acetosyringone (MgCl$_2$(AS)), data not shown). This suggested that the A. tumefaciens strain used, A. tumefaciens GV3101 (AtGV3101), was affecting either plant signal transduction or pathogen behaviour. To systematically investigate whether disease or HR-associated necrosis induced by P. syringae was attenuated by the presence of AtGV3101, we inoculated AtGV3101 into N. tabacum leaves at 10$^5$ cfu/ml and 48 hours (h) later challenged the leaves with Pta or the non-host pathogen P. syringae pv. tomato DC3000 (Pto) at 10$^5$ cfu/ml. After 24 hours, we observed confluent necrosis in leaf panels pre-treated with MgCl$_2$(AS) and then inoculated with Pto or Pta. However, pre-treatment with AtGV3101 resulted in a significant reduction in the amount of necrosis observed (Fig. 1A), indicating that AtGV3101 inhibits both the HR caused by Pto, and the necrosis associated with inoculation of high densities of Pta.

The effect of AtGV3101 on P. syringae growth was measured in experiments in which Pta and Pto were infiltrated into N. tabacum leaves at 10$^5$ cfu/ml, 48 h after infiltration of AtGV3101. As expected, the population density of the non-host pathogen Pto did not increase significantly over time in either MgCl$_2$(AS) or AtGV3101-pre-treated leaves (Fig. 1B). However, we observed at least a 10-fold reduction in Pto population density in AtGV3101-pre-treated leaves relative to MgCl$_2$(AS)-pre-treated leaves at 5 and 11 days post-inoculation (Fig. 1B). In contrast, the population density of Pta did increase over time, however, AtGV3101 pre-treatment resulted in a 100-fold and 40-fold reduction in population density at 5 and 11 days post-inoculation respectively (Fig. 1B). Consistent with this, the symptoms produced by Pta in AtGV3101-pre-treated leaves were severely reduced or absent (Fig. 1A, bottom panel). The population density of AtGV3101 remained relatively constant throughout the experiment (data not shown). Collectively, these results show that infiltration of AtGV3101 at densities typically used for transient expression, prior to inoculation of P. syringae, inhibits Pta growth, and reduces the viability of Pto, even though the HR elicited by Pto is attenuated.

**A. tumefaciens Suppresses P. syringae-Elicited SA Production and SA-Mediated Defence Responses**

The reduced necrosis and pathogen growth observed in AtGV3101-pre-treated leaves suggested that other aspects of P. syringae-plant interactions, such as plant defence responses, might also be affected by pre-treatment with AtGV3101. One defence response known to be up-regulated during the interaction of Pto and Pta with N. tabacum is the production of the defence signal SA [26]. We therefore used the SA biosensor Acinetobacter baylyi ADPWH-lux (ADPWH-lux) [26] to examine whether AtGV3101 suppressed P. syringae-elicited production of apoplastic SA. When
plants were inoculated with MgCl2(AS) followed by *P. syringae*, both the host pathogen Pta and the non-host pathogen Pto elicited a significant increase in apoplastic SA 24 h after inoculation (77.4±12.4 and 57.5±17.6 μM, respectively, Figure 2B), which is consistent with previous results [26]. However, SA levels were 4- to 5-fold lower in leaves that had been pre-treated with AtGV3101 for 24 or 48 h (Figures 2A and B; 48 h data shown). SA-spiking experiments in which ADPWH-lux was co-infiltrated with 100 μM SA confirmed that inhibition could not be explained in terms of a suppressive effect of AtGV3101 on biosensor activity (data not shown).

To examine whether the effect of AtGV3101 on SA elicitation by *P. syringae* was confined to the apoplastic pool of SA, we also examined SA levels in whole leaf extracts (Figure 2C). Although the values obtained from whole leaf samples are expressed as μg SA/g of freeze-dried tissue and thus cannot be directly compared to values obtained with the biosensor, there was a comparable reduction in SA in both apoplast and whole leaf samples (a 4- and 3-fold reduction in Pta- and Pto-challenged leaves, respectively), which suggests that the effect of AtGV3101 on *P. syringae*-induced SA is associated with an overall reduction in SA synthesis rather than apoplast-specific processes such as inhibition of SA transport or degradation of SA. Consistent with this, we were able to use ADPWH-lux to confirm that AtGV3101 did not deplete SA concentrations when incubated in synthetic medium or in *N. tabacum* apoplast extracts supplemented with 100 μM SA (Figures S1A and B).

It is well established that pathogen-induced accumulation of SA in *N. tabacum* leaves leads to the induction of pathogenesis-related (PR) proteins and the establishment of SAR [11]. As expected, both Pta and Pto elicited *PR1a* expression after 24 h in MgCl2(AS)-pre-treated leaves, but *PR1a* expression was reduced between 3- and 2-fold respectively in leaves that had been pre-treated with AtGV3101 and later inoculated with Pta and Pto, compared to the MgCl2(AS) pre-treatment (2.69±0.71 fold in leaves inoculated with Pta and 2.21±0.35 fold in leaves inoculated with Pto; average of three independent experiments ± standard error; see Figure 2D for a representative experiment). Collectively, the two independent measurements of SA in the apoplast and in whole leaves, and measurement of *PR1a* expression demonstrate that pre-treatment with AtGV3101 suppresses both SA synthesis and SA-mediated responses elicited by *P. syringae*.

Pre-Treatment of Leaves with High Inoculum Densities of Living *A. tumefaciens* Is Needed to Obtain Maximum Suppression of *P. syringae*-Elicited SA Production

Having shown that pre-treatment with AtGV3101 suppressed pathogen growth, symptom development, defence-associated cell death (HR) and SA production, we performed a series of experiments to define the conditions in which AtGV3101 was able to suppress *P. syringae*-elicited symptoms and SA production. Using the same experimental set up as described above, we observed that a suspension of heat-killed AtGV3101 retained some
ability to suppress *P. syringae*-elicited SA, but was less able to suppress SA than live bacteria (Figure 3). This is in accordance with results published by Pruss and collaborators [17], which showed that heat-killed *A. tumefaciens* was less effective at protecting tobacco leaves from infection by TMV than live bacteria. Nevertheless, the observation that heat-killed bacteria retain some ability to suppress SA elicitation provides some support for the hypothesis that plant responses to *A. tumefaciens* PAMPs contribute to the effect of *A. tumefaciens* on *P. syringae*-plant interactions.

Minimum inoculation densities of $10^7$ and $10^6$ cfu/ml of AtGV3101 were needed to obtain consistent suppression of the SA production elicited by Pta and Pto respectively (Figure S2). However, co-inoculation of AtGV3101 at $10^7$ cfu/ml with *P. syringae* at $10^5$ cfu/ml failed to suppress SA production (Figure S3). This argues against the hypothesis that suppression is due to direct inhibition of *P. syringae* as a result of nutrient competition or antagonism, and in favour of the hypothesis that inoculation of high levels of AtGV3101 prior to inoculation with other pathogens elicits host responses that make plant tissues more resistant to subsequent infection.

*A. tumefaciens* Does Not Suppress SA Production Elicited by HrpZ

The observation that AtGV3101 suppressed both the HR and SA production could indicate that AtGV3101 has a repressive effect on plant signal transduction pathways leading to defence-associated cell death. We used the type III secreted protein, HrpZ, which has been shown to elicit both SA production and the HR in *N. tabacum* [27], to examine the effect of AtGV3101 in the absence of AtGV3101-*P. syringae* interactions. As expected, at 1 mg/ml, HrpZ elicited a macroscopic HR (not shown). The tissue collapse associated with the macroscopic HR prevents infiltration of the SA.
biosensor, so we infiltrated HrpZ at 0.1 mg/ml to observe the SA response. AtGV3101 did not have a significant effect on the amount of necrosis elicited by 1 mg/ml HrpZ, or on the amount of SA elicited by 0.1 mg/ml HrpZ [2.623 ± 0.506 μM (MgCl2) and 4.415 ± 1.127 μM (AtGV3101)]; 3 replicates, p-value > F = 0.1783; df = 5) (Figure 4). However, SA production in response to 0.1 mg/ml HrpZ was notably lower than SA production in response to Pto or Pta. These results suggest that the effect of AtGV3101 on the P. syringae-elicited HR and on P. syringae-elicited SA production involves pathways or mechanisms that are not active during plant responses to HrpZ.

Suppression of P. syringae-Elicited SA Production by A. tumefaciens Is Not Dependent on the Type IV Secretion System

AtGV3101 is a non-oncogenic derivative of A. tumefaciens C58 that carries the plasmid mP90, which lacks the T-DNA region but contains the vir genes that encode the type IV secretion system [28]. To investigate whether the type IV secretion system or other Ti plasmid-borne genes contribute to the mechanism by which AtGV3101 suppresses P. syringae-elicited SA production, we tested three derivatives of A. tumefaciens C58 using the SA biosensor assay described earlier: A. tumefaciens A136 is a derivative of C58 cured of its octopine Ti plasmid and carrying the single cryptic plasmid pATC58 [29], A. tumefaciens A348 is isogenic with A136, and carries the Ti plasmid pTiA6 [30]. A. tumefaciens A348 (virG−) lacks a functional copy of virG, which encodes a positive regulator of type IV protein secretion [31]. All three strains displayed a similar ability to suppress SA synthesis induced by Pta and Pto, although in some experiments, the plasmid-cured strain A136 was marginally less effective at suppressing SA synthesis in response to Pta (Figure 5). This is consistent with the observation that a plasmid-cured strain of A. tumefaciens was less able to suppress infection by TMV [17], and suggests that plasmid-borne genes play a minor, but non-essential role in SA suppression.

Infiltration of A. tumefaciens into N. tabacum Leaves Results in Reduced ABA Levels and Primes Callose Deposition

One mechanism by which pathogens suppress plant defence responses is by manipulation of plant hormones [32]. Both the wound and developmental signal JA and the abiotic stress hormone ABA have been implicated in suppression of plant defences during P. syringae infection [32–36]. However, relatively little is known regarding the effect that A. tumefaciens has on ABA or JA levels in plants. To assess whether pre-treatment with AtGV3101 modified hormone levels in N. tabacum we measured both ABA and JA in the experimental conditions described here and found that, as expected, inoculation of MgCl2(AS)-pre-treated plants with Pto or Pta resulted in a significant increase in ABA. However, we observed a marked decrease in ABA in leaves that were pre-treated with AtGV3101, to the extent that ABA levels in AtGV3101-treated leaves were lower than in untreated control leaves (Figure 6A). Interestingly, we did not observe detectable amounts of JA in leaves challenged with Pta or Pto (data not shown), in contrast to the increased JA levels previously reported in Pto-infected Arabidopsis leaves [32]. This is consistent with a previous study by Huang and collaborators [37], who reported that N. tabacum plants inoculated with Pto or Pta produced the volatile SA derivative MeSA, but found that neither Pto or Pta elicited significant production of JA.

The observation that P. syringae grew to lower population densities in AtGV3101 pre-treated leaves compared to control leaves suggested that although AtGV3101 suppressed some defence responses, other defences were enhanced by the presence of AtGV3101. SA-independent defence responses such as callose deposition have been shown to exhibit a strong correlation with plant resistance to P. syringae [38], so we examined the number of callose deposits in infected and healthy N. tabacum leaves pre-treated with AtGV3101 or MgCl2(AS). Neither Pto nor Pta elicited callose deposition in MgCl2(AS)-treated N. tabacum leaves, which...
indicates that basal defences such as callose deposition were effectively suppressed by the pathogenicity and virulence factors produced by these two strains of \textit{P. syringae} \cite{39,40}. However, callose deposits were observed in leaves pre-treated with \textit{AtGV3101}, and were present in high numbers in \textit{Pto-} and \textit{Pta}-infected leaves that had been pre-treated with \textit{AtGV3101} (Fig. 6B and C). This indicates that pre-treatment of \textit{N. tabacum} with \textit{AtGV3101} primes callose deposition, or reduces the ability of \textit{Pto} and \textit{Pta} to suppress the basal immune response.

**Discussion**

The observation that infiltration of \textit{N. tabacum} leaves with \textit{AtGV3101} reduced the symptoms caused by \textit{P. syringae} prompted us to investigate the mechanistic basis of this phenomenon. Our results show that at inoculum densities typically used for transient expression, \textit{AtGV3101} reduces ABA levels, elicits low levels of callose deposition and alters plant responses to virulent and non-host strains of \textit{P. syringae}, suppressing necrosis, SA synthesis and \textit{PR1a} expression, but increasing callose deposition.

This is not the first study to report that \textit{A. tumefaciens} modifies plant responses to \textit{P. syringae}, as Robinette and Matthysse \cite{41} have previously reported that pre-treatment with wild type \textit{A. tumefaciens} \textit{C58} eliminated the HR elicited by \textit{Pseudomonas syringae pv. phaseolicola (Pph)} in \textit{N. tabacum}. However, they concluded that \textit{vir}-independent auxin biosynthesis by the products of genes located in the T-DNA region of the Ti plasmid was required for HR inhibition. In our experiments, a disarmed strain of \textit{A. tumefaciens} that lacked the ability to transform plant cells with auxin biosynthesis genes suppressed \textit{P. syringae}-induced SA synthesis when inoculated at densities typically used for transient gene expression, and similar results were obtained with strains that lacked the Ti plasmid or carried a mutation in \textit{virG}. This demonstrates that Ti-plasmid dependent auxin synthesis is not required for \textit{A. tumefaciens} to modify plant responses to \textit{P. syringae}, and suggests that the results described in this paper illustrate a mechanistically distinct phenomenon. There are clear parallels between our work and that of Pruss and collaborators \cite{17}, who...
showed that agroinfiltration of tobacco leaves with disarmed strains of *A. tumefaciens* resulted in increased resistance to TMV and SA-independent chlorosis and inhibition of leaf expansion over several days. Interestingly Pruss et al. also showed that agroinfiltration of wild-type *A. tumefaciens* elicited a T-DNA dependent increase in miR393, which has been linked to PAMP-dependent repression of auxin signalling, suggesting that T-DNA-dependent effects on plant defence responses, such as those reported by Robinet and Matthysse [41], could be linked to either increased or decreased auxin, depending on the stage of infection. The possible mechanisms by which disarmed strains of *A. tumefaciens* alter plant interactions with *P. syringae* are outlined in Figure 7 and discussed below.

First, it is possible that the presence of high densities of *A. tumefaciens* in the plant apoplast has a direct effect on the proliferation or virulence of *P. syringae* by occupying the intercellular space, producing anti-microbial chemicals and extracellular signalling molecules and competing with *P. syringae* for nutrients, all of which could reduce the ability of *P. syringae* to grow and induce host responses (Figure 7a). However, AtGV3101 was unable to suppress *P. syringae* plant-interactions when co-infiltrated with *P. syringae* (data not shown), and we observed detectable inhibition even when using heat-killed cells (Figure 3), which indicates that *A. tumefaciens* does not need to be alive to exert an inhibitory effect. This argues against direct antagonism and competition as the sole reason for the effect of AtGV3101 on interactions between *N. tabacum* and *P. syringae*. Nevertheless, as the highest level of inhibition was observed with living *A. tumefaciens* cells containing a disarmed Ti plasmid, it seems likely that *A. tumefaciens* needs to become established in plant tissues and to express plant-induced genes before it can most effectively interfere with *P. syringae* growth and *P. syringae*-plant interactions.

An alternative explanation for the effect of AtGV3101 on plant interactions with *P. syringae* is that it could be priming basal defence responses (Figure 7b,c). We observed that at concentrations normally used for transient expression assays, AtGV3101 caused a significant decrease in ABA levels and elicited moderate callose deposition, and when plants were subsequently inoculated with *P. syringae*, we observed reduced SA synthesis and high levels of callose deposition. The hypothesis that *A. tumefaciens* primes plant defences is consistent with reports that signal transduction, gene expression and defence responses are altered upon infection of *N. tabacum* by non-oncogenic *A. tumefaciens* [3,17,18], and that pre-treatment with non-oncogenic *A. tumefaciens* induces plant resistance to TMV, a phenomenon in which direct interactions between apoplastic *A. tumefaciens* and cytoplasmic TMV are unlikely to affect the outcome of pathogen infection [17]. However, in their study, Pruss and collaborators [17] observed a significant increase in *PR1a* expression in response to *A. tumefaciens* ASE/pPZP211, while in our experiments *A. tumefaciens* treatment alone did not alter SA production and *PR1a* expression. This discrepancy may be linked to the fact that they used a significantly

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**Figure 6. Agroinfiltration decreases ABA levels and primes callose deposition in *N. tabacum*.** Leaves were inoculated with *A. tumefaciens* GV3101 (AtGV3101) at 10^7 cfu/ml or with 10 mM MgCl2 (AS), followed 48 hours later by *P. syringae* pv. tabaci 11528 (Pta) or *P. s. pv. tomato DC3000 (Pto) at 10^8 cfu/ml or 10 mM MgCl2. A. ABA levels were determined for whole leaf samples by LC/MS/MS. The graphs show average values from five independent experiments. The bars indicate the standard error of the mean. One way ANOVA revealed statistical differences between treatments (F = 11.1058; p < 0.0001; df = 6). Means with the same letter were not significantly different at the 5% confidence level based on Student’s t-test. B. Leaf sections were excised 24 hours after infiltration with *P. syringae*, stained with aqueous aniline blue, and imaged under ultraviolet excitation at 370 nm. Pictures are representative of five areas of 1.3 mm^2 taken from a leaf section. Two sections from two leaves were stained in each experiment and the experiment was performed twice with similar results. Scale bar, 100 μm. C. Average callose deposits per field of view (1.3 mm^2). The bars indicate the standard error of the mean. GLM revealed statistical differences between treatments (F = 58.7652; p < 0.0001; df = 5). Means with the same letter were not significantly different at the 5% confidence level based on Tukey’s HSD Test.

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Figure 7. Potential mechanisms underlying the effect of *A. tumefaciens* on *P. syringae*-plant interactions. (A) *A. tumefaciens* inhibits proliferation or virulence gene expression in *P. syringae*. The presence of high densities of *A. tumefaciens* in the plant apoplast could have a direct inhibitory effect on proliferation and virulence gene expression in *P. syringae*. This, in turn, could enhance PAMP-triggered immunity (PTI) elicited by *P. syringae* MAMPs and suppress SA production. Infiltration of AtGV3101 also enhances some basal defence responses and ABA levels are reduced. (B) *A. tumefaciens*-mediated priming of the basal immune response inhibits proliferation or virulence gene expression in *P. syringae*. The observation that heat-killed AtGV3101 partially suppresses *P. syringae*-elicited SA argues against a direct interaction between AtGV3101 and *P. syringae* as the sole cause for SA suppression and enhanced expression of basal defences. An alternative explanation would be that AtGV3101 primes for an enhanced basal immune response, which has a negative effect on the ability of *P. syringae* to suppress PTI, and on the ability of *P. syringae* to elicit salicylic acid (SA) synthesis as a result of effector-triggered immunity (ETI) or disease. (C) *A. tumefaciens*-mediated priming of the basal immune response suppresses SA and ABA synthesis. In addition to inhibiting the activities of *P. syringae*, an enhanced basal immune response could directly suppress both SA and ABA synthesis. CW: cell wall; PM: plasma membrane; ABA: abscisic acid; MAMPs: microbe-associated molecular patterns.

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The infiltration of susceptibility to biotrophic pathogens [21,50,51] and been shown to promote increased ABA levels in mock inoculated leaves that had been primed for basal defences. Furthermore, Klement and collaborators [45] demonstrated that induction of local induced resistance (IR) prior to infiltration of P. syringae can suppress expression of the hrp-encoded type III secretion system (TTSS), which is known to be involved in both pathogenesis and HR elicitation.

Interestingly, when we used HrpZ instead of Pto to elicit the HR, we did not observe a decrease in SA accumulation or necrosis (Figure 4 and data not shown). This suggests that HrpZ is able to by-pass any effect of A. tumefaciens on plant cells. Analyses of the signal transduction mechanisms involved in HR elicitation by HrpZ have shown that the HR elicited by HrpZ is SA-dependent, while the P. syringae HR is potentiating, but is not dependent on SA [27,46–48]. It is therefore conceivable that the increase in SA synthesis elicited by HrpZ and P. syringae is SA-dependent, and to proliferate in plant tissues (Figure 7b). For example, Oh and Collmer [44] showed that the HR elicited by Pto DC3000 was suppressed in N. benthamiana leaves that had been primed for basal defences. Furthermore, Klement and collaborators [45] demonstrated that induction of local induced resistance (IR) prior to infiltration of P. syringae can suppress expression of the hrp-encoded type III secretion system (TTSS), which is known to be involved in both pathogenesis and HR elicitation.

Concluding Remarks

Collectively, our results show that infiltration of N. tabacum leaves with Agroinfiltration and ABA

higher density of A. tumefaciens (8×10^9 cfu/ml) as opposed to 1×10^7 cfu/ml), which could have had a more pronounced impact on PRR1a expression. Pruss and collaborators [17] speculated that stimulation of SA-dependent defences by A. tumefaciens could contribute to increased resistance to TMV, but in our system this seems less likely.

The identity of the A. tumefaciens factor or factors that prime callose synthesis and suppress P. syringae-elicited SA responses in N. tabacum remains unclear, but the observation that some suppression was observed with heat-killed cells indicates that one or more MAMPs present in heat-killed bacteria contribute to priming and SA suppression by triggering PTI. SA suppression occurred in response to both a virG mutant and a Ti-plasmid-cured strain of A. tumefaciens C58, which indicates that at least one trait involved in this phenomenon is encoded by a chromosomal, Vir-independent gene. Robinette and Matthisis [41] observed that A. tumefaciens strains carrying a mutation in the chromosomal β-D-1,2-glucan synthesis gene chcB, which disables the ability of bacteria to attach to plant cells and produce high levels of extracellular polysaccharide showed a reduced ability to inhibit a Pph-elicited HR. This suggests that recognition of A. tumefaciens by plant cells benefits from close contact between host cells and bacteria, and raises the possibility that defences are elicited by surface-associated MAMPs.

As shown in Figure 7c, it is possible that callose production in response to A. tumefaciens MAMPs could result in callose-dependent suppression of SA synthesis in response to P. syringae [42,43]. Alternatively, the enhanced basal defences observed in response to A. tumefaciens could have a negative effect on the ability of P. syringae to express or use virulence factors to suppress basal defences, and to proliferate in plant tissues (Figure 7b). For example, Oh and Collmer [44] showed that the HR elicited by Pto DC3000 was suppressed in N. benthamiana leaves that had been primed for basal defences. Furthermore, Klement and collaborators [45] demonstrated that induction of local induced resistance (IR) prior to infiltration of P. syringae can suppress expression of the hrp-encoded type III secretion system (TTSS), which is known to be involved in both pathogenesis and HR elicitation.

Interestingly, when we used HrpZ instead of Pto to elicit the HR, we did not observe a decrease in SA accumulation or necrosis (Figure 4 and data not shown). This suggests that HrpZ is able to by-pass any effect of A. tumefaciens on plant cells. Analyses of the signal transduction mechanisms involved in HR elicitation by HrpZ have shown that the HR elicited by HrpZ is SA-dependent, while the P. syringae HR is potentiating, but is not dependent on SA [27,46–48]. It is therefore conceivable that the increase in SA synthesis elicited by HrpZ and P. syringae involves distinct pathways, and that the HrpZ-dependent pathway is not subject to interference arising from pre-treatment with A. tumefaciens. Alternatively, our results could indicate that the low level of PTI elicited by Agroinfiltration and ABA

The reduction in ABA levels observed in N. tabacum leaves after infiltration of A. tumefaciens (Figure 6A) poses interesting questions. The role of ABA in plant-pathogen interactions is complex and dependent on the pathogen’s lifestyle [32,42,49]. Several studies have shown that increased ABA levels can contribute to increased susceptibility to biotrophic pathogens [21,50,51] and P. syringae has been shown to promote increased ABA levels in Arabidopsis leaves. This increase in ABA levels aids the infection process by suppressing basal defences [32,34]. As expected, we observed increased ABA levels in mock inoculated N. tabacum leaves that were later inoculated with P. syringae, along with little or no callose deposition. The role of ABA in suppressing basal defences in Arabidopsis is known to include the attenuation of callose deposits and suppression of PAMP-induced genes (PIGs) [32,52]. It is therefore conceivable that the reduction in ABA levels in Agroinfiltrated leaves could contribute to priming of plant defences and enhanced callose deposition (Figure 6B and C). In support of this, Asselbergh and collaborators [53] have recently reported that the ABA-deficient tomato mutant Solanum lycopersicum s收到R shows an enhanced basal immune response to the soft-rot pathogen Dickeya dadantii (Erwinia chrysanthemi).

We were unable to detect significant levels of JA in any of our treatments, which is consistent with a previous study by Huang and collaborators [37], and suggests that JA plays a relatively minor role in the hormonal interplay that occurs during A. tumefaciens and P. syringae interactions with N. tabacum. This contrasts with the central role of JA signalling in P. syringae-tomato and P. syringae-Arabidopsis interactions [54–56]. We did not examine the impact of disarmed strains of A. tumefaciens on auxin, which might also be expected to have a significant impact on plant defence responses if altered, and to suppress SA signalling if increased [21]. However, as noted above, both this study and previous work by Pruss and collaborators [17] have shown that the ability of A. tumefaciens to suppress pathogen-induced necrosis and SA synthesis is independent of T-DNA dependent auxin synthesis and T-DNA dependent elicitation of the auxin-repressing miRNA miR393. Nevertheless, it would be interesting to examine whether miR393 or auxin levels are altered in Agroinfiltrated leaves following inoculation of P. syringae, as increased expression of miR393 might be expected to occur in conjunction with enhanced expression of basal defence responses.

Materials and Methods

Plant Material and Bacterial Strains

Tobacco (N. tabacum L. cv Xanthi) seeds were sown in a 3:1 soil:vermiculite mix and grown in a cooled incubator (LMS Ltd, Kent, UK) under a 16 h photoperiod and 21°C temperature. When cotyledons were visible, seedlings were transferred to 1 l pots and grown under the same conditions for 6 to 8 weeks.

The bacterial strains used were: P. syringae pv. tomato DC3000 [57], P. syringae pv. tabaci ATCC11528 (American type culture

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were heated at 70°C prior to inoculation with P. syringae, an overnight liquid culture of live or heat-killed AgtGV3101 was re-suspended in 10 mM MgCl₂ with 0.5 mM acetosyringone (MgCl₂/AS) at approximately 10⁷ cfu/ml and infiltrated into leaves. Ten millimolar MgCl₂/AS was used as a negative control. Plants were incubated in the growth conditions described above, and after 48 h, overnight cultures of Pta and Pto were re-suspended in 10 mM MgCl₂ and infiltrated at 10⁷ cfu/ml. For SA biosensor assays leaves were infiltrated with the SA biosensor ADPWH_lux 24 h after inoculation of Pta and Pto, as discussed below. For HR assays Pta and Pto were infiltrated at 10⁷ cfu/ml. Experiments were also carried out in which leaves were co-inoculated with AgtGV3101 and P. syringae, or pre-treated with AgtGV3101 for 24 h prior to inoculation with P. syringae, in order to examine whether pre-treatment with AgtGV3101 was necessary for modulation of plant defences. To heat-kill A. tumefaciens, the resuspended cells were heated at 70°C for 30 min, then cooled at room temperature and diluted to 10⁷ cfu/ml. No colonies were recovered after plating 100 μl of heat-treated cells. HrpZ was prepared and purified as described by Lee and collaborators [60]. Crude protein extracts of 1 mg/ml and 0.1 mg/ml were used to infiltrate tobacco leaves.

For population counts each replicate consisted of two leaf discs of 0.5 cm², which were collected from leaves at 0, 3, 5 and 11 days after inoculation of P. syringae. Leaf discs were homogenised in 10 mM MgCl₂ and the resulting suspension was dilution plated on LB agar supplemented with Gentamicin (10 μg/ml) for selection of AgtGV3101 and Nitrofurantoin (50 μg/ml) for selection of P. syringae strains and incubated at 28°C overnight.

Salicylate Utilisation Assays

An overnight culture of AgtGV3101 was resuspended at OD₆₀₀ 0.02 in 100 μl of LB, M9 minimal medium [59], or N. tabacum apoplast extract [61], or in media and apoplast extracts supplemented with 100 μM SA, and incubated for 24 h in a 96-well plate at 28°C with shaking. Media and apoplast extracts supplemented with SA were used as internal standards. AgtGV3101 growth was measured after 24 h at OD₆₀₀ using a Tecan Infinite® 2000 plate reader (Tecan Ltd, Männedorf, CH). To measure SA levels, the plate was centrifuged at 3000 rpm for 5 min and 20 μl of supernatant were mixed with 50 μl of A. baylyi ADPWH_lux [OD₆₀₀ 0.4] and 30 μl of LB in a black clear-bottom 96-well plate. Bioluminescence and OD₆₀₀ were measured after incubation for 1 hour at 28°C with shaking. Relative bioluminescence was obtained by dividing bioluminescence of OD₆₀₀ to estimate the SA concentration as described previously [26,58].

SA-induced luminescence was imaged in the dark using a photon-counting camera (Photek Ltd., East Sussex, UK) one hour after infiltration of the ADPWH_lux biosensor. Leaves were acclimated to the dark for at least 100 seconds to allow photosynthesis-associated autoluminescence to disappear, and then luminescence counts were recorded every second for 200 seconds. In vitro SA concentration ladders were included with each imaged leaf to allow comparison of bioluminescence between separate images and were prepared as described previously [26] to yield final SA concentrations of 0, 0.02, 0.05, 0.1, 0.2, 0.3, 0.4, 0.5, 1.0 and 2.0 μM in a final volume of 100 μl. The total pixel intensity for each inoculated area was obtained using Photek IFS32 image processing and data acquisition software (Photek Ltd.) and normalized luminescence values were obtained by dividing the total counts by number of pixels in the inoculated area. Numerical SA values were calculated using a calibration curve as described in [26]. Briefly, N. tabacum leaves were used to produce in vivo SA concentration ladders by mixing various concentrations of SA with the biosensor prior to infiltration into the leaf. Images of the in vivo SA concentration ladders from at least two separate leaves were used to construct a calibration curve.

Measurement of SA, ABA and JA in N. tabacum Leaf Extracts by LC/MS/MS

To confirm the SA measurements obtained with ADPWH_lux and to obtain whole leaf data for SA, ABA and JA, plants were inoculated with AgtGV3101 (10⁵ cfu/ml) and P. syringae (10⁵ cfu/ml) as described above. Twenty four hours after inoculation of P. syringae, plant material was harvested in liquid nitrogen and freeze-dried. To obtain homogeneous samples, samples from three leaves of 6 to 8 week-old N. tabacum plants were pooled for each treatment. Ten milligrams of freeze-dried tissue were processed for each sample and contents of SA, ABA and JA levels were analysed as described previously [62].

Analysis of PR1a Defense Gene Expression

To study the effect of AgtGV3101 on SA-dependent gene expression, plants were inoculated with AgtGV3101 (10⁵ cfu/ml) and P. syringae (10⁵ cfu/ml) as described above, and expression was analysed 24 h after inoculation of P. syringae by quantitative real-time RT-PCR (qRT-PCR), using EF-1α and actin as internal controls. Leaf samples were homogenised in liquid nitrogen and RNA was isolated using the RNeasy® Plant Mini kit (Qiagen GmbH, Hilden, Germany) according to the manufacturer’s instructions. RNA was treated with RNase free DNase (Applied Biosystems/Ambion, Austin, USA) and random primed first-strand cDNA was generated using the ImProm-II™ Reverse Transcription System (Promega Corp., Madison, WI, USA). The primers used to detect the relative abundance of PR1a were PR1aF
(5′-GCCAAAATATGCTTCCCA-3′), PR1aR (5′-CCGT-CATGAAATGCACCATTT-3′), EF1αF (5′-CTGTGGATGACCAAGACTTC-3′), EF1αR (ACAAAAGCGCTTGAGACATG-3′), NtActinF (5′-CCAGTGCGCTAAGACGGTAT-3′) and NtActinR (5′-CCACCGAAGATTTGCTAG-3′). Real-time PCR was performed with the SYBR Green PCR Master Mix kit (Applied Biosystems). Each reaction was performed in triplicate plus a negative control using a 7300 Real Time PCR system (Applied Biosystems) and three independent experiments were performed. The comparative CT method [63] was used to quantify the relative expression ratio of PR1a in leaves infected with P. syringae that had been pre-treated with AtGV3101 or MgCl2.

**Callose Staining**

To assess basal defence responses, leaves were inoculated with AtGV3101 (107 cfu/ml) and P. syringae (107 cfu/ml) as described above and leaf sections were excised 24 hours after infiltration with P. syringae and stained with aqueous aniline blue [64]. Leaf sections were imaged under ultraviolet excitation at 370 nm using an Olympus BX50 epifluorescence microscope. Two sections from two leaves were stained in each experiment and the experiment was performed twice.

**Supporting Information**

**Figure S1**  *A. tumefaciens* does not degrade SA. A. The ability of *A. tumefaciens* AtGV3101 (AtGV3101) to degrade SA was examined in plant extracts and synthetic media. AtGV3101 was inoculated into LB or tobacco apoplastic extract (APOPLAST) supplemented with 100 µM SA. After 24 h of growth, 20 µl of the supernatant of the AtGV3101 cultures was mixed with 50 µl of the SA biosensor ADPWH-lux (grey bars) and SA-induced luminescence was measured after 1 hour. As control, the SA biosensor was mixed with uninoculated media containing 100 µM SA (black bars). General Linear Model (GLM) analysis did not reveal statistical differences between treatments (F = 0.7589; p = 0.5479; df = 11). The experiment was performed twice with similar results. B. The inability of *A. tumefaciens* AtGV3101 to degrade SA was not due to inhibition of bacterial growth by SA or plant extracts. *A. tumefaciens* AtGV3101 was inoculated into LB, tobacco apoplastic extract and the same media supplemented with 100 µM SA. Growth was assessed after 24 hours. General Linear Model (GLM) analysis revealed statistical differences between treatments (F = 1394.993; p<0.0001; df = 11). Means with the same letter were not significantly different at the 5% confidence level based on Tukey’s Honestly Significant Mean Differences (HSD) Test. Error bars in A and B show standard error of the mean.

**Figure S2** High densities of *A. tumefaciens* are needed to suppress *P. syringae*-elicited SA. Tobacco leaves were inoculated with *A. tumefaciens* GV3101 (AtGV3101) at 107, 106, 105 and 104 cfu/ml or 10 mM MgCl2/AS (MgCl2) followed by inoculation with *P. syringae* pv. tabaci 11528 (Pta) or *P. syr. pv. tomato* DC3000 (Pto) (107 cfu/ml) after 48 hours. The SA biosensor ADPWH-lux was inoculated into leaves 24 hours after inoculation with *P. syringae*. SA-induced luminescence was measured one hour after biosensor inoculation using a photon-counting camera and numerical SA values were calculated using a calibration curve as described in Huang et al. [26]. The chart shows average SA values from at least three leaves from different plants. Bars show standard error of the mean.

**Figure S3** Co-inoculation of *A. tumefaciens* with *P. syringae* does not suppress *P. syringae*-elicited SA. Luminescence was examined 24 h after co-inoculation of 105 cfu/ml AtGV3101 with 105 cfu/ml Pta or Pto by infiltrating the ADPWH-lux SA biosensor in the inoculated zones. SA-induced luminescence was measured as described in Figure 2. The Y axis is shown in log scale. General Linear Model (GLM) analysis did not reveal statistical differences between treatments (F = 0.4547; p = 0.7293; df = 7). The experiment was performed twice with similar results. Error bars show standard deviation.

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**Author Contributions**

Conceived and designed the experiments: AR GP. Performed the experiments: AR MB SF. Analyzed the data: AR MB SF. Contributed reagents/materials/analysis tools: MB SF WH. Wrote the paper: AR GP.

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