

BRIEF REPORT

The plant host environment influences competitive interactions between bacterial pathogens

Hanareia Ehou-Taumaunu¹  | Kevin L. Hockett^{1,2,3} 

¹Department of Plant Pathology and Environmental Microbiology, The Pennsylvania State University, University Park, Pennsylvania, USA

²Center for Infectious Diseases Dynamics, The Pennsylvania State University, University Park, Pennsylvania, USA

³The Huck Institutes of the Life Sciences, The Pennsylvania State University, University Park, Pennsylvania, USA

Correspondence

Kevin L. Hockett, 316 Buckhout Lab., University Park, PA 16802, USA.
Email: kh450@psu.edu

Funding information

College of Agricultural Sciences, The Pennsylvania State University; Fulbright New Zealand; Indigo Agriculture Phytobiomes Fellowship; Maori Education Trust; Ministry of Education- New Zealand; Penn State Microbiome Center, The Pennsylvania State University; The Huck Institutes for the Life Sciences, The Pennsylvania State University; USDA National Institute of Food and Agriculture and Federal Hatch Appropriations, Grant/Award Number: PEN04648 (accession no.1016871)

Abstract

Bacteria compete for resources in diverse environments using an array of antagonistic strategies, including the production of narrow-spectrum protein antibacterials termed bacteriocins. Although significant research has focused on bacteriocin-mediated dynamics in culture environments, little research has explored bacteriocin-mediated dynamics within a host context, particularly in plant environments. Here, we show that a bacterial plant pathogen, *Pseudomonas syringae* pv. *syringae* (*Psy*), expresses a bacteriocin both in culture and in leaf apoplast when co-inoculated with a bacteriocin-sensitive competitor, *P. syringae* pv. *phaseolicola* (*Pph*). Although there is an observable negative effect of the bacteriocin on the *Pph* population at most time points both in culture and in the leaf apoplast, a bacteriocin-mediated benefit to *Psy* was only observed when the producing strain was co-infiltrated at a low population frequency (1:9) into the leaf apoplast. At 6 days post-infiltration, *Psy* achieved an eightfold population increase compared to a bacteriocin-deficient mutant in the apoplast. No bacteriocin-mediated benefit for *Psy* was observed under the culture conditions tested. Additionally, we found that the bacteriocin-mediated benefit for *Psy* was dependent on the Type III Secretion System. Taken together, our results demonstrate that the fitness benefit of bacteriocin-mediated antagonism is influenced by interactions within the host plant.

INTRODUCTION

Microbial competition is pervasive throughout the microbiological world. Populations with overlapping niche requirements often engage in both direct (interference) and indirect (exploitation) competition (Ghoul & Mitri, 2016; Granato et al., 2019). Outcomes of microbial competition have numerous consequences at the population and community levels and can also be harmful for plant and animal hosts by causing disease. For bacteria, one of the most intensively studied forms of competition is interference competition mediated by the production of antimicrobials, including protein toxins called bacteriocins (Ghequire & De Mot, 2015; Kommineni et al., 2015; Majeed et al., 2011; Riley &

Wertz, 2002). Most bacteriocins are narrow spectrum within an individual species, where they target strains closely related to the producer (Mills et al., 2017; Riley & Chavan, 2007). A group of bacteriocins known as tailocins are multi-protein complex bacteriocins that are morphologically and evolutionarily related to the tails of Caudovirales bacteriophages (Ghequire & De Mot, 2014; Hockett et al., 2015; Scholl, 2017). Due to their large size, tailocins particles must be released via cell lysis resulting in a cost to the individual producing bacterium (Scholl, 2017).

In silico and *in vitro* work has demonstrated the benefit of bacteriocin production is dependent on the environmental context. Initial studies examined competition with two populations of a producer and a sensitive

This is an open access article under the terms of the [Creative Commons Attribution-NonCommercial-NoDerivs](https://creativecommons.org/licenses/by-nc-nd/4.0/) License, which permits use and distribution in any medium, provided the original work is properly cited, the use is non-commercial and no modifications or adaptations are made.

© 2022 The Authors. *Environmental Microbiology Reports* published by Society for Applied Microbiology and John Wiley & Sons Ltd.

strain in a physically unstructured environment such as liquid broth (Chao & Levin, 1981). When both populations are at equal frequencies, they can reach equilibrium as the sensitive population is able to exploit the resources made available from the killing activity by the producer (Durrett & Levin, 1997). In this scenario, there is no fitness benefit for the producer as its population remains the same. However, when the environment is spatially structured (e.g. agar plate) the two species form microcolonies, which results in local interactions where bacteriocins only affect cells that are physically close to the producing cells, as well as resources made available by killing (Chao & Levin, 1981; Kerr, 2007). This allows the producer to preferentially gain a fitness benefit, an increase in population size, from the available resources and space.

Competition within host environments can add an additional level of complexity compared to a static lab culture setting, given the host's potential to sense and respond to microbial invaders. In mice models, bacteriocin production by the human pathogen *Salmonella enterica* in the gut provides a competitive advantage against *Escherichia coli*, but only if *S. enterica* is able to induce inflammation (Nedialkova et al., 2014). The change in the gut environment results in both increased bacteriocin production by *S. enterica* as well as expression of the bacteriocin surface receptor in *E. coli*. Other in vivo studies in animal systems have also shown that bacteriocin production can result in the reduction or elimination of targeted populations (Corr et al., 2007; Kommineni et al., 2015; Sassone-Corsi et al., 2016; Yu et al., 2020), though it is not clear to what extent interaction with the animal host was important for the competitive outcomes in these studies. Indeed, there have been very few studies that have explicitly assessed the role of the host in affecting bacteriocin-mediated interactions.

Bacteriocins have also been shown to promote invasion of the producer into another population. Invasion into a sensitive population by six *E. coli* colicin producers was positive-frequency dependent (Riley & Gordon, 1999). When the number of *E. coli* producing cells increased, resulting in a higher toxin particle number, the time needed for invasion decreased. Whereas the growth of another *E. coli* bacteriocin producing population was negative-frequency dependent, in that bacteriocin production was beneficial when the producer's population was small relative to the competitor (Chao & Levin, 1981). Furthermore, higher bacteriocin production rates aided invasion into a bacteriocin-sensitive population, especially at low initial cell frequencies (Ghazaryan et al., 2019). Contrary to these findings, modelling predicts a different outcome where the benefits for the producer at low frequency are reduced as resources liberated by bacteriocin killing will be as likely

to benefit the sensitive population as it will the bacteriocin producing population (Inglis et al., 2009; Weber et al., 2014).

Beyond basic ecological questions, understanding the fitness benefit of bacteriocins has implications for the creation of biological control agents. For plant health, we have relied on antagonist mechanisms such as toxins and antibiotics; however, it could be useful to also think about when and where it is beneficial for the agent to antagonize a target pathogen. Bacteriocins are of increasing interest as they could reduce non-target effects as observed with chemicals (McEvoy, 1996; Montesinos, 2007). On the surface of plant cells within the apoplast, bacteria can form microcolonies of single or multiple species where they can interact and compete (Bogino et al., 2013; Morris & Monier, 2003). In addition, bacteria have to evade host defences to successfully populate using virulence factors or by *in trans* effector-mediated plant defence suppression for example (Dodds & Rathjen, 2010; Rufian et al., 2018; Singh & Singh, 2018; Xin et al., 2018). To date, there are few studies that have investigated the role of bacteriocin-mediated antagonism in a plant context, let alone in the apoplast, thus it is not clear how much dynamic interaction there was with the host (Dorosky et al., 2018; Godino et al., 2016; Hert et al., 2005; Li et al., 2020).

To investigate plant-associated bacteriocin-mediated competition, we used *Pseudomonas syringae* as a model, as it is possible for multiple distantly related strains of this species to infect the same plant host and many can produce bacteriocins to antagonize competitors (Hirano & Upper, 2000; Holtsmark et al., 2008). Here, we performed a series of in vitro co-inoculations and *in planta* co-infiltrations over an 8-day period with *P. syringae* pv. *syringae* (*Psy*) and *P. syringae* pv. *phaseolicola* (*Pph*), both virulent plant pathogens that cause bacterial brown spot and halo blight in *Phaseolus vulgaris* (Common bean), respectively (Burkholder, 1926; van Hall, 1902). *Psy* is a generalist pathogen that can infect multiple hosts, whereas *Pph* is a specialist with a narrow host range of legume species (Baltrus et al., 2011; Morris et al., 2019). Key to this interaction is that *Psy* encodes a bacteriocin that targets *Pph* (Hockett et al., 2015). In this study, we sought to answer two related questions. First, under what ecological conditions is bacteriocin production beneficial for the producer? Second, how do host interactions influence the fitness benefits of bacteriocin production (i.e. an increased population size when in competition)? This work highlights the importance of understanding how host structure and activity influence microbial competition and is a critical step to improve disease suppression in plant and animal hosts.

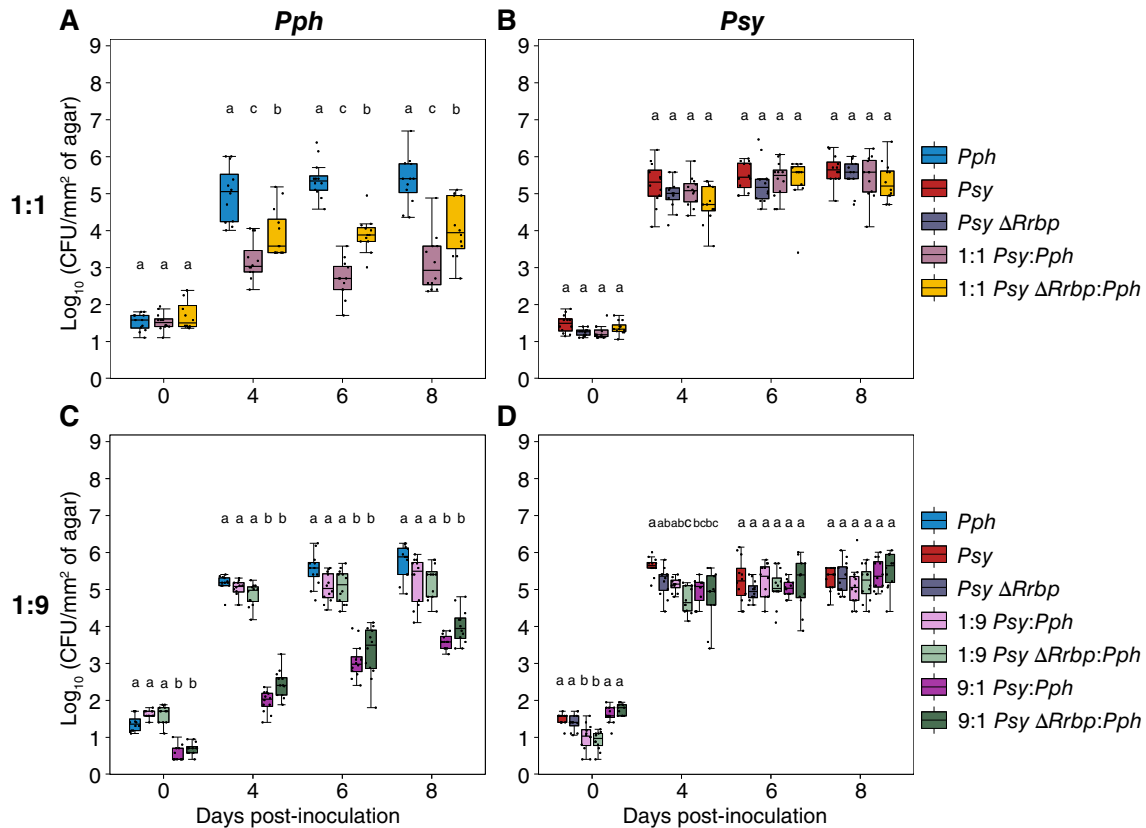


FIGURE 1 Bacteriocin-mediated competition in vitro detrimental for sensitive strain, yet no fitness benefit for producer. Bacterial populations of *Pph* in (A) 1:1 and (C) 1:9 initial starting frequencies (*Psy:Pph*), and *Psy* strains in (B) and (D), respectively, are shown across 8 days post-inoculation. Data points represent four replicates from three independent experiments and error bars indicate standard error of the mean. Average values with the same letter are not significantly different by Tukey's HSD test ($p \leq 0.05$)

RESULTS

Bacteriocins are expressed in vitro but there is no detectable fitness benefit for *Psy* at 1:1 and different co-inoculation ratios

To investigate whether bacteriocin-mediated antagonism provides a fitness benefit within an agar environment, we spotted individual or mixed strains (1:1) of either *Psy* (bacteriocin-producer) or *Psy* $\Delta Rrbp$ (bacteriocin-deficient mutant; Hockett et al., 2015), and *Pph* (bacteriocin-sensitive) on KB agar. At several time points post-inoculation, the growing culture was sampled, and strains were enumerated (Table S1). *Pph* populations were reduced in co-inoculation with either *Psy* or *Psy* $\Delta Rrbp$ at all time points in comparison to *Pph*-only [Figure 1(A)]. The population reduction was greater for *Pph* co-inoculated with *Psy* compared to *Psy* $\Delta Rrbp$, suggesting the bacteriocin was expressed and active under these culture conditions. Bacteriocin production did not provide a detectable fitness benefit in 1:1 co-inoculation in an agar setting due to no significant fitness differences between *Psy* or *Psy* $\Delta Rrbp$

populations in individual or co-inoculations [Figure 1(B)]. Individual inoculation of the bacteriocin-deficient complement strain *Psy* $\Delta Rrbp:Rrbp$ at 4 dpi seemed to be greater by threefold compared to *Psy* $\Delta Rrbp$; however, the statistical difference is only true in one of three independent experiments [$p \leq 0.04$; Figure S2(B)].

As bacteriocin production in 1:1 co-inoculation did not result in a fitness benefit for *Psy*, we sought to assess whether the population frequency influenced competition, as has been previously shown in other systems (Chao & Levin, 1981; Gordon & Riley, 1999; Inglis et al., 2009; Kerr, 2007). We altered the inoculation frequency between *Psy* strains and *Pph* to either 1:9 or 9:1. For example, 'Psy minority' would represent 1:9 (*Psy:Pph*), and vice versa for 'Psy majority' at 9:1 (*Psy:Pph*). In vitro *Pph* minority populations were reduced in co-inoculation with *Psy* or *Psy* $\Delta Rrbp$ compared to *Pph*-only across all time points [$p \leq 0.0001$; Figure 1(C)]. In *Pph* majority competitions, *Pph* growth with *Psy* and *Psy* $\Delta Rrbp$ was not different to *Pph*-only [Figure 1(C)]. Somewhat unexpectedly, *Psy* minority reached a population equivalent to the *Psy* majority treatments by 4 dpi [$p \leq 0.0001$; Figure 1(D)]. This

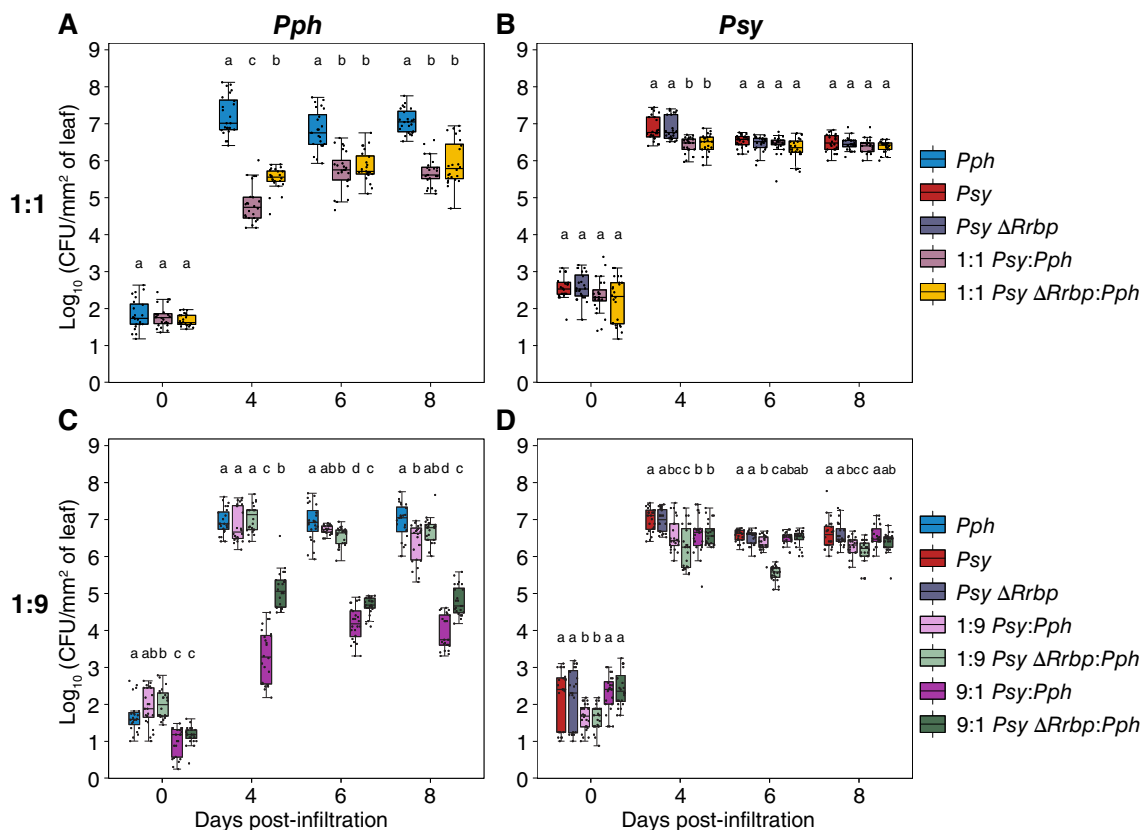


FIGURE 2 Initial low frequency provides fitness benefit for bacteriocin producer at select timepoint. Initial inoculation frequencies of 1:1 are shown in (A) and (B) and 1:9 in (C) and (D) for both *Pph* and *Psy* strain populations (*Psy*:*Pph*). Black dots represent eight replicates across three independent assays over 8 days. Statistically, differences from Tukey's HSD test ($p \leq 0.05$) represented by letters and error bars indicate standard error of the mean

trend was maintained at 6 and 8 dpi. As expected, the *Psy* $\Delta Rrbp$:*Rrbp* population behaved similarly to *Psy* and *Psy* $\Delta Rrbp$ in both individual and mixed inoculations [Figure S2(D)]. Taken together, in vitro, there was no detectable benefit to bacteriocin production for *Psy*, yet a negative effect of bacteriocins on *Pph* is clearly observed.

***Psy* is competitively superior to *Pph* in 1:1 plant co-infiltration regardless of bacteriocin production**

To determine whether the leaf apoplast environment affects the competitive interactions between *Psy* and *Pph*, we infiltrated common bean leaves with either individual or 1:1 mixed inocula of the same strains used in the in vitro assay. Similar to our in vitro results, we found that co-infiltration with *Psy* resulted in a 10-fold greater reduction of the *Pph* population than co-infiltration with *Psy* $\Delta Rrbp$ [$p \leq 0.0001$; Figure 2(A)]. This effect, however, was only observed at 4 dpi. There were no statistical differences after this timepoint, where the *Pph* population was suppressed by a similar

amount by both *Psy* and *Psy* $\Delta Rrbp$. The complement strain *Psy* $\Delta Rrbp$:*Rrbp* showed a similar reduction of *Pph* at 4 dpi, suggesting the *Pph* reduction in co-infiltrations is due to bacteriocin production [$p \leq 0.0001$; Figure S3(A)].

For *Psy*, there was no difference in population growth for *Psy*-only compared to *Psy* $\Delta Rrbp$ -only at all dpi [Figure 2(B)]. There was, however, a sevenfold reduction in population size of *Psy* and *Psy* $\Delta Rrbp$ in 1:1 co-infiltration at 4 dpi compared to *Psy*-only and *Psy* $\Delta Rrbp$ -only, respectively ($p \leq 0.0001$). Therefore, 1:1 co-infiltration with a sensitive strain shows that bacteriocin production provided no fitness benefit for *Psy*.

At low frequency, bacteriocin production provides *Psy* a fitness benefit when competing with *Pph* in the apoplast

The negative effect of the bacteriocin production on *Pph* minority was apparent at all time points, with no effects on *Pph*-majority populations [Figure 2(C)]. Notably, there was an 80-fold reduction for co-infiltrated *Pph*

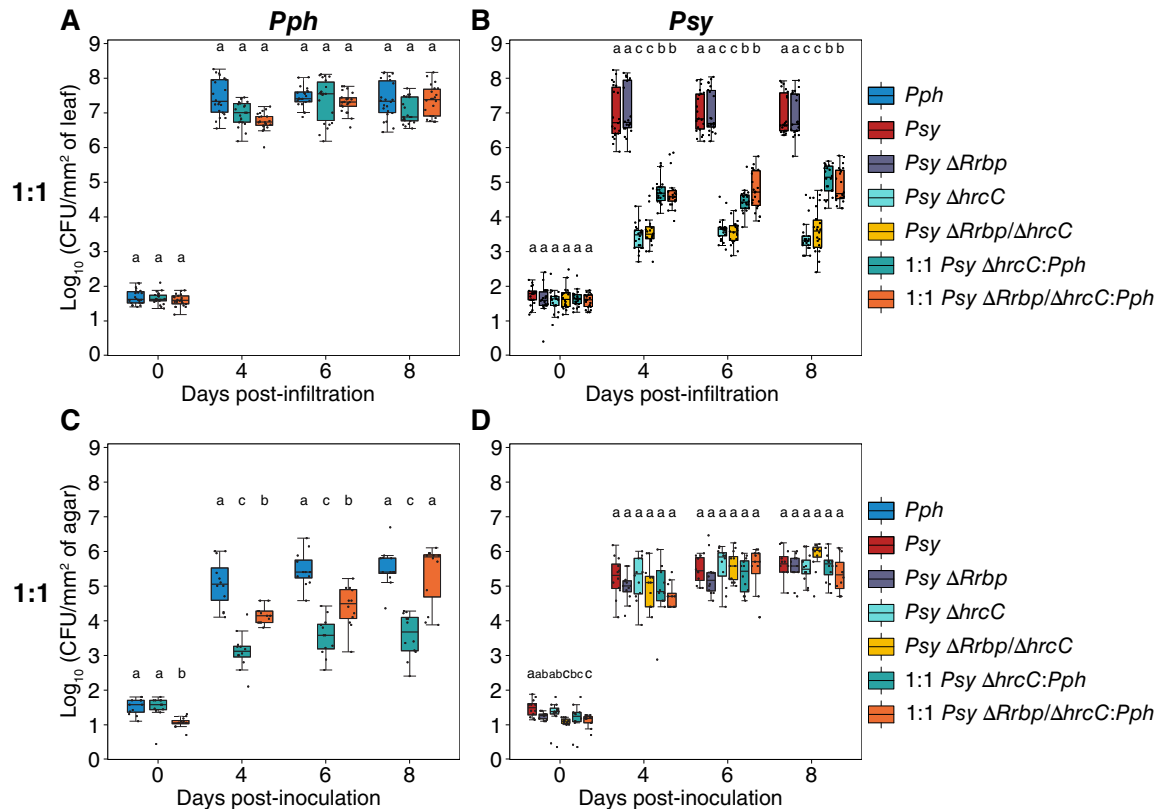


FIGURE 3 Avirulent *Psy* $\Delta hrcC$ cannot reduce sensitive population 1:1 *in planta* competition. Bacterial growth of *Pph* strains is presented (A) in culture and (C) *in planta*, along with *Pph* (C) *in vitro* and (D) *in planta* from initial 1:1 frequency (*Psy*:*Pph*). Data points across 8 days post-inoculation represent eight and four replicates, for *in planta* and *in vitro* respectively, from three independent experiments and error bars indicate standard error of the mean. Average values with the same letter are not significantly different by Tukey's HSD test ($p \leq 0.05$)

minority with *Psy* at 4 dpi compared to *Pph* minority co-infiltration with *Psy* $\Delta Rrbp$ ($p \leq 0.0001$). The differences in population to *Pph*-only were greater for *Pph* minority with *Psy* (6000-fold) in comparison to co-infiltration with *Psy* $\Delta Rrbp$ (80-fold; $p \leq 0.0001$). The differences between co-infiltrated *Pph* minority populations decrease at 6 and 8 dpi, yet the population sizes of co-infiltrated compared to *Pph*-only remained fairly similar to levels at 4 dpi ($p \leq 0.0001$).

When *Psy* minority is co-infiltrated with *Pph* there is a statistical eightfold increase compared to co-infiltrated *Psy* $\Delta Rrbp$ minority at 6 dpi [Figure 2(D); $p \leq 0.0001$]. Both *Psy* minority and *Psy* $\Delta Rrbp$ minority performed equivalently at 4 or 8 dpi, indicating this was the first observation that bacteriocin production is beneficial when faced with a dominant sensitive population. The *Psy* $\Delta Rrbp$ minority presented between a 5- to 10-fold reduction at all time points to *Psy* $\Delta Rrbp$ -only ($p \leq 0.0001$ for all comparisons). The complement *Psy* $\Delta Rrbp$:*Rrbp* minority population level was similar to *Psy* minority indicating the population increase is due to bacteriocin production [Figure S3(B)]. There were no differences between *Psy* majority and *Psy* $\Delta Rrbp$

majority when co-inoculated with *Pph* [Figure 2(D)]. Additionally, *Psy* majority strains exhibited a decrease in population relative to the *Psy*-only infiltration at 4 dpi but maintained roughly equivalent populations at 6 and 8 dpi. Overall, *Psy* starting at a low cell frequency (in the minority) provided a bacteriocin-mediated fitness benefit.

***Psy* virulence is required for *in planta* bacteriocin-mediated effects**

To investigate the role of virulence in pathogen–host interactions, a mutation of the *hrcC* gene, a structural component of the Type III Secretion System (T3SS), was introduced into *Psy*. $\Delta hrcC$ mutants are impaired in their ability to suppress plant defences, and thus incapable of causing disease (Deng et al., 1998; Hirano et al., 1999). The *Pph* population *in planta* was not affected during 1:1 co-infiltration with either *Psy* $\Delta hrcC$ or *Psy* $\Delta Rrbp/\Delta hrcC$ and was able to maintain populations comparable to *Pph*-only infiltration for all dpi [Figure 3(A)]. This was opposite to *in vitro*

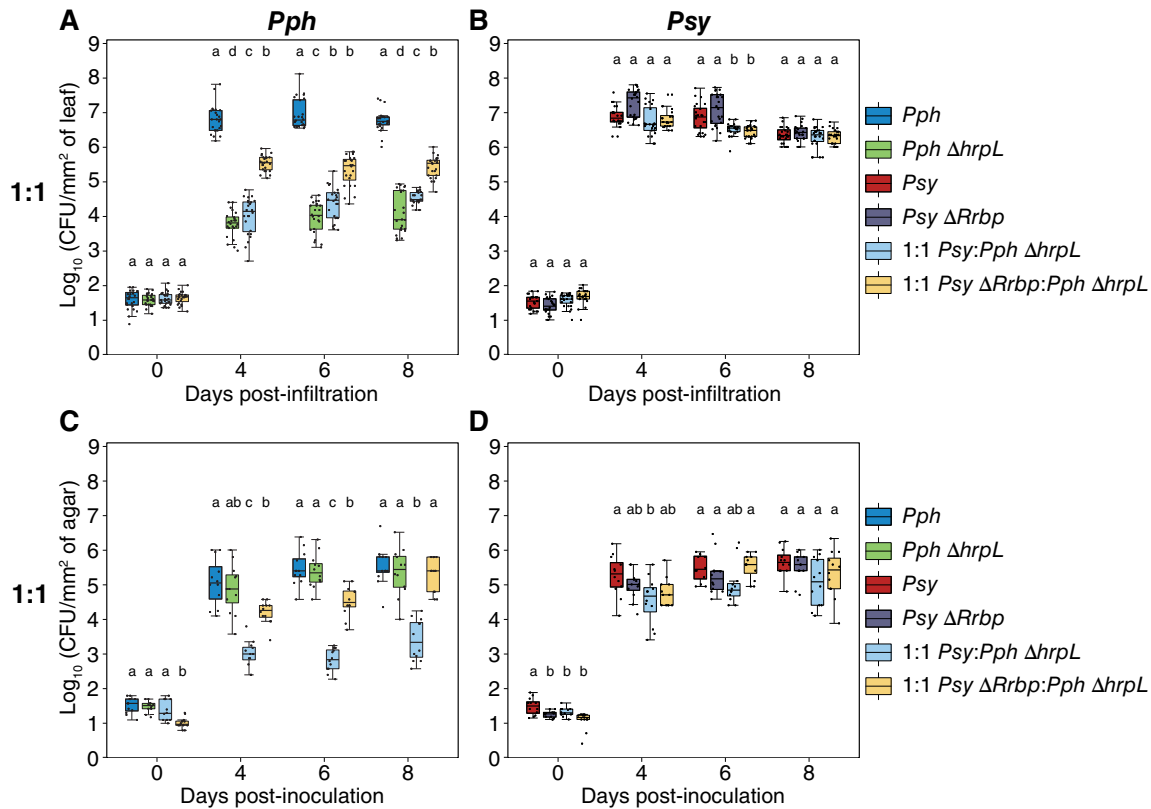


FIGURE 4 No difference in populations of *Psy* strains with avirulent *Pph* $\Delta hrpL$ populations. In 1:1 competition across 8 dpi bacterial growth is shown for (A) *Pph* and (B) *Psy* in planta, and (C) *Pph* and (D) *Psy* in culture (*Psy*:*Pph*). Data points represent eight and four replicates, for *in planta* and *in vitro*, respectively, from three independent experiments over 8 days and error bars indicate standard error of the mean. All treatments were compared by a Tukey's post hoc test (ANOVA) and significant differences are represented by letters ($p \leq 0.05$)

competition, where *Pph* co-inoculated with *Psy* $\Delta hrcC$ at 4 and 6 dpi was reduced by 10-fold compared to the 100-fold reduction at 8 dpi with *Psy* $\Delta Rrbp/\Delta hrcC$ [-Figure 3(C)]. In comparison to *Pph*-only, a 100-fold and 10-fold detriment occurred for *Pph* in co-inoculation with *Psy* $\Delta hrcC$ and *Psy* $\Delta Rrbp/\Delta hrcC$, respectively.

No differences were observed between *Psy* $\Delta hrcC$ and *Psy* $\Delta Rrbp/\Delta hrcC$ populations *in planta* during 1:1 co-infiltration with *Pph* [Figure 3(B)]. Yet, these populations were partially rescued when co-infiltrated with *Pph* by an average increase of 8- to 20-fold at 6 and 8 dpi compared to *Psy* $\Delta hrcC$ -only and *Psy* $\Delta Rrbp/\Delta hrcC$ -only ($p \leq 0.0001$). Both individual and co-infiltrations of *Psy* $\Delta hrcC$ and *Psy* $\Delta Rrbp/\Delta hrcC$ were reduced by 50- to 100-fold from 4 to 8 dpi compared to *Psy*-only infiltration ($p \leq 0.0001$). Whereas *in vitro* competition showed no differences between any individual and mixed *Psy* strains [Figure 3(D)]. These results show that virulence is required for *Psy* to dominate the co-infection environment, as well as to gain a fitness benefit from bacteriocin production.

No bacteriocin-mediated fitness benefit for *Psy* in co-infiltration with avirulent sensitive strain

To identify if *Psy* could gain a fitness benefit in co-inoculations with an avirulent sensitive strain, the *hrpL* gene, required for the activation of the *hrp/hrc* locus responsible for T3SS expression, was knocked-out in *Pph* (Hockett et al., 2015; Ortiz-Martín et al., 2010). The *Pph* $\Delta hrpL$ populations *in planta* were reduced in both individual (1000-fold) and co-infiltrations (between 10- to 100-fold) compared to *Pph*-only from 4 dpi [$p \leq 0.0001$; Figure 4(A)]. For all timepoints the co-infiltration of *Pph* $\Delta hrpL$ with *Psy* resulted in a 9- to 30-fold increase to *Pph* $\Delta hrpL$ -only ($p \leq 0.0001$). However, the population of *Pph* $\Delta hrpL$ co-infiltration with *Psy* was lower than co-infiltration with *Psy* $\Delta Rrbp$ at 4 and 8 dpi ($p \leq 0.0302$). *In vitro* competition resulted in a similar trend with co-inoculated *Pph* $\Delta hrpL$ presenting a 20-fold reduction with *Psy* compared to *Psy* $\Delta Rrbp$ at 4 dpi and reduced by 70- to 80-fold at 6 and 8 dpi [$p \leq 0.0001$; Figure 4(C)]. By 8 dpi, the population

of *Pph* Δ *hrpL* co-inoculated with *Psy* Δ *Rrbp* is equal to *Pph* Δ *hrpL*-only.

The co-infiltration of *Psy* with *Pph* Δ *hrpL* was not statistically different *in planta* to co-infiltrated population of *Psy* Δ *Rrbp*, and both *Psy*-only and *Psy* Δ *Rrbp*-only [Figure 4(B)]. However, *in vitro* co-inoculated *Psy* was statistically reduced by threefold compared to *Psy*-only [$p \leq 0.033$; Figure 4(D)]. Together these results suggest a virulent producer does not benefit from bacteriocin-mediated competition with an avirulence sensitive population.

DISCUSSION

In this study, we sought to understand how interactions with a host plant affect bacteriocin-mediated competition between two bacterial plant pathogens. Overall, our *in vitro* results did not show a bacteriocin-mediated fitness benefit (i.e. an increased population size) for the bacteriocin producer, *Psy*, at any starting frequency when competing with the sensitive strain, *Pph*. Conversely, *in planta* co-infiltrations did show a bacteriocin-mediated fitness benefit for *Psy* when at an initially low frequency. *Pph*, however, suffered from bacteriocin-mediated inhibition during both *in vitro* and *in planta* co-inoculation. Intriguingly, the *in planta* benefit to *Psy* and detriment to *Pph* occurred at specific time points and were not maintained consistently across all time points. Additionally, virulence aided *Psy* bacteriocin-mediated suppression of *Pph* in the plant environment. These results indicate that bacteriocin-mediated interactions within a host plant are influenced by host physiology and pathogen virulence over the course of an infection.

Previous bacteriocin antagonism studies have been performed using computer models or laboratory systems, showing that a fitness benefit for the toxin-producing population is dependent on the environment (Chao & Levin, 1981; Kerr, 2007; Majeed et al., 2011). Our results suggest that when at parity or in the majority there was no bacteriocin-mediated fitness benefit *in vitro* for *Psy* compared to *Psy* Δ *Rrbp* across 8 dpi. It is likely that the sampling of the entire colony is a global measurement of the cumulative effects of local interactions between cells that might mask a fitness benefit that is localized to the colony periphery. The detriment was not a complete elimination of *Pph* as it is hypothesized that primarily sensitive cells at the edges are affected, with the cells near the centre of the microcolony being able to persist (Kerr, 2007). Both *Psy* minority strains are also able to overcome the initial low frequency (regardless of bacteriocin killing) compared to the *Pph* minority indicating *Psy* possesses some additional method of competitive advantage over *Pph*.

Differences in apoplast spatial structure, available resources and host compatibility create a dynamic host environment for two pathogens to compete that is more

complex than an agar plate (Dangl & Jones, 2001; Farvardin et al., 2020; O'Leary et al., 2016; Rico & Preston, 2008). In 1:1 co-infiltration, the outcome of *in planta* competition was similar to the *in vitro* competition where there was no benefit for the bacteriocin producer but there was a bacteriocin mediated detriment to *Pph*, similar to the effects Li et al. (2020) showed for competition between *P. syringae* pv. *tomato* and *P. syringae* pv. *lachrymans* (Li et al., 2020). However, the detriment was not maintained from 6 dpi onwards, potentially indicating the effect of bacteriocin production in the apoplast is limited either by changes in behaviour of *Pph* or changes in the apoplast environment, or both. Previous work has suggested that both sensitive and producing populations are able to coexist through spatial partitioning (Czárán & Hoekstra, 2003; Kerr, 2007), which may occur in the leaf apoplast.

Our results indicate that bacteriocin production is beneficial in a negative-frequency dependent manner, where production is favoured when the population is low (Kerr, 2007; Müller et al., 2019). In our case, this fitness benefit was not observed immediately post-inoculation and occurred once the initial *Psy* minority had a high population level. Importantly, our results showed that the *Pph* population was significantly reduced by bacteriocin-mediated killing at 4 dpi and that the *Psy* population is suppressed when co-inoculated with *Pph* compared to *Psy*-only at the same time point. Taken together, these results indicate that it should have been possible to observe a bacteriocin-mediated benefit for *Psy* at 4 dpi. We also considered the use of the competitive index (CI) to present the fitness benefit, since we have paired populations of *Pph* and *Psy* strains for each treatment. We believe, however, that such calculations would be misleading as the CI will certainly show *Psy* performing better compared to *Psy* Δ *Rrbp*, but this difference would, in nearly all cases, result from less killing of *Pph* by *Psy* Δ *Rrbp* rather than any increase in the *Psy* population compared to *Psy* Δ *Rrbp*. This observation also occurs about the same time that disease symptoms for *Pph* were distinctly identifiable in individual infiltration (e.g. water-soaking and yellowing). Therefore, we hypothesize that during co-infiltration *Pph* can gain greater access to host nutrients at the height of disease progression increasing its population, and the bacteriocin production of *Psy* is able to overcome this growth whereas *Psy* Δ *Rrbp* cannot resulting in the reduction of *Psy* Δ *Rrbp* population at 6 dpi.

Bacterial pathogens must contend with the host plant defences to enable establishment and proliferation of their populations. Pathogenic bacteria can use the T3SS encoded by the *hrp* and *hrc* genes to suppress the plants response in nearby plant cells (Alfano et al., 2000; Arnold et al., 2003). During *in planta* infiltrations both *Psy* Δ *hrcC* and *Psy* Δ *Rrbp*/ Δ *hrcC* populations were greatly reduced compared to *Psy*.

When co-infiltrated, however, T3SS mutant strains received an *in trans* benefit from the virulent *Pph* strain. This has been observed in other work with avirulent strains of *P. syringae* which relied on the proximity of virulent strains to reduce the plant cells effector-triggered immunity (Macho et al., 2007; Omer & Wood, 1969; Rufian et al., 2018). Similarly, *Pph* Δ hrpL benefitted from co-infiltration with virulent *Psy*. There was also no detriment for *Pph* in the presence of *Psy* Δ hrcC, indicating that *Psy* virulence is required for a bacteriocin-mediated effect on *Pph*.

While previous research has focused mainly on the outcomes of bacteriocin-mediated antagonism *in vitro*, we show that such outcomes may not be directly translated into a host plant environment. Our findings show that under certain frequency and temporal conditions bacteriocin production can promote *Psy* fitness while targeting the sensitive strain population. Further research is needed to elucidate the exact spatial distribution of the infiltrated bacteria, such as the use of fluorescent microscopy, alongside measuring the rates of bacteriocin production in the apoplast. Bacteriocin-mediated killing does not necessarily equate to a bacteriocin-mediated fitness benefit. We suggest that this dichotomy applies to past and current biological control research, where the objective typically is to limit the effects of the pathogen but there was no examination of the benefit of the agent to proliferate and maintain present in the field (Fravel, 1988).

ACKNOWLEDGEMENTS

This work was supported by the Indigo Agriculture Phytobiomes Fellowship, the Penn State Microbiome Center, the Fulbright New Zealand Science and Innovation Graduate Award, the Rose Hellaby Postgraduate Scholarship, the Ngārimu VC and 28th (Māori) Battalion Memorial Doctoral Scholarship to H.E. Additional support for K.L.H. came from the USDA National Institute of Food and Agriculture and Federal Hatch Appropriations PEN04648 (accession no. 1016871) and start-up funds through The Huck Institutes for the Life Sciences and the College of Agricultural Sciences at Penn State. Our appreciation to Brian Kvitko from the University of Georgia for gifting pDONR1k18ms and David Baltrus from the University of Arizona for gifting the *Psy* Δ hrcC strain. We acknowledge that The Pennsylvania State University campuses are located on the original homelands of the Erie, Haudenosaunee (Seneca, Cayuga, Onondaga, Oneida, Mohawk and Tuscarora), Lenape (Delaware Nation, Delaware Tribe, Stockbridge-Munsee), Shawnee (Absentee, Eastern and Oklahoma), Susquehannock and Wahzhazhe (Osage) Nations.

CONFLICT OF INTEREST

All authors declare no conflict of interest.

DATA AVAILABILITY STATEMENT

Data available on request from the authors.

ORCID

Hanareia Eha-Taumaunu  <https://orcid.org/0000-0002-6915-4986>

Kevin L. Hockett  <https://orcid.org/0000-0002-0997-4712>

REFERENCES

- Alfano, J.R., Charkowski, A.O., Deng, W.-L., Badel, J.L., Petnicki-Ocwieja, T., van Dijk, K. et al. (2000) The *Pseudomonas syringae* Hrp pathogenicity Island has a tripartite mosaic structure composed of a cluster of type III secretion genes bounded by exchangeable effector and conserved effector loci that contribute to parasitic fitness and pathogenicity in plants. *Proceedings of the National Academy of Sciences of the United States of America*, 97(9), 4856–4861. <https://doi.org/10.1073/PNAS.97.9.4856>
- Arnold, D.L., Pitman, A. & Jackson, R.W. (2003) Pathogenicity and other genomic islands in plant pathogenic bacteria. *Molecular Plant Pathology*, 4(5), 407–420. <https://doi.org/10.1046/J.1364-3703.2003.00187.X>
- Baltrus, D.A., Nishimura, M.T., Romanchuk, A., Chang, J.H. & Mukhtar, M.S. (2011) Dynamic evolution of pathogenicity revealed by sequencing and comparative genomics of 19 *Pseudomonas syringae* isolates. *PLoS Pathogens*, 7(7), 1002132. <https://doi.org/10.1371/journal.ppat.1002132>
- Bogino, P.C., de las Mercedes Oliva, M., Sorroche, F.G. & Giordano, W. (2013) The role of bacterial biofilms and surface components in plant-bacterial associations. *International Journal of Molecular Sciences*, 14(8), 15838–15859. <https://doi.org/10.3390/IJMS140815838>
- Burkholder, W.H. (1926) A new bacterial disease of the bean. *Phytopathology*, 16(12), 915–927.
- Chao, L. & Levin, B.R. (1981) Structured habitats and the evolution of anticompensator toxins in bacteria. *Proceedings of the National Academy of Sciences of the United States of America*, 78(10), 6324–6328.
- Corr, S.C., Li, Y., Riedel, C.U., O'Toole, P.W., Hill, C. & Gahan, C.G. M. (2007) Bacteriocin production as a mechanism for the anti-infective activity of *Lactobacillus salivarius* UCC118. *Proceedings of the National Academy of Sciences of the United States of America*, 104(18), 7617–7621. <https://doi.org/10.1073/pnas.0700440104>
- Czárán, T.L. & Hoekstra, R.F. (2003) Killer-sensitive coexistence in metapopulations of micro-organisms. *Proceedings of the Royal Society B: Biological Sciences*, 270(1522), 1373–1378. <https://doi.org/10.1098/RSPB.2003.2338>
- Dangl, J.L. & Jones, J.D. (2001) Plant pathogens and integrated defence responses to infection. *Nature*, 411(6839), 826–833. <https://doi.org/10.1038/35081161>
- Deng, W.L., Preston, G., Collmer, A., Chang, C.J. & Huang, H.C. (1998) Characterization of the hrpC and hrpRS operons of *Pseudomonas syringae* pathovars *syringae*, *tomato*, and *glycinea* and analysis of the ability of hrpF, hrpG, hrcC, hrpT, and hrpV mutants to elicit the hypersensitive response and disease in plants. *Journal of Bacteriology*, 180(17), 4523–4531. <https://doi.org/10.1128/JB.180.17.4523-4531.1998>
- Dodds, P.N. & Rathjen, J.P. (2010) Plant immunity: towards an integrated view of plant–pathogen interactions. *Nature Reviews Genetics*, 11(8), 539–548. <https://doi.org/10.1038/nrg2812>
- Dorosky, R.J., Pierson, L.S. & Pierson, E.A. (2018) *Pseudomonas chlororaphis* produces multiple R-tailocin particles that broaden the killing spectrum and contribute to persistence in rhizosphere

- communities. *Applied and Environmental Microbiology*, 84(18), e01230-18. <https://doi.org/10.1128/AEM.01230-18>
- Durrett, R. & Levin, S. (1997) Allelopathy in spatially distributed populations. *Journal of Theoretical Biology*, 185(2), 165–171.
- Farvardin, A., González-Hernández, A.I., Llorens, E., García-Agustín, P., Scalschi, L. & Vicedo, B. (2020) The apoplast: a key player in plant survival. *Antioxidants*, 9(7), 1–26. <https://doi.org/10.3390/ANTIOX9070604>
- Feil, H., Feil, W.S., Chain, P., Larimer, F., DiBartolo, G., Copeland, A. et al. (2005) Comparison of the complete genome sequences of *Pseudomonas syringae* pv. *syringae* B728a and pv. *tomato* DC3000. *Proceedings of the National Academy of Sciences of the United States of America*, 102(31), 11064–11069. <https://doi.org/10.1073/pnas.0504930102>
- Fravel, D.R. (1988) Role of antibiosis in the biocontrol of plant diseases. *Annual Review of Phytopathology*, 26(1), 75–91. <https://doi.org/10.1146/ANNUREV.PY.26.090188.000451>
- Ghazaryan, L., Giladi, I. & Gillor, O. (2019) The effects of colicin production rates on allelopathic interactions in *Escherichia coli* populations. *Microorganisms*, 7(11), 564. <https://doi.org/10.3390/microorganisms7110564>
- Ghequire, M.G.K. & De Mot, R. (2014) Ribosomally encoded antibacterial proteins and peptides from *Pseudomonas*. *FEMS Microbiology Reviews*, 38(4), 523–568. <https://doi.org/10.1111/1574-6976.12079>
- Ghequire, M.G.K. & De Mot, R. (2015) The tailocin tale: peeling off phage tails. *Trends in Microbiology*, 23(10), 587–590. <https://doi.org/10.1016/j.tim.2015.07.011>
- Ghoul, M. & Mitri, S. (2016) The ecology and evolution of microbial competition. *Trends in Microbiology*, 24(10), 833–845. <https://doi.org/10.1016/j.tim.2016.06.011>
- Godino, A., Príncipe, A. & Fischer, S. (2016) A ptsP deficiency in PGPR *Pseudomonas fluorescens* SF39a affects bacteriocin production and bacterial fitness in the wheat rhizosphere. *Research in Microbiology*, 167(3), 178–189. <https://doi.org/10.1016/j.resmic.2015.12.003>
- Gordon, D.M. & Riley, M.A. (1999) A theoretical and empirical investigation of the invasion dynamics of colicinogeny. *Microbiology*, 145(3), 655–661. <https://doi.org/10.1099/13500872-145-3-655>
- Granato, E.T., Meiller-Legrand, T.A. & Foster, K.R. (2019) The evolution and ecology of bacterial warfare. *Current Biology*, 29(11), R521–R537. <https://doi.org/10.1016/J.CUB.2019.04.024>
- van Hall, C.J.J. (1902) *Bijdragen tot de kennis der bakterieele plantenziekten*. Amsterdam: University of Amsterdam.
- Hert, A.P., Roberts, P.D., Momol, M.T., Minsavage, G.V., Tudor-Nelson, S.M. & Jones, J.B. (2005) Relative importance of bacteriocin-like genes in antagonism of *Xanthomonas perforans* tomato race 3 to *Xanthomonas euvesicatoria* tomato race 1 strains. *Applied and Environmental Microbiology*, 71(7), 3581–3588. <https://doi.org/10.1128/AEM.71.7.3581-3588.2005>
- Hirano, S.S. & Upper, C.D. (2000) Bacteria in the leaf ecosystem with emphasis on *Pseudomonas syringae*-a pathogen, ice nucleus, and epiphyte. *Microbiology and Molecular Biology Reviews*, 64(3), 624–653.
- Hirano, S.S., Charkowski, A.O., Collmer, A., Willis, D.K. & Upper, C. D. (1999) Role of the Hrp type III protein secretion system in growth of *Pseudomonas syringae* pv. *Syringae* B728a on host plants in the field. *Proceedings of the National Academy of Sciences of the United States of America*, 96(17), 9851–9856.
- Hockett, K.L., Renner, T. & Baltrus, D.A. (2015) Independent co-option of a tailed bacteriophage into a killing complex in *Pseudomonas*. *MBio*, 6(4), e00452. <https://doi.org/10.1128/mBio.00452-15>
- Holtsmark, I., Eijsink, V.G.H. & Brurberg, M.B. (2008) Bacteriocins from plant pathogenic bacteria. *FEMS Microbiology Letters*, 280(1), 1–7. <https://doi.org/10.1111/j.1574-6968.2007.01010.x>
- Inglis, R.F., Gardner, A., Cornelis, P. & Buckling, A. (2009) Spite and virulence in the bacterium *Pseudomonas aeruginosa*. *Proceedings of the National Academy of Sciences of the United States of America*, 106(14), 5703–5707. <https://doi.org/10.1073/pnas.0810850106>
- Joardar, V., Lindeberg, M., Jackson, R.W., Selengut, J., Dodson, R., Brinkac, L.M. et al. (2005) Whole-genome sequence analysis of *Pseudomonas syringae* pv. *phaseolicola* 1448A reveals divergence among pathovars in genes involved in virulence and transposition. *Journal of Bacteriology*, 187(18), 6488–6498. <https://doi.org/10.1128/JB.187.18.6488-6498.2005>
- Kerr, B. (2007) The ecological and evolutionary dynamics of model bacteriocin communities. In: *Bacteriocins*. Berlin, Heidelberg: Springer, pp. 111–134. https://doi.org/10.1007/978-3-540-36604-1_6
- Kommineni, S., Bretl, D.J., Lam, V., Chakraborty, R., Hayward, M., Simpson, P. et al. (2015) Bacteriocin production augments niche competition by *enterococci* in the mammalian gastrointestinal tract. *Nature*, 526(7575), 719–722. <https://doi.org/10.1038/nature15524>
- Li, J., Zhou, L., Peng, Y. & Fan, J. (2020) *Pseudomonas* bacteriocin syringacin M released upon desiccation suppresses the growth of sensitive bacteria in plant necrotic lesions. *Microbial Biotechnology*, 13(1), 134–147. <https://doi.org/10.1111/1751-7915.13367>
- Macho, A.P., Zumaquero, A., Ortiz-Martín, I. & Beuzón, C.R. (2007) Competitive index in mixed infections: a sensitive and accurate assay for the genetic analysis of *Pseudomonas syringae*-plant interactions. *Molecular Plant Pathology*, 8(4), 437–450. <https://doi.org/10.1111/j.1364-3703.2007.00404.x>
- Majeed, H., Gillor, O., Kerr, B. & Riley, M.A. (2011) Competitive interactions in *Escherichia coli* populations: the role of bacteriocins. *The ISME Journal*, 5(1), 71–81. <https://doi.org/10.1038/ismej.2010.90>
- McEvoy, P.B. (1996) Host specificity and biological pest control. *BioScience*, 46(6), 401–405. <https://doi.org/10.2307/1312873>
- Mills, S., Ross, R.P. & Hill, C. (2017) Bacteriocins and bacteriophage; a narrow-minded approach to food and gut microbiology. *FEMS Microbiology Reviews*, 41, S129–S153. <https://doi.org/10.1093/FEMSRE/FUX022>
- Montesinos, E. (2007) Antimicrobial peptides and plant disease control. *FEMS Microbiology Letters*, 270(1), 1–11. <https://doi.org/10.1111/j.1574-6968.2007.00683.x>
- Morris, C.E. & Monier, J.-M. (2003) The ecological significance of bio-film formation by plant-associated bacteria. *Annual Review of Phytopathology*, 41, 429–453.
- Morris, C.E., Lamichhane, J.R., Nikolić, I., Stanković, S. & Moury, B. (2019) The overlapping continuum of host range among strains in the *Pseudomonas syringae* complex. *Phytopathology Research*, 1(4), 1–16. <https://doi.org/10.1186/s42483-018-0010-6>
- Müller, J., Spriewald, S., Stecher, B., Stadler, E. & Fuchs, T.M. (2019) Evolutionary stability of *Salmonella* competition with the gut microbiota: how the environment fosters heterogeneity in exploitative and interference competition. *Journal of Molecular Biology*, 431(23), 4732–4748. <https://doi.org/10.1016/j.jmb.2019.06.027>
- Nedialkova, L.P., Denzler, R., Koeppel, M.B., Diehl, M., Ring, D., Wille, T. et al. (2014) Inflammation fuels colicin lb-dependent competition of *Salmonella* Serovar *Typhimurium* and *E. coli* in Enterobacterial blooms. *PLoS Pathogens*, 10(1), e1003844. <https://doi.org/10.1371/journal.ppat.1003844>
- O’Leary, B.M., Neale, H.C., Geilfus, C.M., Jackson, R.W., Arnold, D. L. & Preston, G.M. (2016) Early changes in apoplast composition associated with defence and disease in interactions between *Phaseolus vulgaris* and the halo blight pathogen *Pseudomonas syringae* pv. *phaseolicola*. *Plant Cell and Environment*, 39(10), 2172–2184. <https://doi.org/10.1111/pce.12770>
- Omer, M.E.H. & Wood, R.K.S. (1969) Growth of *Pseudomonas phaseolicola* in susceptible and in resistant bean plants. *Annals of Applied Biology*, 63(1), 103–116. <https://doi.org/10.1111/j.1744-7348.1969.tb05471.x>

- Ortiz-Martín, I., Thwaites, R., Macho, A.P., Mansfield, J.W. & Beuzón, C.R. (2010) Positive regulation of the Hrp type III secretion system in *Pseudomonas syringae* pv. *phaseolicola*. *MPMI*, 23(5), 665–681. <https://doi.org/10.1094/MPMI>
- Rico, A. & Preston, G.M. (2008) *Pseudomonas syringae* pv. *tomato* DC3000 uses constitutive and apoplast-induced nutrient assimilation pathways to catabolize nutrients that are abundant in the tomato apoplast. *Molecular Plant-Microbe Interactions*, 21(2), 269–282. <https://doi.org/10.1094/MPMI-21-2-0269>
- Riley, M.A. & Chavan, M.A. (2007) *Bacteriocins: ecology and evolution*. Berlin, Heidelberg: Springer-Verlag.
- Riley, M.A. & Gordon, D.M. (1999) The ecological role of bacteriocins in bacterial competition. *Trends in Microbiology*, 7(3), 129–133. [https://doi.org/10.1016/S0966-842X\(99\)01459-6](https://doi.org/10.1016/S0966-842X(99)01459-6)
- Riley, M.A. & Wertz, J.E. (2002) Bacteriocins: evolution, ecology, and application. *Annual Review of Microbiology*, 56(1), 117–137. <https://doi.org/10.1146/annurev.micro.56.012302.161024>
- Rufian, J.S., Macho, A.P., Corry, D.S., Mansfield, J.W., Ruiz-Albert, J., Arnold, D.L. et al. (2018) Confocal microscopy reveals in planta dynamic interactions between pathogenic, avirulent and non-pathogenic *Pseudomonas syringae* strains. *Molecular Plant Pathology*, 19(3), 537–551. <https://doi.org/10.1111/mpp.12539>
- Sassone-Corsi, M., Nuccio, S.P., Liu, H., Hernandez, D., Vu, C.T., Takahashi, A.A. et al. (2016) Microcins mediate competition among *Enterobacteriaceae* in the inflamed gut. *Nature*, 540(7632), 280–283. <https://doi.org/10.1038/nature20557>
- Scholl, D. (2017) Phage tail-like bacteriocins. *Annual Review of Virology*, 4(1), 453–467. <https://doi.org/10.1146/annurev-virology-101416-041632>
- Simon, R., Priefer, U. & Pühler, A. (1983) A broad host range mobilization system for *in vivo* genetic engineering: transposon mutagenesis in gram negative bacteria. *Bio/Technology*, 1(9), 784–791. <https://doi.org/10.1038/nbt1183-784>
- Singh, A. & Singh, I.K. (2018) *Molecular aspects of plant-pathogen interaction*. Singapore: Springer Singapore, pp. 1–351. <https://doi.org/10.1007/978-981-10-7371-7>
- Weber, M.F., Poxleitner, G., Hebisch, E., Frey, E. & Opitz, M. (2014) Chemical warfare and survival strategies in bacterial range expansions. *Journal of the Royal Society*, 11(96), 20140172. <https://doi.org/10.1098/rsif.2014.0172>
- Xin, X.-F., Kvitko, B. & He, S.Y. (2018) *Pseudomonas syringae*: what it takes to be a pathogen. *Nature Reviews Microbiology*, 16(5), 316–328. <https://doi.org/10.1038/nrmicro.2018.17>
- Yu, H., Wang, Y., Zeng, X., Cai, S., Wang, G., Liu, L. et al. (2020) Therapeutic administration of the recombinant antimicrobial peptide microcin J25 effectively enhances host defenses against gut inflammation and epithelial barrier injury induced by enterotoxigenic *Escherichia coli* infection. *The FASEB Journal*, 34(1), 1018–1037. [10.1096/FJ.201901717R](https://doi.org/10.1096/FJ.201901717R)

SUPPORTING INFORMATION

Additional supporting information can be found online in the Supporting Information section at the end of this article.

How to cite this article: Eha-Taumaunu, H. & Hockett, K.L. (2022) The plant host environment influences competitive interactions between bacterial pathogens. *Environmental Microbiology Reports*, 14(5), 785–794. Available from: <https://doi.org/10.1111/1758-2229.13103>