

## Article

# Evaluation of the Antifungal Potential of Different *Photorhabdus* Species Against *Monilinia laxa* and *Colletotrichum fioriniae*

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## Abstract

*Monilinia laxa* and *Colletotrichum fioriniae* are major fungal pathogens causing brown rot and anthracnose in stone fruits and shell fruits, leading to significant economic losses. Chemical fungicides are widely applied but can result in resistance development, environmental contamination, and food safety concerns. Biological control using entomopathogenic bacteria (EPB) of the genus *Photorhabdus* has emerged as an eco-friendly alternative. This study evaluated the in vitro antifungal activity of selected *Photorhabdus* species (*P. kayaii* 1723B, *P. temperata* 3017, *P. cinerea* 3086, *P. laumondii* 3196, and *P. thracensis* 3210) against *M. laxa* (M3) and *C. fioriniae* (VV081) using drop-to-drop confrontation and poisoned agar assays. Effects of fermentation time, preparation mode (original vs. centrifuged and filtered), and concentration (5, 10, 20%) were examined. Species-specific inhibition was observed, with Median Inhibition Index values indicated relatively higher antifungal activity for *P. thracensis* 3210 against *M. laxa* (0.718) and *C. fioriniae* (0.552), followed by *P. cinerea* 3086 (0.643 and 0.552) and *P. kayaii* 1723B (0.629 and 0.541). Fermentation time and preparation mode influenced antifungal activity in a strain-dependent manner, with longer fermentation periods and original culture preparations generally showing stronger inhibitory trends. Higher concentrations, especially 20%, were often associated with increased inhibition, although the magnitude of these effects varied among strain-pathogen combinations. Overall, these findings demonstrate that the strain- and pathogen-specific nature of antifungal responses in *Photorhabdus*, supporting their potential as components of targeted biological control strategies rather than uniform broad-spectrum agents.

**Keywords:** *Photorhabdus*; entomopathogenic bacteria; *Monilinia laxa*; *Colletotrichum fioriniae*; antifungal effect



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## 1. Introduction

*Monilinia* spp. are necrotrophic phytopathogenic fungi belonging to the phylum Ascomycota and are the causal agents of brown rot disease (BRD), one of the most destructive diseases of stone fruits such as cherry, sour cherry, peach, and plum [1]. BRD affects fruits, twigs, leaves, and buds, leading to severe yield and postharvest losses. Disease development is favored by warm temperatures, high humidity, and rainfall, and BRD has been reported across almost all continents, including Europe, America, Africa, and Australia [2–4]. *Monilinia laxa* survives in mummified fruits and infected plant tissues, serving as a primary inoculum source for new infections during the following growing season [5].

Anthrachnose is another economically important disease affecting fruit crops, including walnut and stone fruits, and is mainly caused by species of the genus *Colletotrichum*. Among these, *Colletotrichum godetiae* and *C. fiorinae* have been identified as dominant pathogens in Europe, where yield losses of up to 70% have been reported [6]. These fungi are highly adaptable and capable of causing fruit rot and lesions on multiple hosts [7]. Under favorable environmental conditions, *Colletotrichum* conidia spread primarily through rainfall and splashing, resulting in significant economic damage to various fruit crops [8,9].

Management of *Monilinia* and *Colletotrichum* species has traditionally relied on chemical fungicides, including sterol biosynthesis inhibitors (triazoles), benzimidazoles, and contact fungicides [10–12]. However, the widespread and repeated use of these chemicals has led to the development of fungicide resistance, particularly in *M. laxa*, posing a serious challenge for sustainable disease control [13]. In addition, excessive pesticide use contributes to food contamination, environmental degradation, and aquatic ecosystem pollution [14]. Notably, *Monilinia* isolates from fungicide-treated orchards have been shown to exhibit broader fungicide tolerance compared to those from untreated areas [15].

Due to these concerns, biological control has gained increasing attention as an environmentally friendly alternative. Several biological control agents have shown promising results against *Monilinia* species, particularly *Bacillus* spp. and antagonistic fungi. For example, *Bacillus subtilis* and *B. amyloliquefaciens* have been reported to suppress *Monilinia* infections and enhance fruit tolerance [16,17]. Other microbial antagonists, including *Aureobasidium pullulans* and *Trichoderma* species (*T. viride*, *T. harzianum*, and *T. longibrachiatum*), have also been successfully incorporated into integrated pest management strategies [18,19]. Commercial biopesticides such as Serenade and BioPK are already in use for controlling *Monilinia* spp. [20]. For the management of *C. fiorinae*, chemical control using demethylation-inhibiting (DMI) fungicides such as difenoconazole and propiconazole has been reported as highly effective, although their efficacy may vary depending on cultivar and environmental conditions [21,22]. In addition, biocontrol agents such as *Bacillus subtilis*, producing volatile organic compounds, have shown strong inhibitory effects on the mycelial growth of *C. fiorinae*, providing a promising non-chemical alternative [23].

Entomopathogenic bacteria (EPB) of the genera *Photorhabdus* and *Xenorhabdus* are obligate symbionts of entomopathogenic nematodes and are known for producing a wide range of bioactive secondary metabolites [24–26]. These bacteria secrete extracellular toxins, enzymes, and antimicrobial compounds that inhibit competing microorganisms [27–29]. Numerous in vitro studies have demonstrated their antifungal activity against major plant pathogens such as *Botrytis cinerea*, *Fusarium oxysporum*, *Rhizoctonia solani*, and *Sclerotinia sclerotiorum* [30–33]. However, studies targeting *Monilinia* and *Colletotrichum* species remain limited, despite some promising reports showing strong antifungal effects of EPB-derived metabolites [34–41].

Despite these findings, information on the antifungal efficacy of EPB-derived metabolites against *M. laxa* and *C. fiorinae* remains scarce. Therefore, the present study aimed to evaluate the in vitro antifungal activity of crude and bacterial preparations derived from selected *Photorhabdus* species against these economically important pathogens. Bacterial strains obtained from *P. thracensis*, *P. cinerea*, *P. kayaii*, and *P. laumondii* were tested using dual-culture confrontation and poisoned agar assays under different experimental conditions. By comparing fungal growth inhibition across fermentation times, concentrations, and preparation methods, this study seeks to identify effective EPB-based strategies for sustainable biological control.

## 2. Materials and Methods

### 2.1. Sources of EPB and Fungal Isolate

Bacterial species were selected from the microbiological strain collection at Újfehértó Research Station of University of Debrecen (national collection identifier HUNGB810). EPB strains of the collection were isolated from Hungary [42]. The collection is maintained as cryopreserved material (Table 1).

**Table 1.** List of bacterial species and their corresponding strain codes.

Species	Strain Code
<i>Photorhabdus cinerea</i>	3026, 3086, 3176, 3186
<i>Photorhabdus kayaii</i>	3167, 3209, 1723B
<i>Photorhabdus laumondii</i>	3196, 1720B
<i>Photorhabdus temperata</i>	3017, 3047, 3179
<i>Photorhabdus thracensis</i>	3210

Fungal pathogen strains *Colletotrichum fioriniae* (VV081) and *Monilinia laxa* (M3) were selected from the strain collection of the Research Centre for Fruit Growing (Institute of Horticultural Science, Érd, Hungary). The strains were isolated by Prof. László Vajna and Virág Varjas, and are maintained as slant agar culture at 4 °C.

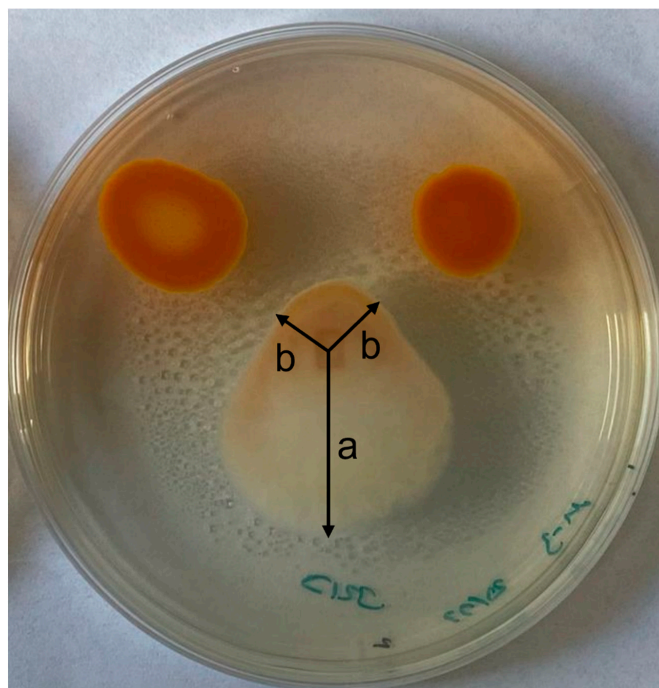
### 2.2. Production of Bacterial Suspension

To obtain bacterial suspension containing natural products, bacteria were cultured on Tryptic Soy Agar (TSA) plates. The TSA medium was prepared using 17 g Tryptone, 3 g Soy Peptone, 2.5 g Glucose, 2.5 g di-Potassium Hydrogen Phosphate, 5 g Sodium Chloride, and 15 g Bacteriological Agar (all from Biolab Inc., Hungary) per 1000 mL of distilled water. The plates were incubated at 25 °C for 48 h. A single colony was then collected from these plates and inoculated into 100 mL Erlenmeyer flasks containing 25 mL of sterile Tryptic Soy Broth (TSB). The TSB medium was prepared using 17 g Tryptone, 3 g Soy Peptone, 5 g Sodium Chloride, 2.5 Glucose, and 2.5 g di-Potassium Hydrogen Phosphate per 1000 mL of distilled water. The cultures were then incubated in a shaker (25 °C, 120 rpm, in the dark). Based on preliminary experiments conducted with the bacterial strains used in these experiments, under the specified conditions (25 mL of culture medium in a 100 mL Erlenmeyer flask, 25 °C, 120 rpm), the bacterial CFU was  $0.8\text{--}1.2 \times 10^9$  in all cases after 48 h. Longer incubation (72 h) did not increase the CFU, meaning that the culture entered a stationary phase.

### 2.3. Direct Interaction Between Bacterial and Fungal Strains (Drop to Drop Technique)

Potato Dextrose Agar (PDA) was prepared following the manufacturer's instructions (Biolab Inc., Hungary) and sterilized at 121 °C for 15 min, and poured in 90 mm plastic Petri dishes. After solidification, a 5-mm mycelial plug was aseptically taken from actively growing fungal cultures of *Monilinia laxa* and *Colletotrichum fioriniae* using a sterile loop and placed at the center of each Petri dish. EPB strains were cultured as described in Section 2.2. and were applied by placing 10 µL (left) and 20 µL (right) drops with an automatic micropipette onto the agar surface near the fungal plug. Radial mycelial growth toward the bacterial colony (the 'treatment' direction, 'b' in Figure 1), and toward the opposite side of the fungal colony (the 'control' direction 'a' in Figure 1) was monitored and measured daily for six days. The inhibition index was calculated as  $1-b/a$ , where b was calculated as the mean of measurements taken from two opposite points toward the

bacterial colony. The index value was 0 if no inhibitory effect was observed, and 1 if the fungal mycelia did not grow at all in the direction of the bacterial colony.



**Figure 1.** Image illustrating the interaction between bacteria and fungal colonies (*Photorhabdus thracensis* strain 3210 vs. *Monilinia laxa* strain M3). The inhibitory effect of bacteria on the growth of fungal colonies is expressed by the Inhibition Index  $(1 - /a)$ .

#### 2.4. Effect of Living-Cell-Free Bacterial Preparation on Fungal Growth (Poisoned Agar Technique)

##### 2.4.1. Production of Living-Cell Free-Bacterial Preparation

EPB species were initially transferred from well-growing stock cultures maintained on Tryptic Soy Agar (TSA) into 25 mL of Tryptic Soy Broth (TSB) and incubated on a GFL 3005 orbital shaker at 25 °C in the dark, initially at 150 rpm for 24 h. Two types of bacterial preparations were obtained: (i) Original (O)—the crude fermentation broth used directly without centrifugation or filtration, and (ii) Centrifuged and Filtered (CF)—the bacterial preparation obtained by centrifugation followed by sterile filtration through a 0.22 µm membrane filter. Both preparations were incorporated into PDA medium prior to autoclaving. Autoclaving was intentionally applied to ensure complete inactivation of bacterial cells while preserving heat-stable secondary metabolites. This approach allowed the evaluation of antifungal activity attributable solely to bacterial metabolites and ensured experimental consistency and sterility across treatments.

##### 2.4.2. Effect of Heat Treatment of Bacterial Preparations

In this study, we aimed to evaluate the antifungal activity of bacterial preparations devoid of viable cells. As heat treatment represents an effective approach for eliminating bacterial viability, heat-treated preparations were employed throughout the experiments. To assess whether heat treatment affected the efficacy of the bacterial preparations, a comparative experiment was conducted using cell-free preparations from multiple bacterial strains.

The PDA medium was prepared according to the manufacturer's instructions. 48 h old bacterial cultures were centrifuged (10,000 rpm, 5 min) and the supernatants were filtered through sterile membrane filters (0.2 µm, FisherBrand X50) to remove bacterial cells. The resulting cell-free preparations were incorporated into the PDA medium at final

concentrations of 10% and 20% (*v/v*) under two different conditions: before autoclaving and after autoclaving.

For the heat-treated preparations, the centrifuged and filtered bacterial preparations were added to the PDA medium prior to autoclaving, and the entire medium was then heat-sterilized (121 °C, 20 min.). For the non-heated treatments, the centrifuged and filtered preparations were added aseptically to the PDA medium after autoclaving, once the medium had cooled to approximately 50–55 °C. The amended media were mixed thoroughly, poured into Petri dishes, and allowed to solidify before inoculation with the fungal isolates.

#### 2.4.3. Dose Effect

To evaluate the dose-dependent antifungal activity of bacterial metabolites, fermentation broths of EPB species were prepared by incubating the cultures in TSB for 24 and 48 h at 25 °C under shaking conditions (GFL 3005 orbital shaker, Burgwedel, Germany, 180 rpm, in the dark). Crude fermentation broth (O) was incorporated into the PDA medium at different concentrations: 0% (control), 5%, 10%, and 20%.

Following medium solidification, a 5-mm mycelial plug was aseptically taken from actively growing cultures of *M. laxa* and *C. fioriniae* using a sterile loop and placed at the center of each Petri dish. Mycelial growth was monitored and measurements were recorded on the sixth day after inoculation. The inhibitory effect of the bacterial material on fungal growth was expressed by an Inhibition Index similar to that defined in Section 2.3, but 'a' was the diameter of fungal colony grown on control plates (without bacterial product), and 'b' was the diameter of the fungal colony grown on PDA plates amended with bacterial product.

#### 2.4.4. Effect of Bacterial Fermentation Time

To investigate the impact of bacterial fermentation duration on antifungal activity, selected *Photorhabdus* strains (*P. cinerea*-3086, *P. thracensis*-3210, and *P. kayaii*-1723B) were cultured in 20 mL of TBA-modified TSB medium under shaking conditions described in Section 2.2 (25 °C, 150 rpm, in the dark) for three different time intervals: 24, 48, and 72 h. For each time point, one type of preparation was generated: original (O) (crude culture) used without processing. The preparation was incorporated into PDA medium by mixing 5 mL of the bacterial sample with 1.95 g of PDA powder and 45 mL of distilled water, followed by autoclaving. This procedure ensured bacterial cell inactivation while retaining heat-stable metabolites, allowing a standardized assessment of metabolite-mediated antifungal effects. After cooling and solidification, Petri dishes were inoculated with a 5-mm mycelial plug taken from actively growing cultures of *M. laxa* (M3) or *C. fioriniae* (VV081). Radial mycelial growth was measured on the sixth day after inoculation to determine the effects of fermentation duration and treatment type on fungal development.

#### 2.5. Statistical Analysis

The Inhibition Index (II) was calculated separately for each Petri dish, and each plate was treated as one independent experimental unit for all statistical analyses. Normality and homogeneity of variance were assessed for each dataset using the Shapiro–Wilk and Levene's tests using R version 4.5.0 (2025-04-11 UCRT). As the datasets did not meet the assumptions of normality, non-parametric statistical analyses were applied. Overall differences among groups were evaluated using the Kruskal–Wallis test, and for datasets related to fermentation and concentration dependency, statistically significant outcomes were further examined via Dunn's post hoc test with Bonferroni correction. For the Preparation Mode dataset, pairwise comparisons were conducted using the Mann–Whitney U test. In the case of the Drop-to-Drop assay, only the Kruskal–Wallis test was applied; since the re-

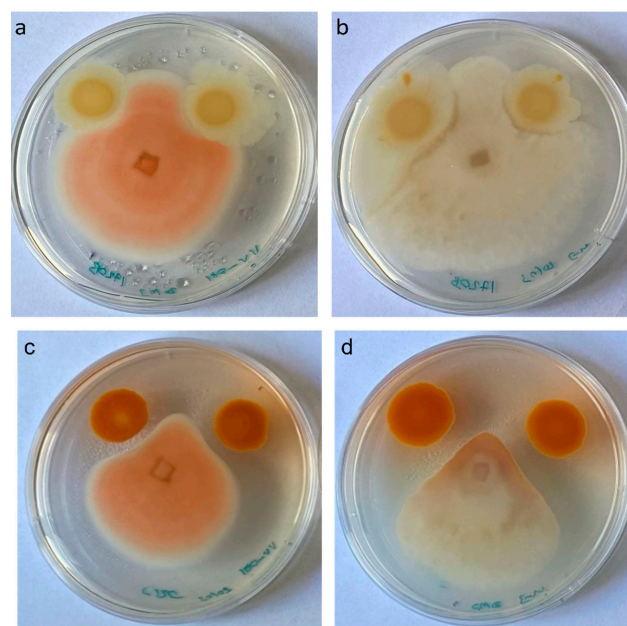
sulting  $p$ -value was below 0.05, non-parametric post hoc tests were subsequently performed to identify significant differences between groups. Data import and preprocessing were performed with the `readxl` and `dplyr` packages, while graphical representations—including bar plots with error bars and faceted layouts—were generated using `ggplot2`, and multiple plots were arranged as required using `patchwork`.

### 3. Results

#### 3.1. Pre-Selection

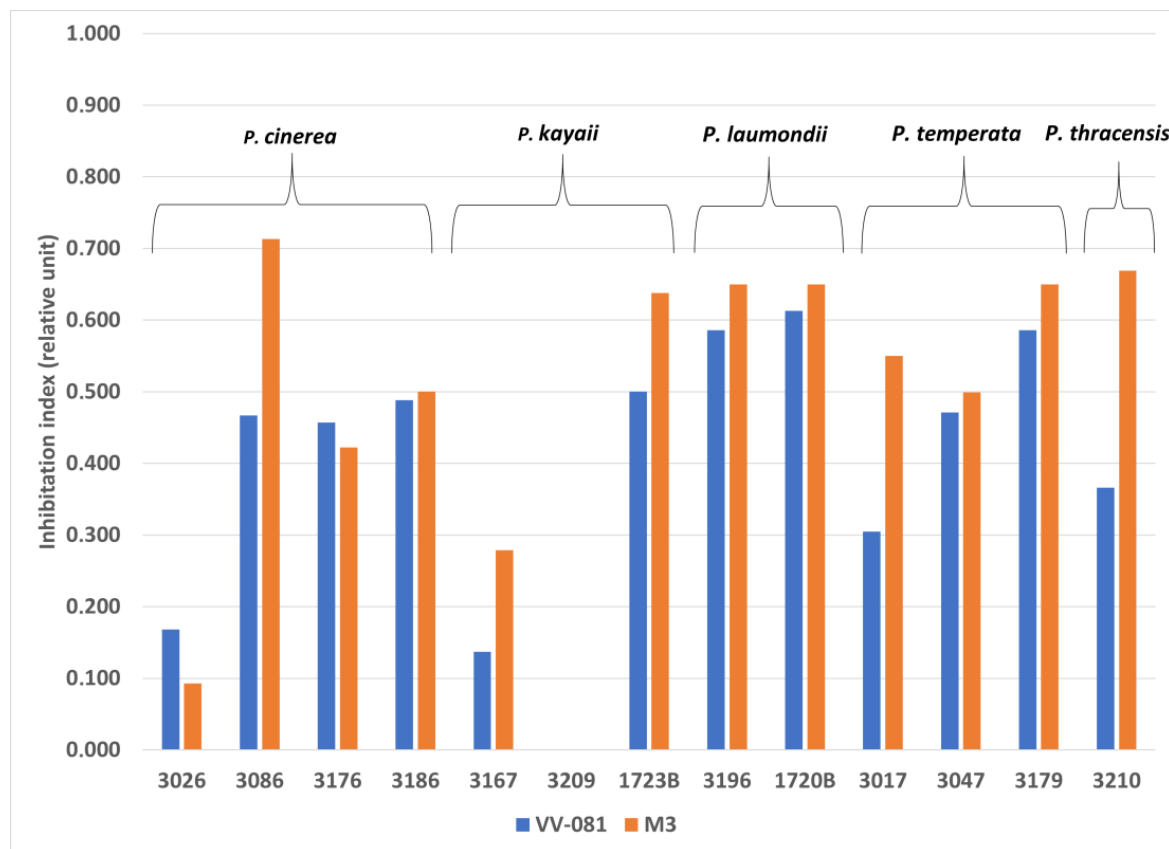
For the pre-selection process, a total of 13 bacterial species from five *Photorhabdus* species were initially considered: four strains from *P. cinerea* (strains 3026, 3086, 3176, and 3186), three from *P. kayaii* (strains 3167, 3209, and 1723B), two from *P. laumondii* (3196 and 1720B), three from *P. temperata* (3017, 3047, and 3179), and one from *P. thracensis* (3210). Based on the inhibition results, only five strains were selected as the most effective against the two fungal isolates (*M. laxa* M3 and *C. fioriniae* VV081). Each bacterial strain was tested against two fungal strains, and each bacterium–fungus combination was performed with three technical replicates. All replicates originated from the same bacterial culture, and therefore represented technical rather than biological replicates.

Strain selection was guided by predefined experimental objectives and was based on reproducible inhibition patterns observed across independent replicates, rather than inhibition index values alone. For instance, although the inhibition indices of strains *P. laumondii* 1720B and *P. kayaii* 1723B were similar, 1720B was not prioritized for subsequent analyses because fungal growth restriction occurred primarily at sites of direct bacterial–fungal contact, suggesting a contact-dependent interaction. (Figure 2). Strains were preferred when they could inhibit fungal growth without extensive physical contact, reflecting a presumed chemical compound-based antifungal effect. Such contact-mediated antagonism is recognized as a legitimate biocontrol mechanism; however, the primary objective of the present study was to prioritize diffusible, contact-independent antifungal activity with greater relevance for living-cell-free preparations and formulation-based applications.



**Figure 2.** Pre-selection process for *Photorhabdus* species based on inhibition and colony interaction *P. laumondii* 1720B against *C. fioriniae* VV081 (a), and *M. laxa* M3 (b). Interaction *P. thracensis* 3210 against *C. fioriniae* VV081 (c), and *M. laxa* M3 (d).

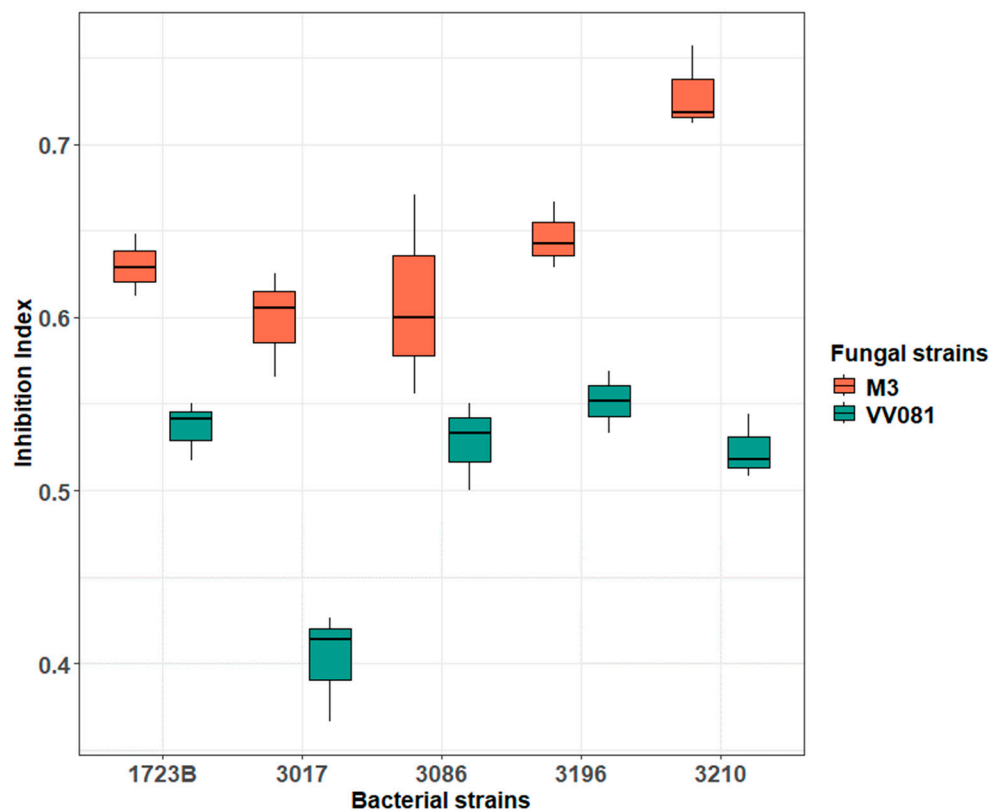
Among the tested strains within each species, the inhibitory activity varied. Therefore, one representative strain per species was selected for subsequent experiments: 1723B (*P. kayaii*), 3017 (*P. temperata*), 3086 (*P. cinerea*), 3196 (*P. laumondii*), and 3210 (*P. thracensis*) (Figure 3).



**Figure 3.** Inhibition Index values of various *Photorhabdus* bacteria against *M. laxa* M3 and *C. fioriniae* VV081 fungal strains.

### 3.2. Drop-to-Drop Assay

The Kruskal-Wallis test indicated a borderline effect of bacterial species on the inhibition of the *M. laxa* M3 fungal isolate ( $\chi^2 = 9.12$ ,  $df = 4$ ,  $p = 0.058$ ). Median inhibition indices varied among strains, with *P. thracensis* 3210 showing the highest inhibition (0.718) and *P. cinerea* 3086 the lowest (0.600). Strains *P. laumondii* 3196 and *P. kayaii* 1723B also demonstrated relatively high inhibition (0.643 and 0.629, respectively), while *P. temperata* 3017 displayed a moderate effect (0.606) (Figure 4). These results suggest a tendency for differences in antifungal efficacy among strains, although variability in the data prevented statistical significance at the 0.05 level. Notably, *P. thracensis* 3210 consistently showed the strongest inhibition, indicating a potentially higher biocontrol potential against *M. laxa* M3. Overall, these results indicate a trend toward strain-dependent differences in antifungal activity, although variability in the data prevented statistical significance at the  $p < 0.05$  level. Five bacterial species were tested against two fungal strains. Each bacterium–fungus combination was tested with three technical replicates and the experiment was independently repeated at two different time points, which were considered as biological replicates.



**Figure 4.** Inhibition Index values of one representative strain from each of the five *Photorhabdus* species against *M. laxa* M3 and *C. fioriniae* VV081 fungal strains. See Table 1 for the designation of bacterial strains.

For the *C. fioriniae* VV081 fungal isolate, the Kruskal-Wallis test again indicated a borderline effect of bacterial strain ( $\chi^2 = 8.84$ ,  $df = 4$ ,  $p = 0.065$ ). Median inhibition indices ranged from 0.414 (*P. temperata* 3017) to 0.552 (*P. laumondii* 3196), with *P. kayaii* 1723B, *P. cinerea* 3086, and *P. thracensis* 3210 showing intermediate values (0.541, 0.533, and 0.518, respectively). The data suggest that *P. laumondii* 3196 and *P. thracensis* 3210 may provide relatively higher inhibition of *C. fioriniae* VV081, whereas *P. temperata* 3017 exhibits the lowest effect (Figure 4). As observed for *M. laxa* M3, the data suggest strain-specific trends in antifungal activity, without reaching conventional statistical significance.

When comparing *M. laxa* M3 and *C. fioriniae* VV081, some strains (notably *P. thracensis* 3210) consistently inhibited fungal growth across both isolates, whereas others (e.g., *P. temperata* 3017) were less effective (Figure 4). The drop-to-drop assay enabled the assessment of direct bacterial–fungal interactions under localized contact conditions, providing a comparative framework to identify strains with consistent inhibitory effects.

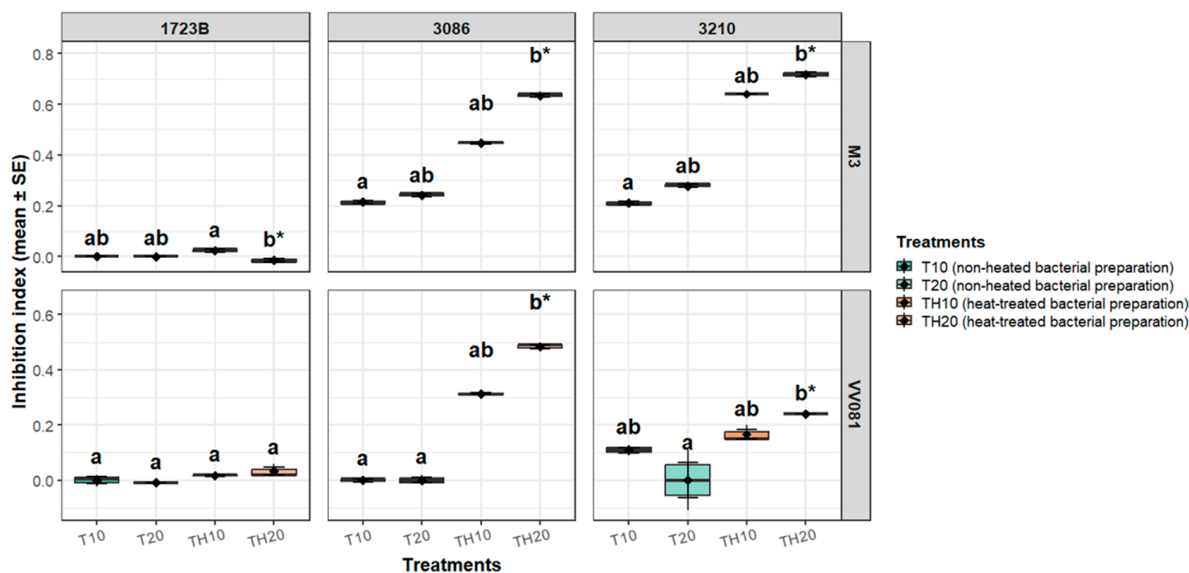
### 3.3. Effect of Heat Treatment of Bacterial Preparations on Fungal Growth Inhibition

Normality of the data distribution was assessed using the Shapiro–Wilk test. Both datasets significantly deviated from normality (heat-treated:  $W = 0.8866$ ,  $p = 0.001495$ ; non-heat treated:  $W = 0.8378$ ,  $p = 0.000101$ ). Therefore, all subsequent analyses were performed using non-parametric statistical tests.

Regarding heat-treated preparations (TH), the Kruskal–Wallis test revealed highly significant differences among bacterial strains for both fungi (M3 *M. laxa*:  $\chi^2 = 14.3$ ,  $df = 2$ ,  $p = 0.000769$ ; VV081 *C. fioriniae*:  $\chi^2 = 15.4$ ,  $df = 2$ ,  $p = 0.000446$ ). At the strain level, comparisons between TH10 and TH20 showed significant dose-dependent effects in five out of six bacterial–fungal combinations. Significant differences were detected for strain



1723B *P. kayaii* against M3 *M. laxa* ( $\chi^2 = 4.09$ ,  $df = 1$ ,  $p = 0.0431$ ), for strain 3086 *P. cinerea* against both M3 *M. laxa* ( $\chi^2 = 4.09$ ,  $df = 1$ ,  $p = 0.0431$ ) and VV081 *C. fioriniae* ( $\chi^2 = 4.09$ ,  $df = 1$ ,  $p = 0.0431$ ), and for strain 3210 *P. thracensis* against both M3 *M. laxa* ( $\chi^2 = 4.35$ ,  $df = 1$ ,  $p = 0.0369$ ) and VV081 *C. fioriniae* ( $\chi^2 = 4.50$ ,  $df = 1$ ,  $p = 0.0339$ ). Dunn’s post hoc test confirmed significant differences between TH10 and TH20 in these combinations (adjusted  $p = 0.0339$ – $0.0431$ ). In contrast, no significant difference was observed for strain 1723B *P. kayaii* against VV081 *C. fioriniae* ( $\chi^2 = 1.67$ ,  $df = 1$ ,  $p = 0.197$ ) (Figure 5).



**Figure 5.** Effect of heat-treated and non-heated bacterial preparations of different *Potorhabdus* strains against *M. laxa* M3 and *C. fioriniae* VV081, expressed as inhibition index. T10 and TH10 indicate a 10% concentration, while T20 and TH20 indicate a 20% concentration. Bars represent mean  $\pm$  SE. See Table 1 for the designation of bacterial strains. Different letters indicate statistically significant differences among treatments within each panel. Asterisks indicate significant pairwise differences (Dunn’s post hoc test with Bonferroni adjustment; \*  $p < 0.05$ ).

Regarding non-heated preparations (T), significant differences among bacterial strains were detected only for M3 *M. laxa* ( $\chi^2 = 11.9$ ,  $df = 2$ ,  $p = 0.00261$ ), whereas no overall effect was observed for VV081 *C. fioriniae* ( $\chi^2 = 3.19$ ,  $df = 2$ ,  $p = 0.203$ ). At the strain level, significant differences between T10 and T20 were found only for strain 3086 *P. cinerea* against M3 *M. laxa* ( $\chi^2 = 4.09$ ,  $df = 1$ ,  $p = 0.0431$ ) and strain 3210 *P. thracensis* against M3 *M. laxa* ( $\chi^2 = 4.09$ ,  $df = 1$ ,  $p = 0.0431$ ). Dunn’s post hoc test supported these results (adjusted  $p = 0.0431$  for both). No significant dose-dependent effect was detected for any strain against VV081 *C. fioriniae* or for strain 1723B *P. kayaii* against either fungus ( $p > 0.18$ ) (Figure 5).

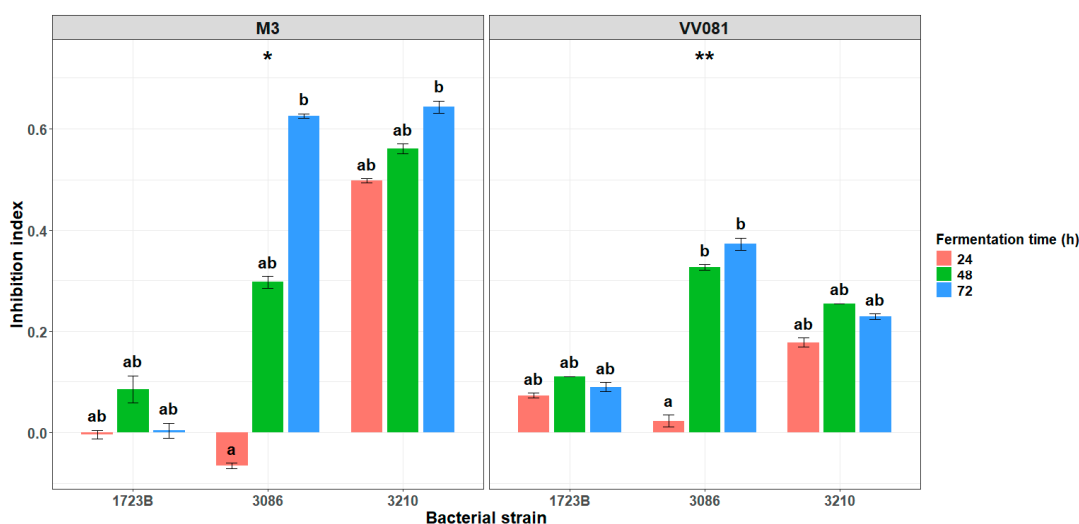
Direct comparisons between heat-treated and non-heated treatments using paired Wilcoxon tests demonstrated that application mode significantly affected antifungal activity for strains 3086 *P. cinerea* and 3210 *P. thracensis* against both fungal isolates. Significant differences were observed for 3086 *P. cinerea* against M3 *M. laxa* ( $W = 21$ ,  $p = 0.0355$ ) and VV081 *C. fioriniae* ( $W = 21$ ,  $p = 0.0313$ ), and for 3210 *P. thracensis* against M3 *M. laxa* ( $W = 21$ ,  $p = 0.0355$ ) and VV081 *C. fioriniae* ( $W = 21$ ,  $p = 0.0313$ ). In contrast, no significant differences between autoclaved and non-autoclaved applications were detected for strain 1723B *P. kayaii* against either M3 *M. laxa* ( $W = 10$ ,  $p = 0.586$ ) or VV081 *C. fioriniae* ( $W = 20$ ,  $p = 0.0585$ ) (Figure 5).

Taken together, the results indicate that heat-treated bacterial preparations did not exhibit reduced antifungal activity. While non-heat treated applications showed more

variable and fungus-dependent responses, inhibitory effects were consistently maintained under heat-treated conditions across strains and concentrations. Based on this statistical consistency, heat-treated treatments—particularly at the higher concentration—were selected for subsequent experiments. Therefore, all further assays were performed using centrifuged and filter-sterilized bacterial cultures incorporated into PDA prior to autoclaving, ensuring methodological consistency while retaining reliable antifungal activity.

### 3.4. Fermentation Time Dependency

The temporal effects of incubation time on the biological activity of *P. cinerea* (3086), *P. kayaii* (1723B), and *P. thracensis* (3210) were analyzed against two different fungal strains (*M. laxa* M3 and *C. fiorinia* VV081) (Figure 6). Three bacterial species were tested against two fungal strains at three different fermentation durations. Each bacterium–fungus–fermentation time combination was performed with three technical replicates, and the entire experiment was independently repeated at two different time points, which were considered as biological replicates.



**Figure 6.** Effect of fermentation time of bacterial cultures on the inhibitory effect of different *Photorhabdus* strains against *M. laxa* M3 and *C. fiorinia* VV081 strains, expressed as Inhibition Index. See Table 1 for the designation of bacterial strains. Different letters denote statistically significant differences among groups. Asterisks indicate significant differences (Dunn’s test, Bonferroni-adjusted; \*  $p < 0.05$ , \*\*  $p < 0.01$ ).

Shapiro–Wilk’s test indicated that inhibition index values significantly deviated from normality ( $W = 0.921$ ,  $p = 0.00157$ ). Therefore, all subsequent analyses were conducted using non-parametric methods.

When fermentation time was considered as a single factor across the full dataset, a significant overall effect was detected (Kruskal–Wallis:  $\chi^2(2) = 11.20$ ,  $p = 0.00370$ ). This temporal effect remained significant when the datasets were analyzed separately for *M. laxa* M3 ( $\chi^2(2) = 7.53$ ,  $p = 0.0232$ ) and *C. fiorinia* VV081 ( $\chi^2(2) = 10.30$ ,  $p = 0.00580$ ), indicating that incubation time influenced antifungal activity in both fungal systems.

In addition, bacterial strain had a significant overall effect on inhibition indices ( $\chi^2(2) = 20.89$ ,  $p = 2.91 \times 10^{-5}$ ), whereas fungal strain alone did not ( $\chi^2(1) = 0.704$ ,  $p = 0.401$ ). These results suggested that antifungal activity was primarily shaped by bacterial identity and fermentation time, and prompted further analysis of their combined effects.

To better resolve fermentation time dependency within each fungal system, bacterial strain  $\times$  fermentation time combinations were analyzed separately for each fungus. For *M. laxa* M3, the Kruskal–Wallis test revealed a significant overall effect among combinations

( $\chi^2(8) = 25.39, p = 0.00133$ ). Dunn's post hoc test (Bonferroni-adjusted) showed that only a limited number of comparisons were statistically significant. Specifically, 3086 *P. cinerea* at 24 h differed significantly from 72 h (3086\_24 vs. 3086\_72,  $p_{\text{adj}} = 0.0295$ ), and from 3210 *P. thracensis* at 72 h (3086\_24 vs. 3210\_72,  $p_{\text{adj}} = 0.0113$ ). No other pairwise comparisons remained significant after correction. These results indicate that, under M3 conditions, temporal variation was mainly driven by 3086 *P. cinerea*, whereas 1723B *P. kayaii* and 3210 *P. thracensis* exhibited relatively stable antifungal activity across fermentation times.

Similarly, for *C. fioriniae* VV081, a significant overall effect was detected among bacterial strain  $\times$  fermentation time combinations ( $\chi^2(8) = 25.73, p = 0.00117$ ). Dunn's post hoc analysis identified significant differences only within 3086 *P. cinerea*, where the 24 h treatment differed from both 48 h ( $p_{\text{adj}} = 0.0420$ ) and 72 h ( $p_{\text{adj}} = 0.00744$ ). No significant temporal shifts were observed for 1723B *P. kayaii* or 3210 *P. thracensis* after Bonferroni correction. Thus, fermentation time dependency in the VV081 system was again primarily associated with 3086 *P. cinerea*.

Overall, these findings demonstrate that fermentation time-dependent changes in antifungal activity were strain-specific rather than universal. In both fungal systems, only 3086 *P. cinerea* showed consistent and statistically supported temporal variation, with the most pronounced differences occurring between early (24 h) and later (48–72 h) fermentation stages. From an applied perspective, this supports the selection of 48 h as a practical and reproducible fermentation time for subsequent experiments, capturing early metabolic changes while representing a stabilized activity phase.

### 3.5. Preparation Mode Dependency

Prior to statistical testing, the distribution of inhibition index values was assessed using the Shapiro–Wilk normality test for the complete dataset and for subsets grouped by bacterial isolate, fungal species, and preparation mode. The data significantly deviated from normality in the overall dataset ( $W = 0.891, p = 0.002$ ) and in most subgroups ( $p < 0.05$ ). Therefore, non-parametric statistical approaches were applied.

To examine differences among bacteria–preparation mode combinations, inhibition index values were grouped into six categories (three bacterial isolates  $\times$  two preparation modes). Kruskal–Wallis tests were conducted separately for the *M. laxa* M3 and *C. fioriniae* VV081 datasets. When significant overall effects were detected, Dunn's post hoc multiple comparison test with  $p$ -value adjustment was used to identify pairwise differences.

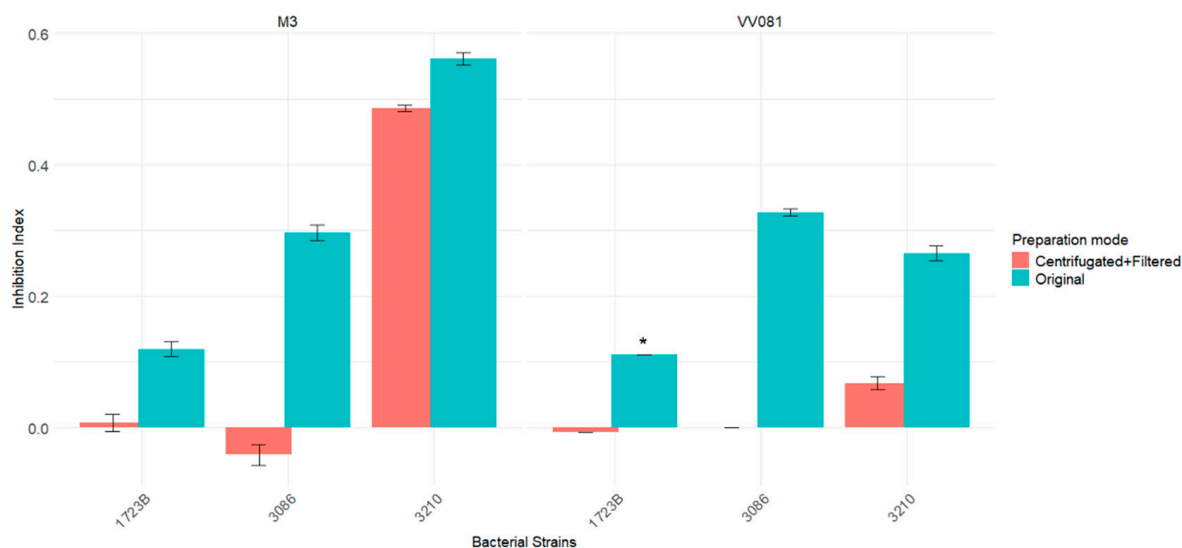
Shapiro–Wilk tests showed that inhibition index data were not normally distributed in the overall dataset ( $p = 0.002$ ) and across most subgroupings, justifying the use of non-parametric tests.

For the *M. laxa* M3 dataset, the Kruskal–Wallis test revealed a significant overall difference among the six bacteria–preparation mode groups ( $\chi^2 = 16.44, df = 5, p = 0.0057$ ). However, Dunn's post hoc analysis with correction for multiple testing detected only one significant pairwise difference, between *P. cinerea* 3086 (Centrifuged + Filtered) and *P. thracensis* 3210 (Original) ( $p_{\text{adj}} = 0.0113$ ). All remaining pairwise comparisons were not statistically significant after adjustment ( $p_{\text{adj}} > 0.05$ ).

Similarly, for the *C. fioriniae* VV081 dataset, the Kruskal–Wallis test also indicated a significant overall group effect ( $\chi^2 = 16.82, df = 5, p = 0.0049$ ). Dunn's post hoc test identified a single significant difference, between *P. kayaii* 1723B (Centrifuged + Filtered) and *P. cinerea* 3086 (Original) ( $p_{\text{adj}} = 0.0079$ ). No other pairwise comparisons reached statistical significance after correction ( $p_{\text{adj}} > 0.05$ ).

Overall, although significant global differences among bacteria–preparation mode combinations were detected for both fungal species, post hoc analyses demonstrated that

statistically supported effects were limited to isolated group comparisons rather than reflecting a consistent effect of preparation mode (Figure 7).



**Figure 7.** Effect of preparation mode of bacterial fermentation broth on the inhibitory effect of different *Photothabdus* strains against *M. laxa* M3 and *C. fiorinia* VV081 strains, expressed as Inhibition Index. See Table 1 for the designation of bacterial strains. Statistical differences between groups were assessed using the Mann-Whitney U test. Significance levels are indicated as: \*  $p < 0.05$ .

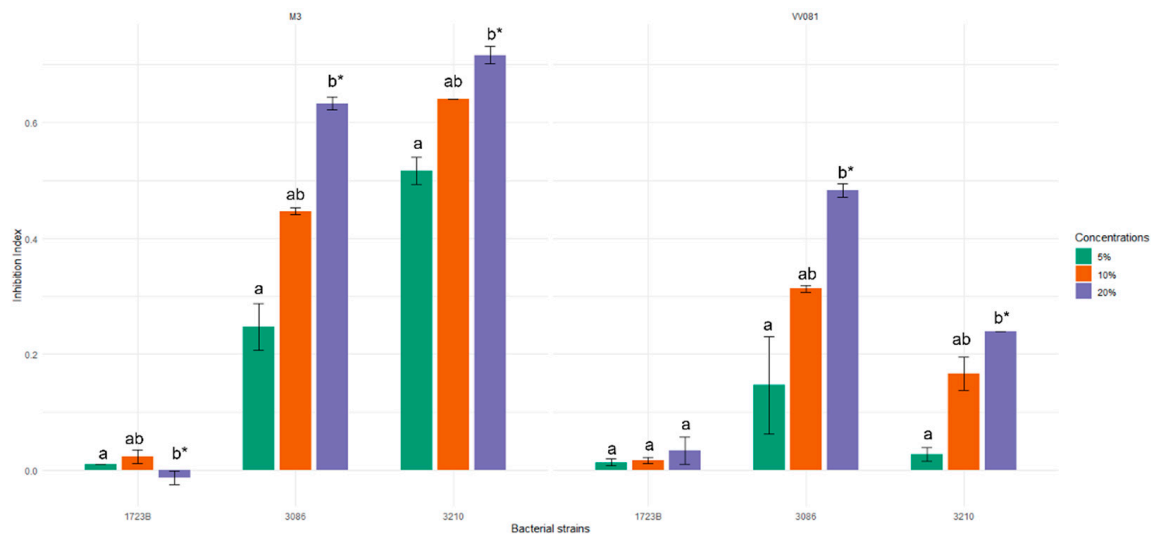
### 3.6. Concentration Dependency

Kruskal–Wallis analysis indicated concentration-dependent differences in antifungal activity for some bacterium–fungus combinations in *M. laxa* M3 and *C. fiorinia* VV081 (Figure 7). Three bacterial species were tested against two fungal strains at three different concentrations. Each bacterium–fungus–concentration combination was performed with three technical replicates, and the experiment was independently repeated at two different time points, which were considered as biological replicates.

For *M. laxa* M3, *P. kayaii* showed  $\chi^2 = 6.889$ ,  $p = 0.03192$ , with Dunn’s post-hoc test indicating that 20% concentration was significantly more effective than 10% ( $Z = 2.593$ ,  $p_{\text{adj}} = 0.014$ ), while 10% vs. 5% and 20% vs. 5% differences were not significant. Similarly, *P. cinerea* 3086 exhibited  $\chi^2 = 7.322$ ,  $p = 0.02571$ , with 20% vs. 5% showing a significant effect ( $p_{\text{adj}} = 0.020$ ), and *P. thracensis* 3210 had  $\chi^2 = 7.513$ ,  $p = 0.02336$ , with 20% vs. 5% also significant ( $p_{\text{adj}} = 0.018$ ). For *M. laxa* M3, higher inhibition values were generally observed at the 20% concentration; however, statistically significant differences were limited to specific pairwise comparisons depending on the bacterial strain.

For *C. fiorinia* VV081, *P. kayaii* 1723B showed no significant differences among concentrations ( $\chi^2 = 3.444$ ,  $p = 0.179$ ), whereas *P. cinerea* 3086 ( $\chi^2 = 7.322$ ,  $p = 0.02571$ ) and *P. thracensis* 3210 ( $\chi^2 = 7.579$ ,  $p = 0.0226$ ) demonstrated significantly higher values at 20% compared to 5% ( $p_{\text{adj}} = 0.020$  and 0.018, respectively). For *C. fiorinia* VV081, a concentration-dependent response was detected for *P. cinerea* 3086 and *P. thracensis* 3210, whereas no significant concentration effect was observed for *P. kayaii* 1723B. These results suggest a tendency toward higher inhibition at increased concentrations, particularly for *P. cinerea* (3086) and *P. thracensis* (3210), while responses remained limited for *P. kayaii* (1723B). The statistical analysis indicated that higher concentrations, particularly 20%, were often associated with increased antifungal activity; however, the magnitude and significance of this effect varied among bacterial strains and fungal targets. While 10% concentration had moderate effects and 5% the lowest, inhibition efficacy differed among fungi. *P. kayaii* showed limited response, whereas *P. cinerea* and *P. thracensis* exhibited strong antifungal

activity at 20%, highlighting that higher concentrations are more effective and that activity depends on both bacterial and fungal strains (Figure 8).



**Figure 8.** Effect of the concentration of bacterial preparation in the fungal culture media on the inhibitory effect of different *Photorhabdus* strains against *M. laxa* M3 and *C. fiorinia* VV081 strains, expressed as Inhibition Index. See Table 1 for the designation of bacterial strains. Different letters denote statistically significant differences among groups, as determined by Dunn's test with Bonferroni-adjusted  $p$ -values \*  $p < 0.05$ .

#### 4. Discussion

To the best of our knowledge, there are currently no studies reporting the antifungal interactions of *Photorhabdus cinerea* and *Photorhabdus thracensis* against *Monilinia* spp. and *Colletotrichum* spp. Under in vitro conditions, the present study provides initial evidence for antifungal activity of these two *Photorhabdus* species.

*Photorhabdus* spp. are considered promising eco-friendly biopesticide candidates due to their ability to inhibit the growth of various phytopathogenic fungi, including *Colletotrichum* spp., *Pythium* spp., *Rhizoctonia* spp., *Fusarium* spp., *Botrytis* spp., *Armillaria* spp. and fungal-like Oomyceta microorganisms, as *Phytophthora* spp. [33,43–49]. Focusing on the pathogens included in this study, *P. akhurstii* has been proposed as a potential biological control agent against *C. gloeosporioides* [41]. In addition, *Photorhabdus luminescens* has been reported to exhibit strong antifungal effects against *C. gloeosporioides*, *C. fragariae*, and *C. acutatum* [50]. Similarly, another study also confirmed the high antifungal activity of *P. luminescens* against *Colletotrichum* spp. [37]. The present study adds two species to the list of bacteria effective against *Colletotrichum* species, as both *P. thracensis* and *P. cinerea* had a significant inhibitory effect on *C. fiorinia*.

Previous studies on the antifungal activity of *Photorhabdus* species against *Monilinia* pathogens have reported generally low to moderate inhibitory effects. For instance, *P. kayaii* was found to be weakly effective against *M. laxa* [34], while *P. temperata* and *P. luminescens* exhibited moderate inhibition against *M. fructicola* [36]. Consistent with these reports, our strain *P. kayaii* 1723B showed weak antifungal activity against *M. laxa*, and *P. temperata* showed limited antifungal activity against either of the tested fungal species. In contrast, our results revealed that *P. cinerea* 3086 exhibited moderate, and *P. thracensis* 3210 pronounced inhibitory effects against *M. laxa* M3, representing novel findings for these species. Previous literature also indicates that *P. luminescens* strains can display strong and strain-specific antifungal activity, completely inhibiting a range of plant pathogenic fungi, including *M. fructicola*, *B. cinerea*, *Ceratocystis* spp., *Mucor piriformis*, and *Pythium* spp. [31,51].

Together, these findings are consistent with earlier reports on variable antifungal activity among *Photorhabdus* strains against different fungal pathogens. Our findings demonstrate that *P. cinerea* 3086 displayed moderate inhibitory activity, whereas *P. thracensis* 3210 exerted pronounced suppression of *M. laxa* M3, constituting previously unreported antifungal activity for these species.

Several studies have highlighted the potential of *Photorhabdus* and *Xenorhabdus* species as biological control agents. For instance, their antifungal activity against *Dothiorella* sp. has been confirmed [52], while strong inhibition of *Moniliophthora roreri*—even surpassing that of a commercial fungicide—has also been reported [53]. The present study shows that entomopathogenic bacteria of these genera, along with their secondary metabolites, effectively inhibit human and animal pathogenic fungi such as *Aspergillus fumigatus* and *Candida albicans*, providing insights into novel antifungal strategies against drug-resistant pathogens [54]. In addition, *Photorhabdus* species have been shown to suppress a range of plant pathogens, including *Magnaporthe grisea*, *Fusarium oxysporum* f. sp. *niveum*, *Fusarium oxysporum* f. sp. *cucumerinum*, *B. cinerea*, and *Phytophthora capsici* [55]. Importantly, the inhibitory effect of a given bacterial species appears to be highly dependent on the fungal target, suggesting that multiple bioactive components may contribute to suppression and that individual fungi exhibit variable sensitivity. This pattern is also evident in our own results. For instance, *P. thracensis* 3210 consistently showed stronger inhibition against *M. laxa* M3 than against *C. fioriniae* VV081, whereas *P. laumondii* 3196 exhibited the opposite trend, displaying higher inhibition values on *C. fioriniae* VV081. From a practical perspective, this specificity implies that *Photorhabdus*- or *Xenorhabdus*-based preparations are unlikely to serve as broad-spectrum antifungal agents, necessitating tailored applications against specific pathogens.

The preparation method of bacterial fermentation broth critically influences its antifungal efficacy. Multiple studies have shown that crude extracts or unfiltered fermentation broth of *Photorhabdus* and *Xenorhabdus* species exhibit significantly stronger inhibition against diverse fungal pathogens, including *Fusarium graminearum*, *B. cinerea*, and *S. sclerotiorum*, compared to filtered or autoclaved preparations [45,56,57]. These observations indicate that the bioactive compounds responsible for antifungal activity are largely associated with bacterial cells and are only partially released into the culture medium. For most species, removal of the cell mass markedly reduces antifungal potency, underscoring the contribution of cell-associated metabolites and volatile compounds. This pattern was also evident in our own results, where the Original (non-centrifuged and non-filtered) preparations consistently exhibited higher inhibition than the Centrifuged + Filtered samples for all tested strains (*P. kayaii* 1723B, *P. cinerea* 3086, and *P. thracensis* 3210) on both fungal targets (*M. laxa* M3 and *C. fioriniae* VV081). Notably, *P. thracensis* represents an exception; although its antifungal activity was diminished upon cell removal, the reduction is less pronounced than in other species, suggesting a higher release or stability of active metabolites in the extracellular milieu. Furthermore, fermentation conditions—including medium composition, aeration, and solid versus liquid culture—substantially affect metabolite yield and activity, highlighting the necessity of optimizing preparation protocols to maximize biocontrol potential [58,59]. Consistent with these observations, our data demonstrate that although the Original (non-centrifuged and non-filtered) preparation exhibited the highest inhibitory activity across all bacterial species, *P. thracensis* 3210 showed the least reduction in antifungal efficacy following centrifugation and filtration. This suggests that, relative to *P. kayaii* 1723B and *P. cinerea* 3086, a greater proportion of the bioactive metabolites of *P. thracensis* 3210 is either more readily released into, or more stable within, the cell-free fraction. Collectively, these findings emphasize that both the presence of bacterial biomass and the culture conditions are key determinants of the efficacy of *Photorhabdus*-

and *Xenorhabdus*-based antifungal preparations, which has important implications for their practical application as targeted biocontrol agents.

The present study represents an initial *in vitro* screening conducted on a limited number of fungal isolates. While the observed antifungal activity highlights the potential of selected *Photorhabdus* strains and their metabolites, the findings should be interpreted within this experimental context. At this stage, conclusions are limited to contact-independent inhibition under controlled *in vitro* conditions, and no inference can be made regarding efficacy in planta or under field conditions. Translation of these findings into practical biocontrol applications will require additional validation steps, including assays on detached fruits or whole plants, evaluation of potential phytotoxic effects, assessment of metabolite stability under storage and environmental conditions, and examination of possible non-target impacts. Although these aspects were beyond the scope of the present study, the current results provide a necessary foundation for such downstream investigations.

To address potential physicochemical effects, we measured the pH of PDA amended with bacterial fermentation broths at all tested concentrations, both before and after autoclaving. Native PDA showed pH values of 5.61 (before) and 5.78 (after autoclaving), whereas amended media exhibited only moderate increases (approximately 0.2–1.1 units), with final pH values not exceeding ~6.7.

The present results demonstrate that heat treatment has no negative effect on the antifungal efficacy of the bacterial preparations applied in this study; in fact, in some cases, heat-treated preparations produced stronger and more consistent inhibitory effects than non-heated ones. While non-heated treatments showed limited and fungus-dependent dose responses, heat treated preparations resulted in significant differences across most bacterial–fungal combinations, indicating a more robust antifungal activity. This suggests that heat-stable metabolites or denatured bacterial components generated at high temperature during autoclaving may play a major role in fungal growth inhibition. The significant differences observed between heated and non-heated preparations for strains 3086 *P. cinerea* and 3210 *P. thracensis* further support the conclusion that the mode of application is a critical determinant of antifungal efficacy. Importantly, these values fall well within the reported growth ranges of both *M. laxa* (approximately pH 2.4–8.8) and *C. fioriniae* (optimal around pH 4–6, with broader tolerance) [60–62]. Therefore, based on both our measurements and published data, the strong inhibitory effects observed cannot be explained solely by pH shifts or simple physicochemical changes of the medium, but are most plausibly associated with biologically active components present in the bacterial preparations. Accordingly, our results should be interpreted as inhibitory effects of heat-treated bacterial preparations rather than unequivocal biological effects. Future work should include uninoculated broth controls, pH-matched PDA controls, and non-autoclaved filter-sterilized preparations to disentangle physicochemical effects from biological activity.

## 5. Conclusions

This study systematically evaluated the antifungal potential of selected *Photorhabdus* strains against *M. laxa* (M3) and *C. fioriniae* (VV081) through a series of pre-selection, drop-to-drop, fermentation time, preparation mode, and concentration-dependent assays. Among the 13 initially screened strains, five representatives of different species (*P. kayaii* 1723B, *P. temperata* 3017, *P. cinerea* 3086, *P. laumondii* 3196, and *P. thracensis* 3210) exhibited the most promising antifungal activity. Notably, *P. thracensis* 3210 consistently demonstrated the strongest inhibitory effects across both fungal isolates, indicating high biocontrol potential.

Temporal analyses revealed that bacterial antifungal activity is dependent on fermentation time, with most strains achieving optimal or stabilized activity after 48 h, although some strains displayed gradual or delayed responses depending on the fungal host. Fur-

thermore, preparation mode significantly influenced antifungal efficacy; Original bacterial cultures containing heat-killed bacterial cells generally outperformed cell-free (centrifuged and filtered) samples, emphasizing the importance of maintaining native metabolic components during application. Concentration-dependent assays highlighted clear concentration-dependent antifungal effects in the studied range of 5–20% in the culture media, with variability observed among both bacterial and fungal strains.

In conclusion, this study demonstrates that heat treatment has no negative effect on the antifungal activity of the bacterial preparations applied in this study. Based on the statistical outcomes, the heat treated, living-cell-free bacterial preparations were therefore preferred for subsequent experiments, and centrifuged and filter-sterilized bacterial cultures were incorporated into PDA prior to autoclaving. These findings highlight the importance of treatment mode in shaping antifungal efficacy and provide a methodological framework for future studies investigating bacterial-derived antifungal factors.

Overall, these findings demonstrate that the antifungal potential of *Photorhabdus* strains is strain-specific, concentration-dependent, and influenced by both incubation time and sample preparation under controlled in vitro conditions. The present study provides a foundation frame work for the future developments, indicating that targeted strain collection, fermentation duration, and application concentrations are important parameters to be systematically explored in subsequent in planta and applied studies aimed at evaluating the biocontrol potential of *Photorhabdus*-derived metabolites against horticultural pathogens.

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## Abbreviations

The following abbreviations are used in this manuscript:

BRD	Brown rot disease
DMI	Demethylation-inhibiting
EPB	Entomopathogenic Bacteria
EPN	Entomopathogenic Nematodes



## References

1. Byrde, R.J.W.; Willetts, H.J. *The Brown Rot Fungi of Fruit. Their Biology and Control*; Elsevier: Amsterdam, The Netherlands, 1977.
2. Abate, D.; Pastore, C.; Gerin, D.; De Miccolis Angelini, R.M.; Rotolo, C.; Pollastro, S.; Faretra, F. Characterization of *Monilinia* spp. populations on stone fruit in South Italy. *Plant Dis.* **2018**, *102*, 1708–1717. [[CrossRef](#)]
3. EFSA Panel on Plant Health (PLH). Pest risk assessment of *Monilinia fructicola* for the EU territory and identification and evaluation of risk management options. *EFSA J.* **2011**, *9*, 2119. [[CrossRef](#)]
4. Rungjindamai, N.; Jeffries, P.; Xu, X.M. Epidemiology and management of brown rot on stone fruit caused by *Monilinia laxa*. *Eur. J. Plant Pathol.* **2014**, *140*, 1–17. [[CrossRef](#)]
5. Holb, I.J. The brown rot fungi of fruit crops (*Monilinia* spp.). II. Important features of their epidemiology. *Int. J. Hortic. Sci.* **2004**, *10*, 17–35. [[CrossRef](#)]
6. Da Lio, D.; Cobo-Díaz, J.F.; Masson, C.; Chalopin, M.; Kebe, D.; Giraud, M.; Verhaeghe, A.; Nodet, P.; Sarrocco, S.; Le Floch, G.; et al. Combined metabarcoding and multi-locus approach for genetic characterization of *Colletotrichum* species associated with common walnut (*Juglans regia*) anthracnose in France. *Sci. Rep.* **2018**, *8*, 10765. [[CrossRef](#)]
7. Tan, Q.; Schnabel, G.; Chaisiri, C.; Yin, L.F.; Yin, W.X.; Luo, C.X. *Colletotrichum* Species Associated with Peaches in China. *J. Fungi* **2022**, *8*, 313. [[CrossRef](#)]
8. Dowling, M.E.; Peres, N.A.; Villani, S.; Schnabel, G. Managing *Colletotrichum* on fruit crops: A “complex” challenge. *Plant Dis.* **2020**, *104*, 2301–2316. [[CrossRef](#)] [[PubMed](#)]
9. Dowling, M.; Schnabel, G. Understanding plant diseases using art and technology. *Int. J. Fruit Sci.* **2020**, *20*, 959–966. [[CrossRef](#)]
10. Holb, I.J.; Schnabel, G. A detached fruit study on the post-inoculation activity of lime sulfur against brown rot of peach (*Monilinia fructicola*). *Australas. Plant Pathol.* **2008**, *37*, 454–459. [[CrossRef](#)]
11. Malandrakis, A.A.; Kavroulakis, N.; Chrysikopoulos, C.V. Copper nanoparticles against benzimidazole-resistant *Monilinia fructicola* field isolates. *Pestic. Biochem. Physiol.* **2021**, *173*, 104796. [[CrossRef](#)]
12. Calabro, J.M.; Spotts, R.A.; Grove, G.G. Biology of Sweet Cherry Powdery Mildew. Ph.D. Thesis, Oregon State University, Corvallis, OR, USA, 2007.
13. Egüen, B.; Melgarejo, P.; de Cal, A. Sensitivity of *Monilinia fructicola* from Spanish peach orchards to thiophanate-methyl, iprodione, and cyproconazole: Fitness analysis and competitiveness. *Eur. J. Plant Pathol.* **2015**, *141*, 789–801. [[CrossRef](#)]
14. Cech, R.; Zaller, J.G.; Lyssimachou, A.; Clausen, P.; Hertoge, K.; Linhart, C. Pesticide drift mitigation measures appear to reduce contamination of non-agricultural areas, but hazards to humans and the environment remain. *Sci. Total Environ.* **2023**, *854*, 158814. [[CrossRef](#)] [[PubMed](#)]
15. Tran, T.T.; Li, H.; Nguyen, D.Q.; Jones, M.G.K.; Sivasithamparan, K.; Wylie, S.J. *Monilinia fructicola* and *Monilinia laxa* isolates from stone fruit orchards sprayed with fungicides displayed a broader range of responses to fungicides than those from unsprayed orchards. *Eur. J. Plant Pathol.* **2018**, *1*, 985–999. [[CrossRef](#)]
16. Avan, M.; Kotan, R.; Albastawisi, E.M.; Erarslan, G. Biological Control of Blossom Blight and Brown Rot Caused by *Monilinia laxa* by Using a *Bacillus subtilis* Strain TV-6F. *Erwerbs-Obstbau* **2023**, *65*, 2399–2405. [[CrossRef](#)]
17. Tsalgatidou, P.C.; Papageorgiou, A.; Boutsika, A.; Chatzidimopoulos, M.; Delis, C.; Tsitsigiannis, D.I.; Paplomatas, E.; Zambounis, A. Insights into the Interaction between the Biocontrol Agent *Bacillus amyloliquefaciens* QST 713, the Pathogen *Monilinia fructicola* and Peach Fruit. *Agronomy* **2024**, *14*, 771. [[CrossRef](#)]
18. Rungjindamai, N.; Xu, X.M.; Jeffries, P. Identification and characterisation of new microbial antagonists for biocontrol of *Monilinia laxa*, the causal agent of brown rot on stone fruit. *Agronomy* **2013**, *3*, 685–703. [[CrossRef](#)]
19. Lv, H.; Zhang, S.; Ma, N.; Boamah, S.; Xu, B. *Trichoderma longibrachiatum* (T6) Peptaibols Inhibiting the *Monilia yunnanensis* Growth and Inducing Pear Fruit Resistance in Its Infection. *Antioxidants* **2024**, *13*, 1517. [[CrossRef](#)]
20. Rungjindamai, N.; Jeffries, P.; Xu, X. Efficacy of biopesticides and fungicides against brown rot on cherry (*Prunus avium*) and plum (*Prunus domestica*). *Biocontrol Sci. Technol.* **2025**, *35*, 479–499. [[CrossRef](#)]
21. Chen, S.N.; Luo, C.X.; Hu, M.J.; Schnabel, G. Sensitivity of *Colletotrichum* species, including *C. fioriniae* and *C. nymphaeae*, from peach to demethylation inhibitor fungicides. *Plant Dis.* **2016**, *100*, 2434–2441. [[CrossRef](#)]
22. Munir, M.; Amsden, B.; Dixon, E.; Vaillancourt, L.; Gauthier, N.A.W. Characterization of *Colletotrichum* species causing bitter rot of apple in Kentucky orchards. *Plant Dis.* **2016**, *100*, 2194–2203. [[CrossRef](#)]
23. Ling, L.; Zhao, Y.; Tu, Y.; Yang, C.; Ma, W.; Feng, S.; Lu, L.; Zhang, J. The inhibitory effect of volatile organic compounds produced by *Bacillus subtilis* CL2 on pathogenic fungi of wolfberry. *J. Basic Microbiol.* **2021**, *61*, 110–121. [[CrossRef](#)]
24. Boemare, N.; Akhurst, R. The genera *Photorhabdus* and *Xenorhabdus*. In *The Prokaryotes*; Dworkin, M., Falkow, S., Rosenberg, E., Schleifer, K.-H., Stackebrandt, E., Eds.; Springer: New York, NY, USA, 2006; pp. 451–494.
25. Challinor, V.L.; Bode, H.B. Bioactive natural products from novel microbial sources. *Ann. N. Y. Acad. Sci.* **2015**, *1354*, 82–97. [[CrossRef](#)]
26. Bode, H.B. Insects: True Pioneers in Anti-Infective Therapy and What We Can Learn from Them. *Angew. Chem. Int. Ed.* **2009**, *48*, 6394–6396. [[CrossRef](#)]

27. Akhurst, R.J. Antibiotic activity of *Xenorhabdus* spp., bacteria symbiotically associated with insect pathogenic nematodes of the families Heterorhabditidae and Steinernematidae. *J. Gen. Microbiol.* **1982**, *128*, 3061–3065. [[CrossRef](#)]
28. Fodor, A.; Gualtieri, M.; Zeller, M.; Tarasco, E.; Klein, M.G.; Fodor, A.M.; Haynes, L.; Lengyel, K.; Forst, S.A.; Furgani, G.M.; et al. Type strains of entomopathogenic nematode-symbiotic bacterium species, *Xenorhabdus szentirmaii* (EMC) and *X. budapestensis* (EMA), are exceptional sources of non-ribosomal templated, large-target-spectral, thermotolerant-antimicrobial peptides (by both), and iodinin (by EMC). *Pathogens* **2022**, *11*, 342. [[CrossRef](#)]
29. Forst, S.; Neelson, K. Molecular biology of the symbiotic-pathogenic bacteria *Xenorhabdus* spp. & *Photorhabdus* spp. *Microbiol. Rev.* **1996**, *60*, 21–43. [[PubMed](#)]
30. Chacón-Orozco, J.G.; Bueno, C., Jr.; Shapiro Ilan, D.I.; Hazir, S.; Leite, L.G.; Harakava, R. Antifungal activity of *Xenorhabdus* spp. and *Photorhabdus* spp. against the soybean pathogenic *Sclerotinia sclerotiorum*. *Sci. Rep.* **2020**, *10*, 20649. [[CrossRef](#)]
31. Chen, G.; Dunphy, G.; Webster, J. Antifungal Activity of Two *Xenorhabdus* Species and *Photorhabdus luminescens*, Bacteria Associated with the Nematodes *Steinernema* Species and *Heterorhabditis megidis*. *Biol. Control* **1994**, *4*, 157–162. [[CrossRef](#)]
32. Cimen, H.; Touray, M.; Gulsen, S.H.; Erincik, O.; Wenski, S.L.; Bode, H.B.; Shapiro-Ilan, D.; Hazir, S. Antifungal activity of different *Xenorhabdus* and *Photorhabdus* species against various fungal phytopathogens and identification of the antifungal compounds from *X. szentirmaii*. *Appl. Microbiol. Biotechnol.* **2021**, *105*, 5517–5528. [[CrossRef](#)]
33. Shi, D.; An, R.; Zhang, W.; Zhang, G.; Yu, Z. Stilbene Derivatives from *Photorhabdus temperata* SN259 and Their Antifungal Activities against Phytopathogenic Fungi. *J. Agric. Food Chem.* **2017**, *65*, 60–65. [[CrossRef](#)]
34. Uluğ, D. Bacterial allies in agricultural defense: Evaluating *Xenorhabdus* and *Photorhabdus* supernatants against *Phytophthora infestans* and *Monilinia laxa*. *Düzce Univ. Sci. Technol. J.* **2024**, *12*, 2131–2138. [[CrossRef](#)]
35. Shapiro-Ilan, D.I.; Reilly, C.C.; Hotchkiss, M.W. Suppressive effects of metabolites from *Photorhabdus* and *Xenorhabdus* spp. on phytopathogens of peach and pecan. *Arch. Phytopathol. Plant Protect.* **2009**, *42*, 715–728. [[CrossRef](#)]
36. Hazir, S.; Shapiro-Ilan, D.I.; Bock, C.H.; Hazir, C.; Leite, L.G.; Hotchkiss, M.W. Relative potency of culture supernatants of *Xenorhabdus* and *Photorhabdus* spp. on growth of some fungal phytopathogens. *Eur. J. Plant Pathol.* **2016**, *146*, 369–381. [[CrossRef](#)]
37. Alforja, S.I.R.; Rico, P.M.B.; Caoili, B.L.; Latina, R.A. Two Philippine *Photorhabdus luminescens* strains inhibit the in vitro growth of *Lasioidiplodia theobromae*, *Fusarium oxysporum* f. sp. *lycopersici*, and *Colletotrichum* spp. *Egypt. J. Biol. Pest Control* **2021**, *31*, 108. [[CrossRef](#)]
38. Park, S.J.; Jun, M.H.; Chun, W.S.; Seo, J.; Yi, Y.K.; Kim, Y.G. Control effects of benzylideneacetone isolated from *Xenorhabdus nematophila* K1 on the diseases of red pepper plants. *Res. Plant Dis.* **2010**, *16*, 170–175. [[CrossRef](#)]
39. Zhang, J.; Tian, C.; Su, L.; Sun, H.; Ding, Y.; Li, M.; Li, L. Metabolites of symbiotic bacteria from *Steinernema feltiae* SRP 18-91 and their antibacterial activity. *Chin. J. Biol. Control* **2024**, *40*, 1128–1134. [[CrossRef](#)]
40. Bussaman, P.; Sa-uth, C.; Rattanasena, P. Comparison between antifungal activities of *Xenorhabdus stockiae* PB09 cell-free supernatants derived from shake-flask cultivation and fermentation. *J. Sci. Technol. Mahasarakham Univ.* **2020**, *39*, 200–205.
41. Tu, P.-W.; Chiu, J.-S.; Lin, C.; Chien, C.-C.; Hsieh, F.-C.; Shih, M.-C.; Yang, Y.-L. Evaluation of the antifungal activities of *Photorhabdus akhurstii* and its secondary metabolites against phytopathogenic *Colletotrichum gloeosporioides*. *J. Fungi* **2022**, *8*, 403. [[CrossRef](#)]
42. Tóth, T.; Lakatos, T.; Kaskötő, Z. Identification and typing of *Photorhabdus* isolates from entomopathogenic nematodes in Hungarian soils. *IOBC/wprs Bull.* **2008**, *31*, 138–143.
43. Vicente-Díez, I.; Dueñas-Hernani, J.; Campos-Herrera, R. Antifungal activity of *Xenorhabdus* and *Photorhabdus* against aerial and soilborne grapevine pathogens: Varying efficiencies and non-target effects. *Biol. Control* **2025**, *205*, 105759. [[CrossRef](#)]
44. Dominelli, N.; Platz, F.; Heermann, R. The insect pathogen *Photorhabdus luminescens* protects plants from phytopathogenic *Fusarium graminearum* via chitin degradation. *Appl. Environ. Microbiol.* **2022**, *88*, e00645-22. [[CrossRef](#)]
45. Kgosiemang, J.L.; Ramakuwela, T.; Figlan, S.; Cochrane, N. Antifungal effect of metabolites from bacterial symbionts of entomopathogenic nematodes on *Fusarium* head blight of wheat. *J. Fungi* **2024**, *10*, 148. [[CrossRef](#)]
46. Vicente-Díez, I.; Carpennero, E.; Pou, A.; Campos-Herrera, R. Exploring bacterial cell-free supernatants, unfiltered ferments and crude bacteria uses of *Xenorhabdus* and *Photorhabdus* (Morganellaceae) for controlling *Botrytis cinerea* (Helotiales: Sclerotiniaceae). *Biol. Control* **2023**, *183*, 105259. [[CrossRef](#)]
47. Lalramchuan, M.; Lalramnghaki, H.C.; Vanlalsangi, R.; Lalhmingliani, E. Characterization and screening of antifungal activity of bacteria associated with entomopathogenic nematodes from Mizoram, North-Eastern India. *J. Environ. Biol.* **2020**, *41*, 942–950. [[CrossRef](#)]
48. Palmieri, D.; Portillo, E.; Sulbarán, Y.; Guerra, M.; San-Blas, E. Biocontrol of *Phytophthora* root and stem rot disease in papaya (*Carica papaya*) plants by *Photorhabdus*, the symbiont bacterium of *Heterorhabditis amazonensis*. *BioControl* **2019**, *64*, 595–604. [[CrossRef](#)]
49. Shapiro-Ilan, D.I.; Bock, C.H.; Hotchkiss, M.W. Suppression of pecan and peach pathogens on different substrates using *Xenorhabdus bovienii* and *Photorhabdus luminescens*. *Biol. Control* **2014**, *77*, 1–6. [[CrossRef](#)]

50. Bock, C.H.; Shapiro-Ilan, D.I.; Wedge, D.E.; Cantrell, C.L. Identification of the antifungal compound, trans-cinnamic acid, produced by *Photorhabdus luminescens*, a potential biopesticide against pecan scab. *J. Pest Sci.* **2014**, *87*, 155–162. [[CrossRef](#)]
51. Maher, A.M.D.; Asaiyah, M.; Quinn, S.; Burke, R.; Wolff, H.; Bode, H.B.; Griffin, C.T. Competition and co-existence of two *Photorhabdus* symbionts with a nematode host. *Microb. Ecol.* **2021**, *81*, 223–239. [[CrossRef](#)]
52. San-Blas, E.; Parra, Y.; Carrillo, Z. Effect of *Xenorhabdus* and *Photorhabdus* bacteria (Enterobacteriales: Enterobacteriaceae) and their exudates on the apical rotten fruit disease caused by *Dothiorella* sp. in guava (*Psidium guajava*). *Arch. Phytopathol. Plant Prot.* **2013**, *46*, 2294–2303. [[CrossRef](#)]
53. San-Blas, E.; Carrillo, Z.; Parra, Y. Effect of *Xenorhabdus* and *Photorhabdus* bacteria and their exudates on *Moniliophthora roreri*. *Arch. Phytopathol. Plant Prot.* **2012**, *45*, 1950–1967. [[CrossRef](#)]
54. Lalramchuan, M.; Ramliana, L.; Lalramnghaki, H.C.; Chawngthu, A.L.; Ramliana, V.; Lalhmingliani, E. Antifungal potential of entomopathogenic bacteria, *Photorhabdus*, and *Xenorhabdus* (Morganellaceae) against pathogenic fungi. *J. Appl. Biol. Biotech.* **2025**, *13*, 76–88. [[CrossRef](#)]
55. Zhang, P.-J.; Dou, Z.-G.; Wang, H.; Bao, H.-R.; Zhang, K.-Y. Isolation, identification and screening of antifungal spectrum of nine entomopathogenic nematode symbiotic bacterial strains. *J. Nanjing Agric. Univ.* **2021**, *44*, 487–496.
56. Zhang, S.; Han, Y.; Wang, L.; Han, J.; Yan, Z.; Wang, Y.; Wang, Y. Antifungal activity and mechanism of xenocoumacin 1, a natural product from *Xenorhabdus nematophila* against *Sclerotinia sclerotiorum*. *J. Fungi* **2024**, *10*, 175. [[CrossRef](#)] [[PubMed](#)]
57. Yang, X.; Qiu, D.; Yang, H.; Wang, J.; Li, Z. Antifungal activity of xenocoumacin 1 from *Xenorhabdus nematophilus* var. *pekingensis* against *Phytophthora infestans*. *World J. Microbiol. Biotechnol.* **2011**, *27*, 523–528. [[CrossRef](#)]
58. Nishanth, S.K.; Nambisan, B.; Mohandas, C. Antifungal activity of crude extract produced by *Bacillus* sp. associated with entomopathogenic nematode from media formulated by six nitrogen sources with fructose against *Penicillium expansum*. *Arch. Phytopathol. Plant Prot.* **2013**, *46*, 1222–1229. [[CrossRef](#)]
59. Booyesen, E.; Rautenbach, M.; Stander, M.A.; Dicks, L.M.T. Profiling the production of antimicrobial secondary metabolites by *Xenorhabdus khoisanae* J194 under different culturing conditions. *Front. Chem.* **2021**, *9*, 626653. [[CrossRef](#)] [[PubMed](#)]
60. Obi, V.I.; Barriuso, J.J.; Gogorcena, Y. Effects of pH and titratable acidity on the growth and development of *Monilinia laxa* (Aderh. & Ruhl.) in vitro and in vivo. *Eur. J. Plant Pathol.* **2018**, *151*, 781–790. [[CrossRef](#)]
61. Zhou, C.; Chen, J.; Liu, Y.; Luo, N.; Guo, W.; Shi, M.; Li, H. First Report of the Anthracnose Pathogenic Agent on Walnut Fruits in China and Exploration of Its Biological Characteristics. *Horticultrae* **2025**, *11*, 339. [[CrossRef](#)]
62. Li, R.; Cui, L.; Jiang, F.; Ziyang, L.; Chachar, S.; Fan, Z.; Jidi, X.; Guan, C. Identification and biological characteristics of *Colletotrichum* species causing persimmon anthracnose in China and screening of *Colletotrichum horii* antifungal agents. *Plant Dis.* **2025**. [[CrossRef](#)]

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