Antimicrobial effects of symbiotic bacteria from entomopathogenic nematodes for use in biorational control of *Pythium myriotylum*

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Abstract

*Pythium myriotylum* is a destructive soil-borne pathogen, which causes severe yield losses in many crops. The pathogen is a major economic issue for the production of ginger. Due to environmental and regulatory concerns, it is necessary to find biological alternatives to chemical pesticide for *P. myriotylum* control. In the present study, *trans*-cinnamic acid (TCA) and the fermentation broth of symbiotic bacteria from eight species of entomopathogenic nematodes alone and the combination of TCA with the fermentation broth of each symbiotic bacterium, were tested for their effects on mycelial growth and zoospore germination of *P. myriotylum*. Results showed that TCA significantly inhibited mycelial growth. Fermentation broths from seven of the eight strains of symbiotic bacteria directly inhibited mycelial growth, especially symbiotic bacteria isolated from *Steinernema feltiae* (SN strain) and *S. riobrave* (7-12 strain). Moreover, adding TCA significantly increased the extent of the inhibitory effect of mycelial growth by fermentation broths of seven tested bacteria except that associated with *S. rarum*. All bacteria fermentation broths showed inhibitory effects on zoospore germination. However, TCA alone did not show an apparent inhibitory effect on zoospore germination as it did on mycelial growth. Antimicrobial effects on mycelial growth and zoospore germination were proportional to the concentration of symbiotic bacteria isolated from *S. feltiae* (SN strain). This research provides new options for biorational control of *P. myriotylum* using TCA and symbiotic bacteria of entomopathogenic nematodes and will facilitate the sustainable production of ginger and other crops affected by this oomycete pathogens disease.

**Keywords:** ginger, bacterial broth, entomopathogenic nematode, biological control, *trans*-cinnamic acid
Introduction

Ginger (*Zingiber officinale* Rosc, Zingiberaceae) is a perennial monocotyledonous herb, which can be used as both a vegetable and a medicinal plant (Afzal *et al*., 2001). It is also a cash crop for growers in many countries including China, India, Indonesia, Fiji Australia etc (Kavitha and Thomas, 2008). In 2011 the total ginger production in China, the second largest ginger producing country in the world, was 426,032 tons from 36,007 hectare (Li *et al*., 2014). However, ginger is cultivated, which can lead to the outbreak of soil-borne diseases such as ginger blast, damping-off and anthracnose. These diseases result in severe damage and yield loss.

Soft rot (rhizome rot) is one of the most important diseases of ginger. The disease is generally caused by the pathogens *Pythium* sp. Soft rot is reported to be most destructive disease of ginger globally (Rai *et al*., 2018). *Pythium* sp. can infect roots, collar, and succulent parts of the rhizome (Stirling *et al*., 2009), leading to 5 to 30 % yield and even losses, up to 100 % in the case of water logging and high temperatures (Stirling *et al*., 2009; Wang *et al*., 2003). Once ginger fields have been infested with *Pythium* spp., the pathogens could persistent in the soil and lead to ginger soft rot in subsequent replanting (Li *et al*., 2014). To control ginger soft rot and other soil-borne phytopathogenic diseases, farmers repeatedly apply synthetic, plant protection products to the soil, such as Mefenoxam. Mefenoxam, which causes environmental pollution, and also negatively affects soil quality, including a decrease in soil organic matter content and repression of microbial activity (Liu *et al*., 2017). One particular species of *Pythium* species that is extremely harmful to ginger as well as various other crops is *Pythium myriotylum* (Liu *et al*., 2017). Conceivably, a biological control approach would be a viable alternative to the use of chemical control for this oomycete pathogens disease. In this study, we explored the potential of developing a biorational tactic for *P.*
Biocontrol organisms (or their natural by-products) have shown promise in controlling various plant diseases. For example, seven bacterial isolates, belonging to *Bacillus* and *Paenibacillus*, from New Zealand soils cause zoospore germination and germ-tube growth inhibition in *Aphanomyces euteiches* (Wakelin et al., 2002). A bacterial isolate, obtained from mature strawberry fruit and determined to be *Paenibacillus polymyxa* exhibited antagonistic potential against *Botrytis cinerea*, the causal agent of grey mold in strawberries (Helbig, 2001). *Fluorescent pseudomonadse* showed antagonistic activity against *Phytophthora capsici*, a major pathogen of black pepper (Tran and Kruijt, 2008).

Entomopathogenic nematodes (EPNs) are obligate and lethal parasites of insects in the natural environment (Griffin et al. 2005). Some EPN species with high virulence have being produced commercially as biological control agents against insect pests (Acharya et al., 2019). The bacterium *Xenorhabdus* spp. is a species-specific mutualist of nematodes in the genera *Steinernema* (Stilwell et al., 2018), and *Photorhabdus* spp. is a species-specific mutualist of that in genera of *Heterorhabditis*. The *Photorhabdus* and *Xenorhabdus* bacteria are located in the intestine of the infective juvenile nematode (IJs), which is the only free-living stage that survives outside the insect host. IJs penetrate insect hosts through natural openings or through the cuticle and release their symbiotic bacteria into the hemocoel causing septicemia, which eventually leads to host death in 48 hr. The nematodes feed upon the bacterial cells and liquefying host tissues, mature, mate and produce progeny which emerge from the insect cadaver as IJs in search of new target hosts.

Entomopathogenic nematodes have long been identified as an important biocontrol agent against
various above- and below-ground pests and have been fairly well characterized (Ferreira and Malan, 2014; Shapiro-Ilan et al., 2018) whereas their symbiotic bacteria as biological control agents have received less attention. Previous research has shown that the EPN’s symbiotic bacteria could be used in biological control. The bacteria’s secondary metabolites containing toxins, lipases, proteases, antibiotics and lipopolysaccharides, are complex, and can inhibit the growth of mites (Eroglu et al., 2019) and of various microorganisms, including fungi, Gram-positive Micrococcus, Staphylococcus and Bacillus, as well as Gram-negative bacteria (Bi et al., 2018; Caldas et al., 2002; Furgani et al., 2008; Richards et al., 2008; Shi et al., 2017). Some of the secondary metabolites of Photorhabdus luminescens subsp. luminescens and P. temperata have been examined for antibiotic activity. Anthraquinone pigments and trans-stilbenes were determined as antibacterial, whereas trans-stilbenes and trans-cinnamic acid (TCA) were defined as antimicrobial compounds (Bock et al., 2014).

Exploring the EPN’s bacteria (and their metabolites) for biorational control of plant pathogens has received increasing attention recently, and new species and strains of EPNs/bacteria with varying activity have been increasingly reported. Indeed, a wide variety of fungal and oocyte genera have been suppressed using EPN bacteria or their byproducts (Shapiro-Ilan et al., 2014; Hazir et al., 2016). Although many studies have documented the antagonistic properties of bacteria associated with EPNs against fungi, the potential of these bacteria to control P. myriotylum remains unknown. In this paper, we evaluated the antimicrobial activity of EPN symbiotic bacteria against P. myriotylum. Also, the addition of trans-cinnamic acid (TCA) to EPN symbiotic bacteria fermentation broth was assessed considering its antimicrobial characters as one of main metabolites of EPN symbiotic bacteria (Bock et al., 2014). The aim of our study is to exploit antimicrobial
properties of bacteria associated with EPNs, and provide insight into utilizing bacterial metabolites
to control *P. myriotylum* in an integrated pest management system.

**Materials and Methods**

**Symbiotic Bacterial strains**

Eight entomopathogenic nematode species were provided by Southeastern Fruit and Tree Nut
Research Lab in USDA-ARS (Byron, GA) including *Steinernema carpocapsae* (Cxd strain),
*Heterorhabditis bacteriophora* (VS strain), *S. carpocapsae* (All strain), *H. bacteriophora* (HB1
strain), *S. feltiae* (SN strain), *S. rarum* (17 C&E strain), *H. indica* (HOM1 strain), and *S. riobrave*
(7-12 strain). Bacteria that are species-specific and associated with the corresponding nematode
isolates were used here. Details of the strains and their symbiotic bacteria were shown in Table 1.

**Preparation of bacterial fermentation broth**

Symbiotic bacteria from the entomopathogenic nematodes were obtained after inoculating last-
instar *Galleria mellonella* with 15μL of nematode suspension (approximately 100 infective
juveniles per larvae) in a 24-well cell culture plate covered with filter paper. Infected larvae were
stored at 25 °C for 36-48 h, then, all dead larvae were surface-sterilized with 75% alcohol for 1
min. A drop of haemolymph was next added onto NBTA medium (45 g nutrient agar, 25 mg
bromothymol blue and 40 mg triphenyl tetrazolium in 1 L distilled water) in 9 cm Petri dishes and
incubated at 25 °C (Kaya and Stock, 1997). Symbiotic bacteria have two variants. Primary variant
symbiotic bacteria absorbs bromothymol blue in NBTA medium so the colony turns blue. Secondary
variant symbiotic bacteria do not absorb bromothymol blue so the colony is red. We used only
primary variant symbiotic bacteria to do the experiment. After 48 h pure colonies of the bacteria
were inoculated on trypticase soy yeast (TSY) medium (4% tryptic soy broth, 0.5% yeast extract) at 22 °C, 150 rpm for 36h in an incubator shaker (HNY-2102C, Tianjin Honour Instrument Co., Ltd) (Ansari et al., 2003). Then fermentation broth was filtrated through a 0.22 µm filter and finally stored at 4°C until experimentation.

**Antimicrobial effects of cell free fermentation broths of different symbiotic bacteria and TCA**

*Pythium myriotylum* was isolated from diseased ginger grown in Shandong province, China. The isolated fungal colonies were stored at 4 °C on PDA medium (potato dextrose agar) and routinely sub-cultured on PDA in Petri dishes (90 mm in diameter) in the dark at 27°C until the colonies were big enough to use. As indicated in previous research, trans-cinnamic acid (TCA) is one of the major metabolic components of EPN symbiotic bacteria showing antimicrobial properties (Bock et al., 2014) Therefore, TCA (ACROS Organics™) was evaluated as well in the present study; the product was dissolved in acetone (Bock et al., 2014).

Assessments of test solutions on oomycete growth were carried out on PDA plates, including the following 17 treatments: fermentation broth of each of the eight bacteria (10% v/v) with TCA (0.2% v/v), and fermentation broth of each of eight bacteria (10% v/v) without addition of TCA; TCA (0.2% v/v) was also added as a single treatment (without bacterial broth). Additionally there were four controls: TSY medium (10% v/v) + acetone (0.2% v/v), TSY medium (10% v/v), acetone (0.2% v/v), and sterile distilled water (SDW) (10% v/v). The cell-free supernatants and TCA were incorporated into PDA at 10 % (v/v). Prior to autoclaving PDA, the amount of prescribed distilled water was reduced by 10 % to allow for subsequent addition of treatment suspensions. When the media cooled to 45-50 °C, 10 % bacterial supernatant or 0.2% TCA was added and mixed
thoroughly before pouring the plates.

After the media plates had set, an oomycete colony (5 mm diameter) was inoculated in the center of each dish and incubated at 27 °C in the dark. After 2 days of incubation, the diameter of the oomycete colonies was measured in two perpendicular directions using the cross method (Hazir et al., 2016). All experiments had 6 replications and were performed twice in time. The inhibition rate of oomycetes was calculated by the formula (The inhibition rate = 100 × (the diameter in SDW control treatment – the diameter in treatments)/the diameter in SDW control treatment).

Zoospore germination was also assessed. Methods used for zoospore isolation was as outlined in Mbarga et al. (2012) with some modifications. A single oomycete colony (5 mm diameter) was inoculated in the center of PDA medium and incubated at 27 °C in the dark. After 7 days of incubation, oomycete culture was sequentially mixed with 10 ml of sterile distilled water and ground using a mixer. The zoospores were then counted, adjust to 3×10^6 CFU/ml, and immediately used in the bioassay. Zoospore germination assays were carried out on 24-well plates. Each well contained 100μl zoospore suspension and 800μl Potato Dextrose Broth (PDB) and the corresponding test solution. The experiment consisted of the same treatments as the aforementioned oomycete growth bioassay: fermentation broth of each of eight tested bacterial broths (100μl) + TCA (2μl); fermentation broth of each of eight tested bacterial broths (100μl); TCA (2μl); TSY medium (100μl) + acetone (2μl); TSY medium (100μl); acetone (2μl); sterile distilled water (100μl). The plates were incubated at 27 °C. Zoospore germination was assessed after 12 h using a concave slide and light microscope. We counted the number of germinated zoospores of 100 zoospores (Abdelzaher et al. 1994). A zoospore was considered germinated when the germ tube length was 1.5 times as much the zoospore diameter (Plascencia-Jatomea et al. 2010). All experiments had 6 replications and were
performed twice in time. The inhibition rate of oomycete was calculated by the formula (The
inhibition rate = 100 × (the number in SDW control treatment–the number in treatments)/ the
number in SDW control).

165  Antimicrobial effects of different concentrations of X. bovienii fermentation broth

In the previous experiment, the fermentation broth of X. bovienii (the symbiotic bacteria of S. feltiae
SN strain) exhibited a particularly promising level of antimicrobial activity considering mycelial
growth and zoospore germination. Therefore, we further evaluated the antimicrobial activity of X.
bovienii at different dose levels via mycelial growth and the zoospore germination assay. Five levels
of X. bovienii fermentation broth were included: 0.1%, 0.5%, 1%, 5%, 10% (v/v). TSY medium was
used as a control. The procedures of mycelial and zoospore preparation, as well as assessment of
mycelial growth and zoospore germination were the same as described above. All experiments had
6 replications and were performed twice in time.

175  Statistical analyses

Before conducting the statistical analysis, the data from two trials were tested for the significant
differences. For the data of two trials on the mycelial growth, the data were combined as there was
no significant differences. For the data on the zoospore germination, the data were used separately,
as there were significantly differences between the two trials. The antimicrobial effects on mycelial
growth and zoospore germination of P. myriotylum were analyzed using a two-way analysis of
variance, where TCA and the strains of symbiotic bacteria were used as fixed factors. Data for
mycelial growth and zoospore germination in all experiments were also analyzed by one-way
analysis of variance (ANOVA) to evaluate the antimicrobial effects of different solutions on P.
myriotylum. Means were compared by the Tukey’s multiple range test ($P \leq 0.05$). Data were presented as means ± standard error. The data analysis was performed by SPSS for windows version 21.0.

Results

Treatment effects on mycelial growth

The results of two way ANOVA showed that the main effects bacteria strain and TCA significantly impact mycelial growth and there was a significant interaction effect as well (Strain: $F_{8,222} = 409, p < 0.001$; TCA: $F_{1,222} = 1516, p < 0.001$; Strain×TCA: $F_{8,222} = 93.5, p < 0.001$). The treatments had significant effects on mycelial growth based on the results of one-way ANOVA ($F_{19, 239} = 365, P < 0.001$) (Fig. 1). Results showed that TCA alone significantly inhibited mycelial growth. The maximum inhibition rate of mycelial growth (47.5%) was observed in the treatment of the combination of TCA and Xb. Significant inhibition rate of mycelial growth was observed in the TCA treatment (31.5%) as compared with three controls : TSY+Acetone (0.47%), TSY (0.97%), Acetone (2.15%) whereas the treatment of Xb (35.98%) and Xc (33.91%) alone had the comparable inhibitory effect as TCA (Fig. 1). Moreover, we observed that the addition of TCA significantly enhanced the effect (growth inhibition) in all bacteria except not in Xs. With regard to fermentation broth, Xs had a stronger negative influence on mycelial growth than TCA.

Treatment effects on zoospore germination

Both trials showed similar patterns of inhibition rates of zoospore germination as shown in Fig. 2. The inhibition rates of zoospore germination were enhanced both by the addition of bacteria fermentation broth (Trial 1: Strain: $F_{8,102} = 1488, p < 0.001$; Trial 2: Strain: $F_{8,102} = 1638, p < 0.001$;
Fig. 2) and TCA (Trial 1: TCA: $F_{1,102} = 911, p = 0.001$; Trial 2: TCA: $F_{1,102} = 1059, p < 0.001$; Fig. 2). Xb alone showed the maximum inhibition rate of zoospore germination as compared to other treatments with bacteria alone, with 34.78% and 38.47% in trial 1 and trial 2, respectively. However, the effects TCA addition on inhibition rate of zoospore germination were dependent on the bacteria strain. Treatments had significant effects on the inhibition rate based on one way ANVOA (Trial 1: $F_{19,119} = 10.1, p < 0.001$; Trial 2: $F_{19,119} = 29.8, p < 0.001$). Bacteria fermentation broth could inhibit zoospore germination significantly, whereas TCA alone didn’t show any inhibition activity with the low value of 1.02% in trial 1 and 6.76% in trial 2.

**Effects of different *X. bovienii* fermentation broth dose levels on mycelial growth and zoospore germination**

The colony size of *P. myriotylum* was inversely proportional to the dose of fermentation broth of *X. bovienii* (Fig. 3) ($F_{4,59} = 217, p < 0.001$). The maximum value of inhibition rate (31.18%) was observed at the level of 10%. Bacteria fermentation broth, with concentrations of 0.5% and 0.1%, did not show clearly antimicrobial activity.

Similarly, the inhibition rate of zoospores germination was directly proportional to the high dose of *X. bovienii* fermentation broth(10%, 5%, 1%). whereas low dose of fermentation broth (0.5%, and 0.1%) showed the pattern of stimulating the germination of zoospores (Fig. 4) ($F_{4,59} = 60.5, p < 0.001$).

**Discussion**

Ginger soft rot is the most devastating diseases to ginger (Wang *et al.*, 2003). Control of soft rot is difficult because *Pythium* spp. can persist in the soil for years once introduced (Le *et al.*, 2014). In
the present experiment, we evaluated the antimicrobial activity of eight species of symbiotic bacteria isolated from the corresponding eight species of EPNs against *P. myriotylum* and the effects of TCA addition. Results showed that all the fermentation broths inhibited mycelial growth and zoospore germination of *P. myriotylum*, especially *Xenorhabdus bovienii*. Furthermore, we found that *X. bovienii* fermentation broth at doses of 10%, 5%, 1% significantly inhibited mycelial growth. In addition, *trans*-cinnamic acid, as one of the major bioactive ingredients of EPN symbiotic bacteria, also inhibited mycelial growth, but not zoospore germination. These results indicated that *X. bovienii* and TCA could be potentially used as a biorational solution for *P. myriotylum*. This approach could be a replacement of chemical fungicides in the suppression of soil-borne diseases caused by *P. myriotylum*, and thereby promote the sustainable production of ginger and other crops. Previous research reported that compounds produced by *Trichoderma viride* inhibit the growth of *P. myriotylum* and other fungi (Jeerapong *et al.*, 2015). In addition, *T. harzianum* and *T. saturnisporum* also showed strong antagonism against *P. splendens* in vitro (Jeerapong *et al.*, 2015). Seed coating treatments of ginger with *Trichoderma* spp. significantly alleviated ginger soft rot disease compared to the control without *Trichoderma* spp. coating (Ram *et al.*, 2000). Singh (2011) found that *T. harzianum* was able to suppress soft rot at the same level as Ridomil. Interestingly, seeds dipped in a suspension of *Pseudomonas fluorescens* or *Bacillus* sp. combined with inoculation with mycorrhizae *Glomus* sp. during the transplantation process showed substantial suppression of soft rot in ginger (Bhai *et al.*, 2005). Shanmugam *et al.* (2013) also reported that though the efficacy of biological control agents were lower in the fields, better control was achieved when several biological control agents were incorporated together. Conceivably, *Trichoderma* spp. could be combined with the TCA, *Xenorhabdus* or *Photorhabdus* treatments for improved control.
Antimicrobial activity of symbiotic bacteria against *P. myriotylum* depends on the species of symbiotic bacteria from entomopathogenic nematodes. In the present study, *X. bovienii* exhibited the highest level of suppression of *P. myriotylum* mycelial growth as compared to other symbiotic bacteria species. In contrast, inhibition of zoospore germination did not vary as markedly among the bacteria species.

In a previous study, *X. cabanillasii* showed antifungal activity to the plant pathogen *Fusicladium carpophilum* relative to the negative control and *X. bovienii* (Hazir et al., 2016). In the same paper, the supernatants of *X. bovienii* and of *Photorhabdus* spp. did not cause any suppression of growth for the plant pathogen *Monilinia fructicola* relative to the control whereas *X. cabanillasii* caused significant suppression (Hazir et al., 2016). Thus, the antimicrobial activity of various symbiotic bacteria clearly varies with specific plant oomycete pathogens. Similar results were in line with (Bőszörményi et al., 2010) who tested the efficacy of cell-free filtrates of 18 strains of *Photorhabdus* and *Xenorhabdus* against the plant pathogenic bacteria *Erwinia amylovora* and the oomycete *Phytophthora nicotianae*. Vanitha et al. (2010) also observed that antimicrobial activity against *Fusarium oxysporum* (Vanilla), *Alternaria solani* (Tomato), *Sclerotium rolfsii* (Brinjal) and *Aspergillus niger* varied with the bacterial strain of symbiotic bacteria among *Xenorhabdus* spp.

Given the diversity of effects found among bacterial strains and species, in future studies, we might collect more EPN species and evaluate their potential antimicrobial activity against *P. myriotylum*, especially additional strains of *S. feltiae*. With regard to *X. bovienii*, it will be beneficial to trace the antimicrobial activity of the potential antimicrobial compounds (Houard et al., 2013) in their metabolites, and to explore their commercial potential to control plant disease in the field.

*Trans*-cinnamic acid (TCA) in combination with different strains of symbiotic bacteria showed
different antimicrobial activity effects. The compound indicated varying effects among the strains of symbiotic bacteria. TCA was discovered in the metabolites of *Photorhabdus luminescens* (Bock *et al.*, 2014). In the present study, TCA also showed clear suppression on *P. myriotylum*. Mycelial growth was about half as much as the control.

Moreover, we also found the addition of TCA to symbiotic bacteria metabolites of *X. nematophila*, *X. cabanillasii*, *X. szentirmai* and *P. akhurstii* significantly increased their antimicrobial activity. This result is in line with a previous study which demonstrated the combination of TCA with Xs metabolites lead to lower vegetative growth of *Monilinia fructicola* as compared to either of the two treatments alone (Hazir *et al.*, 2017). Similarly, Letsididi *et al.* (2018) found that TCA could inhibit the growth of *Staphylococcus aureus*, *Salmonella typhimurium*, *Pseudomonas aeruginosa*. Additionally, TCA was also reported as an antibacterial compound of *Mycobacterium tuberculosis* (Chen *et al.*, 2011).

*Xenorhabdus* spp. and *Photorhabdus* spp. have been reported to produce various compounds besides *trans*-cinnamic acid (TCA), which shown various functionalities for agricultural and pharmaceutical purposes (Araniti *et al.*, 2018; Chen *et al.*, 2011; Letsididi *et al.*, 2018; Lupini *et al.*, 2016). With regard to plant pathogens, there is a need to find more compounds from symbiotic bacteria associated with various EPNs from natural habitats, and subsequently the antimicrobial properties of these compounds can be evaluated systemically (this should be done with other bacteria that may have suppressive potential as well).

In conclusion, we found that *X. bovienii* metabolites and TCA showed strong antimicrobial activities against *P. myriotylum*, which causes ginger soft rot and disease in other crops. The supernatant of
X. bovienii has the potential to be used in a large scale for ginger seed treatment, such as dipping the ginger seed in the supernatant during the transplant stage. The present study provides insight into a potential integrated management system for reducing plant pathogenic fungi. Further studies should be performed to clarify the function of X. bovienii metabolites and TCA alone, or their combination, to control P. myriotylum. With regard to the broad array of plant pathogenic fungi that threaten various crops, we still need to assess suitable symbiotic bacteria (associated with EPN) for possible suppression tactics. Given the regulatory status of many chemical pesticides and their threat to the environment, it is necessary to build an EPN/bacterial resource depository at a national scale or even a worldwide scale to cope with the increasing challenge of various plant pathogenic fungi and insects pests.

Acknowledgments

This work was jointly supported by National Key R&D Program of China (2017YFE0130400 and 2018YFD0201002) and Natural Science Foundation of China (31470495 and 31170412) and Yunnan Provincial Company of National Tobacco Corporation (2017YN15 and 2014YN21), the Fundamental Research Funds for the Central Universities to Dr. Ruan and 111 project (B08011). We gratefully acknowledge anonymous reviewers for valuable comments on the manuscript.

Compliance with Ethical Standards

The authors declare that ethical standards have been followed and that no human participants or animals were involved in this research.

Competing interests

The authors declare that they have no competing interests.
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compound, trans-cinnamic acid, produced by *Photorhabdus luminescens*, a potential biopesticide against pecan scab. *Journal of Pest Science, 87*(1), 155-162


Kavitha, P. G., & Thomas, G. (2008). Expression analysis of defense-related genes in Zingiber (Zingiberaceae) species with different levels of compatibility to the soft rot pathogen Pythium aphanidermatum. Plant Cell Reports, 27(11), 1767-1776


Table 1. The origin and abbreviation of entomopathogenic nematodes and their symbiotic bacteria

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<thead>
<tr>
<th>EPN</th>
<th>Symbiotic bacteria</th>
<th>Abbreviation</th>
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<tr>
<td><em>Steinernema feltiae</em> (SN strain)</td>
<td><em>Xenorhabdus bovienii</em></td>
<td>Xb</td>
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<tr>
<td><em>S. carpocapsae</em> (All strain)</td>
<td><em>X. nematophila</em></td>
<td>Xna</td>
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<td><em>S. carpocapsae</em> (Cxrd strain)</td>
<td><em>X. nematophila</em></td>
<td>Xnc</td>
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<td><em>S. rarum</em> (17 C&amp;E strain)</td>
<td><em>X. szentirmaii</em></td>
<td>Xs</td>
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<tr>
<td><em>S. riobrave</em> (7-12 strain)</td>
<td><em>X. cabanillasi</em></td>
<td>Xc</td>
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<td><em>Heterorhabditis bacteriophora</em> (VS strain)</td>
<td><em>Photorhabdus luminescens</em></td>
<td>Plv</td>
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<td><em>H. bacteriophora</em> (HB1 strain)</td>
<td><em>P. luminescens</em></td>
<td>Plh</td>
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<tr>
<td><em>H. indica</em> (HOM1 strain)</td>
<td><em>P. akhurstii</em></td>
<td>Pa</td>
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Fig. 1. Effects of fermentation broth of different entomopathogenic nematode symbiotic bacteria on mycelial growth of *Pythium myriotylum*. Both fermentation broth and *trans*-cinnamic acid (TCA) inhibited the mycelial growth of *P. myriotylum*. Moreover, the addition of TCA significantly enhanced the effect (growth inhibition) in all bacteria but Xs (Xb: *Xenorhabdus bovienii*, Xna: *X. nematophila*, Xnc: *X. nematophila*, Xs: *X. sertirmaii*, Xc: *X. cabanillasii*, Plv: *Photorhabdus luminescens*, Plh: *P. luminescens* and Pa: *P. akhurstii*). TSY: trypticase soy yeast). ‘TCA+’ indicate the addition of TCA. ‘TCA-’ indicate no TCA treatment. ‘CK’ indicate control treatments. Different letters above bars indicate significant differences between treatments.

Fig. 2. Effects of fermentation broth of different EPN symbiotic bacteria on zoospore germination of *Pythium myriotylum*. Both fermentation broth and *trans*-cinnamic acid (TCA) could inhibit the zoospore germination of *P. myriotylum*. However, TCA and fermentation broth show antagonistic effect on inhibition of germination. ‘TCA+’ indicate the addition of TCA. ‘TCA-’ indicate no TCA treatment. ‘CK’ indicate control treatments. Different letters above bars indicate significant differences between treatments.

Fig. 3. Effect of *Xenorhabdus bovienii*, symbiotic bacteria (isolated from *Steinernema feltiae*) on mycelial growth of *Pythium myriotylum*. Fermentation broth at a concentration of 10%, 5% and 1%(v/v) shown antimicrobial effects against *P. myriotylum* which was inversely proportional to the dose. Different letters above bars indicate significant differences between treatments.

Fig. 4. Effect of *X. bovienii*, symbiotic bacteria (isolated from *Steinernema feltiae*) on zoospore germination of *Pythium myriotylum*. Fermentation broth at a concentration of 10%, 5% and 1%(v/v) shown antimicrobial effects against *P. myriotylum* which was inversely proportional to the dose. However, low concentration (0.5%, 0.1%) of fermentation broth could promote zoospore germination. Different letters above bars indicate significant differences between treatments.
Fig. 1. Effects of fermentation broth of different entomopathogenic nematode symbiotic bacteria on mycelial growth of *Pythium myriotylum*.
Fig. 2. Effects of fermentation broth of different EPN symbiotic bacteria on zoospore germination of *Pythium myriotylum*. 
Fig. 3. Effect of *Xenorhabdus bovienii*, symbiotic bacteria (isolated from *Steinernema feltiae*) on mycelial growth of *Pythium myriotylum*.
Fig. 4. Effect of *X. bovienii*, symbiotic bacteria (isolated from *Steinernema feltiae*) on zoospore germination of *Pythium myriotylum*. 