



Biochemical Analysis of Banana Plants in Interaction between Endophytic Bacteria *Kocuria rhizophila* and the Fungal Pathogen *Fusarium oxysporum* f. sp. *cubense* Tropical Race (*Foc* TR4) [†]

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+ Presented at the 2nd International Electronic Conference on Plant Sciences—10th Anniversary of Journal Plants, 1–15 December 2021; Available online: https://iecps2021.sciforum.net/.

Abstract: Fusarium wilt, caused by *Fusarium oxysporum* f. sp. *cubense* Tropical Race 4 (*Foc* TR4), is one of the most severe banana diseases in the world. In this study, banana plants treated with endophytic bacteria *Kocuria rhizophila* showed increased PO enzyme activity, reaching the highest activity 72 h after inoculation in the shoots ($0.1640 \pm 0.0335 \mu mol/min$) and 24 h after inoculation in the roots ($0.0129 \pm 0.0024 \mu mol/min$). PPO enzyme activity increased significantly 24 h after inoculation in roots ($0.0131 \pm 0.0026 \mu mol/min$) and 6 h after inoculation in shoots ($0.0201 \pm 0.0065 \mu mol/min$). PAL enzyme activity on roots ($1.776 \mu mol/min$) and shoots ($1.2170 \mu mol/min$) inoculated with endophytic bacteria showed the highest value at 24 h. The highest total phenolic content in shoots treated with endophytic bacteria was at 72 h in roots (41.15384 mg GAE/g samples) and shoots (39.6102 mg GAE/g samples).

Keywords: fusarium wilt; endophytic bacteria; banana; biochemical assay

1. Introduction

The banana plant (*Musa* spp.) is a plant that can live in tropical and subtropical climates. This plant originates in Indomalayan realm and has many benefits, so the demand for bananas is very high. Based on the latest data from FAO 2019, banana exports were recorded at 20.2 million tons worldwide [1]. Unfortunately, banana production was declining due to fusarium wilt disease caused by *Fusarium oxysporum* f. sp. *cubense* Tropical Race 4 (*Foc* TR4). The use of endophytic bacteria as an alternative for controlling fusarium wilt has not been widely studied and could be an environmentally friendly alternative for controlling fusarium wilt. The endophytic bacteria *Kocuria rhizophila*, which has been isolated and investigated in previous research (unpublished data), was further analyzed at the biochemical level for its ability to induce systemic resistance.

2. Materials and Method

2.1. In Vitro Triple Culture

In vitro triple culture of banana, endophyte, and fusarium was performed based on the previous method [2]. Banana plantlets were obtained from a local commercial plant tissue culture laboratory. Banana plantlets that were at least seven days old after subculture were removed from the tube to be inoculated with an endophytic bacteria culture, using an insectarium needle, inserted perpendicularly to the pseudostem axis at a position about 0.5 cm above the root. Plantlets were subcultured to fresh MS0 medium and co-cultivated for seven days. On the seventh day after co-cultivation (7 hpi), the banana plantlets were



Citation: Candra, R.T.; Prasasty, V.D.; Karmawan, L.U. Biochemical Analysis of Banana Plants in Interaction between Endophytic Bacteria *Kocuria rhizophila* and the Fungal Pathogen *Fusarium oxysporum* f. sp. *cubense* Tropical Race (*Foc* TR4). *Biol. Life Sci. Forum* 2022, *11*, 84. https://doi.org/10.3390/ IECPS2021-11990

Academic Editor: Dimitris L. Bouranis

Published: 4 May 2022

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injured again with insectarium needles before being infected with *Foc* TR4. A piece of Potato Dextrose Agar (PDA), covered with 5–7 days culture of *Foc* TR4 mycelium, was placed near the point of injury. As a control for the effect of endophytic bacteria on banana plantlets, banana plantlets that were not pre-inoculated with endophytic bacterial isolates and not challenged with *Foc* TR4 were used.

2.2. Endophyte Bacteria Inoculation and Sampling Time Points

Samples used for biochemical assay were banana plantlets inoculated with endophyte bacteria and *Foc* TR4 (in vitro triple culture assay) with specific time points and harvested at 0, 6, 24, and 72 h of endophyte post-inoculation (0 hpi, 6 hpi, 24 hpi, 72 hpi, respectively). Controls used banana plantlets infected with *Foc* TR4 (in vitro triple culture assay) with specific time points and harvested at 0, 6, 24, and 72 h with composite triplicates sampling from three banana plantlets for roots and three banana plantlets for shoots, for PO and PPO assay.

2.3. Crude Enzyme Extraction

Crude enzyme extraction was carried out using the previously described method [3]. At this stage, crude enzyme extraction will be carried out for biochemical assay, including the peroxidase (PO), polyphenol oxidase (PPO), and phenylalanine ammonia-lyase (PAL) enzyme, and analysis of total phenolic content.

2.4. Peroxidase (PO) Enzyme Activity Assay

PO enzyme activity assay was carried out using the previously described method [3]. A total of 750 μ L of 0.1 M phosphate buffer solution (pH 6.5), 100 μ L of 0.05 M pyrogallol solution in buffer, and 240 μ L of 0.08% hydrogen peroxide solution were prepared and then resuspended until homogeneous. Then, the mixture was added with 20 μ L of the enzyme extract sample and the absorbance was measured at a wavelength of 430 nm with a microplate reader Infinite 200 PRO Nanoquant Microplate Reader (Tecan) (Switzerland) for 3 min by recording the absorbance every 30 s (0, 30, 60, 90, 120, 150, and 180 s) and averaged at the end. The activity was expressed as μ mol/min.

2.5. Polyphenol Oxidase (PPO) Enzyme Activity Assay

PPO enzyme activity assay was carried out using the previously described method [4]. A total of 200 μ L of the enzyme extract sample was added with 1500 μ L of phosphate buffer solution pH 6.5, then homogenized. The mixture was then incubated for 1 min at 25 °C. The absorbance was measured at a wavelength of 495 nm with a microplate reader Infinite 200 PRO Nanoquant Microplate Reader (Tecan) (Switzerland) for 3 min by recording the absorbance every 30 s (0, 30, 60, 90, 120, 150, and 180 s) and averaged at the end. The activity was expressed as μ mol/min.

2.6. Phenylalanine Ammonia Lyase (PAL) Enzyme Activity Assay

PAL enzyme activity assay was carried out using the previously described method [5]. A mixture of sample solutions, blanks, and standard solutions was prepared. After that, the solution mixture was heated using a water bath at 37 °C for 60 min. Then, 0.4 mL of 5 M HCl was added to stop the enzymatic reaction of the solution. The absorbance was measured at a wavelength of 290 nm using the microplate reader Infinite 200 PRO Nanoquant Microplate Reader (Tecan) (Switzerland). The standard curve equation was determined using the absorbance value of the standard solution. The activity was expressed as μ mol/min.

2.7. Measurement of Total Phenolic Content in Plant Tissue

The total phenolic content in the tissue was measured by the previously described method [6]. The 0.5 g samples were ground using liquid nitrogen. The obtained powder was mixed with 5 mL of 80% methanol (MeOH) and incubated for 2 h in a dark room at

room temperature. Then, the extracted solution was centrifuged at 10,000 rpm for 10 min to form a supernatant which would then be used as a sample for the content of plant phenolic compounds. A sample of 200 μ L of phenolic compounds was taken and put into a test tube, which was then added with 200 μ L of Folin-Ciocalteu reagent and homogenized. The solution was incubated for 10 min at room temperature and then added with 2 mL of 20% Na₂CO₃ solution. The solution mixture was homogenized and incubated again for 90 min in the dark at room temperature. The absorbance was measured by UV Vis Shimadzu UV-1280 spectrophotometry at a wavelength of 750 nm. For the standard, gallic acid solutions with concentrations of 0, 20, 40, 60, 80, and 100 g/mL were used. The blank used was distilled water with the same treatment as the sample. The total phenolic content in the tissue was determined based on the calculation of the standard gallic acid curve, which was further described as micrograms of gallic acid equivalent per gram (mg GAE/g) of the wet weight of the sample based on the Equation (1):

C (total phenolic content)(mg GAE/g) =
$$\frac{c (mg mL - 1) \times V (mL)}{mass (g)}$$
 (1)

2.8. Statistical Analysis

Statistical analysis was performed using IBM SPSS Statistics version 25. Statistical tests were considered significant if p < 0.05. All data were tested for normal distribution according to the Kolmogorov–Smirnov test and for homogeneity of variance among groups using Levene's test. After passing the normality test, the analysis of variance between groups was conducted using one-way ANOVA with Duncan posthoc test. When data failed the normality test and Levene's test, they were analyzed by the non-parametric Kruskal–Wallis test, followed by Kruskal–Wallis stepwise step-down multiple comparison posthoc test.

3. Results

Biochemical assay of different enzyme and phenolic content showed different results among root and shoot samples. Detailed results of each assay are described below.

3.1. PO Enzyme Activity Assay

It was observed that PO activity increased and reached its highest activity 72 h after inoculation in shoots ($0.1640 \pm 0.0335 \,\mu$ mol/min) and 24 h in the root ($0.0129 \pm 0.0024 \,\mu$ mol/min) (Figure 1). PO enzyme activity showed a significant difference between plants treated with endophyte treatment and control, across organ and sampling time points (P: 0.00 Kruskal–Wallis test, followed by a stepwise step down multiple comparison posthoc test). PO enzyme activity in roots treated with endophytic bacteria showed a decrease in PO enzyme activity at 6 h, then an increase at 24 h and a decrease at 72 h. The increase in PO enzyme activity at 24 h was thought to be due to the plant starting to produce systemic resistance. PO enzyme activity in shoots treated with endophytic bacteria showed a significant increase in enzyme activity. The highest PO enzyme activity was obtained at 72 h; this could have happened because the systemic resistance occurs in the roots and then spreads to the shoots [7], but, overall, the PPO enzyme activity was higher in shoots than in roots. This is not in accordance with the literature, which states that the activity of the PPO enzyme should fluctuate in the roots of banana plants, while the activity of the peroxidase enzyme tends to decrease in the leaves. This difference might have happened because the root tissue is a point of entrance and colonization of microorganisms, meaning that the focus of the physiological response of banana plants is prioritized in the roots [8].



Figure 1. PO enzyme activity at 0, 6, 24, and 72 hpi in control and treatment of roots and shoots of banana plants. Bars represent \pm SE of the mean. Different letters indicate significant differences among treatments within a time point (p < 0.05).

3.2. PPO Enzyme Activity Assay

PPO activity in banana plants treated with *K. rhizophila* was significantly increased when challenged with the pathogen and reached 24 hpi in roots $(0.0131 \pm 0.0026 \,\mu\text{mol/min})$ and 6 hpi in shoots $(0.0201 \pm 0.0065 \,\mu\text{mol/min})$ (Figure 2). PPO enzyme activity showed a significant difference between treatment and control across organ and sampling time points (P: 0.00 Kruskal-Wallis test, followed by a stepwise step down multiple comparison posthoc test). PPO enzyme activity in roots and shoots treated with endophytic bacteria and control had relatively the same trend, but overall, PPO enzyme activity was higher in shoots than in roots. This is not in accordance with the literature, which states that the activity of the PPO enzyme should fluctuate in the roots of banana plants, while the activity of the peroxidase enzyme tends to decrease in the leaves.



Figure 2. PPO enzyme activity at 0, 6, 24, and 72 hpi in control and treatment of roots and shoots of banana plants. Bars represent \pm SE of the mean. Different letters indicate significant differences among treatments within a time point (p < 0.05).

3.3. PAL Enzyme Activity Assay

PAL enzyme activity was higher in roots treated with endophytic bacteria. In general, roots and shoots treated with endophytic bacteria at 0, 6, 24, and 72 h were generally slightly higher than the control treatment (Figure 3), which showed the systemic induction of plant-defense enzymes. PAL enzyme activity on roots (1.776 μ mol/min) and shoots (1.2170 μ mol/min) inoculated with endophytic bacteria showed the highest value at 24 h. The activity of PAL enzymes in roots and shoots treated with endophytic bacteria and control had relatively the same trend. PAL activity in roots and shoots treated with endophytic bacteria at 6 h decreased enzyme activity, increased at 24 h, and decreased at 72 h. This is presumably due to the fact that banana plants treated with endophytic bacteria had synthesized PAL enzymes since 0 hpi, so they were able to induce other defense systems [9].



Figure 3. PAL enzyme activity in the control treatment and samples of roots and shoots of banana plants.

3.4. Total Phenolic Content

Roots and shoots treated with endophytic bacteria showed the total phenolic content tended to be stable in both roots and shoots, which could be seen from a similar pattern (Figure 4). The total phenolic content in roots and shoots treated with endophytic bacteria decreased at 24 h and increased at 72 h. Total phenolic content in roots (41.15384 mg GAE/g samples) and shoots treated with endophytic bacteria at 72 h (39.6102 mg GAE/g samples) was greatly increased, which indicated that the plant was under stress, resulting in necrosis [10].



Figure 4. Total phenolic content in the control treatment and samples of roots and shoots of banana plants.

4. Discussion

In general, roots and shoots treated with endophytic bacteria showed PO and PPO enzyme activities higher than the control treatment. This happened because the systemic resistance occurs in the roots and then spreads to the shoots. Therefore, the presence of high peroxidase enzyme activity in all treatment groups indicated that the plant response system in recognizing the presence of pathogens was running well; the induction of peroxidase activity indicated that there was stimulation of plant defenses to prevent the entry of pathogens [7].

PO is a component of an early response in plants to pathogen infection and plays a key role in the biosynthesis of lignin, which limits the extent of pathogen spread [7]. When the PO level increases due to the induced systemic resistance, a quick synthesis of reactive oxygen derivatives by oxidative burst leads to cell death and inhibits pathogenic activities that were observed [11]. Obtained results were supported by earlier studies that demarcated the induction of PO in plants infected by pathogens, resulting in faster and stronger resistance against them [12]. They experimentally supported the idea that peroxidase plays a defensive role against attacking pathogens [13].

In plant–pathogen interactions, PPO enzymes play a role in the oxidation process of phenol compounds into quinine compounds, which are quinone toxic compounds and will spread to injured plant tissues. In addition, these compounds will also cause environmental conditions that are not suitable for the development of pathogens, so the activity of this enzyme becomes important in plant resistance [14]. PO and PPO can catalyze the formation of lignin and other oxidative phenols and contribute to the formation of defense barriers by changing the cell structure defense system that gets actuated against pathogens [15]. In the present experiment, PPO activity was significantly enhanced by *K. rhizophila* treated banana plants. Moreover, PPO activity helps in disease resistance as it oxidizes the phenolic level increase during this stage to toxic molecules such as quinones, leading to the invasion of the pathogen [14].

PAL activity is an extremely sensitive indicator of stress conditions and fungal challenges, and elicitor treatment elevates the levels of the flux through the general phenylpropanoid pathway, thereby supplying the carbon skeletons for secondary products such as phenolics, which are the precursor molecules for lignin. Increased PAL enzyme activity can be induced in various ways, including elicitor, plant-pathogen interactions, and plant-biocontrol agent interactions. The observed increase in PAL activity in elicited roots is presumably related to the lignification process [9]. Figure 3 showed that the suspected banana plants treated with endophytic bacteria had synthesized PAL enzymes since 0 hpi, so they were able to induce other defense systems than the control.

PAL enzyme activity (Figure 3) and total phenolic content (Figure 4) showed that, between control and treatment, most of the endophytic effects were less visible because it was shown from previous research that *K. rhizophila* maintains primary metabolism instead of secondary metabolism under *Foc* TR4 infection [2]. *K. rhizophila* inoculation on soybeans also increased plant biomass [16]. Meanwhile, *K. rhizophila* inoculation on maize enhanced salt stress tolerances by regulating phytohormone levels, nutrient acquisition, redox potential, ion homeostasis, photosynthetic capacity, and stress-responsive gene expression [17].

Author Contributions: Conceptualization and methodology, L.U.K.; validation, L.U.K. and V.D.P.; formal analysis and investigation, R.T.C.; writing—original draft preparation, R.T.C.; writing—review and editing, L.U.K. and V.D.P.; supervision, L.U.K. and V.D.P.; project administration, R.T.C. All authors have read and agreed to the published version of the manuscript.

Funding: This research was funded by Atma Jaya Research and Community Service Center 2020, proposal number 1634.

Acknowledgments: We would like to thank Indonesian Tropical Fruit Research Institute for providing *Foc* TR4 culture.

Conflicts of Interest: The authors declare no conflict of interest.

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