

Phialomyces macrosporus REDUCES *Cercospora coffeicola* SURVIVAL ON SYMPTOMATIC COFFEE LEAVES

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ABSTRACT: Brown eye spot is among the most important coffee diseases, it is caused by a necrotrophic fungal *Cercospora coffeicola*. Saprobe fungi have potential in reducing the survival of necrotrophic pathogens and can act through competition of nutrients, mycoparasitism, antibiosis and resistance induction. We have screened saprobe fungi for the ability to reduce *C. coffeicola* sporulation and viability and determined the possible mechanisms involved in the biocontrol. The selected saprobe fungus, *Phialomyces macrosporus*, reduced the germination of *C. coffeicola* conidia by 40%. *P. macrosporus* produced both volatile and non-volatile compounds that inhibited *C. coffeicola* growth, sporulation and viability. The production of antimicrobial substances was the main mode of action used by the saprobe fungi. Therefore, *P. macrosporus* is a promising biological agent for the integrated management of brown eye spot.

Index terms: *Coffea arabica*, competition, biological control, brown eye spot, antibiosis.

Phialomyces macrosporus REDUZ A SOBREVIVÊNCIA DE *Cercospora coffeicola* EM FOLHAS DE CAFÉ SINTOMÁTICAS

RESUMO: A cercosporiose é uma das mais importantes doenças do cafeeiro e tem como agente etiológico o fungo necrotrófico *Cercospora coffeicola*. Os fungos sapróbicos têm potencial na redução da sobrevivência de patógenos necrotróficos e podem atuar por meio da competição de nutrientes, micoparasitismo, antibiose e indução de resistência. Nós selecionamos fungos sapróbios para a capacidade de reduzir a esporulação e viabilidade de *C. coffeicola* e determinamos os possíveis mecanismos envolvidos no biocontrole. O fungo saprobio selecionado, *Phialomyces macrosporus*, reduziu a germinação de conídios de *C. coffeicola* em 40%. *P. macrosporus* produziu compostos voláteis e não voláteis que inibiram o crescimento, esporulação e viabilidade de *C. coffeicola*. A produção de substâncias antimicrobianas foi o principal modo de ação utilizado pelos fungos sapróbicos. Portanto, *P. macrosporus* é um potencial agente biológico que pode ser usado no manejo da cercosporiose do cafeeiro.

Termos para a indexação: *Coffea arabica*, competição, controle biológico, mancha de olho pardo, antibiose.

1 INTRODUCTION

Coffee (*Coffea arabica* L.) is an important commodity worldwide. However, diseases are responsible for many losses in this crop. Brown eye spot (BES), caused by the necrotrophic fungus *Cercospora coffeicola* Berkeley & Cooke is considered one of the most important coffee diseases and affects both leaves and berries, causing yield losses of approximately 50% (Zambolim et al., 2016).

Although effective fungicide-based alternatives are available for the disease management (Patricio et al., 2008), it should not be the exclusive tool but part of the integrated brown eye spot management such as plant nutrition (Cardoso et al., 2013) and biological control (Sirinunta and Akarapsan, 2015).

Considering that conidia of *C. coffeicola* remain viable for up to nine months in coffee stubble (Zhang and Bradley, 2014) until favorable

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conditions for germination, the reduction of the initial inoculum should be taken into account as a strategy for the integrated management of the disease. This can be achieved by compost-accelerating products which encompasses both fast-growing fungi and cell wall degrading enzymes (Bellotte et al., 2009; Hauptman, 2014).

Because *C. coffeicola* is a necrotrophic fungus, its suppression by microbial antagonists may occur through a combination of mechanisms, such as the production of antimicrobial compounds, mycoparasitism and/or competition for nutrients. Competition for nutrients is likely to occur since hydrolytic enzymes produced by both pathogen and antagonist target the saprophytic survival on decaying leaves (Jurado et al., 2015; Alvarez-Rodriguez et al., 2016). Among these enzymes are pectinases, which are composed by an heterogeneous group of enzymes that degrades pectin (Falmy et al., 2008).

One such pectinase producers group are saprobic fungi. They play a significant role in the self-maintenance of a balanced ecosystem, mostly through the decomposition of organic matter, using a system of digestive enzymes that are secreted from the cells to the environment (Lin et al., 2015). Furthermore, they can withstand sudden variations in temperature and humidity and sustain biological control in such environment, which turn them promising biological control agents (Köhl et al., 1995). Currently, saprobe fungi have been used as a biological agent for coffee disease control (Botrel et al., 2018; Rodríguez et al., 2016). Therefore, we hypothesize that saprobe fungi might inhibit *C. coffeicola* growth in crop residues and its production antifungal compounds and cell-wall degrading enzyme may play a role in the pathogen displacement.

2 MATERIALS AND METHODS

2.1. Culture condition

The isolate LFP 37 of *Cercospora coffeicola* was obtained from the Culture Collection of *Cercospora coffeicola* at the Plant Pathology Department, Federal University of Lavras, Minas Gerais State, Brazil. Isolate LFP 37 was grown for 10 days on solid V8 medium (V8S) (200 mL of V8 juice, 800 mL of distilled water and 20 g of agar) at 25°C. A modified version of the methodology proposed by Souza et al. (2005) was used to sporulate the isolate LFP 37. The pathogen spore suspension was adjusted to 3×10^4 conidia mL⁻¹.

Three saprobe fungal strains were used in this study: *Phialomyces macrospores* (0053/07), *Curvularia inaequalis* (0005/06), and *Curvularia eragrostidis* (0047/06), obtained from the Culture Collection of the Microorganisms from Bahia, State University of Feira de Santana, Bahia, Brazil.

The strains were grown for 7 days in a medium of carrot maize agar (CMA) at 27°C. After incubation, one disk (7 mm) containing conidia and mycelium was suspended in 100 mL of CM liquid medium, grown in an orbital shaker at 120 rpm and 27°C. The suspension was shredded, homogenized in a blender, and adjusted to 3.4×10^7 CFU x mL⁻¹. Antagonistic fungi were inoculated by spraying the leaves until runoff, unless otherwise stated.

2.2. Coffee seedlings

Coffee seeds (Cultivar Mundo Novo IAC 376-4) were purchased from Agricultural Research Company of Minas Gerais (EPAMIG) and were seeded in plastic trays containing white sand for pre-germination. Two-month old seedlings were transferred to polyethylene bags (250 x 90mm) filled with substrate made from a dystroferic Red Latosol (700 L m⁻³) taken from a gully, and fertilize was by the addition of natural phosphate (5 kg m⁻³) and potassium sulfate (0.5 kg m⁻³) (MATIELLO et al., 2010). Plants were kept in a greenhouse at 25 °C –28 °C during the entire experiment.

2.3. Screening test

Pathogen inoculation consisted of spraying the inoculum suspension on the abaxial side of the coffee leaves (7-month-old) until runoff. Dark wet plastic bags were placed during 3 days to simulate a moist micro-chamber. At 15 days after inoculation (dai), when BES symptoms appeared, the three abovementioned antagonists were sprayed onto the leaves until runoff. Soil Set® (0.75 mL L⁻¹) associated with compost Aid® (0.75 mL L⁻¹) were used as positive control, as described by Bellotte et al. (2009). Distilled water was applied as a negative control. After saprobe application, 10 leaves from each treatment were stored in plastic bags at 28°C for 7 and 14 d. Number and germination of LFP 37 conidia were determined.

To determine the number of conidia, lesions of each leaf were excised and transferred to 10 mL tubes with 1 mL of distilled water with 300 µL of Tween 20% (v/v). Aliquots of each suspension

were counted using a haemocytometer. For the viability spore test, a 400 μ L aliquot of the suspension of each treatment was transferred to 90mm Petri dishes containing 15 mL of water-agar at 2% (w/v). The plates were incubated at 25°C for 6 h, and the number of germinated spores was counted under light microscope at 40 \times . Conidia were considered germinated when the germ tube was equal to or greater than half of the spore length (Beckman and Payne, 1983). The experiment was performed twice, and the fungus with the highest potential to reduce pathogen survival was selected to continue the study. The experimental design was a randomized block with five treatments (*C. inaequalis*, *C. eragrostidis*, *P. macrosporus*, Compost Aid+ Soil Set and water), five plants per experimental unit and three replications.

2.4. Assessment of antagonist drought-tolerance

Drought tolerance was assessed to study the ability of *P. macrosporus* to reduce *C. coffeicola* isolate LFP 37 survival in plant lesions under humidity variation to simulate field conditions (Köhl et al., 1995). Fungal growth, plant cultivation, and fungal spray were performed as described above. Immediately after antagonist application, wetness was interrupted. Thereafter, 10 leaves per treatment were placed on two layers of sterile dry filter paper in open Petri dishes in laminar flow cabinet for 9 h. The leaves were incubated for two different periods of interrupted wetness (0 and 61 h) after the antagonist application. (Köhl et al., 1995). After the dry period, the filter paper was wetted with 2 mL of sterile tap water. Then, the Petri dishes were closed and kept under the same conditions, as described above. After 7 days, number and germination of pathogen conidia were determined, as previously described. The experiment was set up in a completely randomized block design, using a factorial scheme (1 x 2) with *P. macrosporus* fungus and two different periods of interrupted wetness (0 and 61 h). Each treatment was represented by three replicates and each replicate was composed with 10 leaves per plot.

2.5. Antibiosis and volatile production test

A method of measuring the direct activity of antagonists against pathogens was used in the *in vitro* antibiosis and volatile production test (Prasad et al., 2011). These assays were performed by cultivating the fungi on regular and bipartite Petri for antibiosis and volatile production, respectively.

The putative ability of *P. macrosporus* to produce different molecules or enzyme toxic to the pathogen was assessed by the confrontation assay. Plates were incubated at 25°C and 12 h in the light for 5 days. The assay was evaluated by cultivating *P. macrosporus* on one side of the Petri dish and *C. coffeicola* on the opposite side. The fungi were cultivated on Potato Dextrose Agar (PDA, 39 g L⁻¹), CMA and V8 media. The negative control contained only the pathogen in one side of the plates. The effect of toxic molecules or enzyme produced by *P. macrosporus* on *C. coffeicola* was quantified by comparing the mycelia growth and number of conidia. The experiment was distributed in a randomized block design with four replicates and was performed twice.

The production of fungitoxic volatiles compounds by *P. macrosporus* was evaluated by cultivating *P. macrosporus* in one side of a bipartite Petri and *C. coffeicola* in the opposite side. Plates were incubated at 25°C and 12 h in the light for 5 days. Saprobe fungi were cultivated on PDA, CMA and V8. The negative control contained only the pathogen in one side of the plates. The effect of fungitoxic volatiles compounds by *P. macrosporus* on *C. coffeicola* was quantified by comparing the mycelial growth, number and germination (%) of conidia. The experiment was distributed in a randomized block design with four replicates and was performed twice.

2.6. Pectinase production

The ability to produce hydrolytic enzyme in solid medium was assessed by growing *P. macrosporus* on mineral medium (pH 7.2) (2.0 g L⁻¹ KH₂PO₄; 7.0 g L⁻¹ K₂HPO₄; 1.0 g L⁻¹ (NH₄)₂SO₄; 1.0 g L⁻¹ MgSO₄·7H₂O; yeast extract 0.06% (p/v); and 13 g L⁻¹ agar). The medium was supplemented with 3 g L⁻¹ citrus pectin and incubated at 27°C for 5 days. Agar discs (7 mm diameter) containing mycelia from the colonies were transferred to Mac McIlvaine buffer (0.2 M NaHPO₄, 0.1 M citric acid, 13 g L⁻¹ agar-pH 6.0) with 0.25% (w/v) citrus pectin and incubated at 40°C for 48 h. A solution of I₂/KI (1 g of I₂ and 5 g of KI in 330 mL of H₂O) was added to detect the clearing zones.

2.7. Fungal cultivation for Polygalacturonase (PG) assays

P. macrosporus was cultivated on solid minimum medium containing 6,0 g L⁻¹ NaNO₃; 1,5 g L⁻¹ KH₂PO₄; 0,5 g L⁻¹ KCL; 0,5 g L⁻¹ MgSO₄·7H₂O; 0,01 g L⁻¹ FeSO₄; 0,01 g L⁻¹ ZnSO₄;

10,0 g L⁻¹ glucose; 15 g L⁻¹ agar at 6,8 pH for 7d, to a final concentration of 10⁶ esporos mL⁻¹. Then, one disk (7 mm) containing conidia and mycelium was inoculated into erlenmeyer flasks (125 mL) containing 50 mL of liquid mineral medium (6.8 g K₂HPO₄; 3.8 g KH₂PO₄; 1.0 g (NH₄)₂SO₄; 1.1 g MgSO₄·7H₂O; 0.6 g yeast extract, 3 g pectin 2% in one liter of distilled water) with one disk (7 mm) of mycelium and shaken at 150 rpm at 27°C for 5 days. The culture filtrates were used to directly estimate PG activity and the mycelia were collected daily to quantify total mycelial dry mass.

2.8. Assessment of Fungal growth and PG activity

The fungal growth was quantified by dry weight for 5 days. For this, the mycelium was collected by filtration through nylon cloth under vacuum and dried at 60°C for 48 h. The dried mycelium mass was measured using a precision balance (0.01 g precision).

The PG activity was determined by mixing 1 mL of culture filtrate in 1 mL of substrate solution (100 mM sodium acetate buffer; [pH 4.5]; 0.2 mM NaCl, 1.2% polygalacturonic acid [w/v]) and incubated at 40°C for 20 min. The reducing sugars released in the medium were measured by using the 3,5-dinitrosalicylic acid methods with galacturonic acid as a control. One PG unit was defined as micromoles of galacturonic acid released per minute per millilitre.

2.9. Assessment of each fungal derivate to pathogen viability reduction

To determine the ability of each fungal-derived product or enzyme to displace the pathogen from lesions were carried out *in vitro* and *in vivo* assays. For the *in vitro* assay, coffee leaf samples naturally infected by isolate LFP 37 were collected from field-grown fungicide-free plants. The leaves were distributed in a randomized-block design with four replicates and five to eight injured leaves per treatment per plot. Treatments were: i) water; ii) Conidia suspension (10⁶ spores mL⁻¹) of *P. macrosporus*; iii) polygalacturonase enzyme produced by the saprobe iv) conidia +supernatant of the saprobe and v) the supernatant of the saprobe growth.

In every case, the fungi were grown for the previously determined minimum time necessary for the maximum activity of polygalacturonase, 120 hours. After 7 days of incubation, the number of conidia of LFP 37 and leaf weight were assessed.

For the *in vivo* assay, coffee seedlings cv. 'Mundo Novo IAC 376-4' were inoculated with isolate LFP 37 and from the onset of symptoms (25 dai), each fungal-derived product or enzyme mentioned above was sprayed onto the leaves until runoff. Then, 10 leaves from each treatment were used to determine the pathogen conidia count and initial germination. After 7 days of incubation, the number and final germination of conidia were assessed. The plants were distributed in a randomized-block design with four replicates and one seedling per plot. The treatments were the same used for the *in vitro* assay. The *in vivo* assay was performed twice.

2.10. Experimental design and statistical analysis

Data from different experiments were tested for normality and submitted to an analysis of variance (ANOVA), Tukey and Scott-Knott multiple range tests ($P=0.05$). Data related to Polygalacturonase activity and dry mycelial weight of *P. macrosporus* were submitted to regression equations and subjected to a parallelism (*F*-test) test. The goodness of fit of the models was evaluated by coefficient of determination (R^2). Data analysis was performed in the R software version 3.1.1.

3 RESULTS AND DISCUSSION

To the best of our knowledge, for the first time a biological control option for BES targeted its saprophytic phase was best achieved by the use of *P. macrosporus* (Figure 1). In the screening assay, there was no significant interaction between the treatments and the time when the experiment was carried out for either the conidia count ($P=0.46$) or the viability of LFP 37 conidia ($P=0.14$). Regarding LFP 37 conidial germination, these *in vivo* assays showed that the application of *P. macrosporus* to the injuries caused by the pathogen reduced the viability of LFP 37 by 40% in 7 days ($P=0.0003$) (Figure 1). This effect persisted for at least 14 days after application of the antagonist under study (Figure 1), showing that not only the saprobe was able to reduce the viability of the pathogen, but this effect also occurs at the earliest considered time point (7 days after treatment). Odile-Mathieu and David Daniel (2006) studied isolates of *Microsphaeropsis ochracea* in the lesions of onion leaves during senescence and reported a reduction in the sporulation of *Botrytis squamosa* from 10 days after the application of the antagonist.

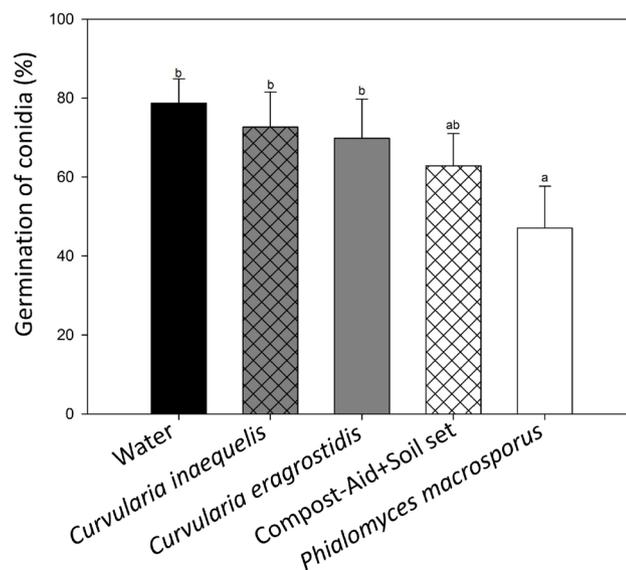


FIGURE 1 - Germination of *Cercospora coffeicola* conidia evaluated at 0, 7, and 14 days after the application of water, *Curvularia inaequalis*, *Curvularia eragrostidis*, Compost-Aid+Soil set and *Phialomyces macrosporus*. Bars with the same letter are similar at the 5% level according to Tukey multiple range test. CV=24.85%. The line on each bar represents \pm SE.

In our study, the contribution of the reduced inoculum of *Cercospora* to the occurrence of BES in the field was not evaluated, but a similar study by Galletti et al (2008) showed that when *Trichoderma* sp. was applied to the leaves of sugar beet infected with *Cercospora beticola*, the viability of the pathogen spores decreased, resulting in a delay of the epidemic in the subsequent crop.

The performance of antagonists under field conditions is determined mainly by their ecological competence. Regardless water stress condition ($P=0.82$ and $P=0.95$), there was a reduction of conidial viability of up to 37.72%. This result is in agreement with Köhl et al. (1995), who showed suppression of sporulation of *Botrytis aclada* in dead onion tissue treated with the antagonist *Ulocladium atrum* in bioassays with repeated wet-dry-cycles. In addition to evaluating the potential of the saprobe to control the pathogen regardless of the stress to which it is exposed, which aims to determine the sustainability of the control technique, it is necessary to evaluate the action mechanisms of the biocontrol agent. One of the most common mechanisms of reduced survival of pathogens is antibiosis. Hiradate et al. (2002) observed a decrease in the survival of *Colletotrichum dematium* on mulberry leaves infected when treated with *Bacillus amyloliquefaciens*, confirming that the

production of antifungal compounds was the primary mechanism of action that was used by the antagonists under study.

For the antibiosis test, interaction was observed between medium and treatments ($P=0.0090$). A positive interaction was observed between culture media CMA ($P=0.0018$), V8 ($P=\leq 0.001$), and production of non-volatile compounds, resulting in greater pathogen control (Figures 2 and 3).

Reduction in the mycelium growth, sporulation and spore germination was observed when *C. coffeicola* was cultured in the presence of the *P. macrosporus* (Figure 4 A, B e C). Thus, it is suggested that such culture media induce the production of a greater amount or range of compounds with an inhibitory action on isolate LFP 37. Similar result was obtained by Abe et al. (2015) who isolated a class of secondary metabolites, polyketides from the fungus *P. macrosporus*, which is known to provide survival advantages to microbial hosts. Some of the benefits could be the production of antibiotics, as reported by Zhang et al. (2015) who studied the antibiosis functions during interactions of *Trichoderma* spp. with plant pathogenic *Rhizoctonia* and *Pythium*. The authors confirmed that polyketide positively correlated to the production of antibiotics.

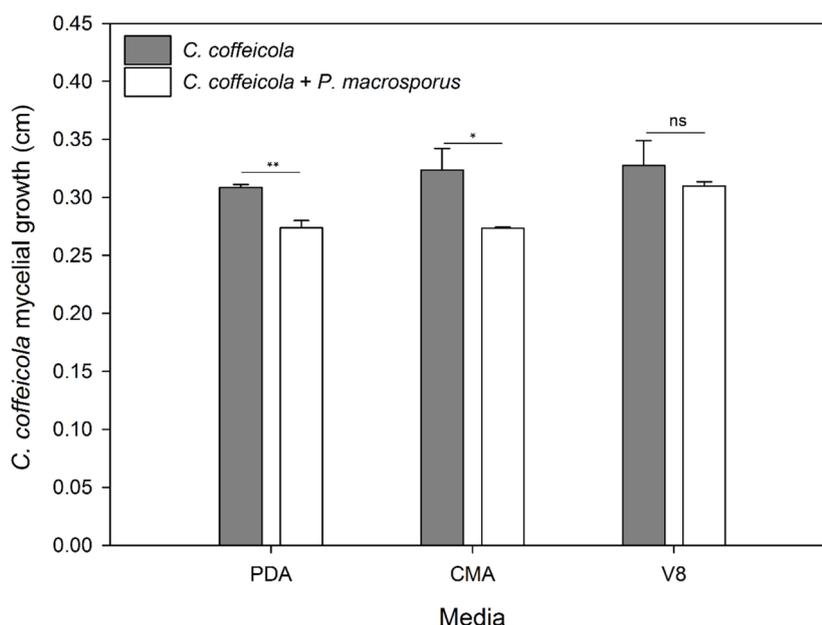


FIGURE 2 - *Cercospora. coffeicola* mycelial growth in confrontation test against *P. macrosporus* on PDA, CMA, and V8 media. Related graphic analysis of the unfolding of the treatments (*C. coffeicola* and *C. coffeicola* + *P. macrosporus*) in each culture medium. ^{ns}Non significant by Student’s t-test. CV=10.60%. Each bar represents the mean of four replicates (Petri dishes) per treatment per culture medium.

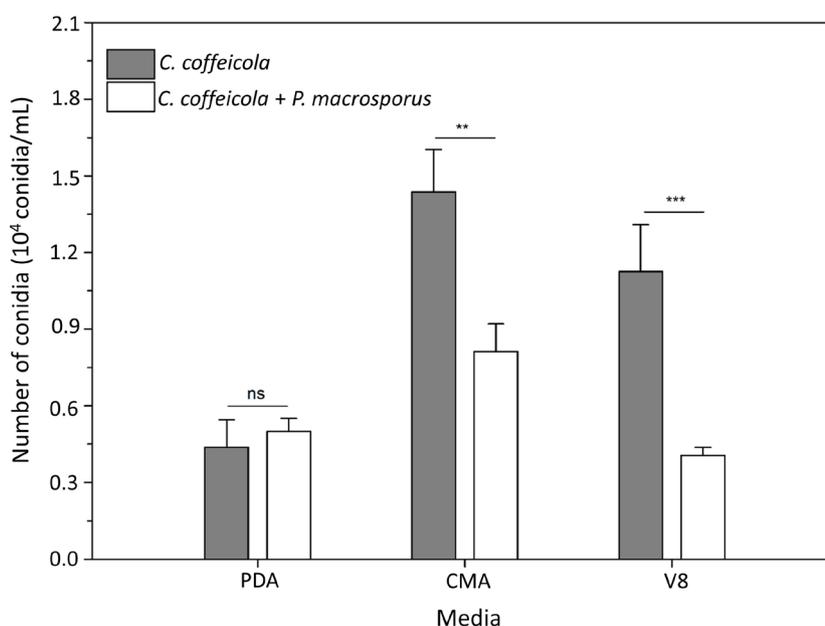


FIGURE 3 - Number of conidia of *Cercospora coffeicola* in volatile test against *P. macrosporus* on PDA, CMA, and V8 media. Graphic analysis of the unfolding of the treatments (*C. coffeicola* and *C. coffeicola* + *P. macrosporus*) within each culture medium. Significantly different according to the Student’s t-test **($P \leq 0.01$) or ***($P \leq 0.001$). ^{ns}Non significant by the Student’s t-test. CV=30.66%. Each bar represents the mean of four replicates (Petri dishes) per treatment per culture medium.

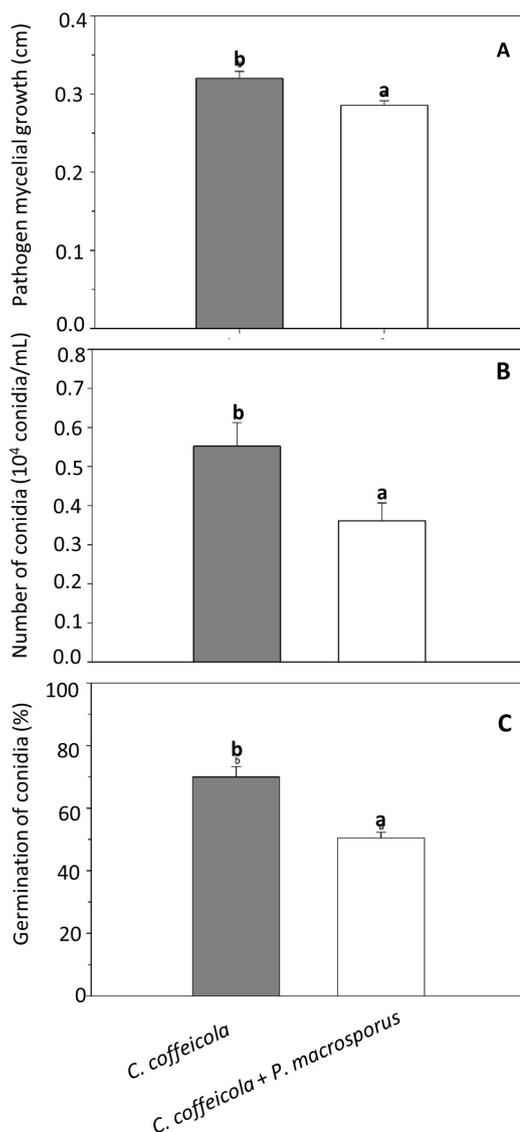


FIGURE 4 - Effect of volatile production by *Phialomyces macrosporus* on mycelium growth (A), sporulation (B) and spore germination (C) of *Cercospora coffeicola*. The bars with the same letter are similar at the 5% level according to the Student's t-test ($P \leq 0.05$). CV = 13.23%, 39.11%, and 7.80% for graphs A, B, and C, respectively. The line on each bar represents \pm SE.

The largest dry micelial weight was observed at 120 hours (Figure 5A). The enzymatic activity of PG, detected in the culture supernatant, showed a gradual increase over the growing time, with the highest activity in 120 hours (Figure 5B). The competition mechanism between microorganisms in a settling necrotic area was tested by determining and quantifying pectinolytic enzymes. These enzymes degrade dead plant tissue and release nutrients for both the pathogen and saprobe. Thrane et al. (2000) monitored GFP-

tagged *Trichoderma* spp. saprobes in cucumber plants infected with *Pythium ultimum* and found that the biocontrol isolates were able to take up the nutrients in the plant tissue more efficiently than was the pathogen, confirming that competition for nutrients is an important mechanism in this biocontrol pathosystem. Rodríguez et al. (2016) recently confirmed this hypothesis by determining that the niche overlap index between *P. macrosporus* and *Colletotrichum gloeosporioides* was 100%, i.e., all nutrients the

pathogen uses, the biocontrol agent also does and surviving in the necrotrophic tissue provoked by the pathogen.

Furthermore, in addition to externally colonizing plants, one important feature of fungi is the colonization of live plant organs and exerting benefits to the plant over the long term (Barahona et al., 2011). Reports by Ahmad et al. (2003) showed that *P. macrosporus* colonizes coffee berries of robusta and arabica. Therefore, similar to *Cladosporium cladosporioides* isolates, *P. macrosporus* growth may acts as a barrier to the entry of other harmful fungi quality of coffee. Furthermore, *C. cladosporioides* is characterized as a saprophyte fungus and PG enzyme producer (Rezende et al., 2013).

In this study, we identified the production of PG on solid medium through the formation of the corresponding halo enzyme production. The average diameter of the formed halo was 0.93 cm. However, when we compared the *P. macrosporus* activity of PG (744.2685 U/mL) with microorganisms that have the ability to secrete large amounts of this protein, such as *Penicillium griseoroseum* (1600 U/mL), we cannot consider this saprobe a significant PG producer (Minussi et al. 1998).

Although, we suggest a synergistic activity of this enzyme and antibiosis act on the pathogen displacement from the lesion hence in reduction of its survival on detached leaves and in vivo test. Further studies are necessary to investigate the role of PG in the observed biocontrol.

Contribution of each fungal derivate, such as conidia and any metabolite, in reducing the number of pathogen conidia and lesion degradation were evaluated by *in vitro* and *in vivo* assays. From the *in vitro* experiment, this supernatant was not effective in the degradation of lesion tissue in the short-term evaluation (7 days after treatment) ($P=0.91$), suggesting longer cultivation time to achieve a significant decrease in weight of leaf and therefore treatment effectiveness. Additionally, there was no difference between the treatments for the number of conidia of LFP 37 ($P=0.44$). On the other hand, *P. macrosporus* consistently reduced LFP 37 conidia viability in detached leaves and in coffee plants. For the *in vivo* assay, not only the conidia of the saprobe fungus but also all of the fungal-derived products or enzymes reduced the germination rate by 40% ($P=0.00003$ and 0.00) (Figure 6).

Since there was no significant difference between these treatments, either one could be used as a control, which gives more versatility to the use of this fungus as a future biological product. In addition, other authors have reported the use of *P. macrosporus* as a biocontrol agent in coffee diseases. Botrel et al. (2018) observed a reduction in the severity of coffee halo blight of up to 72% when the seedlings were previously treated with the fungus *P. macrosporus*. Rodriguez et al. (2016) observed similar results in the control of *C. gloeosporioides* in coffee seedlings treated with *P. macrosporus*.

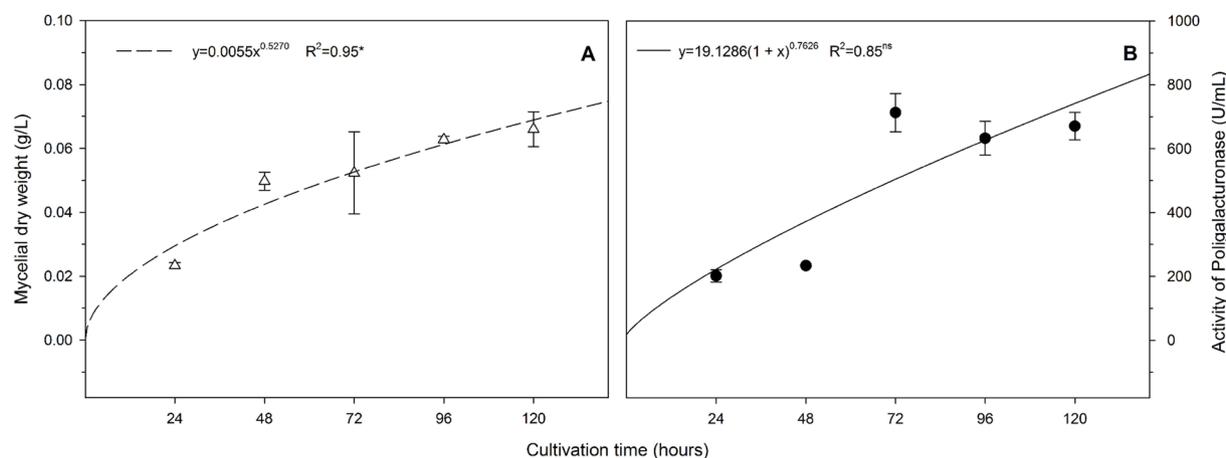


FIGURE 5 - Dry mycelial weight (A) and polygalacturonase activity in the culture supernatant (B) of *Phialomyces macrosporus* at 24, 48, 72, 96, and 120 h of cultivation in liquid medium with pectin. The line represents \pm SE. *Angular coefficients differ significantly by *F*-test.

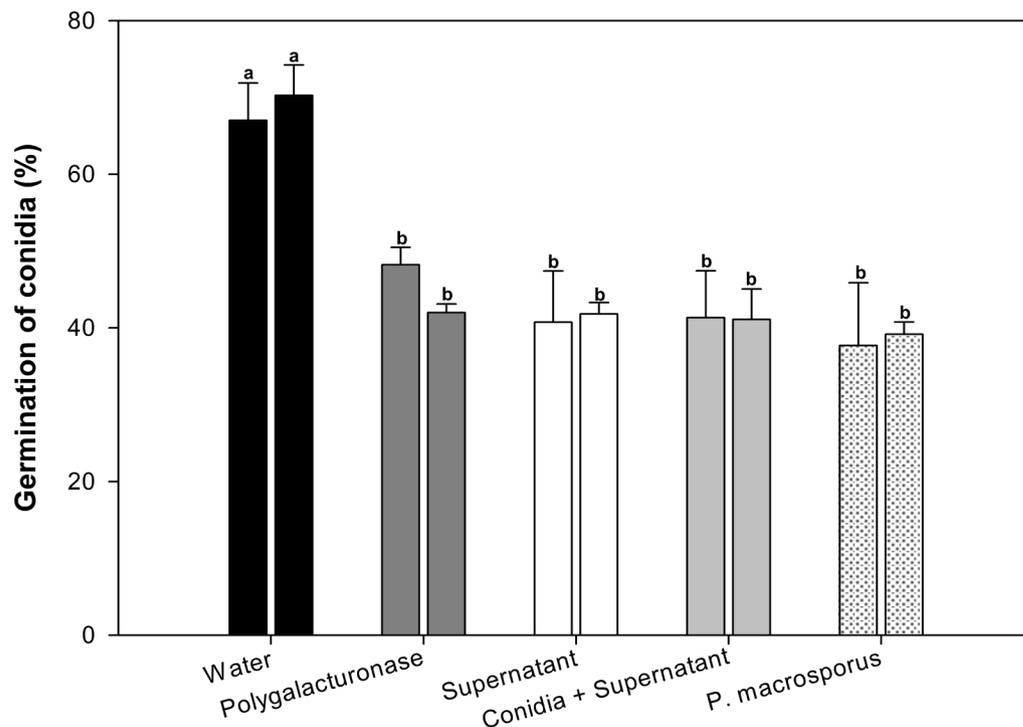


FIGURE 6 - Germination of *Cercospora coffeicola* conidia after treatment with water, polygalacturonase, supernatant alone, conidia+supernatant, *Phialomyces macrosporus* conidia (10^7 conidia mL⁻¹). The graph represents two assays conducted in different periods. The bars with the same letter are similar at the 5% level according to Scott-Knott's multiple range test. CV=14.48% and 12.01%, respectively. The line on each bar represents \pm SE.

4 CONCLUSIONS

P. macrosporus inhibited *C. coffeicola* growth, sporulation and viability. *P. macrosporus* produced antimicrobial substances and is a promising biological agent for the integrated management of brown eye spot.

5 ACKNOWLEDGEMENTS

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