

Effect of biochar and inoculation with *Trichoderma aureoviride* on melon growth and sandy Entisol quality

Erika Valente de Medeiros^{1*}, Marcele de Cássia Henriques dos Santos Moraes¹, Diogo Paes da Costa¹, Gustavo Pereira Duda¹, Julyana Braga de Oliveira¹, Jenifer Sthephanie Araujo da Silva¹, José Romualdo de Sousa Lima¹, Claude Hammecker²

¹Laboratory of Microbiology, Federal Rural University of Pernambuco, Garanhuns, PE, Brazil

²IRD/UMR Eco & Sol, place Pierre Viala, 2, 34060 Montpellier, France

*Corresponding author: evmbio@gmail.com; marcele-moraes@hotmail.com

Abstract

The melon belongs to the family of commercially important cucurbitaceous in the world. However, the production of this crop can be very problematic in some places due to management practices and the climatic instability. Amongst the different options available to overcome these obstacles, the use of biochar often promoted for providing multiple benefits to crops, could contribute in holding more water and nutrients in soil and therefore improve the plant growth. A second way to try to improve the plant development was to use *Trichoderma* (TRI) known as aiding in seed germination, and being an excellent biological control agent against plant pathogenic pests. So, the objective of this study was to evaluate the benefits of the association of biochar and TRI on the initial growth of melon and the effects on the quality of a sandy Entisol. We quantified the effects of these associations through biometric growth in melon plants and chemical, microbial, and enzymatic activities of the biogeochemical cycles in the soil. An experiment in a completely of randomized design was performed in a factorial scheme (3 x 2 + 1) with three sources of biochar (bean husk (BH), coffee ground (CG), and coffee husk (CH)) inoculated with (T+) or without (T-) TRI and additional controls. When the coffee grounds (CG) and bean husks (BH) biochar with T+ soil was inoculated, the fresh weight (number of leaves), dry weight, length (of roots and branch), soil acid and alkaline phosphatase, total organic carbon, phosphorus, magnesium, potassium, and pH were all increased. Moreover, *T. aureoviride* inoculated CG biochar compared to the control increased the shoot length and dry biomass of the melon plant in 30 and 22% between 22 and 30 %. The soil that received coffee husks (CH) biochar and T+ showed higher microbial biomass carbon. However, the melon plants responded more to the type of biochar than to the *T. aureoviride* inoculation, possibly due to the short growth time of melon. Results of BH biochar inoculated with *T. aureoviride* in sandy soil showed improved efficiency on melon growth and increased soil quality.

Keywords: Biocarbon; Cycling of nutrients; Plant growth; Route of inoculation; Soil fertility.

Abbreviations : CG_coffee grounds; CH_coffee husks; BH_bean husks; +T_with inoculation of *Trichoderma aureoviride*; -T_without inoculation of *Trichoderma aureoviride*; RDB_root dry biomass; RFB_root fresh biomass; SDB_shoot dry biomass; SFB_shoot fresh biomass; SLEN_shoot length; RLEN_root length; DIA_diameter; NL_number of leaves; ChloA_Chlorophyll a; ChloB_Chlorophyll b; Ure_soil Urease; Bet_β-glycosidase; P.Alc_alkaline phosphatase; P.Aci_acid phosphatase; TOC_total organic carbon; MBC_microbial biomass carbon; SBR_soil basal respiration; qCO2_soil metabolic quotient.

Introduction

Melon (*Cucumis melo* L.) is one of the most important species of the *Cucurbitaceae* family, which is cultivated worldwide, because its pulp is refreshing with a high nutritional value and pleasant aroma, besides having an extensive range of medicinal properties (Mallek-Ayadi et al., 2018). Melon production is carried out in semi-arid regions with excessive use of agricultural inputs, especially synthetic fertilizers and water for irrigation (Deus et al., 2015). The high costs of fertilizers have triggered the development of alternative products able to fulfil partial, or total nutrient demand of the crop.

In this sense, studies have shown that biochar was an alternative agricultural input for crops, being a low-cost, nutrient-concentrated product obtained through the pyrolysis of different biomasses, including agro-industrial

residues, under limited oxygen (O₂) conditions (Lima et al., 2018; Yang et al., 2018). Biochar application to soil mitigates the emission of gases, increases C sequestration, and improves the structure, fertility, activity of beneficial microorganisms, soil water retention, and agricultural productivity (Weber and Quicker, 2018). Biochar is a soil conditioner that can promote plant growth by improving retention of nutrients and soil fertility but also soil physical attributes, like density and soil water retention (Paneque et al. 2016). It can also modify the composition and abundance of the soil biological community. These changes may also alter nutrient cycling (Anyanwu et al., 2018).

Biochar was successfully used as an inoculation vehicle of microorganisms beneficial to the soil and plants (Medeiros et al. 2019). In this sense, *Trichoderma* spp. is essential for

its versatility in promoting plant development and controlling soil pathogens (Swain et al., 2018). These fungi are mycoparasites secreting secondary metabolites, phytohormones, and enzymes known for assisting root growth and controlling viruses, pathogenic bacteria, and insect pests (Mendoza-Mendoza et al., 2018).

This study raised the hypothesis that biochar could be a potential vehicle for inoculation of *Trichoderma* spp., improving several soil attributes that increase the initial development of melon. As little information about this subject was available, the objective of the present study was to evaluate the effects of biochar from different sources with and without *T. aureoviride* applied to sandy Entisol on the initial melon plant growth and to assess the changes in chemical, microbiological, and enzyme activities regarding the soil attributes.

Results

Beneficial effects of the inoculation of *T. aureoviride* via biochar on initial melon growth and soil quality

Non-metric multidimensional scaling (nMDS) showed that the variations in melon plants responded more to the type of applied biochar (higher separation between points) than to the presence or absence of *T. aureoviride* (closest; Fig. 1). Each type of biochar generated a distinct soil biochemical pattern, primarily due to changes in pH, β -glucosidase, and urease activities and the Mg^{2+} , K^+ and P levels in the soil, which was the most sensitive variables in the applied treatments.

However, the inoculation of *T. aureoviride* revealed significant increases of plant development with responses being modulated as a function of the biochar type. For example the treatment with inoculated coffee husks biochar (CH + T) was different from all other treatments (Fig. 1). This treatment showed a high increase in soil pH (~ 7.0), Ca^{2+} , K^+ , and MBC levels, being significantly higher than in other biochar sources (Table 1).

Analysis of variance (ANOVA) and comparisons between averages

Coffee ground biochar with *T. aureoviride* (CG + T) applied to the sandy soil cultivated with melon showed an increase in β -glucosidase activity, while for the others the β -glucosidase activity was reduced compared to the control (Table 1). The most significant improvements with this treatment were for RDB, RFB, SDB, and SFB, showing a higher potential for promoting plant development. Bean husks with *T. aureoviride* (BH + T) applied to the soil showed a significant increase in RLEN (Table 1).

Interactions between analyzed variables

Some soil metabolic and chemical variables showed significant correlations with the biometric variables of melon plants. Among these, β -glucosidase activity showed a positive correlations with the diameter of the main branch and RFB, SFB, and, SDB (Fig. 2A). Urease activity was positively correlated with the shoot fresh matter and showed negative correlations with the shoot length and with the content of Chlorophyll b.

The melon shoots were longer in soil with higher P and Mg^{2+} levels, as illustrated by the positive correlations between shoot length versus P and Mg concentration in soil (Fig. 2B).

We found other positive and significant relationships between the contents of H^+ + Al^{3+} with diameter, root length, root fresh matter, and shoot fresh and dry matter.

Regarding the interactions between soil metabolic and chemical attributes (Fig. 2C), some strong positive and significant relationships were observed, highlighting correlations: (i) between total organic carbon and soil Mg^{2+} and P contents; (ii) between urease activity and Ca^{2+} ; (iii) between acid phosphatase activity and soil P content.

Discussion

Application of biochar to the soil provides numerous improvements, such as increasing fertility, promoting aggregates formation, and increasing carbon stock (Yang et al., 2018). In the present study, each type of biochar showed different chemical composition, microbial activity, and enzymatic expression (Fig. 1 and Table 1).

Soil pH was one of the most affected parameter by biochar addition especially when soil had been amended with CH. This was mainly attributed to negatively charged chemical groups on the biochar surfaces (Lima et al., 2018), such as carboxylic, phenolic, and hydroxylic radicals, as well as silicates, carbonates, and bicarbonates that bind to H^+ ions, making the pH of the soil more alkaline (Gul et al., 2015).

One of the main advantages of increasing pH in tropical soils is the increase in P availability for plants, mainly as mono- and di-calcium phosphate, which are most soluble for an optimum pH ranging between 6.0 and 7.0 (Buss et al., 2018). In this case, the effect of adequate soil pH for bean husk biochar amendments, probably contributed to the higher levels of P for this biochar (Table 1). These results are analogous to those of Bornø et al. (2018) demonstrating that the type of biochar influenced the adsorption and bioavailability of P when added to Ca^{2+} , Mg^{2+} , and K^+ oxides rich soil.

The simultaneous increase of pH and carbonate availability with biochar amendment favored the formation of organo-mineral complexes that stimulated microbial activity and promoted soil organic carbon stability (Sheng and Zhu, 2018). Results obtained in the present study corroborated these findings since the soils with CH + T and a pH of 7.0 showed a significant increase of the MBC. These results also revealed the active participation of *T. aureoviride* in this treatment, as the optimal conditions where reached for this fungus to increase the MBC of the soil significantly.

The soil amended with CG and *T. aureoviride* (CG + T) displayed active participation of *T. aureoviride* in carbon cycling as this treatment has significantly increased the activity of β -glycosidases, enzymes produced by microorganisms degrading cellulose and other carbohydrates present in the cell walls (Strahsburger et al., 2017). Similarly, Wu et al. (2018) demonstrated that the addition of biochar substantially increased the activity of β -glycosidase, but also cellobiohydrolase, and chitinases in sandy soils, especially in an acidic pH. Different soils showed similar effects. For example, Teutscherova et al. (2018), in the various Mediterranean regions, verified that acidic soils treated with biochar showed higher responses through increased activities of dehydrogenase, urease, pH, MBC, and SBR during the initial six weeks of incubation.

The increase of RFM and SFM of melon plants, promoted by the association between CG and *T. aureoviride* obtained in the present study, showed that the benefits of this fungus for the plants was strictly related to the use of a suitable substrate. Galletti et al. (2015) demonstrated that the

inoculation of *Trichoderma harzianum* in melon seeds promoted the increase of microbial biomass and nutrient cycling in the soil, resulting in a significant increase in dry

matter and the root length of melon plants. Root elongation directly reflects the overall development of the entire plant (Prasad et al., 2018). However, in the study of Galletti et al.,

Table 1. Enzymatic/metabolic attributes of carbon (A), soil chemicals (B), and biometrics of plants (C) treated with different biochars from coffee grounds (CG), coffee husks (CH), and bean husks (BH) with (+ T) and without (-T) inoculation of *Trichoderma aureoviride* under cultivation of melon plants of the variety *BRS Araguaia*.

| A. Metabolic Attributes | | | | | | B. Chemical Attributes | | | | | | C. Biometric Attributes | | | | | | | | | | | |
|--|-------|----|-------|----|-------|--|------|----|------|----|------|--|------|----|------|----|------|----|------|-----|------|---|--|
| CG | | CH | | BH | | CG | | CH | | BH | | CG | | CH | | BH | | | | | | | |
| a. Urease ($\mu\text{g NH}_4\text{-N g}^{-1}\text{dwt } 2\text{h}^{-1}$) | | | | | | a. pH | | | | | | a. Number of leaves | | | | | | | | | | | |
| +T | 0.036 | | 0.040 | A | 0.042 | | 5.64 | cA | 7.07 | ab | 6.85 | b | 7.3 | a | 4.2 | bb | 5.7 | ab | | | | | |
| -T | 0.035 | b | 0.022 | bb | 0.056 | a | 5.53 | cB | 7.31 | aA | 6.87 | b | 6.7 | | 6.3 | A | 6.0 | | | | | | |
| C | 0.029 | | | | | | 5.01 | | | | | | 7.3 | | | | | | | | | | |
| b. β -glucosidase ($\mu\text{g p-Nitrof. g}^{-1}\text{soil h}^{-1}$) | | | | | | b. P (mg kg^{-1}) | | | | | | b. Diameter of branch (cm) | | | | | | | | | | | |
| +T | 47.59 | aA | 31.29 | bb | 25.90 | cB | 0.79 | cB | 1.87 | bA | 2.47 | ab | 6.5 | a | 3.9 | bb | 6.1 | a | | | | | |
| -T | 41.00 | aB | 35.70 | ba | 31.86 | ba | 1.25 | cA | 1.75 | bb | 2.68 | aA | 6.2 | | 6.0 | B | 5.9 | | | | | | |
| C | 41.53 | | | | | | 0.53 | | | | | | 6.6 | | | | | | | | | | |
| c. Acidic Ph. ($\mu\text{g p-Nitrof. g}^{-1}\text{soil h}^{-1}$) | | | | | | c. Ca^{2+} ($\text{cmol}_c\text{dm}^{-3}$) | | | | | | c. Length of branches (cm) | | | | | | | | | | | |
| +T | 3.33 | a | 1.37 | b | 1.90 | b | 0.29 | c | 0.48 | a | 0.37 | b | 18.5 | bb | 10.4 | cB | 20.8 | a | | | | | |
| -T | 3.01 | a | 2.80 | b | 2.27 | b | 0.29 | c | 0.49 | a | 0.42 | b | 23.1 | aA | 17.2 | ba | 22.2 | a | | | | | |
| C | 2.53 | | | | | | 0.46 | | | | | | 16.2 | | | | | | | | | | |
| d. Alkaline Ph. ($\mu\text{g p-Nitrof. g}^{-1}\text{soil h}^{-1}$) | | | | | | d. Mg^{2+} ($\text{cmol}_c\text{dm}^{-3}$) | | | | | | d. Length of root (cm) | | | | | | | | | | | |
| +T | 0.72 | a | 0.65 | b | 0.66 | ab | 0.26 | c | 0.39 | b | 0.56 | a | 14.7 | ab | 7.2 | bb | 27.0 | aA | | | | | |
| -T | 0.68 | ab | 0.63 | b | 0.69 | a | 0.26 | c | 0.40 | b | 0.57 | a | 10.9 | | 25.7 | A | 13.2 | B | | | | | |
| C | 0.64 | | | | | | 0.25 | | | | | | 16.5 | | | | | | | | | | |
| e. TOC ($\text{g kg}^{-1}\text{ soil}$) | | | | | | e. K^+ ($\text{cmol}_c\text{kg}^{-1}$) | | | | | | e. FW. of branches (g) | | | | | | | | | | | |
| +T | 5.24 | B | 4.49 | | 4.93 | B | 0.55 | c | 1.89 | aA | 1.47 | b | 26.0 | a | 16.2 | b | 17.8 | | | | | | |
| -T | 6.10 | aA | 4.01 | b | 6.08 | aA | 0.54 | c | 1.78 | ab | 1.47 | b | 24.9 | a | 16.1 | b | 16.9 | | | | | | |
| C | 5.19 | | | | | | 0.24 | | | | | | 21.3 | | | | | | | | | | |
| f. MBC ($\text{g kg}^{-1}\text{ soil}$) | | | | | | f. Na^{2+} ($\text{cmol}_c\text{kg}^{-1}$) | | | | | | f. FW. of roots (g) | | | | | | | | | | | |
| +T | 0.40 | a | 0.61 | aA | 0.17 | b | 3.05 | a | 3.34 | a | 3.05 | a | 14.7 | a | 5.4 | b | 2.4 | b | | | | | |
| -T | 0.25 | | 0.24 | B | 0.22 | | 3.16 | c | 4.16 | a | 3.60 | b | 13.7 | a | 6.6 | b | 7.6 | b | | | | | |
| C | 0.23 | | | | | | 3.05 | | | | | | 8.0 | | | | | | | | | | |
| g. $\text{qCO}_2(\text{mgC-CO}_2\text{g}^{-1}\text{BMS-C h}^{-1})$ | | | | | | g. $\text{H}^+ + \text{Al}^{3+}$ ($\text{cmol}_c\text{dm}^{-3}$) | | | | | | g. DW. of branches (g) | | | | | | | | | | | |
| +T | 3.35 | | 0.74 | | 11.19 | | 1.62 | a | 0.56 | ba | 0.47 | b | 3.47 | a | 0.63 | cB | 1.68 | b | | | | | |
| -T | 4.53 | | 5.06 | | 6.09 | | 1.57 | a | 0.08 | cB | 0.58 | b | 3.18 | a | 2.00 | ba | 1.94 | b | | | | | |
| C | 12.96 | | | | | | 1.24 | | | | | | 2.71 | | | | | | | | | | |
| | | | | | | | | | | | | h. DW. of roots (g) | | | | | | | | | | | |
| | | | | | | | | | | | | 0.98 | | | | | | a | 0.73 | a | 0.17 | b | |
| | | | | | | | | | | | | 0.98 | | | | | | a | 0.87 | a | 0.60 | b | |
| | | | | | | | | | | | | 1.12 | | | | | | | | | | | |
| | | | | | | | | | | | | i. Chlorofile A ($\mu\text{g g}^{-1}$) | | | | | | | | | | | |
| | | | | | | | | | | | | 34.8 | | | | | | aB | 32.9 | abB | 32.7 | b | |
| | | | | | | | | | | | | 38.1 | | | | | | aA | 36.7 | aA | 32.3 | b | |
| | | | | | | | | | | | | 35.4 | | | | | | | | | | | |
| | | | | | | | | | | | | j. Chlorofile B ($\mu\text{g g}^{-1}$) | | | | | | | | | | | |
| | | | | | | | | | | | | 9.4 | | | | | | B | 8.6 | | 9.4 | | |
| | | | | | | | | | | | | 10.9 | | | | | | aA | 9.6 | ab | 8.3 | b | |
| | | | | | | | | | | | | 10.0 | | | | | | | | | | | |

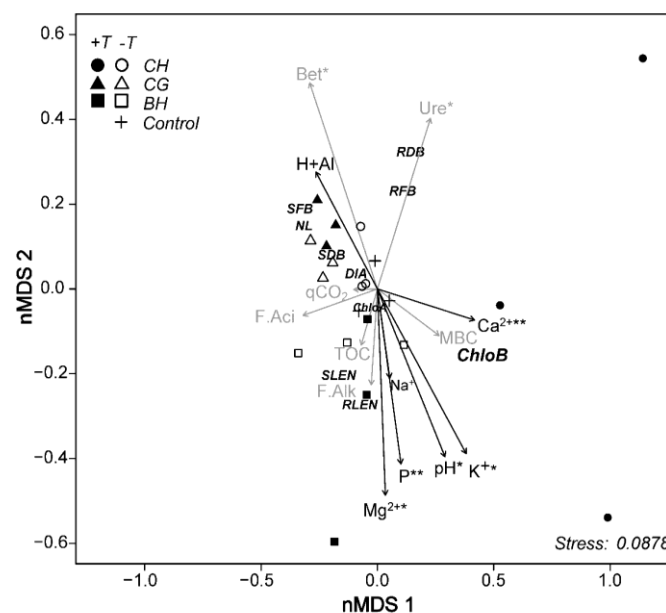


Fig 1. Dissimilarities between the metabolic activities of microorganisms in soils cultivated with melon enriched with biochar from bean husks (BH), coffee grounds (CG), and coffee husks (CH) with (+T) and without (-T) *T. aureoviride*. Non-metric multidimensional scaling (nMDS) revealed a reasonable fit of the explanatory model, according to the stress measure ($S_m = .087$, $K(\text{dimension}) = 3$) suggested by Kruskal (1964). The vectors corresponding to the chemical (black) and metabolic (in gray) attributes followed by asterisks (*) and (**) have significant correlations ($p \leq .05$ and $.10$) with the ordering configuration about the axes. Root dry biomass (RDB), root fresh biomass (RFB), shoot dry biomass (SDB), shoot fresh biomass (SFB), shoot length (SLEN), root length (RLEN), diameter (DIA), number of leaves (NL), Chlorophyll a (ChloA), Chlorophyll b (ChloB) of melon plants; soil Urease (Ure), β -glucosidase (Bet), alkaline phosphatase (P.Alc), acid phosphatase (P.Aci) activities, total organic carbon (TOC), microbial biomass carbon (MBC), soil basal respiration (SBR), soil metabolic quotient (qCO_2).

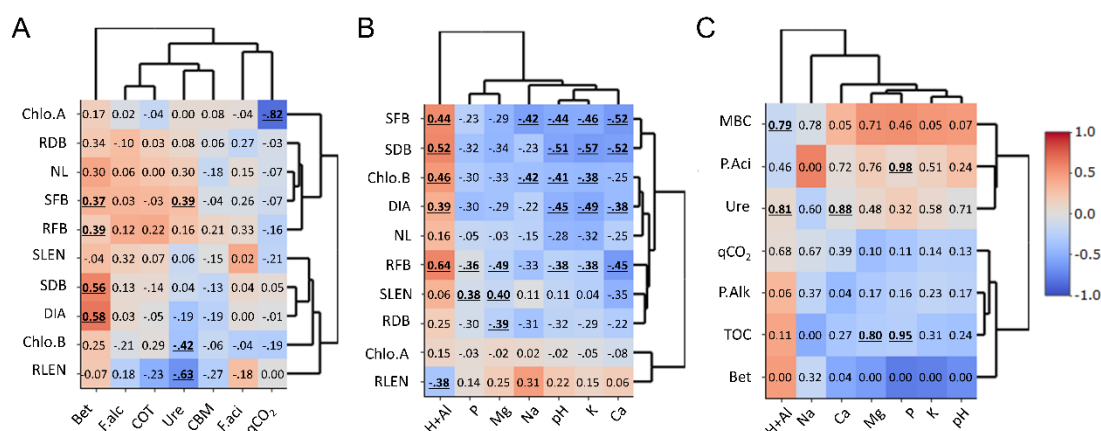


Fig 2. Correlations between biometric data of melon plants, soil enzymatic activities, and chemical attributes on soils with different treatments of biochar with and without *Trichoderma aureoviride*. The heatmap and Pearson correlation coefficients (R_p) station were used to associate the paired samples. The underlined correlations are significant ($p \leq .05$). In general, the Mantel test revealed significant positive correlations between plant biometric and soil metabolic attributes (A, $r = .39$, $p = .001$), biometric and chemical attributes (B, $r = .16$, $p = .023$), and metabolic and chemical (C, $r = .26$, $p = .001$). Root dry biomass (RDB), root fresh biomass (RFB), shoot dry biomass (SDB), shoot fresh biomass (SFB), shoot length (SLEN), root length (RLEN), diameter (DIA), number of leaves (NL), Chlorophyll a (ChloA), Chlorophyll b (ChloB) of melon plants; soil Urease (Ure), β -glucosidase (Bet), alkaline phosphatase (P.Alc), acid phosphatase (P.Aci) activities, total organic carbon (TOC), microbial biomass carbon (MBC), soil basal respiration (SBR), soil metabolic quotient (qCO_2).

(2015), *Trichoderma* did not affect significantly the activities of β -glucosidase, unlike in the present study. In this context, enzymatic activity of β -glucosidase correlated strongly with the main biometric attributes of the melon: DIAM, RFB, SFB, and SDB (Fig. 2A). It showed a strong relation between these variables and the CG biochar with *T. aureoviride*. This treatment provided general

improvements in the soil since β -glucosidase, urease and phosphatase soil activities are efficient indicators of soil quality (Adetunji et al., 2017). In these conditions, it is also worth mentioning the significant positive correlations between urease and the SFM plants (Fig. 2B) and between phosphatase and labile P in the soil (Fig. 2C).

Among the main functions of β -glucosidase and soil urease, the equilibrium of the C: N ratio was also representative of the metabolic activity. In a previous study using the same substrates, they acquired the C: N ratio of 32.73 for the biochar of CH and 16 for the biochar of CG (Lima et al., 2018). Song et al. (2018) studied the metabolic activity of soils treated with maize straw biochar and found that C: N ratio, K^+ and MBC content were the dominant factors affecting soil microbial community. These authors also found that biochar amendements with formulated NPK fertilizer, as adopted in the present study, favoured fast-growing bacteria (r-strategists) on fungi. In this case, N mineralization of organic matter occurred to compensate for the high C: N ratios after the addition of biochar.

In this context, as different types biochar contribute distinctly towards the metabolic activity of soils, the addition of alternative mineral sources may also influence the behavior of soil microorganisms differently. This result is due to the different nutritional needs of each microbial taxon, and this difference is even higher among fungi and bacteria, for which the optimal C: N ratios are 8.3 and 4.4, respectively (Mouginot et al., 2014).

Moreover, biochar does not only serve as an inoculation pathway for *Trichoderma* spp. but also serves as a substrate for inoculation of different microorganisms. For example, in the study conducted by Tripti et al. (2017), tomato plants inoculated with strains of *Bacillus* sp. and *Burkholderia* sp. via biochar showed significant increases of dry and fresh biomass, plant length, and the number of flowers. An increase of dehydrogenase activity and soil fertility was also recorded, leading to a significative tomato productivity increase and therefore replacing high-cost mineral fertilizers. These data reinforce the need to better explore biochar as a new alternative for managing green manure fertilization especially considering the rising prices and the scarcity of mineral fertilizers.

Positive and significant correlation of soil chemical attributes with the biometric variables, highlight the positive relations between the shoot length of melon plants with the contents of P and Mg^{2+} (Fig. 2B) as well as those with TOC (Fig. 2C). These comparisons were relevant, as they pointed to the primary factors associated with the elongation of the main branch of the melon, especially in treatments with CG and BH. In the case of P, acidic soil conditions could justify its low availability for plants. Moreover as biochar contributed to increase P content along with pH. It could be seen as a natural source of this element, improving plant efficiency in the use of organic and inorganic P fertilizers (Muhammad et al., 2017). At the same time biochar also contributed to increase the availability of Mg^{2+} , Ca^{2+} , and K^+ with a raising pH (Bornø et al., 2018), characterizing the first relation between these attributes (Fig. 2C).

The results of this study highlighted the importance of this research which points out promising alternatives management to improve plant development and soil chemical, microbiological, and enzymatic attributes. In the present study, *T. aureoviride* isolates demonstrated high versatility both in plant development and in the promotion of enzymatic activity, thus indicating that this fungus becomes a strong candidate for alternative organic input and application in the field using biochar as a route of inoculation.

Materials and Methods

Plant materials

For this experiment we used the melon hybrid 'BRS Araguaia' that is well appreciated on local market and suitable to exportation for its excellent quality. Three seeds were deposited in each cell and after seven days thinning took place, and only the most vigorous plant was kept and grown at an average room temperature of 28 °C. Trays received daily irrigation, keeping moisture in the range of soil field capacity.

Experiment

The agronomical experimentation consisted of testing the effect of three types of biochar (BH, CG, and CH) with or without inoculation of *T. aureoviride* URM 5158. We used industrial bean husk residues (BH), coffee grounds (CG), and coffee husks (CH) to produce different types of biochar. This biomass was charred for 12 h under limited oxygen conditions in a slow pyrolysis process in which the temperature reached 530°C in a metallic kiln, and they were sieved using a 2-mm sieve. The chemical analysis and specific surface area (SSA) measurements of the different types of biochar can be found in (Lima et al., 2018)

The choice of *T. aureoviride* URM 5158 fungus was based on its physiological and functional versatility mentioned in previous studies (Silva et al., 2016) and was obtained from the Micoteca URM Collection of the Mycology Department of the Biological Sciences Center of the Federal University of Pernambuco (<https://www.ufpe.br/micoteca/>). The fungus conidia was reactivated through three successive sub-cultures in an Erlenmeyer flask containing 50 ml of potato dextrose agar (PDA) liquid medium, grown at $26 \pm 2^\circ\text{C}$ for eight days. Each treatment with *T. aureoviride* was sprayed with 100 ml (1×10^6 conidia per ml) applied to each type of biochar. The different sources of biochar and *T. aureoviride* were mixed and homogenized to the soil before planting.

The experiment was conducted at the Federal Rural University of Pernambuco (UFRPE) in Garanhuns in Brazil. The soil used was topsoil (0–20 cm layer) collected in a patch of natural forest area in São João, Brazil ($08^\circ 48' 34.2'' \text{ S}$, $36^\circ 24' 29.3'' \text{ W}$) at an elevation of 705 m AMSL. The soil is a typical Entisol with almost 90% sand and less than 5% clay and with low cation exchange capacity (CEC) and SSA. The chemical attributes of soil and biochar before the experiment are shown in Lima et al. (2018). This soil was previously treated with formulated fertilizer NPK (6:24:12) to prevent variations related to plant nutritional deficits.

Biochar was added at a rate of 32 t ha^{-1} in the first 20 cm of soil. The experimental design was completely randomized in a double factorial scheme with additional treatment ($3 \times 2 + 1$). The first factor represented the three types of biochar (CG, CH, and BH), while the second considered the inoculation (T+) and absence (T-) of *T. aureoviride* on these substrates. The additional treatment represented absolute control, without biochar nor *T. aureoviride*.

Then, 100 mL of the *T. aureoviride* suspension ($1.46 \text{ conidia mL}^{-1}$) with different types of biochar was added to the soil, followed by the planting of three seeds of the hybrid BRS Araguaia melon variety in each pot. After seven days, only

the most vigorous plant was kept and grown at an average ambient temperature of 28°C under daily manual irrigation, maintaining moisture at the range of soil field capacity.

Biometric analyses of melon plants and soil samples

Melon plants were removed at the end of the experiment (45 days after emergency) to measure the number of leaves (*NL*), diameter of the plant base (*DIAM*), root dry biomass (*RDB*), root fresh biomass (*RFB*), shoot dry biomass (*SDB*), shoot fresh biomass (*SFB*), shoot length (*SLEN*), root length (*RLEN*), Chlorophyll a (*ChloA*), and Chlorophyll b (*ChloB*).

Soil chemical attributes

After removal of the plants from the pots, the soil was collected to measure pH in water (1:2.5) and the content of P, Na⁺, K⁺, Ca²⁺, Mg²⁺, Al³⁺, and H+Al³⁺, according to the methodology of Silva (2009). The inorganic labile P, Na⁺, and K⁺ were extracted using Mehlich's solution 1 (H₂SO₄ .0125 mol L⁻¹ + HCl .05 mol L⁻¹). The P quantified by spectrophotometer colorimetry, and the Na⁺ and K⁺ determined by flame photometry. Ca²⁺, Mg²⁺, and Al³⁺ were extracted with KCl 1.0 mol L⁻¹ solution, where the first two were determined by atomic absorption spectrometry (Raij et al., 2001). Potential acidity (H+Al³⁺) was determined using calcium acetate (Ca(CH₃COO)₂ H₂O) at pH 7.0. Also, Al³⁺ was determined by titration with NaOH 0.025 mol L⁻¹, using bromothymol blue as an indicator. For all extractions, 5.0 g of soil for 50 mL of extractive solution (relation 1:10) was used.

Total organic carbon content (TOC) was obtained using the C oxidation method in the wet form with potassium dichromate followed by determination of remaining Cr₂O₇²⁻ by titration with ammonium iron (II) sulfate (Yeomans and Bremner 1988).

Soil microbial attributes

The microbial biomass carbon (MBC) was determined by irradiation method (Mendonça and Matos 2005), with the extraction carried out by the addition of 80 mL of K₂SO₄ 0.5 M to each 20 g of soil (Tate et al., 1988). The C content was determined by colorimetry (Bartlett and Ross, 1988).

The evolution of CO₂ released by microbial respiration was measured for calculation of soil basal respiration and soil metabolic quotient (*q*CO₂) by the alkaline adsorption method. For this purpose, at the end of the experiment, the soil was maintained at 60% of field capacity based on gravimetry (Anderson and Domsch, 1985). Then, 30 g of each sample were hermetically sealed in containers to capture CO₂ with 10 mL of NaOH .5 M solution. After 72 h, 2 mL of BaCl 10% (m/v) solution was added for total precipitation of CO₂, followed by two drops of phenolphthalein diluted in 100 mL of ethyl alcohol 95% (v/v) the determinations being carried out by titration with HCl 0.25 M.

Soil enzymatic activities

We determined the enzymatic soil activities by estimating the gene expression of the main processes related to acid (P.Aci) and alkaline phosphatases (P.Alk) (E.C. 3.1.3), β-glucosidase (Beta) (E.C. 3.2.1.21), and the urease (Ure) (E.C. 3.5.1.5) according to the colorimetric analysis of the products released by the samples subjected to incubation in

an adequate substrate (Sigma-Aldrich). The activities of P.Aci and P.Alk were estimated according to the methodology of Eivazi and Tabatabai (1977). Beta was estimated by the methodology of Eivazi and Tabatabai (1988), and Ure was estimated using the method by Kandeler and Gerber (1988). The absorbance of the products was measured by spectrophotometer (Libra S22, Biochrom, Cambridge, UK).

Statistical analyses

Statistical and multivariate analyses were conducted using the R language platform (v.3.4.3, 2017) based on the data of the biometric attributes of the plants, chemicals, and enzymatic activities of the soils. Multivariate exploratory analyzes, including non-metric multidimensional scaling (nMDS), analysis of similarity (ANOSIM), heatmaps, and Mantel tests were based on the tools of the *vegan* and *heatmaply* libraries. After adjusting the models and removing the outliers, the means, standard deviations, and variation coefficients were calculated using the *doBy* library. Homogeneity tests, analysis of variances (ANOVA), and comparisons between averages were done with the balanced data, according to the tools contained in the *stats*, *AxpDes*, *multicomp*, and *agricolae* libraries.

Conclusions

Biochar from coffee and bean residues inoculated with *T. aureoviride* and applied to sandy soil are efficient in the initial growth of melon plants in increasing the chemical, microbial, and enzymatic activities of the soil. However, melon plants responded more to the source of biochar than to the *T. aureoviride* inoculation, possibly due to the short growth time of melon. Therefore, it is necessary to perform further studies with more time and more cycles of the crop to verify whether the biochar inoculated with *Trichoderma* and applied to the soil can promote higher plant production.

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