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Proceedings Isolation and characterization of agricultural soil bacteria with biotechnological and biological control potential applications⁺

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Abstract: Unsustainable agricultural practices eventually have an impact on soil conditions and the 12 microbiological diversity. To regain balance, ecologically sound strategies can be an alternative. In 13 this work, a collection of bacteria was isolated from agricultural soil and characterized to evaluate 14 their capacity for phosphorus and iron biofertilization, exoenzyme production, and biocontrol of 15 several phytopathogenic fungi. Bacterial identification pointed out to a majority of Bacillus spp. 16 along with other several minority genera. Isolates globally displayed a high proportion of the bio-17 logical activities tested, especially concerning production of hydrolytic enzymes. Inhibition on fun-18 gal growth was variable among the soil bacterial isolates by production of diffusible compounds 19 and/or VOCs (volatile organic compounds). Evidence from this work provides promise for the ap-20 plication of soil bacteria to improve agricultural soil management and crop production. 21

Keywords: agrosystem; agrochemical; biological activity; sustainability; biofertilization; exoenzyme22production; plant pathogen; biocontrol; diffusible compounds; volatile organic compounds23

1. Introduction

Agricultural activities have a direct influence in a range of fundamental areas such 26 as the environment, the public health, and the global economy [1]. Since many years, ben-27 efits have been achieved through unsustainable agricultural practices, among which the 28 indiscriminate use of chemical compounds as fertilizers, pesticides and/or herbicides in 29 the field, with the aim to increase crop yield and face an increasing demand for worldwide 30 production. However, these conventional practices eventually have a significant impact, 31 mainly on soil conditions and food quality [2]. The continued application of agrochemi-32 cals causes pollution in the environment. Their degradation produces chemical residues, 33 which can remain in the field for a long time [1,2]. Treatments with fertilizers derived 34 from nitrogen result in increased amount of nitrates in the field and disturbance of the 35 natural nitrogen fixation process [3]. Treatments with pesticides and/or herbicides for pest 36 chemical control favour the appearance of resistances to these products, as well as new 37 opportunistic pests due to removal of competitors. Moreover, the global climate change 38 has an additional effect, since it alters the distribution of crops, weeds, pests and diseases, 39 and then, agrochemical use [1]. 40

To regain balance and improve the efficient use of the natural resources, ecologically sound strategies as those microorganism-based can be an alternative. The beneficial interactions and activity of plant and soil microorganisms can be considered to improve agricultural development, crop production and environmental sustainability [4].

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Copyright: © 2023 by the authors. Submitted for possible open access publication under the terms and conditions of the Creative Commons Attribution (CC BY) license (https://creativecommons.org/license s/by/4.0/). In this work, a collection of bacteria was isolated from agricultural soil and characterized to evaluate their properties for biotechnological and/or biocontrol applications, mainly capacity for phosphorus, nitrogen and iron biofertilization, exoenzyme production, and biological control of several species of phytopathogenic fungi.

2. Materials and Methods

2.1. Bacterial isolation

Agricultural soil was mixed with phosphate buffered saline (PBS) buffer [pH 7.4; 137 7 mM NaCl, 10 mM Na₂HPO₄, 2.7 mM KCl, 1.9 mM KH₂PO₄] at 1:10 (w/v), shaken at 200 8 rpm for 30 min to isolate bacteria by serial ten-fold dilutions, and plated onto Nutrient 9 Agar (NA). Isolates were purified, PCR-identified by 27F/1492R and 27F/1525R primer 10 sets [5], and cryopreserved. To perform the tests, they were grown on NA for 24 h at 25°C. 11

2.2. Biofertilization activity and exoenzyme production tests

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Assays were performed according to [6,7].

Phosphate solubilization: growth onto the Pikovskayas medium (PVK) [yeast extract 14 0.5 g/l, dextrose 10 g/l, phosphate calcium 5 g/l, ammonium sulfate 0.5 g/l, potassium chloride 0.2 g/l, magnesium sulfate 0.1 g/l, manganese sulfate 0.0001 g/l, iron sulfate 0.0001 g/l, 16 agar 15 g/l].

Siderophore production: growth onto the CAS medium [100 ml MM9 salt solu-
tion/750 ml ddH2O, 32.24 g piperazine-N,N'-bis-(2-ethanesulfonic acid) PIPES, 15 g agar;18after autoclaving: 30 ml Casamino acid solution, 10 ml 20% glucose solution, 100 ml Blue20Dye solution]. Blue Dye [Solution 1: 0.06 g CAS/50 ml ddH2O, Solution 2: 0.0027 g FeCl3-21 $6H_2O/10$ ml 10 mM HCl, Solution 3: 0.073 g HDTMA/40 ml ddH2O; Solution 1 + 9 ml So-22ution 2 + Solution 3].23

Proteolytic activity: growth onto Skim milk agar [50 g/l milk powder, 15 g/l agar].

Gelatinase activity: growth onto Gelatin medium [0.25% yeast extract, 0.5% Bacto Peptone, 0.5% glucose, 0.1% MgSO₄.7H₂O, 12% gelatin].

Lipolytic activity: growth onto Tween 20 and Tween 80 media [peptone 1%, CaCl_{2.2}H₂O 0.01%, agar 2%, Tween 20 or Tween 80 1%].

Amylolytic activity: growth onto Starch agar [potato starch 10 g/l, KNO₃ 0.5 g/l, K₂HPO₄ 1 g/l, MgSO₄.7H₂O 0.2 g/l, CaCl₂ 0.1 g/l, FeCl₃ traces, agar 15 g/l], and addition of Lugol to the plates for 1 min.

DNase activity: growth onto DNase agar [tryptose 20 g/l, deoxyribonucleic acid 2 g/l, sodium chloride 5 g/l, agar 12 g/l], and flooding with 1N HCl for a few min.

Plates were incubated at 25 °C for all media, and the activities were monitored after 24 h, 48 h and/or 72 h, according to the test.

2.3. Biological control tests

Diffusible compound production: the soil bacterial isolates and phytopathogenic fungi of reference were cocultured onto Potato Dextrose Agar (PDA) in dual culture plate assays [8].

Volatile organic compound (VOC) production: the soil bacterial isolates and phytopathogenic fungi of reference were cocultured onto PDA in plates divided into two sections to prevent physical contact between them [9]. Plates were subsequently sealed to allow the VOCs accumulate in the inside during the experimental period.

Plates were incubated at 25 °C for both types of assays, with daily monitoring of the cocultures. Tests were performed in triplicate for each bacterial isolate.

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3. Results

3.1. Bacterial identification

Molecular identification by partial *16S rRNA* gene amplification pointed out to a majority of species of Gram-positive bacteria, mainly belonging to the genus *Bacillus*, specifically to Group I, along with species from other several minority genera, as *Brevibacterium* 5 spp. and *Enterococcus* spp., which globally accounted for more than 75 % of the total of bacterial species isolated from agricultural soil. 7

3.2. Biofertilization activity and exoenzyme production tests

Isolates globally displayed a high proportion of the biological activities tested, especially concerning production of hydrolytic enzymes such as proteases, lipases, amylases, gelatinases and DNases but, also production of siderophores, and other activities like solubilization of phosphates (Table 1).

Table 1. Potential of the tested bacterial isolates for biotechnological applications.

Biofertilization tests	Bacillus spp.	Brevibacterium spp.	Enterococcus spp.	Other genera	Global isolates (%)
Phosphate solubilization	not detected	not detected	detected	detected/not detected	15
Siderophore production	detected	detected	detected	detected/not detected	49
Exoenzyme production tests					
Proteolytic activity	detected	detected	detected	detected/not detected	70
Gelatinase activity	detected	detected	detected	detected	
Lipolytic (on Tween 20)	detected	not detected	not detected	detected	45
activity (on Tween 80)	detected	not detected	not detected	detected/not detected	
Amylolytic activity	detected	not detected	not detected	not detected	23
DNase activity	detected	detected	not detected	detected/not detected	24

3.3. Biological control tests

Inhibition on fungal growth was also displayed among the soil bacterial isolates by16production of diffusible compounds and/or VOCs (Figure 1) against the phytopathogenic17Verticillium dahliae, Fusarium pseudograminearum, F. oxysporum, Neofusicoccum parvum and18Diplodia seriata. Diffusible compounds were produced against the five fungal species19ranging 15-30 of the isolates. Two of the isolates inhibiting V. dahliae also produced VOCs20against this pathogen. Inhibitory effect against both V. dahliae and F. pseudograminearum21was obtained with three of the isolates, against V. dahliae and F. oxysporum with three of22the isolates, and against F. pseudograminearum and F. oxysporum with four of the isolates.23Scarce production of VOCs was observed against N. parvum and D. seriata.24



Figure 1. Potential of the tested bacterial isolates for biological control applications. Representative33pictures of VOC production assays in: (a,b,c,d) against *Fusarium pseudograminearum* (a,b) and34*Verticillium dahliae* (c,d); (a,c) fungal control without bacteria, and (b,d) bacterial inhibition.35Representative pictures of a diffusible compound assay in: (e,f) against *Neofusicoccum parvum*; (e)36fungal control, and (f) bacterial inhibition.37

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4. Discussion

The activities observed in the soil bacterial isolates indicate their potential to be de-2 veloped and used as beneficial bacteria to contribute to improvement of soil quality and 3 reduction in the application of agrochemicals such as fertilizers and/or pesticides to reach 4 sustainable environmental agrosystems, similarly to observed in previous works [6,7]. Bi-5 ofertilizers based on microorganisms can naturally provide crops with nutrients, mainly 6 related to phosphorous, nitrogen, or even in smaller quantities to iron, by either increasing 7 their efficient uptake or their availability [3]. Large-scale production in bioreactors of the 8 exoenzymes tested could be of interest in agrifood industry. Antagonism against a path-9 ogen can be a treatment to be included in integrated pest management programmes. 10

Evidence from this work provides promise for the application of soil microbiome to attain sustainable agriculture, as proposed elsewhere [10].

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