

Article

The Effect of Banana Rhizosphere Chemotaxis and Chemoattractants on *Bacillus velezensis* LG14-3 Root Colonization and Suppression of Banana Fusarium Wilt Disease

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Abstract: *Fusarium oxysporum f. sp. cubense* (Foc) causes banana Fusarium wilt disease, which is a destructive soil-borne disease. Many plants can recruit rhizosphere microorganisms using their root exudates, thereby shaping the rhizosphere microbiome to resist pathogen infection. Therefore, this study was conducted to explore the role of root exudates in the process of biocontrol strain colonization and resistance to pathogens. In this study, the banana root exudates used as chemoattractants were obtained by hydroponics. *Bacillus velezensis* strain LG14-3 was isolated from the infected area of the root system of banana and showed significant chemotaxis to banana root exudates and strong inhibition of *Fusarium oxysporum f. sp. cubense*. Further analysis found that LG14-3 showed chemotaxis toward the components of banana root exudates, such as citric acid, succinic acid, glycine, D-galactose and D-maltose, and glycine and citric acid, which resulted in more significant chemotaxis of LG14-3. Moreover, banana root exudates enhanced the swarming motility and biofilm formation of LG14-3. Pot experiments showed that glycine and citric acid enhanced the colonization ability of *Bacillus velezensis* LG14-3 in the banana rhizosphere and reduced the disease severity index of banana fusarium wilt. Glycine and citric acid enhanced the growth-promoting ability of LG14-3 under pathogen stress. Our results showed that the addition of chemotactic substances enhanced the biocontrol potential of *Bacillus velezensis* LG14-3 to prevent banana Fusarium wilt.

Keywords: chemotaxis; *Bacillus velezensis*; root exudates; citric acid; glycine; banana Fusarium wilt disease; biocontrol



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1. Introduction

Banana Fusarium wilt, whose pathogen is *Fusarium oxysporum f. sp. cubense* (Foc), is one of the most devastating diseases of banana, which limits the development of the global banana industry [1]. Four physiological races of the pathogen have been identified [2]. Race 4 (Foc4) is the most harmful and infects nearly all banana varieties [1,2]. In recent years, much progress has been made in discovering antagonistic microorganisms to prevent and control banana wilt, but only a few strains have been commercialized. The control effect of most strains on Fusarium wilt of banana and other hosts is unstable and unsatisfactory, and performance in field conditions is often very impoverished compared to that in laboratory conditions [3].

Root exudates are the main cause of antagonistic bacterial colonization in the rhizosphere. Root exudates play a major role in the interaction between plants and plant growth-promoting rhizobacteria (PGPR) and between plants and pathogens [4]. As signal molecules, root exudates can directly mediate plant–microbe interactions, plant–plant interactions, and microbe–microbe interactions in the rhizosphere [5]. Root exudates are

organic materials secreted by plant roots in the rhizosphere. These organic compounds can often be separated into two classes: low-molecular-weight compounds, which include amino acids, organic acids, sugars, phenolics and an array of secondary metabolites, and high-molecular-weight compounds, including mucilage and proteins [6]. The chemotaxis of rhizosphere bacteria to root exudates determines their effect on colonization in the rhizosphere. Chemotaxis is the process by which bacteria travel to higher concentrations of attractants or lower concentrations of repellents [7]. This property allows bacteria to receive beneficial stimulation and escape harmful stimulation [8]. Both *Escherichia coli* and *Bacillus subtilis* use a modified two-component system involving the CheA histidine kinase and CheY response regulator for transmitting signals from the receptors to the flagellar motors [9,10], triggering changes in motion. Most motile bacteria can sense and respond to low concentrations of organic compounds in the surrounding environment by chemotaxis [11].

Motile bacteria are attracted by a wide variety of chemical stimuli, including sugars, amino acids, organic acids, inorganic phosphate, and aromatic compounds [12]. In recent years, an increasing number of researchers have begun to pay attention to the interaction of root exudates and their components with bacteria in vitro. Biofilm formation and swarming motility are also important for bacterial colonization in plant roots. Bacteria achieve chemotaxis through movement. Biofilm and bacterial chemotaxis are recognized as part of the colonization mechanisms [13], and biofilms can protect bacteria from external stress. Malic acid and fumaric acid in banana root exudates attract NJN-6 and promote biofilm formation by activating relevant genes [4]. Organic acids and sugars enhance the biofilm formation of *Pseudomonas* sp. and *Pseudomonas monteilii* [14].

Bacillus spp. are common soil microorganisms that can colonize the rhizosphere and benefit plants [15]. Many *Bacillus* species have also been shown to have chemotactic effects on root exudates. Banana rhizosphere bacterium *B. subtilis* N11 showed significant chemotaxis toward banana root exudates [16]. *Bacillus cereus* YL6 showed chemotaxis to malate, oxalic acid, citric acid, lactic acid, and succinic acid [17]. Root exudates promote the *Bacillus* colonization effect on the plant rhizosphere and thus suppress pathogens. The phenolic acids of banana root exudates may have the potential to enhance root colonization and pathogen suppression abilities of *Bacillus amyloliquefaciens* NJN-6 [4].

Biological antagonistic bacteria are an important control measure for banana Fusarium wilt disease. The biological control of *Bacillus* on banana Fusarium wilt disease has been studied [18–21]. Endophytic *Bacillus subtilis* TR21 improved banana plant resistance to *Fusarium oxysporum* f. sp. *cubense* and promotes root growth [22]. *Bacillus siamensis* had strong Fusarium wilt control and growth-promoting effects on banana [23]. The combinations of *Bacillus* and different amendments are used to control banana Fusarium wilt disease. Biocontrol bacterium *B. velezensis* HN03 combined with wormcast could induce plant resistance to Fusarium wilt and suppress disease [24]. The combination of *B. velezensis*, *Bacillus subtilis* and *Penicillium* sp. could suppress banana disease [25].

Chemotaxis is considered an important characteristic of a successful antagonistic bacterium candidate for soil-borne pathogens [26,27]. The antagonistic microorganisms move toward plant rhizosphere chemoattractants and successfully colonize the rhizosphere to exert more effective biocontrol against pathogens [15,17,26,28]. Therefore, it is very important to optimize the selection of antagonistic bacteria sources with the chemotactic potential to act as biocontrols against soil-borne disease [26]. For example, strong chemoattractant (maize root exudates) of *Bacillus amyloliquefaciens* OR2-30 induced its colonization in maize roots to inhibit *Fusarium graminearum* infection [15]. However, little is known regarding how an antagonistic microorganism affects the suppression mechanisms of banana Fusarium wilt disease when combined with different chemoattractants.

Therefore, the present study focuses on the role of root exudates in biocontrol strain colonization in roots and resistance to pathogens. Therefore, in this study, *Bacillus velezensis* LG14-3 was screened based on its significant chemotactic behavior toward banana root exudates, and the effect of chemotaxis on LG14-3 biocontrol-related traits was investigated.

Finally, pot experiments were conducted to assess the ability of LG14-3 to colonize banana roots and its biocontrol potential against banana wilt disease.

2. Materials and Methods

2.1. Plant Materials, Bacterial and Fungal Strains, and Culture Conditions

Tissue-cultured seedlings of Brazilian banana (*Musa acuminata* Cavendish cv. Brail). *Fusarium oxysporum* f. sp. *ubense* race 4 (Foc4) strain B2 (BioProject No.: PRJNA174275) and other fungal strains were cultured at 28 °C on Potato Dextrose Agar (PDA) medium (200 g/L potato, 20 g/L glucose). Bacterial strains were cultured in a Luria Bertani (LB) medium at 37 °C (10 g/L tryptone, 5 g/L yeast extract, 10 g/L NaCl, 15 g/L agar, pH 7.0). The biocontrol strains were tetracycline-resistant (400 µg/mL). All strains utilized in this study were obtained from the Environment and Plant Protection Institute, Chinese Academy of Tropical Agricultural Sciences.

2.2. Collection of Banana Root Exudates

Healthy banana seedlings were rinsed 3 times using ultrapure water. Each seedling was transferred into 100 mL deionized water in a 100 mL conical flask wrapped in tin foil. The water was changed every day. Seedlings were cultivated to 5–6 leaves and washed with sterile water 3 times. After washing, 50 mL deionized water was added to the conical flask. Seedlings were cultivated for 72 h at room temperature, and the liquid (the root exudates) was collected. A total of 500 mL of root exudates were frozen in a −70 °C refrigerator and dried with a freeze-dried dryer at low temperature until the powder was obtained. The powder was dissolved using 25 mL ultrapure water (20RE), sterilized using a 0.22 µm water phase microtool-filter membrane, and stored at −70 °C. Different concentrations of root exudates were prepared: (1) 20RE; (2) 10RE (diluted 2-fold 20RE); (3) RE (diluted 20-fold 20RE).

2.3. Isolation and Purification of Bacteria

Samples were collected from healthy banana roots in the field of the epidemic area in Lingao County of Hainan Province, China. A total of 0.1 g of banana root system was washed briefly using sterile water and ground with a mortar in 2 mL of sterile water. The suspension was heated in a water bath at 80 °C for 10 min to isolate *Bacillus*. The suspension was diluted, inoculated on LB medium and incubated in a constant temperature incubator at 37 °C for 12 h. Bacterial colonies with different morphologies and colors were picked and passaged, and the pure cultures obtained were stored in 30% glycerol in a −80 °C refrigerator for later experiments.

2.4. Chemotaxis Assay

Banana root secretions were used for chemotaxis analysis. Seven sugars (D-maltose, D-galactose, D-xylan, D-alginate, sucrose, D-fructose and glucose), four amino acids (L-tryptophan, histidine, glycine and D-glutamic acid) and five organic acids (succinic acid, fumaric acid, malonic acid, citric acid and L-malic acid), which are common components of banana root secretions, were also used for chemotaxis assay. The sample to be analyzed was dissolved in chemotactic buffer to a concentration of 10 mM and then sterilized by filtration.

The chemotaxis assay is based on the “drop” assay [29] with minor modifications. Briefly, cells were cultured overnight in 150 mL of LB medium. The culture was centrifuged to remove the supernatant and suspended using 100 mL chemotactic buffer ($K_2HPO_4 \cdot 3H_2O$ 13.9 g/L, KH_2PO_4 5.3 g/L, EDTA 0.0067 g/L, pH 7.0). A 25 mL solution of 1% hydroxypropylmethylcellulose was added to the cell suspension. Then, a 15 mL bacterial suspension was transferred to a Petri dish. A total of 20 µL of 20RE or each substance at a concentration of 10 mM was tested. The plates were incubated at room temperature for 0.5 to 2 h and then observed for the appearance of transparent circles.

A quantitative capillary assay was performed [17,30]. Bacterial strains were cultured in LB medium until the absorbance value of the solution at wavelength 600 nm (OD600)

reached 1.0, and then, the suspensions were centrifuged and resuspended in chemotaxis buffer. Then, 100 μL of the cell suspension was transferred to a 200- μL pipette tip. Instead of a capillary, a 1-mL syringe was used for the quantification assay, and 100 μL of the sample to be tested or chemotaxis buffer was aspirated separately. Then, the syringe needle was placed into the bacterial suspension in the pipette to bring the two solutions into full contact. After two hours of incubation at room temperature, the solution in the syringe was diluted with chemotaxis buffer and coated on LB plates and the number of colonies on the plates was counted after 12 h. The assay was tested 3 times.

$$\begin{aligned} \text{Relative chemotaxis index (RCI)} \\ = \text{the bacterial number toward the sample tested} \\ \div \text{bacterial number toward chemotaxis buffer} \end{aligned}$$

A value of RCI less than or equal to 1 indicates no chemotaxis, while a value of RCI greater than 1 indicates chemotaxis. Additionally, a value of RCI greater than or equal to 2 indicates a significant chemotactic response.

2.5. Identification of the Strain

The physiological and biochemistry characteristics of the bacterium were identified [31], such as Gram staining, endospore staining and methyl red (MR), catalase, citrate utilization, starch hydrolysis, nitrate reduction, protease production and cellulase production tests.

Molecular identification was performed. The genomic DNA of bacteria was extracted using an Ezup Column Bacteria Genomic DNA Purification Kit (Sangon Biotech, Shanghai, China). The 16S rRNA gene and *gyrB* gene were amplified by PCR. For the amplification of the 16S rRNA gene fragment, the 27F/1492R primer set was used. For the amplification of the *gyrB* gene fragment, the UP-1/UP-2r primer set was used. The 50 μL reaction mixture consisted of 2 μL DNA template, 2 μL 27F/UP-1, 2 μL 1492R/UP-2r, 25 μL PCR mix (0.05 units μL^{-1} Taq DNA polymerase; 4 mM MgCl_2 , and 0.4 mM dNTPs), and 19 μL nuclease-free water. The reaction conditions of the 16S rRNA gene fragment were 95 $^{\circ}\text{C}$ for 5 min, followed by 35 cycles of 95 $^{\circ}\text{C}$ for 30 s, 60 $^{\circ}\text{C}$ for 30 s and 72 $^{\circ}\text{C}$ for 2 min, with a final extension step at 72 $^{\circ}\text{C}$ for 10 min. The reaction conditions of the *gyrB* gene fragment were 95 $^{\circ}\text{C}$ for 5 min, followed by 35 cycles of 95 $^{\circ}\text{C}$ for 30 s, 55 $^{\circ}\text{C}$ for 30 s and 72 $^{\circ}\text{C}$ for 2 min, with a final extension step at 72 $^{\circ}\text{C}$ for 10 min. The amplification products were detected by horizontal electrophoresis through 1% agarose gels. PCR amplification products were sequenced at Hainan Nanshan Biotechnology Company (Haikou, China). Sequences were BLAST searched against the NCBI database. The reference 16S and *gyrB* nucleotide sequences were obtained from GenBank. The 16S and *gyrB* sequences of each strain were aligned using MEGA-X, and characters were weighted equally. Then, the two single-sequence datasets were concatenated and aligned using ClustalX (1.7). The neighbor-joining method with 1000 bootstrap replicates was used to build a phylogenetic tree.

2.6. Plant Growth-Promoting Properties of the Strain

The production of auxin was tested [32]. Briefly, bacteria were cultured in a liquid LB medium supplemented with 5 mM L-tryptophan. After 5 days of incubation at 37 $^{\circ}\text{C}$ with shaking at 180 rpm, the bacterial cultures were centrifuged at 6000 $\times g$ for 10 min, and the supernatant was mixed (1:1, *v/v*) with Salkowski reagent (12 g of FeCl_3 per liter in 7.9 M H_2SO_4) and incubated in the dark at room temperature for 30 min. The mixture turned red.

The production of siderophores was tested [33,34]. The isolate was cultured in LB liquid medium for 14 h. Bacterial liquid was inoculated into Chrome azurol S (CAS) plates for 2–3 days at 37 $^{\circ}\text{C}$; the strain producing an orange iron chelating circle was further tested. The isolate was inoculated into LB culture medium at 37 $^{\circ}\text{C}$ for 2 days. After incubation, the cultures were centrifuged for 10 min at 10,000 $\times g$; the same volume of supernatant was mixed with CAS detection solution and reacted for 20 min. The absorbance was determined at 630 nm (i.e., As). The same volume of LB medium was mixed with CAS

detection solution and reacted for 20 min as a control, and the absorbance was determined at 630 nm, which is the absorbance of Ar. The amount of iron carrier was calculated according to the formula $(As - Ar)/Ar \times 100\%$.

Nitrogen fixation was tested. One hundred microliters of bacterial suspension was inoculated in Ashby's nitrogen-free medium (mannitol 10.0 g/L, KH_2PO_4 0.2 g/L, $MgSO_4 \cdot 7H_2O$ 0.2 g/L, NaCl 0.2 g/L, $CaSO_4 \cdot 2H_2O$ 0.2 g/L, $CaCO_3$ 5.0 g/L, pH 7.0–7.2). The bacteria were cultured at 28 °C for 7 days and observed for single colony production.

2.7. Effect of Chemotaxis on the Growth of the Strain In Vitro

The bacterial cells were cultured overnight in LB medium and then adjusted to an OD600 of 1.0, resuspended 3 times with minimal medium without a carbon source ($(NH_4)_2SO_4$ 2.0 g/L, $NaH_2PO_4 \cdot H_2O$ 0.5 g/L, K_2HPO_4 0.5 g/L, $MgSO_4 \cdot 7H_2O$ 0.2 g/L, $CaCl_2 \cdot H_2O$ 0.1 g/L), and then added at 1% to minimal medium containing different concentrations of chemoattractant as the sole carbon source. Cultures were cultured at 37 °C with 180 rpm shaking for 48 h, and finally, the OD600 was determined at different time points for each treatment.

2.8. Swarming Motility Assay

The effect of root exudates on the swarming ability of strains was measured as follows [15]. The bacterial cells were cultured overnight in an LB medium and then adjusted to an OD600 of 1.0. After centrifugation of the cultures, the cells were resuspended in sterile water (control) and different concentrations of root exudates (RE, 10RE and 20RE). Ten microliters of different bacterial suspensions were cultured in LB medium with 0.7% agar and incubated at 37 °C for 12 h. Strain LG11-2 was used for comparison. The assay was repeated three times.

2.9. The Effect of Chemotaxis on Bacterial Biofilm Formation

Four kinds of biofilm mediums were prepared separately: Mmsgg medium (control); Mmsgg medium and root exudates (final concentration of root exudates: 20RE); Mmsgg medium and citric acid (final concentration of citric acid: 0.25 mM); and Mmsgg medium and glycine (final concentration of glycine: 0.25 mM). The bacterial cells were cultured overnight in LB medium and then resuspended in Mmsgg medium and adjusted to an OD600 of 1.0. Then, 20 µL of bacterial suspension was added to a 24-well plate, and each well with 2 mL of biofilm medium, and the plate was incubated for 72 h at 30 °C in a constant temperature incubator. For quantitative analysis of biofilm formation [15], the cultures were carefully removed from wells. The remaining cells were gently washed with PBS buffer three times and fixed with 1 mL methanol for 15 min. Then, 1% crystal violet solution was added to each well for 5 min, and the excess cells were rinsed off with sterilized water. Finally, 33% glacial acetic acid was added to extract the dye, and the absorbance of the solution was measured at 570 nm. The assay was repeated three times.

2.10. Colonization Assay

Bacterial cells were cultured in LB medium at 37 °C with 180 rpm shaking for 48 h. The cells were resuspended in sterile water, and the concentration of the bacterial suspension was adjusted to 5×10^7 colony-forming units/mL (cfu/mL). Forty milliliters of bacterial suspension was added to each seedling, and 6 seedlings were included in each treatment. The banana roots were taken on the 5th, 10th, and 15th days. Banana seedlings were pulled out of the soil. Banana plant roots were washed with sterile water for 3 min, soaked in 75% ethanol by volume for 1 min, rinsed with sterile water 3–5 times and ground with a sterile mortar. The ground solution was spread in tetracycline-LB solid medium, and colonies on the medium for 12 h were checked. The assay was repeated three times.

Three kinds of solutions were prepared as follows: 0.25 mM citric acid; 0.25 mM glycine; sterile water (CK). The banana roots were rinsed three times using sterile water, immersed in a solution for 1 min, placed in a conical flask (bacterial suspension

5×10^7 cfu/mL) and incubated at room temperature for 72 h. Six seedlings were included in each treatment. Banana plant roots were taken, washed briefly with sterile water, ground into homogenate, coated in tetracycline-LB solid plate. After being cultured at 37 °C for 12 h, and the colonies of LG14-3 were checked. The experiment was repeated three times.

2.11. Antagonistic Assays

For antifungal bioassays, the 5 mm pathogenic fungi cake was inoculated in the center of the PDA medium. A total of 5 µL of fresh bacterial culture (OD600 = 1.0) was inoculated evenly at 3 cm from the center of the Petri dishes in all four directions; the Petri dishes were incubated at 28 °C, and the inhibition rate was calculated after 7 days. An equivalent volume of sterile water was used as a control in place of bacterial culture. *The inhibition rate (%) = (Diameter of transparent circle) ÷ Diameter of colony × 100%*. The experiment was repeated three times.

The inhibition effect of strain LG14-3 metabolite on Foc was evaluated. The bacterial cells were cultured in the LB medium at 37 °C with 180 rpm shaking for 3 days and centrifuged to obtain the supernatant. Then, 100 µL fresh culture of Foc4 (10^7 cfu/mL) was evenly distributed into Petri dishes, and the holes were punched evenly on the PDA medium. An equal amount of filtered and debacterized supernatant was added to the holes, and the transparent circle around the holes was observed by incubation at 28 °C for 2–3 days. The experiment was repeated 3 times.

For the antibacterial bioassay, pathogenic bacteria were evenly distributed in LB medium. Five microliters of the tested culture was inoculated on Petri dishes and incubated at 37 °C for 3–5 days. The diameters of antibacterial zones were measured, and the inhibition rate was calculated. *Inhibition rate (%) = (diameter of antibacterial zone – diameter of colony) ÷ diameter of antibacterial zone × 100%*. The experiment was repeated 3 times.

2.12. Biocontrol Experiment

The strain was cultured in LB liquid medium for 48 h, centrifuged to remove the supernatant and resuspended in sterile water, and the concentration of the bacterial solution was adjusted to 5×10^7 cfu/mL. Foc4 was cultured on PDA liquid medium for 5 days, centrifuged to remove the supernatant and resuspended in sterile water, and the concentration of the suspension was adjusted to 1×10^6 (conidia/mL). Four experimental treatments were carried out as follows: CK1 treatment group plants were treated with Foc; CK2 treatment group plants were treated with LG14-3 and Foc; T1 treatment group plants were treated with glycine, LG14-3 and Foc; and T2 treatment group plants were treated with citric acid, LG14-3 and Foc. First, T1 treatment group plants were soaked in 0.25 mM glycine solution for 1 min. T2 treatment group plants were soaked in 0.25 mM citric acid solution for 1 min. CK1 and CK2 treatment group plants were soaked in sterile water for 1 min. Then, plants were planted in sterilized soil. A 100 mL bacterial suspension (5×10^7 cfu/mL) was added around the root of each seedling in the T1, T2 and CK2 treatment groups, and 100 mL sterile water was added around banana roots in the CK1 treatment group. Seven days later, 100 mL of Foc4 suspension was added to each pot of banana seedlings. The growth status of banana seedlings was observed daily. After 60 days, disease incidence was counted, the disease severity index was calculated [35], and plant height (distance from the ground to the tip of the uppermost spreading leaf), pseudostem girth, and pseudostem height (distance from the ground to the intersection of the petioles of the top two leaves) were measured.

Disease severity index

$$= \sum(\text{number of disease plants at all levels} \\ \times \text{representative value of the level}) \\ \div \text{total number of plants} \\ \times \text{representative value of the highest level} \times 100$$

2.13. Extraction and Determination of Chlorophyll Content

A 0.1 g mass of freshly washed banana leaves was placed into a 10 mL mixture of acetone and anhydrous ethanol (1:1, *v/v*), soaked for 48 h in the dark at room temperature. The absorbance of the extracted solution was measured at 645 nm and 663 nm with a mixture of acetone and anhydrous ethanol as the reference. The contents of chlorophyll a and chlorophyll b were calculated by Amon's formula [36]: Ca (mg/g) = $(12.7 \times A^{663} - 2.69 \times A^{645} \times V)/(1000 W)$; Cb (mg/g) = $(22.7 \times A^{645} - 4.68 \times A^{663}) \times V/(1000 W)$; C (mg/g) = $Ca + Cb$. Absorbance values at 645 nm and 663 nm are represented by A^{645} and A^{663} , respectively. Ca and Cb are the contents of chlorophyll a and chlorophyll b, respectively. V is the volume of the extraction solution, and W is the mass of the leaf sample.

2.14. Statistical Analysis

The data were subjected to one-way analysis of variance (ANOVA) and Tukey's test using IBM SPSS Statistics 26. Figures were generated using GraphPad Prism 8.

3. Results

3.1. LG14-3 Exhibited Prominent Chemotactic Behavior toward Banana Root Exudates, Glycine, and Citric Acid

A chemotaxis assay was used to screen 11 bacterial strains from the banana rhizosphere that have chemotaxis to root exudates (Table 1). LG14-3 showed significant chemotaxis to banana root exudates (Figure 1b), whose relative chemotactic index (RCI) was 3.62.

Table 1. Relative chemotaxis index (RCI) to root exudates in strains.

<i>Bacillus</i> Strains	Relative Chemotaxis Index (RCI)
<i>Bacillus velezensis</i> strain LG11-2	1.93 ± 0.02
<i>Bacillus velezensis</i> strain YH-6	1.25 ± 0.11
<i>Bacillus velezensis</i> strain C3	1.91 ± 0.05
<i>Bacillus aerophilus</i> strain YC-6	1.56 ± 0.02
<i>Bacillus aerophilus</i> strain CM13	1.71 ± 0.08
<i>Bacillus aerophilus</i> strain HZ-B	1.09 ± 0.02
<i>Bacillus aerophilus</i> strain JS-5	1.36 ± 0.07
<i>Bacillus velezensis</i> strain X2-1	1.2 ± 0.05
<i>Bacillus megaterium</i> strain Y1R3	1.44 ± 0.04
<i>Bacillus velezensis</i> strain LG14-3	3.62 ± 0.09
<i>Bacillus velezensis</i> strain A3	1.58 ± 0.06

Note: Values are the means ± standard errors.

The chemotaxis of LG14-3 to components of banana root exudates was further evaluated. The "Drop" assay showed that LG14-3 had chemotaxis to five compounds (10 mM), including glycine, D-galactose, D-maltose, citric acid, and succinic acid (Figure 1c–g). Quantitative capillary assays showed that glycine and citric acid had particularly marked effects (RCI ≥ 2) on LG14-3 chemotaxis at various tested concentrations. The strongest chemotactic response was for citric acid (0.25 mM), with an RCI value of 8.37. The chemotactic response of LG14-3 to glycine, citric acid, and succinic acid followed an inverted "U" shape, with a maximum attractive concentration of 0.25 mM. The chemotactic response toward D-maltose and D-galactose was biphasic, with attractive concentration maxima of 0.05 mM (Figure 2).

3.2. Morphological and Molecular Identification and Plant Growth-Promotion Research of LG14-3

The strain was cultured on LB agar medium at 37 °C for 12 h and produced round and creamy white colonies with dry wrinkles on the surface and a rounded depression in the middle. LG14-3 is a Gram-positive bacterium that is able to produce endospores. LG14-3 can decompose hydrogen peroxide and liquefied gelatin, utilize citrate, reduce nitrate, and produce cellulase, protease and amylase (Table 2).

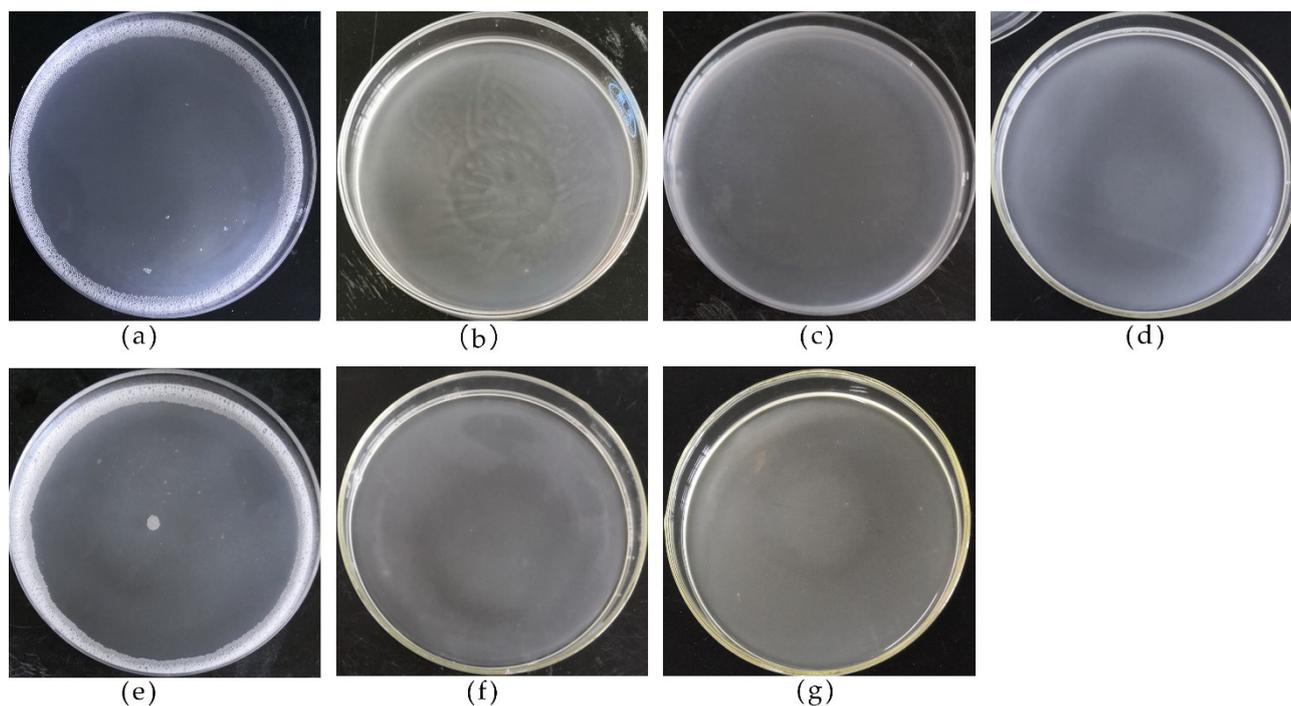


Figure 1. The “drop” assay was conducted. Drops added to the center of the plate contained, (a) chemotactic buffer; (b) root exudates; (c) citric acid; (d) glycine; (e) succinic acid; (f) D-maltose; (g) D-galactose.

For *Bacillus*, it is not possible to rely on 16S sequences alone to identify strains due to the high degree of sequence similarity between species [37]. Therefore, the phylogenetic tree was established using the combination of 16S and *gyrB* genes. The phylogenetic tree showed that strain LG14-3 and *Bacillus velezensis* NN89 clustered in the same large branch with 99% bootstrap (Figure 3). Based on morphological, physiological and biochemical characteristics, and phylogenetic analyses, LG14-3 was identified as *Bacillus velezensis*.

LG14-3 can produce indoleacetic acid (IAA) and siderophores, and the synthesis rate of siderophores is 37%. LG14-3 produced elastic and sticky, oil-drop-like single colonies on Ashby’s nitrogen-free medium, indicating that LG14-3 has nitrogen fixation (Table 2).

Table 2. Morphological and Biochemical characterizations of LG14-3.

Morphological Characters		Morphological Characters		Carbon Source	
Shape	Round	Gram staining	+	Sorbitol	+
Color	Creamy white	Methyl red test	+	D-maltose	+
Gram staining	+	Catalase test	+	Mannitol	+
Endospore production	+	Nitrate reduction	+	Sucrose	+
		Citrate test	+	D-galactose	+
		Starch hydrolysis	+	D-xylan	+
		protease production	+	Glucose	+
		Cellulase production	+	Glucose	+
		Nitrogen fixation	+		
		Phosphate solubilization	–		
		Iron carrier production	+		
		Indoleacetic acid production	+		

Note: +, Positive; –, Negative.

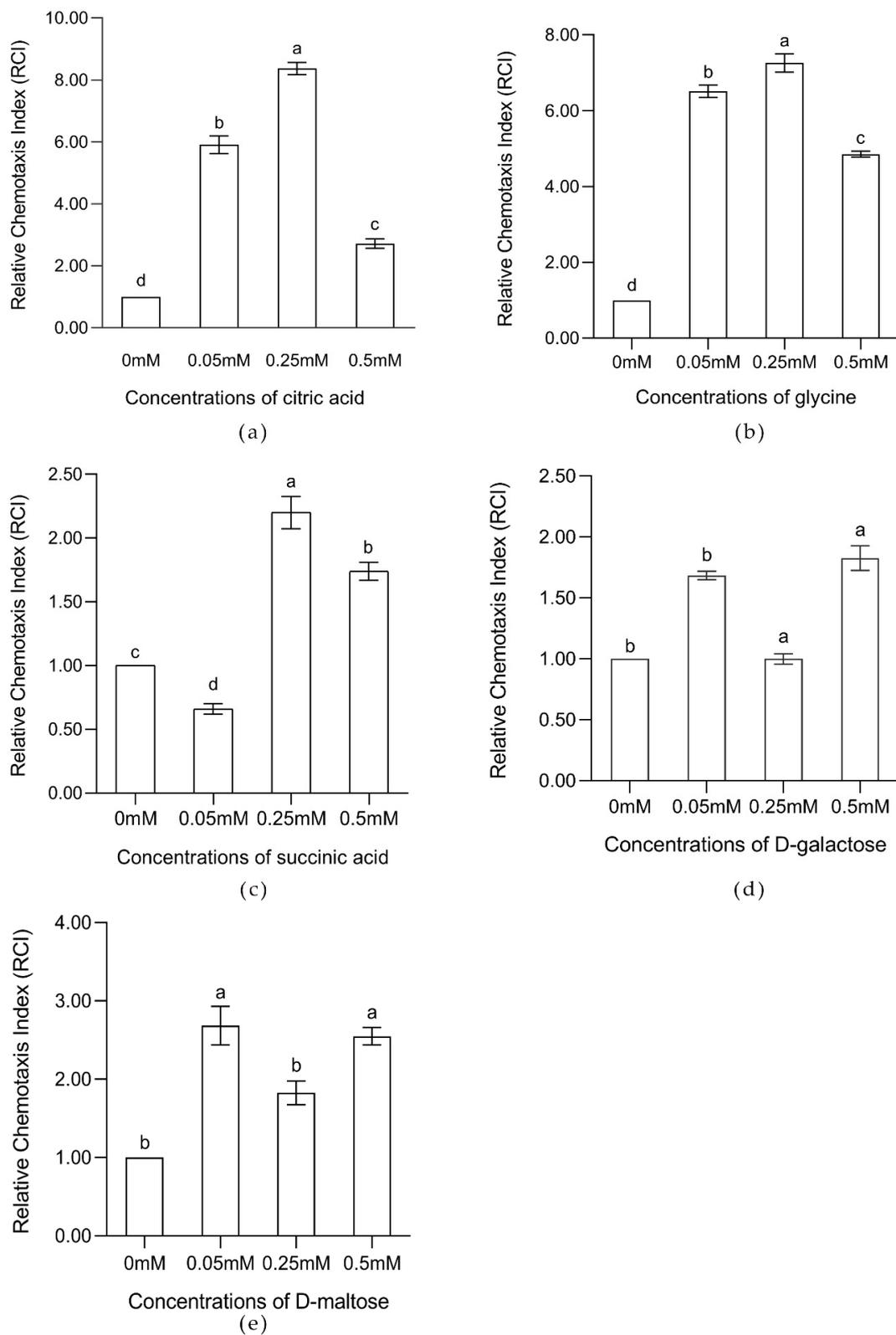


Figure 2. Chemotactic responses of LG14-3 to different chemotaxis at final concentrations of 0.05 mM, 0.25 mM, 0.5 mM by capillary assay. The chemotaxis buffer was supplemented with (a) citric acid; (b) glycine; (c) succinic acid; (d) D-galactose; and (e) D-maltose. Bars indicate standards errors from the means of three replicates. Letters above the columns represent significant difference for each treatment according to Duncan's multiple range test ($p < 0.05$).

3.3. Banana Root Exudates Promoted the Swarming Ability of LG14-3

The swarming analysis showed that compared with the control (CK, sterile water), strain LG14-3 showed stronger swarming ability under root exudates, and the higher the concentration of root exudates was, the larger the area of swarming (Figure 4a,c). However, compared with the control (sterile water), 20RE could enhance the swarming ability of LG11-2, while RE and 10RE could not, and RE inhibited the swarming ability of LG11-2 (Figure 4b,d). The results showed that root exudates enhanced the swarming ability of LG14-3.

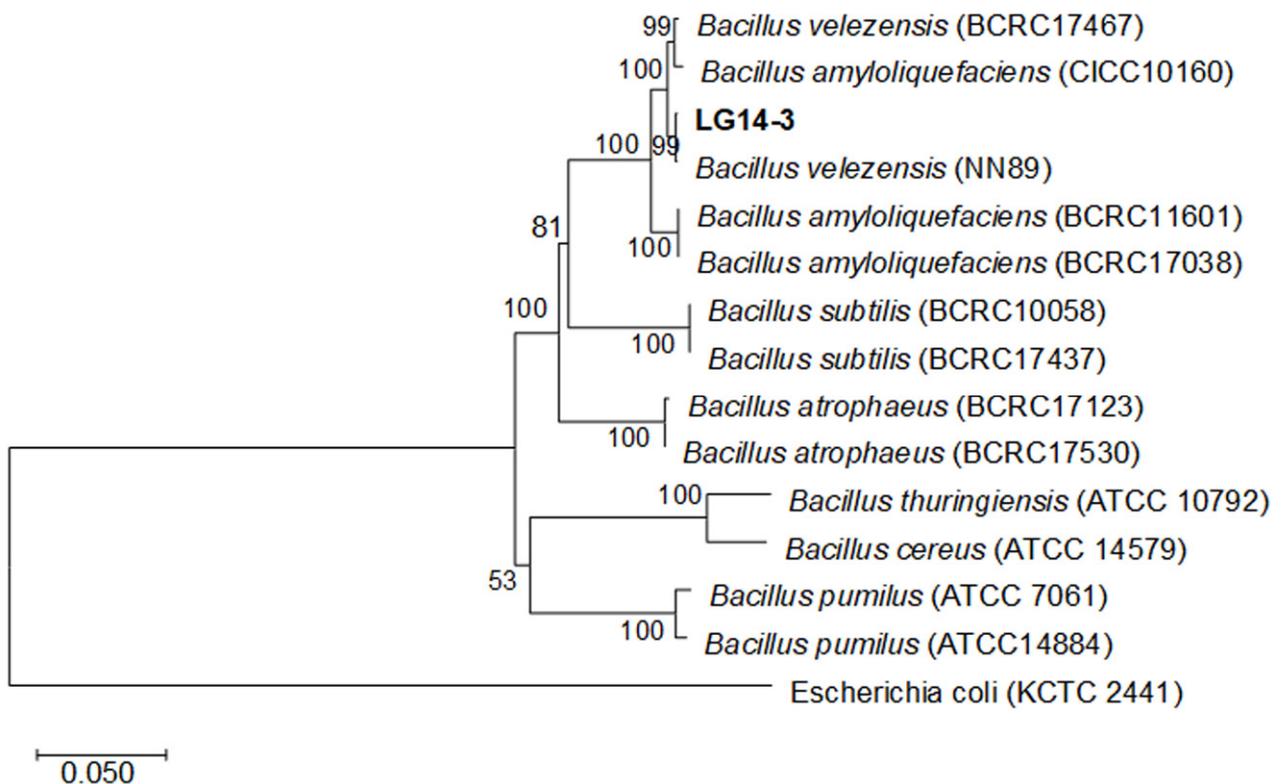


Figure 3. Phylogenetic tree based on 16S and *gyrB* genes.

3.4. Effects of Citric Acid and Glycine on LG14-3 Growth and Biofilm Formation

Compared to the control (0 mM), the concentrations of 0.05 mM, 0.25 mM, and 0.5 mM citric acid supported cell growth up to 0.066, 0.136, and 0.247 of OD600 values, respectively. However, glycine at concentrations of 0.05 mM, 0.25 mM, and 0.5 mM did not support cell growth in the basic medium, and the OD600 values were close to or smaller than the control (Figure 5B(a)). Glycine and citric acid at 0.25 mM did not support the growth of LG11-2 (Figure 5B(b)).

Biofilm formation in plant roots indicates the successful colonization of PGPR [4]. Studies have shown that biofilms can protect rhizosphere bacteria against pathogens and are conducive to enhancing rhizosphere bacteria to obtain nutrients [13,14]. LG14-3 showed the most significant chemotactic response to 0.25 mM citric acid and 0.25 mM glycine, so the concentration used for those two compounds was 0.25 mM; In contrast, LG11-2 had no chemotactic response to 0.25 mM citric acid or glycine. The results showed that the root exudates (20RE), 0.25 mM citric acid, and 0.25 mM glycine enhanced the biofilm formation of LG14-3. The treatment groups of 0.25 mM glycine and 0.25 mM citric acid had more obvious folds and thicker biofilms than the control and root exudates treatments. Crystal violet quantitative analysis showed that the absorbances of biofilms formed by adding chemoattractants to Mmsg medium were higher than the control at 570 nm. Among them, the biofilm formed in the citric acid treatment group had the highest value at OD570 nm (Figure 5A(a,b)).

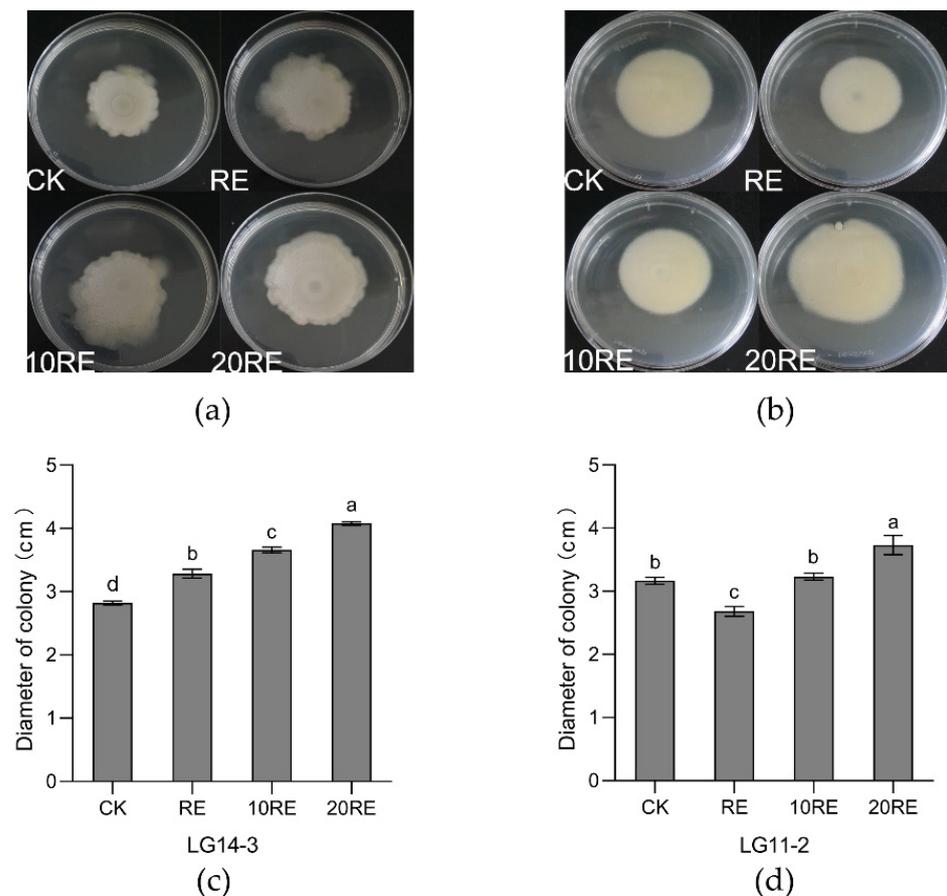


Figure 4. The effects of root exudates on the swarming motility. 5 μ L *Bacillus* suspensions (10^7 cfu/mL) mixed with different concentrations of banana root exudates (RE, 10RE, and 20RE) or sterilized water (CK) were inoculated in the LB medium with 0.7% agar. After incubating at 37 °C for 12 h, the diameters of colonies were measured. (a,c), LG14-3; (b,d), LG11-2. Bars indicate standard errors from the means of three replicates. Letters above the columns represent significant difference for each treatment according to Duncan's multiple range test ($p < 0.05$).

The biofilm morphology of LG11-2 showed that the biofilms formed in the treatment groups treated with glycine and citric acid were thinner. Crystal violet quantitative analysis of biofilms showed that root exudates, glycine, and citric acid did not promote the biofilm formation of LG11-2. The root exudates did not affect biofilm formation, while glycine and citric acid inhibited biofilm formation (Figure 5A(a,c)).

3.5. LG14-3 Has Colonization Advantages in Banana Roots, and Citric Acid can Promote Its Colonization Ability in the Banana Rhizosphere

The antagonistic activities of *Bacillus velezensis* LG11-2 and *Bacillus velezensis* FSB23-1 against Foc4 were similar to that of LG14-3 in vitro, but LG11-2 and FSB23-1 had no significant chemotactic response to banana root exudates. The colonization number of LG14-3 in banana roots reached 2.12×10^4 cfu/g on the 10th day. Compared to LG11-2 and FSB23-1, the colonization amount of LG14-3 in the root was 2.6 and 3.7 times greater on the 5th day, 5.03 and 3.72 times greater on the 10th day and 6.41 and 3.63 times greater on the 15th day, respectively (Figure 6a).

The results of hydroponic experiments showed that the addition of exogenous glycine and citric acid promoted the colonization of strain LG14-3. The citric acid (0.25 mM) significantly promoted colonization by strain LG14-3 (Figure 6c).

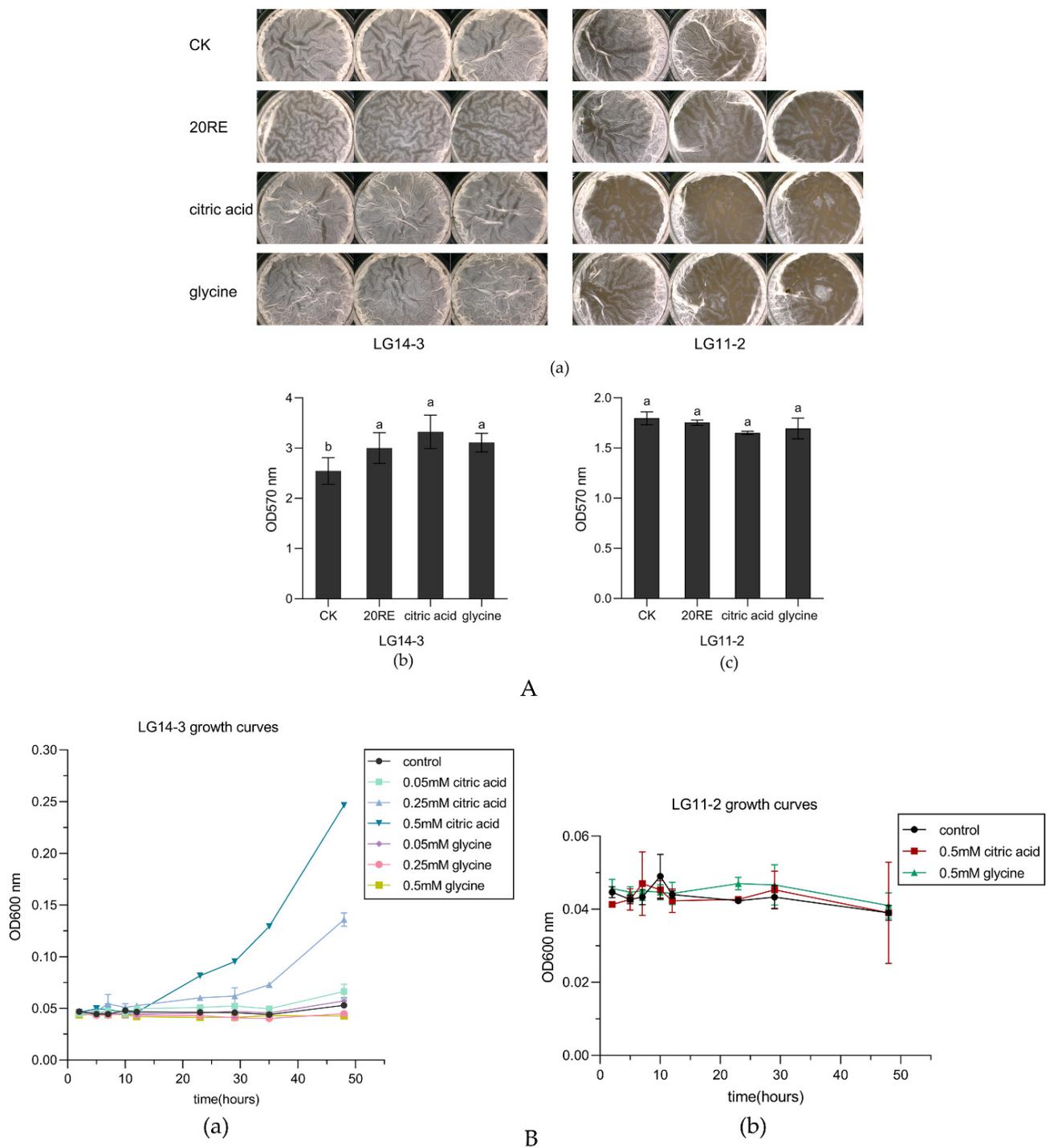


Figure 5. (A) Influence of chemoattractant on the biofilm formation. (a) biofilm phenotype of *B. velezensis* LG14-3 and LG11-2 in biofilm medium; (b,c), crystal violet staining was used to quantify the biofilm. Bars indicate standard errors from the means of three replicates. Letters above the columns represent significant differences for each treatment according to Duncan's multiple range test ($p < 0.05$); (B) Growth curves of *B. velezensis* LG14-3 and LG11-2 in minimal medium containing citric acid and glycine at the concentration indicated in the text as a carbon source. Bars indicate standard errors from the means of three replicates.

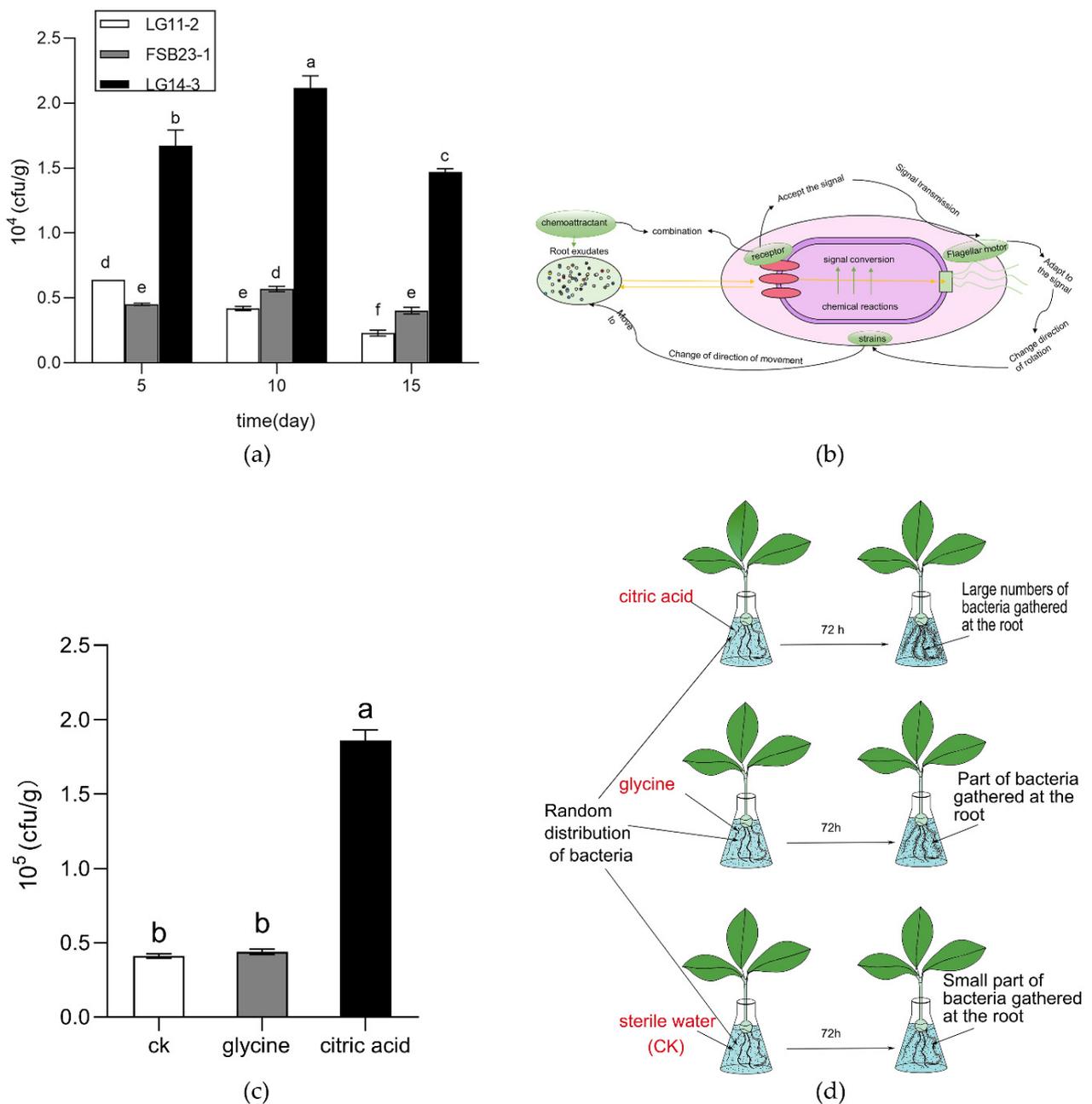


Figure 6. Colonization assay. (a) Effect on the colonization of *B. velezensis* LG14-3, LG11-2, and FSB23-1 in the banana root. (b) Schematic diagram of chemotaxis of strains to root exudates. (c,d) The effect of glycine and citric acid on the colonization of LG14-3 in the rhizosphere of banana. (a,c) Bars indicate standard errors from the means of three replicates. Letters above the columns represent significant difference for each treatment according to Duncan's multiple range test ($p < 0.05$).

3.6. The LG14-3 Strain Exhibited Broad-Spectrum Antagonistic Activity

The LG14-3 strain has broad-spectrum antagonistic activity against *Xanthomonas campestris* pv. *mangiferaeindicae*, *Pestalotiopsis*, *Diaporthe*, *Gliomastix murorum*, and Foc4 (Figure 7a). LG14-3 had a 78% inhibitory rate against Foc4 (Table 3). The inhibition rates of LG14-3 on *Xanthomonas campestris* pv. *mangiferaeindicae*, *Pestalotiopsis*, *Diaporthe*, *Lasiodiplodia*, and *Gliomastix murorum*, were 80.99%, 76.47%, 63.81%, 75.19%, and 76.6%, respectively (Table 3).

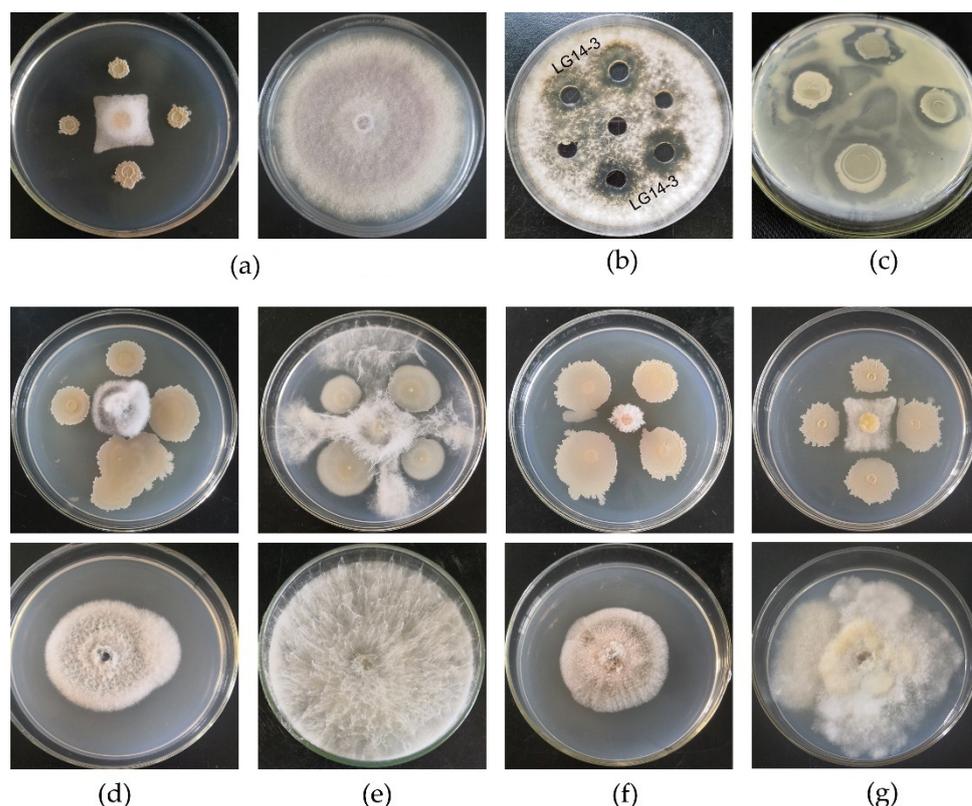


Figure 7. The antagonistic effect of strain LG14-3. Pathogens, (a), (b), Foc4, ((b), the antagonistic effect of metabolites of LG14-3); (c) *Xanthomonas citri* pv. *mangiferaeindicae*; (d) *Diaporthe*; (e) *Pestalotiopsis*; (f) *Lasiodiplodia*; (g) *Gliomastix murorum*.

Table 3. Inhibition rates of LG14-3 on pathogens.

Pathogens	Inhibition Rates (%)
Foc4	78.00 ± 0.94
<i>Xanthomonas campestris</i> pv. <i>mangiferaeindicae</i>	80.99 ± 4.55
<i>Diaporthe</i>	63.81 ± 1.90
<i>Lasiodiplodia</i>	75.19 ± 1.70
<i>Gliomastix murorum</i>	76.60 ± 2.13
<i>Pestalotiopsis</i>	76.47 ± 2.35

Note: Values are the means ± standard errors.

3.7. Biological Control Experiment

Compared with the CK1 (Foc) group, the blackened area of banana seedling corms in the CK2 (LG14-3 + Foc), T1 (glycine + LG14-3 + Foc), and T2 (citric acid + LG14-3 + Foc) groups was significantly reduced, and the T2 treatment group had the smallest blackened area of corms (Figure 8a) and the lowest disease severity index (22.92) (Figure 8b). The disease severity indices of CK1, CK2, and T1 were 93.75, 60.42, and 33.33, respectively. The measured data of plant height, pseudostem height, pseudostem girth, and chlorophyll content of banana seedlings of each group showed that LG14-3 promoted plant growth and increased chlorophyll content, and the exogenous substances (glycine and citric acid) enhanced the growth-promoting ability of LG14-3 on the plants (Figure 8c,d).

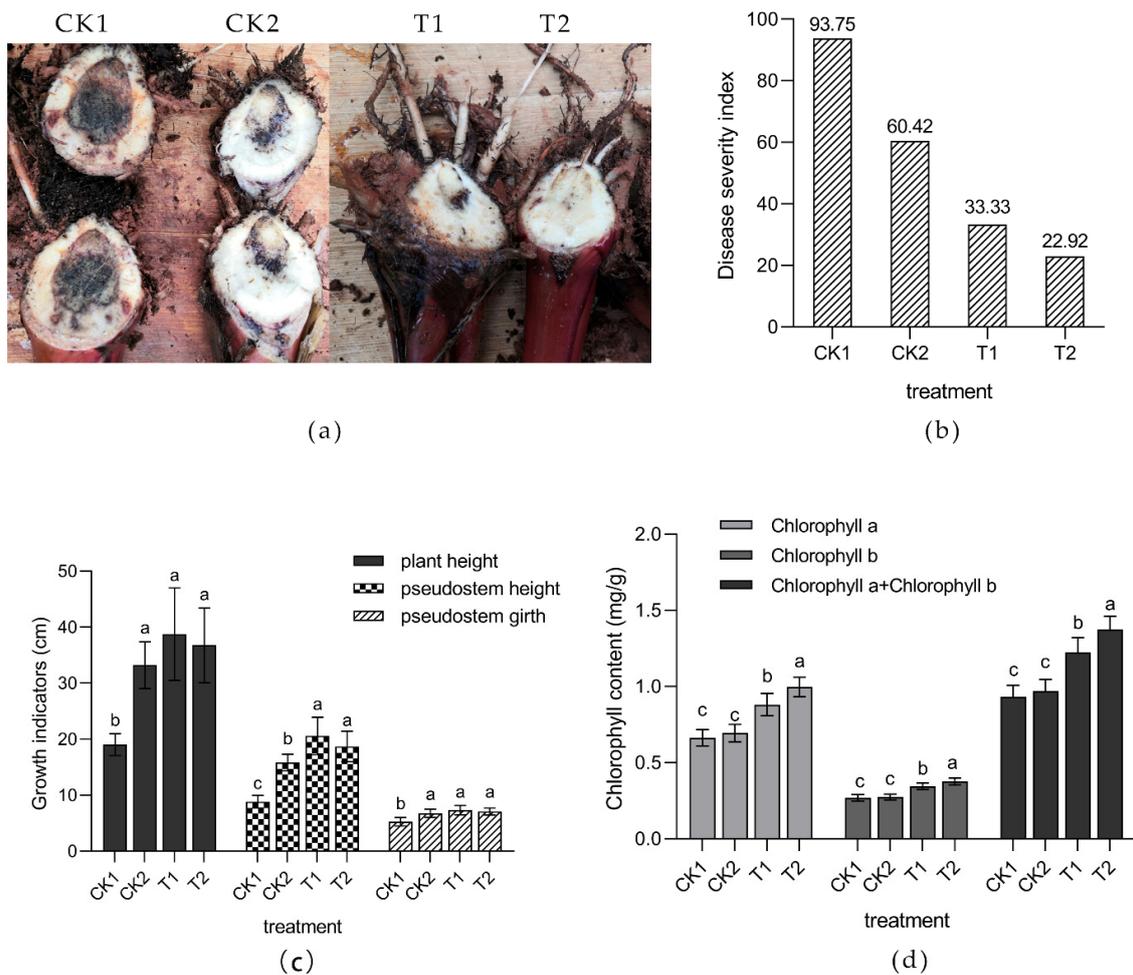


Figure 8. The effects of LG14-3 strain on banana Fusarium wilt disease. (a) the lesion of corms of CK1 (Foc), CK2 (LG14-3 + Foc), T1 (LG14-3 + glycine + Foc) and T2 (LG14-3 + citric acid +Foc); (b) disease severity index of banana Fusarium wilt disease, letters above the columns represent disease severity index; (c) after the 60 days of Foc4 inoculated, the plant height, pseudostem girth, and pseudostem height were measured. (d) after the 60 days of Foc4 inoculated, chlorophyll content was measured. (c,d) bars indicate standard errors from the means. Letters above the columns represent significant difference for each treatment according to Duncan's multiple range test ($p < 0.05$).

Overall, LG14-3 reduced the degree of banana wilt disease, and LG14-3 promoted the growth of plants under pathogenic stress (Table 4).

Table 4. Effects of chemoattractant on *Bacillus velezensis* LG14-3 suppressing of banana Fusarium wilt disease.

		CK1	CK2	T1	T2
Disease severity index		93.75	60.42	33.00	23.00
Growth indicators (cm)	Plant height	19.04 ± 1.93 b	33.21 ± 4.17 a	38.78 ± 8.25 a	36.75 ± 6.68 a
	Pseudostem height	8.86 ± 1.11 c	15.84 ± 1.48 b	20.61 ± 3.32 a	18.70 ± 2.73 a
	Pseudostem girth	5.28 ± 0.76 b	6.79 ± 0.75 a	7.35 ± 0.83 a	7.09 ± 0.62 a
Chlorophyll content (mg/g)	Chlorophyll a	0.66 ± 0.05 c	0.70 ± 0.06 c	0.88 ± 0.07 b	1.00 ± 0.06 a
	Chlorophyll b	0.27 ± 0.02 c	0.28 ± 0.02 c	0.35 ± 0.02 b	0.38 ± 0.02 a
	Chlorophyll a + Chlorophyll b	0.93 ± 0.08 c	0.97 ± 0.08 c	1.23 ± 0.09 b	1.37 ± 0.09 a

Note: Values are the means ± standard errors. Letters represent significant differences for each treatment according to Duncan's multiple range test ($p < 0.05$).

4. Discussion

Microorganisms that grow in the rhizosphere are ideal as biocontrol agents [38]. In this study, *Bacillus velezensis* LG14-3, which has a significant chemotactic effect on banana root exudates, was isolated from the healthy rhizosphere of banana in the banana Fusarium wilt disease epidemic area. Auxin (indole-3-acetic acid (IAA), which can promote plant growth, was the first hormone identified in plants [39]. The research by Patten et al. showed that bacterial IAA stimulated the development of the host plant root system [40]. For a long time, the competition of siderophores produced by bacteria for iron has been considered one of the important mechanisms by which biocontrol bacteria antagonize plant pathogens. *Bacillus subtilis* CAS15, which produces a siderophore, can effectively control Fusarium wilt and has the ability to promote growth [38]. *Pseudomonas aeruginosa* A7 produces siderophores combined with insoluble iron ions, thereby inhibiting the growth of plant pathogens and helping to control plant diseases [41]. Nitrogen (N) is an essential element for microbial and plant life [42]. Biological nitrogen fixation is the process by which nitrogen-fixing microorganisms reduce atmospheric nitrogen to ammonia available to bacteria and plants [43]. Our research results showed that LG14-3 has the ability to produce IAA and iron carriers and to perform nitrogen fixation, so it is a biocontrol strain with good potential.

Chemotaxis and biofilm formation determine the strain colonization in plant roots [13]. Malate and citric acid are the main chemoattractants of the tomato rhizosphere bacterium WCS365 [29]. *Pseudomonas* B4 experienced chemotaxis to biphenyls and benzoates and used them as carbon sources [11] 1-Aminocyclopropane-1-carboxylarboxylate is a strong chemoattractant for the plant-beneficial rhizobacterium *Pseudomonas putida* UW4 [44]. In this study, LG14-3 showed chemotaxis to the root exudate of banana and its common components (citric acid, glycine, succinate, D-galactose, and D-maltose). In previous studies, organic acids and sugars have been reported to induce and enhance the formation of bacterial biofilms in the rhizosphere [14]. In this study, glycine (0.25 mM) and citric acid (0.25 mM) enhanced the biofilm formation of LG14-3. Banana root exudates were also found to improve the swarming motility of LG14-3. Similarly, research by Tan et al. showed that five organic acids significantly increased the swarming ability of T-5 [13].

Several studies have shown that root exudates and organic acids can promote the colonization of *Bacillus* in roots [13,17,45,46]. As with the above results, LG14-3 has a better colonization advantage within banana roots than the other two biocontrol strains. Moreover, the addition of exogenous glycine or citric acid can increase the colonization of LG14-3 in the banana rhizosphere. Some carbon metabolites in plant root exudates can serve as food and as an energy source for rhizosphere microorganisms [5,47,48]. Similarly, citric acid supports the growth of LG14-3.

Bacillus velezensis LG14-3 showed broad-spectrum antagonistic activity against *Xanthomonas campestris* pv. *mangiferaeindicae*, *Pestalotiopsis*, *Diaporthe*, *Gliomastix muro-rum*, and *Fusarium oxysporum* f. sp. *cubense*. The in vitro inhibition rate against Foc4 was more than 70%, and its metabolites also had antagonistic activity. Biocontrol using antagonistic microorganisms is a safer alternative to reduce the use of chemicals in agriculture, and it is considered a promising approach for the management of soil-borne diseases [49]. A large number of bacterial and fungal strains, as well as viruses, nematodes, and insects, have been employed as biological control agents (BCAs) in the management of soil-borne crop pathogens for decades [50], including *Bacillus*, *Pantoea*, *Streptomyces*, *Trichoderma*, *Clonostachys*, *Pseudomonas*, *Burkholderia*, and certain yeasts [51]. Among them, *Bacillus* is one of the most popular because it can form heat-resistant and dry-resistant endophytic spores [52], which are easy to store and apply [53]. *Bacillus* spp., especially *Bacillus subtilis*, *Bacillus cereus*, and *Bacillus amyloliquefaciens*, are effective for the control of plant soil-borne diseases and foliar and postharvest disease [38]. Both *Bacillus amyloliquefaciens* HSB1 and FZB42 were found to inhibit fungal mycelial growth in vitro and in plants and to promote the growth of wolfberry seedlings [49]. Rhizosphere *Bacillus* Y6 and F7 showed strong antagonistic activity against *Fusarium oxysporum* under laboratory and greenhouse

conditions, and their lipopeptides effectively reduced the incidence of banana Fusarium wilt [53].

The results of this study demonstrated that *Bacillus velezensis* LG14-3 was chemotactic to banana root exudates and that citric acid and glycine in root exudates enhanced the abilities of LG14-3 root colonization and pathogen inhibition. Sustaining chlorophyll content in plants upon pathogen invasion is vital, as it will permit plant cells to continue photosynthesis [54]. Moreover, pot experiments showed that LG14-3 was efficient in increasing the chlorophyll a and chlorophyll b contents of bananas, which may be one of the reasons for the increase in plant height and weight after the LG14-3 application. In conclusion, these results demonstrate the biocontrol potential of *Bacillus velezensis* LG14-3 and provide data for exploring the role of root exudates in the interaction between biocontrol *Bacillus*, plants and soil-borne pathogens.

5. Conclusions

In this study, LG14-3 was isolated from the healthy rhizosphere of banana in the banana Fusarium wilt disease epidemic area and identified as *Bacillus velezensis* by morphological and molecular biology. Our research results also showed that LG14-3 has the ability to produce IAA and iron carriers and perform nitrogen fixation. A chemotaxis assay showed that LG14-3 had significant chemotactic effects on banana root exudates, glycine and citric acid. Swarming motility assays and biofilm formation assays showed that root exudates, glycine (0.25 mM) and citric acid (0.25 mM) enhanced the biofilm formation and swarming ability of LG14-3. *Bacillus velezensis* LG14-3 showed broad-spectrum antagonistic activity against *Xanthomonas campestris* pv. *mangiferaeindicae*, *Pestalotiopsis*, *Diaporthe*, *Glomastix murorum* and *Fusarium oxysporum* f. sp. *cubense*. Colonization assays and biocontrol experiments showed that citric acid and glycine in root exudates enhanced the abilities of LG14-3 root colonization and pathogen inhibition.

In summary, banana root exudates as chemoattractants were used to screen antagonistic microorganisms, and the colonization of *Bacillus velezensis* LG14-3 was promoted in banana roots through the dual induction of strong Chemoattractants (citric acid and glycine) and root exudates. *Bacillus velezensis* LG14-3 exerted their effective biocontrol effect, and thus targeted the control of banana wilt. Therefore, the use of antagonistic microorganisms and chemoattractants to control banana Fusarium wilt disease against soil-borne pathogens is a preferable approach. This study provides a new idea and method to optimize the selection of antagonistic microorganisms to control plant diseases.

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