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Article

# Nematicidal Activity of *Moringa oleifera* Water Extracts and a Novel Rhizobacterium for the Suppression of *Meloidogyne incognita* in Tomato

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**Abstract:** Root-knot nematodes (*Meloidogyne* spp.) are major pests of vegetable crops, causing significant yield losses in numerous species worldwide. Their widespread distribution and polyphagous nature make them particularly challenging to manage effectively. The use of biological agents and botanical extracts has emerged as a sustainable alternative to chemical nematicides. The present study evaluated the effectiveness of aqueous extracts from *Moringa oleifera* and a novel strain of plant growth-promoting rhizobacteria, *Bacillus australimaris* strain LWD73, native to Florida, in reducing *Meloidogyne incognita* infection and enhancing plant growth. The combined application of *M. oleifera* extracts and *B. australimaris* LWD73 significantly reduced the number of galls, eggs per root system, second-stage juveniles (J2) per 60 cc of soil, and the nematode reproduction factor. Moreover, it improved key growth parameters in tomato cultivar HM1824. Fourier Transform Infrared Spectroscopy (FTIR) and Gas Chromatography-Mass Spectrometry (GC-MS) analyses of *M. oleifera* extracts identified antimicrobial and nematicidal phytochemicals. Similarly, solid-phase microextraction GC-MS (SPME-GC/MS) analysis of *B. australimaris* LWD73 revealed the presence of volatile organic compounds (VOCs), including 2-nonanone and 2-undecanone, which are known for their potent nematicidal properties. The synergistic application of *M. oleifera* and *B. australimaris* LWD73 demonstrated enhanced nematode suppression and improved plant growth, offering a promising, eco-friendly approach to managing root-knot nematodes in vegetable crops.

**Keywords:** root-knot nematodes; biological control; *Meloidogyne incognita*; *Moringa oleifera*; rhizobacteria; bacterial VOCs; FTIR; GC-MS

## 1. Introduction

Root-knot nematodes (*Meloidogyne* spp.) have a devastating impact on vegetable crops globally, causing significant yield losses and compromising overall plant health. These polyphagous nematodes have an extensive host range, infecting over 3,000 plant species [1,2]. Upon infection, they induce the formation of root galls, which disrupt normal root functions and impair the ability of plant to efficiently absorb water and nutrients from the soil. This leads to stunted growth and reduced crop yields. Moreover, root-knot nematode infections often predispose plants to secondary infections by soil-borne pathogens, exacerbating the damage and further diminishing plant health [3,4].

The severity of the damage caused by root-knot nematodes depends on various factors, including environmental conditions, nematode population density, species virulence, and the

susceptibility of the host plant. Managing these nematodes is particularly challenging due to their high reproductive potential and remarkable adaptability [5]. Although chemical nematicides have proven effective in controlling root-knot nematodes, their adverse environmental impacts have driven a shift toward developing alternative, eco-friendly management strategies [6,7].

The use of biological control agents, primarily bacteria and fungi, offers a promising and sustainable approach to suppressing nematode populations [5]. Likewise, plant-based natural products have also been reported to be effective against a wide range of pathogens [8–12]. Botanical management of root-knot nematodes using plant-based formulations has also demonstrated significant potential. Incorporating macerated leaves of *Cannabis sativa* and *Azadirachta indica* into the soil significantly reduced *Meloidogyne javanica* populations and infection severity in peach plants in a dose-dependent manner [13]. Similarly, the combined application of dried neem leaves and the fungus *Trichoderma harzianum* on tomato plants resulted in a greater reduction in root galls, egg mass production, and egg hatching compared to their individual applications. These combined effects led to increased plant height and fresh shoot weight, with outcomes strongly influenced by both dosage and exposure time [14].

Plant growth-promoting rhizobacteria (PGPR) play a vital role in mitigating plant stress and enhancing plant growth, either directly or indirectly. PGPR offer numerous benefits, including increased mineral availability, production of phytohormones, mitigation of heavy metal stress, and biocontrol of plant pathogens. These attributes contribute to improved nutrient uptake, enhanced plant growth, and higher crop yields [15–17].

For sustainable rice and sugarcane production, bacteria belonging to the genera *Bacillus*, *Rhizobium*, *Comamonas*, *Cyanobacteria*, *Nodosilinea*, *Levinella*, and *Pseudomonas* have been identified as efficient producers of nitrogen and solubilizers of inorganic phosphate, potassium, and other macro- and micronutrients [15].

Moreover, a consortium of *Bacillus cereus* AR156, *Bacillus subtilis* SM21, and *Serratia* sp. XY21 has shown significant efficacy in reducing the severity of root-knot disease while also increasing the yield and fruit quality of cucumber in field trials [18].

Innovative approaches, such as gene editing to confer resistance in crops against nematode infestations, the exploration of bacteria, fungi, and botanicals as potential natural nematicides, and the implementation of cultural practices, are all sustainable strategies for suppressing nematode populations in soil without relying on chemical nematicides [5]. These methods aim to minimize environmental impact while effectively managing plant-parasitic nematodes.

*Moringa oleifera* has emerged as a promising natural alternative for managing root-knot nematodes. The bioactive compounds in *M. oleifera*, such as alkaloids, flavonoids, and phenolic compounds, exhibit strong nematicidal properties, disrupting the life cycle of nematodes by impairing egg hatching and larval development. Studies have shown that both aqueous and methanolic extracts of *M. oleifera* leaves and seeds significantly reduce nematode population densities in the soil and gall formation on roots. Furthermore, its use improves plant vigor and yield by minimizing nematode-induced stress. When combined with other biocontrol agents, such as beneficial rhizobacteria, *M. oleifera* further enhances the suppression of root-knot nematodes, providing a synergistic and eco-friendly approach to nematode management in sustainable agriculture.

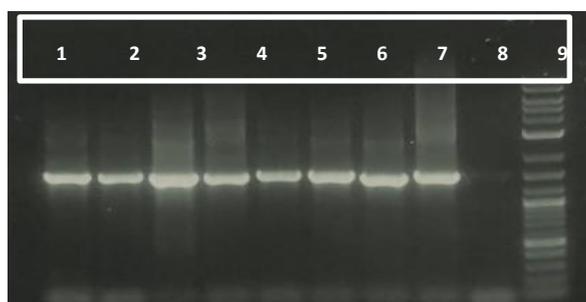
The current study was, therefore, conducted to evaluate the nematicidal potential of *M. oleifera*, both individually and in combination with various plant-growth-promoting rhizobacteria. The primary objective was to identify an effective and eco-friendly solution for managing plant-parasitic nematodes, thereby contributing to sustainable agricultural practices.

## 2. Results

### 2.1. Molecular Characterization of Rhizobacteria

The molecular characterization of rhizobacterial isolates was conducted using the 16S rRNA gene region amplified with universal primers 27F and 1492R. The amplification successfully yielded amplicons of 1500 base pairs, as visualized in gel electrophoresis (Figure 1). This fragment size corresponds to the expected region of the 16S rRNA gene, widely used for bacterial identification and phylogenetic studies.

The sequencing data obtained from the amplified products were compared against the National Center for Biotechnology Information (NCBI) database using the BLAST algorithm to confirm the identity of the isolates at the species level. The rhizobacterial isolates, along with their respective GenBank accession numbers, are detailed in Table 1. The high sequence similarity (>99%) with reference sequences in the database validated the taxonomic assignment of the isolates.



**Figure 1.** Molecular validation of rhizobacterial isolates using 16S rRNA Primers. Lane 1-8 represents the rhizobacterial isolates, 9 as control and L represents 1kb Ladder.

**Table 1.** Rhizobacteria Identity with Accession Numbers.

Sr. No.	Isolate	Rhizobacteria Identified as	Accession No.
1	W2	<i>Bacillus australimaris</i> strain LWD73	<a href="#">OQ366704</a>
2	W13	<i>B. cereus</i> strain HR001	<a href="#">OQ372951</a>
3	W44	<i>B. thuringiensis</i> strain WAG41	<a href="#">OQ370579</a>

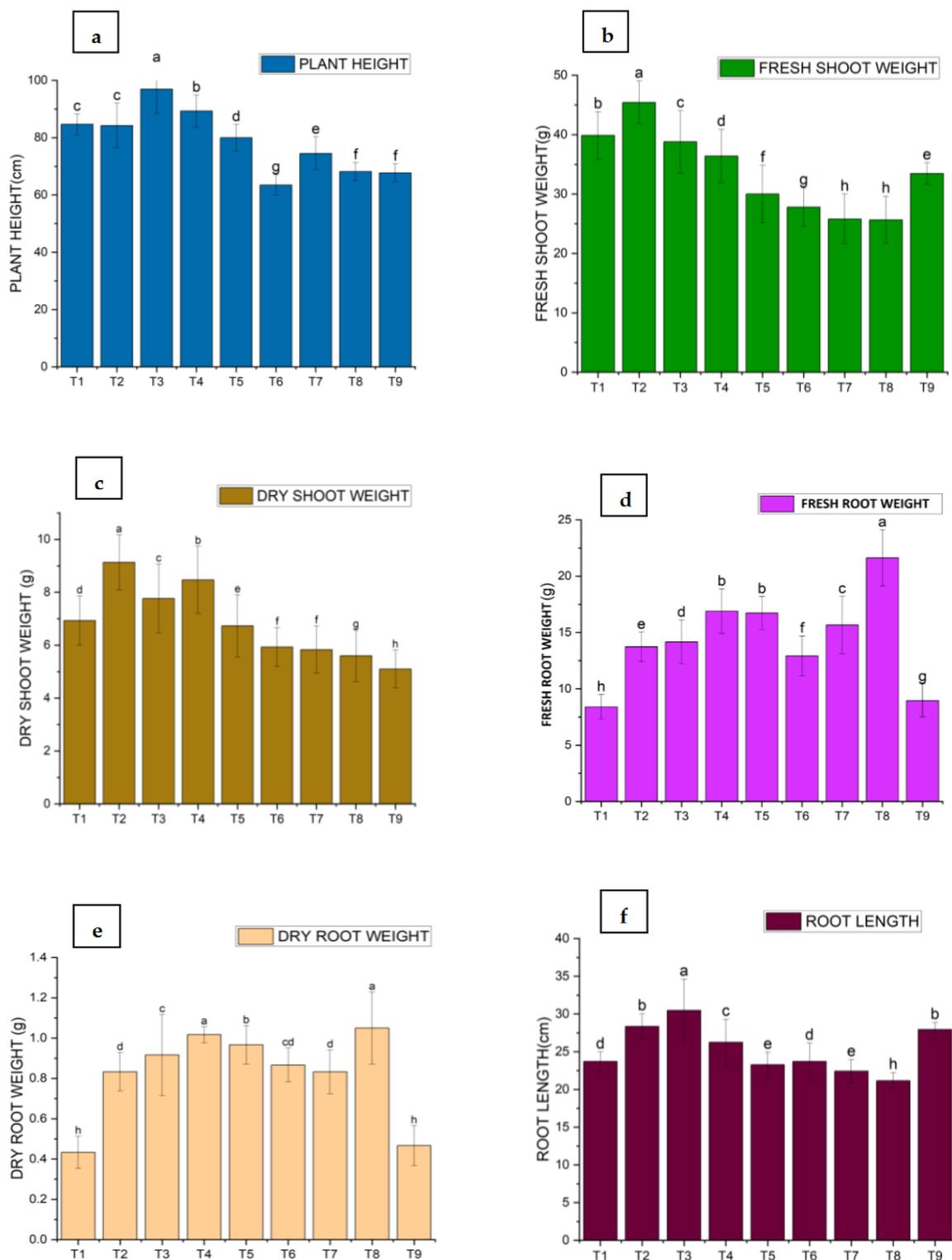
### 2.2. Effect of Rhizobacteria and Plant Extracts on Plant Growth of Tomato

The effect of different treatments of *M. oleifera*, both singly and in combination, was evaluated against *M. incognita* on plant growth parameters (plant height, root length, fresh shoot weight, dry shoot weight, fresh root weight, dry root weight) and *M. incognita* infection parameters (root gall ratings, eggs per root system, J2/60cc soil, reproduction factor).

Plant height and root length were significantly increased in nearly all treatments compared to the control (Figure 2). The maximum plant height and root length were recorded as 96.94 cm and 30.48 cm, respectively, in T3 (BA-LWD73 plus *M. oleifera*), while the lowest plant height and root length (63.50 and 22.44 cm, respectively) were observed in T7 (BT-WAG41 plus *M. oleifera*).

For fresh shoot weight and dry shoot weight, the maximum fresh shoot weight and dry shoot weight were observed in T2, which is the single treatment of BA-LWD73, with values of 45.43 g and 9.13 g, respectively. The lowest fresh shoot weight was recorded in T7 (BT-WAG41 and *M. oleifera* combined treatment) at 25.80 g, while the lowest dry shoot weight (5.93 g) was recorded in both T6 and T7 (BT-WAG41 single and combined with *M. oleifera*).

Fresh root weight and dry root weight were significantly lower in all treatments compared to the control. The minimum fresh root weight was 8.40 g in T1, which is the single treatment of *M. oleifera*. The minimum dry root weight was 5.83 g in T7, which is the combined treatment of BT-WAG41 and *M. oleifera* (Figure 2).

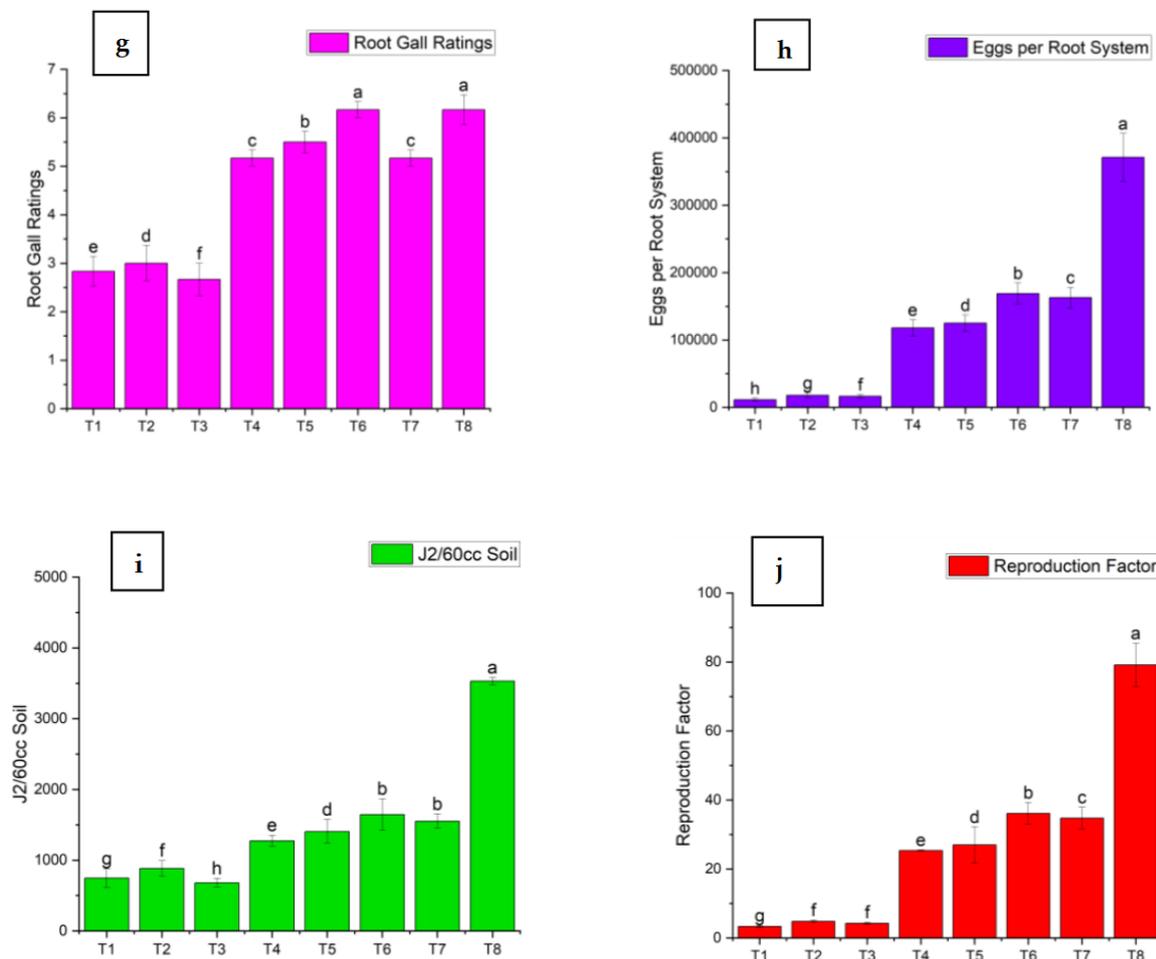


**Figure 2.** Effect of *M. oleifera* and rhizobacteria on Plant Growth Parameters of Tomato: (a) Plant Height; (b) Fresh Shoot Weight; (c) Dry Shoot Weight; (d) Fresh Root Weight; (e) Dry Root Weight and (f) Root Length.

### 2.3. Effect of Rhizobacteria and Plant Extracts on *M. incognita* Infection

Root gall ratings were significantly lower in most of the treatments compared to the positive control (Figure 3). The minimum root gall rating was 2.67 in T3, which is the combined treatment of BA-LWD73 and *M. oleifera*, while T8 (*M. incognita* only, control) showed the highest root gall rating. The single and combined applications of *M. oleifera* and BA-LWD73 resulted in 11,500, 18,000, and

16,500 eggs per root system, respectively. The lowest numbers of second-stage juveniles were recovered from the single and combined applications of *M. oleifera* and BA-LWD73 (746.83, 884.33, and 680 J2/60 cm<sup>3</sup> of soil, respectively). Similarly, the lowest reproduction factors were observed in the single and combined applications of *M. oleifera* and BA-LWD73, with values of 3.35, 4.84, and 4.25, respectively (Figure 3).



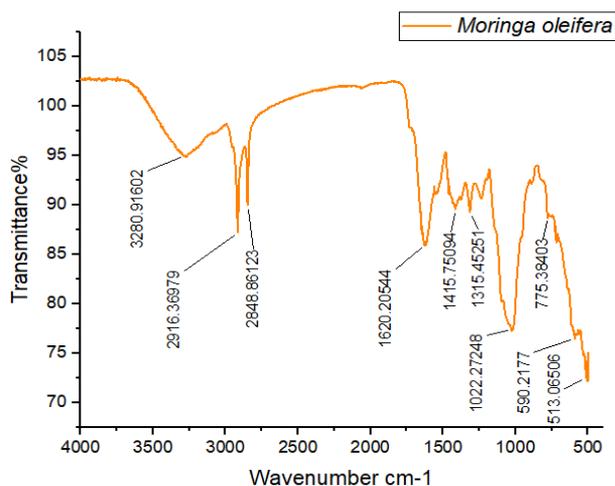
**Figure 3.** Effect of *M. oleifera* and rhizobacteria on Infection parameters of *M. incognita*: (a) Root Gall Ratings; (b) Eggs per Root System; (c) J2/60cc Soil and (d) Reproduction Factor. There was no nematode infection in T9.

#### 2.4. FTIR and GC-MS Analysis of Aqueous Extracts of *M. oleifera*

The FTIR spectra of *M. oleifera* are shown in Figure 4. The first and second peaks observed in the spectra were at wavenumbers 2916.36 cm<sup>-1</sup> and 2848.86 cm<sup>-1</sup>, indicating strong and broad stretching vibrations of hydroxyl groups, predominantly associated with alcohols. The third wavenumber, observed at 1620.20 cm<sup>-1</sup>, exhibited medium stretching and predominantly indicated the presence of a C=C functional group, typical of alkenes. The fourth wavenumber, at 1415.53 cm<sup>-1</sup>, showed strong stretching vibrations primarily due to S=O bonds, indicating the presence of sulfates. The fifth wavenumber, ranging from 1315.45 cm<sup>-1</sup> to 1022.27 cm<sup>-1</sup>, exhibited strong stretching vibrations attributable to C-O functional groups, predominantly signifying tertiary alcohols. The final wavenumber observed at 513.06 cm<sup>-1</sup> was characteristic of halo compounds.

The GC-MS chromatogram of the aqueous extract of *M. oleifera* revealed six distinct peaks, identified as six phytochemicals through comparison with the NIST library: 4(1H)-Pyrimidinone (3.02%), 2,6-dimethyl, Acetic acid [(aminocarbonyl)amino]oxo (3.01%), Maltol (9.58%), 4H-Pyran-4-one, 2,3-dihydro-3,5-dihydroxy-6-methyl (10.30%), 5-Hydroxymethylfurfural (71.76%), and 4(1H)-

Pyrimidinone, 2-(methylthio) (2.32%) (Table 2). All of these phytochemicals are reported to exhibit antimicrobial and nematicidal properties.



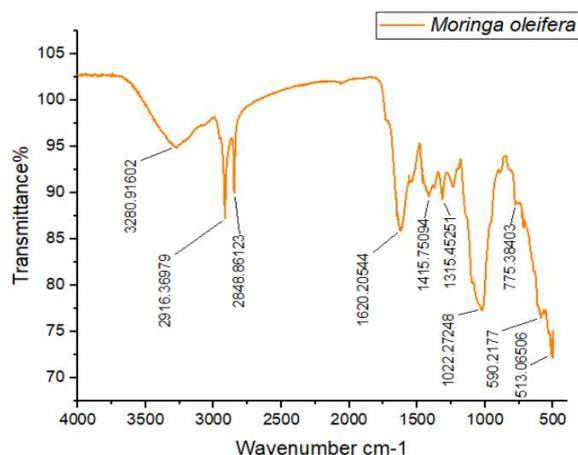
**Figure 4.** FTIR Spectra of *M. oleifera* leaves.

**Table 2.** Compounds identified from aqueous extract of *M. oleifera* L. through GCMS analysis.

Peak No.	Compound Name	Molecular Formula	Molecular Weight (g/mol)	Retention Time (Min)	Peak Area (%)
1	4(1H)-Pyrimidinone, 2,6-dimethyl	$C_{24}H_{30}N_2O$	362.5	5.310	3.02
2	Acetic acid, [(aminocarbonyl)amino]oxo	$C_3H_4N_2O_4$	132.08	5.561	3.01
3	Maltol	$C_6H_6O_3$	126.11	5.705	9.58
4	4H-Pyran-4-one, 2,3-dihydro-3,5-dihydroxy-6-methyl	$C_6H_8O_4$	144.12	6.155	10.30
5	5-Hydroxymethylfurfural	$C_6H_6O_3$	126.11	7.636	71.76
6	4(1H)-Pyrimidinone, 2-(methylthio)-	$C_5H_6N_2OS$	142.18	11.364	2.32

### 2.5. Detection of VOCs Using SPME-GCMS

Volatile organic compounds of *B. australimaris* strain LWD-73 were identified based on comparison of GC-MS results with the NIST library (Figure 5, Table 3). Four peaks were detected including 2-Nonanone (6.54%), 1H-Indole (87.46%), Tetradecanol (0.99%) and 9-Hexadecenoic acid (5.02%).



**Figure 5.** HS-SPME GC-MS chromatogram profiles of VOCs emitted from *B. australimaris* LWD-73.

**Table 3.** Volatile Organic Compounds of *B. australimaris* strain LWD-73 detected by solid phase micro extraction gas chromatography.

No.	Compound	Molecular Formula	Retention Time (min)	Molecular Weight (g/mol)	Peak Area	Percent
1	2-Nonanone	<a href="#">C<sub>9</sub>H<sub>18</sub>O</a>	8.35	142.24	215618130	6.54
2	1H-Indole	<a href="#">C<sub>8</sub>H<sub>7</sub>N</a>	11.50	117.15	2882910162	87.46
3	Tetradecanol	<a href="#">C<sub>14</sub>H<sub>30</sub>O</a>	13.73	214.38	32564023	0.99
4	9-Hexadecenoic acid	<a href="#">C<sub>16</sub>H<sub>32</sub>O<sub>2</sub></a>	19.00	256.42	165322864	5.02

### 3. Discussion

Numerous plant species possess nematicidal constituents that suppress nematode populations and improve plant health. In this study, the aqueous extract of *Moringa oleifera*, alone and in combination with rhizobacterial strains, effectively reduced *Meloidogyne incognita* populations and enhanced plant growth. Treatments combining *M. oleifera* and *Bacillus australimaris* strain LWD-73 demonstrated superior performance compared to other treatments, significantly reducing nematode infection and promoting plant vigor. All parts of *M. oleifera* can be utilized as biopesticides due to their ability to suppress pathogens and enhance crop health [20,21].

*M. oleifera* contains diverse bioactive compounds, including aldehydes, flavonoids, alcohols, phenols, and terpenoids, which may act individually or synergistically to affect nematodes. These compounds disrupt nematode feeding and reproduction, inhibit egg hatching, and exhibit juvenile toxicity. However, the precise mechanisms underlying their nematicidal activity remain unclear [20,21].

Microbes also play a pivotal role in managing polyphagous root-knot nematodes in various crops and soil conditions. They enhance plant health, induce systemic resistance, and combat a wide range of biotic stresses. Rhizobacteria are particularly valuable for their contributions to nutrient uptake, phytohormone production, mitigation of heavy metal stress, and increased crop yields [22,23]. Certain bacterial volatile organic compounds (VOCs), such as dimethyl disulfide, 2-nonanone, 1H-indole, tetradecanol, and 9-hexadecenoic acid, exhibit strong nematicidal activity against RKN [22,24,25].

FTIR and GC-MS analyses of *M. oleifera* extracts confirmed the presence of flavonoids, alcohols, and phenols, which are known for their antimicrobial and nematicidal properties. The nematicidal activity of *M. oleifera* was positively correlated with the abundance of these phytochemicals [26,27].

Among the rhizobacteria tested, *B. australimaris* strain LWD-73 demonstrated the most potent effects against *M. incognita*. SPME-GC-MS analysis identified key VOCs produced by *B. australimaris*, including 2-nonanone, 9-hexadecenoic acid, tetradecanol, and 1H-indole, which possess strong antimicrobial, antifungal, and nematicidal properties. For instance, 2-nonanone exhibits significant nematicidal activity by inhibiting egg hatching, disrupting feeding, and reducing nematode populations, thereby limiting root damage over time [28,29]. This compound also shows antifungal activity against pathogens such as *Verticillium longisporum* and *Botrytis cinerea* [30,31].

Indole, another prominent VOC, disrupts nematode egg-laying and survival, induces oxidative stress at high concentrations, and triggers methuosis at low concentrations, ultimately causing nematode mortality [32,33]. Moreover, indole acts as a key chemical signal, promoting plant growth and influencing auxin signaling. For example, indole emitted by *Escherichia coli* has been shown to enhance the root architecture of *Arabidopsis thaliana* [25,34,35].

9-Hexadecenoic acid (palmitoleic acid) exhibits nematicidal activity by causing toxicity to larvae and eggs, acting as a repellent, and interfering with nematode feeding and reproduction. It may also stimulate plant defenses and influence soil microbial communities, offering a promising approach to integrated nematode management [36].

Tetradecanol, emitted by *Paenibacillus polymyxa* strain J2-4, has demonstrated strong fumigant activity against *M. incognita*, further highlighting the potential of VOCs in nematode management strategies [37]. These findings underscore the significance of plant-derived phytochemicals and microbial VOCs as sustainable tools for controlling nematodes and improving crop health.

## 4. Materials and Methods

### 4.1. Inoculum Preparation of Nematode

The *Meloidogyne incognita* culture was initiated from a single egg mass on a susceptible tomato host, cv. HM1824, and identified as *M. incognita* based on perineal pattern morphology [38]. Small pieces of infected roots were vigorously shaken in a tightly sealed container with 0.5% sodium hypochlorite (NaOCl) solution to release the eggs from the egg masses. The resulting suspension was filtered through a 200-mesh sieve to remove root debris and then through a 500-mesh sieve to collect the eggs, following the described method. The inoculum density was quantified. The eggs were rinsed three times with tap water to remove residual bleach. Freshly hatched second-stage juveniles were obtained by incubating the eggs in extraction trays at 25°C for 48 hours. These juveniles were used for the experiment.

### 4.2. Preparation of Aqueous Extracts of *M. oleifera*

Fresh leaves of *M. oleifera* were carefully washed, blended in a Waring blender with sterile water, and left to stand for 12 hours. The mixture was then filtered sequentially: first through muslin cloth, followed by Whatman filter paper No. 1, and finally through a Millipore filter. Different concentrations (0% to 100%) of the extract were prepared by diluting the standard extract with the requisite amount of distilled water [21].

### 4.3. Isolation and Purification of Rhizobacteria

Soil samples were collected from the rhizosphere of tomato plants by carefully uprooting the plants and gently shaking the roots to remove excess soil. The rhizobacteria were recovered using the serial dilution method [26]. Dilutions of 10<sup>-5</sup>, 10<sup>-6</sup>, and 10<sup>-7</sup> were spread onto sterile Petri plates containing solidified autoclaved nutrient agar (NA). The Petri plates were sealed with parafilm and incubated at 26 ± 2°C. After 24 hours, individual bacterial colonies were purified on NA using the streaking method.

### 4.4. Molecular Characterization of Rhizobacteria Isolates

Genomic DNA was extracted from rhizobacterial isolates using a DNA extraction kit. The extracted DNA was then used for Polymerase Chain Reaction (PCR) analysis. For amplification, the universal primers 27F (5'-AGAGTTTGATCMTGGCTCAG-3') and 1492R (5'-GGTACCTTGTTACGACTT-3'), targeting the 16S rRNA molecular marker, were employed [19]. PCR amplification was conducted under the following conditions: initial denaturation at 94°C for 3 min; followed by 35 cycles of 94 °C for 40 s, 60°C for 50 s, and 72°C for 1 min; with a final extension at 72°C for 10 min

The DNA bands were visualized under UV light, and images were captured using a gel documentation system. The amplified DNA was purified using a Qiagen DNA purification kit and quantified with a NanoDrop spectrophotometer. Finally, the amplified products were sent to Eurofins Genomics, USA, for sequencing.

### 4.5. Sub-Culturing and Preparation of Rhizobacteria Suspensions

The identified rhizobacterial strains were initially cultured on NA plates and subsequently transferred to Luria Bertani broth. The cultures were incubated at 25°C in a rotary shaker for 48 hours to facilitate mass culturing. Cell suspensions of the rhizobacterial strains were prepared by measuring

optical density (OD) at 600 nm using a spectrophotometer. The OD was adjusted to 1.0, equivalent to 109 CFU/mL. The cell suspensions were centrifuged at 5000 rpm for 15 minutes at 4°C, after which the supernatant was discarded. The resulting pellet was gently washed with autoclaved distilled water. Following the wash, additional autoclaved distilled water was added to the falcon tube, and the contents were mixed thoroughly using a vortex mixer. The final concentration of each rhizobacterial suspension was adjusted to 109 CFU/mL using a spectrophotometer [39].

#### 4.6. Effect of Rhizobacteria and Plant Extracts on Plant Growth and *M. incognita* Infection

The soil drench method was employed to assess the effects of rhizobacteria and plant extracts on plant growth parameters and *M. incognita* infection. To inoculate the plants, 5 mL of suspension containing 5000 freshly hatched second-stage juveniles of *M. incognita* was applied to 21-day-old tomato plants by creating three small holes around each plant. After 48 hours, 50 mL of rhizobacterial suspension (109 CFU/mL) and 50 mL of an aqueous extract of *M. oleifera* were applied as a soil drench. Six replicates were conducted for each treatment, with water serving as the control. For the control treatment, 5 mL of *M. incognita* filtrate was applied to ensure a consistent environment for all plants.

After 45 days, the plants were uprooted to evaluate growth and infection parameters. Plant growth parameters included plant height, tap root weight, dry root weight, shoot length, fresh shoot weight, dry shoot weight, and root length. The *M. incognita* infection parameters assessed were root gall ratings, eggs per root system, J2 population per 100 cc of soil, and the nematode reproduction factor.

**Table 4.** Treatments for the Effect of Rhizobacteria and Moringa Leaf Extracts on Plant Growth Attributes and *M. incognita* infection.

T1	<i>M. oleifera</i> + <i>M. incognita</i>
T2	<i>Bacillus australimaris</i> LWD73 + <i>M. incognita</i>
T3	<i>B. australimaris</i> LWD73 + <i>M. oleifera</i> + <i>M. incognita</i>
T4	<i>Bacillus cereus</i> HR001+ <i>M. incognita</i>
T5	<i>B. cereus</i> HR001 + <i>M. oleifera</i> + <i>M. incognita</i>
T6	<i>Bacillus thuringiensis</i> WAG41 + <i>M. incognita</i>
T7	<i>B. thuringiensis</i> WAG41 + <i>M. oleifera</i> + <i>M. incognita</i>
T8	<i>M. incognita</i> only
T9	Healthy Control

#### 4.7. Detection of Volatile Organic Compounds from Rhizobacteria

*B. australimaris* LWD73 was cultured overnight at 37°C in LB broth. The bacterial culture was adjusted to an optical density (OD) of 0.132 at 600 nm (equivalent to 0.5 McFarland units) using a spectrophotometer. A 5 mL aliquot of the bacterial culture was transferred to a headspace (HS) vial, and an SPME fiber was inserted into the vial. The HS vial containing the SPME fiber was placed in a water bath at 60 °C for 60 minutes. After incubation, the SPME fiber was immediately exposed to the hot GC injection port for 28 minutes. Each treatment was replicated nine times [40].

#### 4.8. Compound Identification from Plant Extracts

Nematicidal compounds were identified from aqueous extracts of *M. oleifera* by Fourier Transform Infrared Spectroscopy [34] and Gas Chromatography and Mass Spectrometry [35].

#### 4.9. Statistical Analysis

The data obtained from egg hatch inhibition and juvenile mortality was analyzed by 3-factor factorial in CRD and the data obtained from plant growth parameters and infection parameters was analyzed by 2-factor factorial in CRD. The means were separated by Fischer's unprotected LSD test, Gen Stat statistical software was used at significance level ( $P < 0.05$ ).

## 5. Conclusions

The study identified a new bacterial strain, *Bacillus australimaris* strain LWD-73, with significant nematicidal activity, offering potential for the effective management of *Meloidogyne incognita*. The integration of this bacterial strain with moringa leaf extract has demonstrated compatibility, enhancing the efficacy of rhizobacteria for nematode control. This combination has yielded improved results in reducing nematode populations. Both moringa and the selected rhizobacteria are rich in phytochemicals with potent nematicidal and antimicrobial properties. These bioactive compounds provide promising solutions for nematode management and can be incorporated into integrated pest management strategies. Small-scale vegetable growers can benefit from using aqueous extracts of moringa leaves as a cost-effective and accessible method for nematode control, contributing to the improved health and productivity of vegetable crops.

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