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The potential of *Bacillus* species isolated from *Cinnamomum camphora* for biofuel production

Noura Sh.A. Hagaggi^{1*}  and Eman A. El Rady²

Abstract

Background Increasing concerns about climate change and global petroleum supply draw attention to the urgent need for the development of alternative methods to produce fuels. Consequently, the scientific community must devise novel ways to obtain fuels that are both sustainable and eco-friendly. Bacterial alkanes have numerous potential applications in the industry sector. One significant application is biofuel production, where bacterial alkanes can serve as a sustainable eco-friendly alternative to fossil fuels. This study represents the first report on the production of alkanes by endophytic bacteria.

Results In this study, three *Bacillus* species, namely *Bacillus atrophaeus* Camph.1 (OR343176.1), *Bacillus spizizenii* Camph.2 (OR343177.1), and *Bacillus aerophilus* Camph.3 (OR343178.1), were isolated from the leaves of *C. camphora*. The isolates were then screened to determine their ability to produce alkanes in different culture media including nutrient broth (NB), Luria–Bertani (LB) broth, and tryptic soy broth (TSB). Depending on the bacterial isolate and the culture media used, different profiles of alkanes ranging from C₈ to C₃₁ were detected.

Conclusions The endophytic *B. atrophaeus* Camph.1 (OR343176.1), *B. spizizenii* Camph.2 (OR343177.1), and *B. aerophilus* Camph.3 (OR343178.1), associated with *C. camphora* leaves, represent new eco-friendly approaches for biofuel production, aiming towards a sustainable future. Further research is needed to optimize the fermentation process and scale up alkane production by these bacterial isolates.

Keywords Endophyte, *Bacillus*, *Cinnamomum camphora*, Alkane, Biofuel, Production

Background

Fossil fuels have been used for decades to produce liquid fuels such as diesel, gasoline, and kerosene. However, it is predicted that petroleum reserves will be depleted within 40 years [1]. This has raised concerns about the global petroleum supply, and environmental issues such as

global warming and climate change. As a result, there is growing interest in exploring alternative fuel sources [2]. Consequently, a significant focus has been on developing alternative biosynthesis methods for sustainable and eco-friendly biofuel.

Alkanes are hydrocarbons that are essential for biofuel production. They are the key building blocks of renewable biodiesel. Alkanes offer a sustainable alternative to fossil fuels, aligning with global efforts to reduce climate change and improve environmental sustainability [3]. The stable chemical structure of alkanes also helps biofuels retain their quality and performance in long-term storage. This stability is critical for meeting the requirements of individual consumers as well as

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commercial and industrial sectors [4]. The traditional commercial production of alkanes constantly increases production costs, non-renewable energy consumption, and gaseous pollutants [5]. The microbial biosynthesis of alkanes can be a promising sustainable alternative for chemical production [6]. Incorporating microorganisms into the global future of green energy can achieve a distributed and sustainable supply chain that is safe, reliable, and responsive to ever-changing global demand [7, 8].

The biosynthesis of alkanes by bacteria has attracted significant attention in recent years due to its potential for biofuel production [9]. Bacterial alkanes possess desirable properties, including high energy content and low freezing points compared to other biofuel sources, making them well-suited for specific applications in the aviation or automotive industries [10].

Cinnamomum camphora (L.) J. Presl., popularly known as the camphor tree, is a member of the Lauraceae family. It is native to China, Korea, and Japan and is extensively cultivated in Asia, Africa, North America, and Australia [11]. Every part of the plant contains volatile organic compounds that have medicinal properties [12]. Previous studies have shown that it is possible to utilize endophytes that inhabit plant tissues to synthesize compounds similar to those produced naturally by the host plants [13]. This approach avoids the risk of over-harvesting or the negative effects of climate change on plants, which can affect the production of these compounds [14]. Although alkane biosynthesis has been recognized in various microorganisms, including cyanobacteria, genetically modified bacteria, yeasts, and fungi, no studies have reported the production of alkanes by endophytic bacteria [15]. Therefore, this study aims to isolate endophytic bacteria from *C. camphora* leaves and screen the production of alkanes by the isolates in different culture media. The study is an attempt to find renewable sources for bio-alkanes that may be promising for sustainable biofuel production.

Materials and methods

Plant material

Leaves from healthy trees of *C. camphora* were collected from Aswan City, Egypt (24° 5′ 20.1768″ N, 32° 53′ 59.3880″ E) and brought directly to the Aswan University bacteriology laboratory for the isolation of endophytic bacteria.

Isolation and identification of endophytic bacteria

The surfaces of the leaves were sterilized using 5% NaClO, 70% CH₃CH₂OH, and autoclaved distilled water, respectively [16]. In a 9 mL sterile saline solution, 1 g of the leaves was mashed well. One milliliter of the resulting

suspension was then inoculated in trypticase soy and nutrient agar plates. Plates were incubated for 72 h at 37 °C. Three isolates coded as Camph.1, Camph.2, and Camph.3 were subjected to molecular identification by partially sequencing their 16S rRNA genes. The amplification primers 27F and 1492R were used [17]. The separation of PCR products was performed using 1% (w/v) agarose gel. The sequencing of the obtained bands was commercially performed at SolGent Co., Korea. The sequence similarity and identity percentages were determined using the NCBI website (<https://www.ncbi.nlm.nih.gov/>). An accession number was gained for each isolate after submitting its 16S rRNA gene partial sequence into the NCBI database. The phylogenetic relationship among the present isolates and the other close members of NCBI was constructed using neighbor-joining analysis in MEGA X 10.1.7 software [18].

Determination of bacterial growth curves

The growth curves of the bacterial isolates grown in nutrient broth (NB), Luria–Bertani (LB) broth, and tryptic soy broth (TSB) were determined using the turbidimetric method [19]. In 250 mL conical flasks, 50 mL of each medium was prepared and autoclaved. The flasks were inoculated with 100 µL of each bacterial inoculum (1.5×10^8 CFU/mL, OD₆₀₀=0.1). The flasks were incubated at 37 °C under shaking (150 rpm). The optical density was read at 600 nm at intervals of 10 h until the stationary phase was reached. The flasks without bacterial inoculums served as controls. The experiment was conducted three times.

Fermentation conditions

The freshly prepared inoculum (100 µL of 1×10^7 CFU/mL) of each bacterial isolate was inoculated in 500 mL flasks containing 100 mL of three different broth culture media: NB, LB, and TSB. The flasks were incubated for 48 h at 37 °C and 150 rpm. Flasks containing media without bacterial inoculum were used as controls. Triplicates were made for all fermentations.

Extraction and GC/MS analysis of alkanes

Hexane was added to the bacterial cultures in a ratio of 1:1 (v/v) and homogenized well. The solvent layers were then separated and concentrated using a rotary evaporator at 40 °C and 130 rpm under pressure. Each extract (1 mg) was redissolved in 10 mL of hexane. The GC/MS system used in this study was the Agilent Technologies 7890A GC/5977A MSD supplied with a TR-5MS GC column (30 m, 0.25 mm ID, and 0.25 µm film). The sample (1 µL) was injected into the column, and the oven temperature was initially set at 30 °C for 1 min. The temperature was then increased at a rate of 10 °C/min until

reaching 200 °C, where it was held for an additional 1 min. The carrier gas, helium, was used at a flow rate of 20 mL/min. The retention times of the sample peaks were compared with NIST11.L standard reference compounds. The alkane standard mixture (C₇-C₄₀, Millipore Sigma™ Supelco™) was used to quantify the alkanes in the samples.

Effect of carbon sources on alkane production

In conical flasks, the basal medium consisted of the following components per liter: KH₂PO₄ (1.3 g), MgSO₄·7H₂O (0.2 g), NaCl (5 g), (NH₄)₂SO₄ (1 g), and yeast extract (5 g) was supplemented with different carbon sources including glucose, sucrose, and sugar cane molasses, each at a concentration of 10 g/L. Flasks were then inoculated with 100 µL of a freshly prepared inoculum containing 1×10⁷ CFU/mL. Flasks were incubated for 48 h at 37 °C and 150 rpm. Flasks containing media without bacterial inoculum were used as controls. Triplicates were prepared for all fermentations. Alkanes were extracted and analyzed using the method described above.

Results and discussion

The use of biofuels has become crucial in addressing the worldwide concerns of the energy crisis and climate change. Microbial alkanes provide a renewable, eco-friendly, and promising source for the sustainable production of biofuels [20]. Unlike fossil fuels, biofuels derived from microbial alkanes not only decrease carbon emissions but also mitigate the effects of global warming [21]. The low toxicity and biodegradability of bacterial

alkanes make them eco-friendly alternatives to synthetic alkanes, serving various applications [22].

Isolation and identification of endophytic bacteria

In this study, three endophytic bacteria were isolated from the leaves of *C. camphora* and coded as Camph.1, Camph.2, and Camph.3. Based on 16S rRNA gene sequence analysis, the isolates Camph.1, Camph.2, and Camph.3 were found to be quite similar to *Bacillus atrophaeus* (NR024689.1), *Bacillus spizizenii* (NR112686.1), and *Bacillus aerophilus* (NR042339.1), respectively (Fig. 1). The NCBI accession numbers of the isolates Camph.1, Camph.2, and Camph.3 are OR343176.1, OR343177.1, and OR343178.1, respectively. It was observed that the genus *Bacillus* was dominant among endophytes, this may be attributed to the ability of *Bacillus* spp. to form spores and tolerate extreme temperatures in the Aswan region. This finding agreed with previous studies which reported the isolation of *Bacillus* spp. from different plants grown in Aswan [23–25].

Determination of bacterial growth curves

From a commercial perspective, the growth of microorganisms is a significant challenge in the industrial production of valuable chemicals [26]. Therefore, the growth curve for each bacterial isolate was determined in each culture medium. It was observed that the exponential phase of the three isolates began after 20 h of incubation and extended until 50 h. The stationary phase continued for 20 h, after which the growth rate declined (Fig. 2). Generally, the growth rate was higher in

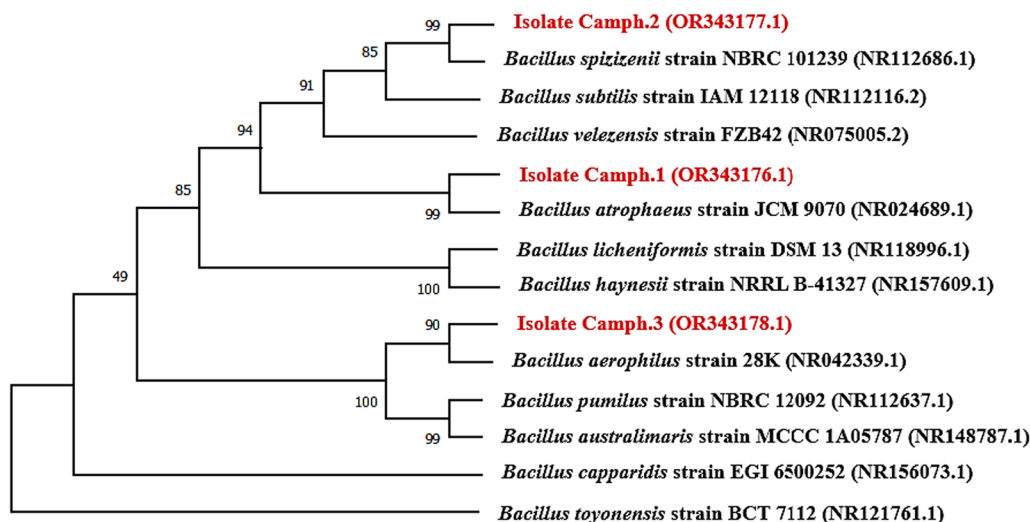


Fig.1 The phylogenetic relationship among the isolates Camph.1, Camph.2, Camph.3, and the closely related species from the NCBI database using the neighbor-joining method in MEGA X10.1.7 software

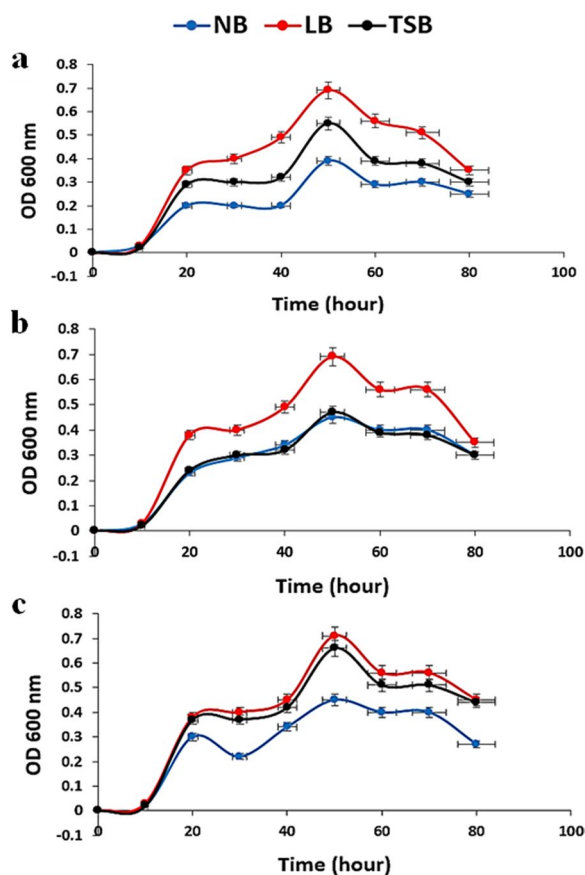


Fig. 2 The bacterial growth curves for *B. atrophaeus* Camph.1 (a), *B. spizizenii* Camph.2 (b), and *B. aerophilus* Camph.3 (c), in NB, LB, and TSB media. The bars represent the standard errors of the means

LB followed by TSB. The NB, on the other hand, had the lowest growth rate for all isolates (Fig. 2).

GC/MS analysis of alkanes

The production of alkanes was detected by GC/MS analysis after growing the bacterial isolates in three different culture media: NB, LB, and TSB. It was interesting to note that the alkane profiles vary depending on the growth medium and the bacterial strain. For *B. atrophaeus* Camph.1, the major number of alkanes was detected in the LB medium, where fourteen alkanes were evaluated, including Heptane-2,2,4,6,6-pentamethyl, Decane-2,4,6-trimethyl, Octadecane-1-iodo, Tetradecane, Tridecane-3-methyl, 10-Methylnonadecane, Hexacosane, Tetracosane, Eicosane-2-methyl, Undecane-2,9-dimethyl, Heptadecane-9-octyl, Heptadecane-2-methyl, Nonadecane-2-methyl, and 2-methyloctacosane. TSB medium contained ten alkanes, which were Nonane-2,2,3-trimethyl, Octane-2-methyl, Tetradecane-4-ethyl,

Decane-3-methyl, Heptadecane-2-methyl, Pentacosane, Octadecane, Hexacosane, Cyclobutane-1,2-diethyl, and Eicosane. NA medium contained seven alkanes, which included Heptane-2,2,4,6,6-pentamethyl, Decane-2,4,6-trimethyl, Hexadecane-3-methyl, Hexacosane, Octadecane-1-iodo, 1,3,5,7,9-Pentaethyl-1,9-dibutoxypentasiloxane, and Hentriacontane.

Fourteen alkanes were produced by *B. spizizenii* Camph.2 in LB medium (Table 5). In comparison, eleven alkanes were detected in both NB and TSB (Tables 4 and 6) and (Fig. 4). LB medium included Heptane-2,2,4,6,6-pentamethyl, Undecane-3,9-dimethyl, Decane-3,8-dimethyl, Tridecane-1-iodo, Hexadecane-2,6,11,15-tetramethyl, Pentacosane, Octadecane, Decane-3-methyl, Hexadecane, Hentriacontane, Eicosane, Heneicosane, Heptacosane, and 2-Bromo dodecane (Table 5). NB medium contained Nonane-2,2,3-trimethyl, Dodecane, Eicosane, Hexadecane, 2,2-Dimethyleicosane, Octacosane, Heptadecane-2-methyl, Hexadecane-8-hexyl-8-pentyl, 5-Ethyl-5-methylnonadecane, Cyclobutane-1,2-diethyl-trans, and Octane-2,5,6-trimethyl (Table 4). On the other hand, TSB medium included Heptane-2,2,4,6,6-pentamethyl, Undecane-4,7-dimethyl, Hexadecane, Heneicosane, Hexacosane, Hentriacontane, Heptadecane-2-methyl, Heptadecane-9-octyl, Octacosane, Octadecane-1-iodo, and Pentadecane-2-methyl and (Table 6).

On the other hand, *B. aerophilus* Camph.3 produced eleven alkanes when grown in NB medium: Heptane-2,2,4,6,6-pentamethyl, Decane-3,8-dimethyl, Eicosane, Hexacosane, Pentadecane, Tetracosane, Hexadecane, Heptadecane, Heneicosane, Hentriacontane, and Octacosane (Table 7 and Fig. 5). Fourteen alkanes were produced in both LB and TSB media: Heptane, 2,2,4,6,6-pentamethyl, 1-Iodo-2-methylnonane, Hexadecane, Tetradecane-2,6,10-trimethyl, 10-Methylnonadecane, Octacosane, Pentacosane, Heptacosane, Heptadecane, Heptadecane-2-methyl, Hentriacontane, Octadecane, Pentadecane-2-methyl, and Hexacosane (Table 8 and Fig. 4) and Heptane-2,2,4,6,6-pentamethyl, Nonane-4,5-dimethyl, Heptadecane-2-methyl, Eicosane, Dodecane-2,6,11-trimethyl, Heptacosane-1-chloro, Dodecane, Tetracosane, Hexadecane, Pentadecane, Hexacosane, Octacosane, Decane-3-methyl, and Decane-4-methylene (Table 9 and Fig. 5), respectively.

Interestingly, the profiles of alkanes released by the three bacterial isolates in the three tested culture media differed. For all isolates, the highest number of alkanes was detected in the LB medium (Fig. 3, 4 5). This finding aligns with previous studies that have reported a significant effect of medium composition on

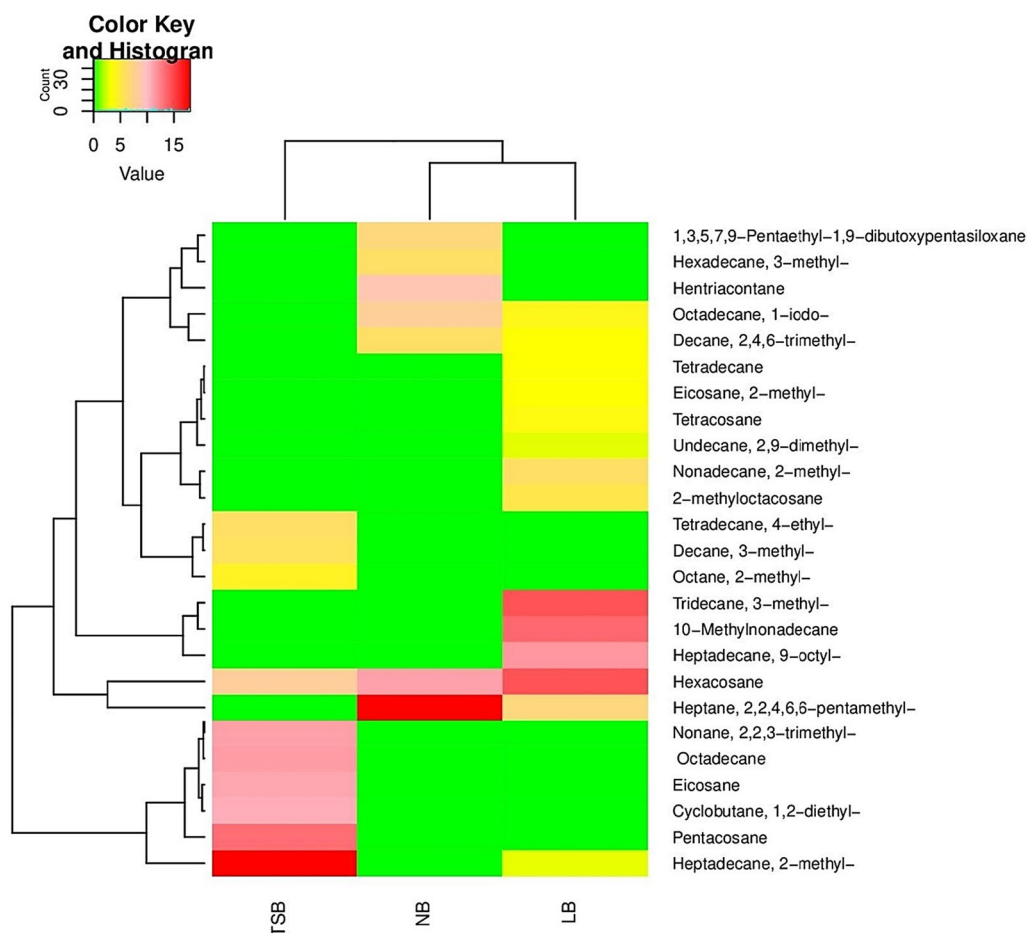


Fig. 3 The heatmap displays the amounts of bio-alkanes (mg alkane/L culture) produced by *B. atrophaeus* Camph.1 in NB, LB, and TSB media

the profiles of volatile organic compounds released by microorganisms [27].

Effect of carbon sources on alkane production

Interestingly, various alkanes were produced by the three bacterial isolates using glucose, sucrose, and sugar cane molasses as carbon sources. This is consistent with previous studies that reported significant differences in hydrocarbon profiles produced by microorganisms based on carbon sources [28]. *B. atrophaeus* Camph.1 produced fifteen different alkanes using glucose as a carbon source which are Tetradecane, 2,2-dimethyl (3.8 mg/L), Undecane, 2-methyl (2.38 mg/L), Eicosane (12.47 mg/L), Decane, 3,8-dimethyl (2.29 mg/L), Heptadecane, 4-methyl (12.03 mg/L), Hentriacontane (3.2 mg/L), Octadecane, 2-methyl (3.1 mg/L), Heneicosane (2.9 mg/L), Hexadecane (2.8 mg/L), Hexacosane (13.8 mg/L), Hexadecane, 2,6,10,14-tetramethyl (2.6 mg/L), Octacosane (7.39 mg/L), Pentacosane (6.08 mg/L), 2-methyloctacosane (2.15 mg/L), and Heneicosane, 3-methyl (2.85 mg/L). On the other

hand, the GC/MS analysis revealed a total of twelve alkanes produced by *B. atrophaeus* Camph.1 when grown in a medium supplemented with sucrose, which were Heptane, 2,2,4,6,6-pentamethyl (8.36 mg/L), Undecane, 3,7-dimethyl (4.08 mg/L), Decane, 2,9-dimethyl (4.15 mg/L), Decane, 2-methyl (13.19 mg/L), Heptadecane, 8-methyl (14.02 mg/L), Nonadecane, 3-methyl (3.74 mg/L), Hentriacontane (3.25 mg/L), Hexadecane (3.49 mg/L), Heneicosane (15.21 mg/L), 2-methyloctacosane (11.18 mg/L), Octacosane (7.57 mg/L), and Octadecane, 1-iodo (7.22 mg/L). Ten alkanes were detected in a medium supplemented with sugar cane molasses including Decane, 2,2,3-trimethyl (3.35 mg/L), Octadecane (2.76 mg/L), Heptadecane, 2-methyl (9.69 mg/L), Dodecane (9.23 mg/L), Eicosane (2.46 mg/L), Hexadecane (2.86 mg/L), Hexacosane (11.02 mg/L), Nonane, 4,5-dimethyl (2.19 mg/L), Heneicosane (8.96 mg/L), and 2,2-Dimethyleicosane (1.57 mg/L).

For *B. spizizenii* Camph.2, thirteen alkanes were produced in a glucose-based medium, which are Heptane, 2,2,4,6,6-pentamethyl (6.63 mg/L), Undecane,

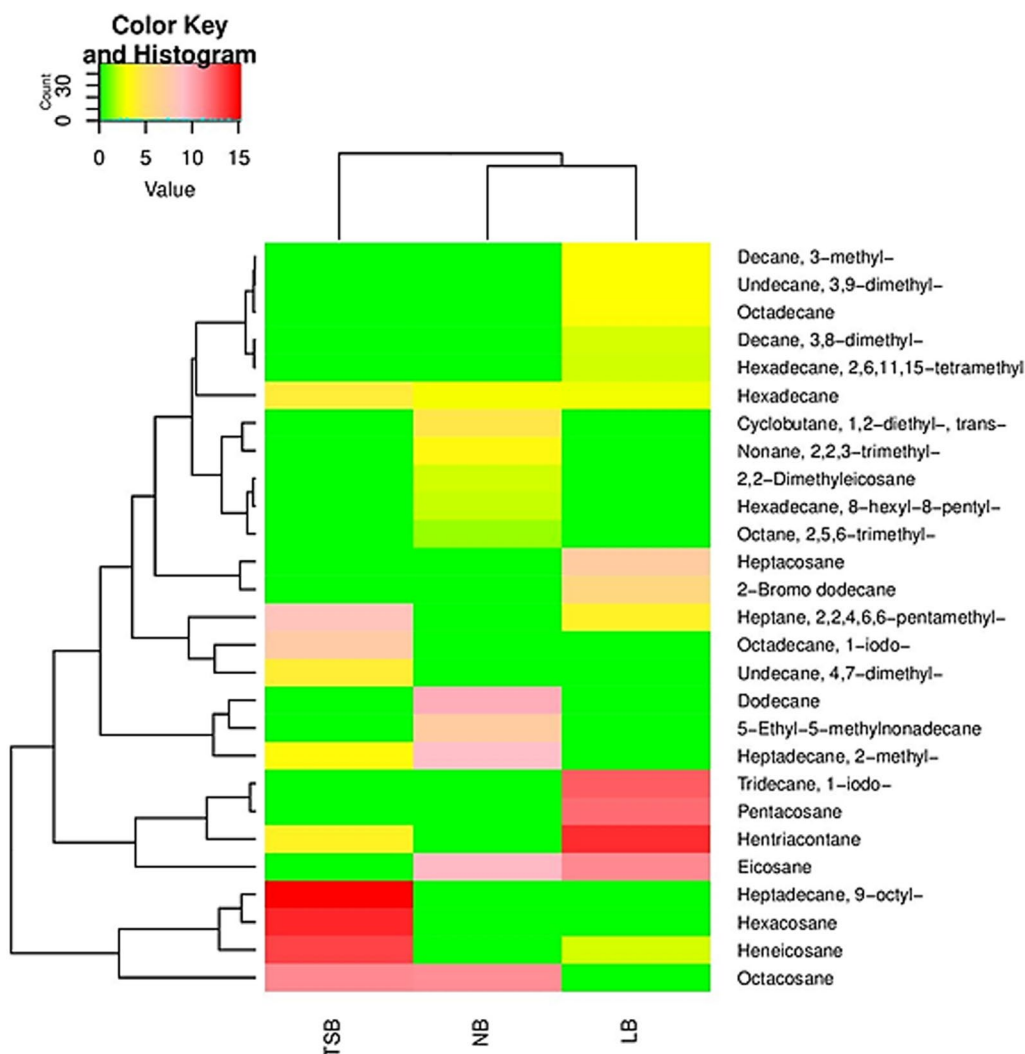


Fig. 4 The heatmap displays the amounts of bio-alkanes (mg alkane/L culture) produced by *B. spizizenii* Camph.2 in NB, LB, and TSB media

4,7-dimethyl (3.18 mg/L), Hexadecane (4.19 mg/L), Decane, 2-methyl (3.26 mg/L), Tridecane, 1-iodo (14.82 mg/L), Heptadecane, 8-methyl (11.21 mg/L), 10-Methylnonadecane (2.98 mg/L), Pentadecane (4.54 mg/L), Octacosane (9.71 mg/L), Hentriacontane (8.95 mg/L), Octadecane (4.55 mg/L), Hexadecane, 2-methyl (10.52 mg/L), and Pentadecane, 3-methyl (6.25 mg/L). Ten alkanes were detected in a sucrose-based medium, including Heptane, 2,2,4,6,6-pentamethyl (7.25 mg/L), Decane, 3,6-dimethyl (3.04 mg/L), Octane, 2,4,6-trimethyl (4.13 mg/L), Pentacosane (2.98 mg/L), Heneicosane (11.80 mg/L), Tridecane, 1-iodo (10.44 mg/L), Hexadecane (3.49 mg/L), Eicosane (3.05 mg/L), Hexacosane (6.85 mg/L), and 2-methyloctacosane (13.55 mg/L). On the other hand, 2,2,7,7-Tetramethyloctane (5.81 mg/L), Decane, 3-methyl (2.56 mg/L), Undecane, 3-methyl (4.75 mg/L),

Octadecane, 2-methyl (12.99 mg/L), Hexacosane (11.38 mg/L), Eicosane (2.97 mg/L), Heptadecane (3.53 mg/L), 2-Bromo dodecane (10.37 mg/L), Triacontane (7.06 mg/L), Heneicosane (7.48 mg/L), were produced in sugar cane molasses-based medium by *B. spizizenii* Camph.2.

Seven alkanes including Heptane, 2,2,4,6,6-pentamethyl (17.92 mg/L), Decane, 3,6-dimethyl (6.01 mg/L), Heptacosane (6.05 mg/L), Tetracosane (11.25 mg/L), Pentacosane (8.02 mg/L), Heptadecane, 2-methyl (9.11 mg/L), and Eicosane (5.75 mg/L) were produced in glucose-based medium by *B. aerophilus* Camph.3. Moreover, Decane, 2,2,3-trimethyl (11.32 mg/L), Undecane, 5-methyl (4.37 mg/L), Eicosane (5.98 mg/L), Nonane, 4,5-dimethyl (5.78 mg/L), Heptadecane, 8-methyl (17.89 mg/L), Octacosane (13.33 mg/L), Octadecane (11.47 mg/L), and Octane, 2-methyl (11.01 mg/L) were

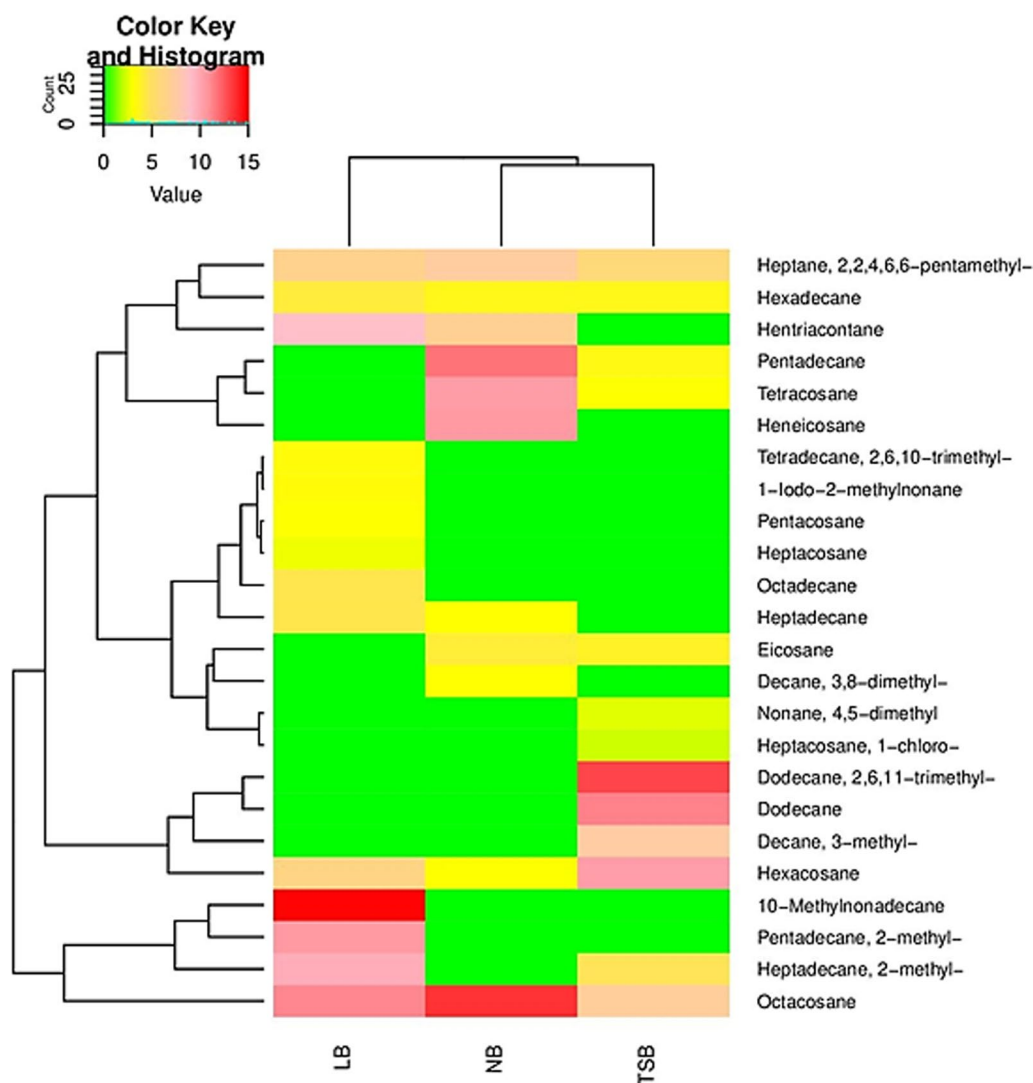


Fig. 5 The heatmap displays the amounts of bio-alkanes (mg alkane/L culture) produced by *B. aerophilus* Camph.3 in NB, LB, and TSB media

detected in sucrose-based medium. Sugar cane molasses-based medium achieved the production of twelve alkanes by *B. aerophilus* Camph.3 which are Heptane, 2,2,4,6,6-pentamethyl (6.98 mg/L), Hexacosane (3.36 mg/L), Nonane, 4,5-dimethyl (4.09 mg/L), Decane, 3-methyl (3.38 mg/L), Heneicosane (14.51 mg/L), 10-Methylnonadecane (2.88 mg/L), Docosane (14.53 mg/L), Octadecane (3.77 mg/L), Octadecane, 1-iodo (13.61 mg/L), Hentriacontane (2.78 mg/L), 2-methyloctacosane (11.65 mg/L), and Heptadecane, 9-octyl (5.96 mg/L).

As stated above, the chain length of alkanes produced by the present isolates ranged from C_8 to C_{31} (Tables 1, 2, 3, 4, 5, 6, 7, 8, 9). Previous studies have reported that bacterial alkanes typically have chain lengths ranging

from C_{10} to C_{36} , although this can vary depending on the bacterial strain and environmental conditions [29]. The biosynthesis of n-alkanes by various bacteria including *Desulfovibrio* sp., *Clostridium* sp., *Pseudomonas fluorescens*, *Vibrio furnissii* M1, and Engineered *Escherichia coli* has been reported [30–33]. Although endophytic bacteria were known within the biotechnology field for their ability to produce a great variety of sustainable safe, eco-friendly products, there are no reports about their ability to produce alkanes [34]. Therefore, this study is the first documentation of alkane production by endophytic bacteria.

The alkanes are preferred as clean fuels, because they burn cleanly and easily, releasing a lot of heat and light energy [35]. In the present study, the three

Table 1 Bio-alkanes produced by *B. atrophaeus* Camph.1 in NB medium

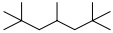
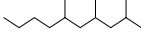
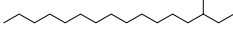

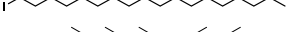
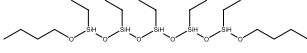
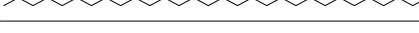
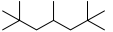
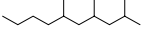
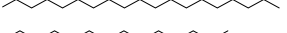
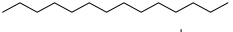
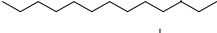
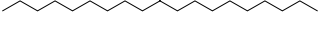
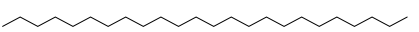



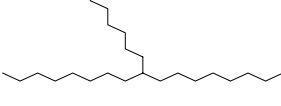

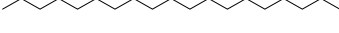

Alkane name	Structure	Formula	Molecular weight	Retention time	Peak area	mg alkane/L culture
Heptane, 2,2,4,6,6-pentamethyl-		C ₁₂ H ₂₆	170	4.432	1,863,723	17.923
Decane, 2,4,6-trimethyl-		C ₁₃ H ₂₈	184	5.215	625,432	6.015
Hexadecane, 3-methyl-		C ₁₇ H ₃₆	240	7.378	629,978	6.058
Hexacosane		C ₂₆ H ₅₄	366	9.089	1,170,008	11.252
Octadecane, 1-iodo-		C ₁₈ H ₃₇ I	380	9.495	834,492	8.025
1,3,5,7,9-Pentaethyl-1,9-dibutoxypentasiloxane		C ₁₈ H ₄₈ O ₆ Si ₅	500	10.789	722,371	6.947
Hentriacontane		C ₃₁ H ₆₄	436	11.378	947,253	9.110

Table 2 Bio-alkanes produced by *B. atrophaeus* Camph.1 in LB medium

Alkane name	Structure	Formula	Molecular weight	Retention time	Peak area	mg alkane/L culture
Heptane, 2,2,4,6,6-pentamethyl-		C ₁₂ H ₂₆	170	4.426	1,316,286	6.984
Decane, 2,4,6-trimethyl-		C ₁₃ H ₂₈	184	5.215	634,388	3.366
Octadecane, 1-iodo-		C ₁₈ H ₃₇ I	380	7.378	772,529	4.099
Tetradecane		C ₁₄ H ₃₀	198	7.733	638,078	3.386
Tridecane, 3-methyl-		C ₁₄ H ₃₀	198	9.089	2,736,279	14.519
10-Methylnonadecane		C ₂₀ H ₄₂	282	11.378	2,565,585	13.613
Hexacosane		C ₂₆ H ₅₄	366	9.495	2,738,304	14.530
Tetracosane		C ₂₄ H ₅₀	338	9.587	710,911	3.772
Eicosane, 2-methyl-		C ₂₁ H ₄₄	296	9.684	624,411	3.313
Undecane, 2,9-dimethyl-		C ₁₃ H ₂₈	184	11.487	524,687	2.784
Heptadecane, 9-octyl-		C ₂₅ H ₅₂	352	11.939	2,196,338	11.654
Heptadecane, 2-methyl-		C ₁₈ H ₃₈	254	9.169	543,692	2.885
Nonadecane, 2-methyl-		C ₂₀ H ₄₂	282	14.531	1,124,351	5.966
2-methyloctacosane		C ₂₉ H ₆₀	408	15.240	1,008,527	5.351

studied endophytic bacteria produced a variety of alkanes as mentioned above. Many of these alkanes are used in biofuel production. Octane and decane are the main constituents of gasoline. Octane is used in internal combustion engines. Nonane, decane, undecane, tetradecane, pentadecane, and hexadecane

make up the majority of diesel, kerosene, and aviation fuel. Heptadecane, octadecane, eicosane, pentacosane, hexacosane, heptacosane, octacosane, and heneicosane are the main components of lubricating oil [36].

Table 3 Bio-alkanes produced by *B. atrophaeus* Camph.1 in TSB medium

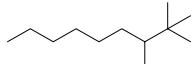
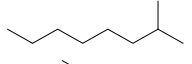
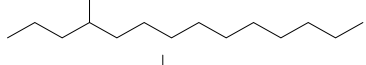
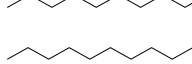
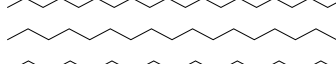

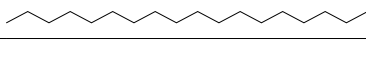
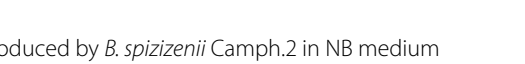
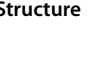

Alkane name	Structure	Formula	Molecular weight	Retention time	Peak area	mg alkane/L culture
Nonane, 2,2,3-trimethyl-		C ₁₂ H ₂₆	170	4.426	1,709,131	11.321
Octane, 2-methyl-		C ₉ H ₂₀	128	5.215	661,119	4.379
Tetradecane, 4-ethyl-		C ₁₆ H ₃₄	226	7.378	903,832	5.987
Decane, 3-methyl-		C ₁₁ H ₂₄	156	7.739	873,751	5.787
Heptadecane, 2-methyl-		C ₁₈ H ₃₈	254	9.089	2,701,569	17.894
Pentacosane		C ₂₅ H ₅₂	352	9.495	2,013,320	13.336
Octadecane		C ₁₈ H ₃₈	254	11.378	1,731,725	11.470
Hexacosane		C ₂₆ H ₅₄	366	11.939	1,224,642	8.112
Cyclobutane, 1,2-diethyl-		C ₈ H ₁₆	112	14.531	1,615,566	10.701
Eicosane		C ₂₀ H ₄₂	282	15.240	1,662,776	11.014

Table 4 Bio-alkanes produced by *B. spizizenii* Camph.2 in NB medium

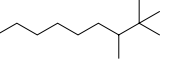
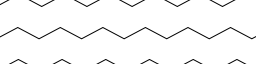
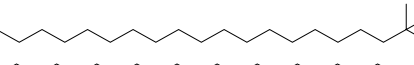
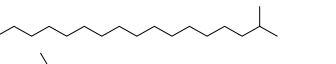
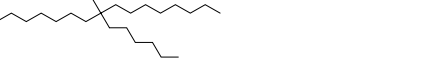
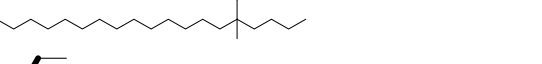

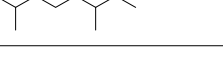



Alkane name	Structure	Formula	Molecular weight	Retention time	Peak area	mg alkane/L culture
Nonane, 2,2,3-trimethyl-		C ₁₂ H ₂₆	170	4.420	1,262,471	3.354
Dodecane		C ₁₂ H ₂₆	170	9.089	3,648,878	9.694
Eicosane		C ₂₀ H ₄₂	282	9.495	3,475,442	9.233
Hexadecane		C ₁₆ H ₃₄	226	10.062	1,079,883	2.869
2,2-Dimethyleicosane		C ₂₂ H ₄₆	310	11.206	855,681	2.273
Octacosane		C ₂₈ H ₅₈	394	11.378	4,149,590	11.024
Heptadecane, 2-methyl-		C ₁₈ H ₃₈	254	11.939	3,373,007	8.961
Hexadecane, 8-hexyl-8-pentyl-		C ₂₇ H ₅₆	380	12.059	789,803	2.098
5-Ethyl-5-methylnonadecane		C ₂₂ H ₄₆	310	14.531	2,794,080	7.423
Cyclobutane, 1,2-diethyl-, trans-		C ₈ H ₁₆	112	14.571	1,681,555	4.467
Octane, 2,5,6-trimethyl-		C ₁₁ H ₂₄	156	19.091	594,154	1.579

Table 5 Bio-alkanes produced by *B. spizizenii* Camph.2 in LB medium

Alkane name	Structure	Formula	Molecular weight	Retention time	Peak area	mg alkane/L culture
Heptane, 2,2,4,6,6-penta-methyl-		C ₁₂ H ₂₆	170	4.420	1,516,557	3.802
Undecane, 3,9-dimethyl-		C ₁₃ H ₂₈	184	7.373	1,186,604	2.975
Decane, 3,8-dimethyl-		C ₁₂ H ₂₆	170	7.733	952,014	2.387
Tridecane, 1-iodo-		C ₁₃ H ₂₇ I	310	9.089	4,977,726	12.479
Hexadecane, 2,6,11,15-tetra-methyl		C ₂₀ H ₄₂	282	9.170	913,946	2.291
Pentacosane		C ₂₅ H ₅₂	352	9.496	4,801,726	12.038
Octadecane		C ₁₈ H ₃₈	254	9.685	1,243,122	3.116
Decane, 3-methyl-		C ₁₁ H ₂₄	156	9.810	1,196,272	2.999
Hexadecane		C ₁₆ H ₃₄	226	10.062	1,118,006	2.803
Hentriacontane		C ₃₁ H ₆₄	436	11.378	5,539,119	13.886
Eicosane		C ₂₀ H ₄₂	282	11.939	4,466,188	11.196
Heneicosane		C ₂₁ H ₄₄	296	12.029	938,724	2.353
Heptacosane		C ₂₇ H ₅₆	380	14.531	2,948,890	7.393
2-Bromo dodecane		C ₁₂ H ₂₅ Br	248	15.235	2,428,083	6.087

Table 6 Bio-alkanes produced by *B. spizizenii* Camph.2 in TSB medium

Alkane name	Structure	Formula	Molecular weight	Retention time	Peak area	mg alkane/L culture
Heptane, 2,2,4,6,6-pentamethyl-		C ₁₂ H ₂₆	170	4.426	1,354,335	8.360
Undecane, 4,7-dimethyl-		C ₁₃ H ₂₈	184	5.215	662,187	4.088
Hexadecane		C ₁₆ H ₃₄	226	7.373	672,525	4.152
Heneicosane		C ₂₁ H ₄₄	296	9.089	2,137,083	13.192
Hexacosane		C ₂₆ H ₅₄	366	9.495	2,272,655	14.029
Hentriacontane		C ₃₁ H ₆₄	436	9.593	606,217	3.742
Heptadecane, 2-methyl-		C ₁₈ H ₃₈	254	9.684	527,376	3.256
Heptadecane, 9-octyl-		C ₂₅ H ₅₂	352	11.378	2,465,248	15.218
Octacosane		C ₂₈ H ₅₈	394	11.939	1,812,518	11.189
Octadecane, 1-iodo-		C ₁₈ H ₃₇ I	380	14.531	1,227,028	7.575
Pentadecane, 2-methyl-		C ₁₆ H ₃₄	226	15.235	1,170,809	7.228

Table 7 Bio-alkanes produced by *B. aerophilus* Camph.3 in NB medium

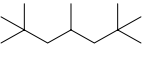
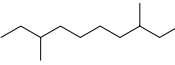
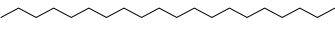
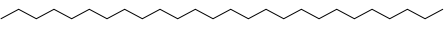
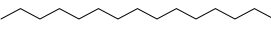
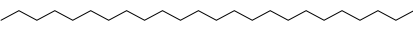
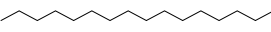
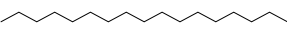

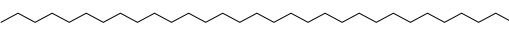
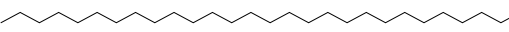
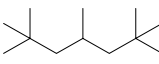
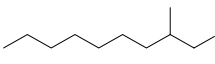
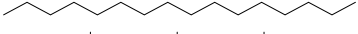
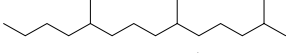
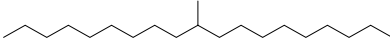
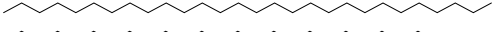



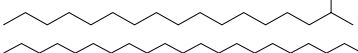
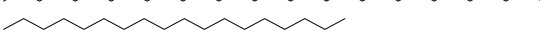

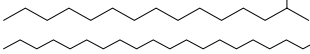
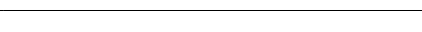
Alkane name	Structure	Formula	Molecular weight	Retention time	Peak area	mg alkane/L culture
Heptane, 2,2,4,6,6-penta-methyl-		C ₁₂ H ₂₆	170	4.432	1,397,729	7.257
Decane, 3,8-dimethyl-		C ₁₂ H ₂₆	170	5.216	586,155	3.043
Eicosane		C ₂₀ H ₄₂	282	7.379	796,408	4.135
Hexacosane		C ₂₆ H ₅₄	366	7.733	574,953	2.985
Pentadecane		C ₁₅ H ₃₂	212	9.089	2,274,449	11.808
Tetracosane		C ₂₄ H ₅₀	338	9.496	2,011,445	10.443
Hexadecane		C ₁₆ H ₃₄	226	10.062	673,346	3.496
Heptadecane		C ₁₇ H ₃₆	240	11.207	588,982	3.058
Heneicosane		C ₂₁ H ₄₄	296	11.378	2,036,079	10.571
Hentriacontane		C ₃₁ H ₆₄	436	11.939	1,319,359	6.850
Octacosane		C ₂₈ H ₅₈	394	14.531	2,611,310	13.557

Table 8 Bio-alkanes produced by *B. aerophilus* Camph.3 in LB medium

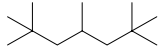
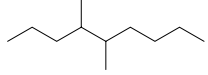
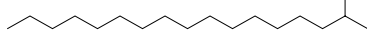
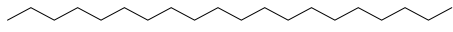
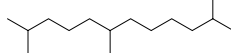
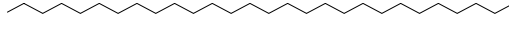
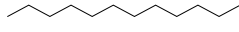
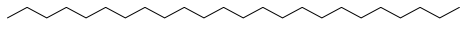
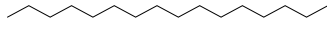
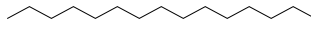
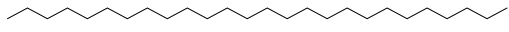
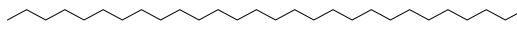
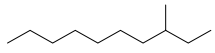
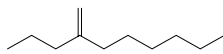
Alkane name	Structure	Formula	Molecular weight	Retention time	Peak area	mg alkane/L culture
Heptane, 2,2,4,6,6-penta-methyl-		C ₁₂ H ₂₆	170	4.432	1,226,975	6.634
1-Iodo-2-methylnonane		C ₁₀ H ₂₁ I	268	5.215	588,941	3.184
Hexadecane		C ₁₆ H ₃₄	226	7.378	776,012	4.196
Tetradecane, 2,6,10-tri-methyl-		C ₁₇ H ₃₆	240	7.739	603,654	3.264
10-Methylnonadecane		C ₂₀ H ₄₂	282	9.089	2,740,948	14.820
Octacosane		C ₂₈ H ₅₈	394	9.495	2,074,784	11.218
Pentacosane		C ₂₅ H ₅₂	352	9.587	552,547	2.988
Heptacosane		C ₂₇ H ₅₆	380	9.684	515,221	2.786
Heptadecane		C ₁₇ H ₃₆	240	11.201	840,206	4.543
Heptadecane, 2-methyl-		C ₁₈ H ₃₈	254	11.378	1,796,476	9.714
Hentriacontane		C ₃₁ H ₆₄	436	11.939	1,656,367	8.956
Octadecane		C ₁₈ H ₃₈	254	12.545	842,691	4.556
Pentadecane, 2-methyl-		C ₁₆ H ₃₄	226	14.531	1,946,339	10.524
Hexacosane		C ₂₆ H ₅₄	366	15.235	1,156,936	6.256

Conclusion

Using microorganisms is a fantastic new starting point for sustainable biofuel production. The study's findings, which were not reported previously, identified three species of bacteria as effective and environmentally benign sources for the production of different alkanes. Three

endophytic bacteria were isolated from the leaves of *C. camphora* and were molecularly identified as *Bacillus atrophaeus* Camph.1 (OR343176.1), *Bacillus spizizenii* Camph.2 (OR343177.1), and *Bacillus aerophilus* Camph.3 (OR343178.1). These isolates showed great potential in producing various alkanes when grown in NB, LB, and TSB

Table 9 Bio-alkanes produced by *B. aerophilus* Camph.3 in TSB medium

Alkane name	Structure	Formula	Molecular weight	Retention time	Peak area	mg alkane/L culture
Heptane, 2,2,4,6,6-pentamethyl-		C ₁₂ H ₂₆	170	4.426	1,284,688	5.810
Nonane, 4,5-dimethyl		C ₁₁ H ₂₄	156	5.216	566,578	2.562
Heptadecane, 2-methyl-		C ₁₈ H ₃₈	254	7.379	1,050,595	4.751
Eicosane		C ₂₀ H ₄₂	282	7.733	836,152	3.782
Dodecane, 2,6,11-trimethyl-		C ₁₅ H ₃₂	212	9.089	2,872,460	12.991
Heptacosane, 1-chloro-		C ₂₇ H ₅₅ Cl	414	9.170	489,970	2.216
Dodecane		C ₁₂ H ₂₆	170	9.496	2,518,180	11.388
Tetracosane		C ₂₄ H ₅₀	338	9.593	657,646	2.974
Hexadecane		C ₁₆ H ₃₄	226	10.062	780,927	3.532
Pentadecane		C ₁₅ H ₃₂	212	11.201	752,401	3.403
Hexacosane		C ₂₆ H ₅₄	366	11.378	2,293,290	10.371
Octacosane		C ₂₈ H ₅₈	394	11.939	1,562,255	7.065
Decane, 3-methyl-		C ₁₁ H ₂₄	156	14.537	1,654,373	7.482
Decane, 4-methylene-		C ₁₁ H ₂₂	154	14.577	967,388	4.375

media. Numerous of the produced alkanes, such as octane, nonane, decane, undecane, tetradecane, pentadecane, and hexadecane are used in biofuel production, such as gasoline, diesel, kerosene, and aviation fuel. Therefore, these endophytic bacteria may be promising and sustainable sources for alkane biofuel production.

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Author contributions

N.Sh.A.H. study design, material preparation, isolation of bacteria and experiments, data collection and analysis, and wrote the main manuscript and E. A. El-R. GC/MS analysis and reviewed the manuscript. Both authors approved the final manuscript.

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Availability of data and materials

The dataset supporting the conclusions of this article is available in the [NCBI] repository [<https://www.ncbi.nlm.nih.gov/nuccore/OR343176.1/>], [<https://www.ncbi.nlm.nih.gov/nuccore/OR343177.1/>], and [<https://www.ncbi.nlm.nih.gov/nuccore/OR343178.1/>].

Declarations

Ethics approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

Competing interests

The authors declare no competing interests.

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