

# Optimization, Characterization and Quantification of Indole Acetic Acid Produced by a Potential Plant Growth Promoting Rhizobacterium *Bacillus safensis* YKS2 from Yercaud Hills, Eastern Ghats

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## Abstract

Plant growth-promoting rhizobacteria (PGPR) have proved to be an effective solution for enhancing growth of various plant species. Five different bacterial isolates extracted from rhizosphere soil were extensively studied for the production of indole acetic acid (IAA) and among those *Bacillus safensis* YKS2 strain was found to produce substantial quantities of IAA. *B. safensis* YKS2 strain was characterized and submitted to National Centre for Biotechnology Information (NCBI) (Gen Bank No. MH539636). Optimization of IAA production with varying pH and temperature revealed that IAA production was maximum at pH 7 and at a temperature of 37°C. The production of IAA was confirmed and quantified by Fourier-transform infrared spectroscopy (FTIR), Thin-layer chromatography (TLC), Gas chromatography-mass spectrometry (GC-MS). The PGPR inoculum showed significant ( $p < 0.05$ ) shoot increase (60.00 - 89.00%) and root increase (30.00 - 90.00%) relative to the controls in *Vigna radiata*. This study showed that IAA producing ability of *B. safensis* YKS2 can be used in the large-scale production of IAA for plant growth promotion.

**Keywords:** Rhizobacteria, Indole Acetic Acid, *Bacillus safensis*, *Vigna Radiata*, PGPR

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## INTRODUCTION

Plant growth-promoting rhizobacteria (PGPR) play a vital role in farming industries by greatly reducing the use of synthetic fertilizers. Rhizosphere bacteria promote plant growth directly or indirectly through many mechanisms. One of them is by synthesis of plant growth-promoting compounds that are vital for direct plant growth. Phytohormones are chemical messengers that regulate the ability of plants to respond to external environmental stimuli.<sup>1-5</sup> Numerous bacterial and fungal species are capable of producing phytohormones. The ability to produce phytohormones is widely distributed with bacterial species linked to soil and plants.<sup>6</sup>

Indole Acetic Acid (IAA) is an active heterocyclic phytohormone and is referred as plant auxin.<sup>7-9</sup> The rhizosphere bacteria that produce IAA synergistically associate with plants by enhancing plant growth; in-return, the plant root exudates are utilized by bacteria as a carbon energy source.<sup>10,11</sup> Indole acetic acid actively stimulates plant growth by breaking apical dominance that induces growth of the main stem. It is essential for plant growth and development including cell expansion or division, flowering, fruit abscission, fruit ripening, gravitropic and phototropic responses, root initiation, leaf abscission, leaf senescence and vascular tissue differentiation.<sup>12-14</sup>

A wide range of bacteria, including species of *Arthobacter* sp., *Alcaligenes* sp., *Bacillus* sp., *Enterobacter* sp., *Klebsiella* sp., *Pseudomonas* sp. and *Serratia* sp., has been documented to stimulate plant growth. Plant growth promotion involves direct mechanisms, such as nitrogen fixation, iron and phosphorous solubilization from the soil, and production of phytohormones viz., cytokinins, gibberellins and IAA.<sup>4</sup> The rhizosphere is a nutrient-rich ecological habitat with diverse microbial species. Analyzing these rhizospheric microbes can provide some vital information on selecting effective PGPR that can be studied as bioinoculants in order to enhance the growth and yield of crops. *Klebsiella* sp., have been found to be present in the rhizosphere and also reported to exhibit essential PGP characteristics. They have been isolated from the rhizosphere of Kentucky bluegrass, soybean, sugarcane, pearl millet and

rice,<sup>10,15,16</sup> and investigated extensively for the production of IAA. The present investigation is focused on the secretion and characterization of IAA from rhizosphere soil bacteria *Bacillus* sp., and to study the effect of pH and temperature on IAA production. The work is also intended to study the effect of IAA produced on root and shoot elongation of *Vigna radiata*.

## MATERIALS AND METHODS

### Soil Collection and Bacterial Isolation

The rhizosphere soil samples were obtained from Yercaud Hills in Eastern Ghats (11°12'56.62'N to 78°12'36.56'E). Sixteen soil samples from four different locations were obtained with a sampling depth of 10 cm. All the samples were cautiously taken with a clean scoop by scrapping, transferred into sterilized polythene bags kept in an icebox and transported to the laboratory. Each sample was serially diluted in the range 10<sup>-3</sup> to 10<sup>-10</sup> and spread onto sterile nutrient agar plates followed by incubation at 37°C for 24 h. After incubation, distinct bacterial colonies were identified by morphological analysis.<sup>17</sup>

### Biochemical and Molecular Identification of IAA Producing Bacteria

Indole acetic acid secretion, catalase, citrate utilization and oxidase activities, Methyl Red-Voges Proskauer (MR-VP) and motility tests of the bacterial strains were examined according to the procedure developed by Cappuccino and Sherman.<sup>18</sup> All the above-mentioned biochemical assays were used for preliminary identification and characterization of bacterial strains. The genomic DNA of IAA producing bacterial strain was extracted as per the protocol of Kalaimurugan et al.<sup>19</sup> Universal primers 27F (5' AGAGTTTGATCMTGCTCAG 3') and 1525R (5'AAG GAG GTG ATCCAGCGCAGCA 3') were used to amplify the partial 16S rRNA gene. The PCR reaction mix was planned for amplification consisting of a total volume of 25 µL comprising 17.3 µL of de-ionized autoclaved water, 2.5 µL of Taq buffer (10x), 1 µL of forward and reverse primers (1 µM/µL), 2 µL of dNTPs (10 mM/µl), 0.2 µL of Taq polymerase (3 U/µL) and 1 µL of DNA template. The isolated bacteria were further

described by the use of 16S rRNA gene sequencing following the protocol reported by Kalaimurugan et al.<sup>19</sup> The bacterial strains were identified by analyzing the sequences in the BLAST tool (NCBI). Neighbour-joining approach was performed for phylogenetic tree analysis in MEGA 6.0 software.

#### **Screening of Bacterial Isolates for IAA Production**

The isolated bacterial cultures were tested for the production of IAA.<sup>20</sup> Test inoculated bacterial cultures in LB broth with tryptophan (5mg/ml) were incubated at 28± 2°C for a week. The bacterial cultures after incubation were subjected to centrifugation at 10,000 rpm for 15 min and approximately 2 mL of clear supernatant was combined with 4 mL of Solawaski's reagent (15 mL H<sub>2</sub>SO<sub>4</sub> + 25 mL of double distilled water + 0.81 g of ferric chloride) and observed at 520 nm for IAA determination.<sup>21</sup>

#### **Production of IAA**

After the screening test, the positive cultures were taken for the production of IAA. The isolated bacteria cultures were inoculated with nutrient broth. Three media such as; LB broth (sodium chloride (NaCl)- 5.00 g/L, tryptone-10.00 g/L, yeast extract- 5.00 g/L), tryptophan broth (casein enzymic hydrolysate-10.00 g/L, sodium chloride (NaCl)-5.00 g/L, peptone-10.00 g/L), nutrient broth (beef extract- 1.00 g/L, yeast extract- 2.00 g/L, peptone- 5.00 g/L, sodium chloride (NaCl)- 5.00 g/L) were supplemented by tryptophan (5 mg/ml) and incubated at 28±2°C for a week. After incubation, the culture medium was taken for extraction and characterization studies.<sup>22</sup>

#### **Extraction of IAA**

In 100 mL of tryptophan-fortified nutrient broth (1-5 mg/mL), IAA positive bacterial culture was inoculated and incubated at 28±2°C for a week by shaking at 150 rpm in incubator shaker. The bacterial cells were discarded and the supernatant alone was collected by centrifuging the culture at 10,000 rpm for 30 min. The supernatant was acidified from pH 2.5 to 3.0 using 1 N HCl and it was extracted twice with double the amount of ethyl acetate. The fraction extracted from ethyl acetate was evaporated to dryness at 40°C using a rotatory evaporator. Finally, the extract was dissolved with methanol and stored at -20°C for further studies.<sup>8</sup>

#### **Optimization of IAA Production**

The IAA production was optimized by pH and temperature. IAA production with respect to pH range of 4 to 9 and temperature at 25, 37, 40 and 45°C were tested using the respective NB medium. The IAA production was analyzed using a UV-Vis Spectrophotometer at 400-800 nm.

#### **Characterization of IAA**

##### **Thin Layer Chromatography**

The ethyl acetate fraction was placed on TLC plates (Silica gel 254, thickness 0.25 mm). The silica gel was activated on a hot plate for 5 min before use. The acidic extract was concentrated to dryness and suspended in methanol and spotted in duplicate along with a standard IAA sample. The solvent system used was (heptane: acetone: glacial acetic acid (50:50:1)). Ascending chromatography was performed in a chromatographic glass chamber at room temperature (37°C). After the development of chromatogram, it was sprayed with ferric perchloric nitrate (FPN) solution, which is composed of 5 mL of ferric chloride (5%), 50 mL of nitric acid 50% (v/v) and 45 mL of perchloric acid 20% (v/v).<sup>23</sup>

##### **GC-MS Analysis of IAA**

The secondary metabolites obtained from bacteria were analyzed by GC-MS (QP2010 plus Equipment) using the protocol described by Cheng et al.<sup>24</sup> The Polaris Q Ion Trap GC/FID column was used, and the temperature was initially maintained at 50°C for 2 min. It was then slowly increased to 280°C for 2 min. Helium was used as the carrier gas at a flow rate of 0.9 mL /min, and the mass spectrum was measured at a voltage of 1.07 K ionizing. The individual compounds were identified by the analyzer Wiley: Registry of a mass spectral database the NIST (V 3.0). Further, the values of retention time (RT) and retention index for the isolated compounds were identified by comparing several authentic references.<sup>25</sup>

##### **FTIR Analysis of IAA**

Approximately 1 mg of the sample was finely powdered with 10 mg of pure anhydrous potassium bromide crystals (IR grade) and made into a pellet for IR examination. The IR spectrum of the sample was measured at the optimum

conditions for *Bacillus* sp. and compared with pure IAA.<sup>26</sup>

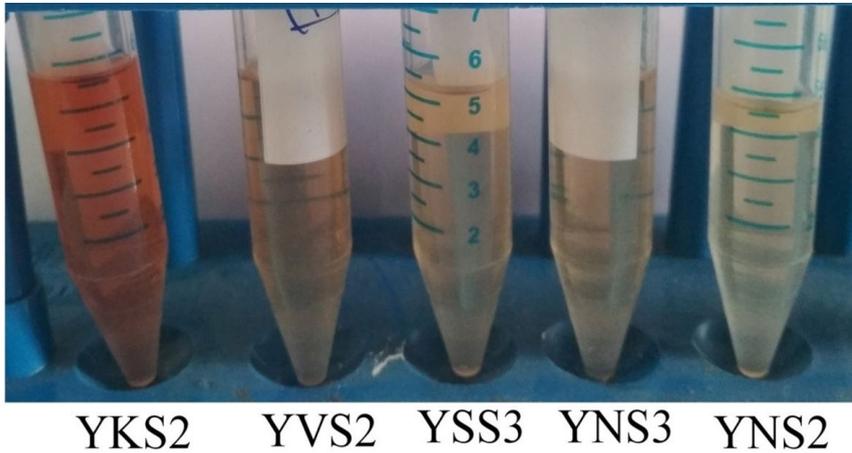
### Response to Plant Growth

*Vigna radiata* seeds from the local market were chosen for the experiment. The seeds were soaked in tap water overnight prior to study to enhance the germination rate. About 25 mL of bacterial suspension was inoculated in each experimental unit. The experimental plants were grown in the greenhouse garden in earthen pots filled with 2 kg of sterilized soil along with 5

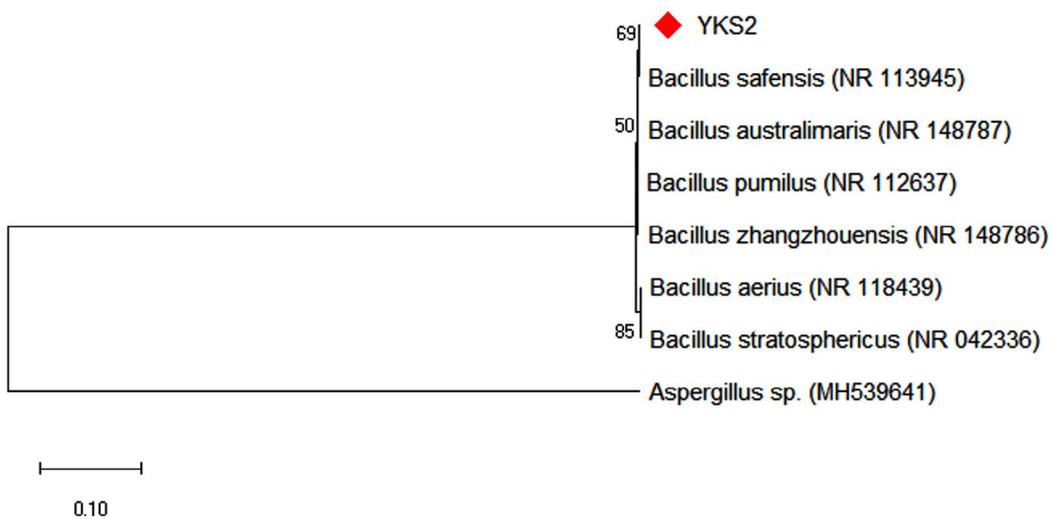
different concentrations of IAA in triplicates. After 7 weeks, the plant growth was examined and the total plant height, root length and shoot length were measured using Imagin tool 3.0 software in centimeter.<sup>27</sup>

### Statistical Analysis

The data collected were analyzed using SPSS (V 16.0) software and the comparison of treatment means was done by Duncan's Multiple Range Test (DMRT) (P<0.05).



**Figure 1.** Bacterial strains of *Bacillus* sp., isolated from rhizosphere soil having the ability to produce IAA in high quantities.



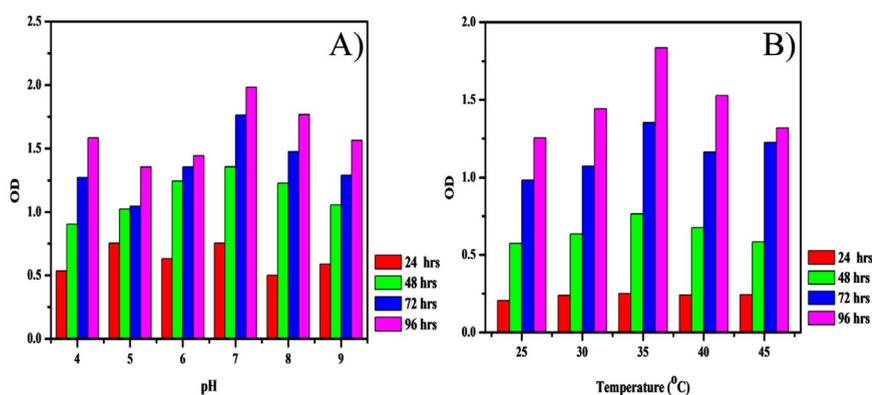
**Figure 2.** Neighbour-Joining Tree represents the phylogenetic placement of the YKS2 strain IAA with similar reference strains.

## RESULTS

### Isolation and Screening of Bacterial Isolates for IAA Production

Bacterial species that produce IAA were isolated from the rhizosphere soil sample. The bacterial isolates were cultured at 37°C for 24 h. After incubation, the samples containing the bacterial cells were subjected to various morphological, biochemical, and molecular tests.

Five strains were isolated from the rhizosphere soil to determine IAA production. Salkowski reagent was used to identify the secretion of IAA by the bacterial cultures. A colour change from pale yellow to pink is an indication of IAA production by the capability of bacterial isolates. The intensity of the colour confirms high concentration of IAA (Figure 1) present in the sample. This is due to the IAA produced by bacteria is released into the medium at a continuous slow phase. The



**Figure 3.** Effects of different parameters A) pH and B) temperature on the IAA producing YKS2 strain.



**Figure 4.** TLC to analyze the band separation in the solvent system of chloroform: methanol (ratio of 9:1, 8:2 and 7:3).

*Bacillus* strain was then subjected to various morphological and biochemical tests (Table 1). The presence of *Bacillus safensis* strain YKS2 (Gen Bank No. MH539636) was confirmed by 16S rRNA gene sequence analysis, which shared 99- 100% similarity with *Bacillus safensis*. The phylogenetic tree was constructed by neighbour-joining approach (Figure 2) using Kimura 2-parameter (K2P) model by MEGA 6.0 software.

### Optimization of IAA Production

Production of IAA by *B. safensis* strain YKS2 was optimized at various pH (4 to 9) values,

**Table 1.** Biochemical tests for *Bacillus* strain

Biochemical Test	Observations Strain YKS2
Methyl red test	Negative
Voges Proskauer test	Positive
Citrate utilization test	Positive
Catalase test	Positive
Oxidase test	Positive

and temperatures (25, 37, 40 and 45°C) to find the maximum production conditions. From the results, it was inferred that the optimal pH was 7.00 and the optimal temperature was 37°C (Figure 3). The IAA produced by *B. safensis* incubated at 37°C in an orbital shaker at 150 rpm after 24, 48, 72 and 96 h was quantified. After incubation, the samples were taken from flask and centrifuged at 8000 rpm for 15 min. The concentration of IAA was determined by Salkowski reagent.

#### Extraction of Crude IAA

Solvent extraction technique was used to extract IAA from the medium. Ethyl acetate was

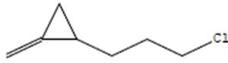
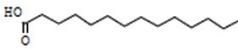
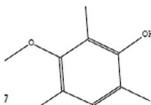
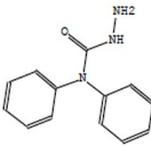
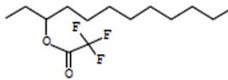
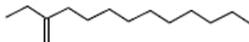
found to be the best performing extraction solvent. The dry crystals were then dissolved in methanol and stored for further analysis.

#### Characterization of IAA

##### Thin Layer Chromatography (TLC) Analysis

Ethyl acetate extract of IAA was subjected to TLC analysis using methanol: chloroform mixture at various ratios such as 9:1, 8:2 and 7:3 as a solvent system. In the Iodine vapor saturated tank, the 7:3 ratio solvent system formed a simple band separation clearly visible under UV light. The R<sub>f</sub> value was determined for the separated fractions (Fraction 1- R<sub>f</sub> = 3.4/5.7 = 0.59649 cm, Fraction 2- R<sub>f</sub> = 4.8/5.5 = 0.87273 cm and Fraction

**Table 2.** Major compounds identified by Gas Chromatography-Mass Spectrometry (GC-MS)

Retention Time	Molecular formula	Molecular weight	Compound name	Structure
23.70	C <sub>17</sub> H <sub>36</sub>	240	Heptadecane	
30.98	C <sub>7</sub> H <sub>11</sub> Cl	130	1-(3-Chloropropyl)-2-Methyle	
36.86	C <sub>14</sub> H <sub>28</sub> O <sub>2</sub>	228	Tetradecanoic Acid	
20.50	C <sub>10</sub> H <sub>14</sub> O <sub>2</sub>	166	Phenol, 3-Methoxy- 2,4,6-Trime	
16.17	C <sub>13</sub> H <sub>13</sub> N <sub>3</sub> O	227	Hydrazinecarboxamide, N,N-	
12.17	C <sub>14</sub> H <sub>25</sub> F <sub>3</sub> O <sub>2</sub>	282	3-Trifluoroacetoxydodecane	
8.38	C <sub>14</sub> H <sub>28</sub>	196	1-Dodecene, 2-Ethyl-	

3-  $R_f = 3.8/5.6 = 0.67857$  cm). The fractions were classified as phenolic ( $R_f$  value = 0.54 and 0.74) compounds based on the obtained  $R_f$  values (Figure 4).

### Gas Chromatography-Mass Spectrometry (GC-MS) Analysis

Indole acetic acid is crucial for plant growth. Its additional supply from rhizosphere bacteria can enhance a plant's defence systems

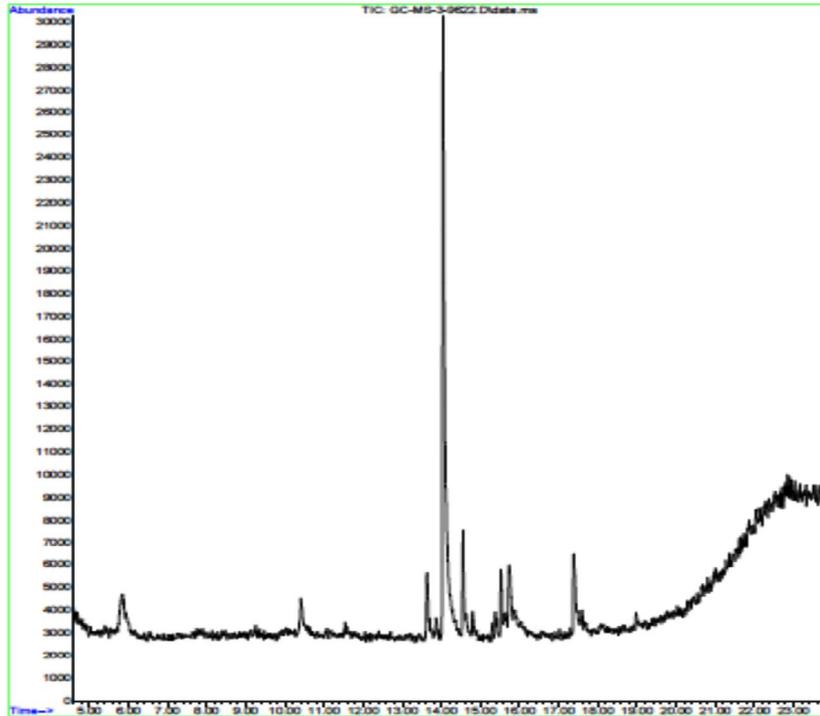


Figure 5. IAA production from bacterial strain on *B. safensis* YKS2 characterized by GC-MS

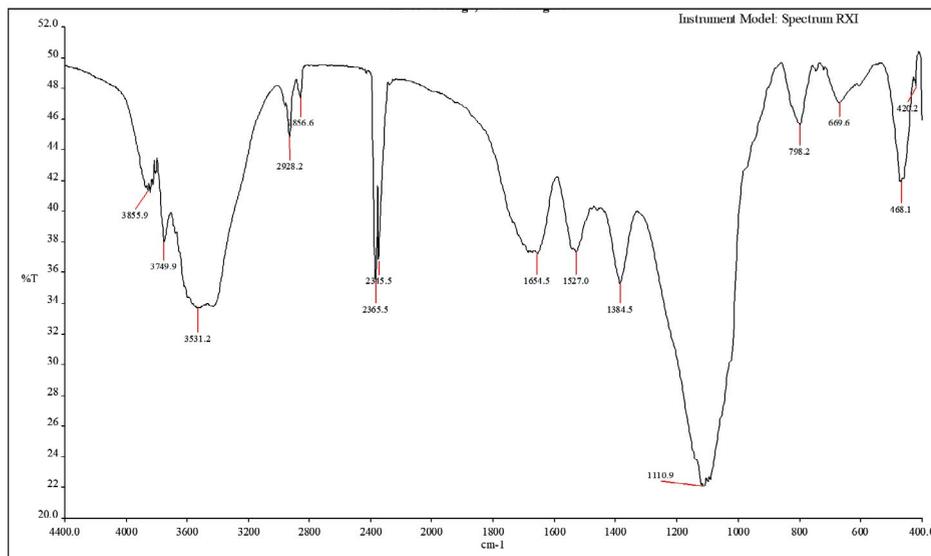
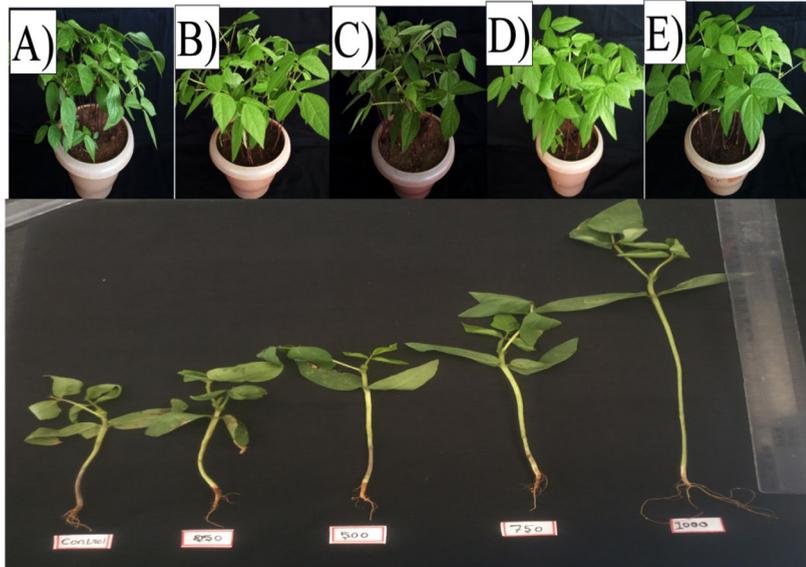


Figure 6. FTIR spectrum of IAA produced by the bacterial strain *B. safensis* YKS2.

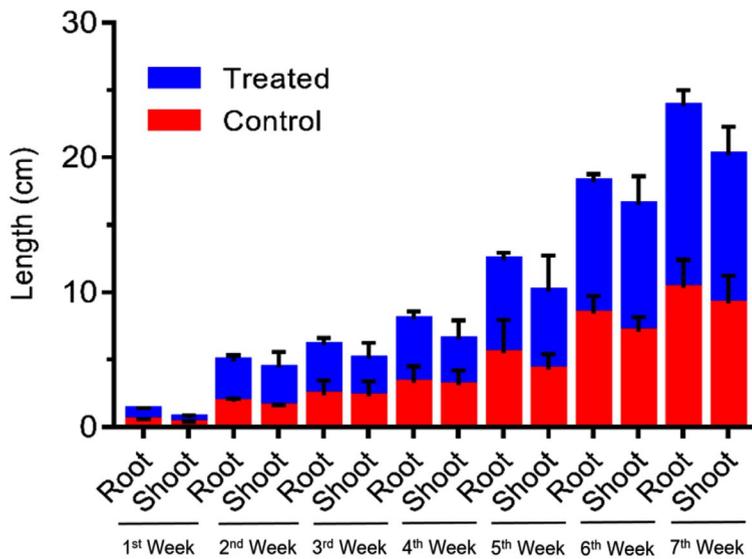
against abiotic as well as biotic stresses. The GC-MS chromatogram of chloroform extracts of IAA produced seven peaks and showed the existence of thirty-six phytochemical compounds. Out of them, five were identified as major compounds in heptadecane. The results showed that *B. safensis* strain YKS2 produced  $85.70 \pm 3.55 \mu\text{g/mL}$  IAA in the culture medium (Figure 5 and Table 2).

#### Fourier-Transform Infrared Spectroscopy (FTIR) Analysis

Indole acetic acid produced by *B. safensis* strain YKS2 was extracted, subjected to FTIR analysis and compared with the spectrum of pure indole acetic acid. Peaks corresponding to C=C, NH, C=O and C-OH vibrations were observed in the IR spectrum of the sample (Figure 6). The



**Figure 7.** Effect of different concentrations of *B. safensis* on plant growth before drought stress (upper panel) and (lower panel) 7 weeks and effect of IAA on plant growth in terms of root and shoot elongation.



**Figure 8.** Effect of different concentrations of *B. safensis* on plant growth before drought on the root and shoot elongation in *Vigna radiate*. The data are expressed as Mean  $\pm$  SD of three replicates. Asterisks (\*) indicate a significant difference between treatment and control at  $*p < 0.05$ .

characteristic N-H (indole) stretching was found at  $3855\text{ cm}^{-1}$ . The bending and wagging vibrations of the N-H bond were detected at  $1654\text{ cm}^{-1}$  and  $669\text{ cm}^{-1}$ , respectively. Indole ring stretching (C-H) bond was recorded at  $1110\text{ cm}^{-1}$ . The C=O stretching peak was observed around  $1700\text{ cm}^{-1}$ . The peak at  $1384\text{ cm}^{-1}$  is due to the C-OH stretching. Thus, the FTIR spectrum confirms the presence of IAA.

### Response to Plant Growth

The plant *Vigna radiata* was exposed to different concentrations of IAA. The initial and the final growth readings were recorded at concentrations of 250, 500, 750 and 1000 ppm. The results showed that IAA enhanced plant growth (Figure 7 and 8) in terms of the shoot length (23.3 cm) and root length (7.4 cm). Thus, it can be concluded that IAA plays a major role in promoting plant growth.

### DISCUSSION

Indole acetic acid is a vital member of the plant hormone group and is generally regarded as a significant natural auxin.<sup>28</sup> It acts as an essential signal molecule for plant growth regulators, cell proliferation, cell differentiation, cell division and gene regulation.<sup>29</sup> Various endophytic bacteria are capable of producing the plant hormone IAA and they develop auxins in the presence of active precursors like L-tryptophan. In the present study, thirteen bacterial isolates from fifteen plant root samples have been analyzed for IAA production capability.

In the present investigation, the bacteria that produce IAA were isolated from rhizosphere soil from Yercaud hills. The effect of *B. safensis* strain YKS2 isolate on the root elongation was analyzed using various concentrations of IAA extracts. Indole acetic acid was also identified to be generally released either on the surfaces or within the plant tissues due to the presence of bacterial growth. Tryptophan is a key intermediate in IAA biosynthesis by microorganisms used as a precursor for IAA production.<sup>27,30</sup> The *Bacillus* sp., isolated in the study produced high amount of IAA. The results correlate well with earlier studies on plant growth following inoculation with IAA producing PGPB.

The growth regulatory effects of indole acetic acid (IAA) have been examined at various concentrations and under different circumstances.<sup>31</sup> In the results presented here, IAA treatment at different concentrations has shown different levels of plant growth in both root and the shoot. In some cases, IAA treatment has shown reduced plant growth in a number of plant species, including flax. Ethylene biosynthetic pathway might be one of the primary suspect,<sup>32</sup> in these cases. Xylem or outer tissue radii of the plant has shown significant improvement due to IAA treatment. These results are in correlation with some reported cases where exogenous IAA treatments stimulate xylem expansion, including one study in flax.<sup>33,34</sup> Treatment conditions such as stems decapitation and defoliation to impair endogenous hormone biosynthesis also plays a role in plant growth promotion. Indole acetic acid has shown significant increase in xylem fiber cell wall thickness. Therefore, the IAA extracts at various concentrations used in this study have shown significant biological responses. Preliminary separation and identification of phenolic compounds, flavonoids, tannins and saponins in alcoholic and water extracts of IAA were performed using chromatography.

A previous study<sup>15</sup> demonstrated that the inoculation of *Rhodobacter* sp., strongly improved the morphological features of cucumber plants due to the production of IAA by the bacterial species. A report by Ahmad et al.<sup>35</sup> stated that the *Azotobacter* sp., and *Pseudomonas* sp., produced high IAA levels when they grow in a nutrient broth amended by tryptophan with 2 and 5 mg/mL. *Azospirillum* sp., is one of the most examined producers of IAA among the PGPR species. The bacterial species belonging to various genera such as *Aeromonas* sp., *Azotobacter* sp., *Bacillus* sp., *Burkholderia* sp., *Pseudomonas* sp., and *Rhizobium* sp.,<sup>36-40</sup> have been isolated from rhizosphere soils. The IAA-producing PGPR bacterial inoculation promotes stem germination and increases root biomass.

The bacteria associated with plants play a vital role in regulating plant physiology. Plant growth promoting bacteria can offer growth-enhancing properties leading to high production and ensuring food and environmental

safety by minimizing the application of harmful agrochemicals.<sup>41</sup> The synthesis of IAA in the rhizosphere by bacteria is an important factor in plant development.<sup>7</sup> In this study, tryptophan (5 mg/mL) was used as a precursor for IAA production. One of the most valuable compounds is IAA, which is a key plant-growth regulator and signaling molecule in the plant response/adaptation to environmental stresses. IAA is produced by the interplay of enzymes that mediate multiple interconnected biosynthesis pathways. The results demonstrated that the plant growth regulators provided by *Bacillus* sp., may also play an important role in promoting plant development.<sup>42,43</sup>

The rhizosphere based bacterial IAA production is an essential property that impacts on the growth of plants. Therefore, the present investigation aimed to evaluate the IAA production by *B. safensis* strain YKS2 was confirmed to produce regulators for plant growth which can also play a crucial role in promoting plant growth.

## CONCLUSION

In this present study, the maximum IAA yield was obtained (85.70±3.55 µg/mL) at pH 7 (83%) and temperature 37°C (91%) by isolated *Bacillus safensis* strain YKS2. It is also evident that *B. safensis* strain YKS2 has emerged as a novel alternate source for the production of IAA, which could aid the development of root and shoot biomass in crops. Therefore, it can also be used as an effective bio-inoculant for promoting plant growth. More research is needed to further examine the rhizosphere-based bacterial IAA production and to study its effect on numerous crop plants that are exposed to both the biotic and abiotic factors in the field.

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## CONFLICT OF INTEREST

The authors declare that there is no conflict of interest.

## AUTHORS' CONTRIBUTION

RL and DK conceptualized the study. DK and PS wrote the original draft. MMP, KP and SV reviewed and edited the manuscript. All authors read and approved the final manuscript for publication.

## FUNDING

None.

## DATA AVAILABILITY

All datasets generated or analyzed during this study are included in the manuscript.

## ETHICS STATEMENT

This article does not contain any studies with human participants or animals performed by any of the authors.

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