

Streptomyces sp. KKU215: Biocontrol of Bacterial Wilt, Tomato Growth Promotion, and Spore Production Optimization

(*Streptomyces* sp. KKU215: Biokawalan Layu Bakteria, Penggalak Pertumbuhan Tomato dan Pengoptimuman Penghasilan Spora)

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ABSTRACT

The development of microbial biocontrol agents and plant growth promoters as alternatives to chemical pesticides is crucial for sustainable agriculture. This study investigated the potential of *Streptomyces* sp. KKU215 for controlling bacterial wilt caused by *Ralstonia solanacearum*, promoting tomato growth, and enhancing spore production. The strain exhibited multiple beneficial traits, including the production of protease, chitinase, lipase, cellulase, and amylase enzymes, as well as siderophores and biofilm. It also demonstrated capabilities for nitrogen fixation, ammonia production, and indole-3-acetic acid (IAA) synthesis. *Streptomyces* sp. KKU215 significantly ($p < 0.05$) promoted the growth of 30-day-old tomato seedlings in tray trials by enhancing shoot and root lengths, fresh and dry weights, and leaf size, while also significantly ($p < 0.05$) reducing the bacterial wilt disease index and improving overall growth in pot trials. Among tested media, modified Wakimoto's and nutrient agar supported the highest levels of spore production after 20 days of incubation. The Plackett–Burman design showed peptone, potato, and incubation time as significant factors enhancing spore production. Subsequently, optimization using response surface methodology with Box–Behnken design determined the optimal conditions of 7 g/L peptone, 700 g/L potato infusion equivalent, and 20 days of incubation, resulting in a significant ($p < 0.05$) increase (46%) in spore yield. The findings of this study suggest that *Streptomyces* sp. KKU215 could serve as a sustainable alternative to harmful chemicals, contributing to enhanced crop yield and quality while supporting the development of industrial-scale biopesticide production for disease management and plant growth promotion in agricultural systems.

Keywords: Actinobacteria; biocontrol agents; Box–Behnken design; plant growth promoters

ABSTRAK

Pembangunan agen biokawalan mikrob dan penggalak pertumbuhan tumbuhan sebagai alternatif kepada racun perosak kimia adalah penting untuk pertanian lestari. Penyelidikan ini mengkaji potensi *Streptomyces* sp. KKU215 untuk mengawal layu bakteria yang disebabkan oleh *Ralstonia solanacearum*, menggalakkan pertumbuhan tomato dan meningkatkan penghasilan spora. Strain ini menunjukkan pelbagai ciri bermanfaat, termasuk penghasilan enzim protease, kitinase, lipase, selulase dan amilase, serta siderofor dan biofilem. Strain ini juga menunjukkan keupayaan untuk mengikat nitrogen, penghasilan ammonia dan sintesis asid indol-3-asetik (IAA). *Streptomyces* sp. KKU215 dengan ketaranya ($p < 0.05$) menggalakkan pertumbuhan anak benih tomato berusia 30 hari dalam ujian dulang dengan meningkatkan panjang pucuk dan akar, berat segar dan kering dan saiz daun di samping mengurangkan indeks penyakit layu bakteria dengan ketara ($p < 0.05$) dan meningkatkan pertumbuhan keseluruhan dalam ujian pasu. Antara media yang diuji, media Wakimoto yang diubah suai dan agar nutrien menyokong tahap penghasilan spora tertinggi selepas 20 hari pengeraman. Reka bentuk Plackett-Burman mendedahkan pepton, kentang dan masa pengeraman sebagai faktor penting untuk meningkatkan penghasilan spora. Seterusnya, pengoptimuman menggunakan metodologi permukaan tindak balas dengan reka bentuk Box-Behnken telah menentukan keadaan optimum 7 g/L pepton, 700 g/L setara infusi kentang dan 20 hari pengeraman menunjukkan peningkatan ketara ($p < 0.05$) sebanyak 46% dalam penghasilan spora. Hasil kajian ini menunjukkan bahawa *Streptomyces* sp. KKU215 boleh berfungsi sebagai alternatif yang mampan kepada bahan kimia berbahaya, menyumbang kepada peningkatan hasil dan kualiti tanaman di samping menyokong pembangunan pengeluaran biopestisid berskala industri untuk pengurusan penyakit dan penggalak pertumbuhan tanaman dalam sistem pertanian.

Kata kunci: Agen biokawalan; aktinobakteria; penggalak pertumbuhan tumbuhan; reka bentuk Box-Behnken

INTRODUCTION

Ralstonia solanacearum (Rs) is a bacterium responsible for bacterial wilt (BW disease), a serious threat to tomato production worldwide, including in Thailand, causing significant yield losses (Akarapisan et al. 2021; Fonseca et al. 2024). Rs produces exopolysaccharides that obstruct xylem vessels, leading to disruption of water and mineral transport, ultimately causing plant wilting and death. Chemical control methods can reduce disease incidence but pose serious risks to human health and the environment (Nicolopoulou-Stamati et al. 2016). Moreover, current management strategies are often ineffective, as Rs can survive in soil and water for extended periods even without a host (Vaillau & Genin 2023). Hence, safer and more efficient alternatives are urgently needed.

Biological control using *Streptomyces* spp. shows promise in managing plant diseases by producing antimicrobial compounds, competing for nutrients and space, and inducing systemic resistance (Khan et al. 2023). They also promote plant growth by fixing nitrogen and producing plant hormones, ammonia, and biofilms (Ajijah et al. 2023; Li et al. 2024). Various *Streptomyces* strains have been shown to reduce BW disease and enhance plant growth (Boukaew, Chuenchit & Petcharat 2011; Devi, Sharma & Manhas 2022; Elsharkawy et al. 2015; He et al. 2024; Kaari et al. 2022a; Ling et al. 2020; Le et al. 2021; Xue et al. 2019; Zhuang et al. 2020). Given *Streptomyces* efficacy in disease suppression and plant growth promotion, this study aimed to discover novel strains for managing BW disease and enhancing tomato growth.

The novel strain of *Streptomyces* sp. K KU215 was isolated from botanical garden soils. The isolate was identified by 16S rRNA gene sequencing (Yottakot & Leelavatcharamas 2019), and the sequence was deposited in GenBank (Accession No. MK835665). Subsequently, its antagonism against *Ralstonia solanacearum* DOA-BCC1954 was tested using a double-layer assay, with preliminary results showing strong antagonistic activity (Figure S1). Building upon these preliminary findings, *Streptomyces* sp. K KU215 appears to be a distinctive strain with a robust combination of multifunctional traits and intrinsic potential for high-yield spore production, a critical requirement for commercial biopesticide development. To enable agricultural applications, it is necessary to investigate the biocontrol and growth-promoting mechanisms of K KU215 and to optimize spore production due to its high inoculum requirements. Techniques such as the Plackett-Burman (PB) design (Liu et al. 2020) and Response Surface Methodology (RSM) with the Box-Behnken (BB) design have been successfully used to optimize microbial metabolite production (Liu et al. 2025).

This study aimed to (1) investigate the characteristics related to disease control and plant growth promotion of K KU215, (2) evaluate its efficacy in controlling BW disease

and promoting tomato growth, and (3) screen factors affecting spore production using PB design, followed by optimization of culture conditions for enhanced spore production through RSM with BB design. The outcomes of this study offer an alternative to harmful chemicals in enhancing crop yield and quality, as well as promoting sustainable agricultural practices.

MATERIALS AND METHODS

MICROBIAL STRAIN AND CULTURE CONDITION

Spore suspension of K KU215, which inhibits RsDOA-BCC1954, was cultured on nutrient agar (NA; Himedia™) at 37 °C for 10 days. This temperature was selected based on preliminary tests indicating optimal and rapid spore production on solid media. Sterile distilled water was added to the plates, and spores were scraped off with a sterile spatula. Spore concentration was adjusted to 1×10^7 spores/mL using a hemacytometer. This concentration was used in all experiments. Cell suspension of RsDOA-BCC1954, obtained from the Plant Protection Research and Development Office (Department of Agriculture, Thailand), was cultured in yeast extract peptone glucose (YPG) medium containing 0.7% peptone, 0.7% yeast extract, and 0.5% glucose (pH 7.0) at 28 °C for 2 days on a rotary shaker (160 rpm). Cells were harvested by centrifugation ($5,635 \times g$, 10 min, 4 °C), washed three times with sterile distilled water, and adjusted to OD₆₀₀ = 0.5.

BIOCONTROL ACTIVITY EVALUATION

Extracellular lytic enzyme production was assessed on selective media: starch agar (amylase; Hasan et al. 2017), carboxymethyl cellulose agar (cellulase; Li et al. 2020), tributyrin agar (lipase; Ilesanmi et al. 2020), skim milk agar (protease; Shaikh et al. 2023), and colloidal chitin agar (chitinase; Xia et al. 2011). A 2 µL spore suspension of K KU215 was spotted onto each medium and incubated at 37 °C. After 2 days, enzyme activity was indicated by clear zones around the colony. Siderophore production was assessed using Chromeazurol S (CAS) agar plates (Schwyn & Neilands 1987), following the method of Fahsi et al. (2021). Similarly, 2 µL spore suspension of K KU215 was spotted onto CAS agar and incubated at 37 °C for 3 days. Siderophore production was indicated by the appearance of orange zones around the colony. Biofilm formation was assessed following Jain et al. (2013). A 100 µL spore suspension of K KU215 was inoculated into 1 mL nutrient broth (NB; Himedia™) in glass tubes and incubated at 37 °C for 7 days without agitation, with an uninoculated control. After incubation, cultures were discarded, tubes washed twice with PBS (pH 7.3), air-dried, stained with 0.1% crystal violet for 5 min, rinsed with sterile distilled water, and examined for staining to confirm biofilm formation.

PLANT GROWTH PROMOTION TRAITS EVALUATION

Nitrogen fixation was assessed on nitrogen-free (NF) agar plates as described by Kifle and Laing (2016), by spotting 2 μ L of a spore suspension of KKKU215 onto the plates and incubating at 37 °C for 5 days, with colony growth indicating the ability to fix nitrogen. Ammonia production was measured following the method of Fahsi et al. (2021), in which 100 μ L of a spore suspension of KKKU215 was inoculated into 100 mL of 4% peptone broth and incubated at 160 rpm and 37 °C for 10 days. After incubation, 0.5 mL of Nessler's reagent was added to 2.5 mL of the culture and allowed to react for 10 min. The ammonia concentration was determined at 420 nm using a spectrophotometer and compared to a standard curve of ammonium sulfate. Indole acetic acid (IAA) production was measured following Fonseca et al. (2024). A 100 μ L spore suspension of KKKU215 was inoculated into NB with L-tryptophan concentrations of 1-5 mg/mL. The culture was incubated at 37 °C with shaking (160 rpm), and samples were collected every 2 days for 15 days. After centrifugation at 5,635 \times g for 10 min, Salkowski's reagent was added to the supernatant and incubated in the dark for 30 min. IAA concentration was measured at 535 nm and compared to a standard curve.

SEEDLING GROWTH

This study was conducted in October 2024 in an open greenhouse with temperatures ranging from 23.2 °C to 32.4 °C and relative humidity between 55.9% and 90.5%. Tomato seeds were surface-sterilized with 1% sodium hypochlorite for 2 min, followed by three rinses with sterile distilled water. Fifteen seeds per treatment were divided into three replicates (trays), with five seeds sown in each tray. Seeds were soaked in a spore suspension of KKKU215 or sterile distilled water (control) for 30 min (Hernandez-Herrera et al. 2016) before being sown in trays with 50 g of sterile peat moss without fertilizer. After 30 days, shoot and root lengths were measured with a ruler, and fresh and dry weights were determined using a digital balance.

POT EXPERIMENTS

The experiment was conducted from November to December 2024 in an open greenhouse with temperatures ranging from 18.9°C to 30.6°C and relative humidity between 48.9% and 90.7%. Twenty-one-day-old tomato seedlings were transplanted into 1.2 kg of sterile peat moss per pot, supplemented with 1.2 g of 15:15:15 fertilizer. Treatments were T1, KKKU215 only; T2, KKKU215+RsDOA-BCC1954 (biocontrol treatment); T3, no KKKU215 and no RsDOA-BCC1954 (uninoculated control); and T4, RsDOA-BCC1954 only (pathogen control).

The pot experiment was arranged in a Completely Randomized Design (CRD) with 4 treatments and 10 replicate pots per treatment. The total number of experimental units was 40 pots, and each pot served as one

experimental unit. A 50 mL KKKU215 spore suspension was applied on days 0, 7, 14, 21, and 35. On day 42, 50 mL of RsDOA-BCC1954 suspension (OD600 = 0.5) was applied to the root zone. At 56 days, disease index (DI) and plant growth parameters (shoot/root lengths, fresh/dry weights) were recorded. The DI was assessed once at 56 days post-transplantation using the scale (0-4) and formula from Wu et al. (2016). The scale defined symptoms as 0 = no symptoms; 1 = 1-25% wilting; 2 = 26-50% wilting; 3 = 51-75% wilting; and 4 = 76-100% wilting or dead. DI was calculated using the formula:

$$DI = \frac{\sum(\text{number of diseased plants in this index} \times di)}{(\text{total number of plants investigated} \times \text{highest di})} \times 100$$

OPTIMIZATION OF SPORE PRODUCTION

Six solid culture media were evaluated for their effectiveness in supporting KKKU215 spore production. All media were prepared with 1.5% (w/v) agar, and their respective compositions (% w/v) were as follows: ATCC Medium: 0.1 g tryptose, 0.1 g beef extract, 0.1 g yeast extract, 1.0 g glucose, and 0.01 g FeSO₄ (Dionigi et al. 1992), half-strength potato dextrose medium (½ PDA) (1.2 g potato dextrose broth powder (Difco™, France)) (Chouychai et al. 2022), inorganic salt starch medium (ISS) (1.0 g starch, 0.1 g K₂HPO₄, 0.1 g MgSO₄·7H₂O, 0.1 g NaCl, 0.2 g (NH₄)₂SO₄, 0.2 g CaCO₃, 0.01 g FeSO₄·7H₂O, 0.01 g MnCl₂·7H₂O, 0.01 g ZnSO₄·7H₂O) (Aoki et al. 2007), NA medium (2.8 g nutrient agar powder (Himedia™) (Yottakot & Leelavatcharamas 2019), glucose yeast-malt medium (GYM) (0.2 g CaCO₃, 1.0 g malt extract, 0.4 g yeast extract, 0.4 g glucose) (Boukaew et al. 2022), and modified Wakimoto's medium (MWM) (30 g potato, 0.5 g peptone, 2 g sucrose, 0.05 g Ca(NO₃)₂, 0.2 g Na₂HPO₄·2H₂O (Adhikari, Mew & Leach 1999). Potato infusion for MWM was prepared by boiling 30 g of diced potato in 100 mL distilled water for 15 min. A 100 μ L suspension of KKKU215 (~10⁶ spores/plate on day 0) was spread onto each medium and incubated at 37 °C; spore numbers were counted on days 5, 10, 15, and 20 using a hemacytometer.

The PB design, based on Liu et al. (2020), evaluated seven variables: beef extract, peptone, potato, sucrose, Ca(NO₃)₂, Na₂HPO₄·2H₂O (selected from MWM and NA media), and incubation time. These variables were assessed for their effects on KKKU215 spore production. The MWM and NA media, as well as incubation time, played a key role in achieving the highest spore production. Each variable was tested at three levels: low (-), central (0), and high (+), to ensure replication and accuracy. The levels were beef extract 1.0, 3.0, and 5.0 g/L; peptone 3.0, 5.0, and 7.0 g/L; potato 100.0, 300.0, and 500.0 g/L; sucrose 10.0, 20.0, and 30.0 g/L; Ca(NO₃)₂ 0.1, 0.5, 0.9 g/L; Na₂HPO₄·2H₂O 1.0, 2.0, and 3.0 g/L; and incubation time 10, 15, and 20 days. A 15-run PB design was used (Table 1), with

each run conducted in triplicate, and statistical analysis (p -value < 0.05) with a Pareto chart determined key factors. Significant factors were further optimized using the BB design. The starter culture and incubation temperature were fixed at 1% spore suspension and 37 °C, respectively.

Box–Behnken design (BB design) in RSM is a widely used technique for optimizing process parameters (Liu et al. 2025). In this study, the BB design was applied to determine optimal parameters for KKKU215 spore production. Three significant variables (p -value < 0.05) determined from the PB design, peptone (X_1), potato (X_2), and incubation time (X_3), were further investigated at low (-), central (0), and high (+) levels corresponding to 3.0, 7.0, and 11.0 g/L for peptone; 300.0, 500.0, and 700 g/L potato infusion equivalent; and 10, 15, and 20 days for incubation time, respectively, for a total of 17 experimental runs. All runs are being performed in triplicate, including five centre point replicates (Table 2). The potato infusion for each run was prepared by boiling the specified amount of fresh, peeled, and diced potato (corresponding to 300, 500, or 700 g) in 1 L of distilled water for 20 min. The resulting infusion was filtered through two layers of cheesecloth, and the volume of the filtrate was adjusted to exactly 1 L with distilled water before being used as the main component of the culture medium for each run. The experiment conditions followed those used in the screening experiment. A quadratic model was constructed from experimental data to describe the effects of these variables on spore production. The predicted optimal conditions were validated experimentally, confirming the model's accuracy.

STATISTICAL ANALYSIS

Data were analysed using SPSS version 29. An independent t-test was used to compare the means of seedling growth parameters, while one-way ANOVA with Tukey's HSD post-hoc test was applied to data from pot experiments and spore production assays. PB and BB design data were analysed using the trial version 23.1.8 (64-bit) of Stat-Ease® 360 (Stat-Ease, MN, USA). A one-sample t-test was conducted to evaluate the difference between the experimental mean spore count under optimized conditions and the corresponding predicted value obtained from the model. Statistical significance was set at $p < 0.05$.

RESULTS AND DISCUSSION

BIOCONTROL AND PLANT GROWTH-PROMOTING TRAITS

KKKU215 exhibited multiple biocontrol and plant growth-promoting traits, including five extracellular lytic enzyme activities, siderophore production, biofilm formation (Figure 1), nitrogen fixation, ammonia production, and IAA production (Figure 2). Maximum IAA and ammonia levels reached 13.03 µg/mL (with 5 mg/mL tryptophan) and 1.43 µmol/mL (after 7 days), respectively. These traits support KKKU215's potential as a biocontrol and growth-promoting agent. Similarly, Awla et al. (2017) reported ammonia, siderophore, amylase, protease, and lipase production by *Streptomyces* UPMRS4, while Kaari et al. (2022a) found biofilm formation in several *Streptomyces* strains. Kaari et al. (2022a) and Vurukonda, Giovanardi and Stefan (2018) further emphasized the

TABLE 1. PB design and results

Std	Run	Be (g/L)	Pe (g/L)	Po (g/L)	Su (g/L)	Ca (g/L)	Di (g/L)	In (day)	Number of spore (spores/plate)
14	1	3.0	5.0	300.0	20.0	0.5	2.0	15	5.012×10^9
13	2	3.0	5.0	300.0	20.0	0.5	2.0	15	4.956×10^9
1	3	5.0	7.0	100.0	30.0	0.9	3.0	10	1.900×10^7
7	4	5.0	3.0	100.0	10.0	0.9	1.0	20	2.100×10^7
10	5	1.0	7.0	500.0	30.0	0.1	1.0	10	2.900×10^9
11	6	5.0	3.0	500.0	30.0	0.9	1.0	10	1.800×10^8
9	7	5.0	7.0	500.0	10.0	0.1	1.0	20	9.950×10^9
3	8	5.0	3.0	500.0	30.0	0.1	3.0	20	2.400×10^8
4	9	1.0	7.0	100.0	30.0	0.9	1.0	20	1.980×10^9
2	10	1.0	7.0	500.0	10.0	0.9	3.0	20	9.898×10^9
12	11	1.0	3.0	100.0	10.0	0.1	1.0	10	1.100×10^7
6	12	1.0	3.0	100.0	30.0	0.1	3.0	20	3.030×10^7
5	13	1.0	3.0	500.0	10.0	0.9	3.0	10	9.800×10^8
15	14	3.0	5.0	300.0	20.0	0.5	2.0	15	5.120×10^9
8	15	5.0	7.0	100.0	10.0	0.1	3.0	10	1.500×10^7

The abbreviations Be, Pe, Po, Su, Ca, Di, and In represent Beef extract, Peptone, Potato, Sucrose, $\text{Ca}(\text{NO}_3)_2$, $\text{Na}_2\text{HPO}_4 \cdot 2\text{H}_2\text{O}$ and Incubation time, respectively

TABLE 2. BB design and results

Std	Run	Peptone (g/L)	Potato (g/L)	Incubation time (day)	Number of spore (spores/plate)	
					Experiment	Predicted
5	1	3.0	500.0	10	1.257×10^9	1.489×10^9
13	2	7.0	500.0	15	2.263×10^9	2.143×10^9
14	3	7.0	500.0	15	2.160×10^9	2.143×10^9
17	4	7.0	500.0	15	1.912×10^9	2.143×10^9
15	5	7.0	500.0	15	2.065×10^9	2.143×10^9
7	6	3.0	500.0	20	9.987×10^9	1.006×10^{10}
16	7	7.0	500.0	15	2.317×10^9	2.143×10^9
3	8	3.0	700.0	15	4.780×10^9	4.447×10^9
1	9	3.0	300.0	15	2.560×10^9	2.579×10^9
10	10	7.0	700.0	10	1.440×10^9	1.541×10^9
2	11	11.0	300.0	15	5.130×10^9	5.463×10^9
11	12	7.0	300.0	20	1.130×10^{10}	1.120×10^{10}
12	13	7.0	700.0	20	1.300×10^{10}	1.325×10^{10}
8	14	11.0	500.0	20	1.510×10^{10}	1.487×10^{10}
6	15	11.0	500.0	10	1.510×10^9	1.428×10^9
4	16	11.0	700.0	15	6.320×10^9	6.300×10^9
9	17	7.0	300.0	10	1.142×10^9	8.901×10^8

role of *Streptomyces* spp. in biocontrol and plant growth promotion via siderophores, lytic enzymes, IAA, ammonia, and nitrogen fixation.

SEEDLING GROWTH

On day 30, tomato seedlings from seeds treated with K KU215 spores showed significantly greater shoot and root lengths (15.21 cm and 16.65 cm), as well as higher fresh and dry weights (29.38 mg and 2.12 mg), compared to the control (Figure 3(A) and 3(B)). The leaf size was also larger (Figure 3(C)). Similarly, *Streptomyces rochei* BT02 has also significantly stimulated shoot and root lengths, and fresh and dry weights of tomato seedlings (Devi, Sharma & Manhas 2022).

POT EXPERIMENT

At 56 days after planting, treatment with K KU215 spores (T2) significantly reduced the severity of bacterial wilt caused by RsDOA-BCC1954, resulting in a disease index of 12.48%, compared to the pathogen control (T4), which exhibited a disease index of 95.75% (Figure 4(A)). In T4, disease symptoms appeared within 7 days, starting with shoot tip wilting and progressing without leaf yellowing (Figure 4(D)). This study suggests that the ability of K KU215 to control BW disease may be attributed to its production of a variety of enzymes, including lipase, amylase, protease, cellulase, and chitinase, as well as

siderophores and biofilms. Kaari et al. (2022a), Lee et al. (2023), and Tan et al. (2006), found that these compounds effectively inhibit the growth of Rs, thereby controlling bacterial wilt disease. These findings align with other studies on *Streptomyces* species, which have shown a role in disease prevention. Enzymes produced by *Streptomyces* species help break down complex molecules and inhibit pathogens, and siderophores sequester iron, depriving pathogens of this essential nutrient (Khan et al. 2023), while biofilms produced by *Streptomyces* species inhibit pathogens by forming protective barriers around plant roots and surfaces, preventing pathogen colonization (Kaari et al. 2022b). Kaari et al. (2022a) found that *Streptomyces* sp. UT4A49 produces amylase, protease, cellulase, and siderophores, which inhibit the pathogen Rs.

K KU215 also significantly promoted tomato growth, as indicated by increased shoot and root lengths, as well as greater fresh and dry biomass in treatments T1 and T2, compared to the controls (T3 and T4) (Figures 4(B), 4(C)). This growth enhancement may be attributed to the abilities of K KU215 in nitrogen fixation, ammonia production, and IAA synthesis, i.e., factors that potentially stimulate plant growth. Similarly, Devi, Sharma and Manhas (2022) found that *Streptomyces* sp. SP5 significantly enhances shoot and root length, as well as the fresh and dry weight of plants through IAA production, while Du et al. (2022) demonstrated that *Streptomyces albidoflavus* St-220 possesses nitrogen fixation, which significantly increases

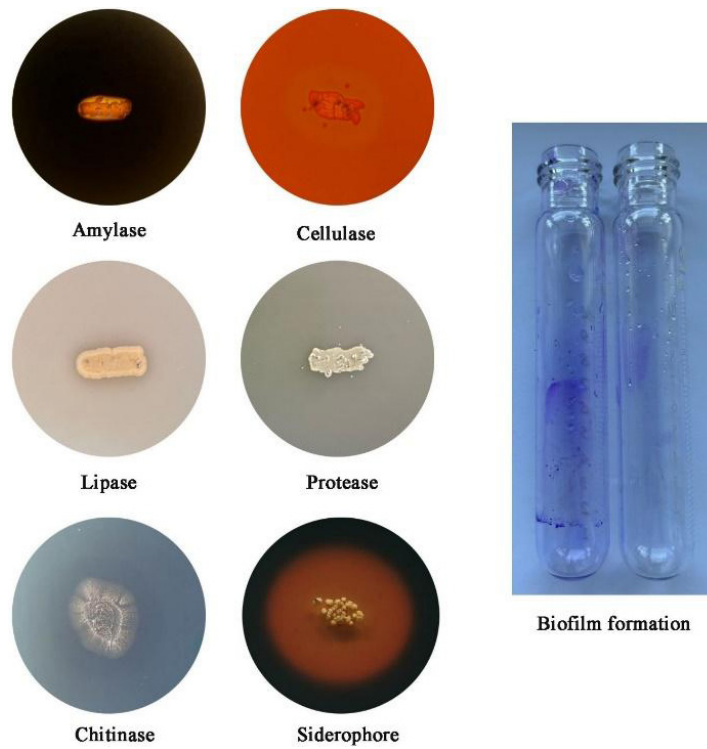


FIGURE 1. Characterization of extracellular lytic enzymes, siderophores, and biofilm production by KKU215

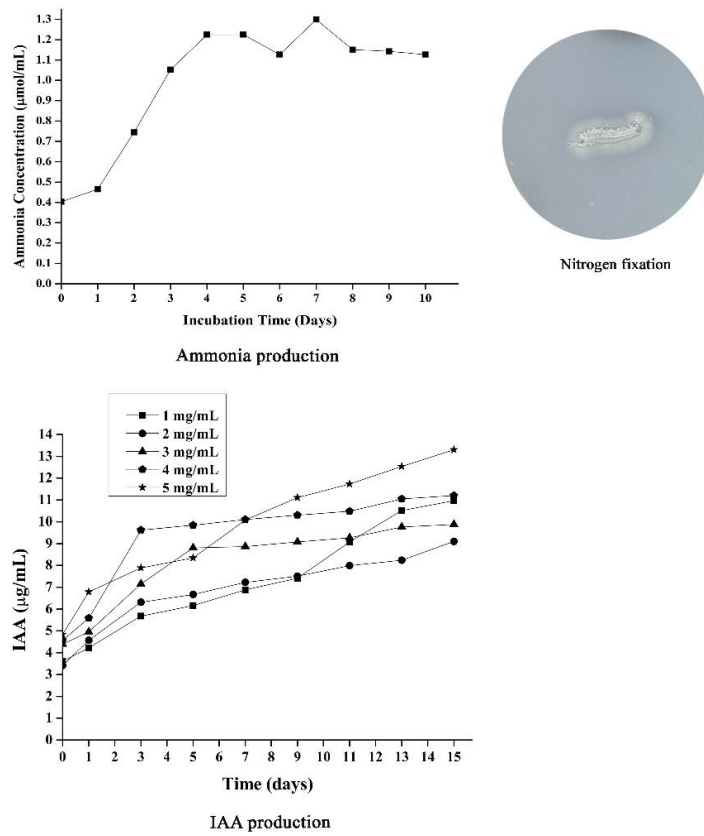


FIGURE 2. Characterization of nitrogen fixation, ammonia, and IAA production by KKU215

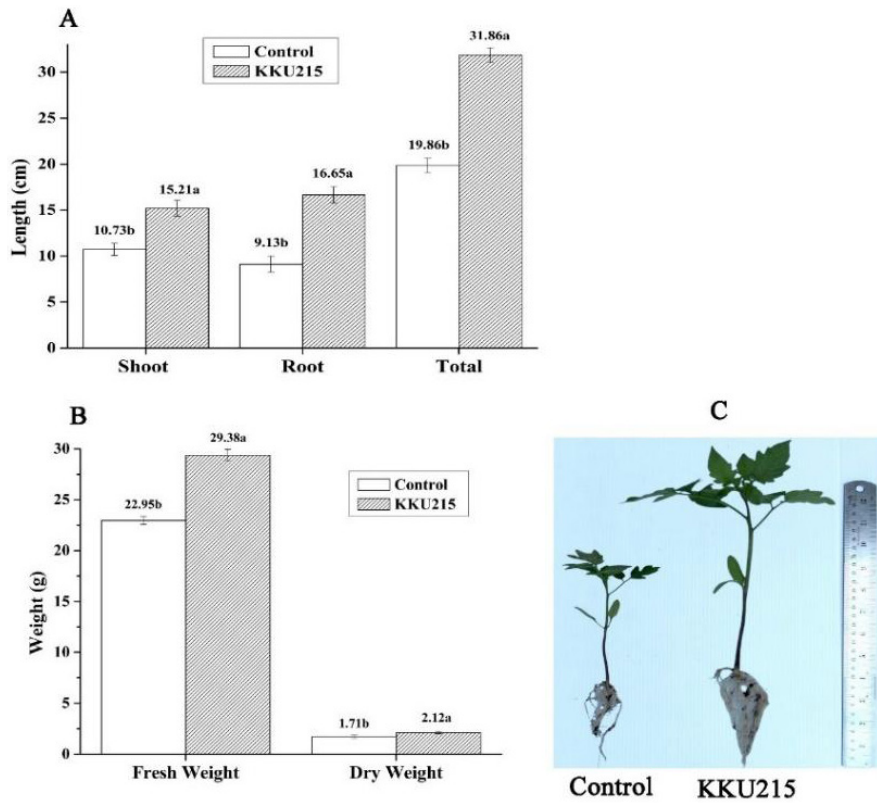


FIGURE 3. Effect of KKU215 on tomato seedling growth 30 days after planting in tray experiment. (A) shoot and root lengths, (B) fresh and dry weights, and (C) photographs of seedlings

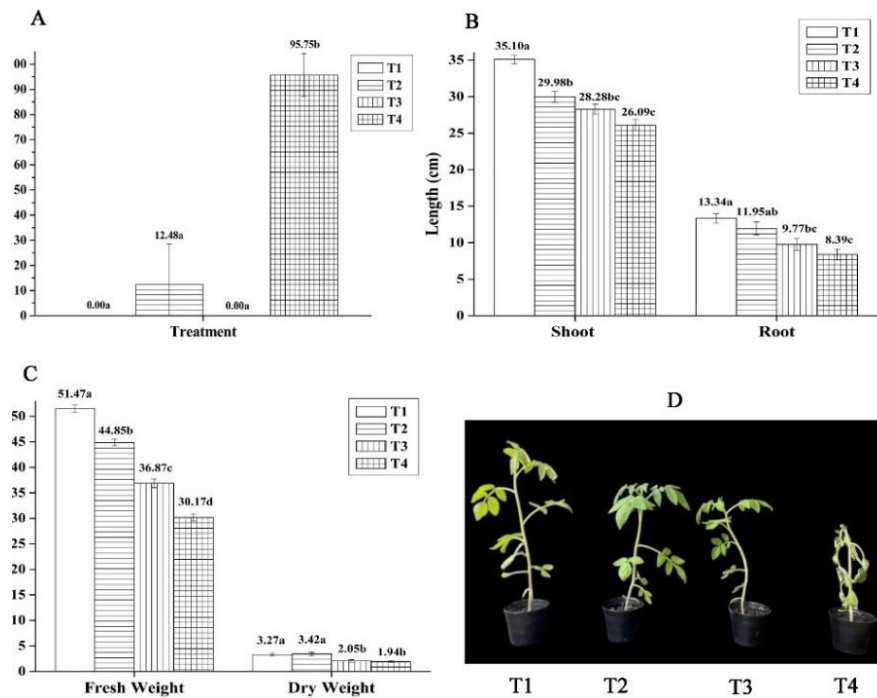


FIGURE 4. Efficacy of KKU215 in controlling BW disease and promoting tomato growth 56 days after planting in a pot experiment. (A) disease index, (B) shoot and root lengths, (C) fresh and dry weights, and (D) photographs of seedlings

the total fresh and dry weight of plant growth. These findings align with other studies that highlight the role of *Streptomyces* species in promoting plant growth. Nitrogen fixation and ammonia production increase the availability of nitrogen, an essential nutrient for plant growth, while IAA promotes plant development by enhancing cell elongation and root growth (Vurukonda, Giovanardi & Stefan 2018).

OPTIMIZATION OF PARAMETERS FOR ENHANCING SPORE PRODUCTION

Spore production by KKU215 was significantly affected by the type of medium and incubation time. Spore production increased rapidly within the first 10 days in most media, except for ISS and ½ PDA, which exhibited minimal increases. This was likely due to the lower nutrient content in these media, which are insufficient to support robust sporulation compared to nutrient-rich media. From day 10 to day 20, spore production continued to rise steadily. After 20 days of incubation, MWM and NA media supported the highest spore yields ($9.51 \pm 0.18 \times 10^9$ and $9.28 \pm 0.21 \times 10^9$ spores/plate, respectively), significantly higher than other media ($p < 0.05$) (Table 3). The study showed that MWM and NA medium, and incubation time all influenced spore production. The six media components (beef extract, peptone, potato, sucrose, $\text{Ca}(\text{NO}_3)_2$, and $\text{Na}_2\text{HPO}_4 \cdot 2\text{H}_2\text{O}$), along with incubation time, were selected for screening using the PB design to identify key factors affecting spore production.

The PB design was used to screen seven factors affecting the culture conditions of KKU215. In Table 4, Analysis of Variance (ANOVA) indicated that peptone (p -value = 0.0066), potato (p -value = 0.0085), sucrose (p -value = 0.0352), and incubation time (p -value = 0.0200) are significant factors (p -value < 0.05). The Pareto chart of standardized effects (Figure 5) showed that peptone, potato, and incubation time significantly exceeded the reference line, confirming their statistical significance ($p < 0.05$) in enhancing spore production. These factors were therefore selected for further optimization. In contrast, sucrose, which had a negative effect, was excluded from the optimization process. Consequently, peptone, potato, and incubation time were selected for further optimization using the RSM with BB design.

Peptone, a rich nitrogen source, plays a crucial role in promoting bacterial growth and spore formation (Hsieh & Labbe 2007; Liggins, Ramirez & Abel-Santos 2023). Potato serves as a source of both carbohydrates and minerals such as Ca, Fe, and Mg (Beals 2019), which are present in adequate amounts to support the transition to sporulation in *Streptomyces* (Yague et al. 2013). Incubation time increases the possibility of sporulation by allowing bacteria to undergo the transition from vegetative growth to the sporulation phase (Aouadhi et al. 2016). Meanwhile, sucrose was excluded from further optimization due to its negative effect on spore production. This may be attributed to the fact that sucrose acts as a single carbon

source, whereas potato provides not only carbohydrates but also essential minerals, and peptone acts as a complex nitrogen source, thereby supporting spore formation more effectively.

Following the PB design results, a BB design was carried out with treatment 5 as the central point. Peptone, potato, and incubation time were selected as independent variables in a three-factor, three-level test, with spore count as the response variable. A total of 17 experimental trials were performed, and the results are presented in Table 2. The regression analysis with quadratic terms, used to describe the relationship between the independent variables and the response variable, resulted in the following equation:

$$Y = 3.20475 \times 10^{10} - 1.67806 \times 10^9 X_1 - 2.85128 \times 10^7 X_2 - 3.605551 \times 10^9 X_3 - 3.21875 \times 10^5 X_1 X_2 + 6.07531 \times 10^7 X_1 X_3 + 3.50475 \times 10^5 X_2 X_3 + 8.74152 \times 10^7 X_1^2 + 28888.90625 X_2^2 + 1.36865 \times 10^8 X_3^2$$

where Y represents the predicted number of spores, and X_1 , X_2 , and X_3 represent peptone, potato, and incubation time, respectively.

Table 5 shows that the ANOVA results indicate the model is highly significant ($p < 0.0001$), with no significant lack of fit ($p = 0.0540$), indicating a good fit between the model and the experimental data. The model's high R^2 (0.9982), Adjusted R^2 (0.9959), and Predicted R^2 (0.9759) values confirm its high accuracy and predictive power. Among the tested variables, peptone, potato, and incubation time had significant linear and quadratic effects on spore production ($p < 0.05$). Interaction between peptone and incubation time, as well as between potato and incubation time, also showed significant influence. In contrast, the interaction between peptone and potato was not significant ($p = 0.1209$), meaning that they were statistically insignificant in affecting the response. This suggests that changes in both factors may not have a significant statistical impact on spore production when assessed together.

The three-dimensional response surface (3D) plots and contour plots are demonstrated in Figure 6. Figure 6(A) presents the 3D and contour plots illustrating the interaction between peptone and potato. The plots show that increasing both components to their highest levels results in optimal spore production. High concentrations of peptone provide the necessary nitrogen, which supports rapid cell division and facilitates the transition into the stationary phase, where sporulation is triggered (Hsieh & Labbe 2007; Liggins, Ramirez & Abel-Santos 2023). Likewise, high levels of potato enhance bacterial proliferation by supplying carbohydrates and nutrients. As the population increases, the nutrients become limited, triggering sporulation. The 3D and contour plots in Figure 6(B) (peptone and incubation time) and Figure 6(C) (potato and incubation time) show that higher levels of either peptone or potato, when combined with longer incubation time, significantly enhance spore production. This prolonged incubation time increases the possibility of sporulation by allowing bacteria to undergo the transition from vegetative growth to the sporulation phase (Aouadhi et al. 2016).

TABLE 3. Spore production of KKKU215 under various media and incubation times

Media	Number of spore (spores/plate)				
	0 Day	5 Day	10 Day	15 Day	20 Day
MWM	1×10^6	$(9.13 \pm 0.19) \times 10^8$ a	$(1.89 \pm 0.03) \times 10^9$ a	$(6.23 \pm 0.14) \times 10^9$ a	$(9.51 \pm 0.18) \times 10^9$ a
NA	1×10^6	$(2.61 \pm 0.12) \times 10^8$ b	$(1.66 \pm 0.09) \times 10^9$ b	$(6.02 \pm 0.09) \times 10^9$ b	$(9.28 \pm 0.21) \times 10^9$ a
GYM	1×10^6	$(1.04 \pm 0.14) \times 10^8$ c	$(3.48 \pm 0.35) \times 10^8$ c	$(5.34 \pm 0.36) \times 10^8$ c	$(6.17 \pm 0.07) \times 10^8$ b
ATCC	1×10^6	$(1.69 \pm 0.12) \times 10^7$ d	$(4.21 \pm 0.46) \times 10^7$ d	$(3.06 \pm 0.22) \times 10^8$ d	$(5.16 \pm 0.10) \times 10^8$ bc
ISS	1×10^6	$(3.38 \pm 0.23) \times 10^6$ d	$(7.84 \pm 0.79) \times 10^6$ d	$(1.25 \pm 0.10) \times 10^8$ de	$(2.30 \pm 0.09) \times 10^8$ cd
½ PDA	1×10^6	$(1.77 \pm 0.15) \times 10^6$ d	$(5.18 \pm 0.13) \times 10^6$ d	$(1.65 \pm 0.16) \times 10^7$ e	$(2.83 \pm 0.13) \times 10^7$ d

Different letters within each column indicate statistically significant differences between means ($p < 0.05$)

TABLE 4. ANOVA for PB design results

Source	Sum of squares	Df.	Mean square	F-value	p-value
Beef extract	2.407×10^{18}	1	2.407×10^{18}	0.8787	0.3847
Peptone	4.523×10^{19}	1	4.523×10^{19}	16.51	0.0066*
Potato	4.060×10^{19}	1	4.060×10^{19}	14.82	0.0085*
Sucrose	2.009×10^{19}	1	2.009×10^{19}	7.33	0.0352*
Ca(NO ₃) ₂	3.831×10^{14}	1	3.831×10^{14}	0.0001	0.9909
Na ₂ HPO ₄ ·2H ₂ O	1.241×10^{18}	1	1.241×10^{18}	0.4530	0.5260
Incubation time	2.703×10^{19}	1	2.703×10^{19}	9.87	0.0200*
Residual	1.644×10^{19}	6			
Corrected total	1.724×10^{20}	14			

*Represents significant, Df: degree of freedom

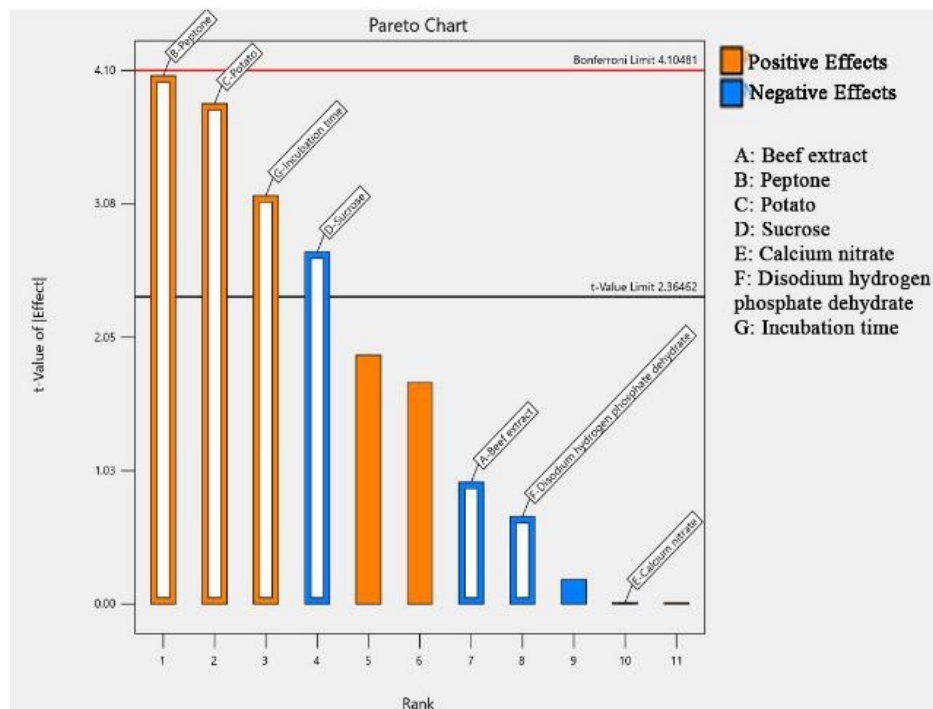


FIGURE 5. Pareto chart of standardized effects on KKKU215 spore production

TABLE 5. ANOVA for BB design results

Source	Sum of squares	Df.	Mean square	F-value	<i>p</i> -value
Model	3.324×10^{20}	9	3.693×10^{19}	433.85	< 0.0001*
Peptone	1.122×10^{19}	1	1.122×10^{19}	131.86	< 0.0001*
Potato	3.656×10^{18}	1	3.656×10^{18}	42.95	0.0003*
Incubation time	2.424×10^{20}	1	2.424×10^{20}	2847.97	< 0.0001*
Peptone*Potato	2.652×10^{17}	1	2.652×10^{17}	3.12	0.1209
Peptone*Incubation time	5.906×10^{18}	1	5.906×10^{18}	69.38	< 0.0001*
Potato*Incubation time	4.913×10^{17}	1	4.913×10^{17}	5.77	0.0473*
Peptone*Peptone	8.237×10^{18}	1	8.237×10^{18}	96.77	< 0.0001*
Potato*Potato	5.622×10^{18}	1	5.622×10^{18}	66.05	< 0.0001*
Incubation time*	4.929×10^{19}	1	4.929×10^{19}	579.13	< 0.0001*
Incubation time					
Residual	5.958×10^{17}	7	8.512×10^{16}		
Lack of Fit	4.915×10^{17}	3	1.638×10^{17}	6.28	0.0540
Pure Error	1.043×10^{17}	4	2.607×10^{16}		
Cor Total	3.330×10^{20}	16			
$R^2 = 0.9982$					
Adjusted $R^2 = 0.9959$					
Predicted $R^2 = 0.9759$					

*Represents significant, Df: degree of freedom

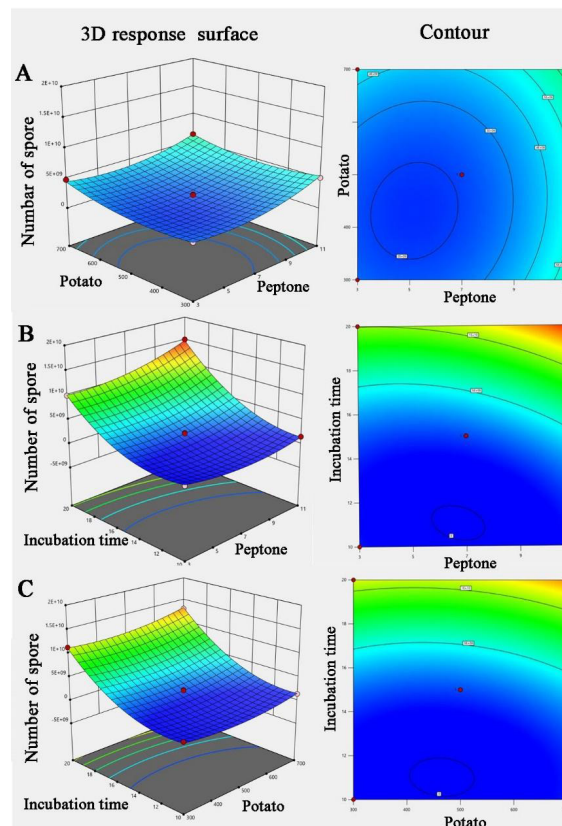


FIGURE 6. The 3D plots and contour plots showing the effect of (A) potato and peptone, (B) incubation time and peptone, and (C) incubation time and potato

CONCLUSIONS

This study highlights the biocontrol and plant growth-promoting potential of K KU215. The strain produced extracellular lytic enzymes, siderophores, biofilms, IAA, and ammonia, and fixed nitrogen, supporting its biocontrol and plant growth-promoting roles. K KU215 reduced BW disease index and improved tomato growth, as shown by increased shoot and root lengths, as well as fresh and dry weights in tray and pot experiments. MWM and NA media yielded the highest spore production of K KU215 ($p < 0.05$), and their components were screened using PB design, showing peptone, potato, and incubation time as key factors. RSM optimization showed the optimal conditions: 7 g/L peptone, 700 g/L potato infusion equivalent, and 20 days of incubation, resulting in a 46% increase in spore yield, closely matching predicted values. This study demonstrates the potential of K KU215 as a sustainable biocontrol agent and plant growth promoter by effectively reducing the BW disease incidence and enhancing tomato seedling growth. Additionally, it showed high spore production under optimized culture conditions. These findings provide a strong foundation for developing K KU215-based bioproducts for agricultural applications. Future research should focus on field trials and commercial formulation development to promote sustainable disease management and reduce chemical pesticide use in crop production.

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