







Influence of Environmental Factors on Biofilm Formation by Heat-resistant Spore-forming *Bacillus cereus* TGS11.1



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Abstract:

Introduction/Objective: This study evaluates the synergistic influence of environmental factors on the biofilm formation capacity of the heat-resistant, spore-forming pathogen *Bacillus cereus* TGS11.1 within milk-processing contexts.

Methods: Response Surface Methodology with a Central Composite Design (RSM-CCD) was implemented to quantify the multifactorial effects of temperature (30-70 °C), pH (4-8), lactose concentration (2-6%), and incubation time (12-60 h) on biofilm development using optical density measurements and phenotypic motility assays.

Results: Temperature emerged as the dominant determinant of biofilm formation, with peak productivity and OD570 values occurring at 40 °C, correlating with maximum swarming motility and the establishment of a dense three-dimensional (3D) architectural network.

Discussion: The observed sensitivity of *B. cereus* to thermal gradients and nutrient availability underscores the need for multifactorial intervention strategies to disrupt the structural resilience of the biofilm matrix during dairy production.

Conclusion: Mitigating recontamination risks in dairy workflows necessitates the strict avoidance of thermal niches near 40 °C to inhibit the optimal physiological conditions for robust biofilm formation by *B. cereus* TGS11.1.

Keywords: *Bacillus cereus*, Biofilm formation, Optical density, Milk processing, Environmental factors, Response surface methodology.

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1. INTRODUCTION

The contamination of raw milk by *Bacillus cereus* constitutes a critical biosafety risk primarily due to the production of heat resistance spores that facilitate enzymatic spoilage and the synthesis of emetic and diarrheal toxins [1]. This microbiological challenge is intensified by the capacity of *B. cereus* for biofilm

formation on dairy contact surfaces, where the protective extracellular matrix confers significantly higher resistance to detergents and disinfectants compared to planktonic populations. Notably, biofilm-associated spores exhibit heightened thermal resistance, leading to post-pasteurization recontamination and diminished product shelf life, which subsequently incurs substantial economic losses for the dairy industry [2]. Characterizing the non-

linear interactions between multiple environmental factors, specifically nutrient concentration, temperature, incubation time, and pH, is imperative for understanding biofilm formation under actual milk processing conditions. While the hazards of *B. cereus* are well-established [3, 4], current literature has predominantly investigated these variables in isolation, leaving their simultaneous synergistic effects largely uncharacterized [4]. Furthermore, the assessment of colony morphology provides essential visual metrics regarding the structural complexity and architectural integrity of the biofilm, facilitating the evaluation of bacterial pathogenicity and risk levels within dairy production workflows [5]. This study utilizes Response Surface Methodology with a Central Composite Design (RSM-CCD) to model the multi-factorial influence of environmental factors on the biofilm formation of *B. cereus* TGS11.1 [6]. The resulting data establish a deterministic framework for optimizing control strategies to mitigate microbial persistence and resistance during the processing and preservation of milk and dairy products [7].

2. MATERIALS AND METHODS

The study utilized the *Bacillus cereus* TGS 11.1 strain,

which was previously isolated from raw milk samples in the Mekong Delta, Vietnam, as reported in our preceding study published in this journal. This specific isolate was selected for its superior heat-resistant, spore-forming capability among the collected strains. The use of TGS 11.1 ensures experimental consistency and allows for a deeper investigation into its biofilm-forming characteristics under the currently defined conditions.

2.1. Experimental Design

Biofilm formation of the *Bacillus cereus* TGS 11.1 strain was quantified using OD values. The investigated factors, including lactose concentration, temperature, incubation time, and pH, were selected based on small-scale industrial production processes to ensure that all parameters are adjustable under current operational conditions. RSM-CCD was applied, comprising 4 factors and 5 levels (refer to Table 1). This approach significantly reduces the number of required experiments compared to a full factorial design. Furthermore, it avoids extreme treatment levels that are often impractical for small-scale laboratory settings. The design aligns with actual production conditions, ensuring that the model accurately reflects its potential for future practical applications.

Table 1. Experimental design matrix and actual results.

Runs	Factors				OD ₅₇₀
	Conc (%)	Temp (°C)	Time (h)	pH	
1	3	40	24	5	0.019
2	5	40	24	5	0.148
3	3	60	24	5	0.020
4	5	60	24	5	0.108
5	3	40	48	5	0.179
6	5	40	48	5	0.242
7	3	60	48	5	0.058
8	5	60	48	5	0.064
9	3	40	24	7	0.052
10	5	40	24	7	0.979
11	3	60	24	7	0.013
12	5	60	24	7	0.077
13	3	40	48	7	0.266
14	5	40	48	7	0.706
15	3	60	48	7	0.033
16	5	60	48	7	0.035
17	2	50	36	6	0.073
18	6	50	36	6	0.066
19	4	30	36	6	0.713
20	4	70	36	6	0.502
21	4	50	12	6	0.260
22	4	50	60	6	0.119
23	4	50	36	4	0.065
24	4	50	36	8	0.074
25	4	50	36	6	0.063
26	4	50	36	6	0.061
27	4	50	36	6	0.072
28	4	50	36	6	0.060
29	4	50	36	6	0.071
30	4	50	36	6	0.069

2.2. Quantification of Biofilm Formation Capacity under Various Experimental Conditions Bacterial culture and Sample Preparation

The *Bacillus cereus* TGS11.1 strain was activated on Luria-Bertani (LB) agar (10 g L⁻¹ tryptone, 5 g L⁻¹ yeast extract, 10 g L⁻¹ NaCl) and pre-cultured in 20 mL Brain Heart Infusion (BHI; Oxoid) medium. Secondary cultures (50 mL BHI supplemented with lactose (2%–6%)) were adjusted to pH levels (4–8) with initial OD₀ = 0.1 and incubated at 37°C and 200 rpm for 3 h. Subsequently, 200 µL aliquots of fermentation broth were transferred to 96-well plates and incubated at temperatures of 30°C–70°C for durations of 12 h–60 h. To ensure biological and technical reproducibility, all experimental conditions were performed in three independent sessions (n=3), with each session comprising triplicate samples.

2.3. Evaluation of Biofilm Formation

Biofilm formation was quantified using the crystal violet staining method [3] with modifications. An aliquot of 200 µL of each sample was transferred into a 96-well plate and incubated statically at 37°C for 24 h. After washing with 200 µL PBS 1X, biofilms were fixed with 100 µL methanol for 15 min, stained with 1% crystal violet (250 µL), and subsequently solubilized with 250 µL of 95% ethanol. OD₅₇₀ was measured, and biofilm production was classified relative to ODC (ODC + 3×SD): non-producer (OD ≤ ODC), weak (ODC < OD ≤ 2×ODC), moderate (2×ODC < OD ≤ 4×ODC), and strong (OD > 4×ODC). ODC (OD control) refers to the mean absorbance of negative control wells; SD denotes standard deviation. All reported values are presented as mean ± standard deviation (SD) of the three independent replicates. Wells containing sterile BHI medium served as negative controls.

2.4. Morphological Characterization of the *Bacillus Cereus* TGS 11.1 Strain's Colony Biofilm under Different Temperature Conditions

To perform this survey, liquid culture medium was prepared by transferring 20 mL Brain Heart Infusion Broth (BHI) into an Erlenmeyer flask and incubated at 37°C and 220 rpm for 24 h. Aliquots (3 µL) of the standardized overnight culture (10⁶ CFU/mL) were spotted onto BHI agar plates (1.5% agar) to develop colony biofilms. The plates were sealed with Parafilm and incubated at different temperatures (30°C, 40°C, 50°C, 60°C, and 70°C) for 24 h. To ensure morphological consistency and statistical reliability, 10 colonies per treatment were monitored across three independent experimental sessions (n=3). Colony morphology at various temperatures was examined and documented using a DZ.1105 stereo zoom microscope equipped with a dedicated digital imaging system (Euromex, Arnhem, The Netherlands). Colony diameters and structural complexity were quantified using ImageJ software, and significant differences between treatments were evaluated using one-way ANOVA ($p < 0.05$).

2.5. Swarming Motility at Different Temperatures

Swarming motility was assessed across a range of temperatures (30°C–70°C). Briefly, 10 µL of a standardized overnight culture (10⁶ CFU/mL) of *Bacillus cereus* TGS11.1 was inoculated at the center of semi-solid BHI agar plates (0.8% agar) supplemented with 5% lactose at pH 7. To ensure statistical reliability, swarming assays were performed in three independent experimental sessions (n=3), with each session utilizing triplicate plates. The plates were incubated at 30°C, 40°C, 50°C, 60°C, and 70°C for 36 h. Swarming zone diameters were measured in two perpendicular directions using a digital caliper. The mean values (mm) were reported as mean ± standard deviation (SD). Statistical differences in swarming migration between temperature treatments were determined using one-way ANOVA followed by Tukey's post-hoc test ($p < 0.05$).

2.6. Statistical Analysis

All experimental sessions were conducted in three independent replicates (n=3), with each treatment performed in triplicate to ensure biological and technical reproducibility. Data from the Response Surface Methodology (RSM) were processed using Design-Expert software (version 13.0). Analysis of Variance (ANOVA) was employed to evaluate the statistical significance of the quadratic model, individual factors, and their synergistic interactions. The fitness of the model was rigorously assessed using the coefficient of determination (R²), adjusted R², and the Lack of Fit test (where $p > 0.05$ indicated a non-significant lack of fit, confirming model adequacy). Furthermore, all graphical representations, including response surface plots, bar charts, and error bars, were generated using R software (version 4.5.1). For all tests, a p -value < 0.05 was considered statistically significant, and results are expressed as mean ± standard deviation (SD) [8–13].

3. RESULTS AND DISCUSSION

In this study, the variables were selected to align with the experimental conditions, enabling an evaluation of both the individual effects and the interaction terms of the variables within the investigated design space. Table 1 details the Central Composite Design (CCD) matrix along with the corresponding actual responses (OD₅₇₀ values) obtained from the 30 experimental runs.

3.1. Experimental Validation of Optimized Conditions for the *Bacillus Cereus* TGS 11.1 Strain's Biofilm Formation

Validation of the CCD experimental matrix established the optimal parameters for biofilm formation in *Bacillus cereus* TGS 11.1 through integrated spectrophotometric quantification and phenotypic microtiter plate analysis. Quantitative assessments confirmed peak biofilm density at 24 h incubation, 40°C, 5% lactose, and pH 7.0, yielding a maximum optical density (OD₅₇₀) of 0.979. Statistical significance ($p < 0.0001$) underscores that this optimal intersection represents a deterministic synergistic interplay

of environmental factors rather than stochastic variation (Fig. 1). Differential biomass accumulation in 96-well microtiter plates corroborated the statistical model through visible turbidity and surface aggregate density at the identified optima. Pre-staining observations revealed a significant contrast in cellular density between the optimal treatment and cultures subjected to acidic stress (pH 5.0) or limited carbon availability (Fig. 2a). Subsequent crystal violet (CV) staining demonstrated intense, uniform coloration in optimal wells, whereas control rows (3% lactose or pH 5.0) exhibited negligible dye uptake, indicating that peak adherence and architectural stability are contingent upon these specific parameters (Fig. 2b). The high congruence between morphological maturation and quantitative data ($p < 0.0001$) identifies the 40°C, pH 7.0, and 5% lactose intersection as a biological “sweet spot” for *B. cereus* TGS 11.1. At this physiological nexus, neutral pH and high carbon availability likely facilitate extracellular polymeric substance (EPS) synthesis by neutralizing cell surface charges and promoting initial attachment, as evidenced by the dense CV-stained pellicles (Fig. 2b). Conversely, the significant biomass reduction at pH 5.0 identifies acidic stress as a primary inhibitor, establishing environmental neutrality as a prerequisite for colonization. These findings extend current frameworks; the 40°C optimum aligns with reported thermal thresholds [14], while the pH 7.0 requirement confirms a physiological preference for neutrality despite a broad tolerance range [14]. Furthermore, the 5% lactose requirement of strain TGS 11.1, exceeding the 3% reported for other *Bacillus* biofilms [15], suggests a more robust carbon-induced EPS induction, underscoring the specialized persistence and resistance of this isolate in high-nutrient milk-processing environments.

3.2. Effects of Individual Factors on Biofilm Formation

The individual influence of each environmental factor on the biofilm formation capacity of *Bacillus cereus* TGS 11.1 was quantified using the Response Surface Methodology (RSM-CCD) framework. Initial ANOVA indicated that while incubation time ($p = 0.8675$) and initial pH ($p = 0.0664$) fell outside the standard 95% significance threshold, the discernible trend exhibited by pH in Fig. (3) necessitated its retention for model refinement. Consequently, the quadratic model was optimized to incorporate three primary variables: lactose concentration (A), temperature (B), and pH (D) (Table 2). The refined model confirmed that the selected parameters significantly modulated the optical density (OD) of the resulting biofilms as detailed in Table 2 and Fig. 3. Lactose concentration (A) exerted a significant positive linear effect ($F = 7.67$; $p = 0.0112$), driving a progressive increase in biofilm density as concentrations rose from 2% to 6%. Conversely, incubation time demonstrated negligible impact on the response, characterized by a stationary trend across the 12–60 h interval. Furthermore, pH (D) exhibited a significant positive linear correlation with OD ($F = 4.74$; $p = 0.0404$), following an upward trajectory from pH 4.0 to 8.0. Among the investigated variables, temperature (B) functioned as the dominant factor governing the biofilm formation process through a profound nonlinear influence (B: $F = 17.90$, $p = 0.0003$; B²: $F = 29.21$, $p < 0.0001$). Biofilm OD₅₇₀ values declined within the 30–55°C range before increasing again up to 70°C. This U-shaped phenotypic response indicates a local optimum and highlights the significant thermal sensitivity of the isolate under milk-processing conditions.

Table 2. Analysis of variance (ANOVA).

Source	Sum of Squares	df	Mean Square	F-value	p-value	Remark
Model	1.29	7	0.1838	11.64	< 0.0001	-
A-Conc	0.1211	1	0.1211	7.67	0.0112	significant
B-Temp	0.2828	1	0.2828	17.90	0.0003	highly significant
C-Time	0.0006	1	0.0006	0.0288	0.8675	not significant (removed)
D-pH	0.0749	1	0.0749	3.92	0.0664	marginally significant (full model)
D-pH	0.0749	1	0.0749	4.74	0.0404	significant
B.B	0.4614	1	0.4614	29.21	< 0.0001	highly significant
A.B	0.1223	1	0.1223	7.74	0.0109	significant
A.D	0.0822	1	0.0822	5.21	0.0326	significant
B.D	0.1419	1	0.1419	8.99	0.0066	highly significant
Residual	0.3475	22	0.0158	-	-	-

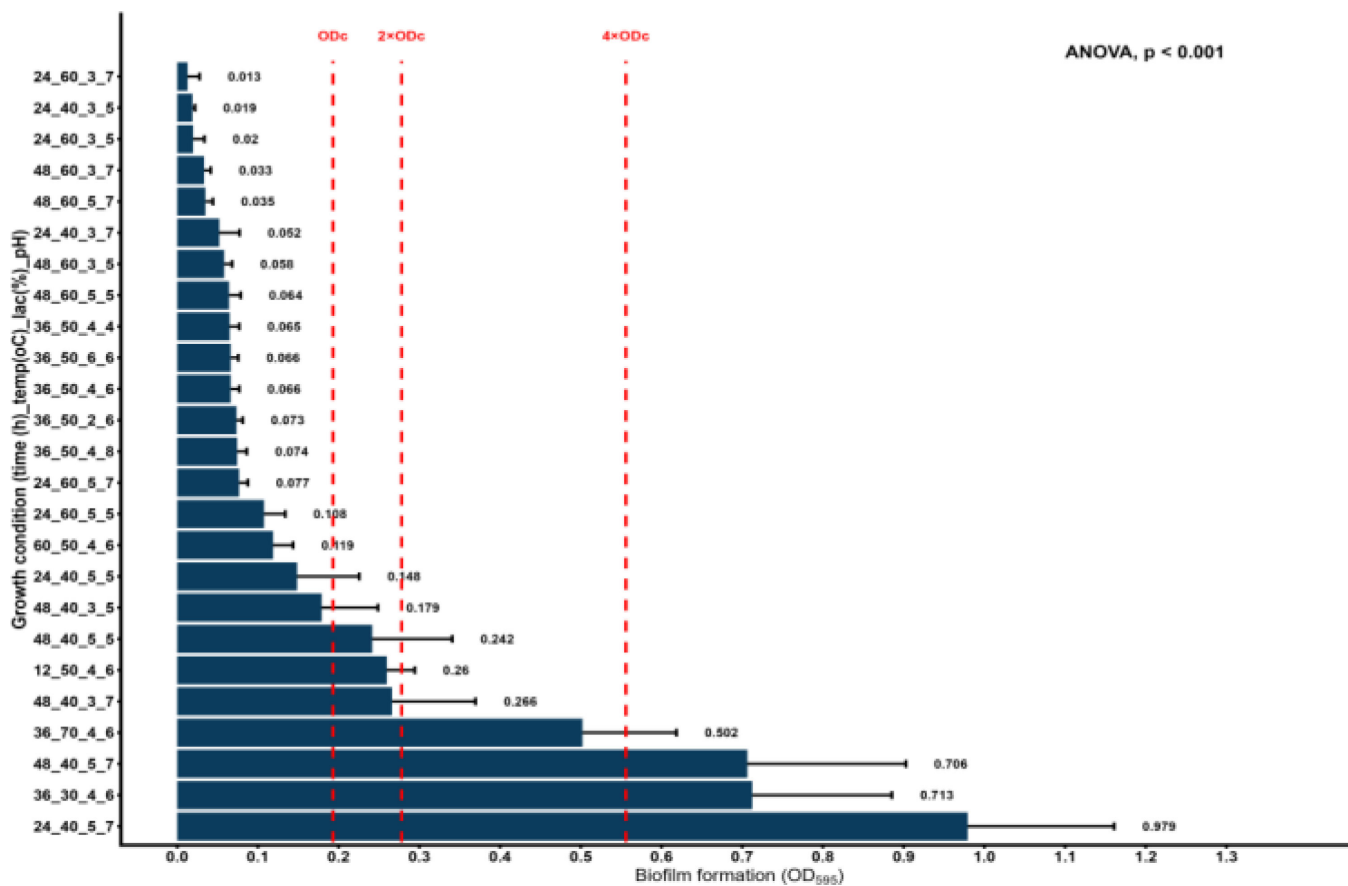


Fig. (1). The optimal conditions for biofilm formation by the *Bacillus cereus* TGS 11.1 strain.

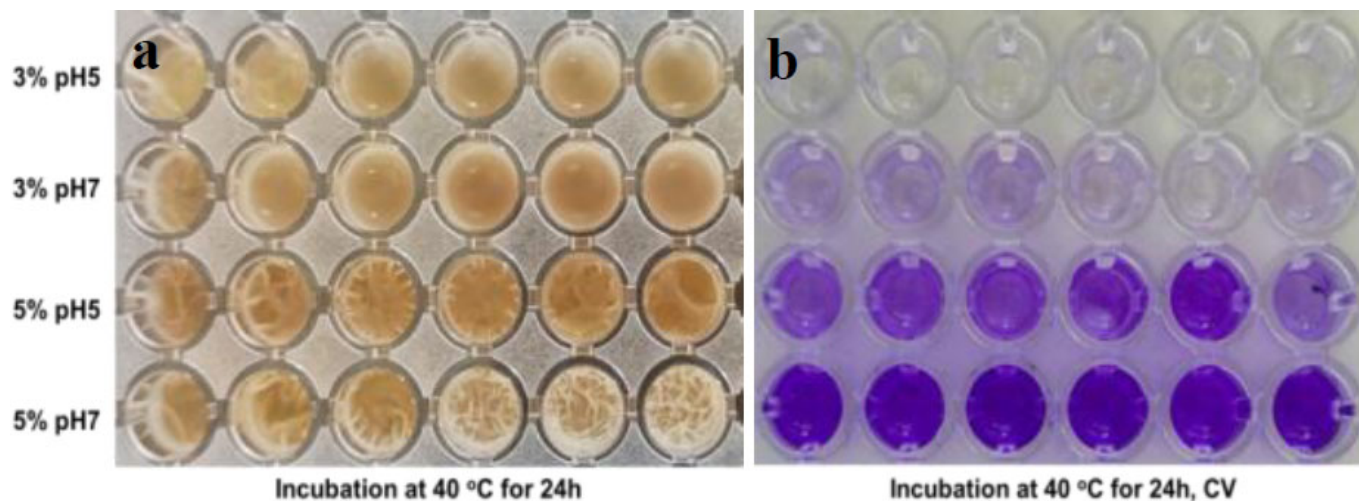


Fig. (2). Biofilm formation under the experimentally optimized condition: a) visually biofilm formation; b) biofilm formation after CV staining.

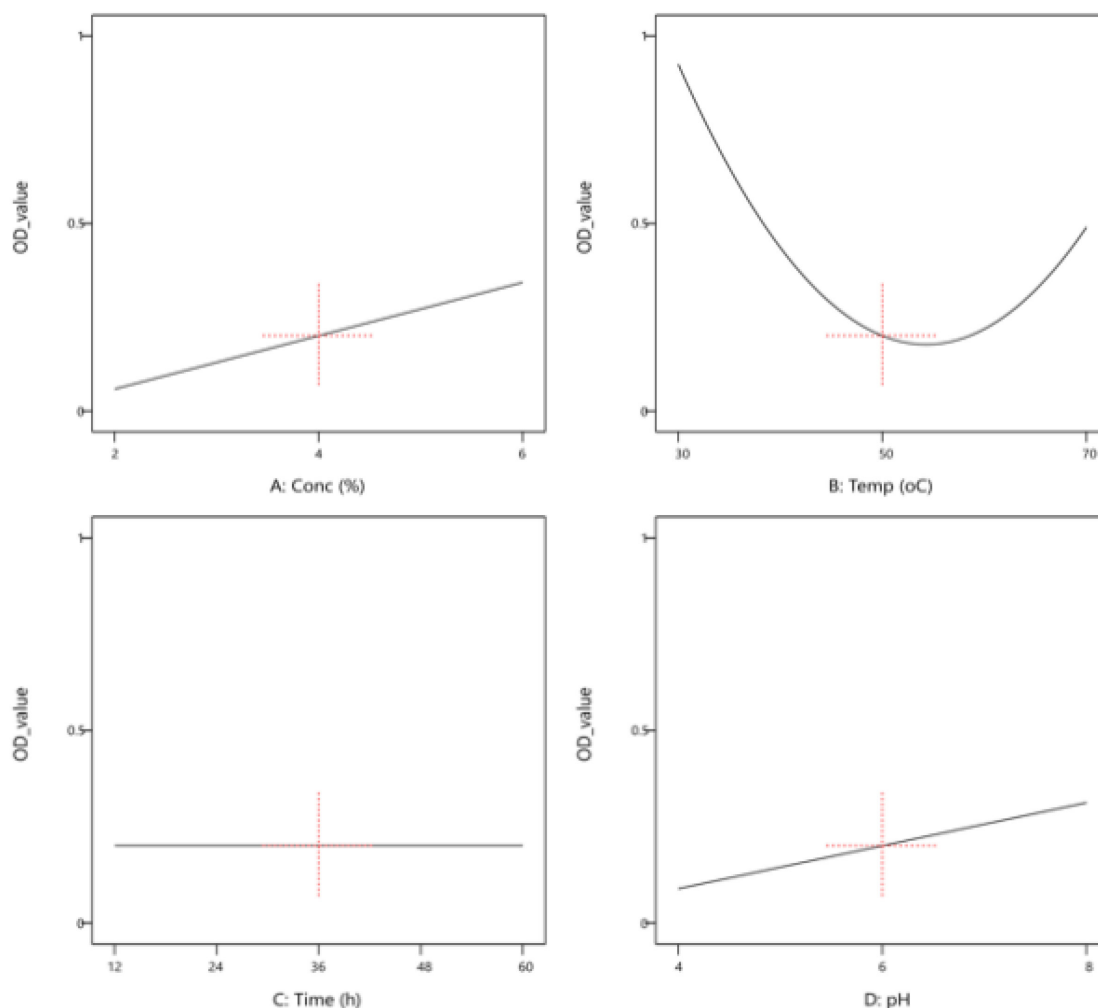


Fig. (3). The impact of individual factors on biofilm formation ability of the *Bacillus cereus* TGS 11.1 strain.

3.3. Interactive Effects of Dual Factors on Biofilm OD Values

Beyond individual parameter analysis, dual-factor interactions were evaluated to elucidate their synergistic influence on the biofilm formation process of *Bacillus cereus* TGS 11.1. ANOVA results (Table 2) confirmed the statistical significance of all three interaction pairs, with the temperature-pH interaction (B×D) exerting the most profound influence ($F = 8.99$; $p = 0.0066$), followed by the temperature-lactose concentration (B×A: $F = 7.74$; $p = 0.0109$) and lactose concentration-pH (A×D: $F = 5.21$; $p = 0.0326$) interactions. The interaction between temperature and pH holds critical biological relevance for microbial persistence in milk-processing environments. At lower temperatures, an alkaline pH facilitates initial bacterial attachment to equipment surfaces by minimizing electrostatic repulsion forces [16]. Conversely, at elevated temperatures ($\text{Temp} \geq 55^\circ\text{C}$), the influence of pH is effectively neutralized, suggesting that thermal stress becomes the primary determinant of heat resistance and survival. Furthermore, the temperature-lactose

concentration interaction underscores the temperature dependency of substrate utilization, where nutrient sources are effectively channeled into biofilm matrix synthesis only in the absence of thermal stress, aligning with established bacterial metabolic energy models under favorable conditions [17].

3.4. Morphological Characterization of the *Bacillus Cereus* TGS 11.1 Strain's Colony Biofilm under Different Temperature Conditions

The RSM-CCD model identified temperature as the primary variable modulating biofilm formation, necessitating morphological characterization of *Bacillus cereus* TGS 11.1 across a thermal gradient of 30–70°C (Fig. 4a). The isolate exhibited viability within the 30–50°C range, while temperatures $\geq 60^\circ\text{C}$ completely inhibited colony formation, thereby defining the upper thermal threshold for this strain. Biofilm radial expansion demonstrated temperature dependency, with diameters measuring 3.0 ± 0.1 mm at 30°C, peaking at 4.0 ± 0.5 mm at 40°C, and declining to 3.0 ± 0.2 mm at 50°C (Fig. 4a).

Morphological transformations across the viable thermal range revealed distinct architectural variations. At 30°C, colonies displayed a smooth, flat architecture with minimal matrix accumulation, whereas 40°C induced a mature biofilm phenotype characterized by a rugose surface and a three-dimensional (3D) network of elevated, opaque aggregates. Conversely, at 50°C, the morphology reverted to a smooth, glassy appearance lacking complex structural features (Fig. 4b). These phenotypic observations identify 40°C as the optimal temperature for architectural maturation and matrix development in *B. cereus* TGS 11.1.

The swarming motility of *B. cereus* TGS 11.1 exhibited

significant variation relative to temperature, reaching a peak diameter of 4.4 ± 0.5 cm at 40°C (Fig. 5). At this optimum, the motility zone displayed vigorous expansion and branching patterns, whereas swarming capacity was significantly impaired at 30°C (3.2 ± 0.3 cm) and 50°C (3.4 ± 0.3 cm). The positive correlation between maximum swarming motility and peak optical density at 40°C suggests that elevated motility facilitates surface coverage, a prerequisite for subsequent biofilm accumulation in milk-processing environments. These results align with established reports regarding optimal temperatures for *Bacillus* motility [18, 19] and surface translocation [20].

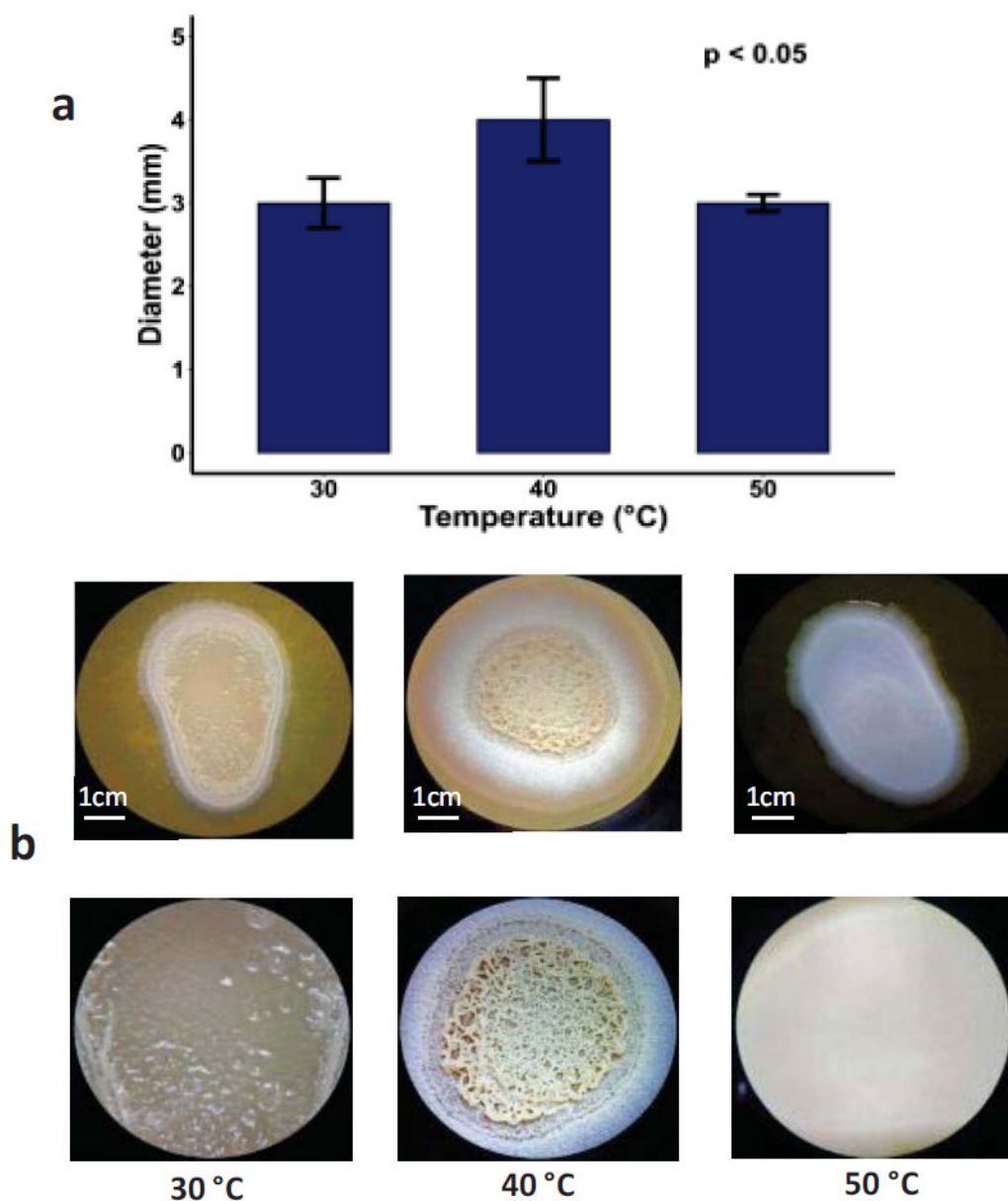


Fig. (4). Diameter (a) and Morphology (b) of *Bacillus cereus* TGS 11.1 colony biofilm grown at different temperatures.

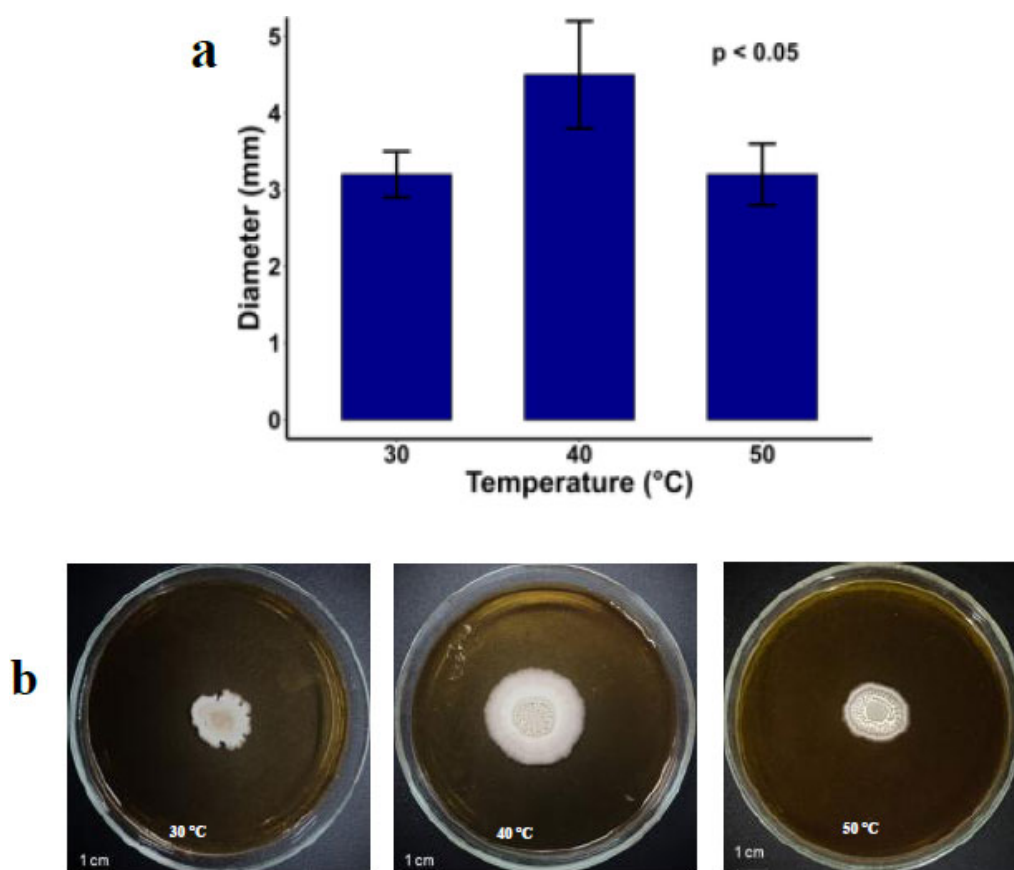


Fig. (5). Motility zone diameter (a) and swarming motility (b) of the *Bacillus cereus* TGS 11.1 strain's colony grown at different temperatures.

CONCLUSION

This study was conducted to evaluate the impact of environmental factors, including temperature (30°C, 40°C, 50°C, 60°C, and 70°C), pH (4, 5, 6, 7, and 8), lactose concentration (2%, 3%, 4%, 5%, and 6%), and incubation time (12, 24, 36, 48, and 60 h), on the biofilm formation of the *Bacillus cereus* TGS 11.1 strain. Among the variables tested, temperature, pH, and lactose concentration significantly influenced biofilm development. Notably, temperature emerged as the most critical factor, with the highest biofilm productivity observed at 40°C. At this temperature, the strain exhibited its most complex biofilm architecture, characterized by a dense three-dimensional (3D) network and peak motility-driven attachment. Based on these findings, it is highly recommended to avoid prolonged exposure to temperatures around 40°C during the processing and storage of milk and dairy products to mitigate the robust biofilm formation of the *Bacillus cereus* TGS 11.1 strain.

AUTHORS' CONTRIBUTIONS

The authors confirm their contributions to this paper as follows: B.T.V.: Responsible for the conception and design of the study; B.H.B.T.: Performed experiments, contributed to methodology, and participated in

manuscript revision; T.H.T.H.: Responsible for data collection, statistical analysis, and participated in manuscript revision; T.V.B.: Contributed to data analysis and assisted in manuscript revision.; T..K.D.H.: Provided support with equipment and infrastructure for the experiments; K.A.T: Supervised the project and provided critical review.

LIST OF ABBREVIATIONS

BHI	=	Brain Heart Infusion
CCD	=	Central Composite Design
CV	=	Crystal Violet
EPS	=	Exopolysaccharides
LB	=	Luria-Bertani
OD	=	Optical Density
ODc	=	Optical Density Control
PBS	=	Phosphate-Buffered Saline
RSM	=	Response Surface Methodology
SD	=	Standard Deviation
3D	=	Three-dimensional

ETHICS APPROVAL AND CONSENT TO PARTICIPATE

Not applicable.

HUMAN AND ANIMAL RIGHTS

Not applicable.

CONSENT FOR PUBLICATION

Not applicable.

AVAILABILITY OF DATA AND MATERIALS

The data and supportive information are available within the article.

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CONFLICT OF INTEREST

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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