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Multi-Response Optimization of Granaticinic Acid Production by Endophytic *Streptomyces thermophilus NT1*, Using Response Surface Methodology

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Abstract: *Streptomyces thermophilus NT1*, an endophytic isolate, was studied for optimization of granaticinic acid production. It is an antimicrobial metabolite active against even drug resistant bacteria. Different media, optimum glucose concentration, initial media pH, incubation temperature, incubation period, and inoculum size were among the selected parameters optimized in the one-variable-at-a-time (OVAT) approach, where glucose concentration, pH, and temperature were found to play a critical role in antibiotic production by this strain. Finally, the Box–Behnken experimental design (BBD) was employed with three key factors (selected after OVAT studies) for response surface methodological (RSM) analysis of this optimization study. RSM analysis revealed a multifactorial combination; glucose 0.38%, pH 7.02, and temperature 36.53 °C as the optimum conditions for maximum antimicrobial yield. Experimental verification of model analysis led to 3.30-fold (61.35 mg/L as compared to 18.64 mg/L produced in un-optimized condition) enhanced granaticinic acid production in ISP2 medium with 5% inoculum and a suitable incubation period of 10 days. So, the conjugated optimization study for maximum antibiotic production from *Streptomyces thermophilus NT1* was found to result in significantly higher yield, which might be exploited in industrial applications.

Keywords: *Streptomyces thermophilus*; granaticinic acid; antimicrobial; optimization; response surface methodology; Box–Behnken design

1. Introduction

Microbial secondary metabolites are extremely species-specific, and sometimes the same species isolated from different ecological statuses produce significant variations in secondary metabolite profile [1]. Microbial cell factories are influenced by physical and nutritional parameters to generate subtle-to-significant variation in biosynthetic pathways that originate precious compounds for mankind [2,3]. Most of such compounds have unique structural features and possess versatile bioactivities. Often, such compounds are accumulated in substantial amounts, which is a key factor in their industrial interest. In biotechnological applications, nevertheless, microbial metabolites used as antimicrobials still continue to represent important research and development of new therapeutic agents [4]. The evaluation and optimization of potential, lesser-known, novel microbial bioactive compounds are of increasing interest to the industries [2]. Among bacteria, the order *Actinomycetales*

has proved its excellence in the production of a diverse range of secondary metabolites, especially with antimicrobial properties with extensive industrial interests [5]. Specifically, the *Streptomyces* are outstanding in the production of large numbers of antibiotics and other bioactive compounds [6].

Streptomyces thermophilus is known to produce granaticins [7], a pigmented antibiotic. Granaticins, produced by many *Streptomyces*, are members of a broader class of pyranonaphthoquinone microbial metabolites [8] and are generally produced as a mixture of different derivatives (granaticin A, granaticin B, dihydrogranaticin, granaticinic acid) in bacterial fermented broth [9], but *Streptomyces thermophilus* NT1 was found to produce solely granaticinic acid in culture filtrate [10]. Granaticins are reported for antibacterial, antitumor and cytotoxicity against KB cells [11], but granaticinic acid was found to be non-mutagenic and to have strong antibacterial activity against both gram-positive and gram-negative bacteria, including the deadly *Pseudomonas aeruginosa*. It is also potentially inhibitory against MRSA (methicillin-resistant *S. aureus*) and PRSA (penicillin-resistant *S. aureus*) strains [10]. This endophytic strain NT1 certainly draws biotechnological importance for the production of future antibiotics which have a potential anti-infective property, even against resistant bacteria, and due to its production of a single granaticin derivative in fermented liquor.

The present communication explains the enhanced yield of granaticinic acid obtained after thoroughly examining different influencing parameters from single and multidisciplinary approach after significant statistical analysis.

2. Experimental Section

2.1. Microbial Strains, Maintenance, and Chemicals

The endophytic *Streptomyces thermophilus* NT1 (Gen Bank Acc. No. KJ486841) was taken for this study. The strain is one of our isolated endophytic actinomycetes strains, which was reported for granaticinic acid production [10]. The strain is maintained in ISP2 broth with 30% glycerol at -20°C . For the assessment of maximum granaticinic acid production, two pathogenic strains were selected; *Bacillus cereus* (ATCC 14579) and *Escherichia coli* (clinical isolate) as test pathogens. The pathogenic strains were maintained in tryptic soy slant culture at 4°C . All the media components were of analytical grade and procured from HiMedia Laboratories Pvt Ltd. (Mumbai, India).

2.2. Inoculum Preparation for Fermentation

Strain NT1 was transferred from stock culture on solid ISP2 medium. It was then incubated at 35°C for 7 days. Agar blocks containing full-grown strain were cut (1 cm^2 surface area) and transferred in Erlenmeyer flask (150 mL) containing 30 mL ISP2 liquid medium. The strain was allowed to further grow for 5 days with shaking (150 rpm) at the same conditions up to biomass generation (dry weight) of 50 mg/100 mL medium. This culture was used as inoculum at different optimization steps.

2.3. Production of Granaticinic Acid

Production of granaticinic acid was carried out in 250 mL Erlenmeyer flasks. The endophytic *S. thermophilus* NT1 was cultured in 50 mL ISP2 medium having pH 7.0 at 28°C inside a rotary shaker incubator at 150 rpm shaking speed for 12 days. Unless otherwise specified, during optimization, culture conditions and media compositions were used as optimized in the previous experimental step.

2.4. OVAT Optimization for Production of Granaticinic Acid

Following parameters were optimized by one-variable-at-a-time (OVAT) to determine granaticinic acid production from this strain at increased levels. Production media, initial media pH, incubation temperature, incubation period, inoculum size, and glucose concentration were selected parameters that were analyzed for maximum granaticinic acid production in optimum conditions.

2.5. Optimization of Media

Few well-known formulated media have been tested to find the most suitable one for yielding maximum granaticinic acid. For this purpose, the liquid media used were as follows; ISP5, ISP2, SCN, TYG and MS broth. Initial pH of all media was adjusted at 7.0. Fermentation was initiated with 2% inoculum in each 50 mL media inside an Erlenmeyer flask (250 mL). Incubation was carried out at 35 °C for 12 days with shaking at 150 rpm. Cell-free culture broth was analyzed for antimicrobial activity.

2.6. Determination of Optimum Glucose Concentration

Optimum glucose concentration for maximum antimicrobial activity was also determined. For this, ISP2 media (devoid of glucose) was taken in separate Erlenmeyer flasks and ameliorated with different glucose concentrations (g%) (0, 0.2, 0.4, 0.6, 0.8, and 1 g%). Each fermentation media was inoculated with 5% (*v/v*) seed culture of NT1. Production was carried out at pH 7.0, 35 °C, for 12 days with shaking at 150 rpm. After completion of fermentation, culture filtrates of each media were evaluated for antimicrobial activity.

2.7. Determination of Initial Media pH

The endophytic strain NT1 was fermented in a series of individual ISP2 media of different pH (4, 6, 7, 8, and 10) at 35 °C. Fermentations were carried out for up to 12 days. Culture filtrates were analyzed.

2.8. Determination of Incubation Temperature

The endophytic strain NT1 was fermented in a series of individual ISP2 media adjusted to pH 7.0 under different incubation temperatures (20, 30, 35, 40 and 45 °C). Fermentations were stopped at the 12th day and culture filtrates were analyzed.

2.9. Determination of Fermentation Period

Liquid fermentation was allowed in ISP2 media with pH 7.0 at 35 °C for fifteen successive days with shaking at 150 rpm. Samples were taken out at each day. Antimicrobial activity as well as dry cellular weight was recorded each day.

2.10. Determination of Most Suitable Inoculum Size

Suitable inoculum size plays a critical role in antibiotic production by *Streptomyces* [12]. Optimum inoculum concentration was determined after transferring different inoculum size (0.5%, 1%, 2%, 5%, and 10%; *v/v*) in ISP2 media, and production was carried at pre-optimized conditions.

2.11. Optimization of Granaticinic Acid Production Using Box–Behnken Design

A more applicable optimization for antibacterial production by this strain was performed with response surface methodology (RSM). The Box–Behnken experimental design (BBD) was employed with three key factors selected from OVAT studies. The experimental design was conducted with the three most influencing independent factors, each at three different levels (−1.0, 0, and +1.0), encompassing a total of 17 experiments (Table 1). Antibacterial production was evaluated, adopting a quadratic polynomial Equation:

$$Y = \beta_0 + \sum \beta_i x_i + \sum \beta_{ii} x_i^2 + \sum \beta_{ij} x_i x_j$$

The equation implies that the input variables x_i and x_j influence the predicted response Y . β_0 is the intercept of the regression equation, β_i is the linear coefficient, β_{ii} is the quadratic coefficient, and β_{ij} is the interaction coefficient. A functional model was generated by the multiple regression analysis

of the responses, and its efficiency was verified by ANOVA and Fisher's F-test. The mutual interactions among independent variables were represented by three-dimensional response surface plots.

Table 1. Experimental design and results of the Box-Behnken design for optimization of antimicrobial production by *Streptomyces thermophilus* NT1.

Run	Independent Variables			Response (Growth Inhibition, mm)	
	A: Glucose (g%)	B: pH	C: Temperature (°C)	Observed	Predicted
1	0.6	7.0	30	5	5.0625
2	0.2	8.0	35	7.3	6.9625
3	0.2	7.0	30	5.3	5.4125
4	0.4	8.0	30	7.2	7.425
5	0.6	6.0	35	5	5.3375
6	0.2	7.0	40	9.3	9.2375
7	0.4	7.0	35	14.2	14.24
8	0.4	8.0	40	9.2	9.6
9	0.4	7.0	35	13.9	14.24
10	0.4	7.0	35	14.6	14.24
11	0.4	6.0	40	10.2	9.975
12	0.2	6.0	35	5.3	5.5875
13	0.6	8.0	35	5	4.7125
14	0.4	7.0	35	14.5	14.24
15	0.4	6.0	30	6.7	6.3
16	0.4	7.0	35	14	14.24
17	0.6	7.0	40	7.2	7.0875

2.12. Validation of RSM Predictions

Strain *S. thermophilus* NT1 was inoculated (5%) in three separate Erlenmeyer flasks (250 mL) containing 50 mL ISP media. Initial media pH, incubation temperature, and glucose concentration was maintained as predicted optimum by the RSM study. Fermentation was carried out for 10 successive days, and antimicrobial activity of culture filtrates against *B. cereus* (ATCC 14579) was determined on MHA media by agar well diffusion assay in three replicates [13]. Inhibition zone produced was compared with RSM predictions.

2.13. Statistical Analysis

All experiments were carried out in triplicate, and data are represented here as mean \pm SEM. The RSM experimental design and concomitant regression analysis of experimental data were executed using Design-expert 8.0 software (STAT-EASE Inc., Minneapolis, MN, USA). The response surface curves, as well as subsequent contour plots from the results of RSM experiment, were also drawn with the same software. The OVAT data analysis was performed using Microsoft Excel 2007 software.

2.14. Estimation of Granaticin at Optimization Phases

The relative amount of granaticin produced was determined after comparing zone of inhibitions produced by different concentrations of 2-(4-Hydroxy-7-methoxy-5,8-dioxo-3,4,5,8-tetrahydro-1H-isochromen-3-yl)-acetic acid ethyl ester (similar class to the antimicrobial compound pyranonaphthoquinone [14]) and granaticin acid at different phases of optimization.

Concentration of granaticin acid in culture filtrates was determined as amount of granaticin acid (volume of culture filtrate) loaded into wells of MHA agar plates.

Volume of well was calculated from the equation:

$$V = \pi r^2 h, \text{ where } r = \text{radius and } h = \text{height of well}$$

$$= (\pi) (0.25)^2 0.5 \text{ [well diameter} = 0.5 \text{ cm and height} = 0.5 \text{ cm]}$$

3. Results

To make the production of the antibiotic feasible, it is necessary to develop the optimum production conditions, as product yield greatly varies between pre- and post-optimized conditions [15]. The strain *Streptomyces thermophilus* NT1 was previously reported for production of granaticin acid, a broad spectrum pyranonaphthoquinone antibiotic.

3.1. OVAT Optimization of Granaticin Acid Production

The endophytic *S. thermophilus* NT1 was fermented in an Erlenmeyer flask for the production of granaticin acid. Its growth with granaticin acid production and other optimization results are presented in Figure 1. An increase in the zone of inhibition always indicates an increased yield. It was observed that granaticin acid production initiates at the 5th day of incubation, after which it exponentially increased and reached its highest level at the 10th day. Biomass generation was also noticed exponentially after a 4 day lag phase. After 8 days of incubation, the organism entered in stationary phase, while antimicrobial production was markedly increased after that (Figure 1b). ISP2 media was found to be the most-preferred production media, and considerable production was also achieved in SCN broth (Figure 1a). Highest antimicrobial production was observed with 5% seed (Figure 1c) as starter of the said fermentation. Study of the relationship between granaticin acid production with initial media pH and incubation temperature revealed that pH 7.0 and 35 °C temperature is the optimum condition for maximum antimicrobial yield (Figure 1d,e respectively). Initial media pH lower than 6 (or slightly alkaline environment) is not at all suitable for granaticin acid production by this strain. There was a sharp relationship between glucose concentration and granaticin acid production. It was noticed that 0.4% glucose is the most effective concentration (Figure 1f) for antimicrobial production by *S. thermophilus* NT1. Almost similar antimicrobial activity was observed against the gram-positive *B. cereus* and the gram-negative *E. coli* during production optimization through OVAT fashion (Figure 1).

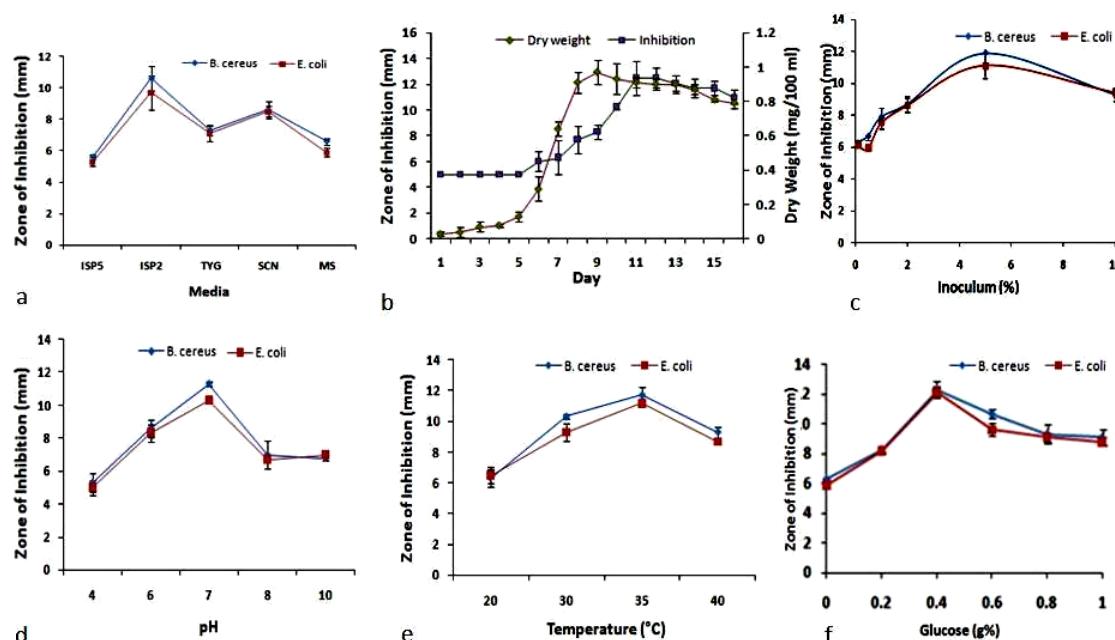


Figure 1. Effect of different influencing parameters in granaticin acid production (antimicrobial activity) by one-variable-at-a-time (OVAT) study; (a) influence of media on activity; (b) growth curve vs. antimicrobial production; (c) influence of inoculum concentration; (d) influence of initial media pH; (e) influence of incubation temperature; (f) influence of glucose concentration on antimicrobial production.

3.2. Box–Behnken Optimization for Granaticin Acid Production

The conventional OVAT approach faces some restrictions, especially when multiple significantly influencing parameters work together. So, a conjugated OVAT–RSM approach is always preferable [16]. In this optimization experiment, a three level Box–Behnken design of three factors (initial pH of the fermentation medium, incubation temperature during fermentation, and glucose concentration) with five replicates at the center point of each factor was implicated (Table 1) as the model for the analysis of granaticin acid production. Experimental design, together with predicted and measured responses of antibacterial production, is presented in Table 1. Significant variations in antibacterial production by both strains were observed at different production conditions, whereas the center point condition was found to always yield maximum antibacterial production compared to others. Multiple regression analysis of predicted response Y for antibacterial production by individual strain was described as:

$$Y_{\text{Granaticinate}} = 14.24 - 0.625A + 0.1875B + 1.4625C - 0.5AB - 0.45AC - 0.375BC - 5.1075A^2 - 3.4825B^2 - 2.4325C^2$$

where $Y_{\text{Granaticinate}}$ implies yield of granaticin acid by strain NT1, and A, B, C are coded factors of glucose concentration (g%), initial media pH, and temperature, respectively. The goodness of fit of the conducted RSM was evaluated with the experimental outputs by performing the regression study (Table 2). According to the constructed model, three-dimensional response surface plots (Figure 2) were generated by Design Expert 8.0 software to better understand the effects of the variables and their interactions on the yield of granaticin acid. The 3D plots (Figure 2) showing interactions between the selected variables aided rapid evaluation of the critical values of each variable for maximum production. It was observed that all of the quadratic surfaces, rather than linear planes, were significant in the 3D plots.

Table 2. ANOVA for response surface quadratic regression model of antimicrobial production by *Streptomyces thermophilic* NT1.

Source	Sum of Squares	df	Mean Square	F-Value	p-Value	Prob>F
Model	228.0028529	9	25.33365033	145.416607	<0.0001	
A	3.125	1	3.125	17.93767938	0.0039	
B	0.28125	1	0.28125	1.614391144	0.2445	
C	17.11125	1	17.11125	98.2195572	<0.0001	
AB	1	1	1	5.740057401	0.0478	
AC	0.81	1	0.81	4.649446494	0.0680	
BC	0.5625	1	0.5625	3.228782288	0.1154	
A²	109.8381316	1	109.8381316	630.47718	<0.0001	
B²	51.06444737	1	51.06444737	293.112859	<0.0001	
C²	24.91392105	1	24.91392105	143.0073369	<0.0001	
Residual	1.2195	7	0.174214286			
Lack of fit	0.8475	3	0.2825	3.037634409	0.1556	
Pure error	0.372	4	0.093			
Cor total	229.2223529	16				
R-Squared	0.994679838					
Adj R-Squared	0.98783963					
Pred R-Squared	0.938307718					
Adeq Precision	29.76196049					

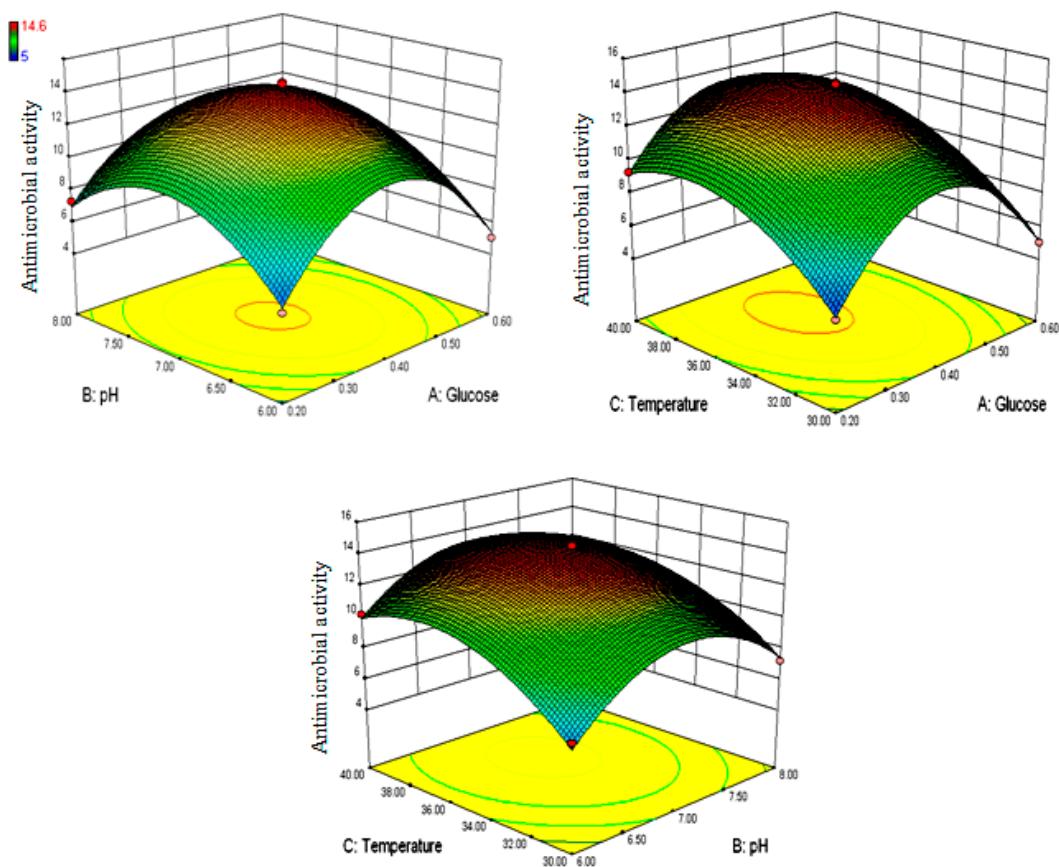


Figure 2. The 3D plot and 2D projections showing most important interactions for maximum antimicrobial production by TNT1, a; between glucose concentration (g%) and initial media pH at temperature 35 °C; b: glucose concentration (g%) and temperature (°C) at initial media pH 7.0; c: between temperature (°C) and initial media pH 7 at glucose concentration 0.4 (g%)

3.3. Validation Study

Validation of these predictions related to granaticinic acid production by this endophytic *Streptomyces* was assessed by laboratory trials in Erlenmeyer flask culture. Antimicrobial activity of culture filtrates grown in RSM optimized conditions were compared (against *B. cereus* (ATCC 14579)) with the predicted stipulation of the RSM study. Increased antimicrobial activity was observed at each stage of optimization (Figure 3).

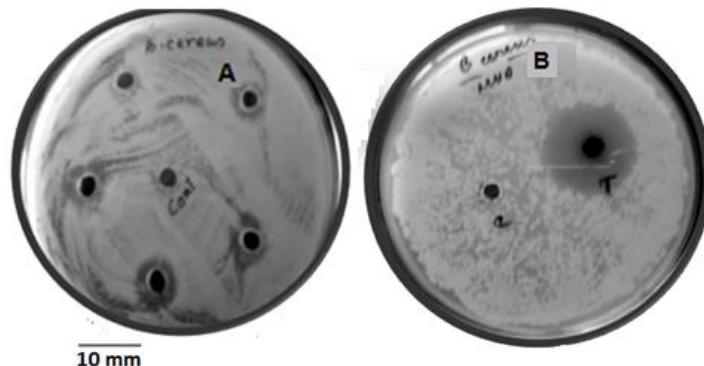


Figure 3. Growth inhibition of *B. cereus* by granaticinic acid at different stages of optimization; (A) un-optimized; (B) RSM optimized.

3.4. Yield of Granaticinic Acid

Culture filtrates of NT1 produced at different phases of optimization of granaticinic acid production were compared with standard data provided by a similar compound synthesized by Lagorio, et al., (2006). From the standard curve (Figure 4), the most probable presence of granaticinic acid in culture filtrates are presented in Table 3. Antibiotic production increased by up to 3.30-fold, which is equivalent to almost 61.35 mg/L, when the three most significant influencing parameters worked cooperatively at their critical value.

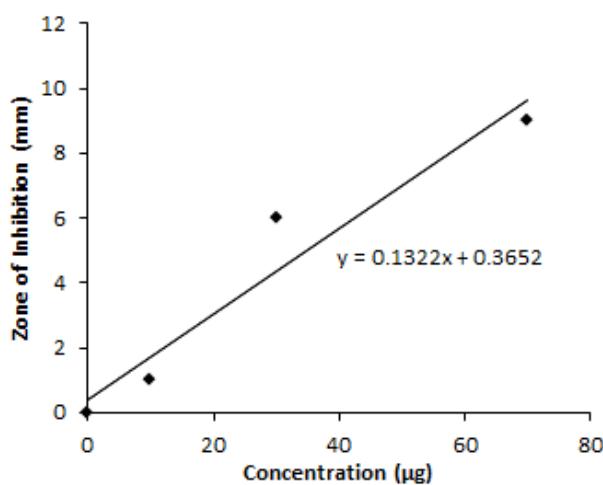


Figure 4. Standard curve of pyranonaphthoquinone antimicrobial to determine granaticinic acid yield.

Table 3. Comparative yield of granaticinic acid at different optimization steps.

Sample Tested	Zone of Inhibition (mm)	Amount of Granaticinic Acid (mg/L)	Fold Increased
Crude	8.83	18.64	0
OVAT*	11.73	40.58	2.17
RSM	14.5	61.53	3.30

* Average zone of inhibition was considered in case of OVAT optimization.

4. Discussion

In this study, optimization of culture conditions for the production of granaticinic acid from endophytic strain *S. thermophilus* NT1 has been studied through OVAT, factorial design, and RSM analysis. The production of secondary metabolites with antimicrobial property is a common phenomenon among the genus *Streptomyces*. This group of bacteria has been isolated so much from traditional ecotopes like water or soil that isolation of *Streptomyces* with unusual secondary metabolites profile has become rare [17]—the quest for endophytic microorganisms indicates some new leads. The endophytic *S. thermophilus* NT1 produces granaticinic acid, which is considered to be significantly pharmaceutically important due to its anti-*Pseudomonas* properties and effectiveness against resistant pathogens. It is well known that the production of antibiotics varies substantially from strains, species, and certainly with the cultural conditions. Growth temperature, media pH, media compositions, limiting substrate concentration, and obviously incubation time influence the microbial growth, which might have effects on onset and intensity of secondary metabolism [18]. In this experiment, the approach successfully allowed the determination of critical combinatorial values of the most influencing parameters that give the highest antibiotic yield. Any change in the yield of antimicrobial production during OVAT or RSM optimization was determined by measuring zone of

inhibition produced by culture filtrates. ISP2 media, which was found to be the most suitable antibiotic production media [19,20], contains malt extract, which consists of various malt hydrolyzed products as carbon source and yeast extract provides an organic nitrogen source and vitamin precursors to facilitate microbial growth. Aside from that it contains glucose as the principal carbon source for growth. The other media employed for granaticin acid production in this experiment did not support its production by the strain; it might be that the nutritional components were not favorable. Less production was observed in SCN broth, indicating that organic carbon and nitrogen played critical role in the production of active compound. In OVAT optimization, the critical concentration of glucose was found to be 0.4 g% for maximum antimicrobial yield. A high glucose concentration always supports higher biomass generation, but secondary metabolites are produced under stressed conditions. In this experiment, it was found that higher glucose concentrations were not satisfactory for this purpose, as it might avoid nutritional stress. It is also assumed that higher levels of glucose may also inhibit some essential enzyme in the biosynthetic pathway of granaticin acid. Earlier study shows the use of glutamate as sole carbon source for granaticin production from *Streptomyces thermophilus* sub sp. *Thermophilus* NCIB 10076 [21]. This substrate could be used both as nitrogen and carbon sources for cellular demand, and high yield of biomass generation was found with it. Use of glucose in this study avoids the generation of ammoniacal nitrogen that may reduce antibiotic yield, as in the case of *S. fradiae* [22]. Besides, glucose easily generates a large amount of acetate, which is the principle building block of granaticin acid via a polyketide-derived secondary metabolic pathway. The NT1strain showed a maximum granaticin acid production at 35 °C in the OVAT study. It explained the well-characterized “tropophase” up to 7 days, and then the “idiophase” was initiated when antibiotic concentration sharply reached its highest point. This nature is highly expected in commercial batch production of valuable products, as complete utilization of substrate is achieved [23]. More or less granaticin production is reported at pH 7.0 from all actinomycetes strains [21], which is collinear to our study.

Nowadays, before going to microbial fermentation, RSM-based optimization is widely accepted due to its higher efficiency, reasonable design for integrated analysis of variants, and mathematical modeling with recommended values of most possible optimum conditions, along with corresponding product yield. The approach is techno-economically more precise and viable than univariate strategies, and more acceptable industrially [24]. In most cases, OVAT followed by RSM is strongly recommended to find out the critical values of those significant factors [16]. The *F* test data was checked, and the model *F*-value of 145.416607 indicated that the models were significant. It was found that there was only a 0.01% chance that an *F*-value this large could occur due to noise. The adjusted determinant coefficient (R^2_{ADJ}) of the polynomial model was measured to test the goodness-of-fit of the regression equation. The value of R^2_{ADJ} was calculated as 0.98783963. It shows that there existed a high degree of correlation between the measured and predicted values, and more than 96% variations in the response could be explained by that second order polynomial equation. Adequate precision measured the signal-to-noise ratio with a desired value of 29.76—i.e., greater than 4—which in turn indicated an adequate signal, and the model could be used to navigate the design space. The lack of fit *F*-value of 0.84 and 0.75 for both the models implies that the lack of fit was not significant relative to the pure error, and hence the fitness of the model was good. The resulted model *p*-value ($p < 0.0001$) shows that the model equation was suitable to explain the response of experiment pertaining to antimicrobial production by the strain NT1. The *p*-value for lack of fit for the equations (0.1556) is significantly higher than 0.05, indicating the high degree of accuracy and consistency of the experimental values. The *p*-value of probability $>F$ less than 0.05 also indicated that the model terms were significant. In these models, the linear and the quadratic effects of glucose concentration (g%), initial media pH, and temperature (°C) were significant ($p < 0.05$). The most effective interaction was obtained between initial media pH and glucose concentration (g%, $p < 0.05$).

The 3D model predicted a maximum response of 14.49 mm (zone of inhibition against *B. cereus* ATCC 14579) when the uncoded levels of the three most significant variables for strain NT1 were

initial media glucose concentration 0.38 g%, pH 7.02, temperature 36.53 °C within the experimental range. Validation of these predictions related to antimicrobial production was assessed by laboratory trials in Erlenmeyer flask culture with the predicted stipulation and 14.4 ± 0.031 mm (inhibition zones were observed after 10 days of fermentation at optimized conditions against strain ATCC 14579). This finding certainly established a significant correlation between expected and investigational antimicrobial production. Experimental verification established the fitness and validation of the model, and the best possible conditions were finally selected through the least number of experiments compared to any other design.

The relative yield of granaticinic acid (61.53 mg/L) revealed the importance of such statistical multi-response optimization in the commercial production of antibiotics. The high degree of similarity between prediction and experimental output certainly encourages such process optimization in antibiotic production. About 3.30-fold enhanced production in any pharmaceutical industry is no doubt considerable. So, this strain could certainly be expedient to its pharmaceutical development.

5. Conclusions

Production of the antimicrobial compound is found to be critically dependent on few culture conditions as well as nutritional status. In this study, any endophytic *S. thermophilus* NT1 is first employed for optimization of antimicrobial production by OVAT followed by RSM using Box–Behnken design. The critical value of three crucial growth parameters (glucose concentration, media pH, and temperature) were determined when interacting associatively in fermentation for the highest yield of granaticinic acid. Such multi-parameter integrated optimization is practically important because during fermentation, microorganisms have to face all influential parameters at same time. Validation experiments for the proposed model were also carried out to verify the adequacy and accuracy of the generated model, and results showed that the predicted value agreed with the experimental values well. The best combination resulted after RSM analysis provided a three-fold increased production of granaticinic acid in 10 days with 5% starter in Erlenmeyer flask culture. The optimum culture conditions achieved in this experiment set the basis for further study with large scale batch fermentation in a fermenter for granaticinic acid production from *S. thermophilus* NT1.

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Conflicts of Interest: The authors declare no conflict of interest.

References

1. Bibb, M.J. Regulation of secondary metabolism in Streptomyces. *Curr. Opin. Microbiol. Microbiol.* **2005**, *8*, 208–215. [[CrossRef](#)] [[PubMed](#)]
2. Wang, X.; Huang, L.; Kang, Z.; Buchenauer, H.; Gao, X. Optimization of the fermentation process of actinomycetes strain Hhs015T. *J. Biomed. Biotechnol.* **2010**, *141876*. [[CrossRef](#)]
3. Jose, P.A.; Santhi, V.S.; Jebakumar, S.R.D. Phylogenetic affiliation, antimicrobial potential and PKS gene sequence analysis of moderately halophilic *Streptomyces* sp. inhabiting an Indian saltpan. *J. Basic Microbiol.* **2011**, *51*, 348–356.
4. Bérdy, J. Bioactive microbial metabolites. *J. Antibiot.* **2005**, *58*, 1–26. [[CrossRef](#)] [[PubMed](#)]
5. Nett, M.; Ikeda, H.; Moore, B.S. Genomic basis for natural product biosynthetic diversity in the actinomycetes. *Nat. Prod. Rep.* **2009**, *26*, 1362–1384. [[CrossRef](#)] [[PubMed](#)]
6. Roy, S.; Banerjee, D. Bioactive endophytic actinomycetes of *Cinnamomum* sp.; isolation, identification, activity guided purification and process optimization of active metabolite. *Am. J. Microbiol.* **2015**, *6*, 4–13. [[CrossRef](#)]

7. St. Pyrek, J.S.; Achmatowicz, O.; Zamojski, A. Naphtho- and anthraquinones of *Streptomyces thermophilicaceus* WR14, structure and model synthesis. *Tetrahedron* **1977**, *33*, 673–680. [[CrossRef](#)]
8. Arnone, A.; Camarda, L.; Cardillo, R.; Fronza, G.; Merlini, L.; Mondelli, R.; Nasini, G.; St. Pyrek, J.S. ^{13}C -NMR analysis of dihydrogranaticin methyl ester. A case of mixed biogenesis. *Helv. Chim. Acta* **1979**, *62*, 30–33. [[CrossRef](#)]
9. Fleck, W.F.; Strauss, D.G.; Prauser, H. Naphthoquinone antibiotics from *Streptomyces lateritius*, fermentation, isolation and characterization of granatomycins A, C and D. *Z. Allg. Mikrobiol.* **1980**, *20*, 543–551. [[CrossRef](#)] [[PubMed](#)]
10. Roy, S.; Banerjee, D. Broad spectrum antibacterial activity of granaticinic acid, isolated from *Streptomyces thermophilicaceus* NT1; an endophyte in *Catharanthus roseus* (L.) G. Don. *J. Appl. Pharm. Sci.* **2015**, *5*, 6–11. [[CrossRef](#)]
11. Bartholomäus, R.; Bachmann, J.; Mang, C.; Haustedt, L.O.; Harms, K.; Koer, U. Synthesis of the AB-Ring pyranolactone substructure of granaticin A. *Eur. J. Org. Chem.* **2013**, *180*, 190. [[CrossRef](#)]
12. El-Enshasy, H.A.; Farid, M.A.; El-Sayed, A. Influence of inoculum type and cultivation conditions on natamycin production by *Streptomyces natalensis*. *J. Basic Microbiol.* **2000**, *40*, 333–342. [[CrossRef](#)]
13. Bauer, A.W.; Kirby, W.M.; Sheriss, J.C.; Turc, M. Antibiotic susceptibility testing by standardized single disc method. *Am. J. Clin. Pathol.* **1996**, *45*, 493–496.
14. Lagorioa, S.H.; Bianchia, D.A.; Sutichb, E.G.; Kaufman, T.S. Synthesis and antimicrobial activity of pyranobenzoquinones related to the pyranonaphthoquinone antibiotics. *Eur. J. Med. Chem.* **2006**, *41*, 1333–1338. [[CrossRef](#)] [[PubMed](#)]
15. Banerjee, D.; Jana, M.; Mahapatra, S. Production of exopolysaccharide by endophytic *Stemphylium* sp. *Micol. Aplicada. Int.* **2009**, *21*, 57–62.
16. Feng, Y.L.; Li, W.Q.; Wu, X.Q.; Cheng, J.W.; Ma, S.Y. Statistical optimization of media for mycelia growth and exo-polysaccharide production by *Lentinus edodes* and a kinetic model study of two growth morphologies. *Biochem. Eng. J.* **2010**, *49*, 104–112. [[CrossRef](#)]
17. Stierle, A.; Strobel, G.; Stierle, D. Taxol and taxane production by *Taxomyces andreanae*, an endophytic fungus of Pacific Yew. *Science* **1993**, *260*, 214–216.
18. Zhang, Y.; Huang, H.; Xu, S.; Wang, B.; Ju, J.; Tan, H.; Li, W. Activation and enhancement of Fredericamycin A production in deep sea-derived *Streptomyces somaliensis* SCSIO ZH66 by using ribosome engineering and response surface methodology. *Microb. Cell Fact.* **2015**, *14*, 64. [[CrossRef](#)] [[PubMed](#)]
19. Rateb, M.E.; Yu, Z.; Yan, Y.; Yang, D.; Huang, T.; Vodanovic-Jankovic, S.; Kron, M.A.; Shen, B. Medium optimization of *Streptomyces* sp. 17944 for tirandamycin B production and isolation and structural elucidation of tirandamycins H, I and J. *J. Antibiot.* **2014**, *67*, 127–132. [[CrossRef](#)] [[PubMed](#)]
20. Siddique, S.; Syed, Q.; Adnan, A.; Qureshi, F.A. Isolation, characterization and selection of avermectin-producing *Streptomyces avermitilis* strains from soil samples. *Jundishapur J. Microbiol.* **2014**, *7*, e10366. [[CrossRef](#)] [[PubMed](#)]
21. James, P.D.A.; Edwards, C. The effects of temperature on growth and production of the antibiotic granaticin by a thermotolerant Streptomycete. *J. Gen. Microbiol.* **1989**, *135*, 1997–2003. [[CrossRef](#)] [[PubMed](#)]
22. Vu-Trong, K.; Gray, P.P. Influence of ammonia on the biosynthesis of the macrolide antibiotic tylosin. *Enzym. Microb. Technol.* **1987**, *9*, 590–593. [[CrossRef](#)]
23. Bommarius, B.; Jenssen, H.; Elliott, M.; Kindrachuk, J.; Pasupuleti, M.; Gieren, H.; Jaeger, K.E.; Hancock, R.E.W.; Kalman, D. Cost-effective expression and purification of antimicrobial and host defense peptides in *Escherichia coli*. *Peptides* **2010**, *31*, 1957–1965. [[CrossRef](#)] [[PubMed](#)]
24. Mahapatra, S.; Banerjee, D. Optimization of a bioactive exopolysaccharide production from endophytic *Fusarium solani* SD5. *Carbohydr. Polym.* **2013**, *97*, 627–634. [[CrossRef](#)] [[PubMed](#)]

