

MATHEMATICAL MODELING OF BIOPROCESS VARIABLES FOR IMPROVED PRODUCTION OF RHAMNOLIPID FROM *Pseudomonas aeruginosa* STRAIN JQ

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Recent developments and experimental methodologies have sparked an interest in exploring multifaceted nature of surfactants of biological origin. Rapidly accumulating scientific data is emphasizing on the extensive nature of the biosurfactants produced by various microbial strains having a wider spectrum of biological activities. Confluence of information also provided evidence about the role of rhamnolipid in the development of sustainable agriculture and soil remediation. In present research an environmental isolate of *Pseudomonas aeruginosa* JQ was evaluated for rhamnolipid production under the effect of some nutritional parameters. Four nutritional factors were investigated viz NaNO₃, KH₂PO₄, MgSO₄ and FeSO₄ by the application of response surface methodology. On the basis of regression analysis, 3.92 g/L NaNO₃, 2.38 g/L KH₂PO₄, 0.26 g/L MgSO₄ and 0.0028 g/L of FeSO₄ were found to be optimum nutritional conditions where 5.67 g/L of the rhamnolipid was produced. In conclusion, *P. aeruginosa* JQ showed excellent metabolic potential to produce biosurfactants that could be effectively used in agriculture, environmental and medical applications.

Keywords: Biosurfactants, Rhamnolipids, response surface methodology, *P. aeruginosa*

INTRODUCTION

Surfactants are one of the most important products of the chemical industry for being extensively used in a wide variety of agrochemical formulations including growth promoters, insecticides, herbicides and fungicides (Deleu and Paquot, 2004). Beside, their wide ranging implications, synthetic surfactants pose serious environmental and health challenges (Burns *et al.*, 2013). In compliance with current environmental legislation, use of sustainable chemicals in agriculture and agrochemical industries has been considerably evoked. Biosurfactants are among those natural products that have gained substantial scientific attraction due of their potential role in agriculture, pharmaceutical, cosmetics and environmental remediation (Das *et al.*, 2010). Chemically, biosurfactants are characterized as glycolipids, lipopeptides, lipoproteins, lipopolysaccharides-protein and polysaccharides-protein-fatty acid complexes (Cameotra *et al.*, 2010). Considering the performance of biosurfactants as an alternative to synthetic homologues, they have several advantages, including their high substrate specificity, less toxicity, biodegradability, durability at extreme operational conditions and ease of production using renewable resources (Banat *et al.*, 2010; Silva *et al.*, 2014). Biosurfactants can be produced by wide variety of microorganisms found in different ecological habitats. In soil, production of these biomolecules have been known to facilitate plant-microbe

interactions, bioavailability of nutrients for beneficial microbes, elimination of pathogens, and improving soil quality (Sachdev and Cameotra, 2013; Yan *et al.*, 2014). The glycolipids containing L-rhamnose and β -hydroxyalkanoic acid moieties are termed as rhamnolipids (RLs). They are characterized as the most effective class of biosurfactants mainly produced by different strains of *Pseudomonas aeruginosa* under limited growth conditions (Chen *et al.*, 2007). The physiochemical properties associated with rhamnolipid enable these to be exploited in wide variety of applications. In agriculture, they are very effective in improving the quality of agriculture soil by removing organic and inorganic pollutants (Pacwa-Plociniczak *et al.*, 2011). Similarly, their excellent antimicrobial activity against variety of plant pathogens engraves a place of themselves as an effective biocontrol agent. Moreover, accumulating information has substantiated its role and efficacy as compared to commercial pesticides for controlling resistant plant pathogens (Sha *et al.*, 2011; Krzyzanowska *et al.*, 2012). Rhamnolipids are also involved in improving plant immunity and growth thus naturally reducing plant infections (Vatsa *et al.*, 2010). Their production in the rhizosphere facilitates establishment of biofilm thereby, improving plant microbe interactions and enhance bioavailability of hydrophobic compounds (Ron and Rosenberg, 2011; Gudina *et al.*, 2015).

The synthesis of rhamnolipids in *P. aeruginosa* is an intricate metabolic process involving a number of physiochemical factors such as carbon substrates, nitrogen source, multivalent ions, pH, temperature, agitation and feeding rate (Nitschke *et al.*, 2005; Kumar *et al.*, 2012). Therefore, optimization of rate limiting factors typically media components have been considered crucial for improved rhamnolipid yield (Gilmour *et al.*, 2006). Statistical design strategies, particularly related to response surface methodology (RSM) are becoming more useful tool in improving the targeted efficiencies of rhamnolipid bioprocess. RSM is an effective statistical technique used for designing experiments, building models, evaluating process variables, and probing minor reaction conditions pertaining to maximum yield of cellular products (Kammoun *et al.*, 2008).

Recently, it has been recognized that self assembly of rhamnolipid monomers generate variety of potent microstructures that could possibly expand their role in various novel applications (Kiran *et al.*, 2011). This has catalyzed a renewed interest in rhamnolipid production and applications. However, current wide spread applications of rhamnolipid are impeded owing to their low cellular productivity and cost of fermentation media. Realizing the future potential of these biomolecules, efforts are underway to improve rhamnolipid production rate by focusing on different nutritional parameters. Therefore, present research is riveted to stimulate the production of rhamnolipids from an indigenous isolate of *P. aeruginosa* by optimization of four critical nutritional factors viz., NaNO_3 , KH_2PO_4 , MgSO_4 and FeSO_4 using response surface methodology.

MATERIALS AND METHODS

Microorganism: *P. aeruginosa* strain JQ was originally isolated from crude oil contaminated soil and characterized for the production of rhamnolipid mixture. The strain was collected from Biochemistry Lab, G.C University, Lahore. The strain was sub-cultured on nutrient agar plates used in subsequent experiments.

Media and inoculum preparation: The basal media with the following compositing (g/L) was used for the inoculum preparation: NaNO_3 5 g, K_2HPO_4 3 g, NaH_2PO_4 2 g, $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$ 0.4 g, FeSO_4 0.004 g and 3% of glycerol was added as sole source of carbon. The concentration of these salts was changed according to the experimental design. This medium was fortified by adding 1 ml of micronutrients' solution containing; $\text{ZnSO}_4 \times 7\text{H}_2\text{O}$, $\text{CuSO}_4 \times 5\text{H}_2\text{O}$, $\text{CoCl}_2 \times 6\text{H}_2\text{O}$, $\text{NaMoO}_4 \times \text{H}_2\text{O}$, $\text{MnSO}_4 \times \text{H}_2\text{O}$, and KCl. The pH of the medium was adjusted at 7 and after sterilization 2.8% inoculum was added into 250 ml flasks which were kept under continuous shaking at 160 rpm and 34°C.

Rhamnolipids estimation: The estimation of rhamnolipid was carried out by the method of Chandrasekaran and Be-

Miller (1980). Briefly, 333 μL of the culture supernatant was taken and extracted using diethyl ether. The organic phase was extracted in triplicate followed by the evaporation of the solvent. The resultant material was mixed with 0.5 μL deionized water. One hundred (100) μL of this mixture was then treated with freshly prepared Orcinol reagent and allowed to react at 80°C for 30 min. After cooling the reaction mixture at room temperature, the absorbance was measured at 421 nm using Spectrophotometer. The results were then analyzed by comparison with a standard curve of the L-rhamnose.

Design of experiment and statistical analysis: Central Composite Design (CCD) strategy was employed in order to formulate an empirical model to determine optimum values of four nutritional factors viz, NaNO_3 , KH_2PO_4 , MgSO_4 and FeSO_4 influencing rhamnolipid production. A full length factorial design was generated with 30 experimental runs using *MiniTab.15* software. Concentration of rhamnolipid (g/L) was selected as response (Y_1) to probe the effect of aforementioned parameters. The process variables and their respective levels are presented in Table. 1 and layout of the central composite design including values of response (Y_1) are expressed in the Table 2. The results obtained after experimentation were subjected for analysis of variance (ANOVA) to validate the accuracy of observations and possible role of each parameter. In addition, multiple regression analysis was applied to find out linear and quadratic effect of these process variables. The interaction of variables in the given system was explained by the following quadratic polynomial equation:

$$\hat{Y} = b_0 + b_1X_1 + b_2X_2 + b_3X_3 + b_{11}X_{21} + b_{22}X_{22} + b_{33}X_{23} + b_{12}X_1X_2 + b_{13}X_1X_3 + b_{23}X_2X_3$$

Where \hat{Y} refers to the predicted response and X_1 , X_2 , X_3 and X_4 are the coded variables. The term b_0 corresponds to the intercept, whereas, b_1 , b_2 , b_3 and b_4 represents linear effects, b_{11} , b_{22} , b_{33} , and b_{44} denote squared effects, and b_{12} , b_{23} , b_{13} interaction terms.

The 3-D response surface plots were made using trial version of *MiniTab 15* to describe individual and cumulative effects of the process variables.

Table 1. Factors with their respective levels.

| Variables | Codes | Levels of the Variables (g/L) | | | | |
|--------------------------|-------|-------------------------------|-------|-------|-------|-------|
| | | -2 | 1- | 0 | +1 | +2 |
| NaNO_3 | A | 0 | 1 | 3 | 5 | 7 |
| KH_2PO_4 | B | 0 | 1 | 2 | 3 | 4 |
| MgSO_4 | C | 0 | 0.1 | 0.3 | 0.5 | 0.7 |
| FeSO_4 | D | 0 | 0.001 | 0.003 | 0.005 | 0.007 |

RESULTS AND DISCUSSION

Process economics has always been the major concern associated with most of the biotechnological products. The

effect of four media components on the production of rhamnolipid was studied using a full length central composite design consisting of 30 experiments. Second order polynomial equation was generated to describe possible role of independent process variables on rhamnolipid production.

Table 2. Experimental Design and Results of Central Composite Design (CCD).

| Runs | NaNO ₃ X ₁ | KH ₂ PO ₄ X ₂ | MgSO ₄ X ₃ | FeSO ₄ X ₄ | Rhamnolipid Y ₁ |
|------|-------------------------------------|---|-------------------------------------|-------------------------------------|-------------------------------|
| 1 | 5 | 1 | 0.1 | 0.005 | 2.31 |
| 2 | 1 | 3 | 0.1 | 0.005 | 1.99 |
| 3 | 5 | 3 | 0.5 | 0.001 | 2.87 |
| 4 | 5 | 3 | 0.1 | 0.005 | 3.24 |
| 5 | 1 | 1 | 0.5 | 0.005 | 1.05 |
| 6 | 5 | 3 | 0.1 | 0.001 | 3.67 |
| 7 | 1 | 3 | 0.1 | 0.001 | 2.16 |
| 8 | 3 | 2 | 0.3 | 0.003 | 4.52 |
| 9 | 1 | 1 | 0.1 | 0.001 | 0.88 |
| 10 | 3 | 0 | 0.3 | 0.003 | 0.23 |
| 11 | 3 | 2 | 0.3 | 0.003 | 4.61 |
| 12 | 1 | 1 | 0.5 | 0.001 | 1.12 |
| 13 | 1 | 3 | 0.5 | 0.001 | 1.64 |
| 14 | 3 | 2 | 0.3 | 0.007 | 1.15 |
| 15 | 3 | 4 | 0.3 | 0.003 | 2.83 |
| 16 | 1 | 3 | 0.5 | 0.005 | 1.11 |
| 17 | 1 | 1 | 0.1 | 0.005 | 0.63 |
| 18 | 5 | 1 | 0.1 | 0.001 | 2.48 |
| 19 | 3 | 2 | 0.3 | 0.001 | 1.47 |
| 20 | 5 | 3 | 0.5 | 0.005 | 1.7 |
| 21 | 5 | 1 | 0.5 | 0.005 | 0.98 |
| 22 | 3 | 2 | 0.7 | 0.003 | 1.4 |
| 23 | 3 | 2 | 0.3 | 0.003 | 4.49 |
| 24 | 3 | 2 | 0.3 | 0.003 | 4.4 |
| 25 | -1 | 2 | 0.3 | 0.003 | 0 |
| 26 | 3 | 2 | 0.3 | 0.003 | 3.84 |
| 27 | 5 | 1 | 0.5 | 0.001 | 1.99 |
| 28 | 3 | 2 | 0.3 | 0.003 | 4.63 |
| 29 | 3 | 2 | -0.1 | 0.003 | 2.15 |
| 30 | 7 | 2 | 0.3 | 0.003 | 2.37 |

Tab. 2: Experimental design and corresponding results of

rhamnolipid production in (g/L) under the influence of X₁, X₂, X₃, and X₄. The values of response (Y₁) were obtained after running experiments in triplicate.

Analysis of variance (ANOVA) test was used to find out the fitness of the experimental design. Results indicated an F-value of 16.90 predicting that our model was significant. Moreover, *P*-value <0.0001 of the model further validated the significance of the experimental procedure. The linear and cumulative interactions between the variables were ascertained by relevant *p*-values of the process variables (Table 3). It was noted that X₁ (NaNO₃), X₂ (KH₂PO₄), X₃ (MgSO₄), X₄ (FeSO₄), X₁ × X₃ (NaNO₃ × MgSO₄) were identified as significant model terms (Table 3). The regression equation for the test model was:

$$Y_1 = +4.42 + 0.56 X_1 + 0.51 X_2 - 0.27 X_3 - 0.18 X_4 - 0.21 X_1 X_3 - 0.16 X_2 X_3 - 0.74 X_1^2 - 0.65 X_2^2 - 0.59 X_3^2 - 0.71 X_4^2$$

Where Y₁ is the estimated response (rhamnolipid) and X₁, X₂, X₃, and X₄ were coded factors for NaNO₃, KH₂PO₄, MgSO₄ and FeSO₄, respectively.

The adequacy of the model was evaluated with reference to the coefficient of variance (C.V) and coefficient of regression (R²). The C.V value of 14.6 % indicates a high degree of precision and reliability of the experimental data (Table 3). Similarly, R² value 0.94, suggested that both predicted and experimental values were in close conformity to each other and 94 % of the variability in the response can be explained by the model (Fig. 1). The adequate precision measures the signal to noise ratio which was 13.76 % in case of present research indicating an adequate signal. The lack of fit was found to be non-significant which depicted that production of rhamnolipids can be further improved by selecting more closer ranges of nutritional parameters. The interactive effects of nutritional parameters on rhamnolipid production by *P. aeruginosa* was investigated by plotting response surface curves against any two independent variables while keeping other at the constant level.

The results revealed that NaNO₃ has a notable (*p*-value <0.0001) effect on rhamnolipid production by *P. aeruginosa* (Table 3). The strain JQ showed variable metabolic responses under the effect of nitrate ions in the medium.

Table 3. ANOVA of Selected Model Terms for Rhamnolipid [Y₁].

| Source | Sum of squares | Degree of freedom | Mean square | F-value | P-value |
|-----------------------------------|----------------|-------------------|-------------|---------|----------|
| Model | 52.43 | 10 | 5.24 | 38.81 | < 0.0001 |
| A-NaNO ₃ | 7.48 | 1 | 7.48 | 55.38 | < 0.0001 |
| B-KH ₂ PO ₄ | 6.18 | 1 | 6.18 | 45.71 | < 0.0001 |
| C-MgSO ₄ | 1.69 | 1 | 1.69 | 12.48 | 0.0022 |
| D-FeSO ₄ | 0.82 | 1 | 0.82 | 6.06 | 0.0236 |
| AC | 0.73 | 1 | 0.73 | 5.42 | 0.0310 |
| A ² | 14.99 | 1 | 14.99 | 110.96 | < 0.0001 |
| B ² | 11.66 | 1 | 11.66 | 86.31 | < 0.0001 |
| C ² | 9.59 | 1 | 9.59 | 70.98 | < 0.0001 |
| D ² | 13.72 | 1 | 13.72 | 101.56 | < 0.0001 |

R² = 0.9533, Adj. R² = 0.9288, Coefficient of Variance = 16.22 %, Adeq Precision = 18.30

Maximum biosurfactant yield was observed at 3.92 g/L of NaNO_3 , a point at the center of response surface curve. Whereas, synthesis of rhamnolipids was negatively affected when the concentration of NaNO_3 either increased or decreased from central reference point as indicated in (Fig. 2 A, B, C). Basically, role of nitrate has been already established in regulating biosurfactant production via activation of glutamine synthase pathway. Furthermore, inorganic nitrogen in the fermentation media facilitates expression of the aforesaid enzyme system through their transcriptional activators such as RpoN and σ factor (σ^{54}) (Chayabutra, *et al.*, 2001; Totten *et al.*, 1990). It has also been suggested that low carbon to nitrate ratio stimulates lipogenesis via glutamine synthase pathway leading to overproduction of rhamnolipid. In contrast, excessive nitrogen drives the metabolism towards protein synthesis. The results of the present findings are in allinece with some previous reports.

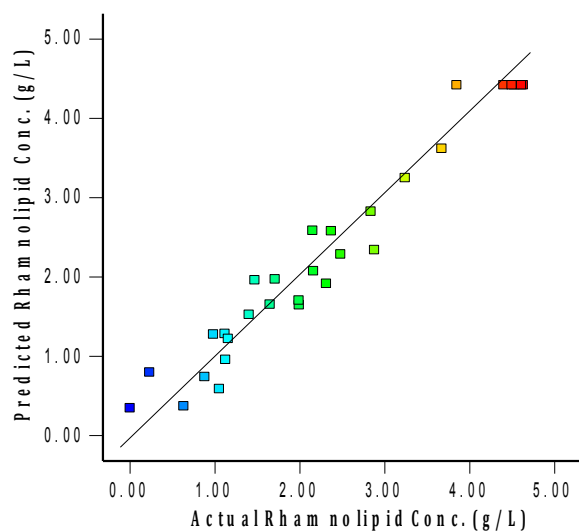


Figure 1. Predicted vs Actual Rhamnolipid Conc. (g/L).

Our results indicated that different Conc. of KH_2PO_4 was positively effecting ($p < 0.0001$) rhamnolipid production by *P. aeruginosa* JQ (Table 3). The optimum Conc. of KH_2PO_4 for highest rhamnolipid production was noted as 2.38 g/L. The production of rhamnolipid was decreased even at slight deviation from the peak area indicated in (Fig. 2 A, D, E). It was observed that production of rhamnolipid from strain JQ was improved significantly at low concentration of KH_2PO_4 in the medium. Phosphate limitation in culture media has been known to influence the expression of rhlR genes through various transcriptional activators such as Vfr, RhlR and σ^{54} (Bazire *et al.*, 2005). It could be suggested activation of RhlR genes under phosphate limitation may resulted in high yield of rhamnolipid by *P. aeruginosa* JQ. These results are in accordance with some previous reports which suggested a multi fold increase in biosurfactants production

under phosphate -limited conditions (Chayabutra *et al.*, 2001; Abalos *et al.*, 2002).

Magnesium is essential micronutrient participate in various enzyme catalyzed reactions such as transfer of phosphate and pyruvate metabolism. MgSO_4 was shown to be a significant model term with p-value of 0.0022 (Table 3). The maximum rhamnolipid was predicted at the center point of response surface plots corresponds to 0.26 g/L of the MgSO_4 (Fig. 2 B, D, F). Less yield of rhamnolipid was recorded while moving beyond this peak area indicating that not only the low concentration of MgSO_4 but also its ratio plays important role in rhamnolipid biosynthesis. It has been previously observed that suboptimal level of Mg^{++} effects not only the growth but also restrict activation of various enzymes involved in metabolic pathway associated with rhamnolipid biosynthesis. The results of present research are comparable with some previous research findings which suggested that Mg^{++} limitation in the media improve rhamnolipid production from *P. aeruginosa* (Roldan-Carrillo *et al.*, 2011).

Iron is a key micronutrient for energy deriving metabolic pathways such as oxidative phosphorylation and a critical parameter for the production of biosurfactants from different bacteria. It was noted that iron was a significant model term (p-value 0.023) for rhamnolipid production (Table 3). The amount of FeSO_4 pertaining to the highest rhamnolipid production was 0.028 g/L predicted at the middle of response surface plots (Fig. 2 C,E,F). The fluctuation from this optimum point hampered the production of rhamnolipid by *P. aeruginosa*. It has been recognized that an inverse relation exists between transcription of rhlAB gene and concentration of iron in the media (Deziel *et al.*, 2003; Glick *et al.*, 2010). Furthermore, limitation of iron has also been associated with stimulation of lasIR and rhlIR genes that increase production of rhamnolipid (Bollinger *et al.*, 2001; Duan and Surette, 2007). Our results are in conformity with the findings of (Persson *et al.*, 1990) indicated the dependency of rhamnolipid biosynthesis under restricted iron conditions.

The cumulative interaction between NaNO_3 and MgSO_4 was found to be significant while all other interactions were of less significance in present study (Table 3). The joint effect of $\text{NaNO}_3 \times \text{MgSO}_4$ is shown in the (Fig. 2 B). The amount of biosurfactant was gradually increased with the increase in NaNO_3 and MgSO_4 concentrations and reached at final optimal values of 3.92 g/L and 0.26 g/L, respectively. However, at further increase in the concentrations of these salts biosurfactant production was negatively affected. The findings of present research suggested that not only the concentration but the relative proportion of each media components affects the cellular productivity of rhamnolipid. It has been described frequently in the literature that low C:N, high C:P and high C:Fe ratios favor the production of biosurfactant from different bacterial strains including *P.*

aeruginosa (Makkar and Cameotra, 2002; Amezcua-Vega *et al.*, 2007; Kumar *et al.*, 2012). Therefore, appropriate ratios of media contents should be investigated for making bioprocess economically more productive.

The optimum concentration of the media components were predicted as 3.92 g/L NaNO₃, 2.73 g/L K₂HPO₄, 0.23 g/L MgSO₄ and 0.0028 g/L FeSO₄. These values were obtained by moving towards the center of the contour plot and validated by performing a conformity experiment in triplicate. In conclusion, 5.67 g/L of the rhamnolipid was produced under these reaction conditions leading a 2 fold increase in yield as compared to unoptimized media.

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