

Original Article

Optimization of Lipase Production with Response Surface Method using *Bacillus subtilis*

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Abstract

The production and application of enzymes can be costly, largely due to the need for inducible production systems. This study aimed to optimize media components for cost-effective extracellular lipase production from soil microorganisms. By using Response Surface Methodology (RSM), the research focused on maximizing lipase yield and efficiency through systematic analysis of optimal parameters. Molasses (g/L), peptone (g/L), and palm oil (% v/v) were selected as key components in the culture medium. Central Composite Design (CCD) was used to optimize their concentrations and evaluate their effects on lipase yield. The experimental design included a three-factor, five-level setup with 20 runs. Statistical analysis involved the correlation coefficient (R), multiple correlation coefficient (R²), and Fischer's F-statistic with its p-value. The highest lipase production, 145.86 U/mL, was achieved with 18 g/L molasses, 27 g/L peptone, and 0.1% v/v palm oil, with a predicted value of 160.24 U/mL. The model was statistically significant, with a p-value of 0.0143 and an F-value of 4.207578. The lack of fit was not significant (p-value of 0.3901), indicating a good model fit. This optimization strategy significantly improves microbial lipase production efficiency while offering a scalable industrial application model.

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1. Introduction:

Lipases (triacylglycerol acyl hydrolases, EC 3.1.1.3) are widespread enzymes present in animals, plants, fungi, and bacteria. They play a crucial physiological role by catalyzing the hydrolysis of triacyl glycerides into glycerides, monoglycerides, glycerol, and fatty acids at the lipid-water interface [1].

Lipases are highly versatile biocatalysts capable of facilitating a variety of bioconversion reactions, including hydrolysis, interesterification, esterification,

alcoholysis, acidolysis, and aminolysis. Due to the catalytic abilities of hydrolyzing triglycerides, esterification, trans-esterification, and enantio-separation, triacylglycerol lipase (EC 3.1.1.3) finds widespread applications across industries such as detergents, pharmaceuticals, food (cheese and tea), pulp and paper, textiles, tanneries, cosmetics, biodiesel production, wastewater treatment, medicine, food, chemistry, textiles, and related industries [2-4]. Microbial lipases are particularly notable for their

extensive industrial applications, ease of cultivation, and scalability in production [5, 6]. Microbial lipases are primarily produced through submerged fermentation (SmF) a well-established method with fully developed engineering aspects.

However, solid-state fermentation (SSF) has demonstrated certain advantages over SmF for enzyme production, even on a commercial scale [4]. The traditional approach of optimizing media for the fermentation process by varying one variable at a time was often time-consuming, costly, and likely to misinterpretation. However, the high production costs of biocatalysts frequently limit their widespread application [7].

To address this challenge, explore various microorganisms, supplements and substrates to identify the most effective combinations for producing high-value lipases. This approach focuses on using substrates and operational conditions that can lower production costs on an industrial scale [8]. To enhance microbial lipase production, various culture parameters are typically examined, including carbon sources, nitrogen sources, initial pH, temperature, and aeration conditions [9].

The high cost of carbon sources contributes to the expense of lipase production. To mitigate this, low-cost substrates such as molasses [10], glycerol [11], olive pomace and wheat bran [12] have been explored as viable alternatives. Edible triacyl glycerides, including olive oil, palm oil, sunflower oil, and soybean oil, have also been used as carbon sources and inducers in lipase production [13].

Optimization of media components for every possible combination of required carbon, nitrogen sources and inducers was impractical due to the large number of required trials. In contrast, statistical experimental methods provide a more efficient and cost-effective way to simultaneously and systematically vary all components [14].

Response Surface Methodology (RSM) is an experimental approach used to determine the optimal conditions in a multivariable system. It is particularly effective for optimizing significant variables identified through factorial designs, such as the central composite

design (CCD). By integrating factorial design and regression analysis, RSM evaluates the impact of various factors and their interactions. Advanced applications of RSM involve mathematical models to analyze experimental data and predict relationships between responses and variables, enabling the creation of predictive contours that facilitate faster and more efficient optimization with fewer experiments [15].

The main purpose of this study was to optimize the media components for cost-effective extracellular lipase production soil microorganisms. Utilizing the statistical optimization tool Response Surface Methodology (RSM), the study was conducted to enhance the efficiency and yield of lipase production. By systematically analyzing and refining the conditions, this approach aims to identify the optimal parameters that will maximize enzyme output and improve overall production processes.

2. Materials and Method:

2.1 Sample collection:

Soil sample was collected from agricultural fields from the Aurangabad district, at a depth of 5-6 cm. Using a sterile spatula, samples were placed in sterile 50 ml tubes to avoid contamination.

2.2. Isolation and screening:

For serial dilution, 1 g of soil sample was mixed with 10 ml of sterile distilled water in a 50 ml Erlenmeyer flask, then agitated at 120 rpm for 30 minutes at 37°C using a rotary shaker. The aqueous slurry was serially diluted up to 10^{-6} using 0.8% saline. Then, 100 μ l from each dilution was spread on tributyrin agar plates containing 0.5% peptone, 0.3% yeast extract, 1% tributyrin, and 2% agar. The plates were incubated at 37°C for 24-72 hours, and lipolytic activity was assessed by observing zones of hydrolysis around bacterial colonies. The isolates showing maximum zone of clearance were selected for further analysis [16].

2.3. Morphological, Biochemical and Molecular identification of the isolates

The morphological and biochemical characterization of lipase-producing bacterial isolates was conducted following guidelines from Bergey's Manual of Systematic Bacteriology [17]. The confirmation of the

isolates was done by molecular identification using 16S rDNA sequencing.

2.4. Response surface methodology

To determine the optimal media component for the lipase production from the isolated bacteria molasses (g/L), peptone (g/L), and palm oil (%v/v) were selected as components of the culture medium. Central Composite Design (CCD) was then utilized to optimize the concentrations of these components and to evaluate their individual effects and interactions on lipase yield[18]. The experimental setup included a three-factor, five-level design ($-\alpha$, -1 , 0 , $+1$, $+\alpha$) with 20 experimental runs, detailed in Table 1. To develop the quadratic model and analyze the multinomial coefficients experiments were conducted using Design Expert v11.1.2.0 software, with lipase production measured in U/mL as the response variable (R1) [19].

2.5. Statistical analysis

Statistical analysis included the evaluation of the correlation coefficient (R), multiple correlation coefficient (R^2) representing the fit of the quadratic model, and Fischer's F-statistic along with its probability p(F). Response surface curves were generated based on the quadratic model[20]. The analysis was done using Statease Design Expert version 7.

3. Result

3.1. Isolation and screening

Colonies exhibiting maximum zone on clearance were selected for quantitative measurement of lipase activity, as colonies with a larger zone of clearance typically indicate higher enzyme activity. Among the isolates, one strain BS-L1 demonstrated the highest zone of clearance and was thus used for further identification and enzyme production and studies.

3.2. Identification of the isolate

The isolated strain was show gram-positive in nature. The selected lipolytic isolate was identified using sequencing techniques, PCR amplification of the 16S rDNA gene. The resulting sequence was then compared with those in GenBank using the BLAST program. As per the results of the isolated bacteria was identified as the *Bacillus subtilis*.

3.3. Optimization of the media component by Response Surface Methodology

The experimental design focused on three variables: molasses, peptone, and palm oil, each tested at specific levels for lipase production using the CCD model. Molasses was assessed within a range of 2 to 18 g/L, with the extreme values of $-\alpha$ and $+\alpha$ set at -3.45434 and 23.45434 g/L, respectively. Peptone concentrations varied from 3 to 27 g/L, with $-\alpha$ and $+\alpha$ levels extending from -5.18151 to 35.18151 g/L. Palm oil was evaluated at concentrations ranging from 0.1% to 0.7% v/v, with $-\alpha$ and $+\alpha$ levels adjusted to -0.10454% and 0.904538%, respectively.

Table 2 and Figure 1, Figure 2 and Figure 3 provides a comprehensive overview of the experimental design and predicted responses for lipase production using Central Composite Design (CCD). The analysis highlights the highest and lowest actual and predicted values from the experiments. The highest actual lipase production occurred in Run 15, where 18 g/L molasses, 27 g/L peptone, and 0.1% v/v palm oil resulted in 145.86 U/mL, with a predicted value of 160.24 U/mL.

A similarly high production was observed in Run 19, which had the same conditions as Run 15, yielding 145.52 U/mL with a predicted value of 120.43 U/mL. Run 16, with 23.45 g/L molasses, 15 g/L peptone, and 0.4% v/v palm oil, achieved 141.89 U/mL compared to a predicted 127.27 U/mL. In Run 17, using 2 g/L molasses, 3 g/L peptone, and 0.7% v/v palm oil, the actual production was 140.80 U/mL, while the predicted value was 133.98 U/mL. Run 18, featuring 10 g/L molasses, 35.18 g/L peptone, and 0.4% v/v palm oil, resulted in 140.53 U/mL of lipase, with a predicted production of 132.61 U/mL.

On the lower end, run 13 recorded the lowest actual lipase production with 2 g/L molasses, 27 g/L peptone, and 0.7% v/v palm oil, yielding 49.62 U/mL, while the predicted response was 74.35 U/mL. Another lower production run was Run 3, which utilized 10 g/L molasses, -5.18 g/L peptone, and 0.4% v/v palm oil, resulting in an actual production of 98.48 U/mL and a predicted value of 108.25 U/mL.

Table 3 evaluated the impact of molasses, peptone, and palm oil, along with their interactions, on lipase production. The overall model was statistically significant, with a p-value of 0.0143 and an F-value of

4.207578, indicating that the combined effects of these variables and their interactions significantly influenced lipase production. Individually, molasses (A), peptone (B), and palm oil (C) didn't show significant effects on lipase production, as indicated by their respective p-values of 0.3492, 0.1597, and 0.6671.

However, the interaction between molasses and peptone (AB) was highly significant, with a p-value of 0.0073 and an F-value of 10.08944, demonstrating a strong synergistic effect on lipase production. Similarly, the interaction between peptone and palm oil (BC) was also significant, with a p-value of 0.0108 and an F-value of 8.830535, suggesting that these two factors together had a considerable impact on enzyme production. The interaction between molasses and palm oil (AC), although showing some effect, was not statistically significant, with a p-value of 0.1088.

The model's residuals, with a sum of squares of 3778.784 and a mean square of 290.6757, represented the unexplained variation, while the lack of fit test indicated that the model fit the data well, as the lack of fit was not significant (p-value of 0.3901). The total variation in the data was represented by the total sum of squares, which was 11117.03.

The Model F-value of 4.21 implies the model is significant. There is only a 1.43% chance that a "Model F-value" this large could occur due to noise. Values of "Prob > F" less than 0.0500 indicate model terms are significant. In this case, AB and BC are significant model terms. Values greater than 0.1000 indicate the model terms are not significant. If there are many insignificant model terms (not counting those required to support hierarchy), model reduction may improve your model.

Following is the equation obtained after optimization of lipase production using Molasses, Peptone and Palm oil.

$$\text{lipase Production} = +125.4732157 - 4.133280508 * \text{Molasses} + 0.598093203 * \text{Peptone} + 24.34416869 * \text{Palm oil} + 0.199444324 * \text{Molasses} * \text{Peptone} + 4.324480234 * \text{Molasses} * \text{Palm oil} - 4.975656267 * \text{Peptone} * \text{Palm oil}$$

4. Discussion:

This research successfully identified the most effective combinations of carbon sources, nitrogen sources, and inducers for enhancing lipase production by BS-L2. The central composite design (CCD) model, analyzed using

ANOVA, revealed significant interactions, particularly between molasses and peptone ($p = 0.0073$) and between peptone and palm oil ($p = 0.0108$). These findings underscore the synergistic effects of these components in optimizing lipase production. Molasses, an economical and readily available carbon source, supports microbial growth due to its high sugar content and essential trace elements [21]. Peptone, rich in proteins, amino acids, and vitamins, provides a favorable environment for microbial cell growth and enzyme production [22]. The interaction between molasses and peptone not only enhances enzyme yield but also reduces production costs, making it a practical choice for large-scale applications. Similar findings were reported by Zhu, Liu [18], where the interaction of these two substrates significantly improved lipase activity ($p = 0.0289$).

The role of peptone combined with low concentrations of coconut oil in maximizing lipase production was highlighted by Patel, Ray [23], where the optimized medium achieved significant enzyme yield. Coconut oil, unlike palm oil, does not act as a strong inducer [24]. However, the cost limitations of ammonium oxalate as a nitrogen source in their study underscore the importance of finding cost-effective alternatives, such as the molasses-peptone combination explored in the present research.

Tanyol, Uslu [22] demonstrated optimal lipase production using peptone, ammonium sulfate, and sunflower oil cake. However, the rapid consumption of inorganic nitrogen sources like ammonium sulfate may lead to enzyme repression [25], a limitation that highlights the advantage of organic sources such as peptone in sustaining enzyme synthesis. Although promising, their reported enzyme yield of 10.8 U/mL was notably lower than the current study's maximum production of 160 U/mL.

Vasiee, Behbahani [26] reported a superior lipase activity of 343 U/mL using a medium containing coriander seed extract, yeast extract, olive oil, and MgCl₂. Olive oil, as a less saturated lipid substrate compared to palm oil, likely served as a better inducer and substrate for lipase synthesis [27]. While the present study achieved lower enzyme activity, it

prioritized cost-effectiveness by utilizing low-cost substrates like molasses and palm oil. The choice of substrate underscores the trade-off between maximizing yield and maintaining economic viability. Future studies could explore alternative lipid sources and their potential to further enhance production.

Additionally, magnesium ions (MgCl₂) were identified as essential cofactors in Vasiee et al.'s study, stabilizing enzyme structure and enhancing catalytic efficiency [28]. Incorporating such cofactors in subsequent optimizations could significantly improve the outcomes of this research.

5. Conclusion:

The optimization of the media components for extracellular lipase production by *Bacillus subtilis* isolated from soil using Response Surface Methodology (RSM) was successfully performed. The Central Composite Design (CCD) model demonstrated that molasses, peptone, and palm oil significantly influence lipase production, with a notable synergistic effect observed between molasses and peptone. The statistical analysis ANOVA confirmed that these interactions play a critical role in enhancing enzyme yield. The study findings suggest that utilizing molasses and peptone, both cost-effective and readily available substrates, can reduce production costs while maximizing lipase yield.

This optimization approach not only enhances the efficiency of microbial lipase production but also provides a scalable model for industrial applications. The significant interactions identified between the media components highlight the potential for further optimization and refinement in commercial enzyme production, offering a valuable framework for improving the economic viability of microbial lipase production processes.

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Author contribution: SG: Developed an idea and verified the content, SD: Conducted experimental work and wrote the manuscript

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APPENDICES

Table 1: Variables and their levels for response surface methodology

| Name | Units | -1 Level | +1 Level | -alpha | +alpha |
|----------|---------|----------|----------|----------|----------|
| Molasses | (g/L) | 2 | 18 | -3.45434 | 23.45434 |
| Peptone | (g/L) | 3 | 27 | -5.18151 | 35.18151 |
| Palm oil | (% v/v) | 0.1 | 0.7 | -0.10454 | 0.904538 |

Table 2: Experimental design and Predicted responses using CCD for lipase production.

| Run | Molasses (g/L) | Peptone (g/L) | Palm oil (% v/v) | Actual responses (U/mL) | Predicted responses (U/mL) |
|-----|----------------|---------------|------------------|-------------------------|----------------------------|
| 1 | 10 | 15 | 0.4 | 126.0847 | 120.4309027 |
| 2 | 18 | 3 | 0.1 | 88.8071 | 71.6337116 |
| 3 | 10 | -5.181513966 | 0.4 | 98.47786 | 108.2472113 |
| 4 | 10 | 15 | 0.4 | 98.40901 | 120.4309027 |
| 5 | 18 | 27 | 0.7 | 140.2274 | 141.5405228 |
| 6 | 10 | 15 | 0.904537849 | 135.1034 | 117.3759794 |
| 7 | 10 | 15 | 0.4 | 116.4857 | 120.4309027 |
| 8 | -3.454342644 | 15 | 0.4 | 126.5568 | 113.588399 |
| 9 | 10 | 15 | 0.4 | 116.3874 | 120.4309027 |
| 10 | 10 | 15 | 0.4 | 115.04 | 120.4309027 |
| 11 | 18 | 3 | 0.7 | 106.5917 | 124.5830053 |
| 12 | 2 | 3 | 0.1 | 116.2989 | 122.5473728 |
| 13 | 2 | 27 | 0.7 | 49.61758 | 74.35255347 |
| 14 | 10 | 15 | -0.104537849 | 114.9318 | 123.485826 |
| 15 | 18 | 27 | 0.1 | 145.8629 | 160.2406794 |
| 16 | 23.45434264 | 15 | 0.4 | 141.8895 | 127.2734064 |
| 17 | 2 | 3 | 0.7 | 140.7979 | 133.9816562 |
| 18 | 10 | 35.18151397 | 0.4 | 140.5323 | 132.6145941 |
| 19 | 10 | 15 | 0.4 | 145.5187 | 120.4309027 |
| 20 | 2 | 27 | 0.1 | 144.9974 | 134.5677203 |

Table 3: Analysis of variance

| Source Model | Sum of Squares | df | Mean Square | F-Value | p-value | Prob > F |
|--------------|----------------|----|-------------|----------|---------|-----------------|
| Model | 7338.244 | 6 | 1223.041 | 4.207578 | 0.0143 | Significant |
| A-Molasses | 274.1396 | 1 | 274.1396 | 0.943111 | 0.3492 | |
| B-Peptone | 646.7362 | 1 | 646.7362 | 2.224941 | 0.1597 | |
| C-Palm oil | 56.29722 | 1 | 56.29722 | 0.193677 | 0.6671 | |
| AB | 2932.755 | 1 | 2932.755 | 10.08944 | 0.0073 | |
| AC | 861.748 | 1 | 861.748 | 2.964637 | 0.1088 | |
| BC | 2566.822 | 1 | 2566.822 | 8.830535 | 0.0108 | |
| Residual | 3778.784 | 13 | 290.6757 | | | |
| Lack of Fit | 2575.102 | 8 | 321.8878 | 1.337097 | 0.3901 | Not Significant |
| Pure Error | 1203.682 | 5 | 240.7364 | | | |
| Cor Total | 11117.03 | 19 | | | | |

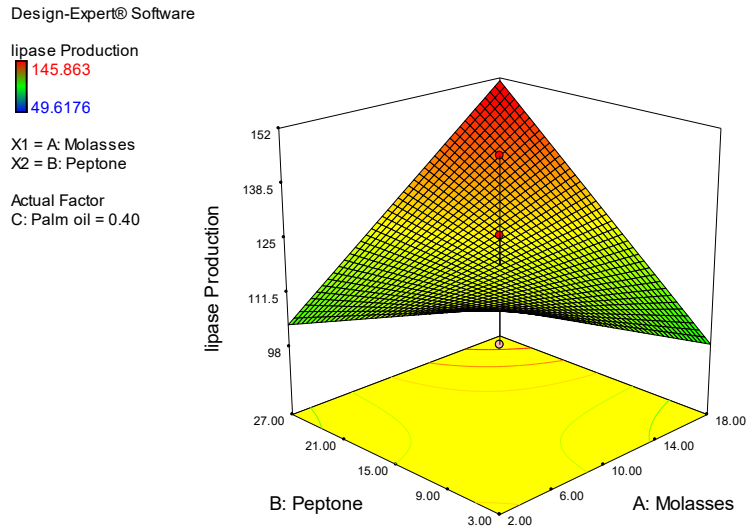


Figure 1: Molasses and peptone at a fixed level of Palm oil

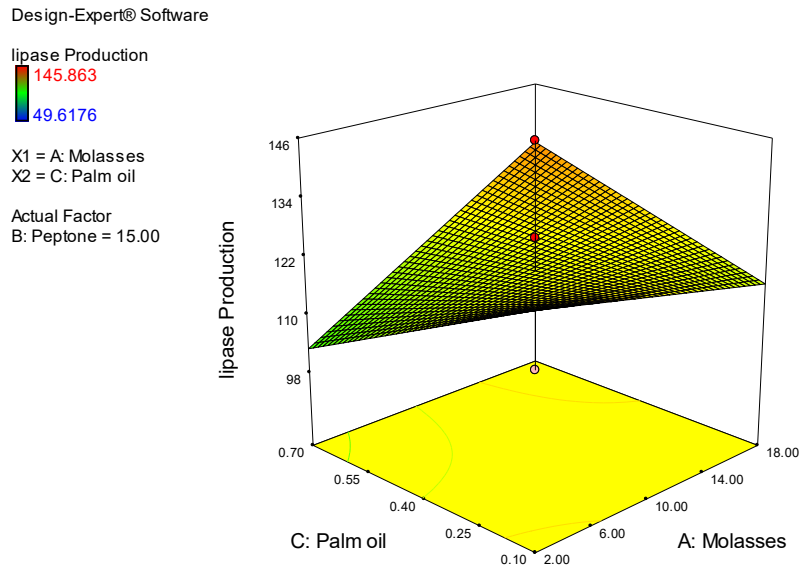


Figure 2: Palm oil and molasses at a fixed level of peptone

Design-Expert® Software

lipase Production



X1 = B: Peptone
X2 = C: Palm oil

Actual Factor
A: Molasses = 10.00

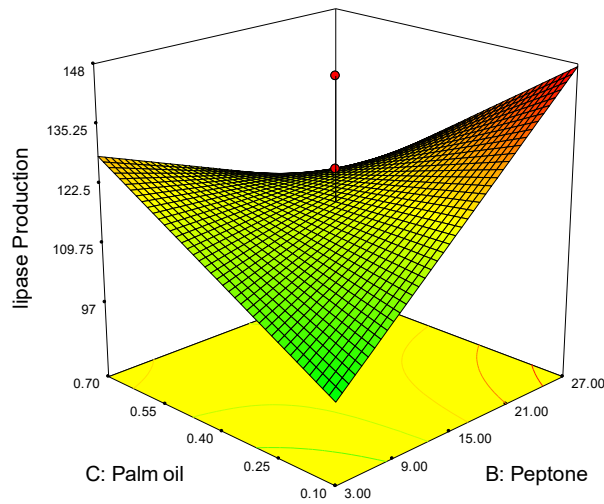


Figure 2: Palm oil and molasses at a fixed level of peptone

Design-Expert® Software

lipase Production



X1 = B: Peptone
X2 = C: Palm oil

Actual Factor
A: Molasses = 10.00

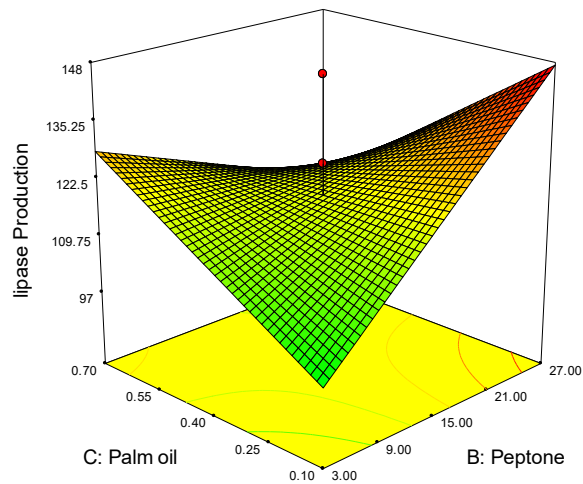


Figure 3: Peptone and palm oil at a fixed level of molasses