

Original Article

OPTIMIZATION OF MEDIUM COMPONENTS FOR ANTIBACTERIAL METABOLITE PRODUCTION FROM MARINE *STREPTOMYCES* SP. PUA2 USING RESPONSE SURFACE METHODOLOGY

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ABSTRACT

**Objective:** The present study is an attempt to optimize the fermentation conditions for the antibacterial compound production from a newly isolated marine *Streptomyces* strain PUA2 by adopting response surface methodology as the statistical tool.

**Methods:** Prior to using the Response Surface Methodology, Plackett Burmann (PB) design was used to explore the effect of variables on the antibacterial compound production. In PB method, high and low values were assigned for the eight variables viz., glucose, glycerol, soybean meal, manganese chloride, calcium carbonate, peptone and pH. Calcium carbonate and peptone were used as dummy variables. Based on the results of combined effects glycerol, soybean meal, manganese chloride and pH were investigated by 2<sup>4</sup> full-factorial central composite design.

**Results:** The results of PB method showed the significant effect of glycerol, soybean meal, manganese chloride and pH on the antibacterial compound production. The results of ANOVA and regression of second order model showed that the linear effects of glycerol and manganese chloride and cross products effects of manganese chloride and pH were more significant. All the critical variables having greatest effect on the production of antibacterial compound from marine *Streptomyces* species PUA2. Optimization of process parameters resulted in increase in antibacterial activity from 7 mm to 14 mm.

**Conclusion:** The factors optimized in the present study were useful for the increased production of antibacterial metabolite from *Streptomyces* sp PUA2. The result confirms the feasibility of medium optimization to improve antibiotic production.

**Keywords:** Antibacterial activity, Marine actinobacteria, Central Composite Design, Plackett Burmann design, Response surface methodology.

INTRODUCTION

Marine actinobacteria are metabolically active and physiologically adaptive to growth in sea water. They are the promising source for novel secondary metabolites with different biological activities like antimicrobial, anticancer, antiviral, anti-inflammatory and antiparasitic, etc. Even with the limited screening efforts dedicated to date to marine actinobacteria, the discovery rate of novel secondary metabolites from marine actinobacteria has recently surpassed that of their terrestrial counterparts, as evident by the isolation of many new chemical entities from marine actinobacteria [1-2]. During the period of 2003-2005, years, 659 marine bacterial compounds have been described in which 256 compounds have originated from actinobacteria [3].

Bioactive metabolites from actinobacteria are commonly produced by submerged or solid state fermentation with one or two commonly used media. Optimizing the suitable conditions and medium components is a prerequisite for better production of antibiotics or any other metabolites. Medium optimization explores a sequence of phases with a specific set of optimal conditions fixed by different methodologies.

Nutritional aspects like carbon source, nitrogen source and minerals and other conditions like time of incubation, pH and temperature are known to have profound effect on the production of antimicrobial metabolites by actinobacteria. Hence optimization of culture conditions is very essential to get maximum yield of antimicrobial metabolites [4-6]. The optimization of the medium by the conventional research technique, the classical one factor at a time method involves changing one independent variable like carbon source and nitrogen source, while fixing others at constant levels. Optimal medium is not always obtainable by the traditional one factor at a time optimization strategy because of potential interactions among medium components [7]. PB design is a well

known and widely used statistical technique for screening and selection of most significant culture variables. PB design offers a good and fast screening procedure and mathematically it computes the significance of a large number of factor variables in one experiment, which saves time and maintains convincing information on each component [8-9]. This method allows the screening of main factors from a large number of experiments that can be retained in further optimization processes.

RSM is a powerful technique for testing multiple process variables owing to fewer experimental trials that are needed compared to the study of one variable at a time. Also, significant interactions between the variables can be identified and quantified by this technique. RSM has been widely used to evaluate and understand the interactions between different physiological and nutritional parameters. A prior knowledge and understanding of these parameters are necessary for achieving a more realistic model [10]. The present study reports the effect of critical medium components on antibacterial metabolite production from a marine *Streptomyces* sp PUA2 using Plackett Burmann design and Response Surface Methodology (RSM).

MATERIALS AND METHODS

*Streptomyces* sp PUA2 strain

*Streptomyces* sp. PUA2 was isolated from marine sediment collected from Andaman ecosystems (Lat. 11.68°N; Long 92.77°E) using starch casein agar medium. It produced dirty white powdery colonies with diffusible pigment production on ISP2 agar medium. Viability of strain PUA2 was maintained on ISP2 agar slants at 4°C. Crude ethyl acetate extract prepared from *Streptomyces* sp PUA2 showed promising activity against the gram positive and gram negative bacterial pathogens. In an earlier study, effect of critical medium components on bioactive metabolite production from *Streptomyces* sp PUA2 was determined by classical one factor at a time method [11].



Fig. 1: Growth of *Streptomyces* sp PUA2 on ISP2 agar

### Inoculum preparation

Well grown culture of *Streptomyces* sp PUA2 was transferred in to 100 ml yeast extract malt extract (YEME) and kept in rotary shaker for 48 h. Basal medium consists of 1% glucose; 1% yeast extract and 0.1% NaCl. The pH of the medium was adjusted to 7.0 prior to sterilization. Each 25 ml of basal medium was prepared and inoculated with different variables in respective concentrations.

### Experimental procedure

Antibacterial activity of cell free supernatant was tested by agar well diffusion method. Previously cultured *Staphylococcus aureus*

MTCC96 was inoculated on nutrient agar plates using sterile cotton swab. A 5 mm diameter well was made on the nutrient agar plates and loaded with 50  $\mu$ l cell free supernatant. The inoculated plates were incubated at 37°C and were kept under observation for 24 hours for determining the zone of inhibition. Dry weight of biomass was estimated using preweighed tube and expressed in mg of biomass per ml of the medium.

### Plackett Burmann Screening Design (PB)

PB design was used for screening the most significant factors affecting the production of antibacterial metabolites. Each independent variable was tested at high and low levels indicated by + and - respectively. Effect of variables on bioactive metabolite production was determined by applying a two level factorial design. Different combinations and sequence of process conditions were investigated to determine the growth conditions best suited for growth and biomass production [12]. High and low values were assigned for the selected variables. Plackett Burmann designs for the seven variables were assigned as  $X_1, X_2, X_3, X_4, X_5, X_6,$  and  $X_7$  at high and low levels in which two factors,  $X_4$  and  $X_6$  were designated as 'dummy' variables. The factors or independent variables considered for design include glucose ( $X_1$ ), glycerol ( $X_2$ ), soybean meal ( $X_3$ ), calcium carbonate ( $X_4$ ), manganese chloride ( $X_5$ ), peptone ( $X_6$ ) and pH ( $X_7$ ) respectively. The results obtained with classical experiments in the previous study [11] were used in the selection of independent and dummy variables.

Table 1: Medium components and test levels for plackett-burmann experiment

Variables	Medium components	Low Level (-)	High Level (+)
$X_1$	Glucose	0.5	1.5
$X_2$	Glycerol	0.5	1.5
$X_3$	Manganese chloride	0.05	0.15
$X_4$	Calcium carbonate	0.5	0.5
$X_5$	Soybean Meal	0.05	0.15
$X_6$	Peptone	0.5	0.5
$X_7$	pH	6	8

### Response surface methodology using central composite design (CCD)

Central composite design is one of the response surface methodologies. For statistical calculations the variables  $X_i$  were coded as  $x_i$  according to Equation 1.

$$Xi = (x_i - x_0) / \Delta x_i, (i=1, 2, 3 \dots k) \dots \dots (1)$$

Where  $X_i$  is (dimensionless) coded value of the real variable  $x_i$ ,  $x_0$  is the real value of  $X_i$  at the center point (zero) level, and the  $\Delta x_i$  is the step change value. Each variable in the design were coded at five levels -2, -1, 0, 1, and 2 with all variables taken at a central coded value of zero. Accordingly, a  $2^k$  factorial experiment design, with an axial point (10mm) and six replicated at the center point, with a total number of 30 experiments were employed. Second degree polynomials, which include all interaction term, were used to calculate the predicted response in equation 2.

$$Y = \beta_0 + \sum \beta_i X_i + \sum \beta_{ii} X_i^2 + \sum \beta_{ij} X_i X_j \dots \dots (2)$$

Where Y represents response variables,  $\beta_0$  is the interception coefficient,  $\beta_i$  coefficient of the linear effect,  $\beta_{ii}$ , the coefficient of quadratic effect and  $\beta_{ij}$ , the coefficient of interaction effect.

## RESULT AND DISCUSSION

### Optimization by Plackett Burmann method

Plackett-Burmann statistical design was applied to reflect the relative importance of various fermentation factors. Seven different variables including fermentation conditions and medium constitution were chose to perform this optimization process. Antibacterial activity was measured after 120 hours of incubation [13]. The design of independent variables coded with the

corresponding medium composition values with high level (H) and low level (L). The high level of each variable was set far enough from the low level to identify which ingredients of the media have significant influence on the antibacterial metabolite production [14]. The main effect of each variable on antibacterial activity was estimated as the difference between both averages of measurements made at the high level and at the low level of that factor. The data given in Table 2 showed a wide variation from 0 to 8 mm of inhibition zone. The variation reflects the importance of medium optimization to attain higher productivity. The main effects of the examined factors on the antimicrobial activity were calculated shown in Table 3. According to the resulting effects of 7 variables on antibacterial activity, glycerol ( $X_1$ ),

### Effect of sources and their interaction on biomass

For each trial carried out experimentally by the Plackett Burmann design for the different combination of high and low values, biomass were determined. Trial 2 recorded highest biomass weight (17.7 mg/mL) while trial 8 has lowest biomass weight (6 mg/mL) [17-18]. The values were shown in Table 2 and data were analyzed graphically in figure 2.

### Optimization by Response Surface Methodology

The optimum concentrations of the individual components in the production medium for antibacterial compound production by *Streptomyces* sp PUA2 were further optimized by the Response Surface Methodology. According to this design, the total number of treatment combinations was  $2^k + 2k + n_0$  where k is the number of independent variables and  $n_0$  is number of repetitions of experiment at centre point. Based on the results of PB design, four critical components of the production medium were selected. For RSM analysis based on CCD, 30 experiments were performed in triplicate. The coded levels of the independent variables are given in

Table 4. A 2<sup>4</sup> factorial CCD was developed by design expert package version 9.0.2 with 8 axial points and 6 replicates at the center points leading to 30 runs [19]. Multiple regression analysis was applied on the experimental data by the CCD design. The Results were fitted

with a second order full polynomial equation. The empirical relationship between antibacterial metabolite production and the four test variables n coded units obtained the application of RSM is given by below equation.

**Table 2: Plackett burmann experimental design for evaluating factors influencing antibacterial metabolite production**

Trial No	Variables							Antibacterial activity against <i>S. aureus</i> MTCC96(mm)	
	X <sub>1</sub>	X <sub>2</sub>	X <sub>3</sub>	X <sub>4</sub>	X <sub>5</sub>	X <sub>6</sub>	X <sub>7</sub>		Biomass(mg/mL)
1	H	H	H	L	H	L	L	6.5	8.3
2	L	H	H	H	L	H	L	6	17.7
3	L	L	H	H	H	L	H	0	8.4
4	H	L	L	H	H	H	L	8	8.4
5	L	H	L	L	H	H	H	7	13.7
6	H	L	H	L	L	H	H	0	10
7	H	H	L	H	L	L	H	6	8.2
8	L	L	L	L	L	L	L	4	6

**Table 3: Statistical calculations for plackett burmann method**

	X <sub>1</sub>	X <sub>2</sub>	X <sub>3</sub>	X <sub>4</sub>	X <sub>5</sub>	X <sub>6</sub>	X <sub>7</sub>
Sum of [H]	20.5	25.5	12.5	20	21.5	21	13
Sum of [L]	17	12	25	17.5	16	16.5	24.5
Effect	0.875	3.38	-3.13	0.63	1.38	1.13	-2.88
MS	1.531	22.78	19.53	0.78	3.78	2.53	16.53
F-test	0.92	13.80	11.83	0.472	2.29	1.53	10.01

MS-mean square;EMS-error mean square = 1.65, soybean meal (X<sub>3</sub>), pH (X<sub>7</sub>) and manganese chloride (X<sub>5</sub>) have significantly influenced antibacterial metabolite production and the dummy variables (X<sub>4</sub> & X<sub>6</sub>) and glucose (X<sub>1</sub>) included in the design have a least effect on the antibacterial compound production based on F-test values. The mean square (variance of effect) showed higher value for glycerol. The highest antibacterial activity was recorded in trial 4 while the lowest antibacterial activity was in Trial 8 and no activity in Trial 5 and 6 [15-16].

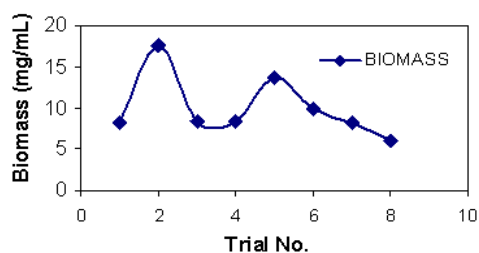
**Table 4: Range and levels of different process variables used in ccd for antibacterial activity**

Variables	-2	-1	0	1	2
Glycerol(X <sub>1</sub> ) g/mL	5	7.5	10	12.5	15
Soybean meal (X <sub>2</sub> ) g/mL	5	7.5	10	12.5	15
Manganese chloride (X <sub>3</sub> ) g/mL	0.5	0.75	1	1.25	1.5
pH(X <sub>4</sub> ) 1-9	5	6	7	8	9

$$Y = 12.17 + 0.083X_1 + 4.53 \times 10^{-17} X_2 + 0.41X_3 + 0.33X_4 - 1.82 \times 10^{-16} X_1X_2 - 2.29 \times 10^{-16} X_1X_3 - 3.94 \times 10^{-16} X_1X_4 + 0.13X_2X_3 - 0.12X_2X_4 + 0.13X_3X_4 - 1.08X_1^2 - 1.08X_2^2 - 0.96X_3^2 - 0.58X_4^2$$

Where Y is the measured response (antibacterial activity units in mm), X<sub>1</sub>, X<sub>2</sub>, X<sub>3</sub> and X<sub>4</sub> are the coded values of independent variables. CCD results on experiments with different combination for X<sub>1</sub>, X<sub>2</sub>, X<sub>3</sub>, and X<sub>4</sub> are presented along with predicted and experimental values for antibacterial activity shown in table 5.

The coefficient of determination (R<sup>2</sup>) was found to be 0.755 for the antibacterial metabolite production, indicating 75.5 % of variability in the response could be explained by the statistical model. The closer the R<sup>2</sup> is to 1.0, the stronger the model and the better it predicts the response. In this case, the value of the determination of coefficient elucidated the validity of the model. The adjusted R<sup>2</sup> value corrects the R<sup>2</sup> value for the sample size and for the number of terms in the model.



**Fig. 2: Showing Biomass (mg/ml) vs. Trial no in Plackett Burmann Method.**

The value of the adjusted determination coefficient (Adj R<sup>2</sup> = 0.53) was also high supporting the significance of the model [20]. In this case the adjusted R<sup>2</sup> was noticeably smaller than the R<sup>2</sup> value since there are many variables in the model and their sample size was also not very large [21]. The coefficient of variation of model was 4.22% and the PRESS statistic value of 118.32 proved the validity of model.

There is a 38.03% chance that a lack of fit has occurred and this could be due to noise. Non-significant lack of fit is good for an apt model showing excellent optimization [22-23]. Response surface contour plots and their three dimensional graphs are employed to understand the relationship between the response and experimental values of each variable. These plots also showed the type of interaction between test variables and helped to obtain the optimum conditions [24]. A total of six response surface plots were obtained by the central composite design. Maximum zone of inhibition showing the antibacterial activity (12.09 mm) was found during the interaction of soybean meal and Glycerol by fixing the quantity of manganese chloride and pH (Fig.1a). The antibacterial activity was observed in terms of the zone of inhibition, 10.78 mm as shown in

the Fig.1b at pH 7 and glycerol (10g/mL). Similar values of the inhibition zone were observed during the interaction of Manganese chloride (1g/mL) and glycerol (10g/mL). The zone of inhibition was 10.67 mm during this interaction (Fig.1c). A comparable inhibition zone of 12.01 mm (Fig.1d) was obtained at pH 7 and soybean meal (10g/ml),

The interaction of Manganese chloride (1gm/ml) and Soybean meal(10g/ml) showed an intermediate zone of inhibition, whose value was observed as 11mm (Fig 1e). Figure.1f shows the interaction between manganese chloride (1g/ml) and pH 7 and this interaction produced an inhibition zone of 11.5 mm [25-27].

Table 5: Central composite design matrix in coded values and responses

Run	Variables				Antibacterial activity against <i>S.aureus</i> MTCC96(mm)	
	X <sub>1</sub>	X <sub>2</sub>	X <sub>3</sub>	X <sub>4</sub>	Experimental value	Predicted value
1	-1	-1	-1	-1	7	7.726
2	1	-1	-1	-1	7	8.117
3	-1	1	-1	-1	7	8.547
4	1	1	-1	-1	7	10.683
5	-1	-1	1	-1	7	10.131
6	1	-1	1	-1	8	11.762
7	-1	1	1	-1	8	11.102
8	1	1	1	-1	8	7.279
9	-1	-1	-1	1	8	7.534
10	1	-1	-1	1	8	7.359
11	-1	1	-1	1	8	7.415
12	1	1	-1	1	8	11.943
13	-1	-1	1	1	9	11.247
14	1	-1	1	1	9	12.082
15	-1	1	1	1	9	11.187
16	1	1	1	1	9	9.195
17	-2	0	0	0	9	8.70
18	2	0	0	0	9	8.557
19	0	2	0	0	9	9.517
20	0	-2	0	0	9	7.141
21	0	2	0	0	9	14.269
22	0	0	2	0	10	9.141
23	0	0	-2	0	11	9.269
24	0	0	0	-2	11	8.308
25	0	0	0	2	11	8.308
26	0	0	0	0	11	8.308
27	0	0	0	0	12	8.308
28	0	0	0	0	12	8.308
29	0	0	0	0	13	8.308
30	0	0	0	0	14	8.308

Table 6: Analysis of variance (anova), regression coefficient estimate and test of significance for antibacterial activity in mm (r1)

Source	Sum of Squares	Df	F Value	P Value Prob > F
Model	79.13	14	3.30	0.0141
X <sub>1</sub>	0.17	1	0.097	0.7593
X <sub>2</sub>	2.842×10 <sup>-14</sup>	1	1.661×10 <sup>-14</sup>	1.0000
X <sub>3</sub>	4.17	1	2.44	0.1395
X <sub>4</sub>	2.67	1	1.56	0.2310
X <sub>1</sub> <sup>2</sup>	32.19	1	18.81	0.0006
X <sub>2</sub> <sup>2</sup>	32.19	1	18.81	0.0006
X <sub>3</sub> <sup>2</sup>	25.19	1	14.72	0.0016
X <sub>4</sub> <sup>2</sup>	9.33	1	5.45	0.0338
X <sub>1</sub> X <sub>2</sub>	4.263×10 <sup>-14</sup>	1	2.492×10 <sup>-14</sup>	1.0000
X <sub>1</sub> X <sub>3</sub>	2.842×10 <sup>-14</sup>	1	1.661×10 <sup>-14</sup>	1.0000
X <sub>1</sub> X <sub>4</sub>	2.842×10 <sup>-14</sup>	1	1.661×10 <sup>-14</sup>	1.0000
X <sub>2</sub> X <sub>3</sub>	0.25	1	0.15	0.7076
X <sub>2</sub> X <sub>4</sub>	0.25	1	0.15	0.7076
X <sub>3</sub> X <sub>4</sub>	0.25	1	0.15	0.7076
Residual	25.67	15		
Lack of Fit	18.83	10	1.38	0.3803
Pure Error	6.83	5		
Correlation Total	104.80	29		

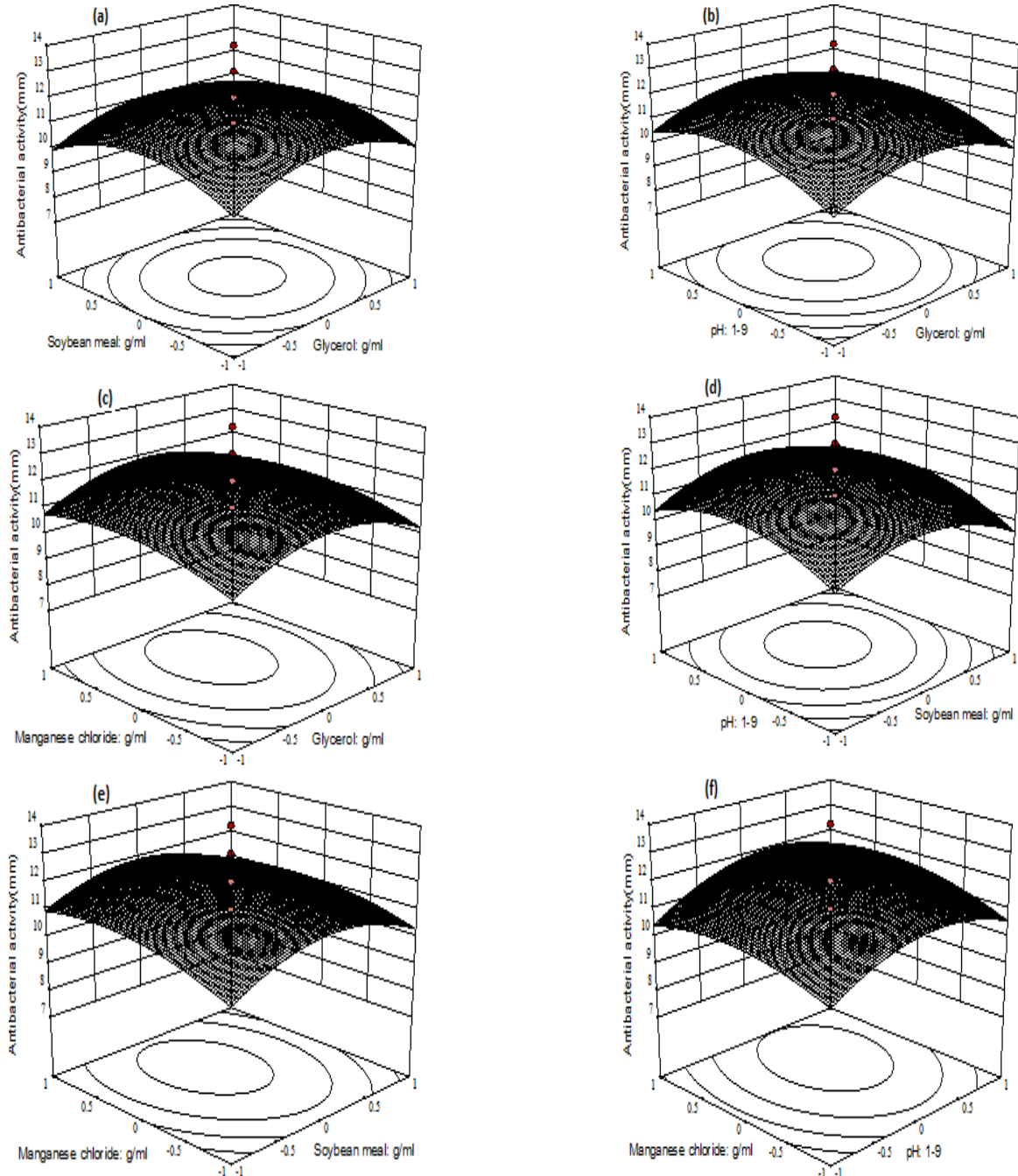
Statistical significance of model was investigated by ANOVA as shown in Table 6. The F value of 3.30 implies the significance of the model. P values of the model (0.0141) indicate the significance of the model terms. In this case X<sub>1</sub><sup>2</sup>, X<sub>2</sub><sup>2</sup>, X<sub>3</sub><sup>2</sup> and X<sub>4</sub><sup>2</sup> are significant model terms. The p value of the Lack of fit term was 1.38 implying its non-significant relative to pure error.

In Response plots and contour diagrams (Fig.3), contour diagrams (Fig.1a,1d, 1e, 1f) were elliptical and tilted, indicating significant

cross product interaction between the factors X<sub>1</sub> and X<sub>2</sub>, X<sub>2</sub> and X<sub>3</sub>, X<sub>2</sub> and X<sub>4</sub>, X<sub>3</sub> and X<sub>4</sub> [28-29]. This is also evident from coefficient

estimate values (Table 6). Contour diagrams of Figure 1b and Figure 1d are elliptical but not tilted, indicating a good deal of negligible interaction between ( $X_1$  &  $X_3$  and  $X_1$  &  $X_4$ ). This implies that the linear effects of glycerol, manganese chloride and pH are significant. The interactive effect of soya bean meal with manganese chloride and pH is more significant than the other factors. These suggest that

the concentrations of glycerol, manganese chloride and pH have a direct relationship with the production of antibiotic and creation of the zone of inhibition. In the present study, the experimental results based on response surface approach clearly showed that the antibiotic production was dependent on the optimized values of the medium components taken into consideration [30-32].



**Fig. 3: Response surface plots for Antibacterial activity: a) Soy bean meal vs. glycerol; b) pH vs. Glycerol; c) Manganese chloride vs. Glycerol; d) pH vs. Soy bean meal; e) Manganese chloride vs. Soybean meal; f) Manganese Chloride vs. pH**

#### CONFLICT OF INTERESTS

Declared None

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