



ISSN: 2230-9926

Available online at <http://www.journalijdr.com>

IJDR

International Journal of
DEVELOPMENT RESEARCH

International Journal of Development Research
Vol. 5, Issue, 04, pp. 4036-4042, April, 2015

Full Length Research Article

OPTIMIZATION OF INDOLE-3-ACETIC ACID PRODUCTION BY *BACILLUS SUBTILIS* TIB6 USING RESPONSE SURFACE METHODOLOGY

^{1,*}Ton That Huu Dat, ²Nguyen Thị Kim Cuc and ¹Pham Viet Cuong

¹Mientrung Institute of Scientific Research, Vietnam Academy of Science and Technology

²Institute of Marine Biochemistry, Vietnam Academy of Science and Technology

ARTICLE INFO

Article History:

Received 07th January, 2015

Received in revised form

11th February, 2015

Accepted 04th March, 2015

Published online 29th April, 2015

Key words:

Bacillus subtilis,

Central Composite Design,

Indole-3-acetic acid,

Plackett-Burman Design,

Response Surface Methodology

ABSTRACT

The influence of carbon, nitrogen sources and tryptophan supplementation on indole-3-acetic acid (IAA) production of *Bacillus subtilis* TIB6 isolated from pepper rhizosphere, Tay Nguyen province of Viet Nam was investigated and obtained results showed that sucrose and yeast extract were the most favorable carbon and nitrogen sources for TIB6's IAA production, respectively, while the presence of L-tryptophan in culture medium stimulated the formation of IAA. The student's t-test of the Plackett-Burman design were used for screening significant factors affecting IAA producing capacity of TIB6 and obtained results showed that L-tryptophan, yeast extract concentrations and inoculation ratio were the main effecting parameters. By using central composite design (CCD) of response surface methodology (RSM), the culture conditions for TIB6 were optimized for maximum IAA production. The optimum values of variables were: L-tryptophan (0.088%), Yeast extract (0.82%) and inoculation ratio (1.65%). At these conditions, the maximum IAA production was estimated to be 120,846 mg/L.

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INTRODUCTION

Plant growth promoting rhizobacteria play a crucial role in agriculture by increasing the exchange of plant nutrients and reducing the use of chemical fertilizers. There are several mechanisms by which rhizosphere bacteria may stimulate plant growth. One of the most commonly reported direct growth promotion mechanism by bacteria is the production of plant growth promoting compounds. IAA (indole-3-acetic acid) is known as the member of the group of phytohormones and is considered the most important native auxin (Ashrafuzzaman et al., 2009), involving in increasing the root growth and length as well as wide root surface area, which enables the plant to easily access to more nutrients from soil (Vessey, 2003). IAA is produced through L-tryptophan metabolism by bacteria, fungus and algae (Sarwar and Kremer, 1995; Stein et al., 1990; Finnie and Van Staden, 1985; Rifat et al., 2010). Bacteria from different genera (*Azospirillum*, *Enterobacter*, *Azotobacter*,

Klebsiella, *Alcaligenes faecalis*), actinomycetes (*Streptomyces olivaceoviridis*, *Streptomyces rimosus*) and fungi (*Colletotrichum gloeosporioides*, *Ustilago maydis*) have shown to stimulate plant growth by the synthesis of IAA (Torres-Rubio et al., 2000; Reineke et al., 2008; Khamna et al., 2010). Root exudates are natural source of L-tryptophan for rhizospheric microflora, which may promote IAA biosynthesis in the rhizosphere (Glick et al., 1995), thus IAA is considered as secondary metabolites (Ahmad et al., 2005). Apart from detecting and screening IAA producing microorganisms, optimization of cultural conditions for IAA production improvement of the isolated strain was essential. Before now, the classical optimization method was used popularly in which "one factor at a time" was studied, i.e. only one factor of a system being variable, whereas the others were fixed during each experiment and therefore, much trials were required (Sasirekha et al., 2012). Additionally, this laborious and time consuming method does not guarantee the determination of optimal conditions (Aravindan et al., 2007). Statistical approach for medium optimization would effectively solve these problems, which involves specific design of trials to minimize the error in determining the effect of variables (Krishnan et al., 1998). Statistical methods have benefits over conventional methods in predicting the accurate

*Corresponding author: Ton That Huu Dat

Mientrung Institute of Scientific Research, Vietnam Academy of Science and Technology

results based on utilization of fundamental principles of statistics, randomization, and replication (Sasirekha *et al.*, 2012). In this study, we applied statistical approach for optimization of culture conditions for IAA production of *Bacillus subtilis* TIB6 isolated from pepper rhizosphere, Tay Nguyen province of Vietnam

MATERIALS AND METHODS

Bacillus subtilis TIB6 strain

Bacillus subtilis TIB6 strain was isolated from pepper rhizosphere in Vietnam and the identification was carried out based on morphological, biochemical characteristics and 16S rDNA sequence analysis. Strain was maintained on nutrient agar slant at 4°C and stored at Vietnam Academy of Science and Technology. Inoculum was prepared in nutrient broth (g/L): glucose 10, peptone 5, KH₂PO₄ 0.3, MgSO₄.7H₂O 0.5, NaCl 3 by transferring a loop of the bacteria from stock culture and incubated at 37°C, shaking 200 rpm for 24 h.

Estimation of IAA production

The IAA produced was estimated by previously described method (Gordon and Weber, 1951). One ml of the cell-free supernatant was added to 4 ml of the Salkovski’s reagent and this preparation was incubated for 30 minutes at room temperature in darkness. Presence of pink color pointed out IAA production. The amount of IAA produced was determined calorimetrically at 540 nm and calculated based on standard graph prepared from authentic IAA

Effects of precursor (l-tryptophan) and carbon, nitrogen sources on IAA production

Effect of nutritional factors on IAA production, such as l-tryptophan, carbon and nitrogen sources was investigated. The bacteria was cultivated in nutrient broth supplemented with 0.1 % tryptophan or 1% different carbon sources like Glucose, Galactose, Lactose, Mannitol, Sucrose and 0.5% different nitrogen sources such Soya bean, beef extract, Yeast extract, peptone and tryptone. After 24 hours cultivation at 37°C and shaking 200 rpm, the amount produced IAA by TIB6 strain was determined.

Screening main parameters affecting on IAA production by Plackett-Burman design

Plackett-Burman design (PBD) is the most commonly used screening design due to its ability to estimate main effects with the same precision (Antony, 2008). It is a fractional factorial design with the advantage of minimizing the experimental runs from large number of variables to smaller most significant factors (El-Refai *et al.*, 2010; Fang *et al.*, 2010). In this design, N factors can be screened with N + 1 trial runs (Fang *et al.*, 2010). In this study, the PBD were applied to evaluate the effect of 10 cultural factors which are considered to be important for IAA production. Each component was tested at two concentration levels, low and high. Table 1 shows the factors used for PBD with 12 experiments for assessing their effects on IAA production. All experiments were carried out in 250 ml Erlenmeyer flasks containing 100 ml media, shaking at 200 rpm.

Table 1. Coded and actual values of the variables used in PBD

Variables	Unit	Low level (-1)	High level (+1)
X1-Sucrose	%	0.1	2
X2-Yeast extract	%	0.1	1
X3-L-tryptophan	%	0.01	0.1
X4-KH ₂ PO ₄	%	0.01	0.05
X5-MgSO ₄ .7H ₂ O	%	0.01	0.05
X6-NaCl	%	0.1	1
X7-Temperature	°C	20	40
X8-pH	-	5	8
X9-Incubation period	Days	2	10
X10-Inoculation ratio	%	1	3

The PBD based on the first order model without interaction effect has been represented by Equation 1.

$$Y = \beta_0 + \beta_i X_i \tag{1}$$

Where Y is the response function, β₀ is the model intercept, i is the variable number, β_i is regression coefficient and X_i represents the independent variables. A 12-experimental run PBD consisting of the 10 factors at 2 levels was designed and the trials were carried out with triplicates. The standard design matrix with the responses is presented in Table 2.

The main effect of each of the variable is given in Equation 2.

$$E_{(x_i)} = (\sum M_{i+} - \sum M_{i-})/n \tag{2}$$

Where E_(x_i) is the effect of the variables, ∑M_{i+} is the summation of the response at high level, ∑M_{i-} is the summation of the response at low level, n is the number of trials (Ndaliman, 2013). When the effect value is positive, the influence of the variable on IAA production is greater at higher concentration, conversely, if the effect value is negative, the influence of the variable is greater at lower concentration. The statistical analysis of the results was performed with the aid of statistical software package Design Expert 7.0 (Sasirekha *et al.*, 2012).

Table 2. PBD with the responses for IAA production

Exp. No.	X1	X2	X3	X4	X5	X6	X7	X8	X9	X10	IAA (mg/L)
1	+	-	+	-	-	-	+	+	+	-	91.91
2	+	+	-	+	-	-	-	+	+	+	105.29
3	-	+	+	-	+	-	-	-	+	+	118.43
4	+	-	+	+	-	+	-	-	-	+	111.82
5	+	+	-	+	+	-	+	-	-	-	93.73
6	+	+	+	-	+	+	-	+	-	-	110.51
7	-	+	+	+	-	+	+	-	+	-	101.19
8	-	-	+	+	+	-	+	+	-	+	105.26
9	-	-	-	+	+	+	-	+	+	-	79.98
10	+	-	-	-	+	+	+	-	+	+	81.81
11	-	+	-	-	-	+	+	+	-	+	87.23
12	-	-	-	-	-	-	-	-	-	-	74.94

Optimization for IAA production using Response Surface Methodology

In order to determine the optimal level of each important independent variable, a central composite design (CCD) of response surface method had been used, which allows to estimate a full quadratic model for each response (Adinarayana and Ellaiah, 2002; Beg *et al.*, 2002; Kristo *et al.*,

2003; Chang *et al.*, 2006). The CCD in 20 runs was adopted (Table 4) for the experiment containing 3 key effect factors at 5 levels each (Table 3). The relationship of the independent variables and the responses was represented in the second-order polynomial equation 4.

$$Y = \beta_0 + \beta_1X_1 + \beta_2X_2 + \beta_3X_3 + \beta_{11}X_1X_1 + \beta_{12}X_1X_2 + \beta_{13}X_1X_3 + \beta_{22}X_2X_2 + \beta_{23}X_2X_3 + \beta_{33}X_3X_3 \quad (4)$$

Where Y is the response; β_0 is the intercept; $\beta_1, \beta_2, \beta_3$ are linear coefficients; $\beta_{11}, \beta_{22}, \beta_{33}$ are squared coefficients; $\beta_{12}, \beta_{13}, \beta_{23}$ are interaction coefficients. The analysis of variance (ANOVA) was used to value the effects and determine regression coefficients of model. The response surface plots were adopted to represent the effect of the independent variables on IAA production. These response curves were then used to predict the optimum level of the factors (Pham *et al.*, 1998; Kwak *et al.*, 2006).

Table 3. Experimental range of the three significant variables used in Central composite design for IAA production

Independent variables	Levels				
	$-\alpha$	-1	0	+1	$+\alpha$
A: L-tryptophan (%)	0.046	0.06	0.08	0.1	0.114
B: Yeast extract (%)	0.33	0.5	0.75	0.1	1.17
C: Inoculation period (%)	0.66	1.0	1.5	2.0	2.34

Table 4. Experimental design and responses of IAA production in Central composite design

Observations	A	B	C	IAA production (mg/L)
1	-1	-1	-1	102.13
2	1	-1	-1	111.41
3	-1	1	-1	109.52
4	1	1	-1	115.63
5	-1	-1	1	109.34
6	1	-1	1	114.75
7	-1	1	1	112.91
8	1	1	1	114.88
9	$-\alpha$	0	0	104.24
10	$+\alpha$	0	0	116.45
11	0	$-\alpha$	0	108.92
12	0	$+\alpha$	0	116.89
13	0	0	$-\alpha$	111.12
14	0	0	$+\alpha$	118.21
15	0	0	0	120.31
16	0	0	0	118.24
17	0	0	0	118.72
18	0	0	0	120.34
19	0	0	0	119.67
20	0	0	0	120.53

RESULTS AND DISCUSSION

Effect of L-tryptophan on IAA production

Obtained results showed that l-tryptophan stimulated IAA production of *B. subtilis* TIB6. TIB6 produced 76.4 ± 2.1 mg/L IAA when 0.1% tryptophan was added into culture medium compared with 28.5 ± 2.8 mg/L IAA in control (without tryptophan). It means that strain *B. subtilis* TIB6 probably synthesized IAA through l-tryptophan dependent pathways. Tryptophan is believed to be the primary precursor for the formation of IAA in plants and rhizobacterium (Monteiro *et al.*, 1988).

The role of tryptophan in the enhancement of IAA production also reported in several earlier reports (Chopade *et al.*, 2008; Khalid *et al.*, 2004). Maximum IAA production of 170 μ g/ml was obtained when 0.5% l-tryptophan was used for culturing *Pseudomonas* sp. (Balaji *et al.*, 2012), while IAA production of *Klebsiella* increased with an increasing of tryptophan up to 0.2%. Similarly, strains *Pseudomonas aeruginosa* produced IAA at 5 fold higher than control (without tryptophan) when culture medium was supplemented with 0.1 g/l-tryptophan (Chaiarn and Lumyong, 2010). IAA synthesis of many bacteria was improved when tryptophan was added into culture medium (El- Khawas and Adachi, 1999; Ali *et al.*, 2010). However, excessive concentration of l-tryptophan may also have adverse effect on IAA production (Nita *et al.*, 2011). This phenomenon may be due to release of IAA degrading enzymes as reported earlier for *Rhizobium* species (Datta and Basu, 2000).

Effect of carbon sources on IAA production

Strain TIB6 can use all investigated carbon sources for growth and IAA synthesis. In culture medium with glucose and sucrose (1%), TIB6 synthesized 76.3 ± 2.7 and 78.3 ± 3.2 mg/L IAA, respectively. Meanwhile, in medium supplemented with other carbon sources, TIB6 produced less amount of IAA (Table 5). The studies regarding correlation between utilization different carbon sources and production of IAA of *Rhizobium* showed that different *Rhizobium* strains prefer different carbon source. The *Rhizobium* strains 12, 16 and 18 required sucrose, whereas *Rhizobium* strain 13 and *Rhizobium* sp. from Cajanus Cajan preferred glucose for maximum production of IAA (Datta and Basu, 2000; Sridevi and Mallaiah, 2007). Sucrose was also the best carbon source for the growth and IAA production of *Acetobacter diazotrophicus* and *Pantoea agglomerans* (Apine and Jadhav, 2011; Nita *et al.*, 2011).

Table 5. Effect of carbon source on IAA production of TIB6

Carbon source	Glucose	Sucrose	Mannitol	Galactose	Lactose
IAA (mg/L)	76.3 ± 2.7	78.3 ± 3.2	63.5 ± 2.3	68.7 ± 2.3	71.2 ± 2.9

\pm standard error.

Effect of nitrogen sources on IAA production

Effect of nitrogen source on IAA production of TIB6 strain was assessed and experimental results pointed out that IAA production reached a peak of 79.1 ± 3.4 μ g/ml when yeast extract was used as nitrogen source. Like carbon sources, bacteria utilize dissimilar nitrogen sources for maximum IAA production. *Rhizobium* can utilize both organic and inorganic nitrogen sources (Sridevi and Mallaiah, 2007), while *Pseudomonas* sp. prefers yeast extract (Balaji, 2011).

Table 6. Effect of nitrogen source on IAA production of TIB6

Nitrogen source	Soya bean	Beef extract	Yeast extract	Peptone	Tryptone
IAA (mg/L)	65.7 ± 2.6	70.1 ± 2.9	79.1 ± 3.4	75.9 ± 3.2	63.9 ± 3.1

\pm standard error.

Beef extract was the most favorite nitrogen source for *Pantoea agglomerans* PVM (Apine and Jadhav, 2011), and NH_4Cl was found to be the most suitable nitrogen source for IAA production of *Acetobacter diazotrophicus* L1 (Nita et al., 2011).

Screening of main factors effecting IAA production by Plackett-Burman design

PBD experiments showed that IAA production by TIB6 varied widely, ranging from 74.94 to 118.43 $\mu\text{g/ml}$. Sucrose, L-tryptophan, yeast extract, KH_2PO_4 , MgSO_4 concentration, and inoculation ratio showed positive effects, while NaCl, temperature, pH and incubation period have shown negative effects on IAA production. The Student's t-test and p-value were used to determine the significance of model at 95% confidence level. On the basis of the Student's t-test and the calculated p-values, l-tryptophan, yeast extract concentration and inoculation ratio were identified as the three significant components affecting IAA production. Based on the PB design, the effect of independent variables on IAA production is given by the first order linear model as equation 5 (actual units). The regression coefficient of the model ($R^2 = 0.9993$) close to 1.0, implying that the given model is well fitted with the trial results. Analysis of variance (ANOVA) given in table 8 showed that the main effect of the independent variables was valid with 93.7% confidence level.

$$\text{IAA} = 72.3 + 2.46X_1 + 13.1X_2 + 215X_3 + 123X_4 + 72.3X_5 - 3.15X_6 - 0.332X_7 - 0.097X_8 - 0.102X_9 + 4.8X_{10} \quad (5)$$

Table 7. Estimated effects and coefficients for analysis of PBD

Variables	Main effect	Coefficients	t-value	p-value
Constant		96.8417	283.44	0.002
X1	4.6733	2.3367	6.84	0.092
X2	11.7767	5.8883	17.23	0.037
X3	19.3567	9.6783	28.33	0.022
X4	5.4067	2.7033	7.91	0.080
X5	2.8900	1.4450	4.23	0.148
X6	-2.8367	-1.4183	-4.15	0.150
X7	-6.6400	-3.3200	-9.72	0.065
X8	-0.2900	-0.1450	-0.42	0.744
X9	-0.8133	-0.4067	-1.19	0.445
X10	9.5967	4.7983	14.04	0.045

Table 8. ANOVA for linear model for effect of independent variables on IAA production

Source	DF	Sum of squares	Mean square	F-value	p-value
Main effects	10	2153.32	215.33	153.72	0.063
Residual error	1	1.40	1.40		
Total sum of squares	11	2154.72			

Plackett-Burman design can be used to find the significant variables in a system and allow them to be ranked in order of importance and to decide which one is to be investigated further so as to determine the optimum values (Liu et al., 2003). Sasirekha et al. (2012) used Plackett-Burman design for screening the most significant effects influencing IAA production by *Pseudomonas aeruginosa*, and yeast extract, tryptophan and EDTA were identified as significant components.

Optimization for IAA production using response surface methodology

CCD has been adopted in determination of optimum cultural conditions and in the analysis of the interaction of three main variables for IAA production by bacteria. All the trials were carried out with triplicates and the average IAA production given in Table 4 was subjected to multiple linear regression analysis using Design-Expert 7.0.0 software. The effect of l-tryptophan, yeast extract concentration and inoculation ratio on IAA production was described in the form of second order polynomial model in coded units.

$$\text{IAA (mg/L)} = 119.66 + 3.17A + 2.10B + 1.84C - 0.83AB - 1.00AC - 0.99BC - 3.47A^2 - 2.57B^2 - 1.95C^2 \quad (6)$$

Where Y is IAA production (mg/L), A is l-tryptophan concentration (%), B is yeast extract concentration (%) and C is inoculation ratio (%). The analysis of variance (ANOVA) for second order polynomial model was given in Table 9. The statistical significance of the second-order polynomial model was assessed by F-test ANOVA. The model F-value of 57.03 implied the model was significant. P-value less than 0.05 will indicate that the model terms are significant at $\alpha=0.05$. In this case, the model had a very low probability value (p-value < 0.0001 < $\alpha=0.05$), implying that the model was highly significant and adequate to represent the relationship between IAA production ($\mu\text{g/ml}$) and l-tryptophan, yeast extract concentration and inoculation ratio. Lack-of-fit of model had a high probability value (p-value = 0.385 > $\alpha=0.05$), which showed that lack of fit is insignificant. Similarly, the lack of fit F-value of 1.32 implied the lack of fit was not significant relative to the pure error. Non-significant lack of fit demonstrated that the model was well fit.

The R^2 value of model represents the correlation between observed and predicted values. In this study, $R^2 = 0.9809\%$ showed good correlation between observed and predicted values and proved that model could explain 98.09% of variability in the response and only 1.91% of the total variance could not be explained by the model. The Pred R-squared of 0.9058 was in reasonable agreement with the Adj R-squared of 0.9637. Adeq Precision measures the signal to noise ratio and a ratio greater than 4 is desirable. In this case, a ratio of 24.707 indicated an adequate signal. The coefficient of variation (CV) indicates the degree of precision with which the experiments are compared. A lower value of coefficient of variation shows the experiments conducted are precise and reliable (Box, 1978). In the present case, a low CV (0.9%) pointed out that the experiments performed are reliable.

Table 9. Analysis of variance (ANOVA) for second order polynomial model of IAA production

Factors	DF	Sum of Squares	Mean Square	F-value	P-value
Model	9	539.6666	59.96295	57.02756	< 0.0001
Lack of fit	5	5.976582	1.195316	1.316964	0.3850
Pure error	5	4.53815	0.90763		
Total	19	550.1813			

$CV = 0.9\%$; $R^2 = 0.9809$; $Pred-R^2 = 0.9058$; $Adj-R^2 = 0.9637$; $Adeq Precision = 24.707$

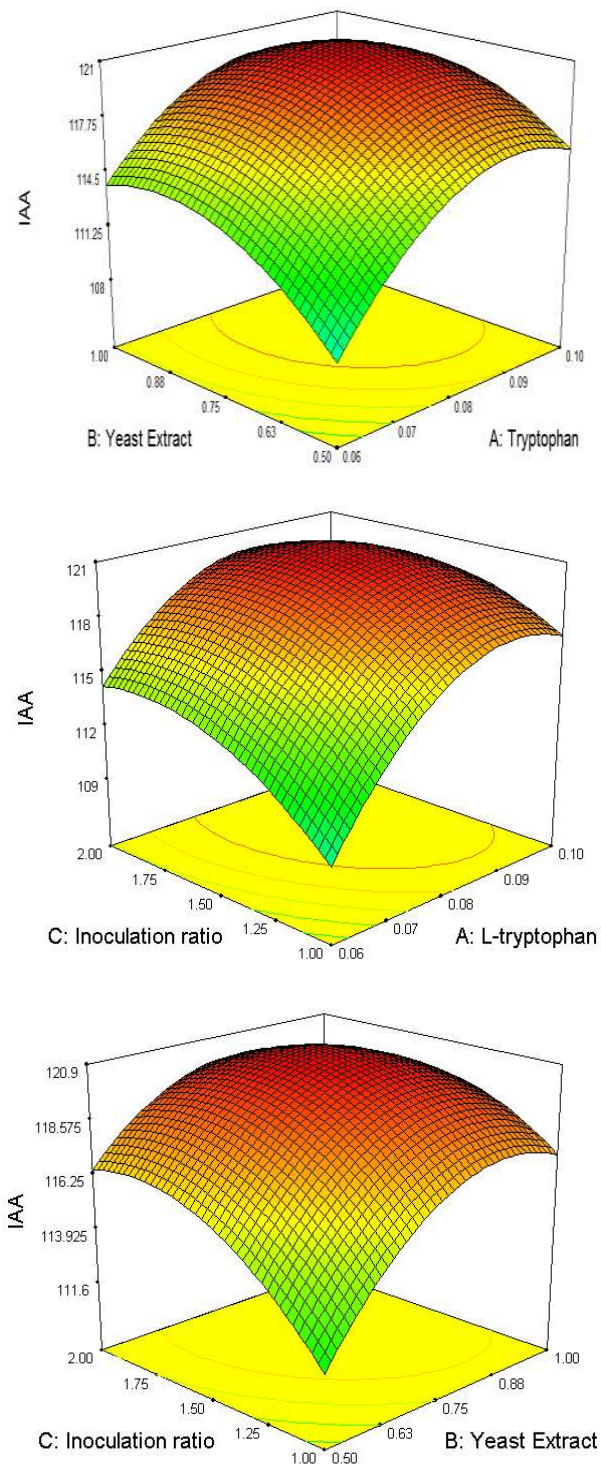


Fig. 1. Optimization for IAA production using response surface method in 3D surface plot

The obtained 3D surface plots represent the effect of each of the three factors and their interactive influence on IAA production. The surface plots (Figure.1) showed that lower and higher level of L-tryptophan, yeast extract and inoculation ratio had less significant effect on the IAA production and increasing of IAA production was observed at middle level of L-tryptophan, yeast extract, as well as inoculation ratio. Results of optimization for culture conditions of IAA production based on response surface method indicated that predicted maximum IAA production of 120.846 (mg/L) was obtained using L-tryptophan concentration of 0.088%, yeast extract

concentration of 0.82% and inoculation ratio of 1.65%. The comparison of predicted (120.846 mg/L) and experimental values (121.35 mg/l) showed good correlation between them, implying that the empirical model derived from RSM can be used to adequately describe the relationship between the cultural conditions and the IAA production. Response surface method has already been successfully adopted for optimization of media and culture conditions in many cultivation process for the production of primary and secondary metabolites (Shirai *et al.*, 2001; Boyaci, 2005), such as amino acids (Jyohi *et al.*, 2005; Xiong *et al.*, 2005), ethanol (Carvalho *et al.*, 2003), flavoring compound (acetoin) (Xian *et al.*, 2007), enzyme (Rao *et al.*, 1993), IAA (Swain and Ray, 2008), and biomass (Sreekumar and Soundarajan, 2010). Swain and Ray (2008) applied RSM-CCD for optimization of IAA production in SSF, and showed that optimum parameters for IAA production were incubation period (6 days), pH (7.0) and moisture holding capacity (70%).

Conclusion

The influence of carbon, nitrogen sources and tryptophan supplementation on IAA production of the TIB6 strain was studied and the experimental results revealed that sucrose and yeast extract were the most favorable carbon and nitrogen sources for TIB6's IAA producing capability, respectively. Besides, the presence of L-tryptophan in culture medium increased amount of IAA produced by TIB6 strain. Plackett-Burman design was used for determination of main culturing factors affecting IAA production by *B.subtilis* TIB6 and obtained results showed that L-tryptophan, yeast extract concentration and inoculation ratio were parameters exhibited significant effect on IAA production. These variables were further optimized using RSM with CCD design and the comparison of predicted and experimental values of IAA production showed good correlation between them. The optimum culture parameters for maximum IAA production (121.35 mg/L) of TIB6 were L-tryptophan (0.088%), yeast extract (0.82%) and inoculation ratio (1.65%).

Acknowledgement

This work was financially supported by Tay Nguyen 3rd Program of Technological and Scientific Development, Vietnam Academy of Science & Technology for the project: *Improving and technological transfer of POLYFA-TN3 product for soil reclamation in Tay Nguyen*. N^o Code: TN3/C10.

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