

Original Research Article

Optimization of Polyhydroxybutyrate (PHB) production by *Azotobacter vinelandii* using experimental design

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A B S T R A C T

Keywords

Azotobacter vinelandii;
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In this study, thirty local isolates of *Azotobacter* spp. were isolated from Egyptian soil and screened for PHB production. Plackett-Burman and Box-Behnken optimization methods have been used to optimize PHB production from the best locally isolated producer. The most producer isolate was identified to the molecular level using 16S rRNA as *Azotobacter vinelandii*. Nine variables represent the medium constituents and the environmental conditions were randomized. The accumulation of PHB granules in cells of *A. vinelandii* significantly depended on the ratio of C- and N- sources in the culture medium, minerals, temperature and aeration. According to the main effect analysis, all the above variables have a positive effect on PHB production. The most effective and promising variables included shaking rate with confidence 90.04%, followed by FeSO₄, sucrose and NaNO₃ with confidence levels 81.16, 78.88 and 77.27%, respectively. The maximum yield of PHB was 0.88 gm/gm dry cells. ANOVA analysis of the regression model showed significant interaction between the various variables that reflect the powerful of Plackett-Burman and Box-Behnken optimization methods. The present data indicated that strategies included in this study could be used for PHB production by this bacterium with high PHB content and adequate properties of the biopolymer

Introduction

Azotobacter vinelandii is an obligate aerobic bacterium that able to fix

atmospheric nitrogen and grow under microaerophilic conditions. This

microorganism is able to synthesize three molecules of important biotechnological and biochemical applications; the extracellular polysaccharide alginate, siderophores compounds and polyhydroxybutyrate (PHB). The latter is biodegradable thermoplastic polyester analogous or better than those of chemically synthesized and petroleum-based polymers such as polyethylene and polypropylene (Byrom, 1987; Doi, 1990). PHB is a biopolymer that has been implicated in supporting nitrogen fixation (Segura *et al.*, 2003), biodegradable thermoplastic material for management including plastics, films, and fiber strategies, and biocompatibility in the medical devices (Steinbüchel, 1995; Gouda *et al.*, 2001; Lenz and Marchessault, 2005). This polymer is accumulated due to depletion of nitrogen, phosphorous or oxygen to form carbon and energy reserve material (Anderson and Dawes, 1990; Aldor and Keasling, 2003; Burns *et al.*, 2007; Galindo *et al.*, 2007; Khanna and Srivastava, 2005; Suzuki *et al.*, 2008).

Production of PHB has been reported using microorganisms such as yeast (Lee *et al.*, 1997), *Alcaligenes latus* (Grothe and Chistri, 2000), *A. eutrophus* (Song *et al.*, 2001), *Pseudomonas* species (Ashby *et al.*, 2002), wide range of bacterial and archaeal species (Aldor and Keasling, 2003), *Rhodobacter sphaeroides* (Chen *et al.*, 2006), *Ralstonia eutropha* (Verlinden *et al.*, 2007) *Bacillus subtilis* and *Bacillus megaterium* (Nur Yüksekdağ *et al.*, 2004; Chaijamrus and Udpuay, 2008), recombinant *Escherichia coli* (Yu *et al.*, 2002; Khanna and Srivastava, 2005; Li *et al.*, 2007) and *Azotobacter* species (Page and Knosp, 1989; Segura *et al.*, 2003; Kishk, 2009; Diaz-Barrera and Soto, 2010)

The microbial large scale production of PHB is dependent on the development of a low cost process that produces biodegradable plastics with properties similar or superior to petrochemical plastics (Doi and Steinbüchel, 2001; Sims, 2003; Apostolis *et al.*, 2006). Various nitrogen-rich media containing casein hydrolysate, yeast extract, tryptone, casamino acids, corn steep liquor and collagen hydrolysate (Lee and Chang, 1995; Bormann *et al.*, 1998; Ghaly, 2003; Khanna and Srivastava, 2005; Chaijamrus and Udpuay, 2008) have been used for PHB production. Carbon-substrate rich media such as molasses, whey, hemicelluloses, palm oil, starch, glucose, fructose, sucrose, maltose, gluconate or glycerol accumulates PHB in bacterial cells (Page, 1989; Page and Knosp, 1989; Quagliano *et al.*, 1994; Alias and Tan, 2005; Chaijamrus and Udpuay, 2008; Kishk, 2009) and have been used as substrates for PHB production. However, unrefined carbon sources such as corn syrup, cane molasses, beet molasses, or malt extract, also supported PHB formation, obtaining yields of PHB comparable to, even better than the refined sugars.

Production of PHB by fermentation has been normally operated as a two stage fermentation fed-batch processes, an initial growth phase followed by a PHB formation stage (Chen and Page, 1997). Among factors influencing production costs is the recovery of the product. In this context, Page and Cornish (1993) reported that beside the substrate cost, the PHA extraction method from inside the cells, and the treatment of the fermentation wastes are cost effective factors. The substrate cost affects the overall cost but the cheapest substrate is not always the ideal choice concerning the downstream

processes. When the PHB productivity increased from 1.98 to 3.2 g/h, the PHB production cost decreased from \$5.37/kg to \$4.91/kg (Lee and Choi, 1998). Aiming to wide range of industrial applications and establishing method for changing PHAs compositions as well as PHAs' overproduction, many researchers have been conducted molecular biology and protein engineering of PhaC synthases, *phbA*, *phbC* and *phbB* genes (Amara *et al.*, 2001; 2002; Taguchi, 2001, 2002; Segura *et al.*, 2003). The optimization method of Plackett-Burman, that still one of the most promising approaches was recommended by Page and Cornish (1993). Recently, Diaz-Barrera and Soto (2010) proposed new experimental strategies based on integration between molecular and bioengineering techniques to enhance the productivity of PHB from *A. vinelandii*.

This study aimed to evaluate the potentiality of *Azotobacter vinelandii*, isolated from soil samples collected from Tanta, Gharbia Governorate, Egypt for PHB production. The advantages of using *A. vinelandii* in PHB production is the easy attainment of this bacterium from soil and fresh water. The most economic parameters, which gave the highest amount of the biopolymer PHB, were undertaken using Plackett-Burman and Box-Behnken designs, in addition to statistical analyses.

Materials and Methods

Microorganisms used

Thirty isolates belonging to the genus *Azotobacter* were isolated previously from cultivated and non-cultivated soil samples, collected from different localities at Tanta, Gharbia Governorate, Egypt. The isolates were purified on nitrogen free medium described by Vancura and Mancura

(1960). The pure isolates were characterized and the most PHB promising *Azotobacter vinelandii* has been used in this study and identified according to the methods described in Tchan (1984) and by sequencing its 16S rDNA genes and comparing the sequences with a data base library using the BLASTN (Altshul *et al.*, 1997) program, a subprogram of Basic Local Alignment Search Tool (BLAST), through the network service of the National Center for Biotechnology Information (NCBI) against nonredundant sequences. The sequences obtained were aligned with *A. vinelandii* 16S rRNA sequences available in the GenBank using the EasyAlign (Miranda, 2002) program that employs the MALIGN algorithm (Wheeler and Gladstein, 1998)

Medium and environmental conditions used

The following enrichment N₂-Free medium after Vancura and Mancura (1960) with modification was used for PHB production. It contained the following components as g/l: Sucrose 20; K₂HPO₄ 0.65; KH₂PO₄ 0.16; NaCl 0.2; MgSO₄·7 H₂O 0.2; CaCO₃ 2; FeSO₄ 0.005; NaBO₄ 0.005; and bacto agar (Law and Slepecky, 1961). The medium constituents and the environmental conditions have been randomized according to Plackett-Burman (1946) design (+1, -1); temperature (37, 30) °C, pH (7.5), sucrose (30, 5) g/l, NaNO₃ (2, 0.1) g/l, KH₂PO₄ (1, 0) g/l, NaCl (0.5, 0) g/l, MgSO₄ (0.5, 0) g/l, FeSO₄ (0.005, 0) g/l, shaking (250, 200) rpm.

Experimental designs

Plackett-Burman

Eighteen experiments containing +1 and -1 values following Plackett-Burman

design have been conducted as in table 1 to optimize ten variables represented at high and low levels, which are denoted by +1 and -1. All the eighteen experiments were conducted using 250 ml Erlenmeyer flasks containing 100 ml media. The media and environmental conditions have been randomized using Plackett-Burman design (+1,-1). The mean of +1 experiment has been calculated using the following formula: $(\sum +1)/n_{(+1)}$. The mean of -1 experiment has been calculated using the following formula: $(\sum -1)/n_{(-1)}$. The main effect of each factor has been calculated from the following formula:

$$\text{Main effect} = \sum (+1)/n_{(+1)} - \sum (-1)/n_{(-1)}$$

The main effect show whether the variable(s) amount is effective positively or negatively on PHB production.

The results of the Plackett-Burman design experiments were applied to linear multiple regression analysis using Microsoft Excel 2002. The statistical analysis of the experimental results in Table 1 has been summarized in Table 3. The variables whose confidence levels % were \geq than 90% were considered to be significantly affected on the PHB. Variables with confidence level% less than 90% till 70% were considered effective (Stowe and Mayer, 1966).

Generating 1st order Model

The model created from the analysis of Plackett-Burman experimental design using multiple regression analysis is based on the 1st order-model $Y = \beta_0 + \sum \beta_i X_i$, Where Y is the predicted response, β_0 model intercept, β_i variables linear coefficient. ANOVA test was generated for each response to determine the

relationship between the variables at the 90% or higher confidence level.

Extraction, purification and determination of PHB

The assay was performed spectrophotometrically (PerkinElmer-UV/VIS Spectrometer Lambda) to determine PHB as crotonic acid. The PHB amount was determined by calculating the amount of crotonic acid derived from the standard curve and calculated as gm PHB/gm cells. *Azotobacter* cultures were grown and inoculated with single colony using tooth pick to 100 ml of modified liquid medium of Vancura and Mancura (1960) in 250 ml Erlenmeyer flasks and incubated in an orbital shaker at different shaking rates for 6 days of incubation. Bacterial cells were separated with centrifugation at 4000 rpm for 15 min and were dried at 40°C for 24 h. The dry weight of the pellets was determined. Bacterial cell walls were lysed by adding sodium hypochlorite, mixing and incubating at 60°C for 1 h. Supernatant was obtained by centrifugation and transferred to a Soxhlet system. Cell lipids and other molecules were treated with 5 ml 96% ethanol and acetone. PHB was extracted by hot chloroform (adding 10 ml chloroform in a water bath). Then chloroform was evaporated to obtain PHB crystals. By adding 10 ml of 98% sulfuric acid at 60°C for 1 h, PHB crystals were converted into crotonic acid. The absorbance of the solution was measured at 235 nm in a UV spectrophotometer (Perkin-Elmer Lambda EZ201-UVvis) against sulfuric acid as blank. The amounts of PHB per gram dry weight of bacterial cells were determined using a standard curve of PHB according to Kuniko *et al.*, (1988).

Result and Discussion

Taxonomic classification and identification of *A. vinelandii*

The morphological, physiological and biochemical characters of the most PHB active isolate (data not shown) were found to be similar with those cited in Bergey's Manual for Systematic Bacteriology for the genus *Azotobacter*. The isolate was also subjected to molecular identification and its DNA was isolated as pure unique band.

The sample of 16S ribosomal DNA contained approximately 1500 base pairs in length (Fig. 1) that was amplified by PCR, and partially sequenced as the following:

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CGCGGTAATACGAAGGGTGCAAGT
AATCGGAATTACTGGGCGTAAACGC
GCTAGGTGGTTCGGCAAGTTGGATG
TGAAAGCCCCGGG CTACCTGGGAAC-
CGCATCCAAA ACT ACTGGGCTAG
A G T A C G G T A G A
GGGTGGTGAATTTCTGTGTAGCG
GTGAAATGCGTAGATATAGGAAGGAACA
CC GTGGCGAAGGCGGCC ACCTGGACC
GATACTGAC ACTGAGGTGCGAAAG
CGTG GGA GCAA CAGGATTAGAT
ACCCTGGTAGTCCA CAGGACCC TG-
GTAGTCCA CGCCGTAAA CGATGT
C G A C T A G C C G T T G G G
CTCCTTGAGAGCTTAGTGGCGCAGC
T AACGCATTAAG TCGAACCGC-
C T G G G G A G T A C G G C C
GCAAGGTTAAA ACTCAATGAATTGA
CGGGGGCCCG ACAAGCGGTGGAGCT-
GTG GTTTAATTTCGAAGCCAACGCGA
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Sequence was then compared to the public database of National Center for Biotechnology Information (NCBI) using

Basic Local Alignment Search Tool (BLAST). Analysis of the sequences data revealed that query sequence showed greatest homology with *A. vinelandii* (95%). Thus identification of the selected and most potent isolate was confirmed to be *A. vinelandii*.

Variables affecting PHB production by *A. vinelandii*

Increasing the productivity of PHB using Plackett and Burman design

The results obtained in Table 1 indicated that there was a wide variation in PHB production from (0 to 0.88 PHB g/g D. wt.), using Plackett and Burman experiments that highlighted the importance of optimizing culture variables in attaining higher PHB production rate. The experiment showed the maximum PHB production rate was at the following conditions: temperature at 37°C, pH 7, sucrose 30 g/l, NaNO₃ 2 g/l, KH₂PO₄ 1 g/l, NaCl 0 g/l, MgSO₄ 0.5 g/l, FeSO₄ 0.005 g/l, CaCl₂ 0.06 g/l, NaMoO₄ 0.05 and shaking rate at 250 rpm. From the results obtained in Table 2 by regression analysis of Plackett and Burman design, it was found that non of the variables has a p=0.05 significant effect on the PHB production which indicate the need for conducting a further optimization. The R-Squared statistic indicates that the model as fitted explains 60.6577% of the variability in PHB. The adjusted R-squared statistic is 16.3977%. The standard error of the estimate shows the standard deviation of the residuals to be 0.229216. The mean absolute error of 0.116353 is the average value of the residuals. The Durbin-Watson statistic tests the residuals to determine if there is any significant correlation based on the order in which they occur in your data file. Since the P-value is greater than 0.05, there is no indication of serial autocorrelation in the residuals at the 95.0% confidence level.

Table.1 Plackett-Burman design for PHB optimization

Experiment No.	Temperature	pH	sucrose	NaNO ₃	KH ₂ PO ₄	NaCl	MgSO ₄	FeSO ₄ **	NaMoO ₄ **	Shaking	PHB gm/gm cells	
Plackett-Burman Experiments	1	-1 (30)	1(7)	1(30)	1(2)	1(1)	1(0.5)	1(0.5)	1(0.005)	1(0.05)	-1(200)	0.199
	2	-1(30)	-1(5)	1(30)	-1(0.1)	1(1)	1(0.5)	1(0.5)	-1(0)	-1(0)	1 (250)	0.000
	3	-1(30)	1(7)	-1(5)	1(2)	-1(0)	1(0.5)	1(0.5)	1(0.005)	1(0.05)	-1 (200)	0.047
	4	1(37)	1(7)	1(30)	1(2)	1(1)	-1(0)	1(0.5)	1(0.005)	1(0.05)	1 (250)	0.880
	5	1(37)	-1(5)	1(30)	-1(0.1)	1(1)	1(0.5)	-1(0)	1(0.005)	1(0.05)	1 (250)	0.600
	6	1(37)	-1(5)	-1(5)	1(2)	-1(0)	1(0.5)	1(0.5)	-1(0)	-1(0)	1 (250)	0.500
	7	1(37)	-1(5)	-1(5)	1(2)	1(1)	-1(0)	1(0.5)	1(0.005)	1(0.05)	-1 (200)	0.083
	8	-1(30)	-1(5)	-1(5)	-1(0.1)	1(1)	1(0.5)	-1(0)	1(0.005)	1(0.05)	1 (250)	0.042
Plackett-Burman Experiments	9	1(37)	1(7)	-1(5)	-1(0.1)	-1(0)	1(0.5)	1(0.5)	-1(0)	-1(0)	1 (250)	0.000
	10	1(37)	-1(5)	1(30)	-1(0.1)	-1(0)	-1(0)	1(0.5)	1(0.005)	1(0.05)	-1 (200)	0.042
	11	-1(30)	-1(5)	-1(5)	-1(0.1)	-1(0)	-1(0)	-1(0)	1(0.005)	1(0.05)	1 (250)	0.000
	12	-1(30)	1(7)	-1(5)	1(2)	-1(0)	-1(0)	-1(0)	-1(0)	-1(0)	-1 (200)	0.000
	13	-1(30)	1(7)	1(30)	-1(0.1)	1(1)	-1(0)	-1(0)	-1(0)	-1(0)	-1 (200)	0.097
	14	1(37)	1(7)	1(30)	-1(0.1)	-1(0)	1(0.5)	-1(0)	-1(0)	-1(0)	-1 (200)	0.02
	15	-1(30)	-1(5)	1(30)	1(2)	-1(0)	-1(0)	1(0.5)	-1(0)	-1(0)	-1 (200)	0.000
	16	-1(30)	1(7)	-1(5)	1(2)	1(1)	-1(0)	-1(0)	1(0.005)	1(0.05)	-1 (200)	0.050
	17	1(37)	1(7)	1(30)	1(2)	1(1)	1(0.5)	-1(0)	-1(0)	-1(0)	1 (250)	0.060
	18	1(37)	-1(5)	-1(5)	-1(0.1)	-1(0)	-1(0)	-1(0)	-1(0)	-1(0)	1 (250)	0.010

** variables have been used as one unit [one variable] during the regression analysis

Table.2 The main effect of the variables

Variables	Main effect Main effect = $\sum(+1)/n_{(+1)} - \sum(-1)/n_{(-1)}$.
Temperature	1.76
pH	0.076
Sucrose	1.166
NaNO ₃	1.008
KH ₂ PO ₄	1.392
NaCl	0.306
MgSO ₄	0.872
NaMoO ₄ and FeSO ₄	1.256
Shaking rate	1.554

Table.3 Multiple regression analysis of the data in Table 1

Parameters	Estimate	P value	Std Error	t Stat	Confidence %
Constant	0.143	0.027	0.054	2.704	97.312
Temperature	0.068	0.292	0.060	1.129	70.85
pH	0.032	0.643	0.066	0.482	35.72
sucrose	0.091	0.211	0.067	1.359	78.88
NaNO ₃	0.091	0.227	0.069	1.308	77.27
KH ₂ PO	0.000	0.998	0.071	0.003	0.21
NaCl	-0.031	0.631	0.062	-0.500	36.93
MgSO ₄	0.013	0.853	0.066	0.191	14.96
FeSO ₄	0.091	0.188	0.063	1.438	81.16
Shaking rate	0.136	0.100	0.073	1.862	90.04

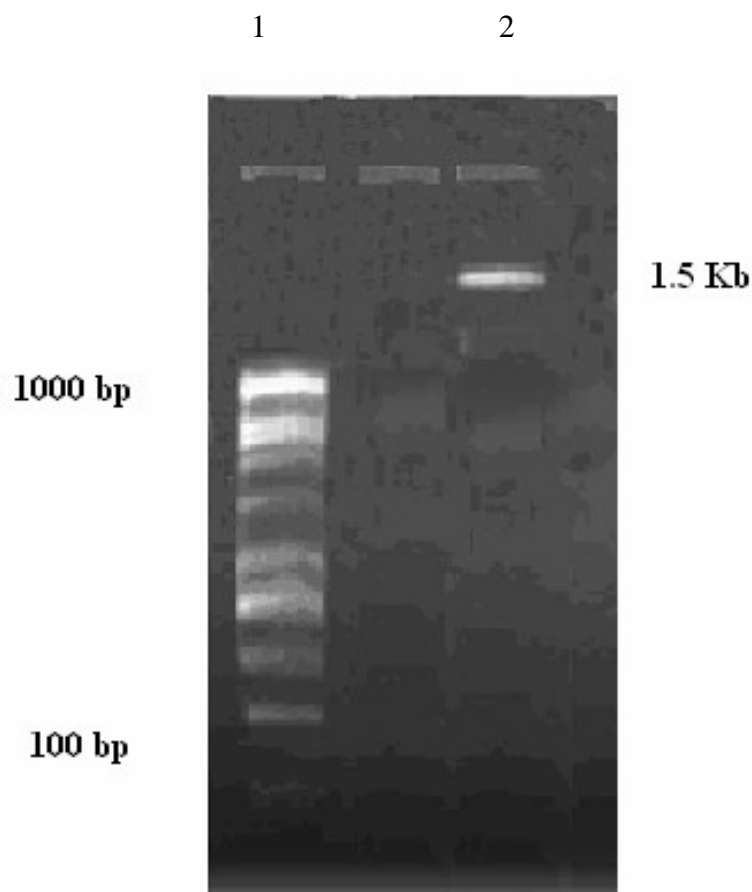
Table.4 Summary of the regression results

Summary	
R-squared	60.65 percent
R ² adjusted	16.39
log 1 residual autocorrelation	0.385
Standard Error	0.229
Durbin-Watson statistic	1.219 (p=0.053)
Mean absolute error	0.116

Table. 5 ANOVA test

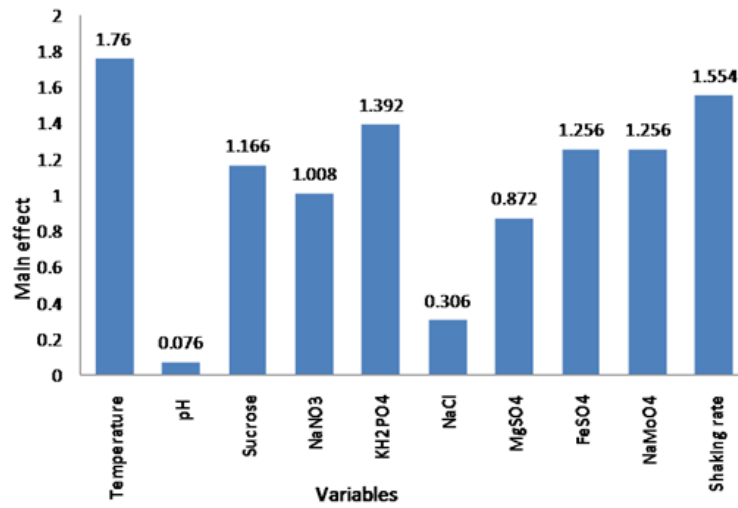
ANOVA					
Source	SS	MS	F	P. Value	df
Regression	0.640	0.072	1.37	0.3338	9
Residual	0.408	0.052			7
Total	1.063				16

Fig.1 Amplification of 16S rDNA PCR product from *A. vinelandii* .



Lane (1) marker DNA 1kb and lane (2) 16s rDNA PCR product with mw (~1.5 kb).

Fig. 2 Main effect o diferent variables used in this study



Optimization of the most significant factors for PHB production using Box-Behnken design

Box-Behnken design was used to optimize the most significant factors affecting PHB production obtained from Plackett and Burman. The results obtained in Table 4 showed the optimum levels of the most significant factors are KH_2PO_4 , pH and agitation rate. The results were analyzed by the second order regression equation provided the levels of PHB production. From the results, it was found that PHB content increased from 0.199 (PHB g/g dry weight) using Plackett and Burman design to 0.7 PHB g/g dry weight using Box-Behnken at the following conditions: sucrose 30, g/l; NaNO_3 , 2 g/l; KH_2PO_4 , 2 g/l; NaCl , 0.5 g/l; MgSO_4 , 0.5 g/l; FeSO_4 , 0.05 g/l; CaCl_2 , 0.06 g/l; NaMoO_4 , 0.005; shaking rate at 225 rpm, pH 7 and 30°C . The most effective variable [data not shown] was the shaking rate with confidence % 88.87, followed by each of FeSO_4 , temperature and sucrose with confidence levels 80.30, 76.41 and 75.84, respectively. Since the P-value in the ANOVA (Table 5) is greater or equal to 0.10, there is no statistically significant relationship between the variables at the 90% or higher confidence level. From the results in Fig. 2, temperature, KH_2PO_4 , NaNO_3 , MgSO_4 , FeSO_4 , shaking, CaCl_2 and NaMoO_4 showed positive effect on PHB production by *A. vinelandii*. Three factors showed negative effect on PHB production represented by NaCl , pH and sucrose. However temperature, NaNO_3 , KH_2PO_4 , FeSO_4 and MgSO_4 are the most effective variables as shown in Fig. 3.

Production of microbial Polyhydroxybutyrate (PHB) is a promising biotechnological area. Optimization the production of biotechnological products is

an aim for many industrial and research sectors (Plackett and Burman, 1946; Amara *et al.*, 2001, 2002; Taguchi *et al.*, 2001, 2002). Basic strategies for economic PHB production based on consideration of good PHB producers and good productivity, as well as PHB recovery affecting overall economics. In spite of great deal of efforts directed towards commercializing PHA, the wide use of PHB remains rare because of its high production cost. One reason for this unfavorable cost for commercial use is that the methods used must select good PHB producers to increase the concentration of PHB, develop the fermentation process, and enhance PHB recovery/purification system. The ideal way to enable PHB to compete with conventional petrochemical plastic is to combine the following methods simultaneously; increase the absolute concentration of PHB per host cell, increase packing of PHB granules into the host, achieve high glucose conversion rate and release the PHB outside the cell to simplify recovery/purification process (Lae *et al.*, 2005).

In this study, Plackett and Burman design was used to investigate significant factors affecting PHB production. Experimental design, as a powerful tool for optimizing different complicated conditions was used to investigate the principal factors for PHB production, and identification of medium components that play a significant role in cell growth and PHB production, as well as their concentrations. Three environmental factors represented in temperature, pH and shaking rate as well as six nutritional factors included sucrose, NaNO_3 , KH_2PO_4 , NaCl , MgSO_4 and FeSO_4 were randomized in eighteen experiments using Plackett-Burman design (1946). The results showed different

responses based on the variation on environmental factors and different nutritional conditions. The variables are heterogeneous as proved by the T-test analysis of variance (ANOVA) where the F significant was > 0.05 (0.3338). This proved the presence of conflict between different factors used in this study. This conflict has been perfectly solved by randomizing the different factors. Plackett-Burman design proved to be a powerful tool for optimizing biotechnological products. However, the study presented here succeeded in producing 88.87% PHB, regarding to the cell dry weight. The variation on the amount of PHB calculated as gm PHB/gm cells proved that Plackett-Burman design is a versatile tool for optimizing complicated conditions. In this context, PHB content in *Azotobacter vinelandii* cells varied from 0.0 to 0.88 g PHB/g dry cell weight (DCW). This mean that the method used herein is successful methods for increasing PHB content. In order to optimize the most significant factors obtained from Plackett and Burman design, Box-Behnken design was applied and maximum PHB production were obtained using 2 g/l KH_2PO_4 , shaking rate at 225 rpm and pH 7. In this respect, Gu *et al.* (2005) reported that response surface methodology provides important information regarding the optimum level of each variable along with its interactions with other variables and their effects on product yield. To validate the exact optimum values of KH_2PO_4 , pH and agitation rate and their interactions, statistical designs were used and the range between the optimum points were selected. Therefore, Box-Behnken design was focused on the interaction between these three factors while other factors were effectively insignificant for PHB production. In this connection, optimization for PHB in *Rhizobium*

meliloti was reported by taking variables such as sucrose, urea, inoculums size and K_2HPO_4 (Lakshman *et al.*, 2004). Similar methods were used by (Khanna and Srivastava, 2005; Lorrungruang *et al.*, 2006). They optimized the concentrations of KH_2PO_4 , Na_2HPO_4 , $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$ and fructose for PHB production in *Wautersia eutropha* using central composite design. Thus, the highest PHB yield (0.88 g PHB/g DCW) using *A. vinelandii* occurred in medium containing NaCl, 0; sucrose, 30 g/l; KH_2PO_4 , 1 g/l; MgSO_4 , 0.5 g/l; FeSO_4 , 0.05 g/l; NaMoO_4 , 0.005 g/l; CaCl_2 , 0.06 g/l; pH 7 and shaking rate at 225 rpm. Thus, the before mentioned constituents and conditions were promising for PHB production. In this connection, Pal *et al.* (1998) reported that PHB accumulation was enhanced when growth was restricted due to unavailability of phosphorus. Under nitrogen deficiency a rise in PHB pool up to 9.5% (w/w of dry cells) was observed. Lugg *et al.* (2008) observed that enterobacteria of natural habitat accumulate 50% PHB of CDW, in medium containing excess carbon source and a deficiency in nitrogen. However, Lee *et al.* (2001) and Li *et al.* (2009) reported that PHB is a NADPH-dependent pathway and increased under high intracellular concentrations of NADPH or high ratio of NADPH/NADP. The possible explanation for the rise in PHB pool under supplementation of glucose, fructose, maltose and ethanol is due to boost in growth. The possible explanation for this rise could be the availability of plenty of precursors i.e., acetate for biosynthesis of PHB, as reported in many bacteria (Zinn *et al.*, 2001; Galino *et al.*, 2007). In some cases, these sugars may be used to enhance cell growth and, in turn, the volumetric productivity as recommended by Quillaguaman *et al.* (2008) for complex nitrogen sources.

The spectrophotometer assay of the polymer that extracted from *A. vinelandii* and the standard (DL- β -hydroxybutyric acid) were analyzed after sulphuric acid digestion and measured at 235 nm which depicted highest degree of similarity with the spectrum of crotonic acid in accordance with Law and Slepecky (1961). Thus, the complete matching of the spectra of the acid-digested sample and the standard with the spectrum of crotonic acid demonstrated the presence of PHB in *A. vinelandii*.

Selecting effective variables is very important in experimental design to reduce the number of the experiments and the number of designs. Not only Plackett-Burman design is able to optimize the nutritional constituents, but also the environmental factors. On the basis of the data obtained in the present study, *A. vinelandii* is capable of PHB accumulation up to 88.89% of dry cell weight. Thus, it can be selected as a candidate for PHB production and commercialization. Molecular studies are needed to increase more PHB content in *Azotobacter* cells. Such studies will be the focus of the nearest future.

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