

Article

Statistical Based Bioprocess Design for Improved Production of Amylase from Halophilic *Bacillus* sp. H7 Isolated from Marine Water

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Abstract: Amylase (EC 3.2.1.1) enzyme has gained tremendous demand in various industries, waste water treatment, bioremediation and nano-biotechnology. This compels the availability of enzyme in greater yields that can be achieved by employing potential laccase producing cultures and statistical optimization. Use of Plackett–Burman design (PBD) that evaluates various medium components and having two level factorial designs help to determine the factor and its level to increase the yield of product. In the present work we are reporting the screening of amylase producing marine bacterial strain identified as *Bacillus* sp. H7 by 16S rRNA. Use of two-stage statistical optimization, i.e. PBD and Response Surface Methodology (RSM) using central composite design (CCD) further improved the production of amylase. A 1.31 fold increase in amylase production was evident at 5.0 L laboratory scale bioreactor. Statistical optimization gives the exact idea of variables that influence the production of enzyme and hence the statistical approach offers the best way to optimize the bioprocess.

Keywords: Amylase; *Bacillus* sp. H7; Optimization; Production; Response surface method

1. Introduction

Currently enzymes have been in huge demand at industrial level due to their eco-friendly, economic advantages catalysis over the chemical in various processing practices [1]. Amylase (EC 3.2.1.1) is starch hydrolyzing enzyme that produces branched and linear oligosaccharides of different chain length. Amylase enzyme has different applications in a wide variety of industries such as textile, food, detergent, paper, sugar industries and pharmaceutical [2-4]. The amylase has covered more than 65% of the world enzyme market [5] and the microbial originated amylase is more in demand due to their wide working range of pH, temperature, broad biochemical diversity, feasibility of mass culture, high enzymatic stability under extreme conditions and ease of genetic manipulation [6-9].

Due to the importance of amylases, isolation of new microbial producers capable of producing amylase provides potential new sources of enzyme [10,11]. Isolates from the extreme environments are considered a rich source for active enzymes that have numer-

ous industrial applications and suitable for harsh condition compared to their counterparts [12]. The comparative analysis of DNA sequences using phylogenetic methods become more significant with the rapid accumulation of molecular sequence detail. Gene sequencing and phylogenetic analysis are helpful to detect the nature and extent of selective forces that shape the evolution of genes and species [13]. For the identification new isolates 16S rRNA gene sequencing has become a most preferable method, because 16S gene is a highly conserved as well as variable in the genomic DNA for each species [14].

Selection of hyper amylase producers and the optimization of a bioprocess are plays the crucial role in the cost management at industrial level [8]. The cell mass and enzymes secretion mainly depends on the physic chemical parameters and their levels. Therefore to increase the production and decrease the production cost, optimization of these parameters is must [15]. The process optimization can be done by One Variable at Time (OVAT) approach but this only has limitation of studying one parameter at time that's why it is tedious, expensive and not suitable for the large numbers of variables [16]. Whereas the statistical approaches like Response Surface methodology (RSM) is facilitates to study the different parameters in combination for the interactive effects on the production and also highlights the significant variables with their optimum level quantifying the relationships between one or more measured responses [17]. Due to the less number of individual experiments the RSM become cost effective [18]. The present study was aimed to ascertain the parameters and the level of individual parameter that enhances the production of amylase at shake flask level and then laboratory scale bioreactor.

2. Results

2.1 Screening and quantitative estimation of amylase production by halophilic isolates

Microorganisms with amylolytic activity were isolated from marine water sample on starch agar medium. Among the morphologically distinct isolates fourteen isolates (H1-H14) showed the good growth in laboratory conditions and so were preliminary selected based on the zone of starch hydrolysis formed on medium. The amylase activity of these isolates was tested by inoculating in APM after 24 h of incubation isolate H1, H2, H4, H5, H6, H10, H11, H13, H14 showed the amylase activity ranging from 37-96 U/mL, while the amylase activity of 108, 120, 102, 110 and 113 U/mL for the isolates H3, H7, H8, H9 and H12 respectively. The highest amylase production was exhibited by isolate H7. As H7 produced highest amylase activity it was selected for further studies.

2.2 Identification of potent isolate

2.2.1. Phenotypic characterization

The colony of isolate H7 was characterized as opaque, milky-white, convex, entire margin while morphologically the isolate H7 was Gram positive, rod shaped, non-motile, non-spore forming bacteria.

2.2.2. Genetic identification of isolate using 16S rRNA Sequencing

The comparison by multiple alignment of 16S rRNA sequence of isolate H7 with the available sequences of gene bank showed the 99.62% similarity with *Bacillus aquimaris* TF-12 (NR_025241) (Figure 1). The isolate was found to belong to *Bacillus* genus and it was identified as *Bacillus* sp. H7. The 16S rRNA gene sequence of the isolate was submitted to gene bank under the name *Bacillus* sp. H7 with Genebank bank accession number MT422535.

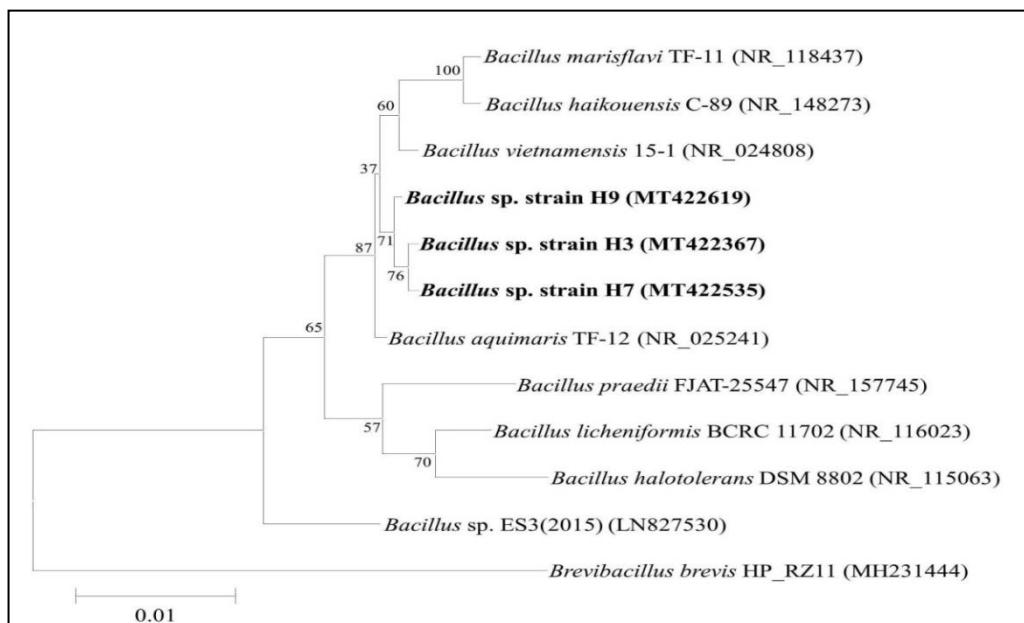


Figure 1 Phylogenetic tree of *Bacillus* sp. H7 drawn using the neighbor-joining method

2.3 Influence of physico-chemical media variables on amylase production

The influence of incubation period on the amylase production revealed the beginning of enzyme production after 6 h in initial lag phase that continued until the late log phase of 96 h. The maximum amylase yield was observed after 37 h (130.53 ± 2.0 U/mL). Good yield of amylase was observed at varying temperature ranging from 20-50 °C, the maximum yield was observed at 35 °C (127.24 ± 2.1 U/mL). The alkaline pH (8.0) of media was found to be the optimum condition for the amylase production. At alkaline pH, the isolate showed the maximum productivity i.e. 133.53 ± 1.2 U/mL. The soluble starch was found as the best carbon source, as the isolate produced maximum (144.64 ± 2.1 U/mL) amylase with starch as a carbon source. Among the nitrogen sources, yeast extract supported the maximum production of amylase (152.96 ± 3.2 U/mL) (Table 1).

2.4. Influence of salt concentrations on growth and amylase production

The effect of salt concentration on growth and amylase production revealed that the organism grows well over the varying concentrations of salt (NaCl). Maximum growth occurred at 1.60 M of NaCl. Amylase production also showed increasing trend with increase in salt concentration. Maximum amylase activity was observed at 1.2 M (154.1 ± 0.8 U/mL). Further increase in the salt concentration did not significantly affect the production of amylase (Figure 2).

2.5 Evaluation of significant production media variable by Plackett–Burman Design (PBD)

Eight Media variables were investigated to determine the optimum medium components suitable for amylase production. The amylase activities from the twelve runs are shown in Table 4. Fractional factorial Plackett–Burman design was used to screen and evaluate the significant variables that can influence enzyme yield because this model does not explain the interaction among various variables [18]. The results (Table 4) indicate a variation in amylase production in the range from 119.17 ± 0.76 to 161.30 ± 0.68 U/mL by *Bacillus* sp. H7. The variations obtained in production revealed the importance of medium optimization to achieve the maximum amylase yield [12]. The ninth run found as the best combination of variables composed of (g/L): KH₂PO₄ (0.1); NaCl (0.1); (NH₄)SO₄ (0.1); Yeast extract (15); Soluble starch (15); Inoculum concentration 1%; pH 9 at 37 °C for the maximum production of amylase.

Table 1 Influence of various physic-chemical variables on amylase production by *Bacillus* sp. H7

INC (h)	Amylase (U/mL)	pH	Amylase (U/mL)	Temp (°C)	Amylase (U/mL)	CS (1%)	Amylase (U/mL)	NS	Amylase (U/mL)	INO (1%)	Amylase (U/mL)
0	0.00	4	56.89±5.3	20	90.1±5	Glucose	144.64±2.1	Peptone	132.6±3.2	0.50%	126.53±0.5
6	59.81±7.5	5	87.53±6.7	25	109.8±4.3	Starch	149.1±2.1	Casein	121.24±5.0	1.00%	133.17±0.6
12	87.24±6.6	6	113.61±0.6	30	121.24±2.1	Fructose	129.81±2.1	Tryptone	111.6±9.0	1.50%	140.03±0.7
18	102.89±1.0	7	128.89±1.0	35	127.24±2.1	Sucrose	121.24±7.5	Yeast ext.	152.96±3.2	2.00%	132.6±3.2
24	117.24±2.8	8	133.53±1.2	40	122.81±1.7	Lactose	109.17±0.6	Urea	88.24±7.0		
30	125.89±1.0	9	131.53±1.5	45	112.8±5.5	Maltose	127.81±4.0	NH4NO3	75.89±7.5		
36	130.53±2.0	10	125.17±1.5	50	101.24±4.9	Dextrose	97.1±4.3	NH4CL2	60.96±4.4		
42	128.89±2.7	11	119.23±4.7								
48	125.53±1.5	12	111.17±4.0								
54	120.53±1.5	13	98.24±5.0								
60	113.89±4.0	14	93.83±7.2								
66	107.53±4.7										
72	100.53±8.4										
78	98.89±7.0										
84	95.17±7.5										
90	92.89±6.2										
96	90.24±7.5										

INC - Incubation; pH - Media pH; NaCL - NaCl concentration (M); Temp – Temperature; CS - Carbon source; NS - Nitrogen Source INO - Inoculum level.

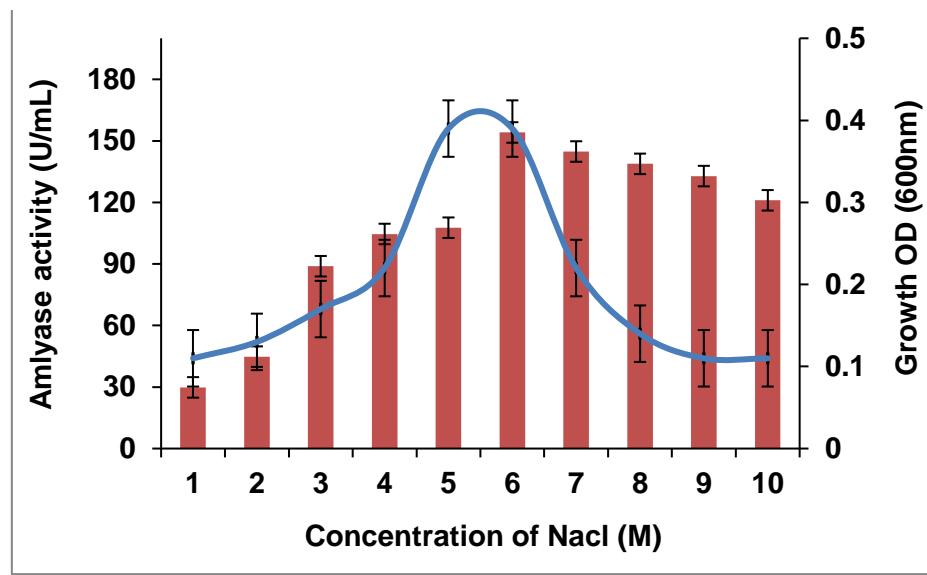


Figure 2. Influence of salt concentration on growth and amylase production in *Bacillus* sp. H7

The resulting production variables were analyzed by multiple linear regression analysis, the estimated *t*- and *p*-values of each media variable. On basis of high *t*-values and *p*-value (*p*<0.05) the significant variables were identified, among the variables soluble starch (*p*=0.002), media pH (*p*=0.008) and incubation period (*p*=0.002) showed the positive effect. The statistical model itself is significant with a *p*-value of 0.008 (Table 2). The Pareto chart illustrates the level of significance of all the media variables on the amylase production (Figure 4). The goodness of fit model was checked by the coefficient of determination (*R*²) which indicated that the model could explain up to 98.88% variation of the data.

Table 2. ANOVA of PBD model for amylase yield (coded units)

Term	Effect	Coef	SE Coef	<i>t</i> -Value	<i>p</i> -Value
Constant		137.630	0.969	141.97	0.000
KH ₂ PO ₄		-4.563	2.281	-2.35	0.100
NaCl		-4.325	2.162	-2.23	0.112
(NH ₄) ₂ SO ₄		-1.673	0.836	-0.86	0.452
Yeast extract		-5.211	2.605	-2.69	0.075
Soluble starch		19.367	9.684	9.99	0.002*
pH		12.301	6.150	6.34	0.008*
Inoculum Concentration		4.879	2.440	2.52	0.086
Incubation period		19.429	9.715	10.02	0.002*

2.5 Estimation of optimization concentration of significant variables using central composite design (CCD) of response surface methodology (RSM)

The optimum levels of significant variables and the effect of their interactions on amylase production were determined by CCD experiments. Soluble starch, media pH and incubation period were selected as variables based on the results of the PBD. The experimental design was carried out to determine the optimum concentrations/levels. The coded and actual values of the three independent variables for amylase production are tabulated in Table 2. The results of 20 runs from CCD experiments for studying the effects

of 3 independent variables on amylase production are represented in Table 2. From the RSM results, the maximum experimental value for amylase production was 196.66 ± 1.09 U/mL. The regression analysis data was fitted to a quadratic model. The second order regression

$$Y = -21011 - 190.3(E) + 212.8(F) + 1138.2(H) - 6.69(E^2) - 10.983(F^2) - 15.866(H^2) - 0.43(EF) + 5.64(EH) - 0.729(FH) \quad (1)$$

Where, Y is the yield of amylase (U/mL),
 E is the Concentration of soluble starch (%),
 F is the media pH and H is the incubation period (h).

The statistical significance was determined by the *f*-test and the analysis of variance (ANOVA) for the response surface quadratic model is presented in Tables 6. The *f*-value of 186.2 from ANOVA for amylase production implies that the model is significant. This is also evident from the model *f*-value and the probability value at $p < f$ value, which was about 0.0 (less than 0.05). The goodness of the model can be determined from the determination coefficient (R^2) and the correlation coefficient (R) [19]. The R^2 value of 0.9941 suggests 99.41% variability in amylase production. The closer the value of R (R = multiple correlation coefficient) to 1, better is the correlation between the experimental and predicted values [20]. The *p*-values correlate with the significance of each coefficient. It is important to indicate the pattern of mutual interaction between the coefficients (Table 3). The smaller the *p*-value, more significant is the corresponding coefficient [21]. The all linear, quadratic coefficients, and one interaction coefficients i.e. E*F were observed to be significant. Since it is a hierarchical model, insignificant coefficients were not omitted. RSM 3D Surface plots (Figure 3 A-C) provide the relation between the response and experimental levels of each variable. These plots are useful in understanding the type of interaction among test variables in order to deduce the optimum conditions [15,16]. The results of the PBD evidenced that an increase in soluble starch concentration (1.5%), interaction with pH at a maximum level of 9 and incubation of 37 h increased the amylase production (Figure 3C).

Table 3 ANOVA of CCD model using RSM of amylase yield in *Bacillus* sp. H70

Source	DF	Coef	SE Coef	Adj SS	Adj MS	t-value	f-Value	p-Value
Constant		196.060	0.821			238.89		0.000
Model	9			6785.68	753.96		186.20	0.000
Soluble starch	1	-2.007	0.545	55.01	55.01	-3.69	13.59	0.004
pH	1	10.431	0.545	1485.93	1485.93	19.16	366.97	0.000
Incubation period	1	-4.298	0.545	252.30	252.30	-7.89	62.31	0.000
Soluble starch*Soluble starch	1	-1.673	0.530	40.31	40.31	-3.16	9.96	0.010
pH*pH	1	-10.983	0.530	1738.31	1738.31	-20.72	429.29	0.000
Incubation period*Incubation period	1	-15.866	0.530	3627.72	3627.72	-29.93	895.90	0.000
Soluble starch*pH	1	-0.217	0.711	0.38	0.38	-0.30	0.09	0.767
Soluble starch*Incubation period	1	2.822	0.711	63.69	63.69	3.97	15.73	0.003
pH*Incubation period	1	-0.729	0.711	4.25	4.25	-1.02	1.05	0.330
Lack-of-Fit	5			38.64	7.73		20.88	0.002
Pure Error	5			1.85	0.37			

Using the above results from RMS analysis the optimum values were predicted for the independent significant variables (Figure 3) the optimized levels of these variables in combination with other media variables the maximum production was predicated to be

199.90 U/mL. The predicted data were validated through confirmatory experiments performed in triplicates. A 1.29-fold increase in amylase activity against un-optimized (OVAT) medium was achieved in the present study authenticating the efficacy of RSM in process optimization (Figure 4).

2.6 Model validation and scale-up at laboratory scale (5L) bioreactor

Once the parameters were standardized in the shake-flasks culture, the experiment was scaled-up to a lab scale bioreactor (5 L). The yield of amylase increased by 1.01 fold (205.69 U/mL), it could be possible because, the enzyme production in a bioreactor is higher than in shake-flasks culture as the various critical variable factors such as the dissolved oxygen (DO) and the pH can be optimally controlled at the desired levels [22].

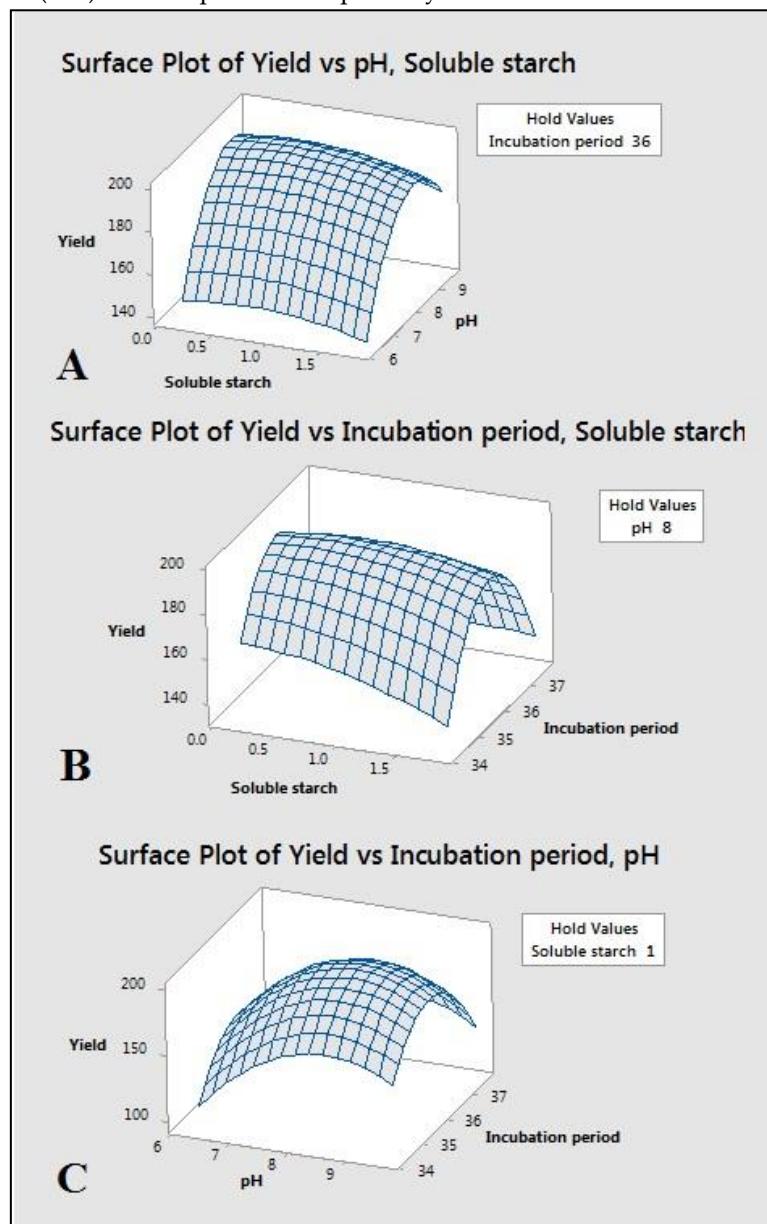


Figure 3. Response surface 3D contour plots representing interaction between variables affecting amylase production (A) soluble starch and pH (B) soluble starch and incubation period (C) incubation period and pH. Other variables were kept constant

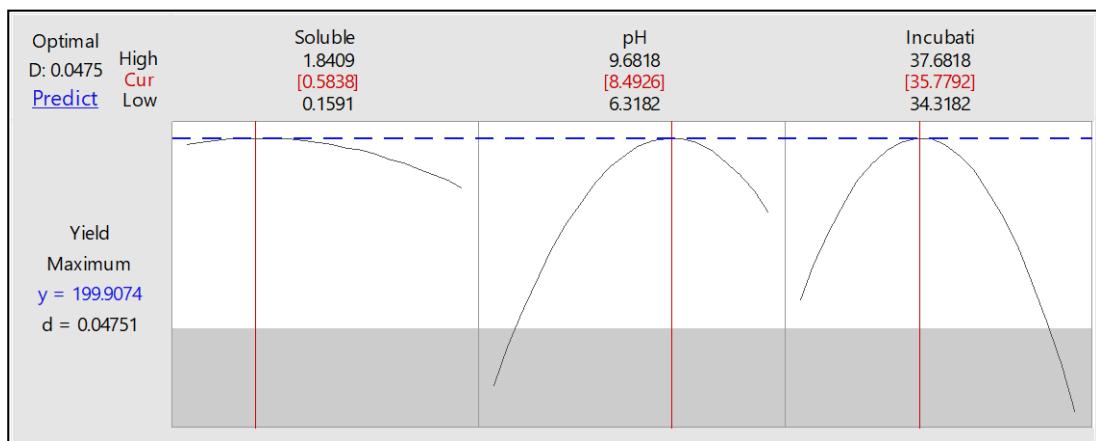


Figure 4 Response optimizer plot illustrating the optimum levels of variables for maximum amylase production by *Bacillus* sp. H7

3. Discussion

The extraordinary metabolic and physiological capabilities of marine isolates to produce metabolites with unique characteristics and tolerance extreme marine environmental conditions like high salt concentration, temperature and pressure which are rarely found in terrestrial microorganisms [23,24].

Incubation period plays an important role in the growth rate of microbial cells as well as on the enzyme production [25]. The yield amylase was observed very less in the initial lag phase of growth as the isolate begins to adapt the in vitro growth conditions, with the increasing incubation period the notable increase in the amylase was observed and the optimum yield was obtained at 36 h of incubation (130.53 ± 2.0 U/mL). Further increment in the incubation period decreased the amylase production. At 72 h of incubation, it was extremely reduced could be due to the microbial cell death, exhaustion of nutrients, accumulation of byproducts in the culture medium, such as toxins and growth inhibitors in addition to the cells showed diminished amylase enzyme biosynthesis during its decline phase of growth [26]. Abdel-Fattah et al. [27] reported the similar results where they observed the maximum amylase at 32 h of incubation in case of *Bacillus licheniformis* AI20. Whereas Rehman et al. [25] reported the maximum production of amylase by *Bacillus cereus* AS2 after 72 h of incubation period.

The Temperature and the pH both are most important variables for to regulate a bioprocess. Optimum amylase yield was recorded at a temperature of 35°C . This is because 35°C is the optimum temperature for both of the responses i.e., bacterial growth as well as enzyme production. Event after the 40°C the enzyme production was not much affected while at 50°C the decrease in production was observed in very less amount which suggests that the isolate has the good thermal stability. In case production media pH the amylase production was observed in pH ranging from 6.00-12.00 and at pH 9.00 the maximum amylase production was achieved. The alkali nature of the isolate reviles from the amylase production at such high alkali condition. The obtained results shows the similarities with the production results of *Bacillus US147* where the incubation temperature was 45°C and pH 9 resulted as the optimum condition for the amylase production [28].

The concentration of Inoculum is a significant parameter in any bioprocess for the enzyme production. The production will be affected by high concentration due to the

increase in moisture contains as well as biomass production whereas the low concentration results in the prolonged production period to transform the substrate into the product [29]. Similarly, Mishra et al. [30] and Nithya et al. [31] have been reported that 1.5% of inoculum concentration was optimum for maximum amylase production for *Bacillus* sp. and *Bacillus licheniformis* KSU-6 respectively.

The source and concentration of nitrogen and carbon are crucial aspects to achieve the maximum production of any enzyme as the various physiological pathways require the carbon as a substrates regulates the enzyme production [32]. In present study, the maximum amylase production was observed 1.0% soluble starch as sole source of carbon when compared to other carbon sources tested for isolate *Bacillus* sp. H7. These results are supported with the previous reports stated that the amylase production was increased when soluble starch used as carbon sources [33-35].

The nitrogen sources have influenced the extracellular amylase enzyme production and also supplement with specific nitrogen source on enzyme production differs from organism to organism, Nitrogen stimulates and down regulates the enzyme production by microorganisms [36].

Yeast extract is a complex nitrogen source, it provides all essential amino acids required for the synthesis of the enzyme and hence it supports higher yields of the enzyme [37]. In case of isolate *Bacillus* sp. H7, yeast extract is dawn out as a best source of nitrogen as compared to the other nitrogen sources tested in presence of 1% yeast extract isolate showed the maximum amylase production. Similarly, Saxena et al. [33] and Teodoro and Martins [38] have been reported that yeast extract influenced the amylase production in *Bacillus* sp.

Salinity of the growth medium strongly influenced the amylases production where in low concentrations the production was moderately low and the increase was observed with increasing concentrations. The isolate *Bacillus* sp. H7 showed the maximum production at high salinity which suggests the halo-tolerant nature of the isolate. Rehman et al. [25] demonstrated the induction of the amylase production in the increasing concentration of NaCl.

The optimum concentrations or levels of the various physicochemical media variables were resulted from the OVAT approach but their interactive effect of amylase production was studied using the CCD and RSM these studies reveled the significant independent variables having important role in the bioprocess as well as their combined effect on the production. The statistical approached resulted in the maximization of amylase production by 1.29 fold as compared to the classical OVAT approach at shake flask level and upon scale up the production was further increase up to 1.01 fold. Elmansy et al. [18] reported the enhance production of α -amylase by the thermo-halophilic bacterial strain *Bacillus* sp. NRC22017 isolated from marine environment where they tested the various physic chemical parameters for the optimum production (15.15 ± 0.47 U/mL). The results obtained by optimization of production process found higher than the previous experimental reporting's by Blanco et al. [39] and Ahmed et al. [23] on the extracellular amylase production by *Bacillus subtilis* sp.(9.26 and 145.4 U/mL respectively). The obtained results clear that the statistical approaches has the significant role in the amylase

production by *Bacillus* sp over the classical optimization approaches for the process optimization [40].

4. Materials and Methods

4.1 Sample collection, Screening and quantitative estimation of amylase production by halophilic isolates

The marine water samples were collected from surface and deep one to two meter from the different locations in the radius of five kilometers of Harihareshwar beach in India (18°00'11.5"N 73°01'01.6"E); samples were collected in sterile plastic containers and mixed to make a composite sample and used for the isolation. Prior to isolation the marine water samples were serially diluted and 0.1 mL of an aliquot (10^{-4}) was spread on nutrient agar medium. Morphologically different bacterial isolates were purified and preserved in 20% (v/v) glycerol with phosphate buffer saline (pH 7.0) until further use. In order to screen of amylase producing isolates, colonies were separately grown on starch agar and incubated at 30 °C for 24 h. The amylolytic activity of isolates was confirmed by the starch hydrolysis test using iodine [41].

Amylase production was carried out in modified amylase production medium (APM) [42] containing (g/L); (NH₄)₂SO₄; 0.5, KH₂PO₄; 0.1, MgSO₄; 0.1, CaCl₂; 0.01, NaCl; 10 and starch; 10.0. Log phase culture of each isolate was grown in the production medium at 30 °C for 24 h. Amylase activity was estimated by centrifugation of culture broth at 10,000 rpm for 10 min and the cell free supernatant was assayed by 3,5-dinitro salicylic acid (DNSA) method [43]. One unit of the amylase activity was defined as the amount of enzyme required to produce 1 μmole of maltose from soluble starch per minute under the assay conditions at 25 °C. The total protein content of the sample was estimated using bovine serum albumin as standard [44].

4.2 Characterization of the potent isolate

4.2.1. Phenotypic characterization

For this, isolates were grown on nutrient agar at 30 °C for 24 h. The morphological characteristics were studied followed by Gram staining, motility and spore staining.

4.2.2 Molecular identification of isolate using 16S rRNA Sequencing

Identification of amylase producing isolate was carried out using 16S rRNA sequencing approach. The Genomic DNA (gDNA) isolation was performed as per the protocol of Sambrook and Russell [45]. By using the gDNA as a template 16S rRNA genomic region was amplified with help of universal primers 27F (5'-GAGTTGATCMTGGCTCAG-3') and 1525R (5'-AAGGAGGTGATCCAGCC-3') in Gene Amplifier PCR System 9700 (Perkin Elmer, USA). For the polymerase chain reaction (PCR) 20–50 ng of template DNA was used for amplification in the PCR condition, initial denaturation at 94 °C for 5 min, 35 cycles of denaturation at 94 °C for 1 min, annealing at 55 °C for 1 min and extension at 72 °C for 1 min, final extension at 72 °C for 7 min with a final hold at 20 °C. Followed by the purification of PCR products on 1.0% agarose gel and sequenced on ABI 3730Xl automated sequencer using a ready reaction kit (Perkin Elmer Applied Biosystems Division, CA). Amplified sequences were identified from NCBI (<http://www.ncbi.nlm.nih.gov>) and EzTaxon (<http://www.eztaxon.org>) database and phylogenetic trees were constructed by using the neighbor-joining method with the help of MEGA5 software [13].

4.3 Influence of physico-chemical media variables of amylase production

The media variable like Incubation period and temperature, media pH, inoculum concentration, carbon and nitrogen sources were examined for the amylase production by *Bacillus* sp. H7 using one variable at a time (OVAT) approach. For the examination of incubation period on the amylase production the culture was incubated for 6-96 h by keeping interval of 6 h. The amylase production was also examined for incubation temperature by incubating the culture at different temperature ranging from 20-50 °C. The effect of media pH on enzyme production was examined by the production of amylase in the varying media pH range 4.0-14.0. The optimum inoculum concentration was estimated by inoculating culture with different inoculum volume (0.5-2.0%). The influence of various carbon source like glucose, starch, fructose, sucrose, lactose, maltose, dextrose and nitrogen sources like peptone, casein, tryptone, yeast extract, urea, NH₄NO₃ and NH₄Cl₂ was tested at 1% concentration.

4.4. Statistical Analyses

All the OVAT experimental result data mentioned is the mean of triplicates followed by the standard deviation analyzed using the Student's - *t* test and the observation having *p* ≤ 0.05 were considered as significant [46].

4.4.. Effect of salinity on growth and amylase production

The effect of various concentrations (0-10 M) of salt (NaCl) on the growth and production of amylase by the isolate was evaluated by separately growing the log phase culture (5x10⁵ cells/mL) of isolate in each production broth containing varying amounts of NaCl salt (0-10 M) at 30 °C for 48 h. Following the incubation, cell growth was measured in terms of absorbance (optical density [OD]) at 620 nm and amylase activity was estimated as described above.

4.5. Evaluation of significant variable of production media by Plackett–Burman Design (PBD)

Among the various variables of production media identification of most significant one is the major process in bioprocess optimization [29], with this regards the investigation was initiated using the eight variables viz., KH₂PO₄ (A), NaCl (B), (NH₄)₂SO₄ (C), yeast extract (D), soluble starch (E), pH (F), inoculum concentration (G) and incubation time (H). These variables were analyzed at their high (+1) and low (-1) levels composing Plackett–Burman design of 12 set of different experiments illustrated in Table 4 [47,48] .

Table 4 Placket-Burman Design (PBD) experimental design for screening and evaluating factors influencing amylase production from *Bacillus* sp. H7

Run	A	B	C	D	E	F	G	H	Amylase activity (U/mL)	
									Predicted	Experimental
1	+1	-1	+1	-1	-1	-1	+1	+1	135.60	133.49 ± 1.22
2	+1	+1	-1	+1	-1	-1	-1	+1	122.86	121.89 ± 0.48
3	-1	+1	+1	-1	+1	-1	-1	-1	130.90	128.60 ± 0.68
4	+1	-1	+1	+1	-1	+1	-1	-1	118.38	119.17 ± 0.76
5	+1	+1	-1	+1	+1	-1	+1	-1	127.68	128.65 ± 0.79
6	+1	+1	+1	-1	+1	+1	-1	+1	158.06	160.36 ± 1.18

7	-1	+1	+1	+1	-1	+1	+1	-1	123.50	122.71 ± 0.34
8	-1	-1	+1	+1	+1	-1	+1	+1	154.32	156.43 ± 0.77
9	-1	-1	-1	+1	+1	+1	-1	+1	163.41	161.30 ± 0.68
10	+1	-1	-1	-1	+1	+1	+1	-1	149.51	148.54 ± 1.04
11	-1	+1	-1	-1	-1	+1	+1	+1	149.81	150.60 ± 1.53
12	-1	-1	-1	-1	-1	-1	-1	-1	117.53	119.82 ± 0.82

The media variables gave the confidence level greater than 95% were considered as significant for amylase production by recording response in the form of amylase activity. The Pareto chart analysis represents the magnitude of each media variable (Figure 5).

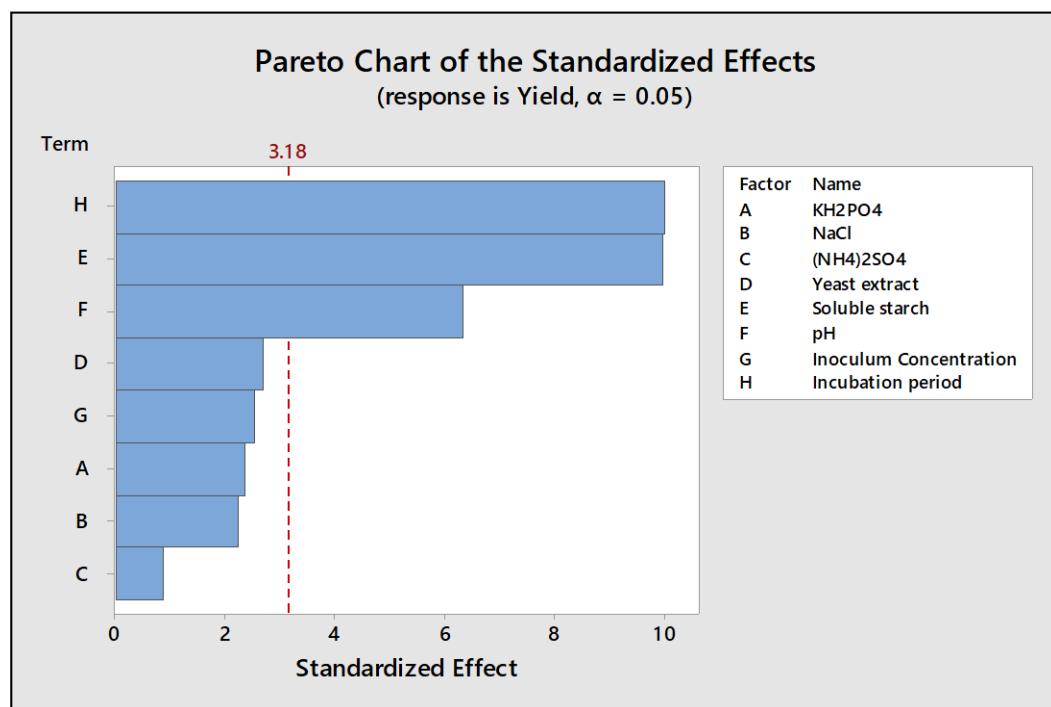


Figure 4 Pareto chart representing the effects of medium variables on amylase production by *Bacillus* sp. H7 ($p<0.05$)

4.4 Estimation and optimization concentration of significant variables using central composite design (CCD) of response surface methodology (RSM)

The PBD results proposed that, incubation period (F), soluble starch (E), and media pH (H) these variables has the significant role in the amylase production by *Bacillus* sp. H7 As a results these three variables were further processed for the estimation of optimum concentration or levels for the maximum production of amylase using the central composite design (CCD) of response surface methodology (RSM) [45]. In CCD these variables were tested in 20 sets of experiments at five different levels consisting axial, factorial and central positions ($-\alpha$, -1 , 0 , $+1$, $+\alpha$) (Table 5 and 6). The experimental results were obtained in the form of amylase activity and correlated with the predicted yield using

Table 5. Range of values for the response surface method- CCD

Independent Variable	Code	Coded level				
		$-\alpha$	-1	0	+1	$+\alpha$
Soluble starch (%)	E	0.16	0.5	1	1.5	1.84
Media pH	F	6.32	7.00	8.00	9.00	9.68
Incubation period (h)	H	34.32	35.0	36	37	37.68

Table 6. Experimental data obtained for significant variables obtained from PBD in CCD.

Run Order	Type	Soluble starch		pH		Incubation period		Amylase activity (U/mL)	
		Coded	Experimental	Coded	Experimental	Coded	Experimental	Predicted	Experimental
1	Factorial	-1	0.50	-1	7.00	-1	35.00	165.29	165.70 ± 0.64
2	Factorial	1	1.50	-1	7.00	-1	35.00	156.06	158.86 ± 0.29
3	Factorial	-1	0.50	1	9.00	-1	35.00	188.04	188.61 ± 0.97
4	Factorial	1	1.50	1	9.00	-1	35.00	177.95	178.85 ± 1.05
5	Factorial	-1	0.50	-1	7.00	1	37.00	152.51	153.79 ± 0.95
6	Factorial	1	1.50	-1	7.00	1	37.00	154.57	156.19 ± 0.64
7	Factorial	-1	0.50	1	9.00	1	37.00	172.34	171.74 ± 1.62
8	Factorial	1	1.50	1	9.00	1	37.00	173.54	175.32 ± 0.75
9	Axial	-1.68	0.16	0	8.00	0	36.00	194.70	194.78 ± 0.88
10	Axial	1.68	1.84	0	8.00	0	36.00	187.95	184.79 ± 0.20
11	Axial	0	1.00	-1.68	6.32	0	36.00	147.45	144.88 ± 0.34
12	Axial	0	1.00	1.68	9.68	0	36.00	182.54	182.02 ± 1.51
13	Axial	0	1.00	0.0	8.00	-1.68	34.32	158.41	156.69 ± 0.69
14	Axial	0	1.00	0.0	8.00	1.68	37.68	143.96	142.59 ± 0.99
15	Central	0	1.00	0.0	8.00	0.0	36.00	196.06	196.25 ± 1.22
16	Central	0	1.00	0.0	8.00	0.0	36.00	196.06	196.62 ± 0.95
17	Central	0	1.00	0.0	8.00	0.0	36.00	196.06	195.68 ± 1.02
18	Central	0	1.00	0.0	8.00	0.0	36.00	196.06	195.15 ± 0.35
19	Central	0	1.00	0.0	8.00	0.0	36.00	196.06	196.53 ± 0.79
20	Central	0	1.00	0.0	8.00	0.0	36.00	196.06	196.66 ± 1.09

analysis of variance (ANOVA) and fitted into the second-order polynomial equation (Equation 1) to represent the confidence of algorithm processed.

The second-order polynomial equation (Equation 2) generated an empirical model that relates to the responses obtained in the independent variable to the experiment. A second-order polynomial equation (Equation 1) was then fitted to the response by multiple regression procedures. This resulted in an empirical model that related the response measured in the independent variables to the experiment.

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

Where

Y_i – predicted response,

$X_i X_j$ - input variables (influence the response variable Y),

β_0 - constant,
 β_i - i^{th} linear coefficient,
 β_{ii} - quadratic coefficient,
 β_{ij} - ij^{th} interaction coefficient.

4.5 Model validation and scale-up at laboratory scale (5L) bioreactor

The amylase production with the optimized levels of the significant media variables was performed at flask level followed by scale up using laboratory bioreactor [Model LF-5 Murhopye Scientific Co., Mysore, India – 5L capacity].

4.6 Software and data analysis

The PBD and RSM modeling and statistical analysis of optimization were performed with software Minitab 18 (Minitab GmbH, Munich, Germany).

5. Conclusions

Bacteria having ability to produce amylase enzyme were isolates from the marine environment. The potent isolate was characterized (morphologically and biochemically) and was identified using 16S rRNA gene sequencing method. The statistical approach facilitates the combinations of experiments to elucidate the significant variables of the production medium for optimum production of extracellular amylase by *Bacillus* sp. H7. The significant variables initially screened by PBD were further optimized by using RSM. The successfully scale up of statistical-based shake flask process to laboratory scale (5L) bioreactor validated the variables and their concentrations studied at the shake flask level. The values of variables namely Incubation period (H), soluble starch (E) and pH (F) predicted at shake flask level further enhanced the enzyme yield by 1.29 fold (from 156.53 ± 2.58 to 203.23 ± 4.0 U/mL). This reflects the usefulness of the application of statistical optimization in amylase production. The experimental results obtained showed a good relationship with the values predicted by Minitab 18 (software). Thus present results illustrated that the statistically optimized bioprocess could be the best suitable approach for enhancing the productivity of extracellular amylase at an industrial scale.

Author Contributions: Conceptualization, Supervision and project administration J.N.B. and R.Z.S.; Data Curation, R.Z.S. and H.P.J.; Formal analysis and investigation, J.N.B., and V.A.T.; Methodology, J.N.B., and V.A.T ; Project administration, J.N.B.; Validation, R.Z.S. and H.P.J.; writing original draft, J.N.B., and V.A.T.; writing-review and editing, R.Z.S.

Institutional Review Board Statement: Not Applicable

Informed Consent Statement: Not Applicable

Data Availability Statement: All the data is available in the manuscript

Acknowledgments: We greatly appreciate Universiti Teknologi Malaysia through UTM Transdisciplinary Research Grant.

Conflicts of Interest: The authors declare no conflict of interest.

Sample Availability: Not applicable

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