Isolation and Characterization of Bacteria for Amylase Production under Solid State Fermentation Using Damaged Wheat as Substrate

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A B S T R A C T

A bacterial isolate (I5) along with Bacillus subtilis and Bacillus circulans was evaluated for extracellular amylase production under SmF and SSF conditions. In SmF (Temperature - 28±2°C, pH -7 and inoculum size-10⁶ cfu ml⁻¹, Time- 96 h), maximum saccharifying activity (SA) was observed with Bacillus subtilis (0.62 IU/ml/min) followed by isolate (I5) (0.58IU/ml/min) and Bacillus circulans (0.46IU/ml/min). Whereas dextrinizing activity (DA) was recorded with Bacillus circulans (1.69U/ml), followed by isolate (I5) (1.65U/ml) and Bacillus subtilis (1.02U/ml). In SSF(1:2 moisture content, incubation time -96 hours, pH-7 inoculum size-10⁶ cfu ml⁻¹ Temp-37°C) using different substrates (potato starch, corn starch, raw wheat flour and rice starch) as carbon sources. Maximum saccharifying activity (23.10IU/ml/min) was expressed by Bacillus circulans in 1% rice starch at 28±2°C. Whereas highest dextrinizing activity was recorded with 1% corn starch (3.31U/ml). On the basis of the present study, Bacillus circulans being the most potential strain for maximum and cost effective alpha amylase enzyme production at appropriate conditions could be utilized as bio-inoculant in fermentation technology in sustainable manner.

K e y w o r d s
Dextrinizing, Saccharifying, Inoculum, Bio-inoculant, Fermentation

Introduction

Profiling microorganisms with high potential for amylase production in submerged fermentation (SmF) which is synthetic media has been widely recognized due to their myriad applicability in bioethanol production. However, the contents of a synthetic medium are very expensive and uneconomical, so there is urgent need to be replaced with more economically available agricultural, industrial and domestic by products, as they are considered to be good substrates for SSF to produce enzymes.

According to an estimate, 12MT (6%) of grains are damaged by post-harvest storage due to poor storage facilities and hence damage by insects, rodents, birds and microbial spoilage (Sharon et al., 2014). These infested spoiled grains though unfit for human and animal consumption, possess high starch content which provides an adequate substrate for saccharification by microbial
alpha- and gluco-amylases for fermentation into ethanol.

SSF holds tremendous potential for the production of enzymes in view of its economic and engineering advantages. It can be of particular relevance in those processes where a crude fermented product may be used as an enzyme source (Pandey et al., 1999). The major critical factors affecting microbial synthesis of enzymes in a SSF system includes selection of a suitable substrate and strains, particle size of the substrate, inoculum concentration, moisture level of the substrate, temperature and pH. Thus it involves the screening of a number of agro-industrial materials for microbial growth and product formation. Currently, a large number of microbial amylases are available commercially and they have almost completely replaced the chemical hydrolysis of starch in the starch processing industry (Pandey et al., 2000). In these respect members of the Bacillus family namely Bacillus licheniformis, Bacillus Stearothermophilus etc. are being best utilized for thermostable alpha amylase production (Rehman and Saeed, 2015). Though, literature widely reports the use of SSF for the production of enzymes and other products has many advantages over submerged fermentation (Lonsane and Ramesh, 1990).

In the present investigation, a soil isolate amylase producing bacteria was selected and compared for enzyme production under submerged and solid state fermentation conditions using wheat flour as substrate.

**Materials and Methods**

**Collection of bacterial cultures**

Bacterial cultures, Bacillus subtilis (MTCC 121) and Bacillus circulans (MTCC 7906) were procured from Department of Microbiology, Punjab University, Chandigarh and Industrial Microbiology Laboratory, Department of Microbiology, Punjab Agricultural University, Ludhiana respectively. All the cultures were being maintained on nutrient agar medium and further was sub-cultured once in a month throughout the period of investigation and stored at 4°C in refrigerator.

**Procurement of agricultural based substrates for SmF and SSF systems**

Damaged wheat grains were procured from different areas of Punjab, milled according to different sieving sizes (1.4 mm, 750 microns, 355 microns, 180 microns and 70 microns) and further subjected to starch analysis by the method of (Clegg, 1956).

**Isolation of amylase producing microorganisms**

The soil samples were collected from vegetable field, PAU, Ludhiana. The bacterial strains were isolated by a serial dilution plate technique and appropriate dilutions (100µl) were placed on starch agar as the carbon source for amylolytic bacteria and incubated at 28±2°C for 3-5 days.

The colonies were selected on the basis of morphological and culture characteristics, then further streaked on their specific media for purification. After purification, bacterial isolates were transferred on nutrient agar slants at 4°C temperature. All the bacterial cultures were maintained on nutrient agar and further sub-cultured once in a month throughout the period of investigation and stored at 4°C in refrigerator.

**Screening of alpha amylase producing microorganisms**

All the bacterial isolates along with two standard cultures (Bacillus subtilis and Bacillus circulans) were screened qualitatively
for the presence of extracellular amylase production in starch agar media containing composition (soluble starch-10.0 g L⁻¹; peptone-10 g L⁻¹; yeast extract-5.0 g L⁻¹; agar-20.0 g L⁻¹). Formation of clear zones of hydrolysis around the vicinity of the isolates was measured in terms of amylolytic activity index. Amylolytic activity index (AI) may be defined as the Ratio between the zone diameter (mm) - colony diameter (mm)/colony diameter (mm).

**Growth kinetics of amylase producing bacterial isolates along with standard cultures**

Amylase producing bacterial isolate 5 (I5) and two standard cultures (*Bacillus subtilis* and *Bacillus circulans*) were studied for their growth profile. 100 ml minimal starch medium (Na₂HPO₄-5.8 g L⁻¹; KH₂PO₄-3.0 g L⁻¹; NaCl-0.5 g L⁻¹; NH₄Cl-1.0 g L⁻¹; MgSO₄ (1M) - 2.5 ml, FeCl₃ (0.01M)-2.5ml, 10% Glucose solution-50 ml, pH-7) was prepared and autoclaved at 121°C (15psi) for 20 minutes. After cooling the medium at room temperature, each flask containing specific medium was inoculated with a loop full bacterial suspension which shows maximum starch hydrolysis. The inoculated flasks were incubated at 28±2°C under shaking at 120 rpm and samples (0-120h) were collected from each flask to study growth at different time intervals and absorbance was measured by using spectrophotometer at 600 nm wavelength.

**Alpha amylase production**

**Submerged fermentation**

On the basis of qualitative screening of amylase production, growth curve and kinetic study of bacterial isolate 5 and two standard cultures (*Bacillus subtilis* and *Bacillus circulans*) were selected for qualitative amylase assessment on the basis of extracellular amylase production under submerged fermentation conditions (Temp-28±2°C, pH-7, inoculum size-10⁶ cfu/ml) after incubating at different intervals of time (24, 48, 72, 96 and 120 hours). The fermentation medium was harvested by centrifugation at 5000 rpm for 20 min at 4°C to obtain the crude extract, which served as enzyme source.

**Solid state fermentation**

The experiments were conducted in 250-ml Erlenmeyer flasks containing 5 g of damaged wheat flour (355 microns) substrate moistened with 10 ml of minimal salt solution containing composition (NaCl - 0.8 g L⁻¹; KCl-0.8 g L⁻¹; CaCl₂- 0.01 g L⁻¹; Na₂HPO₄-2.0 g L⁻¹; MgSO₄-0.2 g L⁻¹; FeSO₄- 0.1 g L⁻¹; Glucose-8 g L⁻¹ and NH₄Cl-2 g L⁻¹. After sterilization, the flasks were cooled at room temperature and inoculated with a 10% (w/w) inoculum and incubated at 37°C for 96 h. After fermentation, the entire contents of the flasks were filtered through muslin cloth. The filtrates were pooled together and centrifuged at 10,000 rpm for 15 min and clear supernatant was used as the crude enzyme source and assayed in terms of saccharifying and dextrinizing activity.

The SA and DA were determined by the methods of (Miller, 1959) and (Fuwa, 1954), respectively.

**Results and Discussion**

**Isolation and screening of alpha amylase producing microorganisms**

Ten bacterial isolates were procured from vegetable farm soil and were coded as isolate I₁ to I₁₀. All the ten isolated cultures along with standard cultures *Bacillus subtilis* (MTCC 121) and *Bacillus circulans* (MTCC 7906) were assayed for extracellular amylase assay (Aneja, 2012). Out of the ten bacterial isolates, I₅ and the two standard cultures
showed maximum zone of starch hydrolysis and amylolytic activity index was calculated for each colony. Amylolytic activity index was found in *Bacillus circulans* (2.98) followed by I₅ (2.58) and *Bacillus subtilis* (2.1). The mechanism of clear zone observation was attributed to the fact that amylase produced during the growth of the microorganisms hydrolysed the starch around the colony, thereby testing negative when flooded with iodine. The un-hydrolysed part of the plate tested positive due to the presence of starch (amylose), hence the blue-black appearance. Our results were also in accordance with (Singh and Kumari, 2016) that observed zone of clearance on starch agar media and among five, *Bacillus* sp. B3 showed the maximum zone of clearance on the starch agar medium i.e., 8 mm. Similarly, Verma *et al.*, (2011) found the maximum amount of amylase production in *B. subtilis* followed by *Bacillus megaterium* among nine strains tested which included *B. cereus*, *B. megaterium* and *B. subtilis*.

**Growth kinetics of amylase producing bacterial isolates**

The three bacterial strains *viz.* Isolate 5 (I₅), *Bacillus subtilis* and *Bacillus circulans*, were studied for their growth profile in minimal starch medium having soluble starch as a sole carbon source. In Figure 1 the cell growth of all the bacterial strains increased exponentially with increase in incubation period up to 48 hours and then declined till 96 hours of incubation under shaking (120 rpm) conditions. The comparative analysis of growth rate (h⁻¹) of Isolate 5 (I₅), *Bacillus subtilis* and *Bacillus circulans* is shown in Table 1.

Our results are well in line with Mishra and Behera (2008) that reported a soil *Bacillus* isolate which manifested maximum growth at 48 h and after that growth declined at 92 h. Similarly, Vyas *et al.*, (2016) also studied the growth kinetics of *B. subtilis* was done with respect to α-amylase production in a basal medium with pH 7.0 at 25±2°C. Chandra *et al.*, (1980) reported that composition, incubation temperature and pH of media greatly affect the growth and production of extracellular amylase production in bacteria. Similarly, (Ho and Ku, 2017) reported that *B. subtilis* possessed the maximum specific growth rate at 48 h with the maximum cell productivity of 1.98 × 10¹⁰ cells/L/h attained during the exponential growth phase.

**Amylase production in submerged fermentation**

In submerged fermentation, highest SA was observed with *Bacillus subtilis* (0.62 IU/ml/min) followed by isolate 5 (0.58 IU/ml/min) and *Bacillus circulans* (0.43 IU/ml/min) at 48, 48 and 72 hours of incubation period respectively as represented in (Fig. 1).

Maximum DA was found with *Bacillus circulans* (1.69 U/ml), followed by isolate 5 (1.65 U/ml) and *Bacillus subtilis*(1.02 U/ml) at 72 hours of incubation period respectively (Fig. 1). Further increase in incubation period did not show any significant results with increase in saccharifying and dextrinizing activities. Our results were coherent with Abd-Elhalem, (2015) who observed that with increase in incubation period, amylase concentration was reduced most probably due to the reduction in nutrients, accumulation of waste product, cell death and catabolite repression. The variation in amylase activities of the isolate could be due to difference in genetic makeup of microorganisms belonging to the same genera and species; it could also be due to the fact that the isolates were obtained from different environments which made them display different amylase activities.
Table 1 Comparative growth profile, saccharifying and dextrinizing activities of different Bacillus strains in minimal starch medium

| Bacterial strains | Time (hours) | Growth rate Per hour | Maximum amylase activity Saccharifying activity IU/ml/min | Dextrinizing activity IU/ml/min |
|-------------------|-------------|----------------------|----------------------------------------------------------|--------------------------------|
| Isolate 5 (I5)    | 48          | 0.11                 | 0.58                                                     | 1.65                           |
| *Bacillus subtilis* | 72          | 0.12                 | 0.62                                                     | 1.02                           |
| *Bacillus circulans* | 48          | 0.22                 | 0.46                                                     | 1.69                           |
| CD (5%)           |             | 0.13                 | 0.49                                                     | 0.58                           |

Table 2 Saccharifying and dextrinizing activities of bacterial strains for alpha amylase production with different substrates under solid state fermentation

| Different substrates (@ 1%) | Time interval (hrs) | Saccharifying activity (μ moles/ml/minute) | Dextrinizing activity (mg/ml) |
|-----------------------------|---------------------|-------------------------------------------|-------------------------------|
|                             |                     | B. subtilis | B. circulans | Isolate (I5) | B. subtilis | B. circulans | Isolate (I5) |
| Raw wheat flour             | 48                  | 9.65±0.46  | 7.32±2.72   | 0.81±0.08   | 0.10±0.02  | 0.09±0.04   | 0.01±0.00   |
|                             | 72                  | 5.96±1.03  | 11.17±1.37  | 0.44±0.09   | 0.14±0.03  | 0.79±0.18   | 0.24±0.05   |
|                             | 96                  | 5.62±0.64  | 12.54±1.31  | 0.27±0.05   | 0.46±0.04  | 1.10±0.15   | 0.15±0.04   |
| Potato Starch               | 48                  | 5.59±0.41  | 13.20±1.56  | 7.20±0.47   | 0.55±0.20  | 0.64±0.08   | 0.21±0.07   |
|                             | 72                  | 5.69±0.53  | 14.30±1.18  | 6.71±0.27   | 0.56±0.09  | 1.31±0.10   | 0.22±0.02   |
|                             | 96                  | 8.00±0.48  | 20.00±2.35  | 6.41±0.41   | 0.63±0.03  | 1.79±0.15   | 0.22±0.06   |
| Corn Starch                 | 48                  | 7.90±0.87  | 9.60±0.71   | 2.19±0.17   | 0.81±0.13  | 0.88±0.29   | 3.24±0.39   |
|                             | 72                  | 4.60±0.64  | 12.06±1.10  | 1.09±0.16   | 0.84±0.07  | 2.25±0.27   | 2.62±0.59   |
|                             | 96                  | 4.21±0.63  | 19.60±2.70  | 1.02±0.09   | 0.92±0.07  | 3.31±0.36   | 2.33±0.29   |
| Rice Starch                 | 48                  | 14.65±2.01 | 22.00±3.63  | 5.45±0.55   | 0.18±0.04  | 0.28±0.07   | 0.76±0.16   |
|                             | 72                  | 10.96±1.44 | 23.09±1.12  | 0.76±0.05   | 0.35±0.06  | 1.36±0.06   | 0.65±0.10   |
|                             | 96                  | 5.36±0.73  | 23.10±2.34  | 0.34±0.09   | 0.48±0.04  | 1.86±0.25   | 0.55±0.05   |
| CD at 5%                    |                     | 1.50±0.14  | 3.19±0.13   | 0.91±0.06   | 0.16±0.02  | 0.27±0.05   | 0.35±0.10   |
Fig. 1 Growth profile, saccharifying and dextrinizing activities of *Bacillus subtilis*, *Bacillus circulans* and isolate 5 (I5) on minimal starch medium

**Fig. 1 (a)**

**Bacillus subtilis**

Log number of cells

Saccharifying activity (IU/ml/min)

Log no. of cells

Saccharifying activity

Dextrinising activity

**Fig. 1 (b)**

**Bacillus circulans**

Log number of cells

Saccharifying activity (IU/ml/min)

Log no. of cells

Saccharifying activity

Dextrinising activity
Similarly, Raplong et al., (2014) reported that maximum amylase activity of 2.56 U/ml was produced by Bacillus cereus SB₂ after 24 hours incubation in submerged fermentation. This might be because after 24hrs, the cells reached the decline phase and displayed low amylase synthesis (Sivakumar et al., 2012). Simair et al., (2016) also investigated the time profile of bacterial growth and amylase production in synthetic medium and it was found that maximum dry cell weight (2.50g/L) and amylase enzyme titer (843 U/ml) was observed at 60 hours after which growth and enzyme titre gradually decreased.

Amylase production in solid state fermentation

In solid state fermentation, starch content of milled and sieved damaged wheat grains was analysed.

The starch content (%) of the sieved damaged wheat grains were 1.4 mm (48.2), 750 microns (48.6), 355 microns (49.2), 180 microns (47.6) and 70 microns (44.89). The relative proportion of starch content was observed to be high in damaged wheat grains, sieved to 355 microns size and was selected for further study.

The three selected bacterial strains viz. Isolate 5 (I₅), Bacillus subtilis and Bacillus circulans were grown on damaged wheat flour (355 microns size) as a substrate. Among the three strains, maximum saccharifying and dextrinizing activities were demonstrated by Bacillus circulans in the different substrates (potato starch, corn starch, wheat flour and rice starch) mentioned in Table 2. Maximum saccharifying activity was observed (@ 1% of different substrates) with Bacillus circulans in rice starch (23.10 IU/ml/min) followed by potato starch (20.00 IU/ml/min), corn starch (19.6 IU/ml/min), and wheat flour (12.54 IU/ml/min). Whereas, highest dextrinizing activity was recorded (@ 1% different substrates) with Bacillus circulans in corn starch (3.308U/ml) followed by rice starch (1.86U/ml), potato starch (1.79U/ml), and wheat flour (1.10U/ml).

Our results are in close agreement with Aassaret al., (1992) who reported that the alpha amylase enzyme obtained from Bacillus lentus was

![Fig.1(c)](image-url)
found to be highly active for hydrolysing dextrin (100%) and starch (96%), less active on glycogen (67.26 %) or amylose (68.92 %), whereas amyllopectin (17.20%) was weakly hydrolysed by the enzyme. Similarly, Farid and Shata (2011) revealed the enzymatic hydrolysis of different carbohydrates (soluble starch, potato starch, corn starch, yellow dextrin and white dextrin) with crude enzyme produced by A. oryzae LS1. Of the substrate tested, soluble starch (100%) was the most hydrolyzed by the enzyme. Corn starch (64.16 %), white dextrin (34.52 %), yellow dextrin (29.02 %) and potato starch (13%) were also hydrolyzed to a lesser extent. Our results are also coherent with (Elegado and Fujio, 1993) who observed that wheat and cassava starches are the most favoured substrates by all strains while potato and corn starches, the least.

In conclusion, the findings of the current study revealed that, Bacillus circulans being the most potential strain for maximum and cost effective alpha amylase enzyme production at appropriate conditions could be utilized as bio-inoculant in solid state fermentation as compared to submerged fermentation technology in sustainable manner.

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How to cite this article:
Nadia Razdan and Kocher, G.S. 2018. Isolation and Characterization of Bacteria for Amylase Production under Solid State Fermentation Using Damaged Wheat as Substrate. Int.J.Curr.Microbiol.App.Sci. 7(07): 946-955. doi: https://doi.org/10.20546/ijcmas.2018.707.114