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Immobilization of Microbial (Wild and Mutant Strains) Amylase on Coconut Fiber and Alginate Matrix for Enhanced Activity

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ABSTRACT

Reports are not available for immobilization of on α -amylase produced by Bacillus subtilis, Agrobacterium tumefaciens, Escherichia coli, Xanthomonas campestris, Staphylococcus aureus, Aspergillus niger, Aspergillus flavus, Metarhizium anisopliae, Azotobacter chroococcum and Rhizopus oryzae on coconut coir. Hence, an attempt was made in present study to immobilize the free amylase on coconut coir. Cell bound amylase of Bacillus subtilis, Escherichia coli, Xanthomonas campestris, Aspergillus niger, Aspergillus flavus and Rhizopus oryzae were immobilized by entrapment in calcium alginate. The immobilized enzyme on cells showed higher α-amylase production compared to free cells. The enzyme was characterized by Thin Layer Chromatography (TLC) and confirmed as alpha amylase. A simple and inexpensive method for immobilizing α -amylase on coconut fiber was also adopted. Highest amylase activity was observed in X. campestris on immobilized state on coconut fibers (1.708±0.103 U/min/g) and also in alginate matrix (0.948±0.082 U/min/100 beads). Hence, the cells of X. campestris were mutated with UV (for different time intervals: 1, 2, 4 and 8 min) and acridine orange (in different concentrations: 1, 2, 4 and 8 ug mL⁻¹). The immobilized catalyst of mutant strain of X. campestris AO₄ on alginate matrix exhibited 66.31% (4.125±0.056 U/min/100 beads) higher than that of the parent strain. It exhibited highest activity, when immobilized on coconut coir. Thus the mutant strain of X. campestris AO₄ could be used in industries for the efficient conversion of starch to oligos accharides.

Key words: Microbial α-amylase, immobilization on coconut fiber, mutant strains, Xanthomonas campestris, Metarhizium anisopliae

INTRODUCTION

Amylases constitute a class of industrial enzymes which alone form approximately 25% of the enzyme market covering many industrial processes such as sugar, textile, paper, brewing, distilling industries and pharmaceuticals. Although amylases can be derived from plants and animals, those obtained from bacterial and fungal sources generally meet industrial demands (Guzman-Maldonato and Paredes-Lopes, 1995; Rao et al., 2007; Ahmed et al., 2008; Djabali et al., 2009; Pascoal et al., 2010). Immobilization increases the stability and catalytic activity of amylases and the resistance of microorganisms against destroying factors. Industrial development of enzymatic reactors requires the use of immobilized enzymes in order to reduce the cost of the biocatalyst. To a large extent this procedure prevents enzyme losses due to washout and at the same time maintains biocatalyst at high concentrations. In previous works, α -amylase has been immobilized into a large variety of supports such as controlled pore glass, nylon, collagen, sephadex, sepharose and polyaminostyrene (Hornby and Goldstein, 1976; Dey et al., 2002; Guisan, 2006; Rao et al., 2007; Yugandhar et al., 2007; Delcheva et al., 2007; El-Banna et al., 2007; Pramod and Lingappa,

2008; Egwim and Oloyede, 2008; Alkhatib et al., 2010). Also the amylase was immobilized on nitrocellulose membrane and zirconium dynamic membrane (Guisan, 2006; Rao et al., 2007). However, reports on utilization of natural fiber such as coconut coir and drumstick fiber for immobilization of amylase are scarce (Dey et al., 2002). Coconut coir is a product of the coconut industry that has numerous industrial applications including automobile upholstery, bedding and mattresses, drainage filters, insulation, packaging, brush and broom manufacturing, reinforcement of thermoplastics, marine cordage and fishnets and in horticulture as a growing medium/soil substrate. Coir has water holding capacity and a cation-exchange capacity of 39-60 meq/100 g (Dey et al., 2002).

These properties and the fact that it is widely available make coconut fiber a potential hydrophilic support that may be used for immobilization of enzymes by physical adsorption. However, its application has been restricted to the mentioned applications and it has tested as an immobilization matrix for adsorption of enzyme by Dey and their colloquies (Dey et al., 2002). However, coconut fiber has been tested as an immobilization matrix only for adsorption of enzyme from Bacillus circulans GRS 313 (Dey et al., 2002). Thus, an attempt was made to test coconut fiber and drumstick fiber as adsorption matrices for immobilization of amylase from Bacillus subtilis, Escherichia coli, Xanthomonas Campestris, Aspergillus niger, Aspergillus flavus and Rhizopus oryzae. The catalytic behaviour of immobilized amylase from different sources will be discussed and compared with the free enzyme.

MATERIALS AND METHODS

Microorganisms used: Pure culture of Bacillus subtilis, Agrobacterium tumefaciens, Escherichia coli, Xanthomonas campestris, Staphylococcus aureus, Aspergillus niger, Aspergillus flavus, Metarhizium anisopliae, Azotobacter chroococcum and Rhizopus oryzae were purified from the mother culture of our laboratory. They were isolated from different sources using selective medium and identified by biochemical and microscopic analysis and stored in collection centre. The cultures were used as reference and sub-cultures were used as wild strains for the present study. The fungal strains were maintained on Potato Dextrose Agar (PDA) agar plates. E. coli, X. campestris, A. tumefaciens and A. chroococcum were maintained on Eosin Methylene Blue (EMB) agar, Luria agar, Agrobacterium agar and nitrogen-free agar, respectively. B. subtilis and S. aureus were grown in nutrient agar (Fig. 1a-g). Study was conducted from September 2008 to August 2010.

Primary screening of amylase producing microorganisms: Primary screening was done by starch agar plate method (Guisan, 2006). Strains of B. subtilis, E. coli, X. campestris, A. niger, A. flavus and R. oryzae showed maximum starch hydrolysis on starch agar medium. These six strains were selected for further investigation.

Preparation of immobilized cell bound enzyme using calcium alginate: Young fungal mycelia or bacterial cells from liquid cultures were centrifuged at 3000 rpm at 10°C for 8 min and the supernatant liquid was discarded. The pellet was resuspended with a previously autoclaved solution of sodium alginate (HiMedia, Mumbai, India). The solution had been adjusted to pH 7.0 before the autoclaving. Unless otherwise stated, a final concentration of 2% (w/v) sodium alginate and 10% (vol/vol) fungal biomass was used. The alginate-fungus mixture was added drop wise from a height of 20 to 50 mm with sterilized 10 mL glass pipette (nominal diameter, 2.0 mm) into an autoclaved solution of calcium chloride (normally 1%, adjusted to pH 7.0), where beads

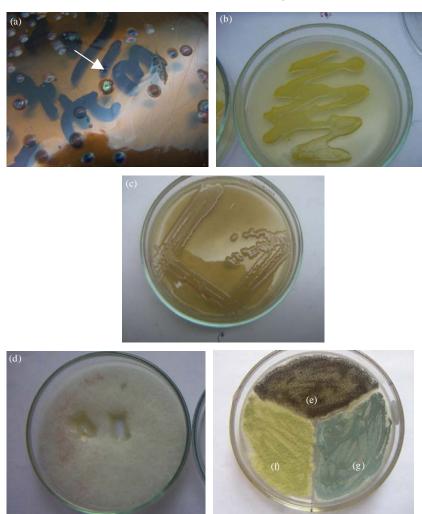


Fig. 1: Microorganisms used in the present study; (a) *E. coli* on EMB agar; (b) *X. campestris* on Luria agar; (c) *A. chroococcum* on nitrogen-free agar; (d) *R. oryzae* on PDA; (e) *A. niger* on PDA; (f) *A. flavus* on PDA and (g) *M. anisopliae* on PDA

formed instantaneously. The beads were left in this gelling solution for 5 min before being harvested by filtration through a 0.18 mm sieve, rinsed with sterilized physiological saline solution and chilled overnight at 4°C and comparative data of their activity was obtained. The preparation of control for immobilized cells was also carried out.

The enzyme production medium contained ammonium dihydrogen phosphate (0.02), manganese chloride (0.05), soluble starch (0.2), beef extract (0.02), distilled water (100 mL). The pH of the medium is adjusted to 6.5 and the flasks were kept on rotary shaker at 27±2°C (for fungi) or 37±2°C (for bacteria) for 72 h. Two sets of production medium were prepared. To one of it, entrapped gel beads of B. subtilis, E. coli, X. campestris, A. niger, A. flavus and R. oryzae were added to the medium. To the second, unimmobilized cells were added. Control was set up by the medium containing calcium alginate without cells. After incubation, the enzyme was assayed.

Amylase assay: Amylase was assayed by adding 0.2 mL of enzyme (fermented broth supernatant) to 0.5 mL of 1% soluble starch and incubated for 10 min at 37±1°C. The reaction was stopped by

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adding 0.75 mL of dinitrosalicylic acid reagent. The final volume was made to 5 mL with distilled water, tubes were kept in boiling water for 6 min, cooled and the absorbency measured at 575 nm with a UV-Visible spectrophotometer.

One unit (U) of amylase was defined as the amount of enzyme that liberates one Mole of reducing sugars, measured as reducing sugar per min under the conditions of assay.

Thin-Layer Chromatography (TLC) analysis: The hydrolyzed products of starch were submitted to TLC on silica gel plates for confirmation as alpha amylase. Enzymes (10 μL) were incubated with 1% of soluble starch (100 μL) in potassium buffer (pH 8.0) at 50°C for 1 h. Aliquots of the reaction mixtures were chromatographed on a silica gel [30 gm of silica gel in 61.5 mL of chloroform: methanol: conc. sulphuric acid (42: 18: 1.5)] with n-butanol, ethanol and water (5: 3: 2) as mobile phase. The products from the incubated content were detected by baking at 120±22°C for 30 min.

Preparation of immobilized enzyme on coconut fibers: The coconut fibers (1 cm) were separated and boiled in water containing 0.01% (w/v) sodium dodecyl sulfate for 1 h. The fibers were dried completely at room temperature. One milliliter of enzyme solution (fermented broth supernatant), along with 0.1 g of the fiber was placed in a shaking water bath at 30°C for a contact time of 2 h. After adsorption the enzyme solution was decanted and saved for subsequent assay. The unbound amylase was washed off with distilled water until no activity was detected in the washing.

Enzyme assay: The reaction mixture, containing 0.5 mL of 1% (w/v) soluble starch solution and 0.05 g of coconut fiber was incubated at 37±1°C in a water bath shaker. Adding 0.75 mL of dinitrosalicylic acid reagent stopped the reaction. The final volume was made to 5 mL with distilled water, tubes were kept in boiling water for 6 min, cooled and the absorbency measured at 575 nm. One unit is defined as the amount of amylase that produced 1 μmol of reducing sugar under assay conditions/g of fiber.

Preparation of mutant strains of *X. campestris*: *X. campestris* which showed maximum starch hydrolysis on starch agar medium and amylase activity on alginate matrix and coconut fibers was selected for further investigation. *X. campestris* cell suspensions (approximately 1.03×10² CFU mL⁻¹ in saline solution) were treated with UV (for different time intervals: 1, 2, 4 and 8 min) or with acridine orange (in different concentrations: 1, 2, 4 and 8 μg mL⁻¹). The treated suspensions were appropriately diluted and plated on to starch agar plates. Colonies, producing large clear zones around them were selected (Strain name-mutagen, *X. campestris* AO1-1 μg mL⁻¹, *X. campestris* AO2-2 μg mL⁻¹, *X. campestris* AO3-4 μg mL⁻¹, *X. campestris* BO4-8 μg mL⁻¹, *X. campestris* UV1-1 min, *X. campestris* UV2-2 min, *X. campestris* UV3-4 min and *X. campestris* UV4-8 min) and tested for amylase activity by culturing in a liquid medium and also by immobilization on alginate matrix, coconut coir and drumstick fiber.

Statistical analysis: All experiments were performed in triplicate and the data presented in this work are the mean value of three different experiments. Standard error was calculated by standard statistics (STATISTICA/w 5.0. software).

RESULTS AND DISCUSSION

Primary screening of amylase producing microorganisms: In the search for microorganisms possessing high amylolytic activity, Bacillus subtilis, Escherichia coli, Xanthomonas campestris, Aspergillus niger, Aspergillus flavus and Rhizopus oryzae exhibited higher amylolytic activity in starch agar medium (Fig. 2, 3a-h) and were selected for further studies. The type of amylase enzyme was identified as α -amylase by analysing the hydrolysed products (maltose and maltotriose) of starch in TLC (Palanivelu, 2001). Dhanasekaran and co-workers adapted the same method to identify the enzyme separated from Bacillus species using in thin layer chromatography. They observed that the Bacillus sp. what they have isolated produced maltose and maltotriose (Dhanasekaran et al., 2006).

The highest (22.00±1.50 mm) activity was observed in *X. campestris* on starch agar plates with clear zone was seen after flooding the plate with iodine as showed in Fig. 2. This result agreed with that of Konsoula *et al.* (2007) on the production of alpha amylase from *Xanthomonas campestris* ATCC 13951.

Amylase immobilization: Number of workers have reported that the addition of various compounds to the enzyme to increase its catalytic activity and stability by preventing spontaneous or heat induced denaturation of the enzymes (Mozhaev et al., 1989; Baldino et al., 2001; Gupta et al., 2003; El-Batal et al., 2005; El-Banna et al., 2007). Effective enzyme immobilization can be achieved using several techniques including adsorption to insoluble materials, entrapment in polymeric matrix encapxsulation, cross linking with a bifunctional reagent or covalent linking to an insoluble carrier (Kara et al., 2005). The most important advantages of immobilization methods are the stability of enzyme activity after immobilization and reuse of the enzyme and support material for different purposes because of reversibility of the method. Enzyme immobilization is a very powerful tool that may be considered to solve this cost problem and saving time (Hamilton et al., 1999; Akgoll and Denizli, 2004). On the other hand, industrial development of the enzyme reactors requires the use of immobilized enzyme in order to reduce the cost of the biocatalyst (Baldino et al., 2002).

In the present study, α-amylase produced by B. subtilis, E. coli, X. campestris, A. niger, A. flavus and R. oryzae was immobilized on different carriers (alginate matrix, coconut coir and drumstick fiber) as an attempt to assess the activity retained upon immobilization in comparison

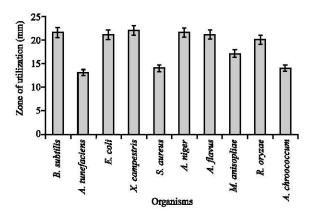


Fig. 2: Zone of starch hydrolysis by different microbes on the starch agar plate C

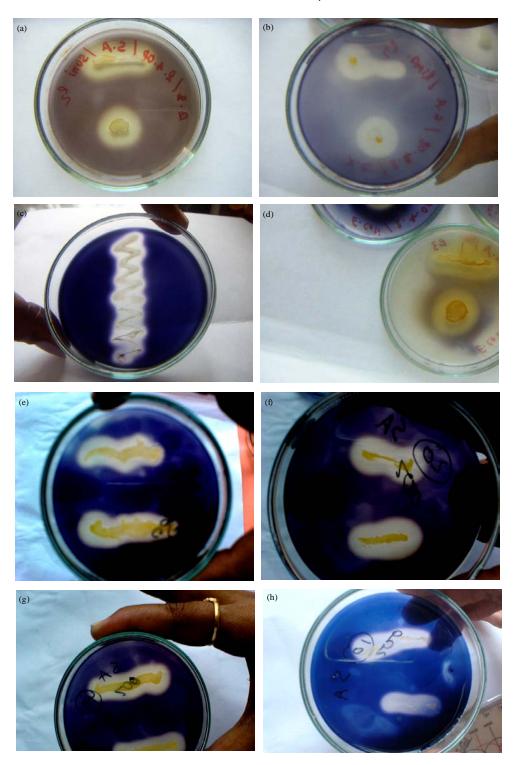


Fig. 3: Screening of the isolates for amylase production in starch agar plate; (a) B. subtilis, (b) X. campestris, (c) A. chroococcum, (d) E. coli, (e) X. campestris AO3, (f) X. campestris AO4, (g) X. campestris AO3 and (h) X. campestris AO2

to the free amylase and to investigate the operational stability of the immobilized enzyme when compare with the free enzyme. The efficiency of enzyme immobilization was evaluated according to the enzyme activity. Immobilization of amylase on alginate matrix showed that X. campestris α -amylase exhibited the highest enzyme activity (0.948±0.082 U/min/100 beads) with 2% calcium alginate. Highest amylase activity was also observed in X. campestris α -amylase on immobilized state on coconut fibers (1.708±0.103 U/min/g) (Fig. 4). On the other hand, the lowest enzyme activity (0.647±0.042 U/min/g) was exhibited by A. flavus α -amylase immobilized on coconut coir. The current study on the α -amylase from six strains immobilized on coconut coir agrees well with the results reported earlier for B. circulans GRS 313 amylase immobilized on coconut coir (Dey et al., 2002). However, all the α -amylase from six isolates showed lower enzyme activity when immobilized on drumstick fibers than that obtained with the other two substrates, alginate matrix and coconut coir (Fig. 4)

Mutagenesis of X. campestris: Among the organisms tested, X. campestris was the most potent strain for amylase production as it showed maximum starch hydrolysis on starch agar medium and amylase activity on alginate matrix, coconut fibers and drumstick fibers (Fig. 4). Thus X. campestris was mutated using ultraviolet irradiation for various time intervals and acridine orange at different concentrations to check its efficacy after mutation.

All the eight mutant strains of *X. campestris* were tested for amylase activity, α-amylase from eight mutant strains of *X. campestris* were analysed in free and in immobilized state on alginate matrix, coconut coir and drumstick fiber. Hyper productive mutants were obtained by these treatments. Under the optimized cultivation conditions, amylase activity of the mutant strain, *X. campestris* AO4 (1.7 μg mL⁻¹) was higher than that of the parent strain (0.52 μg mL⁻¹), *X. campestris* in liquid medium. Immobilized α-amylase from mutant strain, *X. campestris* AO4 on alginate matrix showed higher amylase activity (4.3 μg mL⁻¹) than that of the parent strain, *X. campestris* (0.9 μg mL⁻¹). However, the amylase activity for the immobilized α-amylase from mutant strain, *X. campestris* AO4 on coconut coir and drumstick fibers exhibited lower activity than that of the parent strain (Fig. 5).

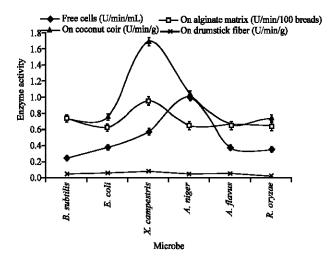


Fig. 4: Activity of free and immobilized α-amylase from bacteria and fungi. Data are given as Means±SD

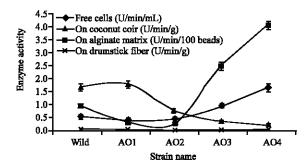


Fig. 5: Activity of free and immobilized α-amylase from mutated X. campestris using acridine orange. Data are given as Means±SD. (AO1-X. campestris 1 μg mL⁻¹; AO2-X. campestris 2 μg mL⁻¹; AO3-X. campestris 4 μg mL⁻¹; AO4-X. campestris 8 μg mL⁻¹)

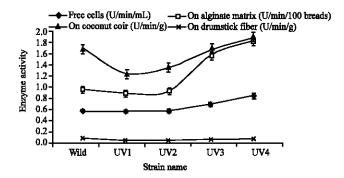


Fig. 6: Activity of free and immobilized α-amylase from mutated X. campestris using ultra violet light (UV1-X. campestris 1 min; UV2-X. campestris 2 min; UV3-X. campestris 4 min and UV4-X. campestris 4 min)

The amylase activity of immobilized α -amylase from mutant strains on alginate matrix declined slightly after 1 min of exposure to ultraviolet irradiation. But the enzyme activity increased slightly after 2 min of exposure to UV irradiation. However, the enzyme activity of immobilized α -amylase from mutant strains on alginate matrix continued to increase reaching the activity of 1.82 μg mL⁻¹ while α -amylase from parent strain (Fig. 6) reached the activity of 0.96 μg mL⁻¹ after 4 min of exposure to UV irradiation.

Thus using mutation, amylase activity was improved by as much as about 50-75% in several of the mutants when compared to the parental strain. This clearly suggests that mutation has enhanced amylase synthesis and secretion (Maruo and Tojo, 1985). A preliminary study with one of these hyper producers showed that thermostability of the α-amylase was maintained which suggests that mutation could have occurred in the regulatory gene (Hamilton et al., 1999; Akgoll and Denizli, 2004) rather than in the structural gene but this must await further investigation. The purpose of the mutation, to make a mutant by altering the regulatory part of the amylase gene was achieved, that can be said from increased production by mutated strains, but gene level studies are needed to confirm the mutation in the regulatory region of the amylase gene.

CONCLUSION

The X. campestris AO4 strain mutated in this study could be used in industries for the efficient conversion of locally produced starch to simple sugars. Furthermore, this mutation technique

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developed to obtain mutants with enhanced overall amylase production is interesting in that the process is simple, effective and economic. This merits consideration in future mutagenic studies, especially if such studies are being conducted in the underdeveloped countries where cost is a major factor.

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