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## Production of Tannase Enzyme by *Penicillium duclauxii*, Screened and Isolated from Soil.

Suman Lata<sup>1\*</sup>, and Km. Poonam Rani<sup>2\*</sup>.

<sup>1</sup>Department of Biotechnology, Meerut Institute of Engineering and Technology, Meerut, Uttar Pradesh, India.

<sup>2</sup>Student of Master of Technology in Biotechnology Department of Biotechnology Meerut Institute of Engineering and Technology, Meerut Uttar Pradesh, India.

### ABSTRACT

Tannase are enzymes that may be used in different industrial sectors as, for example, food and pharmaceutical. They are obtained mainly from microorganisms, as filamentous fungi. However, the diversity of fungi stays poorly explored for tannase production. Most of the research was focused on fungal tannase, as tannin was earlier considered as bacteriostatic. After the discovery of bacterial tannase in 1983, several studies on bacterial tannase were published. Despite the long history and numerous publications, tannase is still considered as one of the costly industrial enzymes. This is due to less titer and long fermentation time of the processes. In view of the growing demand, it is imperative to isolate high productive strains and develop economically feasible processes. A total of 19 samples collected from the tannin-rich soil. The isolated cultures were screened for their tannase producing capability by observing the zone of hydrolysis on tannic acid agar plates. Among the fungal strains selected as tannase producers, the isolate S4RD4 showed largest zone of clearance of 30 mm (diameter) on tannic acid agar plate. Hence, it was selected for further study and identified as *Penicillium duclauxii* (NFCCI, PUNE, INDIA). To enhance the production level of the enzyme different culture conditions were optimized and observed that optimum temperature and pH for tannase production was 30°C and 5.5 respectively. Maximum growth and enzyme production was recorded after 96 hrs of incubation period in the medium (B-modified synthetic medium) containing 1% tannic acid. Malt extract (2%) with NaNO<sub>3</sub> (0.2%) was found to be the best nitrogen source and sucrose found to be a best carbon source for tannase enzyme production. Among the additives, metal ions Cu<sup>2+</sup>, Mg<sup>2+</sup>, Mn<sup>2+</sup>, CO<sup>2+</sup>, K<sup>+</sup>, Ca<sup>2+</sup>, Fe<sup>3+</sup> affected enzyme production. Tween-20, Tween-40, Tween-60, Tween-80, Triton X-100 and SDS detergents tested, these detergents inhibited the production of tannase. The optimization of culture conditions enhanced the production level of tannase (33.41 U/ml) by (1.36) fold. This study reviews the microbial sources, isolation and screening methods, modes of production, substrates and media, temperature and pH of fermentation, duration of fermentation and location of tannase enzyme.

**Keywords:** Tannase enzyme, Gallic acid, Tannic acid, Identification, Tannase Assay.

\*Corresponding author

## INTRODUCTION

The enzyme tannase (E.C. 3.1.1.20) also known as tannin acyl hydrolase, is a hydrolytic enzyme that acts on tannin. It catalyses the hydrolysis of bonds present in the molecules of hydrolysable tannins and gallic acid esters producing gallic acid and glucose [17]. Tannase is an inducible enzyme produced by variety of microorganisms such as fungi, bacteria and yeast [1]. Microorganisms have been the most important source for the production of industrial enzymes due to their biochemical diversity and their technical and economic advantages [4].

Tannins are naturally occurring plant phenolics compounds that have wide ranging effects on animals and microbes. They are polyphenolic secondary metabolites of plants which form hydrogen bonds in solutions, resulting in the formation of tannin-protein complexes. Tannins are present in large number of feed and forages. The formation of complexes of tannins with nutrients, such as carbohydrates, proteins and minerals, has negative effects on their utilization. High concentrations of tannins depress voluntary feed intake and digestive efficiency [3]. The nutrient value of tanniferous feed may be enhanced by various detannification procedures viz., physical, chemical and biological. In biological treatment, various tannase producing microbial strains have been tried for reduction of tannin content and nutritive enhancement of treated material [64]. In this respect, tannase find potential applications in feed, food and beverage industry. Tannase is used as clarifying agent in some wines, juices of fruits and refreshing drinks with coffee flavor [5,12]. The use of tannase helps in overcoming the problem of undesirable turbidity in these drinks which poses the quality problem. Enzymatic treatment of fruit juices reduces bitterness, haze and sediment formation, hence are acclaimed for health benefits and industrial use. Tannase is also being used for production of instant tea preparations. The enzyme has potential uses in treatment of tannery effluents and pretreatment of tannin containing animal feed [2, 54].

One of the major application of tannase is the production of gallic acid. Gallic acid is used for the manufacture of an anti-malarial drug Trimethoprim. Gallic acid is a substrate for the chemical and enzymatic synthesis of propyl gallate, used as anti-oxidants in fats and oils. Applications of tannase include removal of haze formation of beer and wines and detannification of poultry feed to improve the feed efficiency [2].

It is present as intracellular or extracellular enzyme produced by several micro-organisms and plant cells. Tannase has been isolated from number of micro-organisms like fungi, bacteria and yeast. Many fungal species have been reported to produce tannase, including *Aspergillus aculeatus*, *A. aureus*, *A. flavus*, *A. foetidus*, *A. japonicas*, *A. niger*, *A. oryzae*, *Aureobasidium pullulans*, *Fusarium solani*, *F. subglutinans*, *Paecilomyces variotii*, *Penicillium atramentosum*; *P. chrysogenum*, *P. variable*, and *R. oryzae* reviewed by Belur and Mugeraya and Chavez-Gonzalez. The vast majority produce in submerged cultures, while *Aspergillus* and *Penicillium* are the most active microorganisms capable of producing tannase through both submerged and solid state fermentations [19, 10]. The fungal species *Aspergillus* and *Penicillium* are the most active microorganisms capable of producing tannase through submerged and solid state fermentation.

In the last decade, there have been a number of efforts to improve the production, recovery, and purification processes of the enzyme. These efforts include the looking for new sources for tannase enzyme production [5,6], the development of novel fermentation systems [7,8], the optimization of culture conditions [9,10], the production of the enzyme by recombinant microorganism [11,12], the design of efficient protocols for tannase recovery and purification [11]. Technological advances on tannase processing must be supported by basic investigation. Physicochemical properties of several tannase have been characterized [12,13], there have been a special interest in the description of tannase and tannase gene structure [14].

With the rapid advancement in the field of genetic and protein engineering enzymes have found their way into many new industrial processes. Enzymes offers biological alternative to the chemical processes at industrial scale. Biological methods are an important mechanism of organic chemical removal in the natural systems [3].

In view of the growing demand, it is important to isolate high productive strains and develop economically feasible processes. With this view studies on isolation, screening and production of tannase from *Penicillium duclauxii*, NFCCI, PUNE, INDIA by submerged fermentation technique was carried out.

## MATERIAL AND METHODS

### Micro-Organism

#### Fungal strain

The tannase producing fungal strain used in the present investigation was isolated from soil sample collected from Ghaziabad district (UP). The culture was maintained on malt extract medium which consist of malt extract (2%), K<sub>2</sub>HPO<sub>4</sub> (0.1%), NH<sub>4</sub>Cl (2%) and Agar (2%). For enzyme production, this medium was supplemented with 1% tannic acid. The culture was maintained on malt extract agar slant at 4°C and subcultured twice in a month. The strain was identified as *Penicilliumduclauxii* by NFCCI, Pune, INDIA.

#### Collection of soil samples

Nineteen different samples were collected from various locations such as from Meerut, Garh, Delhi, Ghaziabad, by digging the soil up to 5-6 inches deep from the soil surface. Samples were collected (in sterile polythene bags and stored at 4°C until use) from different litter sites such as soil containing decaying material, soil from agricultural field, soil receiving kitchen wastes, soil receiving tea wastes, soil receiving domestic wastes, garden soil etc.

#### Serial dilution technique

In the present investigation, for isolation of tannase producing microorganisms, the soil sample was plated by serial dilution method (Figure 2). Ten gram of soil was suspended in 90 ml of sterile distilled water and shaken well. The sample was serially diluted through a series of dilution to obtain a final concentration of 10<sup>-5</sup>. From each dilution, 200 µl was pipette out and spread on the surface of sterile solid agar plate supplemented with 1% tannic acid.

#### Sprinkling method

A new method for soil sample plating has been used, in which 1 to 2 milligram (mg) of soil sample was simply sprinkled over the tannic acid solid agar plates (supplemented with 1% tannic acid). The plates were incubated at 30°C for 96 hrs. After, different fungal colonies appeared on the plates. These fungal colonies were isolated and purified. They were screened for tannase producing capability qualitatively.

#### Qualitative screening of tannase producing fungi

The tannase producing fungal isolates were screened by observing zone of hydrolysis around colonies. The isolates producing zone around its growth on malt extract medium agar plates (supplemented with 1% tannic acid) were selected as tannase producers. FeCl<sub>3</sub> testing was used for the purpose of screening the potent tannase producer fungi [29]. FeCl<sub>3</sub> reacts with tannic acid present in the medium and produces deep brown colour. The solid agar tannic acid plates with isolated fungal colonies (point inoculated and incubated for 72 hrs) were flooded with FeCl<sub>3</sub> solution. The positive fungal strain hydrolyzed the tannic acid present in the surrounding medium and shown a zone of clearance around the colony. [37].

#### Quantitative estimation of tannase

#### Tannase production

The selected fungal isolate was used for extracellular tannase production in modified malt extract liquid medium. The Erlenmeyer flask containing medium was inoculated with discs (8 mm diameter) of 48 hr grown fungal culture. The flask was incubated at 30°C for 3-5 days. After incubation, the fermentation broth was filtered through Whatman filter paper (No.1) [36]. The tannase activity from the culture supernatant (crude enzyme) was determined by performing tannase assay.

### **Tannase assay**

Tannase was assayed following the method of Mondalet *al.* (2006) using tannic acid (1%) as substrate, prepared in acetate buffer (0.2 M, pH 5.5). The reaction mixture was prepared by the addition of 0.5 ml substrate with 0.1 ml of crude enzyme and incubated at 40°C for 20 min. The enzymatic reaction was stopped by adding 3 ml Bovine serum albumin (BSA) (1 mg/ml). The tubes were centrifuged at 5000 x g for 10 min. The precipitate was dissolved in 2 ml SDS-triethanolamine solution followed by the addition of 0.5 ml FeCl<sub>3</sub> reagent. The contents were kept for 15 min. for stabilizing the colour formed and the absorbance was measured at 530 nm against the blank. One unit of tannase activity can be defined as the amount of enzyme which is able to hydrolyze 1 μmol of substrate tannic acid in 1 min under assay conditions.

### **Optimization of culture conditions for tannase production**

The following culture conditions were optimized to increase the maximum production of tannase from *Penicillium duclauxii*.

#### **Effect of different medium on enzyme production**

To obtain maximum tannase production, different production medium (pH 5.5) such as medium A (Synthetic medium for tannase production), B (Synthetic medium for tannase production), C (Czapek Dox's medium), D (Modified Malt extract Medium) were tested. The composition of these medium was in g/L. The flasks of different medium were inoculated with 72 hrs grown culture of *Penicillium duclauxii* and incubated at 30°C for 5 days and growth as well as tannase activity was determined.

#### **Effect of incubation period on enzyme production**

To evaluate the effect of different incubation period on tannase production, the incubation period of the medium range was varied from 24 hrs to 120 hrs. To study the optimal incubation period for maximum tannase production, the flasks containing production medium B (pH 5.5) were inoculated at 30°C.

#### **Effect of pH on enzyme production**

The effect of initial pH of production medium for tannase production was studied. Flask containing production medium B was adjusted at various pH ranges from 4.5, 5.0, 5.5, 6.0, 6.5 by 1N HCl and 1N NaOH solution and were inoculated at 30°C and assayed for tannase activity as described earlier.

#### **Effect of temperature on enzyme production**

To study the effect of different temperatures on tannase production, the flasks containing medium kept at temperature range was varied from 30-50°C. To study the effect of incubation temperature for maximum tannase production, the flasks with the production medium were inoculated at various temperatures such as 30, 35, 40, 45, 50, 55, 50°C for 72 hrs. The general procedure mentioned earlier was followed for tannase assay.

#### **Effect of carbon source on enzyme production**

To study the effect of different carbon source on tannase production from *penicillium duclauxii*, the medium was supplemented with different sugar (maltose, sucrose, mannose, dextrose, lactose) with concentration 1.0 % (w/v). The production medium (medium B) with 1% tannic acid was kept as control. These flasks were inoculated with fungal culture and incubated at 30°C for 96 hrs and tannase activity was estimated.

#### **Effect of nitrogen source on enzyme production**

To study the different nitrogen sources on tannase production with various organic such as yeast extract, malt extract, beef extract, peptone and inorganic nitrogenous sources such as NaNO<sub>3</sub>, NH<sub>4</sub>Cl, KNO<sub>3</sub>, NH<sub>4</sub>NO<sub>3</sub> and these flask with medium B were inoculated with *penicillium duclauxii* and incubated at 30°C for 3 days and tannase activity was estimated.

### Effect of additives (metals ions and detergents)

The effect of different additives on tannase production in the production medium B was studied by adding different salts such as MgCl<sub>2</sub>, KCl, CuSO<sub>4</sub>, FeCl<sub>3</sub>, CoCl<sub>2</sub> and detergent such as Tween-20, Tween-40, Triton X-100 and SDS inoculated with *penicillium duclauxii* incubated at 30°C. After incubation, filtered the fermentation broth and tannase activity in culture supernatant was determined by performing tannase assay.

### Tannase production under optimized culture conditions

The medium optimized at this stage for tannase production by from *Penicillium duclauxii* was termed as Tannase Production Medium and was used in further studies. The composition or production medium and optimized conditions for tannase production by *Penicillium duclauxii* were also studied.

## RESULTS AND DISCUSSION

### Isolation and screening of tannase producing fungal strains

#### Isolation of tannase producing fungal strains

In the present study, isolation of fungal tannase producers was carried out from nineteen different soil samples. Total eighty six fungal strains were isolated on tannic acid agar plates by using serial dilution technique. On the basis of clear zone formation around the growth of fungal culture, fifty nine fungal isolates showing tannase activity on solid plate (Table 1). Out of fifty nine, only nine fungal isolates were selected as potent tannase producers on the basis of diameter of zone of clearance (Table 1). Further, these cultures were purified and maintained on malt extract agar medium. The fungal species of genus *Aspergillus* and *Penicillium* are the most active micro-organisms capable of producing tannase through submerged and solid state fermentation [19]. There are few reports on tannase production from *Rhizopus* sp. [31,24] and *Fusarium* sp. [23,28]. However, *Trichoderma* sp. is the least explored fungi for tannase production and characterization. *Trichoderma* sp. has been reported as tannase producers [23, 28].

#### Qualitative screening of tannase producing fungi

The selection of promising fungal isolates for tannase production was carried out using screening method. The isolates producing zone around its growth on malt extract medium agar plates (supplemented with 1% tannic acid) were selected as tannase producers. FeCl<sub>3</sub> testing was used for screening the potent tannase producer fungi. Selection of the strains was done on the basis of the degree of zone of clearance around colonies which was due to the tannase activity in the medium. Culture exhibiting zone of clearance after removal of the FeCl<sub>3</sub> were confirmed as tannase producers. The results obtained from this study showed that zone of clearance produced by different fungal isolates vary from 9 mm to 30 mm (Table 1). The isolate S4RD4 showed largest zone of clearance of 30 mm (diameter) on tannic acid agar plate (Fig. 2). The isolate S4RD4 produced tannase activity of (24.68 U/ml). It was identified as *Penicillium duclauxii* by National Fungal Culture Collection of India (NFCCI), PUNE, INDIA.

#### Optimization of culture conditions for maximum tannase production

To study, the optimization of culture conditions for maximum tannase production was carried out using 'one-factor-at-a-time' approach, by changing one control variable at a time while holding the rest parameters constant. The different parameters optimized were as follows:

#### Screening of different medium for tannase production

Four production media such as, Synthetic medium A, Synthetic medium B, medium C (CzapekDox's medium), medium D (Modified Malt extract Medium) was tested in an attempt to improve the tannase production from *Penicillium duclauxii*. Among the different medium tested, growth as well as enzyme production (24.39 U/ml) was found to be maximum in medium B (Synthetic medium supplemented with 1% tannic acid), followed with medium A and C (Fig.3). The minimum enzyme production was observed in medium

D. Several fungal species belonging to genera *Aspergillus* and *Penicillium* such as *Penicillium Charlesi*, *Penicillium variable*, *Aspergillus fumigatus*, *Aspergillus versicolor*, *Aspergillus flavus*, *Aspergillus caespitosum*, *Penicillium Crustosum*, and *Penicillium restrictum* which exhibits tannase producing activity on synthetic medium containing 1% tannic acid [23]. Maximum tannase production from *Paecilomyces variotii* and *Aspergillus foetidus* using synthetic medium were reported by [57, 50]. Many researchers used malt extract broth (supplemented with 1% tannic acid) for maximum tannase production from fungi [41].

#### **Production of Tannase Time course of tannase production**

To determine the optimum incubation time required for maximum tannase production from *Penicillium duclauxii*, the flask was inoculated with fungal culture and incubated at 30°C. Samples were withdrawn at regular intervals of 24 hrs for estimation of tannase activity. The results shown that the tannase activity was initially detected at 48 hr. and increasing with the time. Maximum growth and yield of tannase (27.12 U/ml) was obtained after 96 hrs of incubation (Fig. 4). Decreased enzyme yield on prolonged incubation could also be due to reduced nutrient level of medium. It has been reported that tannase activity decreased after reaching maximum level, due to inhibition of enzyme or degradation of enzyme [64]. Similar observations have been reported for tannase enzyme production by many investigators [28, 8, 29, 14]. There are various reports to study different incubation period for maximal extracellular production of tannase production by fungi. Maximum extra-cellular tannase production by *Aspergillus niger* and *Rhizopus oryzae* at 96 hrs and 120 hrs of incubation respectively [12]. Maximum production of extracellular tannase found in *A. aculeatus* after 72 hrs [27]. However, maximum tannase production from *Trichoderma viride* was reported at 48 hrs [33].

#### **Effect of pH of the culture medium**

The production of tannase enzyme from fungi strongly depends on the extracellular pH because culture pH strongly influences many enzymatic processes and transport of various components across the cell membranes which in turn support the cell growth and product production (45). The pH of medium was adjusted at different pH range (4.5-6.5). Maximum tannase production (29.86 U/ml) from *Penicillium duclauxii* was observed at pH 5.5 (Figure 5). The growth and enzyme production was however, considerable at pH 5.5-6.0. Further increase in pH of production medium, tannase enzyme production was found to decrease at pH 6.5. There are reports describing of the optimum pH as 5.5 for tannase production [27, 13, 14, 33, 23, 19]. Furthermore, the optimal initial medium pH for tannase production by submerged fermentation has also been reported in acidic range of 4.0-4.5 [1, 24]. Tannase production by *Aspergillus niger* and *Aspergillus oryzae* at pH 4.5 respectively, while some researchers reported maximum tannase production at pH 5.0 from *Aspergillus oryzae* [33, 8, 44]. The initial pH of medium 6.0 has also been reported for tannase production from fungus such as *Aspergillus awamori*, *Aspergillus japonicus*, *Aspergillus niger* respectively [23]. Some fungus also produced tannase at pH ranges from 6.0-6.5 [51, 54].

#### **Effect of incubation temperature**

The effect of incubation temperature on tannase production from *Penicillium duclauxii* was studied in the temperature range of 30°C-50°C. The optimum incubation temperature for tannase enzyme production (29.87 U/ml) from fungus was observed at 30°C (Fig. 6). Further rise in temperature, decreased the production of tannase and the minimum tannase activity was observed at temperature 40°C. The fermentation temperature for optimum production of tannase is mostly reported to be 30°C [34, 38, 8, 45, 33, 44]. Some investigators also mentioned tannase production from fungi at temperature 35°C-40°C [45, 24, 35]. Furthermore researcher also focused on tannase production from thermophilic fungi at temperature 50°C-60°C [45, 62, 27].

#### **Effect of substrate concentration**

To determine the optimum concentration of substrate for production of tannase, the flasks of production medium containing different concentrations of tannic acid ranging from 0.5 - 2.0 % were investigated. The maximum tannase activity of 30.42 U/ml was observed at 1.0% concentration of tannic acid in the production medium (Fig. 7). Tannase activity was affected by concentration of substrate in the medium. Maximum tannase activity was observed at a concentration of 1% (w/v) tannic acid used as inducer. Similar observations have been reported that tannase production from *Aspergillus ochraceus* [1] and *Aspergillus*



*niger*[12]. It was also observed that 2% tannic acid was suitable for tannase production, maximum extracellular tannase and gallic acid produced after 36 hrs in liquid submerged fermentation containing 2% tannic acid[27].

### Effect of carbon source

Tannase enzyme production depends on the availability of carbon sources in the medium and carbon sources have been shown to have regulatory effects on tannase enzyme synthesis [23]. To investigate the effect of carbon source on tannase production from *Penicillium duclauxii* was studied (Fig.8). The flasks containing production medium of different carbon sources such as sucrose, maltose, mannose, dextrose, and lactose were inoculated with fungal spores and incubated for 96 hrs at 30°C. The maximum tannase enzyme production was observed with sucrose (33.41 U/ml) followed by maltose (19.44 U/ml) and then with mannose (18.87 U/ml). These results are in agreement with those reported for tannase production from different microorganisms earlier such as *Aspergillus oryzae*[9,38,55] and *Penicillium atramentosum*[36]. Sucrose on degradation will act as a better carbon source for tannase production from *Aspergillus terreus*[40,27].

### Effect of different nitrogen sources

Tannase production depends on the availability of nitrogen sources in the medium. It has been shown to have regulatory effects on enzyme synthesis [23]. The effect of nitrogen source on tannase production from *Penicillium duclauxii* was studied by supplementing the production medium with various organic (2%) such as yeast extract, malt extract, beef extract, peptone and inorganic (0.2%) nitrogenous sources such as NaNO<sub>3</sub>, NH<sub>4</sub>Cl, KNO<sub>3</sub>, NH<sub>4</sub>NO<sub>3</sub>. (Fig.9) However, enzyme production in the presence of sodium nitrate (36.67 U/ml) followed ammonium chloride (24.32 U/ml). Similar results were reported by [27]. Also investigated the effect of different nitrogen sources on tannin degradation by *Aspergillus awamori* and found that nitrogen sources supported tannase production better [54]. Among the inorganic sources sodium nitrate was found to be best [61] obtained results for tannase production and concluded that microorganisms necessitate high level of nitrogen in order to produce enzymes. The effect of supplementation of different inorganic and organic nitrogen sources on tannase production. The organic nitrogen sources such as yeast extract, and malt extract gave considerable enzyme production [63].

### Effect of additives (metals ions and detergents)

The effect of additives (metals ions and detergents) on tannase enzyme production from *Penicillium duclauxii* was studied by adding different salts (0.2% w/v) and detergents (0.1% v/v) in the production medium B. After incubation, samples were withdrawn for analysis and tannase activity was determined. The metal ions Ca<sup>2+</sup>, Na<sup>+</sup>, Mg<sup>2+</sup>, Mn<sup>2+</sup> and Zn<sup>2+</sup> did not affect enzyme production. The maximum tannase enzyme activity was observed in presence of Mg<sup>2+</sup> (30.09 U/ml), followed by K<sup>+</sup> (29.89 U/ml). However, metal ions like Cu<sup>2+</sup>, Co<sup>2+</sup>, Fe<sup>3+</sup> and detergents (Tween-20, Tween-40, Tween-60, Tween-80 and SDS) inhibited tannase enzyme activity (Fig.10 and 11). On the other hand, detergent such as Triton X-100 did not inhibit the tannase enzyme production (29.23 U/ml). Karet *al.* (2003) reported highest tannase production by *Aspergillus foetidus* and *Rhizopus oryzae* in the presence of Mg<sup>2+</sup> in the medium. CaCl<sub>2</sub> to be the most suitable for tannase production by *Aspergillus japonicus* as compared to the other used mineral salts (NaCl, KCl, CuSO<sub>4</sub>, MnSO<sub>4</sub>, FeSO<sub>4</sub>, ZnSO<sub>4</sub> and CdSO<sub>4</sub>) [1]. also studied the effect of different mineral salts on tannase production and found CaCl<sub>2</sub> enhanced both tannase production and tannin degradation by the fungal strains [57,58]. The effect of some metal ions and common chemicals on tannase activity and observed that the enzyme was inhibited by all metal ions tested except Mg<sup>2+</sup> [1,8]. The results indicated that tannase production was stimulated on addition of combination of Mg<sup>2+</sup>, Ca<sup>2+</sup>, Mn<sup>2+</sup> and Zn<sup>2+</sup> in the fermentation media. On the hand, metal ions such as Ba<sup>2+</sup>, Co<sup>2+</sup> and Fe<sup>2+</sup> inhibited tannase biosynthesis from *Aspergillus niger*. [29] reported that tannase production from *Aspergillus niger*, the enzyme showed good stability in presence of detergents such as Tween 20, 80, Triton X-100 and SDS).

Table 1: Isolation and screening of tannase producing fungal cultures

Sample No.	Sample collected	Total no. of Tannase producers	Code of tannase isolates	Zone of clearance (mm)
S1	Mohiddinpur, Meerut (soil from root of jamun tree)	4	S1V1 (brown) S1V2 (white) S1v3 (light green) S1v4v (muddy yellow)	9 17 7 6
S2	Agriculture field area, Meerut (farming soil)	–	–	–
S3	Aminagar, Partapur, Meerut (garden soil).	–	–	–
S4	R.D foundation group of institution, Ghaziabad (garden soil)	4	S4RD1 (black) S4RD2 (white) S4RD3 (green) <b>S4RD4 (light green)</b>	12 7 8 <b>30</b>
S5	Girls hostel (MIET Campus) Meerut, garden soil	3	S5M1 (gray) S5M2 (brown) S5M3 (black)	6 19 9
S6	Puranakila, New-Delhi (garbage site soil)	3	S6PK1 (brown) S6PK2 (green with white ring)	8 9 7
S7	Patiala house court, New Delhi (garden soil)	4	S7PH1 (black) S7PH2 (green with white ring) S7PH3 (dark green) S7PH4 (white)	16 8 10 8
S8	Subharti Hospital Meerut (garbage site)	2	S8SH1 (white) S8SH2 (green with light gray ring)	6 7
S9	Kitchen waste soil sample, Meerut area, Meerut	4	S9KW1 (yellow) S9KW2 (gray) S9KW3 (white) S9KW4 (brown with white ring)	6 7 15 10
S10	Meerut, (soil enriched with guava)	4	S10GT1 (brown with white ring) S10GT2 (white) S10GT3 (yellow) S10GT4 (green)	<b>23</b> 10 6 7
S11	Sugar mill, Mohiddinpur, Meerut (soil from garbage site)	3	S11SM1 (gray with white ring) S11SM2 (yellow with white ring) S11SM3 (green)	18 6 9
S12	Soil from Animal waste, rural area of Meerut	3	S12AW1 (green with light green ring) S12AW2 (yellow) S12AW3 (white)	8 7 19
S13	MIET, Meerut (humus waste soil sample)	4	S13HA1 (green with white ring) S13HA2 (black) S13HA3 (white) S13HA4 (light yellow)	7 11 10 06
S14	T.C Health care, Modinagar (garden soil).	3	S14TC1 (green) S14TC2 (dark brown with light brown ring) S14TC3 (light green)	07 14 10
S15	Padamshri dairy, Modinagar (dairy waste soil)	2	S15DW1 (light brown with white ring) S15DW2 (light yellow)	18 7
S16	Mahrolli village, Meerut (soil enriched with imli)	5	S16IT1 (black) S16IT2 (brown) S15IT3 (green) S15IT4 (white) S15IT5 (gray)	19 10 7 7 8
S17	Soil from legume tree, Meerut (root soil)	4	S17LT1 (white) S17LT2 (yellow with white ring) S17LT3 (black)	9 7 10



			S17LT4 (gray)	8
S18	Partapur hostels area, Meerut (kitchen soil)	4	S18PHA1 (dark green) S18PHA2 (white) S18PHA3 (light green with white ring) S18PHA4 (blue with white ring) S18PHA5 (brown) S18PHA6 (gray)	7 10 8 10 17 8
S19	Heranpur village, Gardh Ganga (agriculture soil)	3	S19GJ1 (black with white ring) S19GJ2 (light green) S19GJ3 (White)	11 7 9

**Comparison of initial and optimized culture conditions for tannase production from *Penicilliumduclauxii***

To improve the production of tannase from *Penicilliumduclauxii*, various culture conditions were optimized including incubation time, incubation temperature, and initial pH of the medium, carbon source and nitrogen source. The optimization of various nutritional and cultural conditions resulted in an increase in the level of tannase production to 1.36-fold as compared to initial conditions (Fig. 12). The results of the time-course study before and after optimization of culture conditions are shown in Figure 2. Under optimized culture conditions such as malt extract medium with 1% carbon source, 4 days incubation at 30°C and pH 5.5, the production level of enzyme increased compared to initial unoptimized conditions (malt extract medium with 1% tannic acid as carbon source, 5 days incubation at 37°C and pH 5.5).

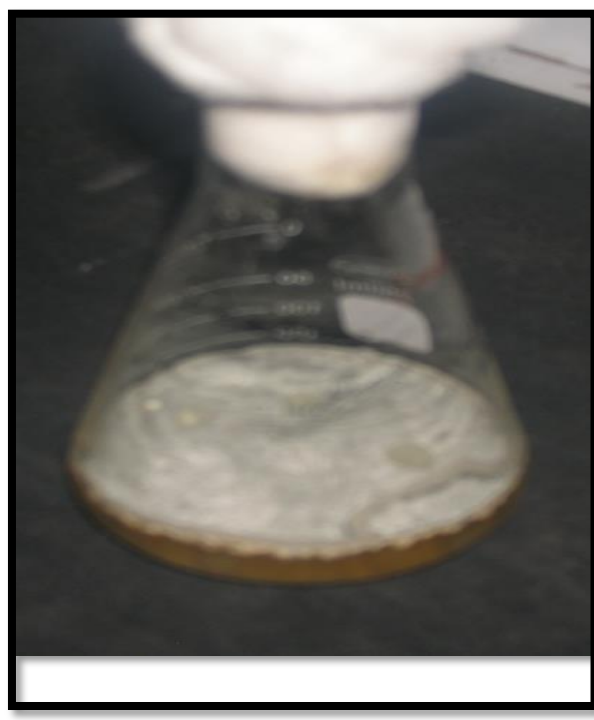


Figure 1: Tannase production (Quantitative estimation)



Figure 2: Fungal culture (S4RD4) showing tannase activity on tannic acid agar plate, (identified as *Penicilliumduclauxii*)

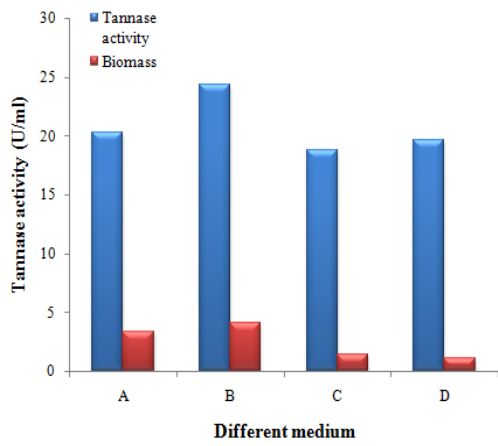


Figure 3: Effect of different medium production

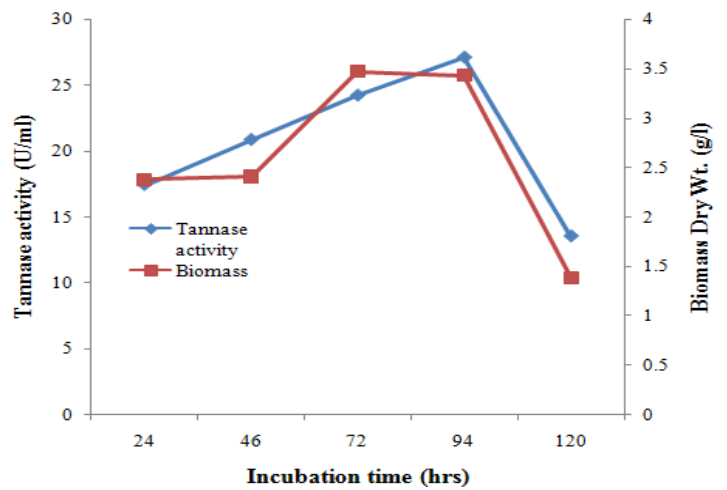


Figure 4: Different time course of tannase on tannase for maximum tannase production

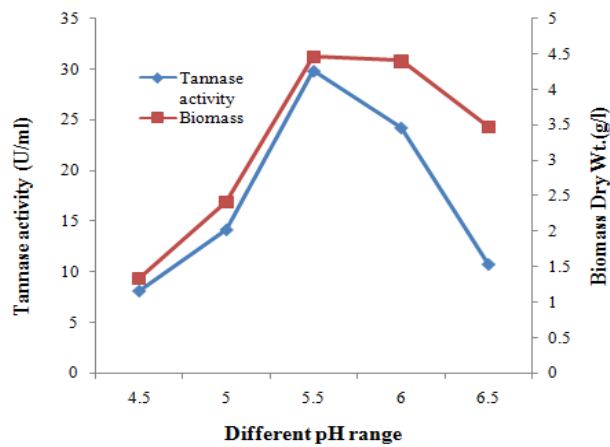


Figure 5: Effect of different pH

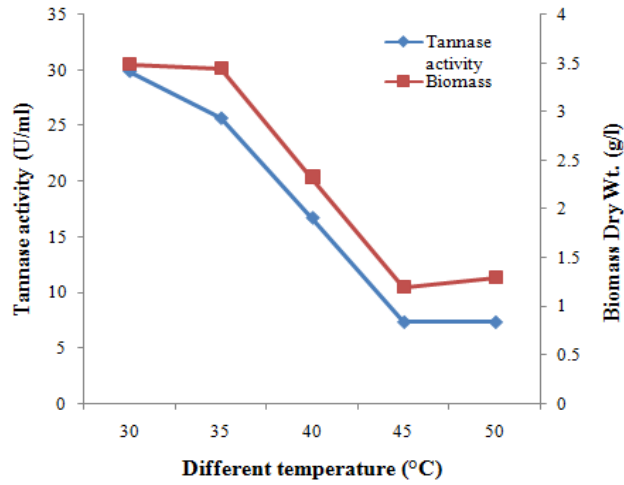


Figure6:Effect of different incubation temperature

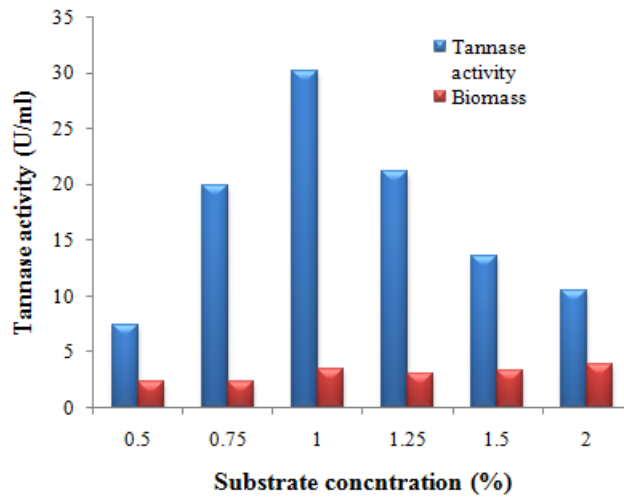


Figure 7:Effect of Substrate concentration on tannase production

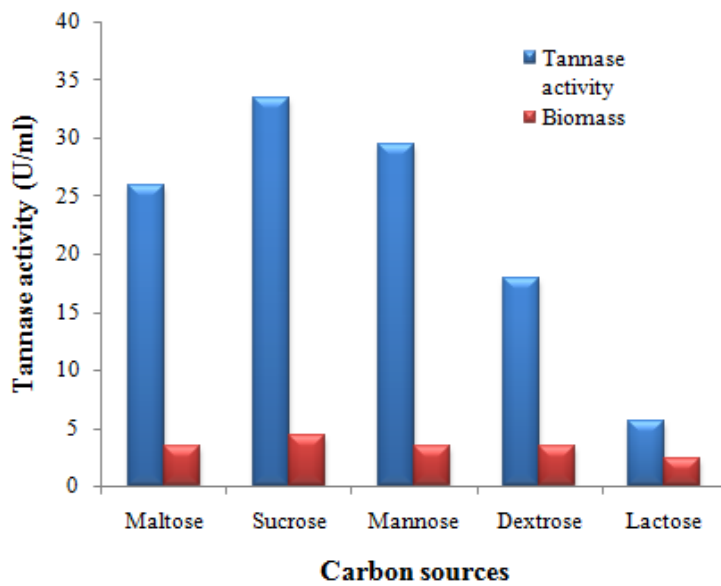


Figure 8: Effect of different carbon sources

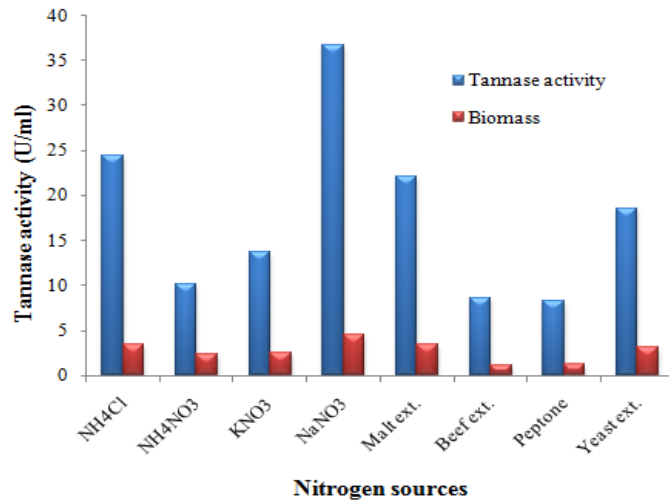


Figure 9: Effect of different nitrogen sources

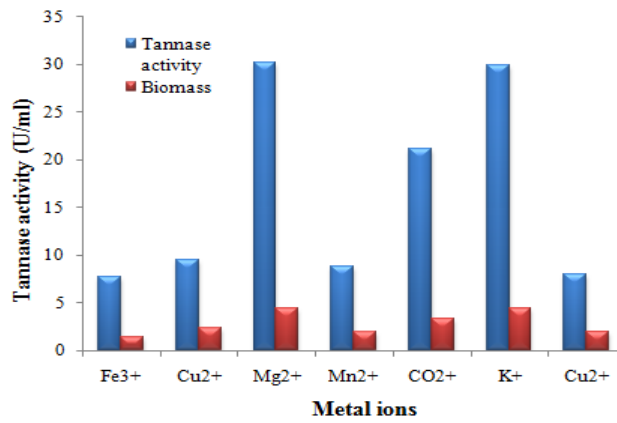


Figure 10: Effect of different metal ions

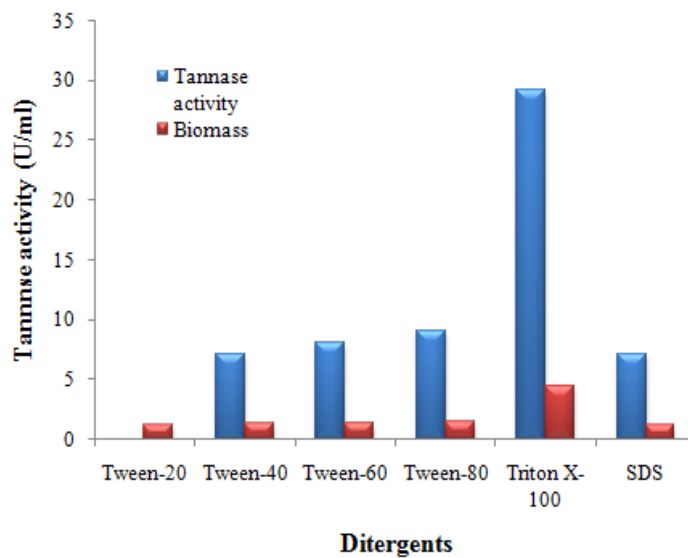


Figure 11: Effect of different salts

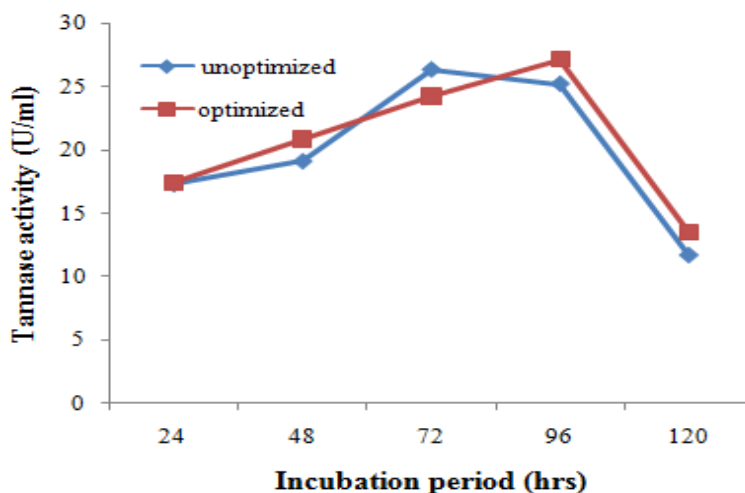


Figure 12: Tannase production from *Penicillium duclauxii* under optimized and unoptimized culture conditions.

### CONCLUSION

Tannase has the potential for a wide range of application, but due to higher production cost & lower yield, they currently have limited uses. This review article suggests the type of fermentation method & optimum conditions for the tannase production. Generalization of the conditions cannot be done for the production of tannase as it varies from organism to organism. In view of growing demand for the tannase for industrial application, it is very important to develop a high yielding & cost-effective process. Some natural tannin sources proved to be better substrates than commercial tannic acid for production of tannase. These agro-residues substrates can be substituted for costly tannic acid in the production medium for economic production of the enzyme at commercial level. Thus, there is a growing interest on basic and applied aspects of tannase.

### REFERENCES

- [1] Arulpani I, Sangeetha R, and Kalaichelvan PT. *Zaffius Biotechnology*, 2008; 3:11-101.
- [2] Gupta R, Bradoo S, and Sexena RK. *Letters in Applied Microbiology*, 1997; 24: 253-255.
- [3] Hernandez MC, Augur C, Rodriguez R, Esquivel JCC, and Aguilar CN. *Food Technology and Biotechnology*, 2006; 44(4): 541-544.
- [4] Abdel-Naby MA, Sharief AA, and Tanash AB. 2011; 10:149-158.
- [5] Aguilar CN, Rogrigues R, Gutierrez-Sanchez G, Augur C, Favela-Torres E, Prado-Barragan LA, Ramiewz-Coronel A and Contreras-Esquivel J. *Microbial Applied Microbiology and Biotechnology*, 2007;76: 47-59.
- [6] El-Fouly1 MZ, El-Awamry Z, Azza AM, Heba A, El-Bialy1, Naeem E, and Ghadeer EEI-Sa. *Journal of American Science*, 2010; 6(12) 709-721.
- [7] Belur P, Dand Mugeraya G. *Research Journal of Microbiology*, 2011; 61: 25-40.
- [8] Mahendran B, Raman N, and Kim DJ. *Applied Microbiology and Biotechnology*, 2006; 70:444-450.
- [9] Mondal KC, Banerjee D, Banerjee R, and Pati B R. *Journal of General and Applied Microbiology*, 2001; 47:263-267.
- [10] Aguilar CN, Augur and Sanches G. *Food Sci. Technol. Int.*, 2001; 7: 373-382.
- [11] Banerjee D, Mondal KC, and Patil BR. *Journal of Basic Microbiology*, 2001; 41: 313- 318.
- [12] Hadi TA, Banerjee R, and Bhattacharyya BC, *Bioprocess Engineering*, 1994; 11:239-234.
- [13] Banerjee D, Mahapatra S, and Pati BR. *Research Journal of Microbiology*. 2007; 2: 462-468.
- [14] Bajpai B, and Patil S. *Technol*, 1997; 20: 612-615.
- [15] Barthomeuf C, Regeat F and Pourrat H. *Journal of Fermentation and Bioengineering*, 1994; 77: 320-323.
- [16] Belmares R, Contreas EJC, Rodriguez HR, Coronet AR, and Aguilar CN. *Lebensm-Wiss U-Techno*, 2004; 37:857-864.
- [17] Beverini M, and Metehe M. *Science Aliments*, 1990; 10: 807-816.

- [18] Bhat TK, Singh B, and Sharma OP. a current perspective, 1998;25:43-357.
- [19] Costa A M, Ribeiro WX, Kato E, Monteiro ARG, and Peralta R.M. Brazilian Archiae of Biology and Technology, 2008; 51: 399-404.
- [20] Costa AM, Kadowaki MK, Minozzo MC, Braeht A, and Peralta RM. African Journal of Biotechnology, 2012; 11: 391-398.
- [21] Baodi DK, and Neufeld RJ. Encapsulation Enzyme and Microbial Technology, 2001; 28: 590-595.
- [22] Batra A and Saxena RK. Process Biochemistry, 2005; 40:1553-1557.
- [23] Bradoo S, Gupta Rand Saxena RK. Process biochemistry, 1997;32:135-139.
- [24] Aguilar C N and Gutierrez-S´anchez G. Food Science and Technology International, 2001; 7:373–382.
- [25] Chaterjee S, Chaterjee BY, and Guha AK. Process Biochemistry, 2004; 39:2229-2232.
- [26] Chhokar V, Sangwan M, Beniwal V, Nehra K and Nehra KS. Applied Biochemistry and Biotechnology, 2010; 160:2256-2264.
- [27] Darah I, Sumathi G, Jain K and Hong L S. Biotechnology Research International, 2011;2011:1-7.
- [28] Enemuor SC, and Odibo FJC. African Journal of Biotechnology, 2009; 8(11): 2554-2557.
- [29] EL-Tanash AB, Sherief A, and Nour A. Innovative Romanian Food Biotechnology, 2012; 10: 709-217.
- [30] Ayed L and Hamdi M. Plantrum Biotechnology letter, 2002; 170:446-1765.
- [31] Abdel-Nabs MA, Sharif AA, El-Tanash AB and Mankarios AT. Journal of Applied Microbiology, 1999; 87:108-114.
- [32] Farias GM, Gorbea C, Elkins JR, and Griffin GJ. Physiological and Molecular Plant Pathology, 1994; 44: 51-63.
- [33] Beniwal V, and Chokar V. International Journal of Science and Technology (IJST), 2010; 2(1): 46-52.
- [34] Garcia-Conesa MT, Kauppinen S, and Williamson G. Carbohydrate Polymers, 2001; 44: 319-324.
- [35] Bajpai B, and Patil S. World Journal of Microbiology and Biotechnology, 1996; 12: 217- 220.
- [36] Hota SK, Dutta JR and Banerjee R. Indian Journal of Biotechnology 2007; 6:200-204.
- [37] Hatamoto O, Watarai T, Kikuchi M, Mizusawa K and Sekine H. Gene, 1996; 175: 215-22 1.
- [38] Kar B, Banerjee R and Bhattacharyya BC. Process Biochemistry, 2003; 38: 1285-1293.
- [39] Kar B, and Banerjee R. Journal of Industrial Microbiology and Biotechnology, 2000; 25: 29-38.
- [40] Iacazio G, Perissol C, and Faure B. A Journal of Microbiological Methods, 2000; 42: 209-214.
- [41] Kumar R A, Nagpal R, Sharma J and Kumari A. Annual Microbiology, 2010; 60:177-179.
- [42] Lokeshwari N, Jaya- Raju K, Pola S and Bobbarala V. International Journal of Chemical and Analytical Science, 2010;15: 106-109.
- [43] Kumar R, Sharma J and Singh R. Microbiol. Res., 2007; 162: 384-390.
- [44] Jana A, Maity C, Halder SK, Mondal KC, Pati BR and Mohapatra PK. Applied Biochemistry and Biotechnology, 2012; 6: 111-117.
- [45] Libuchi S, Minoda Y, and Yamada K. Agricultural and Biological Chemistry, 1997;31: 513-518.
- [46] Lekha PK and Lonsane BK. Process Biochemistry, 1994;29: 497-503.
- [47] Mohapatra S, Pradeep KD, Mondal KC and Pati BR. Polish Journal of Microbiology, 2006; 5(4): 297-301.
- [48] Lokheshwari N, and Raju KJ. Electronic Journal of Chemistry, 2007; 4: 287-293.
- [49] Mukherjee G and banerjee R. World Journal of Microbiology and Biotechnology, 2006; 22:207-212.
- [50] Lekha PK and Lonsane BK. state of art, Advance in Applied Microbiology, 1997; 44: 215-260.
- [51] Mondal KC and Pati BR. Journal of Basic Microbiology, 2000; 40: 223-232.
- [52] Mondal KC, Samanta S, Giri S and Patil BR. Acta microbiol. Pol., 2001;50: 75- 82.
- [53] Mukherjee G and Banerjee R. Basic Microbial, 2004; 44: 42-48.
- [54] Nishitani Y and Osawa R. Journal of Microbiological Methods, 2003; 54: 281-284.
- [55] Osawa R, Kuroiso K, Goto S, and Shimizu A. Applied and Environmental Microbiology, 2000;66: 3093-3097.
- [56] Paranthaman R, Vidyalakshmi R, Murugesh S and Singaravadivel K. Advanced Biological Research, 2009; 3(1-2): 34-39.
- [57] Sharma S and Gupta MN. Bioorganic and Medicinal Chemistry Letters, 2003;13: 395-397.
- [58] Srivastava A and Kar R. Brazilian Journal of Microbiology, 2009; 40: 782-789.
- [59] Pandey A, Soccol CR, Roudriguez-Leon JA and Nigam P. Asiatech Publishers, 2001; 36: 1153-1170.
- [60] Reddy BS and Rathod V. Asian Journal of Biochemical and Pharmaceutical Research, 2012;21: 59-62.
- [61] Rout S and Banerjee R. Indian J. Biotechnol, 2006;5: 346-350.
- [62] Schons PF, Rezender FC, Battestin V and Macedo GA. Journal of Microencapsulation, 2011;28.3:211-214.





- [63] Selwal MK, Yadav A, Selwal KK, Aggarwal N K, Gupta R and Gautam S.K. Brazilian Journal of Microbiology, 2011;42: 374-387.
- [64] Sabu A, Kiran GS, and Pandey A. Food Technology and Biotechnology, 2005 ;432: 133-138.