

Original Research Article

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Optimization of Extraction Techniques for the Release of Intracellular L-Asparaginase from *Serratia marcescens* MTCC 97 and its Characterization

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ABSTRACT

L-asparaginase acts as an efficient agent in curing certain sorts of lymphoma and leukemia by catalyzing the deamination of L-asparagine to L-aspartate and ammonia. Microorganisms are better source of L-asparaginase, as their culturing, extraction and purification is more convenient than plants and other sources. As most of L-asparaginases are intracellular in nature, so the selection of a suitable method for its release with maximum recovery was become more important. In present study, the resting cells of *S. marcescens* MTCC 97 were disintegrated by different enzymatic (lysozyme), chemical (alkali lysis, acetone powder, guanidine-HCl and triton X-100) and physical (motor and pestle, vortex, bead beater and sonicator) methods. Among all methods explored, sonication was found best method with 0.05 U/mg specific activity and minimum loss of enzyme (8%). Different reaction parameters were also optimized for the characterization of released L-asparaginase. The extracted L-asparaginase showed maximum activity (0.985 U/ml) in 0.05M sodium phosphate buffer (pH 7.5) with L-asparagine (8mM) as substrate at 40°C incubation for 20 min. Moreover, different metal ions, additives, chelating agents and protease inhibitors showed negative effects on L-asparaginase activity of resting cells and cell free extract obtained from *S. marcescens* MTCC 97.

Keywords

L-asparaginase,
Serratia marcescens
MTCC 97,
disintegration,
sonication.

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Introduction

L-asparaginases are the enzymes that catalyse the hydrolysis of L-asparagine into L-aspartate and ammonia. The L-asparaginases can be specific for L-asparagine, with negligible activity against glutamine (EC

3.5.1.1), or catalyze both asparagine and glutamine conversion (Sanchez M *et al.*, 2007). These enzymes act as important precursor in the treatment of Acute Lymphoblastic Leukemia in children due to antineoplastic activity (Umesh K *et al.*, 2007). The malignant cells are differentiated from

normal cells due to their nature in slow synthesis of L-asparagine, which causes starvation for this amino acid, while normal cells can produce this amino acid (Prakasham RS *et al.*, 2009). The cancer cells have diminished expression of L-asparagine and mainly utilize the L-asparagine circulating in plasma pools (Manna S and Gram C, 1995; Swain AI *et al.*, 1993).

The *Escherichia coli* and *Erwinia chrysanthemi* asparaginases are useful anti-leukaemic agents (Hill JM, 1967). Some asparaginases are also known to cause hemorrhage in the central nervous system, coagulation abnormalities, thrombosis and hypersensitivity reactions which are treatable upto 80% (Hourani R *et al.*, 2008; Menon J *et al.*, 2008). Clinical trials of L- asparaginase suggest this enzyme as a promising agent in treatment of neoplastic cell diseases in man with very low (1–2%) risk of cerebral venous thrombosis (Oettgen HF *et al.*, 1967; Erbetta A *et al.*, 2008).

L-asparaginases are reported from various sources like plants, animals and microorganisms but the microorganisms are better source of L-asparaginase. It is easy to culture and extract the microbial sources and the purification of enzymes is also convenient from microorganism. A very active form of L-asparaginase was found in *C. glutamicum* under lysine producing fermentation conditions (Mesas JM *et al.*, 1990). Most of L-asparaginases are intracellular in nature and need to be released from the cells for further applications. However, some extra cellular expression was also being exploited in recombinant DNA technology (Khushoo A *et al.*, 2004). This enzyme was isolated from variety of sources such as *Vibrio succinogenes*, *Proteus vulgaris* and *Pseudomonas fluorescens*, which are toxic to Lymphoblastic Leukaemia cells (Pritsa A and Kyriakidis DA, 2001). L-asparaginase

conjugated with poly ethylene glycol approved in year 1994 in United States for the treatment of Acute Lymphocytic Leukemia with trade name Oncaspar®). In the biosynthesis of the aspartic amino acids, L-asparaginases play a very critical role. In addition the role of L-asparaginases in amino acid metabolism and their antitumor properties makes this enzyme of great therapeutic interest.

Number of methods for cell disintegration has been developed in order to release the intracellular products and enzymes from the cells. For the extraction of intracellular materials from the cells, it must be disintegrated either by physical (mechanical) or chemical methods but the selected method of disruption must ensure the protection of labile cell content from denaturation or thermal deactivation. There are some other methods involving genetic engineering of the microorganism to release enzymes to the external medium, but its scope is limited due to high production cost.

Although, in the past few years various intracellular enzymes have been produced by the industries like as: glucose oxidase for food preservation, penicillin acylase for antibiotic conversion and L-asparaginase for possible cancer therapy (Wang B *et al.*, 2003). Chemical methods of cell disruption to release the cellular material may be advantageous as they employ use of acid, alkali, surfactants and solvents in some cases, but are generally avoided due to the limitation imposed by high cost at larger scale and damage due to acid/alkali, contamination of product with these chemicals, which further add more problems to downstream processing.

Mechanical/physical methods of cell disruption include both liquid (high pressure homogenizer) and solid shear (bead mill). The

most commonly method used in large scale to small scale production of intracellular proteins from microorganisms is bead agitation or bead milling which involves the vigorously agitation of harvested cells with beads in a closed chamber (Kula MR and Schutte H, 1987). Sonication, an another method of mechanical disruption had been previously employed for obtaining the cell free extract from *Erwinia carotovora* but there was biggest loss of enzyme occur during extraction (Krasotkina J *et al.*, 2004). But still sonication has been found most effective method for release of intracellular L-asparaginase among chemical and other physical methods used for cell disruption in earlier reports (Singh RS, 2013).

Despite of many cell disintegration methods are available for laboratory scale, only limited number from these methods have been used for large scale applications. The high cost of products by manufacturer is due to necessity of harvesting the cells and extracts the required internal constituent (Kirsop BH, 1981). In order to meet the requirements of L-asparaginases in therapeutics and the intracellular nature of this enzyme makes it necessary to search for a suitable cost effective method for its release from the microbial biomass. So, the present study was designed for the optimization of different extraction techniques for the release of intracellular L-asparaginase from *Serratia marcescens* MTCC 97 and its characterization.

Materials and Methods

Microorganism

The culture of *Serratia marcescens* MTCC 97 used in this study was procured from the Department of Biotechnology, Himachal Pradesh University, Shimla. This culture was maintained in medium containing (% , w/v):

malt extract 1.0, peptone 1.0, NaCl 0.5 and L-asparagine 0.1 (pH 7). After 24h of incubation, the culture was harvested by centrifugation at 10,000 rpm for 15 min at 4°C and the resting cells were used for the release of the L-asparaginase.

Estimation of cell mass

The 24 h old culture broth was centrifuged at 10,000 rpm for 15 min at 4°C and wet weight of cells was estimated. The wet cell pellet was placed in Oven at 80°C over night for drying. Dried cell pellets were cooled in desiccators and their weight were taken. The dried cell weight corresponding to their known amount of wet cell weight and their corresponding optical density was recorded and a standard graph was plotted between dry cell weight and A_{600} .

Assay of L-asparaginase activity

Asparaginase activity was assayed according to the method of Meister A *et al.*, (1955) and ammonia liberated was estimated by Fawett JK and Scott JE (2007) and the calorimetric Bradford assay was used for estimation of protein (Bradford MM, 1976). The L-asparaginase activity is expressed in terms of Unit (U).

For whole cells

The L-asparagine unit (U) has been defined as the μ moles of ammonia released / mg of dcw/ min under standard assay conditions.

For cell free enzyme

The L-asparaginase unit (U) has been defined as the μ moles of ammonia released / ml/ min under standard assay conditions.

Specific activity - U/mg of proteins

For whole cells

The L-asparagine unit (U) has been defined as the μ moles of ammonia released / mg of dcw/ min under standard assay conditions.

For cell free enzyme

The L-asparagine unit (U) has been defined as the μ moles of ammonia released / ml/ min under standard assay conditions.

Specific activity - U/mg of proteins

Procedure for enzyme assay

Cell suspensions (50 μ l) of known A_{600} (25; equivalent to 10.75mg/ml dcw) cells were taken in test tubes and 1.45 ml of buffer was added to make the volume to 1.5 ml. The reaction is started by adding 0.5 ml of 10mM substrate (L-asparagine) and the reaction mixtures were incubated at 45°C for 20 min. The reaction is stopped by adding 0.5 ml of trichloroacetic acid (15 %, w/v). In control tubes, 50 μ l cell suspensions were added after the addition of trichloroacetic acid. One ml reaction mixture was withdrawn from each tube (test and control) and released ammonia was measured. For the estimation of released enzyme, 50 μ l cell free extract was added in test and control. Rests of the conditions were similar to the assay procedure with resting cells.

Disintegration of resting cells of *S. marcescens* MTCC 97 for the release of L-asparaginase

The intracellular nature of the L-asparaginase in *S. marcescens* MTCC 97, Make mandatory to disintegrate the cells to release the L-asparaginase enzyme. Various enzymatic, chemical and physical methods were used for extraction of L-asparaginase from fresh biomass. The resting cells of *S. marcescens*

MTCC 97 were suspended in 0.05M sodium phosphate buffer (pH 7.5) with a cell concentration of 10.75mg/ml after washing twice with the same buffer. After the release of L-asparaginase from the resting cells, calculations were made by using following formulas:

$$\text{Recovery (\%)} = \frac{\text{Amount of released enzyme}}{\text{Maximum enzyme activity}} \times 100$$

$$\text{Loss (\%)} = \frac{\text{Maximum enzyme activity} - (\text{Amount of released enzyme} + \text{Amount of unreleased enzyme})}{\text{Maximum enzyme activity}} \times 100$$

Enzymatic method

Lysozyme treatment (Schutte H and Kula MR, 1993)

In this method, cell pellet obtained from 100 ml of culture broth was suspended in 2 ml of solution A (Glucose: 50mM, EDTA: 10mM, Tris buffer: 25mM, pH 8) and 0.5 ml of solution B (Lysozyme: 50mg/ml, Dissolved in solution A). Mixing was done by vortexing and mixture was incubated in ice for 10 min. To the reaction contents 0.5 ml of solution C (NaOH: 0.2M w/v, SDS:1% w/v) was added, mixed and placed again in ice. The cell slurry was centrifuged at 10,000 rpm for 10 min at 4°C. The L-asparaginase activity was measured in the supernatant as well as in cell debris/unlysed cells.

Chemical methods

Alkali lysis (Birnboim HC and Dolt J, 1979)

Cell pellet obtained from 100 ml of culture broth was suspended in 1ml of solution A (Glucose: 50mM, EDTA: 10mM, Tris buffer: 25mM pH 8) and 2 ml of solution B (NaOH:

0.2M w/v, SDS: 1% w/v). The reaction contents were mixed by inverting the tubes 5-6 times and stored in ice. Then 1.5 ml of ice cold solution C (Potassium acetate: 60 ml 5M, Glacial acetic acid:11.5 ml, Water: 28.5 ml) was added and the tubes were vortexed for 10 min. The cell slurry was centrifuged at 10,000 rpm for 10 min at 4°C. The L-asparaginase activity was measured in the supernatant as well as in cell debris/unlysed cells.

Acetone powder method (Somerville HJ *et al.*, 1970)

Cell pellet obtained from 100 ml of culture broth was suspended in 10 ml of anhydrous acetone and placed in ice for 30 min at 10°C. The reaction contents were mixed by vortexing. Cell slurry was centrifuged at 10,000 rpm for 10 min at 4°C.

Cell pellet was suspended in 10mM of sodium borate buffer (pH 6.5) and incubated at 40°C for 10 min. Cell content was again centrifuged at 10,000 rpm for 10 min at 4°C. The L-asparaginase activity was measured in the supernatant as well as in cell debris/unlysed cells.

Triton X-100 and guanidine-HCl treatment for cell disruption (Helenius A and Simons K, 1975)

Cell pellet obtained from the culture broth was suspended in 10 ml of phosphate buffer 0.05M, pH 7.5 (containing 10.75mg/ml dcw) and 4 ml of 2M Guanidine HCl was added to it. To this reaction mixture 0.24 ml of Triton X-100 2% (v/v) was added.

The reaction contents were mixed and incubated at room temperature for 15 min. Cell slurry was centrifuged at 10,000 rpm for 10 min at 4°C. The L-asparaginase activity was measured in the supernatant as well as in cell debris/unlysed cells.

Physical methods

Disruption of cells by crushing with glass beads in pestle and mortar

Cell pellet obtained from the culture broth was suspended in 15 ml of phosphate buffer (containing 10.75mg/ml dcw 0.05M, 0.05M pH 7.5). The PMSF (0.5mM 0.1 ml) was added to cell slurry ($A_{600} = 25$). The cell slurry was crushed continuously with 15 ml glass beads for 25 min with the help of mortar and pestle in ice chamber to avoid loss of activity due to heat generation during crushing. The crushed mixture was centrifuged at 10,000 rpm for 10 min at 4°C. The L-asparaginase activity was measured in the supernatant as well as in cell debris/unlysed cells.

Disruption of cells by Bead Beater (Kula MR and Schutte H, 1987; Chisti Y and Moo-Young M, 1991)

Cell pellet obtained from 300 ml of culture broth was suspended in 40 ml of phosphate buffer (containing 10.75mg/ml dcw). The cell slurry ($A_{600} = 25$) was disrupted by the use of Bead Beater™ for 36 min. Beads of different diameter (Zirconium 0.5mm, Glass beads 0.5mm and 0.1mm) were used for the disruption of cells with a pulse of 1 min on and 2 min off to avoid heat generation. The assembly containing cell slurry was ice jacketed during the cell disruption cycle. The sample was withdrawn after every 1 min for assay of L-asparaginase activity in supernatant and cell debris/unlysed cells.

Disruption of cells by Sonication (Singh RS, 2013)

Cell pellet obtained from the culture broth was suspended in 40 ml of phosphate buffer (containing 10.75mg/ml dcw). The cell slurry ($A_{600} = 25$) was disrupted by the use of

sonicator for 22 min with a pulse of 60 sec (60 sec on and 60 sec off) at 250 W by keeping the probe (diameter 1 inch) above the bottom of vial. The vial was ice jacketed during the sonication. The samples were withdrawn after every 1 min for the assay of L-asparaginase activity in supernatant and cell debris/unlysed cells.

Optimization of parameters for the maximum release of L-asparaginase by sonication

Number of pulse cycles

The 40 ml cell slurry ($A_{600} = 25$) was disrupted with the sonicator for 22 min with a pulse of 60 sec and at 39% amplitude. The sample was withdrawn after every 60sec and centrifuged at 10,000 rpm for 10 min at 4°C. The L-asparaginase activity was measured in the supernatant and pellets both. The cycle which showed the highest activity was selected and used for further studies.

Cell concentration

The cell slurry (40 ml) of different cell concentration (2.15mg/ml, 4.3mg/ml, 6.45mg/ml, 8.6mg/ml, 10.75mg/ml, 10.75mg/ml, 12.9mg/ml and 15.05mg/ml) were lysed by the sonicator for 9 cycles. The released L-asparaginase activity was measured for each cell concentration in cell free extract and cell debris/unlysed cells. The cell concentration which showed the maximum enzyme activity was selected as the optimum concentration of cells to be used for further studies.

Cell volume

Different cell volumes (20 ml, 30 ml, 40 ml and 50 ml) resting cell of selected concentration (10.75mg/ml) was used for cell disintegration. For each cell volume the

released L-asparaginase activity was measured in cell free extract and cell debris/unlysed cells.

Amplitude of sonication

The cell slurry (40 ml) of cell concentration 10.75mg/ml was lysed in sonicator for 9 cycles at different amplitudes (30%, 35% and 39%). The L-asparaginase activity was measured in the cell free extract and cell debris/unlysed cells.

Characterization of L-asparaginase released from the resting cells of *S. marcescens* MTCC 97

The reaction conditions were optimized for the assay of L-asparaginase activity in cell free extract obtained from *S. marcescens* MTCC 97 and compared with the L-asparaginase of the resting cells of *S. marcescens* MTCC 97.

Selection of buffer and optimization of pH

The optimum pH of released L-asparaginase enzyme was evaluated by measuring the L-asparaginase activity in different buffers of 0.1M concentration. The buffers used were; Acetate buffer (pH 4.0-6.0), Sodium phosphate buffer (pH 6.0-8.0), Potassium phosphate buffer (pH 7.0-8.5), Citrate buffer (pH 4.5-6.5), Glycine NaOH buffer (pH 9.0-10.0), Carbonate-bicarbonate buffer (pH 9.5-10.5), Citrate phosphate buffer (pH 2.5-7.0) were used to perform the assay. The same set of experiment was also performed with resting cells of *S. marcescens* MTCC 97.

Optimization of buffer molarity

To study the effect of concentration of buffer on released L-asparaginase, Sodium phosphate buffer (pH 7.5) of different concentration (0.01M - 0.07M) was used for

the assay of L-asparaginase activity in cell free extract and resting cells.

Optimization of reaction temperature

The optimum temperature of the L-asparaginase from *S. marcescens* MTCC 97 was obtained by measuring the L-asparaginase activity in cell free extract and resting cells at different incubation temperature (30°C, 35°C, 40°C, 45°C, 50°C and 55°C) with L-asparagine as substrate and 0.05M sodium phosphate buffer (pH 7.5).

Effect of incubation time

Optimum reaction time was evaluated by incubating the reaction contents for different time intervals (10, 15, 20, 25, 30 and 35 min) and optimum pH and temperature. The L-asparaginase activity was measured in resting cells and cell free extract obtained from *S. marcescens* MTCC 97.

Substrate specificity

To find out the substrate specificity of L-asparaginase of *S. marcescens* MTCC 97, the activity of enzyme was determined at different substrate like L-asparagine, L-glutamine, D-asparagine and DL-asparagine at 10mM concentration. The experiment was performed with resting cells and cell free extract obtained from *S. marcescens* MTCC 97.

Substrate concentration

For the optimization of substrate concentration of released L-asparaginase and resting cells of *S. marcescens* MTCC 97, different substrate concentrations of L-asparagine (2mM-14mM) were used and assay was performed under optimized conditions.

Role of metal ion

The L-asparaginase activity was assayed in presence of 1mM concentration of metal ions, additives, inhibitors and chelating agents (FeCl₃, MgSO₄.6H₂O, ZnSO₄.7H₂O, COCl₂, CuSO₄.5H₂O, NaCl, AgNO₃, BaCl₂, Dithiothreitol, Ethylene diamine tetra acetic acid, Phenyl methyl sulphonyl fluoride, HgCl₂, CaCl₂.2H₂O, Urea, Polyethylene glycol (PEG), Pb(NO₃)₂, MnCl₂.H₂O and KCl) under previously optimized conditions for cell free extract and resting cells of *S. marcescens* MTCC 97.

Determination of K_m and V_{max} of released enzyme

K_m and V_{max} values were determined by plotting a graph between 1/V and 1/S for resting cells and free extract obtained from *S. marcescens* MTCC 97.

Stability profile of purified enzyme

The Stability of enzyme was determined at three different temperatures (4°C, 25°C, 30°C, 40°C and 50°C). The enzymes (cell free extract and resting cells) were incubated at these temperatures and activity was measured at regular interval of 30 min.

Results and Discussion

Optimization of cell disintegration methods for release of L-asparaginase from *Serratia marcescens* MTCC 97

The isolation of intracellular enzymes requires a suitable cell disruption method (enzymatic, chemical or physical) to release its contents into the surrounding medium (Chisti Y and Moo-Young M, 1991). The L-asparaginase from *S. marcescens* MTCC 97 is an intracellular enzyme and can only obtain by cell disruption. There are several methods

of partial or selective disruption of membranes to solubilise bound proteins including the use of chelating agents, adjustment of ionic strength, pH, organic solvents and detergents (Somerville HJ *et al.*, 1970; Helenius A and Simons K, 1975; Marchesi SL *et al.*, 1970; Schnebli HP and Abrams A, 1970). The resting cells of known A₆₀₀ (25; equivalent to 10.75mg/ml dcw) obtained from *S. marcescens* MTCC 97 were disintegrated by different enzymatic (lysozyme), chemical (alkali lysis, acetone powder, Triton X-100 and Guanidine-HCl) and physical (mortar and pestle, vortex, Bead Beater and Sonicator) methods.

Enzymatic method

In enzymatic methods, the amount of enzyme released was found 7.13 U (Table 1). However, 4.04mg/ml protein was found in the supernatant with 0.073 U/mg specific activity. Even after cell lysis, 5.96 U the enzyme activity was remaining in the unlysed cells. Recovery of L-asparaginase was found to be 42% and almost 13% loss in the enzyme activity was observed. Cell lysis of Gram's negative bacteria was aided by the addition of EDTA to chelate the divalent cations (Schutte H and Kula MR, 1993) and lysozyme was used to cleave β (1-4) glycosidic linkage of bacterial cell wall (Bucke C, 1983). However, the process was very costly at large scale economics points of view.

Chemical methods

Alkali lysis method

Less quantity of L-asparaginase release (0.48 U) with specific activity of 0.030 U/mg of protein was observed when the cells of *S. marcescens* MTCC 97 were subjected to alkali lysis (Table 2). However, the amount of protein released was found to be (3.55mg/ml). The decrease in enzyme activity might be due

to the denaturation of enzyme by SDS. The L-asparaginase recovery was found to be 6% with a loss of 38% after the cell lysis.

Acetone powder

The acetone powder was prepared to release the L-asparaginase resting cells of *S. marcescens* MTCC 97. Overall 6.58mg/ml protein was released in the supernatant with enzyme activity of 1.37 U. The specific activity was found to be 0.021 U/mg of protein (Table 3).

Triton X-100 and Guanidine-HCl

On the treatment of resting cells of *S. marcescens* MTCC 97 with Triton X-100 and Guanidine-HCl, 0.26 U/ml L-asparaginase was released in supernatant with 2.47mg/ml yield of protein (Table 4). The specific activity of enzyme was 0.007 U/mg of protein. The overall loss in the enzyme activity was 12% with 3% recovery of enzyme. Therefore, this method was not found to be suitable for lysis as the specific activity of enzyme was very less and recovery was also low. Among the three chemical methods used for the disruption of the resting cells of *S. marcescens* MTCC 97, the treatment of the cells with Triton X-100 and Guanidine-HCl gave maximum yield (0.26 U of released enzyme) with the release of 2.47mg/ml of protein and specific activity of the enzyme was found to be 0.007 U/mg of protein. Furthermore, with very less recovery (3%), this method was found to be unsuitable for the release of L-asparaginase from the resting cells of *S. marcescens* MTCC 97. Resting cells of *S. marcescens* MTCC 97 were also lysed by acetone powder treatment method with 17% recovery of L-asparaginase. Moreover, during this procedure 30% loss in L-asparaginase was also recorded. However acetone treatment was used to increase the permeability of cell wall of *E. carotovora* and

enzyme recovery in cell free extract was reported to 57% (Lee SM *et al.*, 1989).

Physical methods

Disintegration of cells in mortar and pestle

In supernatant 3.09 U enzyme activity and 6.62mg/ml protein was obtained after the cell disruption in mortar and pestle (Table 5). The specific activity of the released L-asparaginase was 0.031 U/mg of protein. The loss in enzyme activity was 8% with overall recovery of 26% of L-asparaginase.

Disintegration of cells by vortexing with glass beads

Disintegration of the resting cells of *S. marcescens* MTCC 97 was also tried by vortexing the cell slurry with glass beads (0.5mm). The amount of L-asparaginase released was found to be 3.45 U and the protein obtained in supernatant was 6.26mg/ml (Table 6). The specific activity of the supernatant was 0.043 U/mg of protein. The L-asparaginase activity in the cells before disruption was 1.095 U and cells retained 0.072 U L-asparaginase after the cell disruption. The overall loss in enzyme activity was 5% with a recovery of 29%.

Disintegration of cells by Bead Beater using Zirconium beads (0.5mm)

The cell slurry (40 ml) of *S. marcescens* MTCC 97 was disintegrated in a Bead Beater by using Zirconium beads of 0.5mm diameter. The activity in supernatant was 9.48 U with 7.20mg/ml of released protein (Table 7). The specific activity was found to be 0.029 U/mg of protein. Activity in cells before disruption was 33.54 U and cell retained 2.58 U of enzyme after the cell disruption. The overall recovery was 28% with 64% loss.

Disintegration of cells by glass beads (0.5mm)

The 40 ml resting cells suspension of *S. marcescens* MTCC 97 was disrupted by using glass beads of 0.5mm diameter in Bead Beater. The released L-asparaginase activity and protein was found to be 8.29 U and 13.28mg/ml, respectively (Table 8). The specific activity of released enzyme was 0.017 U/mg of protein. The overall recovery of L-asparaginase was 24% with 64% loss in the enzyme activity.

Disintegration of cells by glass beads (0.1mm)

The cell slurry (40 ml) of *S. marcescens* MTCC 97 was disrupted by using glass beads of 0.1mm diameter. The L-asparaginase release was found to be 19.20 U with 16.67mg/ml of protein (Table 9). The specific activity of cell free extract was 0.032 U/mg of protein. The recovery of L-asparaginase was better (50%) but the loss in the enzyme activity was also very significant (48%).

Disintegration of cells by sonication

The disintegration of resting cells of *S. marcescens* MTCC 97 was carried out by sonication. After 9th cycle of sonication, 27.0 U of L-asparaginase and 19.06mg/ml of protein were released in the supernatant (Table 10). The specific activity of released L-asparaginase was found to be 0.05 U/mg proteins. The recovery of L-asparaginase was 68% with a little loss (8%) in of enzyme activity.

Optimization of various parameters for the release of L-asparaginase from *S. marcescens* MTCC 97 cells by Sonication

As the recovery of L-asparaginase was maximum with sonication method with very

less loss of enzyme activity, the different parameters of sonication like pulse rate, cell volume and cell concentration for the maximum release of the enzyme were also optimized.

Optimization of pulse rate

The 40 ml cell slurry of *S. marcescens* MTCC 97 was sonicated for 12 cycles of a pulse of 60 sec. The maximum enzyme activity (0.871 U/ml) and specific activity (0.047 U/mg protein) was found at the 9th cycle of sonication (Table 11). The specific activity of enzyme decreased after 9th cycle possibly due to the thermal denaturation. These results suggest that the 9 on/off cycles were optimum for the maximum release of L-asparaginase from the resting cells of *S. marcescens* MTCC 97.

Optimization of cell concentration

The 40 ml cell slurry of *S. marcescens* MTCC 97 containing varying amount of resting cells were sonicated for the release of L-asparaginase (Table 4.12 A, B, C, D, E, F and G). The amount of enzyme released was decreased beyond the cell concentration of 10.75mg/ml. The maximum protein (19.05mg/ml) was released at the cell concentration of 10.75mg/ml with maximum recovery of 68%. Therefore, 10.75mg/ml resting cells were further used for the release of L-asparaginase by sonication.

Optimization of cell volume

Different volumes (20-50 ml) of cell slurry of *S. marcescens* MTCC 97 containing 10.75mg/ml resting cells were lysed for 9 on/off cycles of sonication (Table 13 A, B, C and D). The maximum enzyme (32.0 U) was released when 40 ml of cell slurry was used. There was a decrease in activity when a higher volume of cell slurry was used.

Optimization of amplitude

The 40 ml cell slurry (containing 10.75mg/ml cells) was sonicated at different amplitudes (30, 35 and 39%) for 9 on/off cycles (Table 14 A, B and C). It is important to mention that the maximum amplitude of sonicator should not exceed 39%. The most efficient amplitude was found to be 39%. Below this amplitude the lysis was not very effective as the activity in pellet after lysis was found to be very high.

Characterization of L-asparaginase released from the resting cells of *S. marcescens* MTCC 97

The reaction conditions were optimized for the assay of L-asparaginase activity in resting cells as well as cell free extract obtained from *S. marcescens* MTCC 97.

Selection of buffer and optimization of pH

For the selection of buffer of optimum pH, 7 buffers of 0.1M concentration having different pH range (4-10.5) were tested. The maximum L-asparaginase activity was found with 0.1M sodium phosphate buffer (pH 7.5) in resting cells (0.116 U/mg dcw) and same buffer was found to be most suitable for cell free extract of *S. marcescens* MTCC 97 with maximum L-asparaginase activity 0.558 U/ml (Table 16). This data suggest that the released enzyme had optimum pH similar to that of resting cell preparations. The activity falls in both cases (resting cells as well as in cell free extract) as the pH was altered from the optimum. The reason behind this may be that enzyme was unable to retain its activity at high or low pH due to the fact that active site losses its affinity towards substrate at these pH. The reaction conditions of L-asparaginase produced by *S. marcescens* MTCC 97 were optimized to find out the most favourable conditions for enzyme to exhibit its maximum activity. Various buffers of pH range (4-10.5)

were used to perform enzyme assay. The maximum L-asparaginase activity was obtained with 0.05 M sodium phosphate buffer (pH 7.5) in resting cells as well as for cell free extract of *S. marcescens* MTCC 97. The enzyme from *Erwinia carotovora* has optimum pH 8.0, which was completely different from the whole cell optimum pH, which are 7.3 (Maladkar and George, 1993). However, the commercial preparation of L-asparaginase (Elspar) was found to be stable at wide pH range of 4.5-11.5 (Stecher AL *et al.*, 1999).

Optimization of buffer molarity

Different concentrations (10-70mM) of sodium phosphate buffer (pH 7.5) were used to select the optimum molarity of the buffer. Maximum L-asparaginase activity was obtained with 50mM concentration of sodium phosphate buffer (pH 7.5) in resting cells as well as in cell free extract of *S. marcescens* MTCC 97. In resting cells and cell free extract the L-asparaginase activity was found to be 0.121 U/mg dcw and 0.815 U/ml, respectively (Fig. 1).

Optimization of reaction temperature

The reaction mixture containing cell free extract and resting cells of *S. marcescens* MTCC 97 were separately incubated at different temperature (30°C-55°C). Maximum L-asparaginase activity in resting cells (0.146 U/mg dcw) and in cell free extract (0.754 U/ml) activity was observed at 40°C (Fig. 2). However, with further increase in incubation temperature, L-asparaginase activity decreased in both cases. The optimum reaction temperature was found to be 40°C in resting cells and in cell free extract of *S. marcescens* MTCC 97 which coincide with *C. glutamicum* having the same optimum reaction temperature (Mesas JM *et al.*, 1990).

Effect of incubation time

Optimum reaction time was evaluated by incubating the reaction contents for different time intervals (10, 15, 20, 25, 30 and 35 min) at optimum pH and temperature. The L-asparaginase activity in cell free extract of from *S. marcescens* MTCC 97 obtained was 0.760 U/ml after 20 min of incubation (Fig. 3). Similar incubation time was found to be optimum for the maximum (0.140 U/mg dcw) L- asparaginase activity. Enzyme activity started decreasing when incubation time was increased beyond 20 min in both the cases.

Substrate specificity

To find out the most specific substrate for the L-asparaginase of *S. marcescens* MTCC 97, the activity of enzyme was determined with different substrate (L-asparagine, L- glutamine, D-asparagine and DL-asparagine) at 10mM concentration. It was found that the L-asparagine was most suitable substrate for the L-asparaginase of *S. marcescens* MTCC 97. The resting cells and cell free extract exhibited 0.145 U/mg dcw and 0.826 U/ml of L-asparaginase activity, respectively. Moreover, it also showed very little D-asparaginase activity and L-glutaminase activity (Fig. 4). The most favorable substrate for the L-asparaginase from *S. marcescens* MTCC 97 was L-asparagine but this enzyme showed very little activity towards substrate D-asparagine also. Moreover, this enzyme also exhibit significant L-glutaminase activity.

Substrate concentration

Different concentrations of L-asparagine (2mM-14mM) were used to obtain the optimum substrate concentration for the L-asparaginase of *S. marcescens* MTCC 97. The maximum L-asparaginase activity was found to be 0.154 U/mg dcw with 10mM

concentration of L-asparagine with resting cells (Fig. 5). However, for the cell free extract of *S. marcescens* MTCC 97, the maximum L-asparaginase activity was obtained at 8mM concentration of L-asparagine (0.985 U/ml). A sharp decrease in L-asparaginase activity was observed with further increase in L-asparagine concentration in both cases. These finding suggest the possibility of substrate inhibition at the higher concentration of L-asparagine.

Role of metal ion

The L-asparaginase activity was assayed in presence of 1mM concentration of metal ions, additives, inhibitors and chelating agents under optimized conditions for cell free extract and resting cells of *S. marcescens* MTCC 97. The metal ions AgNO₃ and HgCl₂ inhibited the L-asparaginase activity in resting cells as well as in cell free extract. A slight increase in enzyme activity was observed by the use of BaCl₂, CaCl₂.2H₂O, Ethylene diamine tetra acetic acid (EDTA) and Phenyl methyl sulphonyl fluoride in resting cells and MnCl₂.H₂O in cell free extract. On the basis of insignificant effect of these metal ions on L-asparaginase activity, it can be suggested that the L-asparaginase of *S. marcescens* MTCC 97 is not a metalloprotein (Fig. 6).

Presence of metal ions does not affect L-asparaginase production indicates that it is not a metalloprotein or does not require co-factor. Presence of chelating agents (EDTA) and compounds having thiol protecting groups (glutathione, dithiothreitol, 2-mercaptoethanol etc) markedly enhance the L-asparaginase activity of *Cylindrocarpum obtusisporum* MB-10(Raha SK *et al.*, 1990).

Determination of K_m and V_{max} of enzyme

K_m and V_{max} values of L-asparaginase were determined by plotting a graph between 1/V

and 1/S for cell free extract and resting cells of *S. marcescens* MTCC 97. The values of V_{max} and K_m was found to be 1.65 U and 5.6×10^{-3} M, respectively for the cell free extract of *S. marcescens* MTCC 97 (Fig. 4.7). However, the V_{max} and K_m were 0.19 U and 1.85×10^{-3} M, respectively for the resting cells of *S. marcescens* MTCC 97 (Fig. 8). The high K_m value of cell free extract suggests that the released L-asparagine has less affinity for L-asparagine than the resting cells of *S. marcescens* MTCC 97.

The K_m values obtained for L-asparaginase in resting cells and cell free extract of *S. marcescens* MTCC 97 were 1.85×10^{-3} M and 5.6×10^{-3} M, respectively. The K_m value of a recombinant L-asparaginase ECAR LANS was found to be 1.6×10^{-2} μ M [16]. The minimum K_m value for L-asparaginase so far reported in *Pseudomonas* 7A (4.4×10^{-6} M) by Rozalska M and Mikucki J, 1992).

Stability profile of L-asparaginase

The Stability of L-asparaginase was determined at five different incubation temperatures (4°C, 25°C, 30°C, 40°C and 50°C). The L-asparaginase from *S. marcescens* MTCC 97 (cell free extract and resting cells) were incubated at these temperatures and activity was determined at regular interval of 30 min. The resting cells and cell free extract of *S. marcescens* MTCC 97 was found to be most stable at 4°C. The half-life of L-asparaginase obtained at 25°C and 30°C was 240 min for the resting cells as well as the cell free extract of *S. marcescens* MTCC 97 (Fig. 4.9 and Fig. 10). When the temperature was increased to 40°C, the half-life of L-asparaginase decreased to 210 min in both the cases. Moreover, at higher incubation temperature (50°C) the half-life of L-asparaginase in cell free extract and in resting cells was found to be 180 and 90 min, respectively.

Table.1 Lysis of the resting cells of *S. marcescens* MTCC 97 cells by lysozyme

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
			4.04	0.073	42	13
Before cell disruption	7.13	ND				
After cell disruption	5.96	0.89				

Table.2 Alkali lysis of the resting cells of *S. marcescens* MTCC 97

Conditions	Enzyme Activity (U)		Released Protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
			3.55	0.030	6	38
Before cell disruption	7.74	ND				
After cell disruption	4.30	0.48				

Table.3 Lysis of resting cells of *S. marcescens* MTCC 97 by acetone powder method

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
			6.58	0.021	17	30
Before cell disruption	8.06	ND				
After cell disruption	4.30	1.37				

Table.4 Lysis of resting cells of *S. marcescens* MTCC 97 by Triton X-100 and Guanidine-HCl treatment

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
			2.47	0.007	3	12
Before cell disruption	9.57	ND				
After cell disruption	8.17	0.26				

Table.5 Disintegration of resting cells of *S. marcescens* MTCC 97 in mortar and pestle

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	1.095	ND	6.62	0.031	26	8
After cell disruption	0.072	3.09				

Table.6 Disintegration of resting cells of *S. marcescens* MTCC 97 by vortexing with glass beads

Conditions	Enzyme Activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	1.095	ND	6.26	0.043	29	5
After cell disruption	0.072	3.45				

Table.7 Disintegration of resting cells of *S. marcescens* MTCC 97 by Zirconium beads (0.5mm)

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	33.54	ND	7.20	0.029	28	64
After cell disruption	2.58	9.48				

Table.8 Disintegration of resting cells of *S. marcescens* MTCC 97 by Glass beads (0.5mm)

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	33.40	ND	13.28	0.017	24	72
After cell disruption	1.18	8.29				

Table.9 Disintegration of resting cells of *S. marcescens* MTCC 97 by Glass beads (0.1mm)

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	38.27	ND	16.67	0.032	50	48
After cell disruption	0.86	19.2				

Table.10 Disintegration of resting cells of *S. marcescens* MTCC 97 by Sonication

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	48.16	ND	19.06	0.047	68	8
After cell disruption	18.00	27.00				

Table.11 Disintegration of resting cells of *S. marcescens* MTCC 97 by Sonication at different cycles

Cycle number	Enzyme activity (U)	Protein released (mg/ml)	Specific activity (U/mg)
1	0.21	5.336	0.039
2	0.32	9.032	0.036
3	0.49	13.07	0.037
4	0.56	15.66	0.036
5	0.65	15.36	0.042
6	0.73	16.90	0.043
7	0.73	20.17	0.036
8	0.85	19.77	0.043
9	0.87	19.06	0.047
10	0.86	21.39	0.040
11	0.85	21.71	0.039
12	0.87	22.01	0.039

Table.12 Disintegration of resting cells of *S. marcescens* MTCC 97 by sonication at different cell concentration

A. Cell concentration = 2.15mg/ml

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in Enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	6.71	ND	4.23	0.011	27	46
After cell disruption	1.81	1.80				

B. Cell concentration = 4.30mg/ml

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	14.62	ND	13.12	0.004	65	26
After cell disruption	1.40	9.46				

C. Cell concentration = 6.45mg/ml

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in Enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	20.64	ND	16.60	0.009	29	34
After cell disruption	7.74	5.92				

D. Cell concentration = 8.60mg/ml

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	31.99	ND	18.90	0.024	57	17
After cell disruption	8.60	18.24				

E. Cell concentration = 10.75mg/ml

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	51.60	ND	19.05	0.046	68	8
After cell disruption	12.9	34.84				

F. Cell concentration = 12.09mg/ml

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	62.43	ND	19.61	0.040	53	16
After cell disruption	19.61	33.00				

G. Cell concentration = 15.05mg/ml

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	77.06	ND	22.40	0.039	46	16
After cell disruption	29.50	35.28				

Table 13 Disintegration of different volume of *S. marcescens* MTCC 97 cells by sonication

A. Cell volume = 20 ml

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	25.8	ND	18.59	0.015	21	55
After cell disruption	6.00	5.50				

B. Cell volume = 30 ml

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	38.70	ND	18.21	0.025	24	47
After cell disruption	13.74	9.48				

C. Cell volume = 40 ml

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	51.6	ND	19.01	0.046	68	8
After cell disruption	12.9	34.84				

D. Cell volume = 50 ml

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	38.5	ND	19.45	0.039	60	7
After cell disruption	21.50	35.00				

Table 14 Disintegration of *S. marcescens* MTCC 97 cells at different amplitudes of sonication

A. Amplitude = 30%

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	47.7	ND	17.72	0.02	26	27
After cell disruption	24.51	13.20				

B. Amplitude = 35%

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	47.7	ND	20.56	0.03	39	21
After cell disruption	20.64	19.96				

C. Amplitude = 39%

Conditions	Enzyme activity (U)		Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
	In cells	In supernatant				
Before cell disruption	51.60	ND	19.05	0.046	68	8
After cell disruption	12.9	34.84				

Table 15 Comparison of the various methods used for the cell disintegration of the resting cells of *S. marcescens* MTCC 97

Methods	Treatments	Released protein (mg/ml)	Specific activity (U/mg)	Recovery (%)	Loss in enzyme activity (%)
Enzymatic	Lysozyme	4.04	0.073	42	13
Chemical	Alkali lysis	3.55	0.030	6	38
	Acetone powder	6.58	0.021	17	30
	Triton X-100 and Guanidine-HCl	2.47	0.007	3	12
Physical	Mortar and Pestle	6.62	0.031	26	8
	Vortex	6.26	0.043	29	5
	Bead beater	16.67	0.032	50	48
	Sonicator	19.06	0.047	68	8

Table.16 Selection of buffer and pH for resting cells and cell free extract obtained from *S. marcescens* MTCC 97

Buffers pH	Citrate phosphate buffer		Acetate buffer		Sodium phosphate buffer		Potassium phosphate buffer		Citrate buffer		Glycine NaOH buffer		Carbonate-Bicarbonate buffer	
	Enzyme activity		Enzyme activity		Enzyme activity		Enzyme activity		Enzyme activity		Enzyme activity		Enzyme activity	
	Resting Cells U/mg dcw	Cell free extract (U/ml)	Resting Cells U/mg dcw	Cell free extract (U/ml)	Resting cells U/mg dcw	Cell free Extract (U/ml)	Resting Cells U/mg dcw	Cell free Extract (U/ml)	Resting cells U/mg dcw	Cell free Extract (U/ml)	Resting Cells U/mg dcw	Cell free Extract (U/ml)	Resting Cells U/mg dcw	Cell free extract (U/ml)
4.0	NA	-	0.020	0.173	-	-	-	-	-	-	-	-	-	-
4.5	NA	-	0.040	0.200	-	-	-	-	0.020	0.186	-	-	-	-
5.0	0.038	0.083	0.049	0.441	-	-	-	-	0.062	0.156	-	-	-	-
5.5	0.061	0.325	0.046	0.160	-	-	-	-	0.067	0.508	-	-	-	-
6.0	0.075	0.492	0.041	0.118	0.029	0.185	-	-	0.075	0.396	-	-	-	-
6.5	0.076	0.323	-	-	0.046	0.367	-	-	0.107	0.500	-	-	-	-
7.0	0.035	0.205	-	-	0.100	0.473	0.038	0.246	-	-	-	-	-	-
7.5	-	-	-	-	0.116	0.558	0.084	0.399	-	-	-	-	-	-
8.0	-	-	-	-	0.069	0.366	0.102	0.497	-	-	-	-	-	-
8.5	-	-	-	-	-	-	0.052	0.230	-	-	-	-	-	-
9.0	-	-	-	-	-	-	-	-	-	-	0.006	0.020	-	-
9.5	-	-	-	-	-	-	-	-	-	-	0.007	0.006	0.031	0.276
10.0	-	-	-	-	-	-	-	-	-	-	0.007	NA	0.026	0.246
10.5	-	-	-	-	-	-	-	-	-	-	-	-	0.023	0.213

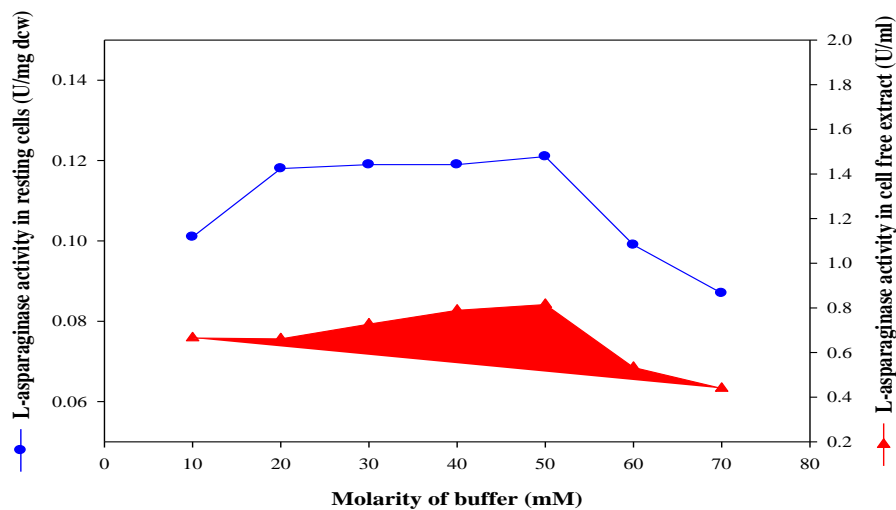


Fig.1 Effect of different concentration of sodium phosphate buffer on L-asparaginase activity of *S. marcescens* MTCC 97

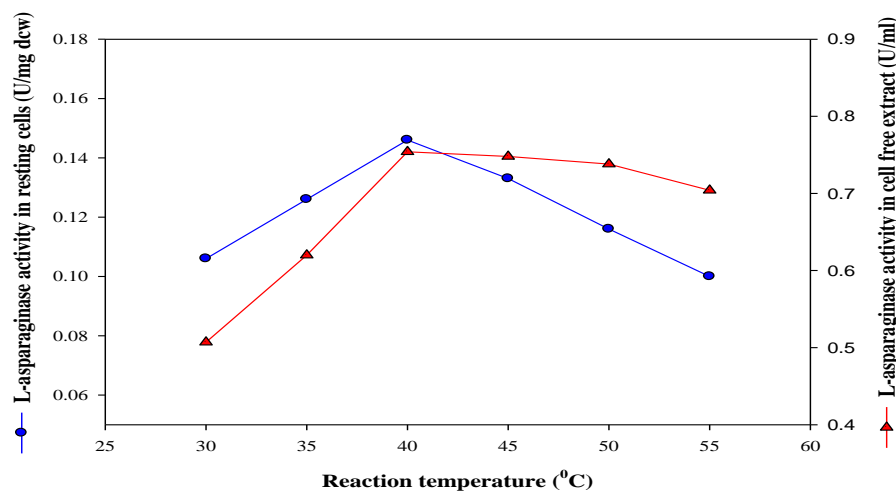


Fig.2 Effect of reaction temperature on L-asparaginase activity of *S. marcescens* MTCC 97

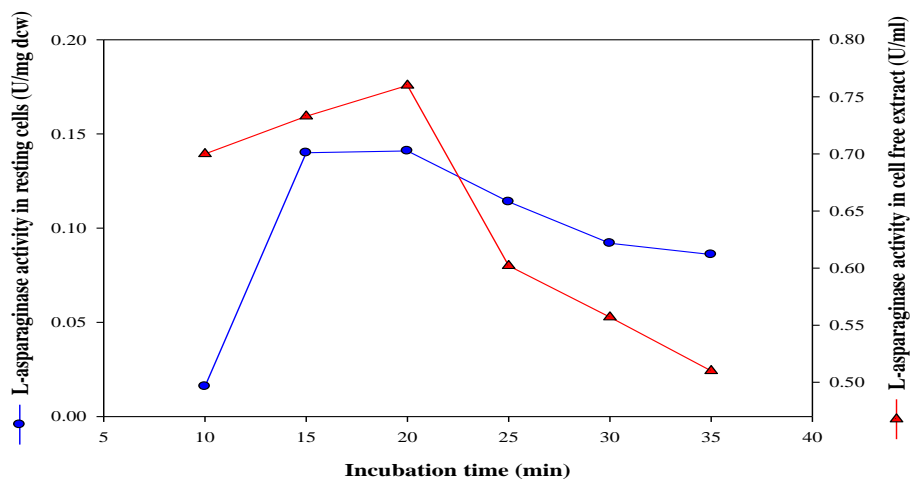


Fig.3 Effect of incubation time on L-asparaginase activity of *S. marcescens* MTCC 97

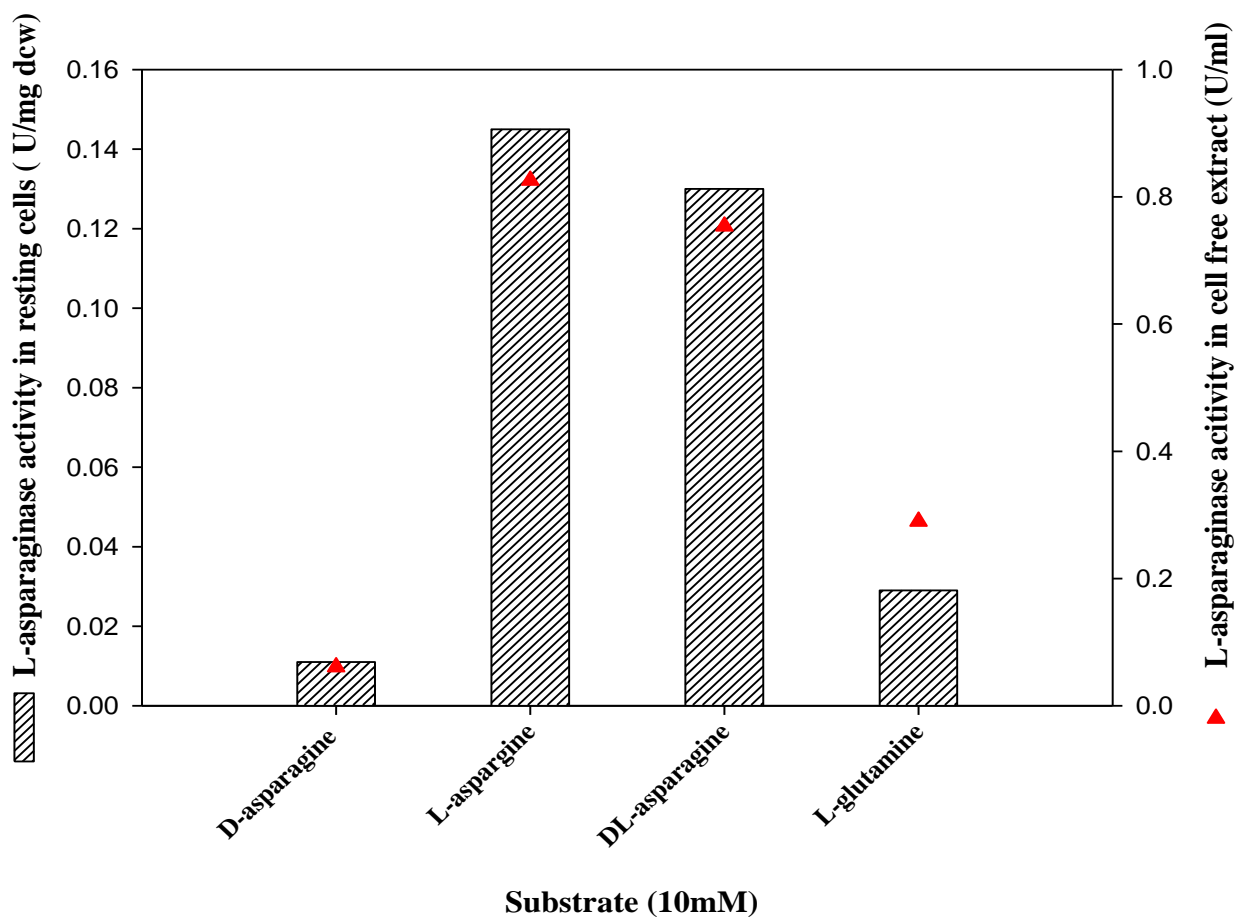


Fig.4 Substrate specificity of L-asparaginase from *S. marcescens* MTCC 97

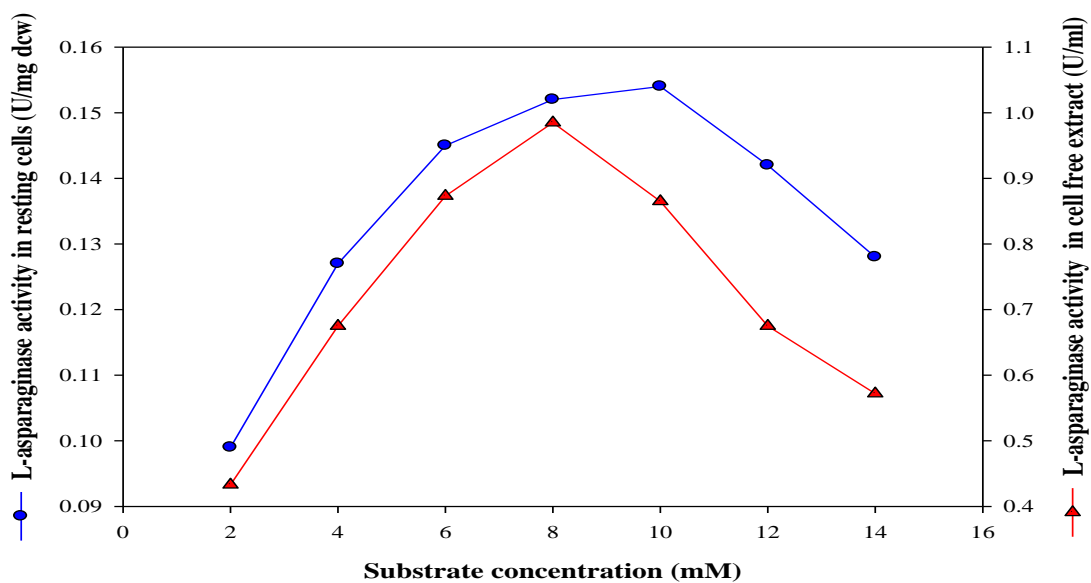


Fig.5 Effect of substrate concentration on L-asparaginase activity of *S. marcescens* MTCC 97

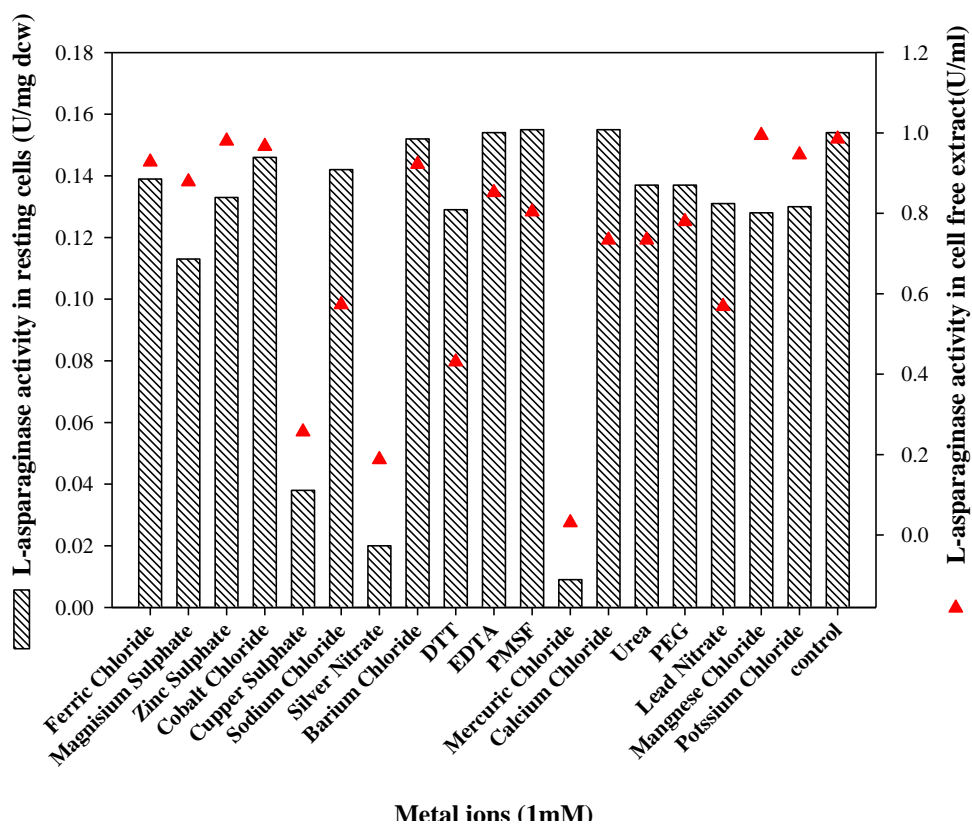


Fig.6 Effect of metal ions, chelating agents and other additives on L-asparaginase activity of *S. marcescens* MTCC 97

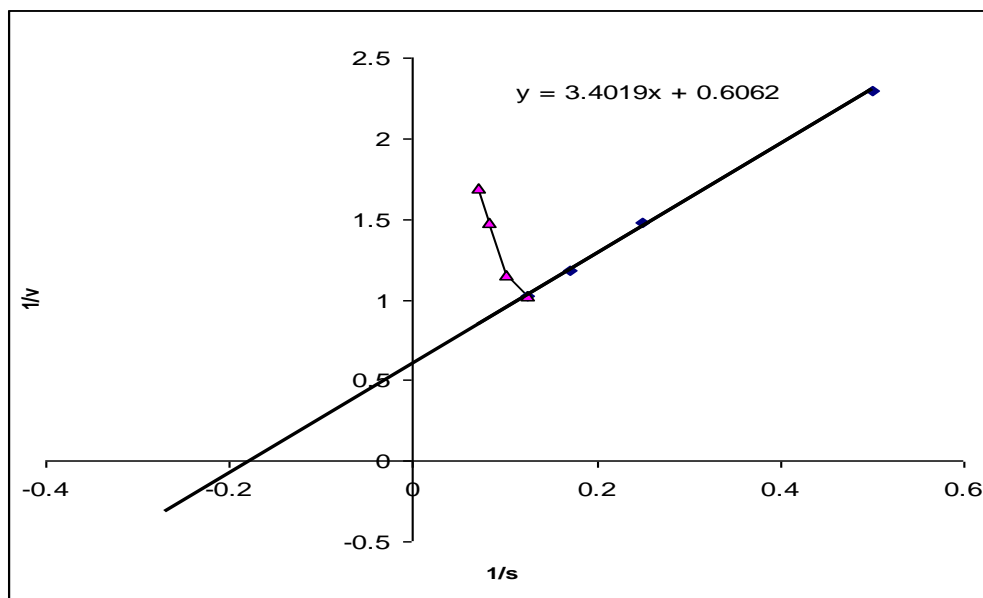


Fig.7 Line Weaver Burke plot for the L-asparaginase activity in the cell free extract of *S. marcescens* MTCC 97

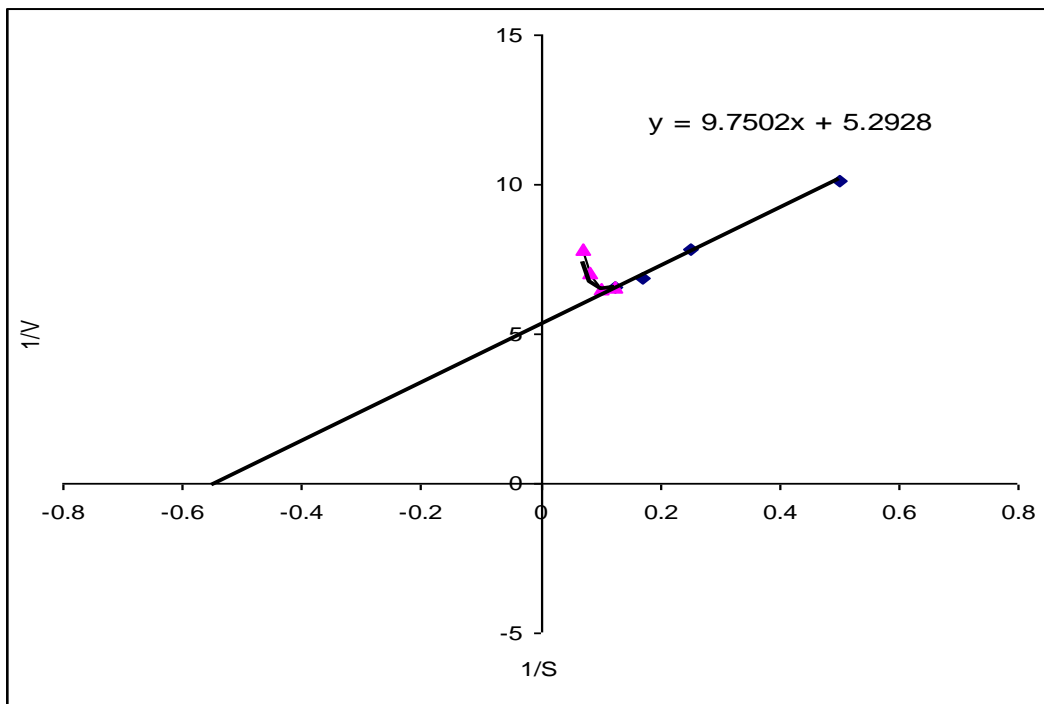


Fig.8 Line Weaver Burke plot for the L-asparaginase activity in the resting cells of *S. marcescens* MTCC 97

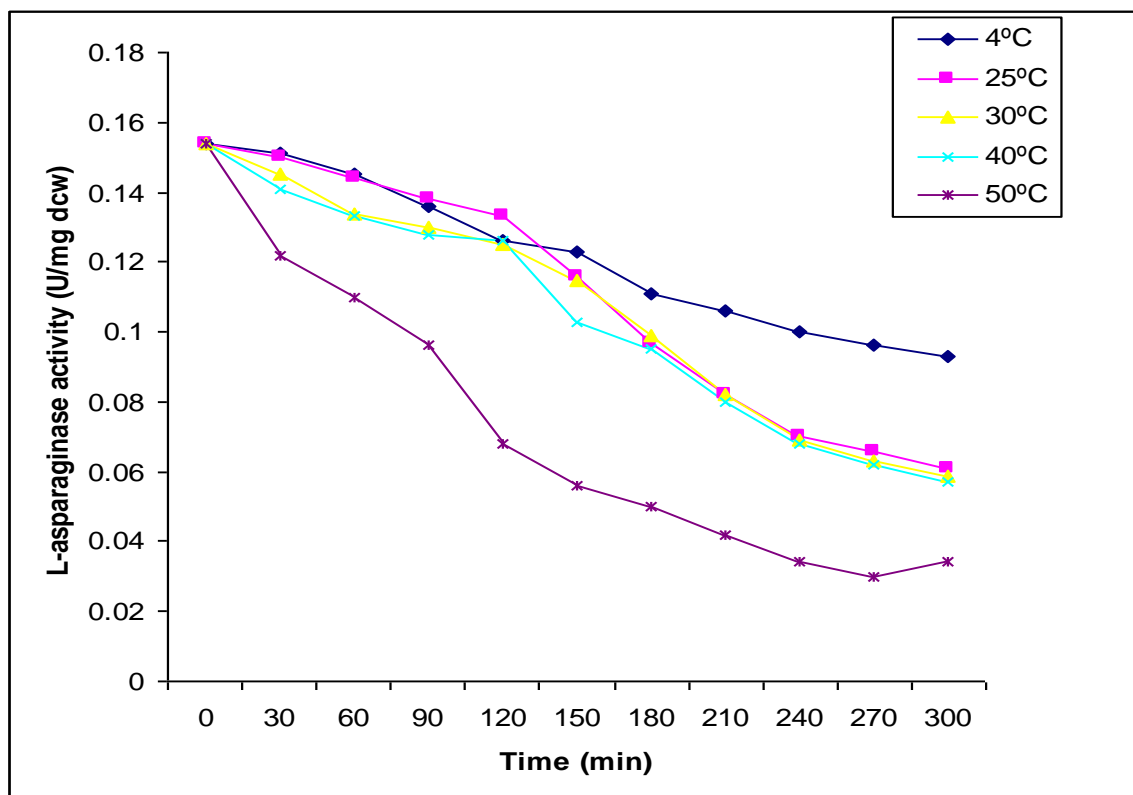


Fig.9 Stability profile of the resting cells of *S. marcescens* MTCC 97

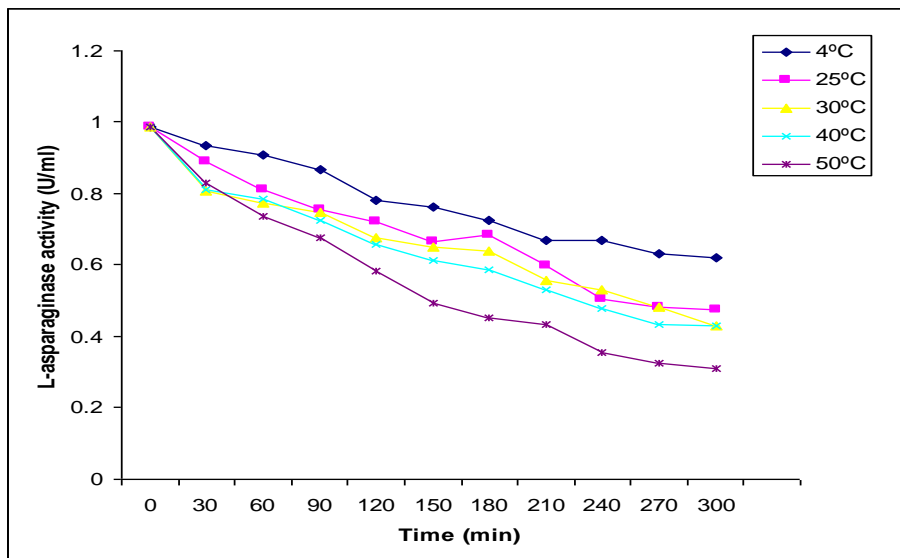


Fig.10 Stability profile of the cell free extract of *S. marcescens* MTCC 97

These findings suggest that the released L-asparaginase was more stable than the resting cells of *S. marcescens* MTCC 97. The thermal stability of L-asparaginase in resting cells and cell free extract of *S. marcescens* MTCC 97 were also established. At higher incubation temperature (50°C) the half-life of L-asparaginase in cell free extract and in resting cells was found to be 180 and 90 min, respectively.

These findings suggest that the released L-asparaginase was more stable than the resting cells of *S. marcescens* MTCC 97. The L-asparaginase activity in *Bacillus sp.* decreased sharply above 40°C and the enzyme was inactivated at 50°C with a half a life period of about 1 h [34]. Commercial preparation of L-asparaginase, Elspar, was found to be very stable at 45°C-55°C [31].

The half-life of Elspar at 60°C is 25 minutes. The L-asparaginase is the first enzyme with antitumor activity to be intensively studied in human beings. The major application of L-asparaginase is as an injectible drug for the treatment of tumors or Lymphoblastic Leukemia in human beings. The sensitivity of application demands high degree of purity of

this enzyme. Most of the microbial L-asparaginase is intracellular in nature except few, which are secreted outside the cells (Mohapatra BR *et al.*, 1995). Hence, the disintegration of resting cells of *S. marcescens* MTCC 97 or any L-asparaginase producing microorganisms seems to be necessary and first step for large scale commercial production of this enzyme.

Discussion

On comparison with the various methods (enzymatic, chemical and physical) used for the cell disintegration of the resting cells of *S. marcescens* MTCC 97, sonication was found to be the most effective method for the release of intracellular L-asparaginase from *S. marcescens* MTCC 97 with the maximum specific activity 0.047 U/mg of protein (Table 15).

The recovery of L-asparaginase was found to be 68% with a loss of only 8%. Amongst all the methods used, the bead beater and sonication were found to be the two most effective methods for the release of intracellular L-asparaginase from *S. marcescens* MTCC 97. The Bead beater was

found to be an efficient means for cell lysis. Disintegration of resting cells of *S. marcescens* MTCC 97 by bead beater using 0.1mm glass beads leads to the 50% recovery of the enzyme. The amount of released protein was found to be 16.67mg/ml. However, the loss of enzyme activity using bead beater was found to be very high (48%) possibly due to the denaturation of enzyme by the generation of heat.

Bead agitation or bead milling has been frequently used in large scale to medium scale preparation of intracellular protein from microorganisms in which harvested cells are vigorously agitated with beads in a closed chamber [15]. Protein release in these devices depend upon the cell disruption caused by shear forces and collision between beads and can be described as first order process (Melenders AV *et al.*, 1992).

The recovery of L-asparaginase using sonicator was found to be maximum (68%) with overall loss of 8% in activity in 9 min. The recovery of L-asparaginase has found to be more than 80% by 10 min sonication in case of a recombinant strain of *E. coli* (Krasotkina J *et al.*, 2004). Sonication being most efficient in the recovery of this membrane bound enzyme, recommended for its extraction from fresh bacterial biomass (Singh RS, 2013).

A remarkable feature of asparaginases is their pronounced antitumor activity, due to which these enzymes have found wide application in medicine as effective antitumor agents for the treatment of Acute Lymphoblastic Leukemia, lymphosarcomas, and reticulum cell sarcomas. Asparaginases are often successfully used when other antitumor drugs are ineffective. The antitumor effect of asparaginases is attributed to their ability to suppress asparagine metabolism, which is necessary for tumor cell growth (Sokolov NN

et al., 2000). The modified asparaginase retained 57% of initial activity. A simple and efficient pegylation procedure can be used for production of asparaginase with improved therapeutic properties (Kuchumova AV *et al.*, 2006).

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Compliance with Ethical Standards and Conflict of interest

The authors declare no conflicts of interest associated with this manuscript.

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