Isolation and screening of \( L \)-asparaginase free of glutaminase and urease from fungal sp.

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Abstract \( L \)-asparaginase is a chemotherapeutic drug used in the treatment of acute lymphoblastic leukaemia (ALL), a malignant disorder in children. \( L \)-Asparaginase helps in removing acrylamide found in fried and baked foods that is carcinogenic in nature. \( L \)-Asparaginase is present in plants, animals and microbes. Various microorganisms such as bacteria, yeast and fungi are generally used for the production of \( L \)-asparaginase as it is difficult to obtain the same from plants and animals. \( L \)-Asparaginase from bacteria causes anaphylaxis and other abnormal sensitive reactions due to low specificity to asparagine. Toxicity and repression caused by bacterial \( L \)-asparaginase shifted focus to eukaryotic microorganisms such as fungi to improve the efficacy of \( L \)-asparaginase. Clinically available \( L \)-asparaginase has glutaminase and urease that may lead to side effects during treatment of ALL. Current work tested 45 fungal strains isolated from soil and agricultural residues. Isolated fungi were tested using conventional plate assay method with two indicator dyes, phenol red and bromothymol blue (BTB), and results were compared. \( L \)-Asparaginase activity was measured by cultivating in modified Czapek–Dox medium. Four strains have shown positive result for \( L \)-asparaginase production with no urease or glutaminase activity, among these \( C_7 \) has high enzyme index of 1.57 and \( L \)-asparaginase activity of 33.59 U/mL. \( L \)-Asparaginase production by \( C_7 \) was higher with glucose as carbon source and asparagine as nitrogen source. This is the first report focussing on fungi that can synthesize \( L \)-asparaginase of the desired specificity. Since the clinical toxicity of \( L \)-asparaginase is attributed to glutaminase and urease activity, available evidence indicates variants negative for glutaminase and urease would provide higher therapeutic index than variants positive for glutaminase and urease.

Keywords \( L \)-Asparaginase · \( L \)-Glutaminase · Urease · Glutaminase-free \( L \)-asparaginase · Urease and glutaminase-free \( L \)-asparaginase

Abbreviations
ALL Acute lymphoblastic leukaemia
BTB Bromothymol blue
\( L \)-Asn \( L \)-Asparagine
\( L \)-Gln \( L \)-Glutamine
MCD Modified Czapek–Dox

Introduction

\( L \)-Asparaginase is an amidohydrolase that catalyses \( L \)-asparagine to \( L \)-aspartate and ammonia. \( L \)-Asparaginase is found to have tumour inhibitory properties. It is mainly used in the treatment of acute lymphoblastic leukaemia (ALL). Normal cells can synthesize \( L \)-asparagine with the help of asparagine synthetase, whereas certain sensitive malignant cells cannot synthesize it by itself and require an external source of \( L \)-asparagine for growth. During the treatment of ALL with \( L \)-asparaginase, all the circulating asparagine in the body of the patient get hydrolysed to aspartic acid and ammonia preventing the absorption of
asparagine by tumour cells thereby depriving the tumour cells of their extracellular source of L-asparagine (Broome 1961). L-Asparaginase is commonly used as a combination chemotherapy drug for the treatment of acute lymphoblastic leukaemia (ALL) in adults and children and non-Hodgkin’s lymphoma in children (Mashburn and Wriston 1964). L-Asparaginase also reduces acrylamide formation in food by selectively hydrolysing asparagine to aspartic acid and ammonia without affecting other amino acids, retaining food quality. Application of L-asparaginase enzyme (2 U/g) successfully reduced acrylamide content by 90% in potato products that have high asparagine content (Friedman 2003; Ciesarová et al. 2006).

L-Asparaginase is widely present in plants, animals and microbes but not in humans. Microbes are a better source for the production of enzyme as they are easy to cultivate and manipulate (Kumar and Sobha 2012). Clinically three asparaginase formulations are available, two from bacterial sources Escherichia coli (E. coli asparaginase) and Erwinia chrysanthemi (Erwinia asparaginase) and PEGylated form of E. coli asparaginase. L-Asparaginase therapy has side effects such as anaphylaxis, coagulation abnormality, thrombosis, liver dysfunction, pancreatitis, hyperglycaemia and cerebral dysfunction, etc. These side effects are either due to the production of anti-asparaginase antibody in the body or L-glutaminase activity of L-asparaginase enzyme (Haskell et al. 1969; Mahajan et al. 2012). Toxicity of L-asparaginase is mainly due to the fact that the enzyme preparations are amidohydrolase, not L-asparaginase. Clinically available L-asparaginase shows notable hydrolysis of L-glutamine and D-asparagine, signifying multiple enzyme activities contaminating enzyme preparation and difficult to eliminate other enzymes (Campbell and Mashburn 1969). Notwithstanding numerous studies on bacterial L-asparaginase, treatment with it sometimes results in hypersensitive reactions such as anaphylactic shock. L-Asparaginase isolated from filamentous fungi, Aspergillus terreus showed a greater carcinostatic effect on static tumour (De-Angeli et al. 1970). Similar effect was observed when L-asparaginase from deuteromycetes Fusarium tricinctum was purified which regressed lymphosarcoma in mice (Scheetz et al. 1971). Later purified extracellular L-asparaginase from A. terreus was conjugated with polyethylene glycol and it did not indicate any glutaminase activity (Loureiro et al. 1971). Later purified extracellular L-asparaginase from A. terreus was conjugated with polyethylene glycol and it did not indicate any glutaminase activity (Loureiro et al. 1971). Later purified extracellular L-asparaginase from A. terreus was conjugated with polyethylene glycol and it did not indicate any glutaminase activity (Loureiro et al. 1971).}

Materials and methods

L-Asparagine was procured from Sigma-Aldrich, India. Other chemicals used were of analytical grade. Aspergillus terreus MTCC 1782 was obtained from Microbial Type Culture Collection Centre and Gene Bank, Institute of Microbial Technology, Chandigarh, India.

Isolation of fungi from collected samples

Soil samples were collected from different locations of Vizag, Kanyakumari and Kerala as mentioned in Table 1. Soil and substrate samples were collected in air-tight containers and kept at room temperature in laboratory. Fungi were isolated by serial dilution of soil and agricultural residue, and plated on modified Czapek–Dox (MCD) agar plates with L-asparagine as a sole nitrogen source and incubated at 30°C for 96 h. Fungal strains showing change were selected and grown on potato dextrose slants.

Screening studies

Semi-quantitative assay for L-asparaginase producing fungi

MCD medium with composition glucose 2 g/L, L-asparagine 10 g/L, KH2PO4 1.52 g/L, KCl 0.52 g/L, MgSO4·7H2O 0.52 g/L, FeSO4·7H2O trace, ZnSO4·7H2O trace, CuNO3·3H2O trace and agar 18 g/L was prepared (Gulati et al. 1997). About 2.5% (w/v) stock solution of the phenol red dye was prepared and MCD medium was supplemented with 0.009% phenol red dye. 0.04% (w/v) stock solution of the bromothymol blue dye was prepared and 0.007% BTB dye was supplemented in MCD medium. Final pH of the media was adjusted to 5.5 using 1 M NaOH (Mahajan et al. 2013). Prepared media was autoclaved and poured into pre-sterilized plates. Control plates were prepared with NaNO3 as sole nitrogen source. MCD plates were inoculated with isolated fungi as test organism and A. terreus MTCC 1782 as positive test. Colony diameter and zone diameter for all the test organisms were measured after 96 h incubation.
organisms were measured and respective zone index was calculated after 72 h of incubation. Morphological observation of positive isolates was done by the method of staining and observing fungal spores using lacto phenol cotton blue staining solution.

Plate assay for L-glutaminase

L-Glutaminase activity of the fungal strains was detected by supplementing MCD medium with L-Gln as sole nitrogen source. Test strains were inoculated and observed for colour change from yellow to pink in case of phenol red dye and yellow to blue for BTB dye.

Plate assay for urease

MCD medium without nitrogen source was autoclaved and 1 % filter-sterilized urea solution was added to MCD media for detection of urease-producing fungi. Test strains were inoculated and observed for change in the colour of the medium.

Quantitative detection of L-asparaginase assay

Quantitative determination of L-asparaginase activity was carried out using selective strains (MTCC 1782, C3, C7, W3 and W5). These strains were cultivated on potato dextrose slants at 30 °C for 96 h. From these, 1 mL of conidial suspension was inoculated into Erlenmeyer flask containing 50 mL of MCD medium with initial pH of 6.2. Flasks were incubated at 30 °C at 180 rpm for 96 h. Samples were withdrawn every 24 h to determine enzyme activity.

Effect of carbon and nitrogen sources

To investigate the effect of different carbon sources on L-asparaginase production, fructose, glucose, maltose, sucrose, lactose and starch were added at concentration of 0.2 % (w/v) to the MCD medium. Influence of nitrogen source on asparaginase production was obtained by substituting asparagine of MCD medium with yeast extract, peptone and sodium nitrate at a concentration of 1 % (w/v). 1 mL of C7 suspension (with 10^9 cells/mL) was inoculated and flasks were incubated at 30 °C at 180 rpm for 72 h. Supernatant was used to determine asparaginase activity and protein content. The effect of inoculum volume at different levels was investigated by employing C7 in MCD medium.

L-Asparaginase activity is obtained by measuring the ammonia liberated using Nesslerization method by spectrophotometric analysis at 425 nm as described by Kumar et al. (2011). Enzyme assay mixture consisted of 900 μL of freshly prepared L-asparagine (40 mM) in Tris–HCl buffer (pH 8.6) and 100 μL of enzyme filtrate, incubated at 37 °C for 30 min and reaction was stopped by adding 100 μL of 1.5 M trichloroacetic acid (TCA). The reaction mixture was centrifuged at 10,000 rpm for 5 min at 4 °C to remove the precipitates. The ammonia released in the supernatant was determined using colorimetric technique by adding 200 μL of Nessler’s reagent into the sample containing 200 μL of supernatant and 1.6 mL distilled water. This mixture was vortexed and incubated at room temperature for 20 min. Absorbance was measured at 425 nm against the blanks that received TCA before the addition of enzyme. The ammonia liberated in the reaction was determined based on the standard curve obtained using ammonium sulfate. One unit (IU) of L-asparaginase activity was defined as the amount of the enzyme that liberates 1 μM of ammonia per min at 37 °C, using asparagine as substrate.

Extracellular protein content was determined using Lowry method (Lowry et al. 1951). Specific activity is expressed as unit enzyme activity per mg of protein.

Results and discussion

Isolation of fungal species

A total of 45 fungal species were isolated on the basis of zone formation from soil, wheat bran, rice husk, cotton.

Table 1 Fungal species isolated from diverse sources for the production of L-asparaginase

| Source          | Region             | Description                                                                 | No. of isolates |
|-----------------|--------------------|-----------------------------------------------------------------------------|-----------------|
| Soil            | Kanyakumari        | Soil samples were collected from different locations of sea shore           | 10              |
|                 | Vizag              | Soil samples were collected from different locations of sea shore           | 8               |
|                 | Kerala             | Soil samples were collected from Western ghats, corresponding to coordinates 9.8403°N, 77.0353°E | 9               |
| Agricultural residues | Cottonseed oil cake | Substrates were collected from local market                               | 18              |
|                 | Rice husk          |                                                                             |                 |
|                 | Wheat bran         |                                                                             |                 |
|                 | Red gram animal feed |                                                                          |                 |
seed oil cake and red gram feed. Among isolated fungi, 34 isolates were able to grow in secondary screening with MCD medium containing different nitrogen sources. Out of 45 fungal isolates, 27 were from soil implying that 60% of isolated fungi were from soil samples, rest from agricultural residues. Aspergillus sp., Penicillium sp., Trichophyton sp. and Onychocola sp. were predominant fungi isolated from the soil samples. Rhizopus sp. and Fusarium sp. were isolated from agricultural residues. Aspergillus sp. was the most dominant species among fungi isolated from soil and agricultural residues. These results were comparable to previously reported studies (Qiao et al. 2008; Tančinová and Labuda 2009).

Screening studies

Current study involved the screening of isolated fungi for the existence of three industrially important enzymes using phenol red and BTB dye. For screening of L-asparaginase, L-glutaminase and urease enzyme, MCD supplemented with L-Asn, L-Gln and urea, respectively, as the sole nitrogen sources are used. These amidohydrolases cleave amine groups and liberate aspartic acid and ammonia in case of L-asparaginase, glutamic acid and ammonia in case of L-glutaminase and carbonic acid and ammonia if urease is produced (as shown in Fig. 1). Ammonia liberated in the medium further reacts with water to produce NH₂OH resulting in increase in the pH of the medium.

Phenol red dye is yellow at acidic pH and turns pink at alkaline pH; presence of pink colour zone around the colonies on MCD plates with different nitrogen sources is due to the liberation of corresponding enzyme (Gulati et al. 1997). Thirty-four isolates showed pink zone around the colonies indicating increase in pH. In Fig. 2, last column shows presence of pink-coloured zone around fungal isolates S₃₄, W₃, W₅, C₃ and C₇ in L-Asn plates indicating L-asparaginase activity. These isolates did not show any colour change in plates containing L-Gln connoting the absence of L-glutaminase. S₃₄ and MTCC 1782 isolates produce the urease enzyme which is confirmed by the pink-coloured zone around the colony in plates with urea as nitrogen source. MTCC 1782 strain showed pink-coloured zone when grown on L-Asn, L-Gln and urea indicating that strain produces three enzymes. Strains W₃, W₅, C₃ and C₇ show pink colour zone only on L-asparagine plate, indicating strains are free of L-glutaminase and urease. To ensure reproducibility, all the isolates were screened with BTB as both the dyes are formulated for screening the hydrolysis of L-Gln, L-Asn and urea. Among phenol red and BTB, 0.007% of BTB dye showed sharp colour contrast zone, ranging from yellow at acidic pH, green at neutral pH to blue at alkaline pH (Mahajan et al. 2013). MCD plates with different substrates supplemented with BTB dye is shown in Fig. 3. After 72 h of incubation, thirty-four isolates showed blue-coloured zone around the colonies indicating increase in pH.
In comparison with phenol red, hydrolysed and unhindered enzymes were clear and precise in MCD supplemented with BTB. Methyl red was incorporated as pH indicator in the recent study to screen L-asparaginase- and L-glutaminase-producing microorganism (Dhale Dhale and Mohan Kumari 2014). Enzyme activity is calculated semi-quantitatively by relative ratio of zone diameter to colony diameter. Level of enzyme production was indicated by zone index. The comparison of zone index values of isolates S 3.4, W 3, W 5, C 3, C 7 and Aspergillus MTCC 1782 strain using phenol red and BTB dye is given in Table 2. Using this qualitative plate assay, rapid screening of the fungi for the synthesis of the enzyme by direct visualization and activity of the enzyme can be measured (Hankin and Anagnostakis 1975). Gulati et al. revealed that equivalent relation exists between zone index and enzyme activity measured from broth. In the current work, enzyme index varied from 0.8 to 4, which is in line with study conducted by Shrivastava et al. (2010). Enzyme index of C 7 is 1.57 with colony diameter of 3.5 cm and zone diameter of 5.5 cm which is lower than that of MTCC 1782 strain with enzyme index of 2.40. Out of 34 isolated fungal species, only 4 isolates showed L-asparaginase free of L-glutaminase and urease as shown in Table 3. Isolated fungi (S3.4, W3, W5, C3 and C7) were cultured in PDA slants, later morphologically identified as Curvularia sp., Rhizopus sp. and Aspergillus sp., respectively (Ellis et al. 2007).

The L-asparaginase activity of the four isolated strains with no glutaminase and urease activity is measured in liquid broth studies along with MTCC 1782 (shown in Fig. 4). MTCC 1782 strain is found to have the highest activity at 72 h with L-asparaginase activity of 34.45 U/mL and specific activity of 71.92 U/mg. Reported activity for optimized Aspergillus terreus MTCC 1782 was 40.186 IU/
Among the four isolated strains, C7 has the highest activity of 33.59 U/mL and specific activity of 64.85 U/mg. Hence, the medium has to be developed and optimized for L-asparaginase production from C7 to enhance L-asparaginase activity. All the strains exhibit the maximum activity at 72 h.

Table 2  L-Asparaginase enzyme index measurement using phenol red and bromothymol blue amended in MCD medium after 72 h incubation and species observed under light microscope

| Isolate   | Phenol red | Bromothymol blue | Species       |
|-----------|------------|------------------|---------------|
|           | Colony diameter (cm) | Zone diameter (cm) | Zone index | Colony diameter (cm) | Zone diameter (cm) | Zone index |
| S3.4      | 4.30       | 6.70             | 1.56         | 3.30          | 6.80             | 2.06       | Curvularia sp. |
| W3        | 2.40       | 2.40             | 1.00         | 3.70          | 4.70             | 1.27       | Rhizopus sp.   |
| W5        | 8.80       | 8.80             | 1.00         | 2.20          | 2.60             | 1.18       | Rhizopus sp.   |
| C3        | 3.00       | 3.00             | 1.00         | 3.80          | 5.50             | 1.45       | Aspergillus sp.|
| C7        | 3.00       | 4.60             | 1.53         | 3.50          | 5.50             | 1.57       | Aspergillus sp.|
| MTCC 1782 | 2.50       | 6.00             | 2.40         | 2.50          | 6.00             | 2.40       | Aspergillus sp.|

mL (Baskar and Renganathan 2012). The assay for screening L-asparaginase-producing fungi amended with different substrates, on plate supplemented with BTB dye. a–d S3.4 isolate grown on plates containing NaNO3, urea, l-Gln and l-Asn; e–h C7 isolate grown on plates containing NaNO3, urea, l-Gln and l-Asn; i–l W3 isolate grown on plates containing NaNO3, urea, l-Gln and l-Asn; m–p C7 isolate grown on plates containing NaNO3, urea, l-Gln and l-Asn; q–t W3 isolate grown on plates containing NaNO3, urea, l-Gln and l-Asn; u–x Aspergillus terreus MTCC 1782 strain grown on plates containing NaNO3, urea, l-Gln and l-Asn; 1 S3.4, 2 C3, 3 W5, 4 C7, 5 W3, 6 MTCC 1782: microscopic images of isolates using light microscope x40 magnification.
Effect of carbon and nitrogen sources

Six different carbon sources (fructose, glucose, maltose, sucrose, lactose and starch) were evaluated for the L-asparaginase production by C_7 which is free of glutaminase and urease. Batch cultivation of C_7 in MCD medium using different carbon sources revealed distinctive variations on L-asparaginase production and specific activity (Fig. 5). In comparison to other carbon sources, C_7 produced maximum asparaginase (16.2 U/mL) when glucose is used as a carbon source; lactose, maltose and starch were the poorest carbon sources. Sucrose and fructose also supported…

| S. no. | Isolation source          | Isolate | Control (NaNO_3) | Urea | L-Asn | L-Gln |
|-------|---------------------------|---------|------------------|------|-------|-------|
| 1     | Soil from Vizag           | V_1     | –                | –    | +     | +     |
| 2     |                           | V_2     | –                | –    | +     | +     |
| 3     |                           | V_3     | –                | –    | +     | –     |
| 4     |                           | V_4     | –                | –    | +     | +     |
| 5     |                           | V_5     | –                | –    | +     | +     |
| 6     |                           | V_6     | –                | –    | +     | –     |
| 7     | Soil from Kanyakumari     | K_1     | –                | +    | +     | +     |
| 8     |                           | K_2     | –                | –    | +     | +     |
| 9     |                           | K_3     | –                | –    | +     | +     |
| 10    |                           | K_4     | –                | –    | +     | +     |
| 11    |                           | K_5     | –                | –    | +     | +     |
| 12    |                           | K_6     | –                | –    | +     | +     |
| 13    |                           | K_7     | –                | +    | +     | +     |
| 14    |                           | K_8     | –                | –    | +     | +     |
| 15    | Soil from Western Ghats   | S_1,1   | –                | –    | +     | +     |
| 16    |                           | S_1,4   | –                | +    | +     | +     |
| 17    |                           | S_2,1   | –                | +    | +     | +     |
| 18    |                           | S_3,4   | –                | +    | +     | +     |
| 19    |                           | S_4,1   | –                | +    | +     | +     |
| 20    | Red gram husk             | P_2     | –                | +    | +     | +     |
| 21    |                           | P_3     | –                | +    | +     | +     |
| 22    | Rice husk                 | R_1     | –                | +    | +     | +     |
| 23    |                           | R_3     | –                | +    | +     | +     |
| 24    | Wheat bran                | W_1     | –                | +    | +     | –     |
| 25    |                           | W_2     | –                | +    | +     | +     |
| 26    |                           | W_3     | –                | –    | +     | –     |
| 27    |                           | W_4     | –                | +    | +     | +     |
| 28    | Cotton seed oil cake      | C_1     | –                | +    | +     | +     |
| 29    |                           | C_3     | –                | –    | +     | –     |
| 30    |                           | C_4     | –                | +    | +     | –     |
| 31    |                           | C_5     | –                | +    | +     | –     |
| 32    |                           | C_6     | –                | +    | +     | +     |
| 33    |                           | C_7     | –                | –    | +     | –     |
| 34    |                           | MTCC 1782| –                | –    | +     | +     |

V (1–6) soil from Visakhapatnam (Vizag), K (1–8) soil from Kanyakumari, S (1.1, 1.4, 2.1, 3.4 and 4.1) soil from Western Ghats, P (2–4) red gram husk, R (1 and 3) rice husk, W (1–5) wheat bran, C (1, 3, 4, 5, 6 and 7) cotton seed oil cake.
L-asparaginase production to a significant degree but glucose acted as good inducer and primary source of carbon for biosynthesis of L-asparaginase using C7. Several reports suggest that glucose serves as a best carbon source for L-asparaginase production and a similar effect was observed for L-asparaginase production using Aspergillus and Fusarium strains (Baskar and Renganathan 2012; Hosamani and Kaliwal 2011). Effect of nitrogen compounds on L-asparaginase production by C7 was studied by supplementing nitrogen sources (asparagine, yeast extract, peptone and sodium nitrate) to MCD medium. C7 amended with asparagine favoured maximum enzyme production indicating L-asparagine itself acts as a nitrogen source and influence L-asparaginase production (Fig. 6). Peptone also supported the production of L-asparaginase to a substantial quantity. A considerable decrease in enzyme activity was observed when C7 was amended with yeast extract. Lower enzyme activity was detected in the media supplemented with sodium nitrate; on the contrary, a study in which Fusarium oxysporum has shown higher enzyme production with sodium nitrate as a nitrogen source (Tippani and Sivadevuni 2012). Effect of inoculum volume on L-asparaginase production by C7 is shown in Fig. 7. At low inoculum concentration, the L-asparaginase production was less, and the enzyme activity increased with increase in inoculum volume. At inoculum volume of 5 × 10^7 cells/mL, enzyme activity is 33.59 U/mL. With further increase in inoculum concentration, the biosynthetic activity decreased due to nutrient depletion.

Most of the L-asparaginase purified from various sources such as chillies and E. coli shows specificity towards both L-Gln and urea. Several studies reveal that specificity of L-asparaginase is important in selective depletion of asparagine-dependent tumour cells (Hill et al. 1967; Durden and Distasio 1981; Distasio et al. 1982). To reduce the toxic effects associated with bacterial L-asparaginase, fungi is preferred as being eukaryotic and evolutionarily closer to human. It can minimize the chances of immunological reactions (Shrivastava et al. 2012). Several fungal endophytes were isolated from various sources and tested for their ability to synthesize L-glutaminase-free L-asparaginase. L-Glutaminase-free L-asparaginase produced by endophytic fungi from seaweed was isolated, later identified as Fusarium, Alternaria sp., Aspergillus sp. and Colletotrichum sp. (Thangavel et al. 2013). Alternaria sp. endophytic fungi isolated from the leaf of Withania somnifera of Western Ghats is reported to show maximum L-asparaginase activity that is free of L-glutaminase (Nagarajan et al. 2014). In the current study, 45 fungi isolates were subjected to screening, with a view to assess the isolates for their ability to utilize different substrates as a nitrogen source. Twenty fungi isolates have shown the presence of urease, L-glutaminase and L-asparaginase
enzyme. Four isolates have shown the presence of L-asparaginase free of urease and L-glutaminase, and six isolates presented L-asparaginase free of L-glutaminase with presence of urease using plate assay. Fungal isolates were selected on the basis of zone formation around the colonies, when grown on MCD with phenol red or BTB as a pH indicator. The change in colour stated the accumulation of ammonia which resulted due to the hydrolysis of amidohydrolase (Singh and Srivastava 2012). Fungi secrete numerous enzymes into the medium and regulation of other contaminating enzymes would make it possibly the preferred drug in the treatment of cancer.

Current preparation of asparaginase used in treatment protocols are E. coli asparaginase, its PEGylated form and Erwinia asparaginase; several studies on different other sources of asparaginase have yielded encouraging outcomes. Further studies and regulatory supports will allow the introduction of new asparaginase drugs with potential benefits to patients. Fungal strain, namely C7, which is an L-asparaginase (free of L-glutaminase and urease)-producing strain has shown highest enzyme activity of 33.59 U/mL with carbon source as glucose; asparaginase as nitrogen source at inoculum volume of $5 \times 10^7$ cells/mL is to be considered for further study on purification and characterization of L-asparaginase enzyme.

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Compliance with ethical standards

Conflict of interest We hereby declare that there has not been any conflict of interest at any point of time during the preparation of this manuscript.

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