

## Asparaginase and Glutaminase Activities of Micro-organisms

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(Received 14 September 1972; revised 28 November 1972)

### SUMMARY

L-Asparaginase and L-glutaminase activities were detected in many micro-organisms and the distribution of these activities was found to be related to the classification of micro-organisms.

Among 464 bacteria, the activities occurred in many Gram-negative bacteria and in a few Gram-positive bacteria. Most members of the family Enterobacteriaceae possessed L-asparaginase. L-Asparaginase and L-glutaminase occurred together in a large proportion of pseudomonads. Among Gram-positive bacteria many strains of *Bacillus pumilus* showed strong L-asparaginase activity. Amidase activities were also observed in several strains in other families.

L-Asparaginase activity was not detected in culture filtrates of 261 strains of species of the genera *Streptomyces* and *Nocardia*, but L-asparaginase and L-glutaminase were detected when these organisms were sonicated.

The amidase activities in culture filtrates of 4158 fungal strains were tested. All the strains of *Fusarium* species formed L-asparaginase. Organisms of the genera *Hypomyces* and *Nectria*, which are regarded as the perfect stage of the genus *Fusarium*, also formed L-asparaginase. Several *Penicillium* species formed L-asparaginase. Two organisms of the family Moniliaceae formed L-glutaminase together with L-asparaginase, and a few ascomycetous fungi formed L-asparaginase or L-glutaminase.

Among 1326 yeasts, L-asparaginase or L-glutaminase occurred frequently in certain serological groups of yeasts: VI (Hansenula) group, *Cryptococcus* group and *Rhodotorula* group. Many strains of *Sporobolomyces* species also showed L-asparaginase activity. Several strains of *Cryptococcus* and *Rhodotorula* group possessed L-glutaminase and L-asparaginase. L-Glutaminase alone was formed in many strains of *Candida scottii* and *Cryptococcus albidus*, both of which are related to Basidiomycetes.

### INTRODUCTION

Kidd (1953) observed that certain transplanted murine leukaemias were suppressed by treatment with guinea-pig serum. Broome (1961) proposed that the antileukaemia activity was attributable to the action of L-asparaginase (EC. 3.5.1.1). Several investigations have confirmed this proposal (Broome, 1963*a, b*; Old, Boyse, Campbell & Daria, 1963; Schwartz, Reeves & Broome, 1966; Yellin & Wriston, 1966*a, b*).

It is well known that certain micro-organisms possess L-asparaginase and some of them show antileukaemia activity. Mashburn & Wriston (1964) proved that partially purified L-asparaginase of *Escherichia coli* suppressed 6C3HED lymphosarcoma of mice. L-Asparaginase of *Escherichia coli* was purified and is now being used experimentally as an anti-leukaemia agent for human leukaemia patients, and has been found to be very effective for acute lymphatic leukaemia of children (Hill *et al.* 1967; Oettgen *et al.* 1967). Besides L-asparaginase of *Escherichia coli*, the enzymes of organisms of Enterobacteriaceae have been found to suppress the growth of 6C3HED lymphosarcoma (Rowley & Wriston, 1967;

Eremenco, Evseev & Nikolaev, 1968; Wade *et al.* 1968; Peterson & Ciegler, 1969*a*; Yamada, 1970; Boyd & Phillips, 1971; Tosa *et al.* 1971). Reddy, Jayaram, Sirsi & Ramakrishnan (1969) found that L-asparaginase from *Mycobacterium tuberculosis* suppressed Yoshida ascites sarcoma in rats. De-Angeli *et al.* (1970) found that L-asparaginase of *Aspergillus terreus* suppressed Walker 256 ascites carcinoma in rats.

Anticancer activity of L-glutaminase (EC 3.5.1.2) has also been proposed. Greenberg, Blumenthal & Ramadan (1964) and El-Asmar & Greenberg (1966) observed that L-glutaminase of a pseudomonad inhibited the initial growth of several murine carcinomas but had little effect on the survival time of animals. Recently, Roberts, Holcenberg & Dolowy (1970) found that L-glutaminase of a Gram-negative rod-shaped bacterium suppressed Ehrlich ascites carcinoma.

L-Asparaginase from *Saccharomyces cerevisiae* (Greenberg *et al.* 1964), *Bacillus coagulans* (Mashburn & Wriston, 1964; Law & Wriston, 1971), *Alcaligenes faecalis* (Yamada, 1970; Tosa *et al.* 1971), *Candida utilis* (Sakamoto *et al.* 1970) and *Fusarium tricinctum* (Scheetz, Whelan & Wriston, 1971) had no antilymphoma activities.

L-Asparaginases from other sources, such as *Penicillium camemberti* (Dox, 1909), *Aspergillus niger* (Bach, 1928; Schmalfuss & Mothes, 1930), *Brucella abortus* (Altenbern & Housewright, 1954), pseudomonads (de Groot & Lichtenstein, 1960; Ramadan, El-Asmar & Greenberg, 1964), *Staphylococcus aureus* (Tsuji, 1957), *Bacillus stearothermophilus* (Manning & Campbell, 1957), *Mycobacterium avium* (Tsuji, 1957), and *Rhodopseudomonas capsulatum* (Tchan, Asano & Kobayashi, 1971), have not been tested for their antilymphoma activities. Recently, Peterson and Ciegler (1969*b*) and Wade, Robinson & Phillips (1971) surveyed the L-asparaginase and L-glutaminase activities of a number of bacteria and found a wide distribution of both activities. Arima, Sakamoto, Araki & Tamura (1972) noticed that many bacteria, fungi and yeasts produced extracellular L-asparaginase.

L-Asparaginase activity occurs in a number of micro-organisms. However, the anti-6C3HED lymphosarcoma activity was shown only by the enzymes from Enterobacteriaceae, and these enzymes are now used for clinical experiments. It may be anticipated that the injection of enzymes for a prolonged period induces the formation of antibody which neutralizes the enzyme activity. Anaphylaxis shock is also supposed to occur. With respect to L-asparaginase of *Escherichia coli*, several unfavourable side actions, including nausea, anorexia and liver dysfunction, have been pointed out (Haskel *et al.* 1969).

The present report shows the distribution of L-asparaginase and L-glutaminase activities among micro-organisms. The relationship between this distribution and the classification of micro-organisms is discussed.

#### METHODS

*Chemicals.* Nutrient broth, yeast extract and malt extract were obtained from Difco Laboratories, Detroit, Michigan, U.S.A.; Polypepton, from Daigo Nutritive Chemicals, Osaka, Japan; corn-steep liquor (CSL), from Corn Product Co., New York, U.S.A.; and Pharmamedia, from Traders Mill Co., Fort Worth, Texas, U.S.A. Other chemicals were purchased from Wako Pure Chemicals Co., Osaka, Japan.

*Organisms.* Most microbial strains were obtained from the Institute for Fermentation, Osaka. Several strains of *Pseudomonas aeruginosa* isolated from clinical sources were obtained from Medical School, Osaka University.

*Culture methods.* CSL-medium of Roberts, Burson & Hill (1968) was used for bacteria. ST-2 medium (pH 7.0) which was used for Streptomyces and Nocardia contained (g/l distilled water): soluble starch, 50; Polypepton, 10; beef extract, 5; and NaCl, 5. M-medium

(pH 6.0) used for yeasts and fungi contained (g/l distilled water): sucrose, 30; Polypepton, 5; beef extract, 5; yeast extract, 2; malt extract, 2;  $\text{KH}_2\text{PO}_4$ , 5;  $\text{K}_2\text{HPO}_4$ , 1.5;  $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$ , 0.5. DP-medium (pH 6.0) used for fungi contained (g/l distilled water): dextrin, 30; Pharmamedia, 40;  $\text{KH}_2\text{PO}_4$ , 5;  $\text{K}_2\text{HPO}_4$ , 1.5;  $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$ , 0.5.

Bacteria, actinomycetes and yeasts were cultivated in 4 ml of media in 14 mm diam. test tubes with shaking. Fungi were cultivated in 8 ml of media in 20 mm diam. test tubes with shaking. Cultivation temperature was 28 °C.

*Enzyme preparation.* The enzyme activities of bacteria were examined by using whole culture broth as the enzyme preparation. In the experiment shown in Table 2, bacterial suspension (bacteria harvested from 1 ml of culture were suspended in 1 ml of distilled water) and culture filtrate were used as the enzyme preparation. The activities of actinomycetes were examined by using culture filtrate or cell-sonicate (actinomycetes harvested from 1 ml of culture were suspended in 1 ml of distilled water, sonically disrupted for 3 min, and cell debris was centrifuged off at 12000 g for 30 min). The activities of fungi were examined by using culture filtrate, and those of yeasts by using whole culture broth as the enzyme preparations.

*Assay of L-asparaginase and L-glutaminase.* L-Asparaginase was assayed as follows. A reaction mixture containing 0.5 ml of 0.04 M-L-asparagine, 0.5 ml of 0.5 M-buffer, 0.5 ml of an enzyme preparation, and distilled water to a total volume of 2.0 ml was incubated at 37 °C for 30 min. The reaction was stopped by adding 0.5 ml of 1.5 M-trichloroacetic acid. Blank tubes were run by adding the enzyme preparation after the addition of trichloroacetic acid. To 3.7 ml of distilled water, 0.1 ml of the above mixtures and 0.2 ml of Nessler's reagent were added. After keeping the mixture at 15 to 20 °C for 20 min, extinction at 450 nm was measured with a Spectronic 20 colorimeter (Schimadzu Bausch & Lomb) using  $\frac{1}{2}$  inch cells, and the amount of released ammonia was determined. One international unit (i.u.) of L-asparaginase is the amount of enzyme which liberates 1  $\mu\text{mol}$  of ammonia in 1 min.

L-Glutaminase was assayed by using L-glutamine in place of L-asparagine.

For the examination of these amidase activities, acetate buffer, pH 5.6, and tris(hydroxymethyl)aminomethane.HCl buffer, pH 8.4, were used for bacterial enzymes and tris(hydroxymethyl)aminomethane.HCl buffer, pH 7.2, for the enzymes of actinomycetes, yeasts and other fungi.

*Measurement of growth.* Growth of bacteria was expressed in extinction at 600 nm of the cultures, which was measured by a Spectronic 20 colorimeter.

## RESULTS AND DISCUSSION

### *L-Asparaginase and L-glutaminase activities of bacteria*

For the examination of bacterial L-asparaginase and L-glutaminase activities, CSL-medium of Roberts *et al.* (1968) was chosen, since it generally gave good growth of a variety of bacteria with comparatively high amidase activities. The cultivation period was set at 40 h, since deviation of the amount of growth of bacteria was less at 40 h cultivation than at 16 h without significant difference in the specific activities, which is the activity per growth (see Table 2). Whole culture broth was employed as the enzyme preparation, because the total complement of the amidase in the cell could be measured without disrupting cells as judged by preliminary experiments using a number of bacteria, e.g. *Escherichia coli* (3 strains), *Proteus vulgaris*, *Serratia marcescens*, *Klebsiella aerogenes*, *Pseudomonas aeruginosa*, *Staphylococcus aureus*, *Sarcina lutea*, *Bacillus subtilis*, *B. cereus*, *B. pumilus*, *B. brevis*,

Table I. *Distribution of L-asparaginase and L-glutaminase among bacteria*

Bacterial strain	Number of strains							
	L-Asparaginase (i.u./ml)				L-Glutaminase (i.u./ml)			
	< 0.10	0.11- 0.50	0.51- 1.00	> 1.01	< 0.10	0.11- 0.50	0.51- 1.00	> 1.01
Pseudomonadaceae								
<i>Pseudomonas aeruginosa</i>	2	81	—	—	2	81	—	—
<i>P. chlororaphis</i>	—	2	—	—	—	2	—	—
<i>P. schuykilliensis</i>	1	—	—	1	1	—	—	1
<i>P. synxantha</i>	—	2	1	—	—	2	1	—
<i>P. fluorescens</i>	1	6	—	—	1	6	—	—
<i>P. pavonacea</i>	2	—	—	—	2	—	—	—
<i>P. iodinum</i>	1	—	—	—	1	—	—	—
<i>P. putida</i>	3	—	—	—	3	—	—	—
<i>P. ovalis</i>	3	1	—	—	3	1	—	—
<i>P. teatrolens</i>	1	—	—	—	1	—	—	—
<i>P. mildenbergii</i>	1	1	—	—	2	—	—	—
<i>P. fragi</i>	1	—	—	—	1	—	—	—
<i>P. putrefaciens</i>	3	2	—	—	5	—	—	—
<i>P. dacunhae</i>	2	4	—	—	2	4	—	—
<i>P. stutzeri</i>	1	—	—	—	1	—	—	—
<i>P. riboflavina</i>	1	—	—	—	1	—	—	—
<i>P. coronafaciens</i>	1	—	—	—	1	—	—	—
<i>P. tabaci</i>	1	—	—	—	1	—	—	—
<i>P. polycolor</i>	—	1	—	—	1	—	—	—
<i>P. marginalis</i>	—	1	—	—	—	1	—	—
<i>P. solanacearum</i>	1	—	—	—	1	—	—	—
<i>P. aureofaciens</i>	—	5	1	—	—	6	—	—
<i>P. multophila</i>	1	—	—	—	1	—	—	—
<i>P. diminuta</i>	1	—	—	—	1	—	—	—
<i>P. nitroreducens</i>	1	—	—	—	1	—	—	—
<i>P. vendrelli</i>	—	1	—	—	—	1	—	—
<i>P. melanogenum</i>	2	—	—	—	2	—	—	—
<i>P. azotoformans</i>	—	1	—	—	—	1	—	—
<i>P. trifolii</i>	1	—	—	—	1	—	—	—
<i>P. alkanolytica</i>	1	—	—	—	1	—	—	—
<i>P. desmolytica</i>	1	—	—	—	1	—	—	—
<i>P. graveolens</i>	—	1	—	—	1	—	—	—
<i>P. convexa</i>	—	1	—	—	—	1	—	—
<i>Aeromonas hydrophila</i>	3	—	—	—	3	—	—	—
<i>A. liquefaciens</i>	1	—	—	—	1	—	—	—
<i>Protaminobacter alboatlavus</i>	1	—	—	—	1	—	—	—
Spirillaceae								
<i>Spirillum lunatum</i>	—	1	—	—	1	—	—	—
<i>S. metamorphum</i>	—	1	—	—	1	—	—	—
Chlamydobacteriaceae								
<i>Sphaerotilus natans</i>	1	2	—	—	2	1	—	—
Azotobacteraceae								
<i>Azotobacter chroococcum</i>	1	—	—	—	1	—	—	—
<i>A. agilis</i>	1	—	—	—	1	—	—	—
<i>A. indicum</i>	1	—	—	—	1	—	—	—
Rhizobiaceae								
<i>Agrobacterium tumefaciens</i>	2	—	—	—	2	—	—	—
<i>A. radiobacter</i>	3	—	—	—	3	—	—	—
<i>Chromobacterium violaceum</i>	1	—	—	—	1	—	—	—
Achromobacteraceae								
<i>Alcaligenes faecalis</i>	1	2	1	—	3	1	—	—
<i>Achromobacter liquidum</i>	—	1	—	—	1	—	—	—

Table 1 (continued)

Bacterial strain	Number of strains							
	L-Asparaginase (i.u./ml)				L-Glutaminase (i.u./ml)			
	< 0.10	0.11- 0.50	0.51- 1.00	> 1.01	< 0.10	0.11- 0.50	0.51- 1.00	> 1.01
<i>A. delmarvae</i>	—	1	—	—	1	—	—	—
<i>Flavobacterium flavescens</i>	2	—	—	—	2	—	—	—
<i>F. suaveolens</i>	1	—	—	—	1	—	—	—
<i>F. esteroaromaticum</i>	1	—	—	—	1	—	—	—
<i>F. aquatile</i>	2	—	—	—	2	—	—	—
<i>F. heparinum</i>	1	—	—	—	1	—	—	—
<i>F. gasgenes</i>	1	—	—	—	1	—	—	—
<i>F. meningosepticum</i>	1	—	—	—	1	—	—	—
<b>Enterobacteriaceae</b>								
<i>Enterobacter cloacae</i>	—	1	—	—	—	1	—	—
<i>Escherichia coli</i>	—	11	5	28	38	6	—	—
<i>Klebsiella aerogenes</i>	6	6	—	—	10	2	—	—
<i>K. pneumoniae</i>	—	1	—	—	1	—	—	—
<i>Erwinia carotovora</i>	—	1	—	—	1	—	—	—
<i>E. aroideae</i>	1	1	—	—	2	—	—	—
<i>E. amylolytica</i>	—	1	—	—	1	—	—	—
<i>E. herbicola</i>	—	1	—	—	1	—	—	—
<i>Serratia marcescens</i>	—	7	—	—	7	—	—	—
<i>S. piscatorum</i>	—	1	—	—	1	—	—	—
<i>S. plymuthicum</i>	1	—	—	—	1	—	—	—
<i>S. indica</i>	—	1	—	—	1	—	—	—
<i>Proteus vulgaris</i>	—	—	4	—	4	—	—	—
<i>P. morganii</i>	1	—	1	—	1	1	—	—
<i>P. mirabilis</i>	—	—	2	—	2	—	—	—
<i>Shigella paradysenteriae</i>	1	—	—	—	1	—	—	—
<b>Brucellaceae</b>								
<i>Brucella bronchiseptica</i>	2	—	—	—	1	1	—	—
<b>Micrococcaceae</b>								
<i>Micrococcus luteus</i>	1	—	—	—	1	—	—	—
<i>M. lysodeikticus</i>	1	—	—	—	1	—	—	—
<i>M. roseus</i>	1	—	—	—	1	—	—	—
<i>M. variabilis</i>	1	—	—	—	1	—	—	—
<i>M. flavus</i>	1	—	—	—	1	—	—	—
<i>M. varians</i>	1	—	—	—	1	—	—	—
<i>M. subflavus</i>	1	—	—	—	1	—	—	—
<i>M. marginata</i>	1	—	—	—	1	—	—	—
<i>M. ureae</i>	1	—	—	—	—	1	—	—
<i>M. rubens</i>	1	—	—	—	1	—	—	—
<i>M. glutamicus</i>	2	—	—	—	2	—	—	—
<i>M. cerificans</i>	—	1	—	—	—	1	—	—
<i>Staphylococcus aureus</i>	3	—	—	—	3	—	—	—
<i>S. epidermidis</i>	1	—	—	—	1	—	—	—
<i>Sarcina lutea</i>	1	—	—	—	1	—	—	—
<i>S. aurantiaca</i>	1	—	—	—	1	—	—	—
<b>Brevibacteriaceae</b>								
<i>Brevibacterium flavum</i>	1	—	—	—	1	—	—	—
<i>B. ammoniagenes</i>	4	1	—	—	4	1	—	—
<i>B. protomorphiae</i>	2	—	—	—	2	—	—	—
<i>B. saperdae</i>	1	—	—	—	1	—	—	—
<i>B. divaricatum</i>	3	—	—	—	3	—	—	—
<i>B. vitarum</i>	1	—	—	—	1	—	—	—
<i>B. stationis</i>	2	—	—	—	2	—	—	—
<i>B. insertum</i>	1	—	—	—	1	—	—	—
<i>B. leucinophagum</i>	—	1	—	—	—	1	—	—

Table 1 (continued)

Bacterial strain	Number of strains							
	L-Asparaginase (i.u./ml)				L-Glutaminase (i.u./ml)			
	<0·10	0·11- 0·50	0·51- 1·00	>1·01	<0·10	0·11- 0·50	0·51- 1·00	>1·01
<i>B. roseum</i>	1	—	—	—	1	—	—	—
<i>B. saccharolyticum</i>	1	—	—	—	1	—	—	—
<i>B. chang-fua</i>	1	—	—	—	1	—	—	—
<i>B. glutamicus</i>	1	—	—	—	1	—	—	—
<i>B. immariophilum</i>	1	—	—	—	1	—	—	—
<i>B. taipei</i>	1	—	—	—	1	—	—	—
<i>B. alkanolyticum</i>	1	—	—	—	1	—	—	—
<i>B. tesaceum</i>	1	—	—	—	1	—	—	—
<i>B. citreum</i>	1	—	—	—	1	—	—	—
<i>B. fuscum</i>	—	1	—	—	1	—	—	—
<i>B. linens</i>	3	1	—	—	4	—	—	—
<i>B. imperiale</i>	1	—	—	—	1	—	—	—
<i>B. pusillum</i>	1	—	—	—	1	—	—	—
<i>B. sulfureum</i>	1	—	—	—	1	—	—	—
<i>B. helvorum</i>	—	1	—	—	—	1	—	—
<i>Kurthia zopfii</i>	1	—	—	—	1	—	—	—
<b>Corynebacteriaceae</b>								
<i>Corynebacterium equi</i>	1	—	—	—	1	—	—	—
<i>C. insidiosum</i>	1	—	—	—	1	—	—	—
<i>C. nephridii</i>	2	—	—	—	2	—	—	—
<i>C. vesiculare</i>	1	—	—	—	1	—	—	—
<i>C. paurometabolum</i>	2	—	—	—	2	—	—	—
<i>C. sepedonicum</i>	1	—	—	—	1	—	—	—
<i>C. flaccumfaciens</i>	1	—	—	—	1	—	—	—
<i>C. hydrocarboclatum</i>	1	—	—	—	1	—	—	—
<i>C. diphtheriae</i>	1	—	—	—	1	—	—	—
<i>C. tritici</i>	1	—	—	—	1	—	—	—
<i>C. xerosis</i>	1	—	—	—	1	—	—	—
<i>C. faccians</i>	2	—	—	—	2	—	—	—
<i>Arthrobacter simplex</i>	2	—	—	—	2	—	—	—
<i>A. globiformis</i>	2	—	—	—	2	—	—	—
<i>A. pascens</i>	1	—	—	—	1	—	—	—
<i>A. aurescens</i>	1	—	—	—	1	—	—	—
<i>A. atrocyaneus</i>	1	—	—	—	1	—	—	—
<i>A. citreus</i>	1	—	—	—	1	—	—	—
<i>A. ramosus</i>	1	—	—	—	1	—	—	—
<i>Microbacterium lacticum</i>	1	—	—	—	1	—	—	—
<b>Bacillaceae</b>								
<i>Bacillus megaterium</i>	6	—	—	—	6	—	—	—
<i>B. cereus</i>	13	—	—	—	13	—	—	—
<i>B. anthracis</i>	2	—	—	—	2	—	—	—
<i>B. licheniformis</i>	6	3	—	—	9	—	—	—
<i>B. subtilis</i>	53	3	—	—	56	—	—	—
<i>B. pumilus</i>	7	3	11	2	22	1	—	—
<i>B. firmus</i>	1	—	—	—	1	—	—	—
<i>B. circulans</i>	1	—	—	—	1	—	—	—
<i>B. brevis</i>	2*	—	—	—	2	—	—	—
<i>B. sphaericus</i>	6†	—	—	—	6	—	—	—

\* One of two strains showed L-asparaginase activity at pH 5·6.

† One of six strains showed L-asparaginase activity at pH 5·6.

*B. licheniformis* and *B. megaterium*. Wade *et al.* (1971) also pointed out that asparaginase activity in bacterial cells could be assayed without disruption.

Thus 464 bacteria were cultivated and L-asparaginase and L-glutaminase activities were measured at pH 5.6 and 8.4. The results, summarized in Table 1, show the distribution of the pH 8.4 activities/ml of culture. The activities at pH 5.6 were generally less than at pH 8.4, but the pattern of distribution was almost identical. The distribution pattern obtained when using the specific activities was more or less the same as that shown in Table 1.

A large proportion of strains of *Pseudomonas* species showed L-asparaginase and L-glutaminase simultaneously. These observations confirm the wide distribution of L-asparaginase and L-glutaminase activities in this genus which has been reported previously by de Groot & Lichtenstein (1960), Greenberg *et al.* (1964), Ramadan *et al.* (1964), El-Asmar & Greenberg (1966), Peterson & Ciegler (1969*b*) and Arima *et al.* (1972). Two *Pseudomonas putrefaciens* strains, like organisms of the Enterobacteriaceae, showed L-asparaginase with little L-glutaminase activities. In this point they differ from other species of *Pseudomonas*. It was also found that they differ from other *Pseudomonas* species in possessing nucleoside *N*-ribosyl transfer activity which occurred rarely in Pseudomonadaceae but was common in Enterobacteriaceae (Imada & Igarasi, 1967). We suspect some relationship between these red *Pseudomonas putrefaciens* and the Enterobacteriaceae, especially *Serratia* species.

Two *Spirillum* strains showed L-asparaginase activity. Two strains of *Sphaerotilus natans* showed L-asparaginase activity and one of the two strains showed L-glutaminase activity. In the family Achromobacteraceae strains of *Alcaligenes* and *Achromobacter* species showed L-asparaginase or L-glutaminase activity, while *Flavobacterium* species showed little activity. Arima *et al.* (1972) indicated the extracellular production of L-asparaginase by *Alcaligenes* species, Wade *et al.* (1971) showed the L-asparaginase activities of *Alcaligenes* species, and Roberts *et al.* (1972) reported the properties of glutaminase-asparaginase of members of the Achromobacteraceae having antitumour activity.

Most members of the Enterobacteriaceae were active in L-asparaginase but not so active in L-glutaminase. *Escherichia coli* strains were extraordinarily active in L-asparaginase, and *Proteus* species also showed strong L-asparaginase activity.

As compared with the above-mentioned Gram-negative bacteria, Gram-positive bacteria were usually less active in the amidases. A few strains among *Micrococcus* and *Brevibacterium* species showed these activities, but none was shown among the *Staphylococcus*, *Corynebacterium* and *Arthrobacter* species.

Several strains of *Bacillus licheniformis*, *B. subtilis* and *B. pumilus*, which are included in the *B. subtilis* group, showed L-asparaginase activity, and for many strains of *B. pumilus* the level was high. *Bacillus megaterium*, *B. cereus* and other *Bacillus* species showed little activity.

Bacteria which showed comparatively high amidase activities were selected and cultivated for 16 and 40 h in 40 ml of CSL-medium in 200 ml Erlenmeyer flasks and the amidase activities in cells and culture filtrates were examined (see Table 2). The activities were always higher in cells cultivated in 200 ml Erlenmeyer flasks than in those cultivated in test tubes. Strains of *Escherichia coli*, *Proteus morganii*, *P. vulgaris*, *P. mirabilis*, *Pseudomonas fluorescens*, *P. schuylkilliensis*, *Alcaligenes faecalis* and *Bacillus pumilus* formed more than 1 i.u./ml of L-asparaginase within the bacteria. Strains of *Pseudomonas aureofaciens*, *P. schuylkilliensis* and *Alcaligenes faecalis* possessed a little L-asparaginase or L-glutaminase activity in their culture filtrates.

Table 2. L-Asparaginase and L-glutaminase activities of selected bacterial strains

Bacterial strain	16 h culture					40 h culture				
	Growth ( $E_{600}$ $\text{nm} \times 10^3$ )	L-Asparaginase* (i.u./ml)		L-Glutaminase* (i.u./ml)		$E_{600}$ $\text{nm} \times 10^3$	L-Asparaginase (i.u./ml)		L-Glutaminase (i.u./ml)	
		Bacteria	Culture filtrate	Bacteria	Culture filtrate		Bacteria	Culture filtrate	Bacteria	Culture filtrate
<i>Escherichia coli</i> F-221 (IFO Iijima)	340	5.23	0	—†	—	300	4.85	0	—	—
<i>Proteus vulgaris</i> IFO3167	330	3.31	0	—	—	340	2.00	0	—	—
<i>P. morgani</i> IFO3848	240	1.15	0	—	—	280	1.46	0	—	—
<i>P. mirabilis</i> IFO12255	310	2.23	0	—	—	320	1.39	0	—	—
<i>Serratia indica</i> IFO3759	280	0.62	0	—	—	270	0.62	0	—	—
<i>S. piscatorum</i> IFO12527	270	0.77	0	—	—	200	0.92	0	—	—
<i>Alcaligenes faecalis</i> IFO12624	145	0.85	trace	—	—	240	2.15	0.26	—	—
<i>Pseudomonas fluorescens</i> IFO3461	240	2.54	0	1.23	0	340	2.39	0	1.92	0
<i>P. aureofaciens</i> IFO3522	260	0.89	trace	0.35	0	290	0.85	0.25	0.69	0.26
<i>P. schuyllkilliensis</i> IFO12055	275	2.39	0	1.08	0	340	2.31	0.07	1.77	0.11
<i>Spirillum metamorphum</i> IFO12012	60	0	0	0	0	200	0.15	0	0.22	0
<i>Micrococcus cerificans</i> IFO12522	240	0.55	0	—	—	270	0.31	0	—	—
<i>Brevibacterium</i> sp. IFO12147	290	0.42	0	0.18	0	200	0.31	0	0.31	0
<i>Bacillus pumilus</i> IFO12093	205	1.54	0	—	—	225	1.62	0	—	—

\* Activities at pH 8.4.

† Not examined.



Table 3. L-Asparaginase and L-glutaminase activities of *Streptomyces* species

Strain	L-Asparaginase (i.u./ml)	L-Glutaminase (i.u./ml)
<i>Streptomyces californicus</i> IFO3386	0·20	0·18
<i>S. globiformis</i> IFO12208	0·18	0·17
<i>S. griseoflavus</i> IFO3428	0·09	0·15
<i>S. griseolus</i> IFO3403	0·09	0·12
<i>S. netropsis</i> IFO3723	0·40	0·44
<i>S. olivochromogenes</i> IFO3178	0·11	0·14
<i>S. rimosus</i> IFO3226	0·22	0·18
<i>S. roseochromogenes</i> IFO3363	0·11	0·09

Table 4. Fungal strains producing L-asparaginase and L-glutaminase in culture filtrates

Fungal strain	L-Asparaginase (i.u./ml)	L-Glutaminase (i.u./ml)
Ascomycetes		
<i>Anixiella reticulata</i> IFO5814	0·42*	0*
<i>Microascus desmosporus</i> IFO7021	0·56	Not examined
	0·28*	0·24*
<i>Dichotomyces albus</i> var. <i>spinosus</i> IFO8655	0·11	0
<i>Nectria haematococca</i> IFO6891	0·46	0
<i>N. elegans</i> IFO7187	0·20	0
<i>N. cinnabarina</i> IFO6821	1·05	0
<i>Hypomyces solani</i> IFO7707	0·66	0
<i>H. solani</i> var. <i>xanthoxyli</i> IFO7710	0·55	0
<i>H. haematococcus</i> IFO5980	0·28	0
Fungi Imperfecti		
<i>Fusarium solani</i> IFO5899	0·29	0
<i>F. solani</i> var. <i>rasinfetum</i> IFO4473	0·12	0
<i>F. oxysporum</i> f.2 IFO5264	0·40	0
<i>F. roseum</i> IFO5421	0·10	0
<i>Tilachlidium humicola</i> IFO5696	0·29	0·31
<i>Verticillium malthoasei</i> IFO6624	0·42	0·38
<i>Penicillium urticae</i> IFO4633	1·32	0·06
<i>P. claviforme</i> IFO4676	0·22	0
<i>P. expansum</i> IFO5453	0·11	0
	0·83*	0*
<i>P. aculeatum</i> IFO7840	0·29	0
<i>P. granulatum</i> IFO5737	0·44*	0*

\* The activities in culture filtrates of 5-day culture in DP-medium in shaking flasks. Others are the activities in culture filtrates of 5-day culture in M-medium in test tubes.

L-Glutaminase activity was also found in cells of *Pseudomonas*, *Spirillum* and *Brevibacterium* species. L-Glutaminase activity was stronger in 40 h-culture bacteria than in 16 h-culture bacteria, while L-asparaginase activity remained unchanged or was slightly weaker in 40 h-culture bacteria. Therefore, it might be suggested that there exist more than two enzymes which deamidate L-asparagine or L-glutamine. Nikolaev, Evseev, Tyul'panova & Abdumalikov (1969) pointed out the presence of isoenzymes in a pseudomonad. One of the isoenzymes deamidated L-asparagine selectively and another deamidated L-asparagine and L-glutamine.

We have observed that the amidases from various bacteria differ from each other with respect to their pH-activity relationship, heat-stability and substrate specificity.

Table 5. *Distribution of L-asparaginase and L-glutaminase activities among yeasts*

Genus and species	Examined	Number of strains			Remarks
		L-Asparaginase active (A/B)*	L-Asparaginase and L-glutami- nase active (A/B)*	L-Glutaminase active (A/B)*	
<i>Eremascus</i>	1	0	0	0	—
<i>Wickerhamia</i>	1	0	0	0	—
<i>Endomyces</i>	5	0	0	0	—
<i>Endomycopsis</i>	19	0	0	0	—
<i>Schizosaccharomyces</i>	29	1	0	0	—
<i>Schizosacch. octosporus</i>	—	(1/2)	—	—	—
<i>Saccharomyces</i>	239	5	0	0	—
<i>S. peka</i>	—	(2/2)	—	—	—
<i>S. mangini</i>	—	(1/1)	—	—	—
<i>S. cerevisiae</i>	—	(1/57)	—	—	—
<i>S. logos</i>	—	(1/3)	—	—	—
<i>Zygosaccharomyces</i>	80	0	0	0	—
<i>Chlamydozoma</i>	3	0	0	0	—
<i>Torulaspora</i>	17	0	0	0	—
<i>Schwanniomyces</i>	2	0	0	0	—
<i>Debaryomyces</i>	89	4	0	0	—
<i>D. kloeckeri</i>	—	(1/14)	—	—	V†
<i>D. hansenii</i>	—	(2/18)	—	—	V
<i>D. nicotianae</i>	—	(1/3)	—	—	V
<i>Hansenula</i>	101	29	2	0	—
<i>H. saturnus</i>	—	(3/8)	—	—	VI
<i>H. anomala</i>	—	(4/28)	—	—	VI
<i>H. suaveolens</i>	—	(2/2)	—	—	VI
<i>H. subpelliculosa</i>	—	(1/1)	—	—	VI
<i>H. schneegii</i>	—	(1/2)	—	—	VI
<i>H. wingei</i>	—	(5/11)	—	—	—
<i>H. canadensis</i>	—	(2/3)	—	—	—
<i>H. jadinii</i>	—	(1/2)	(1/2)	—	VI‡
<i>H. beijerinckii</i>	—	(4/5)	(1/5)	—	VI
<i>H. pettersonii</i>	—	(2/2)	—	—	—
<i>H. fabiani</i>	—	(2/2)	—	—	—
<i>H. miso</i>	—	(2/3)	—	—	—
<i>Pichia</i>	90	2	0	0	—
<i>P. polymorpha</i>	—	(1/4)	—	—	—
<i>P. etchellsii</i>	—	(1/1)	—	—	—
<i>Hanseniospora</i>	10	1	0	0	—
<i>Hanseniospora valbyensis</i>	—	(1/4)	—	—	I
<i>Saccharomycodes</i>	4	0	0	0	—
<i>Nadsonia</i>	3	0	0	0	—
<i>Naganishia</i>	1	0	1	0	CRYP†
<i>Naganishia globosus</i>	—	—	(1/1)	—	—
<i>Metschnikowia</i>	1	0	0	0	—
<i>Nematospora</i>	2	0	0	0	—
<i>Candida</i>	237	19	0	5	—
<i>C. utilis</i>	—	(9/9)	—	—	VI
<i>C. robusta</i>	—	(1/9)	—	—	II§
<i>C. pelliculosa</i>	—	(3/3)	—	—	VI
<i>C. scottii</i>	—	—	—	(5/10)	—
<i>C. humicola</i>	—	(1/3)	—	—	—
<i>C. melnii</i>	—	(1/4)	—	—	III
<i>C. curvata</i>	—	(1/2)	—	—	CRYP†
<i>C. fabiani</i>	—	(2/2)	—	—	—
<i>C. tropicalis</i>	—	(1/10)	—	—	I

Table 5 (continued)

Genus and species	Examined	Number of strains			Remarks
		L-Asparaginase active (A/B)*	L-Asparaginase and L-glutaminase active (A/B)*	L-Glutaminase active (A/B)*	
<i>Lipomyces</i>	4	0	0	0	—
<i>Ashbya</i>	2	0	0	0	—
<i>Petasospora</i>	3	0	0	0	—
<i>Eremothecium</i>	6	0	0	0	—
<i>Sporobolomyces</i>	29	15	0	0	—
<i>S. coralliformis</i>	—	(2/2)	—	—	—
<i>S. holsaticus</i>	—	(2/3)	—	—	—
<i>S. coprophilus</i>	—	(1/1)	—	—	—
<i>S. roseus</i>	—	(6/7)	—	—	—
<i>S. ruber</i>	—	(1/1)	—	—	—
<i>S. pararoseus</i>	—	(2/3)	—	—	RHO†
<i>S. carnicolor</i>	—	(1/1)	—	—	—
<i>Bullera</i>	4	0	2	0	—
<i>B. alba</i>	—	—	(2/4)	—	—
<i>Cryptococcus</i>	44	9	12	7	—
<i>C. laurentii</i>	—	(4/11)	(2/11)	(1/11)	CRYP
<i>C. nitens</i>	—	(1/1)	—	—	—
<i>C. albidus</i>	—	—	(8/16)	(6/16)	CRYP
<i>C. neoformans</i>	—	(1/10)	(2/10)	—	CRYP
<i>C. luteolus</i>	—	(2/2)	—	—	—
<i>C. diffluens</i>	—	(1/1)	—	—	CRYP
<i>Torulopsis</i>	93	6	0	0	—
<i>T. capsuligenes</i>	—	(1/1)	—	—	—
<i>T. aerea</i>	—	(2/2)	—	—	CRYP
<i>T. bacillaris</i>	—	(1/6)	—	—	SCH†
<i>T. colliculosa</i>	—	(1/5)	—	—	VII
<i>T. candida</i>	—	(1/3)	—	—	V
<i>Pityrosporum</i>	3	0	0	0	—
<i>Kluyveromyces</i>	1	0	0	0	—
<i>Pachysolen</i>	1	0	0	0	—
<i>Brettanomyces</i>	10	0	0	0	—
<i>Mycoderma (Candida)</i>	19	0	0	0	—
<i>Kloeckera</i>	28	0	0	0	—
<i>Trichosporon</i>	21	1	1	0	—
<i>T. cutaneum</i>	—	—	(1/9)	—	—
<i>T. pullulans</i>	—	(1/3)	—	—	—
<i>Rhodotorula</i>	108	30	8	0	—
<i>R. flava</i>	—	(2/5)	(1/5)	—	—
<i>R. glutinis</i>	—	(10/27)	—	—	—
<i>R. pilimanae</i>	—	(1/1)	—	—	—
<i>R. rubra</i>	—	(16/39)	(6/39)	—	RHO
<i>R. infirmoniata</i>	—	—	(1/1)	—	—
<i>R. macerans</i>	—	(1/1)	—	—	—
Others	16	1	0	0	—
'Aspergillus Hefe'¶	—	(1/1)	—	—	—

The yeast strains which showed more than 0.03 i.u./ml of the activity were regarded as active ones.

\* (A/B) shows active number (A)/tested number (B).

† Roman number, CRYP (*Cryptococcus*), RHO (*Rhodotorula*) and SCH (*Schizosaccharomyces*) indicate the serological group.

‡ Spore former of *Candida utilis*.

§ Non-spore former of *Saccharomyces cerevisiae*.

|| Heterobasidiomycetous stage is present.

¶ An *Aspergillus* sp. grows like a yeast.

*L-Asparaginase and L-glutaminase activities of Streptomyces and Nocardia*

L-Asparaginase activity was not detected in the culture filtrates of 261 strains of actinomycetes. Since these strains grow well on media which contain L-asparagine as a single nitrogen source, L-asparagine-metabolizing activity was presumed to be present. When 25 strains of *Streptomyces* species were cultivated in 40 ml of ST-2 medium in 200 ml Erlenmeyer flasks for 3 days with shaking, L-asparagine and L-glutamine deamidating activities were found in the sonicated preparations from such cultures. Some of the examples are shown in Table 3. The only other reference to the presence of L-asparaginase in *Streptomyces* is that by Campbell & Mashburn (1969), who showed its presence in *Streptomyces griseus*.

*L-Asparaginase and L-glutaminase activities of fungi*

Culture filtrates of 518 organisms of Basidiomycetes, 440 strains of Phycomycetes, 660 strains of Ascomycetes and 2540 strains of Fungi Imperfecti were examined. Amidase-active strains are listed in Table 4.

Among strains of Fungi Imperfecti, all strains of *Fusarium* species formed L-asparaginase. L-Asparaginase activity was also detected in culture filtrates of a number of *Penicillium* species. In the culture filtrates of strains of *Penicillium claviforme* and *P. expansum*, L-asparaginase occurred frequently. Arima *et al.* (1972) also reported the presence of L-asparaginase in the culture filtrate of *P. claviforme*.

Strains of *Tilachlidium humicola* and *Verticillium malthousei*, organisms of Fungi Imperfecti formed L-asparaginase and L-glutaminase.

Among strains of Ascomycetes, the genera *Hypomyces* and *Nectria*, which are the perfect stage of the genus *Fusarium*, formed L-asparaginase. Therefore the ability of extracellular formation of L-asparaginase is a common property in *Fusarium* and its related genera. Several other species of Ascomycetes also formed L-asparaginase or L-glutaminase.

Basidiomycetes and Phycomycetes showed little growth under our culture conditions and enzyme activities were not detected in their culture filtrates.

*L-Asparaginase and L-glutaminase activities of yeasts*

Among 1326 yeasts, L-asparaginase or L-glutaminase activity occurred in about 12% of them. The distribution of the activities was closely related to the classification, especially to the serological classification by Tsuchiya (1967). The results are summarized in Table 5.

L-Asparaginase activity occurred frequently in yeasts which are serologically grouped into the VI or *Hansenula* group, and included *Candida utilis*, *C. pelliculosa* and many strains of *Hansenula*, the *Cryptococcus* group and the *Rhodotorula* group. Several yeasts of the *Cryptococcus* and *Rhodotorula* groups possessed L-glutaminase together with L-asparaginase. *Sporobolomyces* species which are serologically related to *Rhodotorula* species frequently showed L-asparaginase activity.

That *Hansenula jadini* is regarded as the perfect stage of *Candida utilis* was reflected in the fact that all the strains of these two species possessed L-asparaginase.

Yeast strains forming L-glutaminase alone were found in *Candida scottii*, which has a heterobasidiomycetous stage named *Leucosporidium* (Fell, Statzell, Hunter & Phaff, 1969), in *Cryptococcus albidus* and *C. laurentii*.

Extracellular formation of L-asparaginase by *Candida utilis* and *Rhodotorula rosa* was reported by Arima *et al.* (1972). We have also observed L-asparaginase in culture filtrates of *Hansenula jadini*, *Rhodotorula rubra*, *Cryptococcus albidus*, *Sporobolomyces roseus* and some other strains as well as in the culture filtrate of *Candida utilis*.

*Saccharomyces cerevisiae* is known to possess L-asparaginase (Gorr & Wagner, 1932; Grassmann & Mayr, 1933; Greenberg *et al.* 1964), but under our experimental conditions the activity was detected in only one of 57 strains.

In spite of the presence of a number of investigations on microbial L-asparaginase and L-glutaminase, the correlation between their distribution and microbial classification has not been discussed. Having studied systematically the occurrence of the activities in a wide variety of micro-organisms we have observed that the amidase activities are concentrated in certain taxonomic groups of bacteria, fungi and yeasts. Such evidence may be valuable in understanding to what extent the current classification, which is mainly based on morphology and so some extent on physiology, is correlated with biochemical characteristics. It may also help in selecting microbial sources of enzymes.

We thank Drs S. Tatsuoka and R. Takeda, Central Research Division, Takeda Chemical Industries Ltd, for their continued interest and encouragement throughout the course of our work. We also thank Drs T. Hasegawa, K. Tubaki and I. Banno, Institute for Fermentation, Osaka, for supplying micro-organisms and for rewarding discussions. The technical assistance of Mr H. Ono, Miss H. Sotoma and Miss Y. Nakamura is also appreciated.

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