Enzymatic Hydrolysis of Theanine and the Related Compounds

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Some strains of Pseudomonas were found capable of utilizing L-theanine and D-theanine as a sole nitrogen and carbon source. L-Theanine and D-theanine were hydrolyzed by the enzyme from Pseudomonas aeruginosa to yield stoichiometrically L-glutamate and D-glutamate, respectively, and ethylamine, which were isolated from the reaction mixture and identified. The theanine hydrolase was purified approximately 200-fold. It was shown that the activities of L-theanine hydrolase, D-theanine hydrolase and the heat-stable L-glutamine hydrolase and D-glutamine hydrolase are ascribed to a single enzyme, which may be regarded as a γ-glutamyltransferase from the point of view of the substrate specificity and the properties. This theanine hydrolase catalyzed the transfer of γ-glutamyl moiety of the substrates and glutathione. L-Glutamine and D-glutamine were hydrolyzed by theanine hydrolase and also the heat-labile enzyme whose properties resembled the common glutaminase.

INTRODUCTION

L-Theanine was isolated from tea leaves as an important component of Japanese green tea by Sakatol) in 1950 and the structure was confirmed as N-ethyl-γ-L-glutamine, HOOCCH(NH2)CH2CH2CONHC2H5. The compound was synthesized from L-glutamic acid and ethylamine. Since the discovery of theanine, reports have appeared concerning the isolation and the identification of many derivatives of glutamine and asparagine in which the substituents other than those represented by the common α-amino acids occur at the amide nitrogen atom3-12). Information on metabolism of these amide compounds has never been obtained.

Glutaminase (L-glutamine amidohydrolase) and asparaginase (L-asparagine amidohydrolase) which catalyze the hydrolysis of amide groups of L-glutamine and L-asparagine, respectively, are widely distributed in animals, plants and micro-organism. Studies on these enzymes have been reviewed by several workers13-15). General properties of bacterial glutaminase13-18) and asparaginase19-22) were studied well and discussed. It has been believed so far that the substrate specificity of these enzymes is high from the point of view of the structure and also of the optical activity.

In the present paper, the enzymatic hydrolysis of L-theanine, D-theanine and the related amide compounds, the identification of the products and the properties of the enzyme will be described. The studies on the transfer reaction of the γ-
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glutamyl group of the substrates to hydroxylamine to form the hydroxamates will be also presented. The enzyme which catalyzes the hydrolysis of theanine will be named “theamine hydrolase” tentatively.

EXPERIMENTAL

Materials
L-Theanine, d-theanine and d-glutamine were kindly supplied by Dr. Y. Sakato of Shizuoka University and the late Dr. Y. Tsuchiya of Ajinomoto Co. Inc.. Hydroxylapatite was prepared according to the method of Tiselius et al.23). The amino acids, DEAE-cellulose (Serva) and the other chemicals were obtained commercially.

Microbiological Methods
The bacteria were, unless otherwise stated, cultivated on the L-glutamate medium composed of 10.0 g monosodium L-glutamate, 1.0 g KH₂PO₄, 2.0 g K₂HPO₄, 0.1 g MgSO₄•7H₂O and 0.05 g FeSO₄•7H₂O per liter of tap water. The pH was adjusted to 7.0 by the addition of sodium hydroxide.

The cultures were grown at 30°C in 500 ml flasks or 2 1 flasks containing 100 ml or 500 ml of medium respectively, under shaking for 18 hours. The cells were harvested by centrifugation and washed twice with 0.85 per cent sodium chloride solution.

Preparation of Partially Purified Enzyme
The washed cells were suspended in 0.05 M phosphate buffer, pH 7.4, containing 0.01 per cent 2-mercaptoethanol, and subjected to a 19 Kc Kaijo Denki sonic oscillator for 5 minutes. The intact cells and debris were centrifuged off at 10,000 × g for 30 minutes. The supernatant was dialyzed against 0.001 M phosphate buffer, pH 7.4, containing 0.01 per cent 2-mercaptoethanol for about 15 hours. The dialyzed solution was used as a cell-free extract.

The partially purified enzyme was prepared as follows. To the cell free extract, was added 2 ml of 1 per cent protamine sulfate solution per 100 mg of protein under stirring. The mixture was centrifuged and the bulky inactive precipitate was discarded. The supernatant was adjusted to pH 7.4 with 10 per cent ammonium hydroxide, and brought to 50 per cent saturation by the addition of solid ammonium sulfate. After standing for 30 minutes, the precipitate was removed by centrifugation. To the supernatant, was added solid ammonium sulfate to bring 80 per cent saturation and the precipitate was collected by centrifugation. The precipitate was dissolved in a small volume of distilled water and dialyzed against 0.002 M phosphate buffer, pH 7.4, containing 0.01 per cent 2-mercaptoethanol for 15 hours. The inactive precipitate formed during dialysis was removed by centrifugation. All operations were carried out at 0—5°C. The dialyzed supernatant was employed as a partially purified enzyme, which was purified about ten-fold from the cell-free extract.
Analytical Methods

Amino acids were determined according to the methods described previously as follows. Amino acids were separated by circular paper chromatography, in which n-butanol: acetic acid: water (4:1:1) system was employed. The paper was dried after development, and sprayed with 0.5 per cent ninhydrin solution in 75 per cent ethanol. After heating at 50°C for 30 minutes, the colored zone of the paper was cut off and extracted with 75 per cent ethanol containing 0.005 per cent copper sulfate. The intensity of the color of the extract was measured at 500 m\(\mu\).

Hydroxamic acids were determined according to the method of Lipmann et al. as follows. One ml of the freshly neutralized 2 M hydroxylamine solution and 3 ml of ferric reagent were added to 2 ml of the reaction mixture after incubation. After protein was filtered off from the mixture, the color intensity developed was measured at 540 m\(\mu\). Authentic succinylhydroxamic acid and aspartylhydroxamic acid were employed as the standard for glutamylhydroxamic acid and aspartylhydroxamic acid formed enzymatically, respectively.

L-Glutamic acid was determined manometrically with L-glutamic acid decarboxylase of the acetone dried Escherichia coli cells.

Ethylamine was determined according to the modified Conway's micro-diffusion method.

Protein was determined by the method of Lowry et al.

Measurement of Enzyme Activity

The activity of hydrolase was assayed by measuring the amount of the amino acid formed from the substrate. The standard reaction mixture contained 40 \(\mu\) moles of the substrate, 100 \(\mu\) moles of phosphate buffer, pH 8.0, and the enzyme in a final volume of 2.0 ml. After incubation was carried out at 37°C for 30 minutes, 0.2 ml of 50 per cent trichloroacetic acid was added to the mixture. A 10 \(\mu\)l aliquot of the deproteinized supernatant was used for amino acid analysis. Specific activity was expressed as micromoles of the amino acid formed from the substrate per mg of protein per hour.

The activity of transferase was determined by measurement of the amount of the hydroxamic acid formed. The standard reaction mixture consisted of 40 \(\mu\) moles of the substrate, 400 \(\mu\) moles of freshly neutralized hydroxylamine, 60 \(\mu\) moles of phosphate buffer, pH 7.0, and the enzyme in a final volume of 2.0 ml. After incubation at 37°C for 30 minutes, hydroxamic acid produced was determined as described above. Specific activity was expressed as micromoles of hydroxamic acid formed per mg of protein per hour.

RESULTS

Utilization of Theanine and Hydrolysis of Theanine by Bacteria

Growth of several strains of bacteria on a medium containing L-theanine as a sole carbon and nitrogen source was tested. As shown in Table 1, it was found that
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Table 1. Growth of bacteria on L-theanine medium.
The medium contained L-theanine instead of sodium L-glutamate in the basal medium. Growth of the organisms was expressed as turbidity at 470 mμ of the medium incubated.

<table>
<thead>
<tr>
<th>Strain</th>
<th>Incubation period</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>12 h.</td>
</tr>
<tr>
<td><em>Pseudomonas fluorescens</em></td>
<td>0.120</td>
</tr>
<tr>
<td><em>Pseudomonas fragi</em></td>
<td>0.000</td>
</tr>
<tr>
<td><em>Pseudomonas riboflavinus</em></td>
<td>0.000</td>
</tr>
<tr>
<td><em>Pseudomonas strifaciens</em></td>
<td>0.000</td>
</tr>
<tr>
<td><em>Pseudomonas aeruginosa</em></td>
<td>0.827</td>
</tr>
<tr>
<td><em>Pseudomonas sp. T-1</em></td>
<td>0.232</td>
</tr>
<tr>
<td><em>Escherichia coli</em></td>
<td>0.064</td>
</tr>
<tr>
<td><em>Aerobacter aerogenes</em></td>
<td>0.000</td>
</tr>
</tbody>
</table>

*Pseudomonas aeruginosa*, *Pseudomonas fluorescens* and *Pseudomonas sp. T-1* which had been isolated from the air, utilized L-theanine satisfactorily, while the other strains of *Pseudomonas*, *Escherichia coli* and *Aerobacter aerogenes* could not at all or little grow on the L-theanine medium. *Pseudomonas aeruginosa*, *Pseudomonas fluorescens* and *Pseudomonas sp. T-1* were shown to be able to utilize D-theanine as well as the L-enantiomorph.

L-Theanine hydrolase activity and D-theanine hydrolase activity of the cell-free extracts of the bacteria grown on the media containing peptone as a nitrogen source and sodium L-glutamate as a sole nitrogen and carbon source respectively, were assayed. As demonstrated in Table 2, the highest activity was observed in the cell-free extracts from *Pseudomonas aeruginosa* and *Pseudomonas polycolor* grown on

Table 2. Theanine hydrolase activity of the cell-free extract of various strains.
The cell-free extract was employed and assay of the α enzyme activity was carried out under the standard condition. Peptone medium was composed of 1.0 per cent peptone, 1.0 per cent glycerol, 0.4 per cent K₂HPO₄, 0.2 per cent KH₂PO₄, 0.2 per cent yeast extract and 0.02 per cent MgSO₄·7H₂O. Basal medium: See the text. Specific activity is shown in the table.

<table>
<thead>
<tr>
<th>Strain</th>
<th>Peptone medium</th>
<th>Basal medium</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>L-The*</td>
<td>D-The*</td>
</tr>
<tr>
<td><em>Ps. fluorescens</em></td>
<td>1.76</td>
<td>1.83</td>
</tr>
<tr>
<td><em>Ps. fragi</em></td>
<td>1.63</td>
<td>1.53</td>
</tr>
<tr>
<td><em>Ps. riboflavinus</em></td>
<td>0.36</td>
<td>0.47</td>
</tr>
<tr>
<td><em>Ps. polycolor</em></td>
<td>1.09</td>
<td>1.29</td>
</tr>
<tr>
<td><em>Ps. aeruginosa</em></td>
<td>1.16</td>
<td>1.20</td>
</tr>
<tr>
<td><em>Ps. strifaciens</em></td>
<td>0.24</td>
<td>0.47</td>
</tr>
</tbody>
</table>

* The abbreviation of theanine
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the L-glutamate medium and also the cell-free extracts of the strains which failed to grow on the L-theanine medium showed some extent of activity. *Pseudomonas aeruginosa* was employed in the following experiments.

In the extracts of these organisms, glutamic acid racemase, theanine racemase and D-glutamic acid—pyruvic acid transaminase activities could not be detected. This fact suggests that D-theanine may be hydrolyzed directly to D-glutamic acid, which may be converted to L-glutamic acid or other L-amino acids via certain unknown process.

**Isolation and Identification of Reaction Products**

A large amount of the reaction mixture whose content was described in the methods, was incubated at 37°C for 3 hours. Theanine in the mixture was hydrolyzed completely. The mixture was deproteinized by the addition of 50 per cent trichloroacetic acid and excess trichloroacetic acid was removed by continuous extracting with ether for 10 hours. The solution was passed through Amberlite IR-120 (H⁺ form) column (2.3x25 cm). The column was eluted with 2 M ammonium hydroxide at a flow rate of 30 ml per hour. The effluent containing glutamic acid was concentrated *in vacuo*, and glutamic acid was crystallized from aqueous ethanol. Recrystallization was repeated twice.

<table>
<thead>
<tr>
<th>Analysis</th>
<th>C%</th>
<th>H%</th>
<th>N%</th>
</tr>
</thead>
<tbody>
<tr>
<td>Calcd. for C₅H₈O₄N</td>
<td>40.81</td>
<td>6.16</td>
<td>9.52</td>
</tr>
<tr>
<td>Found</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Product from L-theanine</td>
<td>40.66</td>
<td>6.25</td>
<td>9.73</td>
</tr>
<tr>
<td>Product from D-theanine</td>
<td>40.41</td>
<td>6.32</td>
<td>9.35</td>
</tr>
</tbody>
</table>

Both of these preparations produced from L-theanine and D-theanine were also identified as glutamic acid by a comparison of their Rf values with those of authentic glutamic acid in a several solvent systems and the admixture of the preparations with an authentic sample showed one spot.

Optical activity was determined by manometric method with L-glutamic acid decarboxylase (*Escherichia coli* acetone powder) and by Rudolph photoelectric spectropolarimeter Model 200 S.

<table>
<thead>
<tr>
<th></th>
<th>CO₂ Output µmoles</th>
<th>([\alpha]_20^*)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Calcd.</td>
<td>5.0</td>
<td></td>
</tr>
<tr>
<td>Found</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Authentic L-glutamic acid</td>
<td>4.8</td>
<td>+32.4</td>
</tr>
<tr>
<td>Product from L-theanine</td>
<td>4.8</td>
<td>+32.5</td>
</tr>
<tr>
<td>Authentic D-glutamic acid</td>
<td>0.0</td>
<td>−32.8</td>
</tr>
<tr>
<td>Product from D-theanine</td>
<td>0.08</td>
<td>−32.7</td>
</tr>
</tbody>
</table>

*(1 per cent solution in 1 N HCl)*

These results demonstrate that L-glutamic acid and D-glutamic acid were produced enzymatically from L-theanine and D-theanine respectively.
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**Isolation and Identification of Ethylamine**

Ethylamine was isolated as the ethylamine picrolonate. A reaction mixture in which phosphate buffer, pH 6.4 was employed instead of the same buffer, pH 8.0 in the standard mixture, was incubated at 37°C for 2 hours. The enzyme preparation used in this experiment was dialyzed thoroughly to be free of ammonium ion. After incubation, protein was removed by the addition of 50 per cent trichloroacetic acid. Ammonium was not detected in the deproteinized supernatant by Nessler’s reaction. The supernatant solution which was made alkaline with 40 per cent sodium hydroxide solution, was steam-distilled in a Kjeldahl apparatus for 15 minutes. The tip of the condenser was dipped into 6 ml of ethanol containing picrolonic acid equivalent to ethylamine to be formed. The picrolonate was crystallized. Recrystallization was repeated twice from aqueous ethanol. Authentic ethylamine picrolonate was prepared from pure ethylamine in the same way as described above.

<table>
<thead>
<tr>
<th>Analysis</th>
<th>C %</th>
<th>H %</th>
<th>N %</th>
</tr>
</thead>
<tbody>
<tr>
<td>Calcd. for $C_{12}H_{19}O_5N_5$</td>
<td>46.60</td>
<td>4.89</td>
<td>22.64</td>
</tr>
<tr>
<td>Found</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Authentic ethylamine picrolonate</td>
<td>46.61</td>
<td>4.98</td>
<td>22.58</td>
</tr>
<tr>
<td>Product from L-theanine</td>
<td>46.39</td>
<td>4.95</td>
<td>22.76</td>
</tr>
<tr>
<td>Product from D-theanine</td>
<td>46.56</td>
<td>4.92</td>
<td>23.07</td>
</tr>
</tbody>
</table>

Both preparations of the picrolonate produced from L-theanine and D-theanine showed the same pattern in infrared spectrum as that of the authentic ethylamine picrolonate (Figure 1).

**Fig. 1.** Infrared spectra of ethylamine picrolonate.

I: Authentic ethylamine picrolonate, II: Picrolonate of the product from L-theanine, III: Picrolonate of the product from D-theanine.
Stoichiometry of Hydrolysis Reaction

The formation of glutamic acid and ethylamine from theanine proceeded stoichiometrically as indicated in Figure 2.

![Graph showing the formation of glutamic acid and ethylamine from theanine](image)

**Fig. 2.** Theanine hydrolysis reaction.

The reaction mixture consisted of 40 μmoles of L-theanine in (A) or D-theanine in (B), 100 μmoles of phosphate buffer, pH 6.4 and the enzyme (500 μg) in a final volume of 2.0 ml. Incubation was carried out at 37°C.

- •-•-•-•-•-•: Theanine, -○-○-○-○: D-glutamic acid in (A) and D-glutamic acid in (B), - - - - - - - - : Ethylamine.

Purification of Enzyme

The enzyme was purified from the extract of *Pseudomonas aeruginosa* as described below. All operations, unless otherwise stated, were carried out at 0–5°C.

**Step 1. Extraction.** The washed cells were suspended in 0.05 M phosphate buffer, pH 7.4, containing 0.01 per cent 2-mercaptoethanol and sonicated in a 19 Kc Kaijo Denki sonic oscillator for 10 minutes. The intact cells and debris were removed by centrifugation at 10,000 x g for 40 minutes.

**Step 2. Treatment with protamine.** To the supernatant obtained was added 1 ml of 1.5 per cent protamine sulfate solution per 100 mg of enzyme protein under stirring. The bulky inactive precipitate was centrifuged off. The increase in total activity during this procedure suggests that some inhibitory factor may be removed by protamine treatment.

**Step 3. Heating treatment.** After 1 M acetate buffer, pH 5.3, was added to the supernatant solution to bring the final concentration to 0.04 M, the solution was heated in a water bath of 70°C until the temperature of the solution reached 55°C ± 1°C. The enzyme solution was kept at this temperature for 5 minutes. The enzyme solution became quite milky during this process. The solution was cooled rapidly to 4°C and centrifuged at 10,000 x g for 15 minutes.

(486)
Step 4. Ammonium sulfate fractionation. The clear supernatant solution obtained was adjusted to pH 7.4 by the addition of 10 per cent ammonium hydroxide solution and brought to 70 per cent saturation with ammonium sulfate. After standing for 30 minutes, the precipitate formed was removed by centrifugation at 10,000 × g for 15 minutes. The supernatant was adjusted to 90 per cent saturation with solid ammonium sulfate. After allowing the mixture to equilibrate for 30 minutes, the precipitate was collected by centrifugation at 10,000 × g for 20 minutes, dissolved in a small volume of 0.01 M phosphate buffer pH 7.0, and dialyzed against the same buffer containing 0.01 per cent 2-mercaptoethanol for 15 hours. The inactive precipitate formed during dialysis was centrifuged off.

Step 5. DEAE-Cellulose chromatography. The dialyzed enzyme solution was placed on a column of DEAE-cellulose (2 × 18 cm) equilibrated with 0.01 M phosphate buffer, pH 7.0. Elution was carried out stepwise with 0.01 M phosphate buffer, pH 7.0 containing various concentrations of sodium chloride. The flow rate was 20 ml per hour and 10 ml aliquots of eluate were collected. The elution of the protein was followed by measuring the absorbancy at 280 nm. Figure 3 shows the typical elution pattern of the enzyme. The major protein peaks are designated by letters A to D in the figure. L-Theanine hydrolase activity was found only in fraction C, while L-glutamine hydrolase activity was separated clearly into fractions.
A and C. The finding is in agreement with the result of the heat stability of the partially purified enzyme to be described later, providing the evidence that glutaminase activity in the crude preparation is due to the two different enzymes. In the subsequent studies on the enzyme purification and the properties of theanine hydrolase and glutamine hydrolase, the fraction C was employed unless otherwise noted, although glutamine hydrolase in fraction A was also investigated as described later.

The fractions containing theanine hydrolase activity were pooled and concentrated by the dialysis against solid polyethyleneglycol.

**Step 6. Hydroxylapatite chromatography.** The concentrated enzyme solution was subjected to hydroxylapatite chromatography. Hydroxylapatite was packed into a column (2 x 4 cm) and equilibrated with 0.003 M phosphate buffer, pH 7.0. The enzyme solution was introduced into the column and eluted stepwise with increasing concentration of phosphate buffer, pH 7.0. The buffer was allowed to flow at a rate of 2 ml per hour and eluted fractions of 5 ml were collected. The elution pattern of the enzyme is shown in Figure 4. Theanine hydrolase and glutamine hydrolase activity were found in the same pattern and the protein peak was in parallel with the enzyme activities. It is suggested that the enzyme protein may be chromatographically homogenous.

In Table 3, a summary of a purification of the enzyme was demonstrated. The over-all purification from the extract to DEAE-cellulose chromatography fractions was approximately 200-fold. Since the hydroxylapatite chromatography showed...
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Table 3. Summary of purification of enzyme.

<table>
<thead>
<tr>
<th>Step</th>
<th>Total protein mg</th>
<th>Total activity</th>
<th>Specific activity</th>
<th>Yield %</th>
</tr>
</thead>
<tbody>
<tr>
<td>1. Extraction</td>
<td>3520</td>
<td>39,150</td>
<td>11.1</td>
<td>(100)</td>
</tr>
<tr>
<td>2. Protamine treatment</td>
<td>3240</td>
<td>45,080</td>
<td>13.9</td>
<td>115</td>
</tr>
<tr>
<td>3. Heat treatment</td>
<td>1315</td>
<td>41,900</td>
<td>30.5</td>
<td>107</td>
</tr>
<tr>
<td>4. Ammonium sulfate fractionation</td>
<td>89.3</td>
<td>16,800</td>
<td>188.7</td>
<td>43.1</td>
</tr>
<tr>
<td>5. DEAE-Cellulose chromatography</td>
<td>6.1</td>
<td>13,360</td>
<td>2210</td>
<td>34.1</td>
</tr>
</tbody>
</table>

Enzyme activity was expressed as L-theanine hydrolase activity.

no effect to increase the specific activity of the enzyme, the result of this procedure was not described in the table.

Table 4 shows a summary of comparison of four enzyme activities, i.e. L-

Table 4. Comparison of enzyme activity.

<table>
<thead>
<tr>
<th>Step</th>
<th>L-GHA</th>
<th>D-GHA</th>
<th>L-GHA</th>
<th>D-GHA</th>
<th>L-GHA</th>
<th>D-GHA</th>
</tr>
</thead>
<tbody>
<tr>
<td>1. Extraction</td>
<td>7.56</td>
<td>6.19</td>
<td>0.76</td>
<td>0.77</td>
<td></td>
<td></td>
</tr>
<tr>
<td>2. Protamine treatment</td>
<td>6.89</td>
<td>6.21</td>
<td>0.82</td>
<td>0.74</td>
<td></td>
<td></td>
</tr>
<tr>
<td>3. Heat treatment</td>
<td>6.54</td>
<td>6.10</td>
<td>0.75</td>
<td>0.72</td>
<td></td>
<td></td>
</tr>
<tr>
<td>4. Ammonium sulfate fractionation</td>
<td>1.85</td>
<td>2.26</td>
<td>0.74</td>
<td>0.91</td>
<td></td>
<td></td>
</tr>
<tr>
<td>5. DEAE-Cellulose chromatography</td>
<td>1.67</td>
<td>2.10</td>
<td>0.63</td>
<td>0.74</td>
<td></td>
<td></td>
</tr>
<tr>
<td>6. Hydroxylapatite chromatography</td>
<td>1.62</td>
<td>2.05</td>
<td>0.74</td>
<td>0.83</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

The abbreviation THA and GHA are for theanine hydrolase activity and glutamine hydrolase activity, respectively.

theanine hydrolase, D-theanine hydrolase, L-glutamine hydrolase and D-glutamine hydrolase, during the purification procedures. The ratio of the hydrolase activities of L-theanine and L-glutamine to those of the D-enantiomorphs remained approximately constant and the ratio of L-glutamine hydrolase activity to L-theanine hydrolase activity or of D-glutamine hydrolase activity to D-theanine hydrolase activity decreased remarkably after ammonium sulfate fractionation and thereafter remained almost constant. These facts suggest that L-theanine and D-theanine may be hydrolyzed by a single enzyme or very similar enzymes and that hydrolysis of L-glutamine and D-glutamine may be catalyzed by at least two different enzymes, which were able to be separated by ammonium sulfate fractionation, and also DEAE-cellulose chromatography as illustrated in Figure 3.

Properties of Enzyme. The linear relationship between the amount of the enzyme protein and the amount of L-glutamate or D-glutamate produced from L-theanine or D-theanine, respectively, was obtained under the conditions employed. The formation of glutamate from theanine by theanine hydrolase proceeded as a function of time within 60 minutes under the standard condition.
Fig. 5. Effect of pH on the hydrolysis of L-theanine (I) and D-theanine (II).

The buffers used were acetate for pH 4.6, phosphate for pH 5.25 to 8.0, veronal for pH 8.5 to 9.6 and borate for pH 10.4.

Fig. 6. Effect of pH on the hydrolysis of L-glutamine (I) and D-glutamine (II).

The buffers used were acetate for pH 4.6, phosphate for pH 5.25 to 8.0, veronal for pH 8.5 to 9.6 and borate for pH 10.4.
Effect of pH on Theanine Hydrolase Activity and Glutamine Hydrolase Activity

The effect of pH on the enzymatic hydrolysis of L-theanine, D-theanine, L-glutamine and D-glutamine by the purified enzyme was demonstrated in Figures 5 and 6. Both of these pH-activity curves exhibited a similar tendency and the optimal pH values for these four hydrolase reactions were all in the range of 8.5 to 9.0.

Effect of Heating on the Enzyme Activities

The partially purified enzyme preparation was heated at 55°C for a given period and after being cooled rapidly, the enzyme activities hydrolyzing L-theanine, D-theanine and L-glutamine were determined. As indicated in Figure 7, L-theanine hydrolase and D-theanine hydrolase activities were stable against heating the enzyme under the condition employed here. L-Glutamine hydrolase activity, however, was reduced rapidly to a half of the original activity by heating for 10 minutes and thereafter the activity was no longer influenced by heating. This finding suggests that L-glutamine may be hydrolyzed by two different kinds of enzymes, both of which were contained together in the partially purified enzyme preparation and one of them is heat-stable and another is heat-labile. The result obtained in this experiment is consistent with the separation of the enzyme activity of hydrolyzing L-glutamine into two different fractions by DEAE-cellulose chromatography demonstrated in Figure 3.
Figure 8 shows the effect of the period of heat treatment on the activities of the highly purified enzyme preparation. As indicated in Figure 8, these four kinds of the enzyme activities were all little affected by heating the enzyme at 55°C for 60 minutes. Although the partially purified enzyme preparation may contain two different enzymes catalyzing the hydrolysis of L-glutamine as shown in Figure 7, the highly purified enzyme preparation has probably one enzyme hydrolyzing L-glutamine whose behaviour was similar to those of D-glutamine hydrolase, L-theanine hydrolase or D-theanine hydrolase of the purified preparation during heat treatment. The results obtained here and in Figure 7 indicate that fraction A in Figure 3 contained a heat-labile glutamine hydrolase and that in fraction C containing also L-theanine and D-theanine hydrolyzing enzyme activity, another glutamine hydrolase which is resistant to heating, was found.

**Effect of Substrate Concentration on Enzyme Activity**

The enzyme activities catalyzing hydrolysis of L-theanine and D-theanine with various concentrations of the substrate were investigated. The partially purified enzyme preparation was used in this experiment. The apparent Michaelis constants for L-theanine and D-theanine were calculated by the method of Lineweaver and Burk to be approximately $5.8 \times 10^{-3}$ M and $9.2 \times 10^{-3}$ M, respectively. The apparent Michaelis constants for L-glutamine and D-glutamine were shown to be $4.7 \times 10^{-3}$ M and $6.4 \times 10^{-3}$ M, respectively. It is concluded that affinity of the enzyme to these four substrates is almost the same.
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Formation of γ-Glutamylhydroxamic acid by Transfer Reaction

In the course of these studies, it was observed that incubation of the highly purified enzyme preparation with theanine or glutamine in the presence of hydroxylamine led to significant formation of γ-glutamylhydroxamic acid. Neither metal ion nor nucleotide was required in this reaction. Figure 9 demonstrates the relation-

![Figure 9](image)

**Fig. 9.** Time course of hydrolysis and transfer reaction of L-theanine (A) and L-glutamine (B).

Reaction mixtures contained 50 μ moles of L-theanine or L-glutamine, 400 μ moles of hydroxylamine, 5 μ g of purified enzyme and 60 μ moles of phosphate buffer, pH 7.4, in a final volume 2.0 ml. Incubation for indicated time at 37°C. I: L-theanine (A) or L-glutamine (B); II: Glutamate; III: γ-Glutamylhydroxamic acid.

ship among the formation of glutamate by the hydrolase action, the production of γ-glutamylhydroxamic acid by transferase action, and the disappearance of L-theanine or L-glutamine at various incubation periods. D-Theanine and D-glutamine could be the substrate of the transfer reaction as well as their L-isomers as mentioned later. It is realized that this enzyme catalyzes the hydrolysis reaction of both isomers of theanine and glutamine in the absence of hydroxylamine, and does also the transfer reaction of γ-glutamyl moiety of both isomers of theanine and glutamine to hydroxylamine to form γ-glutamylhydroxamate besides the hydrolysis reaction, in the presence of hydroxylamine. In the latter case, the rate of the formation of glutamic acid by hydrolysis reaction was as high as approximately three times that of the hydroxamic acid by the transfer reaction, and the total amount of these two products coming from both reactions was virtually the same as that of the substrate consumed.

Effect of Inhibitors on Hydrolase Activity and Transferase Activity

The effect of several inhibitors on the hydrolase activity and the γ-glutamyltransferase activity was investigated. Urea and p-chloromercuribenzoic acid...
had no effect on both activities at a concentration of $10^{-3}$ M. Sodium fluoride ($5 \times 10^{-2}$ M) inhibited L-theanine transferase activity in the range of 10 to 20 per cent, but not L-theanine hydrolase activity at all. At a concentration of $10^{-2}$ M, ethylenediamine tetraacetic acid, on the contrary, caused a 30 per cent inhibition for the hydrolase activity, but no change for the transferase activity when L-theanine was employed as a substrate. Mercuric chloride ($10^{-3}$ M), iodoacetic acid ($2 \times 10^{-3}$ M) and copper sulfate ($10^{-2}$ M) inhibited both enzyme activities in the range of 50 to 80 per cent. The inhibition of the activities by iodoacetic acid was protected partially (about 30 per cent) by preincubation of the enzyme with 2-mercaptoethanol at a concentration of $2 \times 10^{-2}$ M for 15 minutes.

**Substrate Specificity**

Table 5 indicates the substrate specificity of the two separate enzyme preparations.

Table 5. Substrate specificity of two enzyme preparations.

<table>
<thead>
<tr>
<th>Substrate</th>
<th>Hydroxamic acid formation</th>
<th>Hydrolysis of amide</th>
<th>Hydroxamic acid formation</th>
<th>Hydrolysis of amide</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>$\mu$ moles</td>
<td>$\mu$ moles</td>
<td>$\mu$ moles</td>
<td>$\mu$ moles</td>
</tr>
<tr>
<td>L-Theanine</td>
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<td>0</td>
<td>1.24</td>
<td>5.27</td>
</tr>
<tr>
<td>D-Theanine</td>
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<td>0</td>
<td>0.31</td>
<td>4.10</td>
</tr>
<tr>
<td>L-Glutamine</td>
<td>2.62</td>
<td>7.20</td>
<td>1.92</td>
<td>6.60</td>
</tr>
<tr>
<td>D-Glutamine</td>
<td>2.22</td>
<td>3.85</td>
<td>1.01</td>
<td>6.09</td>
</tr>
<tr>
<td>L-Asparagine</td>
<td>0.86</td>
<td>3.32</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>D-Asparagine</td>
<td>3.31</td>
<td>4.15</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Glutathione</td>
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<td>–</td>
<td>1.16</td>
<td>–</td>
</tr>
<tr>
<td>Acetamide</td>
<td>0</td>
<td>–</td>
<td>0</td>
<td>–</td>
</tr>
</tbody>
</table>

Enzyme activity was assayed under the standard condition. Experiment I: The enzyme of fraction A on DEAE-cellulose chromatography was used. Experiment II: The enzyme of fraction C on DEAE-cellulose chromatography was used.

rations, i.e. fraction A and fraction C, obtained by DEAE-cellulose chromatography as demonstrated in Figure 3. The hydrolysis of both isomers of asparagine and the formation of $\beta$-aspartylhydroxamic acid were catalyzed by fraction A which was not active for either isomer of theanine but for both isomers of glutamine. D-Asparagine was hydrolyzed and also transferred more rapidly by fraction A than its L-enantiomorph. Fraction C which was not active for L-asparagine and D-asparagine, however, catalyzed the hydrolysis of both isomers of theanine, and the formation of $\gamma$-glutamylhydroxamic acid from them. L-Glutamine and D-glutamine could be hydrolyzed and their $\gamma$-glutamyl moiety was able to be transferred to hydroxylamine by either fraction A or C. The formation of the $\gamma$-glutamylhydroxamic acid from glutathione was catalyzed by fraction C, but not by fraction A at all. Acetamide was inert for both preparations.
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**DISCUSSION**

It is well known that glutaminase and asparaginase catalyzed hydrolysis of the amide group of L-glutamine and L-asparagine, respectively. Groot et al.\textsuperscript{19,20}, described that neither asparaginase from 	extit{Pseudomonas fluorescens} nor that from mammalian liver hydrolyzed both N-ethyl-\(\beta\)-L-asparagine and N-methyl-\(\beta\)-L-asparagine.

It has been also reported that N-ethyl-\(\gamma\)-glutamine (L-theanine) and N-methyl-\(\gamma\)-glutamine are completely inert as a substrate in the enzymatic amide hydrolysis and transfer reactions\textsuperscript{29-32}).

The present studies show that some strains of 	extit{Pseudomonas} can utilize L-theanine and D-theanine as a sole nitrogen and carbon source, and that the cell-free extract catalyzes the hydrolysis of amide group of L- and D-theanine. It is of interest that D-theanine can be utilized by bacteria and hydrolyzed enzymatically to yield D-glutamic acid and ethylamine without conversion into L-theanine via a racemization system. As the enzyme catalyzes the stoichiometric formation of glutamic acid and ethylamine from theanine, it can be named theanine ethylamidohydrolase or theanine hydrolase in this respect.

The highly purified enzyme preparation from 	extit{Pseudomonas aeruginosa} which catalyzed the hydrolysis of L-theanine and D-theanine, and the formation of \(\gamma\)-glutamylhydroxamic acid from these compounds by \(\gamma\)-glutamyl transfer to hydroxylamine, was obtained. Failure to separate these enzyme activities by various procedures suggest that these four reactions may be catalyzed by a single enzyme. Besides L-theanine and D-theanine, L-glutamine and D-glutamine could be used as a substrate for the enzymatic hydrolysis and transfer reactions, but L-asparagine, D-asparagine and acetamide could not. Although the fact that both enantiomorphs of glutamine were potenter substrates than L-theanine and D-theanine appears to show that the enzyme should be considered as a glutaminase which has a wide substrate specificity, the properties of the enzyme in question differ from those of the enzymes of other bacterial and animal sources which have investigated so far. The optimal pH of this enzyme for glutamine hydrolysis was between 8.5 to 9.0, and those of the glutaminases from 	extit{Escherichia coli}\textsuperscript{30}, 	extit{Clostridium welchii}\textsuperscript{37} and some 	extit{Pseudomonas}\textsuperscript{38} were reported to be around 5.0 to 6.6. In this respect, the enzyme is likely to be close to the glutaminases from animal tissue\textsuperscript{31,38-36}, which optimal pH was found in the alkaline, but the present 	extit{Pseudomonas aeruginosa} enzyme differs from the animal glutaminases in many properties. The glutaminases from animal tissues were activated by phosphate ions and were inhibited completely by p-chloromercuribenzoic acid at \(10^{-3}\) M, while this enzyme could not be activated by phosphate ions and was not influenced by p-chloromercuribenzoic acid at the concentration of \(10^{-3}\) M. It is another remarkable difference of this enzyme of 	extit{Pseudomonas aeruginosa} from the usual glutaminases from other sources that the former is able to hydrolyze L-theanine and D-theanine, and to form the \(\gamma\)-glutamylhydroxamate from them and glutathione, and that the latter is inert for other \(\gamma\)-glu-
tamyl derivatives than glutamine.

An enzyme isolated from *Agaricus bisporus* which catalyzes the irreversible hydrolysis of \(\gamma\)-glutamylhydrazide bond and the transfer of \(\gamma\)-glutamyl moiety of agaritine to hydroxylamine, was reported to hydrolyse L-theanine and L-glutamine, and to transfer their \(\gamma\)-glutamyl residue, but the common \(\gamma\)-glutamyl peptides e.g. glutathione, was inactive in these enzyme reactions.

The purified \(\gamma\)-glutamyltranspeptidase from hog kidney which transfers \(\gamma\)-glutamyl residue of \(\gamma\)-glutamyl-\(\beta\)-nitroanilide, hydrolyzed L-glutamine, D-glutamine and L-glutamic acid \(\gamma\)-ester, and catalyzed the formation of \(\gamma\)-glutamylhydroxamate from both isomers of glutamine, glutathione, \(\gamma\)-ethylglutamate and \(\gamma\)-glutamylglycine besides \(\gamma\)-L-glutamyl-\(\beta\)-nitroanilide. L-Asparagine and L-homoglutamine were not used as a substrate in the hydrolysis and transfer of the hog kidney enzyme and it is essential to involve \(\gamma\)-glutamyl residue in the substrate. Williams *et al.* demonstrated that transamidase preparation from *Bacillus subtilis* catalyzed the transfer of \(\gamma\)-glutamyl moiety of glutamine, glutathione and other \(\gamma\)-glutamylpeptide to hydroxylamine.

The behavior of the enzyme in question of *Pseudomonas aeruginosa* for the inhibitors and the substrate specificity serve to differentiate it from an amidase of *Pseudomonas aeruginosa* which is specific for the hydrolysis of several aliphatic amides and the transfer of the acyl moiety of such amides to hydroxylamine. The amidase and transferase activities of the aliphatic amides were reported to be inhibited by urea or potassium fluoride, while our enzyme of the same organisms was not influenced by such compounds and could not hydrolyze acetamide.

Thus, the enzyme in question from *Pseudomonas aeruginosa* seems to be more similar to \(\gamma\)-glutamyltransferases than glutaminases previously reported and is regarded as a \(\gamma\)-glutamyltransferase which has a wide substrate specificity in respect to the kind of the substituted moiety at the amide nitrogen atom of \(\gamma\)-glutamyl compounds and to the stereostructure. The hydrolytic activity of this enzyme can be considered its ability to transfer \(\gamma\)-glutamyl residue to water instead of hydroxylamine.

The fact that the glutamine amidohydrolase activity was separated completely to two different fractions by DEAE-cellulose column chromatography shows the occurrence of two kinds of enzyme catalyzing hydrolysis of glutamine in the cell-free extract of this organisms. One of them is the enzyme to catalyze the hydrolysis of theanine and the transfer reaction of theanine and glutathione as well as glutamine as discussed above. Another enzyme, whose fraction was designated as A in Figure 3 and activity was very unstable, catalyzes the hydrolysis of both enantiomorphs of asparagine and glutamine, and the transfer of \(\beta\)-aspartyl or \(\gamma\)-glutamyl residue of them to hydroxylamine, but none of L-theanine, D-theanine and glutathione was active in such enzymatic hydrolysis and transfer.

Recently, Ramadan *et al.* reported that a highly purified enzyme preparation
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from a strain of *Pseudomonas* which activity was inherently unstable, has both glutaminase and asparaginase activities and d-asparagine has about 20 per cent of the activity of the l-isomer. It was also demonstrated by Ehrenfeld *et al.*, that the enzyme isolated from *Azotobacter agilis* catalyzed hydrolysis of both isomers of glutamine and asparagine, and succinamic acid as well as the formation of hydroxamate from these amides, and that l-theanine, l-glutamic acid γ-ethyl ester, acetamide and other several compounds were not active in both enzymatic reactions.

Therefore, the enzyme found in Fraction A appears to bear the closest resemblance to the common enzyme known as glutaminase or asparaginase and widely distributed. It is of interest that both isomers of glutamine are hydrolyzed and transferred by different two enzymes in the same organisms and both kinds of enzymes have no stereospecificity for their substrates.

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