

# Characteristics of Glutamate Dehydrogenase in Mitochondria Prepared from Corn Shoots<sup>1</sup>

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TOMOYUKI YAMAYA, ANN OAKS\*, AND HIDEAKI MATSUMOTO  
*Institute for Agricultural and Biological Sciences, Okayama University, Chuo 2-20-1, Kurashiki, Okayama 710, Japan (T.Y., H.M.); and Department of Biology, McMaster University, Hamilton, Ontario, L8S 4K1 Canada (A.O.)*

## ABSTRACT

The amination of  $\alpha$ -ketoglutarate ( $\alpha$ -KG) by NADH-glutamate dehydrogenase (GDH) obtained from Sephadex G-75 treated crude extracts from shoots of 5-day-old seedlings was stimulated by the addition of  $\text{Ca}^{2+}$ . The NADH-GDH purified 161-fold with ammonium sulfate, DEAE-Toyopearl, and Sephadex G-200 was also activated by  $\text{Ca}^{2+}$  in the presence of 160 micromolar NADH. However, with 10 micromolar NADH,  $\text{Ca}^{2+}$  had no effect on the NADH-GDH activity. The deamination reaction (NAD-GDH) was not influenced by the addition of  $\text{Ca}^{2+}$ .

About 25% of the NADH-GDH activity was solubilized from purified mitochondria after a simple osmotic shock treatment, whereas the remaining 75% of the activity was associated with the mitochondrial membrane fraction. When the lysed mitochondria, mitochondrial matrix, or mitochondrial membrane fraction was used as the source of NADH-GDH,  $\text{Ca}^{2+}$  had little effect on its activity. The mitochondrial fraction contained about 155 nanomoles Ca per milligram of mitochondrial protein, suggesting that the NADH-GDH in the mitochondria is already in an activated form with regard  $\text{Ca}^{2+}$ . In a simulated *in vitro* system using concentrations of 6.4 millimolar NAD, 0.21 millimolar NADH, 5 millimolar  $\alpha$ -KG, and 5 millimolar glutamate thought to occur in the mitochondria, together with 1 millimolar  $\text{Ca}^{2+}$ , 10 and 50 millimolar  $\text{NH}_4^+$ , and purified enzyme, the equilibrium of GDH was in the direction of glutamate formation.

NADH-GDH<sup>2</sup> is easily solubilized from mitochondria from roots (30), shoots (4, 25), and seeds (28) of various plants. Although it is the only mitochondrial enzyme with the potential for the assimilation of  $\text{NH}_4^+$ , it has been suggested by Miflin and Lea (21) that the  $\text{NH}_4^+$  released by the conversion of glycine to serine in the mitochondria (3) is reassimilated via GS in the cytosol. They have also suggested that the principal function of mitochondrial GDH is the oxidation of glutamate (21). However, a part of intramitochondrially generated  $\text{NH}_4^+$  from glycine is incorporated into glutamate by mitochondria isolated from pea shoots (11). Chou and Splittstoesser (7), Joy (14), and recently Furuhashi and Takahashi (10) have shown that NADH-GDH

(amination reaction), but not NAD-GDH (deamination reaction), is activated by  $\text{Ca}^{2+}$  *in vitro*. According to the work by Bowman *et al.* (4), NAD is the only detectable pyridine nucleotide in mung bean mitochondria. Thus, concentrations of  $\text{Ca}^{2+}$  and NADH in the mitochondria could be important in the regulation of NADH-GDH.

The purpose of the present study was to characterize the activation of NADH-GDH by  $\text{Ca}^{2+}$  and to estimate concentrations of  $\text{Ca}^{2+}$  and NADH for simulation of the GDH reaction in mitochondria isolated from corn shoots.

## MATERIALS AND METHODS

**Plant Materials.** Corn seeds (*Zea mays* L. var W64A  $\times$  W182E) were planted in vermiculite and seedlings were grown in a Toshiba Cold Chain growth chamber, model TGS-13L, with a 12-h photoperiod for 5 d. Light (about 490  $\mu\text{E}/\text{m}^2 \cdot \text{s}$ ) was supplied by Toshiba Yoko lamps DA400 (metal halide lamp) and fluorescent lamps, and the temperature was maintained at 27°C. The plants were watered every morning with a nutrient medium containing 1 mM  $\text{Ca}^{2+}$  as described previously (18).

**Isolation of Mitochondria by Density Gradient Centrifugation.** All reagents and buffer solutions were prepared with double glass-distilled  $\text{H}_2\text{O}$  to minimize  $\text{Ca}^{2+}$  contamination, and all operations were carried out at 4°C. Whole shoots of 5-d seedlings were harvested, washed in water, blotted dry with filter paper, and weighed. The shoots (40 to 60 g in fresh weight) were chopped with razor blades in extraction buffer consisting of 0.4 M mannitol, 0.1 M Hepes-KOH buffer (pH 7.5), 1 mM EDTA, 0.1% (w/v) BSA, and 0.6% (w/v) insoluble PVP, at a ratio of 1 g fresh weight per 3 ml of buffer. The chopped materials were ground with mortar and pestle.

The mitochondrial fraction was isolated by modification of the method described by Nishimura *et al.* (26) as described previously by us (32). The purified mitochondria after Percoll discontinuous density gradient were able to oxidize both malate and succinate with a high P/O and RCR ratios and appear to have about 92% of the outer membrane intact (32). Our preparations were almost free of Chl (chloroplast) or catalase (microbody) contamination. These are criteria used by Nishimura *et al.* for assessing mitochondrial purity (26).

**Preparation of Submitochondrial Fractions.** The purified mitochondrial pellet was treated hypotonically using 1.5 ml of 20 mM Hepes-KOH buffer (pH 7.5). The suspension was mixed with a Teflon-glass homogenizer and incubated in an ice bath for 30 min. The suspension (1 ml) was then centrifuged in the ultracentrifuge equipped with an SW 50-1 rotor at 150,000 g for 1 h. The pellet, resuspended in 1 ml of the same buffer, was designated the mitochondrial membrane fraction. The supernatant remaining above the pellet consisted of the soluble matrix

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<sup>2</sup> Abbreviations: GDH, glutamate dehydrogenase; GOGAT, glutamate synthase; GS, glutamine synthetase;  $\alpha$ -KG,  $\alpha$ -ketoglutarate; 2-ME, 2-mercaptoethanol; MTT, 3-(4,5-dimethylthiazolyl-2)-2,5-diphenyltetrazolium bromide; PES, phenazine ethosulfate.

of the mitochondria.

**Purification of NADH-GDH from the Crude Mitochondrial Fraction.** The crude mitochondrial pellet prepared from 179 g fresh weight of the corn shoots was suspended in about 90 ml of 0.1 M Tris-HCl buffer (pH 8.5) containing 0.1 mM EGTA and 1 mM 2-ME. The suspension was frozen and thawed twice, placed in a salt-ice mixture, and then sonicated with a Branson sonifier, model 200, operating at power setting 5 on a 50% duty cycle for 2 min. Following the disruption treatment, the suspension was centrifuged at 20,000g for 30 min and the supernatant fraction was designated the crude mitochondrial extract. The extract was subjected to 30 to 60%  $(\text{NH}_4)_2\text{SO}_4$  fractionation. The resulting pellet was dissolved in 0.1 M Tris-HCl buffer (pH 8.5) containing 0.1 mM EGTA and 1 mM 2-ME and then dialyzed against the same buffer. The dialyzed preparation was centrifuged and the supernatant fraction was loaded onto a column (1.6 × 10 cm) of DEAE-Toyopearl, equilibrated with 50 mM Tris-HCl buffer (pH 8.5) containing 5 mM  $\alpha$ -KG, 1 mM 2-ME, and 0.1 mM EGTA. The column was washed with the same buffer and the NADH-GDH was eluted with a linear gradient of KCl (0 to 0.5 M). The NADH-GDH activity was recovered in the portion of the effluent containing approximately 0.12 M KCl. Proteins were then sedimented with 70%  $(\text{NH}_4)_2\text{SO}_4$  and the enzyme was further purified by gel filtration chromatography on Sephadex G-200 column (1.6 × 90 cm) equilibrated with 50 mM Tris-HCl buffer (pH 8.5) containing 0.1 M KCl and 1 mM 2-ME. The NADH-GDH activity was observed as a single peak near the void volume of the column, and was free of contaminating EGTA.

When NADH-GDH activity was assayed without purification, the procedures of enzyme extraction from the corn shoots and desalting with Sephadex G-75 were the same as those described by Oaks *et al.* (27).

**Enzyme Assays and Analytical Methods.** Enzyme activities were assayed spectrophotometrically in a double-beam spectrophotometer (Shimadzu, model UV-150-02) at 30°C. NADH-GDH was assayed routinely in 0.2 M Tris-HCl buffer (pH 8.0), containing 0.266 M  $(\text{NH}_4)_2\text{SO}_4$ , 0.16 mM NADH, and 5 mM  $\alpha$ -KG with or without 1 mM  $\text{CaCl}_2$  as described previously (27). However, in critical experiments  $(\text{NH}_4)_2\text{SO}_4$  was used in a concentration range of 2 to 100 mM. NAD-GDH was measured by a modification of the method described by Furuhashi and Takahashi (10). The assay system contained 0.1 M Tris-HCl buffer (pH 8.5), 3 mM NAD, and 50 mM glutamate with or without 1 mM  $\text{CaCl}_2$ . Succinate:Cyt *c* reductase activity was assayed by following reduction of Cyt *c* as described by Douce *et al.* (9). Malate dehydrogenase was assayed by following the oxidation of NADH as described by Priestley and Bruinsma (28). Fumarase was assayed by coupling the reaction with malic enzyme and following the reduction of NADP as described by Hatch (12).

The Ca content in the isolated mitochondrial fraction was determined with an atomic absorption/fluorescence emission spectrophotometer (Shimadzu model AA646, equipped with a Graphite furnace atomizer, model GFA-3). Protein content was determined by the method of Lowry *et al.* (16).

**Extraction and Determination of Pyridine Nucleotides.** Pyridine nucleotide content was determined by the cycling assay method described by Matsumura and Miyachi (19). During extraction and isolation of mitochondria, 10  $\mu\text{M}$  rotenone and 5  $\mu\text{M}$  antimycin A were included in all the buffer solutions and in the Percoll gradient solutions. The final mitochondrial pellet after the Percoll density gradient was dissolved in 1.1 ml of 0.1 M Bicine-NaOH buffer (pH 8.0). The mitochondrial suspension (0.5 ml) was then transferred to preheated NaOH or HCl solutions which had a final concentration of 0.1 N. After an incubation at 100°C for 2 min, the suspension was cooled and centrifuged. NADH is specifically extracted with an alkaline treatment, while NAD is extracted with an acid treatment (19). Each extract

was neutralized by adding an appropriate amount of NaOH or HCl. The extract (50 to 200  $\mu\text{l}$ ) was then incubated in brown test tubes in a reaction system containing 0.1 M Bicine-NaOH buffer (pH 8.0), 4 mM EDTA, 0.42 mM MTT, 1.676 mM PES, and 0.5 M ethanol for 5 min at 37°C. The total reaction volume was 1.0 ml. The reaction was started by adding 10 units of alcohol dehydrogenase and was allowed to run for 30 min at 37°C. The reaction was terminated by adding 12 mM iodoacetate and the absorbance was measured at 570 nm. Amounts of NAD(H) were calculated from the standard curves with authentic NAD(H). A blank test without sample or NAD(H) was also carried out.

**Source of Chemicals.** Bicine, Hepes, Mes, Tris, sucrose, mannitol, PES, Na-glutamate, and rotenone were all purchased from Nakarai Chemicals, Ltd., Japan and were of a specially prepared reagent grade. Fatty acid-free BSA, horse heart Cyt *c* (type III), yeast alcohol dehydrogenase free of NAD(H), chicken liver malic enzyme, and MTT were from Sigma; antimycin A, from Boehringer; Percoll and Sephadex, from Pharmacia. NAD, NADH, and NADP were purchased from Kojin Co. Ltd, Japan and DEAE-Toyopearl was from Toyo Soda MFG Co. Ltd, Japan. All other chemicals were of the highest quality available.

## RESULTS

**Localization of NADH-GDH in Mitochondria Isolated from Corn Shoots.** NADH-GDH is relatively easily solubilized from plant mitochondria and hence the enzyme is thought to be localized in the soluble matrix of mitochondria (4, 28). However, the method of disrupting the mitochondria in those reports (4, 28) involved a relatively vigorous treatment, such as freezing and thawing followed by sonication. To demonstrate the distribution of NADH-GDH in corn mitochondria, we treated mitochondria, purified by centrifugation through a Percoll density gradient, with osmotic shock, a relatively gentle disruption method. The mitochondrial matrix and membrane fractions were then separated by centrifugation (Table I). Most of the fumarase activity (a matrix marker) and succinate:Cyt *c* reductase (a membrane marker) were located in the expected fractions (29). NADH-GDH and malate dehydrogenase, which were thought to be in the mitochondrial matrix (4, 28) were, however, found primarily in the membrane fraction (75%). When the mitochondria were disrupted by harsher treatments (freezing and thawing followed by sonication), about 85% of the NADH-GDH activity was

Table I. Localization of NADH-GDH in Mitochondria Isolated from Corn Shoots

The purified mitochondrial pellet was shocked hypotonically using 1.5 ml of 20 mM Hepes-KOH buffer (pH 7.5), suspended with a Teflon-glass homogenizer, and incubated in an ice bath for 30 min. The lysed mitochondrial suspension (1 ml) was centrifuged at 150,000 g for 1 h. The resulting pellet was resuspended in 1 ml of the same buffer (mitochondrial membrane fraction). The supernatant fraction remaining above the pellet was designated the mitochondrial matrix fraction.

| Fraction                     | NADH-GDH  | Malate Dehydrogenase | Fumarase | Succinate:Cyt <i>c</i> Reductase |
|------------------------------|---|----------------------|----------|----------------------------------|
|                              | <i>μmol substrate utilized · min<sup>-1</sup> · mg<sup>-1</sup> protein</i> |                      |          |                                  |
| Lysed mitochondrial fraction | 0.476   | 2.03                 | 0.024    | 0.116                            |
|                              | <i>% of activity assayed with the lysed mitochondria</i>                    |                      |          |                                  |
| Mitochondrial matrix         | 25.5  | 22.6                 | 70.4     | 1.3                              |
| Mitochondrial membrane       | 77.8  | 74.8                 | 7.4      | 89.2                             |

Table II. Activation of NADH-GDH by  $Ca^{2+}$ 

The amination reaction of GDH was measured using a crude preparation desalted on Sephadex G-75 column (27), a 161-fold purified fraction, a lysed mitochondrial fraction, a mitochondrial matrix fraction, and a mitochondrial membrane fraction from shoots of 5-d-old corn. The NADH-GDH activity was measured in the presence or absence of 1 mM  $CaCl_2$  in the standard assay system containing 0.2 M Tris-HCl buffer (pH 8.0), 0.266 M  $(NH_4)_2SO_4$ , 0.16 mM NADH, and 5 mM  $\alpha$ -KG.

| Treatment   | Desalted<br>Crude<br>Fraction | Source of NADH-GDH |                         |        |          |
|---|-------------------------------|--------------------|-------------------------|--------|----------|
|   |                               | Purified<br>enzyme | Mitochondrial fractions |        |          |
|   |                               |                    | Lysed                   | Matrix | Membrane |
| <i><math>\mu</math>mol NADH oxidized <math>\cdot</math> min<math>^{-1}</math> <math>\cdot</math> mg<math>^{-1}</math> protein</i> |                               |                    |                         |        |          |
| No addition   | 0.01                          | 1.40               | 0.68                    | 0.20   | 0.65     |
| $Ca^{2+}$ (1<br>mM)   | 0.05                          | 10.70              | 0.73                    | 0.21   | 0.73     |

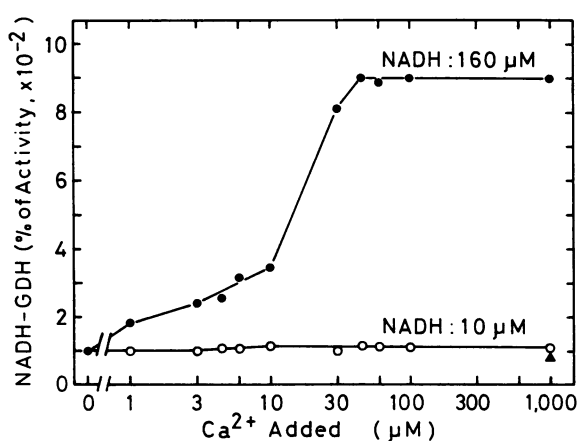


FIG. 1. Effect of  $Ca^{2+}$  on the amination reaction of GDH purified from corn shoots. The assay mixture contained 10 (O) or 160 (●)  $\mu$ M NADH, 5 mM  $\alpha$ -KG, 100 mM  $(NH_4)_2SO_4$ , and 1.2  $\mu$ g of the 161-fold purified enzyme in 0.2 M Tris-HCl buffer (pH 8.0). The NADH-GDH activity was 3.38 and 1.69 nmol NADH oxidized  $\cdot$  min $^{-1}$  with 10 and 160  $\mu$ M NADH, respectively, in the absence of  $Ca^{2+}$ . Effect of 2 mM EGTA was also tested in the presence of 160  $\mu$ M NADH and 1 mM  $Ca^{2+}$  (▲).

solubilized, whereas most of the succinate:Cyt *c* reductase was still associated in the membrane fraction (80 to 88%). These results indicated that both NADH-GDH and malate dehydrogenase are loosely associated with the mitochondrial membrane.

**Activation by  $Ca^{2+}$  of Reductive Amination Reaction of GDH from Corn Shoots.** NADH-GDH was prepared in five different ways from 5-d-old corn shoots: (a) a crude extract desalted by passage of Sephadex G-75, (b) a highly purified enzyme fraction, (c) a lysed mitochondrial fraction, (d) a mitochondrial matrix fraction, and finally (e) a mitochondrial membrane fraction. Activities in the presence of 1 mM  $Ca^{2+}$  in crude mitochondrial preparations and purified enzyme were 0.10 and 16.1  $\mu$ mol NADH oxidized/min  $\cdot$  mg protein. Thus, a 161-fold purification was achieved, 77% of the original activity was recovered, and all of the contaminating NADH oxidizing activity was removed (data not shown).

The effect of addition of 1 mM  $Ca^{2+}$  on the NADH-GDH activity was tested with each of these preparations (Table II). The NADH-GDH in the Sephadex G-75 treated fraction or the highly purified preparation was activated 5.1- and 9.1-fold, respectively, by the addition of  $Ca^{2+}$ . With the GDH obtained from the lysed mitochondria, the mitochondrial matrix, or the membrane fraction,  $Ca^{2+}$  additions had only a minor effect.

The effect of  $Ca^{2+}$  on the NADH-GDH activity was examined further with the highly purified preparation. The response of

$Ca^{2+}$  on the amination reaction was tested using fixed concentrations of 200 mM  $NH_4^+$ , 5 mM  $\alpha$ -KG, and 10 or 160  $\mu$ M NADH. In the presence of 160  $\mu$ M NADH, the activation curve was sigmoidal with significant activation seen with 1  $\mu$ M  $Ca^{2+}$  and maximal activation with 45  $\mu$ M  $Ca^{2+}$  (Fig. 1). The NADH-GDH was thus activated about 9-fold by  $Ca^{2+}$ . Higher concentrations of  $Ca^{2+}$ , up to 10 mM, maintained the activity at the saturated level (results not shown). Addition of 2 mM EGTA completely reversed the activation of the enzyme by 1 mM  $Ca^{2+}$ . When the NADH levels were reduced to 10  $\mu$ M NADH,  $Ca^{2+}$  had no effect on the amination reaction. The oxidative deamination reaction of GDH (NAD-GDH) was not affected by addition of  $Ca^{2+}$ . For example, with purified GDH, 2.58 and 2.41  $\mu$ mol NAD were reduced/min  $\cdot$  mg protein in the absence and presence of 1 mM  $Ca^{2+}$ , respectively.

In animal cells many of the regulatory processes linked to  $Ca^{2+}$  are mediated through their interactions with high-affinity  $Ca^{2+}$ -binding proteins, such as calmodulin (6). Calmodulin also exists in plant cells (17) but its function is not yet known, except perhaps in chloroplasts (13, 23) and the plasma membrane-enriched microsomal fraction (8). Neither bovine brain calmodulin nor the calmodulin antagonist, trifluoperazine, influenced the activity of purified or mitochondrial derived GDH (data not shown).

Saturation curves for the three substrates of NADH-GDH showed the normal Michaelis-Menten kinetics in the presence of 1 mM  $Ca^{2+}$  (Fig. 2). Apparent  $K_m$  values of 63  $\mu$ M for NADH and 2.22 mM for  $\alpha$ -KG were calculated from the curves. The apparent  $K_m$  for  $NH_4^+$  decreased when the concentration of NADH was lowered: it was 12.5 mM in the presence of 160  $\mu$ M NADH and 4.76 mM with 10  $\mu$ M NADH. Thus, the corn shoot-NADH-GDH has kinetic properties similar to the enzyme from *Medicago sativa* (24). With low concentrations of NADH, the apparent  $V_{max}$  for  $NH_4^+$  also decreased. In the absence of  $Ca^{2+}$ , a marked substrate inhibition was observed with the individual substrates and, hence, the  $K_m$  value for each could not be measured.

**Estimation of Concentrations of  $Ca^{2+}$ , Pyridine Nucleotides,  $NH_4^+$ , and Glutamate in Mitochondria Isolated from Corn Shoots.** The concentrations of  $Ca^{2+}$  in purified mitochondria

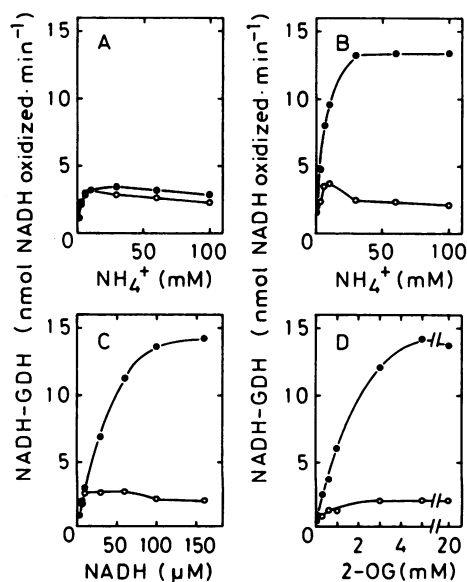


FIG. 2. Substrate saturation curves of the purified NADH-GDH from corn shoots in the presence (●) or absence (○) of 1 mM  $Ca^{2+}$ . A,  $NH_4^+$  with 10  $\mu$ M NADH; B,  $NH_4^+$  with 160  $\mu$ M NADH; C, NADH; and D,  $\alpha$ -KG. In each system the other components were added at the concentrations used in the assay system as described in Figure 1.



either in the standard assay system or in the simulated assay system as shown in Figure 3. The conversion of glycine to serine is catalyzed by two enzymes, glycine decarboxylase and serine hydroxymethyltransferase, both of which are localized in the mitochondrial membrane (22). This conversion is also coupled to the formation of NADH (1). Our finding that the loose association of NADH-GDH with the mitochondrial membrane in corn shoots coupled with the generation of both NADH and  $\text{NH}_4^+$  in that fraction suggest that GDH could participate in the reassimilation of  $\text{NH}_4^+$ .

It has been proposed by Mifflin and Lea (21) that GS should serve in preference to GDH in the assimilation of  $\text{NH}_4^+$  because of its low  $K_m$  for  $\text{NH}_4^+$ . In their scheme, GOGAT allows for the formation of two glutamate molecules from one molecule each of glutamine and  $\alpha$ -ketoglutarate. In normal metabolism, one glutamate molecule would be recycled through glutamine and the other would be available for other biosynthetic reactions. In photorespiration, the two glutamate molecules would be required for the synthesis of two glycine molecules, and a third glutamate molecule from somewhere would be required for the regeneration of glutamine. Our results indicate that levels of  $\text{NH}_4^+$  in the mitochondria are in the 5 to 10 mM range, and that the mitochondria can tolerate these relatively high concentrations of  $\text{NH}_4^+$ . At these levels of  $\text{NH}_4^+$ , there should be a synthesis of glutamate in the mitochondria mediated by GDH. It is our contention that GDH, by virtue of its isolation in mitochondria, can compete successfully with GS for that  $\text{NH}_4^+$ . A GDH fully saturated with  $\text{Ca}^{2+}$  and NADH may not be sufficient to assimilate all the  $\text{NH}_4^+$  released by the conversion of glycine to serine or, for that matter, produced by the action of nitrate reductase. However, its action would permit the net synthesis of glutamate, glutamine and, subsequently, of other amino acids required to sustain both the GS/GOGAT cycle itself and the enhanced rate of protein synthesis seen in light-treated leaves (5).

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