

A Supernodulation and Nitrate-Tolerant Symbiotic (*nts*) Soybean Mutant¹

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ABSTRACT

The nodulation characteristics of soybean (*Glycine max*) mutant *nts382* are described. The mutant nodulated significantly more than the parent cultivar Bragg in the presence and absence of several combined nitrogen sources (KNO₃, urea, NH₄Cl, and NH₄NO₃). The number of nodules on the tap root and on lateral roots was increased in the mutant line. In the presence of KNO₃ and urea, nitrogenase activity was considerably higher in *nts382* than in Bragg. Mutant plants were generally smaller than wild-type plants. Although *nts382* is a supernodulator, inoculation with *Rhizobium japonicum* was necessary to induce nodule formation and both trial strains CB1809 (= USDA136) and USDA110 elicited the mutant phenotype. Segregation of M₃ progeny derived from a M₂ wild-type plant indicated that the mutant character is inherited as a Mendelian recessive. The mutant is discussed in the context of regulation of nodulation and of hypotheses that have been proposed to explain nitrate inhibition of nodulation.

Nodule formation in legumes is tightly regulated. Indeed, symbiotic development is subject to both external factors and internal (or 'autoregulation') control mechanisms. Several environmental conditions, such as light intensity (photosynthate- or non-photosynthate mediated), temperature, pH, and soil moisture, influence nodulation (19). However, under optimum conditions for plant growth, exogenous nitrate represents a major environmental factor controlling the extent of symbiosis (5, 13). Small amounts of nitrate have been demonstrated to stimulate nodulation. Above these minute concentrations, however, nodule fresh weight is inversely related to the level of nitrate in the growth medium (20). To a lesser extent, other sources of combined nitrogen have also been shown to inhibit nodulation (7, 8, 10, 24, 27). The extent of this inhibition does vary considerably with the legume species, the form of combined nitrogen administered, and the experimental system. Some sources of combined nitrogen, for example urea or NH₄⁺, cause acidification of the growth medium (14), and the inhibitory effects of these nitrogen sources may be mediated indirectly through a reduction in pH rather than through the nitrogen status of the plant. To circumvent this complication, Vigue *et al.* (30) controlled pH fluctuations by the inclusion of a pH-buffering carboxy resin in the pots (17). In this system using soybeans, nitrate but not urea suppressed nodule fresh weight per plant at the range of concentrations tested. This was partially explained by reduced uptake of

nitrogen in urea-fed plants. In contrast to the effect of nitrate and urea on nodule mass, rates of acetylene reduction per unit of nodule mass were similar for nitrate and urea treatments (30).

In the absence of externally supplied combined nitrogen, nodulation is tightly regulated with the number of infections greatly exceeding the final number of mature nodules (2). Interruption of invasion is related to the effectiveness of the host-*Rhizobium* association (21) as well as to other internal factors not directly related to the nitrogen status of the plant (23). Generally, ineffective strains of *Rhizobium* form more nodules, especially after the initial stages of nodule formation (21). Nutman also showed that excision of effective (but not ineffective) red clover nodules resulted in a transient increase in the number of nodules subsequently formed (22). In fact, removal of the nodule meristem of effective nodules was sufficient to stimulate subsequent nodule development. The effect of nodule meristem excision on nitrogenase activity was not considered, but the argument that an inhibitory factor emanating from the growing point, and not the bacterial tissue, of the nodule was supported by the finding that excision of the main root tip had the same effect as nodule meristem removal. Clear evidence for internal regulation or autoregulation independent of nitrogen fixation and in the early stages of nodule initiation was reported by Pierce and Bauer (23). Using the spot inoculation technique (26), they showed that inoculation of soybean roots with *R. japonicum* several hours prior to a second inoculation substantially reduced nodulation by the second inoculation. In a split-root system for soybeans, Kossiak and Bohlool (16) showed that autoregulation prior to nodule appearance and nitrogenase activity was not restricted to root tissue immediately adjacent to the inoculated area, since prior inoculation of one side of a split-root suppressed nodulation on the other side (16). Studies by Calvert *et al.* (4) characterized this rapid regulatory phenomenon (23) further and showed that suppression of nodulation due to prior inoculation is mediated through suppression of nodule emergence rather than by inhibition of root hair infection. Clearly, the pending fruition of an infection is subject to internal (or auto-) regulation that exists both prior and subsequent to nitrogen fixation.

Recently, we isolated 15 independent soybean mutants that continued to nodulate in the presence of nitrate (6). These lines were designated *nts* (nitrate-tolerant-symbiosis) mutants. Previously, only four nodulation mutants of soybean had been discovered and all of these naturally occurring mutants are characterized by decreased nitrogenase activity. Soybean plants homozygous for the recessive mutation *rj₁ rj₁* are resistant to nodulation; however, the blockage can be partially circumvented by inoculation with high *R. japonicum* cell densities (18). *Rj₂*, *Rj₃*, and *Rj₄* are strain-specific dominant mutations that condition ineffective nodulation (3, 28, 29).

In this paper we report the nodulation characteristics of mutant line *nts382*. The effects of various nitrogen sources on nodulation, nitrogen fixation, and plant growth are described, as well

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as the influence of two *R. japonicum* strains on the expression of the supernodulation phenotype. Data are presented that indicate that the mutant character is inherited as a Mendelian recessive.

MATERIALS AND METHODS

Soybean (*Glycine max* [L.] Merr.) cv Bragg was used in this study. Mutant line *nts382* was selected for increased nodulation in the presence of high nitrate concentrations. The isolation procedure has been described in detail (6) and is summarized in Figure 1. M_4 plants of mutant line *nts382* were used here. Plants were cultured in either pots of river sand or in Leonard jars (9).

In experiments testing the effect of various nitrogen sources on nodulation and N_2 (C_2H_2) fixation, plants were cultured in 20 cm diameter pots of sand. Three Bragg or *nts382* seeds were planted 1 cm below the surface and inoculated with *Rhizobium japonicum* strain CB1809 (= USDA136). The pots were re-inoculated at day 4. The nutrient solution was as used by Herridge (12), except that all nutrients other than $CaCl_2$ and the nitrogen source were administered at quarter strength for the first 2 weeks. KNO_3 (2.75 and 5.5 mM), urea (5.5 mM, i.e. 11 mM N), NH_4Cl (5.5 mM), NH_4NO_3 (5.5 mM, i.e. 11 mM N) and KCl (as control) were added to the nutrient solution as required. The pots were watered daily with 700 ml of nutrient solution, which was sufficient to flush out residual nutrients from the previous watering.

M_2 family 382 was one of 15 families that segregated for the *nts* phenotype (Fig. 1). Both *nts* variants and wild-type (non-*nts*) siblings were saved to produce M_3 families (i.e. families derived from single M_2 plants). Those M_3 families derived from wild-type M_2 plants were screened for segregation of the *nts* character (Fig. 1). Seeds were planted in 25 cm diameter pots of river sand (12 seeds per pot). The pots were inoculated with *R. japonicum* strain USDA110 at day 0 and day 4. Nutrients were administered as described above, except that all pots received 1.4 L of nutrient solution 3 times a week. After 7 weeks of growth, the plants were screened for the *nts* character.

To control access of *R. japonicum* strains to the roots (9), Bragg and *nts382* were cultured in Leonard jars. Seeds were surface sterilized by rinsing in 95% ethanol followed by immersion of the seeds for 10 min in 3% NaOCl. After several rinses

in sterile distilled H_2O , the seeds were transferred to water agar and germinated at 28°C in the dark. When the radical was approximately 0.5 to 1 cm long (2–3 d after sterilization), the seedlings were transferred to Leonard jars (N-free or supplemented with KNO_3 or urea). The jars either remained uninoculated or were inoculated with *R. japonicum* strains CB1809 or USDA110.

The plants were cultured in a temperature-controlled glasshouse (mean maximum temperature = 26.9°C; mean minimum temperature = 14.8°C) at a latitude of 37° 17'S. Incandescent bulbs supplemented natural light such that the photoperiod was 16 h. At harvest, the plants were measured for nodule number, nodule fresh weight, nitrogenase activity, and plant fresh weight.

Acetylene reduction was used to estimate nitrogen fixation on intact plants (11). Plants were incubated in 1040 ml air-tight jars with 2 to 3 ml of distilled H_2O (to prevent desiccation) at 25 to 27°C. The atmosphere in the jars was 6% acetylene in air. Rates were calculated from integrator units obtained from a Hewlett-Packard Integrator-Recorder 3390 coupled to a HP5590A flame ionization gas chromatograph. Samples were taken at 40 min and rates were determined to be linear over that period. Genotype and combined nitrogen effects were statistically tested by analysis of variance using the general statistical program Genstat (1). The LSD was computed when the *F* statistic was significant (0.05 level of significant). Chi-square analysis was used to statistically test segregation ratios. It was necessary to include Yates correction term in chi-square calculations due to the size of the expected classes (25).

RESULTS

Effect of Combined Nitrogen on Nodulation, Nitrogenase Activity, and Growth. Bragg and *nts382* were inoculated with *R. japonicum* CB1809 and cultured for 4 weeks in the presence and absence of various combined nitrogen sources. The nitrogen-free treatment received 5.5 mM KCl as a control.

Nodule Number. Under all the conditions tested, 4-week-old *nts382* plants had considerably more nodules than wild-type Bragg plants (Fig. 2). In the parent cultivar, all nitrogen sources reduced nodule number (Figs. 2 and 3). In contrast, mutant line *nts382* grown on KNO_3 or urea had increased nodule number per plant over that of the KCl controls (Fig. 2). When data for nodule number are expressed per plant fresh weight, nodule number in *nts382* was unaffected by increasing KNO_3 concentration (Fig. 3). Urea (5.5 mM), on the other hand, caused a small reduction in nodule number per plant biomass in the mutant. The relative degree of inhibition of nodule formation by urea was greater in Bragg than in *nts382* (Fig. 3). Mutant line *nts382* formed nodules on 5.5 mM NH_4Cl and NH_4NO_3 , which totally prevented nodule formation in Bragg (Figs. 2 and 3). Ammonium chloride inhibited growth to a larger extent than did ammonium nitrate (Table I); however, NH_4NO_3 was more inhibitory than NH_4Cl on nodule formation in *nts382* (Fig. 3).

Nitrogenase (Acetylene Reduction) Activity per Plant Biomass. In Bragg, supplementing the nutrient media with nitrate or urea caused a substantial reduction in nitrogenase activity measurable 4 weeks after planting (Fig. 4). In contrast, 2.75 mM KNO_3 significantly stimulated acetylene reduction per gram plant fresh weight in *nts382*. Nitrogenase activity for *nts382* plants cultured on 5.5 mM KNO_3 was not significantly different from the nitrogen-free mutant plants (Fig. 4). For both Bragg and *nts382*, urea was more inhibitory than equimolar KNO_3 concentrations. Regardless of the combined nitrogen supply, *nts382* had higher nitrogenase activity than Bragg, but the difference under N-free conditions was not significant at the 0.05 level of significance. Nitrogenase activity per plant fresh weight in 4-week-old *nts382* plants cultured on 2.75 and 5.5 mM KNO_3 was significantly higher than for Bragg plants cultured in the absence of nitrogen

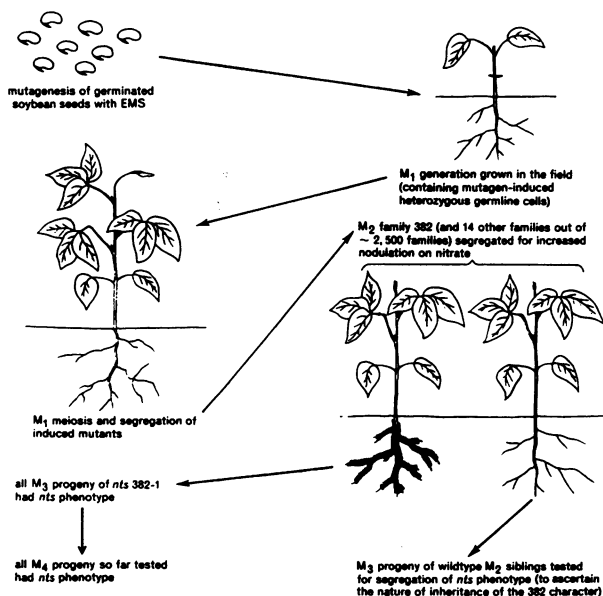


FIG. 1. Schematic representation of the isolation of *nts382* from mutagenized Bragg soybeans.

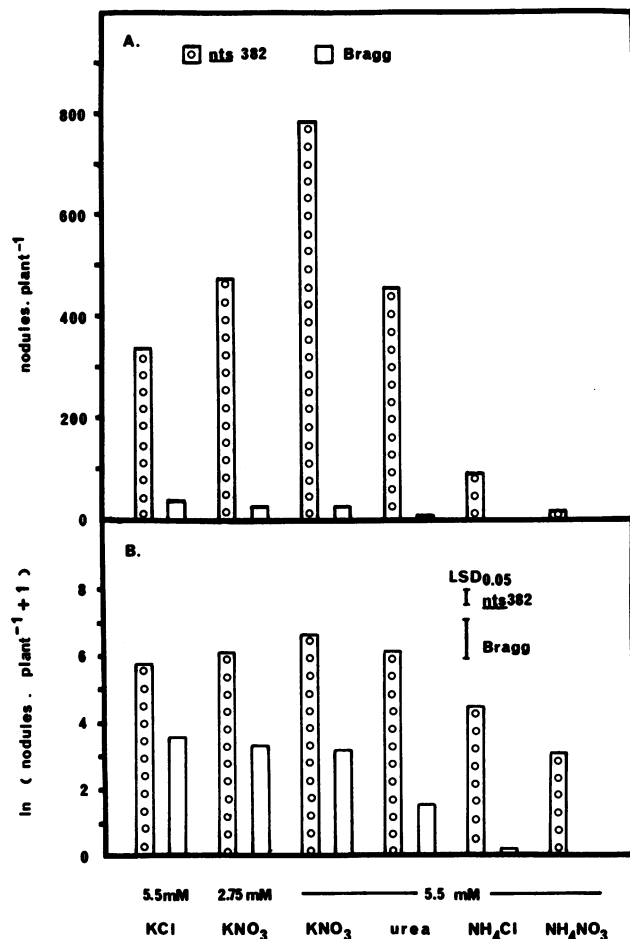


FIG. 2. Effect of nitrogen source on nodule number per plant in Bragg and *nts382* plants. Plants were cultured and harvested as in Table I. Raw data (panel A) required \log_e transformation (panel B) to satisfy assumptions for an analysis of variance. Each column represents the mean of four to eight plants. Genotype, combined nitrogen, and interaction effects were significant.

(Fig. 4).

Visual Characteristics of Nodulation by *nts382*. Figure 5 illustrates a sample of plants that contributed to the data presented above. The photographs indicated that prolific nodulation in *nts382* may occur at the expense of root growth. Mutant line *nts382* continued to nodulate prolifically throughout development. Figure 6 shows the tap root of a *nts382* plant harvested at the podfill stage of development.

Plant Fresh Weight. Nitrate (both concentrations) and urea treatments produced healthy vigorous plants and stimulated growth of Bragg and *nts382* over that of the respective KCl controls (Table I). Regardless of genotype, plants cultured on 5.5 mM NH₄Cl and NH₄NO₃ were unhealthy and stunted. Ammonium nitrate produced no effect on plant fresh weight after 4 weeks of culture, whereas NH₄Cl inhibited fresh weight accumulation in both Bragg and *nts382*. Plant fresh weight was consistently higher in Bragg than in *nts382* (Table I).

Axenic Culture in Leonard Jars. The observations obtained from pots inoculated with *R. japonicum* strain CB1809 and described above were confirmed in Leonard jars supplemented with KNO₃ and urea. Both *R. japonicum* strains tested, namely CB1809 and USDA110, elicited the *nts* phenotype in mutant line 382. Uninoculated controls of *nts382* and Bragg did not nodulate.

Nodulation Pattern. Table II shows the tap root nodulation

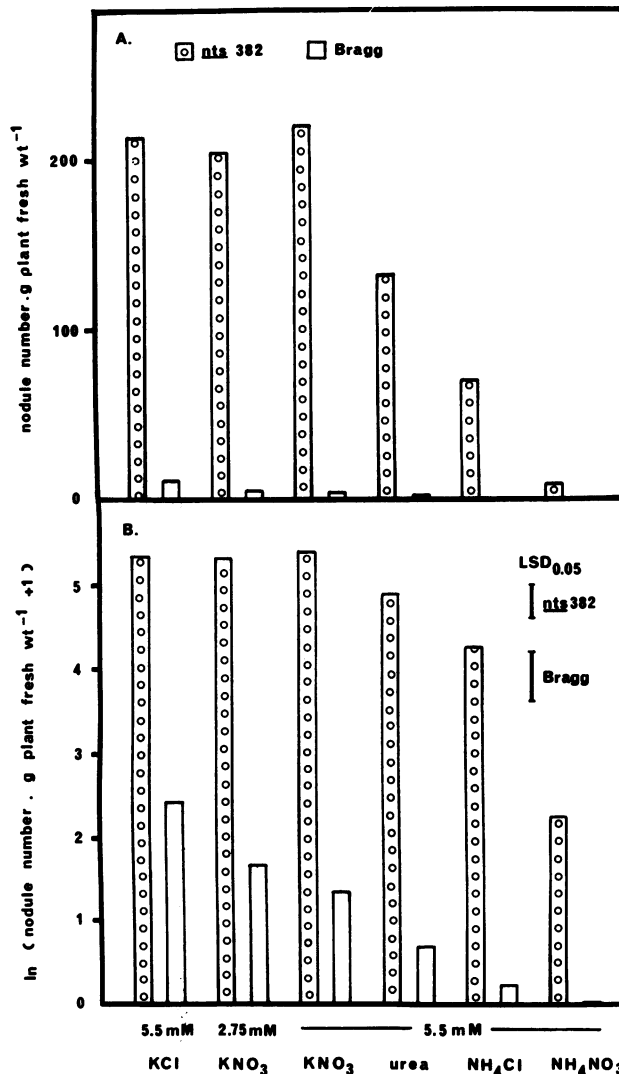


FIG. 3. Effect of nitrogen source on nodule number per plant biomass in Bragg and *nts382*. Plants were cultured and harvested as in Table I. Raw data (panel A) required \log_e transformation (panel B) to satisfy assumptions for an analysis of variance. Each column represents the mean of four to eight plants. Genotype, combined nitrogen, and interaction effects were significant.

pattern of N₂-dependent *nts382* and Bragg plants. Tap root length was less in nodulated *nts382* plants than in Bragg plants cultured under identical conditions (Table II). The tap root nodulation interval is defined as the distance between the uppermost and lowermost nodule on the tap root. This parameter was larger in *nts382*, both in absolute terms (2 times that of Bragg) and when expressed as a percentage of tap root length (3 times that of Bragg). Nodule density (nodules/cm) was also increased in *nts382* (Table II). Nodule density, expressed on the total tap root length was 9 times higher in *nts382* than in Bragg. Within the nodulation interval, the nodule density was 2.5 times higher in the mutant line (Table II). A similar contrast in nodulation pattern was observed on lateral roots and, furthermore, on *nts382* plants cultured on various nitrogen sources (Fig. 5).

Inheritance of the *nts382* Character. Following mutagenesis on M₁ seeds, resultant M₁ plants were grown through to produce M₂ seeds. Seeds from each M₁ plant were grouped together to give M₂ families. Mutant line *nts382* was selected from a M₂ family that segregated for the mutant phenotype (Fig. 1). From 17 plants in M₂ family 382, two expressed the *nts* phenotype and

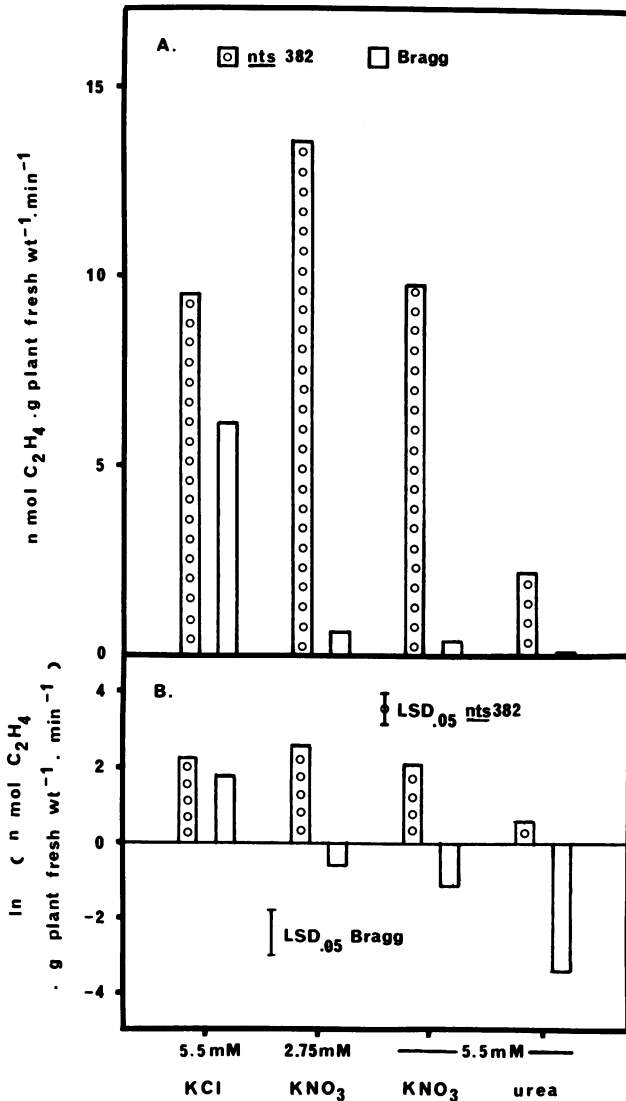


FIG. 4. Effect of potassium nitrate and urea on nitrogenase (acetylene reduction) activity in Bragg and *nts382* plants. Plants were cultured and harvested as in Table I. Raw data (panel A) required \log_e transformation (panel B) to satisfy assumptions for an analysis of variance. Each column represents the mean of four to eight plants. Genotype, combined nitrogen, and interaction effects were significant.

the remaining 15 plants expressed the wild-type phenotype and were indistinguishable from the parent cultivar Bragg for nodulation in the presence of nitrate. In the M_2 screen, plants were harvested 6 weeks after planting and culture on 5 mM KNO₃. The nodule per plant for M_2 *nts* segregants was 146 ± 71 (\pm SD) and for M_2 wild-type (non-*nts*) segregants it was 26 ± 11 . Bragg plants cultured under identical conditions had 19 ± 7 nodules per plant.

M_3 progeny derived from M_2 *nts* plants all expressed the mutant character indicating that these M_2 *nts* plants were homozygous for the mutation. Similarly, all M_4 progeny (used in experiments described above) had the *nts* phenotype.

M_2 wild-type (non-*nts*) segregants were also repotted and grown through to seed. The seeds derived from each M_2 plant were grouped together to give M_3 families. One of six such M_3 families segregated 14 *nts* plants:40 wild-type plants. This ratio closely approximates 1:3, the expected segregation ratio for a recessive character in progeny derived from a self-fertilized heterozygous wild-type plant (chi-square was equal to 0.00 and was

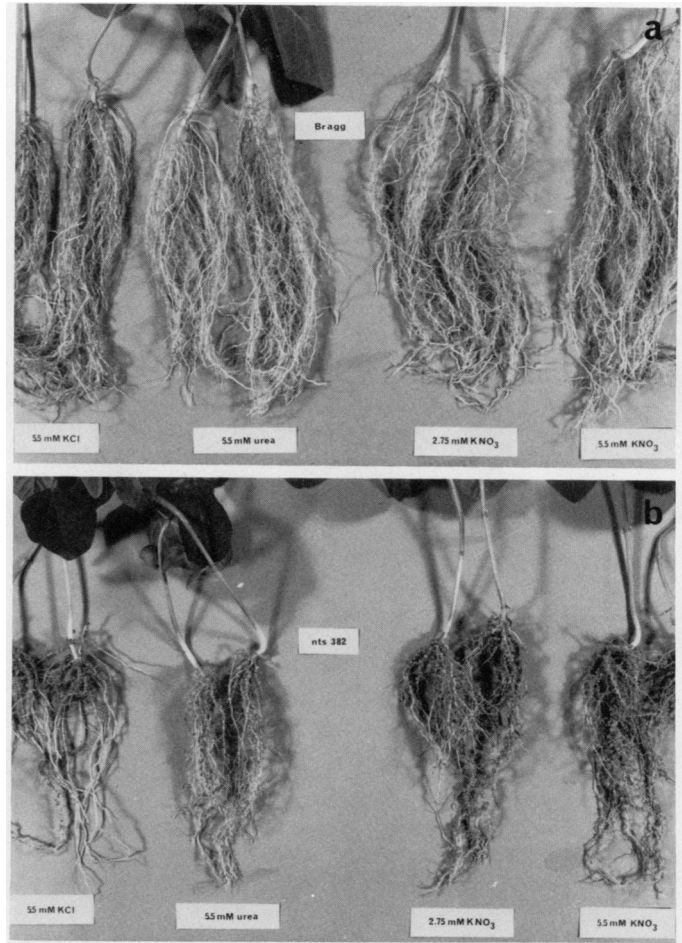


FIG. 5. Effect of potassium nitrate and urea on nodulation in Bragg (a) and *nts382* (b) plants. Plants were cultured and harvested as in Table I.

not significant).

Mutant *nts* plants in the segregating M_3 family qualitatively had a similar phenotype to the original M_2 *nts* selections, and to the M_3 and M_4 progeny derived therefrom. Nodule number per plant for these nitrate-grown *nts* segregants was 789 ± 181 (\pm SD), whereas wild-type segregants had 32 ± 19 nodules per plant. Furthermore, plant fresh weight was lower in *nts* segregants; 8.23 ± 2.33 (\pm SD) g per plant compared to 15.19 ± 3.20 g per plant for wild-type segregants.

After 7 weeks culture on 5 mM KNO₃, *nts* segregants had 23 times the nodule fresh weight per plant biomass of wild-type segregants (Table III). Increased nodulation in *nts* plants was observed on the tap root as well as on the lateral roots (Table III); this trend was consistent with the nodulation pattern shown in Figure 5 for *nts382* grown under a variety of combined nitrogen regimes. As a result of the increased nodule mass, nitrate-grown *nts* segregants had 7 times the nitrogenase (acetylene reduction) activity per plant biomass of nitrate-grown wild-type segregants (Table III).

DISCUSSION

Mutant line *nts382* was one of several *nts* (nitrate-tolerant-symbiosis) mutants selected from an M_2 population of parent cultivar Bragg for increased nodulation in the presence of nitrate (6). Individual M_2 families (resulting from a single mutagenized M_1 seed) were originally screened. M_2 family 382 segregated for the *nts* character (Fig. 1), as did all of the M_2 *nts* mutant families.

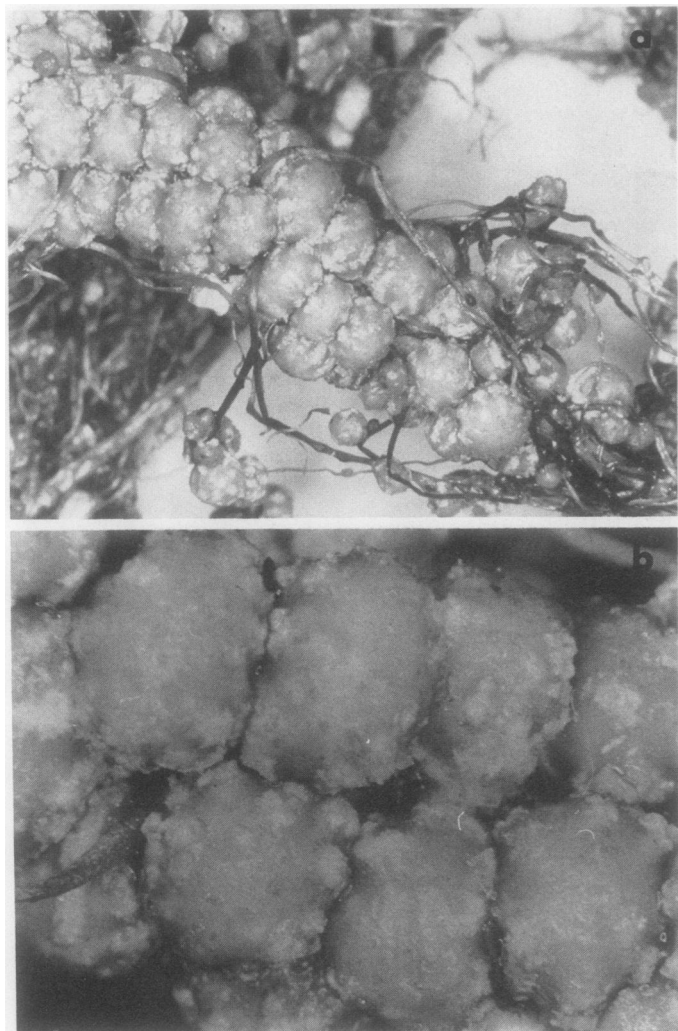


FIG. 6. Nodulation of *nts382* at the podfill stage of development. a, The tap root of a *nts382* plant that had been cultured in a 25-cm pot of vermiculite. This plant had received slow-release nitrogen fertilizer throughout growth. b, Close-up of Figure 6a.

Table I. Plant Fresh Weight of Bragg and *nts382* Plants Cultured for 4 Weeks on Various Nitrogen Sources

The plants were grown in 20-cm sand pots as described in "Materials and Methods."

Nitrogen source	Plant Fresh Wt ^a	
	Bragg	<i>nts382</i>
	<i>g</i>	
0 mM (5.5 mM KCl)	3.24 (100) ^b	1.96 (100) ^b
2.75 mM KNO ₃	5.61 (173)	2.37 (121)
5.5 mM KNO ₃	8.05 (248)	3.68 (188)
5.5 mM urea	4.80 (148)	3.51 (179)
5.5 mM NH ₄ Cl	2.12 (65)	1.27 (65)
5.5 mM NH ₄ NO ₃	3.04 (94)	2.18 (111)

^a LSD_{0.05} = 1.56 for Bragg and 1.08 for *nts382*. ^b Plant fresh weight as per cent of 5.5 mM KCl control.

This observation indicated that each of the mutant families arose from independent mutation events and that the mutants were a result of the mutagenesis program (6). M₂ *nts382* selections were homozygous for the mutation, since all M₃ progeny derived thereof had the *nts* phenotype, as did all M₄ progeny, including the plants used in some experiments described here. M₃ progeny

Table II. Tap Root Nodulation Pattern of N₂-Dependent *nts382* and Bragg Plants

Plants were inoculated with *R. japonicum* USDA110 and harvested 9 weeks after planting. The pots were watered daily with N-free nutrient solution throughout the experiment.

Tap Root Parameter	<i>nts382</i> ^a	Bragg ^b	LSD _{0.05}
Root length (cm)	16.6	28.1	3.1
Nodulation interval (cm)	14.4	7.6	2.1
Nodulation interval (% of root length)	86.5	27.1	6.7
Nodule density on root length (nodules·cm ⁻¹)	4.32 (2.0) ^c	0.47 (0.67)	—(0.15)
Nodule density on nodulation interval (nodules·cm ⁻¹)	5.07	2.10	0.99

^a Each entry in the table for *nts382* represents the mean of seven plants. ^b Each entry in the table for Bragg represents the mean of 28 plants. ^c Raw data required square-root transformation to satisfy assumptions for an analysis of variance; means and LSD of transformed data are shown in parentheses.

Table III. Nodulation and Nitrogenase (Acetylene Reduction) Activity in KNO₃-Grown *nts* and Wild-Type Segregants from a *nts382* M₃ Family

This family was derived from a wild-type M₂ plant. The segregation ratio was 14 *nts*:40 wild type, which approximates 1:3 (chi-square = 0.00 and was not significant). Plants were inoculated with *R. japonicum* USDA110 and harvested after 7 weeks growth on 5 mM KNO₃. Data are expressed per g plant fresh weight.

Symbiotic Parameter	<i>nts382</i> ^a	Wild-type ^b	LSD _{0.05}
Nodule number·g plant fresh wt ⁻¹			
on tap root	9.6 (2.22) ^f	0.7 (-0.45)	(0.20)
on lateral roots	89.2 (9.4) ^d	1.5 (1.1)	(0.4)
mg nodule fresh wt·g plant fresh wt ⁻¹	145.3 (12.0) ^d	6.2 (2.5)	(0.5)
nmol C ₂ H ₄ ·g plant fresh wt ⁻¹ ·min ⁻¹	8.0 (2.05) ^f	1.1 (0.00)	(0.76)

^{a,b} Each entry in the table is the mean of 14 and 18 plants, for *nts* and wild-type, respectively, except that acetylene reduction data are the means of seven *nts* and nine wild-type plants. ^{c,d} Raw data required either log_e (c) or square-root (d) transformation to satisfy assumptions for an analysis of variance; means and LSD of transformed data are shown in parentheses.

of a wild-type M₂ plant segregated 3 wild type:1 mutant, indicating that the *nts382* character is inherited as a Mendelian recessive.

Regardless of the presence or absence of combined nitrogen, *nts382* nodulated more than the parent cultivar Bragg. This trend was consistent over a range of combined nitrogen sources that caused varying degrees of inhibition of nodulation in Bragg. For example, under conducive conditions for nodulation, 4-week-old N₂-dependent *nts382* plants had 9 times the nodule number of N₂-dependent Bragg plants. Similarly, under conditions that totally prevented nodulation in Bragg (5.5 mM NH₄Cl or NH₄NO₃), *nts382* plants were still nodulated. The mutant line also had increased nodulation in Leonard jars, in deep soil pots (soil obtained from soybean field at Breeza, NSW, Australia) and in Georgia (USA) fieldplots. Nodule initiation and nodule growth are coordinated in *nts382*, and *nts382* plants have a considerably larger nodule mass than wild-type plants (6; Table III). We use the term supernodulator to describe *nts382*, since this soybean

genotype has an increased nodule mass under a wide range of environmental conditions.

Although *nts382* is a supernodulator, it is not a constitutive nodulator, since it still requires the inducer (*i.e.* *R. japonicum*) to be present. The two strains of *R. japonicum* used, namely CB1809 and USDA110, elicited the *nts* phenotype in the mutant line. Both these strains form an effective symbiosis with the parent cultivar. A wider spectrum of fast- and slow-growing *R. japonicum* strains, that vary in their ability to nodulate the parent cultivar, were tested on *nts382* to ascertain whether the *nts* character confers a change in the promiscuity of the host plant. Those strains that nodulated Bragg elicited the mutant phenotype on *nts382*, and strains that were nod⁻ on the parent cultivar were also unable to induce nodule formation on *nts382*.

In Bragg, all nitrogen sources caused a significant reduction in the symbiotic parameters that were measured (Figs. 2–5). In contrast, enriching the nutrient solution with KNO₃ did not inhibit any symbiotic parameters in *nts382* and was in some cases stimulatory (Figs. 2 and 4). Urea marginally stimulated nodule number per plant but significantly inhibited nodule number and nitrogenase activity per gram plant fresh weight in the mutant line. Consistent with the comparative effects of KNO₃ and urea on *nts382*, urea was also more severe on symbiotic development in Bragg (Figs. 2–4). This is different to what Vigue *et al.* (30) found with soybean cultivar Steele. Using a carboxy resin to buffer the culture media (17), they demonstrated that nitrate was more inhibitory on nodulation than was urea. To minimize the drop in pH associated with urea and ammonium utilization (14), the pots in our experiments were flushed daily with 700 ml of nutrient solution. This procedure, without a specific buffering agent, produced healthy vigorous urea-fed plants that had significantly stimulated fresh weights in Bragg and *nts382* (Table I). Nevertheless, it is unclear whether pH fluctuations played a role in inhibition of the symbiosis by urea, NH₄Cl, and NH₄NO₃. This, however, is not a contentious issue, since the salient feature of *nts382* illustrated by these experiments is that it consistently nodulated prolifically in comparison to parent cultivar Bragg, regardless of the environmental conditions imposed by the provision of various nitrogen sources.

Since prolific nodulation by *nts382* was not confined to culture on nitrate, it is unlikely that increased nodulation on nitrate resulted from an inability to utilize nitrate. KNO₃ enhanced the growth of *nts382* plants (Table I) and, furthermore, *nts382* has the same nitrate reductase activity as Bragg (6). Indeed, *nts382* is a mutant in the regulation of nodule initiation and nodule

development, and the autoregulation mechanism normally limiting nodulation in wild-type soybeans (23) is anomalous. The nodulation interval is extended and nodule density is increased in *nts382* (Table II; Fig. 5). Although nodule emergence and subsequent nodule growth are coordinately controlled in *nts382*, specific nitrogenase activity (activity/g nodule fresh weight) is not coordinately regulated with these two nodule development parameters in the mutant (6). The fact that *nts382* plants have increased nitrogenase activity per plant biomass under adverse conditions for symbiotic development is a result of increased nodule mass.

In *nts382* mutant plants, regulation of nodulation by environmental conditions (such as by nitrate supply) is coordinated with internal (or auto-) regulation. The supernodulation observed in *nts382* under N-free conditions persisted when combined nitrogen was added to the nutrient media. The increased tolerance of *nts382* to nitrate and other combined nitrogen sources was seen in an accentuation in the magnitude of difference between *nts382* and Bragg as nitrogen sources were added to the nutrient solution (Table IV). For example, in the absence of combined nitrogen *nts382* plants had 9 times as many nodules as Bragg plants (Table IV). As combined nitrogen was added to the nutrient solution, the degree of inhibition in Bragg was not seen in *nts382* and for plants cultured on 5.5 mM urea, for example, *nts382* had 65 times as many nodules per plant as did Bragg. A similar trend was evident for all the other symbiotic parameters measured (Table IV). Clearly, *nts382* is a mutant in the autoregulation pathway and is also less sensitive to regulation by external conditions. Recently, a pea mutant was described to have similar nodulation characteristics (15). Pea mutant *nod₃* nodulated more than the parent cultivar in the absence of nitrate and also nodulation was less affected by nitrate in the mutant (15).

In addition to being useful in studying regulation of nodule development in general, *nts382* is of immense value for critically analyzing and refining some hypotheses that have been put forward to explain nitrate inhibition of nodule development. The carbohydrate deprivation hypothesis argues that nitrate reduction and assimilation deprives the nodule of carbohydrate necessary for growth and development. The alternative hypothesis that has also received support is that products on nitrate reduction, particularly nitrite, are responsible for nitrate inhibition (6). The nodulation characteristics of *nts382* and Bragg indicate that mass flow carbohydrate deprivation or products of nitrate reduction cannot directly cause inhibition of nodule development. Growth of *nts382* was stimulated by potassium nitrate (Table I) and *nts382* had normal nitrate reductase activity (6) indicating the mutant did utilize nitrate and yet nodule development was not inhibited early in plant development (0–4 weeks after planting). Carbohydrate deprivation or products of nitrate reduction may inhibit nodule development in wild type, but it must be mediated through the control mechanism that is anomalous in *nts382*. Further detailed studies on nodulation and nitrate metabolism of *nts382* and related mutants are continuing.

Table IV. Symbiotic Parameters of *nts382* Expressed as a Proportion of Bragg

Each datum in the table was computed by dividing the value of the symbiotic parameter for *nts382* by the value for Bragg cultured under identical conditions (*i.e.* *nts382* + Bragg). Nitrogen sources are listed in order of increasing severity on nodulation in Bragg. Plants were harvested 4 weeks after planting. The inoculant strain was *R. japonicum* USDA110.

Nitrogen Source	Symbiotic Parameter ^a		
	Nodules /plant ^b	Nodules/g plant fresh wt ^c	Nitrogenase activity ^d
0 mM (5.5 mM KCl)	9	21	1.6
2.75 mM KNO ₃	18	51	22.6
5.5 mM KNO ₃	31	73	29.5
5.5 mM urea	65	132	36.0
5.5 mM NH ₄ Cl ^e	∞	∞	—
5.5 mM NH ₄ NO ₃ ^e	∞	∞	—

^a *nts382* + Bragg. ^{b,c,d} Data from Figures 2, 3, and 4, respectively. ^e NH₄Cl and NH₄NO₃ totally prevented nodule formation in Bragg.

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