EVALUATION OF THE EFFECT OF VARIOUS NITROGEN FORMS ON THE *LUPINUS ALBUS* SEED PROTEIN COMPOSITION

DANUTA CIESIOŁKA¹, TADEUSZ ANISZEWSKI², WOJCIECH WYSOCKI¹, RADOSŁAW PILARSKI¹, KRZYSZTOF GULEWICZ^{1*}

^{1*} Laboratory of Phytochemistry, Institute of Bioorganic Chemistry PAS Noskowskiego 12/14, 61-704 Poznań, Poland e-mail: krysgul@ibch.poznan.pl

² Laboratory of Applied Botany, Department of Biology, University of Joensuu PO Box 111, 80101 Joensuu, Finland

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ABSTRACT

The influence of different nitrogen forms: (N_2) , $[N_2+(NH_4^++NO_3^-)]$, (NH_4^+) , (NO_3^-) , $(NH_4^++NO_3^-)$ and $(-NH_2)$ on changes of albumin, globulin, prolamin-glutelin, non-fractioned nitrogen and non-protein nitrogen fractions of the protein seed of alkaloid-low content Lupinus albus L. cv. Butan has been studied. The experiments were performed in a greenhouse on perlite using in all cases the constant P, K, Mg and micronutrients (B, Zn, Mn, Cu, Mo, Fe) fertilization. The control was the treatment without any nitrogen support (Nd). It was clearly shown that nitrogen form has significant effect not only on the seed yield and seed protein content, but also on the composition of protein fractions and on the biological value of lupin protein. The main protein fraction of the seeds from all treatments were albumins (16.73-26.10 mg N/g). Among all the treatments, the highest level of globulin was observed for the seeds of plant growing with the symbiotic nitrogen form (15.26 mgN/g), while the lowest one for the control (Nd) (6.86 mg N/g). Symbiotic nitrogen (N₂) treatment clearly increased the glutelin-prolamin fraction while the addition of mineral nitrogen (NH₄+NO₃-) decreased this fraction from 8.40 to 4.48 mg N/g. The lowest level of the glutelin-prolamin fraction was in the absence of any nitrogen (Nd). Non-protein fraction (Nnp) was highest in the case of plants treated with (-NH2) (9.92 mgN/g), and the lowest in the absence of nitrogen (Nd) (4.90 mgN/g). The level of non-fractioned nitrogen (Nr), with exception of $[N_2+(NH_4^++NO_3^-)]$ and $-NH_2$ treatments, was closest to the start material. The protein fractions (albumins, globulins and glutelins and prolamins) were also electrophoretically characterized. These analysis confirmed the changes in protein composition of particular fractions under the effect of various nitrogen forms used as a fertilizer.

KEY WORDS: lupin, nitrogen forms, protein, fractional composition.

INTRODUCTION

Nitrogen that consists of an indispensable element of protein, nucleic acid and other important nitrogen compounds can be uptaken by plants in the form of NH₄⁺ or NO₃⁻ and, in specific conditions, in the amide form –NH₂ (urea). The preference of plants to the particular form depends mainly on its availability that is conditioned by microbiologic activity dependent on pH, oxygen condition, soil nutrient availability and temperature. Acidofilic and low oxidation-reduction potential tolerating plants uptake easier NH₄⁺ cation while the calciphilous ones prefer anion form, NO₃⁻. The increase of soil pH increases NH₄⁺ and decreases NO₃⁻ relative uptake rates. At pH above 7.5, NH₄⁺ undergoes to free NH₃. This compound is also formed during reduction of N₂ in root nodules of legumes and during hydrolysis of urea used as nitrogen fertilizer. Contrary to

NO₃⁻, ammonia at high concentration is toxic for plants and has to be immediately deactivated by carbon chains. The main source of nitrogen for legumes is atmospheric, gaseous N₂. Although currently we are not able to explain all aspects and experimental results concerning the regulation of nitrogen fixation in legumes (Schulze 2004), it is known that nitrogen content in bacterial lupin nodules has an influence on nitrogen content in seeds. Fixation of nitrogen in biological way is not energetically free. Plants have to contribute significant amount of energy in the form of photosynthates (photosynthesis derived sugars) and other nutritional factors for the bacteria. The symbiotic nitrogen fixation by legumes may be inhibited by the presence in soil of both, nitrate and ammonium (Latimore et al. 1977; Evans 1982; Evans et al. 1987; Malik et al. 1987; Carroll and Mathews 1990; Daimon et al. 1999; Daimon and Yoshioka 2000). As shown by Ciesiołka et al. 2005, different

nitrogen forms used as fertilizer had a significant effect not only on the development and yield of both alkaloid-low and alkaloid-rich lupin cultivars, but also on total raw protein, alkaloid and α -galactoside content in seeds. Proteins can be classified into two groups: globular and fibrous proteins (Wilson and Walker 2001). Globular proteins are generally water soluble and they may contain a mixture of α -helix, β -pleated sheet and random structure. Fibrous proteins are structural proteins, generally insoluble in water and consisting of long cable-like structures build entirely of either helical or sheet arrangements. The native state of a protein is its biologically active form. Lupin seed proteins are composed dominantly by globular proteins, but protein quality depends on lupin species and biotype (Aniszewski 1993a, 1993b).

The objective of our studies was alkaloid-low (sweet) lupin *L. albus* L. cv. Butan. The choice of lupin was not accidental. Lupin could be used as a source of protein or fiber or for supplementation in already existing or new products (Aniszewski 1993a, 1993b; Mohamed and Rayas-Duarte 1995). The feature which distinguishes lupin from among other plants may be determined as follows: *i.* belongs to highest protein plants, *ii.* has modest soil and climatic demands, *iii.* leaves big organic mass in soil, *iv.* improves airwater conditions of soil, *v.* makes accessible macro and micro elements eluted to sublayer of soil, *vi.* lupin protein has good functional properties, i.e. emulsifying power, binding, and foaming properties (Edvin 1974; Gulewicz et al. 1994; Aniszewski 1995a, 1995b; Ciesiołka et al. 2005).

The main aim of our studies was to answer the following questions: How do different nitrogen forms affect the quality of seed proteins? Can various nitrogen sources modify the chemical composition of legume proteins and improve or worsen its biological value? The presented studies were supported by the chemical analysis devised by Osborne (1924), which were based on differential solubility of the seed protein fractions (Nelson 1969; Oomah and Bushuk 1983).

MATERIAL AND METHODS

Greenhouse experiments

The experiments were carried out on Perlite in a greenhouse at determined conditions within a period of April-August, 2005. The seeds of lupin, sterilized with 1.5% sodium hypochlorite, were sowed in 1.5 l capacity pots, 13 cm (diameter) × 13 cm (height) filled with Perlite on April 12. Seven fertilizer treatments with different forms of nitrogen were used in the experiment as follows:

T1 – nitrogen deficiency: (Nd)

 $T2 - N_2 - (Bradyrhizobium lupini): (N_2)$

 $T3 - N_2 - (Bradyrhizobium lupini) + NH_4NO_3$:

 $[N_2 + (NH_4^2 + NO_3^-)]$

 $T4 - NaNO_3$: (NO_3^-)

 $T5 - (NH_4)_2SO_4$: (NH_4^+)

 $T6 - NH_4NO_3$: $(NH_4^+ + NO_3^-)$

 $T7 - NH_2CONH_2$: $(-NH_2)$

Each treatment was done in six repetitions. One pot containing three plants constituted one repetition. During vegetation period for each pot of T4, T5, T6, T7 treatments, the total dose 116 mg N of suitable form was used. The dose was four applications of 29 mg N one per week. The first

nitrogen fertilization was used during the formation of flower buds. In the case of T3 treatment, single dose of 29 mg N per pot in the form (NH₄⁺+NO₃⁻) was applied. P₂O₅, K₂O, MgO and micronutrients (Zn, Mn, Cu, Mo, Fe) were used single in the doses of 22.0; 34.0; 8.0 and 9.0 mg per pot, respectively. During the vegetation period, the plants were irrigated with water in sufficient amount to avoid the lixiviation of nitrogen from perlite.

Seed harvest and determination of protein

The seeds were harvested from each combination and analyzed afterwards. The analysis also involved the seeds sowed (SM – start material). The crude protein in seeds from all the treatments and from the SM was determined by Kjedahl method using a Kjeltec Auto Distillation 2200 apparatus (FOSS TECATOR).

Determination of fractional composition of seed proteins

In general, the fractionation of seeds protein was supported by the method described by Michael and Blume (1961) with some modification (Peretiatkowicz et al. 1988), showed in Figure 1. The total nitrogen amounts of initial material and fractions marked in Figure 1 by a thickened frame (A – albumins, Nnp – non protein nitrogen, G – globulins, Gt+P glutelins and prolamins, Nr-residual nitrogen) were determined by the Kjedahl method.

Sodium dopdecyl sulphate polyacrylamide gel electrophoresis (SDS-PAGE) of protein fractions

The precipitated albumin fraction with TCA (Fig. 1) was resolved in TBS buffer (25 mM NaCl and 25 mM TRIS/HCl pH 7.5) and passed through the Amicon filter (10 kD). The content of protein in albumins and other lupin fractions was determined according to the Bradford (1976) method. Next, 20 μ g of protein were precipitated with four volume of acetone. The samples were frozen for 40 min

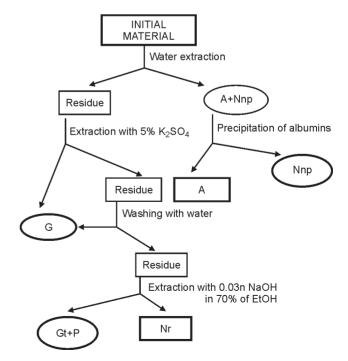


Fig. 1. Scheme of seeds' proteins fractionation. Nnp - non-protein fraction; A - albumin; G - globulin, Gt+P - glute-lin+prolamin; Nr - residual fraction

and centrifugated at 12 000 rpm for 15 min. After removing of aceton, the pellet was dryed in dessicator. The dry samples were solved in 10 μL of water and 10 μL of sample buffer (SB) cointaining: 1 mL glicerol, 0.5 mL β -mercaptanol, 2 mL of 10% SDS, 0.25 mL of 0.05% bromophenol blue (BPB) and 1.25 mL of stocking buffer (3 g of TRIS, 14.4 g of amino acetic acid and 10 mL of 10% SDS-all components in 1000 mL of water). Afterward, the samples were boiled during 3 min and applied onto 10% acrylamide gel. The SDS PAGE was performed on the Biometra Power Pack P25 apparatus at 7.5 mA (concentrating gel), 14 mA (developing gel) for 3 hours (Lammeli 1970). Then, the gels were passed to the fixing agent (10% acetic acid in 50% ethanol) and dyed with Coomassie Brillant Blue.

Statistical analysis

Multi-way analysis of variance (ANOVA) was conducted using MATLAB program, version 6.5 (MathWorks Inc.) available at Poznań Supercomputing and Networking Center of the Institute of Bioorganic Chemistry, PAS, Poznań, Poland.

RESULTS

Yield of seeds, seed protein content and yield of protein

The yield of seeds, content of protein and yield of protein in seeds harvested from the plants, growing under various nitrogen forms are presented in Table 1. The highest yield of seeds (38.08 g) was collected in the treatment, where nitrogen was used as [N₂+(NH₄+NO₃)] form, while the lowest one (2.81 g) in the case of nitrogen deficiency (Nd). Generally, the fixation of nitrogen by lupin increased distinctly the seeds yield in comparison to other nitrogen form used. Similarly, in the case of N₂ forms, the content of protein in seeds and the mean yield of protein was at the level of 32%. The content of protein in seed of treatments: -NH₂, Nd and NH₄++NO₃ was on the level of 30%. The seeds collected from the treatments NH₄+or NO₃- contained lower amount of protein ca. 27%. The yield of protein was also highest for both treatments with symbiotic nitrogen (11.12 and 12.30 g). The relatively lowest yield of protein (0.824 g) for Nd treatment was observed.

Changes of protein fractions

The results of fractional composition of starting material (SM) – the seeds used for sowing and the seeds collected from various nitrogen treatments are presented in Table 2. In contrast to SM where main proteins were globulins (20.16 mg N/g), the main protein fraction of seeds from all treatments were albumins, from 17.71 to 27.16 mg N/g for

(NH₄⁺) and (NO₃⁻) treatments, respectively, and the differences between particular treatments were statistically significant, except the Nd treatment where albumin content was the same as in SM. Among all of the treatments, the highest level of globulin was found for N₂ treatment (15.26) mg N/g), while the lowest one for nitrogen deficiency (Nd) (6.86 mg N/g). The closest level of Gt+P in relation to SM shows N₂ treatment, although statistically significant difference between them was observed. Addition of mineral nitrogen form to the N₂ treatment caused significant decrease of the Gt+P content from 8.40 to 4.48 mg N/g. The lowest level of these proteins for Nd was noted. All non-protein fractions (Nnp), with exception of $[N_2+(NH_4^++NO_3^-)]$ treatment, differ statistically from SM and between particular nitrogen forms. The residue fraction (Nr), with exception for $[N_2+(NH_4^++NO_3^-)]$ treatment, was about 2.2 mg N/g. The lowest statistical difference was observed in these fractions. The comparison of total mg N in protein fractions with total mg N in seeds shows that the used method is very accurate, which confirms high level of recovery from 97.84 to 102.09% (Table 2).

Figure 2 presents elecrophorograms of particular protein fractions of lupin seeds collected from different nitrogen treatments where the particular fractions differ among themselves with protein composition. Also, within the framework of fraction, the effect of nitrogen form is clear. In the case of albumin, the decay of protein ca.76.5 kDa, characteristic for SM (line 8), in case of the treatments N₂ (line 2), N₂+both mineral nitrogen form (line 3), NO₃⁻ (line 4) and -NH₂ (line 7) is observed. Significant difference occurs between proteins with molecular weight ca. 61.56; 49.0; 47.5 and 25.9 kDa. The globulin fraction shows totally different protein composition in relation to albumin fraction. Additionally, in this case the nitrogen form has influence on intensity of some protein bands. It especially concerns the proteins 60.5 and 30.5 kDa. For glutelin and prolamin, high non-migrating molecular protein is observed in all cases of nitrogen forms. Some differences in intensity of bands are showed for line 7, particularly for 60.5 kDa protein. This protein occurs in the highest content in comparison to other nitrogen form. Moreover, some differences between nitrogen forms are visible for proteins with molecular weight of 40 kDa.

DISCUSSION

Storage protein form is an important group of plant proteins which provide nitrogen and amino acids for developing seedlings and constitute a major source for human and livestock consumption. In this regard, main economic im-

TABLE 1. Yield of seeds, content of protein in seeds and yield of protein from proper treatment.

	Nitrogen forms/Treatments									
_	T1*	T2	Т3	T4	Т5	Т6	Т7			
Yield of seeds from treatment (g)	2.81	34.32	38.08	15.66	6.86	11.27	8.96			
Percentage content of protein in seeds	29.34	32.40	32.29	27.52	27.00	29.03	30.40			
Yield of protein from treatment (g)	0.82	11.12	12.30	4.30	1.85	3.27	2.72			

^{*} T1 - Nd; T2 -N₂; T3 - N₂+NH₄++NO₃; T4 - NO₃; T5 - NH₄+; T6 - NH₄++NO₃; T7 - NH₂

The content of protein in the seeds of start material (SM) was 35.55%

TABLE 2. Proteins composition of starting material (SM) and the lupin seeds collected from different treatments (mg N/g of seeds).

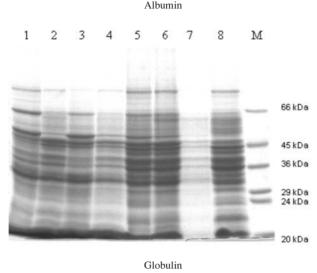
Nitrogen form	Fractions					Total N	Total N in seeds	Recovery
	Nnp	A	G	Gt+P	Nr	in protein fr. (mg/g of seeds)	(mg/g)	(%)
SM	7.21±0.17* a**	18.27±0.1 a	20.16±0.4 a	7.84±0.10 a	2.17±0.31 ae	55.65	56.88	97.84
Nd	4.90±0.10 b	18.62±0.5	6.86±0.27 b	1.82±0.21 b	2.10±0.30 a	34.30	34.86	98.39
$\overline{N_2}$	6.76±0.14 c	19.6±0.13 b	15.26±0.9 c	8.40±0.51 c	1.82±0.01 b	51.84	52.50	98.74
N ₂ +NH ₄ ++NO ₃ ⁻	7.21±0.05 a	23.1±0.92 c	12.04±0.6 d	4.48±0.09 d	4.83±1.19 c	51.66	51.38	100.54
NO ₃ ⁻	6.23±0.09 d	27.16±0.6 d	5.81±0.14 e	2.45±0.27 e	2.38±0.45 ae	44.03	43.13	102.09
NH ₄ ⁺	5.67±0.05 e	17.71±0.0 e	13.30±0.42 f	3.71±0.13 f	2.80±0.06 d	43.19	42.42	101.81
NH ₄ ⁺ +NO ₃ ⁻	4.90±0.21 b	16.73±0.15 f	11.55±0.24 d	4.28±0.53 d	2.59±0.40 ed	40.05	41.58	96.32
-NH ₂	9.92±0.14 f	26.10±0.3	14.56±0.1 c	3.00±0.37 g	1.40±0.34 f	54.96	54.10	101.59

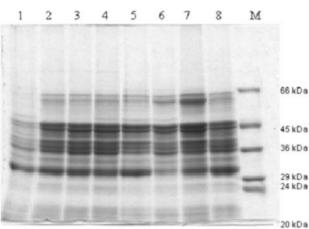
^{*} standard deviation (SD); ** letters a-g indicate statistical significance at p<0.05

Nnp – non-protein fraction; A – albumin; G – globulin, Gt+P – glutelin+prolamin; Nr – residual fraction

portance is attributed to plants of the Graminae, particularly Fabaceae family. Till now, the best studied and utilized legume has been soybean. The great success of soybean as a source of protein initiated intensive investigations on other legumes and particularly lupin that, except high protein content, belongs to unique ecological plants (Gulewicz et al. 1994). Our present study concentrates on determination of how various nitrogen forms used as fertilizer affect composition of seed proteins of L. albus. This question may be especially important for the nutritive value of seed proteins. This statement was confirmed by the results presented in Table 1, where clear effect of nitrogen form on the yield of seeds, content of protein and yield of protein in seeds were observed. These results are in agreement with our earlier studies (Ciesiołka et al. 2005) as well. The highest protein content in seeds was found in the treatments where nitrogen in the forms of N_2 , $[N_2+(NH_4^++NO_3^-)]$ and -NH₂ was used (Table 1). The other nitrogen forms brought much poorer results. It is worth to emphasize that only in the case N_2 and $[N_2+(NH_4^++NO_3^-)]$ treatments, high level of seeds protein content corresponds to high yield of seed protein. In other treatments, the seeds protein content did not correlate with the yield of protein. It shows that at determined conditions of the experiment only fixation of atmospheric nitrogen by lupin may create the best conditions for protein accumulation in seeds. According to Blagrove and Gillespie (1975), Duranti and Cerletti (1983), the major proteins of lupin seeds are a group of globulins called conglutins. The globulin fraction contains two major proteins: conglutin α (11S) and β (7S) (acidic) and a lupin-specific proteins called conglutin γ (7S) (basic) and conglutin δ 2S (Blagrove and Gillespie 1975; Restani et al. 1981; Cerletti et al. 1988; Lambert and Yarwood 1992; Petterson 1998 Sepulveda et al. 1999). In correspondence with this

finding are the results of start material only (Table 2). However, these seeds used in our experiment originated from field cultivation. When the same lupin seeds were cultivated on perlite in greenhouse conditions, the predominant fraction was albumins. It proves that both soil and environmental conditions may have effect on fractional composition of lupin proteins. However, taking into consideration that experimental conditions for all treatments were the same, we have been able to observe the effect of different nitrogen forms on chemical composition of protein fraction. This statement confirms the fact that in the (N_2) treatment, the closest to natural condition of nitrogen fixation, a similar difference exists between albumins and globulins in comparison to start material (SM). The statistical difference of protein level within the framework of the same fraction certifies significant effect of nitrogen form on fractional composition. Similarly, the level of storage protein fractions of all nitrogen forms, except N₂, is lower in comparison to SM and statistical significant difference between the treatments certifies distinct effect of nitrogen form on the formation of this fraction. It is worth to mention that in the case of the treatment where any nitrogen was used (Nd), the level of Gt+P was the lowest. Diversified level of nitrogen in non-protein fractions (Nnp), in particular the treatments, may certify that nitrogen forms affect the incorporation of amino acids to protein structure. As we know, the Nnp fraction is represented mainly by free amino acids and peptides, and total nitrogen of this fraction depends on physiological state of plant and seed. The increased level of this fraction, as in the case of -NH₂ form (Table 2), may be a result of not terminated protein synthesis or disintegration of protein for amino acids. Another explanation is that just –NH₂ form is influenced by nitrates accumulations as it took place in the case of spinach (Gulser 2005). The





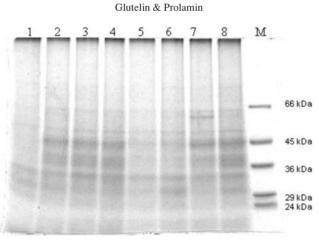


Fig. 2. Sodium dodecyl sulphate polyacrylamide gel electrophoresis (SDS_PEAG).

Lines: 1 - Nd; $2 - N_2$, $3 - N_2$ +both ions NH_4^+ and NO_3^- ; $4 - NO_3^-$; $5 - NH_4^+$; $6 - NH_4^+$ + NO_3^- ; $7 - -NH_2$; 8 - SM; M - marker

decrease of Nnp, as in case of for example Nd, $\mathrm{NH_4^+}$ and $\mathrm{NH_4^+}$ + $\mathrm{NO_3^-}$ forms proves additional incorporation of amino acids of these fractions to protein synthesis. From nutritional point of view, non-protein nitrogen (Nnp) may be of considerable importance (Hill 1977). Non-fractionated nitrogen (Nr) fractions, composed of nucleic acids compounds (purines, pyrimidines), poliamines and non-fractionated protein of wall cells in all treatments, except $\mathrm{N_2}$ and

 $-\mathrm{NH}_2$ forms, are closest to SM. It is worth to note that low level of (Nr) at N₂ and $-\mathrm{NH}_2$ treatments is connected with increase of level of Nnp fraction. It is especially visible in the $-\mathrm{NH}_2$ treatment. The distinct increase of level of GT+P fraction in the case of additional dose of nitrogen in the treatment [N₂+NH₄⁺+NO₃⁻)] may be caused by e.g. increase of polyamine content in the seeds (Aniszewski et al. 2001). It should be also emphasized that biological nitrogen (N₂) clearly increased Gt+P protein fraction, and additional nitrogen [N₂+NH₄⁺+NO₃⁻)] decreased it.

Elecrophorograms of protein fractions collected from different nitrogen treatments confirming that nitrogen forms have effect not only on quantitative change of protein fractions, but also on the quality one. The results presented in Figure 2 are in agreement with literature data of Blagrove and Gillespie (1975) who showed that conglutin α had been found to release three or four types of subunits with an apparent MW in the range 55 000-800 000 Da, each of which may contain a disulphide-bonded moiety with a MW near 20 000 Da and conglutin β yielded four subunits in the range 20 000-80 000 Da with no covalent links. Conglutin γ contains one subunit of MW 40 000 composed of two disulphide-linked peptides of MW 17 000 and 30 000 Da.

In the light of presented results, it is worth to give emphasis to the used protein fractionation method. There are two-way determination of nutritive value of seeds protein: first one is direct, based on nutritional experiment, and the second one is indirect, based on chemical analysis. Although the former gives most reliable results, it is an expensive and labor and time consuming method, and is not suitable for wide use. The chemical method of evaluating nutritive value of proteins is based on determination of general nitrogen that may be counted over on the so-called "crude protein", and determination of its fractional composition. Certainly, we are aware of the possibly existing critical viewpoints connected with the term "crude protein" (Tkachuk 1969; Cherry 1977; Ewart 1981; Boisen et al. 1987; Haque et al. 1989). However, the interpretation of such analysis is based on the assumption that structural protein, e.g. albumins, as a rule has richer amino acid composition and higher biological value than storage protein. So, increase of albumin, Nnp (containing free amino acids, peptide) and decrease of residue fraction Nr, using different nitrogen forms, should be beneficial from nutrition point of view. We observed the case when the treatment -NH₂ was used (Table 2). In this case, we observed increase of albunin fraction from 18.27 to 26.10 mg N/g and Nnp fraction from 7.21 to 9.92 mg N/g and decrease of Nr fraction from 2.17 to 1.40 mg N/g. Unfortunately, using nitrogen in -NH₂ form (urea), similarly to other mineral forms, affects principal yield of seeds, content of protein in the seeds, and yield protein (Table 1).

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