

Long-Term Study of Weather Effects on Soybean Seed Composition

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ABSTRACT: A long-term study initiated in 1989 at Sanborn Field, Columbia, Missouri, was designed to evaluate the affect of environmental factors, nitrogen application, and crop rotation on soybean (*Glycine max* [L.] Merr.) seed composition. Soybeans were grown as part of a four-year rotation which included corn (*Zea mays* L.), wheat (*Triticum aestivum* L.), and red clover (*Trifolium pratense* L.). Results from soil tests made prior to initiation of the study and subsequently every five years, were used to calculate application rates of nitrogen, phosphorus, and potassium necessary for target yield of pursuant crops. In the experimental design, nitrogen was applied to one-half of the plot on which the non-leguminous crop, either corn or wheat was grown. Analysis of soybean seed by near infrared reflectance spectroscopy collected over an 11-year period revealed a linear increase in protein and decrease in oil content. Application of nitrogen fertilizer to non-leguminous crops did not have an apparent effect on total protein or oil content of subsequent soybean crop. Analysis of soybean seed proteins by sodium dodecyl sulfate polyacrylamide gel electrophoresis in conjunction with computer-assisted densitometry revealed subtle changes in the accumulation of seed proteins. Immunoblot analysis using antibodies raised against the β -subunit of β -conglycinin showed a gradual increase in the accumulation of the 7S components during successive years of the experiment. A linear increase in temperature and decrease in rainfall was observed from the onset of data collection. Higher temperatures during the growing season have been linked to increased protein and diminished oil content of soybean, thus changes observed in this study are possibly related to climatic conditions. However, crop rotation and subsequent changes in soil ecology may contribute to these observed changes in the seed composition.

Keywords: crop rotation, seed composition, soybeans

Production economics are factors in the selection of agronomic practices (Katsvairo & Cox, 2000). Although rotation of cereals, such as wheat and corn with legumes is a long-standing cultural method (Campbell *et al.*, 1991; Vanotti & Bundy, 1995), the advent of mechanization has led to pre-

dominance of monoculture systems in today's agricultural production. Research devoted to improving soil and environmental quality while maintaining the cost of production, and enhancing grain yield and quality involves long-term experiments. Increased awareness and knowledge of human and animal nutrition compel plant scientists to develop cultivars whose composition provides optimal nutrition. Selection and breeding, and more recently genetic engineering, are tools being utilized to this end.

Soil ecology is a factor in the mineralization and availability of nutrients to the soybean (Dodor & Tabatabai, 2003) which in turn influences content of seed storage compounds that are important in human and animal nutrition (Sexton *et al.*, 1998). Soil organic carbon (SOC) is one indicator of soil quality and has an influence on the chemistry and biology of soil (McCallister & Chein, 2000; Kristensen *et al.*, 2003; Sparling *et al.*, 2003; Shukla *et al.*, 2004). Since crop rotation influences this parameter (Reeves, 1997; West & Post, 2002), potential exists for a crop rotation system to influence nutrient quality of crops. Higher microbial carbon and nitrogen levels were reported to occur in a four-crop rotation system, which included a forage legume as opposed to a monoculture system (Anderson & Domsch, 1989; Anderson & Gray, 1990).

Enzymatic activity necessary for mineralization of plant nutrients has been found correlated to SOC (Jordan *et al.*, 1995). Mineralized sulfur and nitrogen are essential for the production of amino acids, which are components of seed storage proteins. Availability of both sulfur and nitrogen has been associated with relative accumulation of seed protein and thus quality (Kim *et al.*, 1999; Krishnan *et al.*, 2000; Nakasathien *et al.*, 2000; Paek *et al.*, 2000). Protein and oil content of soybeans are also affected by symbiotic associations with vesicular mycorrhiza and symbiotic bacteria. Protein content was increased in plants that formed symbiotic relationship with arbuscular mycorrhiza (Bethlenfalvay *et al.*, 1997). Rhizobial nodulation factors have been found to enhance the colonization of soybean roots by mycorrhizobia (Xie *et al.*, 1995). Tripartate symbiosis involving plant mycorrhiza and bacteria benefits from an increased soil organic carbon (Xie *et al.*, 1995).

Soybean seed composition is influenced by environmental

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conditions during the growing season (Piper & Boote, 1999; Grieshop & Fahey, 2001; Yaklich *et al.*, 2002; Fehr *et al.*, 2003; Grieshop *et al.*, 2003). In a general sense, soybeans growing at higher temperatures contain more protein while those from cooler conditions contain higher percentage of oil (Wolf *et al.*, 1982; Yaklich *et al.*, 2002). Soybeans of maturity groups II through IV exhibited the highest oil content while those of maturity group VIII were highest in protein (Yaklich *et al.*, 2002). The protein and oil content data were compiled from a multitude of soybean lines submitted for evaluation in uniform tests. Recent work has shown that conditions existing at specific times during the growing season can be correlated with the final protein and oil content of soybeans (Yaklich & Vinyard, 2004). Since variety and growing conditions have been reported to affect seed composition (Piper & Boote, 1999; Grieshop & Fahey, 2001; Yaklich *et al.*, 2002; Fehr *et al.*, 2003), utilizing a single variety at one location in a long-term study will provide information as to the specific effects of environment on seed components. In this study, soybean 'Williams 82' was utilized and the cumulative effects of nitrogen application, environmental conditions, and crop rotation on protein and oil content were investigated.

MATERIALS AND METHODS

Field plot fertility and design

Williams 82 soybeans were grown beginning in 1989 on Sanborn field in Columbia, MO, in a four-year rotation consisting of wheat, red clover, corn, and soybeans. Data for this experiment were collected from 1991 to 2002, exclusive of 1999, for which samples were not available. A split-plot design was selected to ascertain, in part, the effect of nitrogen carryover on seed storage compounds in subsequent soybean crops and the cumulative effect of crop rotation on content of seed storage compounds. Each 30.8 m × 9.4 m plot in the study was surrounded by a grass-seeded alley. Soil tests were conducted prior to the onset of the trial and every five years subsequently. Annual fertilizer application was based upon yield goals of 8.15 t ha⁻¹ for corn, 4.7 t ha⁻¹ for wheat, 4.0 t ha⁻¹ for soybeans, and 9.4 t ha⁻¹ for red clover hay (Buchholz, 1983). Calculated amount of nitrogen was applied to one-half of the plot to be planted in either corn or wheat. Red clover forage was harvested in the fall and re-growth was moldboard-plowed in spring as a green manure crop prior to corn planting.

Near infrared reflectance spectrographic (NIRS) analysis of seed protein and oil

Protein and oil content of seed aliquots was determined by

NIRS using the Infratech 1255 Food and Feed analyzer (Tecator AB, Hoganas, Sweden). The reported protein and oil was based upon normalized moisture content of 135 g kg⁻¹ in each sample. Data from NIRS analysis for protein was corroborated by University of Missouri Experimental Station Laboratory using the nitrogen combustion method, 990.03, as outlined in the 17th edition of Official Methods of Analysis of the International Association of Analytical Communities (AOAC).

Sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) separation of seed protein.

An aliquot of 10 seeds was selected from each year's soybean harvest and ground with mortar and pestle. A 15-mg aliquot of the powdered seed was extracted in 1 mL of a solution comprised of 4% sodium dodecyl sulfate (w/v), 20% glycerol (v/v), and 0.03 mM bromophenol blue solubilized in 125 mM Tris-HCl buffer (pH 6.8). Cell debris was sedimented by centrifugation at 14300 × *g* for 15 min, the supernatant decanted to clean centrifuge tube, and combined with 50 μL 2-mercaptoethanol. Proteins were fractionated by SDS-PAGE (Laemmli, 1970) on a 12.5% resolving gel (w/v) at 20 mA for 1 hr using a Hoefer SE260 minigel apparatus (Amersham Biosciences, Piscataway, NJ). The gels were stained overnight with Coomassie blue.

Densitometry

Proteins were separated on a 12.5% polyacrylamide gel as described in the previous section and the relative densities of the bands were determined by computer-assisted densitometry employing the Gene Wizard System (Syngene, Beacon House Nuffield Road, Cambridge, UK). The ratio of the acidic and basic subunits of glycinin with respect to α', α, and β-subunits of β-conglycinin was determined.

Western blot analysis

Total protein extract was resolved by SDS-PAGE on a 10% gel (w/v) at 20 mA for 1 hr using a Hoefer SE260 minigel apparatus. After 10 min equilibration in transfer buffer (0.02 M Tris, 0.15 M glycine, 200 mL methanol), proteins were electrophoretically transferred to a nitrocellulose membrane, incubated for 1 hr in 1× Tris buffered saline (TBS) (0.02 M Tris, 0.5 M NaCl) which contained 5% non-fat-dehydrated milk to block non-specific binding. Subsequently, the membrane was incubated overnight in 1× TBS solution containing 5% non-fat-dehydrated milk (w/v) to which the primary antibody had been added in a final 1 to 5000 dilution. After three 10 min washes in 1× TBS contain-

ing 0.05% Tween 20 (TBST) the membrane was exposed to a secondary antibody, goat anti-rabbit-HRP conjugate (Pierce Chemical Co., Rockford, IL), for 1 hr and then washed four times in 1× TBST. Relative amounts of proteins present were determined using the chemiluminescent, Supersignal®, HRP substrate (Pierce Chemical Co., Rockford, IL).

Fatty Acid Methyl Ester (FAME) analysis of oil components

An aliquot of seeds was selected from each year of soybean production. The seed coat was cracked and seed was extracted overnight in 1 mL hexane/chloroform/methanol (8:5:2 v/v/v). Fatty acids from a 150 aliquot of the extract were methylated with 75 μ L of sodium methoxide-methanol/methanol/petroleum ether/ethyl ether solution (1:4:2 v/v/v). Methyl ester derivatives of the fatty acids were separated on a 30 mm \times 0.53 mm \times 0.5 μ m AT-Silar capillary column (Alltech, Deerfield, IL) in the Agilent 6890 gas chromatograph (Agilent, Palo Alto, CA). The methylated fatty acids were detected in the effluent stream by flame ionization. A standard containing the methyl ester derivatives of the five fatty acids in soybean, palmitic, stearic, oleic, linoleic, and linolenic acids was used for determining relative amounts of each fatty acid.

Statistical analysis

The effect of years, rainfall, temperature, and nitrogen application on soybean seed protein, protein components, oil, and fatty acids was assessed by analysis of variance (ANOVA). Years, rainfall, and temperature were treated as random effects while nitrogen application was treated as a fixed effect. Relationships between aforementioned seed traits and environmental conditions and between seed traits and nitrogen application were evaluated using regression analysis. Statistical analyses were determined using the 8.2 version of General Linear Model SAS software.

RESULTS

Near infrared reflectance spectroscopy analysis of protein and oil

Accumulation of seed storage components in soybean is influenced by environmental conditions prevailing during the growing season. To monitor the changes in seed composition during the course of this study, we determined the protein and oil content of seeds by NIRS analysis. This analysis revealed that the total protein increased from 35 to 37.5% of

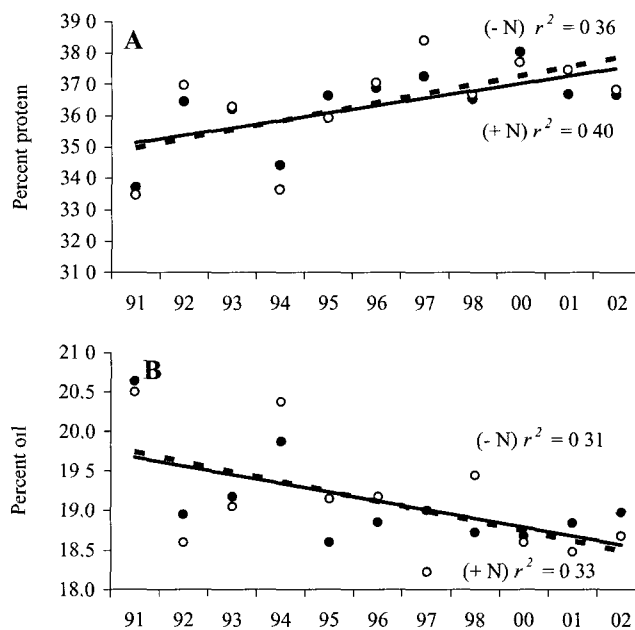


Fig. 1. Change in protein and oil content during an 11-year study. Near infrared reflectance spectroscopy analysis revealed changes in protein (1A) and oil (1B) content. Comparison of seed components from nitrogen fertilized (●) and non-fertilized (○) plots shows the effect of years (1A and 1B). Solid regression lines represent data acquired from plots receiving nitrogen, while broken lines correspond to non-fertilized plots.

seed weight, while that of oil declined from 19.5 to 18.5% over the 11-year period (Figs. 1A and 1B). We also examined the residual effect of exogenous nitrogen provided to non-leguminous crops in the rotation on soybean seed composition. Application of nitrogen fertilizer did not have an apparent effect on accumulation of these seed storage compounds (Figs. 1A and 1B). During the course of this experiment, an increase in protein ($P \leq 0.05$) was noted in seed from plants that did not receive supplemental nitrogen. Seeds from nitrogen fertilized plots did not exhibit a significant response (Figs. 1A and 1B). The year effect putatively reflects a combination of environmental factors and changes soil ecology. Since protein and oil content are inversely correlated, the significant increase in protein should be accompanied by a decrease in oil. Oil content decreased from 19.5% of seed weight to 18.5% over the course of the study. Application of exogenous nitrogen had no significant effect on the seed oil content.

Sodium dodecyl sulfate polyacrylamide gel electrophoresis and immunoblot analysis of seed storage proteins

SDS-PAGE fractionation of proteins supported the observation that an increase in protein content had occurred dur-

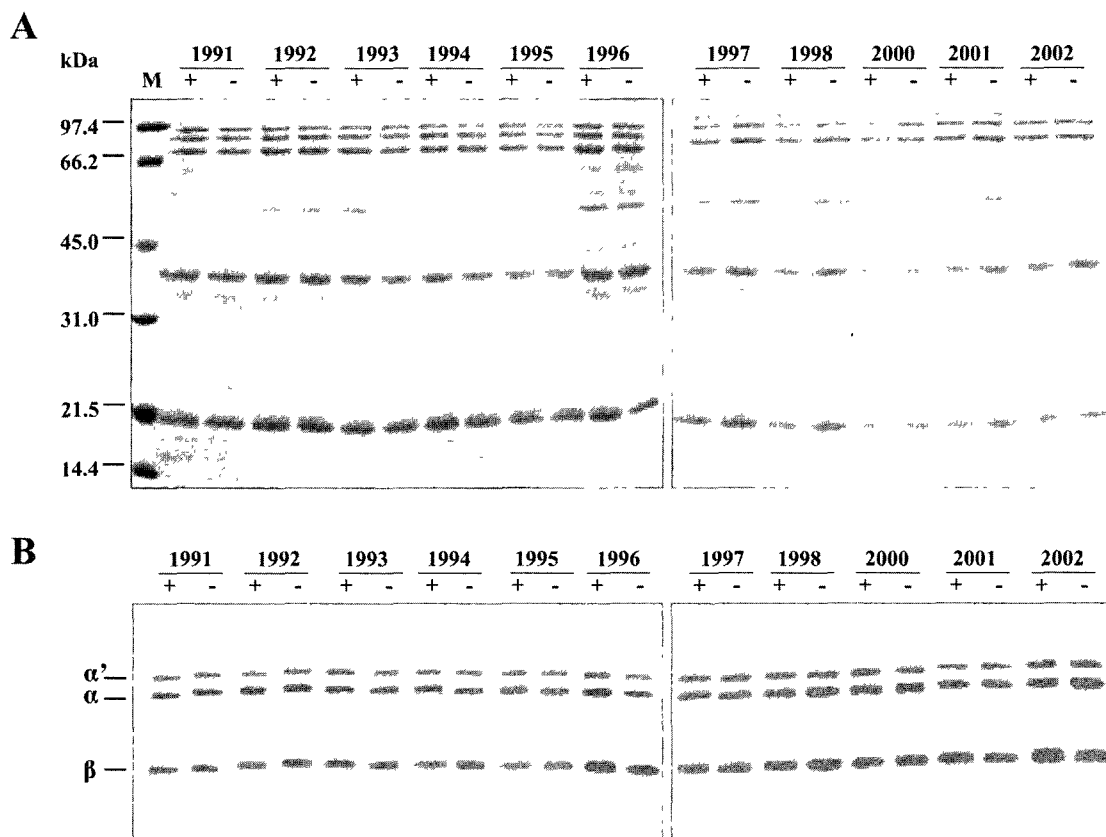


Fig. 2. Sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) and immunoblot analysis of seed storage protein. Seed proteins were fractionated on 12.5% polyacrylamide gel and stained with Commassie blue (1A). Proteins were electrophoretically transferred from a 10% gel to a nitrocellulose membrane for immunoblot analysis (1B). Immunoblot analysis of seed protein shows that the accumulation of the β -subunit of β -conglycinin was notably higher in seeds from nitrogen-fertilized plots. Conversely, application of nitrogen fertilizer was noted to depress the accumulation of the glycinins in several years (Fig. 1B).

ing the course of the experiment (Fig. 2A). The α' , α , and β subunits of β -conglycinin and the A4 subunit of glycinin appeared to increase over the course of the trial in both fertilized and non-fertilized plots (Fig. 2A). Immunoblot analysis was used to assay this putative increase in protein content (Fig. 2B). The use of polyclonal antibodies, raised against the β -subunit of β -conglycinin, which cross-react with the α' and α subunits, confirmed an increase in accumulation of these proteins. Information provided by the western blot analysis also indicated that application of nitrogen fertilizer did not have a consistent effect on accumulation of β -conglycinin in particular the β -subunit, which has been shown previously to respond to nitrogen application (Paek *et al.*, 2000; Ohtake *et al.*, 2002). Application of exogenous nitrogen increased the content of this subunit during only three years of the study 1996, 2001, and 2002 (Fig. 2B). Conversely, plants grown without exogenous nitrogen appeared to accumulate more glycinins than did those from fertilized plots. This was particularly evident in 1992, 1997, 1998, 2001, and 2002 (Fig. 2B). A densitometry analysis revealed

that although total protein increased during the course of the trial, the glycinin /conglycinin ratio did not vary appreciably (data not shown).

Gas chromatographic analysis of fatty acid components of seed oil

Fatty acid composition is integral to soybean oil utilization. Resistance to oxidation is enhanced by increasing the oleic acid content, thus making the oil amendable for cooking and storage. Alternatively, oil containing more of the polyunsaturated linolenic acid is purportedly beneficial to human health. The decrease in oil content of the soybeans observed in this study was not at the expense of one particular fatty acid. Analysis of variance of the fatty acid components, palmitic, stearic, oleic, linoleic, and linolenic acids, indicated a lack of correlation between year and accumulation of these long-chain compounds (Fig. 3). Application of nitrogen fertilizer did not alter the total fatty acid content or profile.

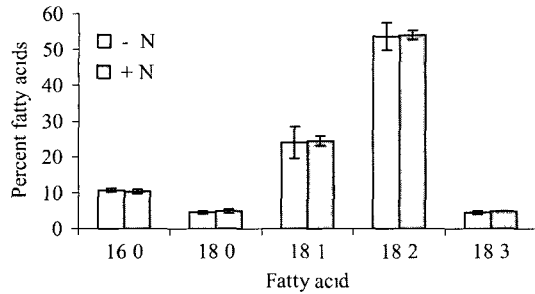


Fig. 3. Gas chromatographic determination of fatty acid content. Fatty acids were solvent extracted and methylated in preparation for separation and quantification by gas chromatography. Error bars represent standard deviation of the mean of three replications of analysis.

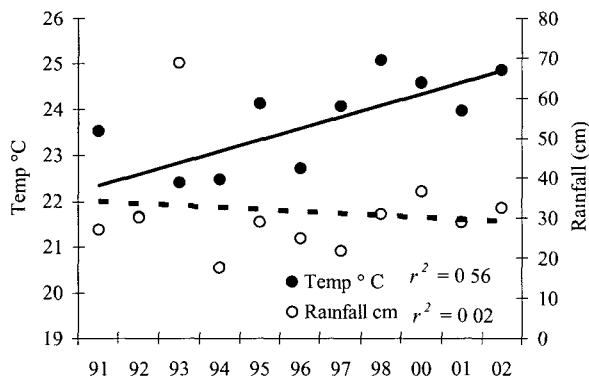


Fig. 4. Environmental conditions during an 11-year study. Average daily temperature and rainfall were recorded over the span of the experiment. Temperature (●) during the span of the trial increased an average 0.25°C per year ($P \leq 0.05$).

Environmental effects on protein and oil

Moisture availability and temperature are crucial parameters in soybean reproduction and ostensibly, are involved in the quality of seed produced. Temperature and rainfall data during the reproductive period of July through September were recorded and the effect on accumulation of seed storage compounds determined. The maximum mean temperature occurring during pod-fill increased 0.25°C year⁻¹ ($P \leq 0.05$) (Fig. 4). Additionally, a positive correlation was noted between year and recorded temperature. Although differences in rainfall during pod fill was not significant over the time span of the experiment, a negative correlation did exist between rainfall and years (Fig. 4). The effect of rainfall and average maximum temperature during pod fill period was examined by analysis of variance, treating these environmental conditions as random factors. Neither parameter was found to influence the accumulation of seed storage compounds. During the 1993 soybean reproductive period, rainfall exceeded 68 cm, which was well above the 30-year

average rainfall of 28 cm for Boone County, Missouri for this time period (<http://agebb.missouri.edu/mass/MO30avg.pdf>). Accumulation of seed storage compounds did not appear to be affected in that year by this inordinate rainfall, but in the subsequent year, protein and oil content showed a dramatic decrease and increase, respectively (Fig. 1).

DISCUSSION

During the course of this field experiment, the total protein component of soybean seed increased from 35 to 37.5% of total seed weight, while oil diminished from 19.5 to 18.5%, regardless of nitrogen treatment. A warming trend during the reproductive stage of plant development was evident from 1991 to 2002, while rainfall diminished slightly. A positive correlation exists between protein accumulation and years. The ratio of protein gain and oil loss observed during this study is worthy of note. An inverse relationship has been shown in the accumulation of these two storage compounds because they compete for carbon and energy necessary for synthesis (Hanson *et al.*, 1961; Simpson & Wilcox, 1983). Since oil contains twice the Kcal g⁻¹ energy as protein, it has been suggested that a change in accumulation of one unit of oil would be accompanied by a two-unit change in protein (Hanson *et al.*, 1961; Chung *et al.*, 2003). This approximate exchange was observed because oil decreased over 1% and protein increased approximately 2.5% over the 11-year period. Assigning specific causes to the increase in protein and decrease in oil is tenuous because both changing environmental conditions and the incipient crop rotation system could be involved. Higher ambient temperature has been associated with increased soybean protein and diminished oil accumulation in both growth-chamber (Wolf *et al.*, 1982) and field studies (Piper & Boote, 1999; Yaklich *et al.*, 2002). In this trial, however, changes in protein and oil content could not be entirely assigned to environmental conditions, which allude to other possible causes for the differences observed in the accumulation of these storage compounds.

Cropping systems such as the rotation protocol selected for this trial are known to affect soil ecology (Lavelle, 2000) which influences plant nutrient availability (Moore *et al.*, 2000; Dodor & Tabatabai, 2003). Whether the chemical, biological, and physical properties of soil (Hsieh, 1992; Haynes & Tregurtha, 1999; Hickman, 2002; Eivazi *et al.*, 2003) have been modified sufficiently as to influence the assimilation of seed storage protein and oil is not certain. Separation of environmental factors from agronomic effects, using statistical methods capable of segregating the influence of these parameters may facilitate determining the specific and underlying reasons for the changes in seed protein and oil content.

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