

Effect of Row Spacing on Seed Isoflavone Contents in Soybean [*Glycine max* (L.) Merr.]

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Received 1 October 2014; revised 3 November 2014; accepted 9 December 2014

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Abstract

Soybean isoflavones compounds such as genistein, daidzein, and glycitein have numerous human health benefits including the reduction of risks of cardiovascular diseases, breast and prostate cancers, and menopause symptoms in women. Understanding the genetic and environmental control of isoflavones accumulation is of great importance for developing new cultivars with high amounts of seed isoflavones. This study was conducted to analyze the effect of row spacing (25 cm vs. 50 cm) on seed isoflavones accumulation using a recombinant inbred line (RIL) population derived from the cross of PI 438489B and "Hamilton" (PIxH, n = 50). The two row spaces generated plant densities of 250,000 plants/ha and 90,000 plants/ha, respectively. Significant differences in soybean seed isoflavones (daidzein, genistein and glycitein) contents have been observed between plants grown in the two different plant densities. The mean daidzein content was $0.03458 \mu\text{g}\cdot\text{g}^{-1}$ in plants grown in 50 cm row spaces (low plant density), which was significantly higher than its content ($0.03019 \mu\text{g}\cdot\text{g}^{-1}$) in plants grown in 25 cm row spaces (high plant density). Similarly, the mean glycitein content in plants grown in 50 cm row spaces ($0.01905 \mu\text{g}\cdot\text{g}^{-1}$) was significantly higher than its content in plants grown in 25 cm row spaces ($0.00498 \mu\text{g}\cdot\text{g}^{-1}$). Also, the mean genistein content in plants grown in 50 cm row spaces ($0.01466 \mu\text{g}\cdot\text{g}^{-1}$) was higher than its content in plants grown in 25 cm row spaces ($0.00831 \mu\text{g}\cdot\text{g}^{-1}$). These preliminary results are important in guiding farmers and breeders on choosing the best row spaces to grow soybean plants in order to optimize isoflavones contents. Further studies are needed to understand the correlation between seed

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isoflavones contents and other agronomic traits such as seed yield, protein, and oil contents.

Keywords

Daidzein, Genistein, Glycitein, Row Space, Plant Density

1. Introduction

Soybeans [*Glycine max* (L.) Merr.] are one of the major cash crops in the United States with a wide variety of benefits for human health. Due to cultural practices and environmental stimuli, farmers are constantly faced with the rise of production costs and the decline on crop yields in many regions. To increase yield, farmers, breeders, and scientists tried tillage system, herbicide technology, and row spacing in several crops including soybean [1]-[7].

Several studies reported that higher yields are obtained in plants grown in low row spaces compared to plants grown in high row spaces in rice [8], cotton [3] [5], Cuphea (*Cuphea viscosissima* Jacq.) [9], sorghum [10], and soybean [2] [6]. In rice, higher biomasses were observed in plants grown under low row spaces 18, 21, and 24 cm compared to high row spaces 27, 30, and 33 cm [8]. Cotton plants grown in 38 cm row spaces had higher yield compared to plants grown in 102 cm row spaces [5]. In another study, cuphea plants were grown in 38, 56, and 74 cm row spaces and data showed that plants grown in low row spaces show higher biomass and seed yield [9]. In sorghum, two cultivars in 2008 and 5 cultivars in 2009 have been grown in 10 - 15, 37.5, and 74 cm row spaces. Data showed no changes in biomass yield (Dry Matter, DM t·ha⁻¹); however, cultivar Goliah showed higher DM yield than all the other tested cultivars. Moreover, higher sugar contents have been found in cultivar Aron grown in the lowest row space, 37.5 cm [10]. In soybean, a study found that plants grown in 45 cm row spaces had higher yields (3997 kg·ha⁻¹) compared to those grown in 60 cm (3130 kg·ha⁻¹) and 90 cm row spaces (2490 kg·ha⁻¹) [2]. Another study found that soybean plants grown in 25 cm row spaces have higher yield (2470 kg·ha⁻¹) compared to plants grown in 100 cm row spaces (2108 kg·ha⁻¹) [1]. In this population, we found that plants grown in 25 cm row spaces had higher 100-seed weight, total seed weight, pod, and seed numbers compared to plants grown in 50 cm [11].

The fact that soybean seed isoflavone contents are genotype and environment dependent is well established [12]-[16]; however, very few studies investigated the effect of row spacing on seed composition including seed isoflavone contents [6] [17]. For example, Rahman *et al.* [6] found that a decrease in row spacing promoted an increase in seed protein contents in two soybean cultivars PB-1 and G-2 while the contents of several minerals decreased with a decrease in row spacing [6]. Al-Tawaha and Seguin [17] studied the effects of row spacing, planting date, and weed on seed isoflavone contents and other seed components. They found that row spacing has no effect on seed isoflavone contents; however, these contents increased by 9% in the presence of weeds [17]. Negative correlations have been found between yield crude protein (CP) and seed yield, and seed isoflavone contents [17]. Isoflavones overall contents are greater in early soybean planting system (ESPS, April) compared to conventional soybean planting system (CSPS, June) [17]. Further studies are needed to investigate the effects of row spacing on seed isoflavone contents as well as the correlations between seed yield and isoflavone contents.

This study was designed to investigate the effect of row spacing on seed isoflavone contents by growing the PI 438489B by “Hamilton” recombinant inbred line (RIL) population in early soybean planting system (ESPS) in two different row spaces, 25 cm and 50 cm which generate 250,000 and 90,000 plants/ha, respectively.

2. Material and Methods

2.1. Plant Material

In this study, soybean line PI 438489B, cultivar “Hamilton”, and their recombinant inbred lines (RILs) (n = 50) were used. The PI 438489B by “Hamilton” recombinant inbred line population [18] was provided to us by Dr. Khalid Meksem of Southern Illinois University. Several soybean RIL populations have been used to genetically map quantitative trait loci (QTL) in soybean and we used this population for both QTL mapping and row spacing effects on isoflavones contents.

2.2. Growing Conditions

In the greenhouse, 4 seeds of parents, PI 438489B and Hamilton, and each RIL were sown in pots (15×14 cm) filled with potting soil at $25^\circ\text{C} \pm 1^\circ\text{C}$ temperature and under natural daylight. After 3 weeks, the plants were transferred into a sandy field in St. Pauls, NC ($34^\circ 48' 26''\text{N}$ $78^\circ 58' 22''\text{W}$ and 170 feet above sea level; Robeson County). Two groups of plants were grown in the greenhouse and transferred randomly into the same field; Group I was planted with 25 cm row spaces between plants and the Group II was planted with 50 cm row spaces between plants. The plants were watered and kept in the field until maturity of all RILs and parents. No pesticide or herbicide was applied in the greenhouse or field.

2.3. Trait Measurements and Statistical Data Analysis

Days to germination (DG) were recorded in the greenhouse and days to flowering (DF) were recorded in the field. Plant height (PH) was recorded at maturity of parents and all RILs, just before harvest (120 days). Plants have been pulled from the ground gently with their roots and brought to the lab to measure pod numbers (PN), seed numbers (SN), 100-seed weight (g) (100SW), and total seed weight (g) (TSW). All these measurements were recorded for both Group I (25 cm row spacing) and Group II (50 cm row spacing). Concentration of seed isoflavones was analyzed using HPLC machine from Qiagen Inc. (Germantown, MD).

2.4. Isoflavones Extraction and Quantification

Three seed isoflavones daidzein, genistein, and glycinein were extracted and quantified in $\mu\text{g}\cdot\text{g}^{-1}$ of seed by a near-infrared reflectance (NIR) diode array feed analyzer (Perten, Springfield, IL, USA) as described earlier [6] [19]. Means, ranges, and standard deviations for seed genistein, glycinein, and daidzein contents were estimated for the parents and RILs based on three replicates per plant.

2.5. Data Analysis

Statistical data analysis was done using SPSS program and descriptive analysis, paired samples test, T-test, and Pearson correlations were performed to evaluate the mean seed isoflavone contents in the population and to see if there are significant differences in these isoflavones in the two groups (Group I and II).

3. Results

Means, standard deviations, ranges and co-efficient of variation for seed isoflavone contents in parents and RILs are shown in **Table 1** and **Table 2**. The results showed that seed daidzein, genistein, and glycinein contents of Group II plants are higher than those of Group I plants. The lowest coefficients of variation (CV) were found for daidzein in Group II plants (0.80%) and glycinein in Group II plants (0.87%) while the highest CV values are found for glycinein in Group I plants and for genistein in Group II plants (**Table 1**). The frequency distributions of seed daidzein, genistein, and glycinein contents in the RILs are not normal and showed skewness towards the low isoflavones contents for all the three isoflavones in the two groups (**Figure 1**).

The RILs showed 1.6 - 6 folds high contents than both parents for all the three isoflavones in plants of both Groups I and II (**Figure 2** and **Figure 3**). The RILs also showed high contents of daidzein then glycinein, then genistein, respectively in both row spaces (**Figure 2** and **Figure 3**). “Hamilton” plants grown in both 25 cm (Group I) and 50 cm (Group II) row spaces showed higher contents of glycinein and genistein (**Figure 2**) than PI 438489B plants grown in the same row spaces. However, “Hamilton” plants grown in 50 cm row spaces showed higher daidzein contents than PI 438489B plants grown in the same row spaces while the opposite is true for plants grown in 25 cm row spaces (**Figure 2** and **Figure 3**).

A paired-samples T-test was used to elucidate whether there was a statistically significant mean difference between the isoflavone concentrations of daidzein, genistein and glycinein in the RILs, PI 438489B, and “Hamilton” planted at 25 cm and 50 cm row spaces. **Table 1** shows that levels of isoflavone elicited a difference of $-0.00439 \mu\text{g}\cdot\text{g}^{-1}$ (95% CI, -0.0208 to 0.0120) when comparing daidzein at 25 cm to 50 cm row spacing ($t(27) = -0.548$, $p < 0.05$). When comparing glycinein at 25 cm and 50 cm, isoflavone levels displayed a difference of $-0.0141 \mu\text{g}\cdot\text{g}^{-1}$ (95% CI, -0.0207 to -0.0074) and a $t(27) = -4.351$, $p < 0.05$. Last, when comparing genistein at (25 cm and 50 cm) row spacing, levels showed a difference of $-0.00635 \mu\text{g}\cdot\text{g}^{-1}$ (95% CI, -0.0143 to 0.0016) with the $t(27) = -1.636$, $p < 0.005$, respectively.

Table 1. Description of mean, standard deviation, range, maximum and minimum of genistein, daidzein, and glycinein in two row spaces (25 cm and 50 cm) of PI 438489B by Hamilton and their recombinant inbred lines (RILs).

Paired differences									
	Mean	Std. deviation	Std. error mean	95% confidence interval of the difference		t	df	Sig. (2-tailed)	
				Lower	Upper				
Pair 1	Dai25 - Dai50	0.0044	0.0424	0.0080	-0.0208	0.0120	-0.548	27	0.588
Pair 2	Gly25 - Gly50	0.01407	0.0171	0.0032	-0.0207	-0.0074	-4.351	27	0.000
Pair 3	Gen25 - Gen50	0.0063	0.0205	0.0038	-0.0143	0.0016	-1.636	27	0.113

Pair 1, 2, and 3; Comparisons of daidzein, glycinein, and genistein contents in Group I and II plant seeds, respectively.

Table 2. Paired samples test. The Pearson correlation of daidzein (Dai), genistein (Gen), and glycinein (Gly) contents between Group I and Group II RILs. Seed isoflavone amounts are expressed in $\mu\text{g}\cdot\text{g}^{-1}$ of seed weight.

	Daidzein 25 cm	Daidzein 50 cm	Glycinein 25cm	Glycinein 50 cm	Genistein 25 cm	Genistein 50 cm
Range	0.1653	0.1033	0.0350	0.0663	0.0280	0.0826
Minimum	0.0007	0.0060	0.0010	0.0017	0.0010	0.0010
Maximum	0.1660	0.1093	0.0360	0.0680	0.0290	0.0836
Mean	0.0302	0.0345	0.0049	0.0190	0.0083	0.0146
Std. deviation	0.0381	0.0277	0.0071	0.0162	0.0072	0.0200
CV%	1.26	0.80	1.43	0.85	0.87	1.36

The (*t*) indicates comparison to the *t*-distribution, the (27) represents the degree of freedom which is (*n* – 1) and the numbers equal the obtained value or the value of the *t*-statistic. The (*p* < 0.05) indicates the probability of obtaining the given *t*-value. A paired sample statistic test was used to determine the effectiveness of planting at different row spacing. Results showed that isoflavone contents are higher in plants of Group II [Daidzein ($0.0346 \pm 0.005 \mu\text{g}\cdot\text{g}^{-1}$), genistein ($0.0147 \pm 0.004 \mu\text{g}\cdot\text{g}^{-1}$), and glycinein ($0.0191 \pm 0.003 \mu\text{g}\cdot\text{g}^{-1}$)] compared to plants of Group I [Daidzein ($0.0302 \pm 0.007 \mu\text{g}\cdot\text{g}^{-1}$), genistein ($0.0083 \pm 0.001 \mu\text{g}\cdot\text{g}^{-1}$) and glycinein ($0.0050 \pm 0.001 \mu\text{g}\cdot\text{g}^{-1}$)] (**Table 2** and **Table 3**; **Figure 2** and **Figure 3**). The Pearson correlation between the two plant groups showed that glycinein and genistein contents were significantly higher in Group II than Group I plants with values of 0.626 and 0.591, respectively (**Table 4**).

4. Discussion

This study was conducted to study the effect of row spacing (plant densities) on seed isoflavone accumulation using the recombinant inbred line (RIL) population derived from the cross of PI 438489B and Hamilton. Previous studies showed that high yields are obtained from plants grown in high plant densities (low row spaces) compared to plants grown in low plant densities (high row spaces) in rice [8], maize [20] [21], cotton [3] [5], Cuphea (*Cuphea viscosissima* Jacq.) [9], sorghum [10], and soybean [2] [6] [11] [22]. However, only one study reported the influence of row spacing on seed isoflavone contents [17]. In an earlier study, we used the “Flyer” by “Hartwig” RIL population ($F \times H$, *n* = 92) and measured days to germination (DG), plant height (PH), days to flowering (DF), 100-seed weight (SW), pod number (PN), seed number (SN), harvest index (HI), root dry weight (RDW), shoot fresh weight (SFW), root fresh weight (RFW), shoot dry weight (SDW), lateral root number (LRN), and maximum root length (MRL) in plants grown in early soybean planting system (ESPS, April) and conventional soybean planting system (CSPS, June) [22]. We found significant differences for all measured traits except RFW and LRN [22]. SW decreased by 74.7%, PH by 41.96%, PN by 65.2%, HI by 73.3%, and SN by 64.9% in plants grown in CSPS compared to plants grown in ESPS [22]. Interestingly, Al-Tawaha and Seguin [17] found that the overall isoflavone contents are greater in ESPS compared to CSPS [17].

Our results showed that there are highly significant differences for genistein and glycinein contents in plants of Group I compared to plants of Group II. These contents are higher in Group II (50 cm row spaces) compared

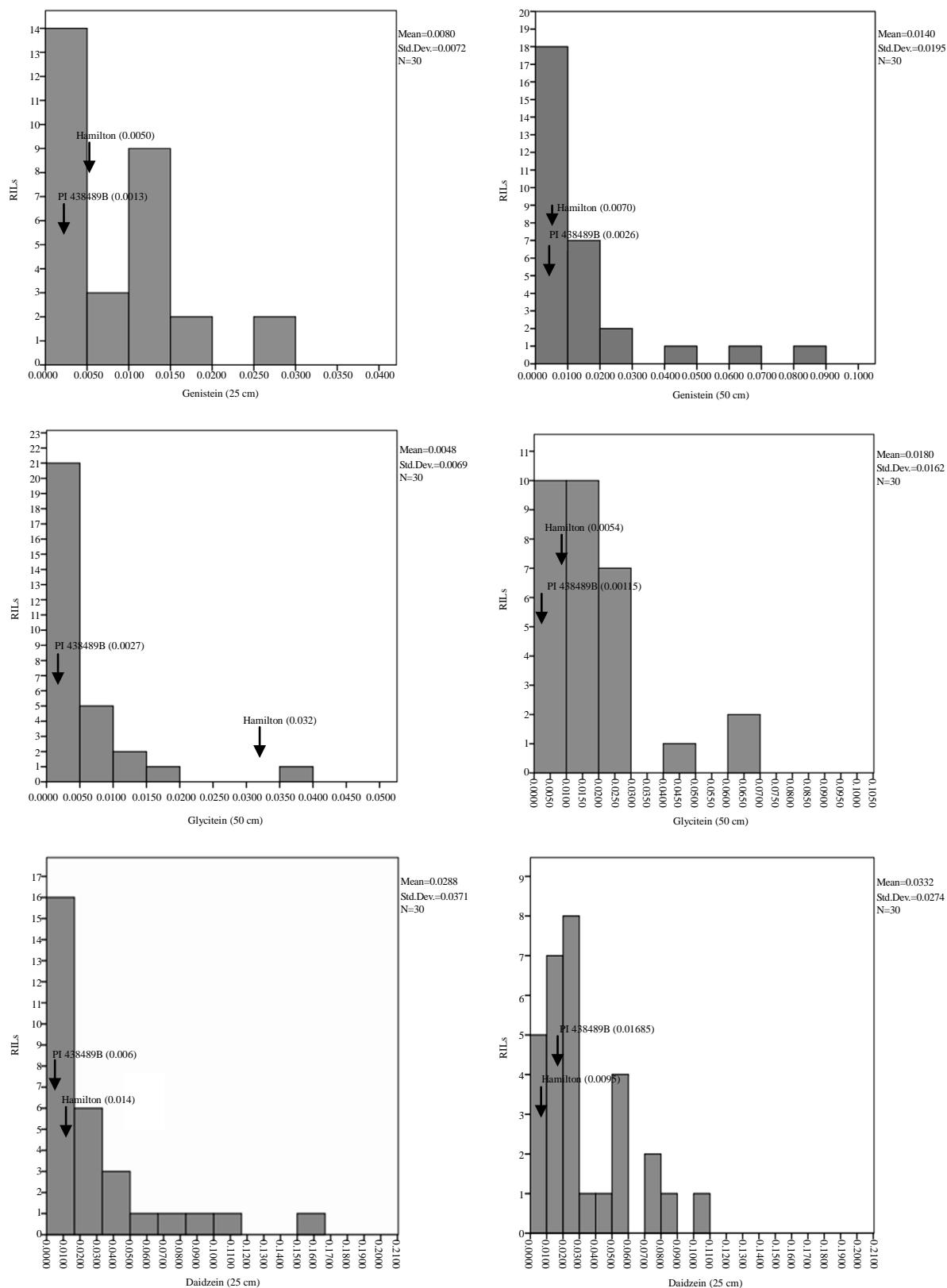


Figure 1. Frequency distribution of seed isoflavone contents of PI 438489B by Hamilton RILs plants grown in two row spaces (25 cm and 50 cm). Seed isoflavone amounts are expressed in $\mu\text{g}\cdot\text{g}^{-1}$ of seed weight.

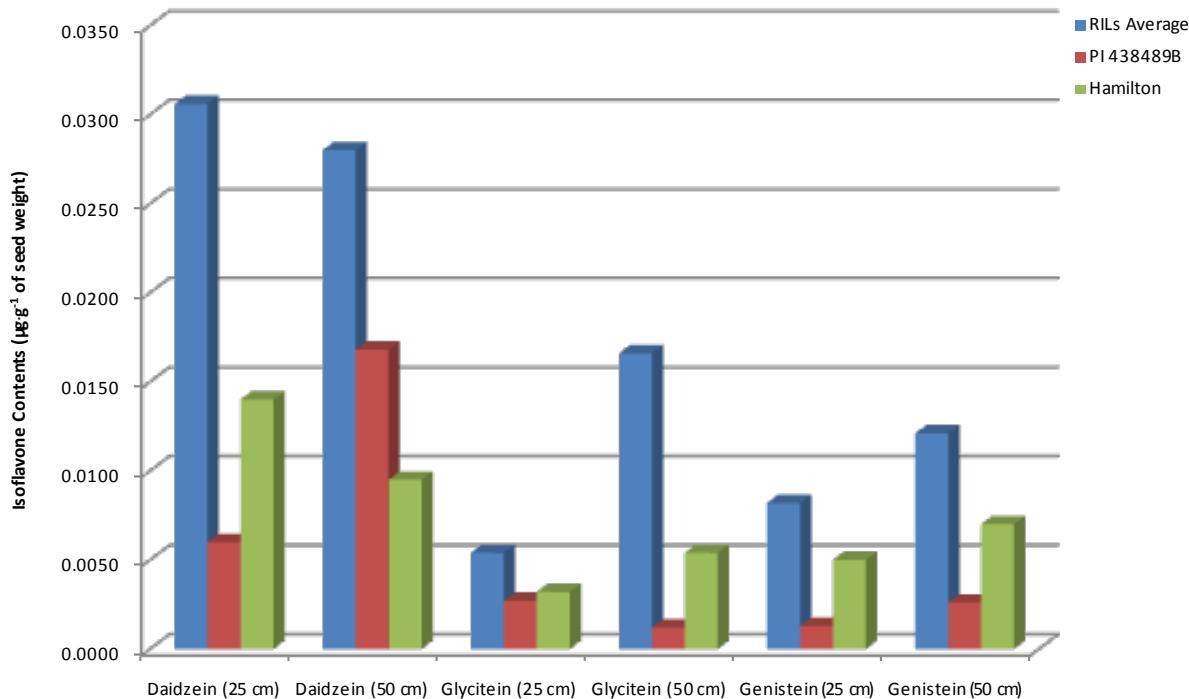


Figure 2. Comparison of levels of isoflavone contents between parent and offspring. Seed isoflavone amounts are expressed in $\mu\text{g}\cdot\text{g}^{-1}$ of seed weight.

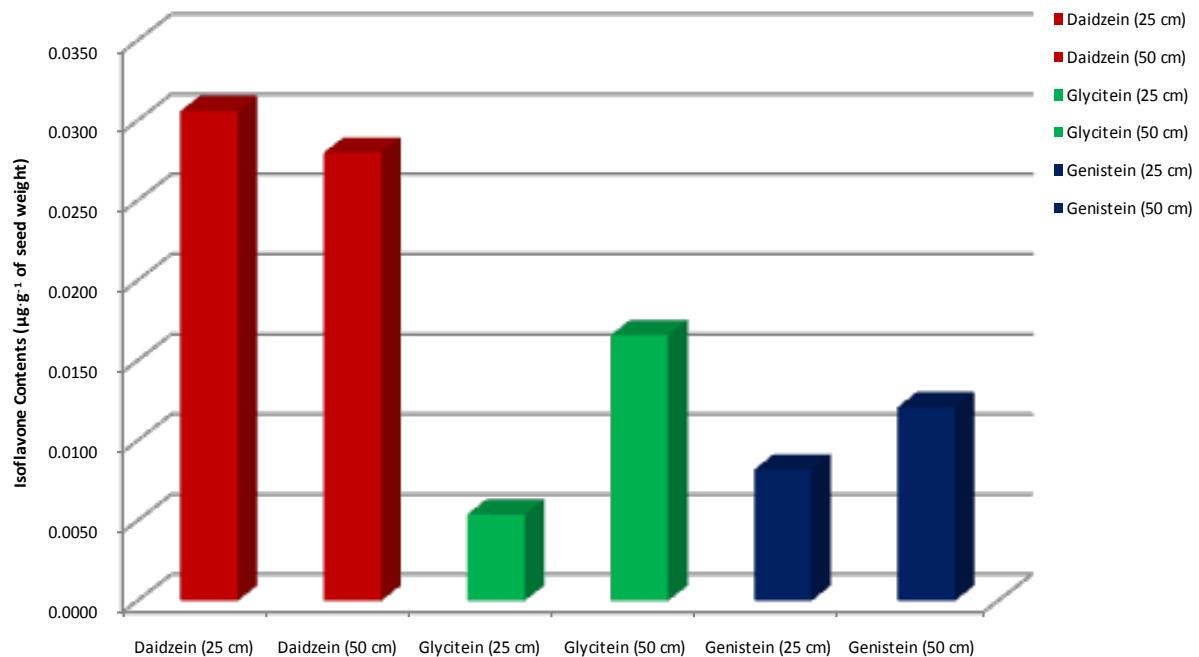


Figure 3. Comparison of isoflavone contents in plants grown at 25 cm vs. 50 cm row spaces. Seed isoflavone amounts are expressed in $\mu\text{g}\cdot\text{g}^{-1}$ of seed weight.

to Group I plants (25 cm row spaces) which is in disagreement with the previous and only study that reported no effect of row spacing on seed isoflavone contents [17]. Another study reported that an increase in protein content was promoted by a decrease in row spacing in two soybean cultivars PB-1 and G-2 while the contents of several minerals decreased [6]. Al-Tawaha and Seguin [17] found negative correlations between yield crude protein (CP)

Table 3. T-test showing summary of paired statistics for the two experimental conditions for seed daidzein (Dai), genistein (Gen), and glycinein (Gly) contents between Group I and Group II RILs. Seed isoflavone amounts are expressed in $\mu\text{g}\cdot\text{g}^{-1}$ of seed weight.

Paired samples statistics				
	Isoflavones	Mean	Std. deviation	Std. error mean
Pair 1	Dai25	0.0301	0.0380	0.0072
	Dai50	0.0345	0.0277	0.0052
Pair 2	Gly25	0.0049	0.0071	0.0013
	Gly50	0.0190	0.0162	0.0030
Pair 3	Gen25	0.0083	0.0072	0.0013
	Gen50	0.0146	0.0200	0.0037

Pair 1, 2, and 3; Comparisons of daidzein, glycinein, and genisten contents in Group I and II plant seeds, respectively.

Table 4. The Pearson correlation of seed isoflavone contents in the two plant Group I (grown with 25 cm row spaces) and Group II (grown with 50 cm row spaces).

Paired samples correlations			
		Correlation	Sig.
Pair 1	Dai25 & Dai50	0.200	0.308
Pair 2	Gly25 & Gly50	0.096	0.626
Pair 3	Gen25 & Gen50	0.106	0.591

Pair 1, 2, and 3; Comparisons of daidzein, glycinein, and genisten contents in Group I and II plant seeds, respectively.

and seed yield, and seed isoflavone contents. A prolonged drought stress during the stages of seed development drastically reduces the accumulation of seed isoflavone contents [23] [24]. However, Caldwell *et al.* [23] found that drought combined with high levels of CO₂ at 23°C promoted the levels of 6'-O-malonygenistin and genistin [23].

While the results presented here shed some light on the effect of row spacing on seed isoflavone contents, further studies are needed using different cultivars and/or populations as well as studies on the correlations between seed yield and isoflavone contents.

Acknowledgements

The authors would like to thank the Department of Defense (DOD) for funding this work through the grant# W911NF-11-1-0178 to MAK and SK. I would also like to thank all the undergraduate students who helped care for plants in greenhouse and fields. Many thanks to AURAK for its partial payment of the publication fee.

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