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INDEPENDENT INHERITANCE OF GENES FOR DHURRIN AND LEUCOANTHOCYANIDIN IN A SORGHUM CROSS

F. A. HASKINS* AND H. J. GORZ

Abstract

Flag leaves of KS8 sorghum [*Sorghum bicolor* (L.) Moench] are low in dhurrin [*p*-hydroxy-(S)-mandelonitrile- β -D-glucoside] content and thus in hydrocyanic acid potential (HCN-p), and they contain little (if any) leucoanthocyanidin (LAC). Comparable leaves of 'Colman' sorghum are intermediate in HCN-p and high in LAC. This study was conducted to investigate the inheritance of HCN-p and LAC in crosses of KS8 $\times$ Colman. Flag leaves from field-grown plants of both parents; the F_1 , KS8 $\times$ Colman; the backcross, KS8 $\times$ (KS8 $\times$ Colman); and the F_2 , (KS8 $\times$ Colman) selfed, were assayed for both HCN-p and LAC. Assays for HCN-p indicated that the backcross values provided a good fit to 1 intermediate:1 low ($X^2 = 0.05$, $P = 0.82$) and F_2 results to 3 intermediate:1 low ($X^2 = 0.96$, $P = 0.33$). For LAC, backcross results were 1 LAC+:1 LAC- ($X^2 = 0.20$, $P = 0.65$) and F_2 values were 3 LAC+:1 LAC- (exact fit). Classification for both traits yielded good fits to 1:1:1:1 for the backcross ($X^2 = 0.30$, $P = 0.96$) and 9:3:3:1 for the F_2 ($X^2 = 3.24$, $P = 0.36$). These results indicated that the difference in HCN-p between Colman and KS8 depended primarily on a single gene, the difference in LAC depended largely on a separate single gene, and the HCN-p gene and the LAC gene were not linked.

A 1986 REPORT (1) indicated that a single gene difference was primarily responsible for the difference in dhurrin content (expressed as hydrocyanic acid potential, HCN-p) of mature leaves between KS8 (low-HCN-p) and N32 (high-HCN-p) lines of sorghum. Similarly, the large difference in flag-leaf leucoanthocyanidin (LAC) content between 'White Collier' (LAC-) and Colman (LAC+) sorghums depended largely on a single pair of genes (2). Both KS8 and N32 were LAC-, and the HCN-p values for flag leaves of both Colman and White Collier were intermediate between the values for KS8 and N32 (F.A. Haskins and H.J. Gorz, 1988, unpublished observations). Therefore, the crosses used in the previous studies were considered inappropriate for investigating the possible linkage of the HCN-p and LAC traits. The objective of the present study was to investigate the segregation of these two traits in a cross of KS8 (low-HCN-p, LAC-) $\times$ Colman (intermediate-HCN-p, LAC+).

Materials and Methods

Colman (male-fertile), AKS8 (cytoplasmic male-sterile), and BKS8 (male-fertile, sterility maintainer) sorghum plants were grown at the Agronomy Farm, Lincoln, NE, during the summer of 1985. The status of Colman with respect to fertility restoration was unknown; therefore, to ensure that male-fertile F_1 plants would be produced, BKS8 plants were hand emasculated and pollinated with Colman pollen. In Novem-

ber, 1985, the F_1 seed was sent to Puerto Rico (Tropical Agricultural Research Station, USDA-ARS, Mayaguez) where F_1 plants were self-pollinated to produce F_2 seed, and were also used as pollen parents in backcrosses to AKS8. AKS8 $\times$ Colman crosses also were made during the summer of 1985 to obtain larger quantities of seed from which F_1 plants could be grown for sampling.

Plants of BKS8, Colman, the F_1 (AKS8 $\times$ Colman), the backcross [AKS8 $\times$ (BKS8 $\times$ Colman)], and the F_2 [(BKS8 $\times$ Colman) selfed] were started in the greenhouse during the spring of 1986 and were transplanted to the Agronomy Farm on 29 May. Plants were placed 0.61 m apart in rows with a 0.76-m spacing. The experiment was planted in four replications, with each replication including one 10-plant row of each parent and the F_1 , two rows of the backcross, and five rows of the F_2 . Thus, the experiment was designed to include 40 plants of each parent and the F_1 , 80 backcross plants, and 200 F_2 plants. Entries were assigned at random to the 10 rows in each replication. Insufficient plants of both parents and the F_1 were available for transplanting; therefore, seeded rows of BKS8, Colman and the F_1 were used in two, three, and one of the replications, respectively.

On 30 July, when panicles were emerging from most plants, the blade of the flag leaf was harvested from each plant. One KS8 and 21 F_2 plants were not sufficiently advanced to allow positive identification of the flag leaf; for these plants, the blade of the youngest leaf with a visible collar was harvested. Midribs were removed from the leaf blades, and the remaining tissue was dried at 70 to 75 °C overnight. The dry tissue was ground through a 1-mm screen and stored in plastic vials at -18 °C prior to extraction for assay.

Extraction and assay for HCN-p were conducted as described previously (1) except that dhurrin was hydrolyzed enzymatically rather than with NaOH. The enzyme preparation was an extract made by soaking defatted almond meal (Sigma Chemical Co., St. Louis, MO)¹ in distilled water (8 mg mL⁻¹) and filtering the suspension through Whatman no. 1 filter paper. To hydrolyze dhurrin, 1 mL of this filtrate was added to 1 mL of leaf extract, and the mixture was incubated in a parafilm-capped tube at room temperature for 1.25 h. Following this incubation, 8 mL of 0.1 M NaOH was added, and a 1-mL portion of the resulting solution was assayed colorimetrically as described previously (1).

The procedure described by Haskins and Gorz (2) was used for extraction and assay of LAC. Absorbance at 540 nm was used as a measure of LAC content.

Results and Discussion

Segregation for HCN-p

Mean HCN-p values for KS8 and Colman were 29 and 135 mg kg⁻¹, respectively, and standard errors were such that these means appeared to be well separated (Table 1). However, ranges in HCN-p for these two parents overlapped slightly, which caused some uncertainty in the classification of backcross and F_2 plants as either low (L) or intermediate (I) in HCN-p. As shown in the table, with 50 mg kg⁻¹ as the dividing line between the classes, only one of 37 KS8 plants was classified as I-HCN-p and only two of 40 Colman plants were classified as L-HCN-p. With this dividing line, 41 of the backcross plants were I-HCN-

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¹ Names of products are included for the benefit of the reader and do not imply endorsement or preferential treatment by the USDA or the Univ. of Nebraska.

Table 1. Level of hydrocyanic acid potential (HCN-p) and leucoanthocyanidin (LAC, as measured by A_{540}) in flag leaves of parental KS8 and Colman sorghum plants and of F_1 (AKS8 $\times$ Colman), BC_1 [AKS8 $\times$ (BKS8 $\times$ Colman)], and F_2 [(BKS8 $\times$ Colman) selfed] plants derived from these parents.

Entry	Phenotypic class†				HCN-p		A_{540}		Chi-square
	HCN-p: LAC:	I +	I -	L +	Mean $\pm$ SE	Range	Mean $\pm$ SE	Range	
					mg kg ⁻¹ dry wt				
KS8			1		29 $\pm$ 2	7-52	0.005 $\pm$ 0.0004	0-0.010	
Colman		38		2	135 $\pm$ 13	31-349	1.33 $\pm$ 0.05	0.770-2.0	
F_1		36		3	86 $\pm$ 6	47-194	0.51 $\pm$ 0.02	0.242-0.72	
BC_1		22			170 $\pm$ 23	61-505	0.41 $\pm$ 0.04	0.090-0.80	for 1:1:1:1
			19		222 $\pm$ 37	51-751	0.006 $\pm$ 0.001	0-0.017	0.30
				20	27 $\pm$ 2	11-44	0.43 $\pm$ 0.04	0.117-0.86	(P = 0.96)
					30 $\pm$ 3	10-48	0.006 $\pm$ 0.001	0-0.013	
F_2		104			204 $\pm$ 19	52-1178	0.64 $\pm$ 0.04	0.063-1.80	for 9:3:3:1
			40		332 $\pm$ 40	52-1247	0.007 $\pm$ 0.001	0-0.017	3.24
				46	31 $\pm$ 2	4-50	0.63 $\pm$ 0.05	0.034-1.40	(P = 0.36)
				10	25 $\pm$ 4	7-38	0.008 $\pm$ 0.001	0.001-0.014	

† Class limits: Intermediate (I) HCN-p—values >50 mg kg⁻¹; Low (L) HCN-p—values ≤ 50 mg kg⁻¹; LAC+— $A_{540} > 0.02$; LAC— $A_{540} \leq 0.02$.

p and 39 were L-HCN-p; and in the F_2 , 144 plants were I-HCN-p and 56 were L-HCN-p. Chi-square values for fits to 1:1 and 3:1 ratios were 0.05 (P = 0.82) and 0.96 (P = 0.33), respectively. These data support the hypothesis that a single gene difference had a major role in determining HCN-p in crosses between KS8 and Colman. The possible relationship of this gene to the gene with a major influence on HCN-p in the KS8 $\times$ N32 cross (1) was not investigated.

Segregation for LAC

The difference in LAC between KS8 and Colman plants was more distinct than the difference in HCN-p. The lowest A_{540} value observed for Colman was about 70 times as large as the highest value for KS8 (Table 1). An A_{540} value of 0.02 was used to separate LAC+ and LAC- classes. The observed backcross segregation was 42 LAC+:38 LAC-, and the F_2 segregation was 150 LAC+:50 LAC-. The former ratio was a good fit to 1:1 (χ^2 = 0.20, P = 0.65), and the latter an exact fit to a 3:1 ratio, indicating that with respect to LAC, KS8 and Colman differed primarily by a single gene. In preliminary qualitative tests, F_1 plants resulting from the cross N32 $\times$ White Collier and also from (KS8 $\times$ N32) $\times$ White Collier were uniformly LAC- (F.A. Haskins and H.J. Gorz, 1988, unpublished observations). Thus, KS8, N32, and White Collier probably were genetically similar with respect to their inability to form LAC.

Independent Segregation for HCN-p and LAC

Classification of the 80 backcross and 200 F_2 plants for both HCN-p and LAC provided good fits to 1:1:1:1 and 9:3:3:1 ratios, respectively (Table 1). Thus, the two traits were inherited independently in the cross of KS8 $\times$ Colman.

In a previous study (3), assay of F_2 plants from a cross of Colman $\times$ White Collier for both HCN-p and LAC revealed a significant but rather weak negative association between the two traits [r = -0.42** (at the 0.01 probability level), n = 119]. The possibility was suggested that this relationship might have resulted from linkage between the major gene for LAC and one or more genes for HCN-p. Alternatively, it

was suggested that the negative association might have arisen from competition for common precursors of dhurrin and LAC, both of which are aromatic compounds. The backcross and F_2 ratios obtained in the present study indicated that in the KS8 $\times$ Colman cross, major genes for HCN-p and LAC were not linked. Furthermore, if such linkage existed, a positive association between HCN-p and LAC in the backcross and F_2 generations would be expected, because the L-HCN-p and LAC- traits came from one parent and I-HCN-p and LAC+ traits from the other parent in the cross. This positive relationship was not observed; the relationship between HCN-p and LAC in the backcross generation was nonsignificant (r = 0.092, n = 80), and in the F_2 there was a weak but significant negative relationship (r = -0.202**, n = 200). The tendency toward a negative relationship between HCN-p and LAC also was apparent in the backcross and F_2 means (Table 1). In the backcross generation, the mean HCN-p value for the I-HCN-p, LAC+ class was 170 mg kg⁻¹ compared to 222 for the I-HCN-p, LAC- class. Similarly, in the F_2 generation, means of 204 and 332 mg kg⁻¹ were obtained for the I-HCN-p, LAC+ and I-HCN-p, LAC- classes, respectively. Also, mean HCN-p values were higher and A_{540} values were lower for the I-HCN-p, LAC+ classes in both backcross (170 and 0.41) and F_2 (204 and 0.64) generations than for the Colman parent (135 and 1.33). However, regardless of any effect LAC level might have had on HCN-p, the two major genes appeared to be inherited independently.

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