Genetic Basis for Varied Levels of Injury to Sweet Corn Hybrids from Three Cytochrome P450-metabolized Herbicides

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ABSTRACT. Some sweet corn (*Zea mays* L.) hybrids and inbreds can be severely injured or killed after postemergence applications of certain P450-metabolized herbicides. Consequently, existing hybrids are regularly evaluated for tolerance to new herbicides, and new hybrids are evaluated for tolerance to existing herbicides. In 2005 and 2006, the University of Wisconsin Cooperative Extension Service coordinated 12 trials in six states in which a total of 149 sweet corn hybrids were evaluated for tolerance to three cytochrome P450-metabolized herbicides: nicosulfuron, foramsulfuron, and mesotrione. Hybrid responses differed substantially within and among locations. The objective of this study was to determine if alleles affecting herbicide sensitivity (e.g., cytochrome P450 alleles) were associated with differences in levels of injury to sweet corn hybrids in these trials. Based on responses of F₂ progeny to nicosulfuron, foramsulfuron, and mesotrione, 95 hybrids were classified as homozygous for alleles conditioning herbicide tolerance; 47 hybrids were classified as heterozygous with one allele each conditioning tolerance and sensitivity; and two hybrids were classified as homozygous for alleles conditioning sensitivity. When trial mean levels of injury after applications of mesotrione, nicosulfuron, and foramsulfuron in the herbicide trials were above 1%, 4%, and 5%, respectively, the response of the three genotypic classes of hybrids followed a consistent pattern. Homozygous-sensitive hybrids were injured most severely and often were killed by the two acetolactate synthase-inhibiting herbicides, nicosulfuron and foramsulfuron. Heterozygous hybrids had an intermediate response to all three herbicides that was more similar to homozygous-tolerant hybrids than homozygous-sensitive hybrids; however, injury to heterozygous hybrids was 1.5 to 2.3 times greater and significantly ($P < 0.05$) different from homozygous-tolerant hybrids based on *t* tests of group means and comparisons of predicted values from regressions of genotypic means on trial means. Based on responses of the 149 hybrids in this trial, the potential for and level of crop injury from use of nicosulfuron, mesotrione, and foramsulfuron on any specific sweet corn hybrid is conditioned largely by alleles at a single locus.

Cytochrome P450 enzymes in corn play an integral role in phase I metabolism of herbicides in at least six chemical families (Barrett 1995, 2000). Some sweet corn hybrids and inbreds can be injured or killed after postemergence applica-

tions of certain P450-metabolized herbicides, including those that inhibit acetolactate synthase (ALS) or 4-hydroxyphenylpyruvate-dioxygenase (HPPD) such as nicosulfuron, foramsulfuron, and mesotrione (Diebold et al., 2003; Masiunas et al., 2004; Morton and Harvey, 1992; O'Sullivan et al., 2002; Robinson et al., 1993; Stall and Bewick, 1992; Williams et al., 2005). The adoption and use of new postemergence herbicides in sweet corn and the use of existing herbicides on new sweet corn hybrids are limited by the potential for crop injury. A better understanding of the basis for variation among sweet corn hybrid response to P450-metabolized herbicides would allow for more informed decisions about the potential for crop injury when these chemicals are used.

Varied responses to P450-metabolized herbicides may be the result of dissimilar rates of metabolism among sweet corn hybrids that differ for alleles that regulate cytochrome P450 monooxygenases. Differences in metabolism rates of nicosulfuron appear to be responsible for different responses of a tolerant sweet corn hybrid, 'Landmark', and a sensitive sweet corn hybrid, 'Merit' (Burton et al., 1994). Sensitivity to

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nicosulfuron is conditioned by a single recessive gene designated as either *nsf1* or *ben1* (Barrett et al., 1997; Green and Ulrich, 1993; Kang, 1993). The dominant allele, *Nsf1*, from a nicosulfuron-tolerant field corn inbred line, B73, was recently sequenced after being located on chromosome 5S. *Nsf1* was the second locus among a cluster of four closely linked genes with homologies to cytochrome P450 genes (i.e., CYP genes) (Williams et al., 2006). Nicosulfuron-sensitive field corn inbred lines, GA209 and W703A, from which the *ben1* and *nsf1* genes were originally identified (Fleming et al., 1988; Kang, 1993), contained the same 392-base pair insertion in the *Nsf1* gene sequence relative to nicosulfuron-tolerant lines. Hence, *nsf1* and *ben1* appear to be the same insertion mutation of a CYP gene.

Some sweet corn hybrids and inbreds are sensitive to multiple P450-metabolized herbicides, including nicosulfuron, foramsulfuron, and mesotrione (Nordby, 2008; Williams et al., 2005). Sensitivity to these and other P450-metabolized herbicides in the sweet corn inbred Cr1 is inherited as a single recessive gene (Pataky et al., 2006) or a group of closely linked genes that appear to be allelic with or very closely linked to the *nsf1/ben1* gene (Nordby et al., 2008). This gene or group of genes appears to be widely prevalent in commercially available sweet corn hybrids and their inbred parents (Pataky et al., unpublished data). Biomass of sweet corn hybrids nearly isogenic for CYP alleles at *Nsf1* or a closely linked locus differed after application of mesotrione in dose-response studies (Volenberg et al., 2006). Biomass of the heterozygous hybrid was intermediate to that of near isogenic homozygous-sensitive and homozygous-tolerant hybrids.

Because of potential injury from postemergence herbicides, existing sweet corn hybrids are regularly evaluated for response to new herbicides, and new hybrids are evaluated for response to existing herbicides to identify sensitive hybrids (e.g., Bollman et al., 2006; Diebold et al., 2003; Edenfield and Allen, 2005; Morton and Harvey, 1992; O'Sullivan et al., 2002; Williams and Pataky, 2008; Williams et al., 2005). In 2005 and 2006, the University of Wisconsin Cooperative Extension Service coordinated a sweet corn hybrid-herbicide trial in which a total of 149 hybrids were evaluated for tolerance to nicosulfuron, foramsulfuron, and mesotrione in 12 trials in Wisconsin, Minnesota, New York, Delaware, Colorado, and Idaho (Bollman et al., 2005). Responses of hybrids differed substantially within and among locations leading to our hypothesis that variation in levels of injury was associated with the presence in some hybrids of one or two mutant CYP alleles (e.g., the *nsf1/ben1* allele).

The objective of this study was to determine if alleles affecting herbicide tolerance and sensitivity (e.g., CYP alleles) were associated with differences in levels of injury to sweet corn hybrids in herbicide trials coordinated by the University of Wisconsin in 2005 and 2006.

Materials and Methods

HERBICIDE TRIALS. Responses of 149 commercial sweet corn hybrids to postemergence applications of nicosulfuron, foramsulfuron, and mesotrione were evaluated in 2005 and 2006 in a total of 12 field experiments at nine locations, including Olathe, CO; Georgetown, DE; Caldwell, ID; Nampa, ID; Stanton, MN; LeRoy, NY; DeForest, WI; Plainfield, WI; and Sun Prairie, WI. The hybrids represent diverse genetic

backgrounds from six different commercial sweet corn breeding programs and include sugary, sugary, enhancer, and shrunken-2 endosperm mutants. These hybrids are grown commercially for fresh market and processing throughout North America and the world. Trials in 2005 and 2006 included 114 and 81 sweet corn hybrids, respectively. Forty-five hybrids were common to trials in both years. Field sites were prepared using conventional production practices. Preemergence herbicides applied after planting were selected to control early-season weeds based on site-specific needs. Each trial included six duplicate ranges of sweet corn hybrids planted in 6-m long rows spaced 76 cm apart. Planting density was 40 seeds per row. Within each pair of ranges, one was treated with a postemergence herbicide and one was a nontreated control. Thus, each trial included one replicate of hybrids for each of three pairs of herbicide-treated and nontreated ranges. Randomization of hybrids differed in each of seven trials (i.e., locations) in 2005 and five trials (i.e., locations) in 2006.

Herbicide treatments were applied at the V3 to V4 growth stage (Ritchie et al., 1997) at twice the normal use or anticipated use rates to maximize differences among tolerant and sensitive hybrids. Nicosulfuron was applied at 70 g·ha⁻¹ a.i. plus 1% v/v petroleum oil concentrate and 2.2 kg·ha⁻¹ ammonium sulfate. Foramsulfuron was applied at 74 g·ha⁻¹ a.i. plus the safener isoxadifen-ethyl at 74 g·ha⁻¹ a.i., 1.8 L·ha⁻¹ methylated seed oil, and 2.2 kg·ha⁻¹ ammonium sulfate. Safeners such as isoxadifen-ethyl enhance the activity of degradative enzymes such as cytochrome P450 monooxygenases, thus protecting plants from injurious herbicides such as the sulfonylureas (Hatzios, 1997). Mesotrione was applied at 210 g·ha⁻¹ a.i. plus 1% v/v petroleum oil concentrate. All three herbicides were applied in 190 L·ha⁻¹ spray volume.

Sweet corn injury was evaluated 7 and 14 d after application of herbicide treatments. Injury from nicosulfuron and foramsulfuron was visually assessed as percent stunting (0% to 100%). Injury from mesotrione was assessed visually as the percentage of the total leaf area bleached or chlorotic (0% to 100%).

GREENHOUSE AND FIELD EVALUATIONS OF F₂ PROGENY. Sweet corn hybrids included in the 2005 and 2006 herbicide trials were grown in 2006 at the University of Illinois Vegetable Crops Farm, Champaign. Each hybrid was grown in a single, 3-m long row of 15 plants. Seeds of F₂ progeny were produced for 144 of the 149 hybrids by self-pollinating five to 10 plants per row.

Phenotypic responses of F₂ progeny to nicosulfuron, foramsulfuron, and mesotrione were assayed to determine if hybrids were homozygous or heterozygous for a gene(s)-conditioning response to these herbicides. F₂ progeny of 144 hybrids were assayed for responses to nicosulfuron and foramsulfuron in greenhouse trials at the University of Illinois Plant Care Facility. Responses of F₂ progeny to mesotrione were assayed in field trials in 2007 at the University of Illinois Crop Sciences Research and Education Center, Champaign.

Greenhouse evaluations of responses of F₂ progeny to postemergence applications of nicosulfuron and foramsulfuron were adapted from the methods used by Nordby et al. (2008). Seventy-two seeds of F₂ progeny of each hybrid were planted in a randomly assigned 30 × 60 × 7-cm flat containing a 1 : 1 : 1 mixture of soil, peat, and perlite supplemented with 15N–3.9P–9.7K Osmocote Plus® (Scotts Co., Marysville, OH) slow-release fertilizer. One flat of an herbicide-sensitive inbred, Cr1,

and one flat of an herbicide-tolerant inbred, Cr2, were included as controls. Herbicides were applied when plants were at the three-leaf stage, ≈ 13 d after planting. Two flats were sprayed simultaneously in an enclosed spray chamber equipped with an 80° flat-fan nozzle delivering 187 L·ha⁻¹ of treatment solutions at 207 kPa. Nicosulfuron was applied at 35 g·ha⁻¹ a.i. with 0.25% v/v nonionic surfactant and 2.5% v/v 28% urea ammonium nitrate. Foramsulfuron was applied at 40 g·ha⁻¹ a.i. plus the safener isoxadifen-ethyl at 40 g·ha⁻¹ a.i. with 1% v/v methylated seed oil and 2.5% v/v 28% urea ammonium nitrate.

In the field trial in 2007, two replicates of F₂ progeny of each hybrid and the sensitive and tolerant inbreds were planted in single, 3-m long rows with ≈ 25 plants per row. The experimental design was a randomized complete block. Mesotrione was applied at the V3 to V4 growth stage at 105 g·ha⁻¹ a.i. plus 1% v/v crop oil concentrate and 2.5% v/v 28% urea ammonium nitrate.

Plants were rated visually ≈ 13 d after nicosulfuron and foramsulfuron were applied and 7 d after mesotrione was applied. Plants sensitive to nicosulfuron or foramsulfuron usually were dead or nearly dead, although some plants sensitive to foramsulfuron occasionally had malformed leaves and symptoms similar to injury caused by a plant growth regulator. Leaf tissue in whorls of plants sensitive to mesotrione was severely chlorotic or “bleached.” The number of sensitive and tolerant F₂ plants was counted. Hybrids with all F₂ progeny killed or severely injured were classified as sensitive and homozygous for an allele-conditioning herbicide sensitivity. Hybrids with uninjured F₂ progeny were classified as tolerant and homozygous for an allele conditioning herbicide tolerance. Hybrids with sensitive and tolerant F₂ progeny were tested by χ^2 analysis for goodness of fit for segregation of tolerant:sensitive F₂ progeny in a ratio of 3 : 1, which would be expected if a single dominant gene conditioned a tolerant phenotypic response (e.g., the *Nsf1* allele). If segregation of F₂ progeny fit a 3 : 1 ratio ($P > 0.01$), hybrids were classified as heterozygous for alleles conditioning tolerance and sensitivity. If more than 10% of the F₂ progeny were sensitive but the ratio of tolerant:sensitive plants did not fit a 3 : 1 ratio ($0.01 < P < 0.001$), hybrids were still classified as heterozygous. When fewer than 5% of the F₂ progeny were sensitive to the three herbicides and segregation did not fit a 3 : 1 ratio ($P < 0.001$), hybrids were classified as tolerant.

COMPARISON OF FIELD RESPONSES OF TOLERANT, SENSITIVE, AND HETEROZYGOUS HYBRIDS. Responses of individual hybrids were not compared within trials because each of the 12 trials included only one replicate of hybrids treated with each of the three herbicides. However, in each trial, an adequate number of hybrids were classified as homozygous-tolerant or heterozygous to allow for statistical comparison of mean injury from each herbicide between the groups of homozygous-tolerant and heterozygous hybrids. Thus, mean levels of injury were compared by *t* tests between 74 homozygous-tolerant and 34 heterozygous hybrids in 2005 and between 54 homozygous-tolerant and 24 heterozygous hybrids in 2006. Homozygous-sensitive hybrids were not included in the statistical analysis because sample size was too small ($n = 2$ and 1 in 2005 and 2006, respectively).

Responses to each herbicide also were compared by regression of mean levels of injury of homozygous-tolerant and heterozygous hybrids from each trial against trial mean levels of injury. Injury to heterozygous hybrids was compared as a

ratio with injury to homozygous-tolerant hybrids based on predicted values from regression.

Results

EVALUATIONS OF F₂ PROGENY AND CLASSIFICATION OF HYBRIDS INTO GENOTYPIC CLASSES. All of the 105 plants of the sensitive inbred, Cr1, were killed or severely injured by applications of nicosulfuron, foramsulfuron, or mesotrione. None of the 110 plants of the tolerant inbred, Cr2, were injured by these herbicides. Two hybrids, ‘Merit’ and ‘EX 08705770’, were classified as homozygous-sensitive because all 373 F₂ progeny of these hybrids were injured or killed after applications of nicosulfuron, foramsulfuron, or mesotrione. These hybrids are homozygous for an allele that conditions herbicide sensitivity. Ninety-five hybrids were classified as homozygous-tolerant (Table 1) based on responses of their F₂ progeny. These hybrids are homozygous for an allele that facilitates metabolism of these herbicides. None of 10,844 F₂ progeny of 73 homozygous-tolerant hybrids were injured after applications of nicosulfuron, foramsulfuron, or mesotrione. An additional 22 hybrids were classified as homozygous-tolerant, although less than 1% (19 of 2534) of the F₂ progeny were sensitive to nicosulfuron or foramsulfuron and less than 4% (117 of 3424) of the F₂ progeny were sensitive to all three herbicides. Forty-seven hybrids were classified as heterozygous based on segregation among F₂ progeny (Table 2). These hybrids have an allele that facilitates herbicide metabolism and an allele that does not. Segregation of F₂ progeny from 41 hybrids was not significantly different ($P \geq 0.01$) from a 3 : 1 ratio of tolerant:sensitive plants (Table 2). The percentage of sensitive F₂ progeny ranged from 14% to 37% for these hybrids. Six hybrids with 10% to 16% sensitive F₂ progeny also were classified as heterozygous, although segregation was significantly different ($P < 0.01$) from 3 : 1 as a result of too few sensitive progeny (Table 2). Possibly, the herbicide-sensitive inbred parent of these hybrids may be segregating at the *Nsf1* locus. This explanation seems more plausible than modifying genes because at least one of these five hybrids is known to segregate for resistant and susceptible responses to maize dwarf mosaic virus (J. K. Pataky, personal observation).

Among the 47 heterozygous hybrids, 10 hybrids had fewer than expected (i.e., 25%) sensitive F₂ progeny for one of the three herbicides (data not shown). Less than 25% of the F₂ progeny of ‘Devotion’, ‘EX 0870 5788’, ‘GH 2042’, ‘Hollywood’, ‘Polaris’, and ‘Welcome TSW’ were sensitive to nicosulfuron. Less than 25% of the F₂ progeny of ‘Bold’, ‘Eliminator’, ‘How Sweet It Is’, and ‘Jubilee’ were sensitive to foramsulfuron. Less than 25% of the F₂ progeny of ‘Jubilee’ were sensitive to mesotrione.

COMPARISON OF FIELD RESPONSES TO NICOSULFURON. Mean injury (percent stunting) at 7 or 14 d after application of nicosulfuron ranged from 2% to 27% and averaged 7.5% among the 12 trials (Table 3). Injury ranged from 15% to 100% and averaged 75% for homozygous-sensitive hybrids. Fourteen days after application of nicosulfuron, homozygous-sensitive hybrids were stunted 80% or more in 10 of 12 trials and were killed in five of these trials. Mean injury to heterozygous hybrids ranged from 0% to 50% and averaged 10% among the 12 trials. On average, heterozygous hybrids were stunted 11% and 10% at 7 and 14 d after application of nicosulfuron, respectively. Mean injury to homozygous-tolerant

Table 1. Sweet corn hybrids classified as homozygous-tolerant for a gene-conditioning herbicide sensitivity based on homogeneous-tolerant responses of F₂ progeny to nicosulfuron, foramsulfuron, and mesotrione.

Hybrid cultivar	Source ^z	Hybrid cultivar	Source	Hybrid cultivar	Source
Alexis ^y	MM	Frosty	Cr	Nantasket	MM
Ambrosia	Cr	Gateway	Rog	Native Gem	MM
Applause	Cr	GH 2171 ^y	Rog	Obsession	Sem
Bliss	SnRv	GH 6462	Rog	Optimum	Cr
Bodacious	Cr	GSS 1303 ^y	Rog	Passion	Sem
Bon Appetit TSW	MM	GSS 1477 ^y	Rog	Polka	Cr
Bon Jour TSW	MM	GSS 2008	Rog	Prelude	SnRv
Breeders Choice	MM	GSS 4165 ^y	Rog	Prime Plus ^y	Sem
BSS 3495	Rog	GSS 6564	Rog	Protégé	Rog
Cameo ^y	Cr	Harvest Gold	Sem	Providence	Rog
Captivate	Cr	HB 1920 OJ	SnRv	Reflection	HM
Cascade	Cr	HB 2162 OL ^y	SnRv	Renaissance	HM
Cavalry	HM	HM 2390	HM	Revelation	HM
Challenger	Sem	HMX 4387WS	HM	Rising Sun	SnRv
Charmed ^y	Cr	HMX 4394	HM	Rocker	Rog
Chase ^y	Cr	HMX 4396 S	HM	Sentinel ^y	HM
Cinderella	Cr	Holiday ^y	Cr	Sheba R	Sem
CNS710 RR	Cr	HW 1336 Ok ^y	SnRv	Spring Treat	MM
Colonial	Cr	HY 0579 OK	SnRv	Sugar Buns	Cr
CSEYP1-3	Cr	HY 17900L	SnRv	Sugar Pearl TSW	MM
CSHYP2-57	Cr	Ice Queen	HM	Surpass	Cr
CSUYP2-28	Cr	Incredible ^y	Cr	Temptation	Sem
Dasher	Cr	Kristine ^y	Cr	Trinity	Cr
Double Up	Rog	Lancaster ^y	SnRv	Tuxedo	MM
Early Gold ^y	HM	Lancelot ^y	MM	UY 0607 OJ	SnRv
Enterprise	SnRv	Legacy	MM	UY 0712 OJ	SnRv
Equinox	Cr	Luscious TSW	MM	UY 1953 OK	SnRv
Everest ^y	SnRv	Maestro ^y	Cr	Valor	Cr
EX 08705353 ^y	Sem	Marvel	Cr	Venus	Cr
EX 08716636	Sem	Miracle ^y	Cr	Whiteout	MM
EX 9381178	Sem	Misquamicut	MM	XP 08705808	Sem
Fleet	Cr	Montauk	MM		

^zCr = Crookham Co., Caldwell, ID; HM = Harris Moran Seed Co., Modesto, CA; MM = Mesa Maize, Olathe, CO; Rog = Syngenta Seeds Inc., Rogers Brands, Boise, ID; Sem = Seminis Inc., Oxnard, CA; SnRv = Snowy River Seeds, Orbst, Australia.

^yHybrids classified as tolerant with less than 1% of F₂ progeny sensitive to nicosulfuron and rimsulfuron and less than 5% of F₂ progeny sensitive to all three herbicides.

hybrids ranged from 0.1% to 20% and averaged 5% among the 12 trials. On average, tolerant hybrids were stunted 5% and 4% at 7 and 14 d after application of nicosulfuron, respectively.

Based on *t* tests, injury was significantly ($P < 0.06$) greater for heterozygous hybrids than for homozygous-tolerant hybrids in 17 of 24 comparisons (Table 3). When the trial mean level of injury was above 4%, heterozygous hybrids were injured significantly ($P < 0.06$) more than homozygous-tolerant hybrids in 15 of 16 comparisons. When the trial mean level of injury was below 4%, heterozygous and homozygous-tolerant hybrids were not significantly different in six of eight comparisons.

When mean levels of nicosulfuron injury to heterozygous hybrids were plotted against the trial mean levels of injury, the slope coefficient from the regression was 1.6 indicating that injury to heterozygous hybrids was $\approx 60\%$ greater than average injury in a trial (Fig. 1A). When mean levels of injury to homozygous-tolerant hybrids were plotted against trial mean levels of injury, the slope coefficient from the regression was 0.7 indicating that injury to tolerant hybrids was $\approx 30\%$ less than average injury in a trial. Slope coefficients for heterozy-

gous and homozygous-tolerant hybrids were significantly different ($P < 0.01$) based on a *t* test. When trial mean injury was above 4%, the ratio of predicted injury to heterozygous hybrids compared with homozygous-tolerant hybrids ranged from 1.7 to 2.2; thus, nicosulfuron injured the group of heterozygous hybrids approximately twice as much as the group of homozygous-tolerant hybrids when conditions were conducive to injury.

COMPARISON OF FIELD RESPONSES TO FORAMSULFURON. Mean injury (percent stunting) at 7 or 14 d after application of foramsulfuron ranged from 2% to 31% and averaged 10.6% among the 12 trials (Table 4). On homozygous-sensitive hybrids, injury ranged from 3% to 100% and averaged 81%. Fourteen days after application of foramsulfuron, homozygous-sensitive hybrids were stunted 90% or more in 10 of 12 trials and were killed in five trials. Mean injury to heterozygous hybrids ranged from 0% to 37% and averaged 13% among the 12 trials. On average, heterozygous hybrids were stunted 15% and 11% at 7 and 14 d after application of foramsulfuron, respectively. Mean injury to homozygous-tolerant hybrids ranged from 0% to 26% and averaged 8% among the 12 trials.

Table 2. Sweet corn hybrids classified as heterozygous for a gene-conditioning herbicide sensitivity based on 3:1 segregation of F₂ progeny for tolerant and sensitive responses to nicosulfuron, foramsulfuron, and mesotrione.

Hybrid cultivar	Source ^z	Plants (no.) ^y		Sensitive (%) ^x	P ^w
		Tolerant	Sensitive		
Accord	MM	112	48	30	0.14
Argent	Cr	120	53	31	0.09
Basin R	Sem	114	50	30	0.10
Bojangles	Cr	130	44	25	0.93
Bold ^v	Rog	162	28	15	0.001
Brocade TSW	MM	81	17	17	0.08
Cahill (GH 6333)	Rog	128	38	23	0.53
Celestial	Cr	119	42	26	0.75
Chief Ouray	MM	118	33	22	0.37
Coho	HM	127	42	25	0.96
Colombus	SnRv	82	24	23	0.57
CSUWP1-7	Cr	112	37	25	0.96
Delectable	Cr	127	43	25	0.93
Devotion ^v	Sem	92	13	12	0.003
Double Gem	MM	110	33	23	0.60
Dynamo	HM	123	57	32	0.04
Eliminator ^v	Cr	155	28	15	0.002
Empire	SnRv	142	31	18	0.03
EX 08705640	Sem	148	39	21	0.19
EX 08705788 ^v	Sem	125	23	16	0.008
EX 08755780	Sem	139	30	18	0.03
GH 2042 ^v	Rog	108	27	20	0.18
GH 2669	Rog	113	44	28	0.38
GH 4927	Rog	138	38	22	0.30
GH 6014	Rog	144	45	24	0.71
GH 6223	Rog	122	19	13	0.002
GH 8267	Rog	139	42	23	0.58
GSS 2914	Rog	73	17	19	0.18
HB 1321 OK	SnRv	65	26	29	0.43
Hollywood ^v	Sem	72	17	19	0.20
How Sweet It Is ^v	Cr	151	27	15	0.002
HY 1901 OL	SnRv	106	27	20	0.21
Jubilee ^v	Rog	70	15	18	0.12
Max	HM	80	25	24	0.78
Mystique	Cr	108	44	29	0.26
Overland	Rog	84	27	24	0.87
Polaris ^v	HM	101	28	22	0.39
Precious Gem	MM	129	36	22	0.35
Punch	SnRv	147	34	19	0.05
Ranger	HM	127	33	21	0.20
SS Jubilee Plus	Rog	57	19	25	1.00
Suregold	HM	91	47	37	0.01
Turbo	HM	117	47	29	0.28
UY 2830 OL	SnRv	144	36	20	0.12
UY 2835 OL	SnRv	138	30	18	0.03
Welcome TSW ^v	MM	95	16	14	0.01
Winstar	Rog	126	39	24	0.69

^zCr = Crookham Co., Caldwell, ID; HM = Harris Moran Seed Co., Modesto, CA; MM = Mesa Maize, Olathe, CO; Rog = Syngenta Seeds Inc., Rogers Brands, Boise, ID; Sem = Seminis Inc., Oxnard, CA; SnRv = Snowy River Seeds, Orbost, Australia.

^yApproximately 50 F₂ progeny each evaluated for tolerant or sensitive responses to nicosulfuron, foramsulfuron, and mesotrione.

^xPercentage of F₂ progeny sensitive to nicosulfuron, foramsulfuron, and mesotrione.

^wProbability associated with a χ^2 goodness of fit test for 3:1 segregation of F₂ progeny for tolerant and sensitive responses to nicosulfuron, foramsulfuron, or mesotrione.

^vHybrids with fewer than expected (i.e., 25%) sensitive F₂ progeny for one of the three herbicides (see text).

Table 3. Mean level of injury on sweet corn hybrids classified as homozygous-sensitive (SS), heterozygous (TS), or homozygous-tolerant (TT) after applications of nicosulfuron in 12 trials in 2005 and 2006.

Location of trial	Yr	Rating (d) ^z	Mean level of injury (%) ^y				<i>P</i> ^v
			Trial mean ^x	SS ^w	TS ^w	TT ^w	
Georgetown, DE	2005	14	27.3	99	49.8	15.7	<0.0001
Georgetown, DE	2005	7	25.5	78	34.9	19.8	<0.0001
Georgetown, DE	2006	7	20.7	90	29.8	15.5	<0.0001
LeRoy, NY	2006	14	13.1	100	19.2	10.5	0.043
Georgetown, DE	2006	14	11.3	100	16.7	8.9	<0.0001
DeForest, WI	2005	7	6.8	60	13.1	2.5	<0.0001
LeRoy, NY	2006	7	6.3	100	7.3	4.3	0.047
Sun Prairie, WI	2005	7	5.9	55	8.0	3.4	<0.0001
Olathe, CO	2005	7	5.7	68	5.8	3.9	NS
Stanton, MN	2006	14	5.6	100	6.5	3.5	0.0350
DeForest, WI	2005	14	5.5	93	8.5	1.8	<0.0001
DeForest, WI	2006	14	5.5	97	5.8	3.6	0.0690
Stanton, MN	2006	7	5.4	85	8.1	2.9	<0.0001
DeForest, WI	2006	7	4.9	45	6.8	3.3	0.0040
Plainfield, WI	2005	7	4.3	73	5.6	2.4	<0.0001
Sun Prairie, WI	2005	14	4.0	90	4.3	1.5	<0.0001
Plainfield, WI	2005	14	3.7	92	2.6	1.4	NS
Caldwell, ID	2005	14	3.0	60	2.7	1.6	NS
Caldwell, ID	2005	7	3.0	25	2.9	2.4	NS
Nampa, ID	2005	7	3.0	15	3.0	2.6	NS
Plainfield, WI	2006	7	2.9	50	3.8	1.6	0.0140
Nampa, ID	2005	14	2.6	55	1.8	1.4	NS
Plainfield, WI	2006	14	2.1	99	1.9	0.5	0.0040
Olathe, CO	2005	14	1.5	80	0	0.1	NS
Mean			7.5	75.4	10.4	4.8	

^zInjury ratings 7 or 14 d after the application of nicosulfuron.

^yInjury measured as stunting as a percentage of the height of nontreated controls.

^xTrial mean injury weighted by number of hybrids in each of three classes. In 2005, *n* = 2, 34, and 74 for SS, TS, and TT hybrids, respectively. In 2006, *n* = 1, 24, and 54 for SS, TS, and TT hybrids, respectively.

^wHybrids classified as SS, TT, or TS for a gene-conditioning herbicide sensitivity based on responses of F₂ progeny.

^vProbability associated with a *t* test comparison of means between TS and TT hybrids.

^{ns}Nonsignificant.

On average, homozygous-tolerant hybrids were stunted 10% and 6% at 7 and 14 d after application of foramsulfuron, respectively.

Based on *t* tests, injury was significantly ($P < 0.06$) greater for heterozygous hybrids than for homozygous-tolerant hybrids in 19 of 24 comparisons (Table 4). When the trial mean level of injury was above 5%, heterozygous hybrids were injured significantly ($P < 0.06$) more than homozygous-tolerant hybrids in 17 of 19 comparisons. When the trial mean level of injury was below 5%, heterozygous and homozygous-tolerant hybrids were not significantly different in three of five comparisons.

When mean levels of foramsulfuron injury on heterozygous hybrids were plotted against the trial mean levels of injury, the slope coefficient from the regression was 1.3 indicating that injury to heterozygous hybrids was ≈30% greater than average injury in a trial (Fig. 1B). When mean levels of injury on homozygous-tolerant hybrids were plotted against trial mean levels of injury, the slope coefficient was 0.88 indicating that injury to homozygous-tolerant hybrids was ≈12% less than average injury in a trial. Slope coefficients for heterozygous and homozygous-tolerant hybrids were significantly different ($P < 0.01$) based on a *t* test. When trial mean injury was above 5%, the ratio of predicted injury to heterozygous hybrids compared with homozygous-tolerant hybrids ranged from 1.5 to 1.8; thus,

foramsulfuron injured the group of heterozygous hybrids ≈50% to 80% more than the group of homozygous-tolerant hybrids when conditions were conducive to injury.

COMPARISON OF FIELD RESPONSES TO MESOTRIONE. Mean injury (percent chlorotic leaf area) at 7 or 14 d after application of mesotrione ranged from 0% to 24% and averaged 4.7% among the 12 trials (Table 5). Injury ranged from 0% to 100% and averaged 33% on homozygous-sensitive hybrids. Seven and 14 d after application of mesotrione, homozygous-sensitive hybrids averaged 37% and 28% chlorotic leaf area, respectively. Mean injury to heterozygous hybrids ranged from 0% to 34% and averaged 7% among the 12 trials. On average, heterozygous hybrids had 10% and 5% chlorotic leaf area at 7 and 14 d after application of mesotrione, respectively. Mean injury to homozygous-tolerant hybrids ranged from 0% to 18% and averaged 3% among the 12 trials. On average, homozygous-tolerant hybrids had 4% and 2% chlorotic leaf area at 7 and 14 d after application of mesotrione, respectively.

Based on *t* tests, injury was significantly ($P < 0.05$) greater to heterozygous hybrids than homozygous-tolerant hybrids in 16 of 23 comparisons (Table 5). When the trial mean level of injury was above 1%, heterozygous hybrids were significantly ($P < 0.05$) more chlorotic than homozygous-tolerant hybrids in 15 of 17 comparisons. When the trial mean level of injury was

Fig. 1. Mean injury in 12 field trials in 2005 and 2006 at 7 and 14 d after application of (A) nicosulfuron, (B) foramsulfuron, or (C) mesotrione on sweet corn hybrids homozygous for alleles conditioning herbicide tolerance or hybrids heterozygous for alleles conditioning tolerance and sensitivity. Injury measured as percent stunting for nicosulfuron and foramsulfuron or percentage of chlorotic leaf area for mesotrione.

below 1%, heterozygous and homozygous-tolerant hybrids were not significantly different in five of six comparisons.

When mean levels of mesotrione injury on heterozygous hybrids were plotted against the trial mean levels of injury, the

slope coefficient from the regression was 1.4 indicating that injury to heterozygous hybrids was $\approx 40\%$ greater than average injury in a trial (Fig. 1C). When mean levels of injury on homozygous-tolerant hybrids were plotted against trial mean levels of injury, the slope coefficient was 0.7 indicating that injury of homozygous-tolerant hybrids was $\approx 30\%$ less than average injury in a trial. Slope coefficients for heterozygous and homozygous-tolerant hybrids were significantly different ($P < 0.01$) based on a t test. When trial mean injury was above 4%, the ratio of predicted injury to heterozygous hybrids compared with homozygous-tolerant hybrids ranged from 2.1 to 2.5; thus, mesotrione injured the group of heterozygous hybrids more than twice as much as the group of homozygous-tolerant hybrids when conditions were conducive to injury.

Discussion

Sweet corn hybrids evaluated for tolerance to three P450-metabolized postemergence herbicides in 12 trials in 2005 and 2006 differed substantially. The degree and type of injury also differed between herbicides with two different modes of action, ALS inhibitors and HPPD inhibitors. When the hybrids were characterized for alleles affecting herbicide metabolism, injury was associated with genotypic classes regardless of the mode of action or the presence of a herbicide safener. Two hybrids that were homozygous for an allele-conditioning herbicide sensitivity (i.e., a mutant CYP allele) were severely injured or killed by all three herbicides. Ninety-five hybrids that were homozygous for an allele-conditioning herbicide tolerance were uninjured or injured least by the three herbicides. Forty-seven hybrids that were heterozygous for alleles conditioning tolerance and sensitivity had responses to all three herbicides that were intermediate to homozygous-sensitive and homozygous-tolerant hybrids. When trial mean levels of injury after applications of mesotrione, nicosulfuron, and foramsulfuron were above minimal levels (i.e., greater than 1%, greater than 4%, and greater than 5%, respectively), the response of the three genotypic classes of hybrids followed a consistent pattern. Homozygous-sensitive hybrids were injured most severely and often were killed by the ALS-inhibiting herbicides. The response of heterozygous hybrids to all three herbicides was more similar to homozygous-tolerant hybrids than to homozygous-sensitive hybrids; however, injury to heterozygous hybrids was ≈ 1.5 to two times greater than injury to homozygous-tolerant hybrids.

Our observation that variation in levels of injury is associated with the presence of one or two nonfunctional CYP alleles is supported further by comparing hybrid responses from previous sweet corn-herbicide trials (Burton et al., 1994; Diebold et al., 2003; Masuinas et al., 2004; Morton and Harvey, 1992; O'Sullivan et al., 2002; Robinson et al., 1993; Stall and Bewick, 1992; Williams et al., 2005). The hybrids 'Merit', 'Del Monte 20-38', and 'Shogun' often have been identified as sensitive to or intolerant of specific, P450-metabolized herbicides. In this study, "Merit" was classified as homozygous-sensitive based on sensitive responses of all F_2 progeny. In a related study (Pataky et al., unpublished data), these three hybrids also were classified as homozygous for CYP alleles conditioning sensitivity. Other hybrids in previous evaluations have been observed to have moderate levels of visible injury or yield reduction resulting from P450-metabolized herbicides, including the hybrids 'Calico Belle', 'Cupola', 'Jubilee',

Table 4. Mean level of injury (%) on sweet corn hybrids classified as homozygous-sensitive (SS), heterozygous (TS), or homozygous-tolerant (TT) after applications of foramsulfuron in 12 trials in 2005 and 2006.

Location of trial	Yr	Rating (d) ^z	Mean level of injury (%) ^y				<i>P</i> ^v
			Trial mean ^x	SS ^w	TS ^w	TT ^w	
Georgetown, DE	2005	7	30.5	73	36.5	26.1	<0.0001
Stanton, MN	2006	7	28.6	97	32.2	25.7	0.0130
Georgetown, DE	2005	14	26.2	95	36.2	18.9	<0.0001
Georgetown, DE	2006	7	21.8	90	26.5	18.2	<0.0001
Stanton, MN	2006	14	15.2	100	21.8	10.6	<0.0001
Georgetown, DE	2006	14	14.0	100	17.2	10.9	<0.0001
LeRoy, NY	2006	14	12.7	100	14.0	10.6	NS
DeForest, WI	2005	7	10.1	68	16.2	5.9	<0.0001
Sun Prairie, WI	2005	7	9.4	78	12.9	6.2	<0.0001
Caldwell, ID	2005	7	8.2	33	8.8	6.8	0.0030
DeForest, WI	2006	7	8.2	40	11.0	6.3	0.0020
Plainfield, WI	2006	7	7.7	60	9.9	5.5	0.0240
DeForest, WI	2006	14	6.8	97	11.2	3.1	0.0010
Plainfield, WI	2005	7	6.6	83	7.8	3.8	0.0020
DeForest, WI	2005	14	6.5	90	9.5	2.6	<0.0001
Sun Prairie, WI	2005	14	5.9	93	7.7	3.0	<0.0001
LeRoy, NY	2006	7	5.7	100	6.3	4.0	0.0470
Caldwell, ID	2005	14	5.5	80	4.8	3.8	0.0580
Olathe, CO	2005	7	5.3	65	4.6	4.1	NS
Plainfield, WI	2006	14	4.8	100	5.3	2.8	0.0250
Plainfield, WI	2005	14	4.1	93	3.6	1.8	0.0200
Nampa, ID	2005	7	4.1	3	4.7	3.9	NS
Nampa, ID	2005	14	3.6	87	2.1	2.2	NS
Olathe, CO	2005	14	1.9	100	0	0.1	NS
Mean			10.6	81	13	8	

^zInjury ratings 7 or 14 d after the application of foramsulfuron.

^yInjury measured as stunting as a percentage of the height of nontreated controls.

^xTrial mean injury weighted by number of hybrids in each of three classes. In 2005, *n* = 2, 34, and 74 for SS, TS, and TT hybrids, respectively. In 2006, *n* = 1, 24, and 54 for SS, TS, and TT hybrids, respectively.

^wHybrids classified as SS, TT, or TS for a gene-conditioning herbicide sensitivity based on responses of F₂ progeny.

^vProbability associated with a *t* test comparison of means between TS and TT hybrids.

NS = nonsignificant.

‘Heritage’, ‘How Sweet It Is’, ‘Rival’, ‘Silver Xtra Sweet’, ‘Silverado’, ‘Snowbelle’, ‘SummerSweet 7201’, ‘SummerSweet 8701’, ‘Supersweet Jubilee’, and ‘Zenith’. Three hybrids among this group that were included in our studies (i.e., ‘Jubilee’, ‘How Sweet It Is’, and ‘Supersweet Jubilee’) were classified as heterozygous based on segregation of their F₂ progeny. Similarly, the hybrid ‘Crisp n Sweet 710’, which had tolerant responses in previous trials, was classified as homozygous-tolerant in this study. Thus, the genetic grouping that explains variation among responses of hybrids to nicosulfuron, foramsulfuron, and mesotrione in these 12 field trials is corroborated by previous evaluations of sweet corn hybrid responses to these and related herbicides.

In some trials, none of the hybrids were substantially injured. Levels of injury on heterozygous and homozygous-tolerant hybrids were not significantly different in 19 of 71 comparisons. Sixteen of those 19 comparisons were from trials in Nampa, ID; Caldwell, ID; and Olathe, CO. In nine of these 16 situations, injury to the homozygous-sensitive hybrids was also substantially less than the mean level of injury for sensitive hybrids. Thus, low levels of injury tended to be associated with trials located in drier climates. Mesotrione has been shown to have greater foliar activity against certain weed species {e.g., common waterhemp (*Amaranthus rudis* Sauer) and large

crabgrass [*Digitaria sanguinalis* (L.) Scop.]} at high relative humidity and temperature (Johnson and Young, 2002). Possibly, injury to sweet corn from these herbicides also is more substantial under these environmental conditions.

Although the genotype of hybrids at a single locus explained a substantial amount of the variation in response of hybrids to nicosulfuron, foramsulfuron, and mesotrione, other factors had an effect on hybrid responses, especially to mesotrione. Some variation was observed within groups of homozygous-tolerant and heterozygous hybrids. For example, injury ratings for a few homozygous-tolerant hybrids were consistently above the regression estimate of mean injury levels for homozygous-tolerant hybrids indicating that those hybrids were consistently injured more than the average homozygous-tolerant hybrid (data not shown). Similarly, a few heterozygous hybrids were consistently injured less than the predicted value from the regression. Hybrids with consistent patterns of deviations from regression may be indicative of other factors that affect phenotype such as minor genes that do not distinctly affect phenotype among segregating F₂ progeny as much as the CYP alleles but, nevertheless, noticeably affect levels of tolerance or sensitivity of hybrids. Thus, a hybrid with a homozygous-tolerant genotype may not always be classified as having a tolerant phenotype and heterozygous hybrids may, in some

Table 5. Mean level of injury on sweet corn hybrids classified as homozygous-sensitive (SS), heterozygous (TS), or homozygous-tolerant (TT) after applications of mesotrione in 12 trials in 2005 and 2006.

Location of trial	Yr	Rating (d) ^z	Mean level of injury (%) ^y				<i>P</i> ^v
			Trial mean ^x	SS ^w	TS ^w	TT ^w	
Georgetown, DE	2005	7	23.5	73	33.3	17.9	<0.0001
Georgetown, DE	2005	14	19.5	70	24.2	16.0	<0.0001
LeRoy, NY	2006	7	14.7	90	29.6	6.9	<0.0001
Georgetown, DE	2006	7	10.8	5	15.1	8.9	0.0090
Stanton, MN	2006	7	6.3	75	14.1	1.7	<0.0001
LeRoy, NY	2006	14	5.8	100	10.6	2.2	<0.0001
Plainfield, WI	2006	7	3.9	60	9.0	0.7	0.0010
Plainfield, WI	2005	7	3.8	65	5.8	0.8	<0.0001
DeForest, WI	2005	7	3.6	43	5.8	1.2	<0.0001
Nampa, ID	2005	7	3.4	3	2.6	3.8	NS
Nampa, ID	2005	14	3.0	0	1.5	4.0	NS
Sun Prairie, WI	2005	7	2.2	20	4.7	0.4	0.0002
Stanton, MN	2006	14	2.0	55	3.4	0.5	0.0010
DeForest, WI	2005	14	1.7	20	3.2	0.3	<0.0001
Plainfield, WI	2005	14	1.4	28	1.7	0.2	0.0070
Sun Prairie, WI	2005	14	1.1	12	2.2	0.1	0.0010
Plainfield, WI	2006	14	0.8	15	1.7	0.1	0.0050
Caldwell, ID	2005	14	0.7	0	0.6	0.7	NS
Caldwell, ID	2005	7	0.7	3	0.5	0.5	NS
DeForest, WI	2006	14	0.1	5	0	0	NS
DeForest, WI	2006	7	0.1	5	0.2	0	0.0430
Olathe, CO	2005	14	0	0	0	0	NS
Olathe, CO	2005	7	0	0	0	0	NS
Mean			5.2	35.5	7.9	3.1	

^zInjury ratings 7 or 14 d after the application of mesotrione.

^yInjury measured as percentage of leaf area bleached and chlorotic.

^xTrial mean injury weighted by number of hybrids in each of three classes. In 2005, *n* = 2, 34, and 74 for SS, TS, and TT hybrids, respectively. In 2006, *n* = 1, 24, and 54 for SS, TS, and TT hybrids, respectively.

^wHybrids classified as SS, TT, or TS for a gene-conditioning herbicide sensitivity based on responses of F₂ progeny.

^vProbability associated with a *t* test comparison of means between TS and TT hybrids.

^{NS}Nonsignificant.

cases, be classified as having tolerant phenotypes (J.D. Bollman, unpublished data). Consequently, field evaluations may be necessary to confirm the phenotype of a hybrid even when the genotype at the *Nsf1* locus or other CYP loci is known.

Segregation of F₂ progeny of hybrids classified as heterozygous was not always consistent among the individual evaluations with nicosulfuron, foramsulfuron, and mesotrione. Most of the hybrids for which the probability from the χ^2 goodness of fit test was low (i.e., 0.001 < *P* < 0.01) were among this group. Additional evaluations of inbred parents and progeny of these hybrids may indicate that metabolism of a specific herbicide may be affected by more than one gene, which would be similar to the genetic basis for the metabolism of bentazon. Bradshaw et al. (1994) reported that tolerance to bentazon in field corn inbreds was conditioned by two independent, dominant genes (i.e., duplicate dominant epistasis), which Barrett et al. (1997) designated as *Ben1* and *Ben2*. The *Ben1* gene, which has been shown to be the same as the *Nsf1* gene (Williams et al., 2006), affected metabolism of bentazon, nicosulfuron, and imazethapyr, whereas the *Ben2* gene was specific for bentazon (Barrett et al., 1997). Also, the activity of *Ben1* was induced by an herbicide safener, naphthalic anhydride, whereas activity of *Ben2* was not. Bentazon metabolism also appears to be conditioned by two independent genes in F_{3:5}

families derived from a cross of sensitive and tolerant sweet corn inbreds, Cr1 and Cr2, respectively (Nordby et al., 2008).

Based on comparisons of predictions from regressions when mean levels of injury in a trial were 7% or greater, injury to heterozygous hybrids from nicosulfuron and mesotrione was 1.9 to 2.3 times greater than injury to homozygous-tolerant hybrids, whereas injury to heterozygous hybrids from foramsulfuron was only 1.5 to 1.7 times greater than injury to homozygous-tolerant hybrids. Although the overall level of injury from foramsulfuron was higher than the other two herbicides (i.e., grand mean levels of injury of 10.6%, 7.4%, and 5.2% for foramsulfuron, nicosulfuron, and mesotrione, respectively), the smaller difference between levels of injury to heterozygous and homozygous-tolerant hybrids from foramsulfuron may have been the result of the inclusion of a herbicide safener, isoxadifen-ethyl, in the commercial formulation of foramsulfuron. Safeners have been shown to enhance the activity of P450 monooxygenases (Kreuz et al., 1996). Differential sensitivity in field corn to P450-metabolized herbicides has been associated with different rates of metabolism (Bunting et al., 2004; Eberlein et al., 1989; Hinz and Owen, 1996). Previously, Bunting et al. (2004) reported increased tolerance in field corn hybrids and increased metabolism of foramsulfuron when the herbicide was applied with the safener, isoxadifen-ethyl. If the normal rate of herbicide

metabolism is slower in heterozygous sweet corn hybrids than in homozygous-tolerant sweet corn hybrids, enhancement of P450 activity from safeners may be of greater consequence in heterozygous hybrids, which would explain a more similar response of heterozygous and homozygous-tolerant hybrids to foramsulfuron.

Although environment and other factors affect crop injury from postemergence herbicides, the potential for injury and the level of injury from nicosulfuron, foramsulfuron, and mesotrione is substantially affected by alleles at a single locus, presumably the *Nsf1/Ben1* locus or closely linked loci on the short arm of chromosome 5. Hybrids can be severely injured or killed after applications of nicosulfuron, foramsulfuron, or mesotrione if hybrids are homozygous for alleles that condition sensitivity to P450-metabolized herbicides. Use of these or other P450-metabolized herbicides on homozygous-sensitive hybrids is likely to result in unwanted, detrimental consequences. Under most conditions, heterozygous or homozygous-tolerant hybrids are uninjured by these herbicides or levels of injury are relatively minor; however, under conditions conducive to crop injury, injury is greater to heterozygous hybrids than homozygous-tolerant hybrids. In the future, as new herbicides are screened for potential injury on sweet corn, and as recommended use rates are established for herbicides, inclusion of hybrids that are homozygous-tolerant, heterozygous, and homozygous-sensitive will provide a more complete assessment of the potential for crop injury. Similarly, by eliminating alleles that condition herbicide sensitivity such as the *nsf1* allele from sweet corn inbred lines, new hybrids will no longer be heterozygous or homozygous-sensitive, and the risk of injury to sweet corn from P450-metabolized herbicides will be reduced.

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