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## RESEARCH ARTICLE

# Abundance and diversity of ground-dwelling arthropods of pest management importance in commercial Bt and non-Bt cotton fields

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## Keywords

Carabidae; Cicindelinae; *Falconia gracilis*; genetically modified cotton; Labiduridae; predatory heteropterans; Staphylinidae.

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## Abstract

The modified population dynamics of pests targeted by the Cry1Ac toxin in *Bacillus thuringiensis* (Bt) transgenic cotton (Bt cotton) and possible reduced insecticide use in these transgenic varieties may exert a variety of effects on ground-dwelling predator communities. A survey of ground-dwelling arthropods was carried out weekly each of 3 years (during the cropping season) in commercial Bt and non-Bt cotton fields. Sixty-five taxa of ground-dwelling arthropods (carabids, cicindelids, staphylinids, dermapterans, heteropterans and araneids) of importance for cotton pest management were recorded in the survey. Species abundance and dynamics across seasons were evaluated with univariate analysis of variance for higher taxa or multivariate principal response curve analysis for the whole community of 65 taxa. Diversity and richness indices and cumulative species curves also were calculated. The analyses demonstrated no differences in the ground-dwelling arthropod communities between cotton types. One araneid species, *Pardosa pauxilla*, comprised ~80% of all araneids, *Labidura riparia* comprised ~96% of all dermapterans, *Megacephala carolina* comprised ~97% of cicindelids and four carabid species (*Stenolophus palliatus*, *Apristus latens*, *Harpalus gravis* and *Anisodactylus merula*) comprised ~80% of carabid species. *M. carolina* outnumbered all other collected species in each of the 3 years. When only predatory carabid species were considered, *A. merula*, *Calosoma sayi*, *Harpalus pennsylvanicus* and *Stenolophus ochropezus* were predominant and numbers trapped were similar between cotton types. The abundance of dermapterans, staphylinids, araneids and heteropterans varied among sample dates and across seasons but did not differ between cotton types. The frequent capture of *M. carolina*, *S. palliatus* and *P. pauxilla* in all fields and seasons in both cottons suggests that these species may be important for monitoring further changes in local communities as result of agricultural practices.

## Introduction

There is considerable interest in determining risks and benefits of agricultural practices for conservation of arthropod communities important for pest control and sustainability. In this context, ground-dwelling arthropod communities, composed of a variety of species with differ-

ent feeding behaviours, have been used to assess the impact of pest management practices and cropping systems on beneficial arthropods (Eyre *et al.*, 1989; Carmona & Landis, 1999). Omnivorous and generalist arthropods are commonly found among ground-dwelling communities, and their role in suppressing pests may be quite significant (Stinner & House, 1990; Breene *et al.*,

1993; Lövei & Sunderland, 1996; Knisley & Schultz, 1997). The conservation of certain groups of ground-dwelling arthropods in crop fields has been attempted through modified agricultural practices, with positive results in several cases (Stinner & House, 1990; Nentwig *et al.*, 1998). Thus, ground-dwelling predatory species can be of substantive economic significance in agricultural systems.

The growing worldwide deployment of *Bacillus thuringiensis* (Bt) transgenic crops may impose risks on communities of ground-dwelling arthropods. Predatory arthropods on the ground may have direct contact with activated Bt toxins released into the rhizosphere through plant root exudates and decaying plant material (Saxena *et al.*, 1999; Zwahlen *et al.*, 2003; Baumgarte & Tebbe, 2005). They may be exposed to toxin through prey that are able to acquire and convey toxins to predators (Saxena & Stotzky, 2001) and may be indirectly impacted through the changes in the herbivore community that serves as the prey base. Despite debates over the amount of Bt toxins that accumulate and remain active in the soil during and after the crop season, and the adequacy of methods used in such studies (Sims & Holden, 1996; Sims & Ream, 1997; Head *et al.*, 2002; Hopkins & Gregorich, 2003; Zwahlen *et al.*, 2003), the results indicate clear differences between laboratory and field experiments. The degradation of the toxin synthesised by plants is expected to co-occur with plant decomposition, which can take days to months depending on environmental conditions, and contact with ground-dwelling species during this period is likely and has been demonstrated (Harwood *et al.*, 2005; Zwahlen & Andow, 2005).

Ground-dwelling arthropods are considered to be important not only for insect pest management but also for managing weeds and other organisms competing with cultivated crops (Stinner & House, 1990; Ball & Bousquet, 2001; Tooley & Brust, 2002); use of Bt transgenic crops can benefit ground-dwelling arthropods by reducing the use of broad-spectrum insecticides. For example, planting Bt potatoes led to increased abundance of ground-dwelling, generalist predator carabids and staphylinids and of araneids in the plant canopy when compared with conventional potato fields treated with broad-spectrum insecticides (Hoy *et al.*, 1998; Reed *et al.*, 2001). On the other hand, Bt toxin may negatively affect epigeal predators through contaminated prey and/or through plants and their products. The omnivorous feeding behaviour of ground-dwelling arthropods can bring them into contact with prey conveying Bt toxin (Saxena & Stotzky, 2001; Zwahlen & Andow, 2005) or with decaying plant material containing the toxin.

Transgenic Bt cotton is completing one decade since the initiation of grower use. Since 1996, the first season of Bt

cotton in grower fields, it has been extensively planted across the US Cotton Belt, with more than 58% of 5.85 million hectares of cotton cultivated with Bt transgenic varieties in 2004 (NASS, 2004). Planting transgenic cotton expressing Cry1Ac toxin from *Bacillus thuringiensis* (Berliner) to manage the bollworm complex (*Helicoverpa*, *Heliothis* and *Pectinophora*) has resulted in reduced insecticide use, with direct benefits to growers and the environment (Betz *et al.*, 2000). Introduction of Bt transgenic varieties coincided with successful eradication of the boll weevil, *Anthonomus grandis* Boheman, throughout the southeastern USA, which contributed to a significant reduction in insecticide use in the region. In Georgia, insecticide was applied to cotton 15–18 times per season prior to boll weevil eradication and was reduced to 4–6 applications after eradication (J. R. R., personal observation). Since the introduction of Bt transgenic varieties, the number of insecticide applications further declined to two to four per season, averaging 2.5 per season (J. R. R., personal observation). Nevertheless, transgenic plants have evoked many questions concerning possible direct and indirect effects on nontarget organisms.

The primary objective of this survey was to determine if Bt cotton and non-Bt cotton, cultivated and managed according to standard grower agricultural practices, support similar abundance, dynamics and diversity of important ground-dwelling arthropod predators. The abundance and diversity of the ground-dwelling arthropod communities in cotton fields present a challenge to selecting an ecologically representative species or group. Therefore in this study, we surveyed a range of taxa, including carabids, cicindelids, staphylinids, dermapterans, heteropterans and araneids – all of which include predators of interest for pest management in cotton. We used commercial cotton fields representative of the size and agronomic practices for the region as the ultimate system to realistically evaluate impacts of transgenic crops on nontarget organisms (Marvier, 2002; O'Callaghan *et al.*, 2005).

## Materials and methods

### Site description and crop management

This study was conducted from 2002 to 2004 in grower fields cultivated with standard agricultural practices and located near Tifton, in southern Georgia, USA. The region is comprised of a mixed mosaic of agricultural habitats and forest remnants. Three pairs of Bt (DPL 458 or DPL 555) and non-Bt cotton (DPL 491) fields, from 5.5 to 15.0 hectares each, were monitored each season. Each pair of fields was separated from the others by 3.2 to 27 km, between the coordinates 31°39'N, 83°54'W and 31°51'N, 83°55'W.

At each location, adjacent fields of Bt and non-Bt cotton were separated from one another by either a water ditch or a field road, and roads separated the study fields from adjacent crops such as peanut, tobacco and watermelon. All fields were planted during the first or second week of May each year and received preventative in-furrow treatments to suppress thrips [aldicarb 560 g (AI) ha<sup>-1</sup>] and foliar insecticide applications as needed during the season to control bollworms [*Helicoverpa zea* (Boddie) and *Heliothis virescens* (Fabr.)], stinkbugs and whiteflies. The Bt cotton varieties used in the study express the Cry1Ac toxin that targets only lepidopteran larvae. The toxin provides absolute control of *H. virescens* and very good control of *H. zea* and a number of other lepidopterans. Based on scouting data, non-Bt fields received insecticide applications to control bollworms, and both Bt and non-Bt fields were treated to control stinkbugs late in the season (Table 1). The non-Bt cotton field at Chula was treated twice for bollworms within 15 days in July 2003 because of rainfall immediately after the first application. The frequency of rain during July 2003 caused variation in dates and frequency of sprays applied to each field.

In 2004, after a 1-year rotation with tobacco in Marchant, the selected Bt and non-Bt fields returned to the same locations used in 2002, only changing crop arrangement inside the cultivated areas. The Chula fields remained in the same location all three seasons, but study fields were rotated inside a cultivated area of approximately 40 ha surrounded by forest remnants. The third pair of fields (Ty Ty, Old House and Frazier) was set in different locations each season.

### Sampling of ground-dwelling arthropods

A convenient pitfall trap was made using 500-ml plastic cups (9-cm diameter × 12-cm depth) (Solo<sup>®</sup> P-16, Solo Cup Company, Urbana, IL, USA). On each side of the cup, we made two holes of 2-cm diameter approximately 5 cm from the bottom and covered with mesh in order to drain excess water resulting from irrigation or rainfall. As retention liquid, we used water mixed with Tween 20 at 0.2% to break surface tension, and as preservative, we used four to five pellets per cup of Diamond Crystal<sup>®</sup> water softener salt (Cargil Co., Minneapolis, MN, USA). Each pitfall cup was installed inside a larger and deeper

**Table 1** Time and insecticide applied to manage pest infestations in Bt and non-Bt cotton fields near Tift County, GA, USA, during 2002–04

Dates (Fields)	Non-Bt Cotton	Bt Cotton	Targeted Pest <sup>a</sup>
2002			
At planting	Aldicarb 15G (560 g ha <sup>-1</sup> ) <sup>b</sup>	Aldicarb 15G (560 g ha <sup>-1</sup> )	Thrips
8–9 July (C, M, T)	Spinosad (100 g ha <sup>-1</sup> )	–	Heliothines
10–12 August (C, M)	Lambda-cyhalothrin (34 g ha <sup>-1</sup> ) + thiodicarb (680 g ha <sup>-1</sup> )	–	Heliothines
14 August (T)	–	Dicrotophos (390 g ha <sup>-1</sup> )	Stinkbugs
5–7 September (C, M)	Pyriproxifen (60 g ha <sup>-1</sup> )	Pyriproxifen (60 g ha <sup>-1</sup> )	Whiteflies
2003			
At planting	Aldicarb 15G (560 g ha <sup>-1</sup> )	Aldicarb 15G (560 g ha <sup>-1</sup> )	Thrips
8 July (O)	Spinosad (100 g ha <sup>-1</sup> )	–	Heliothines
13 July (M)	Spinosad (100 g ha <sup>-1</sup> )	–	Heliothines
14 July (C)	Lambda-cyhalothrin (30 g ha <sup>-1</sup> )	–	Heliothines
21 July (C)	Lambda-cyhalothrin (45 g ha <sup>-1</sup> )	–	Heliothines
3–5 August (M, O)	Lambda-cyhalothrin (45 g ha <sup>-1</sup> )	–	Heliothines + stinkbugs
30 August (O) <sup>a</sup>	Bifenthrin (70 g ha <sup>-1</sup> )	Bifenthrin (70 g ha <sup>-1</sup> )	Stinkbugs
2004			
At planting	Aldicarb 15G (560 g ha <sup>-1</sup> )	Aldicarb 15G (560 g ha <sup>-1</sup> )	Thrips
2–7 July (C, M, F)	Lambda-cyhalothrin (45 g ha <sup>-1</sup> )	–	Heliothines
15 July (C,M)	Spinosad (100 g ha <sup>-1</sup> )	–	
29 July (C)	–	Dicrotophos (420 g ha <sup>-1</sup> )	Stinkbugs
5 August (C)	Zeta-cypermethrin (160 g ha <sup>-1</sup> )	–	Heliothines + stinkbugs
17 August (M, F)	Zeta-cypermethrin (210 g ha <sup>-1</sup> )	Zeta-cypermethrin (210 g ha <sup>-1</sup> )	Heliothines + stinkbugs
31 August (M) <sup>a</sup>	–	Acephate (810 g ha <sup>-1</sup> )	Stinkbugs

Bt, *Bacillus thuringiensis*; C, Chula; F, Frazier fields; M, Marchant; O, Old House; T, Ty Ty.

<sup>a</sup>Insecticide application after terminating experimental sampling; Thrips (*Frankliniella occidentalis*, *Frankliniella fusca* and *Thrips tabaci*), Heliothine (*Helicoverpa zea* and *Heliothis virescens*) and Stinkbugs (*Nezara viridula* and *Euschistus servus*). Thrips threshold (preventative treatment); Heliothine threshold, 8–10% of plants with eggs or small larvae on terminals; stinkbug threshold, 18–20% bolls of ~2.5 cm diameter with internal damage; whitefly threshold, plants infested and honeydew on plants (Guillebeau, 2004).

<sup>b</sup>Rate in grams of active ingredient per hectare.

plastic base cup (10-cm diameter  $\times$  15-cm depth) (Packer Ware<sup>®</sup>, Lawrence, KS, USA) that had no bottom to permit drainage and had been installed previously across the fields. The pitfall cups and their bases were built so that the pitfall cups fit snugly inside the base cup, with the rim of the lining cup held in place slightly below the top edge of the base cup. The upper edge of the base cup was level with the soil surface. Twenty traps were set up from border to border of all fields in 10 equally spaced, prefixed stations with two traps each. The sampling stations were set up immediately after seedling emergence. The first sampling station was set at the fifth cotton row from the border. At each station, two traps were installed within cotton rows and with five rows between traps at the same station. The distance between stations varied as a function of field size. The traps were collected weekly (ca 1 week of exposure) by replacing the cups and using the same base throughout the season. Collected traps were returned to the laboratory, where the contents were washed, removed and stored in 20-ml scintillation vials containing 70% ethanol. The data presented will focus on species representative of arthropod communities relevant to pest management of cotton in the region (e.g. carabids, cicindelinae, dermapterans, staphylinids, heteropterans and araneids).

### Species identification and statistical analysis

In the laboratory, specimens were sorted into morphospecies, and all adult insects were identified to order, family and species as possible. Identification of dermapterans to species was based on Hoffman (1987), Lindroth (1961–1969), Ciegler (2000), the University of Georgia Collection of Arthropods (UGCA), Athens (GA) and the Florida State Collection of Arthropods, Gainesville (FL) were used to identify carabid species; species nomenclature followed Ciegler (2000). Functional group designation of carabid species (carnivore or phytophage) was based on predominant feeding behaviour reported in the literature (Table 2). Cicindelinae were identified to species based on Knisley & Schultz (1997), araneids were identified to species with Kaston (1978) and Breene *et al.* (1993) and Staphylinidae were only sorted to family level. The vials containing the collected material were deposited at the Biological Control Laboratory (University of Georgia-Coastal Plain Experiment Station), in Tifton, GA, and voucher specimens were deposited at the UGCA, Athens, GA.

Prior to analysis, data from individual pitfall traps were pooled within each week and for each field. These totals were standardised as the number of individuals per pitfall trap recovered out of 20 traps per week and per field, discarding traps lost to flooding or other destructive event

and therefore comprising three replications (i.e. fields) for each sampling week. Year-long averages were generated from each weekly average over the number of sample weeks per year (10 weeks in 2002 and 11 weeks in 2003 and 2004). Because the questions of interest were related to overall changes in the species community in Bt cotton relative to non-Bt cotton fields, species were pooled to higher identified taxa (i.e. carabids, cicindelinae, dermapterans, araneids, staphylinids and heteropterans). The data for each taxon were analysed through six independent repeated measures analyses of variance (ANOVA) comparing cotton types within year and across 3 years, which also avoided violation of ANOVA assumptions by considering species occurring only at very low densities. All data were  $\log(x + 1)$  transformed prior to univariate analyses but untransformed averages are presented. The results were submitted to one-way repeated measures ANOVA, with repeated measures on sample weeks within year using fields as a blocking factor because the arthropod sampling was conducted on the same fields over the year. These analyses were carried out using the Proc ANOVA of SAS (SAS Institute, 1999–2001), adapting the PROFILE statement, as suggested by Cody & Smith (1997). For analyses across 3 years, data within year were averaged per year and submitted to ANOVA using the Proc MIXED of SAS (SAS Institute, 1999–2001), with year as a random effect and cotton type as a fixed effect.

Because unequal numbers of pitfall traps were evaluated in each sample period, species accumulation curves were generated to assess the effect of sampling effort (10 weeks in 2002 and 11 weeks each in 2003 and 2004) and numbers of individuals collected (i.e. abundance) on species richness results, allowing comparisons of Bt and non-Bt cotton fields. The software program EstimateS (Colwell, 2004) was used to calculate species accumulation curves for the whole community – species richness through a Jackknife estimator and diversity using the Shannon ( $H'$ ) and Simpson's indices for each field within each season (ca 18 estimations: 9 Bt fields and 9 non-Bt fields), involving 100 randomisations of the samples (Colwell, 2004).

Changes in species abundance of the ground-dwelling community were investigated using multivariate analysis through principal response curves (PRC) and considering each taxon collected in Bt cotton fields relative to non-Bt cotton, which was designated as the control. PRC analysis is a multivariate technique derived from redundancy analysis (RDA) that focuses on the proportion of the variance in variables of interest (in this case, ground-dwelling species of economic interest for pest management collected in Bt cotton field on all sampling weeks throughout the cotton season) explainable by the independent variable, cotton type. Parameters of the PRC were generated using

**Table 2** Totals and means per pitfall trap (Bt = 1569 traps and non-Bt = 1501 traps), functional group, abundance and diversity indices of ground-dwelling arthropods collected in three pairs of Bt and non-Bt commercial cotton fields during the season 2002 ( $n = 10$  sampling weeks) and 2003 and 2004 ( $n = 11$  sampling weeks), Tift County, GA, USA

Taxa	Group <sup>a</sup>	2002			2003			2004		
		<i>n</i>	Bt	Non-Bt	<i>n</i>	Bt	Non-Bt	<i>n</i>	Bt	Non-Bt
Araneae										
Clubionidae										
<i>Castianeira</i> nr. <i>floridana</i>	C <sup>1</sup>	70	0.11	0.12	5	0.003	0.005	9	0.006	0.009
Corinidae										
<i>Falconia gracilis</i> (Keyserling)	C <sup>12</sup>	6	0	0.032	14	0.013	0.010	20	0.017	0.018
Lycosidae										
<i>Hogna</i> sp.	C <sup>1</sup>	15	0.017	0.016	7	0.006	0.004	13	0.012	0.008
<i>Pardosa milvina</i> (Hentz)	C <sup>1</sup>	26	0.027	0.033	35	0.030	0.030	110	0.110	0.068
<i>Pardosa pauxilla</i> Montgomery	C <sup>1</sup>	684	1.16	0.93	476	0.398	0.405	424	0.455	0.250
<i>Schizocosa</i> sp.	C <sup>1</sup>	23	0.026	0.042	8	0.007	0.007	15	0.008	0.016
Oxyopidae										
<i>Oxyopes salticus</i> Hentz	C <sup>1</sup>	14	0.017	0.022	8	0.002	0.016	0	–	–
Salticidae										
<i>Habronattus</i> <i>coecatus</i> (Hentz)	C <sup>1</sup>	19	0.021	0.032	7	0.011	0.003	26	0.025	0.016
Tetragnathidae										
<i>Tetragnatha</i> <i>laboriosa</i> Hentz	C <sup>1</sup>	0	–	–	0	–	–	1	0.002	0
Theridiidae										
<i>Lactrodectus</i> sp.	C <sup>1</sup>	8	0.006	0.012	0	–	–	0	–	–
Thomisidae										
<i>Xysticus</i> sp.	C <sup>1</sup>	3	0.005	0.002	0	–	–	1	0	0.002
Dermaptera										
Forficulidae										
<i>Doru taeniatum</i> (Dohorn)	C <sup>2</sup>	6	0.005	0.017	5	0.001	0.007	23	0.033	0.004
Carcinophoridae										
<i>Euborellia annulipes</i> (Lucas)	C <sup>2</sup>	97	0.13	0.11	2	0	0.006	56	0.036	0.065
Labiduridae										
<i>Labidura riparia</i> (Pallas)	C <sup>2</sup>	2329	3.66	3.49	367	0.04	0.22	2406	2.447	1.547
Coleoptera										
Staphylinidae										
		134	0.19	0.13	200	0.181	0.206	356	0.196	0.419
Cicindelinae										
<i>Megacephala</i> <i>carolina</i> Linnaeus	C <sup>3</sup>	4654	4.59	4.56	6903	6.755	5.207	13 524	11.877	10.462
<i>Cicindela punctulata</i> Oliver	C <sup>3</sup>	259	0.21	0.29	156	0.112	0.157	157	0.157	0.104
<i>Megacephala</i> <i>virginica</i> Linnaeus	C <sup>3</sup>	21	0.20	0.018	99	0.071	0.108	73	0.054	0.071
Carabidae										
<i>Acupalpus testaceus</i> Dejean	C <sup>5</sup>	0	–	–	0	–	–	30	0.024	0.023
<i>Agonum aeruginosum</i> Dejean	U	0	–	–	0	–	–	1	0.002	0
<i>Amara cruceolata</i> Putzeys	P <sup>4,5</sup> –C <sup>5</sup>	10	0.008	0.009	3	0.002	0.004	10	0.005	0.011
<i>Amara impuncticolis</i> (Say)	P <sup>5</sup> –C <sup>5,12</sup>	0	–	–	10	0.007	0.103	6	0.008	0.004
<i>Amara</i> sp.	?	26	0.012	0.018	19	0.017	0.015	10	0.005	0.015
<i>Anisodactylus merula</i> (Germar)	C <sup>5</sup> –P <sup>5</sup>	12	0.019	0.005	50	0.023	0.064	132	0.077	0.014
<i>Apenes sinuatus</i> (Say)	U	0	–	–	1	0.002	0	0	–	–
<i>Apristus latens</i> (LeConte)	U	230	0.16	0.32	88	0.092	0.076	129	0.161	0.048
<i>Ardistomis schaumii</i> LeConte	U	0	–	–	0	–	–	2	0.003	–
<i>Aspidoglossa</i> <i>subangulata</i> (Claudoir)	P <sup>5</sup>	0	–	–	1	0.002	0	0	–	–

Table 2 Continued

Taxa	Group <sup>a</sup>	2002			2003			2004		
		<i>n</i>	Bt	Non-Bt	<i>n</i>	Bt	Non-Bt	<i>n</i>	Bt	Non-Bt
<i>Bembidion semistriatum</i> (Haldeman)	U	2	0	0.002	0	–	–	0	–	–
<i>Calleida decora</i> (Fabricius)	C <sup>5</sup>	1	0.008	0	0	–	–	3	0	0.005
<i>Calosoma sayi</i> Dejean	C <sup>5</sup>	16	0.018	0.003	41	0.012	0.063	26	0.033	0.009
<i>Calosoma scrutator</i> (Fabricius)	C <sup>5</sup>	1	0	0.002	0	–	–	0	–	–
<i>Chlaenius aestivus</i> Say	C <sup>8</sup>	8	0.008	0.014	5	0.004	0.007	2	0.003	0
<i>Chlaenius sericeus</i> sericeus (Foster)	C <sup>5</sup>	0	–	–	0	–	–	1	0.002	0
<i>Chlaenius tricolor</i> Dejean	C <sup>5,8</sup>	0	–	–	5	0.003	0.005	0	–	–
<i>Clivina americana</i> Dejean	U	0	–	–	0	–	–	3	0.003	0.001
<i>Clivina bipustulata</i> (Fabricius)	U	2	0.010	0	13	0.015	0.009	0	–	–
<i>Clivina</i> sp.	?	1	0.002	0	0	–	–	0	–	–
<i>Dicaelus elongatus</i> Bonelli	C <sup>5</sup>	0	–	–	0	–	–	1	0	0.002
<i>Dyschirius filiformis</i> LeConte	U	0	–	–	2	0.005	0	5	0.005	0.003
<i>Dyschirius haemorrhoidalis</i> (Dejean)	U	1	0.002	–	0	–	–	0	–	–
<i>Euryderys grossus</i> Say	P <sup>5,8</sup>	1	0	0.012	0	–	–	0	–	–
<i>Galerita bicolor</i> Drury	C <sup>5</sup>	0	–	–	0	–	–	2	0.002	0
<i>Harpalus caliginosus</i> (Fabricius)	C <sup>5</sup> –P <sup>5,8</sup>	10	0.018	0.013	6	0.008	0.005	6	0.005	0.006
<i>Harpalus gravis</i> LeConte	U	9	0.015	0.013	520	0.621	0.358	20	0.022	0.011
<i>Harpalus pennsylvanicus</i> (De Geer)	C <sup>5</sup> –P <sup>5</sup>	25	0.031	0.025	10	0.003	0.019	6	0.009	0
<i>Lebia analis</i> Dejean	C <sup>5</sup>	0	–	–	1	0.002	–	1	0.002	0
<i>Lebia ornata</i> Say	C <sup>5</sup>	0	–	–	1	0	0.002	0	–	–
<i>Lebia viridis</i> Say	C <sup>2</sup>	6	0.010	0.006	0	–	–	0	–	–
<i>Leptotrachelus dorsalis</i> (Fabricius)	C <sup>5</sup>	0	–	–	0	–	–	3	0.002	0.003
<i>Loxandrus velocipes</i> Casey	U	2	0.005	0	0	–	–	0	–	–
<i>Morion monilicornis</i> (Latreille)	U	1	0	0.002	0	–	–	0	–	–
<i>Nemotarsus elegans</i> LeConte	U	0	–	–	1	0.002	–	0	–	–
<i>Platynus decentis</i> (Say)	C <sup>5</sup>	0	–	–	1	0	0.002	0	–	–
<i>Scarites quadriceps</i> Claudoir	C <sup>6</sup>	0	–	–	0	–	–	3	0	0.005
<i>Scarites subterraneus</i> Fabricius	C <sup>5</sup> –P <sup>8</sup>	2	0.005	0	0	–	–	0	–	–
<i>Selenophorus palliatus</i> (Fabricius)	P <sup>7</sup>	679	0.66	0.94	1669	1.99	1.15	323	0.28	0.26
<i>Semiardistomis viridis</i> (Say)	U	0	–	–	0	–	–	1	0.002	0
<i>Stenolophus conjunctus</i> (Say)	C <sup>5</sup>	0	–	–	1	0	0.002	0	–	–
<i>Stenolophus ochropezus</i> (Say)	C <sup>9,12</sup> –P <sup>5</sup>	5	0.010	0.10	13	0.008	0.020	10	0.012	0.006
<i>Tetragonoderus fasciatus</i> (Haldeman)	U	6	0.007	0.004	0	–	–	0	–	–
<i>Tetragonoderus intersectus</i> (Germar)	U	31	0.032	0.044	43	0.026	0.044	20	0.008	0.029
Heteropteran										
Geocoridae										
<i>Geocoris punctipes</i> (Say)	C <sup>10</sup>	65	0.064	0.082	24	0.016	0.030	55	0.032	0.052
<i>Geocoris uliginosus</i> (Say)	C <sup>10</sup>	94	0.092	0.105	16	0.012	0.012	243	0.164	0.245

Table 2 Continued

Taxa	Group <sup>a</sup>	2002			2003			2004		
		<i>n</i>	Bt	Non-Bt	<i>n</i>	Bt	Non-Bt	<i>n</i>	Bt	Non-Bt
Nabidae										
<i>Nabis</i> spp.	C <sup>11</sup>	17	0.024	0.017	233	0.151	0.272	12	0.004	0.017
Total/seasonal		9631	11.4	11.5	11 069	11.7	9.1	18 280	16.3	14.0
mean per trap			(7.8–15.1) <sup>b</sup>	(8.4–14.6)		(8.9–14.5)	(6.5–13.1)		(13.0–19.6)	(10.5–17.3)
Total species richness			37.5	39.0		37.9	37.5		36.9	38.4
(Jackknife estimator)			(34.3–40.7)	(35.5–42.5)		(35.4–40.4)	(34.9–40.1)		(34.6–39.2)	(34.6–42.2)
Diversity (H') index			1.42	1.56		1.28	1.55		1.03	1.01
			(0.92–1.91)	(1.37–1.74)		(0.40–2.15)	(0.83–2.26)		(0.39–1.67)	(0.33–1.68)
Simpson's index			2.47	2.80		2.41	2.78		1.88	2.05
			(1.99–2.96)	(2.14–3.46)		(2.20–2.62)	(2.19–3.37)		(1.62–2.14)	(1.63–2.47)

Bt, *Bacillus thuringiensis*.

<sup>a</sup>Functional group based on predominant feeding behaviour as carnivorous (C), phytophagous (P), unknown (U) and respective reference source: <sup>1</sup>Breene *et al.* (1993), <sup>2</sup>Hoffman (1987), <sup>3</sup>Knisley & Schultz (1997), <sup>4</sup>Johnson & Cameron (1969), <sup>5</sup>Larochelle & Larivière (2003), <sup>6</sup>Best & Beegle (1977), <sup>7</sup>Lindroth (1961–1969), <sup>8</sup>Ball & Bousquet (2001), <sup>9</sup>Jo & Smitley (2003), <sup>10</sup>Crocker & Whitcomb (1980), <sup>11</sup>Lattin (1989), <sup>12</sup>Frank & Shrewsbury (2004) and <sup>12</sup>Bonaldo (2000).

<sup>b</sup>Values between parentheses stand for 95% confidence intervals of means from untransformed data, and *P* values comparing mean abundance per trap within group between cotton types are offered in the text.

CANOCO 4.5 for Windows (Lepš & Šmilauer, 2003) through RDA least-squares estimates. By plotting values of  $c_{dt}$  for the treatment over sampling time, a PRC diagram is obtained that depicts species abundance changes in the community composition. We compared community abundance changes in Bt cotton fields with the non-Bt cotton community as our standard ( $c_{dt} = 0$ ). For each set of analyses, the null hypothesis that the PRC does not explain significant treatment variance was tested using an *F*-type test obtained by permutating the whole time series in the partial RDA from which the PRC was obtained (Lepš & Šmilauer, 2003). Random permutation through the Monte-Carlo method (999 permutations) was also performed for significant treatment PRCs using CANOCO 4.5 within each sampling date to test the null hypothesis that, on each sampling date, the principal response  $c_{dt}$  did not differ significantly between cotton types. Abundance values (predator species per pitfall trap) were log transformed to reduce the effect of dominant species.

## Results

### Seasonal patterns

Numbers of carabids captured gradually declined throughout the growing season (one-way repeated measures ANOVA,  $F_{10, 40} = 19.08$ ;  $P < 0.0001$ ) but did so equally between Bt and non-Bt cotton fields ( $F_{1, 4} = 0.32$ ,  $P = 0.6027$ ), and there were no cotton and year interactions ( $F_{2, 8} = 0.22$ ,  $P = 0.8049$ ). When only predatory carabids were considered, no changes in abundance were observed in any year over time (sampling week effect,  $F_{10, 40} = 1.26$ ;  $P = 0.2835$ ), nor were differences ob-

served between cotton types ( $F_{1, 4} = 0.42$ ,  $P = 0.5529$ ). Likewise, there was no significant interaction between cotton type and year ( $F_{2, 8} = 0.00$ ,  $P = 0.9983$ ). Cicindelids were more commonly collected in traps as June progressed, and their abundance declined significantly later (sampling week effect,  $F_{10, 40} = 40.98$ ;  $P < 0.0001$ ) but equally between Bt and non-Bt cotton fields ( $F_{1, 4} = 0.40$ ,  $P = 0.5596$ ) and with no interactions between cotton type and year ( $F_{2, 8} = 0.273$ ,  $P = 0.5107$ ). The other four taxa (dermapterans, araneids, staphylinids and heteropterans) were similarly abundant in both cotton types ( $P > 0.05$ ), with high variability among sample weeks throughout each season. No pattern of abundance emerged for these taxa, except in the case of heteropterans ( $F_{10, 40} = 3.0$ ,  $P = 0.0064$ ), which declined in number towards the end of the season.

### Ground-dwelling species abundance

A total of 38 980 ground-dwelling individuals, comprising 65 taxa of interest in cotton pest management, were collected across all seasons and fields during the study period. All the specimens were identified to species, except Staphylinidae, which were sorted only to family level. Species of geocorids, nabids, cicindelids and dermapterans were collected in all fields, as were the most abundant species of carabids and araneids (Table 2). Among the most abundant predatory ground-dwelling taxa (more than 5–10 individuals were collected in a single year), none was unique to cotton genotype or year, except one carabid, *Acupalpus testaceus* Dejean, that comprised 30 individuals collected only in 2004 but was found in both Bt and non-Bt cotton fields (Table 2).



Seasonal averages for pitfall catches ranged from 9.1 to 16.3 ground-dwelling arthropods per trap per sample date across seasons and means always overlapped within 95% confidence intervals between Bt and non-Bt cotton fields (Table 2). Therefore, there is no evidence for differences in relative numbers captured per trap or in predominant species within groups between Bt and non-Bt cotton fields.

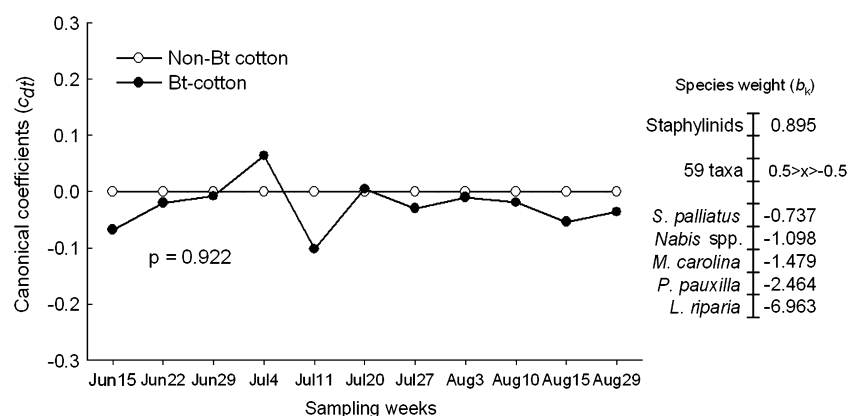
With the exception of heteropterans and araneids, all other groups varied in abundance across seasons (season effect under mixed model of ANOVA). A significant effect of seasons on abundance of dermapterans was observed ( $F_{2, 12} = 5.33$ ,  $P = 0.021$ ), with relatively low numbers of the predominant species, *Labidura riparia*, trapped in 2003 compared with the other years. Likewise, numbers of cicindelids varied across seasons ( $F_{2, 12} = 8.68$ ,  $P = 0.0047$ ), with numbers of *Megacephala carolina* L. increasing late during 2004, similar to what was observed for staphylinids ( $F_{2, 12} = 9.65$ ,  $P = 0.0032$ ), which also were more abundant in 2004. All carabids, including predators and phytophagous species, tended to be trapped in greater numbers in 2003 ( $F_{2, 12} = 5.54$ ,  $P = 0.0198$ ), especially common species such as *Selenophorus palliatus* Fabr. and *Harpalus gravis* LeConte (Table 2). Because populations of each group varied differently in occurrence among seasons, the mean number of individuals captured per pitfall trap did not differ significantly with cotton type for any surveyed group within a season or over the three-season period.

Principal response curve analyses, in accordance with univariate analyses with higher taxa (Table 2), showed no statistically significant impact of Bt cotton compared with non-Bt cotton (the standard reference) on numbers

of 65 taxa of ground-dwelling arthropods trapped in all 3 years pooled ( $F = 2.86$ ,  $P = 0.922$ ; Fig. 1). Sample week (i.e. time) was the major contributor to variance in species abundance, with 73.8% of this variance explained by the first PRC axis. Although the second PRC axis explained an additional 11.2% of variance, the second PRC axis was not significant ( $P > 0.05$ ). The interaction of sample week and cotton type explains 54.7% of the variance, whereas variance because of Bt cotton alone accounted for only 4.2%. This result underscores the apparent lack of effect of Bt cotton on trap capture and dynamics of ground-dwelling arthropods in Bt relative to non-Bt cotton fields (Fig. 1). The contribution of each species to the community changes (response,  $c_{dt}$ ) depicted by PRC diagrams also can be interpreted using the statistical weights ( $b_k$ ) of each species shown on the right side of the diagram (Fig. 1). Species with high weight values are most likely to exhibit population patterns that correspond to the changes in abundance shown in the diagram, while low values contribute little to the overall community response, indicating a weak association or a response pattern different from that displayed in the diagram (Van den Brink & Ter Braak, 1999). Thus, of the 65 taxa, only those that make relatively important statistical contributions are shown on the right side of the diagram.

#### Experimental fields and species abundance and diversity

The outcomes showed no significant effect of field area on number of species collected in either Bt ( $r = 0.12$ ,  $P = 0.7396$ ) or non-Bt cotton fields ( $r = -0.55$ ,

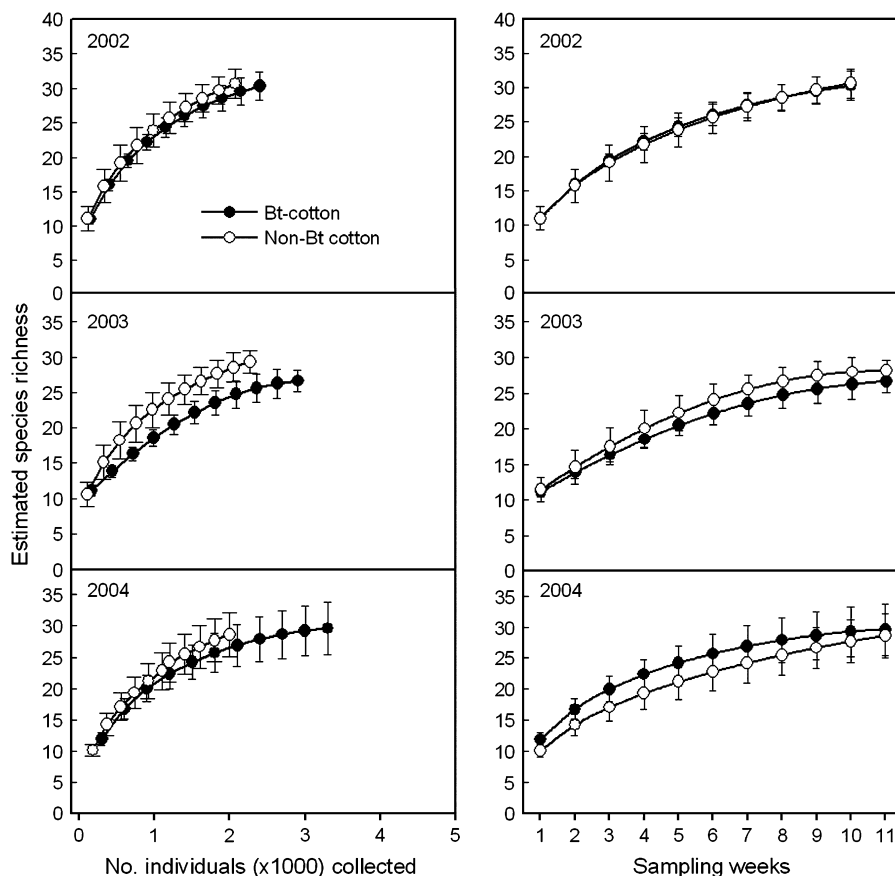


**Figure 1** Principal response curves (PRC) and species weights for species numbers collected in pitfall trap throughout the cotton-growing seasons of 2002–04 in Tift County, GA. The PRC curve shows the main effect of *Bacillus thuringiensis* (Bt) cotton on predator community relative to non-Bt cotton ( $y = 0$  line). The  $P$  value indicates significance of the PRC diagram over all sampling dates based on  $F$ -type permutation test. The higher the species weight, the more likely the actual response pattern of the species follows the pattern of the PRC. For a complete list of all species included in the analyses, see Table 2.

$P = 0.1178$ ). Similarly, the number of species collected was not influenced by numbers of individuals collected in Bt ( $r = -0.05$ ,  $P = 0.8972$ ) or in non-Bt ( $r = 0.33$ ,  $P = 0.3819$ ) cotton fields. Also, the species numbers collected were not a function of the number of pitfall traps recovered from each field (Bt,  $r = -0.17$ ,  $P = 0.6471$  and non-Bt,  $r = 0.16$ ,  $P = 0.6673$ ). These results suggest no interaction between final sampling effort (i.e. number of pitfall traps recovered and the relative abundance and diversity of ground-dwelling predator arthropods sampled) and number of species sampled in each year and cotton type (Fig. 2). These results are supported by the species accumulation curves generated for each cotton type and year (Fig. 2). There is a clear trend in increase of total numbers of individuals collected among years (Table 2) and between Bt and non-Bt cotton fields in 2004 (Fig. 2). However, the increase in species accumulation was not a linear relationship with sampling weeks or abundance (i.e. individuals collected). Thus, increases in number of individuals collected until sampling week 10 reached a plateau and did not result in

significant differences in species richness, which is clearly seen in re-scaling sampling efforts for sampling weeks (Gotelli & Colwell, 2001). Moreover, the means of the estimated total species richness for the respective cotton types overlap within the 95% confidence intervals within seasons and across all years (Table 2).

Although 65 taxa of ground-dwelling arthropods were identified from family to species level, relatively few taxa comprised the majority of the trapped specimens (Table 2). Among araneids, *Pardosa pauxilla* Montgomery was the most commonly captured species, accounting for 68.5–85% of the 11 araneid species collected across years (Table 2). Cicindelinae and Dermaptera were chiefly represented by one species each, *M. carolina* and *L. riparia*, respectively, comprising >94% of all collected individuals of cicindelinae and dermapterans. For example, *M. carolina* comprised 48.3% of specimens of all taxa collected in 2002 and outnumbered all other taxa together (>50%) in 2003 and 2004. Carabidae was the most speciose taxon, with 44 species collected, but only four species – *S. palliatus* Fabr., *Apristus latens* LeConte,



**Figure 2** Comparison of species accumulation curves of ground-dwelling arthropods in *Bacillus thuringiensis* (Bt) and non-Bt cotton fields based on number of individuals collected and number of sampling weeks for each season from 2002 to 2004.

*H. gravis* and *Anisodactylus merula* Germar – accounted for more than 80% of all carabids collected. Of the most abundant carabids, one is omnivorous (*A. merula*) and the other three are predominantly seed feeders (Larochelle & Larivière, 2003). Among those species with a predatory habit, four species were more abundant, and the most abundant species again was the omnivorous *A. merula*. The others were *Calosoma sayi* Dejean, *Harpalus pennsylvanicus* DeGeer and *Stenolophus ochropezus* (Say) (Table 2). Among predatory heteropterans, there was relative constancy in numbers caught among the species, except for *Nabis* spp. and *Geocoris uliginosus* (Say), which were predominant in 2003 (85%) and 2004 (78.4%), respectively.

Diversity, dominance and species richness, as measured by the Shannon ( $H'$ ) and Simpson's indices, and total species richness are shown in Table 2 as means of the three fields within years for each cotton type. The 95% confidence intervals of the estimated means comparing cotton types within year always overlapped. Both Shannon and Simpson indices tended to be lower in 2004 because of large collections of individuals representing few species, as indicated in the species accumulation curves (Fig. 2). Total species richness ranged from 36.9 to 39 species (Table 2), although the number of species per field within a season ranged from 25 species collected at Frazier field in 2004 to 35 at Marchant field in 2003, both in non-Bt cotton fields. Three new state records for Georgia were found: one spider, *Falconia gracilis* (Keyserling), a species accidentally introduced into the USA and already reported in Florida and Texas (Bonaldo, 2000), was collected in all 3 years, with abundance increasing from 2002 to 2004 and two carabid species, *Apristus latens* and *Euryderus grossus* Say, based on the catalogue of Bousquet & Larochelle (1993) and Arnett (2000).

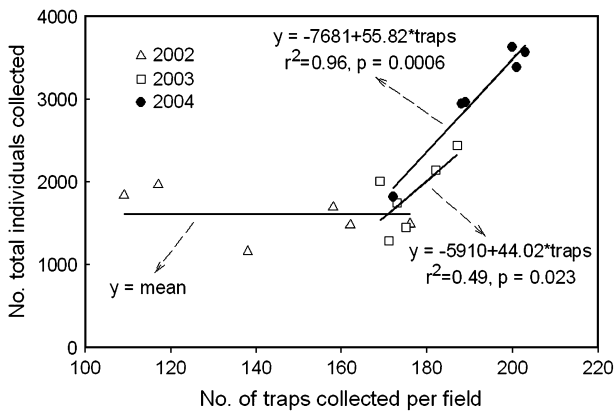
## Discussion

The 3-year data show that Bt cotton fields sustain abundance and species richness of ground-dwelling arthropods of agronomic interest for cotton pest management at levels comparable to non-Bt cotton fields receiving reduced insecticide applications (Table 2 and Figs 1 and 2), despite changes in dynamics of potential prey targeted by Bt cotton. The mean differences for abundance of each species rated per pitfall trap (e.g. non-Bt mean minus Bt cotton mean) produced values of 0.0129, 0.0204 and  $-0.006$  individuals per pitfall trap in 2002, 2003 and 2004, respectively. These values corresponded to an average of 1.29%, 2.03% and  $-0.63\%$  of difference between Bt and non-Bt fields, indicating no significant differences between cotton types.

The cotton field size (areas from 5.5 to 15 ha) and agricultural practices (insect control – Table 1) of the 18 fields studied from 2002 to 2004 reflect typical cotton production in the region. Assessment risks on commercial scales provide the most realistic field experiments for studying nontarget impacts of transgenic crops on natural enemies (Marvier, 2002; O'Callaghan *et al.*, 2005). A concurrent study of the dynamics of foliage-dwelling predatory arthropods on Bt and non-Bt cotton, using the same fields, is reported in Torres & Ruberson (2005). These pooled data reasonably cover most of the possible changes in ground- and foliage-dwelling arthropod communities of importance for pest management in the regional cotton ecosystem under standard grower farming practices.

Variation in total abundance among years was observed. The numbers of individuals collected per season increased from 2002 to 2004 (Table 2), although relative numbers averaged per trap were similar between cotton types and did not correlate with species richness (Table 2 and Fig. 2). This result was not unexpected because in 2002, only 860 traps were recovered over 10 sampling weeks compared with 2003 and 2004, with 11 sampling weeks each. Also the number of individuals captured did not correlate with number of traps recovered in 2002 (Fig. 3), as few traps were lost in 2002 because of limited rainfall (144.5 mm accumulated over 25 rain days) compared with the following years. In 2003 and 2004, 1058 and 1153 traps were evaluated, respectively, and the number of individuals collected was positively correlated with number of traps evaluated (Fig. 3). Differences in total number of individuals between 2003 and 2004 can be explained by the additional 95 pitfall traps recovered in 2004. Rainfall also may have had some effect. In 2003, 470.4 mm of rainfall was accumulated over 43 rain days, whereas in 2004, only 245.0 mm of rainfall accumulated throughout 28 days of raining. Flooding and mud inside pitfall traps were the major causes of trap losses, resulting in the capture of relatively fewer individuals and a trend towards reduced means of individuals caught per trap in 2003 compared with 2004 (Table 2).

Although variation in the community structure was observed within and across years, there was no significant effect of cotton genotype (Fig. 1 and Table 2). The seed-feeding carabids *S. palliatus* and *H. gravis*, predominant species of our communities, were more abundant in 2003 than in 2002 and 2004, but this difference may have been as result of variation in weed availability. The extended rain frequency in 2003 delayed weed control at two of three locations in 2003 and, considering that these species feed on grass seeds (Larochelle & Larivière, 2003), it is possible that weed seed was more abundant in 2003 as a result of limited herbicide use. Despite their



**Figure 3** Effect of trap recovered per field on number of individuals collected (mean pooled per field of Bt and non-Bt cotton during 2002–04).

abundance in our fields, little is known about the life histories of *S. palliatus* and *H. gravis*, and they could be favoured by the high soil moisture resulting from extended rainfall in 2003 and the greater abundance of weeds that same year. In contrast, all dermapteran species were less abundant in 2003 compared with the other seasons, but this appears to have been because of the rotation of the third pair of fields (Table 1). The Ty Ty field and Frazier fields were major sources of *L. riparia* in 2002 and 2004, respectively, whereas the Old House field used in 2003 had relatively low dermapteran abundance. Abundance of predatory heteropterans per pitfall trap decreased over the season, and less evident variation was observed for araneids and dermapterans. Predatory heteropterans sampled are predominantly plant foragers, except *G. uliginosus*, which is predominantly epigeal (Crocker & Whitcomb, 1980). The decline in numbers of predatory heteropterans collected was consistent in Bt cotton fields but abundance was highly variable in non-Bt cotton fields (Fig. 1). This may be a response to the increased plant foliage area with seasonal progression, increasing the area for foliar foraging on both cottons and reducing activity on the ground, thereby lessening chance of capture by pitfall.

The greater variability of population dynamics in non-Bt cotton fields corresponded with insecticide use on plant foliage (i.e. the first weeks of July and August) (Table 1). Therefore, the variability in population dynamics of ground-dwelling taxa, such as heteropterans, araneids and dermapterans, that forage in the plant canopy and ground is likely because of the impact of foliar insecticide. Strict epigeal species, such as most carabids, cicindelids and staphylinids, would likely not be as readily affected by foliar insecticides (Table 1). Disregarding the in-furrow treatment to control thrips, which was made on all fields at planting, insecticides

were applied to control defoliating and sucking pests 22 times in non-Bt cotton versus 8 times in Bt cotton across all fields over 3 years (average of 2.4 applications per non-Bt field and 0.9 applications per Bt field; Table 1). The overall greater abundance of arthropods that might be expected in Bt fields as a result of lower insecticide pressure might have been masked somewhat in our study by use in non-Bt fields of a selective insecticide (spinosad; selective primarily for lepidopteran pests) that has short residual activity (Table 1). However, it is standard practice in the region to use selective insecticides as much as possible early in the season to conserve natural enemies. In addition, as mentioned above, most groups evaluated in this study forage predominantly on the ground; hence, foliar insecticides probably had limited impact. Given the reduced use of insecticides in Bt cotton and the greater use of selective insecticides in recent years, we would expect effective conservation of epigeal predators in cotton systems in the region.

The cicindelid *M. carolina* (Knisley & Schultz, 1997), the carabid *S. palliatus* (Ciegler, 2000), the dermapteran *L. riparia* (Hoffman, 1987) and the araneid *P. pauxilla* (Breene *et al.*, 1993) should be considered as significant species for monitoring of agroecosystem effects in the southeastern USA and for population comparisons representing important ground-dwelling taxa in modified cotton crop systems. All these species were frequently collected, are large and visually apparent, were consistently trapped in all locations and years and are taxonomically well defined.

The adoption of transgenic cotton has been considered detrimental for insect predators and parasitoids by directly eliminating prey/host availability or rendering prey/hosts unsuitable (reviews in O'Callaghan *et al.*, 2005). In this study, abundance and diversity of ground-dwelling arthropods (chiefly predators) in cotton fields were not affected during 3 successive years of growing transgenic Bt cotton using standard grower practices. The hypothesis elaborated for this study – that diversity and abundance of ground-dwelling predator arthropods would be reduced in Bt cotton – was not supported. Indeed, the use of transgenic cotton may generate positive changes in abundance and diversity of arthropods as a result of reduced applications of broad-spectrum insecticides in the Bt cotton ecosystem, making the system more salubrious for communities of ground-dwelling predators, although this difference was not observed in our low-insecticide cotton production system.

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