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Bacillus thuringiensis in caterpillars and associated materials collected from protected tropical forests in northwestern Costa Rica
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the growth of Bt populations (Aronson et al. 1986). On the other hand, some δ -endotoxins have not yet been associated with insecticidal action, but with other interesting biological activities (Misuki et al. 1999). Bt has been isolated from many sources, e.g., from soil (De Lucca et al. 1981, Ohba and Aizawa 1986), live and dead insects (Damgaard et al. 1998), stored grains (Meadows et al. 1992), foliar surfaces (Smith and Couche 1991, Misuki et al. 1999), water bodies (Ichimatzu et al. 2000), marine sediments (Maeda et al. 2000) and activated sludge (Misuki et al. 2001). The widespread distribution of apparently dormant Bt in nature (Martin and Travers 1989), the ecological roles it may play in soil ecosystems and the potential interactions of this microorganism with caterpillars in tropical forests are still poorly understood issues. In preliminary experiments (Sittenfeld et al. 2002), Bt has been isolated from midgut of Automeris zugana caterpillars collected in the Área de Conservación Guanacaste in northwestern Costa Rica (Janzen 2000). This study reports on the isolation of Bt from caterpillar guts, fecal pellets and host plants collected in different protected ecosystems (dry, rain and cloud tropical forests) as a contribution to the better understanding of Bt ecology in relation to lepidopterans in wildland environments.

MATERIALS AND METHODS

Sampling. Caterpillars and their food plants were collected through random search in trees and bushes located in tropical dry forests (Santa Rosa and Los Almendros at 200-300 masl), a cloud forest (Cacao at 1000 masl) and a rain forest (San Cristóbal at 500-600 masl) in the Área de Conservación Guanacaste (ACG). All caterpillars were separately put into a new plastic bag *in situ* with plenty of foliage from the same plant where the specimen was found. Bags were transported by ground to rearing barns in the Administration Area in the sector of Santa Rosa in the ACG and hung for 24-48 hours in order to simultaneously evaluate

the viability of the individuals collected and to obtain larval excrement at the bottom of each bag. All caterpillars appeared active and healthy at the moment of collection and their feeding activity and survival remained constant during the 48 hours they were kept in captivity before sacrifice. Each caterpillar collected from a single host plant was treated as an individual set of gut and fecal material.

Foliar material and larval excrement cultures. Approximately 3-5 leaves and 1-3 g of fecal pellets from each bag were separately inoculated in 10 ml of Luria Bertani (LB) broth. Leaves were aseptically sectioned into ca. 0.25 cm² pieces prior to inoculation, and fecal pellets were macerated inside the tube and homogenized to increase contact of the material with the culture media.

Larval dissection and gut culture. Caterpillars were killed by placing them separately for 15-30 min at –20°C. The external surface was disinfected through sequential washes with 0.2 μm filter-sterilized 70% Ethanol (2 min), a mix of 5.25% sodium hypochloride and 0.1% Tween 20 (2 min) and several washes with sterile distilled water. As a control for this process, each disinfected caterpillar was rolled on the surface of a LB agar plate, which was later incubated at 30°C for five days. Afterwards, the gut from each caterpillar was dissected aseptically in a laminar flow hood, macerated and inoculated along with its contents in 10 ml of LB broth

Bt identification. After incubation in an aerobic atmosphere at 30°C for 48 h, a loopful from each broth was streaked on LB agar plates. Based on colony morphology, a single presumptive Bt colony from each plate was selected and obtained in pure culture on Bacillus cereus selective agar plates (Holbrook and Anderson 1980). As a definitive criteria for identification, sporulated cultures of presumptive Bt isolates were examined by phase contrast microscopy for the synthesis of crystalline parasporal inclusions. The occurrence of Bt isolates was

analyzed considering each type of material as separate (according to sampling site) and in systems defined by the food plant leaves, gut and fecal pellets associated with each caterpillar collected (*Bt*-harboring systems).

Crystal inclusions morphology. Smears from sporulated cultures on LB plates were stained for 3-5 min with Coomassie Blue (3.75% in HOAc-EtOH 95%) and observed under light microscopy (100X). The morphology of the crystals synthesized by each isolate was recorded after at least three observations on different days.

RESULTS

A total of 37 caterpillars and 27 samples of food plants representing 16 Lepidoptera species from five insect families (Nymphaliidae, Hesperiidae, Arctiidae, Sphingidae and Notodontidae) and 15 species of food plants from 10 plant families (Simaroubaceae, Asclepiadaceae, Malphigiaceae, Fabaceae, Myrtaceae, Rubiaceae, Poaceae, Melastomataceae, Clusiaceae and Araliacea) were collected (Table 1).

As confirmed by the presence of parasporal crystal inclusions and the phenotypic reaction on B. cereus selective agar plates, at least one Bt isolate was cultured from 25 (24%) out of 101 samples analyzed. They were isolated from all sample types independently of their origin within the ACG, but mostly from food plant foliage (56%) and fecal pellets (36%) (Table 1). Further differences were detected in the number and percentages of isolates cultured from leaves and caterpillar guts collected in different ecosystems: it was more frequently cultured from foliar material collected from San Cristóbal and Cacao and from guts of caterpillars collected at San Cristóbal and Los Almendros (Table 2).

Bt was isolated from at least one material in 38% (n=14) of the systems constituted by the food plant, gut and fecal pellets corresponding to a single caterpillar. Within a system, it was

more frequently cultured from both the foliar material and fecal pellets (64%) than from gut samples (3%) (Table 1).

Four different morphologies of crystalline inclusions were observed (oval, irregular, bypiramidal and squared), with bipyramidal and irregular crystal morphologies being the most prevalent (20 and 44% respectively). Nine isolates (36%) exhibited a combination of at least two different crystal morphologies (results not shown). Moreover, *Bt* was cultured from the leaves naturally consumed by 3 of 5 monarch caterpillars collected, and in some cases also from their guts or fecal pellets (Table 1).

DISCUSSION

These results are in accordance with reports where foliar surfaces and insect-favoring habitats have been indicated as frequent sources of *Bt* isolation (Ben-Dov *et al.* 1997, Bravo *et al.* 1998, Arrieta *et al.* 2004). However, this is the first report of *Bt* isolation from fecal pellets of caterpillars collected in tropical protected environments.

The results obtained indicate that non-Btsusceptible caterpillars and possibly Bt-susceptible caterpillars, contribute directly to the dispersion of this bacterium in the ACG forests. The presence of Bt in fecal pellets of caterpillars (Tables 1 and 2), which fall or are washed to the foliage and forest floor below, could allow this bacterium to recirculate in these ecosystems (i.e., from the soil, to the foliage, passed through an insect, and back down to the forest floor), making it part of the horizontally transmitted caterpillar-based microbial community. It is plausible to speculate that the Bt population is thinly distributed throughout the habitat, "waiting" to be eaten by a caterpillar (or other insect) in which it will multiply, either lethally or benignly.

The different spatial distribution of Bt on the leaves and the fact that it is not possible to collect the same leaf material that the caterpillar has already cut for ingestion, may partially explain why Bt was not successfully isolated

TABLE 1
Isolation of Bacillus thuringiensis from food plants and caterpillar-derived materials collected in the Area de Conservación Guanacaste

Code ^a	Collection Site (forest type) b,c	Caterpillar	Food Plant	Bt from leaves	Bt from guts	Bt from fecal pellets
00-SRNP-114	SC (RF)	Bungalotis diaphorus	Simarouba amara	-	-	-
00-SRNP-424	SC (RF)	Danaus plexippus Asclepias curassavica		+	+	-
00-SRNP-425	SC (RF)	Danaus plexippus	Asclepias curassavica	+	-	-
00-SRNP-426	SC (RF)	Danaus plexippus	Asclepias curassavica	-	-	-
00-SRNP-427	SC (RF)	Enosis 00-SRNP-427 a	Poaceae DHJ14172 a	+	-	+
00-SRNP-428	SC (RF)	Enosis 00-SRNP-427 a	Poaceae DHJ14172 a	-	-	-
00-SRNP-429	SC (RF)	Enosis 00-SRNP-427 a	Poaceae DHJ14172 a	+	-	+
00-SRNP-437	SC (RF)	Danaus plexippus	Asclepias curassavica	+	-	+
00-SRNP-438	SC (RF)	Danaus plexippus	Asclepias curassavica	-	-	-
00-SRNP-544	SC (RF)	Cephise nuspesez	Hiraea sp.	+	-	+
00-SRNP-545	SC (RF)	Cephise nuspesez	Hiraea sp.	+	-	-
00-SRNP-546	SC (RF)	Morpho peleides	Lonchocarpus oliganthus	-	-	-
00-SRNP-547	SC (RF)	Parelbella macleannani	Eugenia basilaris	-	-	-
00-SRNP-548.01	SC (RF)	Actinote leucomel	Mikania DHJ14177 a	-	-	-
00-SRNP-548.02	SC (RF)	Actinote leucomel	Mikania DHJ14177 a	-	-	-
00-SRNP-548.03	SC (RF)	Actinote leucomel	Mikania DHJ14177 a	-	-	-
00-SRNP-548.04	SC (RF)	Actinote leucomel	Mikania DHJ14177 a	-	-	-
00-SRNP-548.05	SC (RF)	Actinote leucomel	Mikania DHJ14177 a	-	-	-
00-SRNP-548.06	SC (RF)	Actinote leucomel	Mikania DHJ14177 a	-	-	-
00-SRNP-548.07	SC (RF)	Actinote leucomel	Mikania DHJ14177 a	-	-	-
00-SRNP-548.08	SC (RF)	Actinote leucomel	Mikania DHJ14177 a	-	-	-
00-SRNP-548.09	SC (RF)	Actinote leucomel	Mikania DHJ14177 a	-	-	-
00-SRNP-548.10	SC (RF)	Actinote leucomel	Mikania DHJ14177 a	-	-	-
00-SRNP-548.11	SC (RF)	Actinote leucomel	Mikania DHJ14177 a	-	-	-
00-SRNP-2339	LA (DF)	Aellopos fadus	Genipa americana	+	-	+
00-SRNP-2340	LA (DF)	Urbanus proteus	Centrosema sagittatum	+	-	+
00-SRNP-2341	LA (DF)	Urbanus proteus	Centrosema sagittatum	-	-	-
00-SRNP-2343	LA (DF)	Notodontidae 00-SRNP-2343 ^a	Psychotria valeriana	-	+	-
00-SRNP-2344	LA (DF)	Bungalotis erythus	Schefflera morototoni	-	-	-
00-SRNP-2348	LA (DF)	Chioides cattilus	Rhynchosia calycosa	+	-	+
00-SRNP-6102	SR (DF)	Bungalotis quadratum	Inga vera	-	-	-
00-SRNP-6179	SR (DF)	Telemiades fides	Inga vera	-	-	-
00-SRNP-6228	SR (DF)	Telemiades fides	Inga vera	+	-	+
00-SRNP-9031	C (CF)	Coloradia marathusa	Miconia lonchophylla	+	-	-
00-SRNP-9032	C (CF)	Coloradia marathusa	Miconia lonchophylla	+	-	-
99-SRNP-17124	C (CF)	Pyrrhopyge cosyra	Chrysochlamys glauca	+	-	+

a Janzen and Hallwachs, 2002. b SC: San Cristóbal, LA: Los Almendros, SR: Santa Rosa, C: Cacao. c DF: Dry forest, CF: Cloud forest, RF: Rain forest.

TABLE 2
Occurrence of Bt on food plants, caterpillar guts and fecal pellets among dry, cloud and rain forests located at the ACG

	Rain Forest (San Cristóbal)		Dry Forest (Los Almendros)		Dry Forest (Santa Rosa)		Cloud Forest (Cacao)	
	Bt isolates	Samples analyzed	Bt isolates	Samples analyzed	Bt isolates	Samples analyzed	Bt isolates	Samples analyzed
Food plants	7 (50%)	14	3 (43%)	7	1 (33%)	3	3 (100%)	3
Guts	1 (4%)	24	1 (15%)	7	0 (0%)	3	0 (0%)	3
Fecal pellets	4 (17%)	24	3 (43%)	7	1 (33%)	3	1 (33%)	3

from all of the three materials of the system constituted by the food plant, gut and fecal pellets corresponding to a single caterpillar. A similar observation applies for the fecal pellets; in this case the sample used for culturing represented only a portion from both the leaf material digested and the total number of fecal pellets excreted during the time the caterpillars remained in captivity.

The low recovery of *Bt* from gut samples in spite of a higher prevalence in fecal pellet cultures is puzzling. This could be explained by intraluminal competition from other bacterial species, which could somehow keep the *Bt* population low until it passed from the gut to the fecal pellets (Takatsuka and Kunimi 2000), by abiotic inhibitory effects within the gut cavity (Luthy *et al.* 1985, Pang and Grigorten 1998), by simple dormancy of the *Bt* population while passing through the gut, or by an accumulation of *Bt* spores in fecal pellets due to the high speed of transit.

Since molecular-based methods were not considered, the observed frequency of Bt isolates could have been underestimated. In addition, individuals affected by the action of the delta endotoxins were not detected during the search for caterpillars in the foliage and could have caused a misrepresentation of Bt in the samples. In regard to the dynamics of Bt within insect guts, it is still necessary to conduct research in regards to the transit rate of Bt and quantitatively estimate its density and

metabolic stage (UFC/g of gut or UFC/cm² of vegetative cells or endospores) on the different materials through time.

All *Bt* isolates derived from a *Bt*-harboring system synthesized crystals of the same morphology, suggesting that the isolates recovered from the guts or fecal pellets are similar to the ones ingested on the leaves. In addition, the coexistence of more than one crystal type within a strain could have consequences on the constitution and stability dynamics of insect populations, and could eventually be a factor in the appearance of *Bt*-resistant insects.

The use of insect-protected transgenic crops incorporating Bt toxins has provided effective tools for the control of a variety of insect pests. However, populations of non-target insect species belonging to the same order as the target pest might be affected as well. Initial concerns about the effects of pollen from Bt transgenic crops on the larvae of the monarch (Danaus plexippus) and black swallowtail (Papilio polyxenes) butterflies have raised a controversy about the use of this kind of insecticide (Losey 1999, Maeda et al. 2000). In this study, the presence of Bt in tropical protected forests in the ACG, located over a hundred kilometers from areas where Bt is sprayed on crop fields (at present, no transgenic crops are authorized for commercial purposes in Costa Rica) was demonstrated. The apparent lack of toxicity of these isolates to their host caterpillars (collected alive and active) might be explained

by either a low prevalence of *cryI* genes in these wild *Bt* populations, a natural resistance to *Bt*, or by a combination of both. The effects of *Bt*-based pesticides on the survival and physiological traits of caterpillars (such as weight gain) have not been previously studied under field conditions in wildlands of tropical environments, and it remains to be determined, as *Bt* might be one of the many natural mechanisms that maintains a delicate balance among insect populations in tropical forests.

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RESUMEN

Bacillus thuringiensis (Bt) sintetiza inclusiones cristalinas que resultan tóxicas para algunas larvas de lepidópteros y otros órdenes de invertebrados. Su presencia fue examinada en materiales asociados a 37 orugas de mariposas de 16 especies, las cuales fueron colectadas mientras se alimentaban en 15 especies diferentes de plantas hospederas en bosques secos, nubosos y húmedos localizados dentro del Área de Conservación Guanacaste (ACG) en el noroeste de Costa Rica. A partir de un total de 101 muestras se obtuvo 25 aislamientos de Bt: 56% a partir de material foliar de las plantas hospederas, 8% a partir del contenido intestinal de las larvas y 36% a partir de sus excrementos. Esta bacteria fue cultivada a partir de al menos uno de los 3 diferentes tipos de muestra asociados a una oruga particular (planta hospedera, intestino, excremento) en 38% de los casos posibles. En la colección de aislamientos obtenida se observaron cuatro morfologías de inclusiones cristalinas, siendo aquellas bipiramidales e irregulares las más prevalentes.

Palabras clave: *Bacillus thuringiensis*, larvas de mariposa, plantas hospederas, tractos intestinales de insectos, material fecal, áreas protegidas.

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