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Original Research Article

Transgenic corn decreased total and key storage and lipid transport protein levels in honey bee hemolymph while seed treatment with imidacloprid reduced lipophorin levels

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Introduction

In addition to natural seasonal variations that directly interfere with the availability of nectar and pollen for bees, anthropogenic effects on ecosystems and agroecosystems can result in unfavorable conditions for nutritional subsistence due to intensification of food shortage periods; this includes extensive monoculture plantings with floral features that do not meet the nutritional demand of bees (Cock et al., 2013; Winston, 1987). The use of artificial diets is increasing worldwide

Pollen substitutes are currently widely used in apiculture because of a lack of natural forage and an increasing need for bees for pollination services. A major concern in the confection of bee diets is whether specific ingredients, such as GMO grain, or contaminants such as pesticides that are applied to crops, negatively impact bee health. We measured hemolymph protein levels and the lifespan of honey bees fed on diets containing non-hybrid (IAC-Airan), hybrid (AG 7088) or transgenic (AG 8088YG) maize, the seeds of which had been treated with fipronil, imidacloprid or left untreated. Corn from these plantings was harvested, dried and finely ground, and then used for the preparation of corn-based diets. Groups of 100 newly emerged bees were placed in cages, where the diets were offered ad libitum. Honey bee lifespan was similar for all the corn diets, though shorter than for bees fed bee bread and longer than for bees fed only honey. Total protein in the hemolymph of bees fed with transgenic maize was 15% lower than in bees fed on non-hybrid maize; lipophorin and vitellogenin, were more reduced, by over 30%. Irrespective of corn variety, imidacloprid resulted in over 25% lower levels of lipophorin compared with the control. Lower storage and transport protein levels would compromise the ability of the bees to rear brood and develop an adequate immune response to pathogen challenge. Consequently, transgenic maize, as well as any maize produced from seed treated with imidacloprid should not be included as ingredients in honey bee diets.

Maiz transgénico disminuye niveles de proteínas claves de almacenamiento y transporte de lípidos en hemolinfa de abejas melíferas así como el tratamiento de semillas con imidacloprid reduce los niveles de lipoforina

Substitutos del polen actualmente son utilizados en larga escala en la apicultura por causa de la falta de alimento natural y un aumento en la necesidad de abejas para servicios de polinización. Una preocupación importante en la confección de dietas para las abejas es si ingredientes específicos, como granos OGM, o contaminantes como plaguicidas aplicados a los cultivos, impactan negativamente sobre la salud de la abeja. Nosotros medimos los niveles de proteína en la hemolinfa y el tiempo de vida de abejas melíferas alimentadas con dietas conteniendo maíz no-híbrido (IAC-Airan), híbrido (AG 7088) o transgénico (AG 8088YG); durante la siembra de estos cultivos las semillas fueron tratadas con fipronil, imidacloprid o sin ningún tratamiento. Muestra de maíz de estas plantaciones fueron cosechadas, secadas, molidas y después utilizadas para preparar las dietas basadas en maíz. Grupos de 100 abejas recién emergidas fueron colocadas en jaulas y las dietas fueron ofrecidas a libre consumo. El tiempo de vida de las abejas fue similar para todas las dietas con maíz, pero más corto del que para abejas alimentadas con pan de la abeja y mayor en comparación con abejas alimentadas solamente con miel. Los niveles de proteína total en la hemolinfa de las abejas alimentadas con maíz transgénico fue 15% menor en comparación con abejas alimentadas con maíz no-híbrido; lipoforina y vitellogenina presentaron más reducción, cerca de 30%. Independientemente de la variedad de maíz, imidacloprid resultó en más de 25% de la reducción de lipoforina, comparada con el control. Niveles menores de proteínas de almacenamiento y transporte pueden comprometer la habilidad de las abejas en producir cría y desarrollar una respuesta inmunológica adecuada cuando confrontada con un desafío de patógenos. Por lo tanto, maíz transgénico y cualquier maíz producido utilizando semillas tratadas con imidacloprid no deben ser utilizados como ingredientes en dietas de abejas melíferas.

Keywords: Apis mellifera; artificial diet; fipronil; genetically modified organism; neonicotinoid; vitellogenin
in order to maintain the bees and to prepare them for 
honey production and pollination services (Brodtschnei-
der & Crailsheim, 2010; Tomlinson, Dixon, Didham, & 
Bradshaw, 2017).

Although nutritional requirements of honey bees for 
carbohydrates, proteins, lipids, vitamins, minerals and 
water are well known (Haydak, 1970; Herbert, 1992; 
Huang, 2010), artificial diets are often empirically formu-
lated to provide energy and protein requirements for 
honey bee colony maintenance and development. Bee-
keepers have traditionally used refined sucrose, high 
fructose corn syrup and soy-based products (Somerville, 
2005). The choice of ingredients or substitutes is contin-
gent on availability and prices.

In Brazil, approximately 80% of all cultivated areas are 
occupied with soybean, maize and sugarcane crops (IBGE, 
2016). These crops are not utilized for honey and pollen 
production (Knopper, Dan, Reisig, Johnson, & Bowers, 
2016). Nevertheless, artificial diets for bees are prepared 
using the products of these crops. All three cultures are 
susceptible to specific insect pests, such as leafhoppers, 
termites and borers, which are controlled with insecti-
cides such as fipronil and imidacloprid (Andreí, 2013; 
Simon-Delso et al., 2015). These persistent and systemic 
insecticides are especially toxic for bees, manifesting 
lethal and sublethal effects (Nicodemo et al., 2014). Bees 
can be directly affected by insecticide product application 
to plantings when they feed on floral resources of treated 
crops or even when they are fed diets formulated with 
products obtained from these crops (Decourtye et al., 
2004; El Hassani, Dacher, Gauthier, & Armengaud, 2005; 
Iwasa, Motoyama, Ambrose, & Roe, 2004; Johnson, Ellis, 
Mullin, & Frazier, 2010; Mayer & Lunden, 1999; Roat 
et al., 2013).

Maize and soybean are the main genetically modified 
(GMO) crops used worldwide. Herbicide tolerance and 
insect resistance are the principle traits in these GMO 
crops (Qaim, 2009). There is evidence that nectar, pol-
len and products obtained from GMO crops can be 
harmful to bee health (Han, Niu, Lei, Cui, & Desneux, 
2010a; de Maagd, Bravo, & Crickmore, 2001; Ramirez-
Romero, Desneux, Decourtye, Chaffiol, & Pham-
Delégué, 2008; Sabugosa-Madeira, Abreu, Ribeiro, & 
Cunha, 2007). If floral resources of transgenic plants 
negatively impact the health of bees, diets formulated 
with ingredients obtained from these plants also are 
potentially harmful.

We examined whether honey bee diets based on 
transgenic vs. non-transgenic maize treated or not with 
imidacloprid or fipronil at sowing would affect bees by 
measuring their hemolymph proteins and lifespan.

Material and methods

Maize seeding and insecticide application

Maize seeds of non-hybrid (IAC-Airan), hybrid 
(AG7088) and transgenic (AG8088YG) cultivars were 
sown in stands of 55,000 plants per ha with fertilization 
conducted according to Raij, Cantarella, Quaggio, and 
Furlani (1997) in Dracena, Brazil (21°27′37″S latitude, 
51°33′21″W longitude, and 421 m altitude).

The three cultivars were grown in 300 m² fields that 
were at least 600 m away from other maize cultivar 
fields. Each area of a single cultivar was subdivided into 
three plots of 100 m² for insecticide application.

In the first plot for each area, application of a com-
mercial product that included fipronil as its active ingre-
dient (a.i.) was conducted during furrow planting using 
80 g ha⁻¹ of a.i. diluted in water for the application of 
250 L ha⁻¹ of this solution, according to the manufac-
ther’s suggestion for the control of Diabrotica speciosa 
(Coleoptera: Chrysomelidae) (Melhorança et al., 2010; 
Regent 800 WG, 2002).

For the second plot for each area with the same maize 
cultivar, seeds were treated with an insecticide that 
included imidacloprid as its a.i. Before sowing, seeds were 
treated with a proportion of 490 g of a.i. 100 kg⁻¹ of 
seeds to control D. speciosa (Gaucho, 1996; Melhorança 
et al., 2010). In the third plot for each area, no insecticides 
were used. No other pesticides were used during the 
development of the maize plants in the three areas.

Diets and feeding

Nine maize-based diets were prepared (three maize cul-
vivars × two insecticides and a control). Diets consisted 
of 8 parts very finely ground cornmeal and 2 parts 
organic honey. To compare the tested diets with natural 
bee feeding conditions, 8 parts beebread and 2 parts 
organic honey were used to prepare a positive control 
diet. For the protein-negative control diet, only organic 
honey was fed. Diets were offered ad libitum.

Bromatological analysis

Crude protein (CP) content of dietary ingredients was 
determined and expressed in dry matter (DM), with 
three replicates for each ingredient (Association of Offi-
cial Analytical Chemists [AOAC], 1990; Silva & Queiroz, 
2006).

Honey bees

Brood combs were obtained from a single honey bee 
colony housed in a standard Langstroth hive, containing 
sealed brood ready to emerge. This hive was fed previ-
ously for two months with 50% (w/v) sucrose syrup to 
stimulate brood production. Each comb was placed in a 
screened cage and stored in an incubator at 33 °C with 
70% relative humidity for 36 h. Under the same incuba-
tor conditions, each plastic screened cage was stocked 
with 100 newly emerged adult worker honey bees. The 
experiments were repeated three times.
Hemolymph protein levels
After adult workers emerged, a sample of 10 bees was collected from each cage to determine the concentration of proteins in the pool of bee hemolymph on the day of emergence (day 0) and another 10 bees on day 7.

Hemolymph was collected with the aid of micropipettes inserted into a small cut made at the base of the bee's wing with microscissors and transferred to micro-centrifuge tubes with phenylthiocarbamide (PTC) and protease inhibitor, using microcapillary tubes previously washed in a 0.1% (wt/vol) phenylthiourea solution in water.). Hemolymph samples were centrifuged at 1,500 rpm for 4 min at 4 °C. The supernatant was removed and frozen at 20 °C. Total protein quantification was performed by the Bradford method (1976). Quantification of the samples was made using a Beckman Coulter DXT-880-multimode detector spectrophotometer at 595 nm with three replicates, and a standard curve was constructed using bovine serum albumin (Cremonez, De Jong, & Bitondi, 1998).

Analysis of hemolymph protein profiles
For evaluation of newly emerged (day 0) and seven-day-old honey bee hemolymph protein profiles, 1 μl of hemolymph was dissolved in sample buffer with 0.25 M Tris-HCl, pH 6.8, containing 70% sucrose, 0.25% sodium dodecyl sulfate (SDS), 0.1% bromophenol blue and 5% β-mercaptoethanol. Samples were submitted to thermal denaturation, and after cooling, they were applied to separation gels with 7.5% polyacrylamide (Laemmli, 1970).

The proteins myosin (205 kDa), β-galactosidase (116 kDa), phosphorylase b (97), BSA (66 kDa), ovalbumin (45 kDa) and carbonic anhydrase (29 kDa) were used as molecular mass markers. For quantitative evaluation of lipophorin and vitellogenin proteins, SDS-PAGE gels were scanned and individual profiles of each sample were analyzed by densitometry using Adobe Photoshop 7.0 Software, with three replicates. Data was represented as arbitrary optical density units (D.U.).

Evaluation of longevity
One hundred bees per cage were placed in an incubator with controlled temperature (33 °C) and humidity (70%). At day 0, 10 g of protein food was offered to the bees in feeders inside the plastic pots. This diet was renewed every two days. Once a day, dead bees were counted and removed until all the bees had died.

Statistical analysis
Data were submitted to statistical analysis and the effects were considered significant at p ≤ 0.05. Two models were applied to data. The first model comprises the 11 treatments for comparisons with controls (beebread and honey based-diets), as described below:

$$ y_{ij} = \mu + \alpha_i + \epsilon_{ij}, $$

where $y_{ij}$ = response variable of bees fed diets (i) in replication (j), $\mu$ = overall mean values for $y$, $\alpha_i$ = fixed effect of diets, $\epsilon_{ij}$ = error term.

The second model was used to study the interactions between the maize type and insecticide, except for longevity evaluation, as follows:

$$ y_{ijk} = \mu + \alpha_i + \beta_j + (\alpha \times \beta)_{ij} + \epsilon_{ijk}, $$

where $y_{ijk}$ = response variable of bees fed maize types (i) and submitted to different insecticides (j) in replication (k). $\mu$ = overall mean values for $y$, $\alpha_i$ = fixed effect of diets, $\beta_j$ = fixed effect of insecticide. $(\alpha \times \beta)_{ij}$ = interaction of maize type with insecticide. $\epsilon_{ijk}$ = error term.

All response variables were submitted to analysis of variance, and the Tukey test was used for multiple comparisons. To evaluate the longevity of the bees, the Kaplan-Meier non-parametric method was used. Data were processed with SAS software (SAS Institute, 2003).

Results
In the analysis of the 11 dietary ingredients, beebread (28.8% ± 0.7) and honey (0.4% ± 0.03) presented the highest and lowest protein levels ($p < 0.05$), respectively. In the evaluation of maize types, treated with insecticides fipronil and imidacloprid or left untreated, no difference was observed for crude protein content, with a general average of 10.5% ± 0.1 (Table 1).

Total protein concentration in the hemolymph of adult bees on the day that they emerged (day 0) was 16.0 ± 2.1 μg/l. From day 0 to day 7, bees were fed experimental diets. Based on the analyses conducted at the end of the 7-day period, we observed that the highest value ($p < 0.05$) of total proteins in the hemolymph (22.6 ± 2.3 μg/μl) was obtained when beebread was offered to the bees. With the honey-based diet, the mean value of total proteins in the hemolymph (5.5 ± 0.7 μg/μl) did not differ significantly from that of bees fed on any of the corn-based diets, which additionally did not differ from each other ($p > 0.20$).

Considering only the maize-based diets, the possible interactions and the main effects were studied. No difference was found regarding the interactions between maize types and insecticides (Table 2). For the main effect type of insecticide, we found that whether insecticide was used or not used did not influence hemolymph protein concentration for bees fed on maize-based diets. Concerning maize types only, we observed that total hemolymph protein concentration for the bees was lower ($p < 0.05$) when the transgenic cultivar was used compared with the non-hybrid cultivar. Non-transgenic hybrid maize did not differ from the other two corn cultivars ($p > 0.20$).

Through qualitative analysis of the polyacrylamide gels, we found that the highest protein levels occurred...
at day 0. After seven days of the bees being fed the experimental diets, the diet consisting of beebread allowed the bees to obtain the highest protein density profile. On the other hand, along with it being the diet containing the lowest protein content, the honey-based diet provided the lowest protein profile density in the hemolymph among all the diets.

On day 0, the densitometric quantification of lipophorin present in the hemolymph of the bees was 233.7 ± 3.4 D.U. Analyzing the 11 diets on day 7, the concentration of this protein was higher \((p < 0.05)\) when bees were fed a beebread diet (211.1 ± 3.6 D.U.) than when they were fed a honey-based diet (108.0 ± 6.1 D.U.). No difference was observed \((p > 0.20)\) for lipophorin levels of bees fed on the nine maize-based diets.

No effect of interactions between maize types and insecticides was observed for lipophorin levels. Regarding only the maize type effect, lipophorin levels were lower \((p < 0.05)\) in beebread-fed bees compared to those fed on non-hybrid maize.

No effect of interactions between maize types and insecticides was observed for vitellogenin levels. Regarding only the maize type effect, vitellogenin was lower \((p < 0.05)\) in beebread-fed bees compared to those fed on non-hybrid maize.

The densitometric quantification of vitellogenin in the hemolymph of newly emerged bees was 79.2 ± 4.8 D.U. For the analyses on day 7, the concentration of vitellogenin was higher when bees were fed a beebread diet \((162.4 ± 5.4 \text{ D.U.})\), while the lowest concentration values were observed with honey-based diets \((23.6 ± 1.0 \text{ D.U.})\).

No effect of maize and insecticide interactions on vitellogenin protein levels was observed. The main effect type of insecticide also did not promote differences in vitellogenin protein levels \((p > 0.20)\). However, in the analysis of the maize types, the concentration of vitellogenin was lower \((p < 0.05)\) in beebread-fed bees compared to non-hybrid maize. The concentration of vitellogenin found in the hemolymph of bees fed on non-transgenic hybrid maize did not differ from the concentration of vitellogenin found in the hemolymph of bees fed on non-hybrid and transgenic maize-based diets \((p > 0.20)\).

Longevity increased for bees fed on beebread and decreased for bees fed on honey \((p > 0.20)\). No difference \((p > 0.20)\) was observed between bees fed on maize-based diets that were obtained from transgenic, hybrid and non-hybrid corn treated with fipronil or imidacloprid or left untreated.

### Discussion

We did not find a superiority of crude protein content for any maize cultivar. Pollen, the common source of protein in bee food, has 2–4 times more protein than corn \((\text{Herbert et al., 1978})\). Therefore, it is not possible to fully replace pollen with maize as the only ingredient in a diet. However, maize can be used in the composition of bee diets because it is easily found on

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*Results are expressed as the mean ± standard error of the mean. Means followed by different letters differ from each other \((p < 0.05)\) as indicated by the Tukey test.

**Table 1.** Crude protein content as a percentage in dry matter of the maize ingredients used in experimental diets for honey bees produced from seeds that had been treated or not with insecticide at planting.

<table>
<thead>
<tr>
<th>Maize types</th>
<th>No insecticide</th>
<th>Fipronil</th>
<th>Imidacloprid</th>
<th>Overall average</th>
</tr>
</thead>
<tbody>
<tr>
<td>Non-hybrid</td>
<td>10.2 ± 0.3 b²</td>
<td>10.1 ± 0.2 b</td>
<td>10.7 ± 0.2 b</td>
<td>10.3 ± 0.2</td>
</tr>
<tr>
<td>Hybrid</td>
<td>10.6 ± 0.1 b</td>
<td>10.8 ± 0.2 b</td>
<td>10.9 ± 0.5 b</td>
<td>10.8 ± 0.2</td>
</tr>
<tr>
<td>Transgenic</td>
<td>10.4 ± 0.2 b</td>
<td>10.0 ± 0.7 b</td>
<td>11.1 ± 0.1 b</td>
<td>10.4 ± 0.3</td>
</tr>
<tr>
<td>Overall</td>
<td>10.4 ± 0.1 b</td>
<td>10.2 ± 0.2 b</td>
<td>10.9 ± 0.2 b</td>
<td>10.5 ± 0.1 b</td>
</tr>
</tbody>
</table>

*Results are expressed as the mean ± standard error of the mean. Means followed by different letters differ from each other \((p < 0.05)\) as indicated by the Tukey test.

**Table 2.** Concentration of total proteins \((\mu g/\mu l)\) in the hemolymph of seven-day-old adult honey bees fed from day 0 with corn-based diets derived from corn plants treated with fipronil or imidacloprid or left untreated at sowing.

<table>
<thead>
<tr>
<th>Maize types</th>
<th>Without insecticide</th>
<th>Fipronil</th>
<th>Imidacloprid</th>
<th>Overall average</th>
</tr>
</thead>
<tbody>
<tr>
<td>Non-hybrid</td>
<td>7.9 ± 0.6</td>
<td>8.2 ± 0.8</td>
<td>7.3 ± 0.7</td>
<td>7.8 ± 0.4 a²</td>
</tr>
<tr>
<td>Hybrid</td>
<td>6.7 ± 0.1</td>
<td>7.1 ± 0.5</td>
<td>7.9 ± 0.7</td>
<td>7.2 ± 0.3 ab</td>
</tr>
<tr>
<td>Transgenic</td>
<td>6.6 ± 0.6</td>
<td>6.7 ± 0.6</td>
<td>6.4 ± 0.3</td>
<td>6.6 ± 0.3 b</td>
</tr>
<tr>
<td>Overall</td>
<td>7.0 ± 0.3</td>
<td>7.3 ± 0.4</td>
<td>7.2 ± 0.4</td>
<td>7.2 ± 0.2</td>
</tr>
</tbody>
</table>

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the market and typically cheaper than other bee diet ingredients (favoring the development of lower cost diets). Maize has all the essential amino acids for honey bees except tryptophan (De Groot, 1953; Rostagno et al., 2005).

As expected, honey had the lowest protein content of all ingredients used in the diets; therefore, honey cannot be the sole ingredient of a honey bee diet if colonies have insufficient pollen (Somerville, 2005). However, this natural energetic food is important for honey bees when they feed their larvae (Ellis & Hayes, 2009; Herbert & Shimizu, 1978).

As holometabolous insects, the protein need for honey bees is higher for their brood compared with adult insects, although workers can benefit the health and development of the brood they feed by consuming protein (Crailsheim, 1990). Nurse bees need protein to complete the development of hypopharyngeal glands and produce worker or royal jelly (Alqarni, 2006; Maurizio, 1954; Pernal & Currie, 2000) and thoracic musculature (Hersch, Crewe, Hepburn, Thompson, & Savage, 1978).

At seven days old, bees should have high levels of protein in their hemolymph to appropriately feed larvae. Protein levels in bees fed the corn-based diets could have been affected by differences in consumption compared to the bee bread diet. We did not measure consumption; however, Cremonez et al. (1998) found similar consumption by caged bees of a bee bread compared to a corn meal diet. Protein levels in the hemolymph was similar for bees fed on all three varieties of corn, independent of whether the seeds had been treated or not with pesticide (Table 2). The protein titer in the honey bee hemolymph is related to the protein content of the diet (De Jong, da Silva, Kevan, & Atkinson, 2009). Cremonez et al. (1998) observed that honey bees fed on diets based on corn meal had a significant reduction of protein hemolymph titer compared to bees fed on diets based on bee bread, soybean and yeast or pollen, though it was superior to sucrose.

On the market, most maize seeds available for commercialization are transgenic hybrids, which are mainly used because of higher yields, rather than to improve grain quality (Wolfenbarger & Phifer, 2000). However, higher maize protein content, including higher lysine content via the development of transgenic maize varieties, can promote better nutritive value of this dietary ingredient (Tang et al., 2013). In our study, no differences were observed regarding protein content among maize cultivars. Therefore, we did not expect to observe differences in the protein titer of the honey bee hemolymph. As there was no difference between protein crude content in all evaluated maize based diets, we inferred that other factors (not protein content) affected the amount of protein in the hemolymph of the bees.

The YieldGard varieties, such as the transgenic maize cultivar AG8088YG that we evaluated, were developed to provide resistance to lepidopteran and coleopteran pests by producing Bt proteins known as Cry proteins. When these insects attack maize plants, their digestive enzymes activate the toxic form of this protein, which acts quickly to damage the larvae (Vidal, Kuhlmann, & Edwards, 2005). In a meta-analysis consisting of 25 studies that independently assessed potential effects of Bt Cry proteins on honey bee survival, Duan, Marvier, Huesing, Dively, and Huang (2008) did not observe any direct negative effects of Bt crops on honey bees.

Evaluation of lethal effects is very important; however, it cannot be exclusively used to determine whether any product impairs the health of an animal species. Sublethal effects should also be taken into account. In this case, the type of Cry toxin and the vegetable component used in honey bee feeding, such as pollen, nectar, grains, etc., must be considered on a case-by-case analysis (Malone & Pham-Dele`gue, 2001).

Lipophorin is a lipoprotein responsible for the transport of lipids between absorption sites in the intestine and deposition sites in body fat cells or for use in muscles and other tissues (Chino & Downer, 1982). Vitellogenin is a glycolipoprotein whose synthesis peaks when bees are 5–15 days old when the bees normally feed brood. In addition, vitellogenin influences a honey bee’s lifespan, immunity, zinc transport and larval food synthesis (Amdam et al., 2004, 2005; Amdam, Norberg, Hagen, & Omholt, 2003). The amount of vitellogenin present in hemolymph is influenced by the quantity and quality of food that bees have available (Bitondi & Simões, 1996). Concentrations of both of these aforementioned proteins were reduced by more than 30% transgenic maize types, Without insecticide

| Table 3. Densitometric quantification, measured in arbitrary optical density units (D.U.), of the band intensity corresponding to lipophorin on an SDS-PAGE gel of the hemolymph of seven-day-old honey bees fed from emergence with corn-based diets derived from corn plants treated with fipronil or imidacloprid or left untreated at sowing. |
|-----------------------------------------------|------------------|-----------------|------------------|
| Maize types                           | Without insecticide | Fipronil       | Imidacloprid     | Overall average |
| Non-hybrid                            | 188.3 ± 1.4       | 177.9 ± 2.2     | 169.8 ± 3.2      | 178.6 ± 2.9 A   |
| Hybrid                                | 198.6 ± 4.8       | 159.5 ± 8.8     | 109.0 ± 4.3      | 155.7 ± 13.3 A  |
| Transgenic                            | 147.2 ± 8.4       | 113.6 ± 3.3     | 107.6 ± 2.5      | 122.8 ± 6.7 B   |
| Overall average                        | 178.0 ± 8.3 a*    | 150.3 ± 10.0 ab | 128.8 ± 10.4 b   | 152.4 ± 6.6     |

*Results are expressed as the mean ± standard error of the mean. Means followed by different letters in the right column or in the bottom line differ from each other (p < 0.05) as indicated by the Tukey test.
Lipophorin concentration was over 25% lower in honey bees fed on maize whose seeds were treated with imidacloprid compared with honey bees fed on maize whose seeds were not treated with insecticides (Table 3). Imidacloprid acts as a neurotoxin and binds to the nicotinic acetylcholine receptor of insects (Boucias, Stokes, Storey, & Pendland, 1996). As it is a systemic insecticide, bees can consume contaminated food or have direct contact with contaminated food as a consequence of imidacloprid drift due to aerial application drift or mechanical stirring of contaminated soil (Greatti, Sabatini, Barbattini, Rossi, & Stravisi, 2003). Though the pesticides did not affect longevity, there is concern that accumulation in the soil after consecutive plantings could have a greater effect on bees.

Although artificial diets may be prepared from select ingredients to improve a colony’s health, AG8088YG (transgenic) maize must be avoided in addition to any other type of maize treated with imidacloprid at sowing. Additionally, an artificial diet must be prepared considering the nutritional value of the diet’s ingredients, as well as its genetic traits and cultivation mode with respect to the use of agrochemicals.

Conclusions
Maize should not be used as the only supplemental dietary protein ingredient for honey bee colonies.

The use of transgenic maize from cultivar AG8088YG as an ingredient of a protein diet reduced total hemolymph protein concentration, especially affecting the transport and storage proteins lipophorin and vitellogenin. Therefore, the use of transgenic maize for honey bee diets should be avoided.

Disclosure statement
No potential conflict of interest was reported by the authors.

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