



Impact of net houses on the natural regulation of the populations of *Trialeurodes vaporariorum* (Homoptera: Aleyrodidae) and *Tuta absoluta* (Lepidoptera: Gelechiidae), two major tomato pests in Kenya

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Abstract

The abandonment of insecticide treatments to achieve sustainable crop production calls for a combination of methods to obtain satisfactory pest control. To this end, net houses and biological control are two promising methods, though we suspected that most natural enemies are blocked outside the net houses. In Kenya, tomato crops are particularly threatened by *Trialeurodes vaporariorum* (Westwood 1856) and *Tuta absoluta* (Meyrick, 1917) that are targets of most insecticide treatments in this country. We investigated in an on-farm experiment (i) the abundances of *T. vaporariorum* and *T. absoluta* in open fields and in net houses, (ii) the diversity of arthropod natural enemies of these pests, and (iii) the effect of net houses on the natural regulation. To complete, in the laboratory, we checked the capacity of natural enemies to pass through different net types. We identified several natural enemies, mainly the mirid bug *Nesidiocoris tenuis* (Reuter 1895) that was the most abundant predator for both pests. One parasitoid species was also identified for each pest, i.e. *Encarsia formosa* Gahan (1924) and *Copidosoma* sp. for *T. vaporariorum* and *T. absoluta* respectively. Net houses reduced drastically the pest populations. Predators were less abundant under net houses, while parasitoids did not seem to be affected by the nets. *Encarsia formosa* was shown to be able to pass through the different net types, while *N. tenuis* adults were not. These results raise the potential for augmentative biological control under net houses. Using kairomones to attract natural enemies, or introducing natural enemies under the net houses through inoculative releases, are two approaches to increase the abundance of natural enemies under net houses.

Keywords *Nesidiocoris tenuis* · *Encarsia formosa* · Augmentative biological control · Agroecology

Introduction

Pest control methods can interact synergistically or antagonistically. Yet, to avoid pesticide treatments, several integrated agroecological control methods are required to achieve satisfactory control (Gurr and Kverdaras 2010;

Kungu et al. 2019). This could involve multi-tool control programs, e.g. physical crop protection (net houses) and biological control, as already implemented in European greenhouses (Calvo et al. 2012; Chailleux et al. 2012). However, studies are required to determine how this combination could be effectively applied in net houses in Africa, where pesticides are still heavily used (De Bon et al. 2014).

Tomato crops in Africa suffer massive damage caused by large caterpillars such as the cotton bollworm *Helicoverpa armigera* (Diatte et al. 2018). Net houses were successfully developed as a control method to protect tomato plants from such large pests (Achieng'a et al. 2016), yet one drawback has been noted—small arthropod pests are able to penetrate and thrive under net houses and destroy the crops if no insecticides are sprayed (Simon et al. 2014). We hypothesize that small pests escape the

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control of their natural enemies that might be too big to pass through the net (Majumdar et al. 2015). Moreover, larger pests sometimes also ultimately manage to invade the net houses through the entrances during the coming and going of personnel for agricultural practices (Gogo et al. 2014; Simon et al. 2014). Inside net houses, pests are mostly controlled using chemical insecticides, stressing the need for biological control methods adapted to this context (Nordey et al. 2017; Ramasamy and Ravishankar 2018). Some biological control techniques, such as augmentative biological control, would be facilitated by the net-confined environment.

Major pests threatening tomato production under net houses include the invasive tomato leafminer, *Tuta absoluta* (Meyrick 1917) (Lepidoptera: Gelechiidae) and the worldwide whitefly *Trialeurodes vaporariorum* (Westwood 1856) (Homoptera: Aleyrodidae). *Tuta absoluta* is a devastating tomato pest originating from South America (Guillemaud et al. 2015). Since 2004, it has rapidly spread and caused heavy tomato crop loss throughout the Mediterranean Basin (Desneux et al. 2010), Africa (Mansour et al. 2018; Rwomushana et al. 2019) and Asia (Han et al. 2019). *Trialeurodes vaporariorum* is a sucking pest native to Central and South America (Mound & Halsey 1978). As one of the most important crop pests and virus vectors (Jones 2003), it also causes substantial tomato crop loss in Kenya (Gamarra et al. 2016; Kimathi et al. 2020). Whiteflies are small-sized insects and are suspected to be able to pass through most nets, and *T. absoluta* females can oviposit on nets and subsequently the hatched larvae can produce a silk thread (Desneux et al. 2010) that they use to drop onto the tomato plants (AC personal observation). Many different chemical pesticides are applied under net houses in Kenya to control these two crop pests (Njuguna 2000; Nderitu et al. 2018).

Natural enemies are known for both species, particularly predatory mirid bugs for *T. absoluta* (Zappala et al. 2013), and parasitoid wasps for *T. vaporariorum* (Perić and Dimić 1997; Perić et al. 2008), yet very little information is available on the natural enemies of these pests in Kenya. Natural regulation protagonists and how they are affected by net protection is an essential information for the development of biological control in net houses. We hypothesize that several indigenous arthropod natural enemies are already present in Kenya for both pests, although that contrary to pests, they could be blocked outside the net house. We conducted a field experiment in Kenya to collect natural enemies and compare tomato crops in open fields and under net houses. Then we carried out laboratory tests to assess the net-crossing capacity of the natural enemies.

Material and methods

On farm-experiment

Study site and experimental design

In partnership with Kenyan farmers, a set up was established in 16 farms located in Kirinyaga, Kiambu, Machakos, Kajiado, and Migori in Kenya (Table 1). Farmers were asked to grow tomato plants both in their net house (0.9 mm mesh size, not treated net) and in an open field (at least 50 m² for both situations). However, not all the farmers accepted to grow open field tomato crops. Three monitoring sessions were conducted, one in the cool dry season (June to July 2017, 12 farms), one in the hot dry season (February 2017, three farms), and one in the short rainy season (November 2017, one farm). Indeed, many fields were affected by bacterial wilt during the hot season, preventing us from conducting more monitoring at this season. The monitoring session of each field depended on farmers' cropping season habits. Monitoring and sampling were conducted once per field. Tomato crop development at the time of our monitoring ranged from 2 weeks after transplanting to the fruiting stage.

Pest and natural enemy recording and identification

An entomological field aspirator (Improved Prokopack Aspirator—Model 1419 from John W. Hock company) was used to sample the predators and parasitoids on tomato plants. Ten entire plants, randomly selected, were sampled in each open field and each protected net house.

To record whitefly abundance and parasitism, three leaflets were sampled per plant (bottom, middle, and top of the plant) on the 10 plants in each open field and each protected net house. The number of whitefly nymphs was counted using a binocular microscope. Then, all leaflets were stuck into small pieces of humid floral foam in boxes (10 cm diameter, 9 cm deep) closed with a 160 µm mesh netting and stored at 25 °C to check for parasitoid emergence. Leaflets were kept for 2 weeks, with parasitoid emergence checked every 2 days.

To record *T. absoluta* abundance, three leaves per plant (bottom, middle and top of the plant) on the 10 plants per field were monitored. The presence of larvae in mines was confirmed by shining a torch through the leaves. Leaflets containing larvae were collected in each field when available, and kept in aery boxes (10 cm diameter, 9 cm deep) for 2 weeks, with parasitoid emergence checked every 2 days.

Table 1 Description of the Kenyan tomato farms investigated

Field identifier	Location	Gps coordinates	Tomato variety	Open field/Net house	Insecticides used	Crop state
1	Kirinyaga	S 00.618231 E37.369609	Perseo	OF+NH	Metarhizium	Fruiting
2	Kirinyaga	S 00.618231 E37.369609	Perseo	OF+NH	Metarhizium	Fruiting
3	Kirinyaga	S 00.611750 E 037.388580	KILELE F1	OF+NH	Abamectine	Fruiting
4	Kirinyaga	S 00.602460 E 037.364280	KILELE F1	OF+NH	Imidaclopride + Beta-cyfluthrine, Chlorantraniliprole, Abamectine	Started harvesting
5	Kirinyaga	S 00.597070 E 037.380050	ANNA F1	OF+NH	Flubendiamide	Fruiting
6	Kiambu	S 01.233250 E 036.664310	BRAVO F1	OF+NH	Metarhizium	Fruiting
7	Kiambu	S 01.143570 E 036.578370	ANNA F1	OF+NH	Metarhizium	Fruiting
8	Machakos	S 01.318040 E 037.226040	EVA F1	OF+NH	Thiamethoxam, metaxyl + mancozeb, Mefenoxam + Mancozeb	2 weeks after transplanting
9	Machakos	S 01.385300 E 037.249250	EVA F1	OF+NH	Thiamethoxam, metaxyl + mancozeb, Mefenoxam + Mancozeb	3 weeks after transplanting
10	Kajiado	S 01.432100 E 036.682300	TYLKA F1	NH	Neonicotinoides	Flowering
11	Kajiado	S 01.303160 E 036.777720	BRAVO F1	NH	Pyréthroïdes	3 weeks after transplanting
12	Kajiado	S 01.435010 E 036.675950	TYLKA F1, ANNA F1	NH	Metarhizium	Fruiting
13	Kajiado	S 01.434130 E 036.676760	EVA F1	OF+NH	Lambda-cyhalothrin, Imidacloprid 250 + Bifenthrin 50 SC	2 weeks since transplanted
14	Migori	S 00.763980 E 034.585090	RIOGRANDE	OF+NH	No insecticide	Fruiting
15	Migori	S 00.764100 E 034.584900	RIOGRANDE	OF+NH	No insecticide	Flowering
16	Migori	S 00.861280 E 034.610370	CAL J	NH	No insecticide	Fruiting

All pest and natural enemy samples were sorted out by morphotype and placed in ethanol (70%) and sent to the Centre for Biology and Management of Populations (CBGP, Montpellier, France) for identification or confirmation by barcoding. DNA was extracted non-destructively from full specimens using a DNeasy Blood & Tissue kit (Qiagen, Hilden, Germany), according to the manufacturer's protocol. PCR amplifications were performed using standard primers for barcoding (cytochrome c oxidase subunit 1 mitochondrial gene) of invertebrates: LCO1490: 5'-GGTCAACAAATCATAAAGATATTG G-3' (forward primer sequence) and HCO2198: 5'-TAA ACTTCAGGGTGACCAAAAATCA-3' (reverse primer sequence) (Folmer et al. 1994). PCR reactions were carried out on a Mastercycler® Nexus (Eppendorf, Hamburg, Germany) in a 10 µL PCR mix containing 5 µl of Multiplex Master Mix (Qiagen, Hilden, Germany), 0.8 µl primers (forward and reverse at 2 µM) and 2 µl of DNA. The PCR conditions were as follows: initial DNA denaturation at 94 °C for 15 min, followed by 10 cycles of 30 s each at 94 °C, 1 min at 60–50 °C (touchdown of -1 °C per cycle) and 30 s at 70 °C, followed by 30 cycles of 30 s at 94 °C, 1 min at 55 °C, and 1 min at 72 °C with a final 20 min extension at 72 °C (Haran et al. 2018). The PCR

products were sequenced by Eurofins Genomics (<http://www.eurofinsgenomics.eu/>). All voucher specimens were deposited in the general collection at CBGP. Barcode sequences were aligned and checked using CodonCode Aligner V.3.7.1. (CodonCode Corporation, Centerville, MA, USA). Each sequence was then subjected to molecular identification using the BLAST tool implemented in the NCBI platform (<https://www.ncbi.nlm.nih.gov/>).

Laboratory experiments

Laboratory experiments were conducted at 25 ± 2 °C and $65 \pm 5\%$ RH. Six-week-old tomato plants from the Kilele F1 tomato hybrid (Syngenta, Nairobi, Kenya) were used in the experiments.

Individual crossing tests

A device was created using two transparent tubes (10 cm long, 5 cm diameter) to check the capability of the main natural enemies, i.e. *Nesidiocoris tenuis* (Reuter, 1895) (Hemiptera: Miridae) and *Encarsia formosa* Gahan (1924) (Hymenoptera: Aphelinidae), to pass through the nets (Deletre et al. 2015). Three types of nets were tested (A to Z Textile Mills

Ltd, Arusha, Tanzania), two types with different mesh sizes (0.9 mm and 0.4 mm, not treated nets) and one-insecticide treated net (0.9 mm mesh size; net impregnated with deltamethrin during the manufacturing process). One tube contained the insects and was closed at the top with the tested net. A second tube contained a trifoliolate tomato leaf and was fixed on the top of the first tube (on the mesh side). The end of this second tube was closed with 160 µm mesh netting to retain all insects. The tube containing the insect was covered with aluminum foil and black paper, and the light was placed on the edge of the device. The presence of the leaflet and the light contrast would motivate the insects to cross the net. The set-up was monitored after 30 min to check whether the individuals were able to go through the net or not.

For *N. tenuis*, 15 replicates were conducted per instar and per male and female adults. Only one insect was placed per tube, thus altogether, 105 individuals were tested. For *E. formosa*, adults were tested (mix of males and females). Ten individuals were placed in a tube and 10 replicates were carried out, so 100 individuals were tested.

Oviposition crossing test

The capacity of *N. tenuis* females to oviposit on a leaflet in direct contact with the net, by crossing the three types of nets with their ovipositor, was tested. A device was thus created using Petri dishes (9 cm diameter). The bottom and cover of the Petri dishes were separated using the tested mesh type. A piece of young tomato stem was stuck into floral foam and deposited on the mesh (on the cover side). Then one week-old female was introduced at the bottom. After 24 h, the stem was removed and placed in a plastic box closed with a 160 µm mesh netting for 15 days and watered daily. After 15 days, the number of juveniles in the boxes was recorded. Twenty replicates were carried out per net and 20 replicates of the control treatment without net were tested.

Statistical analyses

All analyses were conducted using R software (R Core Team 2020). For the field survey, a principal component analysis (PCA) was performed to describe the link between the two pests and the *N. tenuis* abundance (the main natural enemy detected). For the field survey, the *T. absoluta*, *N. tenuis* and whitefly densities were analyzed with the generalized linear model (GLM) using a Poisson distribution with net presence as factor. Parasitism was analyzed with a GLM using a binomial distribution with the same factor. Net crossing test results of *N. tenuis* were analyzed using a GLM with a quasibinomial law with the net type (mesh size and insecticide treatment) and the instar as factors. In addition, the net crossing proportions per instar and per mesh were compared using chi-squared tests. A GLM with a quasibinomial

law was conducted for the *E. formosa* net cross test, with the net type as factor, and was complemented by pairwise Tukey analyses (post-hoc test using the multcomp package). A value of $p < 0.05$ was considered statistically significant and a value of p between 0.05 and 0.1 were considered marginally significant.

Results

Field survey and identifications

In overall 27 samplings conducted, whiteflies were found 19 times (70%) and *T. absoluta* were found 16 times (59%). Whiteflies were morphologically identified to be *Trialeurodes vaporariorum* (Bruno Michel, CBGP, CIRAD). This identification was confirmed by barcoding, the three specimens sequenced [CBGP vouchers JHAR01877/83/85] showing 99–100% identity with a specimen identified as *T. vaporariorum* [GB accession: MN312161.1]. No other whitefly species were found during this field survey.

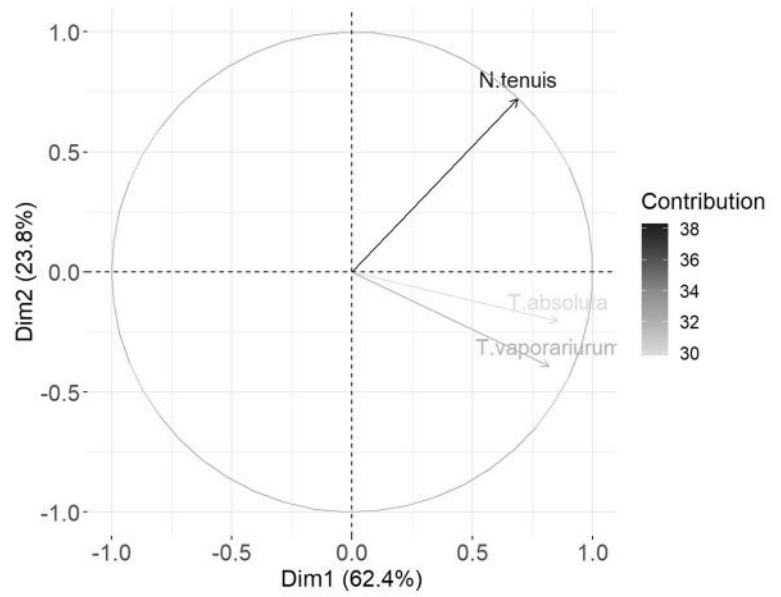
The dominant predator recorded in the aspirator sampling was *Nesidiocoris tenuis* (Hemiptera, Miridae; CBGP voucher JHAR04152, 99% identity with [GB accession: AB587603.1]), which is an efficient zoophytophagous predator that attacks both *T. absoluta* and *T. vaporariorum* (Urbaneja et al. 2009; Hassanpour et al. 2016). It represented 96.5% (335 individuals) of the individuals we caught and suspected were predatory. Nevertheless, individuals of another Miridae species generally considered as a predator, i.e. *Campylomma leucochila* (Reuter 1905) (Hemiptera, Miridae; CBGP voucher JHAR01863, 100% identity with [GB accession: KT201382.2]) were also recorded. This species constitutes 2.59% (9 individuals) of the potential predatory species we identified.

Two parasitoid species were obtained via field sampling and laboratory rearing: *Encarsia formosa* (Hymenoptera, Aphelinidae; CBGP voucher JHAR01894, 99% identity with [GB accession: MG813797.1]) emerged from *T. vaporariorum*, and *Copidosoma* sp. (Hymenoptera, Encyrtidae; CBGP voucher JHAR01876, 92% identity with [GB accession: MF906707.1]) emerged from *T. absoluta*.

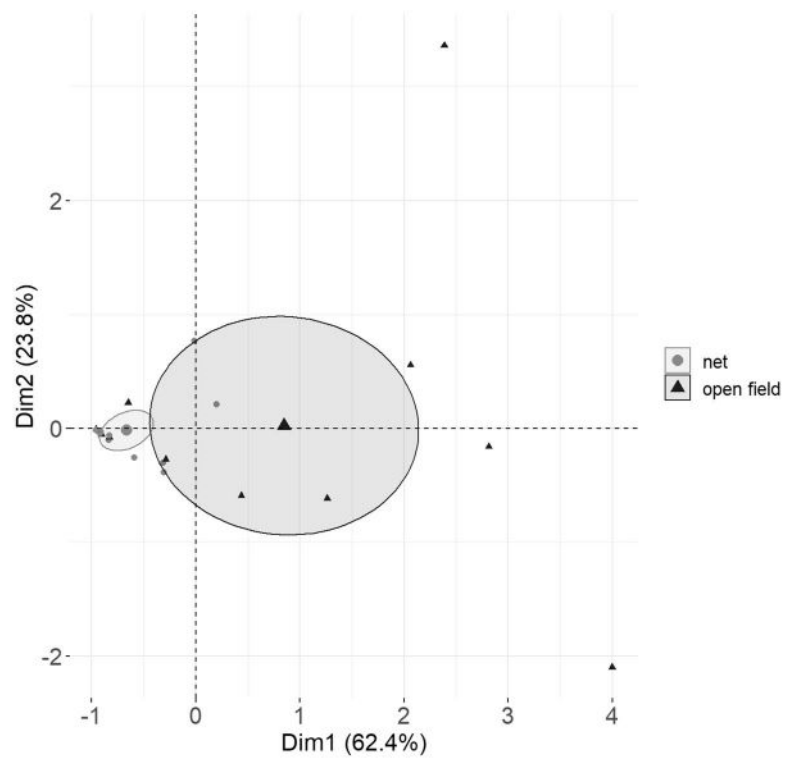
In the aspirator sampling, a few individuals of other parasitoid species were trapped in addition to the previous natural enemies mentioned, probably not attacking *T. vaporariorum* or *T. absoluta*, but rather attacking other minor pests of the sampled crops. We still report this information as a testimony of their presence in this area for further studies. The following were morphologically identified by Gerard Delvare (CBGP, CIRAD, France) to at least the genus level: *Diglyphus isaea* (Walker 1838) (Hymenoptera: Eulophidae), *Opius* sp. (Hymenoptera: Braconidae) and *Maruessa* sp. (Hymenoptera: Eulophidae) that attacks leafminer

Fig. 1 PCA mapping of the fields based on the whitefly and *T. absoluta* infestations and *N. tenuis* abundance. **(A)** represents the contribution of the variables to the axis, and **(B)** is the positioning of the fields on the axes

A.



B.



flies, *Aphidius* sp. (Hymenoptera: Braconidae) that attacks aphids, *Oomyzus* sp. (Hymenoptera: Eulophidae) that attacks *Plutella xylostella*, *Neochrysocharis* sp. (Hymenoptera: Eulophidae) that attacks leafminer flies or microlepidoptera, and *Ceranisus menes* (Walker 1839) (Hymenoptera: Eulophidae) that attacks thrips.

Regarding field infestation and predator presence, PCA mapping of the data collected revealed that *T. absoluta* and whitefly infestation were linked, while *N. tenuis* abundance was not linked to pest infestation (Fig. 1A). We also observed that open field tomato were well segregated from net tomato with respect to the pest infestation and *N. tenuis* abundance (Fig. 1B). PCA shows that: (i) net protected crops hosted fewer pests and predators, (ii) *N. tenuis* was only highly abundant in open fields, and (iii) open fields were more variable in terms of pest and predator abundances (Fig. 1B).

The presence of nets significantly reduced pest infestations ($p < 0.001$ for both pests) but did not completely prevent infestations. Under the nets, 1.02 ± 0.46 (mean \pm SE) whiteflies were found per leaflet, while 7.71 ± 4.07 (mean \pm SE) were found per leaflet in open fields. Similar differences were observed with *T. absoluta*, 0.12 ± 0.05 (mean \pm SE) larvae per leaflet were found under the nets, while 0.69 ± 0.18 (mean \pm SE) were found in open fields.

The parasitism ratio was low and not significantly affected by the nets ($p = 0.781$ and $p = 0.360$, for *T. vaporariorum* and *T. absoluta*, respectively). Regarding whiteflies, we found $3.57 \pm 3.57\%$ (mean \pm SE) parasitism under the net and $0.06 \pm 0.04\%$ (mean \pm SE) in open fields. Regarding *T. absoluta* larvae, we found $2.86 \pm 2.86\%$ (mean \pm SE) parasitism under the nets and $3.50 \pm 2.57\%$ (mean \pm SE) in open fields. However, cases of parasitism were observed under nets for both pests, highlighting that parasitoids were able to enter the net houses.

Nesidiocoris tenuis was the dominant predator species and was therefore the focus of our statistical analyses. Its population was significantly affected by the net ($p < 0.001$).

Populations were higher in open fields, with 2.57 ± 1.24 (mean \pm SE) individuals found per plant, while 0.37 ± 0.26 (mean \pm SE) individuals per plant were found under the nets.

Laboratory experiments

Individual crossing tests

The net type, i.e. the mesh size and the treatment of the net with deltamethrin, did not significantly affect the proportion of *N. tenuis* individuals passing through the mesh ($p = 0.338$) (Table 2). However, the ability of *N. tenuis* instars to cross the different net types significantly varied ($p < 0.001$). The first three *N. tenuis* instars were able to cross the untreated 0.9 mm mesh net, while only the first two instars were able to cross the untreated 0.4 mm net and the treated 0.9 mm mesh net (Table 2).

Encarsia formosa adults were able to cross all the three types of nets, but the proportion that passed through the mesh was marginally significantly affected by the net type ($p = 0.096$), with the lowest crossing rate obtained for the untreated 0.4 mm mesh net (Table 2).

Oviposition test

All net types significantly reduced oviposition by *N. tenuis* females ($p < 0.001$), although they were able to oviposit through all nets without any significant differences noted between net types (Fig. 2).

Discussion

We investigated the impact of net houses on tomato crop infestation by two major pests in Kenya, i.e. *T. vaporariorum* and *T. absoluta*, and their natural enemies. Several natural

Table 2 Percentage of *Nesidiocoris tenuis* and *Encarsia formosa* individuals that crossed nets of two different mesh sizes and treated or not with deltamethrin. Letters indicate differences between nets for each insect stage, that were significant ($p < 0.05$) for *N. tenuis* and

marginally significant ($p < 0.1$) for *E. formosa*. For *N. tenuis*, crossing percentages were compared using a chi-squared tests ($P < 0.05$), for *E. formosa*, mean \pm SE were compared using a Tukey test

Species	Instar	0.4 mm untreated (%)	0.9 mm treated (%)	0.9 mm untreated (%)
<i>N. tenuis</i>	1st nymph instar	80 a	70 a	80 a
	2nd nymph instar	70 a	60 a	60 a
	3rd nymph instar	0 a	0 a	60 b
	4th nymph instar	0	0	0
	5th nymph instar	0	0	0
	Adult females	0	0	0
	Adult males	0	0	0
<i>E. formosa</i>	Adults mixed sex	12 ± 3 a	18 ± 3 ab	23 ± 4 b

enemy species, both predators and parasitoids, were found and identified. In addition, the capacity of the key natural enemies of the two pests to cross the nets was tested.

Nesidiocoris tenuis, the most abundant predator found, is a well-known predatory species that feeds on eggs and larvae of *T. absoluta* (Urbaneja et al. 2009), and on nymphs of *T. vaporariorum* (Hassanpour et al. 2016). It is also known to be present in 17 African countries (CABI 2019a) and to feed on *T. absoluta* in invaded areas (Sylla et al. 2016; Boualem et al. 2012; El Arnaouty and Kortam 2012). In this study, its abundance was linked

neither to *T. absoluta* nor whitefly abundance, which might possibly be explained by its zoophytophagous and polyphagous diet. However, this also highlighted the impact of factors other than the prey resource on its abundance, such as insecticide treatments (Arnó and Gabarra 2011), crop diversity (Thomine et al. 2020), climate or agroecological zones (Sridhar et al. 2012), but these factors could not be tested with our sampling design. In our study, most of the pesticides used were not selective, and it is known that organic insecticides can also negatively affect mirids (Biondi et al. 2012).

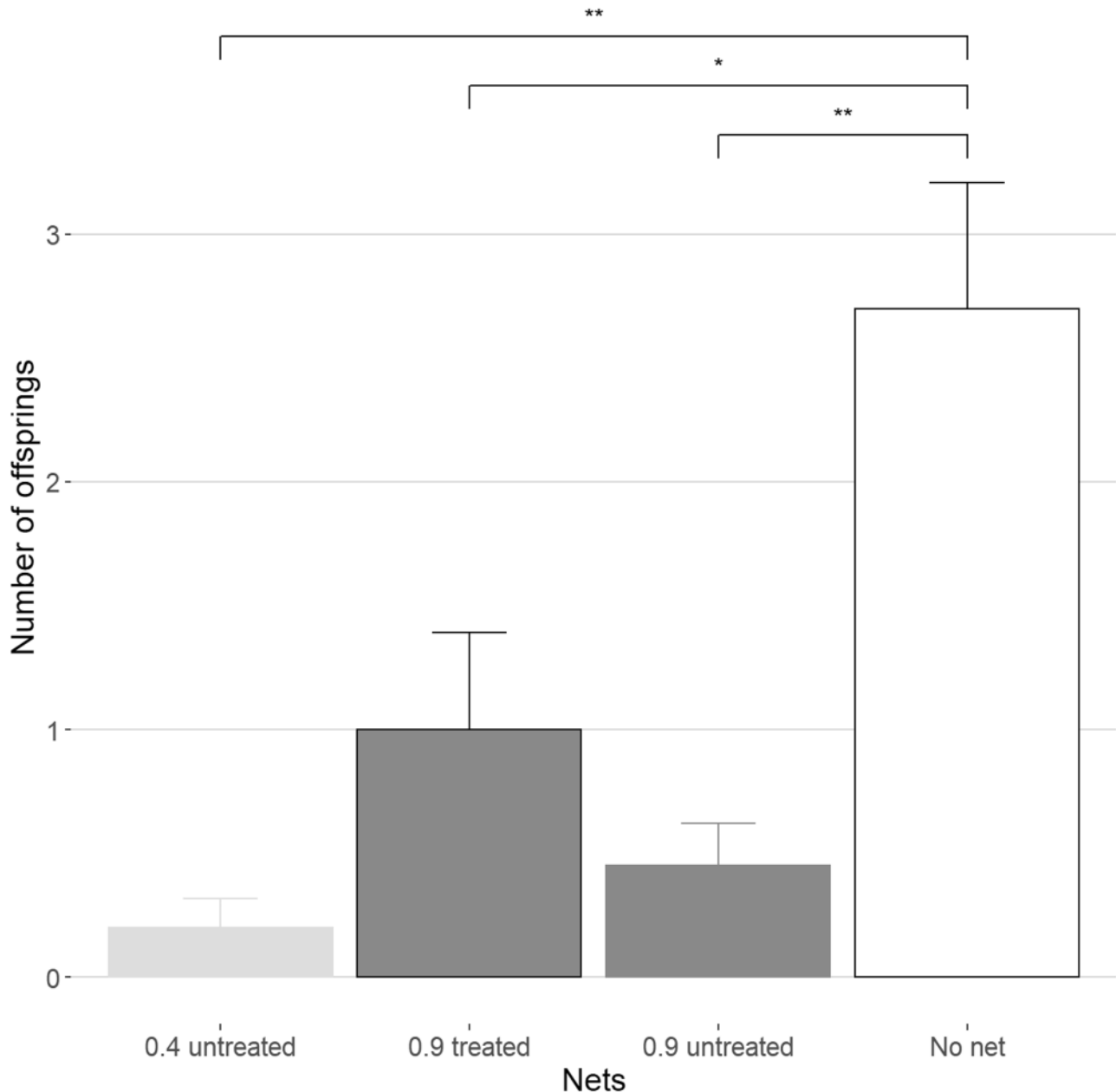


Fig. 2 Oviposition (mean \pm SE) of *N. tenuis* females through nets of two different mesh sizes, treated or not with deltamethrin, and a control (no net). Stars indicate significant differences using a Tukey test (**: $P < 0.01$ and *: $P < 0.5$)

Parasitoids were identified for both pests, despite parasitism being low. Indeed, the natural parasitism of both pests is generally found to be low when no practices are set up to improve it (Kinyanjui et al. 2021; Abbes et al. 2014; Nannini et al. 2006). In addition, farmers were allowed to pursue the practices they usually set up, involving insecticides, yet it is known that parasitoids are susceptible to insecticides (Yarahmadi et al. 2018; Abbes et al. 2015) including entomopathogenic fungus (Oreste et al. 2016). We still identified the parasitoid *E. formosa*, that is widely used for biological control of *T. vaporariorum*, firstly used successfully in England in 1926 (Speyer 1927). By 1988, *E. formosa* was estimated to have been used on 3,800 ha of greenhouse area worldwide for biological control of *T. vaporariorum* (van Lenteren and Woets 1988). It has been authorized for biological control in Kenya since 2003 (Songa 2003). This might explain why this was the only parasitoid we found despite other parasitoids are known to develop on *T. vaporariorum* and to be able to coexist with *E. formosa* when sharing a host (Bográn et al. 2002). This can also owe to the sampling size that might not be sufficient to detect rare species, or do not exhaustively represent whitefly instars, thus biasing the results (*E. formosa* preferentially attack third and fourth instars (Nell et al. 1976)). We also identified *Copidosoma* sp. that is associated with *T. absoluta* in South America (Desneux et al. 2010). Contrary to most of *T. absoluta* parasitoids (Chailleux et al. 2014), *Copidosoma* sp. is a polyembryonic endoparasitoid belonging to the egg-larval parasitoid guild (Salas Gervassio et al. 2016). This genus is known to be present throughout Africa (CABI 2019b) and the species *Copidosoma koehleri* Blanchard (1940) is known to be present in Kenya (Guerrieri and Noyes 2005).

Our study is the first formal report of the presence of *N. tenuis* and *E. formosa* in Kenyan fields. Both of these natural enemies are commercially produced and sold worldwide. It is unclear if these natural enemies are indigenous to Kenya or if they were introduced for biological control in industrial horticultural crops. This is quite likely with regard to *E. formosa*, which has been produced and sold all over the world for decades, while acclimation and establishment in the environment has not been investigated everywhere. Besides, the fact that both natural enemies are easily mass-reared is crucial toward their use in augmentative biological control programs. Indeed, *E. formosa* is mass-reared on *T. vaporariorum* on tobacco plants (Scopes 1969), and plantless rearing of *N. tenuis* fed on *E. kuheniella* eggs is being developed (De Puyssseleyr et al. 2013).

Our results showed that net houses reduced the pest population but did not entirely prevented neither pest infestation nor the establishment of natural enemies. Predators were less abundant under net houses, while parasitoids did not seem to be affected by the nets. This was probably due to differences in the size of the natural enemies, i.e. parasitoids were smaller than the predators found in this

survey. Moreover, our laboratory experiments confirmed this hypothesis, showing that *E. formosa* was even able to pass through the 0.4 mm mesh net, while only *N. tenuis* nymph, which are wingless and therefore less mobile, were able to cross the nets. Still, the 0.4 mm mesh reduced the percentage of parasitoids crossing the net, probably owing to the visual barrier created by the small mesh, that superimpose to the physical barrier. Indeed, *E. formosa* uses visual stimuli to locate its host (Romeis and Zebitz 1997), thus the visual barrier may have decreased parasitoid motivation to cross the net.

Conclusion and recommendations

Net houses are a first step toward pesticide reduction but are not sufficient, while indigenous natural enemies have the potential to complement the tomato protection under nets. Our results highlighted two main needs warranting further use of natural enemies for tomato protection in Kenya. First, the need to identify the factors other than prey abundance affecting *N. tenuis* populations, so as to boost its abundance in open fields and make them more reliable. Second, there is a need to find ways to increase the abundance of natural enemies, either by (i) using kairomones to attract natural enemies (Ayelo et al. 2021), or (ii) introducing natural enemies under net houses through inoculative releases, as has been done for *N. tenuis* (Calvo et al. 2012), or through inundative releases, as has been done for *E. formosa* (van Lenteren et al. 1996).

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Declarations

Conflict of interest statement On behalf of all authors, the corresponding author states that there is no conflict of interest.

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