



Article

Determination of Insecticide Susceptibility of Field Populations of Tomato Leaf Miner (*Tuta absoluta*) in Northern Nigeria

Ibrahim Bala ^{1,2}, Muhammad M. Mukhtar ¹, Habeeb K. Saka ² , Nasiru Abdullahi ¹ and Sulaiman S. Ibrahim ^{1,3,*} 

¹ Department of Biochemistry, Bayero University, PMB 3011 Kano, Nigeria; binbala04@gmail.com (I.B.); muhammadmahemukhtar@gmail.com (M.M.M.); nabdullahi.bch@buk.edu.ng (N.A.)

² Nigerian Stored Products Research Institute, Kano Office, PMB 3032 Kano, Nigeria; saka.habeeb@yahoo.com

³ Vector Biology Department, Liverpool School of Tropical Medicine, Liverpool L3 5QA, UK

* Correspondence: sulaimansadi.ibrahim@lstmed.ac.uk; Tel.: +234-80-2901-0371

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Abstract: In 2016, northern Nigeria experienced a devastating infestation by the tomato leaf miner, leading to soaring in prices of tomatoes across the country. Unfortunately, information on the resistance status of this pest is lacking in northern Nigeria, hampering appropriate control measures. Here, we identified to species level and, using bioassays, characterised insecticide susceptibility profile of a field population of a tomato leaf miner from northern Nigeria. Highest resistance was observed with λ -cyhalothrin (a Type II pyrethroid) with a low mortality (18.52% at 56 h) and LD₅₀ of 7461.474 ppm. Resistance was also established toward propoxur and chlorpyrifos-methyl with average mortalities each of 56% and LD₅₀s of 1023.51 ppm and 106.351 ppm, respectively. Highest susceptibility was observed from abamectin with mortality of 86% and LD₅₀ of 0.034 ppm. Pre-exposure to the synergist piperonylbutoxide significantly recovered λ -cyhalothrin susceptibility ((mortality~90%, $\chi^2 = 98.35$, $p < 0.0001$) and LD₅₀ = 0.92 ppm) implicating P450 monooxygenases. No significant changes were observed on pre-exposure to diethyl maleate and triphenylphosphate-inhibitors of glutathione S-transferases and carboxylesterases, respectively. Sequencing of domain II of the voltage-gated sodium channel established 1014F *kdr* mutation 100% fixed in both λ -cyhalothrin-alive and dead larvae. These findings highlight the challenges for control of this invasive agricultural pest in northern Nigeria.

Keywords: Savanna; Nigeria; tomatoes; *Tuta absoluta*; metabolic; resistance; *kdr*

1. Introduction

The leaf miner, *Tuta absoluta* (Meyrick 1917) (Lepidoptera: Gelechiidae), native to South/Central America [1], is one of the most devastating pests of tomato (*Solanum lycopersicum* L.), potato (*Solanum tuberosum* L.) and other solanaceous plants [2]. Since its first appearance in Spain in 2006 [3] *T. absoluta* has expanded its geographic range outside Americas [4], and is now found in far flung places in Europe [5,6], middle East/Asia [7,8] and Africa [9,10]. Following its first appearance in sub-Saharan Africa at Senegal [11], this pest has expanded southward and, in recent years, has been reported in East [12] and southern African countries [13]. Africa is a continent where the agricultural sector accounts for more than 60% of total labour force [14], and Nigeria is the largest producer of tomato in the continent [15]. However, following the 2016 *T. absoluta* invasion (nicknamed ‘tomato ebola’) more than 80% of the tomato produce was lost in northern Nigeria [16], leading to estimated 10-fold increase in tomato prices [17]. In Kano State alone, farmers lost more than 2 billion Nigerian Naira in the 2016 season [18].

Farmers from northern Nigeria responded by increasing the quantities of pesticides they apply (personal communication), often in mixtures of classes having similar mode of actions. At Kadawa farms in Kano, where collection for this study was conducted, farmers were found to be using the following pesticides for control of the leaf miner: (i) Expert 50 WDG, Emacot 050 WG, and Caterpillar Force, all three made of Emamectin benzoate; and (ii) TEMA, made of emamectin benzoate (60 g/kg) and Teflubenzurone (75/kg). They claimed that TEMA is the most effective of all four formulations.

Unfortunately, scientific evidence on pesticide resistance profile and its underlying molecular mechanisms in *T. absoluta* was non-existent in Nigeria prior to this study. Elsewhere, significant progress has been reported on the bionomics of this pest, its pesticides resistance status, and the underlying molecular mechanisms driving the resistance in the field. Studies in different countries have shown that *T. absoluta* has developed resistance to insecticide classes in use for its control [19,20]. Cases of resistance in *T. absoluta* and its underlying molecular mechanisms have been reported in Southern America and Europe, e.g. in Chile and Brazil [21,22], and are reported in Greece, Italy, Spain, and Portugal [23,24]. In Africa, reliable studies that describe pesticide resistance in this pest are from Ethiopia, e.g. the recent work by Ayalew and Shiberu, with their colleagues [25,26].

Since insecticide resistance could be heterogenous even over short distances, as observed in other species like mosquitoes [27], it is not wise to extrapolate findings from other countries to the local populations in Nigeria and/or Africa. To fill these gaps in knowledge and provide information to the relevant agricultural authorities, we characterised a field population of *T. absoluta* from a single site in the Sudan Savanna of northern Nigeria. Following field collection and morphological identification, subsample of the *T. absoluta* were identified to species level using molecular approach. Insecticides resistance profile was then established and the possible enzymes systems responsible for metabolic resistance interrogated using synergist bioassays. The *T. absoluta* populations were resistant to pyrethroid, carbamate, and organophosphate insecticides, with resistance also suspected to abamectin. Synergist bioassays significantly recovered susceptibility with mortalities increasing fourfold on average; revealing that the P450 monooxygenases are possibly involved in the pyrethroid resistance. Sequencing of the domain II, transmembrane helix 6 fragment of the voltage-gated sodium channel encompassing the 1014 codon identified the presence of the 1014F *kdr* mutation in all larvae alive and dead from exposure to λ -cyhalothrin.

2. Materials and Methods

2.1. Materials

Chemicals and Reagents

The four different insecticides classes used for the bioassays: λ -cyhalothrin (a type II pyrethroid with an α -cyano group, IRAC Class 3A), propoxur (a carbamate, IRAC Class 1A), chlorpyrifos-methyl (an organophosphate, IRAC Class 1B) and abamectin (an ivermectin, IRAC Class 6), were purchased from SIGMA ALDRICH, UK (Dorset, United Kingdom). The triton X-100 and the synergists piperonyl butoxide (PBO) and diethyl maleate (DEM) and triphenyl phosphate (TPP) were all purchased from SIGMA, Dorset, UK. For species identification, the KAPATaq kit (<https://www.kapabiosystems.com/>) was used to amplify the cytochrome oxidase fragment, and for amplification of IIS6 domain of the voltage-gated sodium channel QuantaBio HiFi Accustar II SuperMix (www.quantabio.com) was used. Other chemicals used were to make a LIVAK DNA extraction buffer and are given in the methods section.

2.2. Methods

2.2.1. Field Collection and Rearing of Insects

T. absoluta larvae and eggs at different stages of development were collected from tomato farms at Kadawa ((11.6457 °N, 8.4479 °E), Figure 1) in Garun Mallam Local Government Area, Kano, Nigeria. Collection was done for two days in three farms in April 2017.

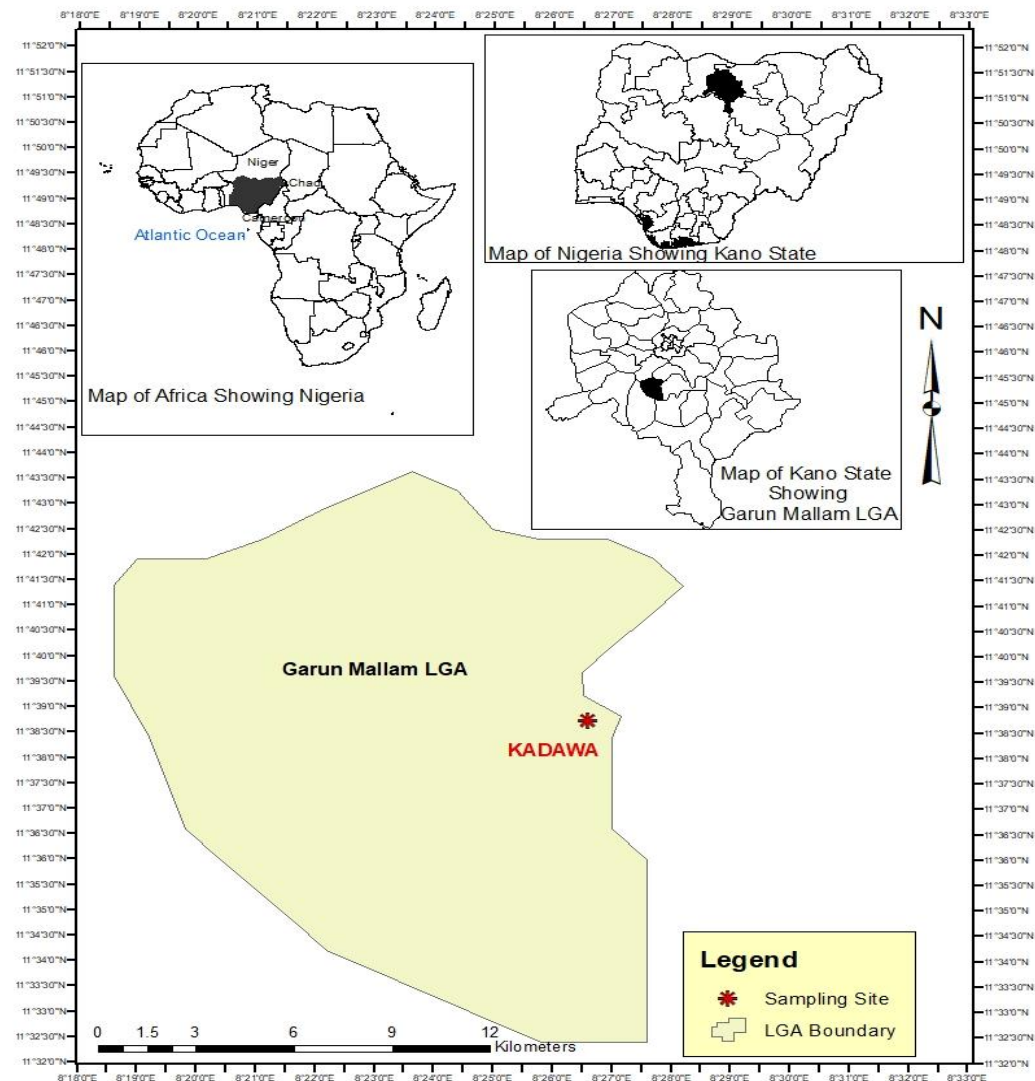


Figure 1. Sampling site, showing the location of Kadawa, Kano State, Nigeria.

Eggs and larvae were placed into large, wet jute bags. Approximately 300 larvae were collected. The samples were transferred to net cages measuring 70 cm × 50 cm × 50 cm (height × width × depth), locally constructed following established procedure [28]. This was done within an hour of collection to avoid stressing the larvae. Cages were maintained at 22–25 °C, relative humidity of 70–75%, and a photoperiod of 12:12 h (light:dark cycle) at Wellcome-Bayero Insectary, at Bayero University Kano, Nigeria. Adults that emerged were provided daily with fresh tomato leaves and allowed to mate. After laying eggs, the F_0 parents were transferred into other cages and killed. Sixteen (16) randomly selected adults were selected for morphological identification and molecular identification. Fresh foliage was provided daily for the newly emerged F_1 larvae and allowed to grow for 3–5 more days until the second instar stage (after second moulting). The tomato leaves were continually sprayed with water to keep them from wilting.

2.2.2. Morphological Identification of *T. absoluta* Life Stages

Identification of eggs, larvae, pupae, and the field-collected adults was carried out morphologically using a stereomicroscope and following the protocol of European and Mediterranean Plant Protection Organisation (EPPO) [2].

2.2.3. Molecular Identification of *T. absoluta* to Species Level

Following morphological identification 16 F₀ parents were used for DNA extraction using previously established protocol [29]. Buffer was made by dissolving 5.48 g sucrose, 1.57 g Tris, 1.6 mL of 5 M sodium chloride in 10.16 mL of 0.5 M EDTA. This was then followed by a 2.5 mL of 20% SDS, and the volume was finally made up to 100 mL in a volumetric flask. The buffer solution was then filtered and sterilised. Then, 5 mL aliquots were stored at −20 °C, which was heated in a water bath and whirled to re-dissolve precipitate before use. Larvae were homogenized individually using a battery-operated mortar and pestle (SIGMA) in 50 µL preheated grind buffer in 1.5 mL Eppendorf. The pestle was rinsed with a further 50 µL of the buffer to a total of 100 µL. Homogenate was incubated at 65 °C for 30 min. Condensation was collected by microfuging and 14 µL of 8 M of potassium acetate added to a final concentration of 1 M. Samples were vortexed and incubated for 30 min on ice. Tubes were centrifuged at 13,000 rpm for 20 min, at 4 °C, after which the supernatant was transferred carefully to a 1.5 mL Eppendorf. At this point, 200 µL of 100% ethanol was added and mixture centrifuged again at 13,000 rpm for 15 min, at 4 °C. Pellets was rinsed in approximately 100 µL ice cold 70% ethanol, air-dried for 1hr and then re-suspended in 100 µL of distilled water. Tubes were finally incubated at 65 °C for 10 min.

Identification to species level was carried out by amplifying Cytochrome Oxidase subunit I (COI) gene using polymerase chain reaction [30,31]. The universal forward and reverse primers: LCO1490 (5'-GGTCAACAAATCATAAAGATATTGG-3' and HCO2198 (5'-TAAACTTCAGGGTGA CCAAAAAATCA-3') were used for PCR using 1 µL each of the genomic DNA in a total reaction volume of 15 µL. Reaction mix comprise 1.5 µL of 10× TaqA Buffer, ~0.4 µM (0.5 µL) of each of forward and reverse primers, 1.25 mM (0.75 µL) of MgCl₂, 0.25 mM (0.15 µL) of dNTP mixes and 0.12 µL of Taq DNA polymerase, in ddH₂O. Amplification was carried out using the following conditions: initial denaturation of 5 min at 95 °C, followed by 35 cycles each of 30 s at 94 °C (denaturation), 30 s at 57 °C (primer annealing) and 1 min at 72 °C (extension). This was followed with 10min final extension at 72 °C. PCR products were separated in a 1.5% agarose gel stained with ethidium bromide. Eight (PCR) products were cleaned with a QIAquick® PCR Purification Kit (QIAGEN, Hilden, Germany) and sequenced on both strands. Sequences were assembled and edited using the BioEdit 7.2.3.0 [32] and the CLC sequence viewer 6.9 (<http://www.clcbio.com/>). Consensus sequences were BLASTED (NCBI Blastn) in the GenBank database (<http://www.ncbi.nlm.nih.gov/>) for identification and comparison with sequences previously deposited. Sequences were deposited at the GenBank with accession numbers MK189155- MK189162.

2.2.4. Insecticides (Leaf-Dip) Bioassay

Initial bioassay was conducted to establish the susceptibility profile of the *T. absoluta* larvae to the four different insecticides classes in use for control of agricultural pests. In total, 270 L2 larvae were utilized, 234 for tests with pesticides and 36 for control. Larvae were sorted using a fine brush according to sizes to select those at the second instar stage. Insecticides were tested using the leaf-dip bioassay protocol of the Insecticide Resistance Action Committee (IRAC test method # 022) [33] with modifications. However, due to high mortality before standardization of insectary conditions low numbers of larvae were used in these bioassays compared with the recommendation of Insecticide Resistance Action Committee. Initially, acetone stock solutions at different concentrations were prepared for the various insecticides: 0.1% abamectin, and 0.5% each of propoxur, chlorpyrifos-methyl and λ-cyhalothrin. These were serially diluted into decreasing concentrations (1:6) in water containing

0.01% Triton X-100. Fresh tomato leaves were cut to equal sizes of 2.5 cm diameter using hand-held punch. For each insecticide, for the six different concentrations, nine tomato leaves (three replicates of three leaves) were dipped individually into insecticide solution, with gentle agitation. The only exception was abamectin with eight different concentrations instead of six. After treatment, the leaves were placed individually on wire net for 60 s to dry. Individual leaves were transferred into bioassay trays (ref: RT32W) containing slightly moistened filter paper. A fine soft brush was used to transfer the second instar larvae (three replicates of three larvae) into cells in the bioassay trays, individually and the trays covered with a lid (ref: RTCV4). Trays were closed carefully, sealing the cells with their lids and then stored at about 25 ± 2 °C, 60–70% relative humidity, and 12:12 h light: dark photoperiod. Leaf damage as well as larvae mortality were then evaluated after 56 h. Larval mortality was recorded with regards to those which were unable to make coordinated movement from gentle stimulus with fine pointed forceps to the posterior body segment (considered dead or seriously affected by the insecticide). To establish antifeeding effect leaf damage was evaluated by physical examination of the leaves. Since uniform leaves (~2.5 cm punched out) at the beginning of the assay the extent of damage was estimated as percentage of total leaf area mined (IRAC method # 022). For control, the same procedure as above was followed with nine larvae sets for each experiment with the different insecticides, excepting that the leaves were dipped into water containing acetone.

2.2.5. Synergists Bioassay

To determine the possible contribution of metabolic resistance in the *T. absoluta* populations, synergist assay was conducted with piperonyl butoxide (PBO: an inhibitor of P450 monooxygenases), DEM (an inhibitor of glutathione S-transferases) and TPP (an inhibitor of esterases) against λ -cyhalothrin (the insecticide to which the larvae exhibited the highest resistance). 108 L2 larvae were used for the synergist test and 27 for control. Three replicates of three larvae per insecticide concentration were placed in cell units containing a leaf dipped into either 4% PBO, 8% DEM or 10% TPP (all synergists prepared in acetone) for 1 hr, as done in some other insects [34,35]. The larvae were then immediately transferred into bioassay cells containing leaves individually dipped into the various concentrations (0.643 ppm, 3.853 ppm, 23.15 ppm, and 138.89 ppm) of λ -cyhalothrin-concentrations at which lowest mortalities were previously observed in the conventional bioassay described above.

For control, 27 larvae were first treated each with either PBO, DEM, or DEF, as above, and then placed in cells containing untreated leaves. Mortality was assessed 56 h after exposure.

2.2.6. Sequencing of Domain II of the Voltage-Gated Sodium Channel

To investigate the presence of the 1014F *kdr* mutation DNA was extracted [29] from eight larvae alive and eight dead following exposure to λ -cyhalothrin. Amplification of the fragment encompassing the 1014 codon was done using a nested PCR with the primers described by Haddi and colleagues [24]: TAF3 and TAR1 in the first PCR and the primers TAF4 and TAR2 in the second reaction. To 12.5 μ L of the 2 \times AccuStartII HiFi PCR SuperMix (QuantaBio, Beverly, Massachusetts) containing optimized concentrations of MgCl₂ and dNTP mixes, 0.2 μ M each of the forward and reverse primer was added, followed by 1 μ L gDNA. 10.5 μ L of dH₂O was added to obtain a total volume of 25 μ L. Amplification was carried out using the following conditions: initial denaturation of one cycle at 94 °C for 3 min; 35 cycles of 94 °C for 30 s (denaturation), 60 °C for 30 s (annealing), and extension at 72 °C for 1 min; and one cycle at 72 °C for 5 min (final elongation).

Nest 2 product was purified, sequenced on both strands and assembled/edited as described above, and NCBI Blastn used to identify similar sequences in the GenBank. Fifteen sequences, seven from the larvae alive from λ -cyhalothrin exposure, and eight from the dead larvae were deposited in the GenBank with accession numbers MK189140-MK189154.

2.3. Data Analysis

The intensity of resistance was estimated by calculating the LD₅₀ for the various insecticides using glm probit analysis as implemented in MASS package of R version 3.5.0 (<https://cran.r-project.org/bin/windows/base/>). All figures were prepared with and the results of synergised and un-synergised tests with λ -cyhalothrin compared using a two-tailed Chi-Square test of independence using GraphPad Prism 7.02 (GraphPad Inc., La Jolla, CA, USA).

3. Results

3.1. Morphological and Molecular Identification of *T. absoluta* to Species Level

The field collected F₀ larvae and adults were morphologically identified as *T. absoluta* (Meyrick 1917) (Lepidoptera: Gelechiidae), based on the following characteristics as explained in previous publications [2,36]: early instars were white/creamy with black heads which changed into greenish from second instar with heads turning to brown/dark brown. The first instar larvae were <1 mm, and there was gradual increases in length until the fourth instar, which was on average 7–8 mm long. Second instar larvae were 4–5.5 mm. Pupae were brown in colour and were folded into leaves singly. Adults were about 9–10 mm long, with filiform antennae, silver-grey scales, and evidence of black spots on anterior wings.

DNA from all the 16 F₀ parents produced ~658 bp characteristic of metazoan cytochrome c oxidase I [37] (Figure 2). Sequencing of eight PCR products revealed that they were 100% identical to KX443111 [38], KT452897, KY212128, and KU565719 deposited from previous studies at the NCBI. The 695bp sequences have been deposited in the GenBank: accession numbers MK189155–MK189162.

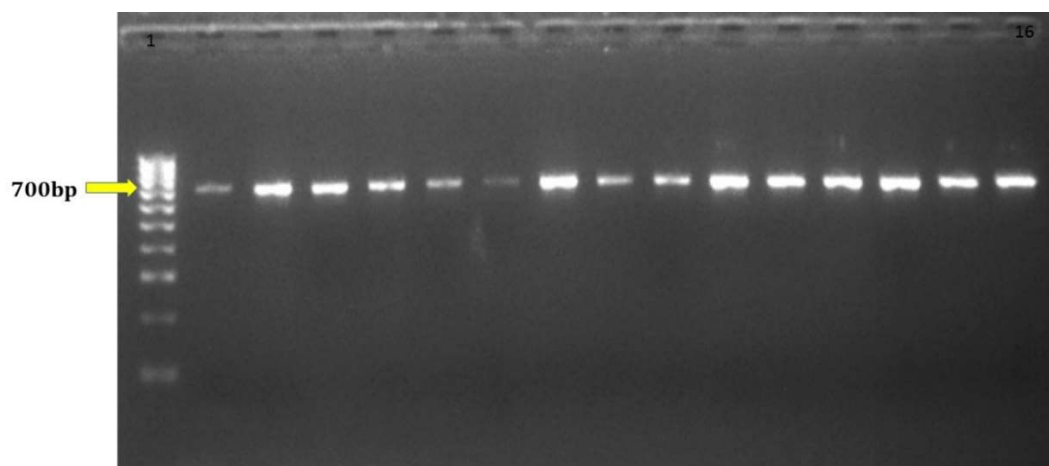


Figure 2. Polymerase chain reaction species identification of *T. absoluta* species. A characteristic band of 658 bp typical to metazoans is evident in lanes 2–16. Lane 1 represents HyperLadder IV from Bioline (~1013 bp).

3.2. Insecticides Resistance Profile of *T. absoluta* Populations

The leaf-dip bioassay revealed highest mortality with the type II pyrethroid λ -cyhalothrin, (average mortality of only 18.52% \pm 2.0 after 56 h) (Figure 3A, Table 1) and LD₅₀ of 7461.474 ppm \pm 1213.793 (Table 2). Highest leaf damage (~15% of the leaves destroyed) was also observed with this insecticide (Figure 3B, and Table 1), though not significantly different from the other insecticides tested.

In contrast with λ -cyhalothrin results, approximately 56% mortalities were recorded respectively, for propoxur and chlorpyrifos-respectively (Figure 3A, Tables 1 and 2). However, the LD₅₀ of propoxur (1023.35 ppm \pm 218.69) was on average ten times higher than obtained with chlorpyrifos-methyl (106.30 ppm \pm 13.09) (Table 1, Table 2) due to the higher mortalities obtained at lower concentrations with the latter.

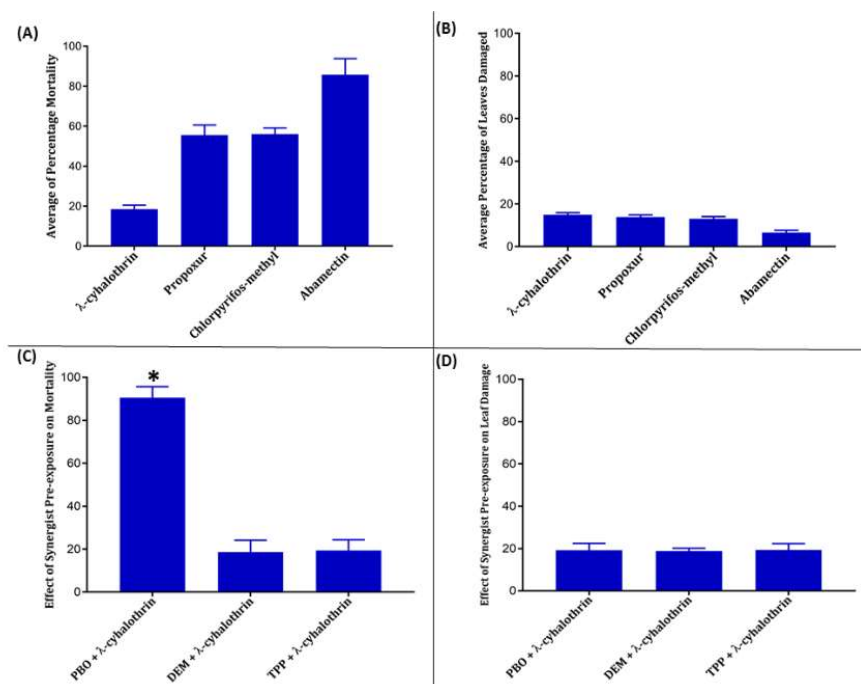


Figure 3. Results of conventional and synergist bioassays with various concentration of insecticides. Average of percentage for six different concentration ranges for each insecticides \pm standard error of mean, for (A) mortalities, (B) leaves damaged, (C) mortalities with pre-exposure to synergists PBO, DEF, and DEM, and (D) leaves damaged from synergist bioassay with PBO. * Significantly different from conventional test with λ -cyhalothrin only, $\chi^2 = 98.35$, $p < 0.0001$.

Table 1. Susceptibility profiles mortalities) and leaf damage from exposure to various insecticides and synergists.

Concentration (ppm)	Population before Treatment	Population after Treatment	Mortality (%)	Leaf Damage (%)
Abamectin				
1000	9	0 (9 dead)	100	0.22
166.67	9	0 (9 dead)	100	2.17
27.78	9	0 (9 dead)	100	4.78
4.630	9	0 (9 dead)	100	6.00
0.7716	9	0 (9 dead)	100	7.22
0.1286	9	1 alive (8 dead)	85.71	6.78
0.0214	9	4 alive (5 dead)	42.86	12.00
0.0036	9	3 alive (6 dead)	57.14	13.44
0.00 (control)	9	9 alive (0 dead)	0.00	16.61
Propoxur				
5000	9	2 alive (7 dead)	77.77	5.78
833.33	9	3 alive (6 dead)	66.66	8.67
138.89	9	4 alive (5 dead)	55.55	9.67
23.15	9	4 alive (5 dead)	55.55	14.22
3.858	9	5 alive (4 dead)	44.44	19.78
0.643	9	6 alive (3 dead)	33.33	20.22
0.00 (control)	9	9 alive (0 dead)	0.00	25.56

Table 1. Cont.

Concentration (ppm)	Population before Treatment	Population after Treatment	Mortality (%)	Leaf Damage (%)
λ-cyhalothrin				
5000	9	6 alive (3 dead)	33.33	7.89
833.33	9	7 alive (2 dead)	22.22	9.17
138.89	9	8 alive (1 dead)	11.11	13.33
23.15	9	7 alive (2 dead)	22.22	16.89
3.858	9	8 alive (1 dead)	11.11	19.22
0.643	9	8 alive (1 dead)	11.11	22.78
0.00 (control)	9	9 alive (0 dead)	0.00	25.56
Chlorpyrifos-Methyl				
5000	9	0 alive (9 dead)	100	4.94
833.33	9	0 alive (9 dead)	100	8.72
138.89	9	4 alive (5 dead)	50	17.78
23.15	9	5 alive (4 dead)	37.5	17.67
3.858	9	5 alive (4 dead)	37.5	16.11
0.643	9	8 alive (1 dead)	11.11	17.78
0.00 (control)	9	9 alive (0 dead)	11.11	20.44
Piperonyl Butoxide (PBO) + λ-cyhalothrin				
138.89	9	0 alive (9 dead)	100	1
23.15	9	0 alive (9 dead)	100	1
3.858	9	1 alive (6 dead)	87.50	3
0.643	9	2 alive (7 dead)	75.00	5
0.00 (control)	27	27 alive (0 dead)	0.00	-
Diethyl Maleate (DEM) + λ-cyhalothrin				
138.89	9	7 alive (2 dead)	22.22	13.33
23.15	9	8 alive (1 dead)	11.11	16.89
3.858	9	8 alive (1 dead)	11.11	19.22
0.643	9	8 alive (1 dead)	11.11	22.75
Triphenyl Phosphate (TPP) + λ-cyhalothrin				
138.89	9	7 alive (2 dead)	22.22	12.66
23.15	9	7 alive (2 dead)	22.22	17.47
3.858	9	7 alive (2 dead)	22.22	23.10
0.643	9	8 alive (1 dead)	11.11	20.85

Highest mortality was obtained with abamectin ($85.71\% \pm 8.1$) with a very low LD_{50} of only 0.034 ppm ± 0.0036 (Figure 3A, Tables 1 and 2). Highest foraging inhibition was also observed with this pesticide with only ~7% of the leaves damaged at the end of the experiment (Figure 3B, Table 1, and Table 2).

Table 2. Results of probit analyses for insecticides bioassay and leaf damage with various concentrations of insecticides.

Insecticides	<i>n</i>	Leaf Damage \pm S.E.M.	% Mortality \pm S.E.M.	LD_{50} (ppm)	LD_{50} (S.E.M)	LD_{50} (FL 95%)
λ -cyhalothrin	54	14.88 \pm 1.05	18.51 \pm 5.70	7461.47	1213.79	5082.47–9840.89
Propoxur	54	13.83 \pm 1.00	55.55 \pm 5.02	1023.35	218.68	594.08–1452.04
Chlorpyrifos-methyl	54	13.06 \pm 1.06	56.01 \pm 3.10	106.30	13.09	80.75–132.15
Abamectin	72	6.58 \pm 1.10	85.71 \pm 8.10	0.034	0.0036	0.01–0.09
Control	36	22.04 \pm 3.05	0.00 \pm 0.00	ND	ND	ND

Results are an average of percentage (9 larvae) for six different concentration ranges (for abamectin, 8 different concentrations, *n* = 72) of insecticides. FL = fiducial limit, S.E.M. = standard error of mean for mortalities, leaf damage and LD_{50} s. ppm = part per million. ND = not determined.

3.3. Identification of Possible Mechanism of Resistance Using Synergists

To establish the possible enzyme systems responsible for the pyrethroid resistance synergist assays were conducted for λ -cyhalothrin with PBO, DEM and TPP. In contrast with the observation from pre-exposure to DEM ($\chi^2 = 0.89$, $p = 0.34$) and TPP ($\chi^2 = 0.03$, $p = 0.86$) (Table S1), pre-treatment with PBO significantly recovered susceptibility ($\chi^2 = 98.35$, $p < 0.0001$) with on average more than four-fold increase in mortalities from 18.5% to ~90% in synergized bioassay (Figure 3C). The LD₅₀ plummeted down to only 0.92 ppm \pm 0.15, more than 8000 times lower than obtained in the bioassay with λ -cyhalothrin alone (Table S1). Thus, a synergistic ratio calculated for PBO+ λ -cyhalothrin was the highest, at 8154.35.

Surprisingly, for all the three synergists tested, no major difference in foraging capability was observed between the synergized tests and conventional treatment with λ -cyhalothrin (Figure 3D, Table 1).

3.4. Identification of 1014F *kdr* Mutation in Domain II of the Voltage-Gated Sodium Channel

The leucine to phenylalanine *kdr* mutation commonly reported in resistant species of diverse arthropods [39] was 100% fixed in seven λ -cyhalothrin-alive and eight dead larvae from Kadawa. Sequencing unambiguously identified TTT codon in the 1014 position (Figure 4B,C) specific to phenylalanine. The 15 sequences, which were 99% similar to the NCBI accessions JQ701800 and KY767010, have been deposited in the GenBank.

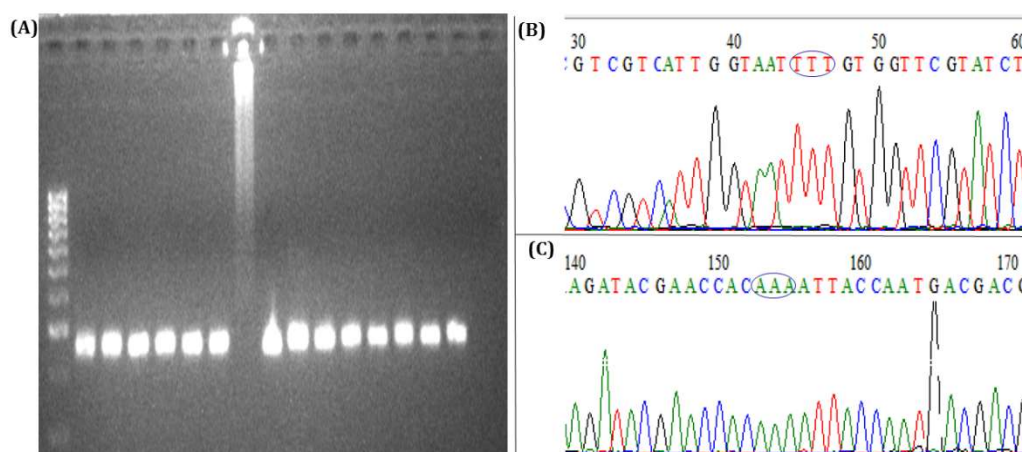


Figure 4. Genotyping of the IIS6 fragment of the voltage-gated sodium channel for the 1014 F *kdr* mutation in *T. absoluta* larvae. (A) Agarose gel of the nest 2 PCR fragment showing a characteristic band of around 270–280 bp; (B) and (C) are chromatogram traces respectively of forward and reverse sequences from a single *T. absoluta* larvae. The 1014 F codon is encircled in blue.

4. Discussion

Tomato is the second most important vegetable crop in the world, next to potato, and Nigeria is its largest producer in Africa (and ranked 14th largest producer in the world). However, *T. absoluta* invasion is threatening the sustainability of tomato farming in Nigeria, with millions of Nigerian Naira lost in the recent years. The abrupt expansion and destruction of tomatoes by the leaf miner caught Nigerian farmers and other stakeholders unprepared. Continuous invasion is reported here and there in northern Nigeria, with no clear plans on ground on how to control this pest [17]. In contrast to studies carried out elsewhere in Europe and Americas, except for few reports from Ethiopia, most of the studies done on *T. absoluta* from sub-Saharan Africa did not describe the insecticides susceptibility status of this pest. For, example, published data from Senegal [11], Burkina Faso [40], Niger [10], Tanzania [12], Angola [41], and Botswana [13] described only ecology and/or bionomics of this pest.

Pyrethroids being cheap and safest insecticides for mammals are the frontline insecticides for use against agricultural pest. However, the high pyrethroid insensitivity resistance in agricultural pests, as observed in this northern Nigerian populations resulted in a shift to other more effective but expensive pesticides. High resistance to type I pyrethroid permethrin and type II pyrethroids, λ -cyhalothrin, deltamethrin and α -cypermethrin have been previously described for the Brazilian and Iranian populations of *T. absoluta* [22,24,42]. Our finding of 1014F *kdr* mutation in all the alive and dead larvae from λ -cyhalothrin exposure was in line with previous findings. This target-site mutation has been described as 100% fixed in *T. absoluta* populations from several European countries [24].

In this study, comparable mortalities observed for the organophosphate chlorpyrifos-methyl and the carbamate propoxur, were lower than obtained with λ -cyhalothrin. Organophosphate and carbamate resistance had been described for *T. absoluta* populations previously, e.g. for chlorpyrifos and methamidophos in Iranian and Brazilian populations [42,43], and towards methamidophos in the Brazilian populations [43]. Just as established in our study the above studies described multiple resistance to pyrethroids, carbamates and organophosphates in the *T. absoluta* populations from different parts of the world [42,43]. The lowest mortalities observed towards abamectin was in keeping with a report from Argentine populations with abamectin exhibiting lowest LD₅₀ in three different populations compared to methamidophos and deltamethrin [44]. In a recent study [26] resistance to emamectin benzoate (a derivative of abamectin) has also been shown to exist in populations from Ethiopia.

Synergists PBO, DEM, and TPP have been used by Sequeira and colleagues [45] to synergize abamectin in bioassays with results implicating CYP450s, in comparison to the other two synergists with much lower synergism. This is in agreement with our finding which suggests that CYP450s could be the major drivers of metabolic resistance to λ -cyhalothrin. However, using biochemical assays of enzyme activities another study conducted with Brazilian populations of *T. absoluta* established greater correlation between pyrethroids resistance and increased levels of both monooxygenases and GSTs [22]. Yet, the presence of the 1014F *kdr* mutation in both alive and dead larvae point to potential metabolic mechanism responsible for the resistance.

5. Conclusions

In northern Nigeria, farmers desperate to control *T. absoluta* often mix and increase the amount and frequency of pesticides they apply. Unfortunately, this unscientific approach could possibly increase selective pressure in this pest populations inducing resistance. The low mortality towards multiple insecticides observed from a population of *T. absoluta* in a single field in Nigeria is of great concern. However, the highest mortality observed with respect to abamectin suggests its possible potency in the field. But this insecticide as well as diamide insecticides like chloranthraniliprole are very expensive and possibly unaffordable by subsistence farmers in Nigeria. The claim by farmers of better kill with a formulation containing a mixture of emamectin benzoate and teflubenzurone could possibly be due to lower or absence of resistance to the teflubenzurone, which is an insect growth regulator benzoylurea (IRAC Class 15, type 0). Further work need to be conducted with more replicates and more insecticides to establish the resistance profile with higher statistical power. Also, in future, sampling should be carried out at different sites spanning northern Nigeria. This will provide a comprehensive picture of the resistance in the field population of this pest and establish the control failure likelihood.

Supplementary Materials: The following are available online at <http://www.mdpi.com/2077-0472/9/1/7/s1>, Table S1: Results of synergist bioassay, for mortalities, probit analyses and leaf damage with λ -cyhalothrin. Figure S1: Results of dose-response bioassay to establish the LD₅₀ for *T. absoluta* L2 larvae with (A) λ -cyhalothrin, (B) propoxur, (C) chlorpyrifos-methyl, (D) abamectin, and (E) PBO pre-exposure with λ -cyhalothrin.

Author Contributions: Conceived and designed by S.S.I., I.B. and M.M.M. collected samples from field and carried out pesticides bioassays. S.S.I. did the molecular analyses with the help of M.M.M., S.S.I. analyzed the data with the help of N.A., I.B. and S.K.H., and wrote the manuscript. All authors read and approved the manuscript.

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