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Spatiotemporal multiple insecticide resistance in *Aedes aegypti* populations in French Guiana: need for alternative vector control

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⁵Institut Pasteur, Paris, France

BACKGROUND *Aedes aegypti* is the sole vector of urban arboviruses in French Guiana. Overtime, the species has been responsible for the transmission of viruses during yellow fever, dengue, chikungunya and Zika outbreaks. Decades of vector control have produced resistant populations to deltamethrin, the sole molecule available to control adult mosquitoes in this French Territory.

OBJECTIVES Our surveillance aimed to provide public health authorities with data on insecticide resistance in *Ae. aegypti* populations and other species of interest in French Guiana. Monitoring resistance to the insecticide used for vector control and to other molecule is a key component to develop an insecticide resistance management plan.

METHODS In 2009, we started to monitor resistance phenotypes to deltamethrin and target-site mechanisms in *Ae. aegypti* populations across the territory using the WHO impregnated paper test and allelic discrimination assay.

FINDINGS Eight years surveillance revealed well-installed resistance and the dramatic increase of alleles on the sodium voltage-gated gene, known to confer resistance to pyrethroids (PY). In addition, we observed that populations were resistant to malathion (organophosphorous, OP) and alpha-cypermethrin (PY). Some resistance was also detected to molecules from the carbamate family. Finally, those populations somehow recovered susceptibility against fenitrothion (OP). In addition, other species distributed in urban areas revealed to be also resistant to pyrethroids.

CONCLUSION The resistance level can jeopardize the efficiency of chemical adult control in absence of other alternatives and conducts to strongly rely on larval control measures to reduce mosquito burden. Vector control strategies need to evolve to maintain or regain efficacy during epidemics.

Key words: *Aedes aegypti* - insecticide resistance - French Guiana - spatiotemporal distribution
companies. As a result, deltamethrin resistance was recorded in Ae. aegypti populations in French Guiana as early as 2000.\(^{(2)}\) Pyrethroid pressure on mosquitoes and particularly on Ae. aegypti resulted in a tremendously high resistance ratio for deltamethrin.\(^{(3,4)}\) The historical succession of insecticide use has most likely contributed to the selection of resistant Ae. aegypti populations.\(^{(3)}\)

Although mechanisms responsible for insecticide resistance are complex, resistance in mosquitoes is mainly associated with target site modification and metabolic changes. Voltage-gated sodium channel trans-membrane proteins (\(\text{Na}_v\)) are essential for proper electrical signaling in the nervous system and other excitable cells. The toxic effect of pyrethroids modifies the function of \(\text{Na}_v\) by extending the activation of sodium channels and interacting with receptor sites. Mutations of pyrethroid receptor target sites in resistant mosquitoes restore this process of electrical signaling. Metabolic resistance, on the other hand, involves more subtle alterations in the expression of a complex array of enzymes and detoxification pathways. To date, studies of the resistance mechanisms in populations of Ae. aegypti in French Guiana have demonstrated that isoleucine amino-acid at the 1016 position (1016I) in the IIS6 region of Na\(_v\) and cytochrome P450 gene over-expression are strongly associated with deltamethrin resistance.\(^{(3,4,5,6)}\) Copy number variations (CNVs) was also associated with gene over-expression. In addition, allelic variations of enzymes demonstrate the presence of isoforms that permit higher levels of detoxification. At a larger scale, DDT and permethrin resistant Ae. aegypti populations exhibit a higher frequency of cysteine amino-acid at the 1534 position (1534C) of Na\(_v\).\(^{(7,8)}\) This mutation is also known to enhance deltamethrin resistance when associated to 1016I in South America.\(^{(9)}\) Furthermore, the two loci are known to be genetically linked and to have co-evolved.\(^{(10,11)}\)

The study described herein presents the results of insecticide resistance monitoring in French Guiana from 2010 to 2015. Spatial and temporal distribution of alleles at the 1016 and 1534 positions of Na\(_v\) protein are reported in wild Ae. aegypti populations to assess the distribution and evolution of these mutations in French Guiana.

**Ethics statement** - Mosquito blood feeding was done on mice. Experiments were authorised by the agreement number B973-02-01 delivered by “La préfecture de Guyane” and renewed on June 6th, 2015. The protocol for the use of mice was approved by the ethical committee C-E-TEA Institut Pasteur (n° 89), report number 2015-0010 issued on May 18th, 2015.

**MATERIALS AND METHODS**


Populations of Culex quinquefasciatus were collected in the larval stage in Kourou, Saint Georges de l’Oyapock and Rémire-Montjoly in 2016. Adults of the F0 generation were reared and allowed to mate. Females were provided a blood meal from mice to collect eggs of the F1 generation. Adults of the F1 generation were then obtained by rearing immature stages concurrently under insectary conditions (28 ± 1°C, 70 ± 10% RH, 12:12h photoperiod) in French Guiana.

Aedes taeniorhynchus females were collected in Cayenne, fed on mice and allowed to lay eggs. Following generations were allowed to reproduce as previously mentioned. The F2 generation of that species was used for testing.

**Assessment of insecticide resistance -** Insecticide resistance bioassays were carried out for each population through tarsal contact tests using filter papers impregnated with technical grade compounds (Sigma-Aldrich, St Louis, MO, USA): deltamethrin (CAS # 52918-63-5) (Sigma-Aldrich, St Louis, MO, USA) at the diagnostic dose of 0.06% for Aedes sp. and 0.025% for Culex sp.,
alpha-cypermethrin (#45806, Sigma-Aldrich) at the diagnostic dose of 0.05%, malathion (#45614, Sigma-Aldrich) at the dose of 0.8 and 5%, fenitrothion (#45487, Sigma-Aldrich) at the diagnostic dose of 0.5%, propoxur (#45644, Sigma-Aldrich) at the diagnostic dose of 0.1% and bendiocarb (#45336, Sigma-Aldrich) at the diagnostic dose of 0.1%. Filter papers were impregnated using acetone solutions of insecticide and silicone oil or olive oil\(^{(13)}\) by dropping 2 mL of the solution onto each paper (12 x 15 cm). Doses are expressed in weight/volume (w/v) percentage of the active ingredient in silicone oil. The papers were dried for 24 h before the test. For each population, four batches of 25 non-blood fed, 2-5 day old females were introduced into holding tubes for one hour then transferred into the exposure tube and placed vertically for 1 h.\(^{(13)}\) Knocked-down (\(kd\)) and dead mosquitoes were recorded after this time (1 h KD) before being transferred back to the holding tubes. A mosquito was recorded as knocked-down if it was lying on its back or side and was unable to instigate flight after a gentle tap. Mortality was recorded 24 h after exposure (24 h M). Controls, made of only acetone and silicone oil delivered to the filter paper, were performed as mentioned above with a total of two replicates of 25 non-blood fed 2-5 day old females. Females from PAEA and New Orleans reference strains were raised under identical insectary conditions and used as susceptible controls. All replicates were conducted at 27 ± 2°C and 60 ± 10% RH. A ten percent sugar solution was provided to the females during the 24-h observation period in holding tubes.

WHO protocols for insecticide resistance monitoring provide a diagnostic threshold of resistance.\(^{(13)}\) Any population is considered susceptible when the percentage of mortality is above 98% after exposure to an impregnated paper at the diagnostic dose. Mortality below 98% is suggestive of resistance; below 90%, resistance is confirmed; between 90% and 97% tests must be triplicated and individuals tested for molecular resistance markers. When several tests were performed during one-year results of mortality or \(kd\) were aggregated by means.

**Genotyping V1016I and F1534C loci** - Each adult mosquito was grinded individually by hand with a sterile grinder. DNA was extracted using either the Purelink Genomic DNA extraction kit (Invitrogen, Carlsbad, CA, USA) or DNeasy Blood & Tissue Kit (Qiagen, Hilden, Germany) according to the manufacturer’s instructions or Collins et al.\(^{(14)}\) The DNA isolated was quantified using a spectrophotometer (Nanodrop 2000c, Thermo Scientific, Waltham, MA, USA) and a 10-fold dilution was used for polymerase chain reaction (PCR). DNA was then stored at -20°C.

Mosquitoes from each locality were genotyped at both 1016 (Val/Leu) and 1534 (Cys/Phenylalanine) positions from genomic DNA by real time PCR allelic discrimination assay (Applied Biosystems, CA, USA). Two primers and two minor groove binding probes were designed for V1016I detection as previously described.\(^{(8)}\) A second amino acid substitution, F1534C, was targeted on the IIS6 region of the Na\(_{a}\) gene according to Yanola et al.\(^{(9)}\) Primers and probe sequences are listed in Supplementary data (Table I). Each Allelic Discrimination assay was carried out with 12.5 µL of 2X TaqMan® Universal Master Mix II (Life Technologies, Gaithersburg, MD, USA), 1.44 µM of primers, 0.4 µM of each of the specific probes and 3 µL of genomic DNA (20 ng), in a total volume of 25 µL. Thermocycling amplification conditions were 10 min at 95°C followed by 45 cycles of 95°C for 15 s and 60°C for 1 min. The amplification reactions were performed in a StepOnePlus Real-Time PCR System (Life Technologies, Gaithersburg, MD, USA) and analyses were done by the StepOne Software version 2.1.

The cor.test function in R 3.2.2 (R Development Core Team, Vienna, Austria) was used to find the \(r\) correlation between percent mortalities and frequency of alleles or genotypes. Dilocus haplotypes 1016/1534 appear in four possible forms: V1016I/F1534C (VF), V1016I/C1534C (VC), I1016/F1534C (IF), I1016/C1534C (IC). In double heterozygotes, the bi-allelic combination is unknown. Therefore, the composite disequilibrium frequencies and tested linkage disequilibrium were determined as described previously.\(^{(11)}\) In addition, variance analysis was performed to evaluate the effect of each genotype on the resistance phenotypes using aov function in R 3.2.2 (R Development Core Team, Vienna, Austria). Data from Dusfour et al.\(^{(9)}\) were used as a baseline for spatio-temporal analyses.

**RESULTS**

**Selection of deltamethrin resistance in French Guiana** - Since 2009, resistance monitoring has been conducted for *Ae. aegypti* populations across their distribution in French Guiana in a total of 15 localities (Fig. 1). Nine of the populations have been monitored two or more times since 2009, resistance monitoring has been conducted for *Ae. aegypti* populations across their distribution in French Guiana in a total of 15 localities (Fig. 1). Nine of the populations have been monitored two or more times.

<table>
<thead>
<tr>
<th>Alleles/genotypes</th>
<th>1 h KD (r)</th>
<th>p-value</th>
<th>CI 95%</th>
<th>24 h M (r)</th>
<th>p-value</th>
<th>CI 95%</th>
</tr>
</thead>
<tbody>
<tr>
<td>1534C</td>
<td>-0.76</td>
<td>0.0001</td>
<td>-0.90; -0.47</td>
<td>-0.76</td>
<td>0.0002</td>
<td>-0.90; -0.46</td>
</tr>
<tr>
<td>1016I</td>
<td>-0.89</td>
<td>&lt; 0.0001</td>
<td>-0.95; -0.72</td>
<td>-0.80</td>
<td>&lt; 0.0001</td>
<td>-0.92; -0.53</td>
</tr>
<tr>
<td>1016I/1534C</td>
<td>-0.80</td>
<td>&lt; 0.0001</td>
<td>-0.92; -0.54</td>
<td>-0.72</td>
<td>0.0007</td>
<td>-0.88; -0.38</td>
</tr>
</tbody>
</table>

CI 95%: confidence interval 95%.
times over that period (Fig. 1). The more populated towns [Saint Laurent du Maroni (SLM), Kourou (KOU), Cayenne (CAY), Saint Georges de l’Oyapock (SGO)] were sampled several times with fewer collections in small and remote villages. Prior monitoring showed that deltamethrin resistance was already established in 2008-2009 in the main coastal towns. Consecutive follow-up in hamlets in littoral zones [Matiti (MAI), Saut Sabbat (SSA)] or in remote inland villages [Loka (LOK), Maripasoula (MPS)] revealed that high resistance was widespread

### TABLE II

Mean percentages of mortality and standard deviation at 24 h for impregnated paper tests per insecticide, *Aedes aegypti* population and year. The total number of exposed mosquitoes are indicated (n). Data from the years 2008 and 2009 are from Dusfour et al. [3]

<table>
<thead>
<tr>
<th>Molecule</th>
<th>Dose</th>
<th>Population</th>
<th>Year</th>
<th>n</th>
<th>Mean 24 h M (%)</th>
<th>SD (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Alphacypermethrin</strong></td>
<td>0.05%</td>
<td>CAY</td>
<td>2016</td>
<td>100</td>
<td>5.00</td>
<td>5.03</td>
</tr>
<tr>
<td></td>
<td></td>
<td>KOU</td>
<td>2016</td>
<td>100</td>
<td>3.00</td>
<td>3.82</td>
</tr>
<tr>
<td></td>
<td></td>
<td>MPS</td>
<td>2016</td>
<td>100</td>
<td>0.00</td>
<td>0.00</td>
</tr>
<tr>
<td></td>
<td></td>
<td>SGO</td>
<td>2016</td>
<td>100</td>
<td>0.00</td>
<td>0.00</td>
</tr>
<tr>
<td><strong>Bendiocarb</strong></td>
<td>0.1%</td>
<td>MPS</td>
<td>2016</td>
<td>100</td>
<td>91.00</td>
<td>5.03</td>
</tr>
<tr>
<td></td>
<td></td>
<td>SGO</td>
<td>2016</td>
<td>100</td>
<td>96.00</td>
<td>3.26</td>
</tr>
<tr>
<td><strong>Propoxur</strong></td>
<td>0.1%</td>
<td>CAY</td>
<td>2011</td>
<td>102</td>
<td>13.69</td>
<td>1.95</td>
</tr>
<tr>
<td></td>
<td></td>
<td>MPS</td>
<td>2016</td>
<td>100</td>
<td>76.00</td>
<td>3.26</td>
</tr>
<tr>
<td></td>
<td></td>
<td>SGO</td>
<td>2011</td>
<td>100</td>
<td>76.00</td>
<td>3.26</td>
</tr>
<tr>
<td><strong>Fenitrothion</strong></td>
<td>0.5%</td>
<td>MPS</td>
<td>2012</td>
<td>98</td>
<td>100.00</td>
<td>0.00</td>
</tr>
<tr>
<td></td>
<td></td>
<td>SGO</td>
<td>2008</td>
<td>321</td>
<td>29.30</td>
<td>18.54</td>
</tr>
<tr>
<td></td>
<td></td>
<td>SLM</td>
<td>2009</td>
<td>199</td>
<td>85.40</td>
<td>7.98</td>
</tr>
<tr>
<td></td>
<td></td>
<td>CAY</td>
<td>2014</td>
<td>196</td>
<td>31.88</td>
<td>26.47</td>
</tr>
<tr>
<td></td>
<td></td>
<td>KOU</td>
<td>2014</td>
<td>192</td>
<td>9.07</td>
<td>7.81</td>
</tr>
<tr>
<td></td>
<td></td>
<td>MAT</td>
<td>2014</td>
<td>124</td>
<td>20.86</td>
<td>11.7</td>
</tr>
<tr>
<td></td>
<td></td>
<td>SGO</td>
<td>2008</td>
<td>99</td>
<td>12.12</td>
<td>3.27</td>
</tr>
<tr>
<td></td>
<td></td>
<td>SLM</td>
<td>2015</td>
<td>98</td>
<td>10.25</td>
<td>7.80</td>
</tr>
<tr>
<td><strong>Malathion</strong></td>
<td>0.8%</td>
<td>CAY</td>
<td>2014</td>
<td>195</td>
<td>99.47</td>
<td>1.47</td>
</tr>
<tr>
<td></td>
<td></td>
<td>KOU</td>
<td>2015</td>
<td>95</td>
<td>4.16</td>
<td>3.40</td>
</tr>
<tr>
<td></td>
<td></td>
<td>MAT</td>
<td>2014</td>
<td>124</td>
<td>20.86</td>
<td>11.7</td>
</tr>
<tr>
<td></td>
<td></td>
<td>SGO</td>
<td>2014</td>
<td>191</td>
<td>8.89</td>
<td>5.52</td>
</tr>
<tr>
<td></td>
<td></td>
<td>SLM</td>
<td>2015</td>
<td>98</td>
<td>10.25</td>
<td>7.80</td>
</tr>
<tr>
<td><strong>Malathion</strong></td>
<td>5%</td>
<td>CAY</td>
<td>2014</td>
<td>199</td>
<td>98.5</td>
<td>2.07</td>
</tr>
<tr>
<td></td>
<td></td>
<td>KOU</td>
<td>2015</td>
<td>94</td>
<td>94.65</td>
<td>5.20</td>
</tr>
<tr>
<td></td>
<td></td>
<td>MAT</td>
<td>2015</td>
<td>95</td>
<td>100.00</td>
<td>0.00</td>
</tr>
<tr>
<td></td>
<td></td>
<td>SGO</td>
<td>2014</td>
<td>100</td>
<td>100.00</td>
<td>0.00</td>
</tr>
<tr>
<td></td>
<td></td>
<td>SLM</td>
<td>2015</td>
<td>98</td>
<td>100.00</td>
<td>0.00</td>
</tr>
</tbody>
</table>

CAY: Cayenne; KOU: Kourou; MPS: Maripasoula; SGO: Saint Georges de l’Oyapock; SLM: Saint Laurent du Maroni; MAT: Matoury. SD: standard deviation.
across the territory with monitoring results showing mortality below 40% mainland. The most noticeable result is the susceptibility observed from a mosquito population on Iles du Salut (IDS) in 2013 with a subsequent loss of susceptibility the following years [Fig. 1, Supplementary data (Table II)]. In the main towns, mortality and knock-down effect dropped below 5% for CAY, KOU and SLM.

**Resistant alleles and double resistant genotypes are reaching fixation** - A total of 621 and 567 samples were genotyped for loci 1534 and 1016 respectively, and 550 with the double genotyping. Overall, observed frequencies of the resistant 1016I varied from 0.52 to 1.00 while those of 1534C allele ranged from 0.76 to 1.00 in continental populations [Supplementary data (Table III), Fig. 2]. As expected, 1016I frequencies were lower on IDS at 0.13 and 0.40 in 2013 and 2014 respectively. Frequencies of 1534C in IDS were 0.31 and 0.56 in 2013 and 2014 respectively.

In 2009, 1534C frequencies were already high, above 0.80, in the main town and have almost reached fixation in SLM, KOU and CAY as of the last recorded assay (Fig. 2). 1016I frequencies were lower than 1534C for those populations and tended to increase over time. SLM populations exhibited high frequencies throughout all periodic assays (> 0.80). In examining the whole dataset, independent frequencies of those alleles correlated negatively with percent mortality and knocked-down mosquitoes in resistance assays (Table I). As expected, significant linkage disequilibrium was observed between the two loci (p < 0.001). Therefore, both loci were combined for further analyses. Biallelic resistant haplotype increased in the same pattern as independent alleles and is also significantly correlated with the phenotypes (Table I).

Double mutant genotype (CCII) is known to confer a higher selective advantage under insecticide pressure [Supplementary data (Table III), Fig. 3]. This highly resistant genotype was maintained or increased over time the last decade, in accordance with deltamethrin phenotype of mortality (r = -0.73, p-value < 0.0001) and knock-down (r = -0.82, p-value < 0.0001) [Supplementary data (Table III)]. IDS and KOU are the only populations exhibiting double homozygote susceptible but whose frequency decreased.

Variance analysis statistically confirmed the dominant effect of CCII genotype but also showed that CCVI, FCVV and FFVV have an effect on kd phenotype of the populations while CCII, CCVI, FCII and FCVV contribute to the mortality phenotypes. Overall genotypes contribute higher to kd phenotypes than to mortality [Supplementary data (Table IV)].

**Ae. aegypti phenotypes for other insecticides** - Over the years, sporadic checks of resistance to other molecules from the main insecticide families have been performed (Table II). 24-h mortalities against alpha-cypermethrin were even lower than the score for deltamethrin mortality. Bendiocarb (carbamate) demonstrated a loss of susceptibility while propoxur from the same insecticide family demonstrated resistance according to WHO recommended thresholds. *Ae. aegypti* populations seemed to regain susceptibility against fenitrothion in CAY and SGO where they were assayed at several time points. Susceptibility for that molecule remained in MPS between 2012 and 2016 while resistance has not changed in KOU.

Inconsistencies occurred in the follow-up on malathion resistance. Indeed, the diagnostic dose of 0.8% did not succeed in killing 100% of the reference strains (data not shown). The diagnostic dose was then complemented by a 5% dose, formerly used for that species and currently applied for other mosquito species. Then malathion resistance was confirmed to be recorded in *Ae. aegypti* populations from French Guiana.

**Resistance in other urban mosquitoes** - In a context of emergent arboviruses, the resistance of two other urban mosquito species was tested against deltamethrin in 2016 and 2017 (Table III). In absence of any reference strain for *Ae. taeniorhynchus*, the diagnostic dose established for *Ae. aegypti* was used. The test detected resistance in the population from Cayenne. Populations of *Cx. quinquefasciatus* from Kourou, Rémiere-Montjoly and Saint Georges de l’Oyapock also presented high resistance to deltamethrin.

**DISCUSSION**

Monitoring resistance to insecticides is a key component in sustainable vector control and the development of a resistance management program. French Guiana is an overseas territory of France, located in South America but subject to European and French biocide regulations. In the absence of a vaccine or treatment for arboviruses, *Ae. aegypti* control had been implemented by reducing

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**TABLE III**

Mean percentage of kd effect and mortality with standard deviation for impregnated paper tests with deltamethrin 0.06% (*Aedes*) and 0.025% (*Culex*) per population. The total number of exposed mosquitoes are indicated (n).

<table>
<thead>
<tr>
<th>Species</th>
<th>Locality</th>
<th>n</th>
<th>% 1 h KD</th>
<th>SD (%)</th>
<th>% 24 h M</th>
<th>SD (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>Aedes taeniorhynchus</em></td>
<td>CAY</td>
<td>101</td>
<td>46.93</td>
<td>20.02</td>
<td>17.14</td>
<td>11.10</td>
</tr>
<tr>
<td></td>
<td>KOU</td>
<td>100</td>
<td>1.00</td>
<td>2.00</td>
<td>1.00</td>
<td>2.00</td>
</tr>
<tr>
<td></td>
<td>REM</td>
<td>97</td>
<td>14.43</td>
<td>3.06</td>
<td>5.17</td>
<td>2.51</td>
</tr>
<tr>
<td></td>
<td>SGO</td>
<td>100</td>
<td>3.00</td>
<td>3.80</td>
<td>9.00</td>
<td>3.82</td>
</tr>
</tbody>
</table>

CAY: Cayenne; KOU: Kourou; REM: Rémiere-Montjoly; SGO: Saint Georges de l’Oyapock. SD: standard deviation.
or treating breeding sites all year, and by insecticide spraying during dengue and other arbovirus transmission. After the reintroduction of DDT-resistant *Ae. aegypti* population in the 1950s, malathion, an organophosphorous, was used for outdoor spatial spraying for decades. Later, deltamethrin, a molecule from the pyrethroid family, was also used for indoor spatial sprays around dengue cases. Temephos (organophosphorous) was used as larvicide succeeded by *Bacillus thuringiensis var israelensis* in 2000 along with larval source reduction measures. In 2011, deltamethrin became the only authorised and available insecticide for indoor and outdoor adult mosquito control. In 2014, a chikungunya outbreak hit the territory and *Aedes* control was intensified with weekly ground-spatial sprays of deltamethrin but also with malathion, exceptionally implemented for six months during this epidemic.\(^{(2)}\) The Zika outbreak followed in 2016, during which only deltamethrin was used for spraying. These periodic outbreaks with intensified insecticide use are the times when heavy selective pressure occurs, but other indirect sources of pyrethroid exposure exist. In fact, control of malaria or pest mosquitoes is conducted with recurring indoor residual spraying, distribution of pyrethroid-impregnated bed nets and spatial spraying which also exert non-negligible insecticide pressures on the mosquito populations. In addition, the use of household sprays is frequent and suspected to play a role in resistance selection by pyrethroids in urban mosquito populations.\(^{(16)}\) No robust data exists to support the foregoing hypothesis in French Guiana before 2009. The selection caused by insecticides in urban environments needs to be studied as a whole to explain resistance patterns of current and potential vectors at local and global scales.\(^{(17,18)}\)

Even though it is difficult to decipher the dominant factor inducing resistance to deltamethrin, high levels were already recorded in *Ae. aegypti* populations at the early stage of its use for spatial spraying in 2011.\(^{(3)}\) At that time, both resistance to deltamethrin and DDT were documented in French Guiana.\(^{(2)}\) Cross resistance is often observed between pyrethroids and DDT\(^{(19,20)}\) due to the same target sites of these molecules, even if recent meta-analysis data in the geographical region does not concur.\(^{(21)}\) The continuous pressure by either one or both molecules before 2011 probably maintained the level of resistance to deltamethrin and other pyrethroids in *Ae. aegypti* populations and other urban mosquitoes in areas where these compounds are used for malaria control by indoor residual spraying (i.e. Saint Georges de l’Oyapock and Saint Laurent du Maroni). However, low gene flow and local selection of resistance should have shaped different resistance patterns amongst populations.\(^{(22,23)}\) In contrast, malathion and temephos were the mainstay of vector and pest mosquito control in the central littoral strip, where malaria is a rare event for the last 40 years.

However, spatio-temporal monitoring of deltamethrin resistance demonstrates the widespread distribution of the phenotypes and its important increase globally, reaching mortality lower than 5% in 2015-2016 with the exception of the population on Iles du Salut, whose resistance development appears later than on the mainland. This archipelago is a unique location where vector or pest control is rarely performed and is connected with Kourou on the continent by boat and helicopter only.

Our monitoring also showed a great variability amongst results probably a consequence of the intensification of domestic, and uncontrolled, use of pyrethroids during periods of high infestation as already been evidenced.\(^{(24)}\)

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Fig. 2: frequencies of resistant alleles at 1534 (C) and 1016 (I) loci separately and combined (IC), per year and per site. MPS: Maripasoula; APA: Apatou; SLM: Saint Laurent du Maroni; SSA: Saut Sabbat; KOU: Kourou; IDS: Iles du Salut; CAY: Cayenne; SGO: Saint Georges de l’Oyapock.
The investigation of molecular markers related to resistance at positions 1016 and 1534 of the Na\textsuperscript{+} gene demonstrated a strong correlation between resistant alleles alone or in combination with percent of knocked-down mosquitoes and mortality. These results are in agreement with previous data observed in French Guiana.\textsuperscript{4,5,6} The resistant 1534C was already at high frequencies, almost to fixation, in 2009 compared to 1016I, whose frequencies were below 50% across the studied populations except for Saint Laurent du Maroni, where frequency was already above 90%. This observation has been made in other countries where 1534C is suspected to have appeared before 1016I.\textsuperscript{11} The appearance of 1534C could be associated with DDT resistance to which that mutation confers a selective advantage greater than for pyrethroids.\textsuperscript{11} The rapid expansion of 1016I is then related to the co-evolution of the two loci, the presence of 1534C and in particular the genotype CCIV in initial populations had most likely been crucial as the double homozygote resistant gives a strong advantage under deltamethrin pressure.\textsuperscript{9}

In addition, resistant alleles are partially recessive and the presence of heterozygous genotypes do not confer full resistance.\textsuperscript{10,25} Four different patterns can be observed in genotypes. First, the population from Saint Laurent du Maroni was already close to fixation in 2009. Then, genotype evolution in Saint Georges de l’Oyapock was composed of more frequent double heterozygotes or 1534 heterozygotes than in other locations. In addition, Cayenne and Kourou, in the centre of the littoral strip, have similar patterns of evolution with the rapid expansion of double resistant homozygotes while C1534 was already in place. Finally, Iles du Salut was drastically different and the double homozygotes susceptible recorded in 2013 decreased to favor heterozygote or homozygote resistant individuals. The primary cause is most likely genetic structure among populations of French Guiana, inducing a local selection of resistance and therefore the development of different patterns.\textsuperscript{22,23} These selective pressures with either pyrethroids or DDT are not easy to unravel. However, the fact that Saint Laurent du Maroni and Saint Georges de l’Oyapock have been subjected to both malaria vector control and \textit{Aedes} control must be noted. In opposite, Kourou and Cayenne were poorly impacted by malaria control due to low transmission. A few malaria hot-spots were sporadically treated in both towns. Iles du Salut is a unique setting without spraying except occasional pest treatment. Another study shows local structure of resistance in \textit{Ae. aegypti}, reinforcing the difficulty to monitor and predict their development.\textsuperscript{26} Even with the high importance of kdr mutation in deltamethrin phenotype resistance, previous investigations have demonstrated the implication of detoxification enzymes in French Guiana populations and particularly genes from the CYP450 gene.\textsuperscript{4,5,6}

Recent data on those particular populations demonstrate that other CYP450 genes are also associated with fenitrothion (organophosphorous) and bendiocarb (carbamates)\textsuperscript{27} resistance in the absence of mutations on the acetylcholinesterase gene,\textsuperscript{4} the target of organo-
phosphorous and carbamates. This result concurs with the low cross-resistance between deltamethrin and bendiocarb and propoxur and the susceptibility recovery for fenitrothion we observed. Cross-resistance was also observed between pyrethroid, malathion and temephos whose resistance selection by one another has been demonstrated.28,29,30 The persistence of malathion resistance in the presence of pyrethroid pressure and vice-versa before 2009 in French Guiana support these observations.

Beyond the Ae. aegypti populations resistance patterns, resistance to pyrethroids occurs for Cx. quinquefasciatus, and Ae. taeniorhynchus, pest mosquitoes in French Guiana but competent to transmit other arboviruses.

Effective urban mosquito control with pyrethroids has come to a dead-end in French Guiana due to high levels of resistance across the country. Alternative methods of vector and pest control should be investigated to eventually regain susceptibility, reinforce larval control and support a resistance management plan.

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AUTHORS’ CONTRIBUTION

AG, PN, JR, LM, PG, JI, RC and YE contributed to field and lab works; NP, FC, YE, RG and ID conceived the experimental plan; ID wrote the initial manuscript and all co-authors read, edited and commented the paper.

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