

Regional Situation of Resistance to Antimalarials

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Molecular Markers of Drug Resistance Surveillance Update -Brazil and Haiti



Update on Roraima, Brazil

Haiti



Molecular Surveillance Study Partners in Roraima State, Brazil



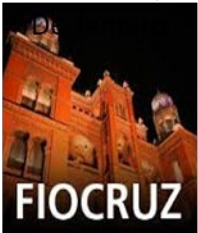
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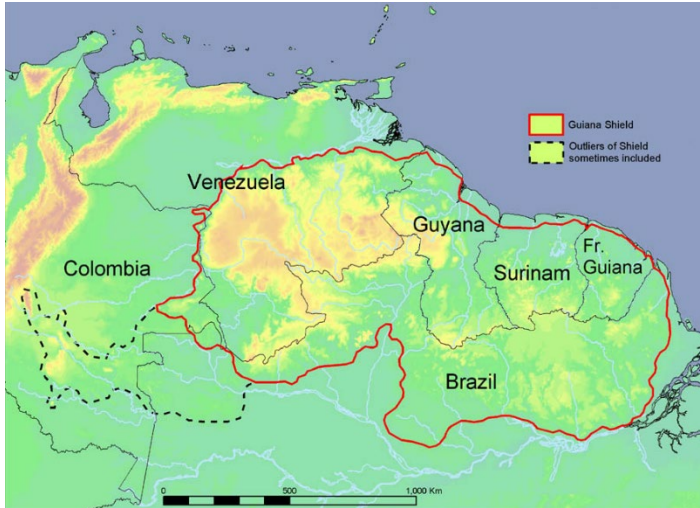
Institute Oswaldo
Cruz- Fiocruz, Rio



Federal University
of Roraima



Is there any migration of artemisinin resistant K13 markers to Roraima State, Brazil?



Molecular Surveillance Study Sites



- Three sites in the study:
 - Pacaraima
 - Boa Vista
 - Rorainópolis
- Collect filter paper samples from enrolled patients:
 - Malaria diagnosis
 - Molecular resistance markers
 - *pfK13*, *pfCRT* sequencing
 - *pfmdr1*: gene duplication and SNP
 - Neutral microsatellite analysis

Total number of samples collected 2016-Feb 2018

751 enrolled patients

346: Pacaraima

348: Boa Vista

57: Rorainapolis

Dried blood spots collected for molecular diagnosis and characterization of drug resistance markers

275 *P. falciparum* positives (including mixed Pf/Pv)

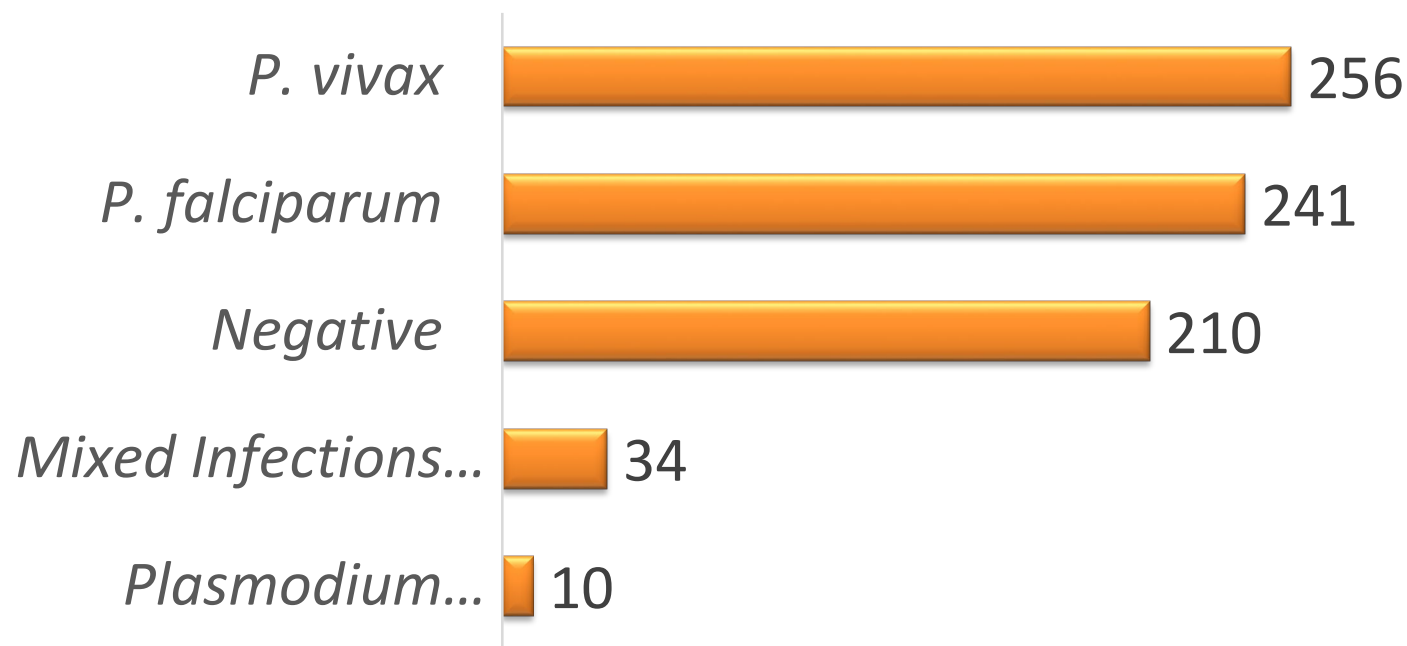
Kelch 13 SNPs

Pfmdr1 SNPs

Pfcrt SNPs

Pfmdr1 CNV

***Plasmodium* Species Detected by PET-PCR in 2016-Feb 2018**



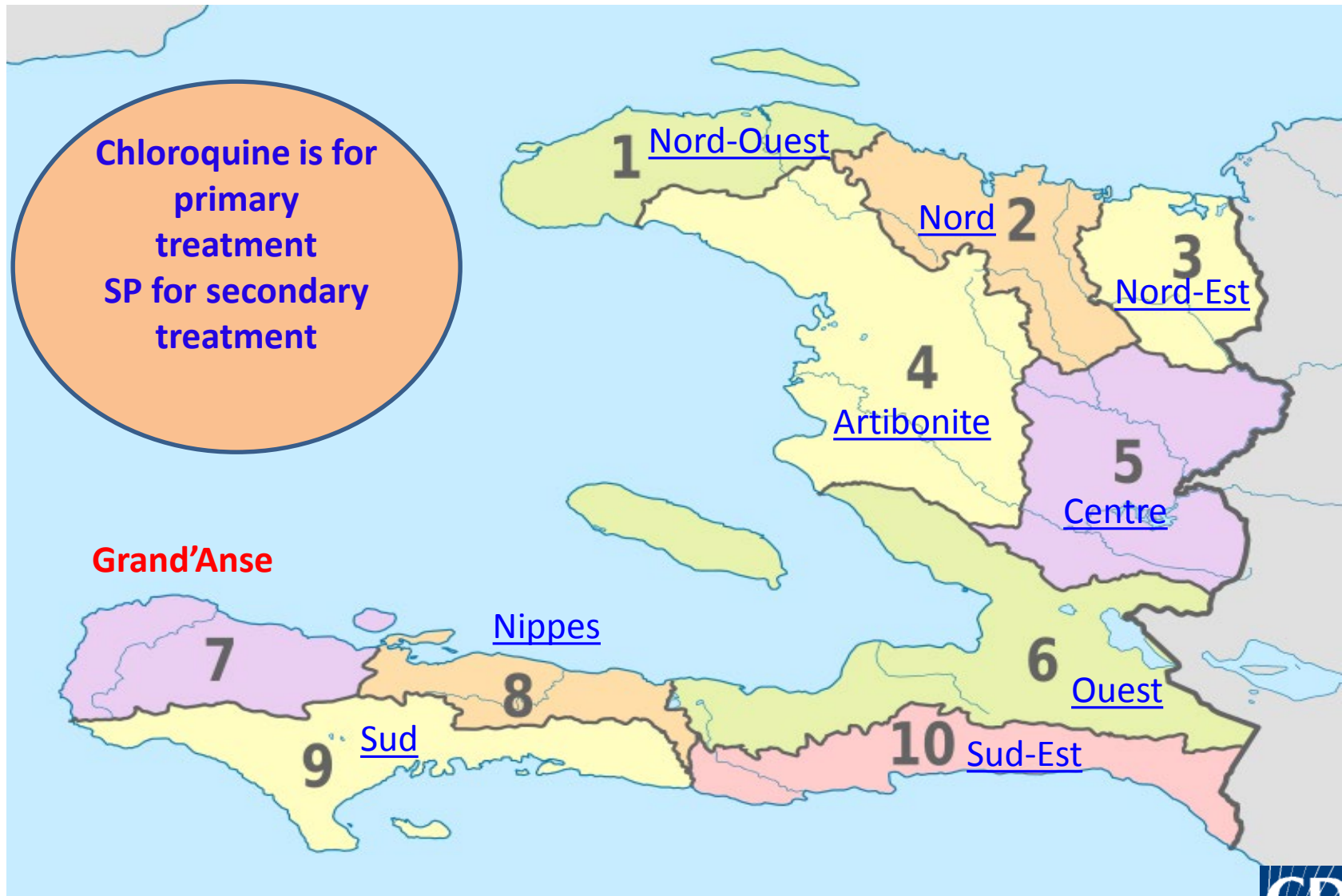
Preliminary results: molecular markers of resistance

Gene/Codons assayed	Number tested successfully	Mutations observed
<i>PfK13</i> C580Y; R539T; Y493H; I543T	238	No mutations
<i>Pfmdr1</i> N86Y; Y184F; S1034C; N1042D; D1246Y	225	184F/1042D/1246Y (45; 20%) 184F/1034C/1042D/1246Y (180; 80%)
<i>Pfcrt</i> C72S; M74I; N75E; K76T	209	SVMNT =200; 96% CVMNK = 4; 2% SVMNK =3; 1% CVIET =2; 1%
<i>Pfmdr1</i> Copy number variation (CNV)	246	6 (2.4%) with two copies

Preliminary results: molecular markers of resistance

- **We did not observe any artemisinin resistance associated mutations in the *pfk13* propeller domain.**
 - These results suggest that artemisinin resistance mutation (C580Y) found in Guyana has not been detected yet in the Roraima State of Brazil.

Surveillance for Drug Resistance in Haiti



Summary of CQ and SP Resistant Markers in Haiti -2016-17

- CQ resistant pfprt mutants (codons 72-76) were 0/741 (Rogier E., et al, unpublished)
- Pyrimethamine resistant dhfr triple (51I, 59R, 108N) mutants were 1/548 and 108N single mutants were 33%
- Sulphadoxine resistant triple (437G, 540E, 581G) mutants were 0/406 and a single 437G mutant were found
- This is consistent with current CQ treatment policy and forms a baseline data for monitoring MDA (using SP) as part of Malaria Zero project

Training -Peruvian National Institute of Health

Mr. Carlos A Bartra More

Responsable de Laboratorio

**Laboratorio de Referencia Supranacional de
Malaria, Instituto Nacional de Salud, Peru**

June 25th-Aug 3rd, 2018

Sequencing methods:

Artemisinin resistant markers- pfK13

Other resistant markers-pfmdr1

Population based markers:

**Microsatellite markers for differentiating
recrudescence from reinfection**



Training -Planned for Colombia 2019

Ms. Angela Patricia Guerra Vega
National Reference Laboratory
Parasitology Group
National Institute of Health
Bogota, Colombia

March 18-April 27th, 2019

Sequencing methods:

Artemisinin resistant markers- pfK13

Other resistant markers-pfmdr1

Population based markers:

**Microsatellite markers for differentiating
recrudescence from reinfection**



Summary, Challenges and Future Directions

- **Molecular markers and tools are complimentary tools for elimination efforts**
- **Valuable in detecting resistant markers, tracking parasite migration, emerging novel strains, HRP2 deletion, evolutionary changes etc**
- **Molecular data bases of population based markers are not available**
- **Long term data collection is critical**
- **Local capacity building is needed; academic, NGO, and MOH partnership is essential**



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