



Yellow Fever Update

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Disclosure

- No conflict regarding YF Vaccine
- Hold a patent regarding pharmacological treatment of YF using MAP Kinases inhibitors
- Honorary payments or grant support regarding dengue vaccine from: Sanofi-Pasteur and/or Butantā-NIH Vaccine
- Thanks to Dr. Marcos Freire - FIOCRUZ and Prof. Betania Drumon -UFMG for sharing some data used in this presentation

**BASIC YF DATA COMES FROM EARLY
XX CENTURY**

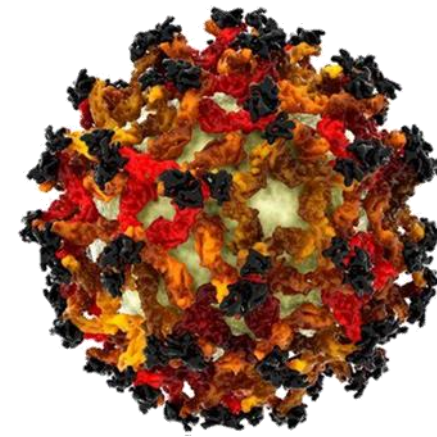


“I THANK GOD that I did not accept anybody’s opinion on this subject, but determined to put it to a through test with human beings in order to see what would happen... actual trial proven that I was right...” - Walter Reed

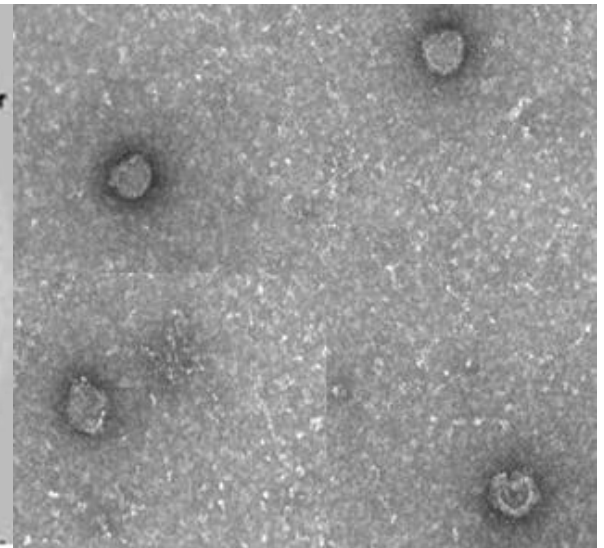
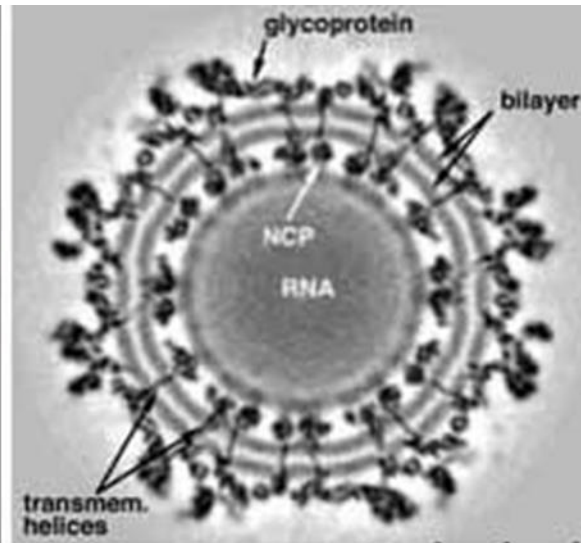
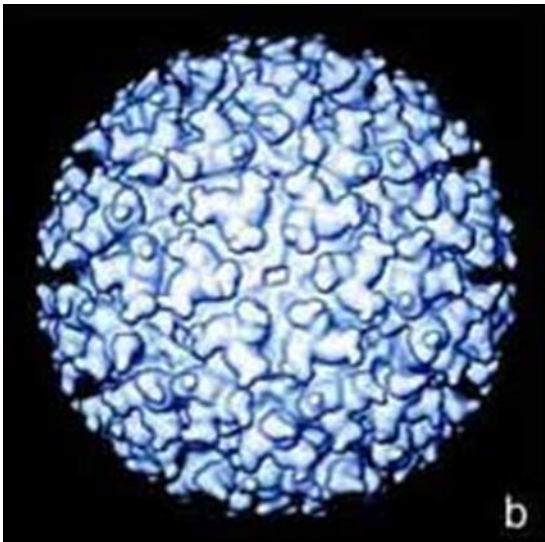
“The Etiology of Yellow Fever an Additional Note,” read before the Pan-American Medical Congress at Havana, in February, 1901

“1. The mosquito – *C. facciatus* – serves at the intermediate host for the parasite of yellow fever.^[SEP]“2. Yellow fever is transmitted to the nonimmune individual by means of the bite of the mosquito that has previously fed on the blood of those sick with this disease.^[SEP]“5. Yellow fever can also be experimentally produced by the subcutaneous injection of blood taken from the general circulation during the first and second days of this disease.^[SEP]“8. Yellow fever is not conveyed by fomites, and hence disinfection of articles of clothing, bedding, or merchandise, supposedly contaminated by contact with those sick with this disease, is unnecessary.^[SEP]“10. The spread of yellow fever can be most effectually controlled by measures directed to the destruction of mosquitoes and the protection of the sick against the bites of these insects.”

YELLOW FEVER VIRUS



Etiological agent:



Transmission Cycles of yellow fever

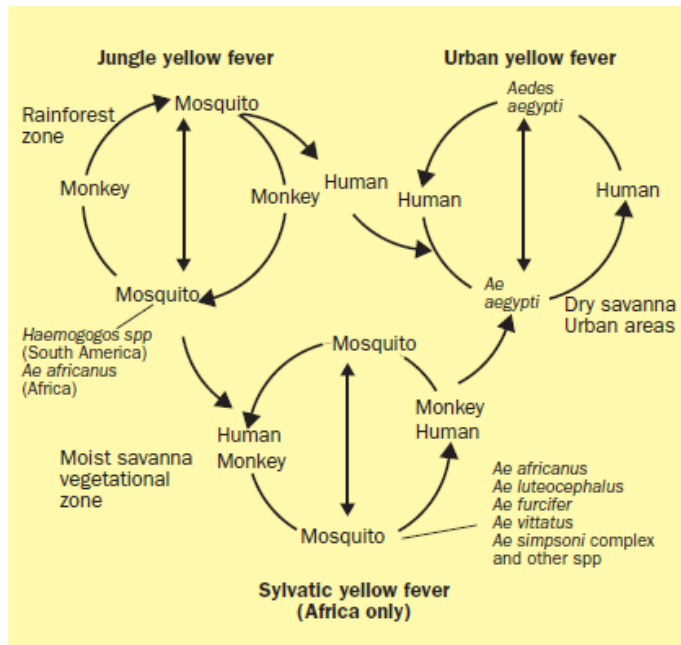
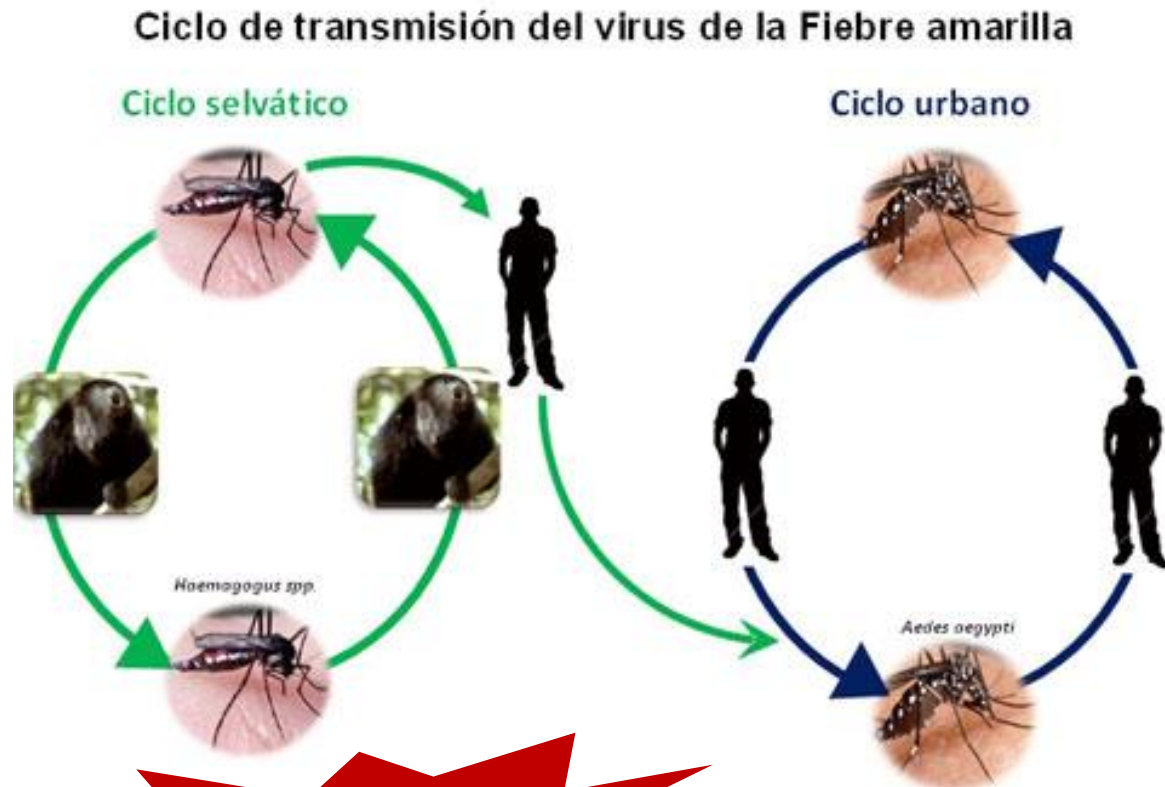


Figure 3. The transmission cycles of yellow fever. The virus is maintained by transmission between monkeys and tree-hole breeding mosquitoes. Human beings acquire "jungle yellow fever" when exposed to the bite of mosquitoes that have previously fed on an infected monkey. The vectors and ecology differ in Africa and South America. In Africa, tree-hole breeding *Aedes* spp reach high densities in the moist savanna vegetational zone and transmit the virus between people. In both continents, *Ae aegypti*, which breeds in and around houses in man-made containers, is responsible for interhuman transmission of "urban" yellow fever virus.



YF: impossible to eradicate



Alouatta sp
(guariba, bugio)

Cebus sp
(macaco prego)



Callithrix sp
(mico, soim)

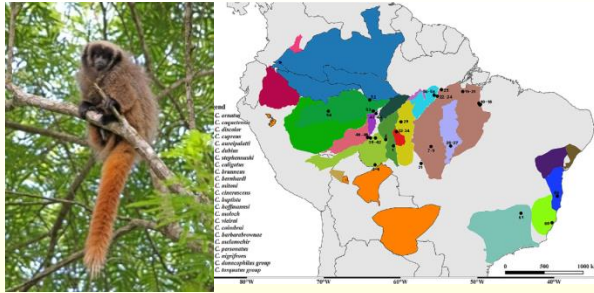


Host

Amplify

Disseminate

Brazil: non-human primates (NHP): hosts for YFV



Callicebus spp. (widow monkey)



Sapajus spp. (tufted capuchins)

Genera: less
susceptible to
YFV



Callithrix spp. (marmosets)
urban centers

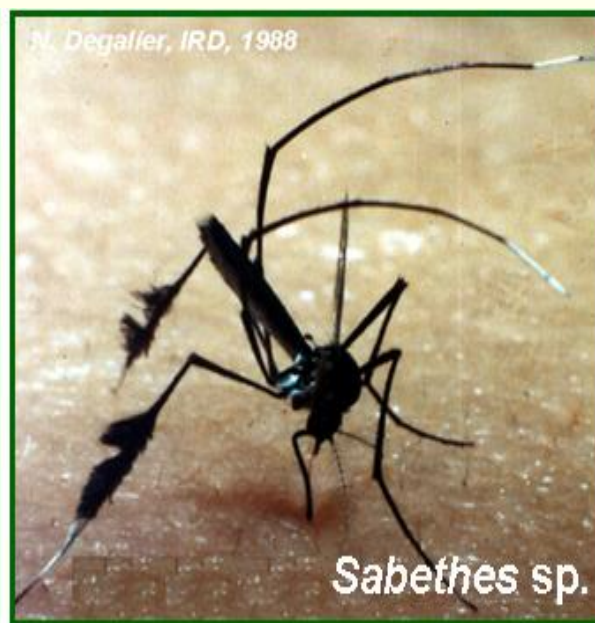
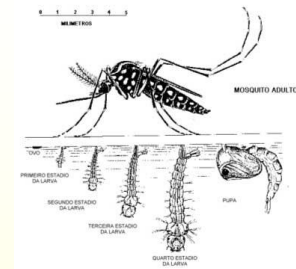


Allouata spp. (howler monkeys)

Genera: more
susceptible to
YFV

Susceptibility and wide occurrence: NHP -
sentinels for YF





Vectors

Reservoirs

Dissemination

THE DISEASE



Figure 5. Yellow fever patient during the period of infection. The patient febrile and acutely ill, with prominent conjunctival congestion. During the pre-icteric phase, the illness is difficult to differentiate from many other infectious diseases. Virus is present in the blood and the patient is infectious for blood-feeding mosquitoes.

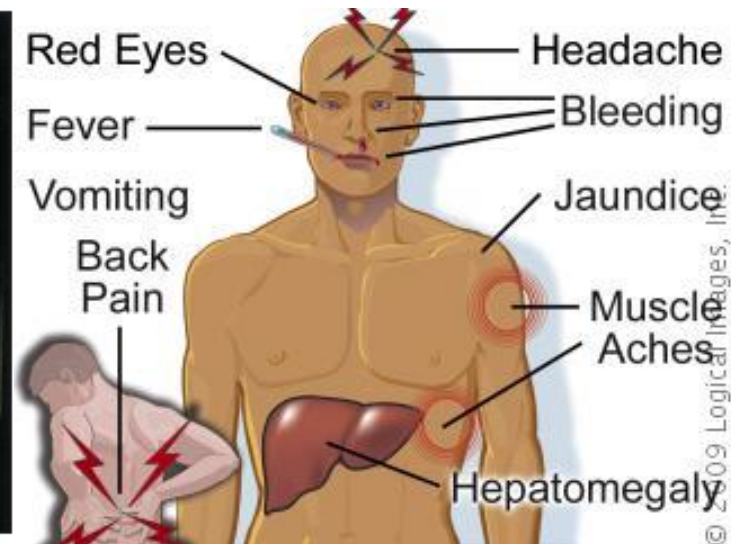
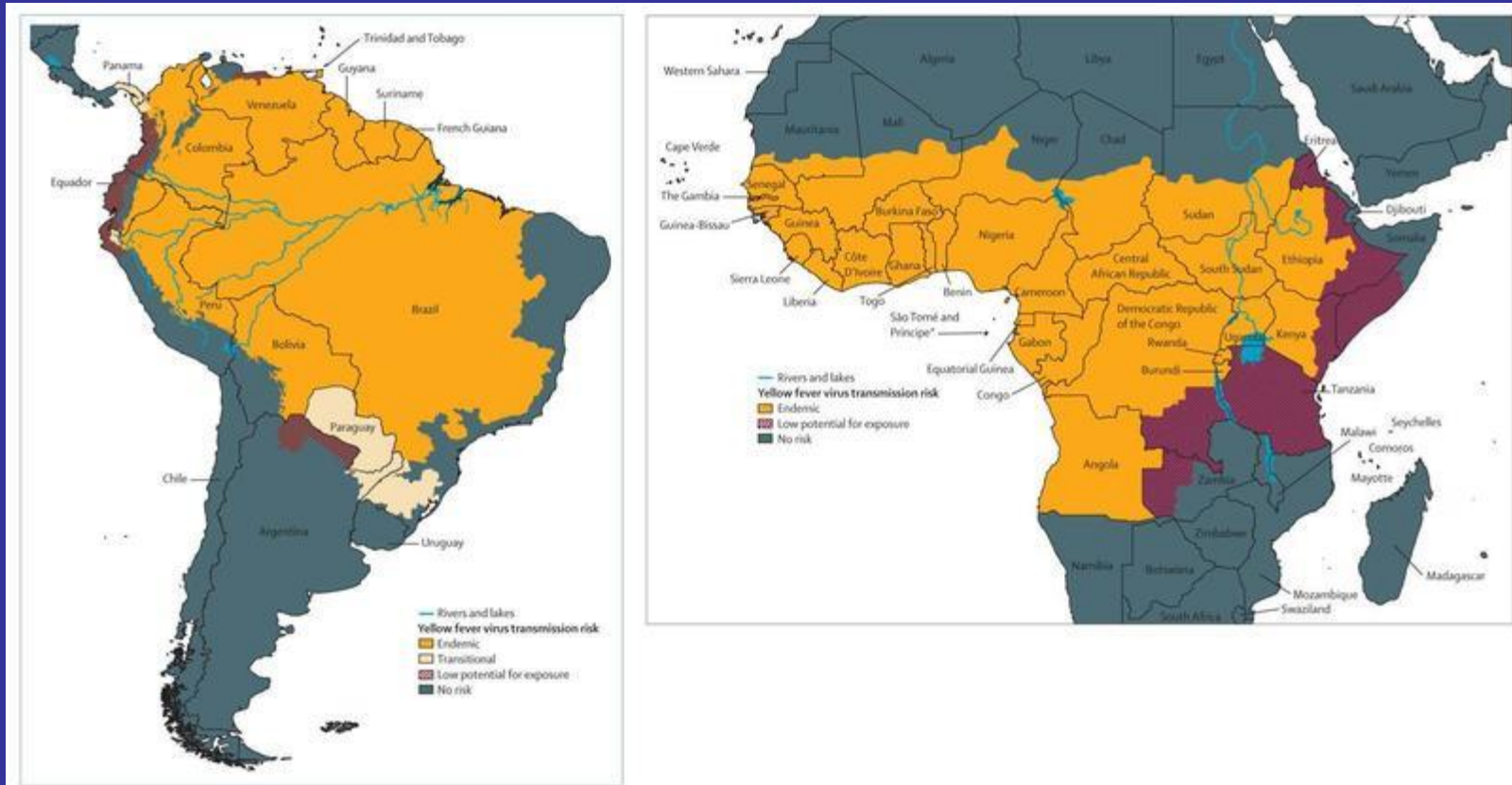


Figura 6 - O "iceberg" da febre amarela. Distribuição das formas clínicas.

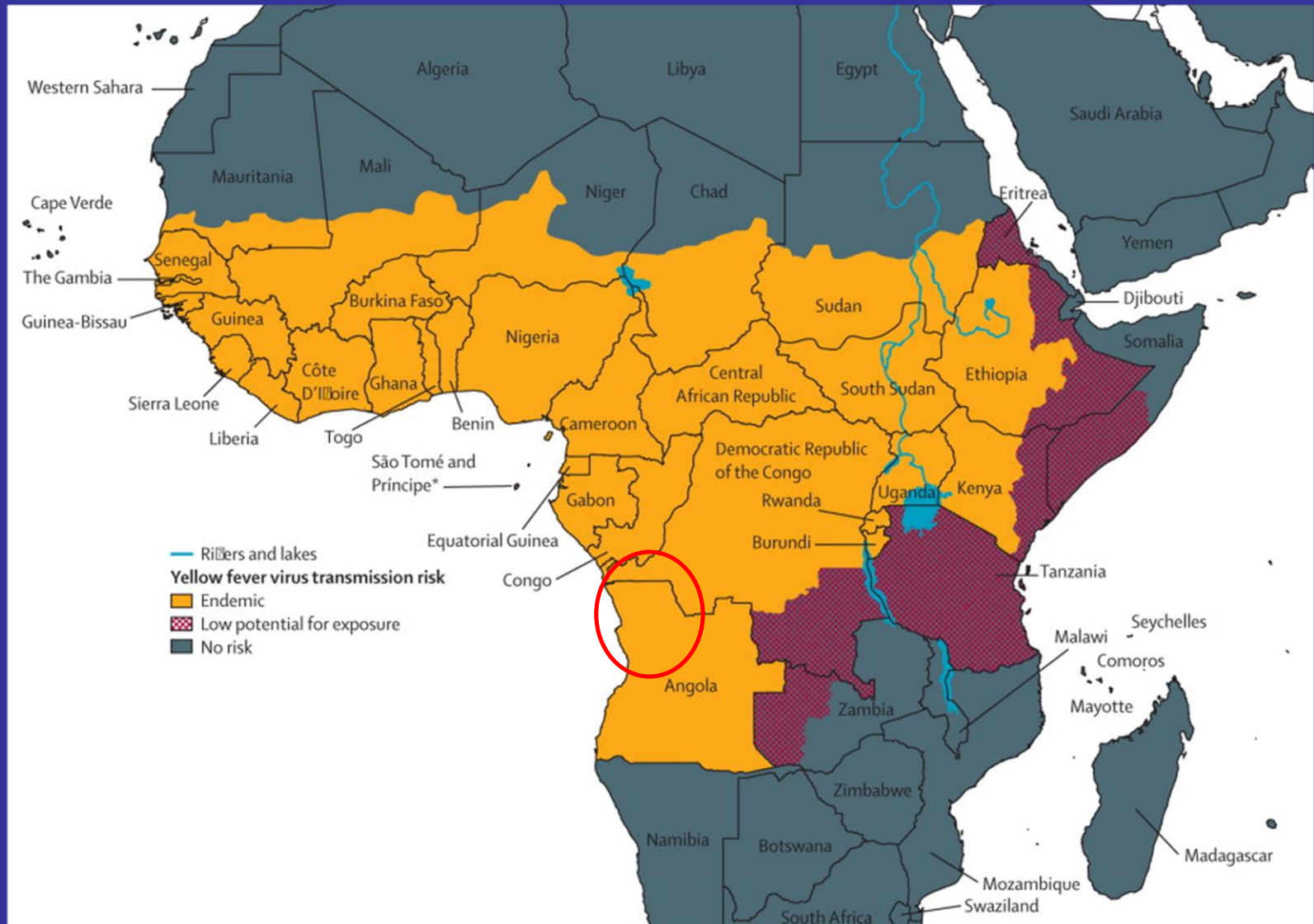
CURRENT SITUATION OF YFV

YELLOW FEVER VIRUS TRANSMISSION RISK



EPIDEMIC IN AFRICA

2016



EPIDEMIC IN AFRICA (DEMOCRATIC REPUBLIC OF THE CONGO AND ANGOLA) 2016

Angola

884 confirmed cases

121 deaths among confirmed cases (case fatality rate, 13.7%)

4347 suspected cases

377 deaths among suspected cases (case fatality rate, 8.7%)

DR Congo

78 confirmed cases (57 imported from Angola, 8 sylvatic, 13 autochthonous)

16 deaths among confirmed cases (case fatality rate, 21.1%)

2987 suspected cases

121 deaths among suspected cases (case fatality rate, 4.0%)

Kenya

2 confirmed cases

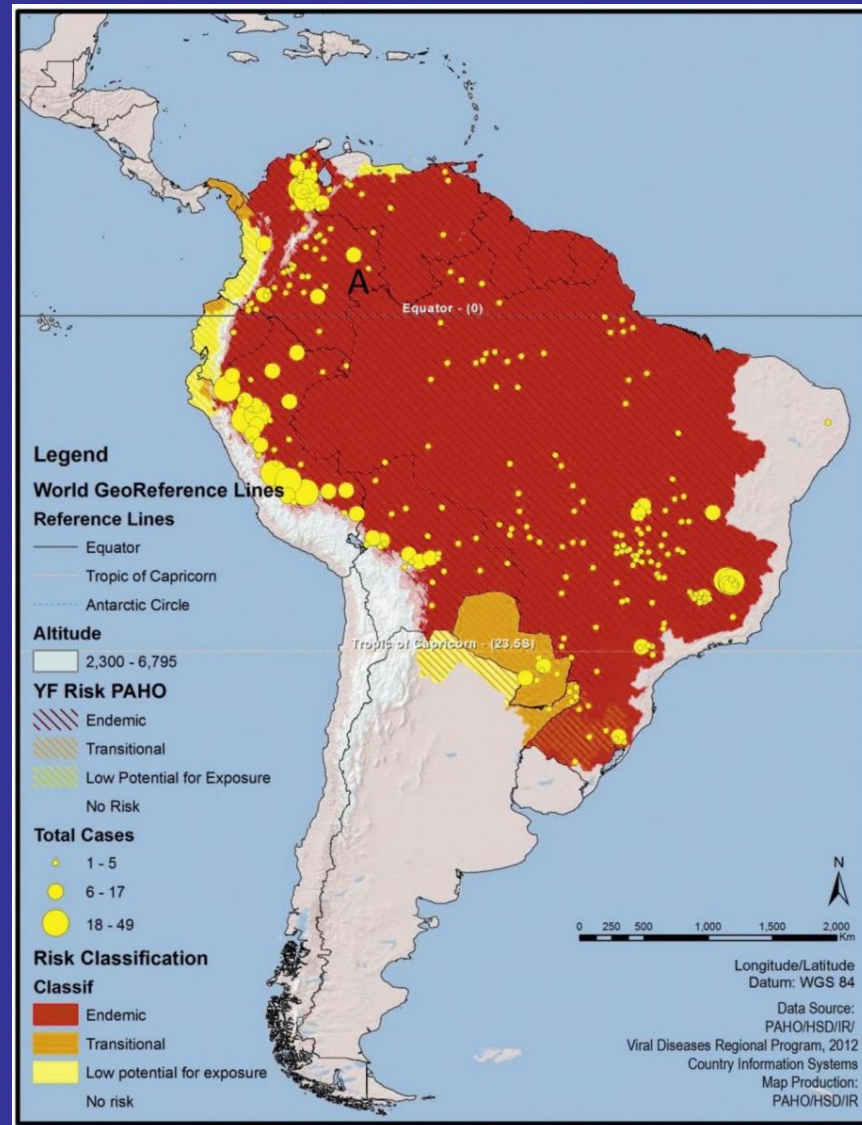
China

11 confirmed cases

Approximately 30 million people were vaccinated in the two countries.

This depleted the WHO/UNICEF and Brazilian stocks

YELLOW FEVER IN THE AMERICAS

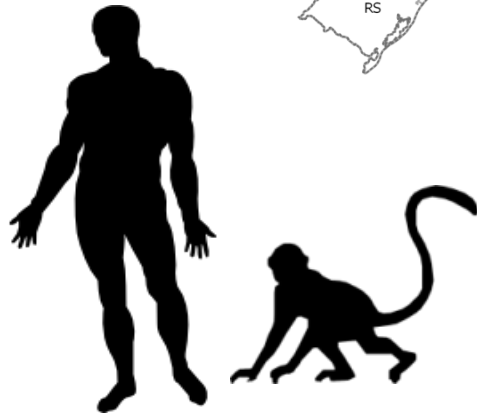
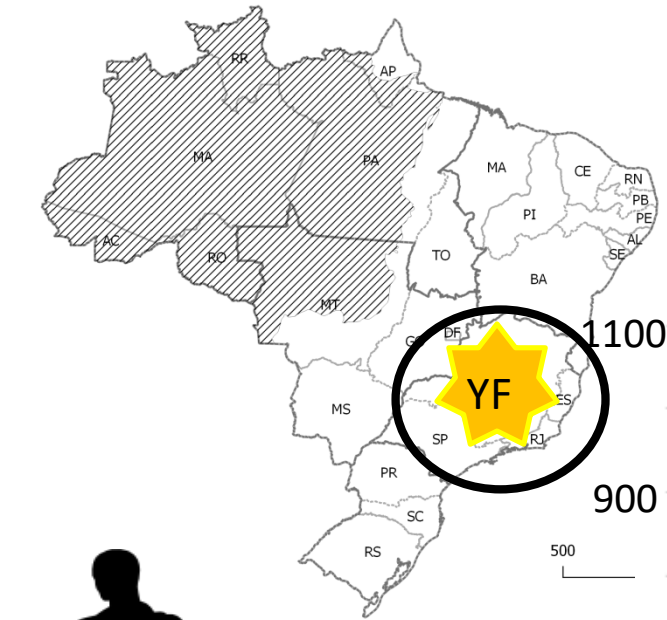


YF in Brazil

2016 to 2018: 2,153 confirmed cases

744 deaths

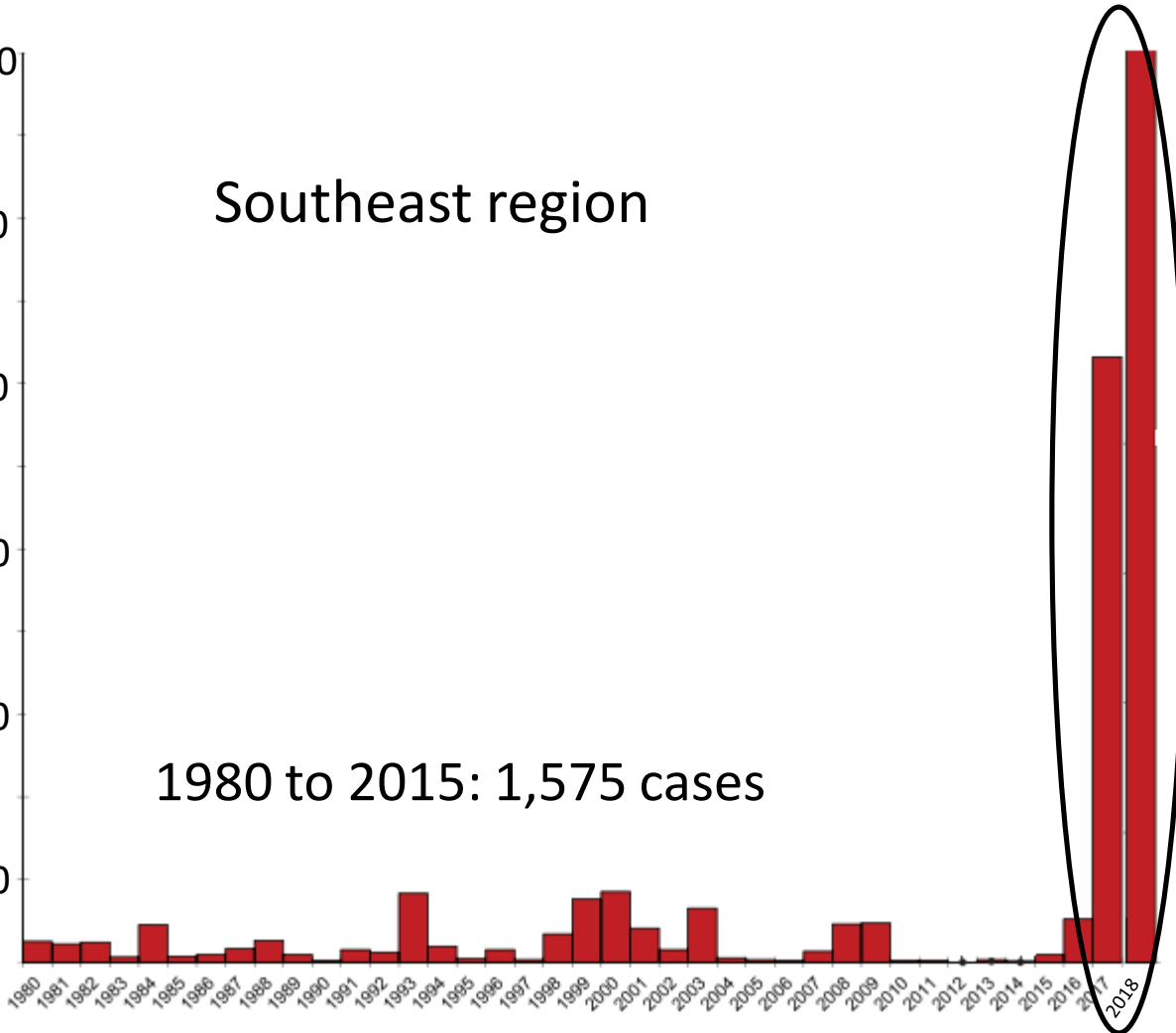
Southeast region



+ 5,000
deaths
NHP

1100
900
700
500
300
100

1980 to 2015: 1,575 cases



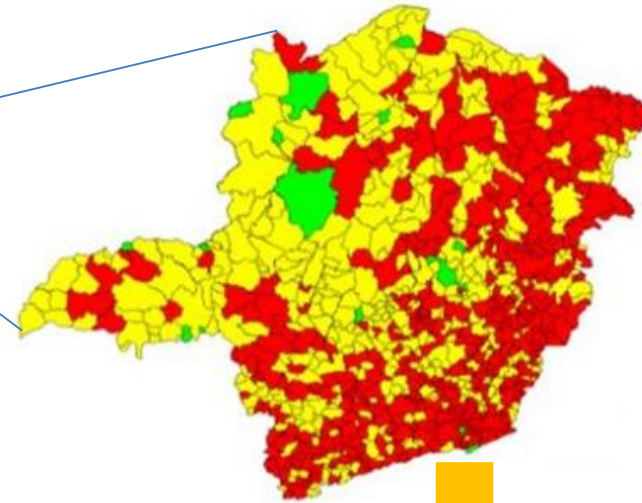
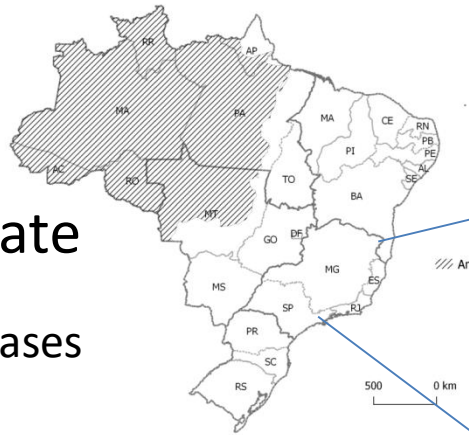
YF: Brazil

Minas Gerais state

2017-2018: 1,003 cases

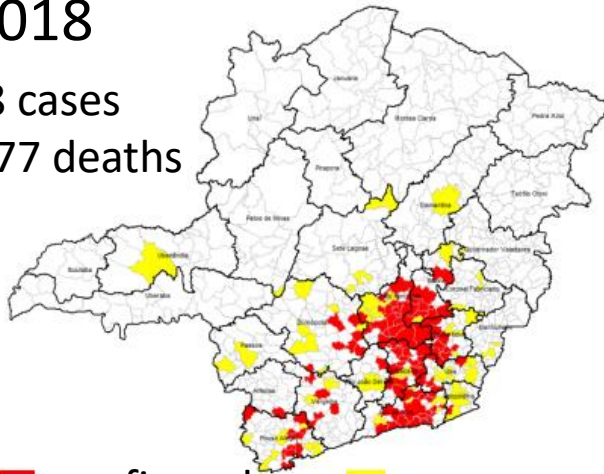
339 deaths (33.9%)

2016 – vaccination coverage



2018

528 cases
– 177 deaths

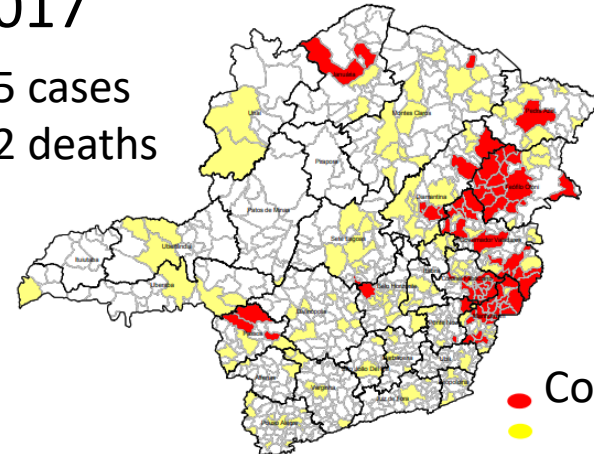


confirmed



2017

475 cases
162 deaths



Confirmed

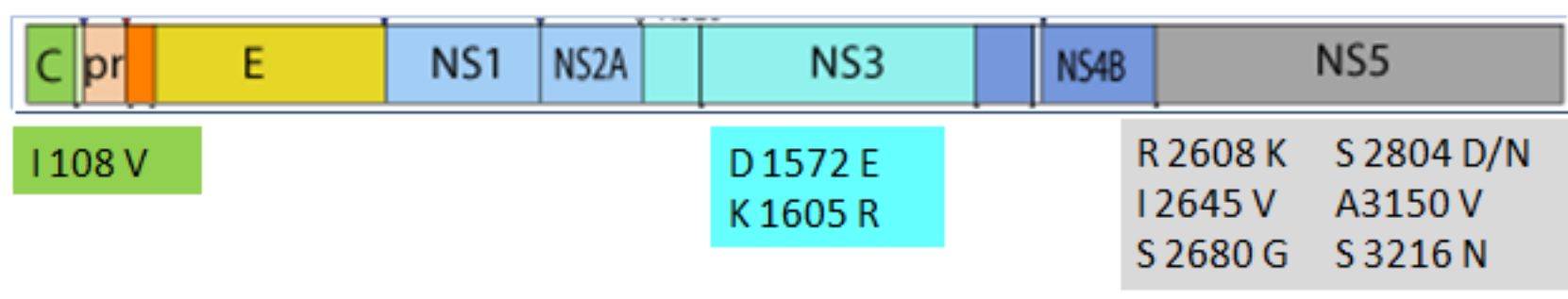


MS-BR, 2017, 2018



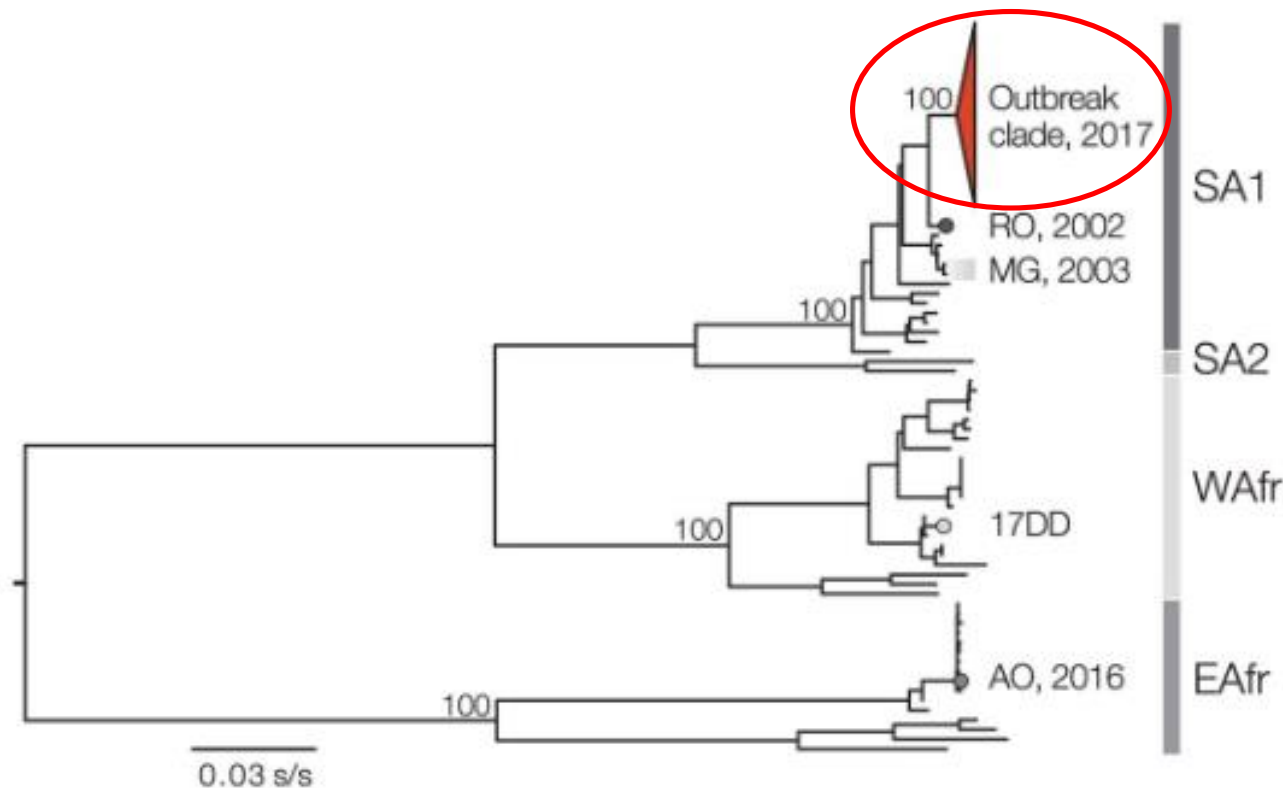
A new lineage of YFV (South American GI) was responsible for outbreaks in Southeast - 2017

9 unique aa substitutions in the polyprotein sequence



Genomic and epidemiological monitoring of yellow fever virus transmission potential

N. R. Faria^{1,*†}, M. U. G. Kraemer^{1,2,3,*}, S. C. Hill^{1,*}, J. Goes de Jesus^{4,*}, R. S. Aguiar^{5,*}, F. C. M. Iani^{6,7,*}, J. Xavier⁴, J. Quick...



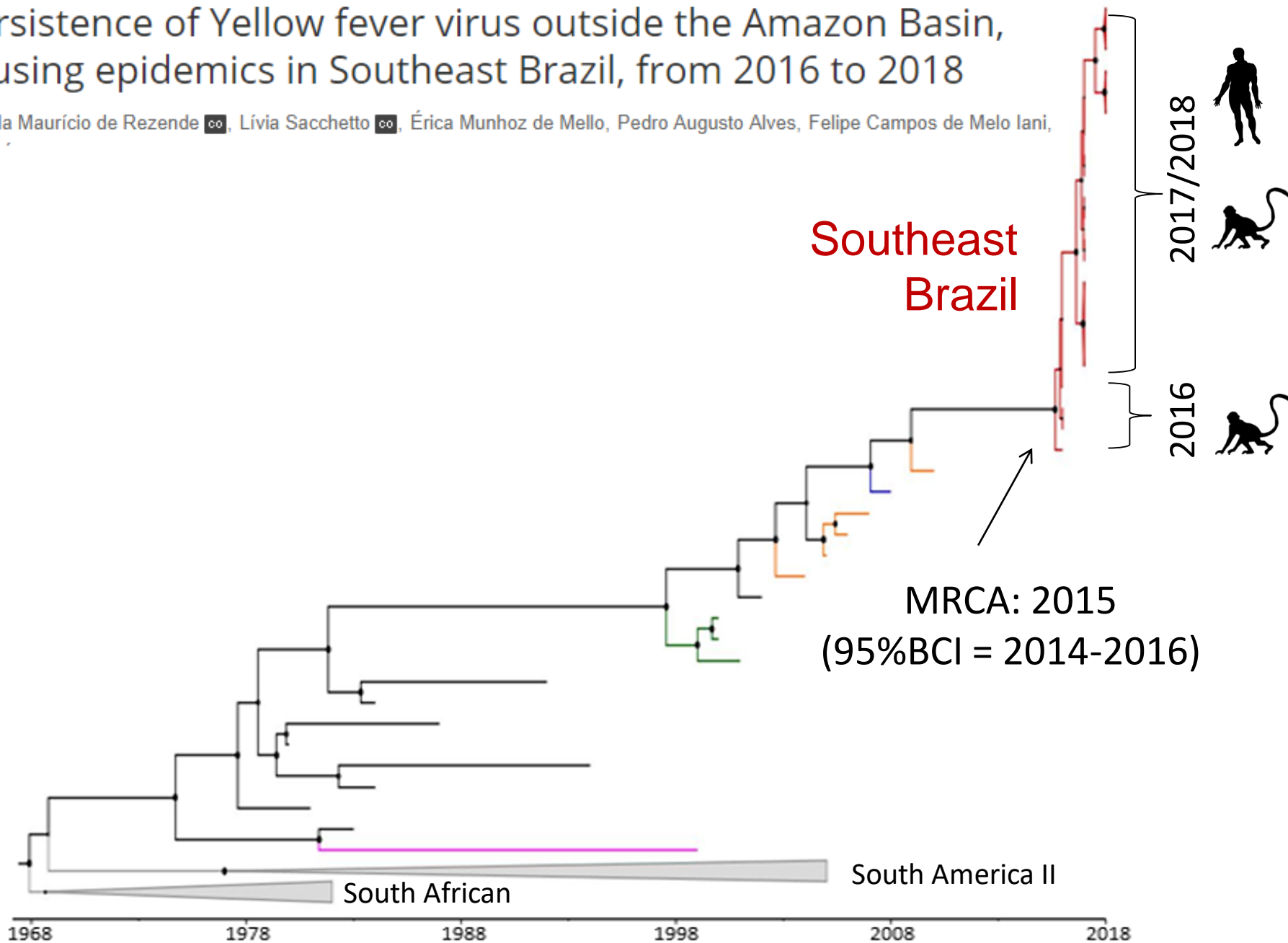
Most recent common ancestor:
July 2016
[95% BCI: Mar - Nov 2016]

Persistence of Yellow fever virus outside the Amazon Basin, causing epidemics in Southeast Brazil, from 2016 to 2018

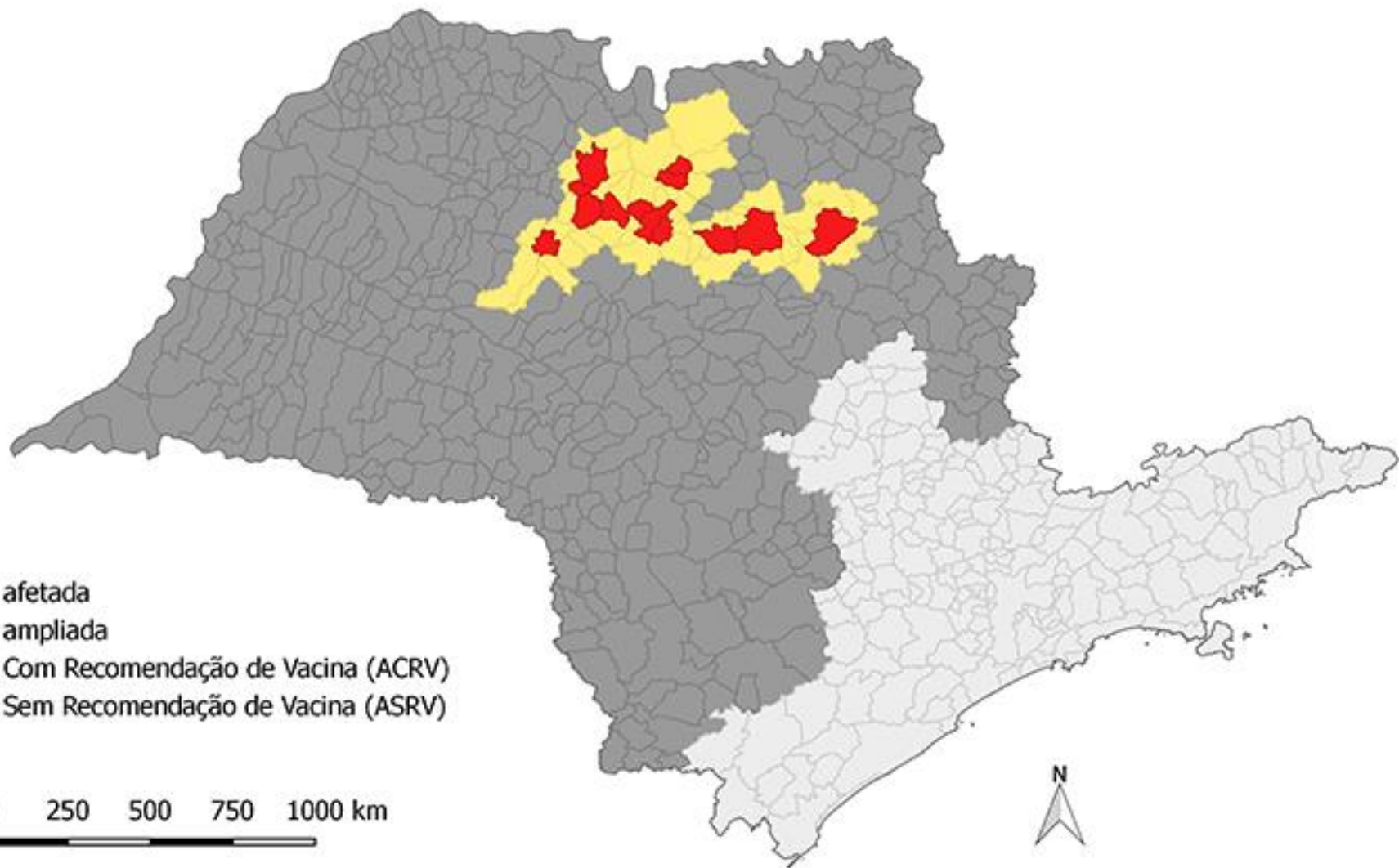
Izabela Maurício de Rezende , Lívia Sacchetto , Érica Munhoz de Mello, Pedro Augusto Alves, Felipe Campos de Melo Iani,

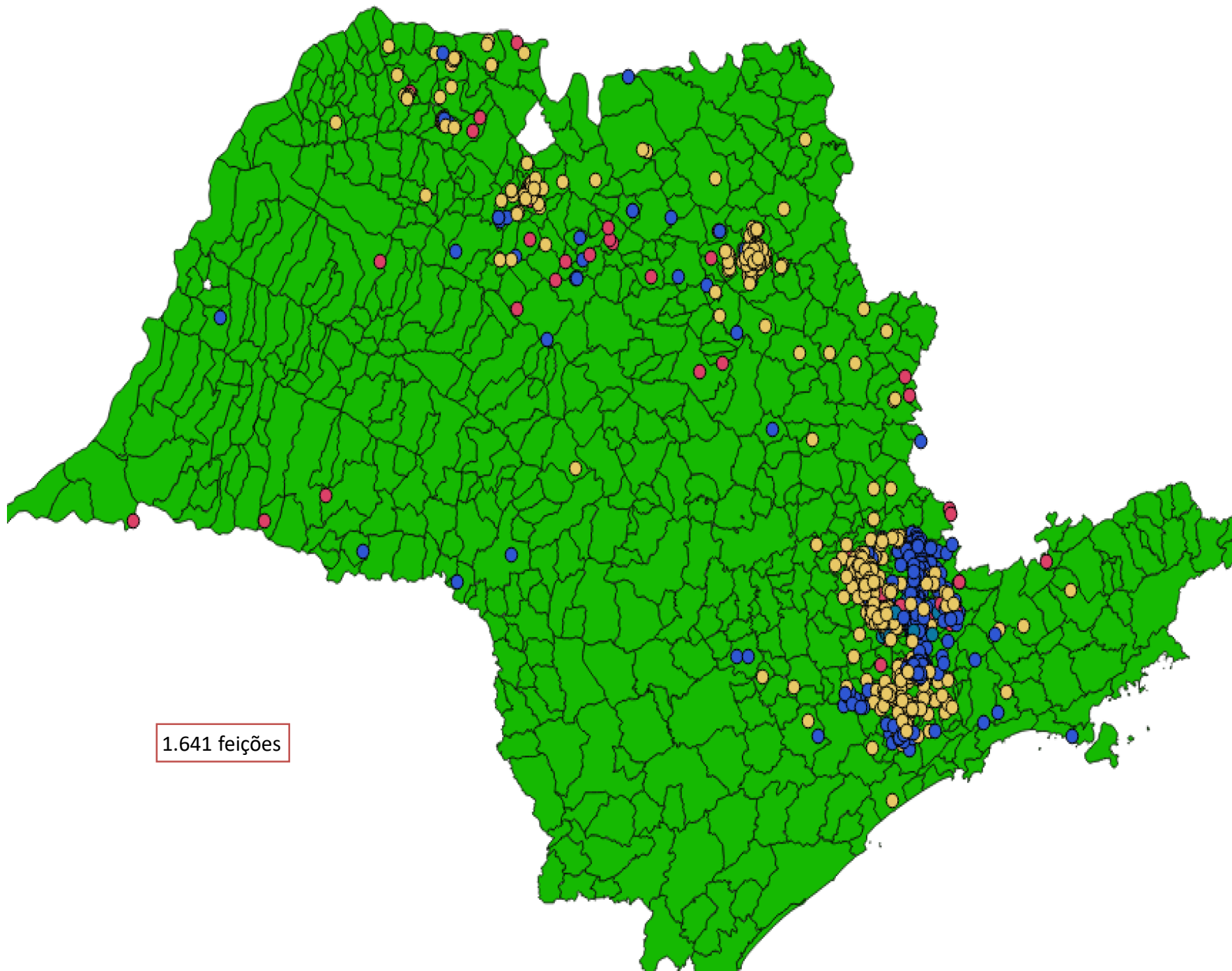
**Southeast
Brazil**

MRCA: 2015
(95%BCI = 2014-2016)



YF Sao Paulo 2016





1.641 feições



States with confirmed locally-acquired cases since July 2017



Area at risk for yellow fever transmission



Area considered at no risk for yellow fever transmission



Federal state

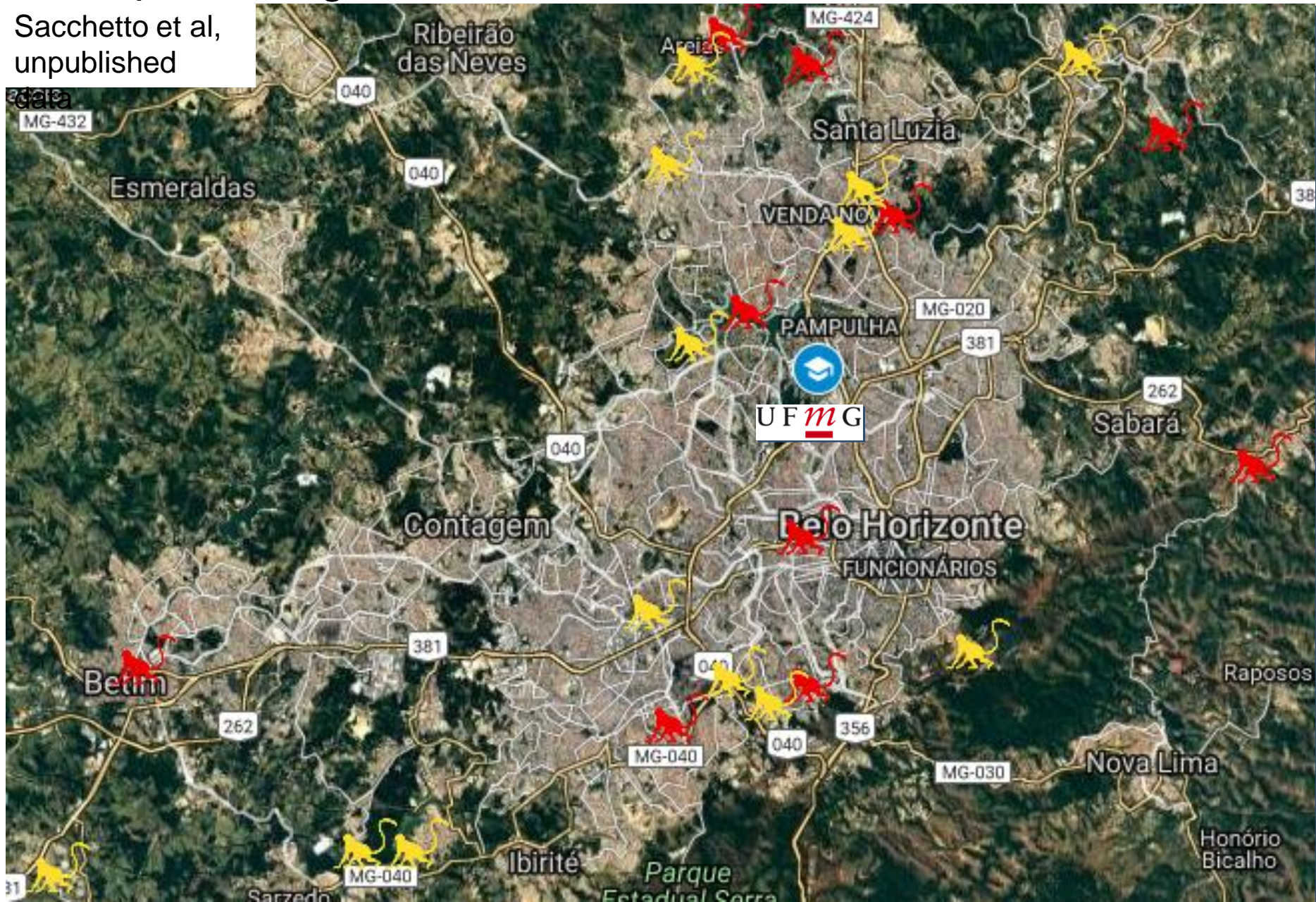


Probable place of infection of the European travel-related yellow fever cases

YFV positive *Callithrix* sp. – 2017 and 2018

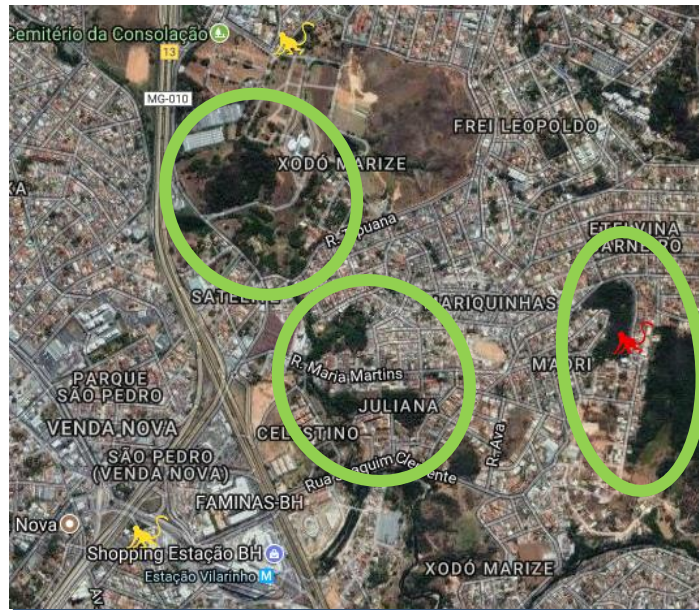
Metropolitan region – Belo Horizonte 5.8 million inhabitants

Sacchetto et al,
unpublished



YFV: *Callithrix* sp. – 2017 and 2018

Metropolitan region – Belo Horizonte 5.8 million inhabitants

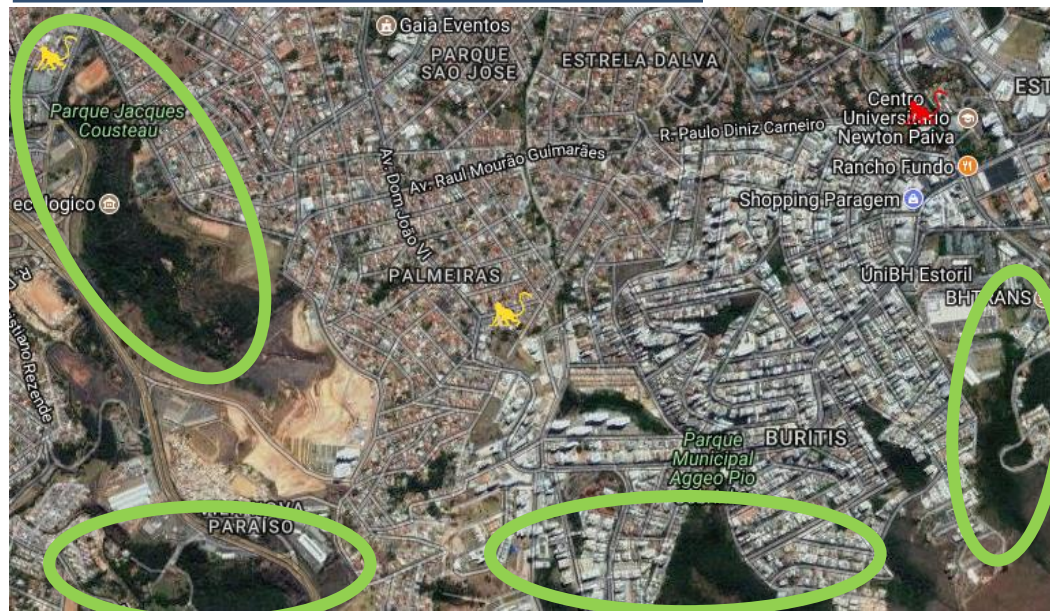


Forest pockets – commonly observed inside/boundaries of Brazilian cities

Marmosets are very common in urban areas – Brazil

Vector surveillance - needed

Sacchetto et al, unpublished data



YELLOW FEVER VACCINE

History of Virus Attenuation of Wild Yellow Fever Asibi Strain

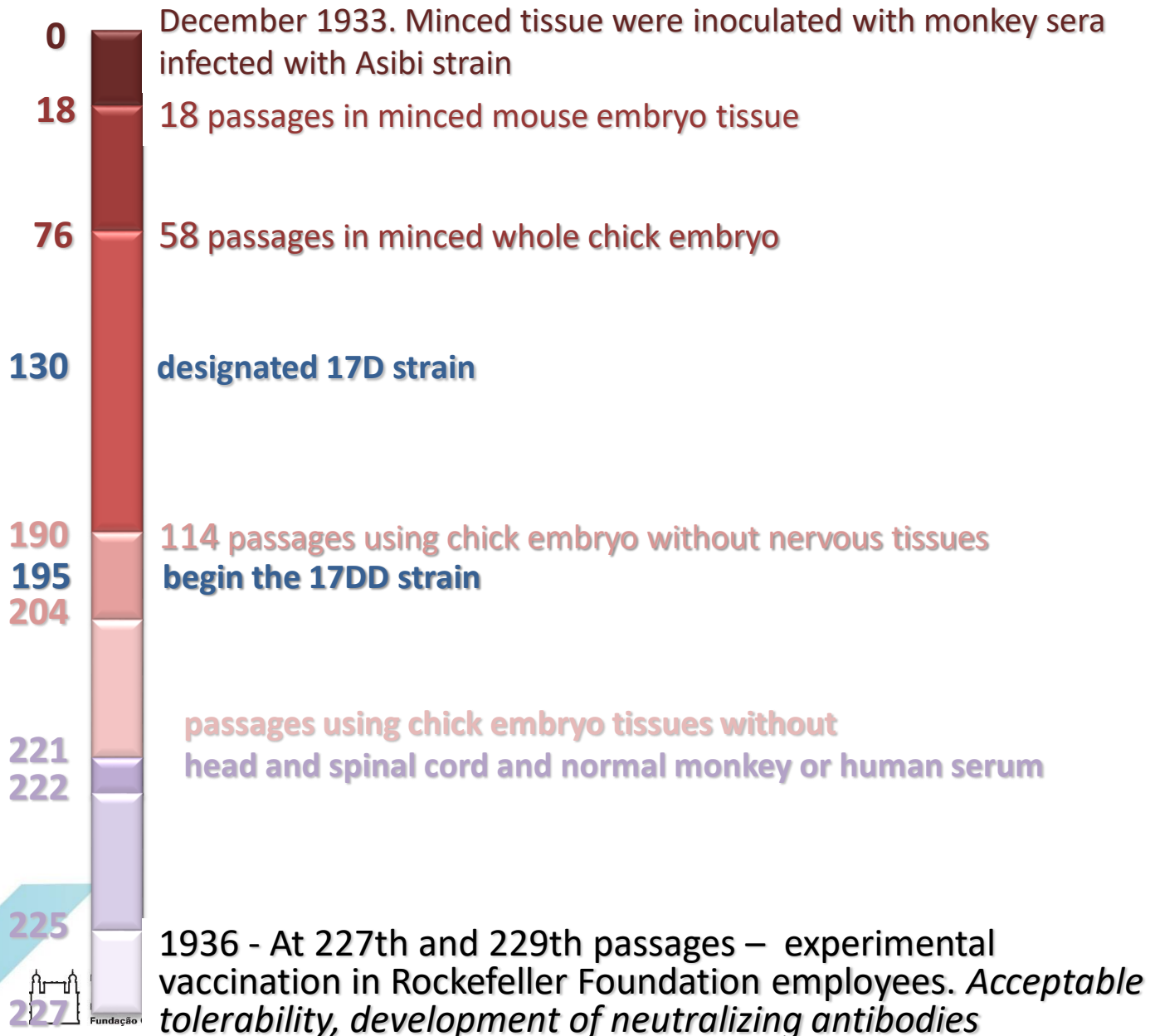


30/06/1927

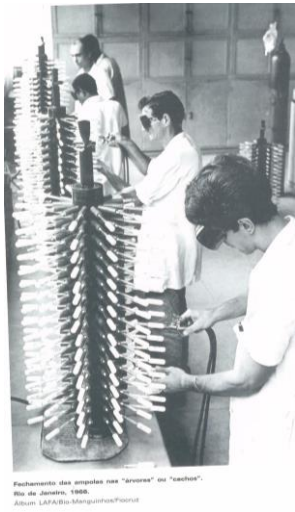
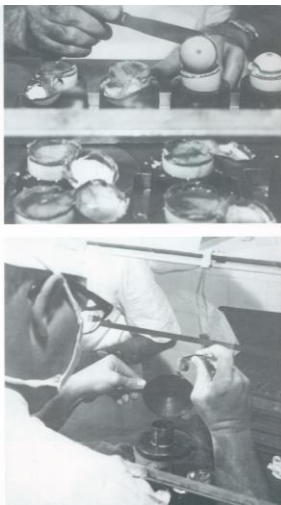
ASIBI virus,

Mahaffi &
Bauer

54 passages
in rhesus
monkeys



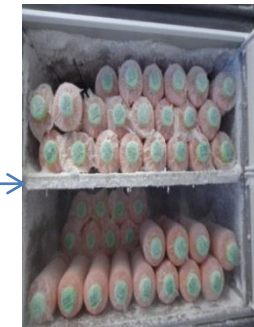
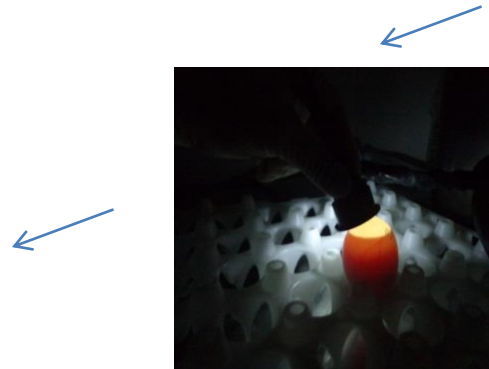
VACCINE PRODUCTION PROCESS (1942)



Fechamento das ampolas nos "arvores" ou "cachos".
Rio de Janeiro, 1960.
Arquivo LAC/Instituto Manguinhos/Fiocruz

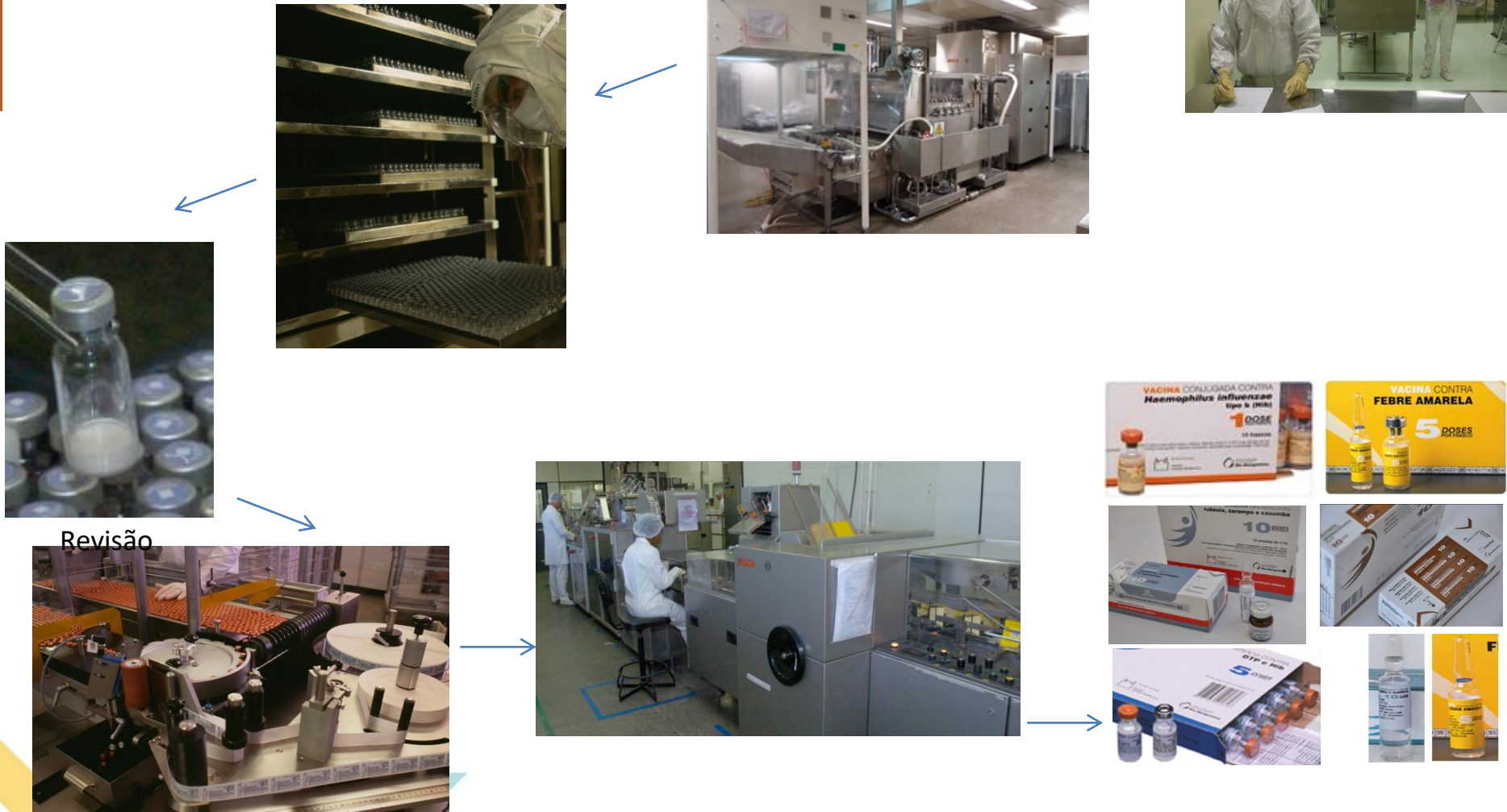


VACCINE PRODUCTION PROCESS (2017).



Suspensão
viral
(IFA)

VACCINE PRODUCTION PROCESS (2018-).



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Yellow fever vaccination booster not needed

News release

17 MAY 2013 | GENEVA - The yellow fever 'booster' vaccination given ten years after the initial vaccination is not necessary, according to WHO. An article published in WHO's Weekly Epidemiological Record (WER) reveals that the Organization's Strategic Advisory Group of Experts on immunization (SAGE) has reviewed the latest evidence and concluded that a single dose of vaccination is sufficient to confer life-long immunity against yellow fever disease.

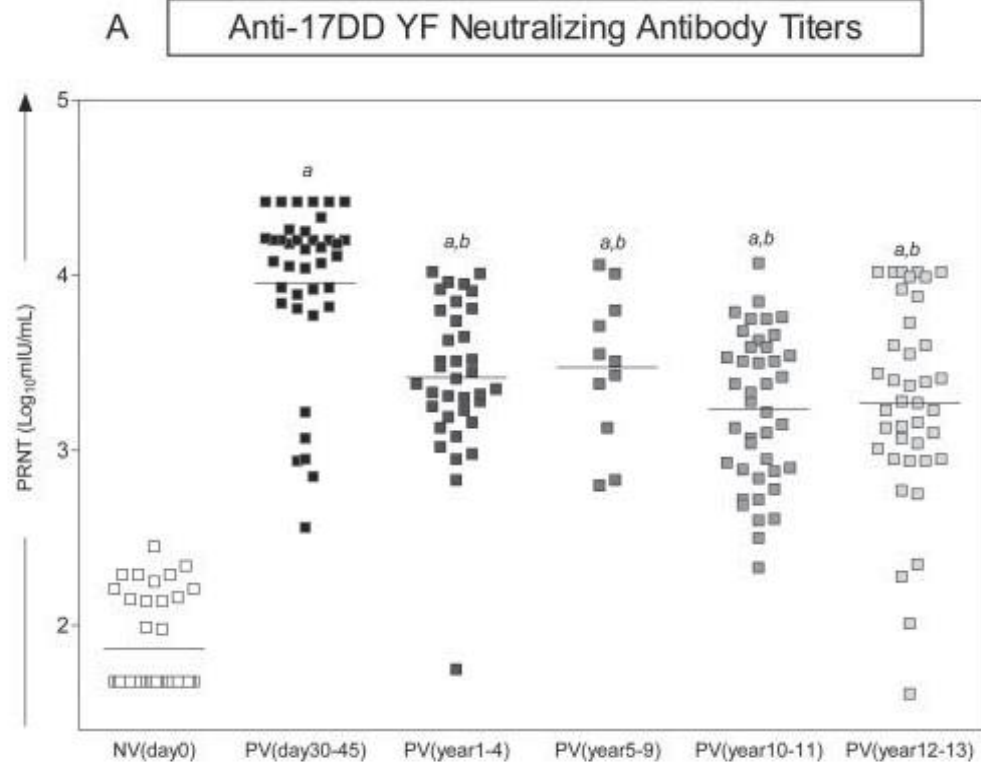
Since yellow fever vaccination began in the 1930s, only 12 known cases of yellow fever post-vaccination have been identified, after 600 million doses have been dispensed. Evidence showed that among this small number of "vaccine failures", all cases developed the disease within five years of vaccination. This demonstrates that



Related links

[Weekly Epidemiological Record \(WER\)](#)
[About SAGE](#)
[Yellow fever fact sheet](#)
[More on yellow fever](#)

DURATION OF IMMUNITY



Successful Use of Fractioned Doses (1/5th)

- Backed up by SAGE
- Strong political buy-in
- 2 months from decision to implementation
- Technical, Operational & Logistical challenges
 - Syringe supply, vaccine reconstitution, training of HCW, social mobilization...
- Coordinated effort among multiple partners (MoH, NGOs, National and International PH agencies, donors, community)
- INRB/CDC immunogenicity study ongoing on 742 individuals
- SAGE will meet mid-October to provide recommendations on FD

~7, 5 m people >2y vaccinated in Kinshasa



ORIGINAL ARTICLE

Immunogenicity of Fractional-Dose Vaccine during a Yellow Fever Outbreak — Preliminary Report

Steve Ahuka-Mundeke, M.D., Ph.D., Rebecca M. Casey, M.B., B.S., M.P.H.,
Jennifer B. Harris, Ph.D., M.P.H., Meredith G. Dixon, M.D.,
Pierre M. Nsele, M.D., Gabriel M. Kizito, M.D., Grace Umutesi, M.P.H.,
Janeen Laven, B.S., Gilson Paluku, M.D., M.P.H., Abdou S. Gueye, M.D., Ph.D.,
Terri B. Hyde, M.D., M.P.H., Guylain K.M. Sheria, M.D., Ph.D.,
Jean-Jacques Muyembe-Tanfum, M.D., Ph.D., and J. Erin Staples, M.D., Ph.D.

ABSTRACT

BACKGROUND

In 2016, the response to a yellow fever outbreak in Angola and the Democratic Republic of Congo led to a global shortage of yellow fever vaccine. As a result, a fractional dose of the 17DD yellow fever vaccine (containing one fifth [0.1 ml] of the standard dose) was offered to 7.6 million children 2 years of age or older and nonpregnant adults in a preemptive campaign in Kinshasa. The goal of this study was to assess the immune response to the fractional dose in a large-scale campaign.

METHODS

We recruited participants in four age strata at six vaccination sites. We assessed neutralizing antibody titers against yellow fever virus in blood samples obtained before vaccination and 28 to 35 days after vaccination, using a plaque reduction neutralization test with a 50% cutoff (PRNT₅₀). Participants with a PRNT₅₀ titer of 10 or higher at baseline were considered to be seropositive. Those with a baseline titer of less than 10 who became seropositive at follow-up were classified as having undergone seroconversion. Participants who were seropositive at baseline and who had an increase in the titer by a factor of 4 or more at follow-up were classified as having an immune response.

RESULTS

Among 716 participants who completed follow-up, 705 (98%; 95% confidence interval [CI], 97 to 99) were seropositive after vaccination. Among 493 participants who were seronegative at baseline, 482 (98%; 95% CI, 96 to 99) underwent seroconversion. Among 223 participants who were seropositive at baseline, 148 (66%; 95% CI, 60 to 72) had an immune response. Lower baseline titers were associated with a higher probability of having an immune response ($P < 0.001$).

CONCLUSIONS

A fractional dose of the 17DD yellow fever vaccine was effective at inducing seroconversion in most of the participants who were seronegative at baseline. These findings support the use of fractional-dose vaccination for outbreak control. (Funded by the U.S. Agency for International Development and the Centers for Disease Control and Prevention.)

Long-Term Protection After Fractional-Dose Yellow Fever Vaccination

Follow-up Study of a Randomized, Controlled, Noninferiority Trial

Anna H.E. Roukens, MD, PhD*; Karlijn van Halem, MD*; Adriëtte W. de Visser, BSc; and Leo G. Visser, MD, PhD

Background: Outbreaks of yellow fever and a frequently depleted vaccine stock increase demand for a dose-sparing strategy. A fractional dose of 17D yellow fever virus (17D-YFV) vaccine has been shown to be noninferior to the standard dose in inducing seroprotection.

Objective: To evaluate whether fractional-dose vaccination can confer long-term immunity.

Design: Ten-year follow-up of a subgroup of a randomized, controlled, noninferiority trial. (Dutch Trial Register: NTR7094 [current study] and ISRCTN46326316 [original study])

Setting: The Netherlands.

Participants: Seventy-five of 155 participants in the original trial provided a blood sample for this study. These 75 participants had received primary vaccination with 17D-YFV vaccine 10 years before. Forty received a 0.1-mL fractional dose intradermally, and 35 received the standard 0.5-mL dose subcutaneously.

Measurements: Virus-neutralizing antibody responses were measured by a plaque reduction neutralization test.

Results: Thirty-nine of 40 (98% [95% CI, 89% to 100%]) participants had protective levels of yellow fever-neutralizing antibodies more than 10 years after receiving a fractional dose of 17D-YFV vaccine compared with 34 of 35 (97% [CI, 87% to 100%]) in the standard-dose group.

Limitation: Only 48% of participants from the original trial participated in this study.

Conclusion: Intradermal administration of a one-fifth dose of yellow fever vaccine induced a protective immune response that lasted for 10 years after vaccination. Persons receiving a fractional dose of yellow fever vaccine do not require a booster vaccination for long-term protection against yellow fever.

Primary Funding Source: Leiden University Medical Center and the International Society of Travel Medicine.

Ann Intern Med. doi:10.7326/M18-1529

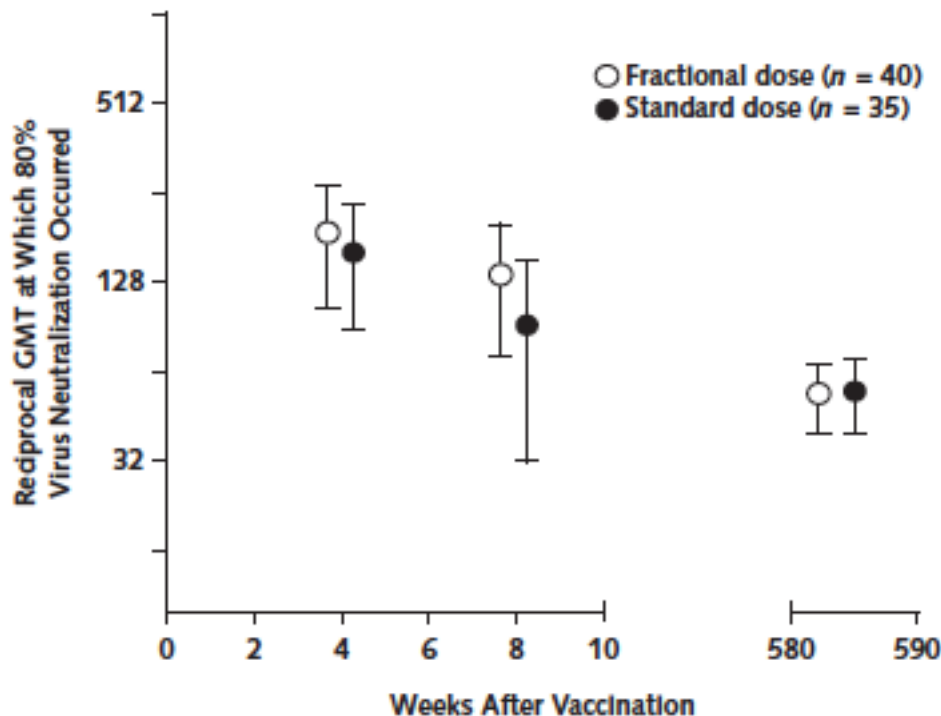
For author affiliations, see end of text.

This article was published at Annals.org on 27 November 2018.

* Drs. Roukens and van Halem contributed equally to this work.

Annals.org

Figure 2. Protective virus neutralization after fractional- or standard-dose vaccination.



Comparison of reciprocal serum dilutions at which 80% of yellow fever virus was neutralized in constant virus-varying serum dilution tests among 75 participants 10 y after primary vaccination with the intradermal fractional dose (0.1 mL) or the subcutaneous standard dose (0.5 mL). Error bars represent 95% CIs. Virus-neutralizing capacity of serum in both groups was evaluated at similar time points, but indicators are juxtaposed for visual enhancement. GMT = geometric mean titer.

SERIOUS ADVERSE EVENTS OF YF VACCINE

Case	Place, year	Age,	Sex	Time after vaccine	Clinical and laboratory surveillance	Outcome
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Virology 290, 309–319 (2001)

doi:10.1006/viro.2001.1168, available online at <http://www.idealibrary.com> on



Phenotypic and Molecular Analyses of Yellow Fever 17DD Vaccine Viruses Associated with Serious Adverse Events in Brazil

R. Galler,*[†] K. V. Pugachev,[†] C. L. S. Santos,[‡] S. W. Ocran,[†] A. V. Jabor,* S. G. Rodrigues,[§] R. S. Marchevsky,[†] M. S. Freire,[†] L. F. C. Almeida,[†] A. C. R. Cruz,[§] A. M. Y. Yamamura,[†] I. M. Rocco,[‡] E. S. Travassos da Rosa,[§] L. T. M. Souza,[‡] P. F. C. Vasconcelos,[§] F. Guirakhoo,[†] and T. P. Monath[†]

**Instituto Oswaldo Cruz and [†]Instituto de Tecnologia em Imunobiológicos, Fundação Oswaldo Cruz, 21045-900, Rio de Janeiro, RJ, Brazil; [‡]Acambis, Inc, Cambridge, Massachusetts; [‡]Instituto Adolpho Lutz, São Paulo, SP, Brazil; and [§]Instituto Evandro Chagas/Fundação Nacional de Saúde, Belém, PA, Brazil*

Received July 6, 2001; returned to author for revision August 20, 2001; accepted August 31, 2001

5	Minas Gerais, 2001	19	F	3	Fever, myalgia, cephalaea. AST and ALT 12 e 6 x. Bilirubin 6,0. Leukopenia with left shift. Coagulation disorder.	Death 10th day
6	Rio Grande do Sul, 2001	4	M	4	Fever, prostration, petechiae. Lymphadenopathy. AST and ALT 20 x; Bilirubin 7,01. Leukopenia with left shift. Renal failure.	Death 10th day
7	Rio de Janeiro, 2002	67	M	4	Fever, asthenia, myalgia, cephalaea and prostration. AST: 2572; TGP: 2525. Leukopenia. Respiratory failure. Yellow fever neutralizing antibodies: 2532 mIU/mL (10 days after	Recovered

SERIOUS ADVERSE EVENTS OF YF VACCINE

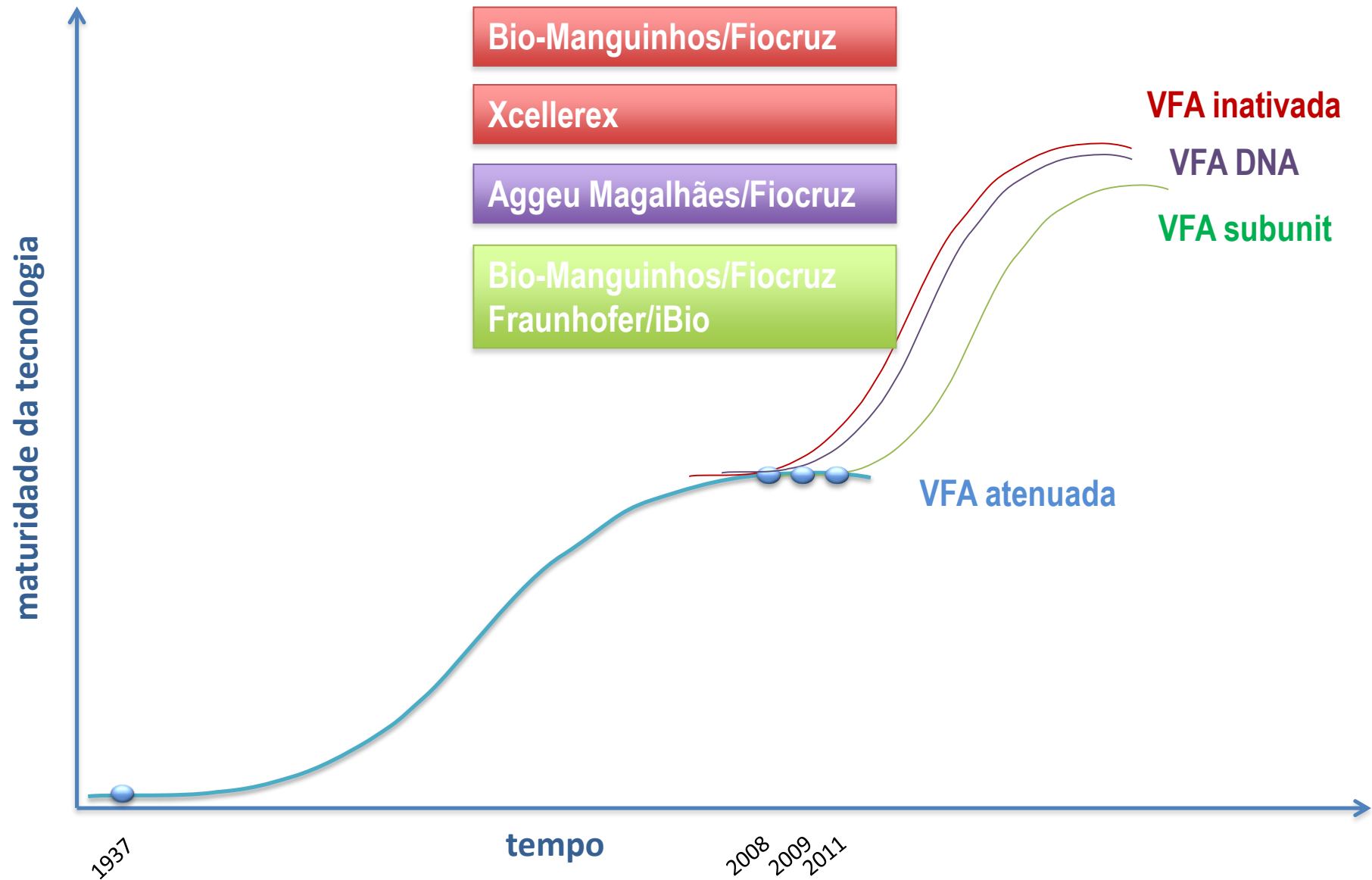
❖ International Data

- Viscerotropic disease (0,3/100.000 doses)
- Neurologic disease (0,4/100.000 doses)
- Allergic reactions (0,8/100.000 doses)

- * death about 0,5/1.000.000. We need to vaccinate 80 Million people in Brazil

YELLOW FEVER VACCINE: TECHNOLOGICAL IMPROVEMENT

YELLOW FEVER VACCINE: TECHNOLOGICAL IMPROVEMENT



AN INACTIVATE VACCINE AGAINST YF



Contents lists available at [ScienceDirect](#)

Journal of Virological Methods

journal homepage: www.elsevier.com/locate/jviromet



Pressure-inactivated yellow fever 17DD virus: Implications for vaccine development

Luciane P. Gaspar^{a,*}, Ygara S. Mendes^b, Anna M.Y. Yamamura^a, Luiz F.C. Almeida^a, Elena Caride^a, Rafael B. Gonçalves^{b,1}, Jerson L. Silva^b, Andréa C. Oliveira^b, Ricardo Galler^a, Marcos S. Freire^a

^a Programa de Vacinas Virais, Instituto de Tecnologia em Imunobiológicos, Fundação Oswaldo Cruz, Rio de Janeiro, RJ 21045-900, Brazil

^b Programa de Biologia Estrutural, Instituto de Bioquímica Médica and Centro Nacional de Ressonância Magnética Nuclear de Macromoléculas Jiri Jonas, Universidade Federal do Rio de Janeiro, Rio de Janeiro, RJ 21941-590, Brazil

Bio-Manguinhos/Fiocruz

AN INACTIVATE VACCINE AGAINST YF

The NEW ENGLAND JOURNAL of MEDICINE

ORIGINAL ARTICLE

An Inactivated Cell-Culture Vaccine against Yellow Fever

Thomas P. Monath, M.D., Elizabeth Fowler, Ph.D., Casey T. Johnson, D.O.,
John Balser, Ph.D., Merribeth J. Morin, Ph.D., Maggie Sisti, B.S.,
and Dennis W. Trent, Ph.D.

From Xcellerex, Marlborough, MA (T.P.M., E.F., M.J.M., M.S., D.W.T.); Johnson City Clinical Trials, Lenexa, KS (C.T.J.); and Veristat, Holliston, MA (J.B.). Address reprint requests to Dr. Monath at Kleiner, Perkins, Caufield, and Byers, 2750 Sand Hill Rd., Menlo Park, CA 94025, or at tmonath@kpcb.com.

N Engl J Med 2011;364:1326-33.

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Xcellerex

DNA VACCINE AGAINST YF



Anais da Academia Brasileira de Ciências (2009) 81(4): 663-669
(Annals of the Brazilian Academy of Sciences)
ISSN 0001-3765
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Membrane and envelope virus proteins co-expressed as lysosome associated membrane protein (LAMP) fused antigens: a potential tool to develop DNA vaccines against flaviviruses

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How to do a clinical trial of YF vaccine?

Cost? Market?

Population?

Markers of protection?

Ethical Considerations



Take Home Lesson

- a) YFV is re-emerging in South America and Africa with higher than usual number of cases
- b) There is a good vaccine
- c) There is not enough vaccine available. The stocks are in record low
- d) There is technology for new vaccines. But is there interest on it?
- e) No drug available

Zika in NPH Primates, Brazil

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Evidence of natural Zika virus infection in neotropical non-human primates in Brazil

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Sample ID	NHPs species	Organs (Ct value)						
		KIDNEY	BRAIN	SPLEEN	LUNG	LIVER	HEART	GONADS
PR 17/02	<i>Callithrix</i> sp.	n/a	n/a	38.37	n/a	n/a	n/a	n/a
PR 17/03	<i>Callithrix</i> sp.	n/a	37.65	n/a	neg	neg	n/a	n/a
PR 17/04	<i>Callithrix</i> sp.	37.84	neg	38.37	38.37	neg	36.19	n/a
PR 17/05	<i>Callithrix</i> sp.	38.19	36	35.14	36.37	37.88	neg	n/a
PR 17/06	<i>Callithrix</i> sp.	37.96	neg	38.23	neg	neg	n/a	n/a
PR 17/07	<i>Callithrix</i> sp.	38.18	37.71	neg	37.92	neg	neg	n/a
PR 17/08	<i>Sapajus</i> sp.	35.98	34.44	38.91	neg	neg	neg	n/a
PR 17/11	<i>Callithrix</i> sp.	Neg	37.36	neg	neg	neg	neg	n/a
PR 17/12	<i>Callithrix</i> sp.	35.66	35.64	neg	n/a	37.27	neg	n/a
PR 17/13	<i>Callithrix</i> sp.	35.57	32.31	37.13	neg	29.25	neg	n/a
PR 17/14	<i>Callithrix</i> sp.	Neg	37.12	37	neg	neg	neg	n/a
PR 17/15	<i>Callithrix</i> sp.	Neg	neg	neg	neg	37.51	n/a	n/a
PR 17/16	<i>Callithrix</i> sp.	30.87	30.4	31.48	27.76	29.18	31.32	n/a
PR 17/17	<i>Callithrix</i> sp.	31.66	31.13	n/a	31.75	32.13	30.36	n/a
PR 17/18	<i>Callithrix</i> sp.	29.17	31.86	32.87	30.47	30.16	29.72	n/a
PR 17/19	<i>Callithrix</i> sp.	32.65	32.38	31.4	31.89	neg	32.32	n/a
PR 17/20	<i>Callithrix</i> sp.	34.68	31.11	neg	34.36	neg	34.41	n/a
PR 17/21	<i>Callithrix</i> sp.	32.89	31.11	33.27	31.9	32.86	32.59	n/a
PR 17/22	<i>Callithrix</i> sp.	29.11	31.86	30.68	34.06	31.56	34.47	n/a
PR 17/23	<i>Callithrix</i> sp.	Neg	neg	n/a	neg	neg	37.85	n/a
PR 17/25	<i>Callithrix</i> sp.	37.79	neg	34.75	37.28	36.44	neg	n/a
PR 17/26	<i>Callithrix</i> sp.	37.69	n/a	35.94	neg	neg	36.97	n/a
PR 17/27	<i>Callithrix</i> sp.	38.47	neg	neg	neg	neg	neg	n/a
MG 17/01	<i>Callithrix</i> sp.	n/a	n/a	n/a	n/a	36.0	n/a	n/a
MG 17/02	<i>Callithrix</i> sp.	n/a	n/a	n/a	n/a	33.9	n/a	n/a
MG17/15	<i>Callithrix</i> sp.	27.1	n/a	n/a	n/a	n/a	n/a	neg
MG 17/16	<i>Callithrix</i> sp.	n/a	n/a	n/a	n/a	35.3	n/a	n/a
MG17/30	<i>Callithrix</i> sp.	neg	n/a	n/a	n/a	35.6	n/a	36.3
MG 17/31	<i>Callithrix</i> sp.	n/a	n/a	n/a	n/a	35.9	n/a	n/a
MG 17/32	<i>Callithrix</i> sp.	n/a	n/a	n/a	n/a	36.3	n/a	n/a
MG 17/45	<i>Callithrix</i> sp.	n/a	n/a	n/a	n/a	35.6	n/a	n/a
MG 17/51	<i>Callithrix</i> sp.	n/a	n/a	n/a	n/a	35.7	n/a	n/a
Mosquitoes (Ct Value)								
Sample ID	Mosquito species	Ct						
17/151	<i>Ae. aegypti</i>	35.17						
17/160	<i>Ae. aegypti</i>	36.89						
17/161	<i>Ae. aegypti</i>	36.48						
17/163	<i>Ae. aegypti</i>	31.87						
17/164	<i>Ae. aegypti</i>	36.77						
17/169	<i>Ae. aegypti</i>	22.23						

Table 1. Non-human primates positive for Zika virus, by RT-qPCR. Positive samples and mosquitoes are indicated by the Ct (cycle threshold) value. n/a: not available. neg: negative. Samples collected in São José do Rio Preto (SP), from January to March 2017 are identified by PR followed by year and sample ID. Samples collected in Minas Gerais, from January to June 2017, are identified by MG followed by year and sample ID. Mosquitoes collected in São José do Rio Preto (SP), in the first trimester of 2017 are identified by year and sample ID.

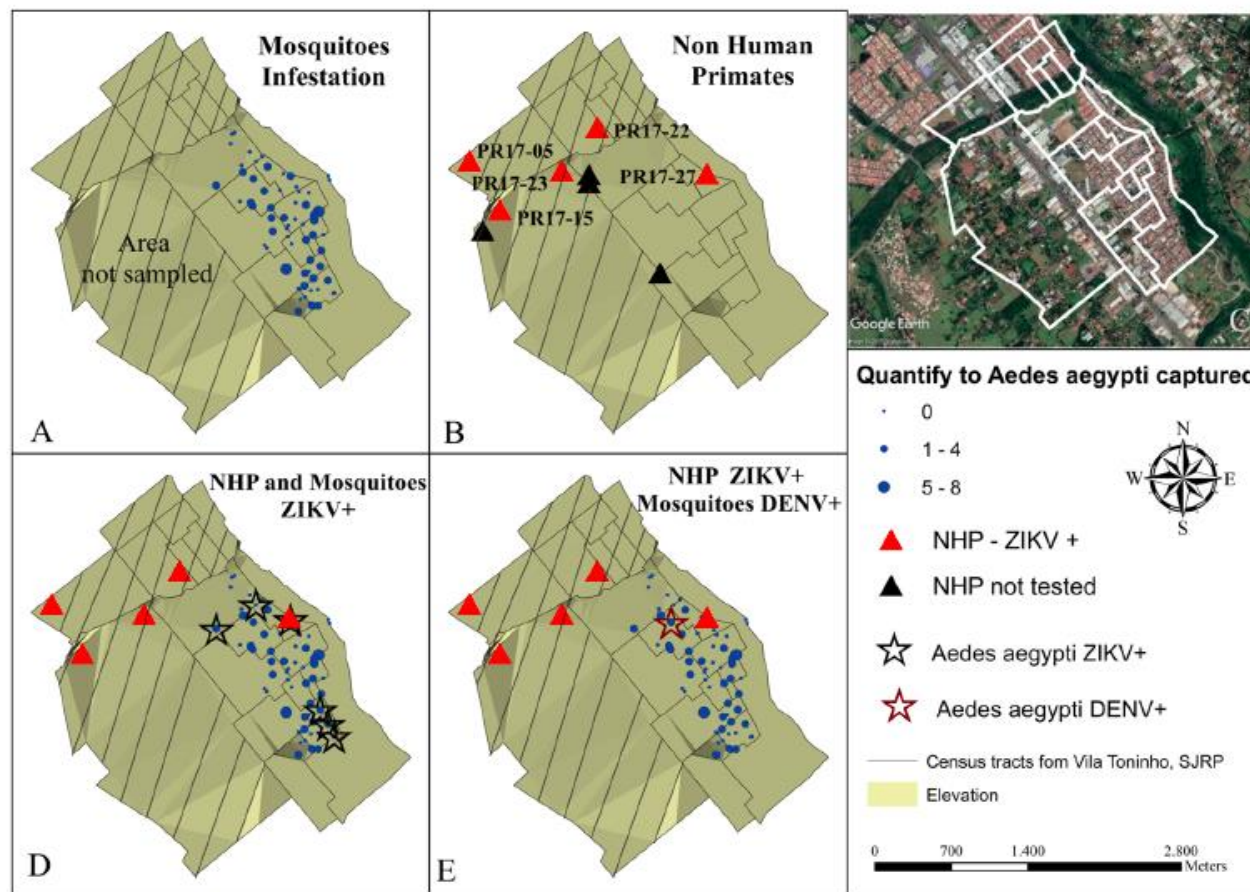


Figure 1. Geoprocessing map of the NHPs and mosquitoes captured in the Vila Toninho neighborhood. (A) Schematic representation of the area where mosquitoes are regularly collected in the Vila Toninho neighborhood. The hatched area represents the area where there is no specimen collection. The blue dots represent the collection points of the mosquitoes and the quantity of specimen collected. (B) Schematic representation of the collection points of the nine NHP found dead. The NHPs identified by ID PR 17-05, PR 17-15, PR 17-22, PR 17-23, PR 17-27 were analyzed and tested positive for ZIKV in one or more tissue samples and are represented by a red triangle. The black triangles represent the NHPs collected but not tested. (C) Satellite image of the Vila Toninho neighborhood. The boundary of the neighborhood is marked in white. Vegetation cover area can be seen in green surrounding the neighborhood. (D) Overlap of the area of the animals and mosquitoes collection. The ZIKV-positive PR 17-27 is overlapping with a ZIKV-positive *Ae. aegypti* mosquito pool. (E) Overlap of the areas of animals and mosquito collections with the presence of the DENV-positive *Ae. aegypti* mosquitoes (Vila Toninho satellite image by Google Earth Pro 7.3.1.4507 (64-bit) software. URL <https://www.google.com/maps/@-20.84677,-49.34063,5682m/data=!3m1!1e3>). Map data: Google, 2018 DigitalGlobe.

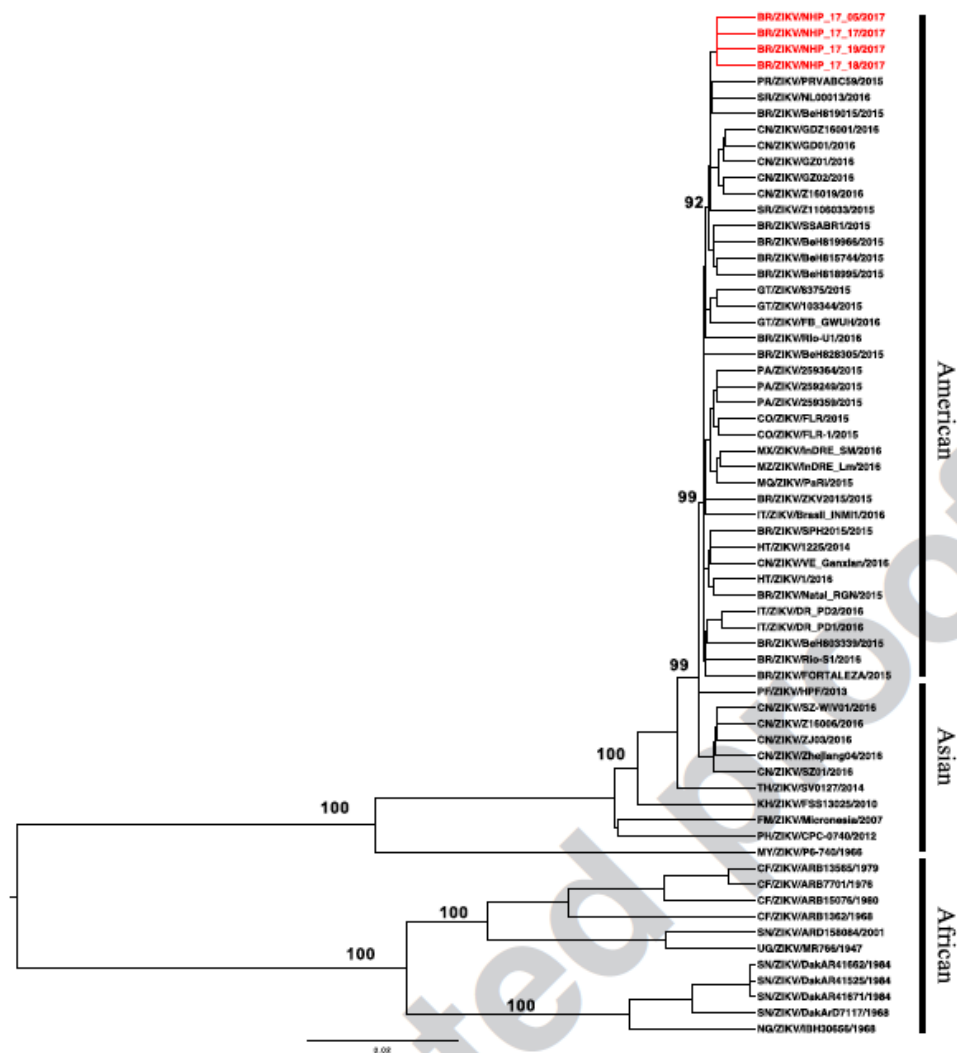


Figure 2. Molecular Phylogenetic analysis of Zika virus by the Maximum Likelihood method. The four strains obtained from NHPs (marmosets) are highlighted in red. Bootstrap values above 90% are shown. Initial tree(s) for the heuristic search were obtained automatically by applying Neighbor-Join and BioNJ algorithms to a matrix of pairwise distances estimated using the Maximum Composite Likelihood (MCL) approach, and then selecting the topology with superior log likelihood value. A discrete Gamma distribution was used to model evolutionary rate differences among sites (5 categories (+G, parameter = 1.7699)). The rate variation model allowed for some sites to be evolutionarily invariable ([+I], 52.6922% sites). The tree was drawn to scale, with branch lengths measured in the number of substitutions per site. There were a total of 10269 positions in the final dataset. Evolutionary analyses were conducted in MEGA7⁶⁴.

Days	ZIKV RNA genome copies/ml			
	NHP 1	NHP 2	NHP 3	NHP 4
-1	Negative	Negative	Negative	Negative
0	Negative	Negative	Negative	Negative
2	200	Negative	1076	204
3	1150	214	238	2191
4	2848	626	786	3296
5	19958	1762	30200	4156
8	Negative	217	Negative	Negative
9	580	229	Negative	9840
12	2208	Negative	Negative	Negative
15	Negative	Negative	Negative	Negative
19	Negative	347	Negative	Negative

Figure 3. Viremia measurement in experimentally ZIKV-infected *Callithrix penicilata* collected from day –1 until 19 dpi. One-step qRT-PCR was used to measure semi quantitatively the ZIKV RNA loads in the serum of four animals at indicated days p.i. and represented as viral RNA copies per mL of sample standard curve. The curve was obtained from a standard sample with known titer after serial dilutions (5×10^1 to 5×10^6 copies/mL) on the plasma of the non-infected marmosets. Values are expressed by RNA genome copies per mL for all the infected marmosets. Viremia was detected in the serum of marmosets 1, 3 and 4 on day 2 p.i. and in all infected marmosets on day 3 p.i. The figure shows that viremia increased on day 5 p.i. when compared to other evaluated days for all the infected marmosets. p.i.: post infection. NHP: non-human primates. Day – 1: day before the infection.

Conclusions - II

- Zika was detected in dead NHP in urban environment
- The sequence is similar to the human cases
- The spatial distribution was assessed
- The virus can infect experimental monkeys, with viremia and IgG seroconversion

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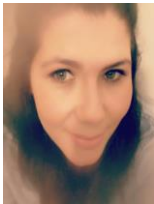


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