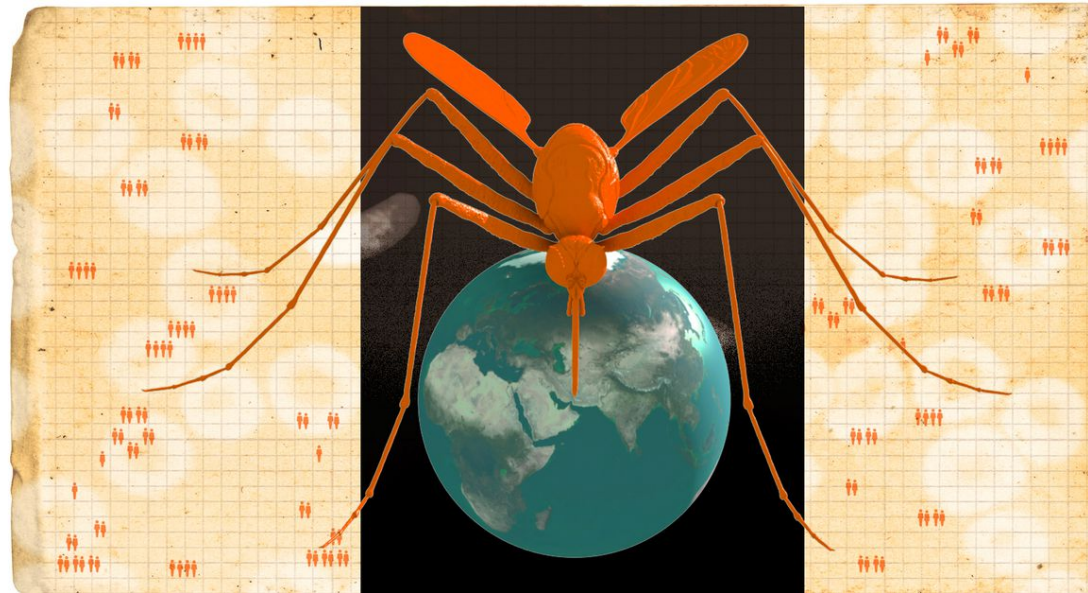


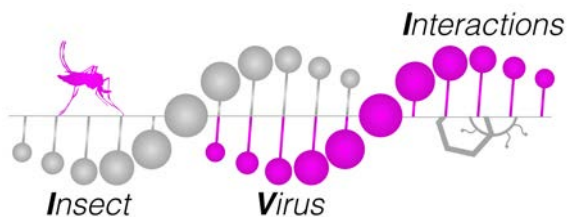
18 décembre 2019 – Collège de France

Des moustiques, des virus et des hommes

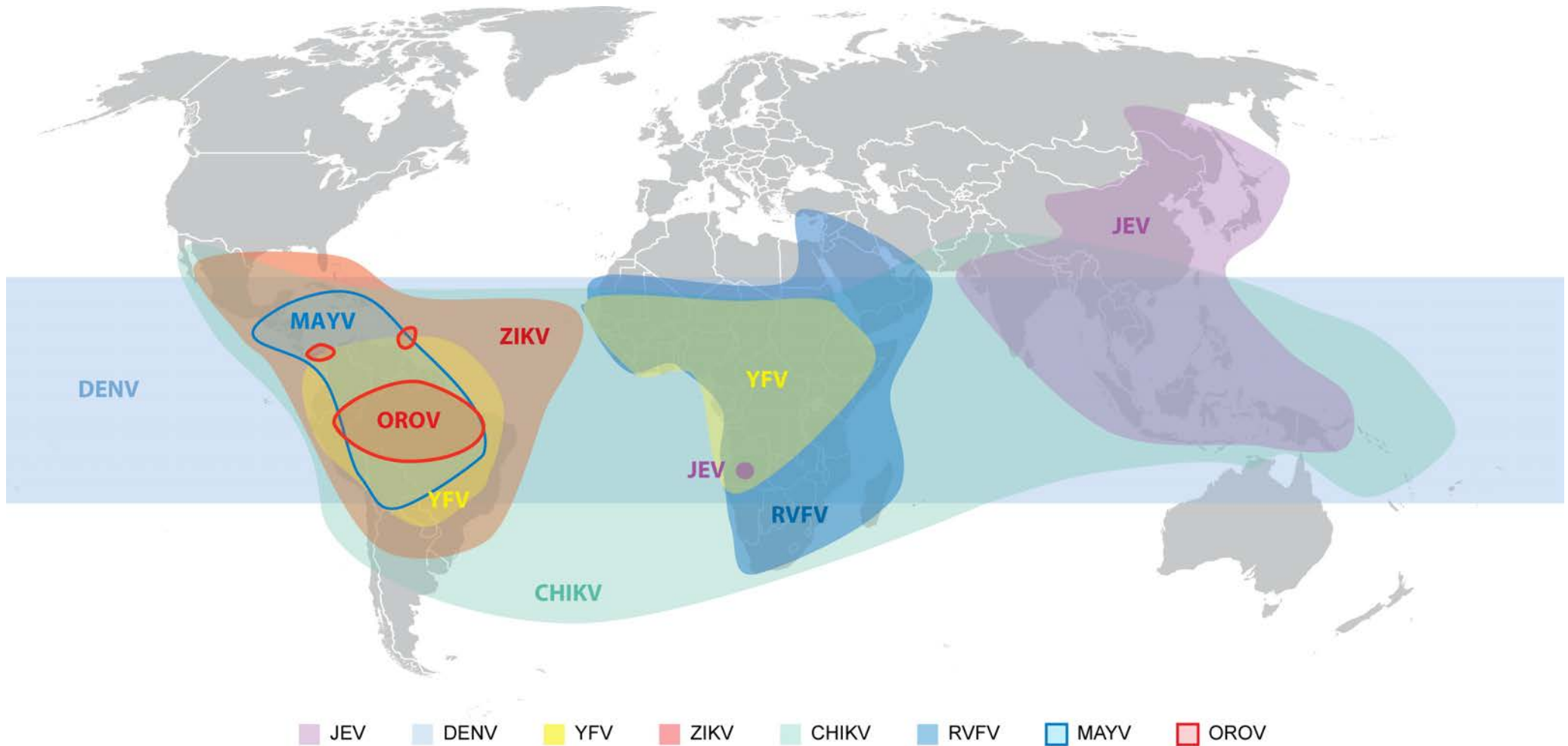


Louis Lambrechts

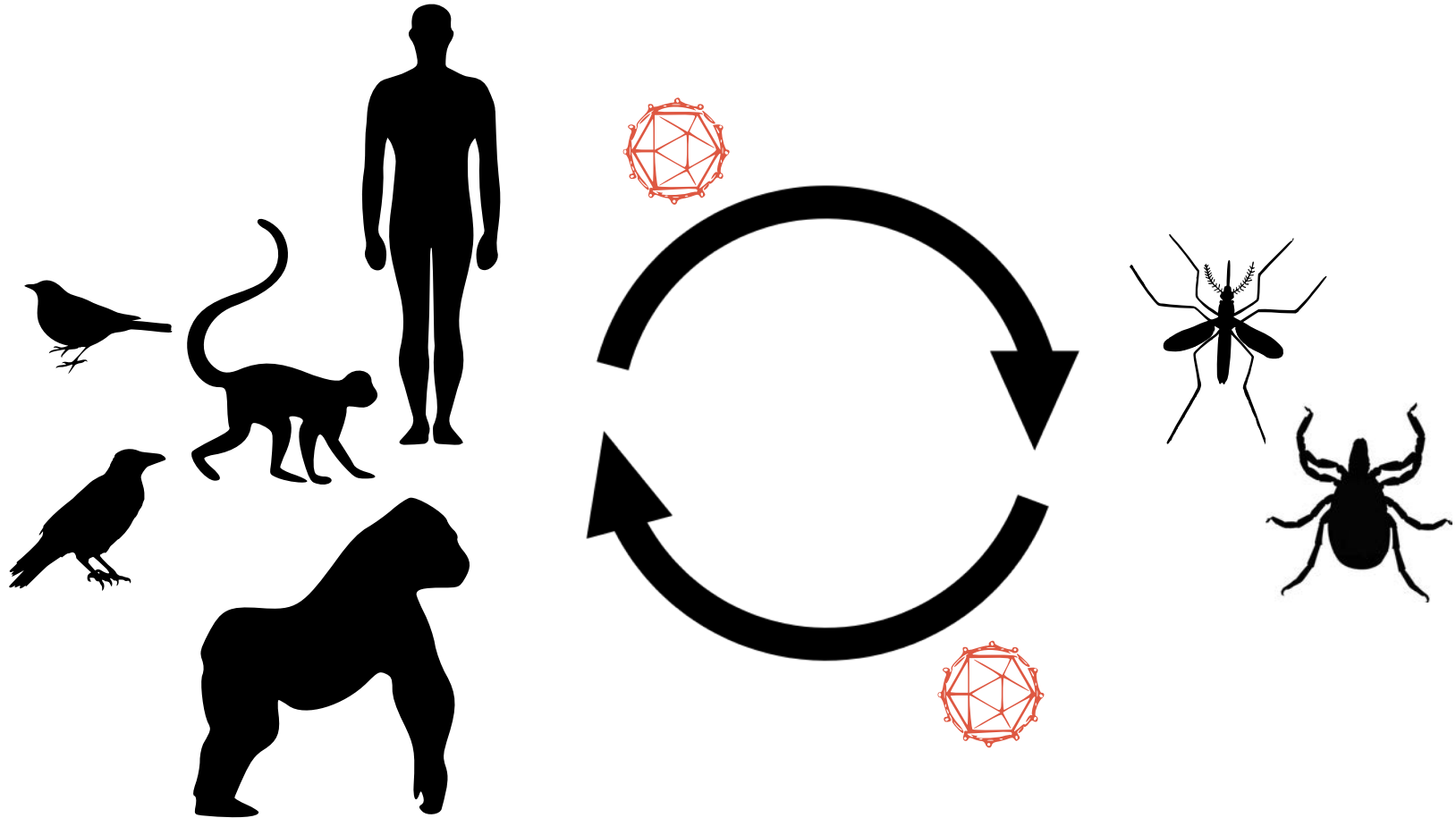
*Unité Interactions Virus-Insectes, Institut Pasteur-CNRS UMR 2000, Paris, France
Laboratoire d'Excellence Biologie Intégrative des Maladies Infectieuses Emergentes*



Emerging mosquito-borne viruses



Arthropod-borne viruses (Arboviruses)



Outline

- 1. Examples of emerging arboviruses**
 - Yellow fever
 - Dengue
 - Zika
- 2. Factors underlying arbovirus emergence**
 - Globalization
 - Urbanization
 - Ineffective control
- 3. Control of emerging arboviruses**
 - Conventional vector control
 - Innovative strategies

Outline

1. Examples of emerging arboviruses

- Yellow fever
- Dengue
- Zika

2. Factors underlying arbovirus emergence

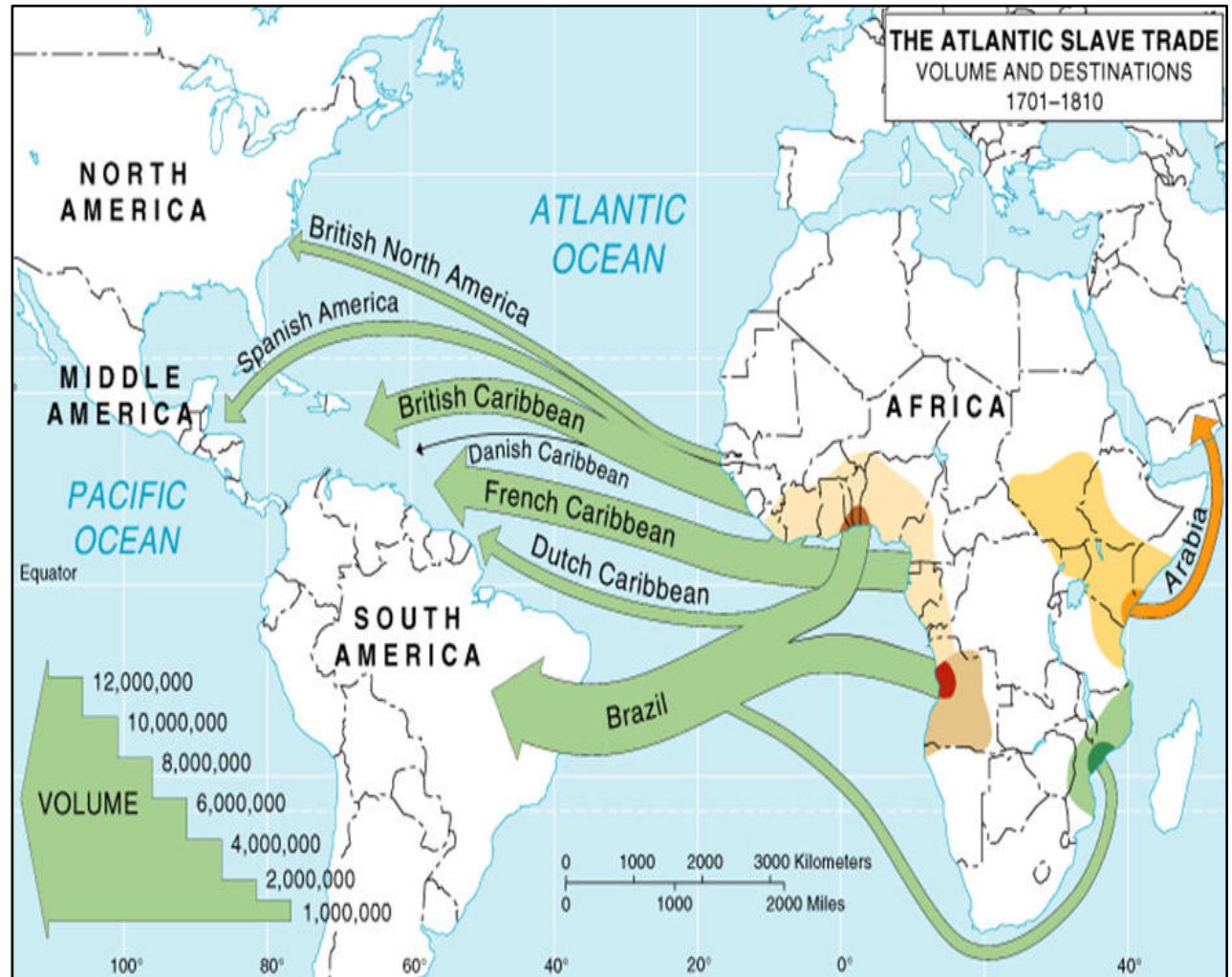
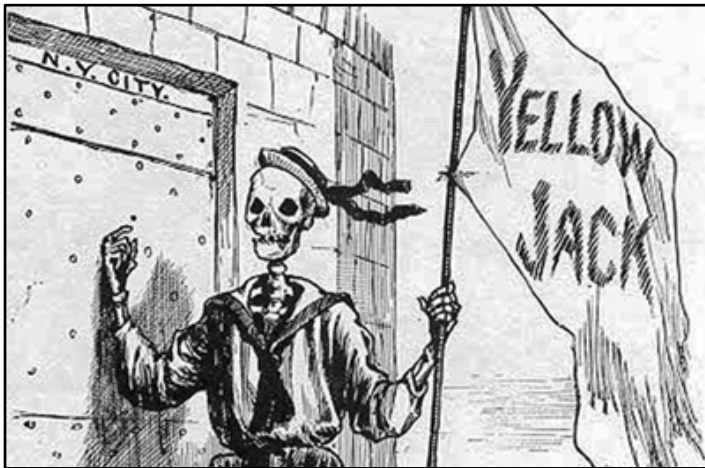
- Globalization
- Urbanization
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3. Control of emerging arboviruses

- Conventional vector control
- Innovative strategies

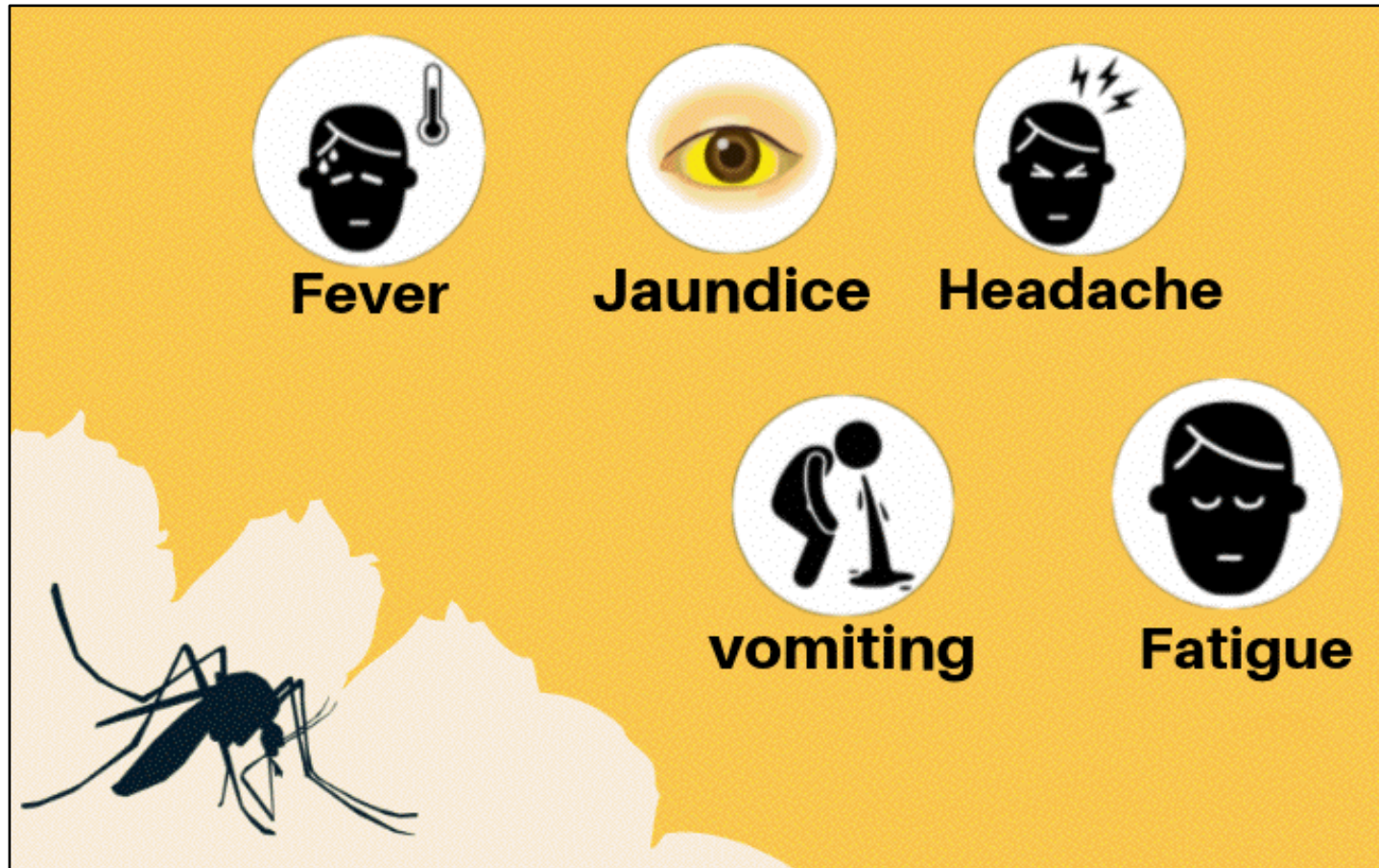
Yellow fever virus (YFV)

- Originally from tropical Africa
- Subsequently established in South America

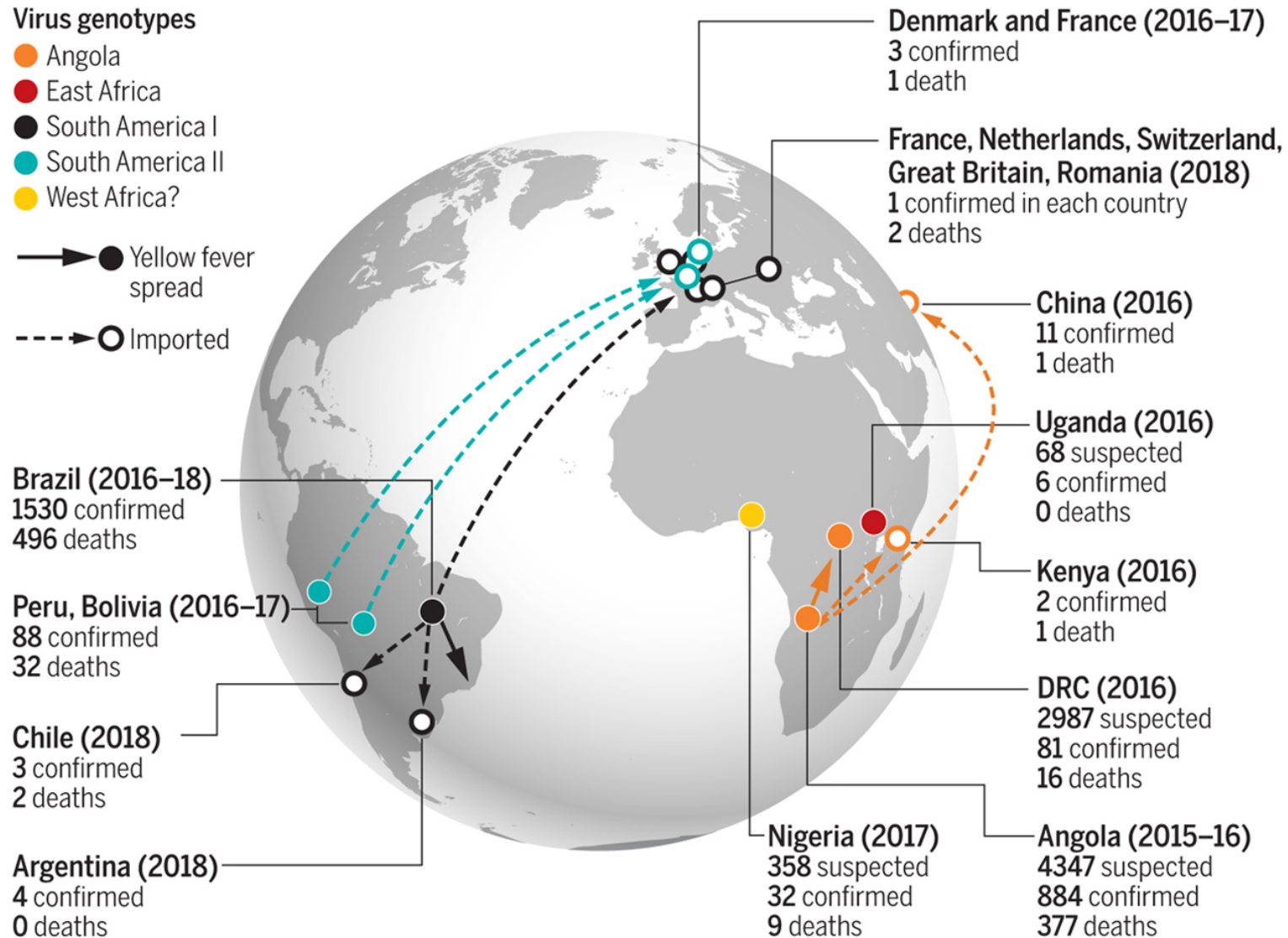


Yellow fever

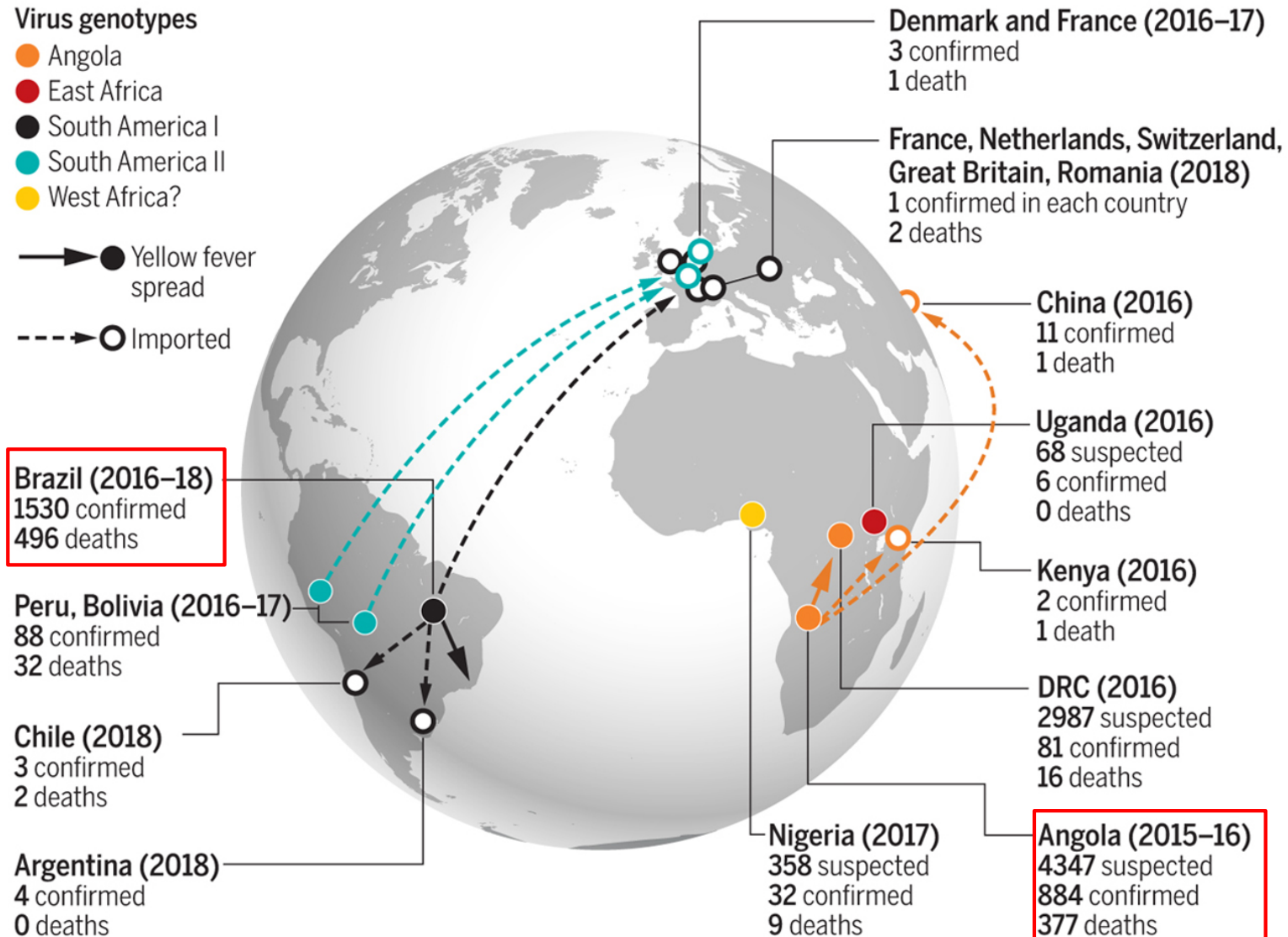
- Hemorrhagic fever
- Case-fatality rate up to 50%
- 200,000 cases /year
- 30,000 deaths /year



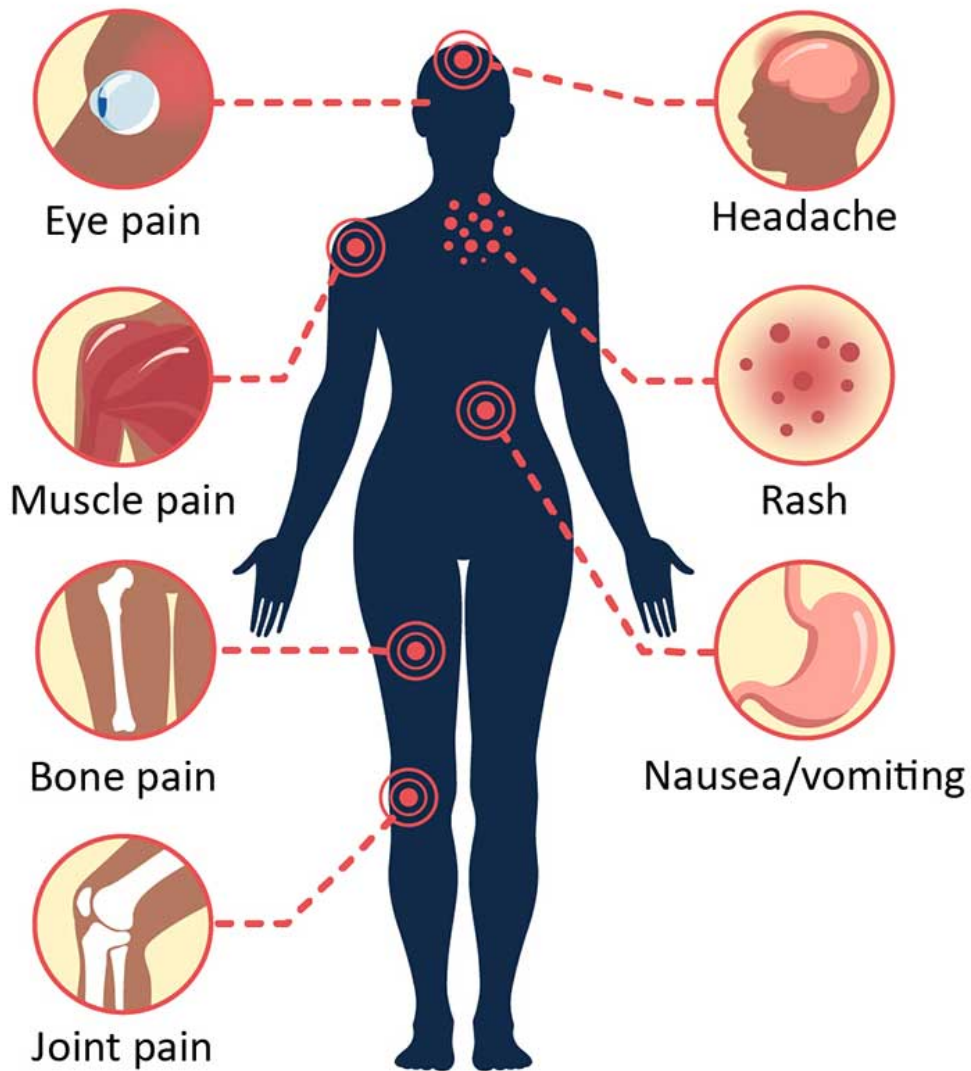
Yellow fever outbreaks 2016-2018



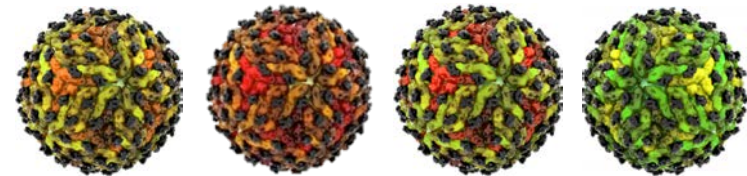
Yellow fever outbreaks 2016-2018



Dengue fever



- 4 billion people at risk
- 100 million cases /year
- 20,000 deaths /year

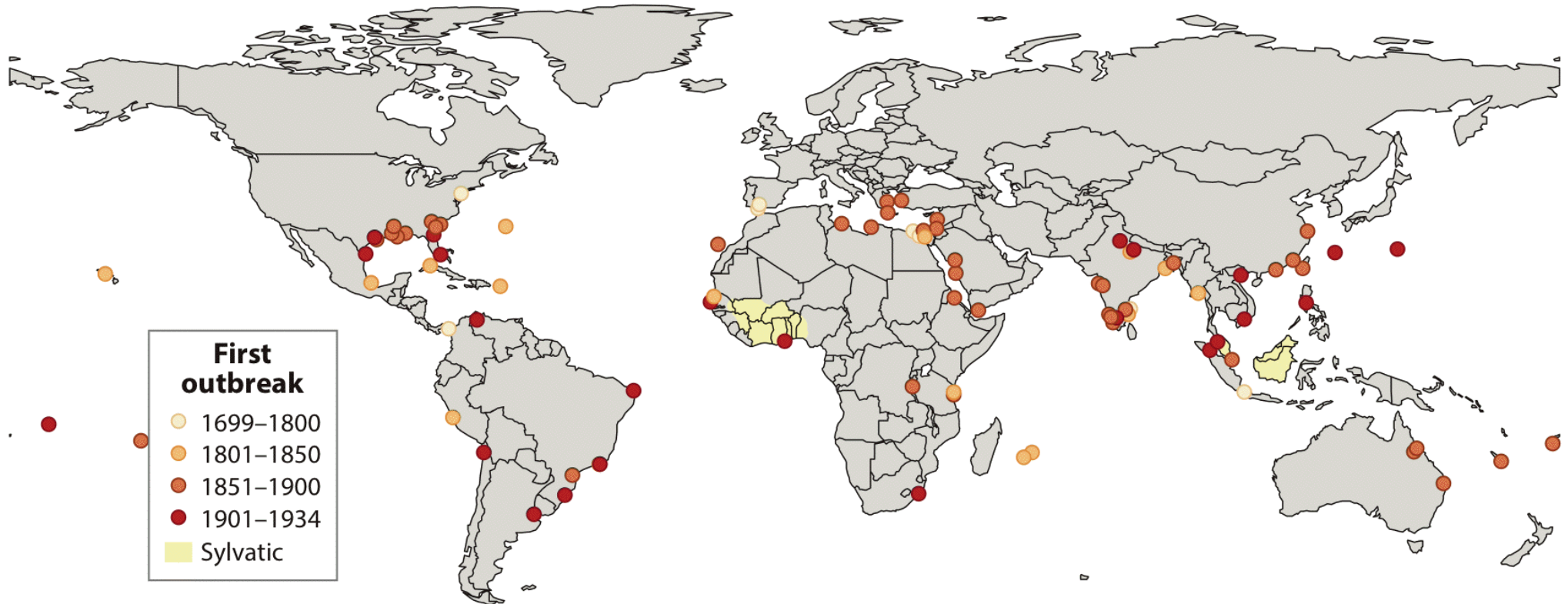


DENV-1, -2, -3, and -4

- Genetically close but distinct
- Immunologically cross-reactive

DENV global emergence

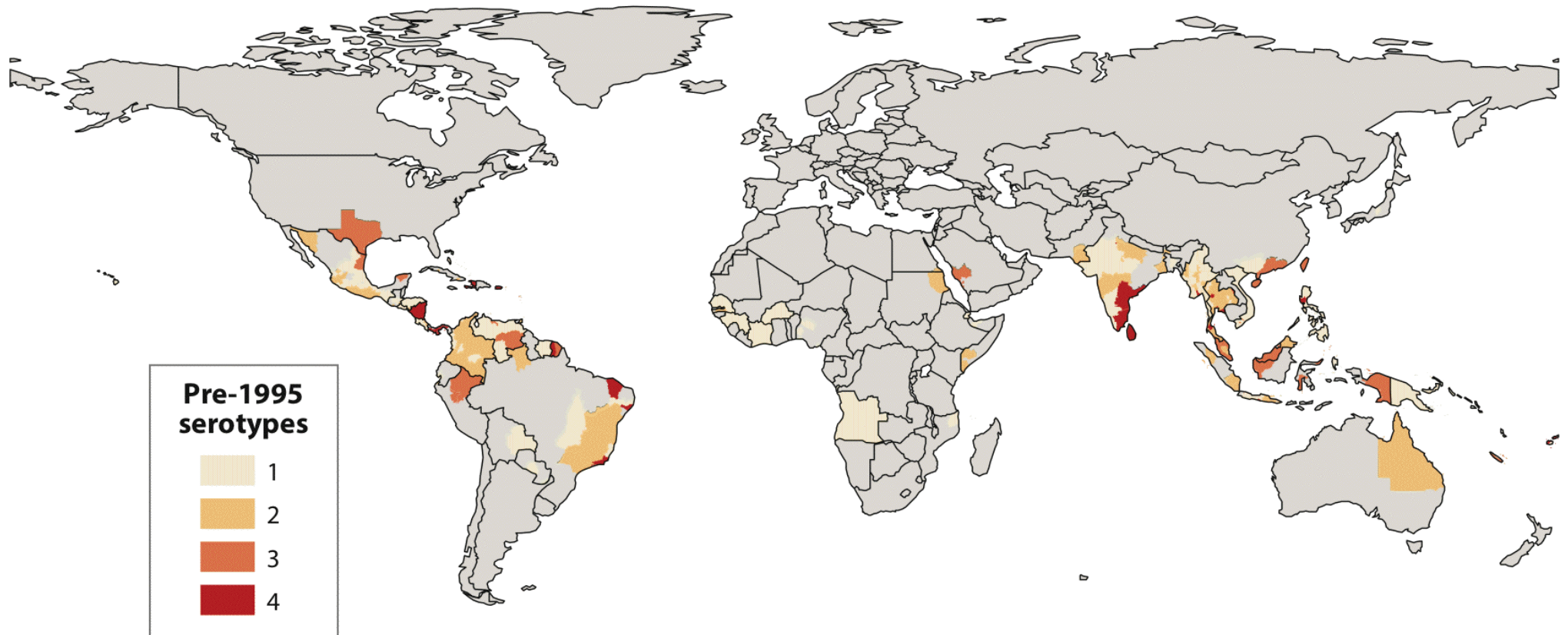
1. First urban outbreaks in port cities increasing after 18th century



Cities with evidence of dengue-like illness prior to 1945

DENV global emergence

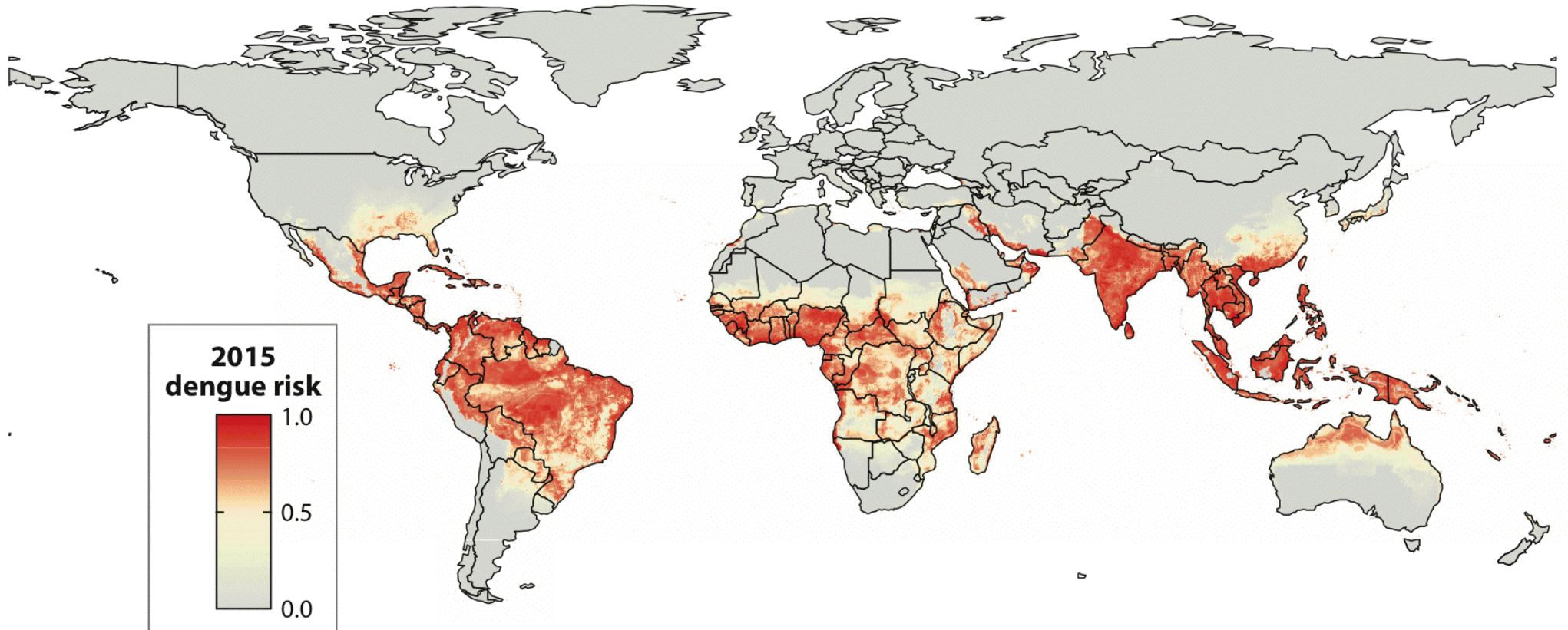
2. Co-circulation of serotypes and emergence of hemorrhagic dengue



Cumulative number of DENV serotypes during 1945-1994

DENV global emergence

3. Consolidation throughout tropical belt and into rural areas

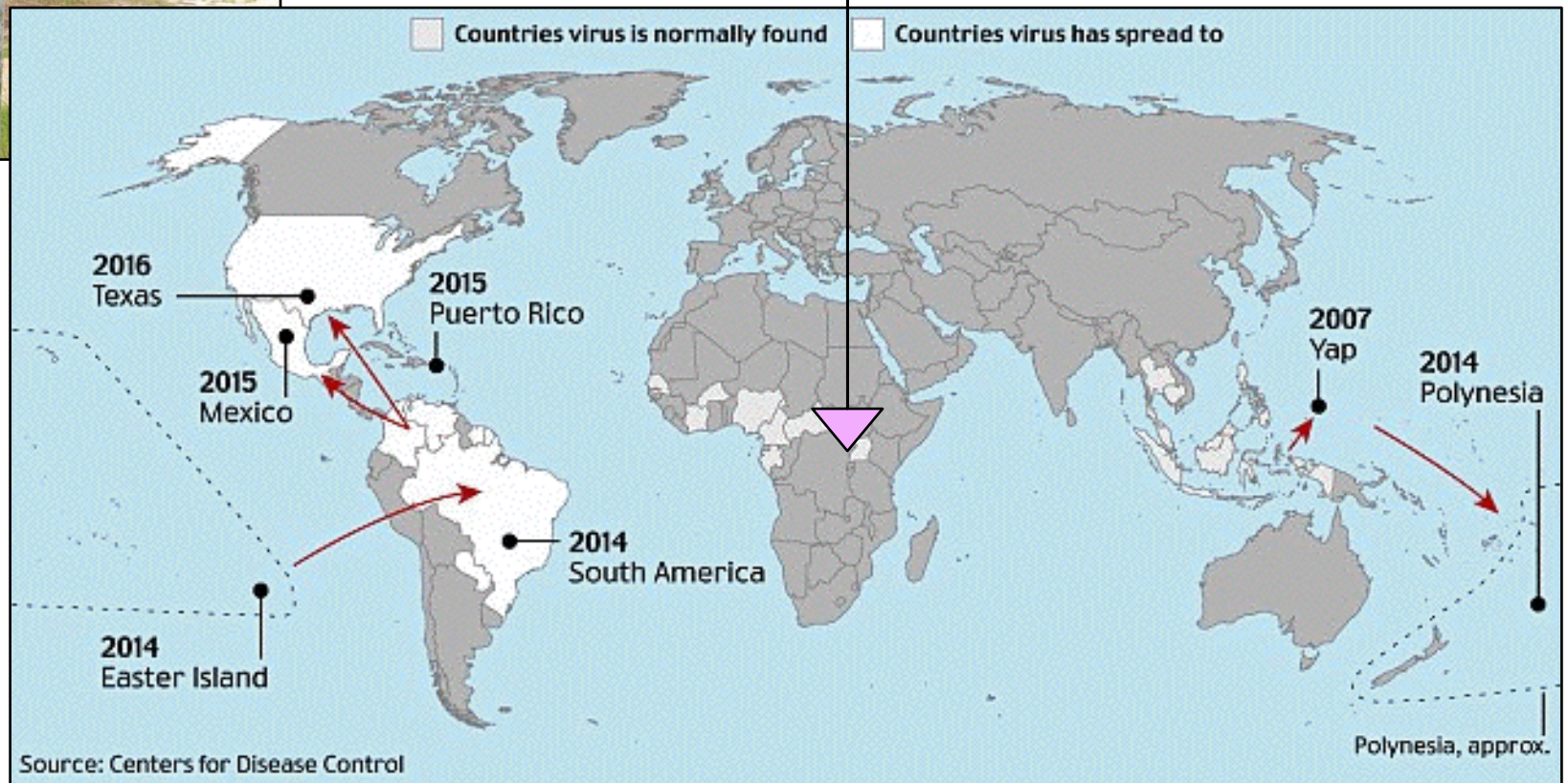
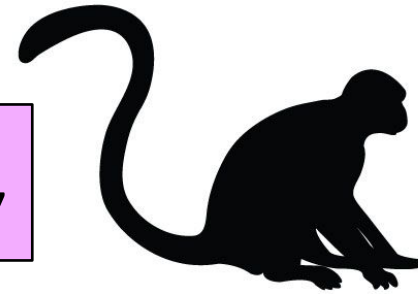


Contemporary predicted global distribution of dengue risk

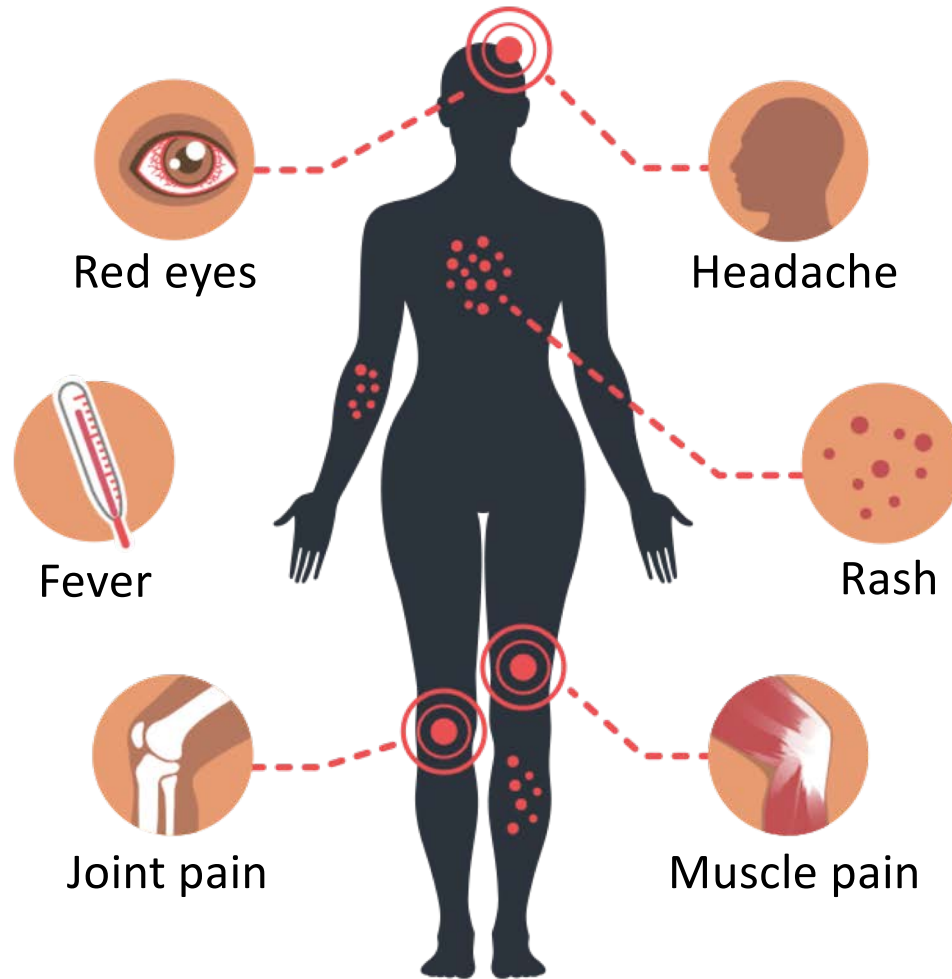
Global Zika virus (ZIKV) emergence



1st isolation
Uganda 1947



ZIKV clinical features



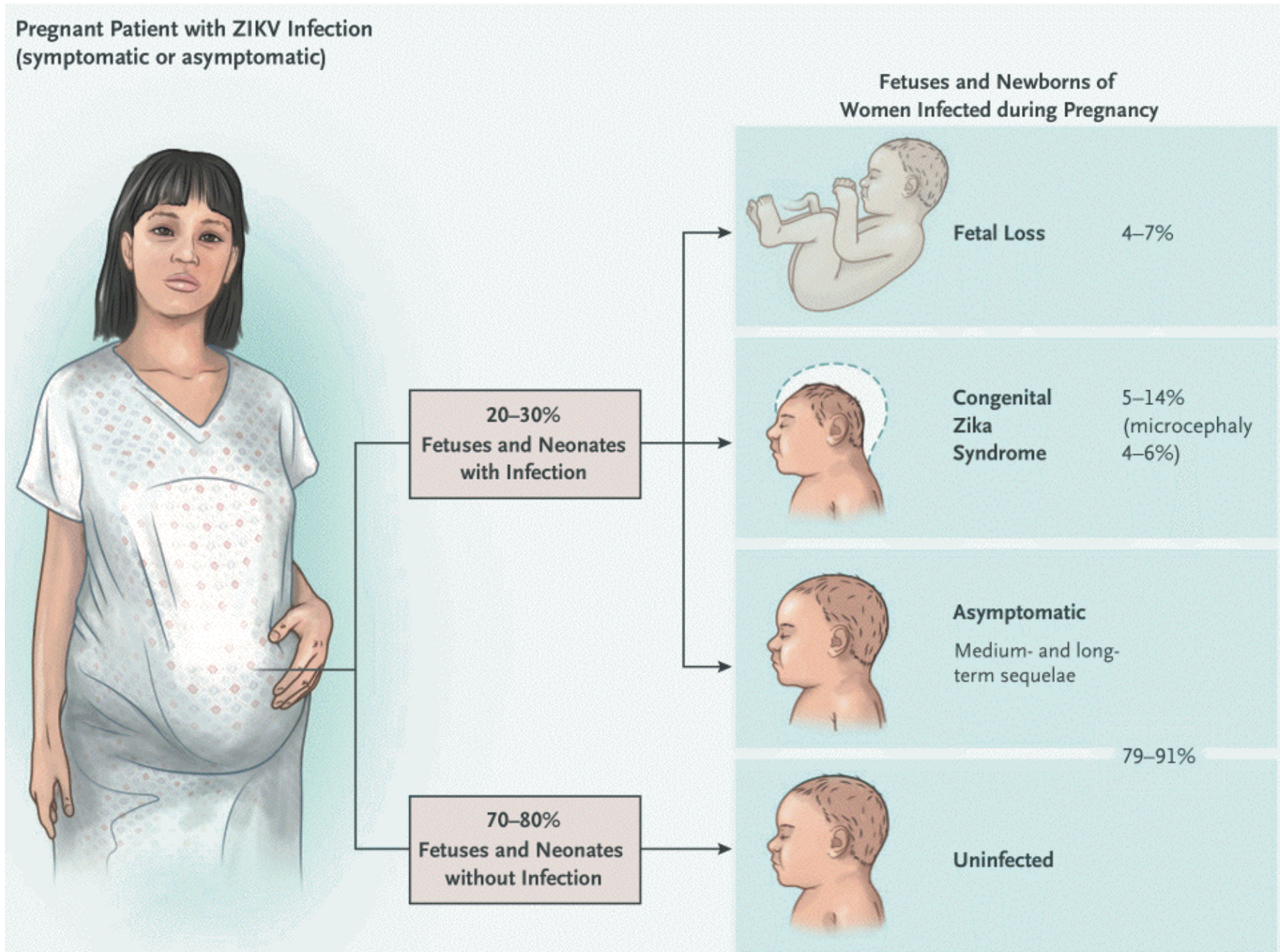
Infection

- Mosquito-borne
- Sexual
- Pregnancy

Disease

- Asymptomatic 50-80%
- Mild disease 20-50%
- Severe complications <1%
- Case-fatality rate <0.01%

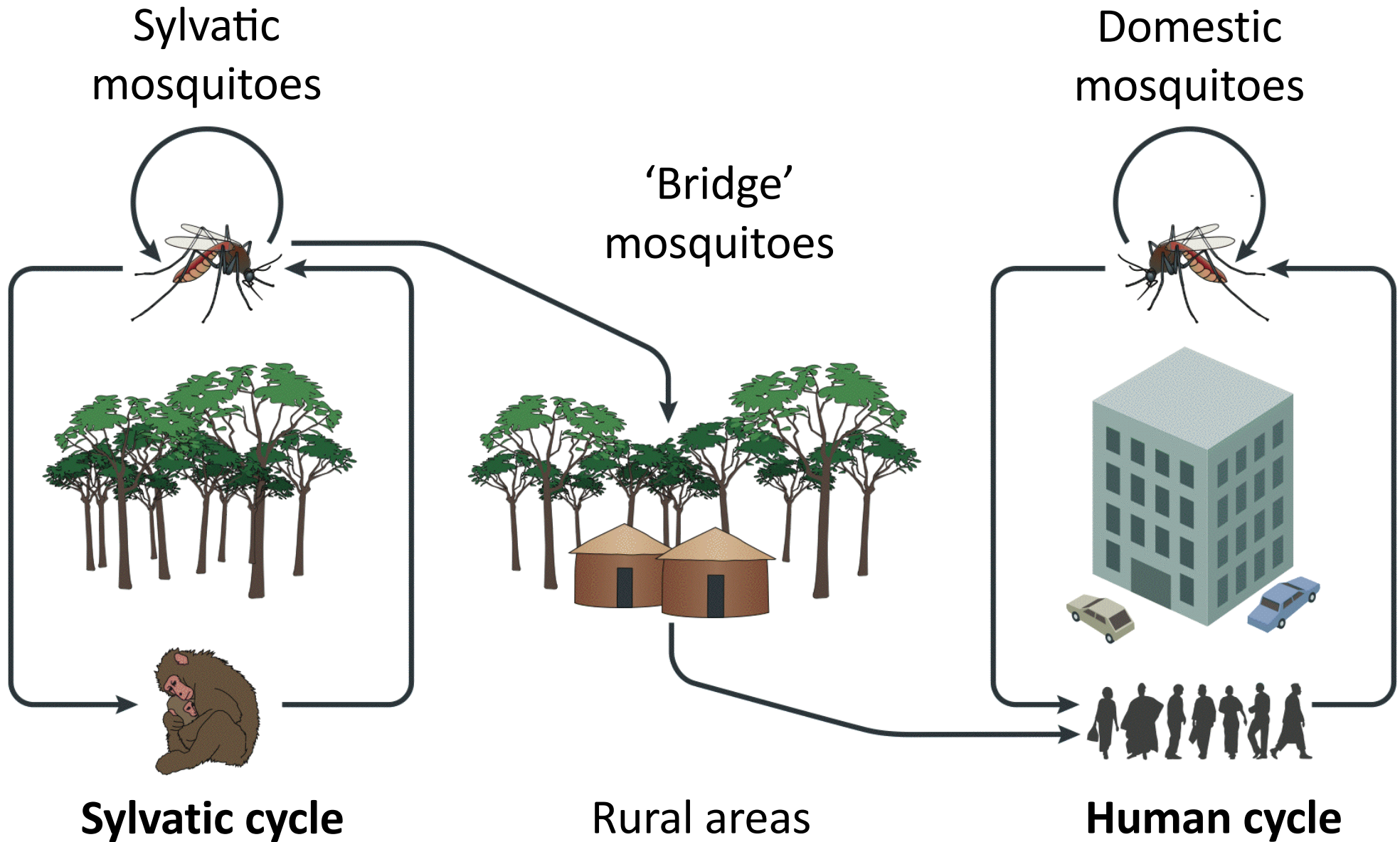
ZIKV-associated fetal and birth defects



Outline

1. Examples of emerging arboviruses
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 - Zika
2. **Factors underlying arbovirus emergence**
 - Globalization
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Arbovirus transmission cycles



Factors driving arbovirus emergence

Globalization

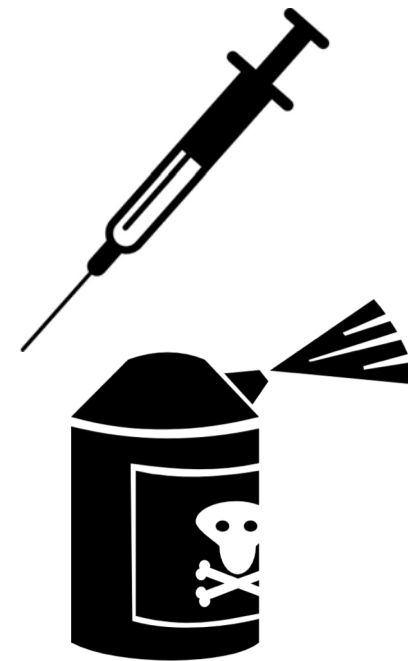


Urbanization

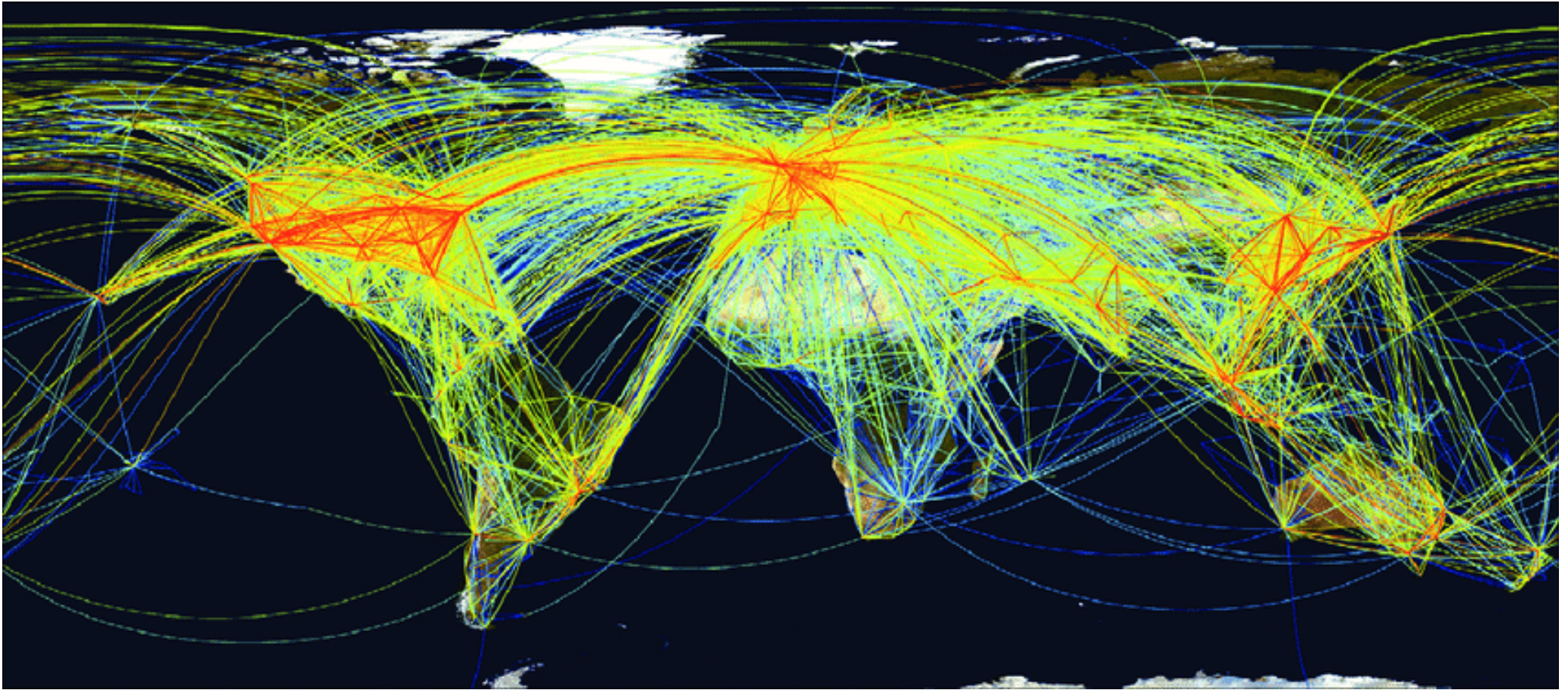


Limited prevention methods:

- Few licensed vaccines
- Pesticide-based control of vector populations difficult to sustain



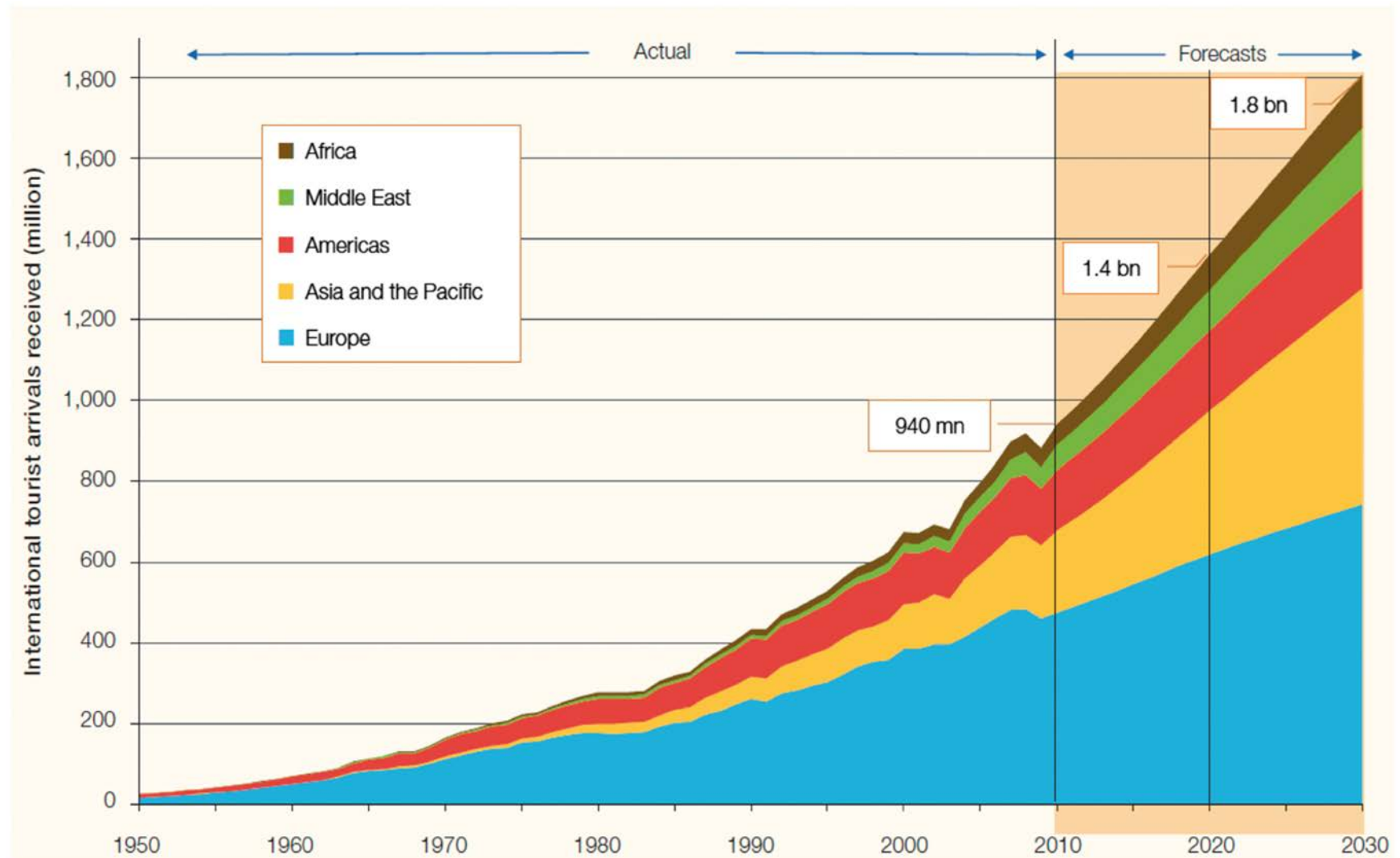
Global aviation network



Lines show direct links between airports and passenger capacity

- Red: >1000 passengers/day
- Yellow: >100 passengers/day
- Blue: >10 passengers/day

International tourist arrivals

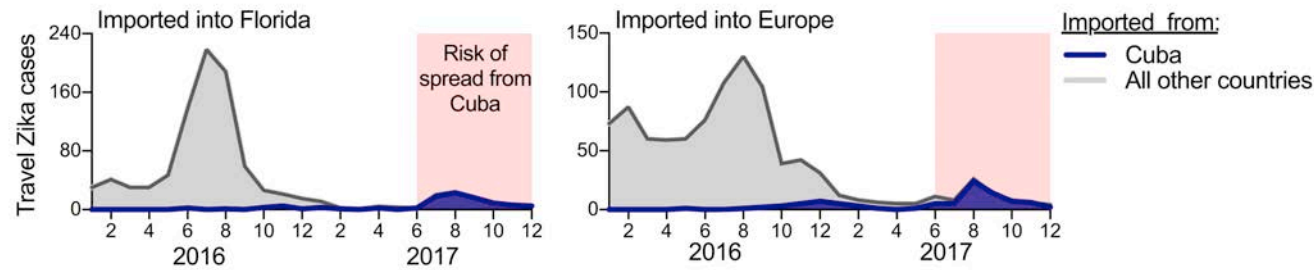


Source: UNWTO (2016a)

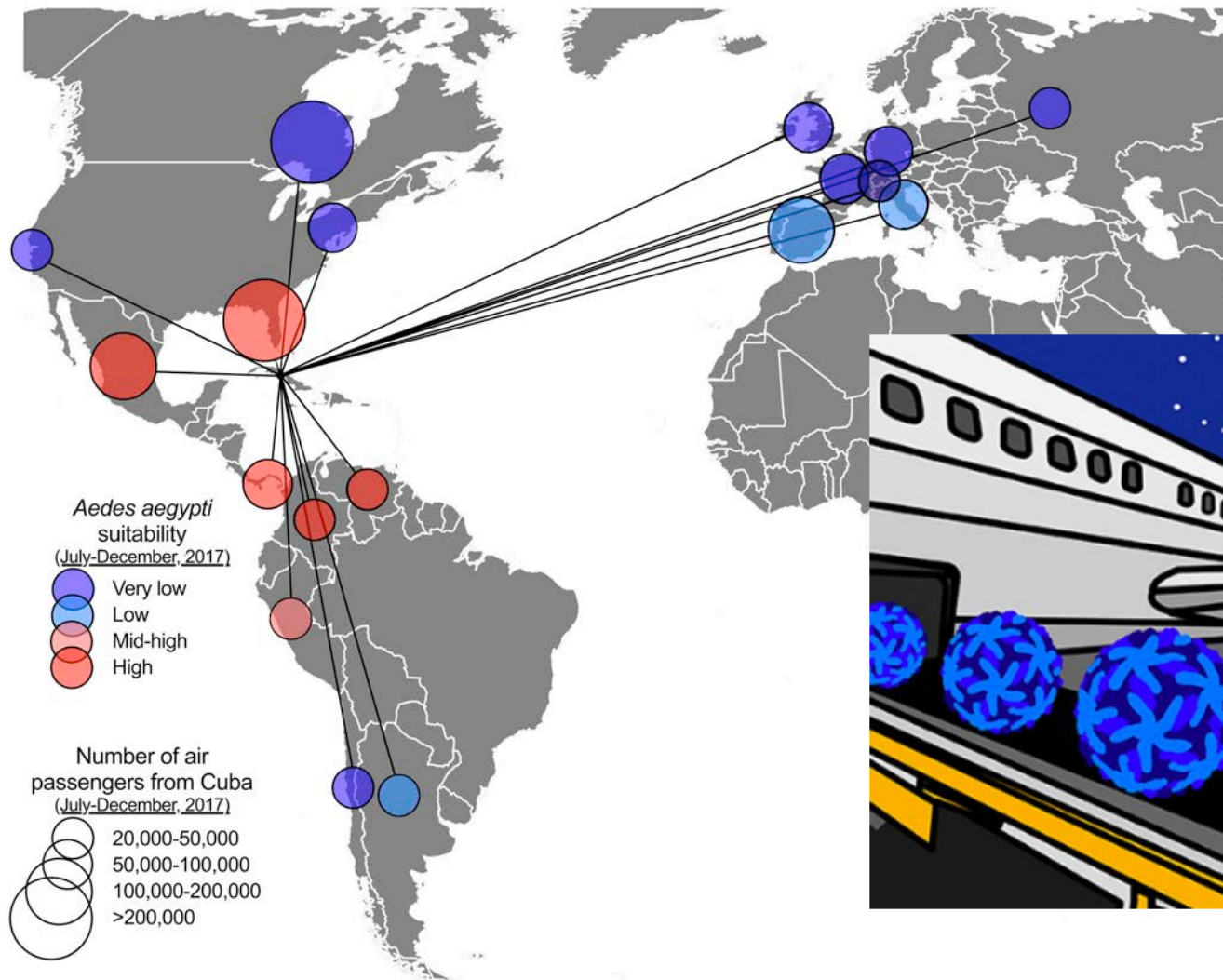
International travel mediates arbovirus dispersal



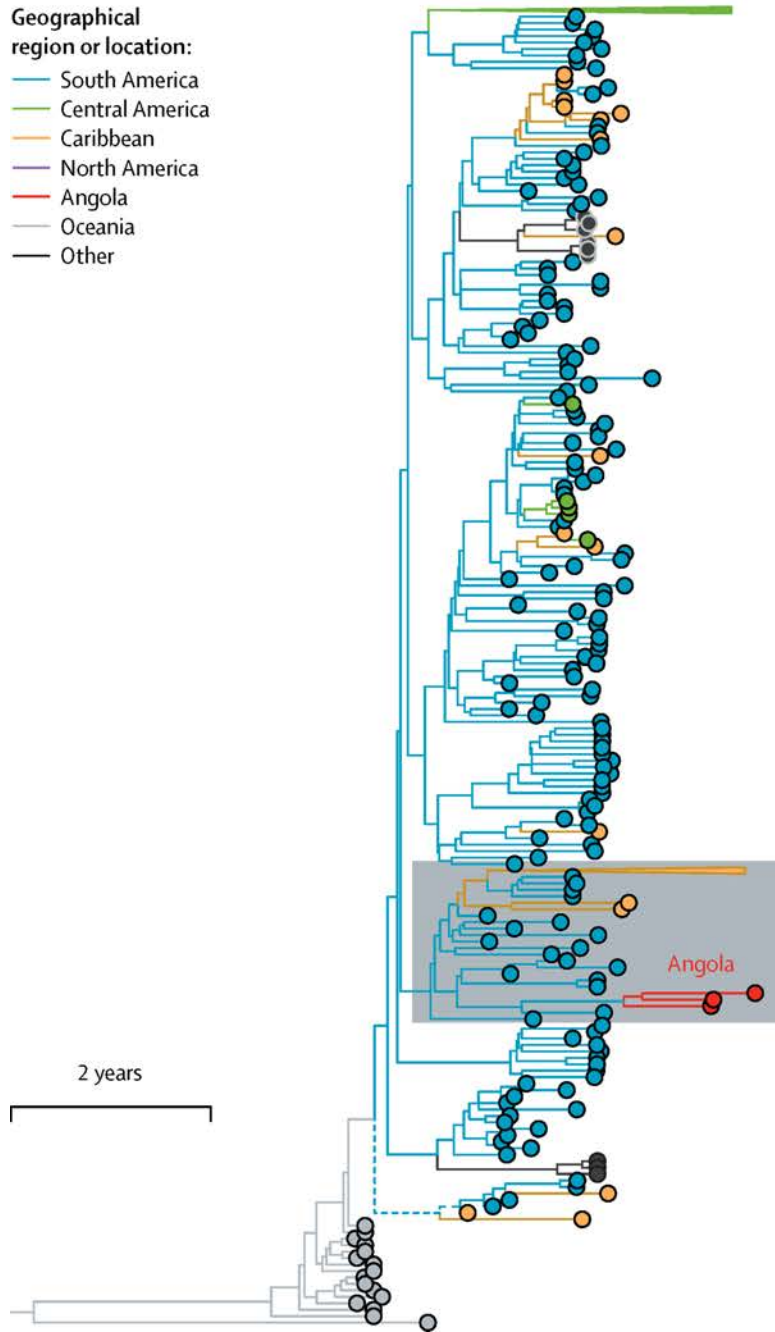
Travel surveillance uncovers ZIKV outbreak in Cuba



>98% of all travel-associated Zika cases during June–December 2017 came from Cuba



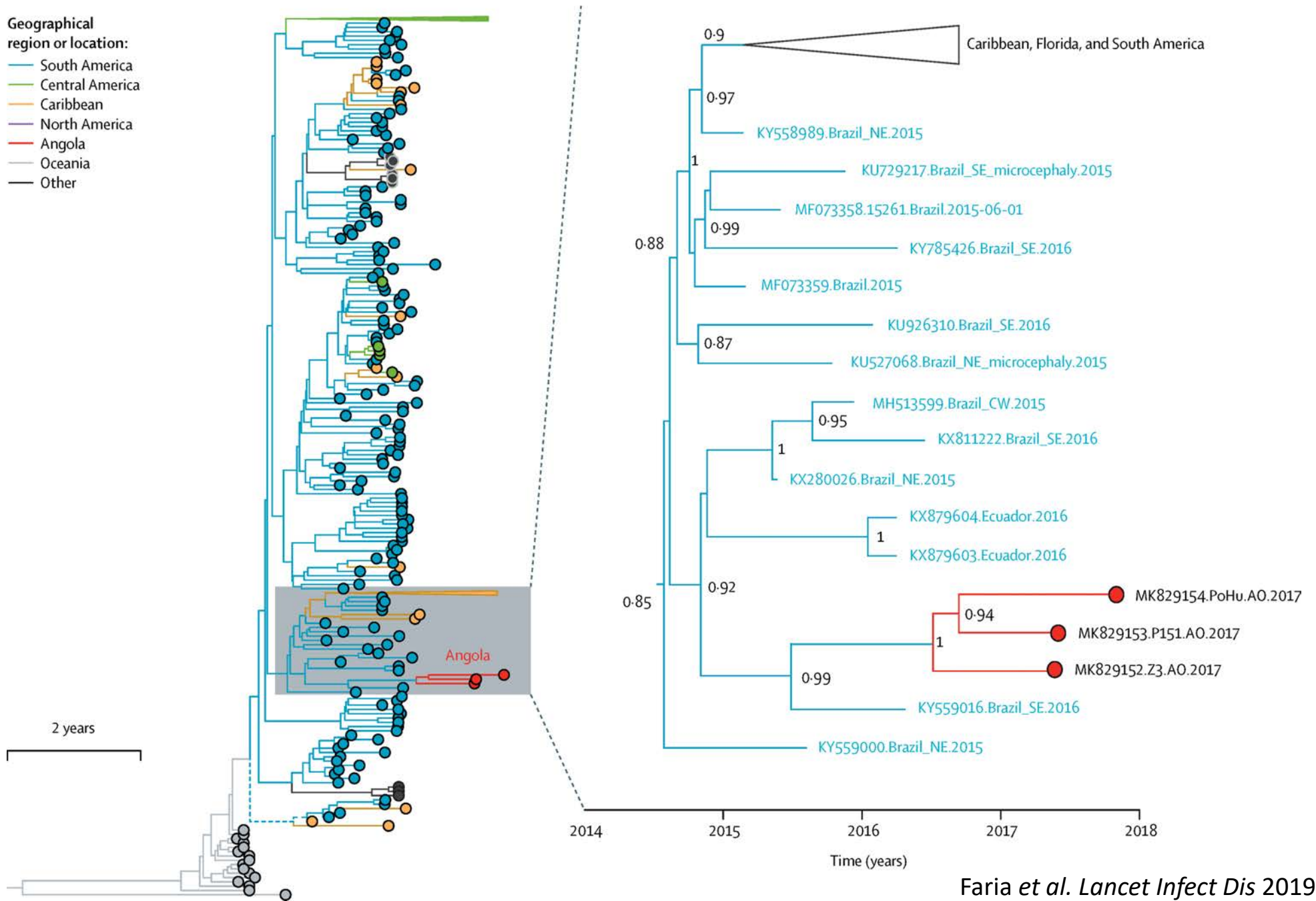
ZIKV introduction into Angola from Brazil



ZIKV introduction into Angola from Brazil

Geographical region or location:

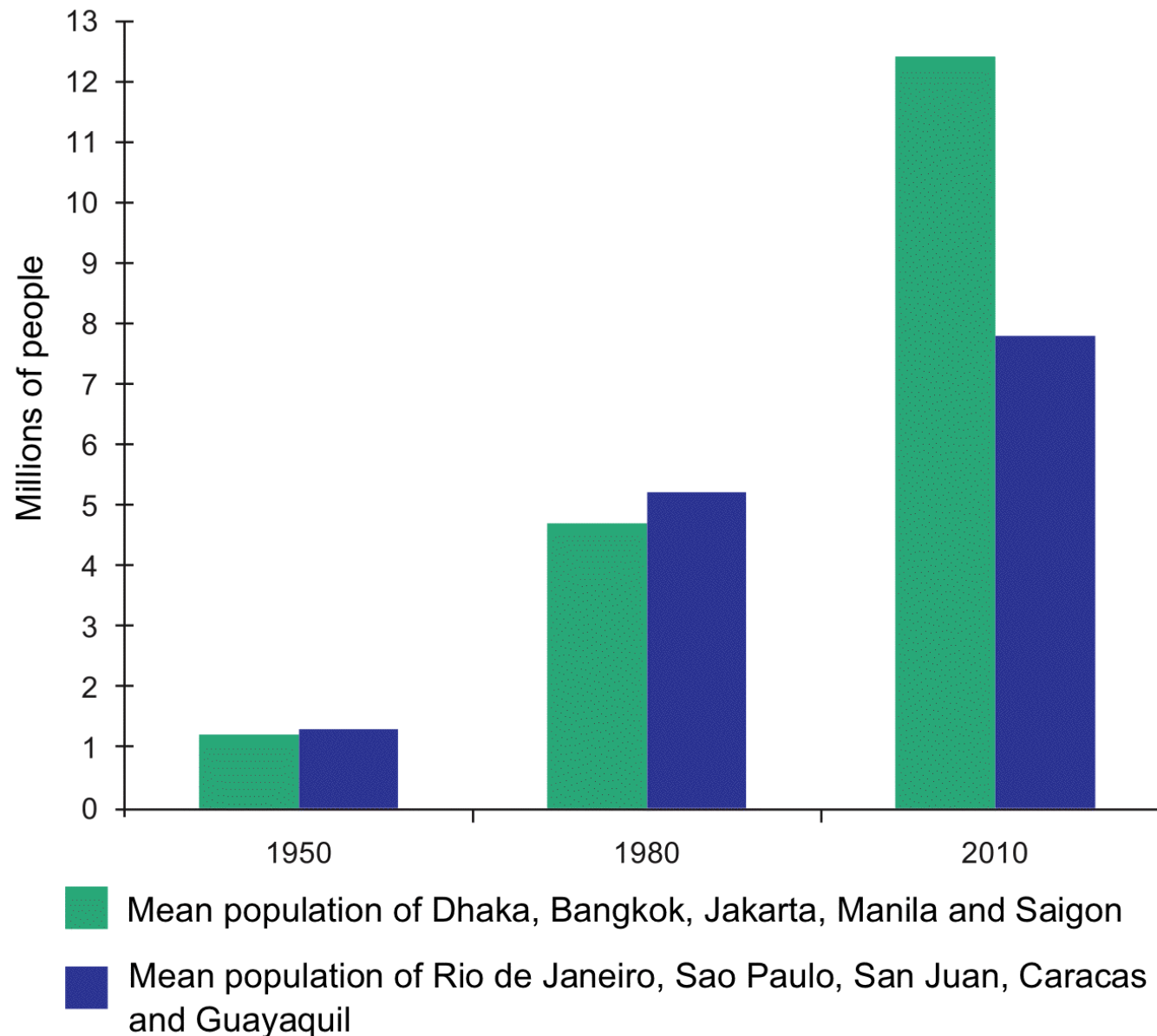
- South America
- Central America
- Caribbean
- North America
- Angola
- Oceania
- Other



Urbanization



Urban growth in tropical megacities



Domestic *Aedes* vectors of arboviruses



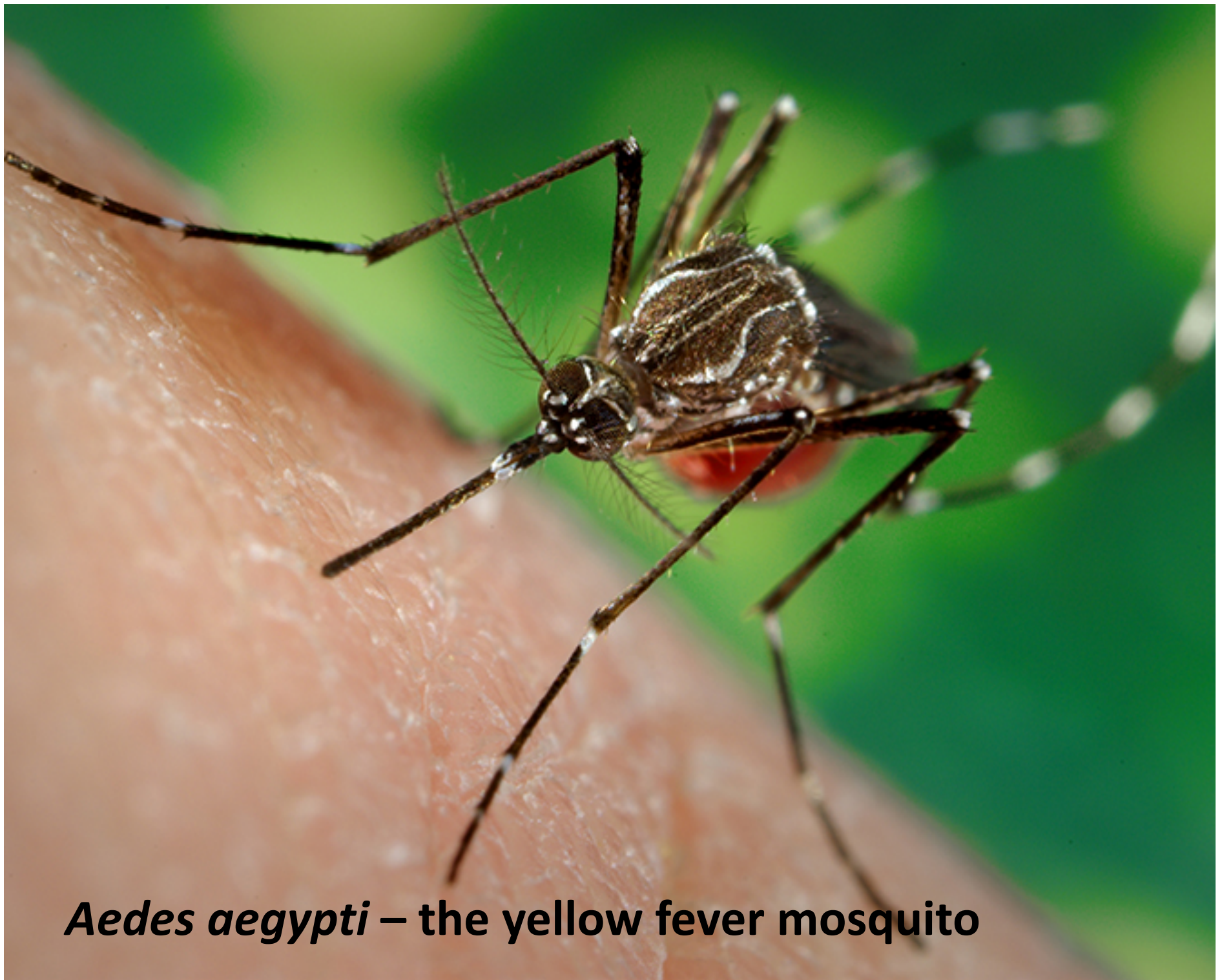
Aedes aegypti



Aedes albopictus



- Thrive in the human environment
- Breed in artificial containers
- Feed on humans preferentially
- Bite during daytime
- Lay desiccation-resistant eggs

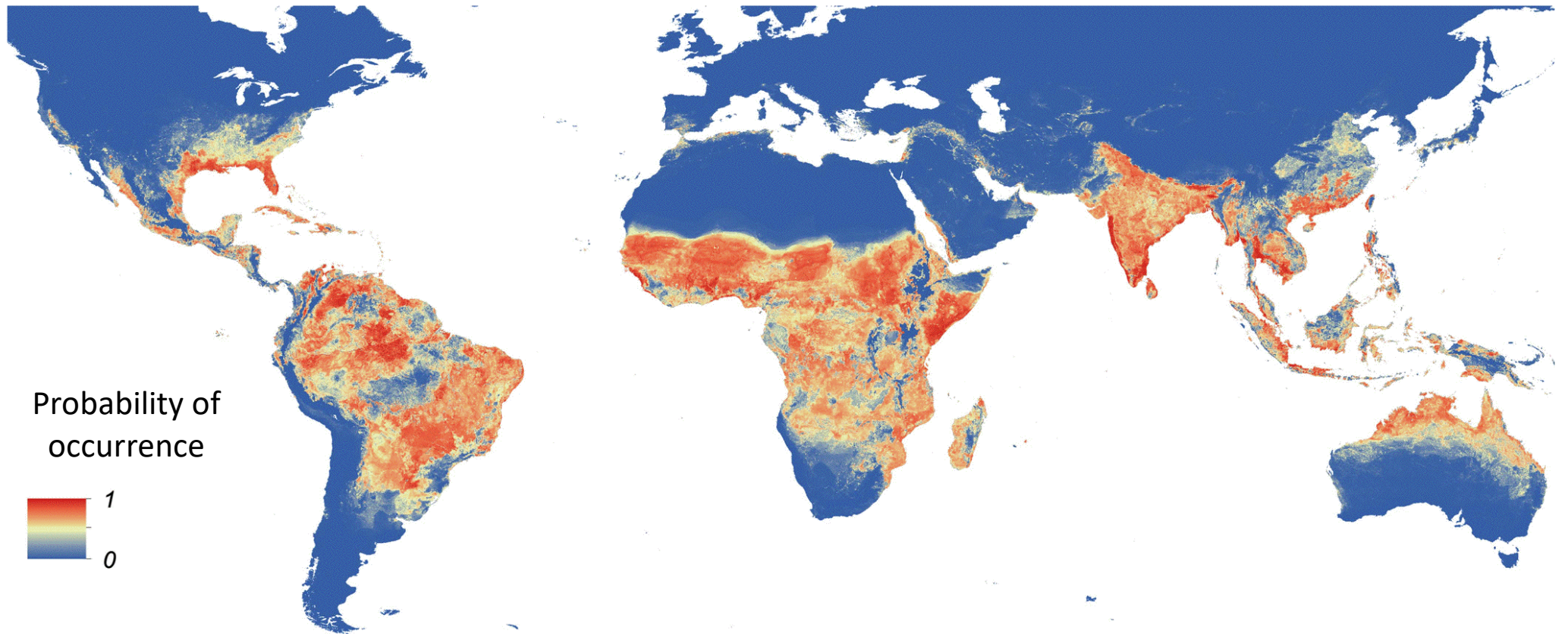


***Aedes aegypti* – the yellow fever mosquito**

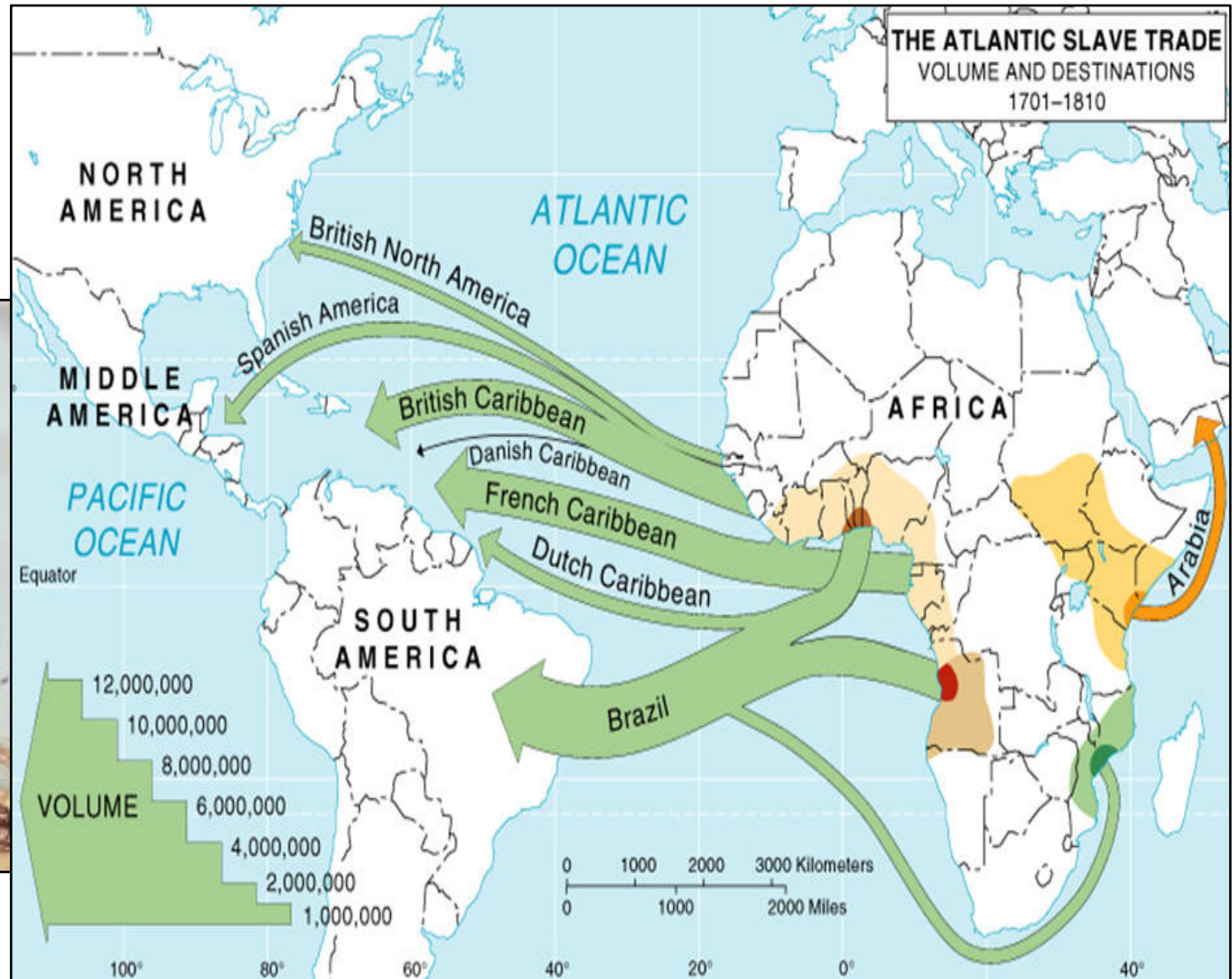
Aedes aegypti



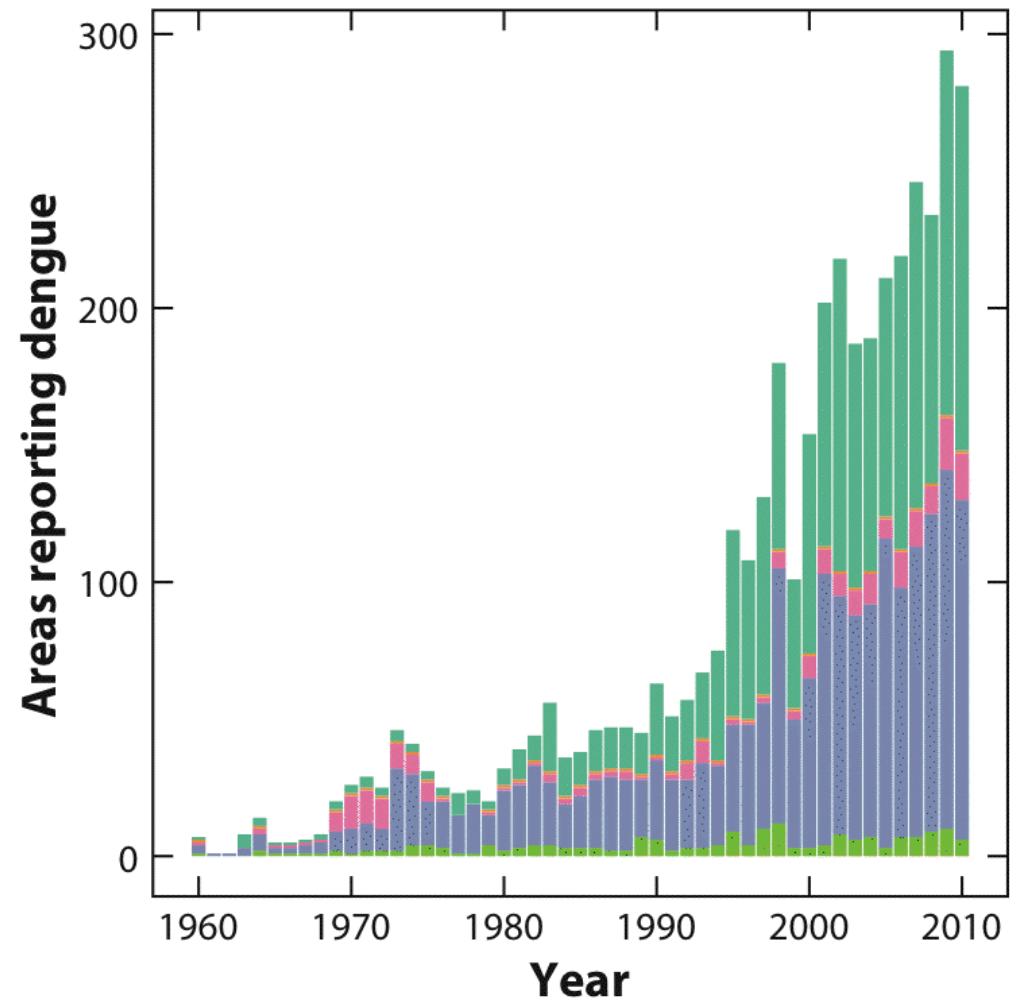
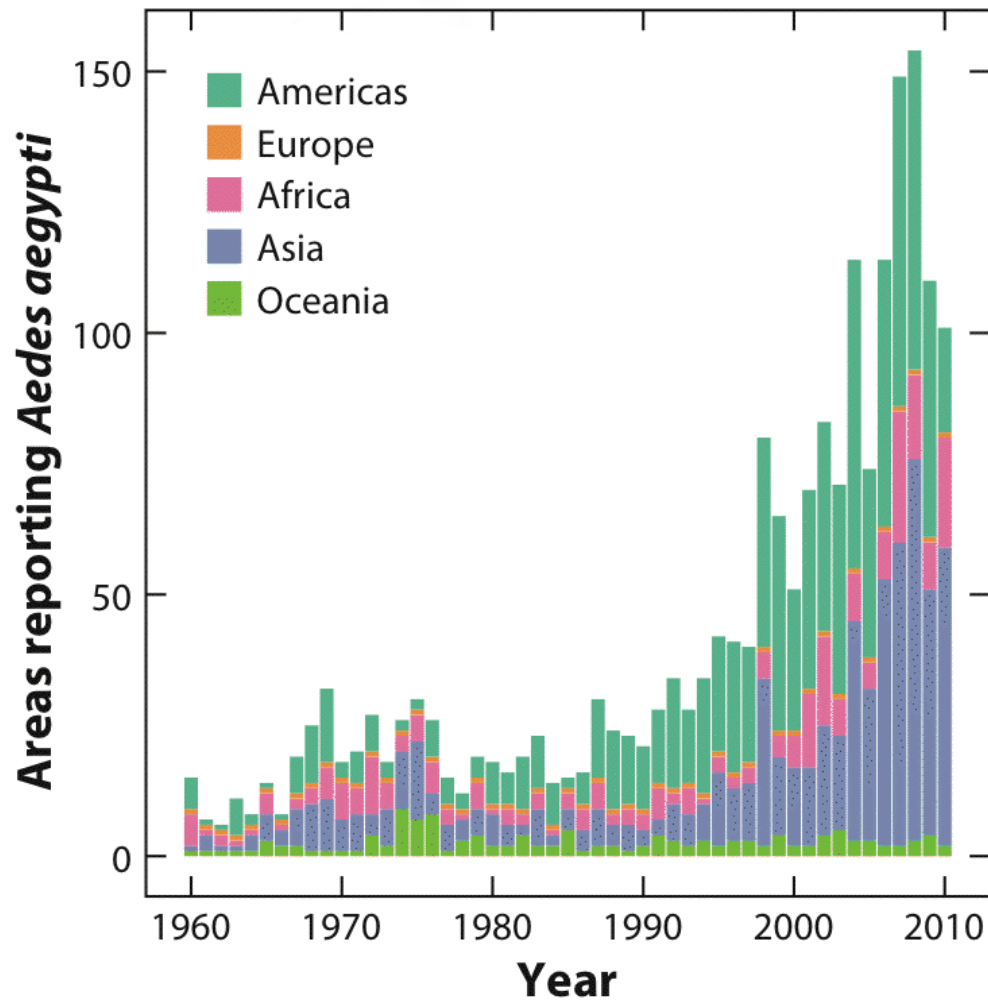
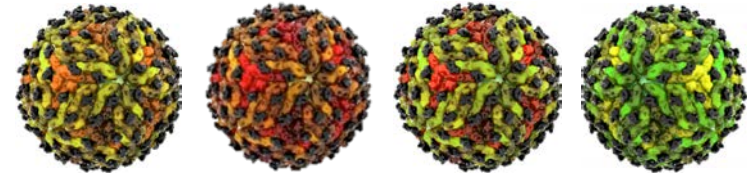
- Originally from Africa
- Spread out in the last few centuries
- Found in tropical and sub-tropical regions



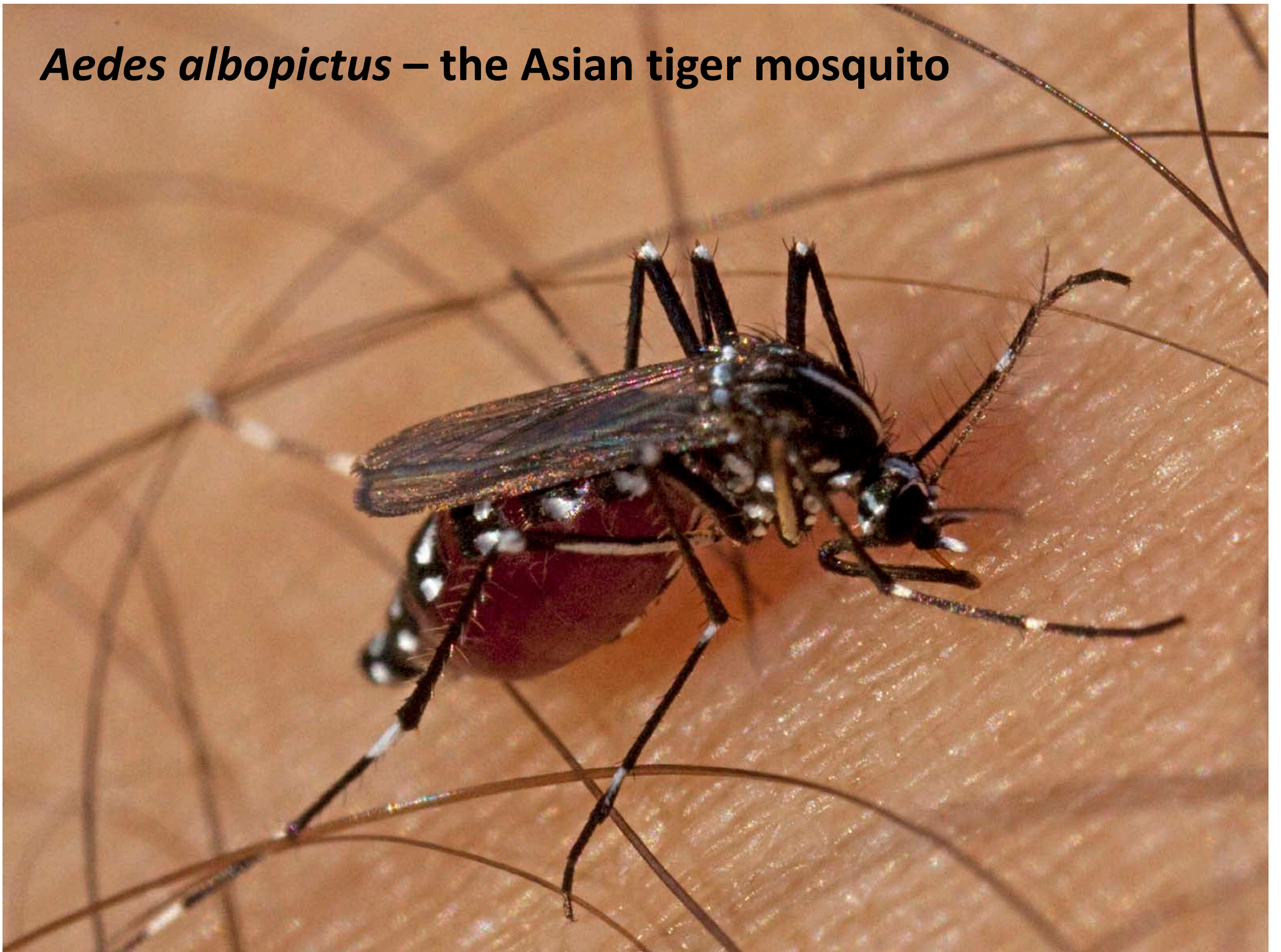
Spread of *Aedes aegypti* out of Africa



Aedes aegypti expansion enabled and possibly drove dengue emergence



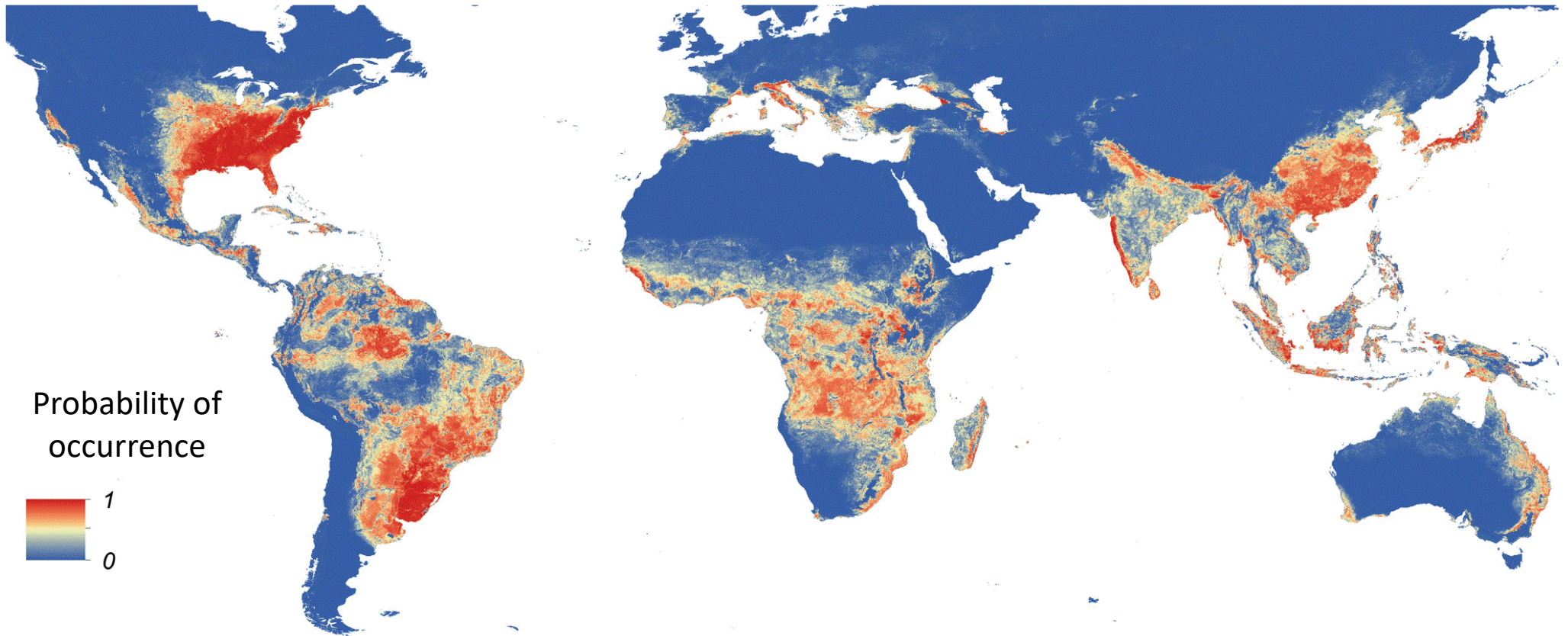
***Aedes albopictus* – the Asian tiger mosquito**



Aedes albopictus



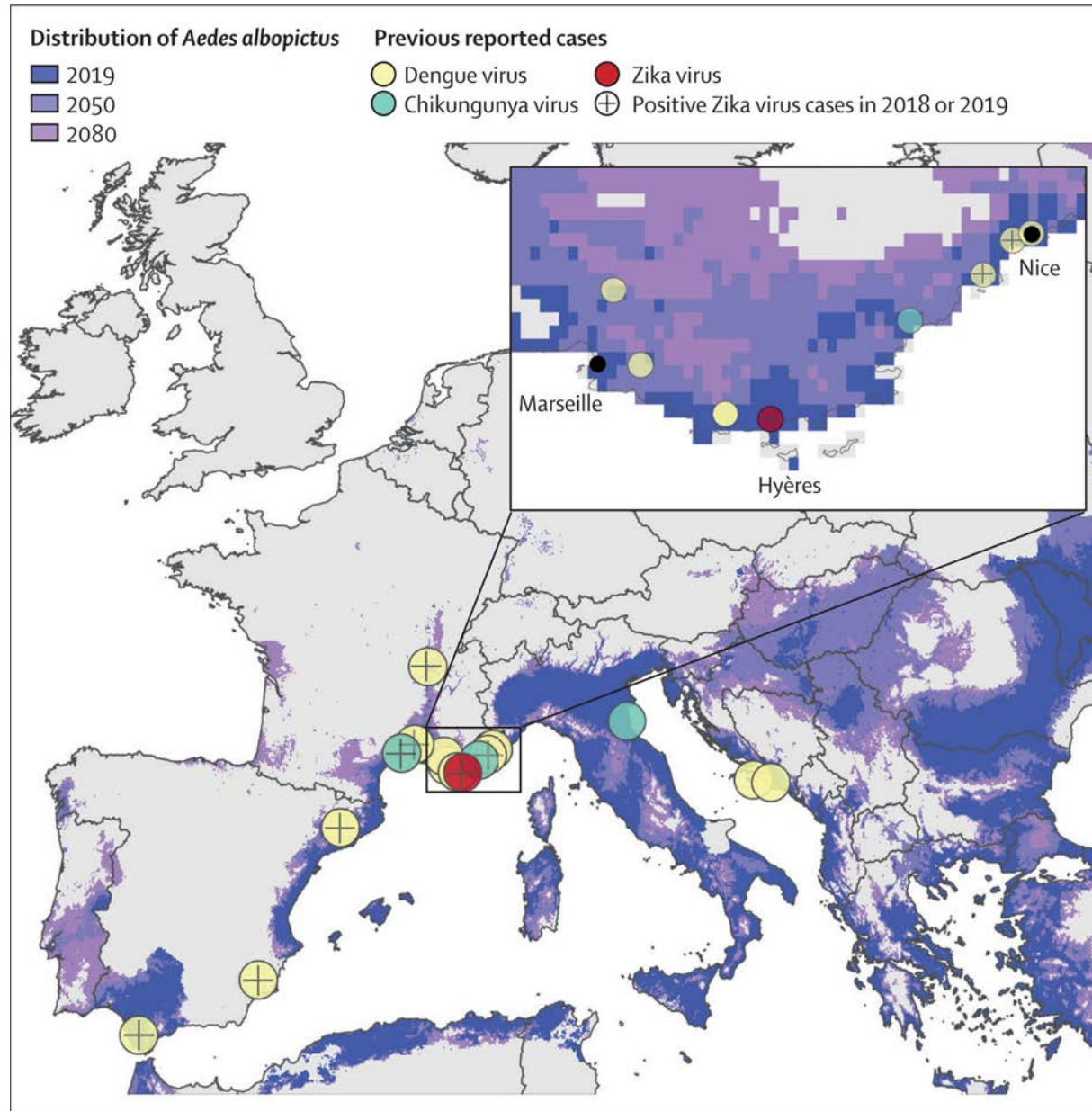
- Originally from Asia
- Spread out in the last few decades
- Found in tropical and temperate regions



Global spread of *Aedes albopictus*



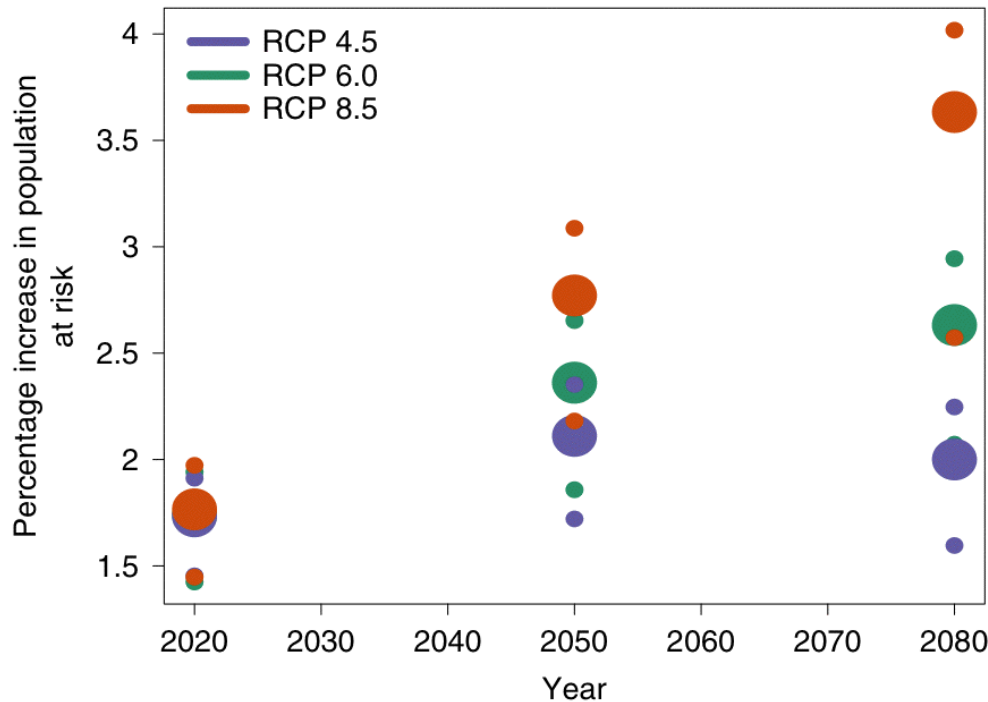
Aedes albopictus is responsible for local transmission of arboviruses in Europe



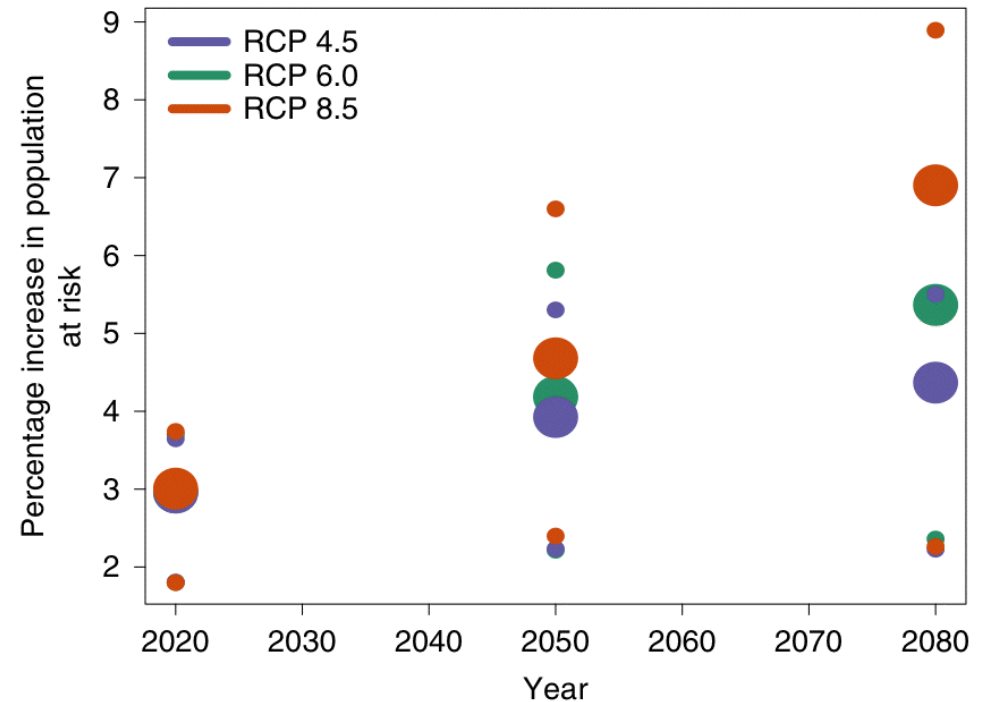
Geographical expansion of *Aedes*-infested areas



Ae. aegypti



Ae. albopictus



Global population predicted to live in *Aedes*-suitable areas for 3 climatic scenarios:

- RCP 4.5 = conservative
- RCP 6.0 = medium
- RCP 8.5 = worst-case

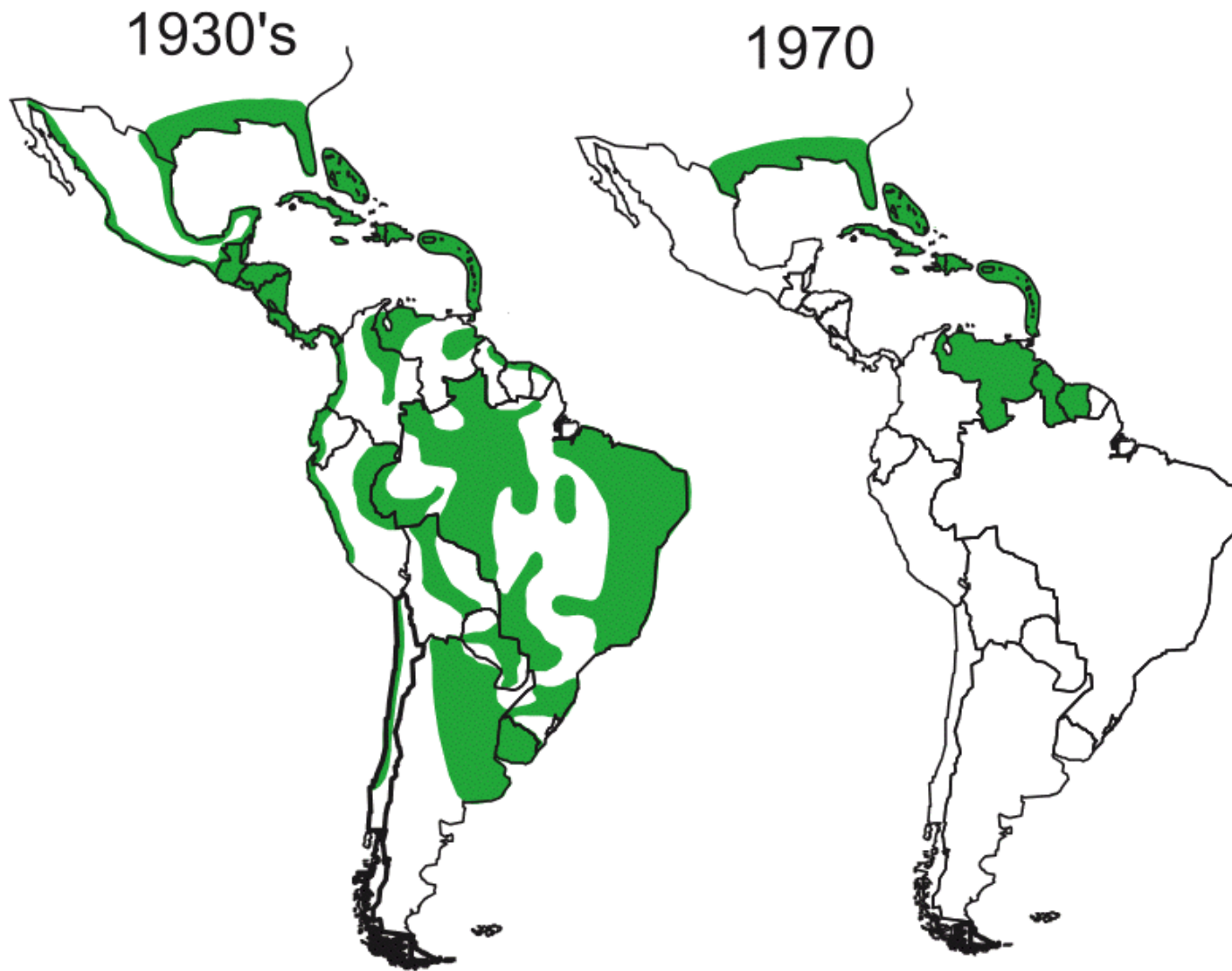
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Conventional vector control

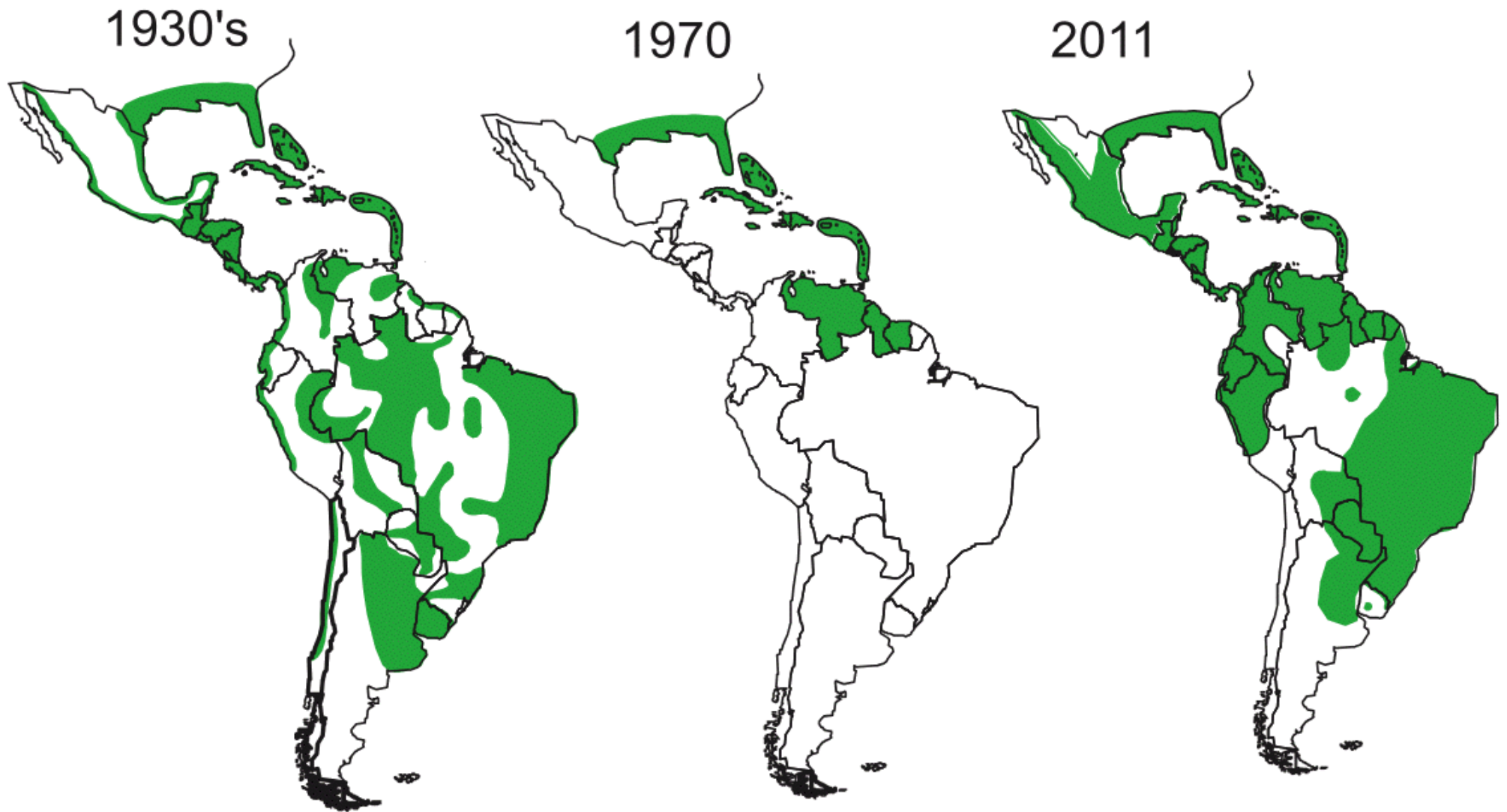


PAHO *Aedes aegypti* eradication campaign



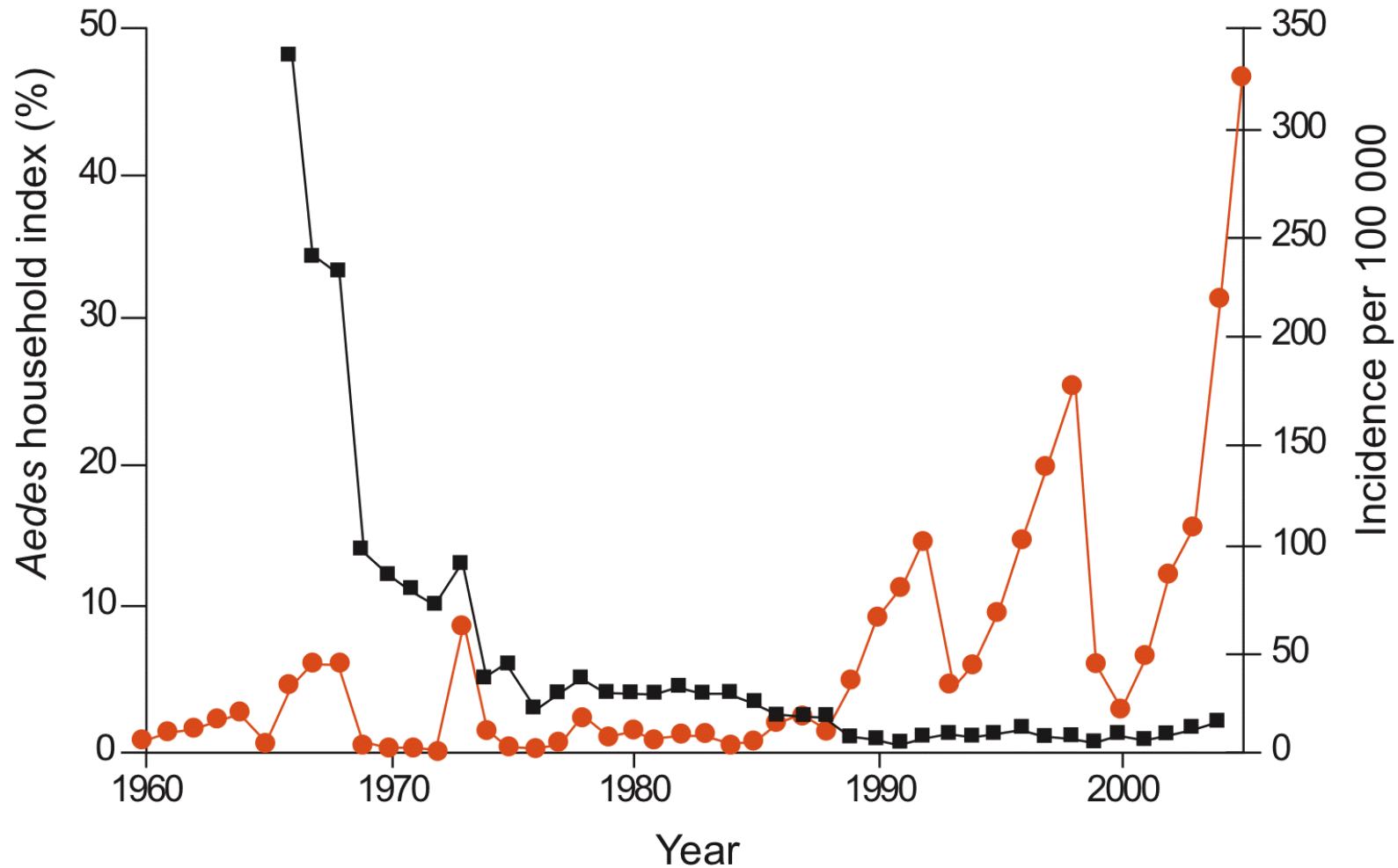
- *Ae. aegypti* eliminated from 20 countries between 1947 and 1962

PAHO *Aedes aegypti* eradication campaign



- *Ae. aegypti* eliminated from 20 countries between 1947 and 1962
- *Ae. aegypti* reinfestation after end of program in 1985

Dengue re-emergence in Singapore despite intensive vector control program



- Observed annual average *Aedes* household index
- Annual clinical incidence of dengue fever

Basic reproduction number of arboviruses



R_0 is the total number of infectious vector bites that arise from one infected person introduced into a susceptible human population

Basic reproduction number of arboviruses

$$R_0 = \frac{ma^2 p^n b}{-\ln(p)} \frac{c}{r}$$

R_0 is the total number of infectious vector bites that arise from one infected person introduced into a susceptible human population

Basic reproduction number of arboviruses

$$R_0 = \frac{ma^2 p^n b}{-\ln(p)} \underbrace{\frac{c}{r}}_{\text{HUMAN}}$$

c = human host competence
 r = daily recovery rate of infected humans

R_0 is the total number of infectious vector bites that arise from one infected person introduced into a susceptible human population

Basic reproduction number of arboviruses

$R_0 = \frac{ma^2 p^n b}{-\ln(p)} \frac{c}{r}$

m = vector density per person

a = daily probability of human biting

p = daily probability of vector survival

n = duration in days of extrinsic incubation period (EIP)

b = vector competence

c = human host competence

r = daily recovery rate of infected humans

$\underbrace{\hspace{10em}}_{\text{VECTOR HUMAN}}$

R_0 is the total number of infectious vector bites that arise from one infected person introduced into a susceptible human population

Medical interventions indirectly target human-related parameters

$$R_0 = \frac{ma^2 p^n b}{-\ln(p)} \frac{c}{r}$$

m = vector density per person

a = daily probability of human biting

p = daily probability of vector survival

n = duration in days of extrinsic incubation period (EIP)

b = vector competence

c = human host competence

r = daily recovery rate of infected humans



R_0 is the total number of infectious vector bites that arise from one infected person introduced into a susceptible human population

Conventional vector control strategies target primarily vector-related factors



m = vector density per person

a = daily probability of human biting

p = daily probability of vector survival

n = duration in days of extrinsic incubation period (EIP)

b = vector competence

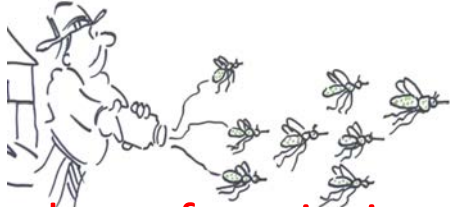
c = human host competence

r = daily recovery rate of infected humans

$$R_0 = \frac{ma^2 p^n b}{-\ln(p)} \frac{c}{r}$$

R_0 is the total number of infectious vector bites that arise from one infected person introduced into a susceptible human population

Vector population modification strategies target vector-arbovirus interactions



$R_0 = \frac{ma^2 p^n b}{-\ln(p)} \frac{c}{r}$

m = vector density per person

a = daily probability of human biting

p = daily probability of vector survival

n = duration in days of extrinsic incubation period (EIP)

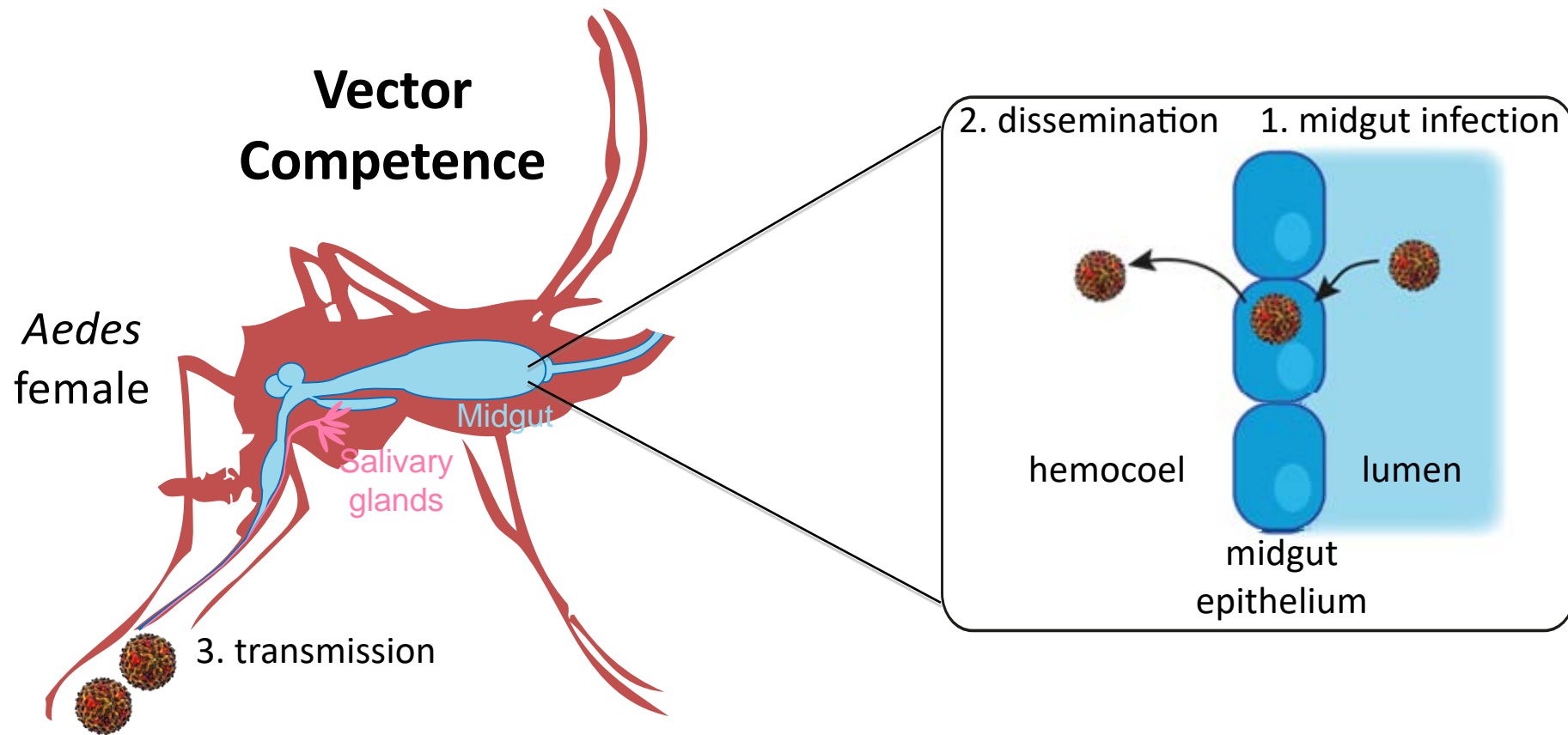
b = vector competence

c = human host competence

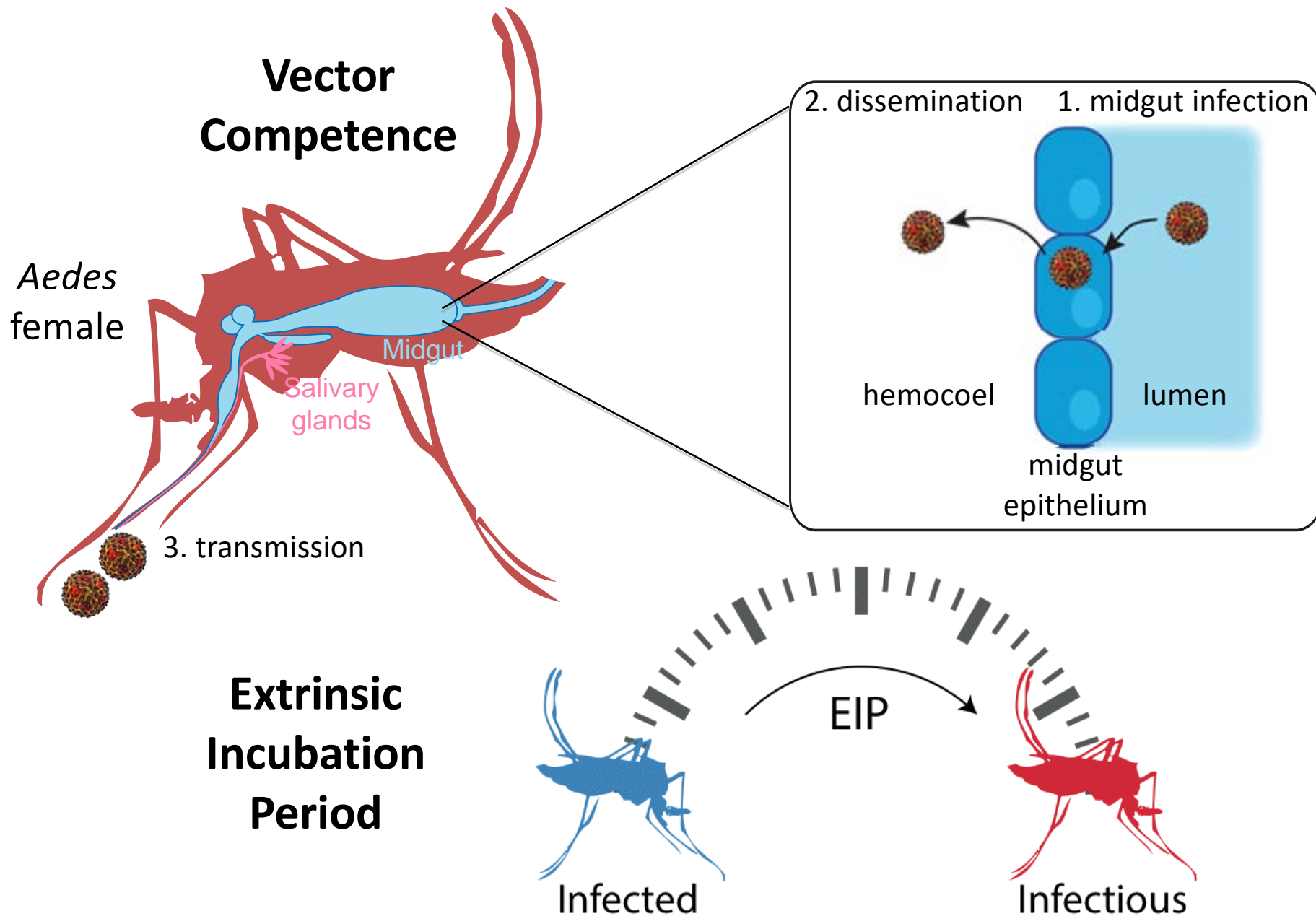
r = daily recovery rate of infected humans

R_0 is the total number of infectious vector bites that arise from one infected person introduced into a susceptible human population

Arbovirus–mosquito interactions

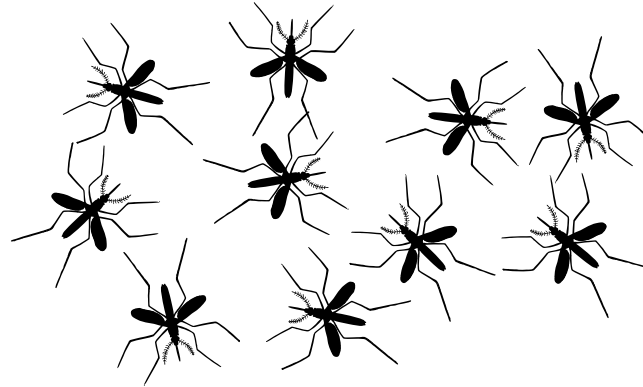


Arbovirus–mosquito interactions



Arbovirus prevention by modification of vector populations

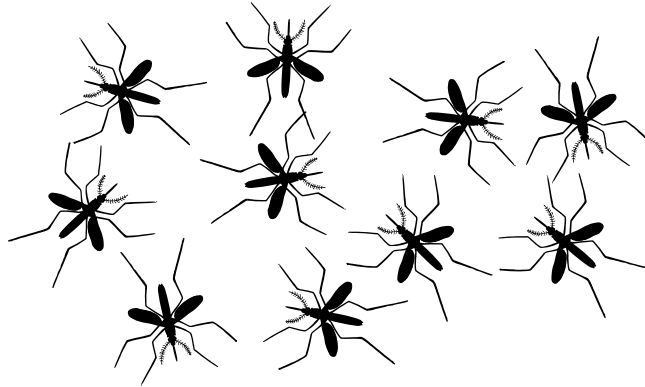
Wild target population



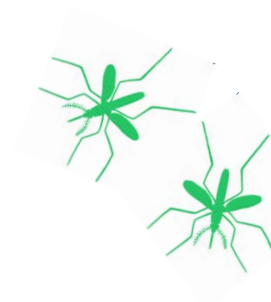
Virus transmission

Arbovirus prevention by modification of vector populations

Wild target population



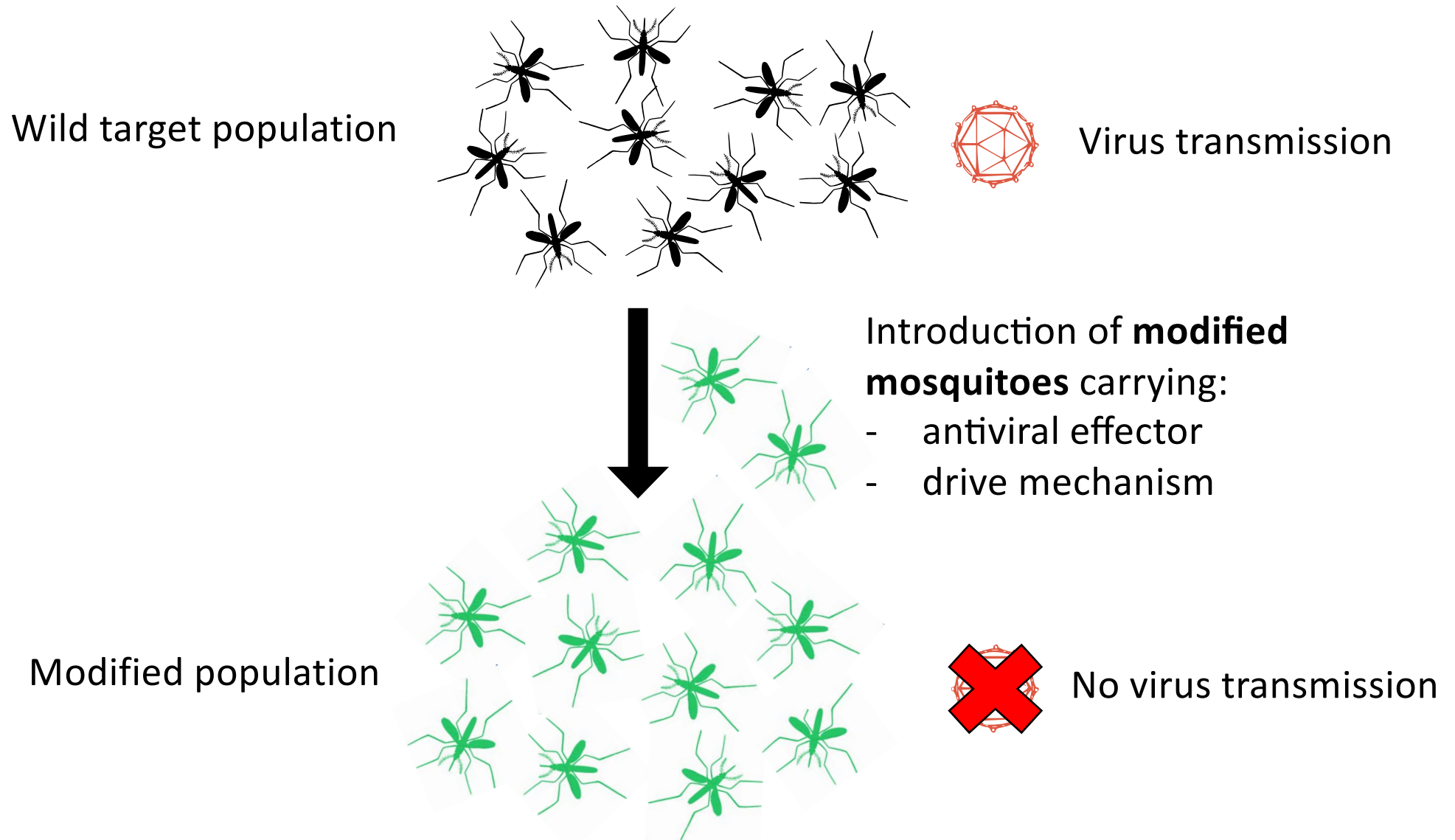
Virus transmission



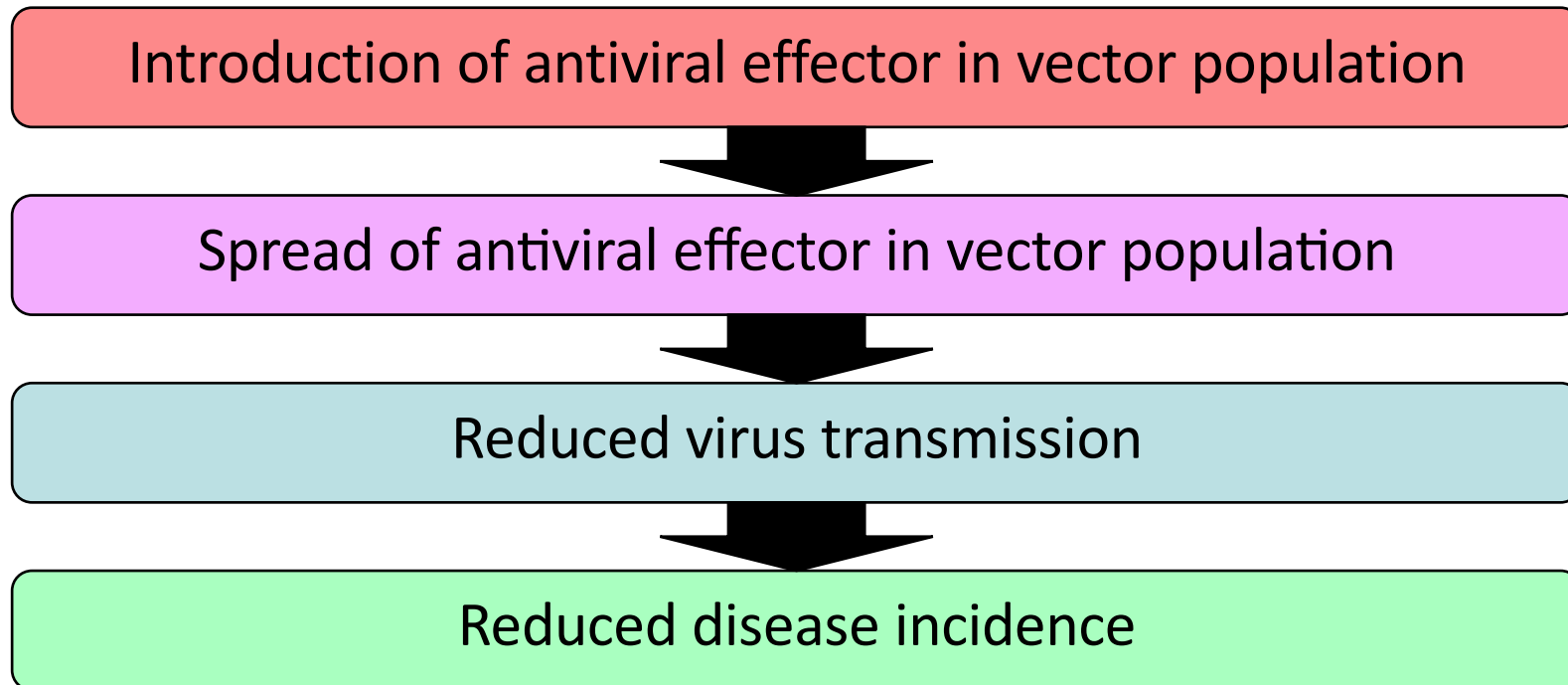
Introduction of **modified mosquitoes** carrying:

- antiviral effector
- drive mechanism

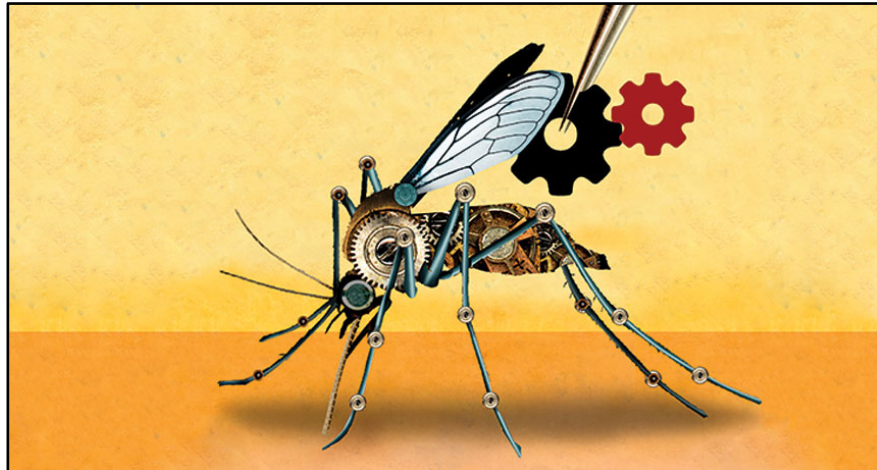
Arbovirus prevention by modification of vector populations



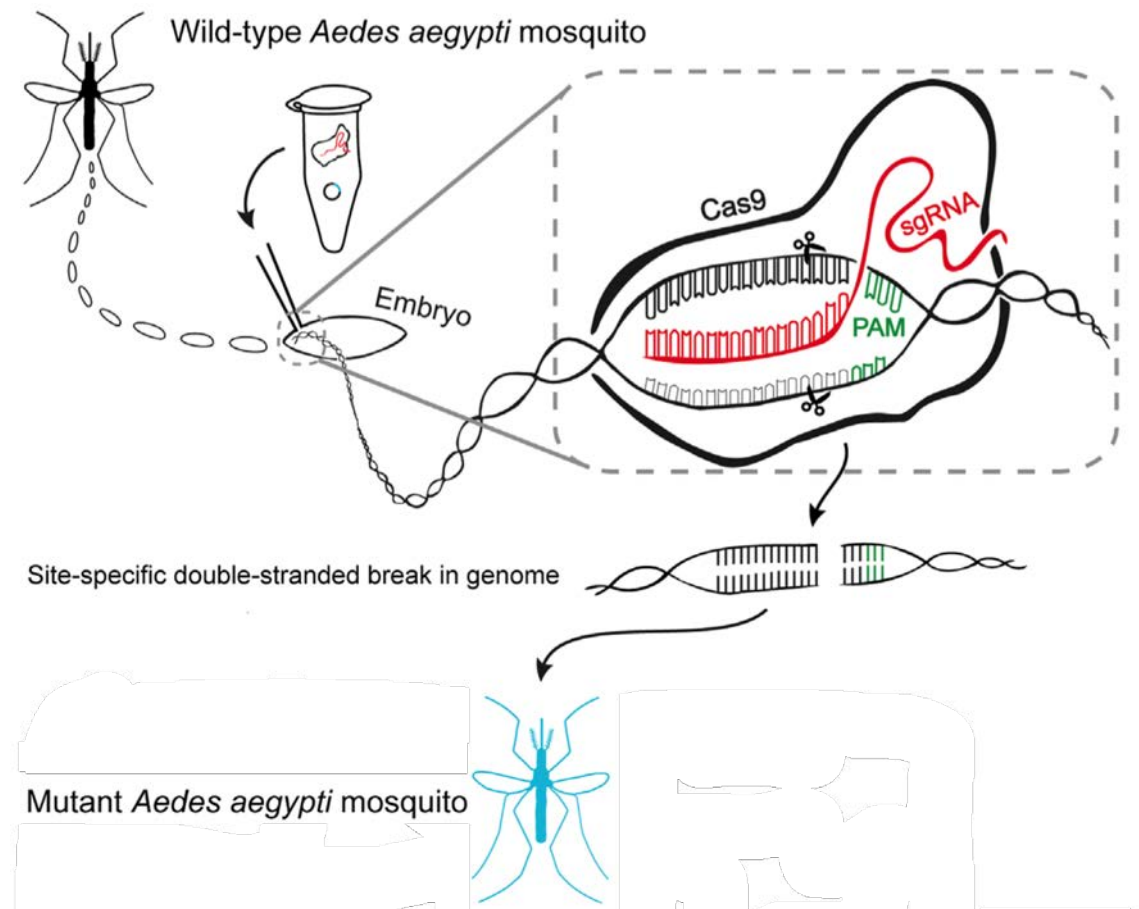
Arbovirus prevention by modification of vector populations



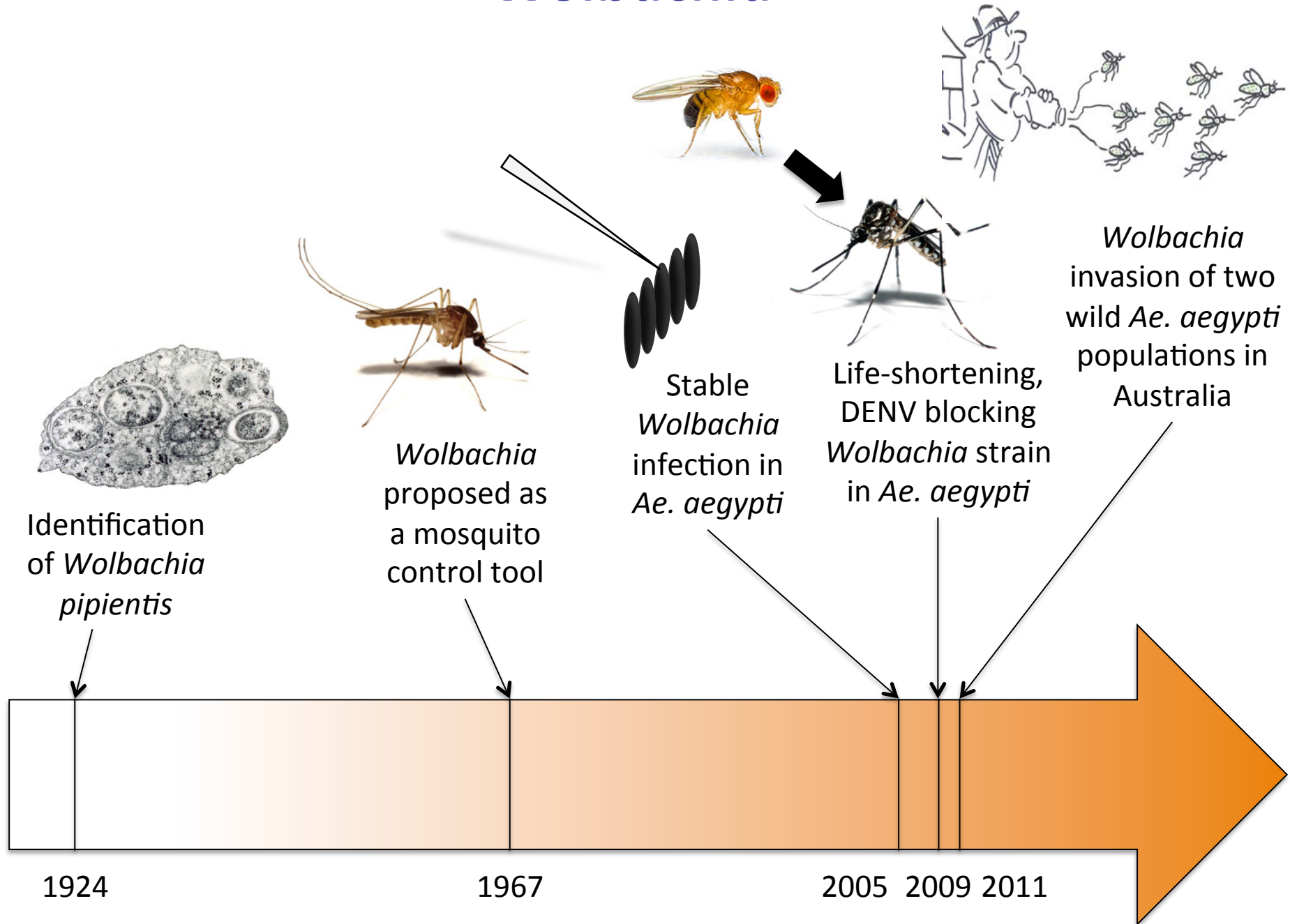
Mosquito genetic engineering



CRISPR/Cas9 gene editing in the mosquito *Aedes aegypti*

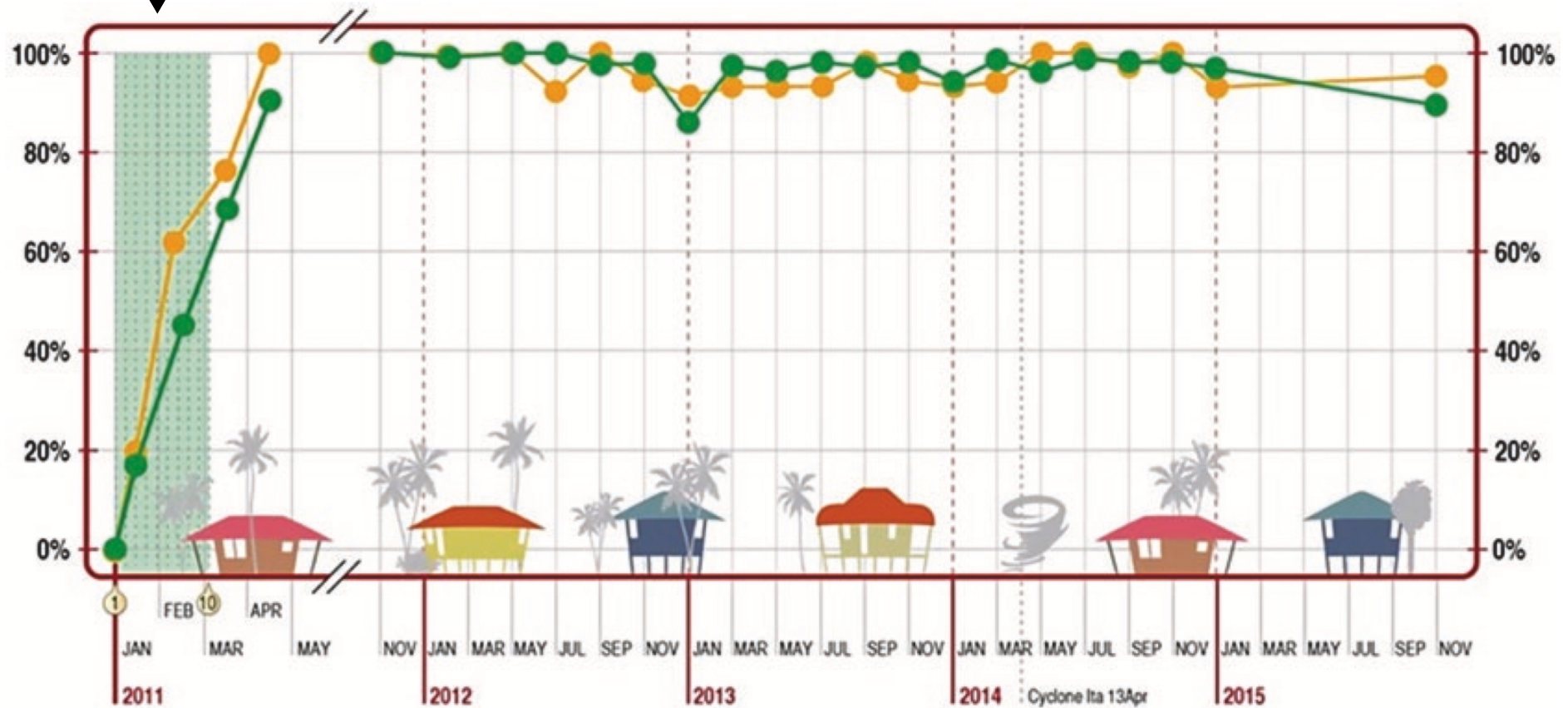


Wolbachia



Wolbachia invades and persists in natural populations of *Aedes aegypti*

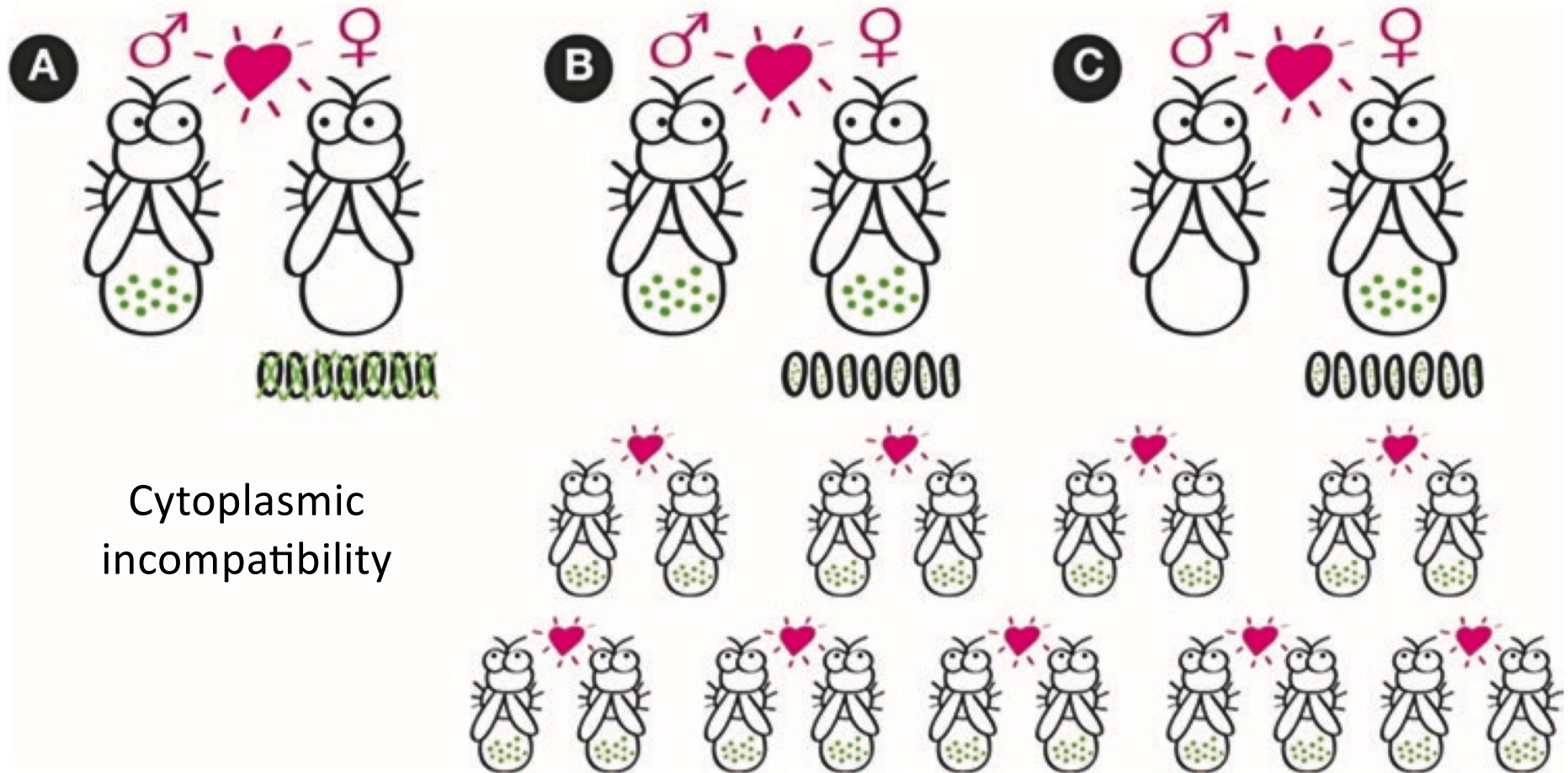
10 weeks of releases



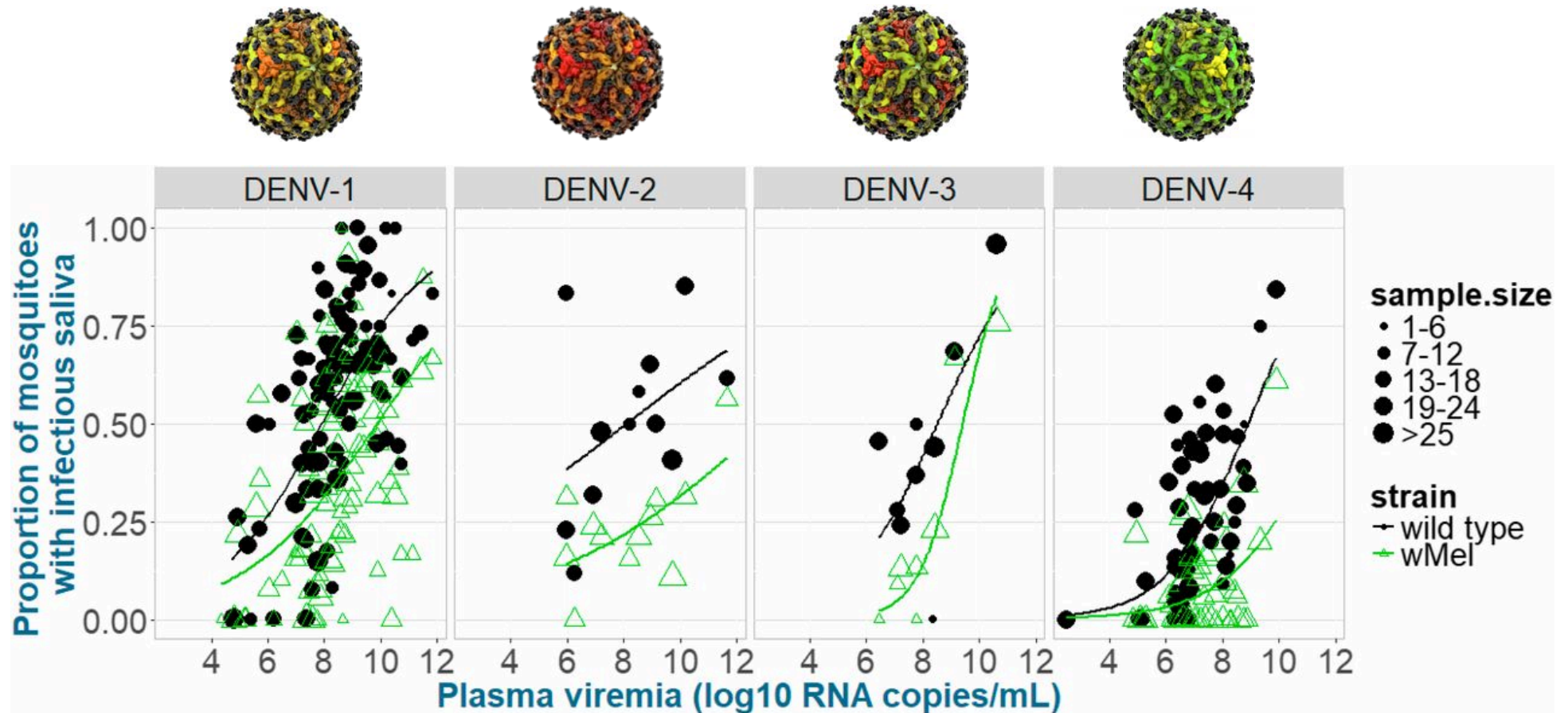
Percentage of mosquitoes with *Wolbachia* in:
— Yorkeys Knob — Gordonvale

— *Wolbachia* mosquitoes released

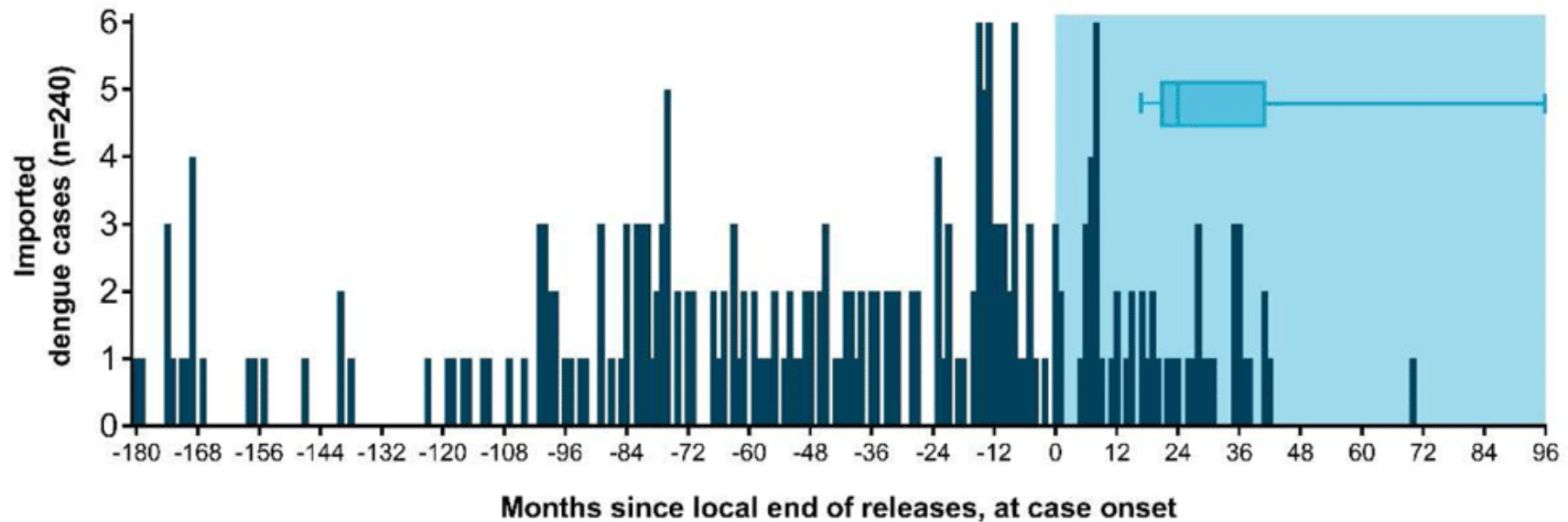
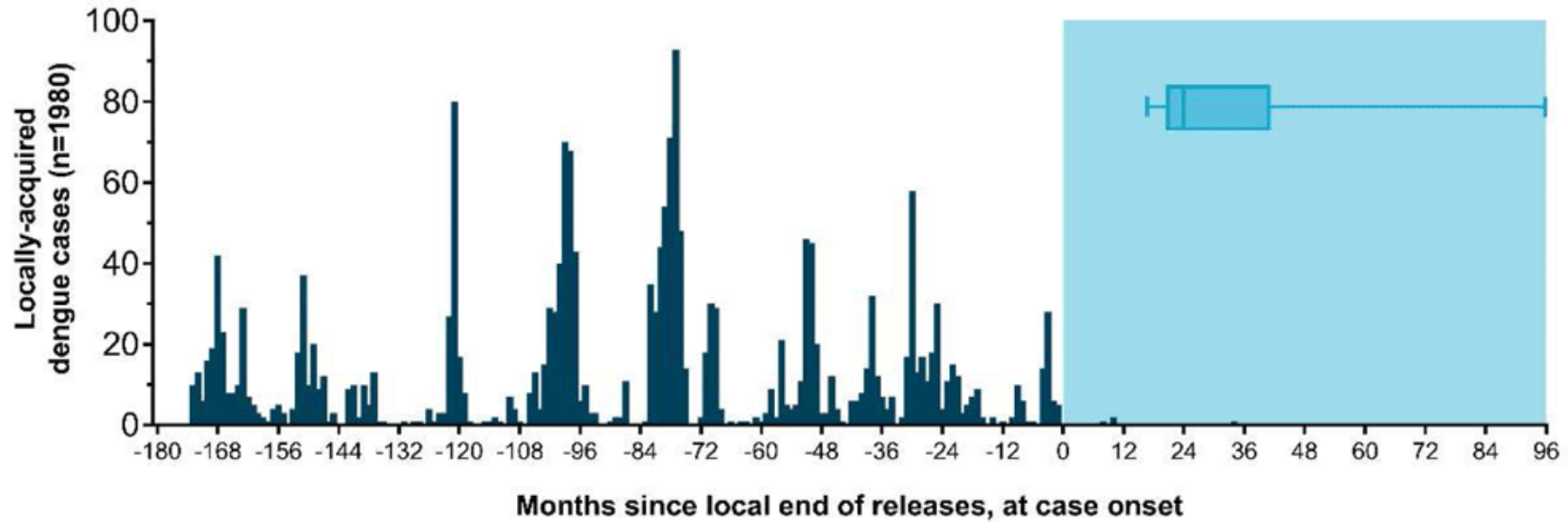
Cytoplasmic incompatibility drives the rapid spread of *Wolbachia* in mosquito populations



Wolbachia reduces DENV transmission by *Aedes aegypti* from naturally-infected people



Dengue case notifications before and after *Wolbachia* releases January 2000 – March 2019



Conclusions

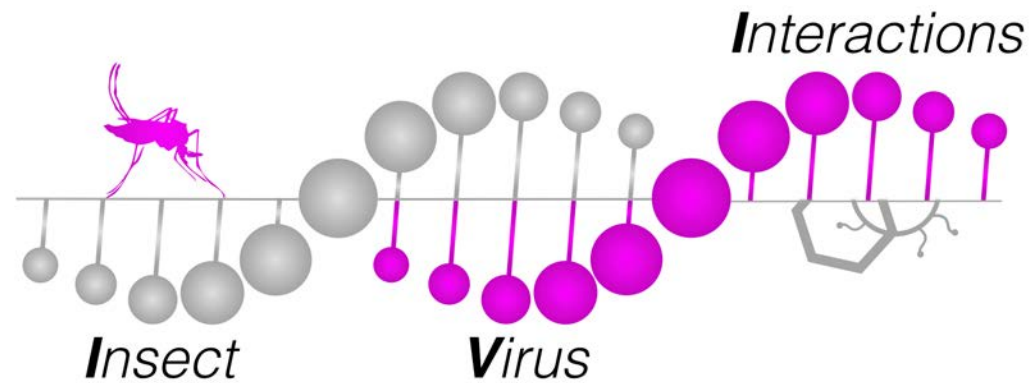
- **Main drivers of arbovirus emergence:**
 - Globalization
 - Urbanization
 - Lack of effective control

- **Other factors:**
 - Lack of political will and resources
 - Decay in public health infrastructure
 - Changing life style
 - Virus evolution

- **Innovative vector control strategies:**
 - Promising results with *Wolbachia*
 - Field trials to measure epidemiological impact



Thank you for your attention



Website: research.pasteur.fr/en/team/insect-virus-interactions

Contact: louis.lambrechts@pasteur.fr

Twitter: @LambrechtsLab

