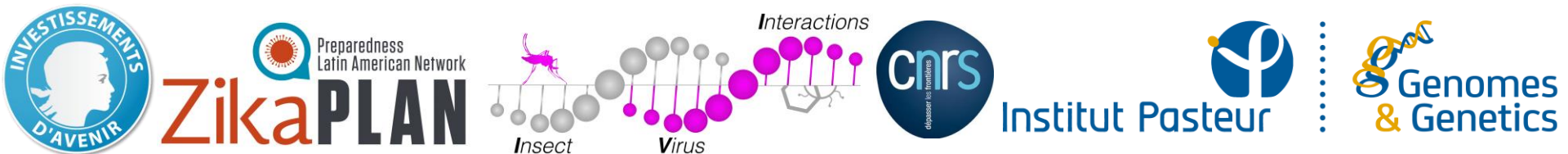


22 June 2018 – *Les Pensières* Center for Global Health

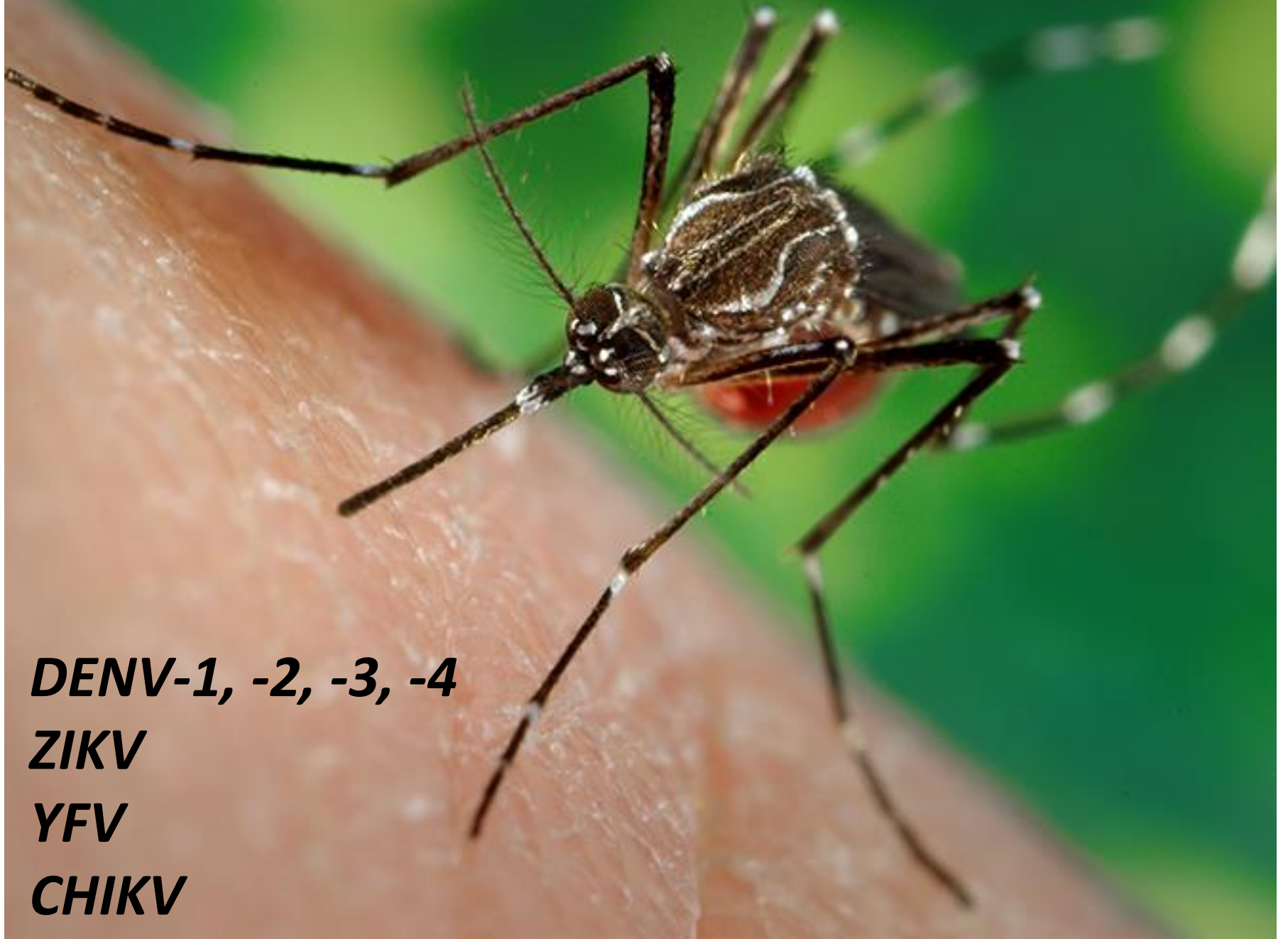
# Overview of *Aedes aegypti* biology and interactions with dengue and Zika viruses

Louis Lambrechts

*Insect-Virus Interactions, Institut Pasteur-CNRS UMR 2000, Paris, France*



***Aedes aegypti*, a major arbovirus vector**



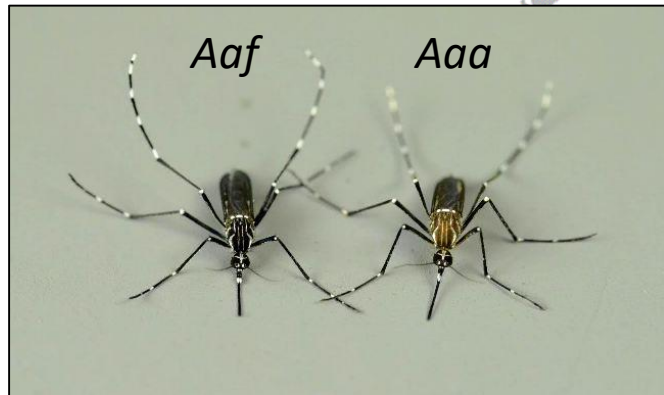
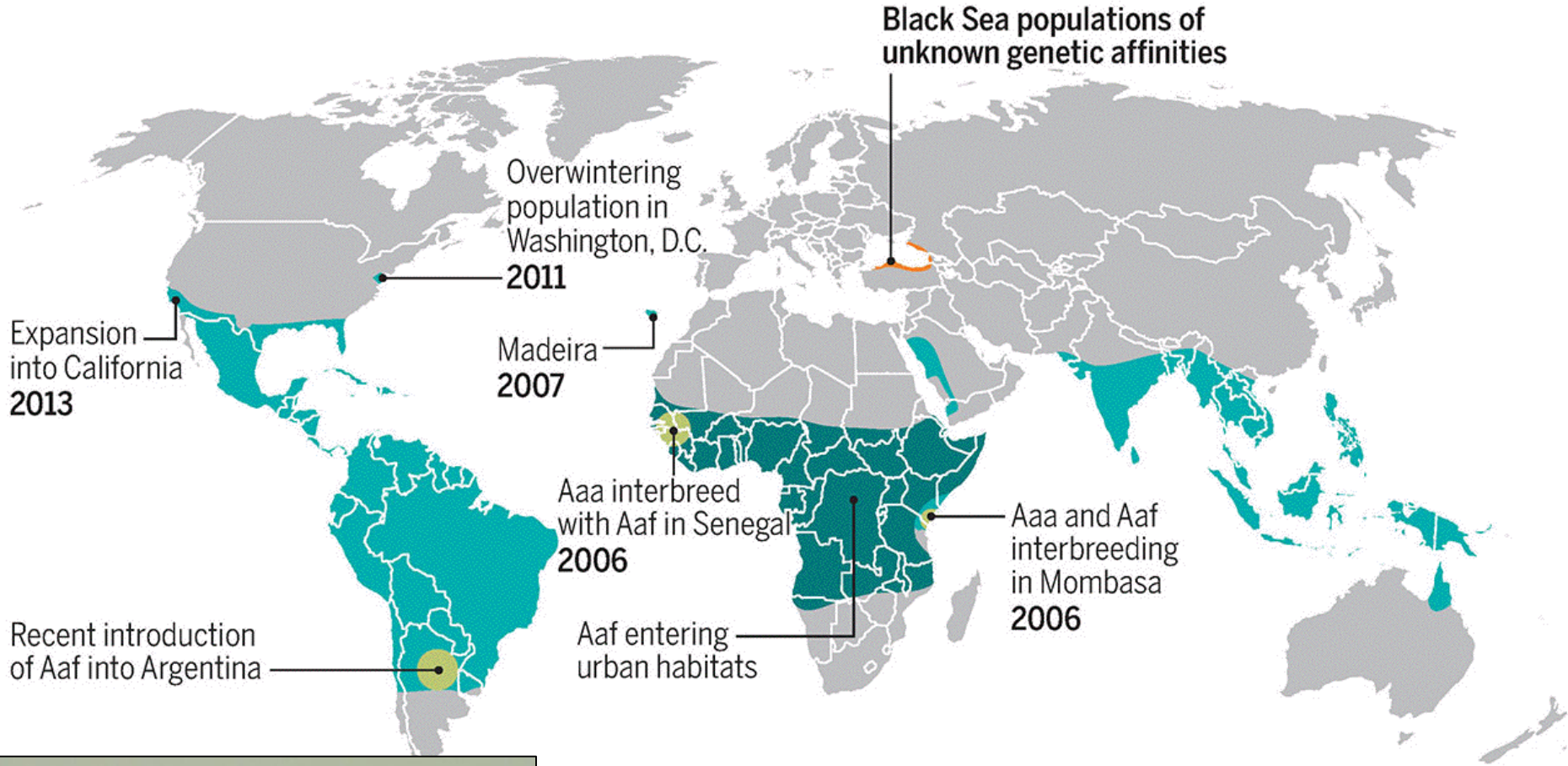
***DENV-1, -2, -3, -4***

***ZIKV***

***YFV***

***CHIKV***

# Worldwide distribution of *Aedes aegypti*



● *Aedes aegypti formosus* (Aaf) ● *Aedes aegypti aegypti* (Aaa)



**What does make  
*Aedes aegypti*  
such a good  
arbovirus vector?**

# Vector-borne pathogen transmission

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

$R_0$  (basic reproduction number) = total number of infectious vector bites that arise from 1 infected person introduced into a susceptible human population

# Vector-borne pathogen transmission

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

$c$  = human host competence

$r$  = daily recovery rate of infected humans

$R_0$  (basic reproduction number) = total number of infectious vector bites that arise from 1 infected person introduced into a susceptible human population

# Vector-borne pathogen transmission

The diagram shows the equation for the basic reproduction number  $R_0$  for vector-borne pathogen transmission. The equation is  $R_0 = \frac{ma^2p^n b}{-\ln(p)} \frac{c}{r}$ . The terms are defined as follows:

- $m$  = vector density per person
- $a$  = daily probability of human biting
- $p$  = daily probability of vector survival
- $n$  = duration in days of extrinsic incubation period
- $b$  = vector competence
- $c$  = human host competence
- $r$  = daily recovery rate of infected humans

The equation is annotated with lines connecting the variables to their definitions. The terms  $ma^2p^n b$  and  $c$  are grouped under brackets labeled "VECTOR" and "HUMAN" respectively. The terms  $-\ln(p)$  and  $r$  are also grouped under brackets, with  $r$  being labeled as the daily recovery rate of infected humans.

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

$R_0$  (basic reproduction number) = total number of infectious vector bites that arise from 1 infected person introduced into a susceptible human population

# Vector-borne pathogen transmission

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

$m$  = vector density per person

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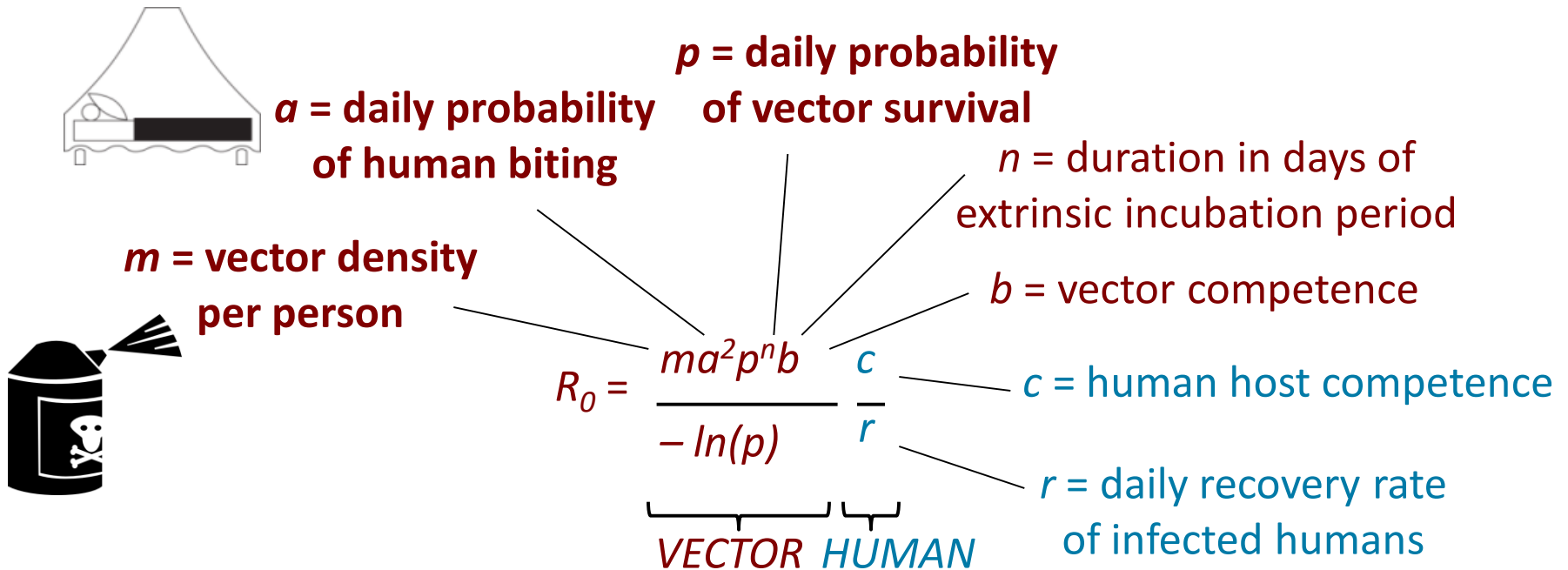
**VECTOR** **HUMAN**



$R_0$  (basic reproduction number) = total number of infectious vector bites that arise from 1 infected person introduced into a susceptible human population



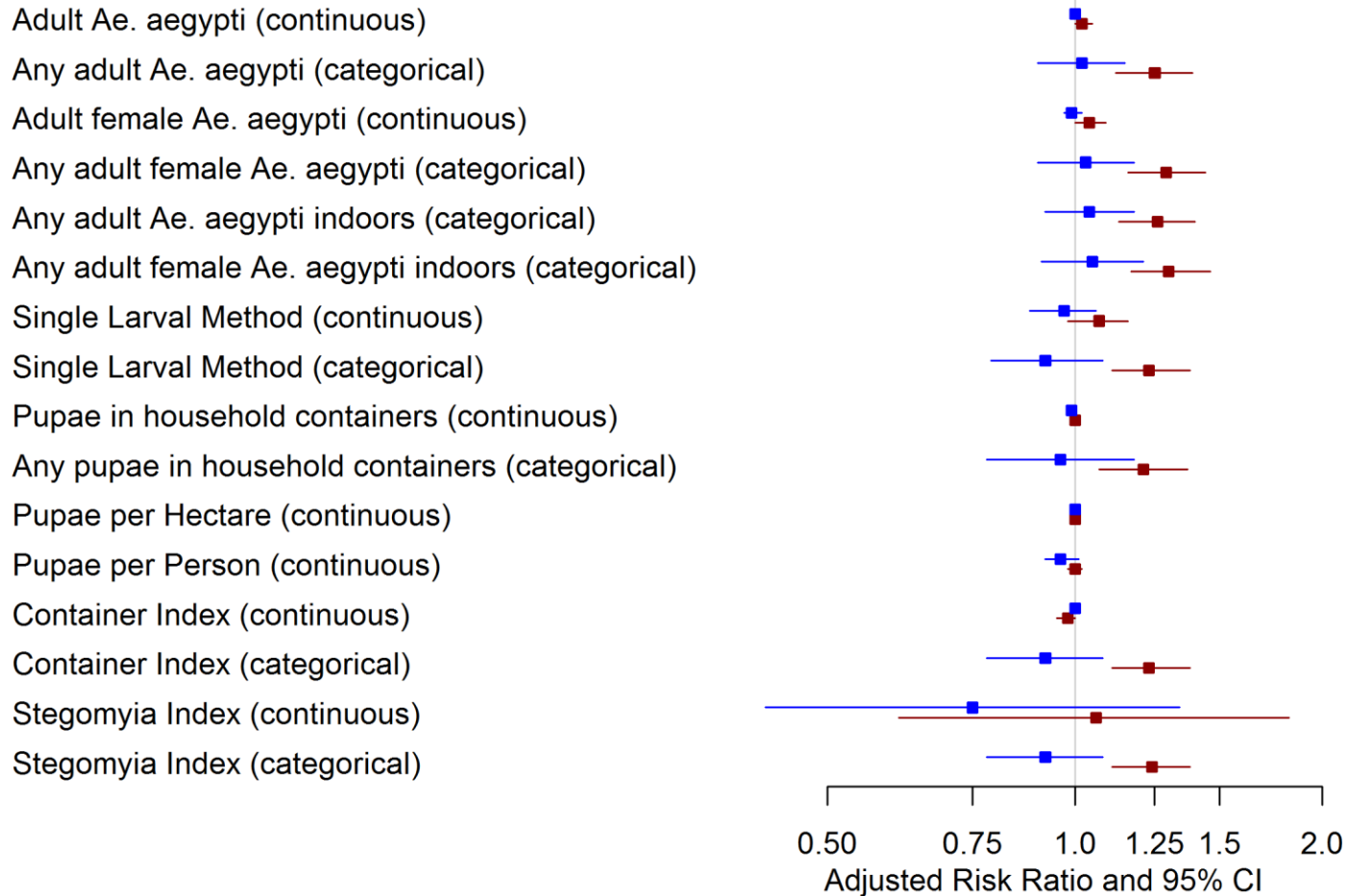
# Vector-borne pathogen transmission



$R_0$  (basic reproduction number) = total number of infectious vector bites that arise from 1 infected person introduced into a susceptible human population

# *Aedes aegypti* abundance and DENV infection risk

■ Cross-sectional ■ Longitudinal



Household-level indicators of *Ae. aegypti* abundance are associated with DENV seroconversion when they are calculated from longitudinal entomological data

# Impact of mosquito blood feeding behavior on vector-borne pathogen transmission

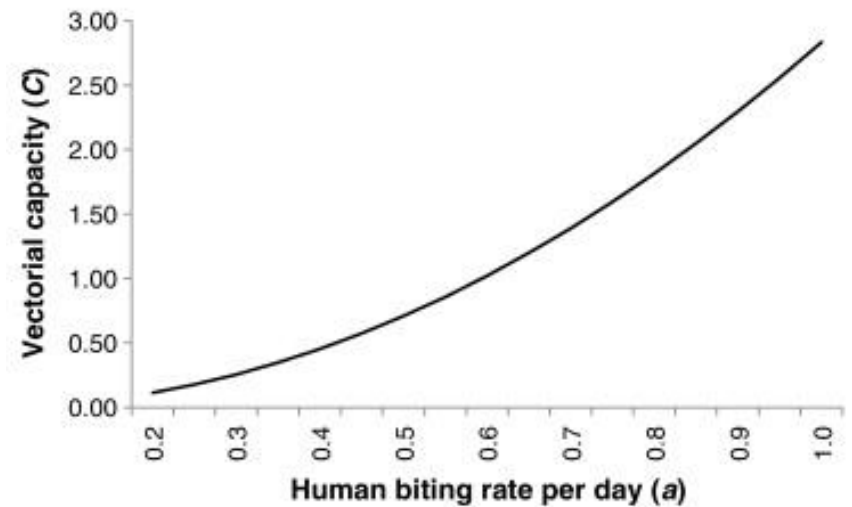
## Most adult female anautogenous mosquitoes

- One vertebrate blood meal per egg-laying cycle
- Consumption of plant carbohydrates



## *Aedes aegypti*

- >90% blood meals taken on humans
- Multiple blood feeding
- 0.63 to 0.76 blood meals per day
- No sugar feeding when human blood available



TRENDS in Parasitology

# Vector-borne pathogen transmission

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

$m$  = vector density per person

$a$  = daily probability of human biting

$p$  = daily probability of vector survival

$n$  = duration in days of extrinsic incubation period

$b$  = vector competence

$c$  = human host competence

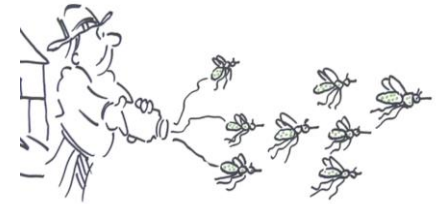
$r$  = daily recovery rate of infected humans

**VECTOR** **HUMAN**

The diagram shows the equation  $R_0 = \frac{ma^2p^n b}{-\ln(p)} \frac{c}{r}$ . Brackets under the numerator group  $ma^2p^n b$  as 'VECTOR' and  $\frac{c}{r}$  as 'HUMAN'. Lines connect each parameter to its definition:  $m$  (vector density per person),  $a$  (daily probability of human biting),  $p$  (daily probability of vector survival),  $n$  (duration in days of extrinsic incubation period),  $b$  (vector competence),  $c$  (human host competence), and  $r$  (daily recovery rate of infected humans).

$R_0$  (basic reproduction number) = total number of infectious vector bites that arise from 1 infected person introduced into a susceptible human population

# Vector-borne pathogen transmission



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

$a$  = daily probability of human biting  
 $p$  = daily probability of vector survival  
 $n$  = duration in days of extrinsic incubation period  
 $b$  = vector competence  
 $c$  = human host competence  
 $r$  = daily recovery rate of infected humans  
 $m$  = vector density per person

$R_0$  (basic reproduction number) = total number of infectious vector bites that arise from 1 infected person introduced into a susceptible human population

# Talk outline



## **Genetic determinants of vector-virus interactions**

- G x G interactions
- *Ae. aegypti* vs. *Ae. albopictus*

## **Non-genetic determinants of vector-virus interactions**

- Temperature
- Larval microbiota
- Vertebrate host factors

# Talk outline



## **Genetic determinants of vector-virus interactions**

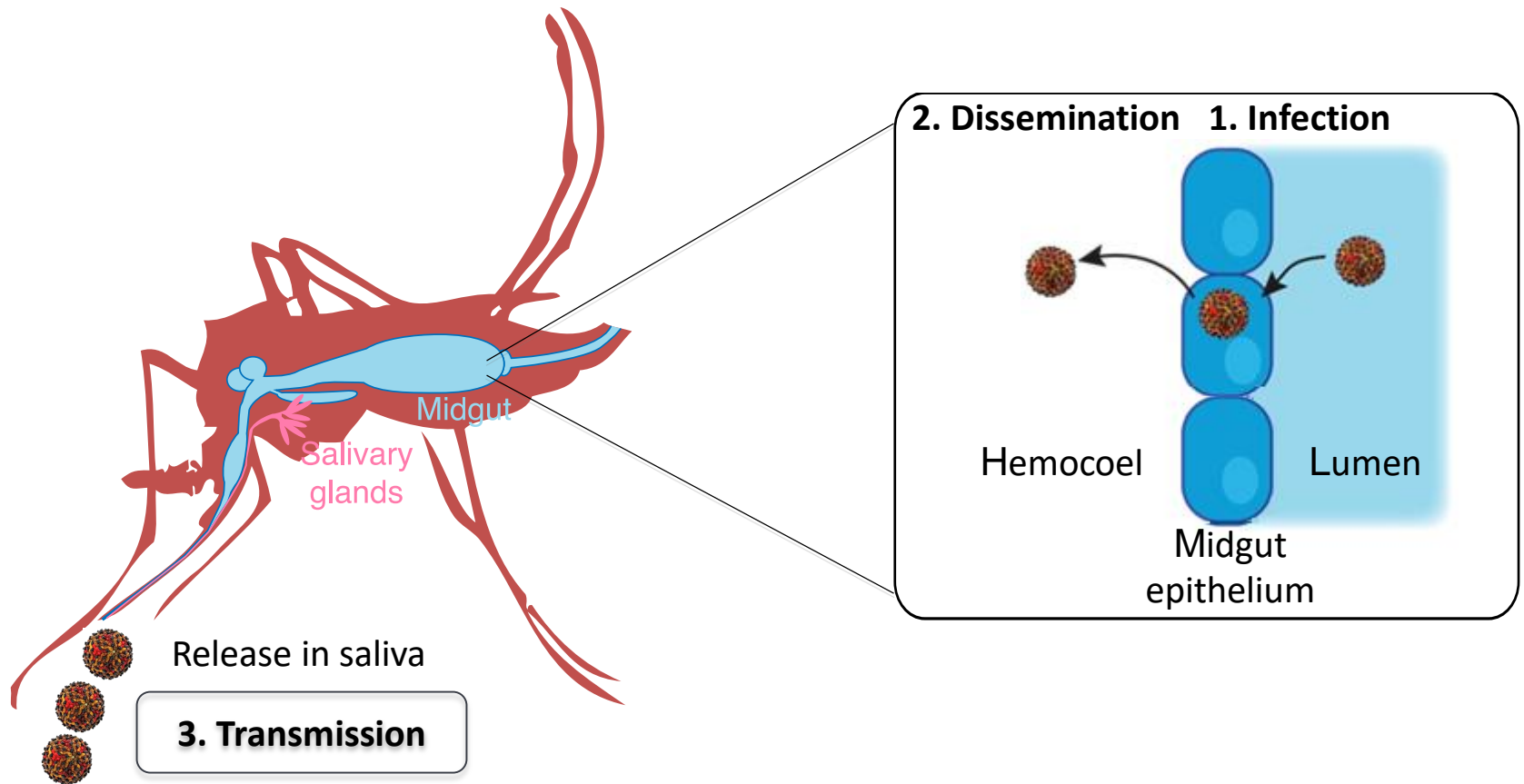
- G x G interactions
- *Ae. aegypti* vs. *Ae. albopictus*

## **Non-genetic determinants of vector-virus interactions**

- Temperature
- Larval microbiota
- Vertebrate host factors

# Vector competence

Intrinsic ability to become infected and subsequently transmit a pathogen



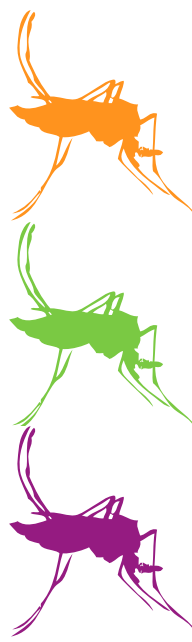


# Genotype-by-genotype (G x G) interactions

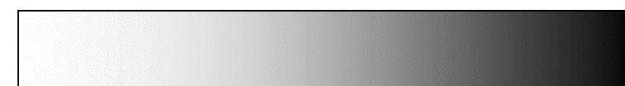
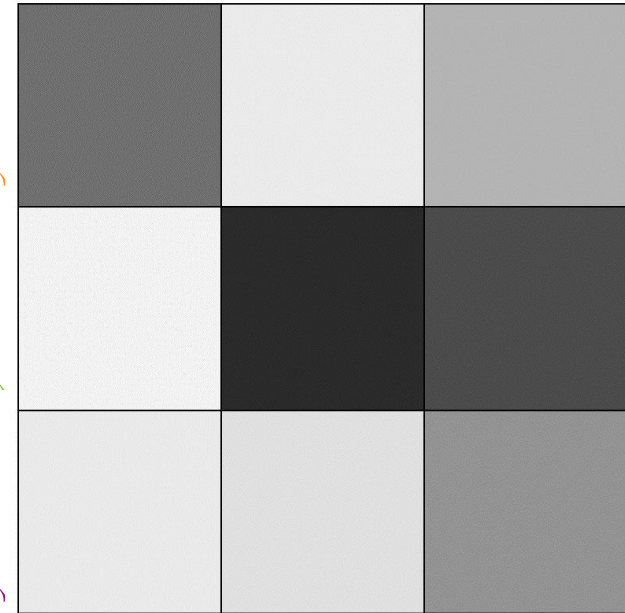
Ratchaburi, Thailand



*Ae. aegypti*  
genotypes

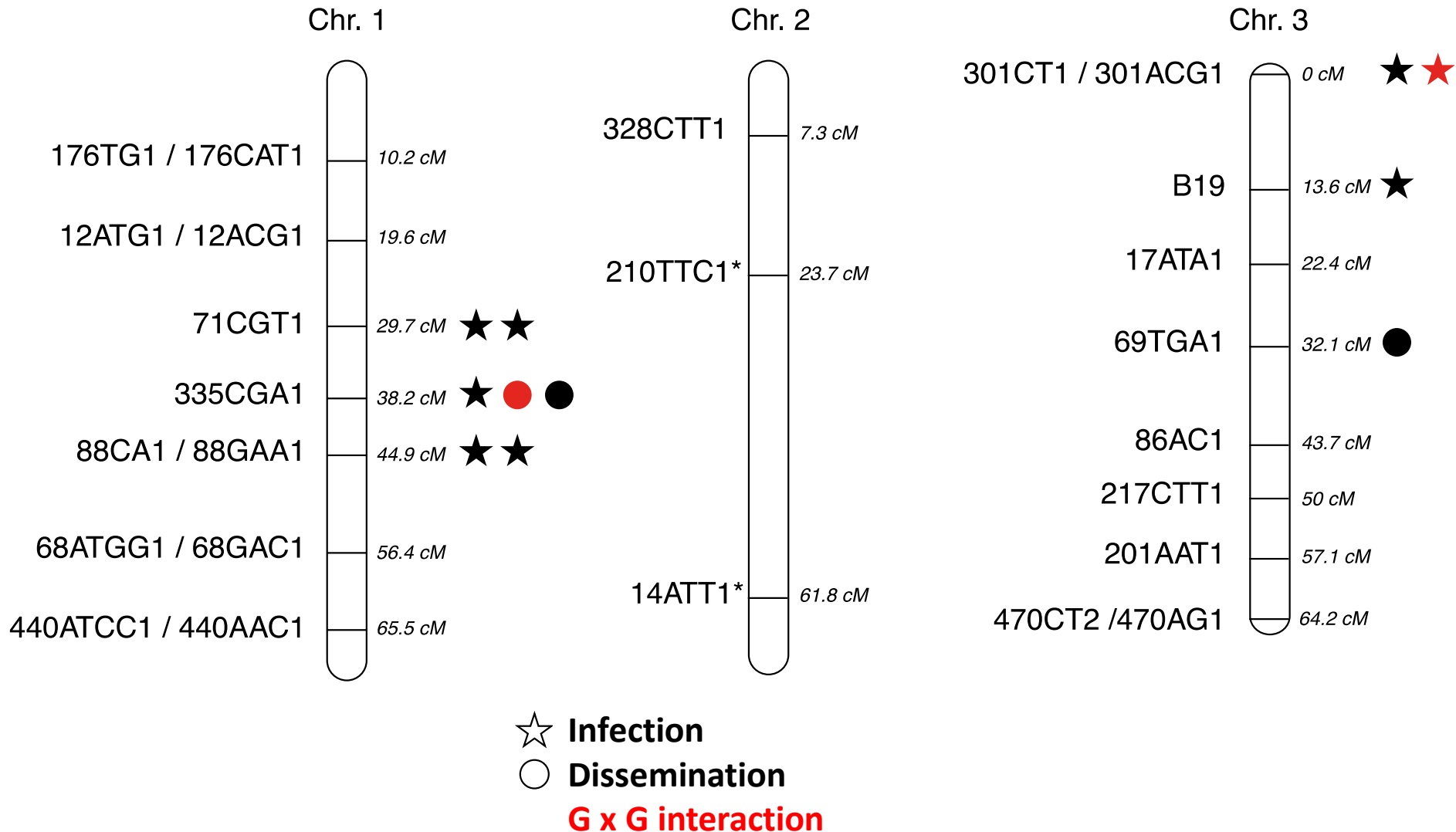


DENV-1 isolates



LOW                      Vector competence                      HIGH

# Genetic mapping of G x G interactions between *Ae. aegypti* and DENV

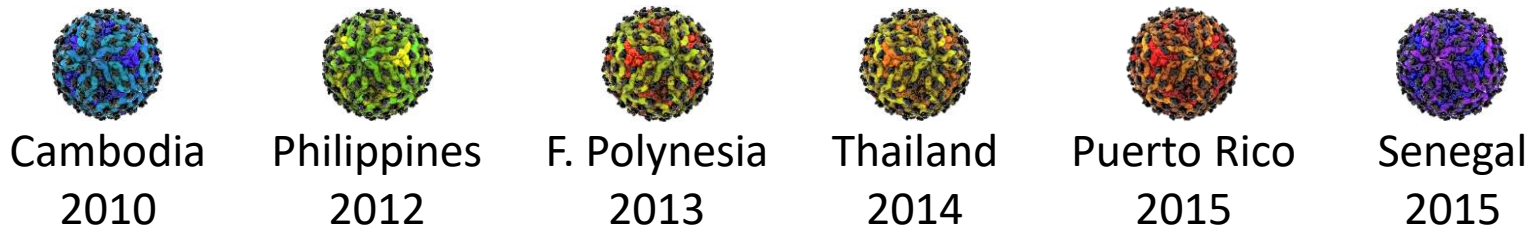


# Worldwide survey of *Ae. aegypti* susceptibility to diverse strains of ZIKV

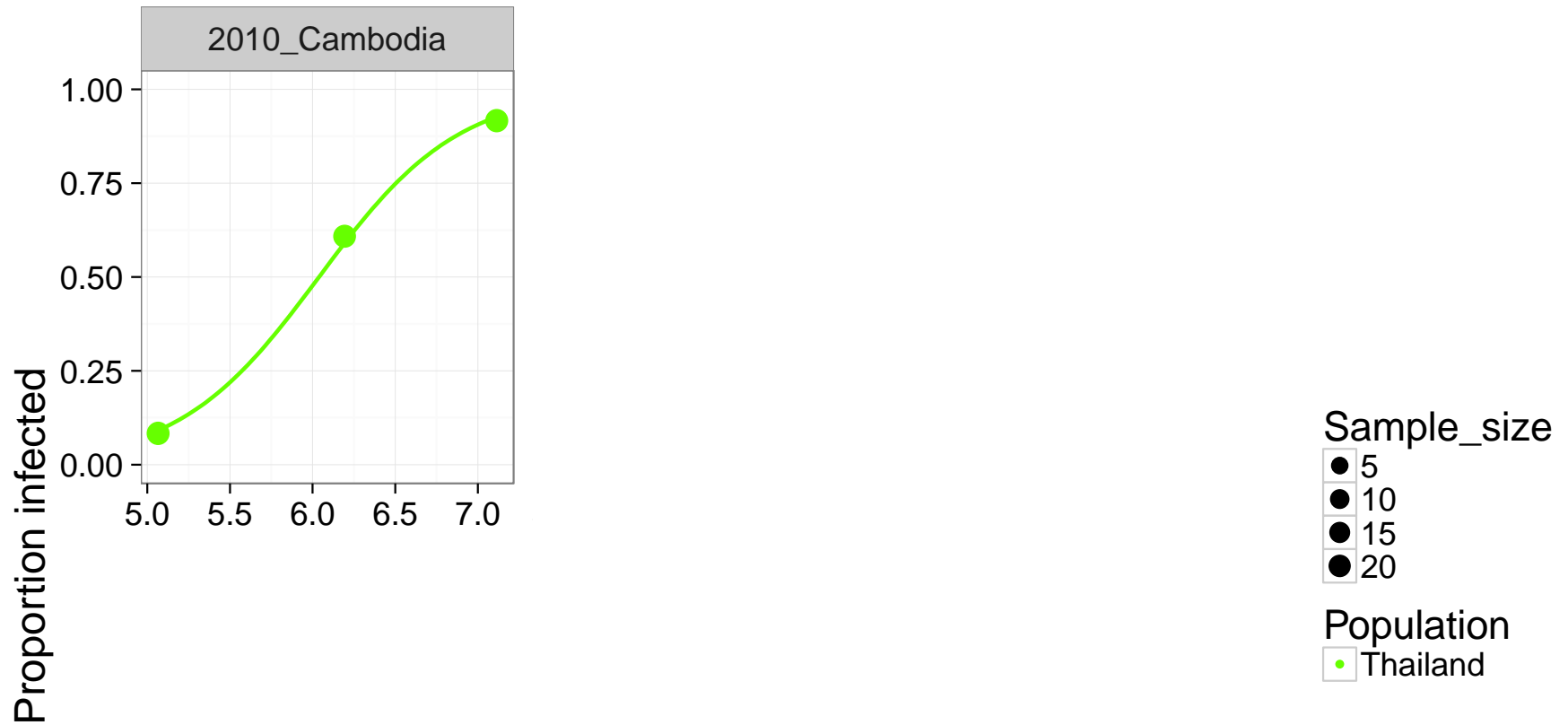
- 8 field-derived *Ae. aegypti* colonies



- 6 low-passage ZIKV strains:

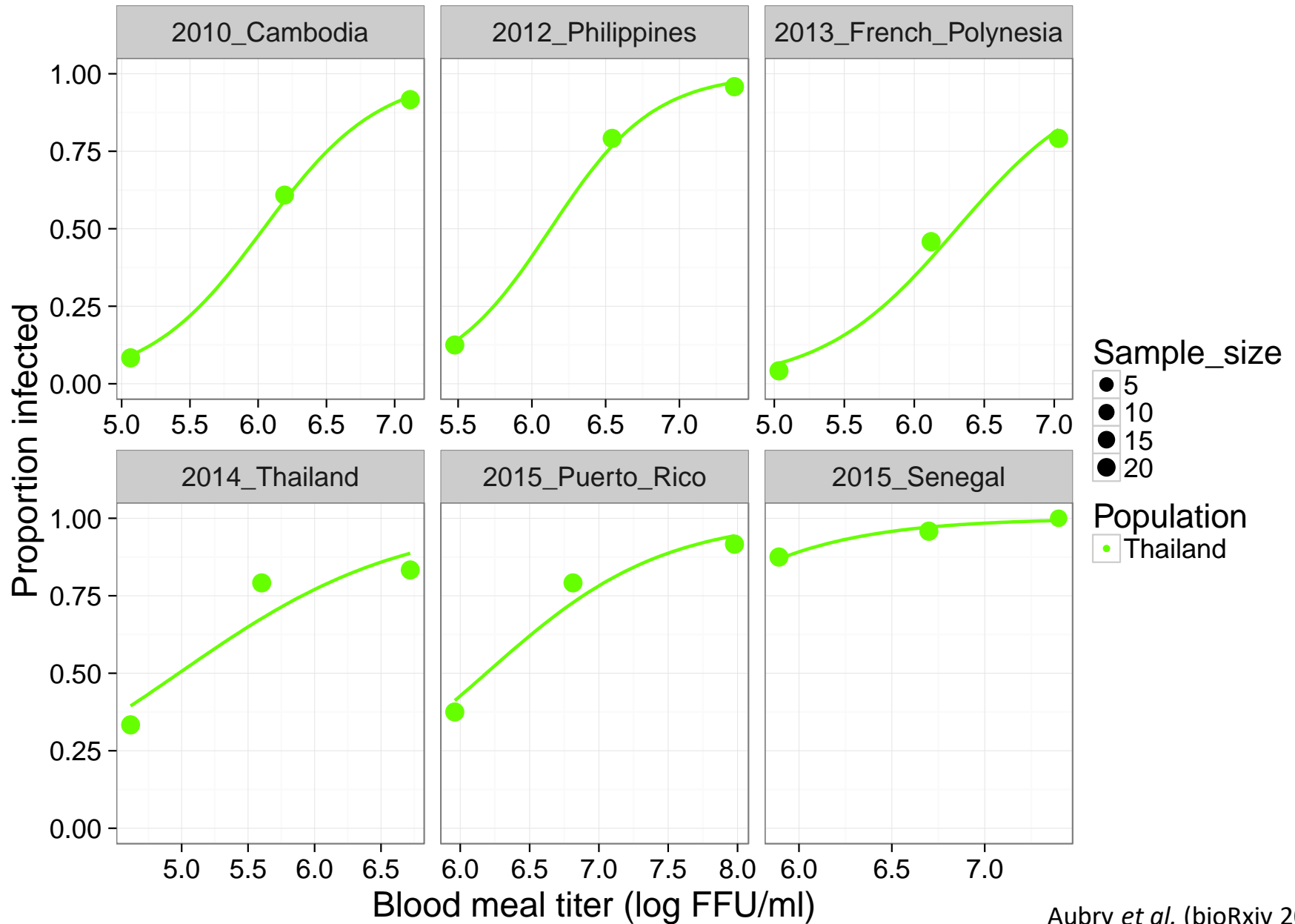


# *Aedes aegypti* susceptibility to ZIKV

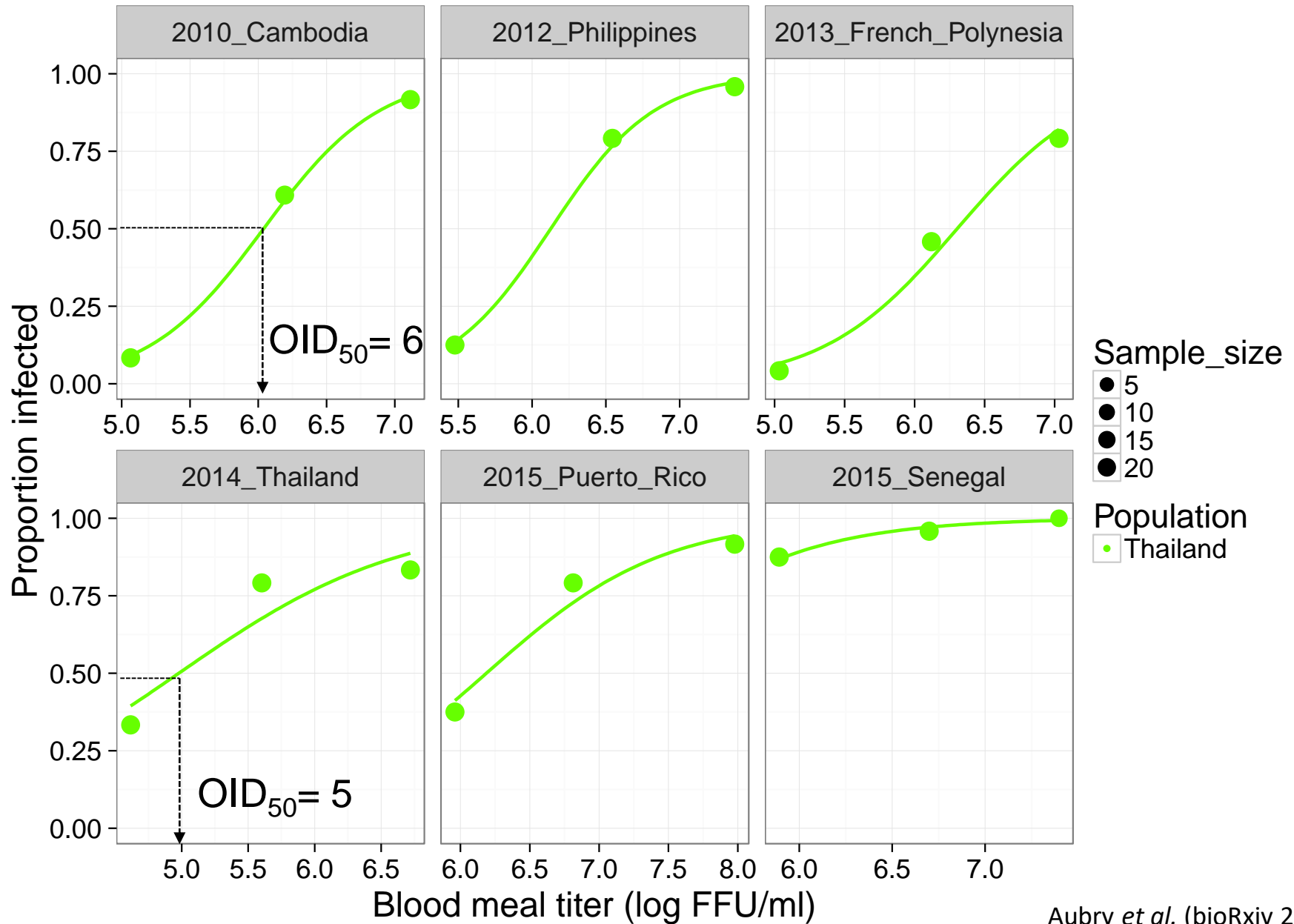


Blood meal titer (log FFU/ml)

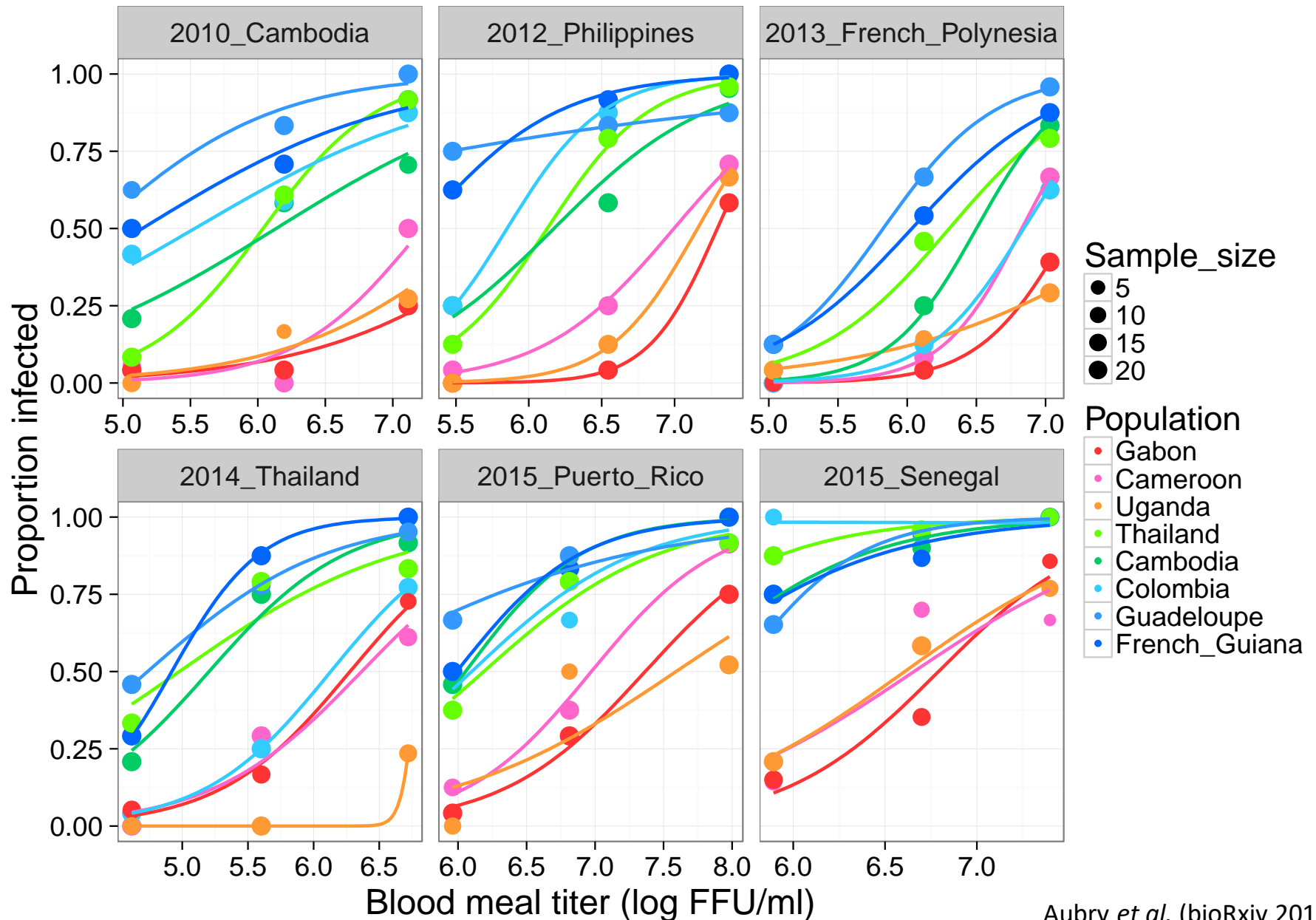
# *Aedes aegypti* susceptibility to ZIKV



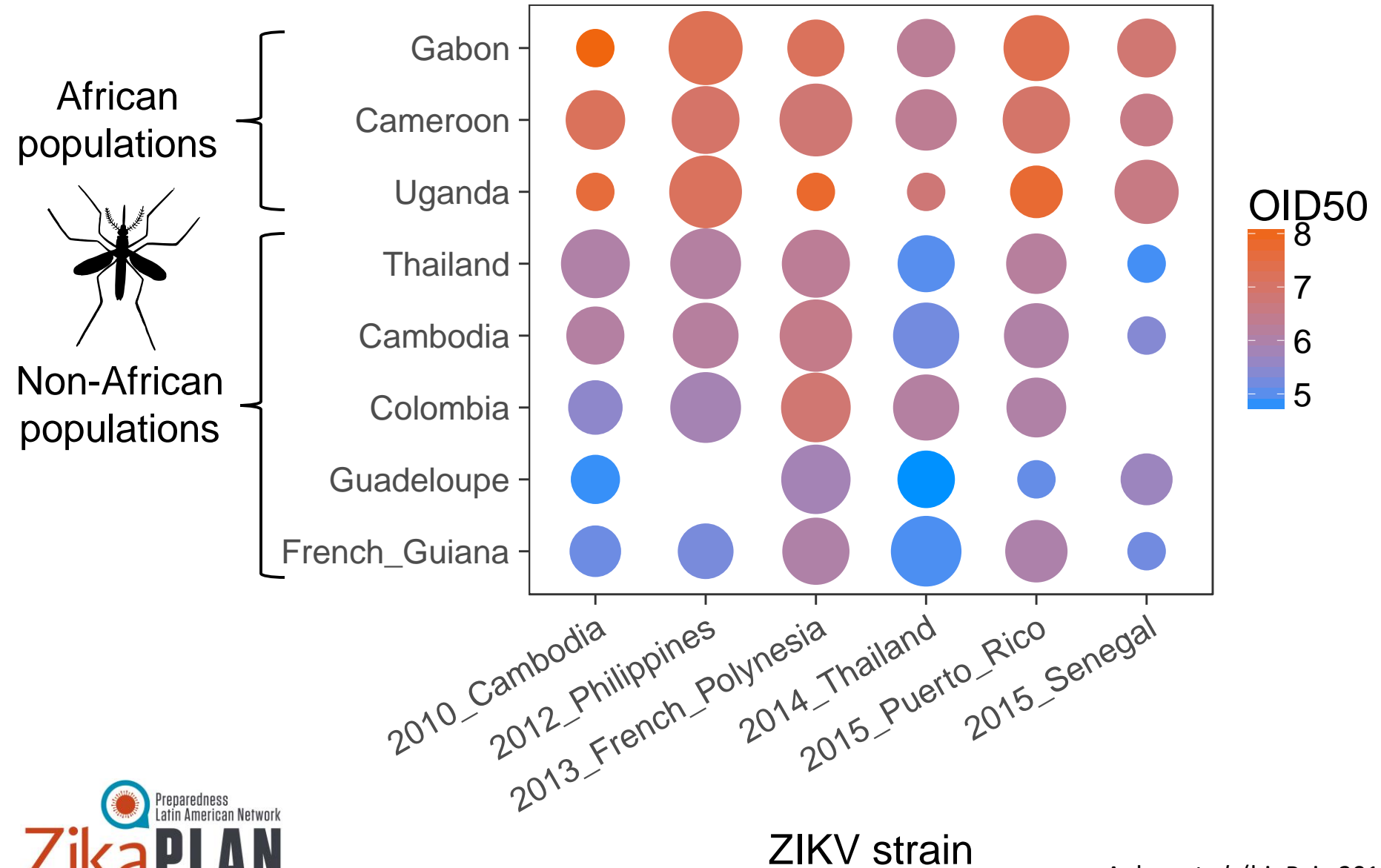
# *Aedes aegypti* susceptibility to ZIKV



# *Aedes aegypti* susceptibility to ZIKV



# African *Ae. aegypti* are less susceptible to ZIKV





# Talk outline



## Genetic determinants of vector-virus interactions

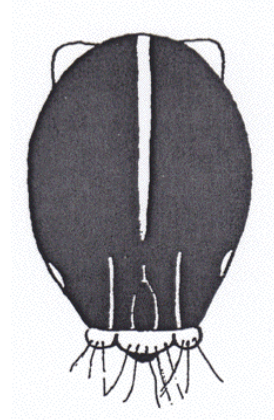
- G x G interactions
- *Ae. aegypti* vs. *Ae. albopictus*

## Non-genetic determinants of vector-virus interactions

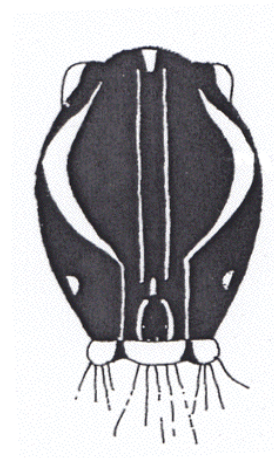
- Temperature
- Larval microbiota
- Vertebrate host factors

# *Aedes aegypti* vs. *Aedes albopictus*

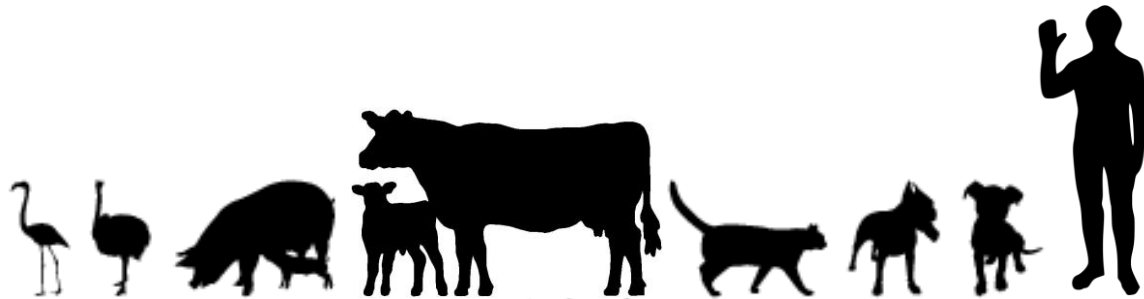
Secondary DENV/ZIKV vector:  
*Aedes albopictus*



Primary DENV/ZIKV vector:  
*Aedes aegypti*



# Feeding behavior of *Ae. aegypti* vs. *Ae. albopictus*



*Aedes albopictus*

- Day biting
- **Opportunistic**



*Aedes aegypti aegypti*

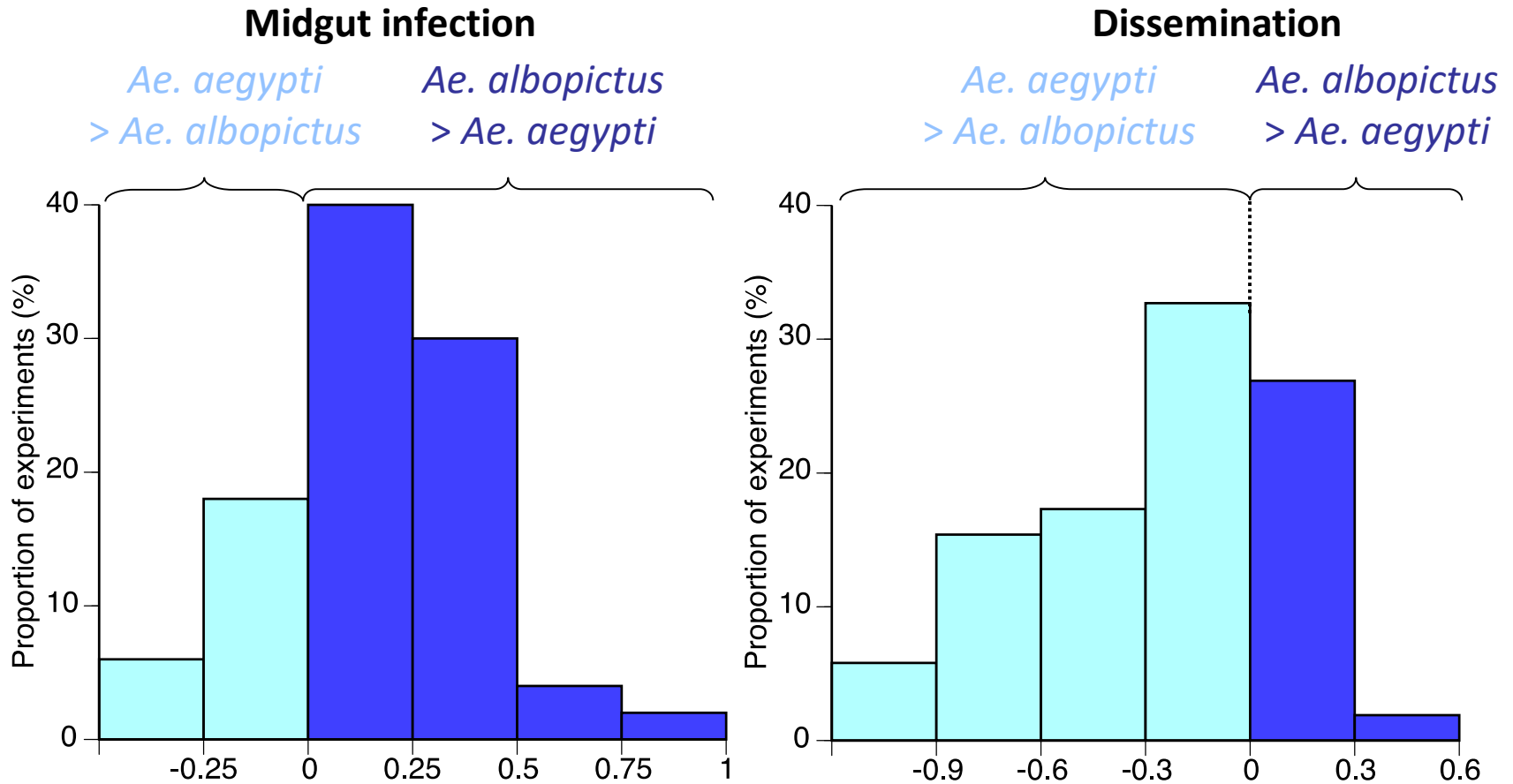
- Day biting
- **Anthropophilic**

However, multiple examples of *Ae. albopictus* preference for humans:

- North Carolina (Richards *et al.* *J Med Entomol* 2006)
- La Réunion Island (Delatte *et al.* *Vector Borne Zoonotic Dis* 2010)
- Southern Thailand (Ponlawat & Harrington *J Med Entomol* 2005)

# DENV competence of *Ae. aegypti* vs. *Ae. albopictus*

Meta-analysis of 91 separate experiments from 14 studies (1971-2009)

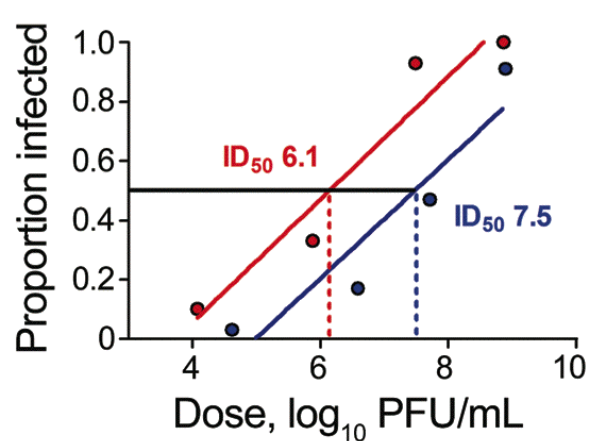


- *Ae. albopictus* 16% more susceptible to midgut infection
- Dissemination 26% less likely than in *Ae. aegypti*

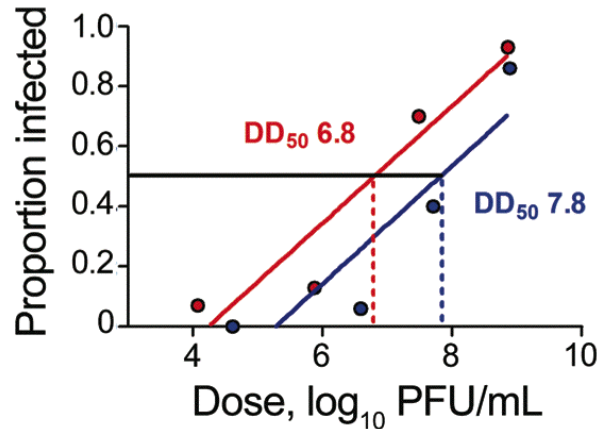
# ZIKV competence of *Ae. aegypti* vs. *Ae. albopictus*

*Ae. albopictus* is more susceptible but less competent than *Ae. aegypti*

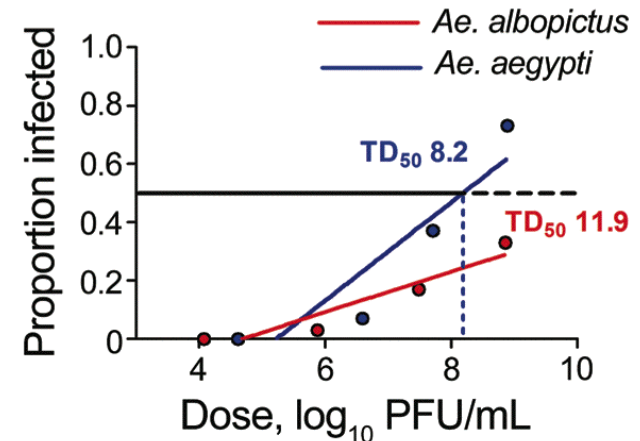
## Infection (body)



## Dissemination (legs)



## Transmission (saliva)



- Data shown for 2016 ZIKV strain from Honduras
- Similar results for 2010 ZIKV strain from Cambodia

# Talk outline



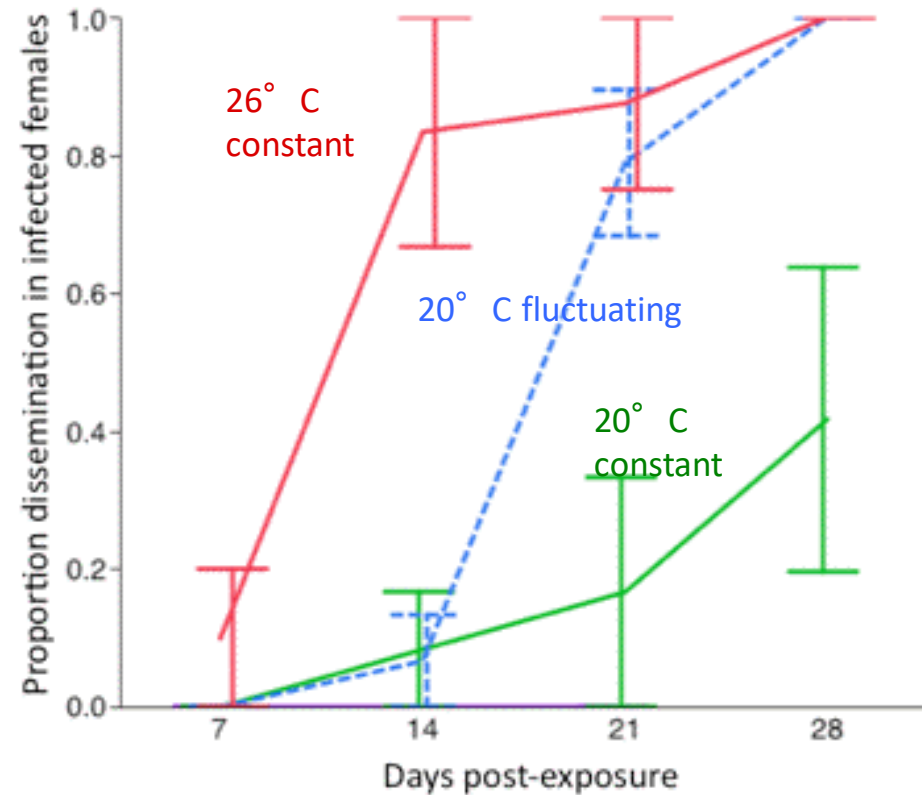
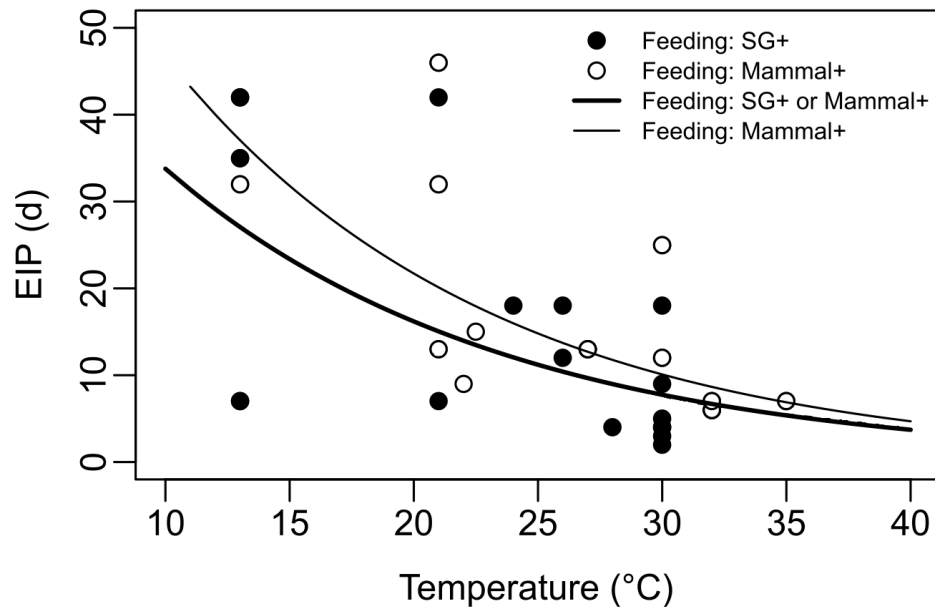
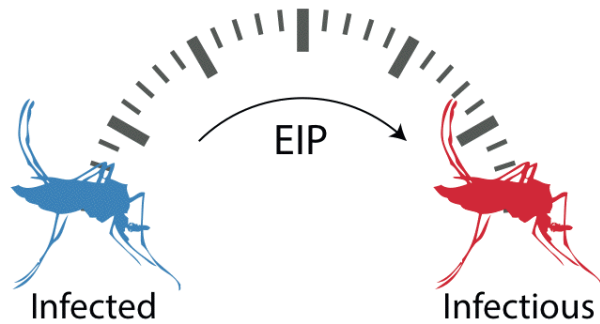
## Genetic determinants of vector-virus interactions

- G x G interactions
- *Ae. aegypti* vs. *Ae. albopictus*

## Non-genetic determinants of vector-virus interactions

- Temperature
- Larval microbiota
- Vertebrate host factors

# Influence of ambient temperature on DENV extrinsic incubation period in *Ae. aegypti*



# Talk outline



## Genetic determinants of vector-virus interactions

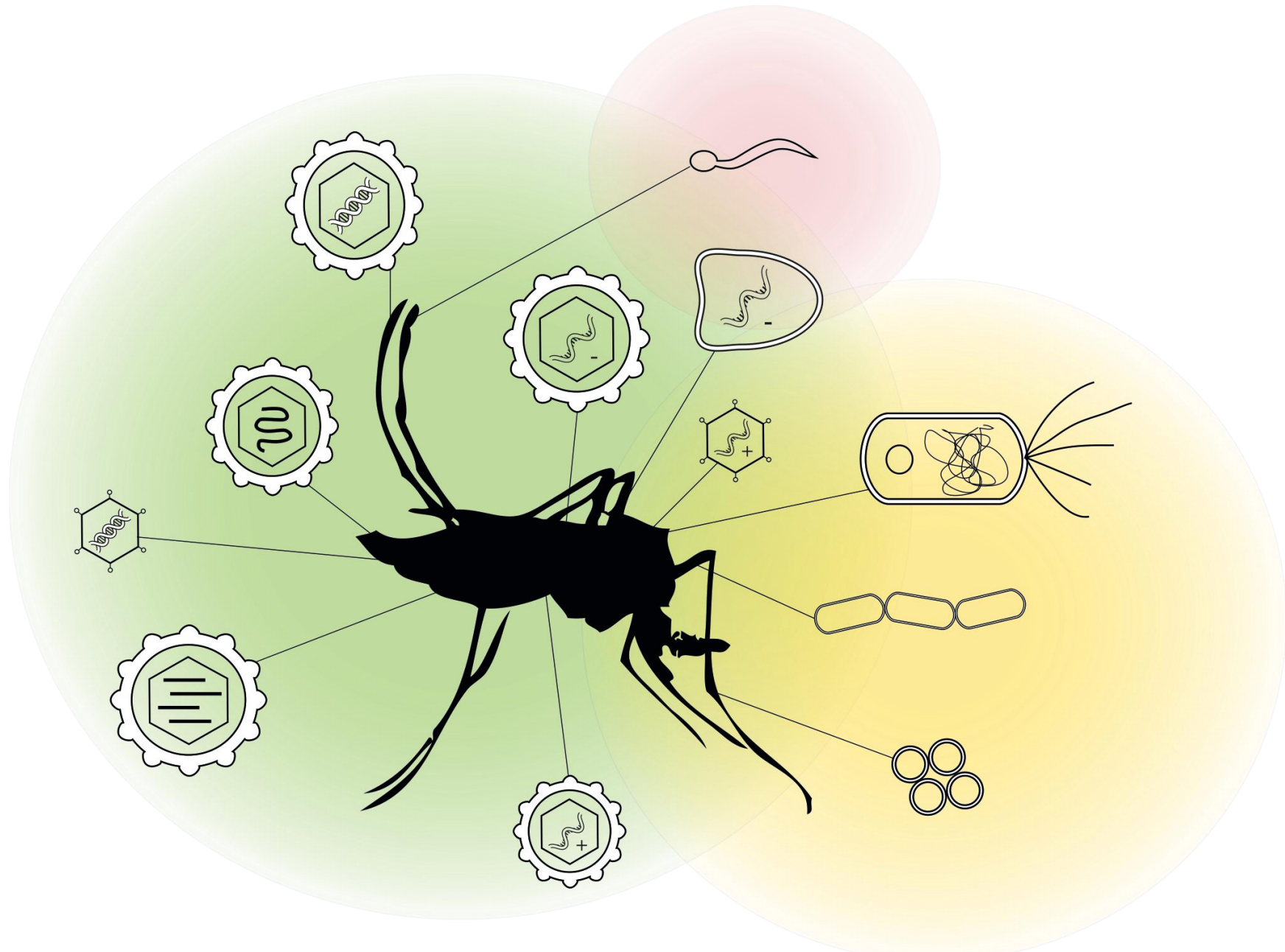
- G x G interactions
- *Ae. aegypti* vs. *Ae. albopictus*

## Non-genetic determinants of vector-virus interactions

- Temperature
- Larval microbiota
- Vertebrate host factors

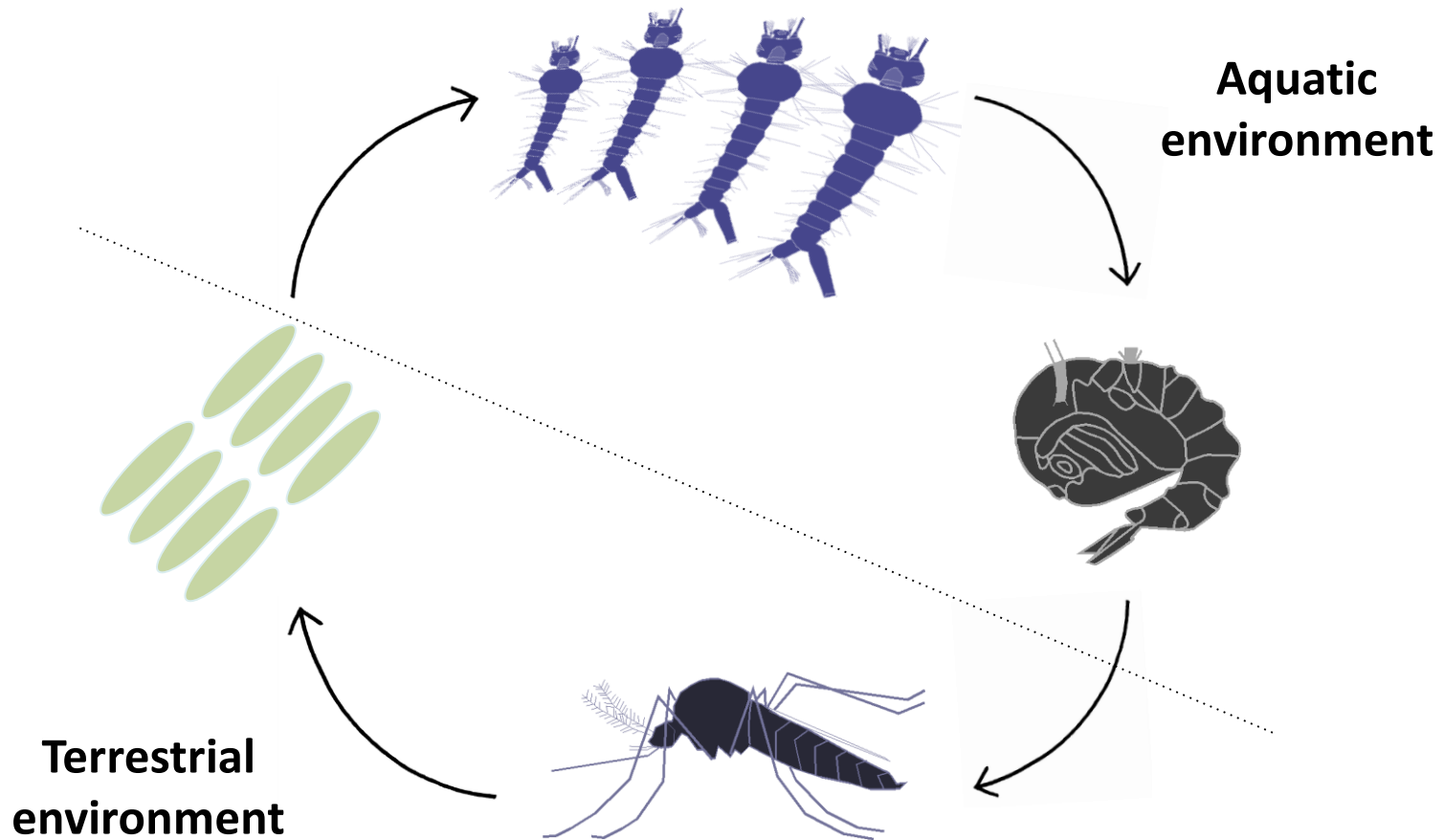


# The mosquito 'holobiont'



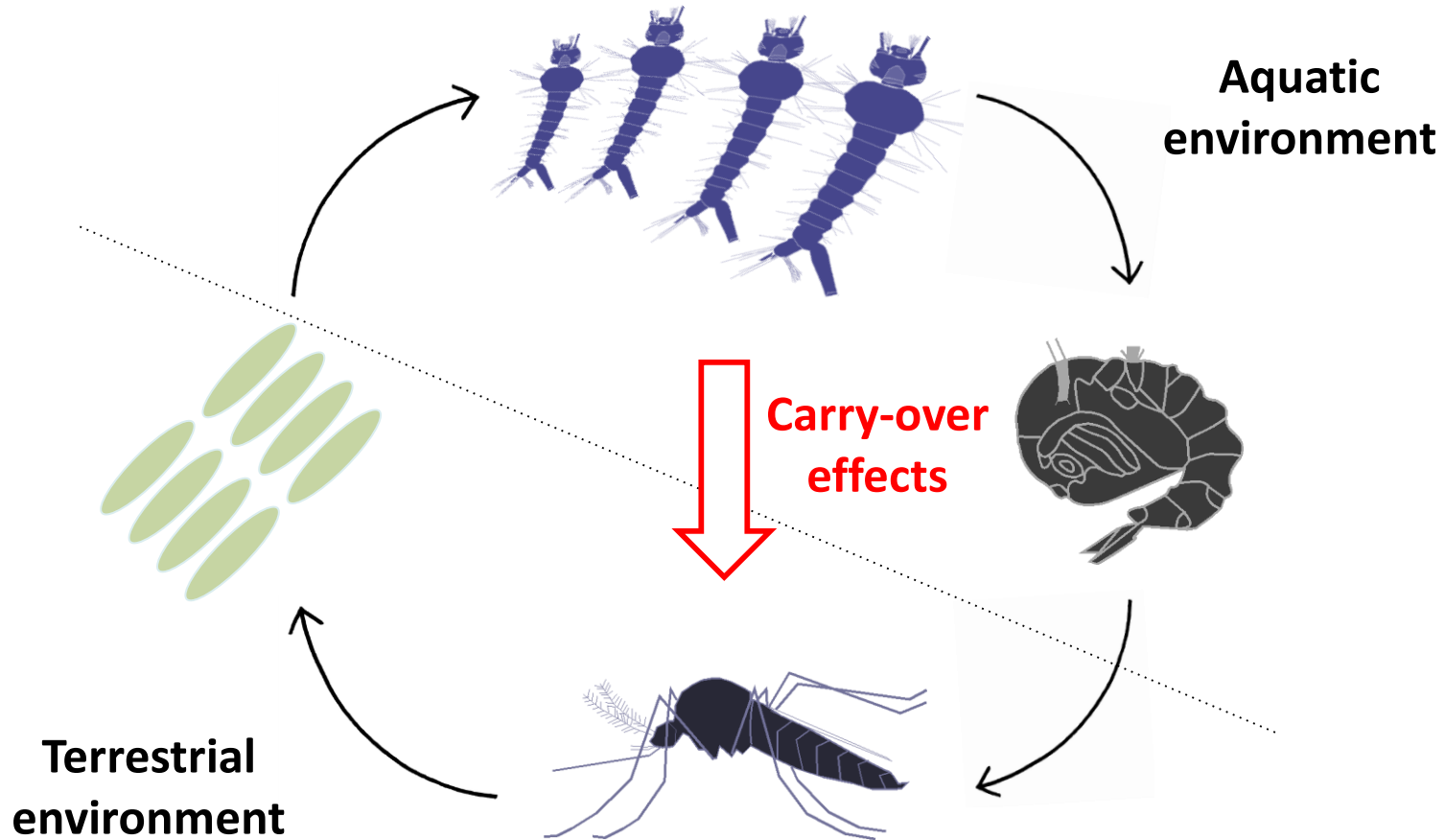
# Carry-over effects of larval microbiota

## Mosquito life cycle



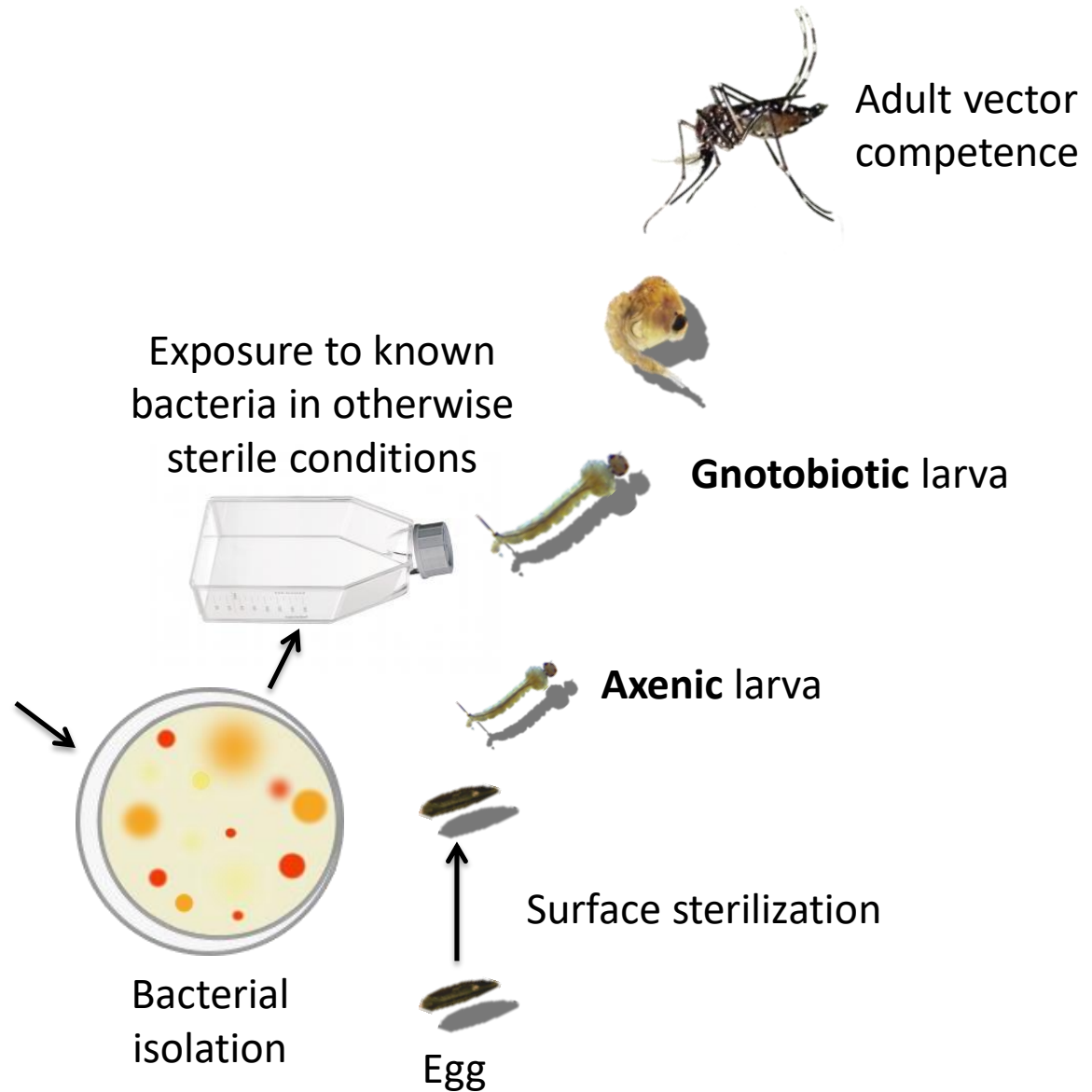
# Carry-over effects of larval microbiota

## Mosquito life cycle

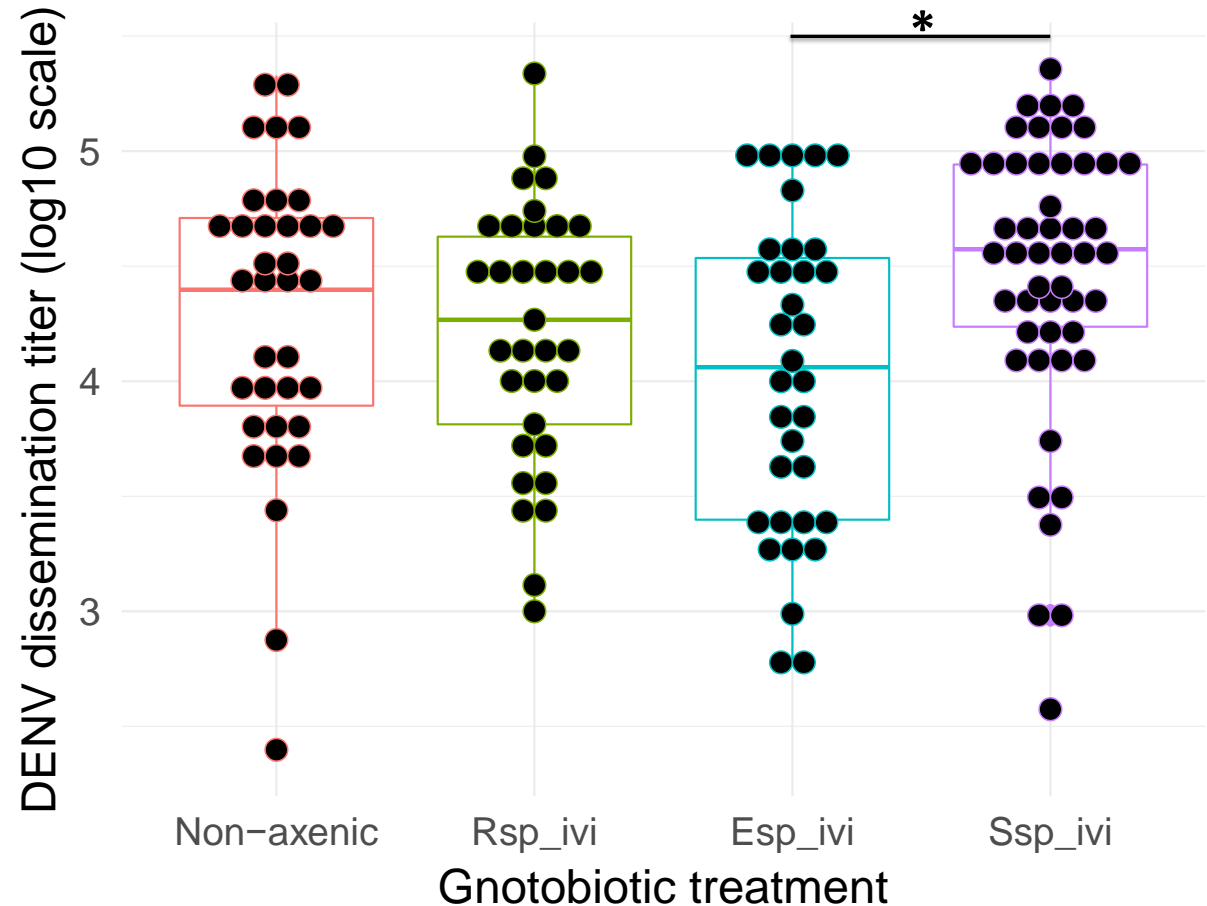
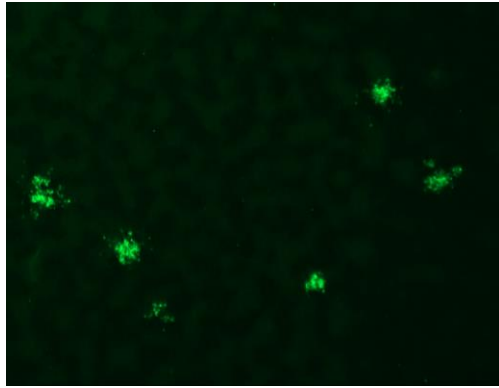


# Gnotobiotic mosquitoes

Water sampling from natural breeding sites in Gabon



# Exposure to different bacteria during larval development affects adult DENV vector competence



Non-axenic = control

Rsp\_ivi = *Rhizobium* isolate

Esp\_ivi = Enterobacteriaceae isolate

Ssp\_ivi = *Salmonella* isolate

# Talk outline



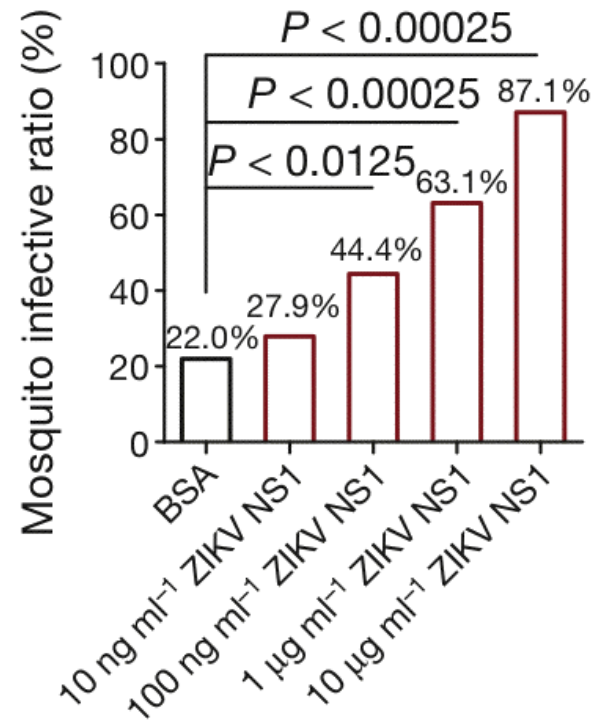
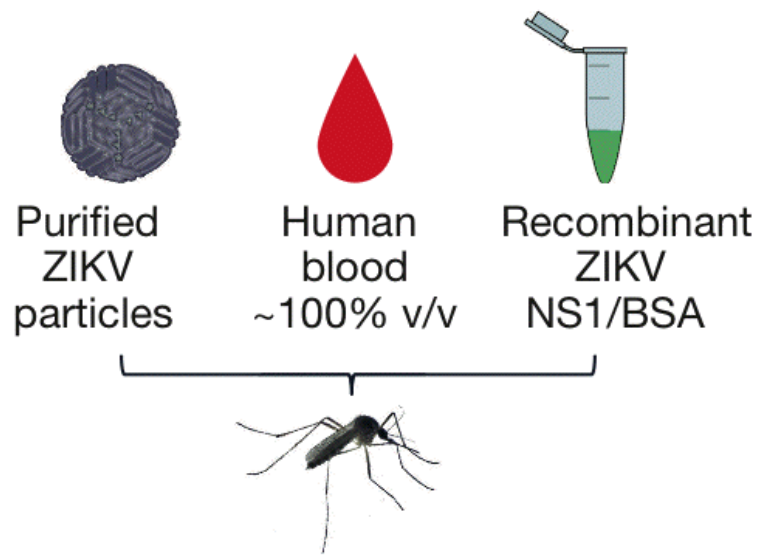
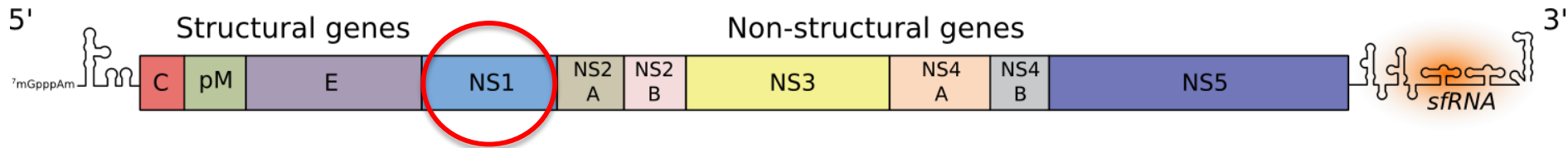
## Genetic determinants of vector-virus interactions

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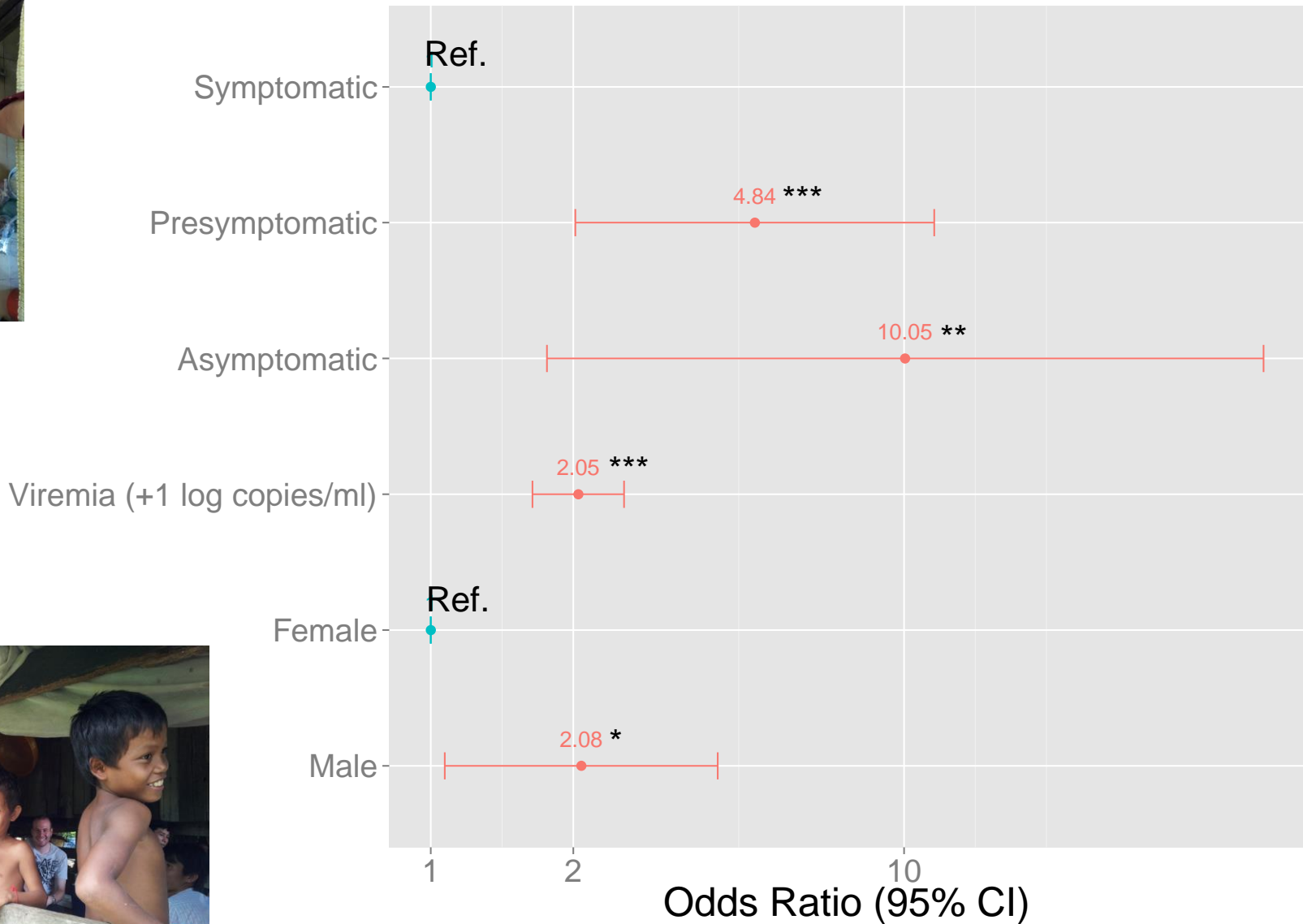
## Non-genetic determinants of vector-virus interactions

- Temperature
- Larval microbiota
- Vertebrate host factors

# NS1 facilitates ZIKV infection of *Aedes aegypti*



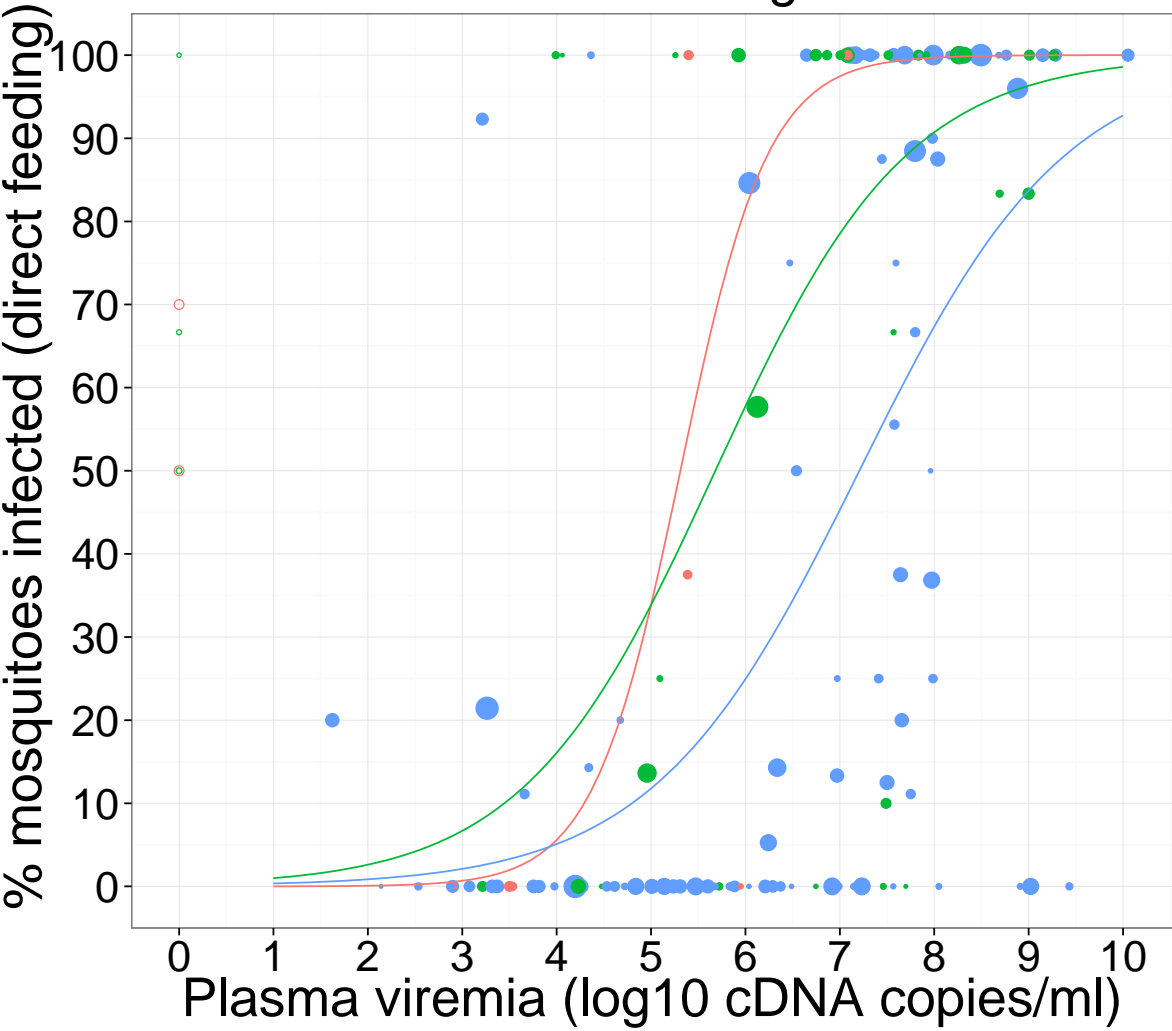
# Predictors of successful human-to-mosquito DENV transmission in Cambodia





# DENV dose response by disease category

*Direct feeding*



Disease category	OID <sub>50</sub> (95% CI)
Asymptomatic	5.31 (4.81–5.80)
Presymptomatic	5.68 (5.30–6.00)
Symptomatic	7.21 (7.05–7.36)

# Take home



- *Aedes aegypti* is a **major arbovirus vector** owing to its widespread distribution, human biting behavior and vector competence
- Vector competence is governed by specific **G x G interactions** between mosquito and arbovirus strains
- Arbovirus transmission by mosquitoes is modulated by manifold **biotic and abiotic factors** of the environment
- Understanding variation in vector-virus interactions provides insights into arbovirus epidemiology and supports development of **innovative strategies** to interrupt arbovirus transmission

# Acknowledgements

## IP Paris

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Veasna Duong

## IP Dakar

Amadou Sall

Cheikh Diagne

## IP Cayenne

Isabelle Dusfour

## IP Guadeloupe

Anubis Vega-Rua

## UniNorte Colombia

Claudia Romero-Vivas

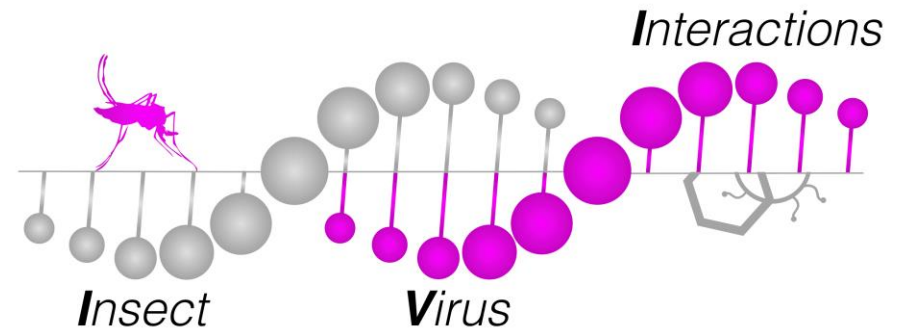
## CVR Glasgow

Alain Kohl

## UVR Entebbe

Julius Lutwama

# Thank you for your attention

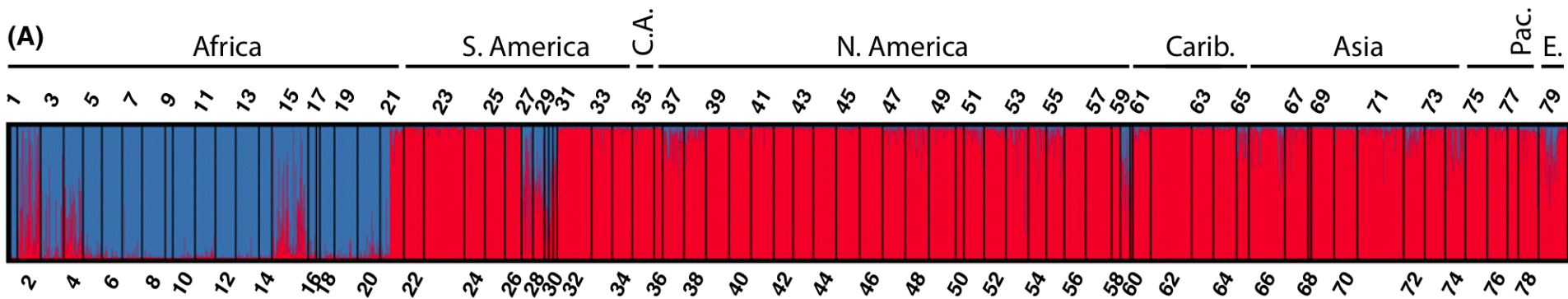
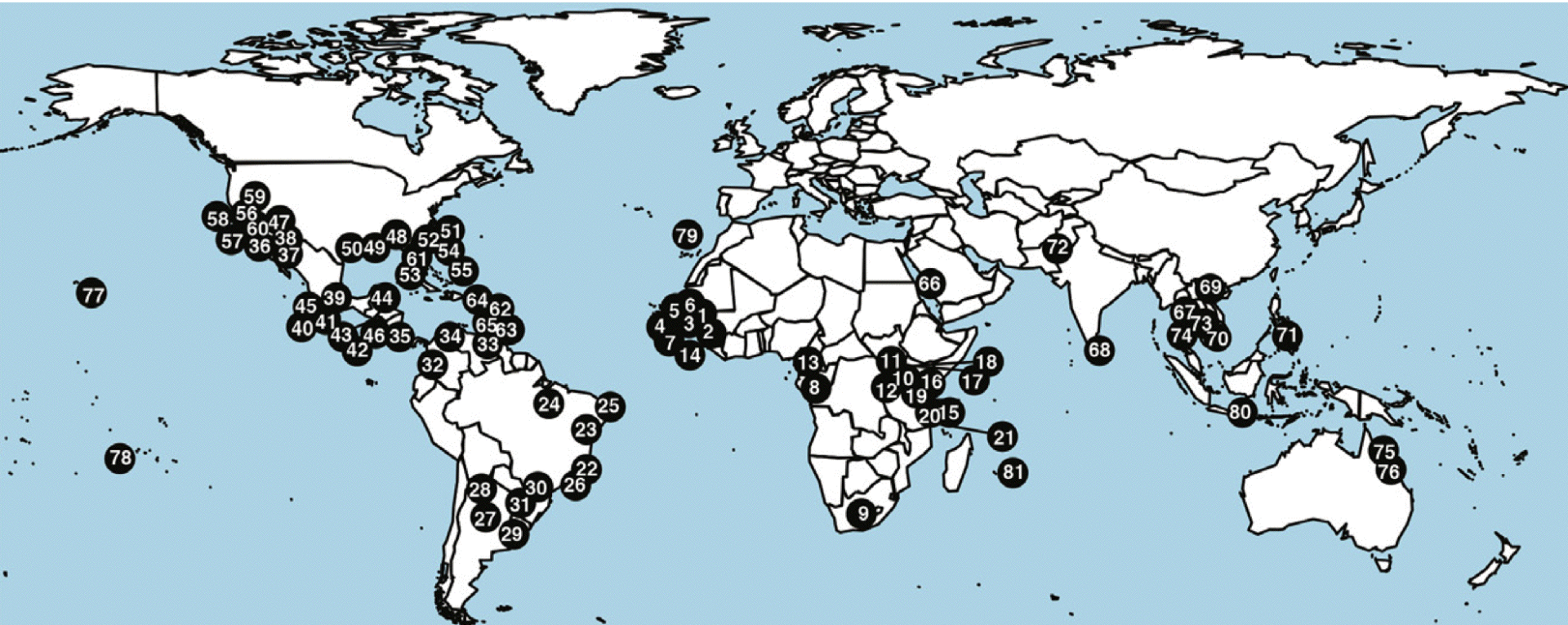


<http://research.pasteur.fr/en/team/insect-virus-interactions>

[louis.lambrechts@pasteur.fr](mailto:louis.lambrechts@pasteur.fr)



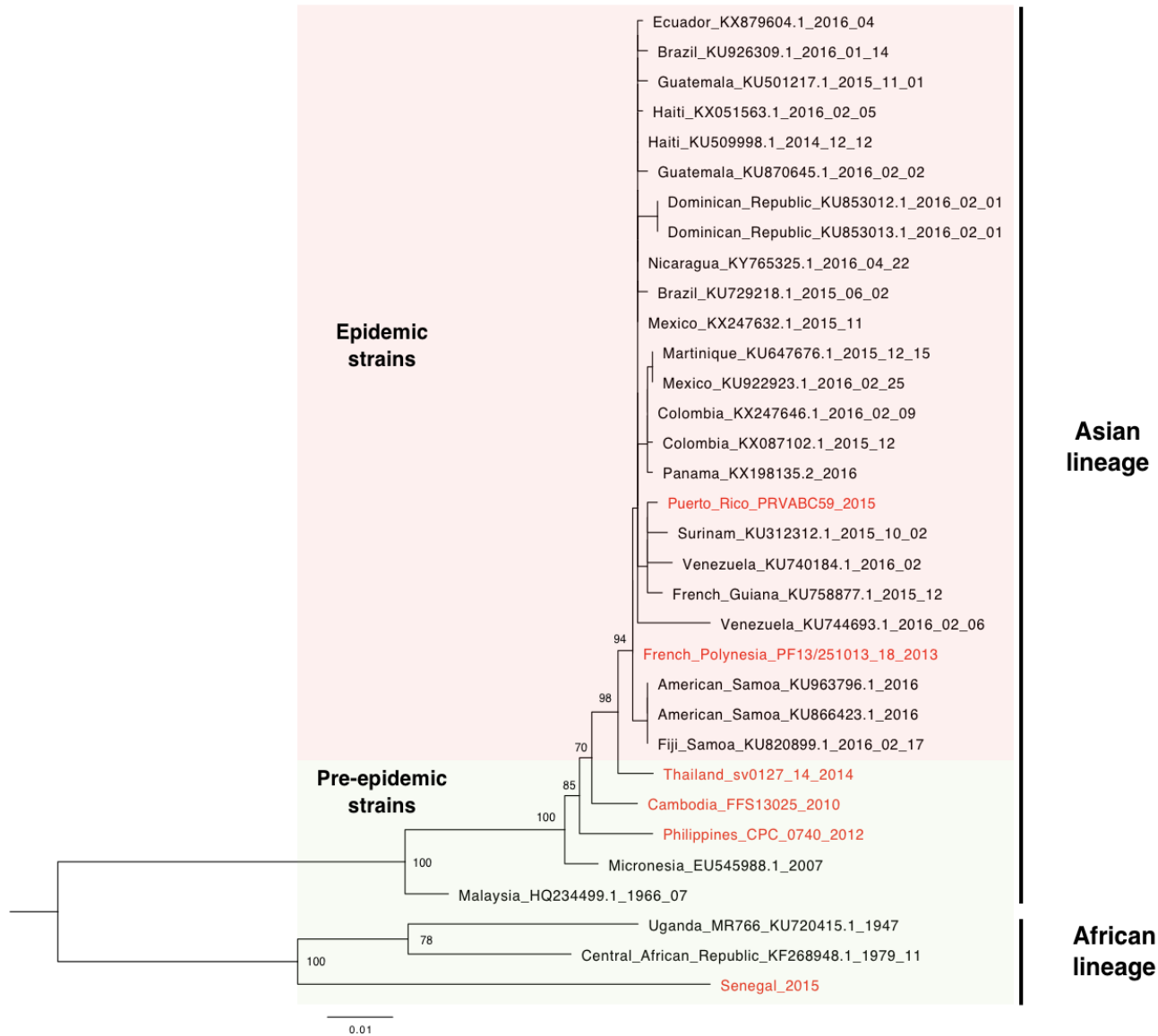
# *Aedes aegypti* subspecies



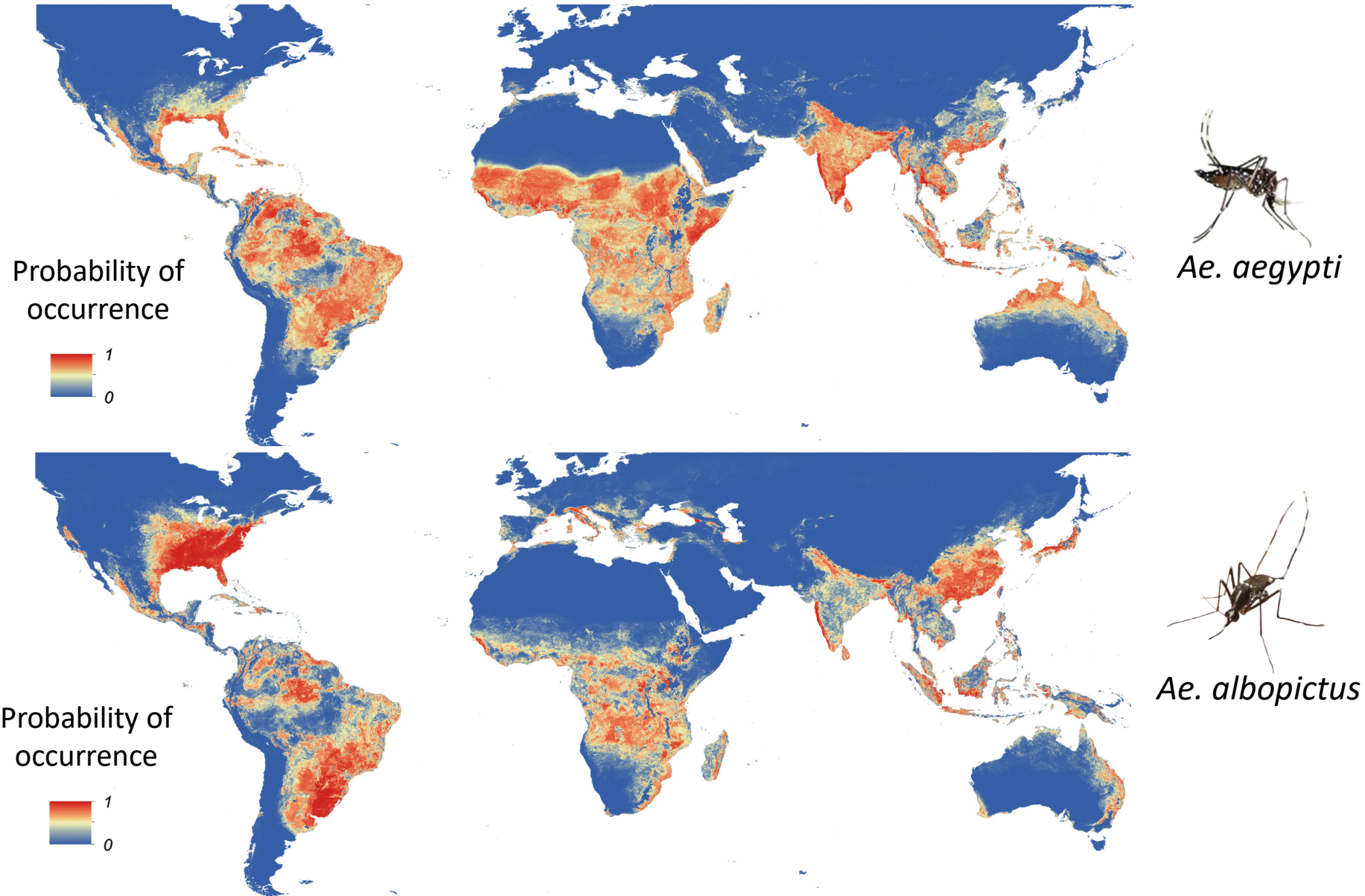
Blue = subspecies *Ae. aegypti formosus*

Red = subspecies *Ae. aegypti aegypti*

# ZIKV strains



# Distribution of *Aedes aegypti* and *Aedes albopictus*

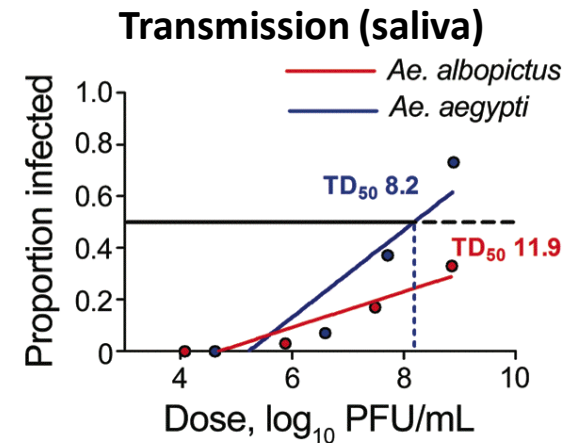
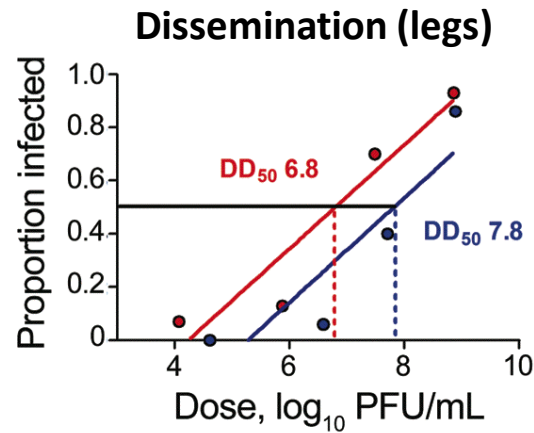
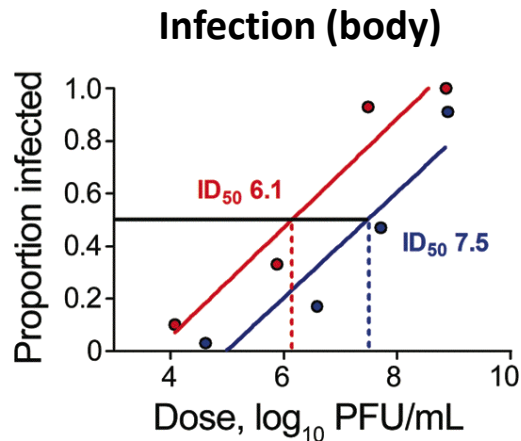


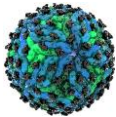


# ZIKV competence of *Ae. aegypti* vs. *Ae. albopictus*

*Ae. albopictus* is more susceptible but less competent than *Ae. aegypti*

  
Honduras  
ZIKV  
strain



  
Cambodia  
ZIKV  
strain

