

# ELIMINATE DENGUE<sup>TM</sup> OUR CHALLENGE



ELIMINAR A  
**DENGUE**  
DESAFIO BRASIL



ELIMINAR EL  
**DENGUE**  
DESAFIO COLOMBIA

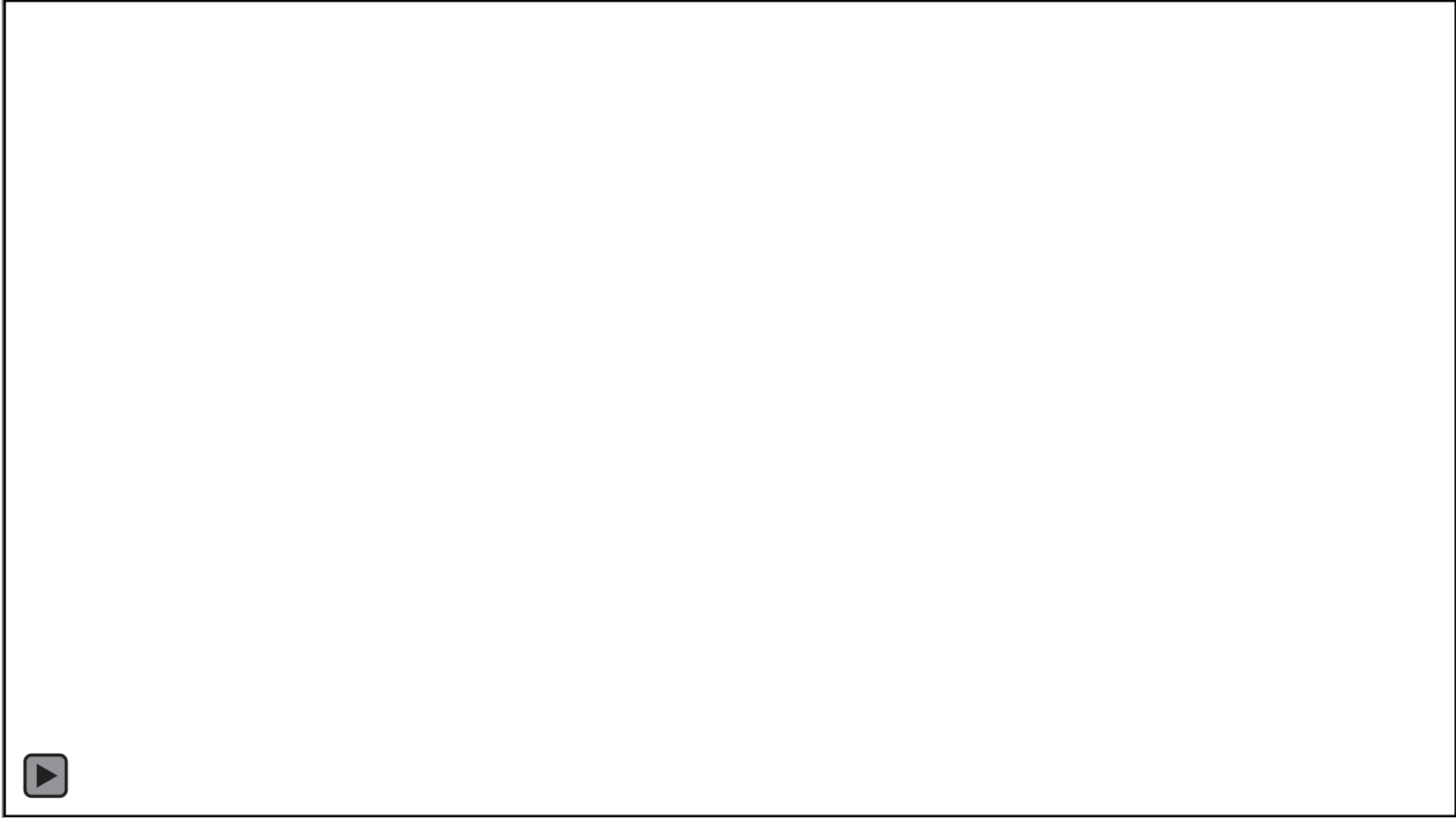


ELIMINATE  
**DENGUE**  
TANTANGAN INDONESIA



HƯỚNG TỚI LOẠI TRỪ  
**SỐT XUẤT HUYẾT**  
TẠI VIỆT NAM





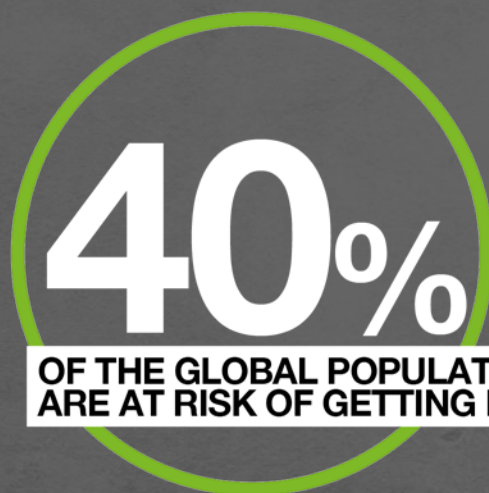


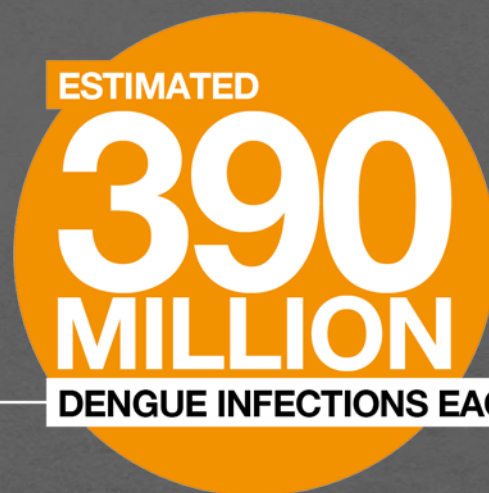
# Global burden of dengue

 **8.9B** global government spend (USD) on dengue each year

**45%** of health costs are borne by the family.

These costs can be upwards of **3x** the family's monthly income.

 **40%**  
OF THE GLOBAL POPULATION ARE AT RISK OF GETTING DENGUE.

 **ESTIMATED 390 MILLION**  
DENGUE INFECTIONS EACH YEAR.

# Zika virus outbreak

# 2,211

reported microcephaly cases in Brazil since February 2016, up on the previous yearly average of 163 cases nation-wide.

ZIKA HAS SPREAD RAPIDLY TO  
**75**  
COUNTRIES  
OVER 60% OF THESE SINCE 2015



# What is Wolbachia?

- Naturally occurring bacteria
- Transmitted from female insect to offspring through the eggs
- Safe for humans, animals and the environment
- **Reduces ability of mosquitoes to transmit disease (dengue, Zika, chikungunya)**

60%

OF ALL INSECT SPECIES  
HAVE *WOLBACHIA*

# Broad range of pathogen interference

Human pathogens that *Wolbachia* has been shown to interfere with in mosquito vectors:

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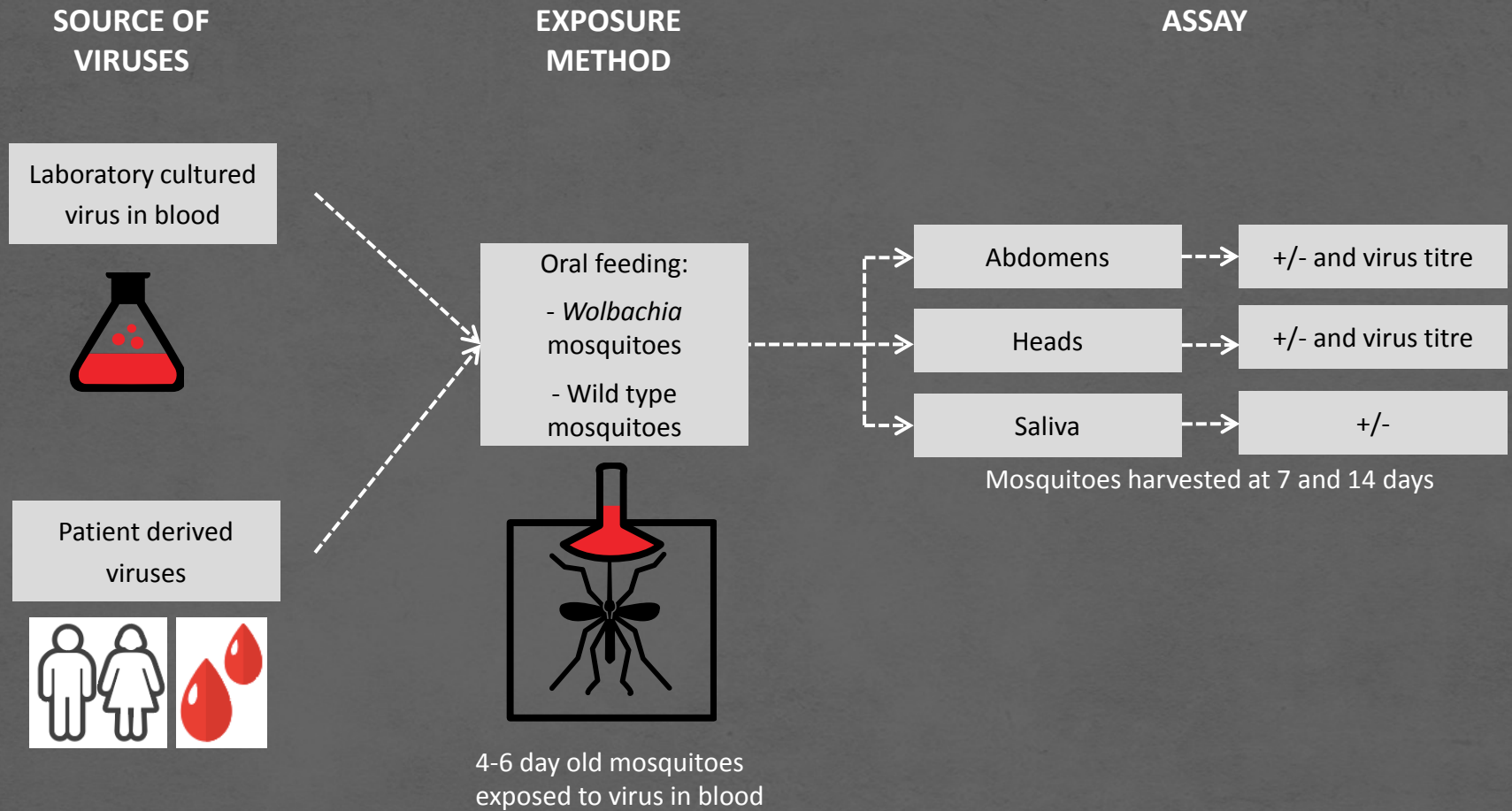
- ✓ Dengue viruses – all serotypes
- ✓ Yellow Fever
- ✓ West Nile
- ✓ Chikungunya
- ✓ Zika
- ✓ Filarial Nematodes – *Brugia malayi*
- ✓ Malaria parasites – *Plasmodium gallanaceum*, *P. falciparum*, *P. berghei*

Human pathogens where *Wolbachia* interference is not yet demonstrated but predicted:

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- Other Flaviviruses – eg Japanese encephalitis
- Other Alphaviruses – eg Semliki Forest virus, Venezuelan Equine encephalitis
- Other species of *Plasmodium* and Filarial nematodes

# Testing of *Wolbachia Aedes aegypti*




































# DENV-1 viremic blood feeds (n=29)

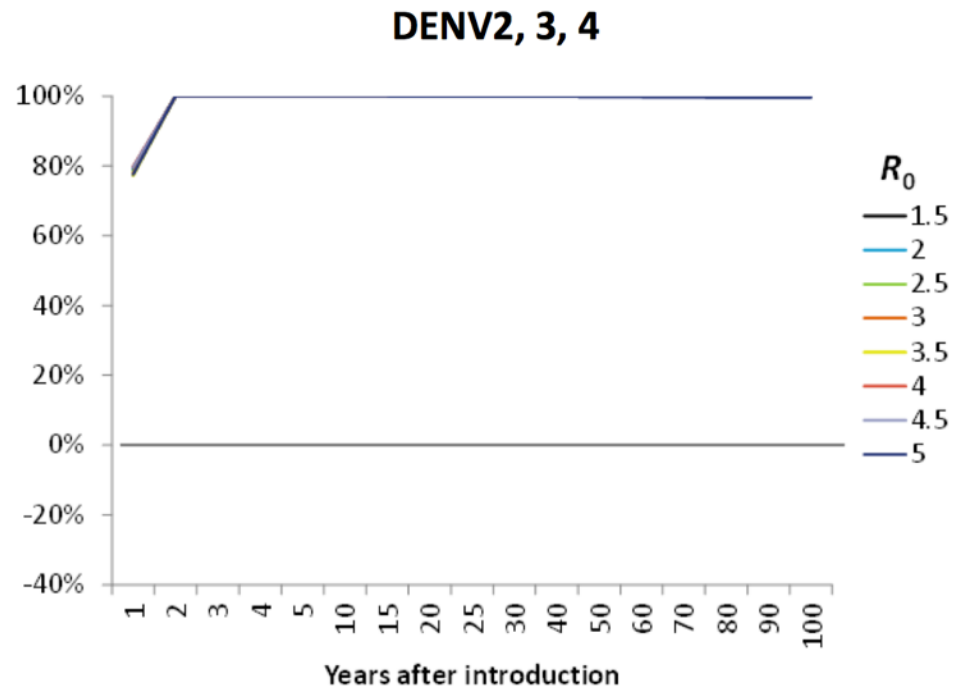
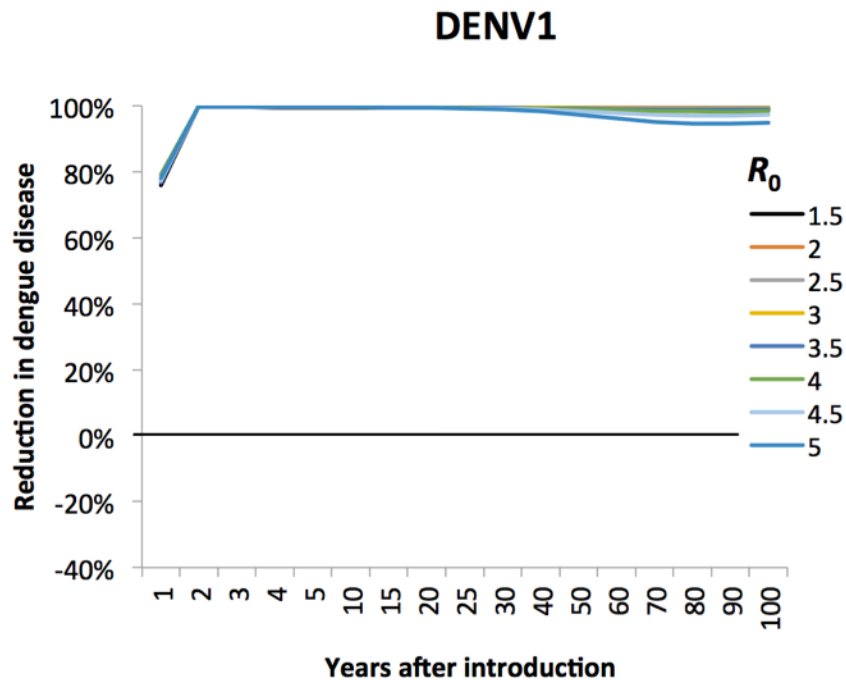
Patient	Serotype	Log10 Viremia	% Saliva +ve DENV			
			Day 10		Day 14	
			WT	wMel	WT	wMel
30DX/69	DENV-1	5.42	0.00	0.00		
30DX/188		5.99	0.00	0.00	0.33	0.00
30DX/68		6.04	0.00	0.00	0.50	0.00
30DX/187		6.95	0.67	0.09	0.88	0.33
30DX/114		7.31	0.13			
29DX/51		7.45	0.77	0.00	0.86	0.00
30DX/191		7.61	0.42	0.00	0.58	0.00
30DX/181		7.64	0.30	0.10	0.75	0.17
30DX/121		7.67	0.20	0.00	0.80	0.00
13DX1/1191		7.73	0.38	0.00	0.75	0.80
13DX16/244		7.78	0.33	0.40	1.00	0.40
30DX/70		7.86	0.30	0.00	0.86	0.00
13DX16/174		7.91	0.25	0.14	0.75	0.20
30DX/113		7.91	1.00	0.00	0.40	0.20
30DX/128		7.91	0.70	0.33	1.00	0.71
13DX1/1183		8.19	0.75	0.00	0.89	0.42
30DX/196		8.25	0.57	0.20	0.67	0.00
30DX/173		8.29	0.50	0.30	0.67	0.00
30DX/175		8.35	0.22	0.00	0.00	
13DX3/863		8.45	0.57	0.20	0.40	0.83
30DX/81		8.57	0.10	0.00	0.10	0.10
30DX/184		8.92	0.75	0.38	0.75	0.33
30DX/75		9.07	0.50	0.00	1.00	0.10
13DX16/172		9.12	0.13	0.20	0.70	1.00
30DX/189		9.28	0.58	0.13	0.80	0.40
30DX/194		9.73	0.67	0.29	0.75	0.22
30DX/193		9.91	0.83	0.00	1.00	0.25
30DX/82		9.93	0.60	0.10	0.60	0.10
30DX/195		10.57	0.91	0.18	1.00	0.17



# DENV-2, -3, -4 viremic blood feeds (n=32)

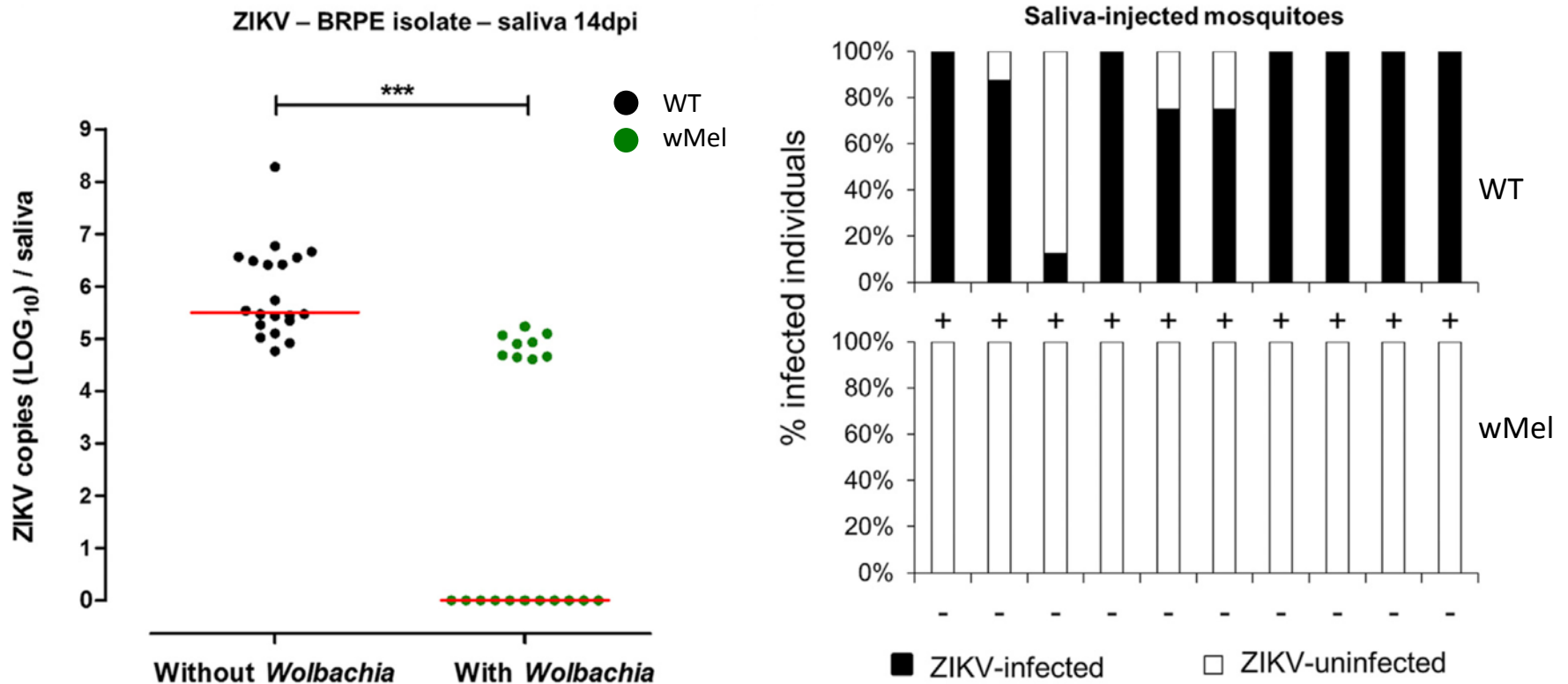
Patient	Serotype	Log10 Viremia	% Saliva +ve DENV			
			Day 10		Day 14	
			WT	wMel	WT	wMel
09DX/749	DENV-2	6.01 	0.00	0.00	0.00	0.00
09DX/747		6.13 	0.17	0.00	0.14	0.00
30DX/182		6.26 	0.00	0.00	0.40	
09DX/750		6.97 	0.00	0.00	0.00	0.00
30DX/83		7.13 	0.20	0.00	0.13	0.00
30DX/92		7.41 	0.00	0.00	0.00	0.00
13DX3/861		8.59 	0.20	0.00	0.67	0.50
29DX/31	DENV-3	8.95 	0.25	0.00		
13DX6/372		6.10 	0.00	0.00	0.10	0.00
29DX/58		6.73 	0.36	0.19	0.69	0.11
30DX/127		8.47 	0.00	0.00	0.00	0.00
09DX/746		8.82 	0.33	0.00	0.33	0.17
30DX/125	DENV-4	5.37 	0.00	0.00	0.00	
13DX3/938		5.42 	0.00		0.00	0.00
30DX/122		5.43 	0.22	0.00	0.50	0.00
30DX/76		6.61 	0.00	0.00	0.00	0.00
09DX/745		6.91 	0.00	0.00	0.00	0.00
30DX/71		6.93 	0.00	0.00	0.00	0.00
30DX/72		7.01 	0.00	0.00		
30DX/202		7.22 	0.00	0.00	0.17	0.00
30DX/174		7.63 	0.10	0.00	0.60	
29DX/59		7.73 	0.21	0.00	0.65	0.00
29DX/30		8.06 	0.00			
30DX/192		8.28 	0.25	0.00	0.57	0.00
30DX/198		8.41 	0.50	0.00	0.60	0.00
30DX/197		8.51 	0.38	0.00	0.50	0.00
13DX4/611		8.58 	0.17	0.00	0.89	0.33
30DX/85		8.68 	0.00	0.00	0.50	0.00
30DX/201		8.68 	0.70	0.00	0.86	0.00
29DX/69		8.82 	0.00	0.00	0.44	0.10
30DX/190		8.86 	0.64	0.00	0.33	0.00

# wMel deployment and dengue disease



90% transmission of wMel from female to progeny;  
15% fitness cost

# Zika virus titre in wMel *Aedes aegypti* (Brazil)

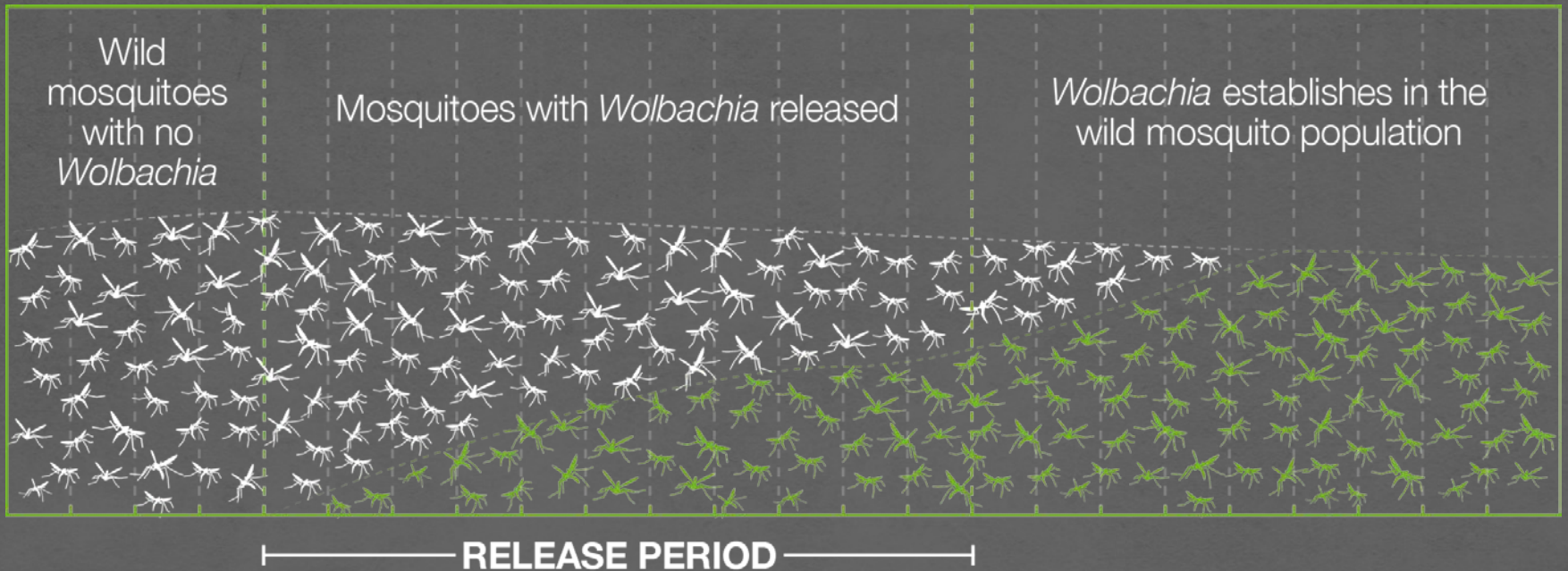


Mosquitoes (Brazil background) exposed to Zika virus  $5.0 \times 10^6$  PFU/ml (BRPE) and  $8.7 \times 10^3$  PFU/ml (SPH)

n = 20 mosquitoes were assayed at each time point

\* P Value <0.0001 (Mann-Whitney U Test)

# Release of Wolbachia mosquitoes



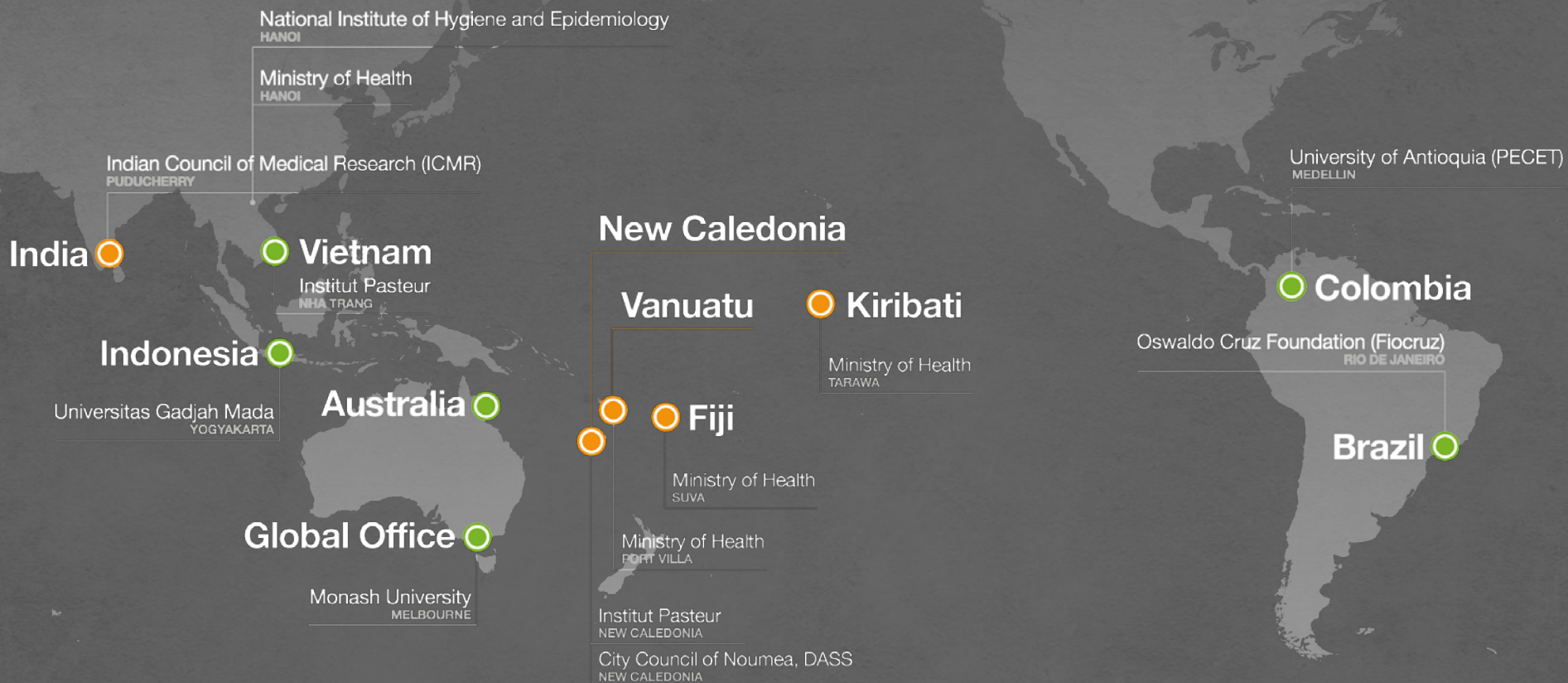
# Townsville

Mozzie box in action





# Eliminate Dengue Project Sites

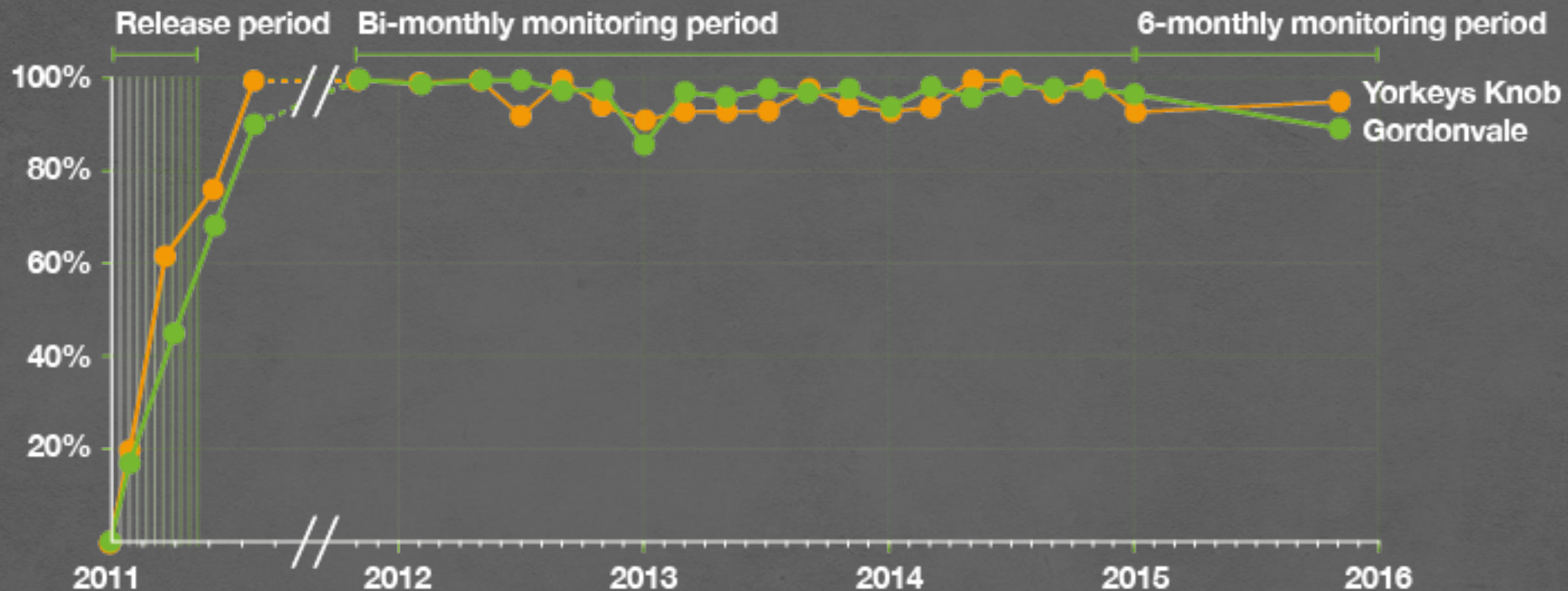




# Current field sites

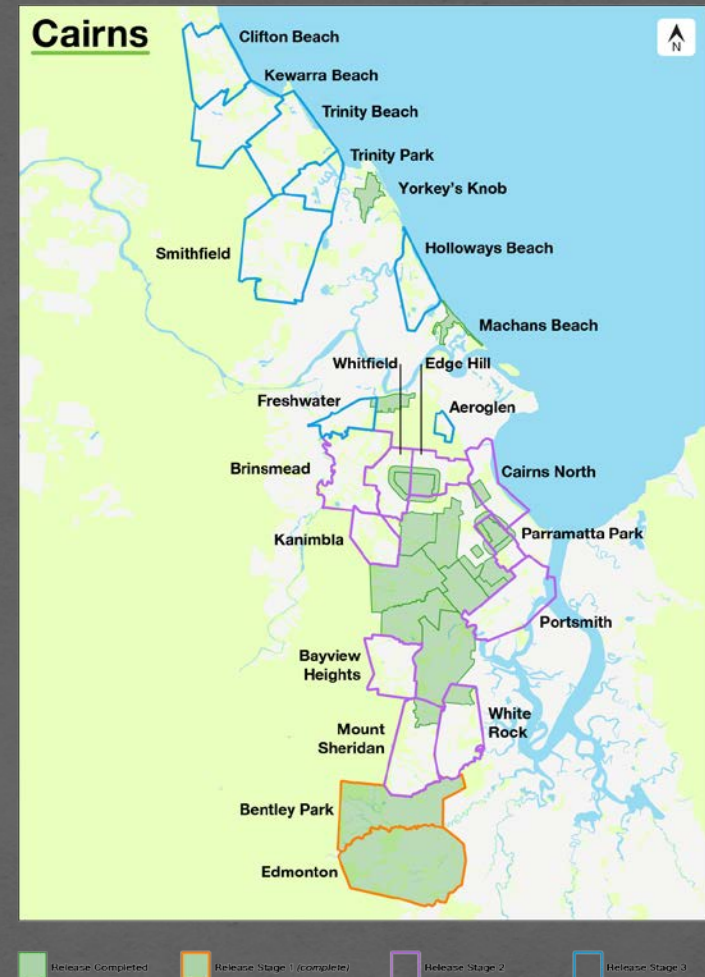


# Self-sustaining intervention



# 2011 – 2017: Cairns, Australia

- 28 suburbs over seven years
- High levels of *Wolbachia* in all areas
- No significant local dengue transmission



41km<sup>2</sup>



93,456  
people



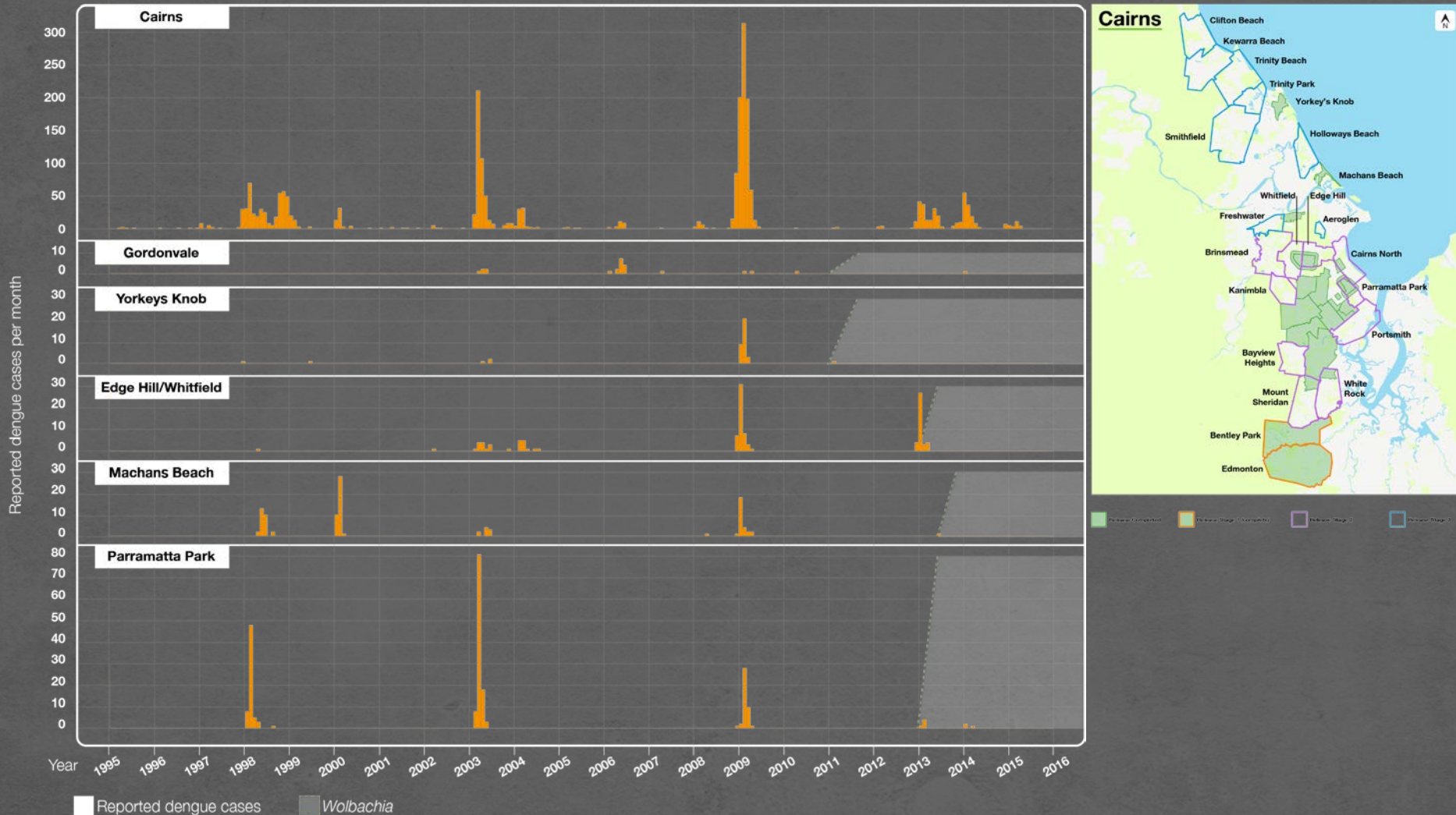
35,000  
households



# Disease surveillance, before vs after

## Cairns, Australia

(*Wolbachia* coverage = population 93,456)

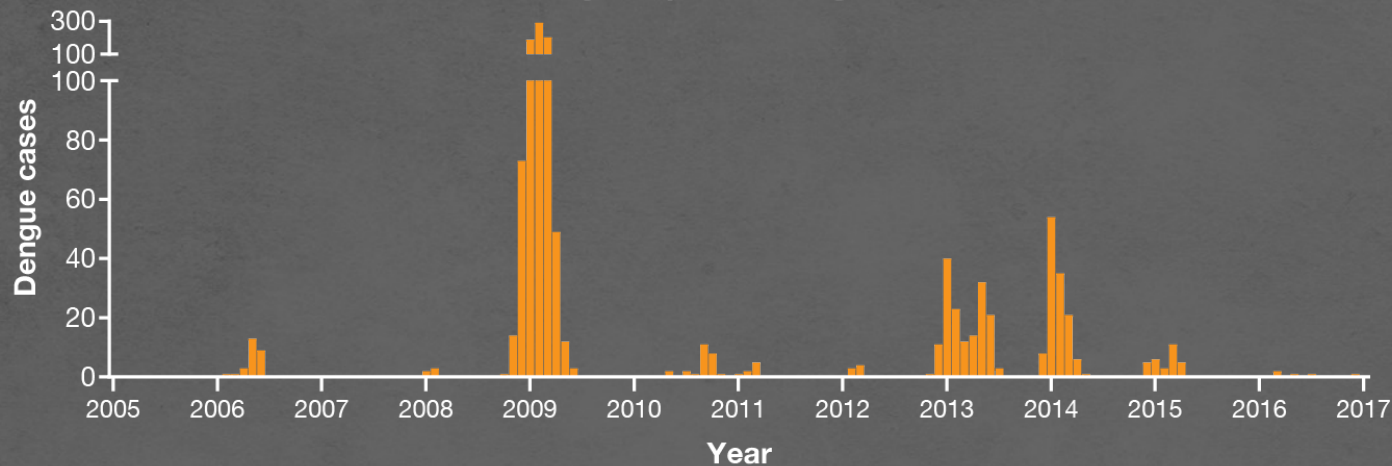


# Disease surveillance, before vs after

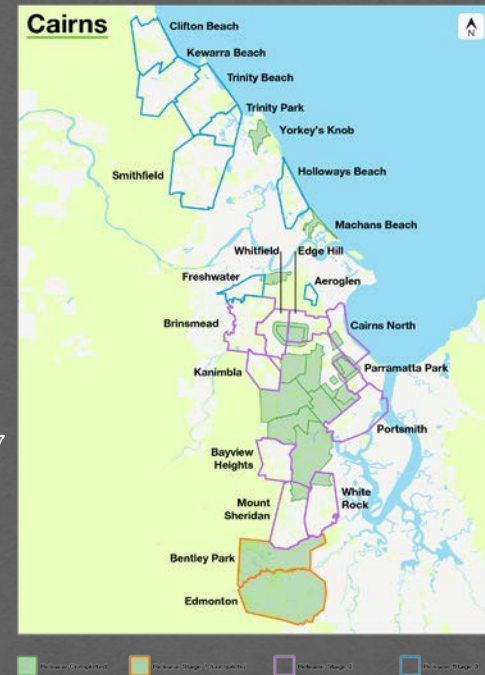
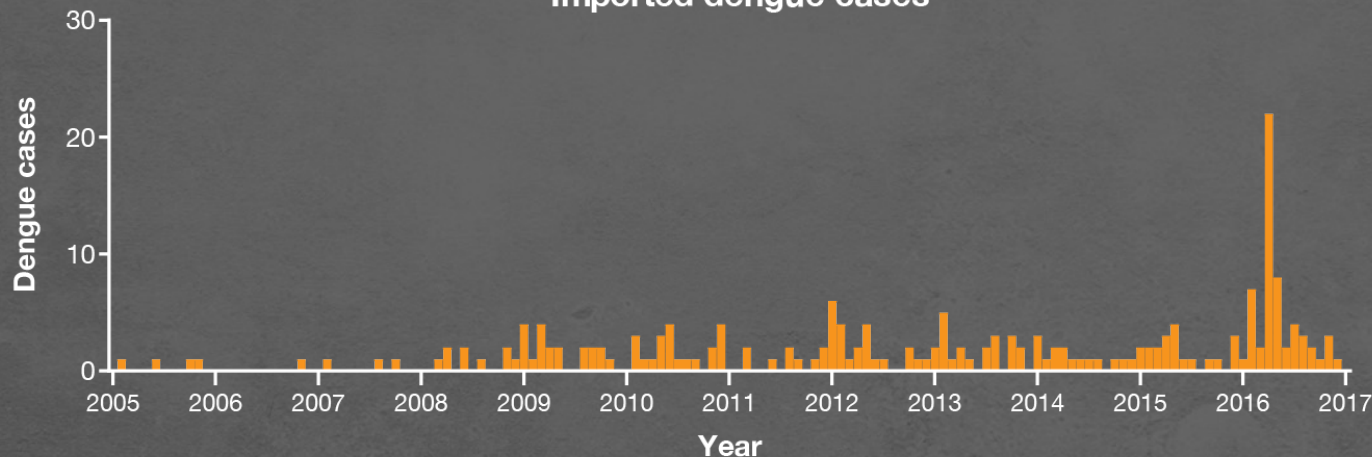
## Cairns, Australia

(*Wolbachia* coverage = population 93,456)

### Locally Acquired dengue cases



### Imported dengue cases



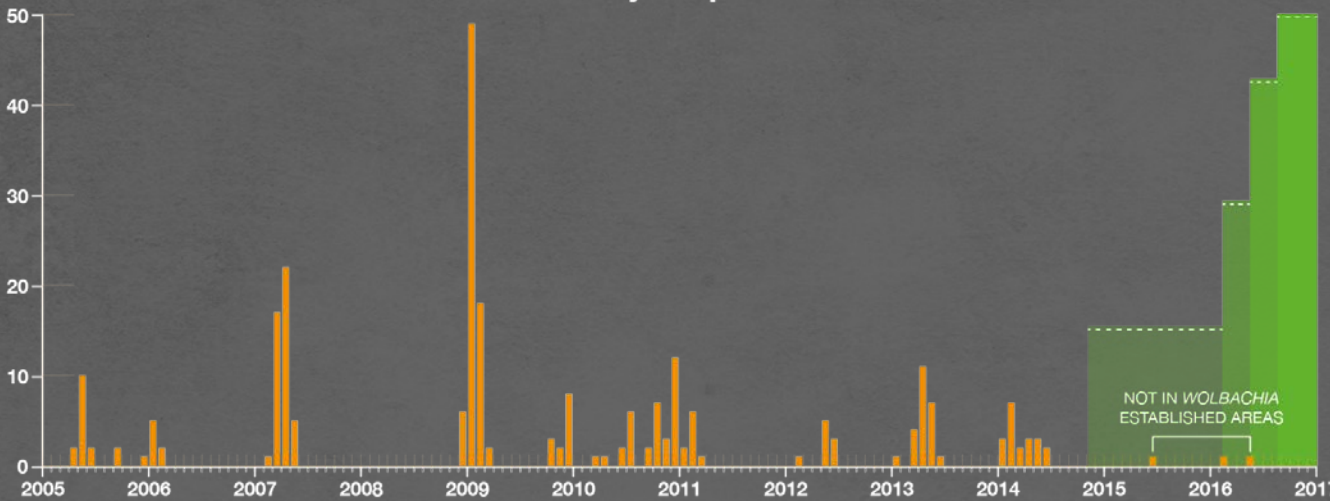


# Disease surveillance, before vs after

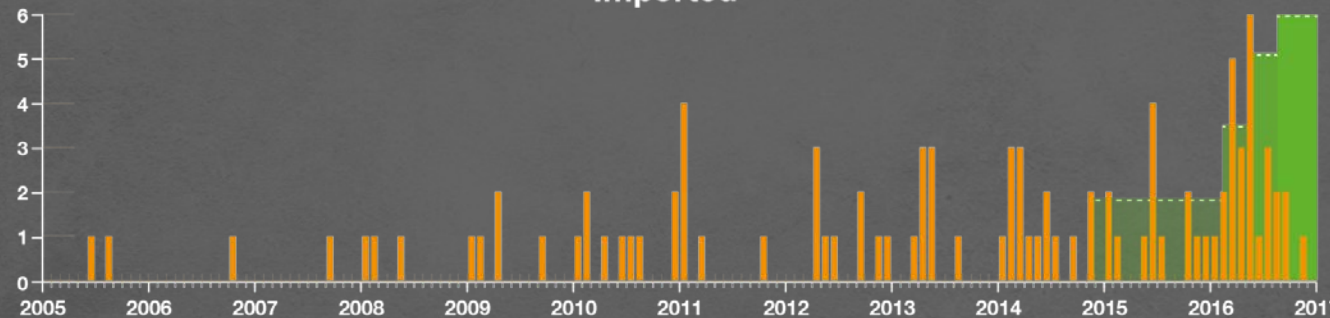
## Townsville, Australia

(*Wolbachia* coverage = population 139,757)

Locally Acquired



Imported



Dengue Cases



*Wolbachia* Release







**18** SUBURBS  
**43** KM<sup>2</sup>  
**12** MONTHS

**US\$16**  
PER PERSON

**50,000** people



**15,000** households



**3,000** release containers



**1.3M** mosquitoes



**180** BG traps (adult capture)



**30%** DEPLOYMENT

**21%** MONITORING

**24%** COMMUNITY  
ENGAGEMENT

**25%** OTHER

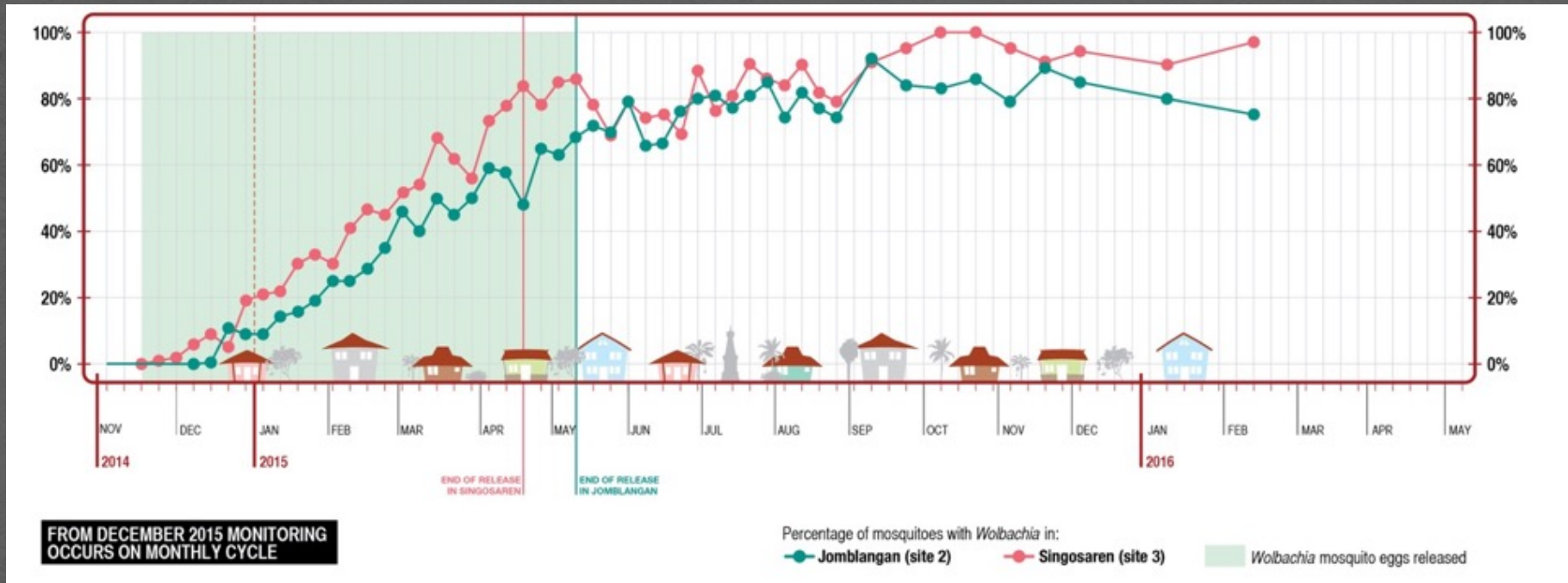


# Indonesia

- **2014:**  
Small scale releases in Nogotirto, Kronggahan, Jomblangan and Singosaren
- **2016 – 2019:**  
Large scale release and impact study in Yogyakarta city

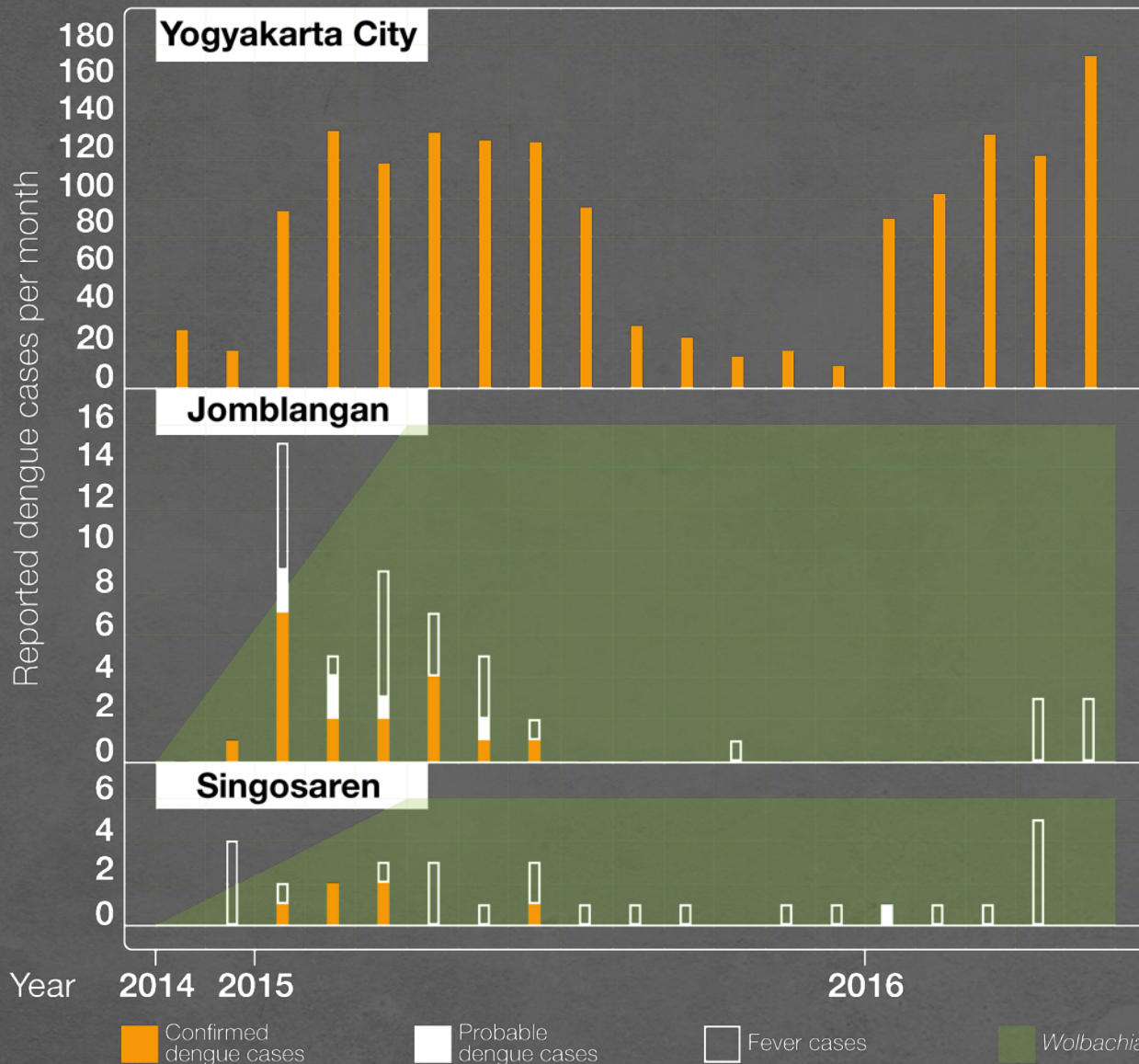


# Jomblangan & Singosaren, Yogyakarta



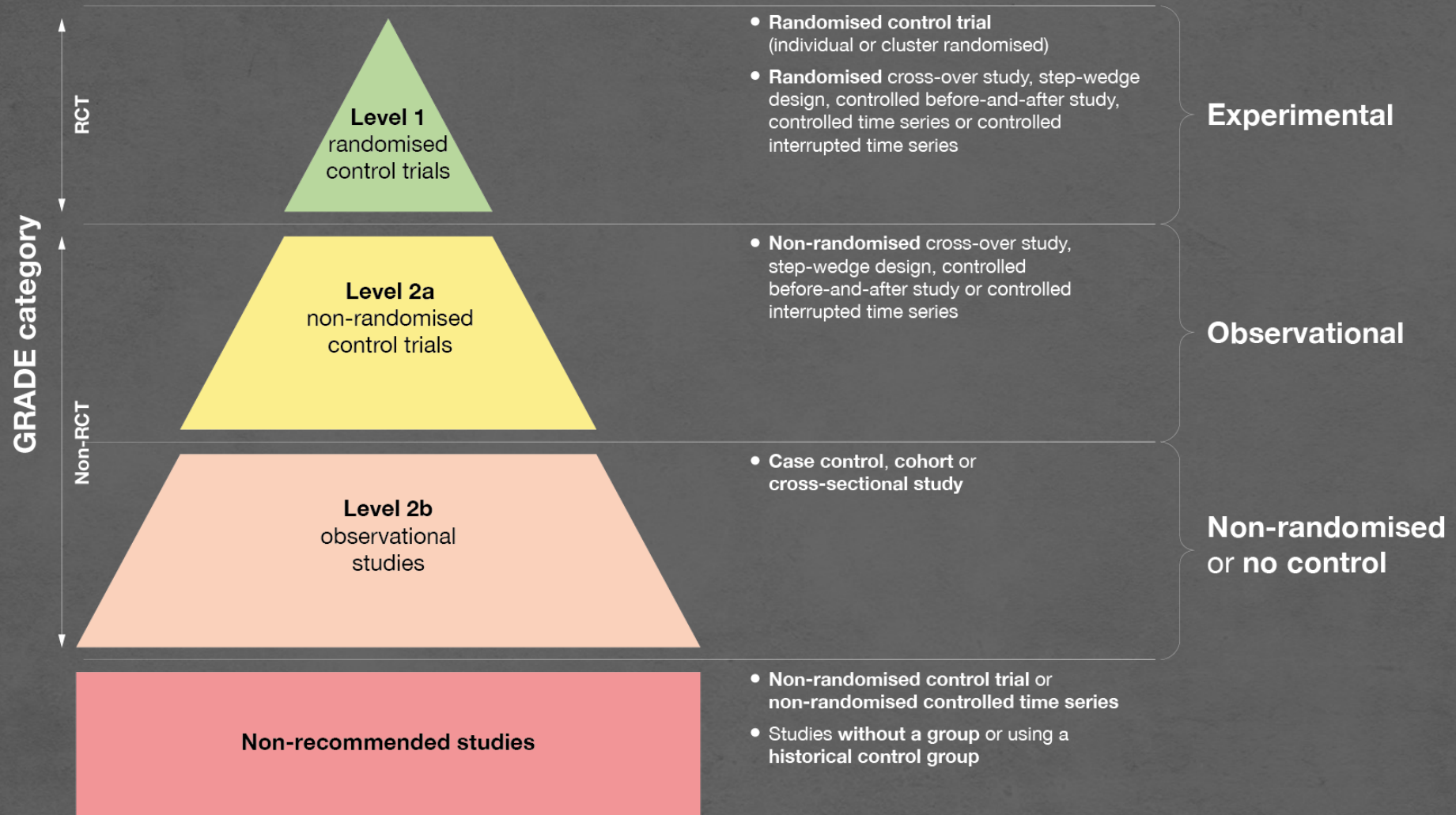


# Dengue cases in Yogyakarta City, Indonesia Jomblangan and Singosaren (Nov 2014 – May 2016)



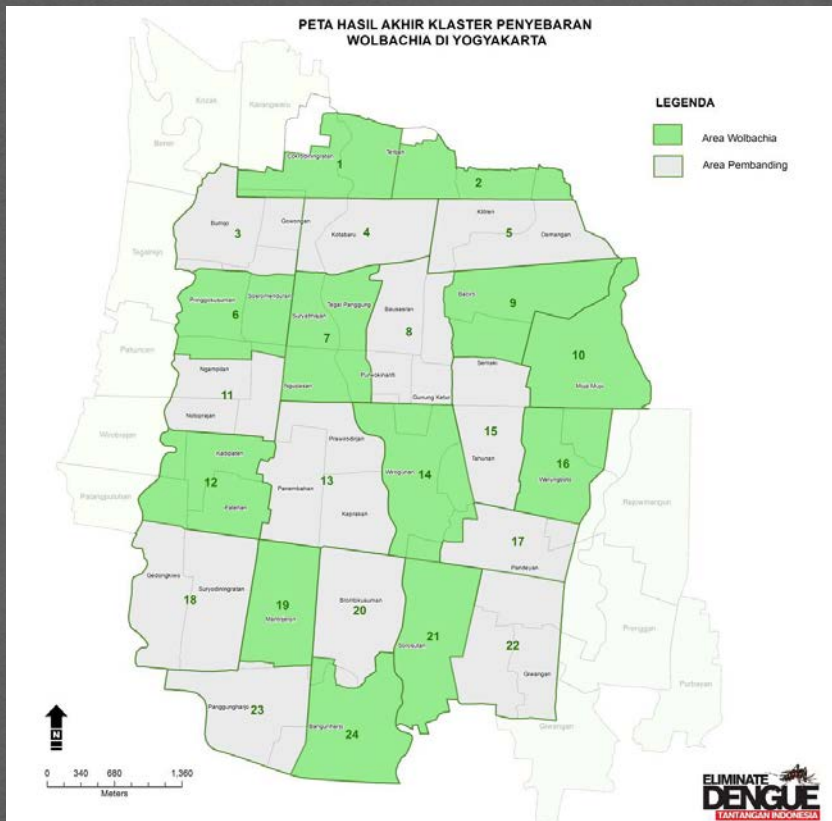
Confirmed and probable cases, identified through active febrile illness surveillance

# Hierarchy of study designs for assessing the efficacy of vector control interventions





# Clustered randomised trial of the effect of *Wolbachia* releases on dengue



## Study area:

- ~26km<sup>2</sup> (Yogyakarta City & Bantul District)
- Population 325,000
- 37 Kelurahan (administrative areas) within study site
- Divided into 24 clusters, using natural borders where possible

## Study arms (each 12 clusters):

- Un-treated: standard practice vector control
- Treated: Deployment of *Wolbachia* mosquitoes in addition to standard practice

## Allocation:

- Constrained randomisation, balancing on demographic characteristics and other covariates associated with dengue risk

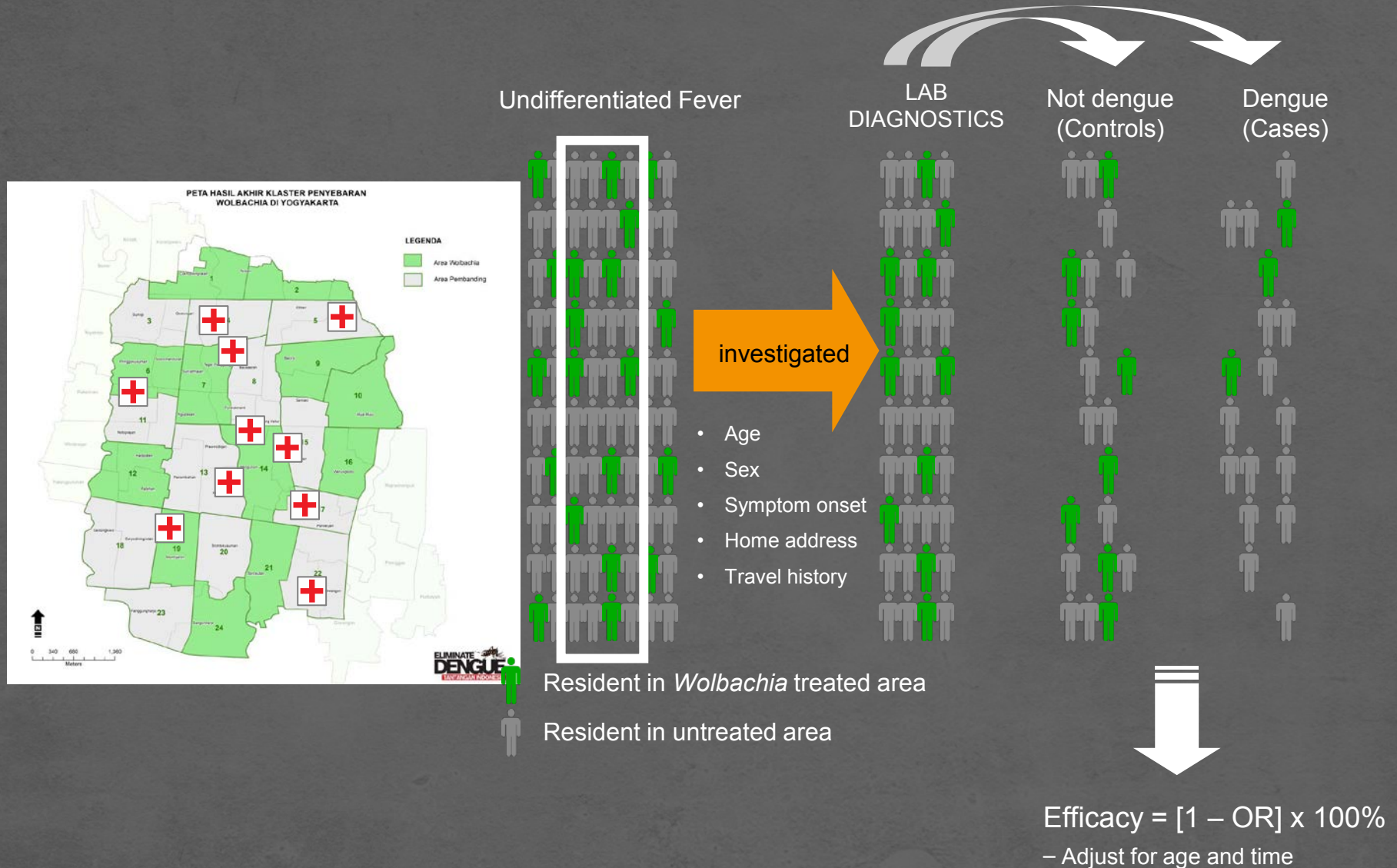


# Randomised Control Trial

## Yogyakarta, Indonesia



# Randomised Control Trial



# WHO Vector Control Advisory Group (VCAG)

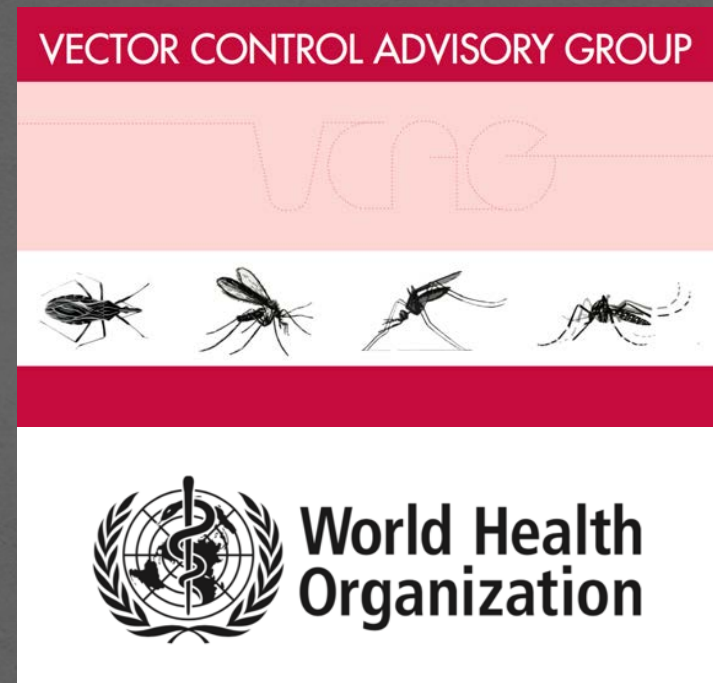
**18 March 2016:** WHO-VCAG reviewed new vector control tools for use in response to the Zika virus outbreak.

## Statements:

- *Wolbachia* reduce the mosquito's ability to transmit arboviruses to humans
- *Wolbachia* infection reduces viral replication of dengue, chikungunya and Zika viruses

## Recommendations:

- Pilot deployments under operational conditions & rigorous independent monitoring and evaluation
- Continue with plans for randomised control trials (RCTs) with epidemiological outcome





# Current field sites

Medellín &  
Bello, Colombia

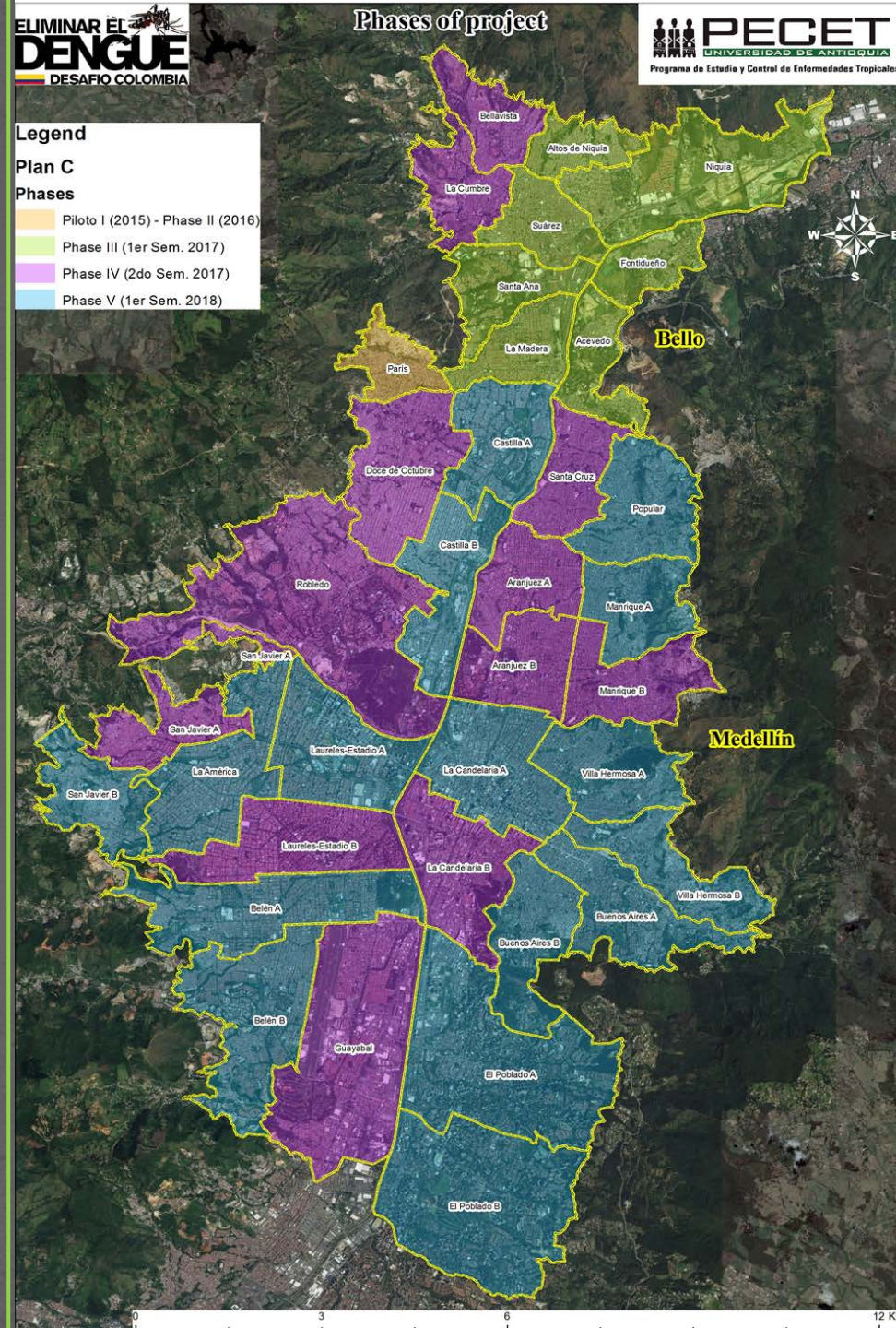
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# Colombia: scale up

2016-2018 – 2.5 M people





# Current field sites



Rio de Janeiro  
& Niterói, Brazil

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# Brazil: scale up

2016-2018 – 2.5 M people

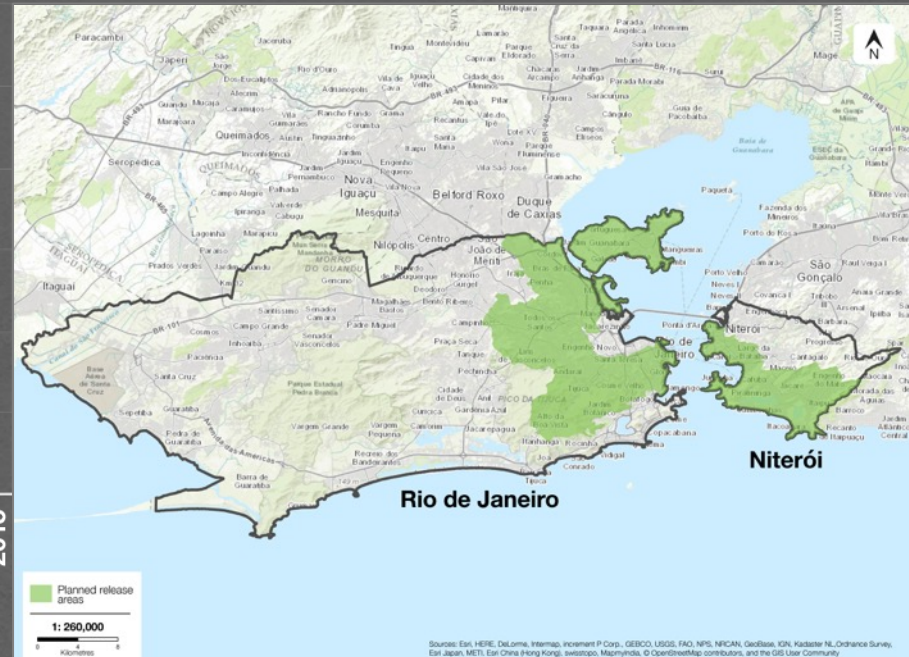
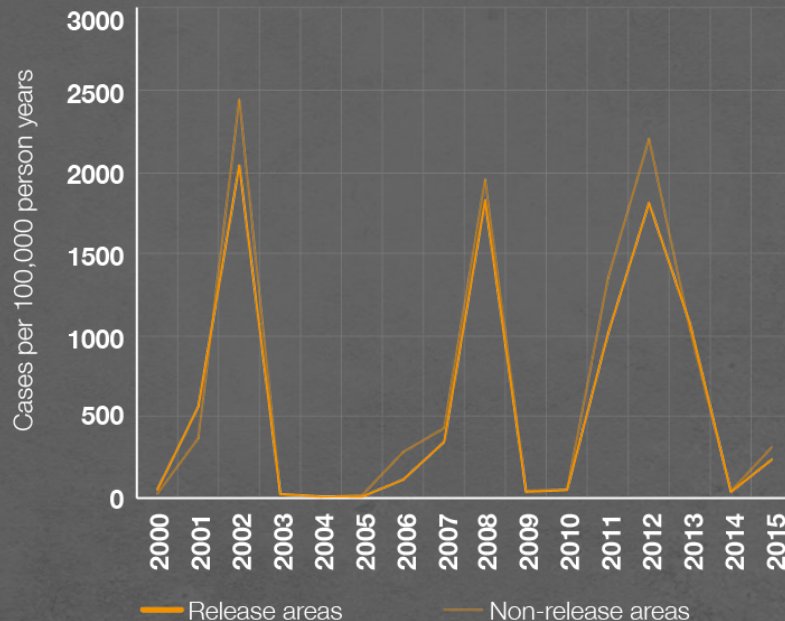




# Disease surveillance, release vs non-release areas Rio de Janeiro, Brazil

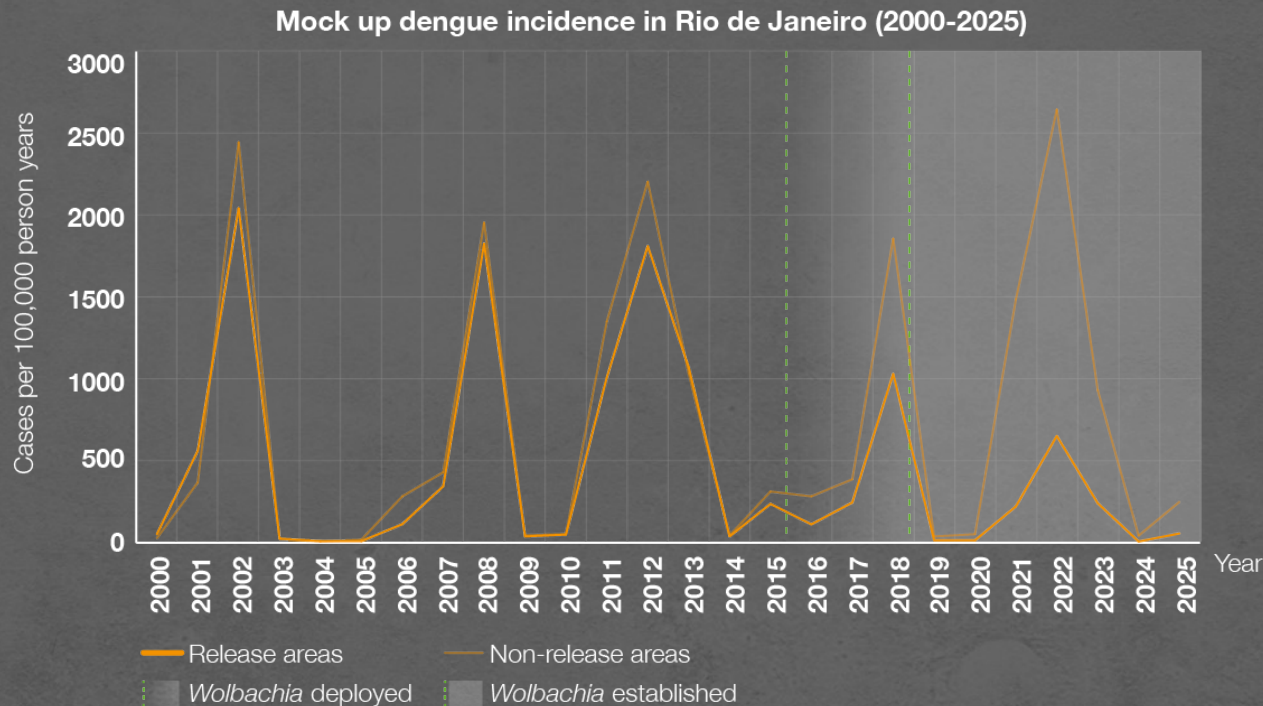
- Compare IRR between planned release & non-release areas, before & after deployment
- Planned release & non-release areas in Rio have comparable historical dengue time series
  - Incidence rate ratio pre-intervention  $\approx 1$

Dengue incidence in Rio de Janeiro (2000-2015)



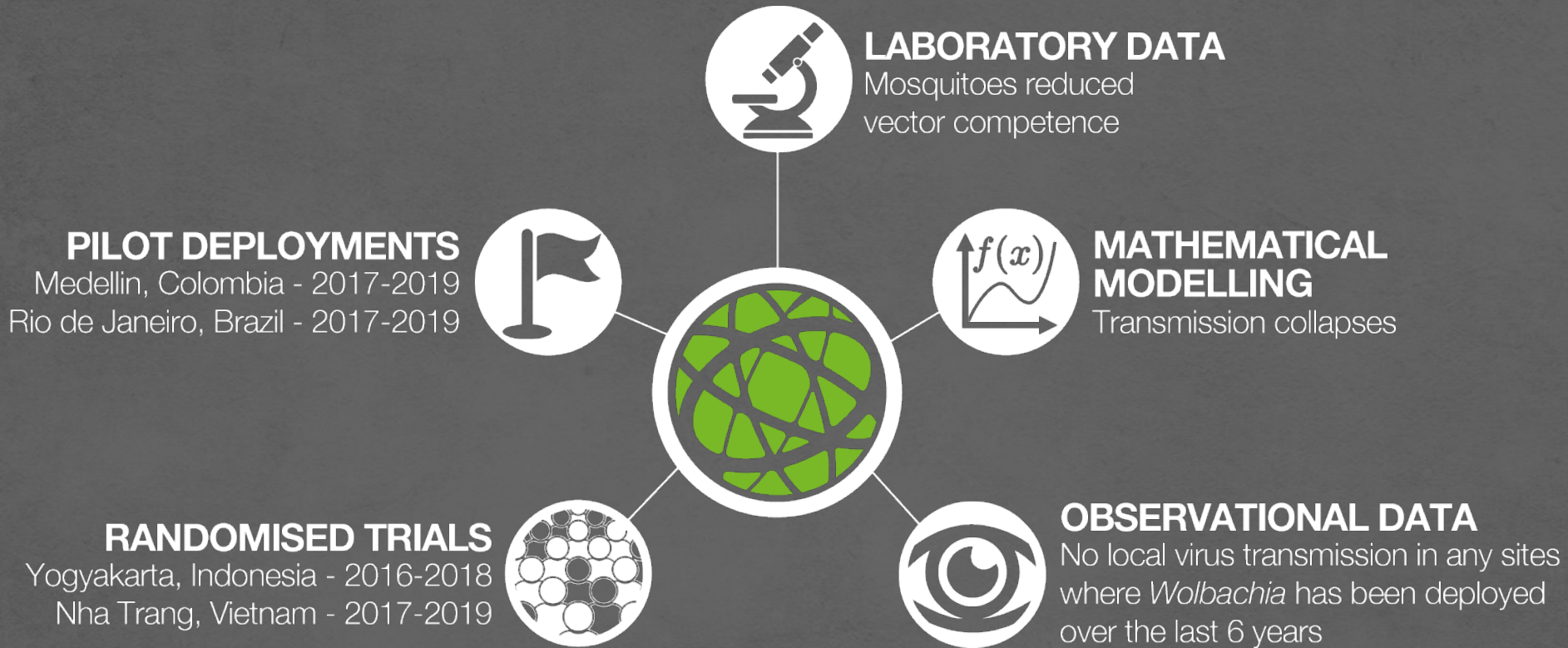
# Disease surveillance, release vs non-release areas Rio de Janeiro, Brazil

- Calculate IRRs stratified by deployment status (municipality level):  
Pre-intervention > During deployment > Post-deployment  
(annually)
- Test null hypothesis of no change in IRR with deployments  
(Mantel Haenszel Chi-Square test of homogeneity)





# Global Evidence Picture



# Future goals

- Continue to accumulate evidence of impact
- Reduce cost of deployment to less than \$1 per person
- Partner with and initiate projects in the top 20 dengue burden countries
- Establish capacity, expertise and infrastructure in those countries
- Develop and assist with National rollout strategies as required in countries according to their wishes

# An international research collaboration

## Scientific collaborators:

- Đặng Đức Anh, Vietnam National Institute of Hygiene and Epidemiology
- Adi Utarini, Universitas Gadjah Mada
- Cam Simmons, Oxford University Clinical Research Unit & Monash University
- Luciano Moreira, FIOCRUZ/Centro de Pesquisas René Rachou
- Ivan Velez & Jorge E. Osorio, PECET Universidad de Antioquia, Colombia
- Michael Turelli, University of California, Davis

## Supporters:

- Bill & Melinda Gates Foundation
- Foundation for the National Institutes of Health
- Yayasan Tahija, Indonesia
- The Wellcome Trust
- DFID
- The Australian and Queensland Governments
- The Brazilian Government
- USAID
- The Gillespie Family Foundation

