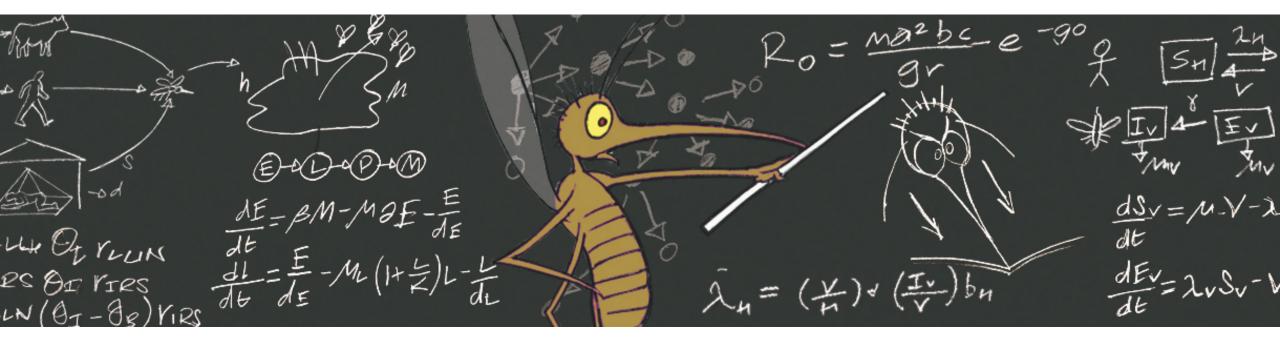
## Modeling novel genetic control strategies for *Aedes aegypti* disease vectors

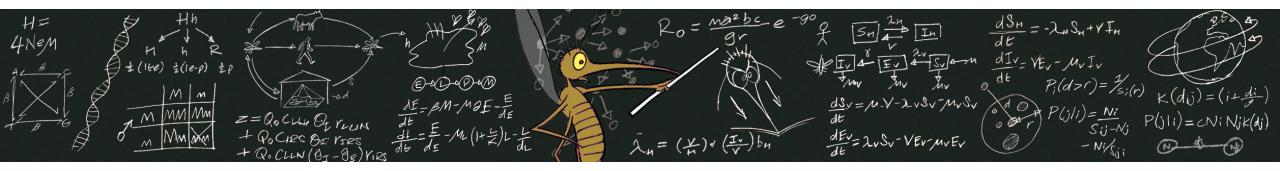
John M. Marshall, Héctor M. Sánchez C., Jared B. Bennett, Sean L. Wu, Tomás M. León, Gordana Rašić, Omar S. Akbari



John Marshall Divisions of Epidemiology & Biostatistics School of Public Health University of California, Berkeley john.marshall@berkeley.edu

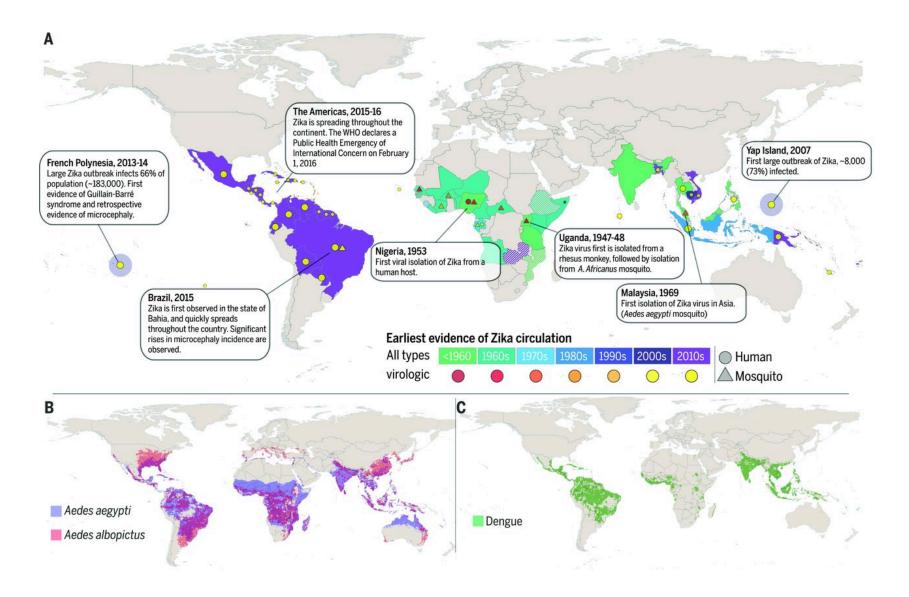
Berkeley Public Health

## Talk outline



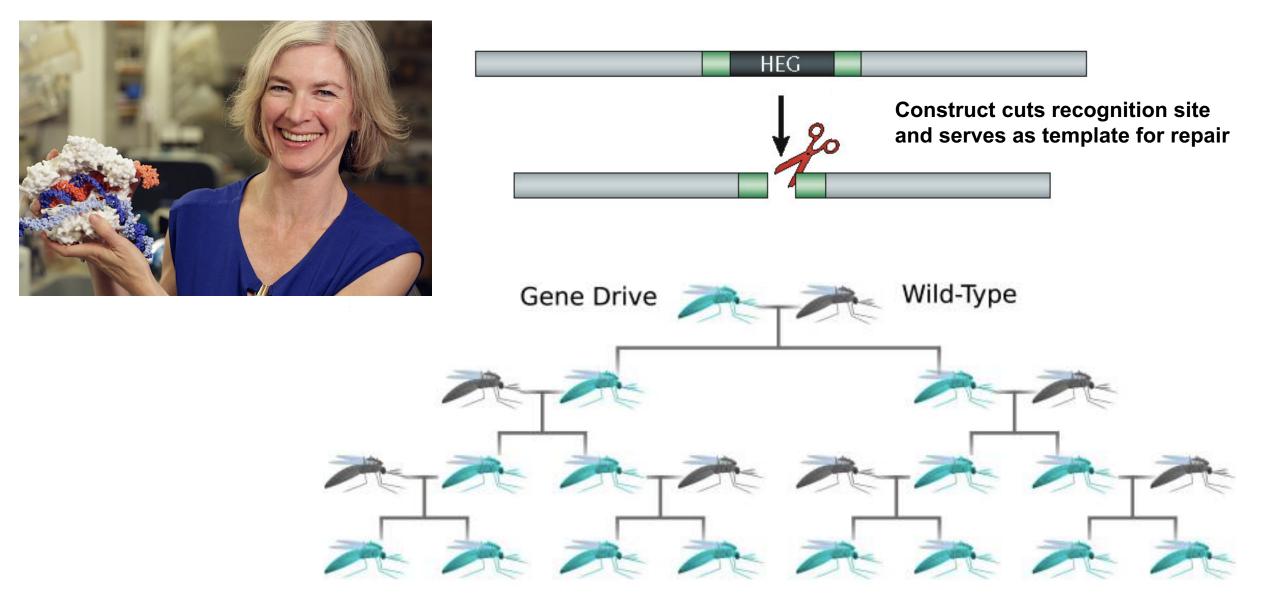
- 1. Why genetic control?
- 2. Modeling framework (Mosquito Gene Drive Explorer)
- 3. Novel genetic control technologies:
  - i. Chromosomal translocations as threshold-dependent systems
  - ii. CRISPR-based split drive as a self-limiting system
  - iii. Precision-guided sterile insect technique (pgSIT)
- 4. Role of these technologies in Ae. aegypti control

#### Arboviruses on the rise worldwide



• Lessler J, Chaisson LH, Kucirka LM, Bi Q, Grantz K et al. (2016) Science 353: aaf8160

### Discovery of CRISPR, applied to gene drive



#### DARPA Safe Genes Program & Team CA

Genetic

Remediation

**Control of Gene** Editing

Enable temporal, spatial, and reversible control of gene editors

Inhibit unwanted gene editing activity

Countermeasures

and Prophylaxis

Remove engineered genes from environments to return to baseline



Omar Akbari





Anthony James

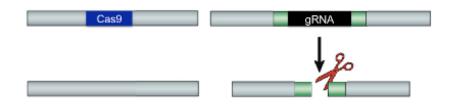
Ethan Bier





**Cinnamon Bloss** 







John Marshall





Sergey Kryazhimskiy





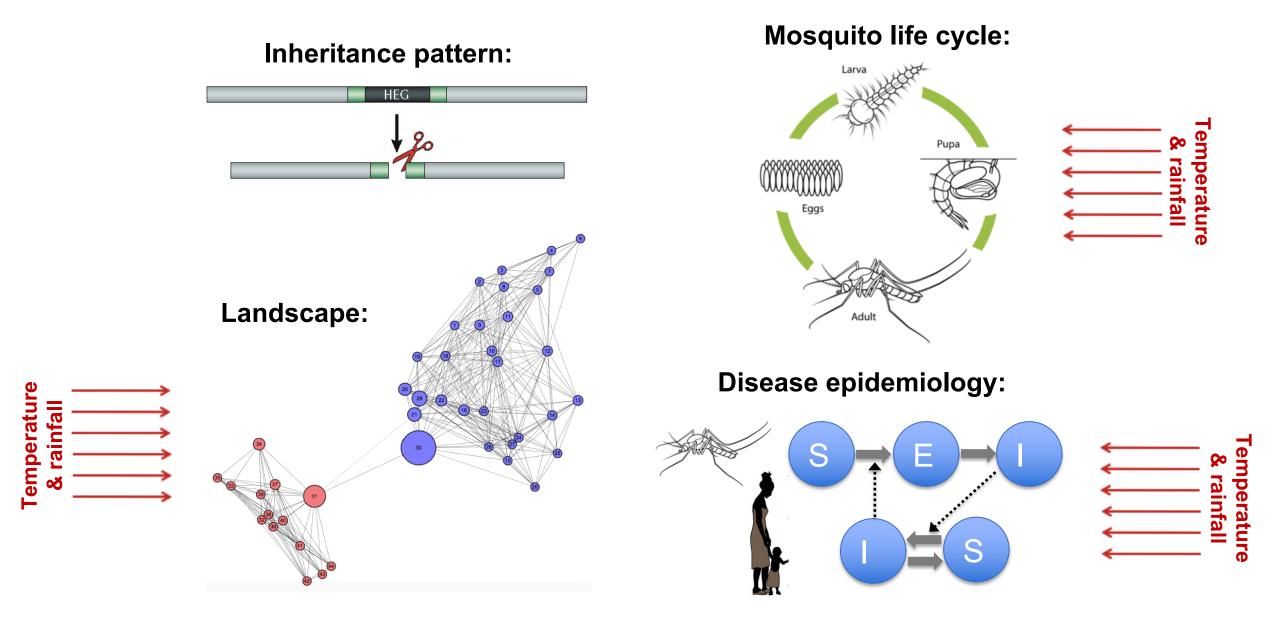






**Greg Lanzaro** 

#### Mosquito Gene Drive Explorer (MGDrivE)



## Mosquito Gene Drive Explorer (MGDrivE)

#### MGDrivE

Mosquitos + Tensors + Genetics + CS + Networks + Math + Coffee

 View

 Releases List

 Browse

 Documentation

 View on

 Youtube

 Fork on

 GitHub

 Download

 ZIP File

Download TAR Ball

#### Developed in John Marshall's Lab by:

-Lead: Héctor M. Sánchez C. -Core Dev: Sean L. Wu,Jared Bennett -Spatial Analysis: Biyonka Liang, Sarafina Smith, Sabrina Wong -Movement Kernels: Partow Imani

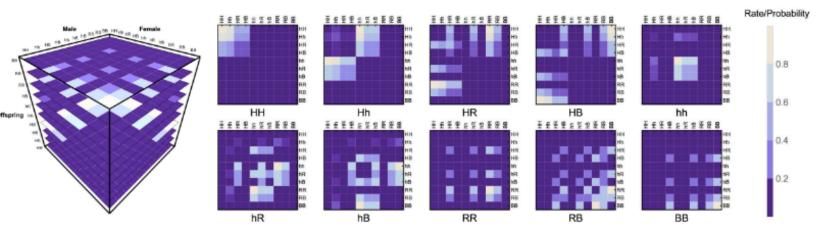
.

...and, of course, our PI; John M, Marshall!

# Mosquito Gene Drive Explorer

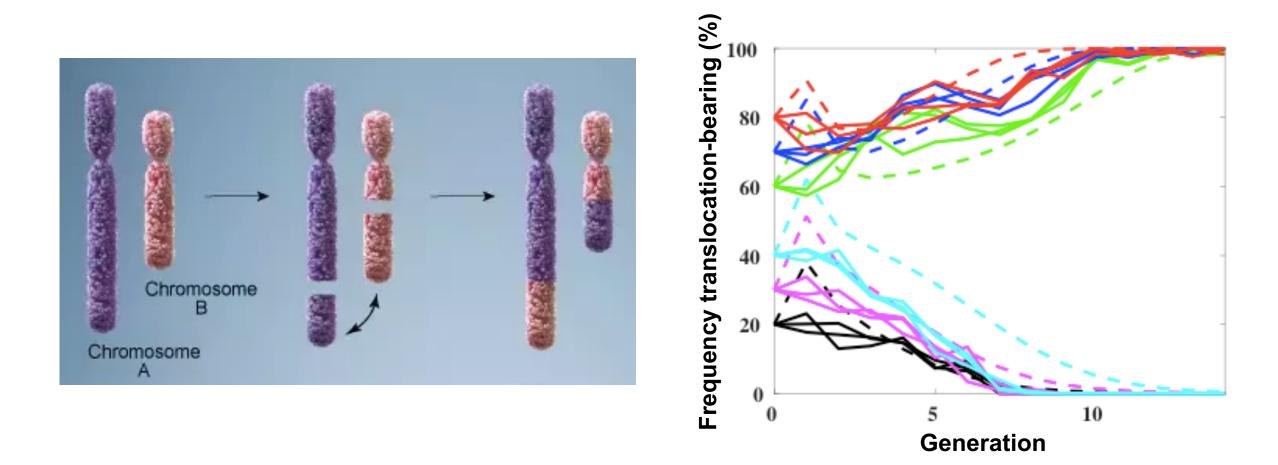
#### **Brief Description**

**MGDrivE** is a framework designed to serve as a testbed in which gene-drive releases for mosquito-borne diseases control can be tested. It is being developed to accommodate various mosquito-specific gene drive systems within a population dynamics model that allows migration of individuals between nodes in a spatial landscape.



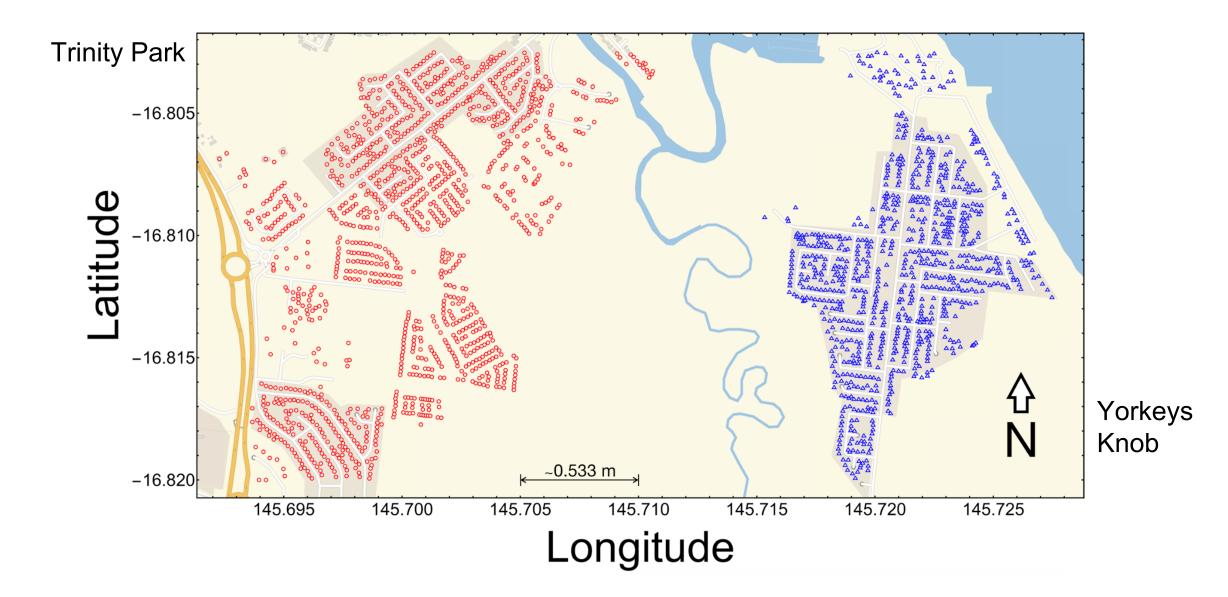
Sánchez HM, Wu SL, Bennett JB, Marshall JM (2019) Methods in Ecology and Evolution

#### Reciprocal chromosomal translocations



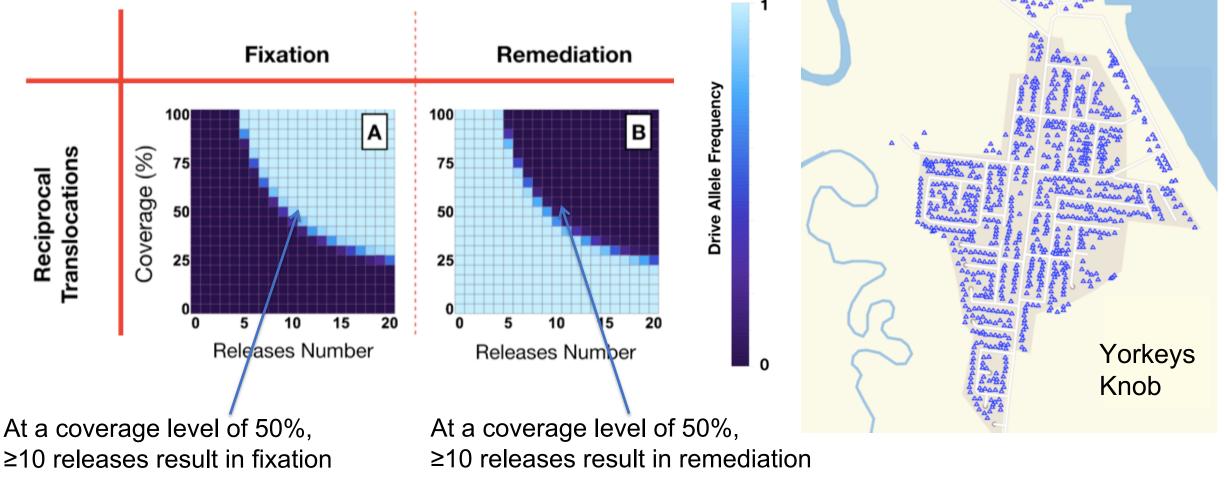
• Buchman A, Ivy T, Marshall JM, Akbari OS, Hay BA (2018) ACS Synthetic Biology

#### Simulations at Yorkeys Knob, Queensland



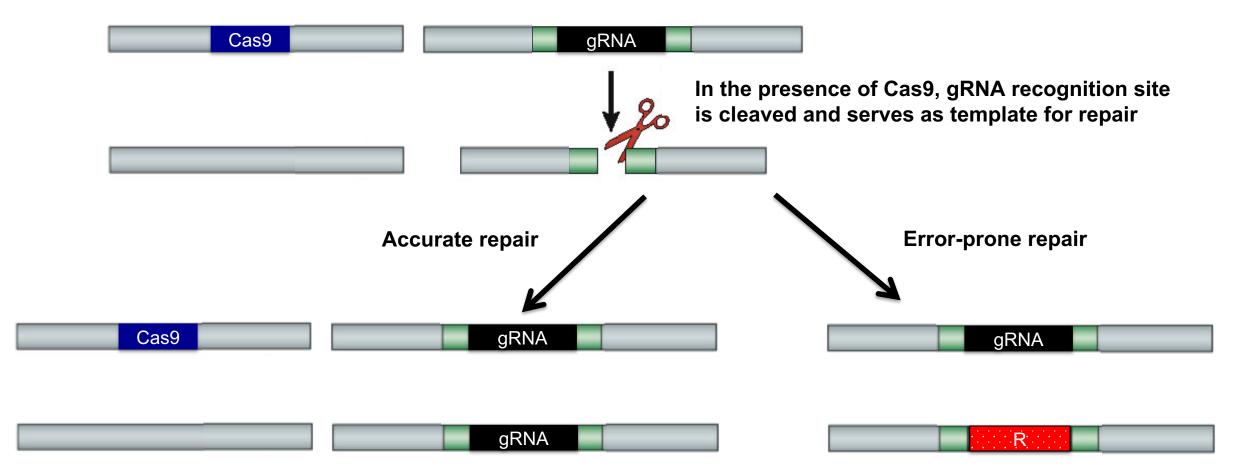
## Fixation & remediation of translocations

- Data suggests there are ~15 adult Ae. aegypti per household
- Weekly releases of 20 adult males having the translocation
- Vary household coverage & number of releases



• Sánchez HM, Bennett JB, Wu SL, Rasic GL, Akbari OS, Marshall JM (2020) BMC Biology

#### **CRISPR-based split drive**

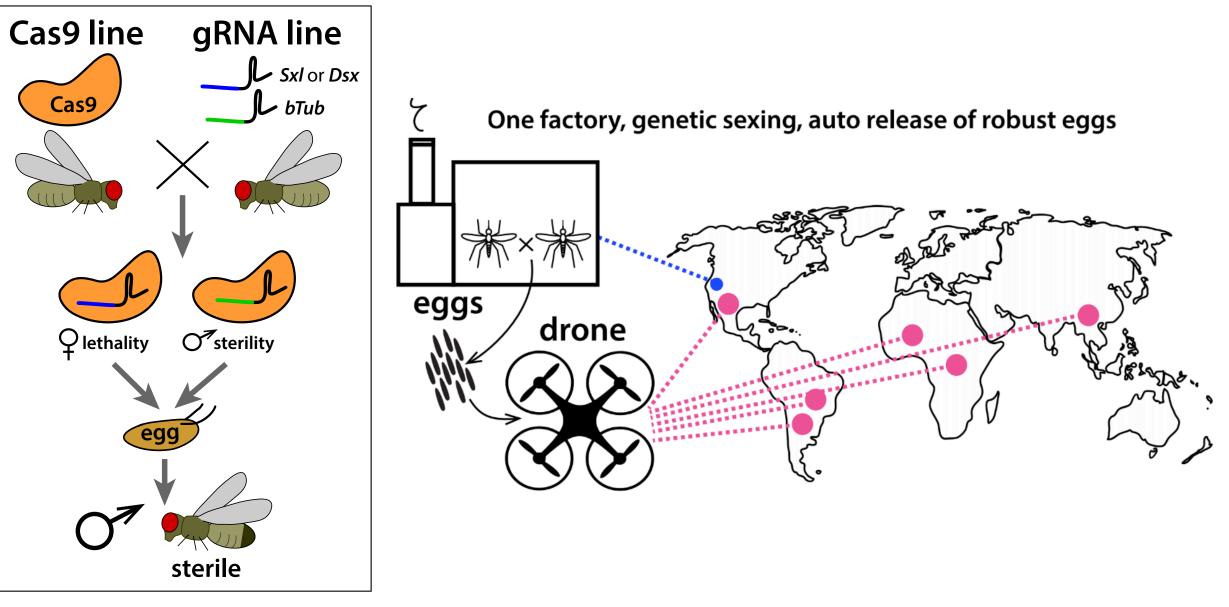


• Li M, Yang T, Kandul N, Biu M, Gamez S, Bennett JB, Sánchez HM, ..., Marshall JM, Akbari OS (2020) eLife

#### Simulations at Yorkeys Knob, Queensland

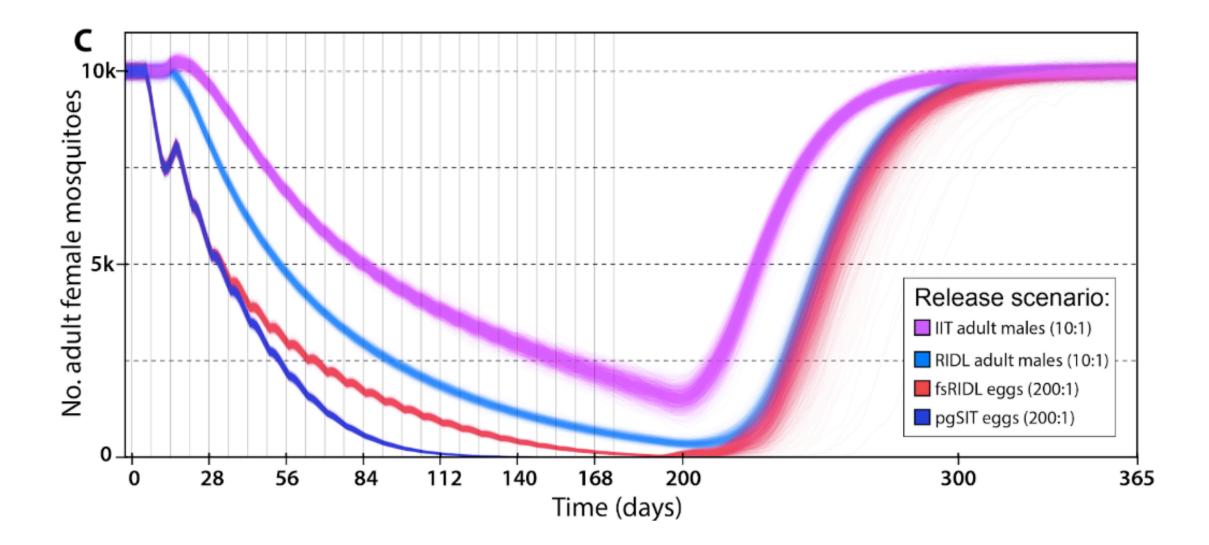


## Precision-guided SIT (pgSIT)



• Kandul NP, Liu J, Sánchez HM, Wu SL, Marshall JM, Akbari OS (2019) Nature Communications

## Modeling pgSIT in Ae. aegypti

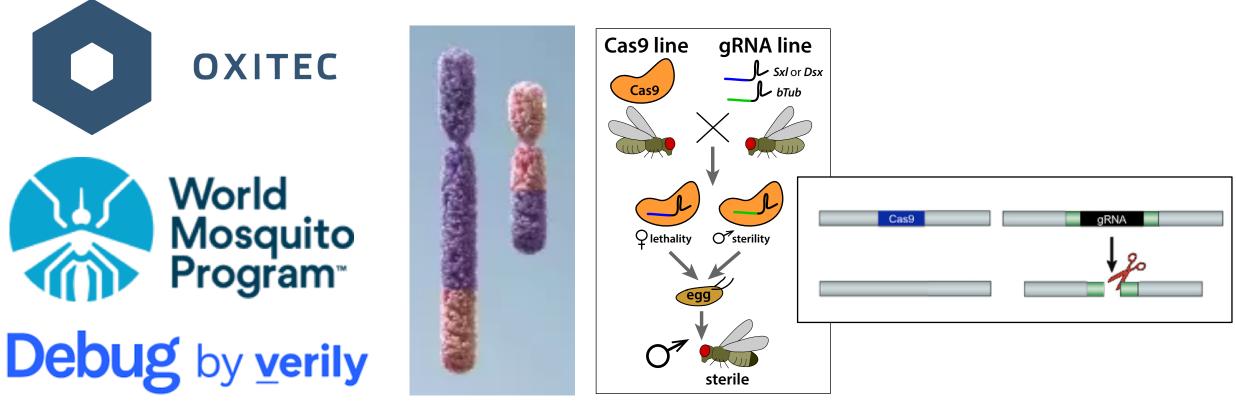


• Kandul NP, Liu J, Sánchez HM, Wu SL, Marshall JM, Akbari OS (2019) Nature Communications

#### Conclusion



- CRISPR-based strategies offers for more tools for the tool kit.
- Ae. aegypti control has been marked by technological innovations through Wolbachia transfection & transgenesis.
- Translocations may offer an opportunity to introduce a diseaserefractory into an *Ae. aegypti* population gene locally.
- Split drive may be an effective precursor to linked drive.
- pgSIT may be an effective suppression system in some locations.



## Acknowledgements

#### LAB MEMBERS:

• Héctor M. Sánchez C. o Tomás León

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• Ameek Bindra

• Chris De Leon

- Sean L. Wu
- Yogita Sharma
- Sanjay Lamba
- Gordana Rašić
- Darpa Anireddy
- Ashley Zhang
- Daniel López
- Priscilla Zhang

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- Akbari Lab @ UC Riverside
- Lanzaro Lab @ UC Davis
- James Lab @ UC Irvine
- Bier Lab @ UCSD
- Hay Lab @ Caltech
- Malaria Elimination Initiative @ UCSF
- Prof David Smith @ IHME, UW
- Dr Samson Kiware @ Ifakara Health Institute
- School of Public Health @ UC Berkeley

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#### Innovative Genomics Institute

**UC IRVINE** MALARIA INITIATIVE





BILL& MELINDA

The Parker FOUNDATION GATES foundation

