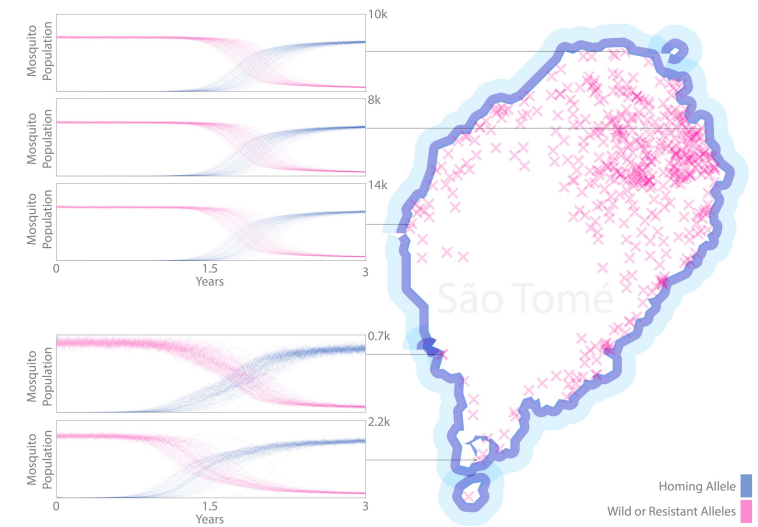
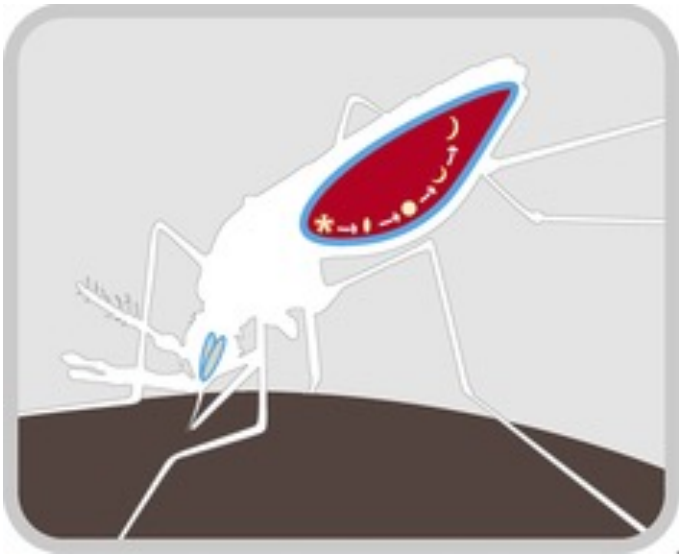


Modeling priorities as gene drive mosquito projects transition from lab to field

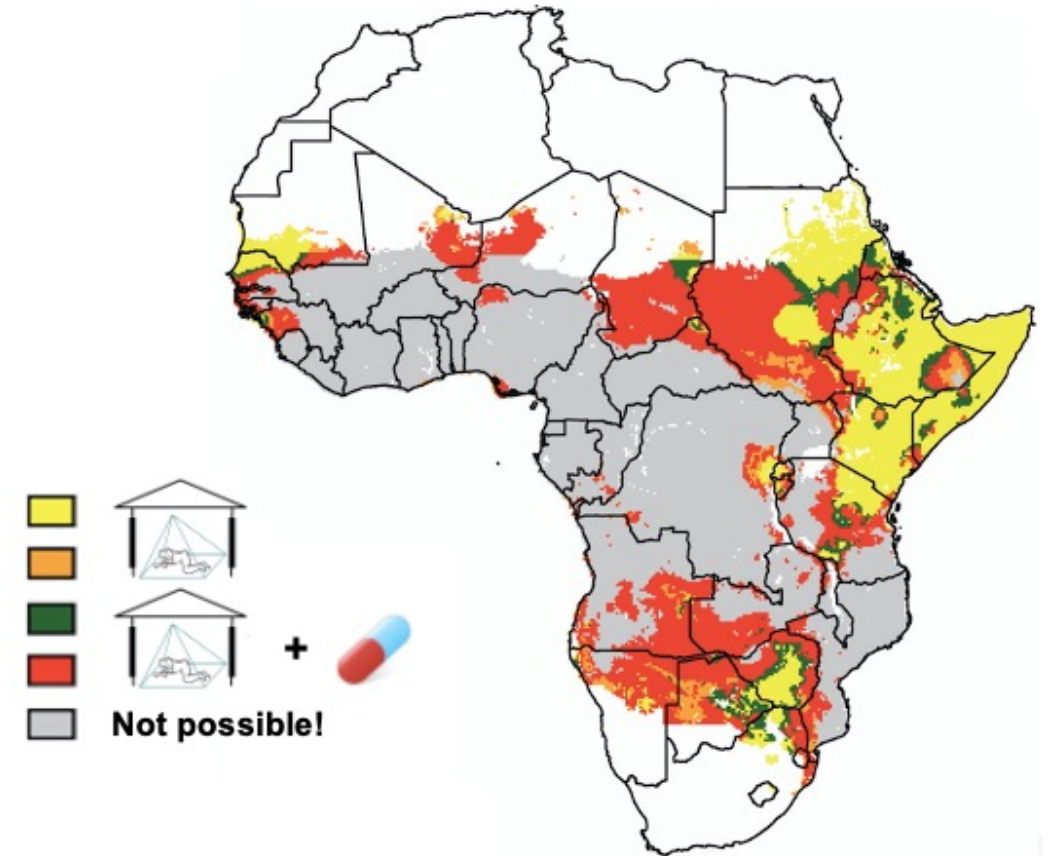
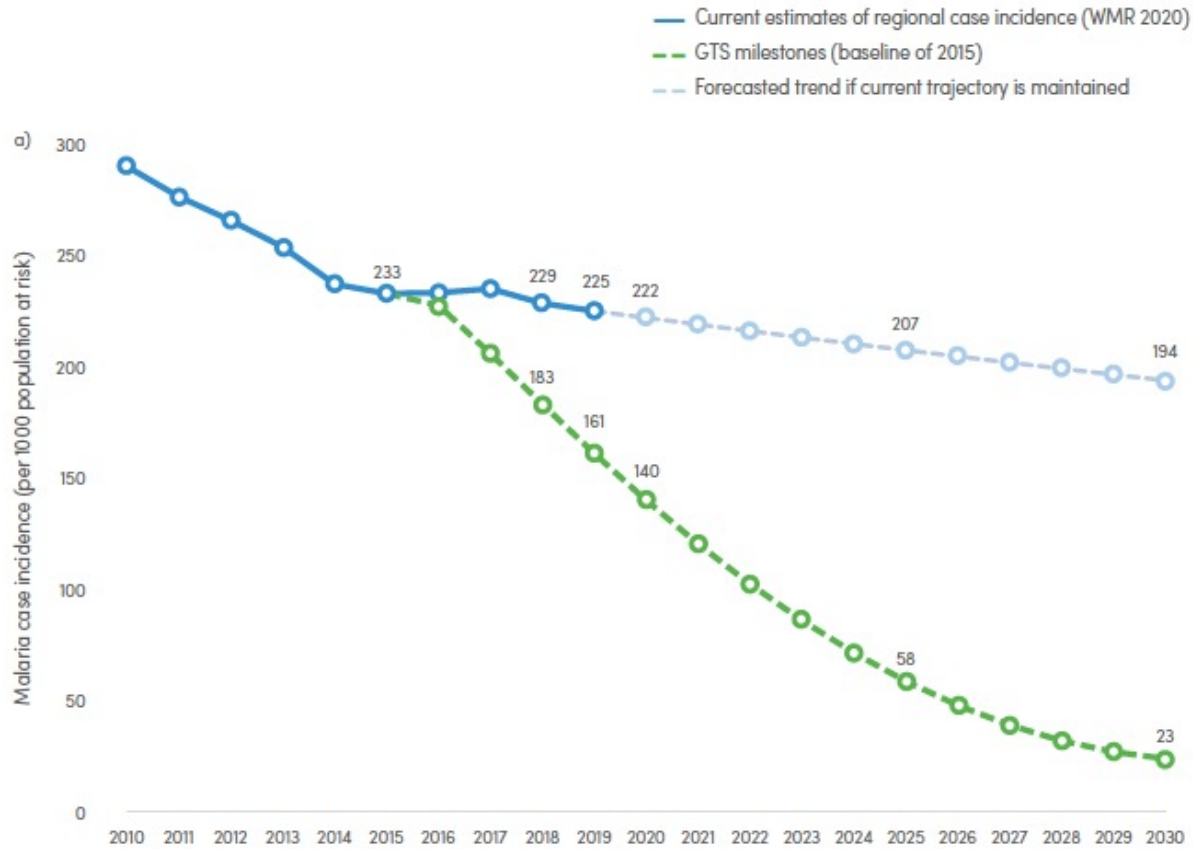
John M. Marshall, Héctor M. Sánchez C., Ace R. North



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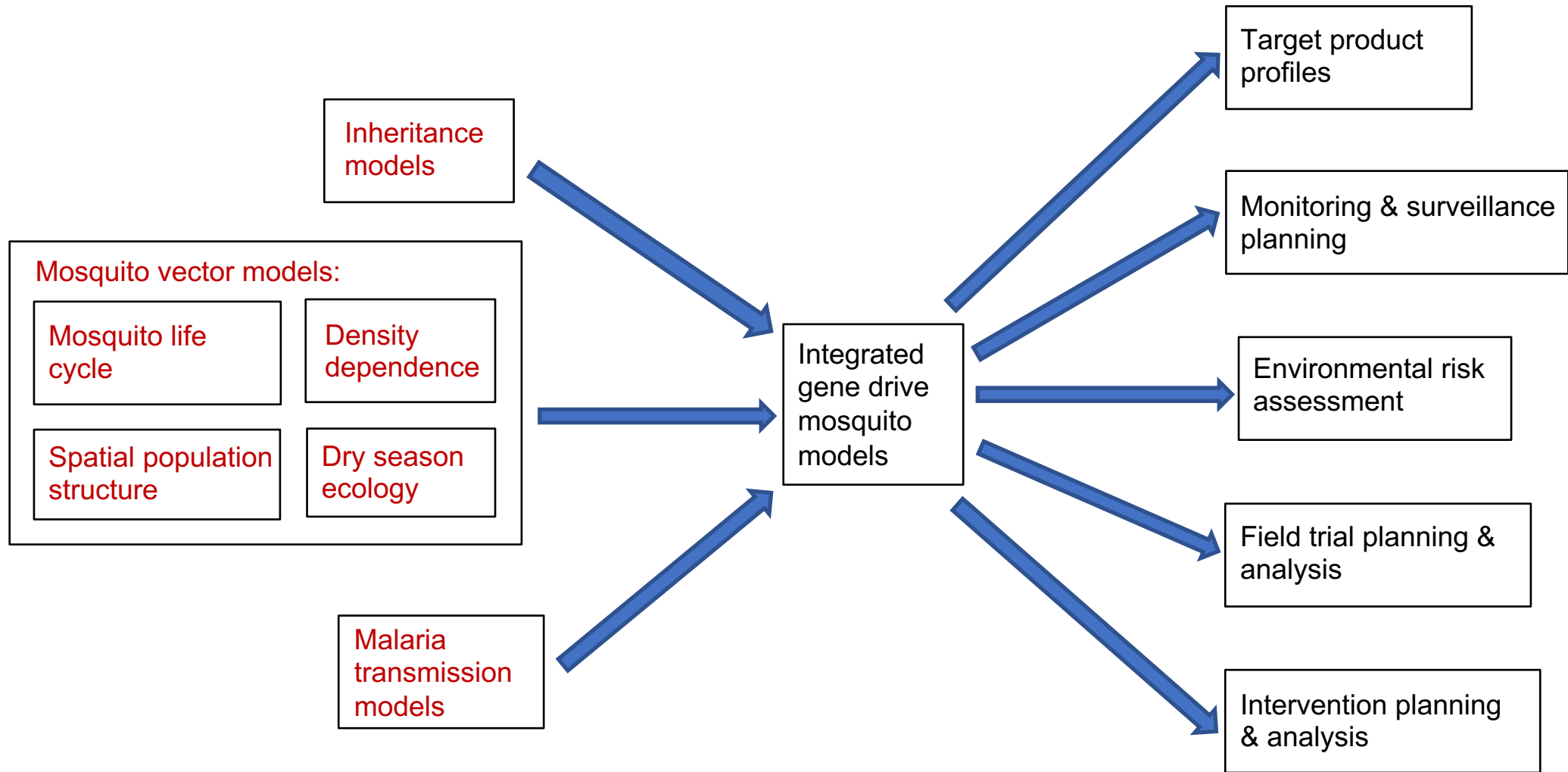
Malaria burden stagnating, elimination difficult



GTS: Global technical strategy for malaria 2016–2030; WHO: World Health Organization; WMR: world malaria report.

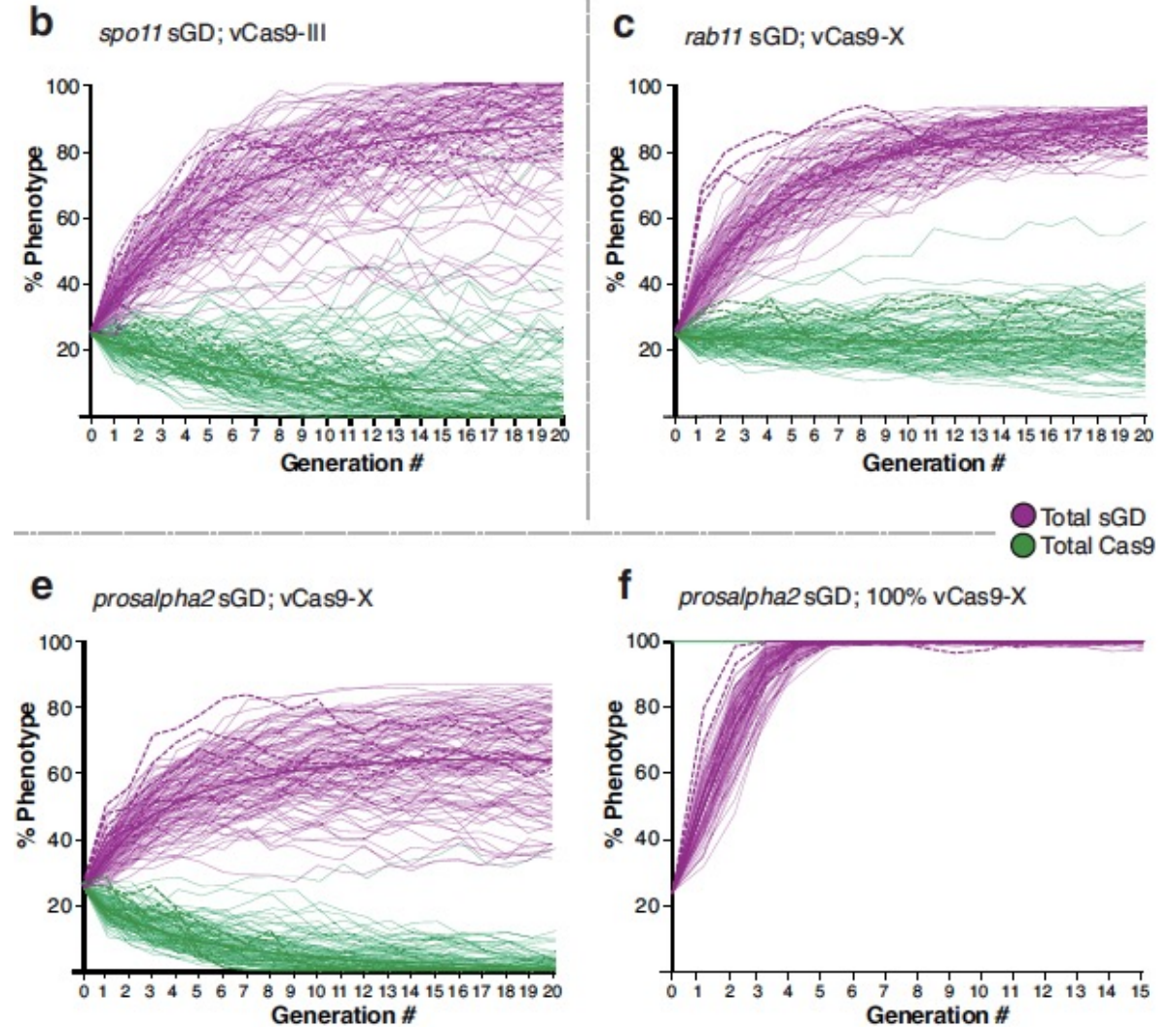
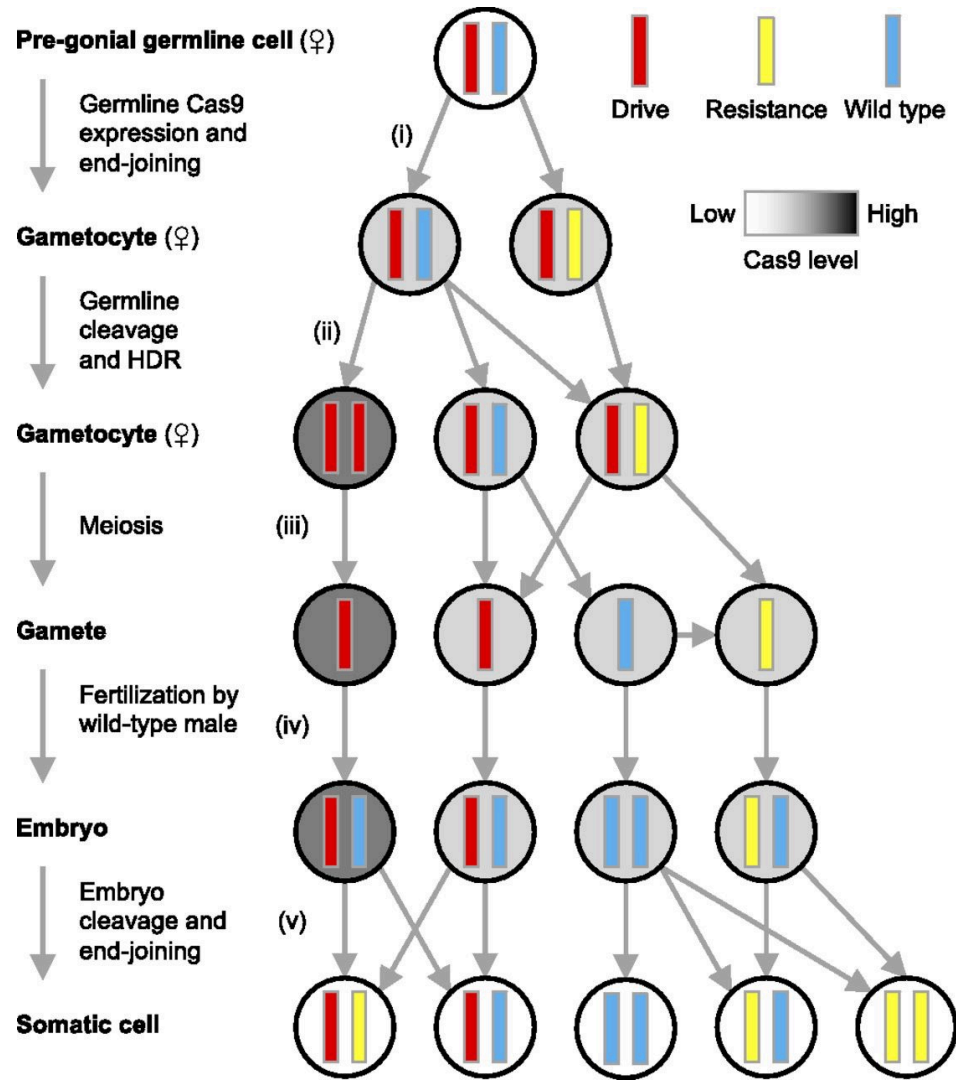
- World Health Organization (2021) World Malaria Report 2020
- Walker PGT, Griffin JT, Ferguson NM, Ghani AC (2016) Lancet Global Health

From lab to field: Model building priorities



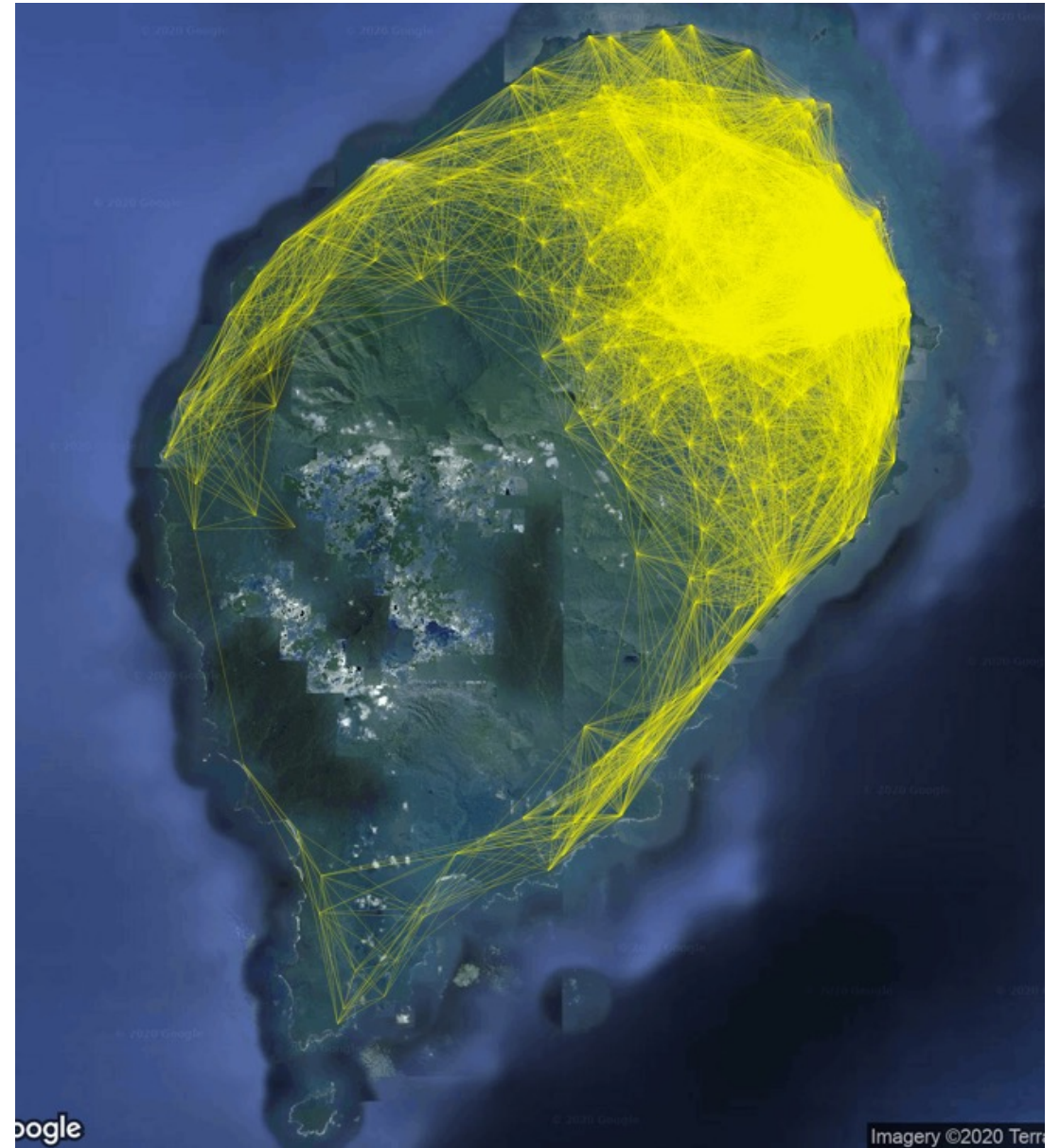
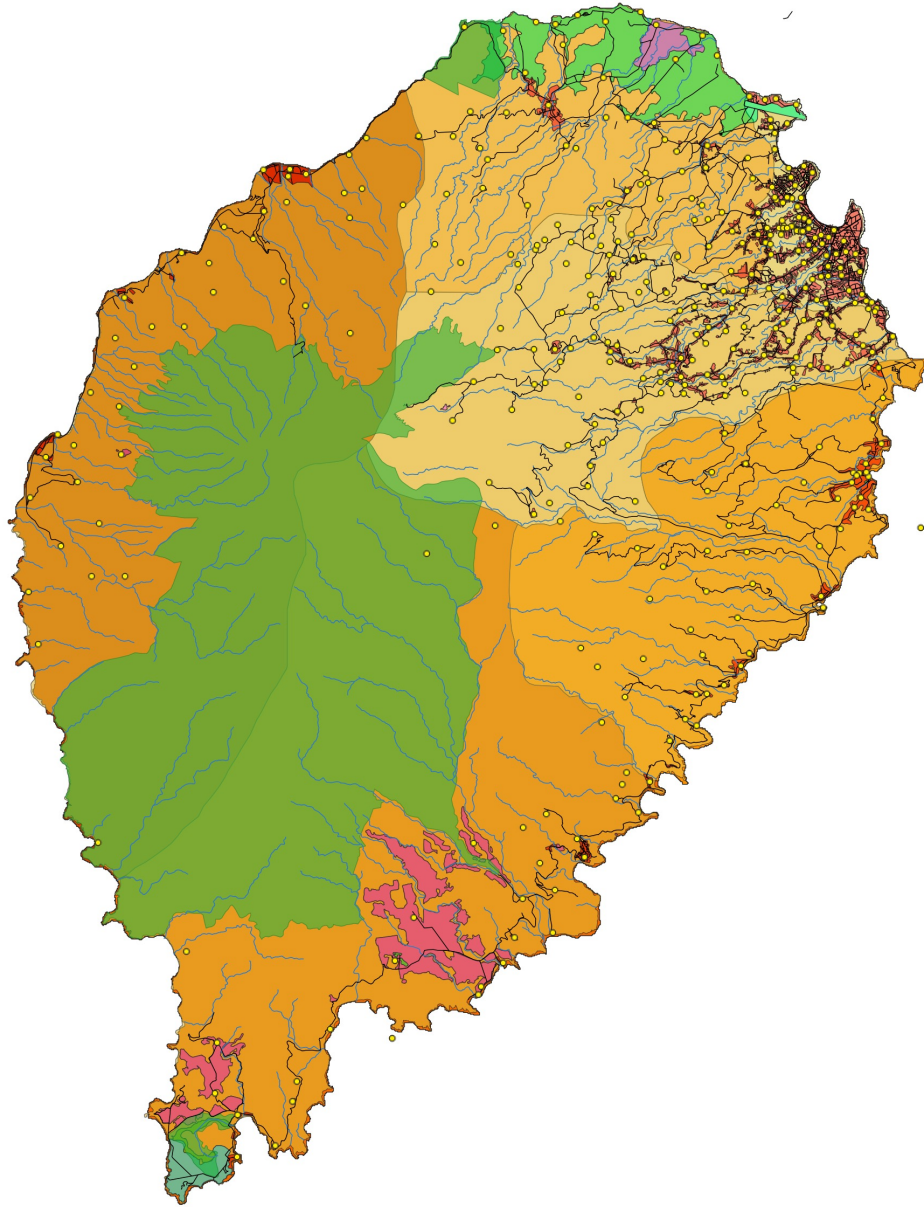
- Marshall JM, North AR (in press) Gene Drives and the Malaria Eradication Agenda

Understanding resistance alleles & Cas9 dynamics



- Champer J, Liu J, Oh SY, Reeves R *et al.* (2018) Proc Natl Acad Sci USA
- Terradas G, Buchman AB, Bennett JB *et al.* (2018) Nature Communications

Spatial population structure

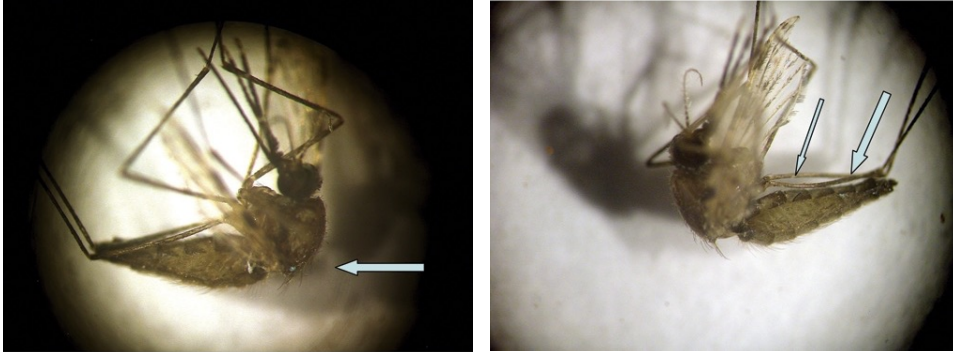


Spatial population structure

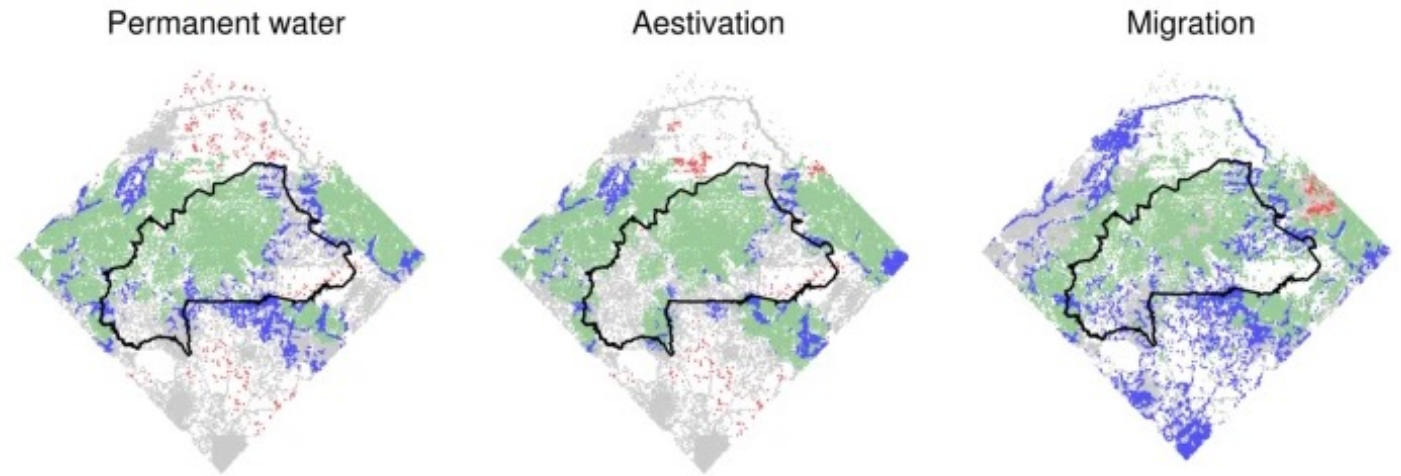
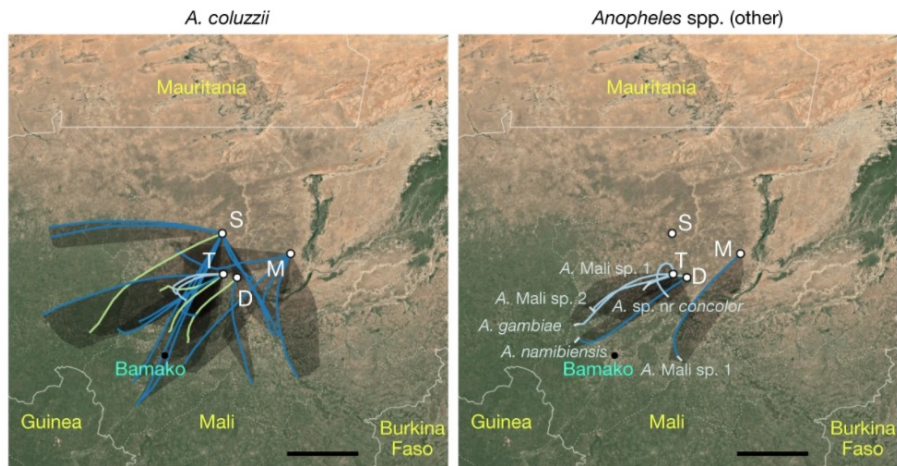


Dry season ecology

Aestivating female caught ~7 months after being marked:

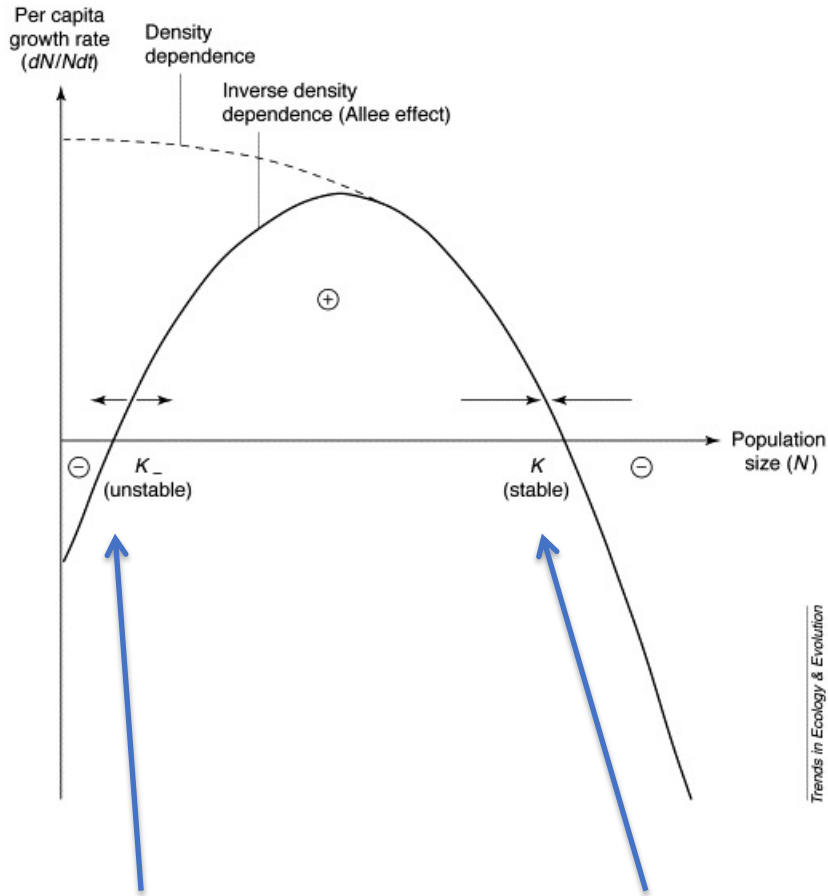


Trajectories of flights for *Anopheles* capture events:



- North AR, Burt A, Godfray HCJ (2019) BMC Biology
- Lehmann T, Dao A, Yaro AS, Adamou A *et al.* (2010) Am J Trop Med Hyg
 - Huestis DL, Dao A, Diallo M, Sanogo ZL *et al.* (2019) Nature

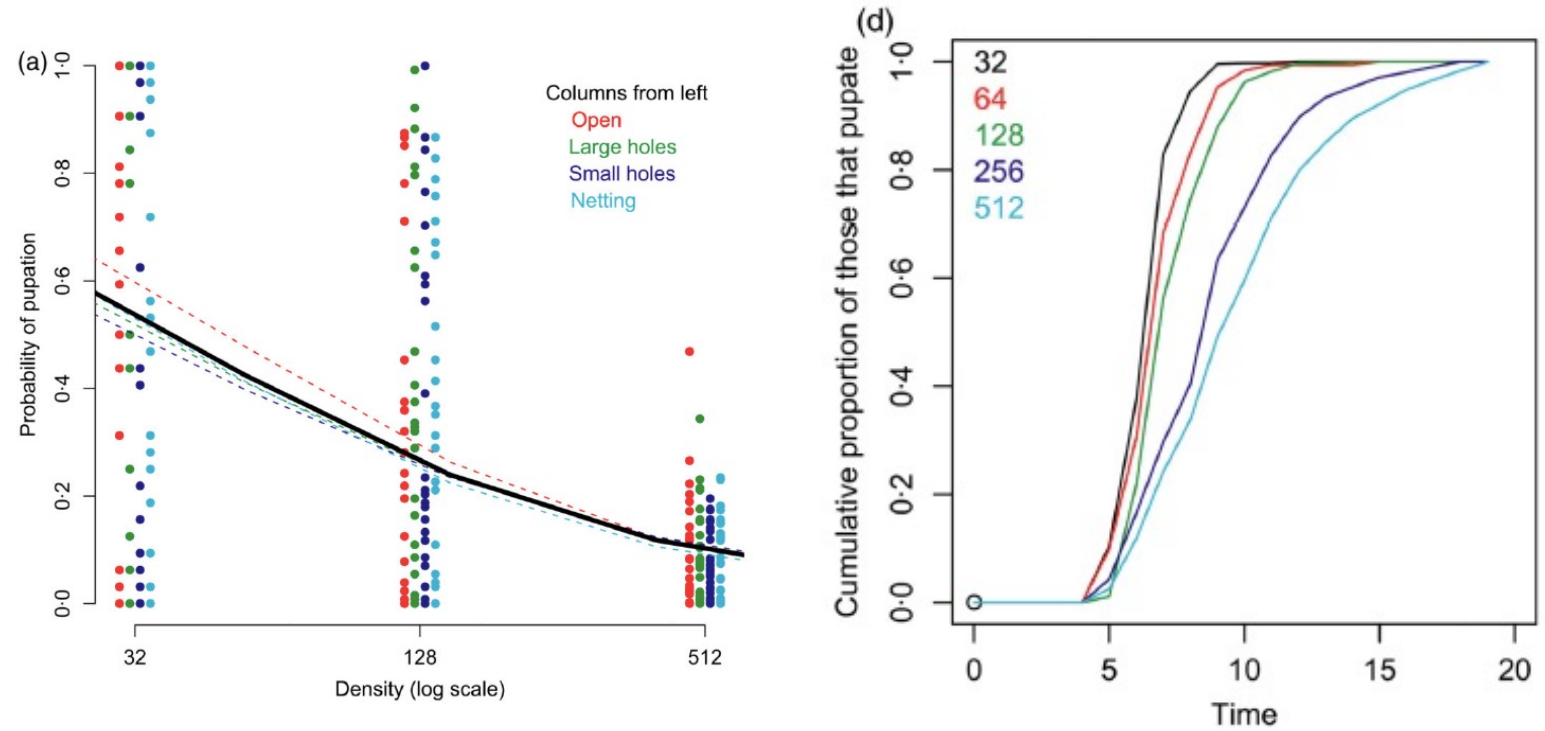
Density dependence



More difficult to form swarms & find mates at low densities

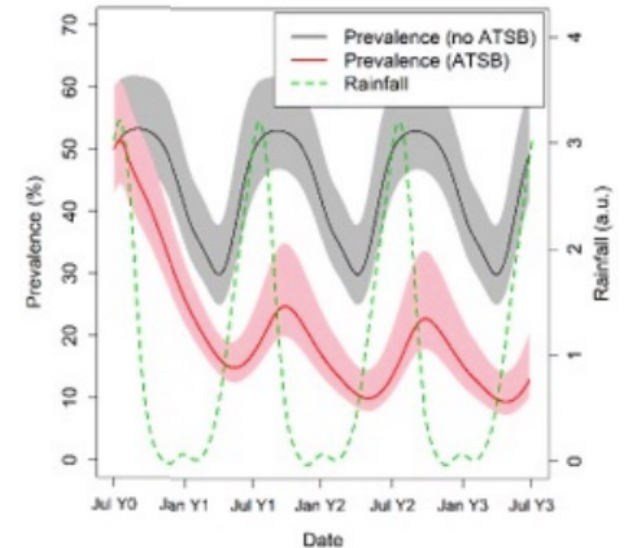
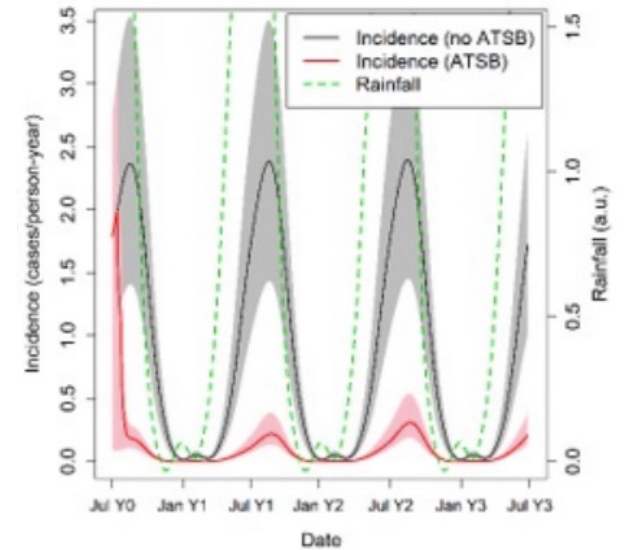
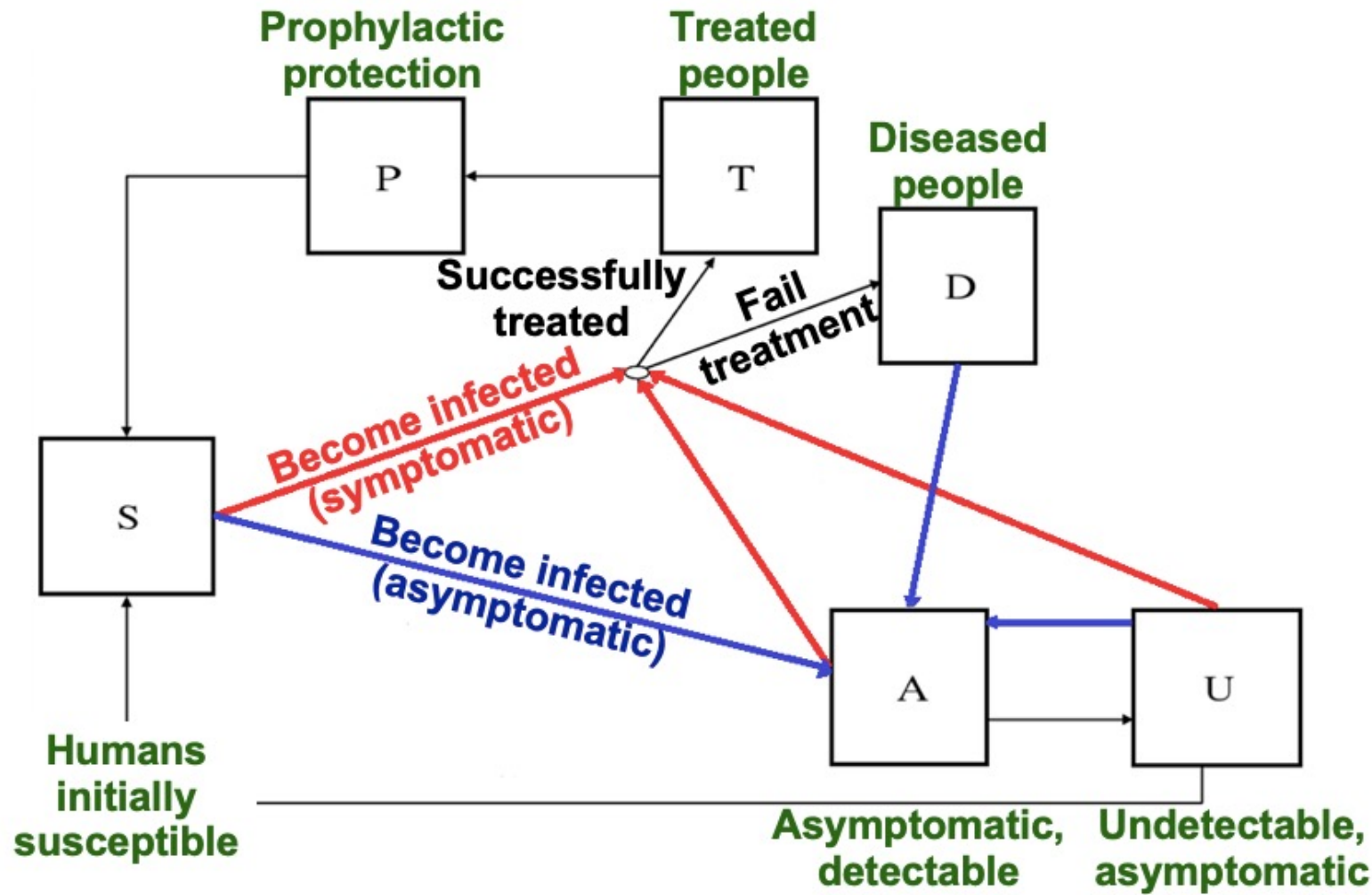
Crowding of larvae leads to increased competition for space & resources

Higher larval density leads to less pupation & longer time to pupation:



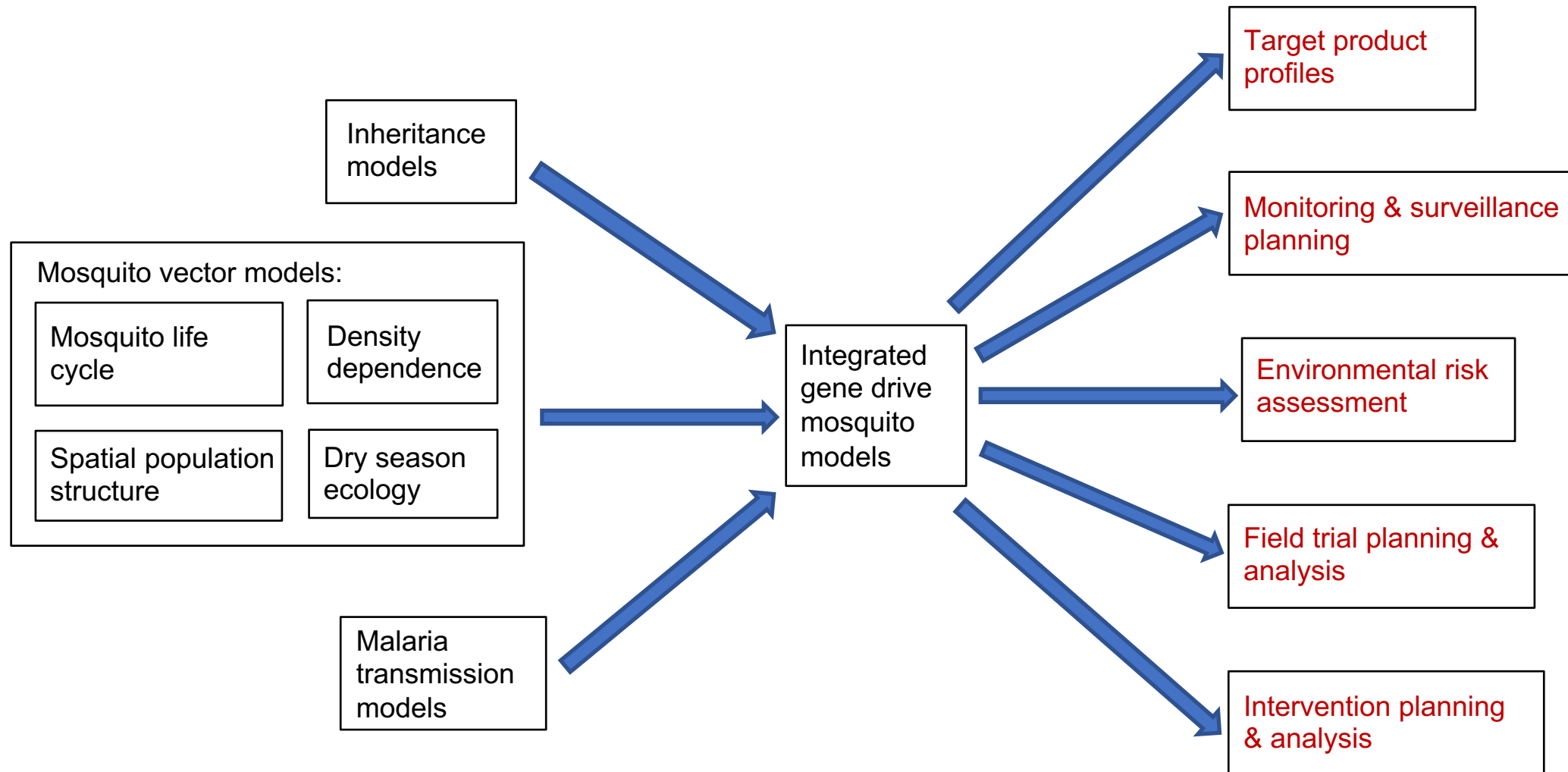
- Courchamp F, Clutton-Brock T, Grenfell B (1999) Trends Ecol Evol
- Muriu S, Coulson T, Mbogo CM, Godfray HCJ (2013) J Anim Ecol

Malaria transmission models



- Griffin JT, Hollingsworth TD, Okell LC, Churcher TS *et al.* (2010) PLoS Medicine
- Fraser KJ, Mwandigha L, Traore S, Traore M, Dombia S *et al.* (2021) Malaria Journal

From lab to field: Model application priorities



- Marshall JM, North AR (in press) Gene Drives and the Malaria Eradication Agenda

Target product profiles

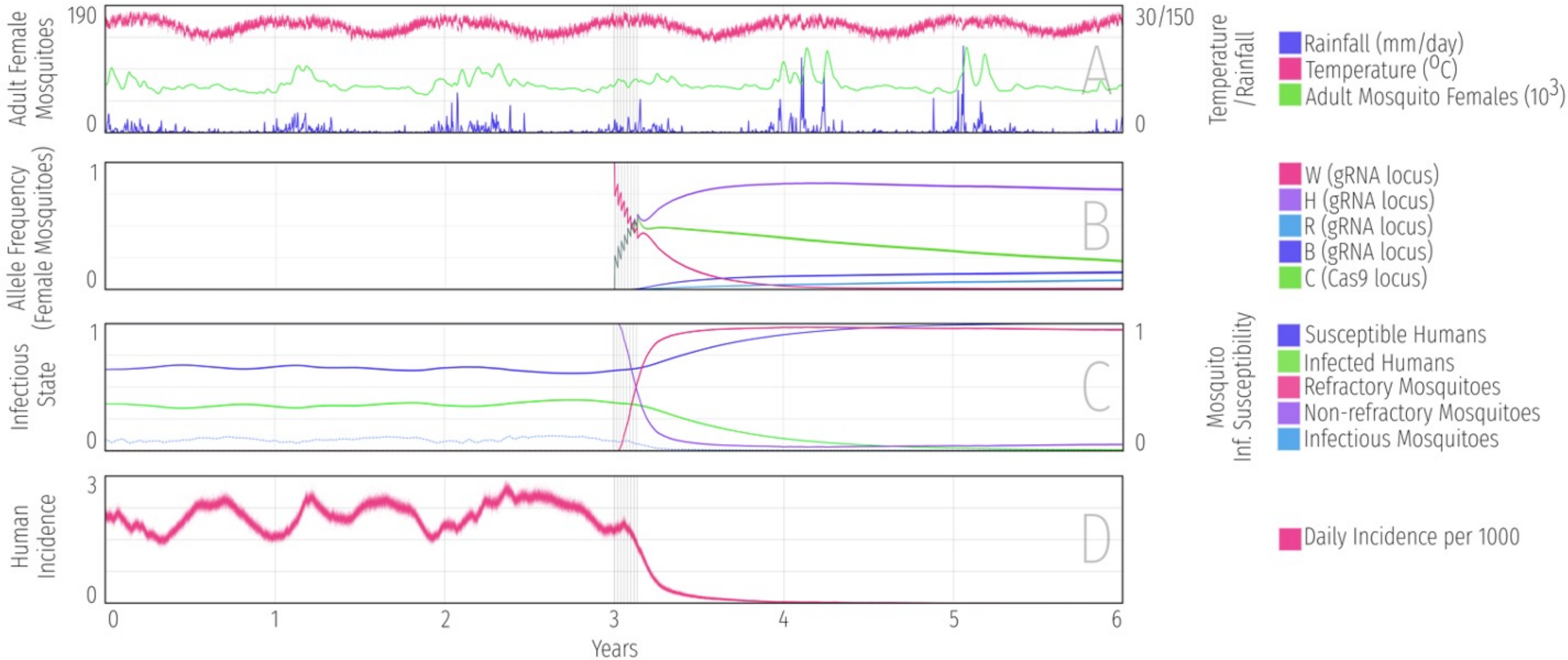
TABLE 1. PROPOSED CRITERIA FOR PREFERRED PRODUCT CHARACTERISTICS OF GENE DRIVE-MODIFIED MOSQUITOES

-
1. Indication—target vector and parasite species
 2. Epidemiological efficacy goal—reduction in clinical incidence of malaria
 3. Entomological efficacy goal—reduction in vectorial capacity commensurate with epidemiological protection goal
 4. Duration of protection—time over which the epidemiological efficacy impact will be evident
 5. Time to impact—time required for the product to achieve epidemiological and entomological goals over a specified area
 6. Safety for human health and the environment
-

TABLE 2. PROPOSED EFFICACY PARAMETERS FOR TARGET PRODUCT PROFILES OF GENE DRIVE-MODIFIED MOSQUITOES

-
1. Homing rate—predictor of rate of spread and time to impact
 - a. Deviation from Mendelian expectation of inheritance in cages
 2. Life history and reproductive success—predictor of rate of spread and time to impact
 - a. Adult longevity
 - b. Adult biting rate
 - c. Mating efficiency
 - d. Egg clutch size and hatching rate
 - e. Sex ratio of progeny
 - f. Development and mortality rate at different life stages
 3. Construct functionality—predictor of entomological and epidemiological efficacy
 - a. Population suppression—population decline in cages
 - b. Population replacement—reduction in carriage of the target parasite species
 4. Functional resistance—predictor of duration of protection
 - a. Population suppression and replacement—functional resistance to the drive
 - b. Population replacement—parasite resistance to the effector(s)
-

Target product profile:



- Wu SL, Bennett JB, Sánchez HM, Dolgert AJ, León TM, Marshall JM (2021) PLoS Computational Biology

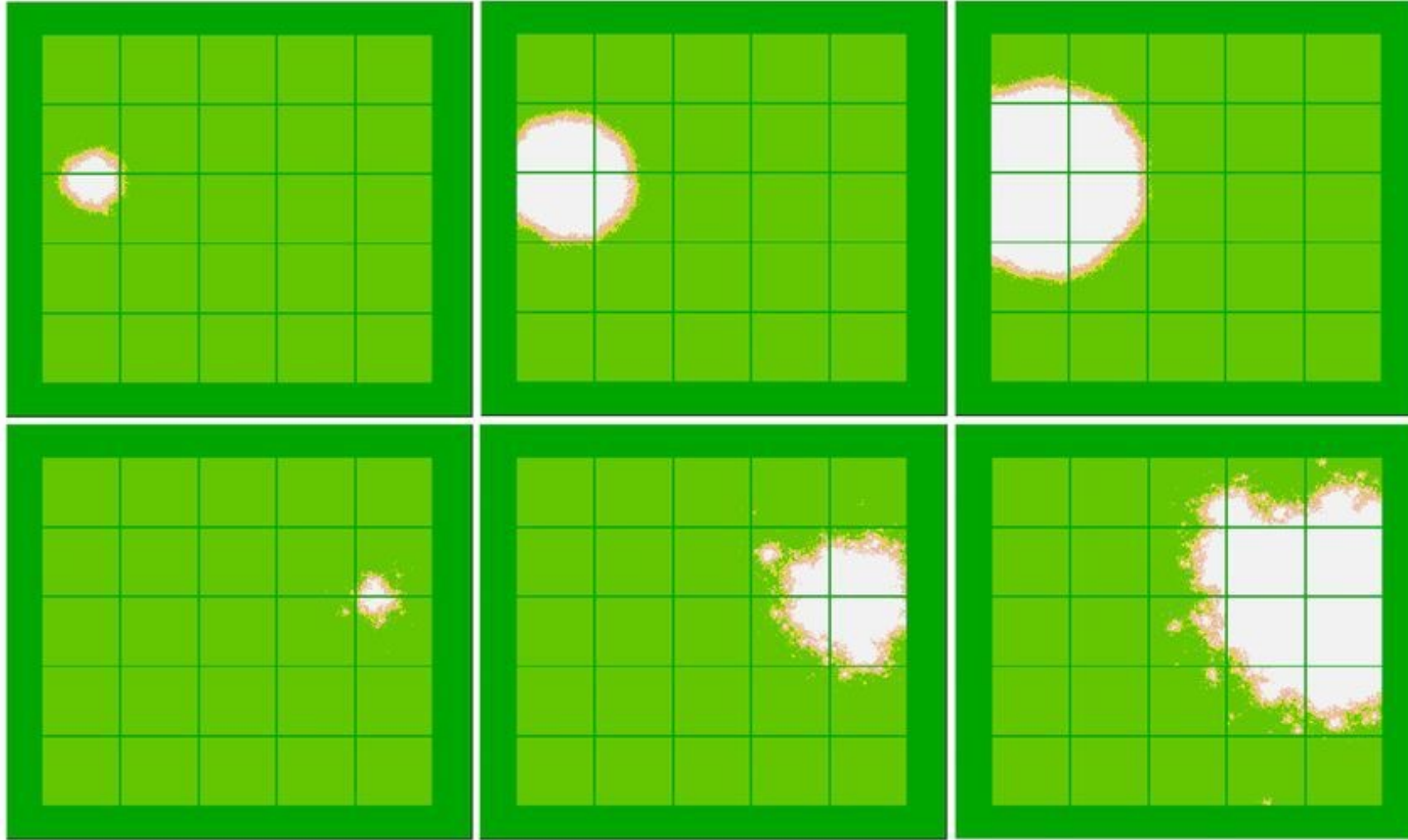
Monitoring needs: Heterogeneity in spread

Wolbachia
frequency in
Yorkeys Knob
& Gordonvale:



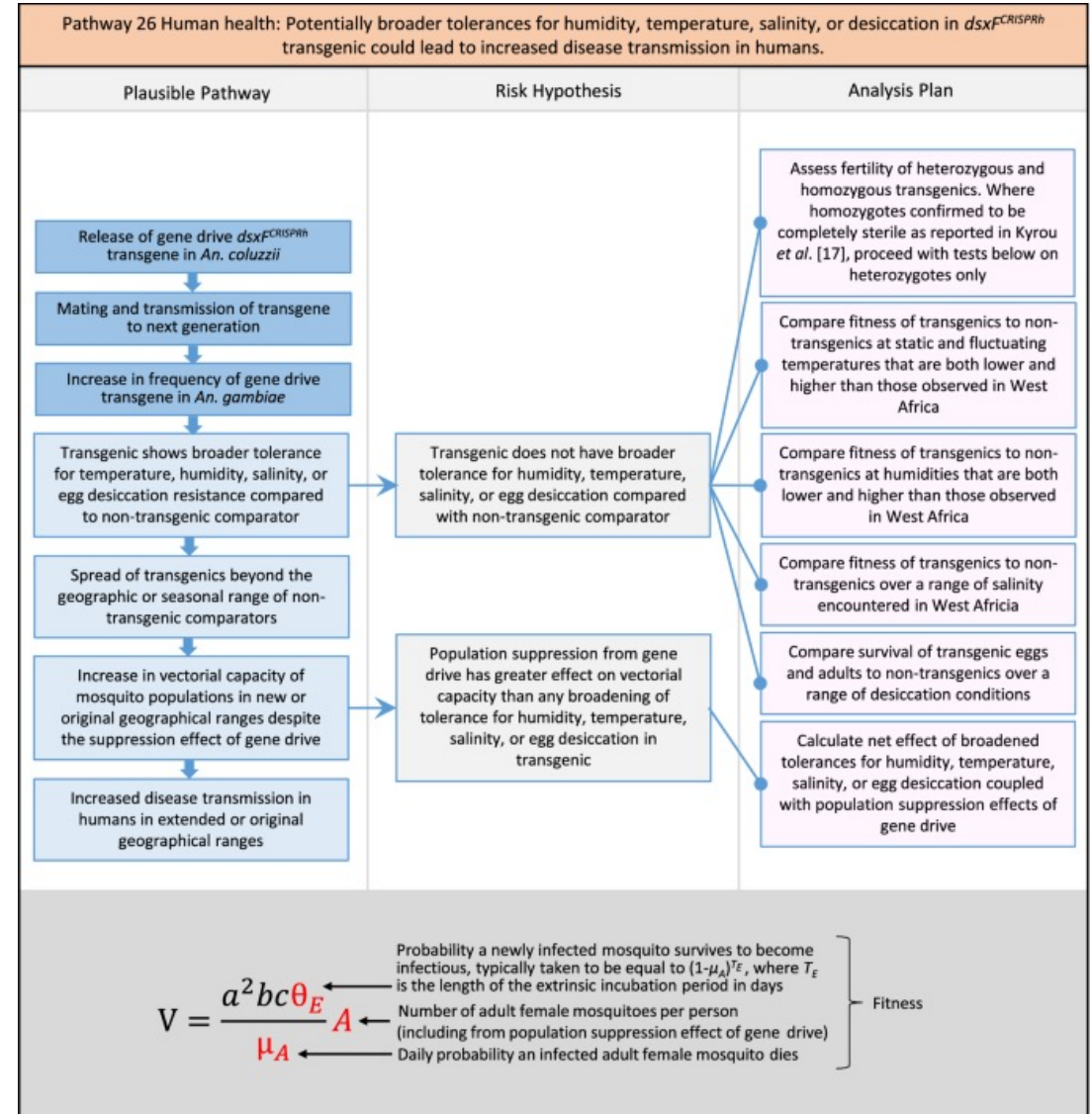
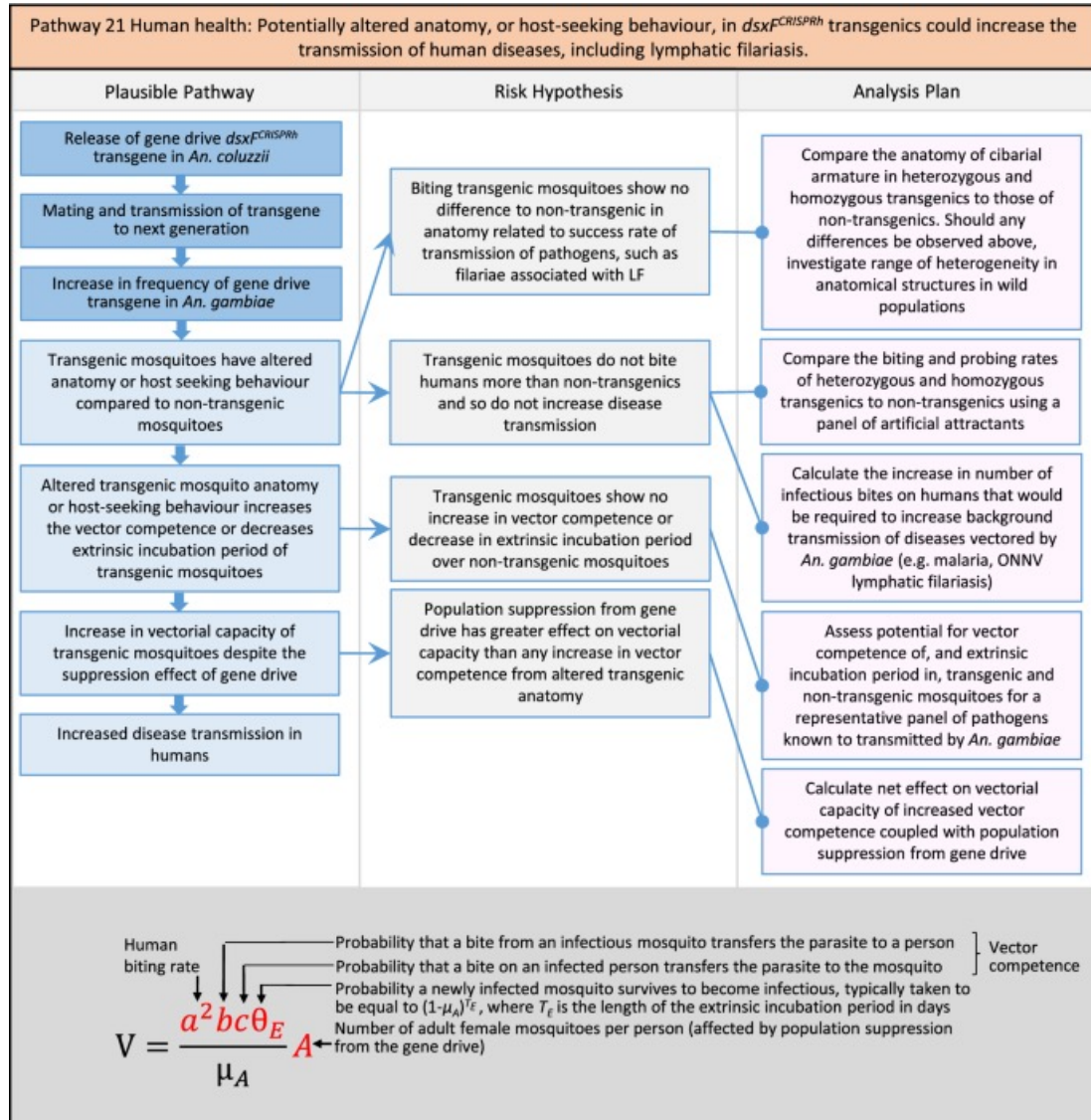
- Hoffmann AA, Montgomery BL, Popovici J, Iturbe-Ormaetxe *et al.* (2011) Nature

Surveillance needs: Detecting resistance alleles / unintended spread

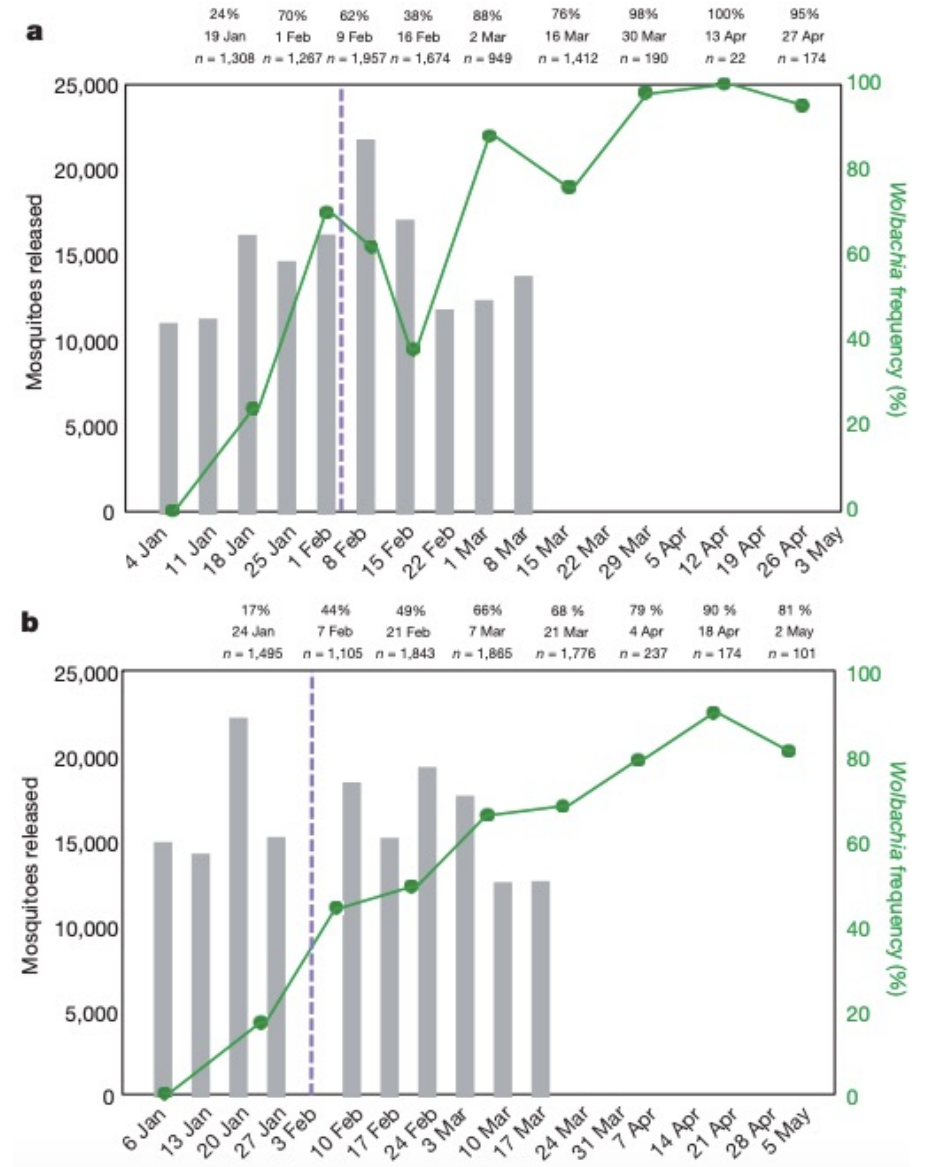
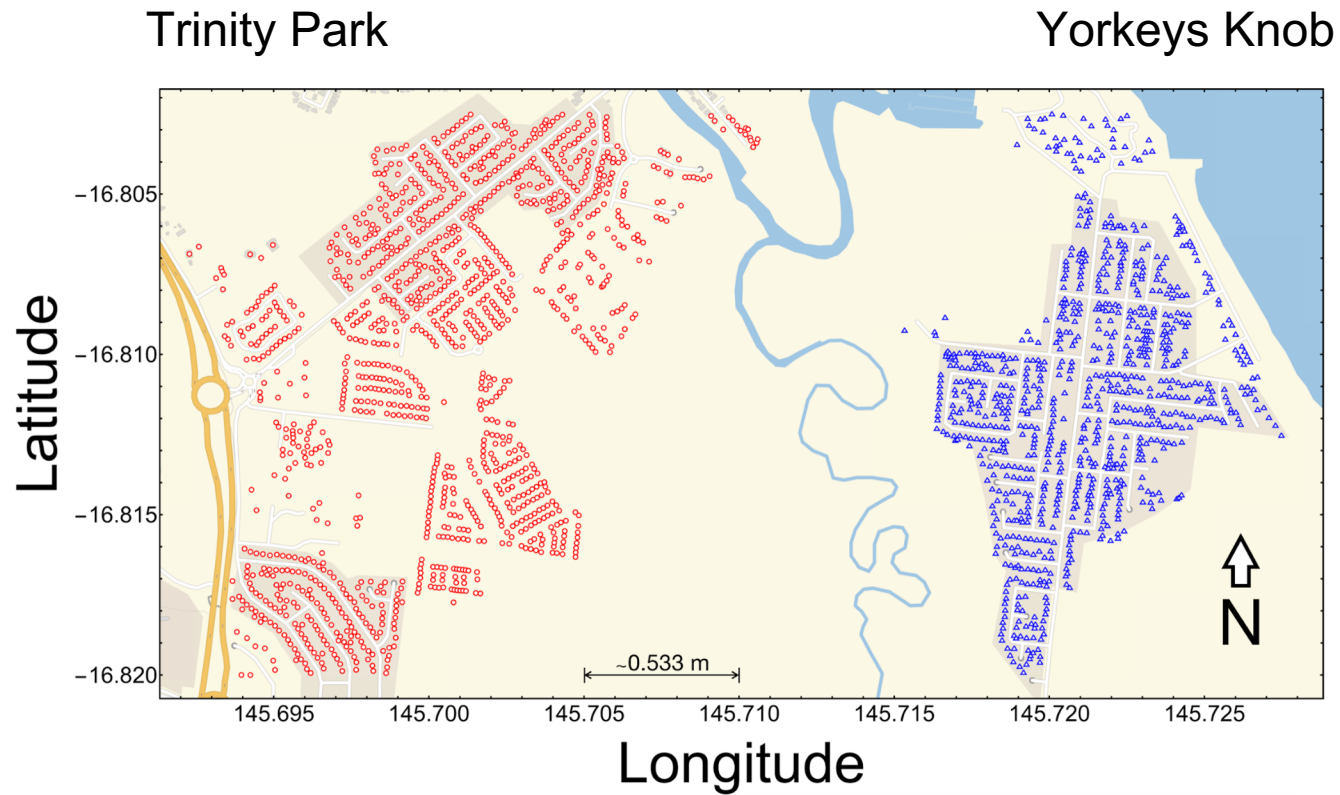


- Triska MD, Renton M (2018) R Soc Open Sci 5: 171784.

Risk assessment

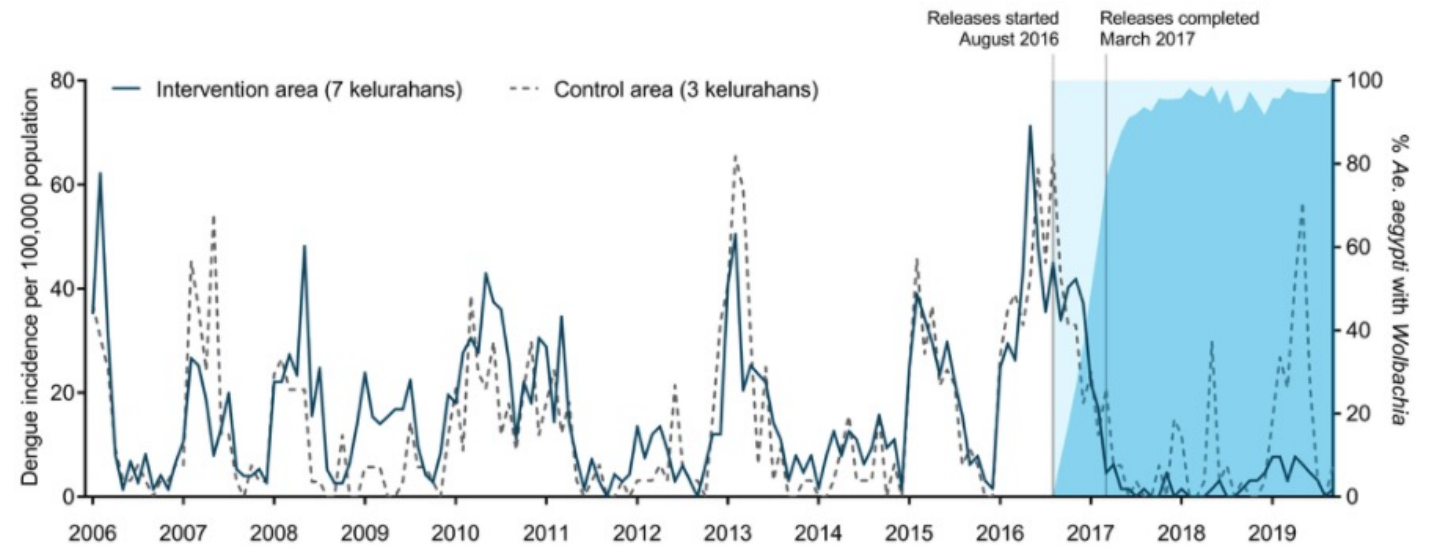
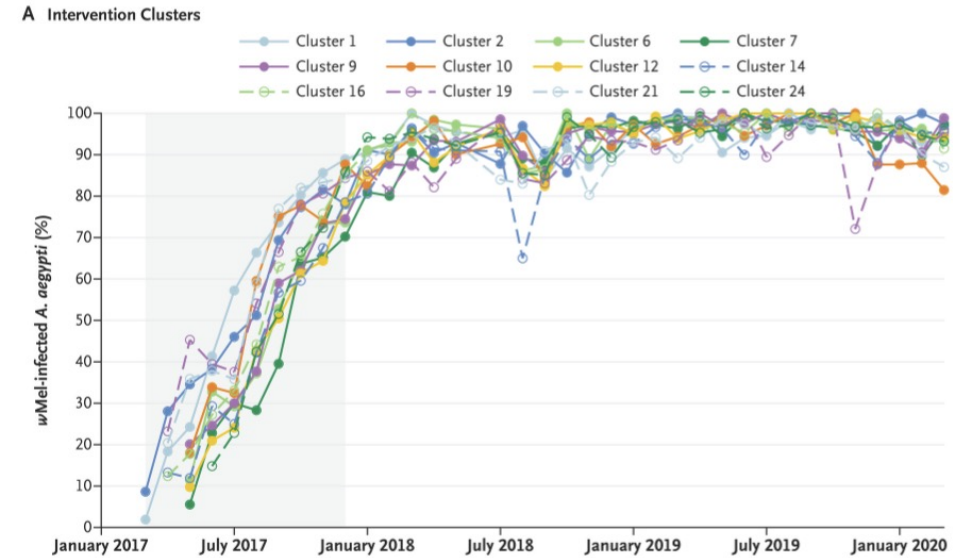
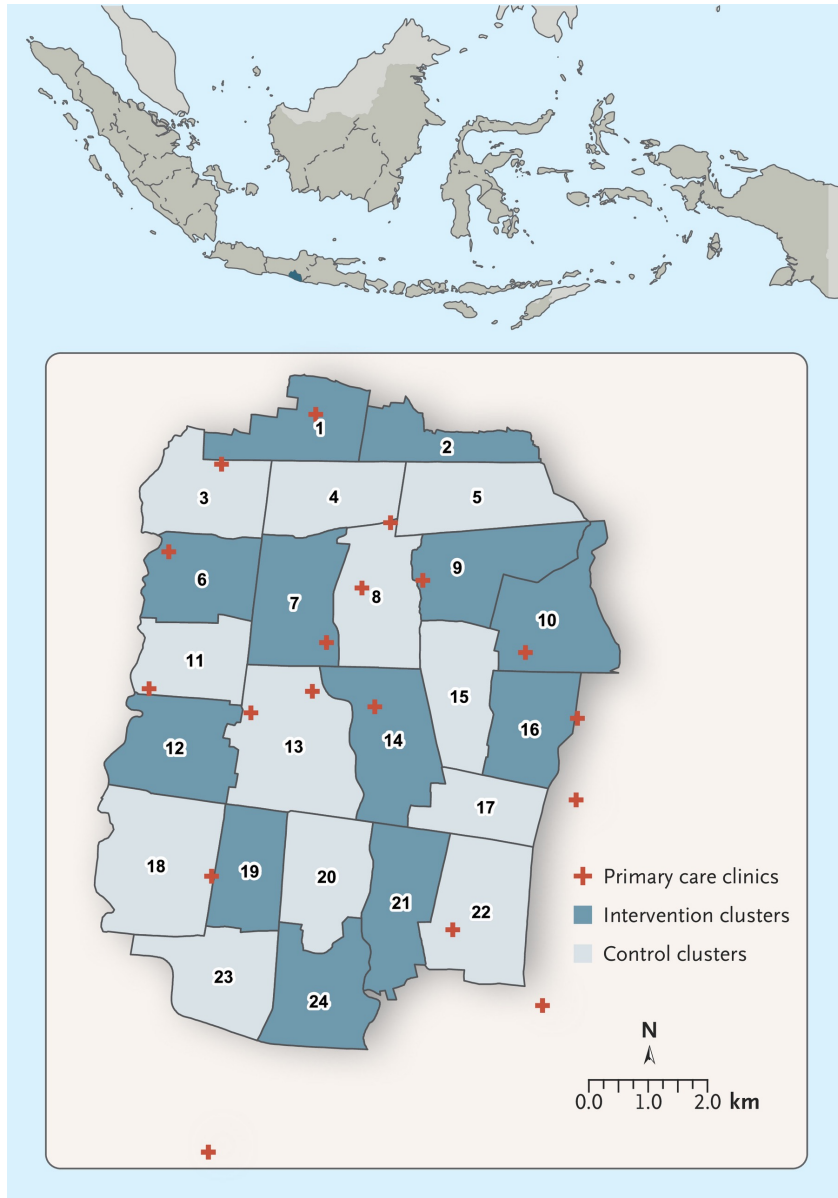


Field trial models: *Wolbachia* as a case study



- Utarini A, Indiani C, Ahmad RA, Tantowiyojo W *et al.* (2021) New England J Medicine

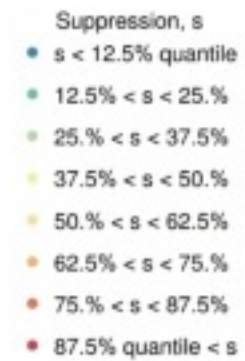
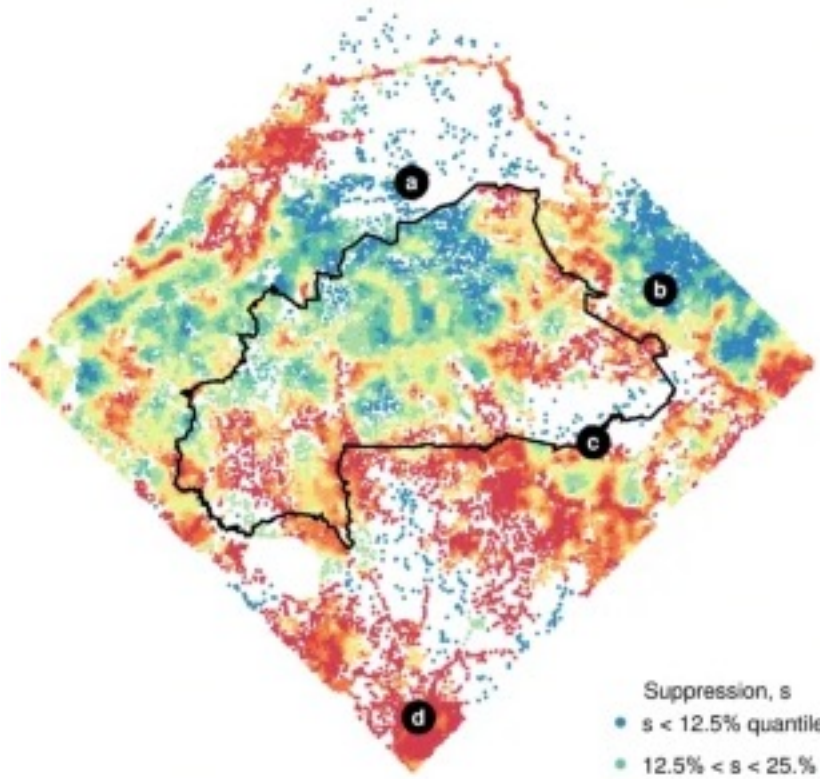
Field trial models: *Wolbachia* as a case study



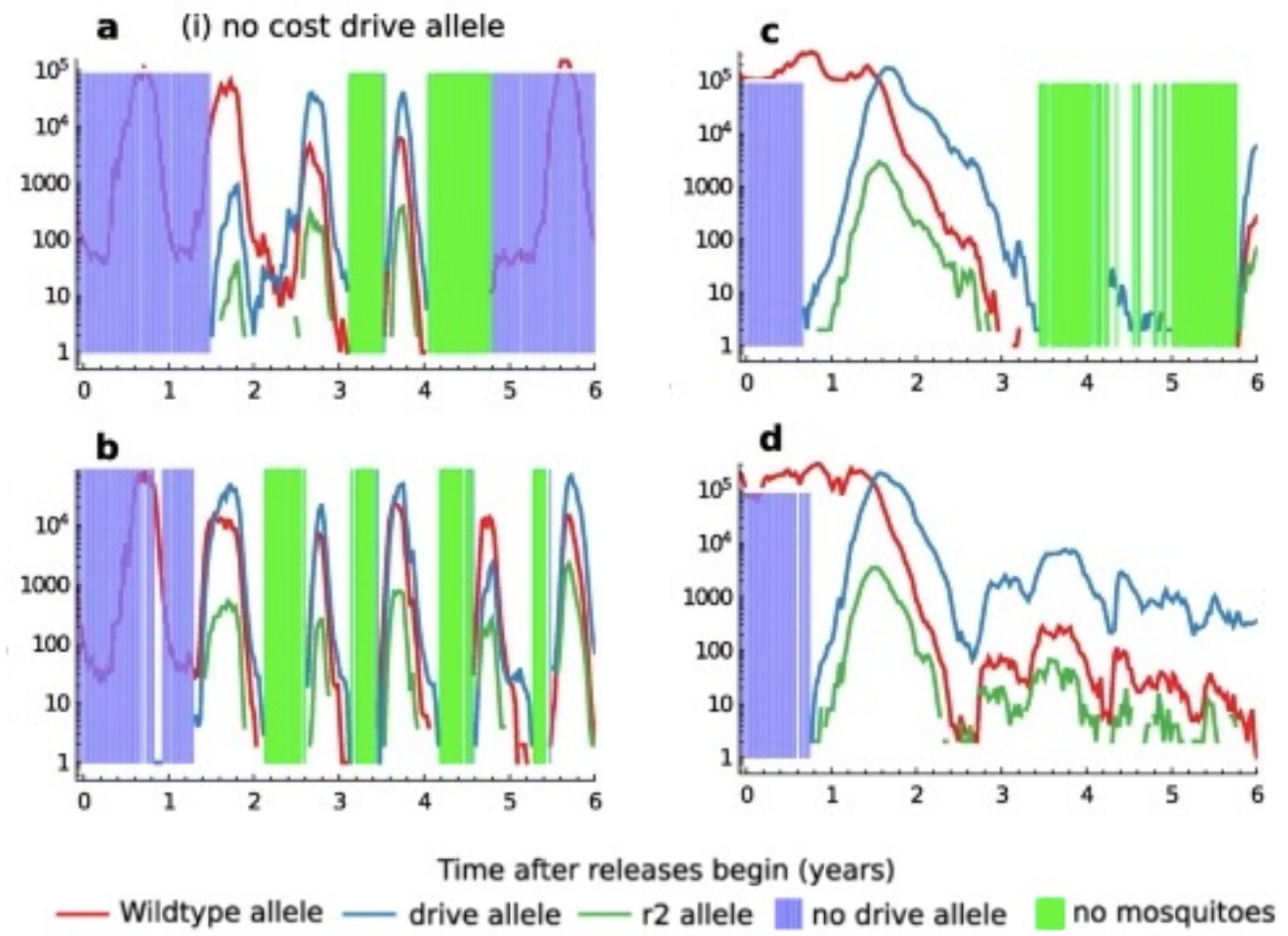
- Utarini A, Indiani C, Ahmad RA, Tantowiyojo W *et al.* (2021) *New England J Medicine*

Intervention models: *dsx* drive in Burkina Faso

No cost drive allele

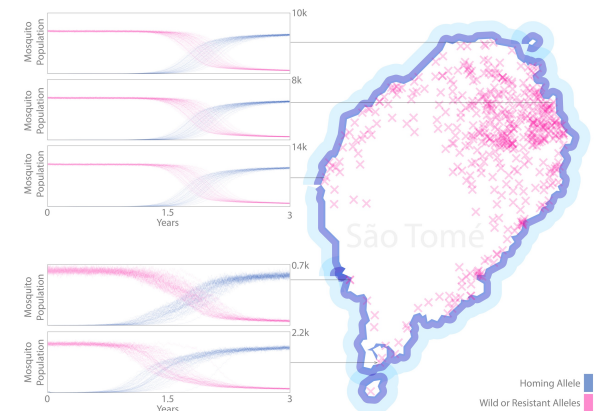
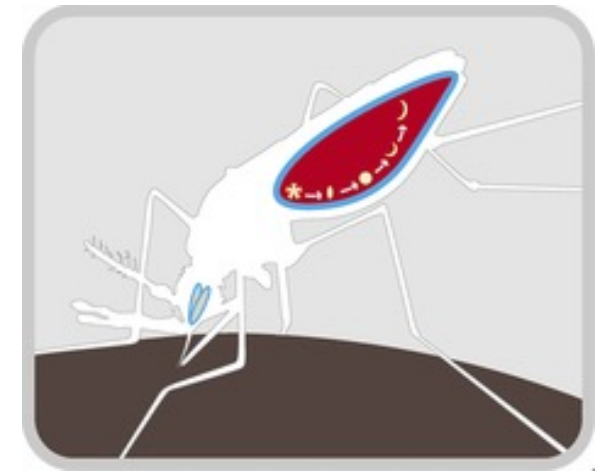


Population size (number of alleles among adult males)



Recap

1. Modeling is expected to play a growing role as gene drive mosquito projects transition from lab to field
2. Data are required to refine models concerning mosquito density dependence, habitat distribution, movement patterns & resistance allele formation
3. Assessing alignment with a target product profile & risk assessment will be important prior to a release
4. Monitoring & surveillance are expected to be cost drivers & modeling can inform cost-efficiency
5. Modeling priorities will then progress from designing cage trials & field trials to wide-scale interventions



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LAB MEMBERS:

- Héctor M. Sánchez C.
- Sean L. Wu
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- Priscilla Zhang
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- Valeri Vásquez
- Agastya Mondal
- Francois Rerolle
- Natasha Harrison
- Hao Wang
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- Chris De Leon

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- **Lanzaro Lab @ UC Davis**
- **James Lab @ UC Irvine**
- **Bier Lab @ UCSD**
- **Hay Lab @ Caltech**
- **Malaria Elimination Initiative @ UCSF**
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- **Dr Samson Kiware @ Ifakara Health Institute**
- **School of Public Health @ UC Berkeley**

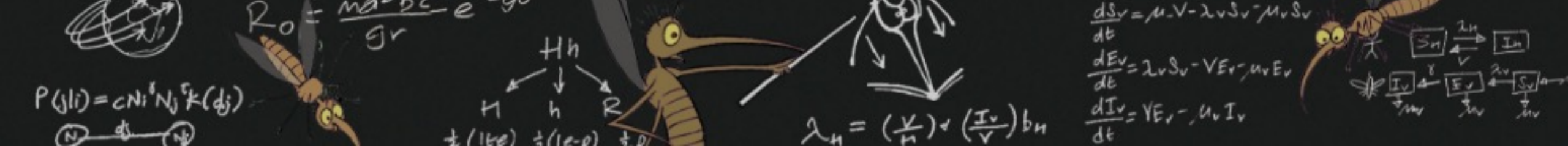
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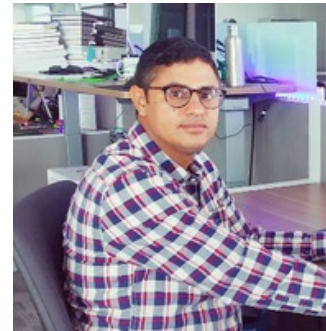
Héctor M. Sánchez C.



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Gordana Rašić



Jared Bennett



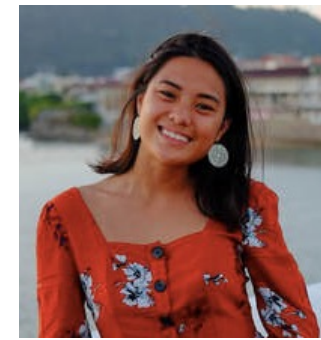
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Thien-An Ha



Darpa Anireddy



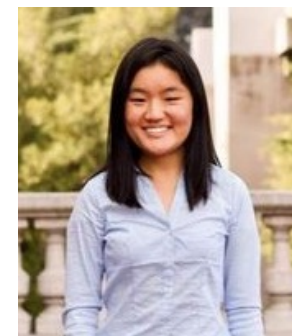
Natasha Harrison



Ameet Bindra



Chris De Leon



Ashley Zhang