



# **INFECTIOUS DISEASE MANAGEMENT UNDER CLIMATE CHANGE: MODELS AND APPLICATIONS**

**VÁLERI N. VÁSQUEZ  
UNIVERSITY OF CALIFORNIA BERKELEY  
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# CONTEXT



# CLIMATE CHANGE

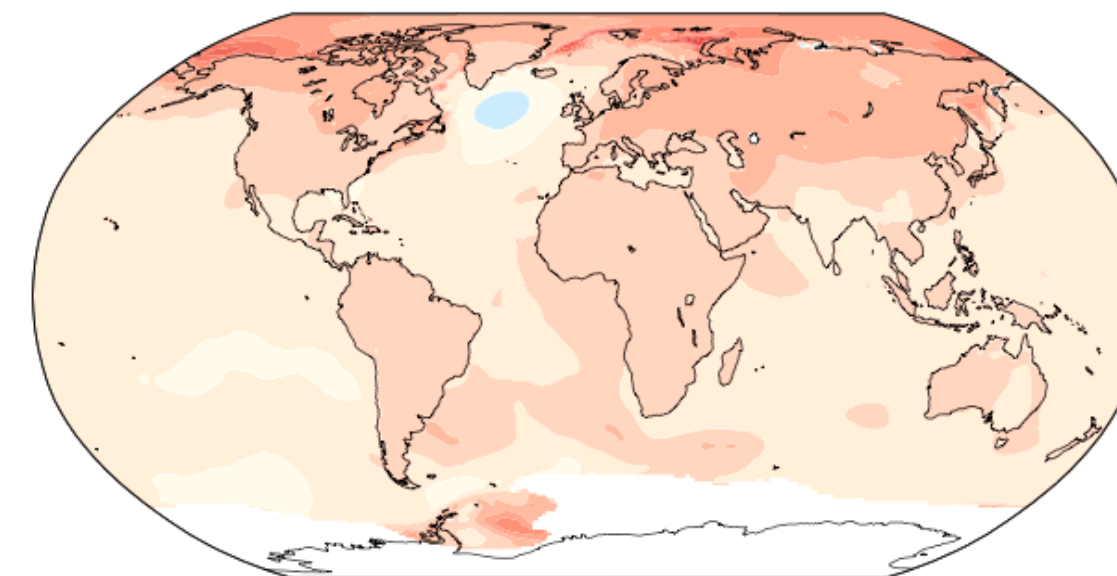
## HIGHER AVERAGES

“With every increment of global warming, changes get larger in regional **mean temperature.**”

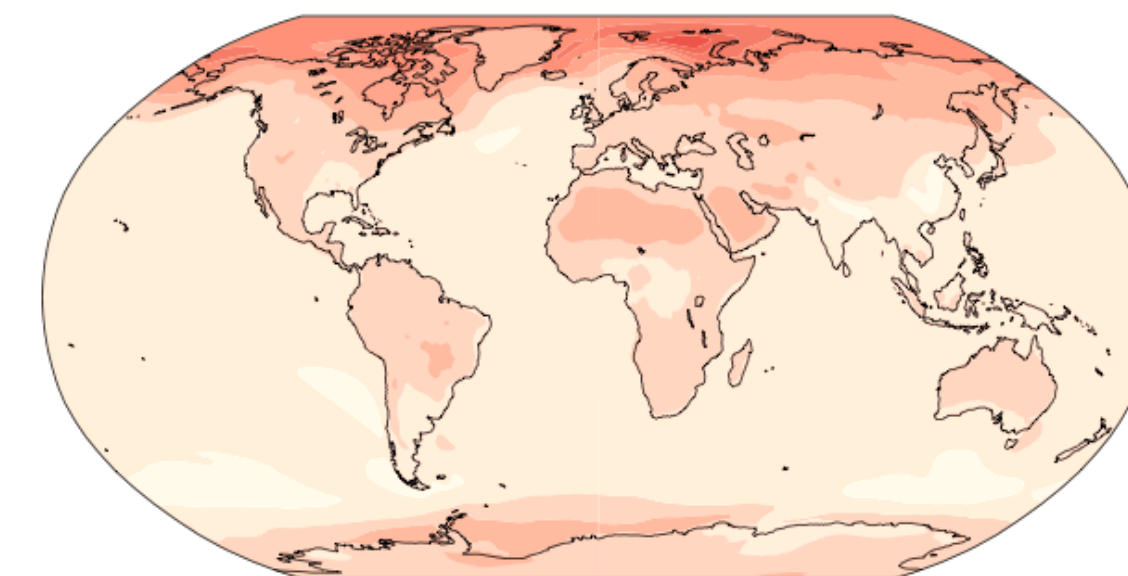
(a) Annual mean temperature change (°C) at 1°C global warming

Warming at 1°C affects all continents and is generally larger over land than over the oceans in both observations and models. Across most regions, observed and simulated patterns are consistent.

Observed change per 1°C global warming



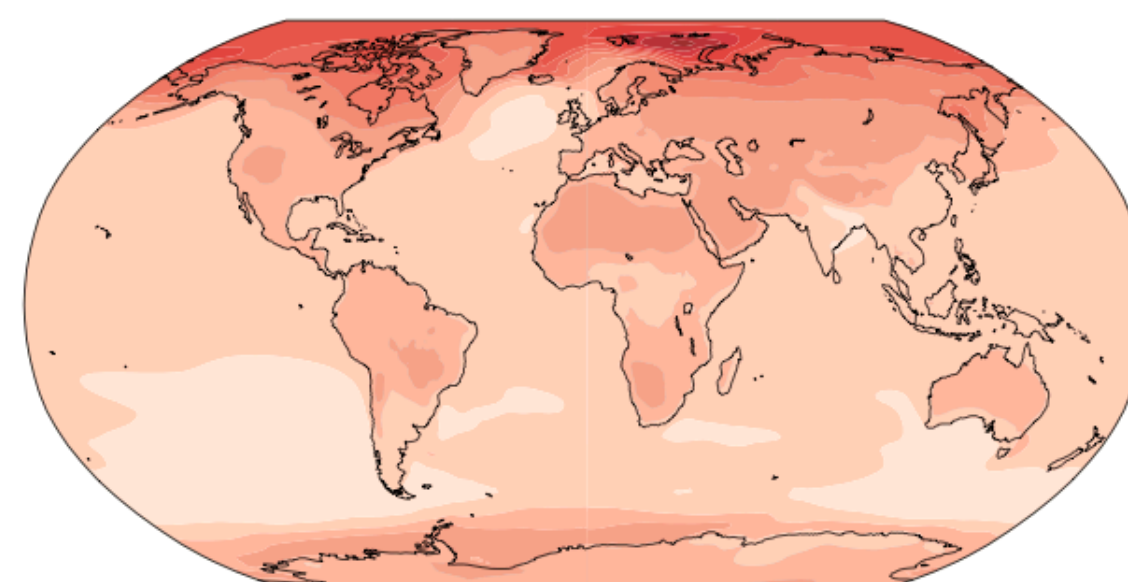
Simulated change at 1°C global warming



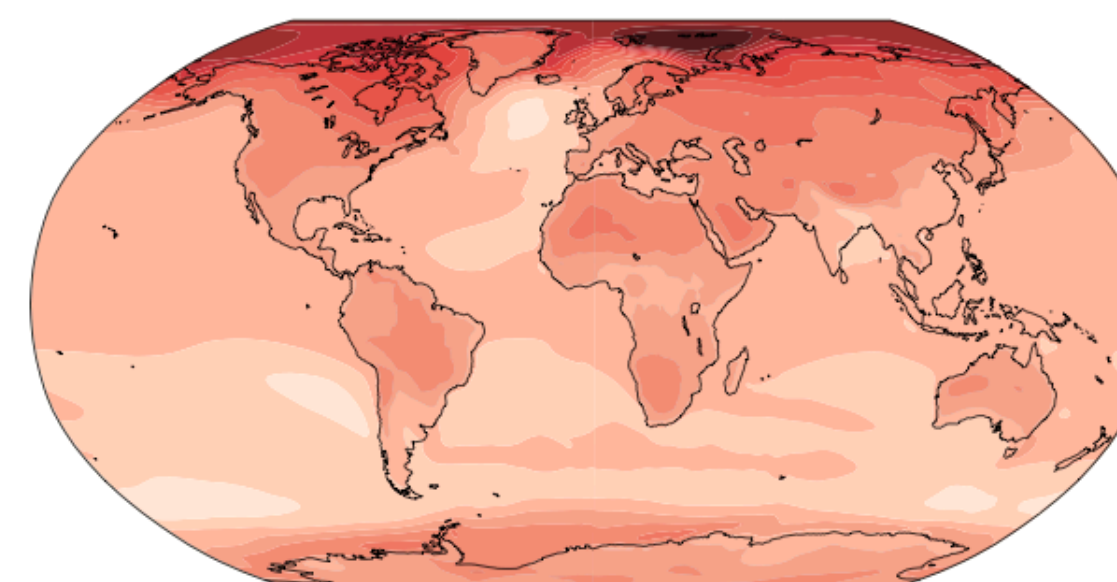
(b) Annual mean temperature change (°C) relative to 1850–1900

Across warming levels, land areas warm more than ocean areas, and the Arctic and Antarctica warm more than the tropics.

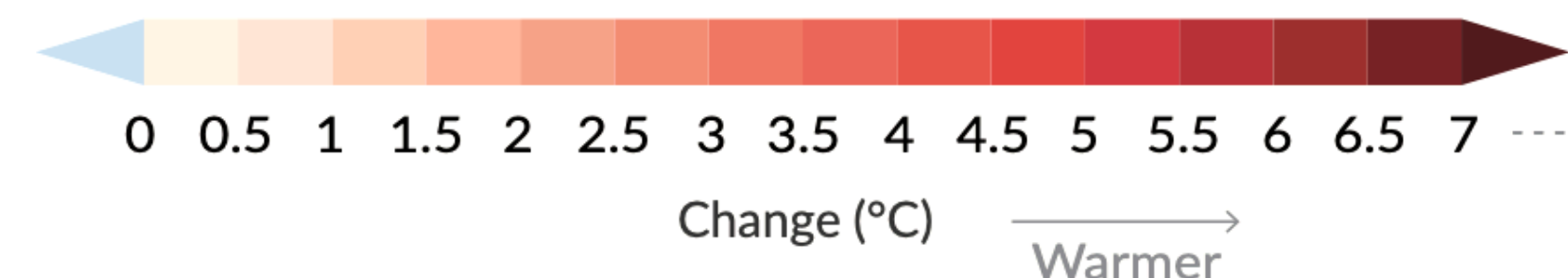
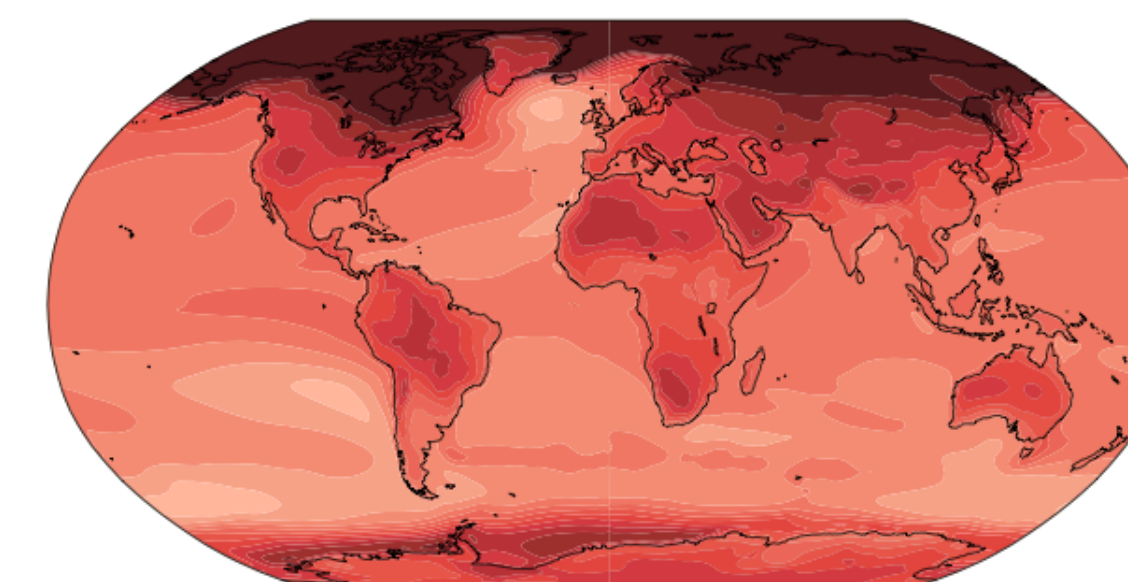
Simulated change at 1.5°C global warming



Simulated change at 2°C global warming



Simulated change at 4°C global warming



# CLIMATE CHANGE

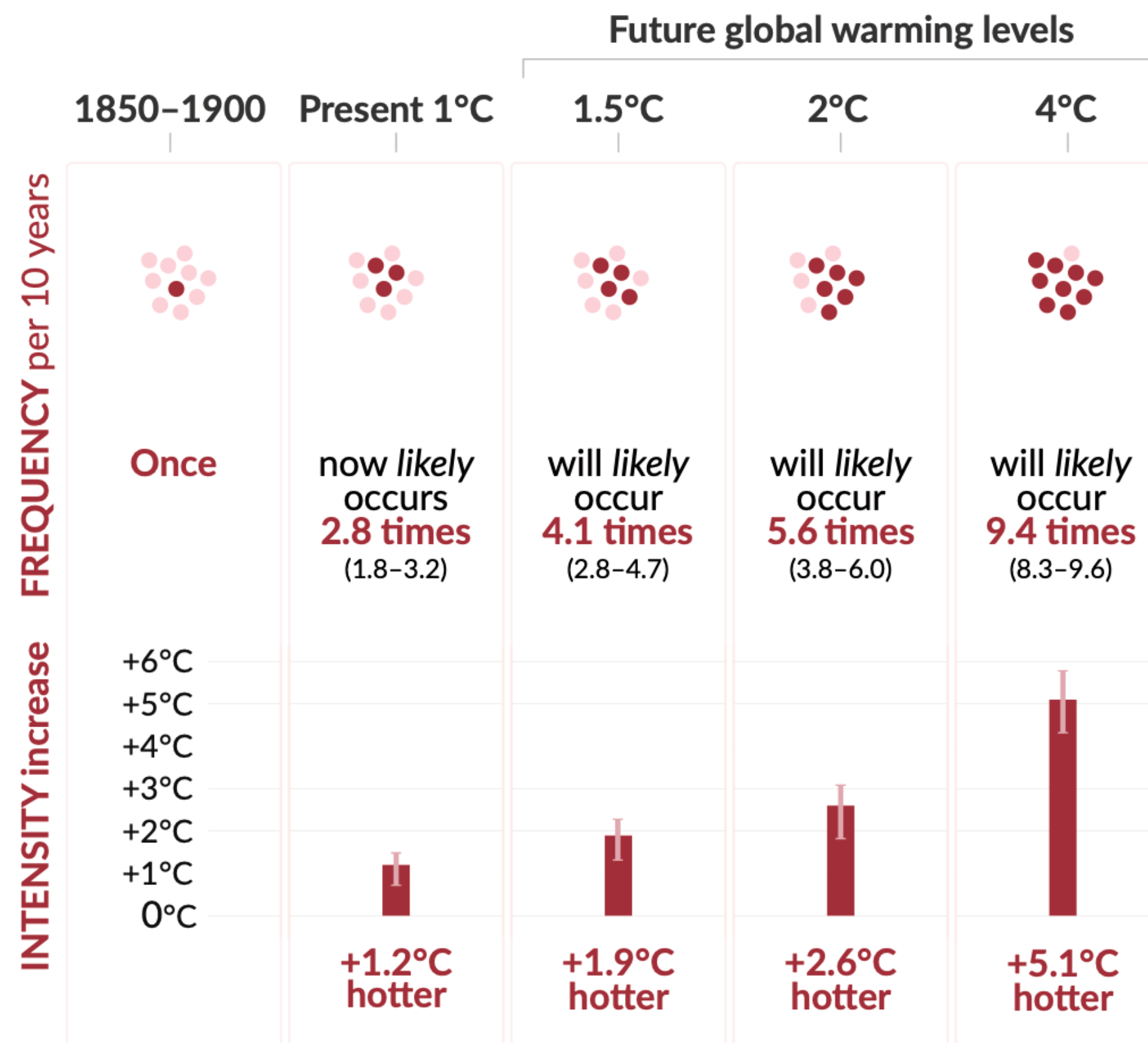
## GREATER EXTREMES

“Projected changes in extremes are larger in **frequency and intensity** with every additional increment of global warming.”

### Hot temperature extremes over land

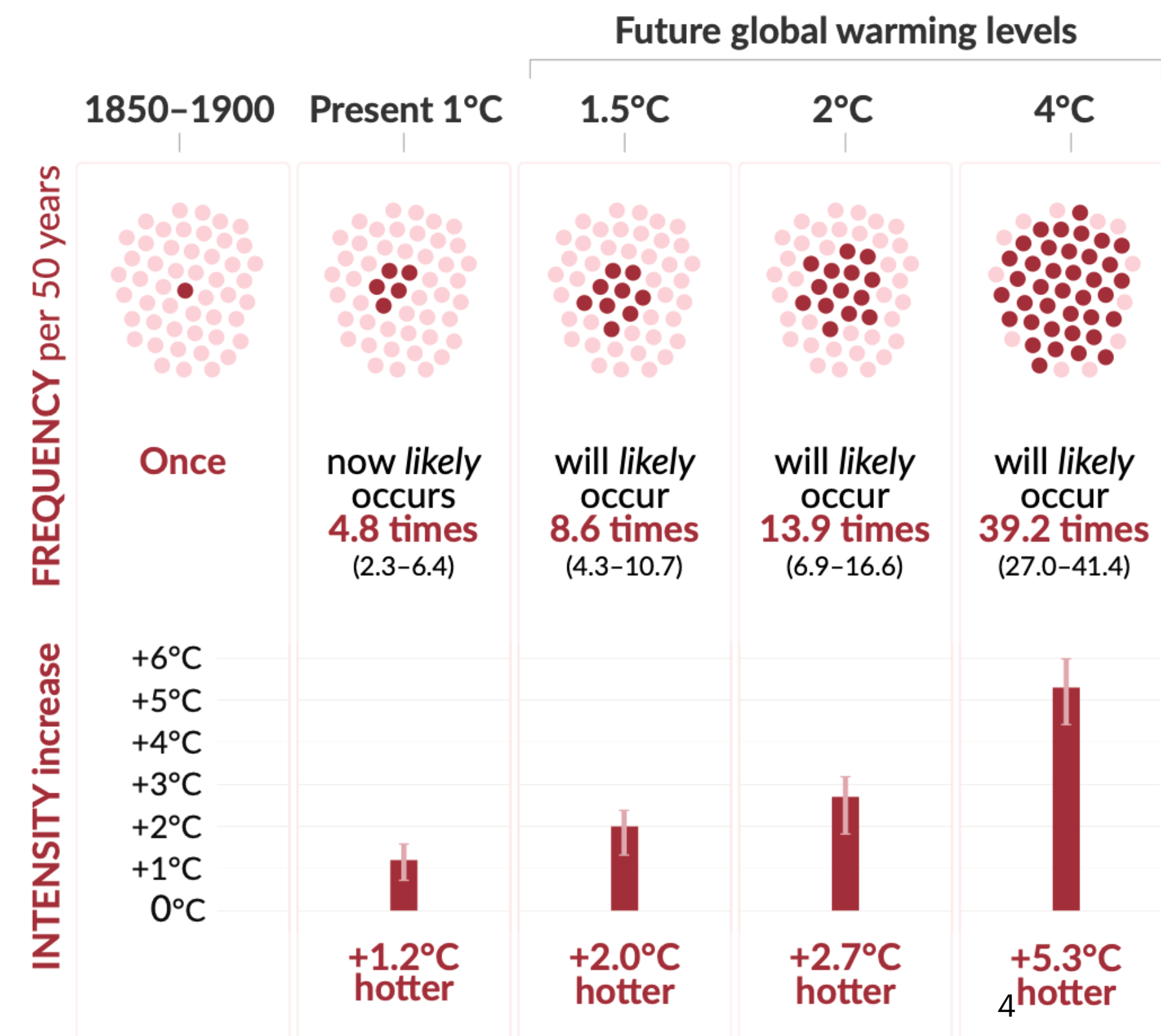
#### 10-year event

Frequency and increase in intensity of extreme temperature event that occurred **once in 10 years** on average in a climate without human influence



#### 50-year event

Frequency and increase in intensity of extreme temperature event that occurred **once in 50 years** on average in a climate without human influence

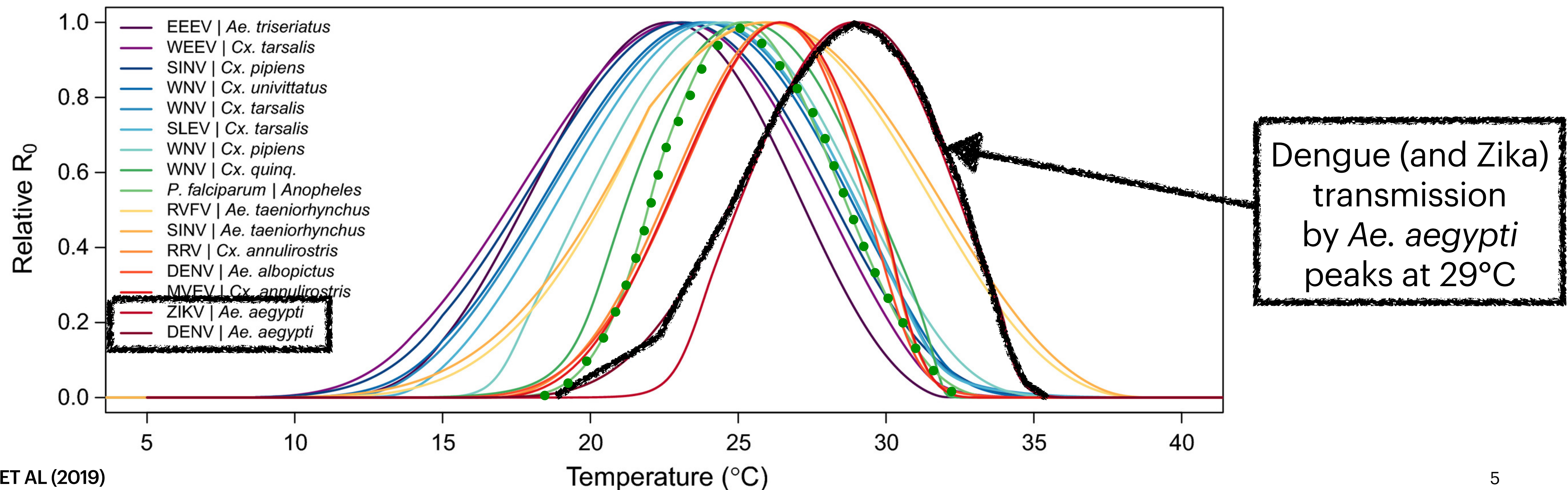




# SPATIOTEMPORAL SHIFT IN DISEASE

## THERMAL BIOLOGY OF VECTORS

While scientists “cannot conclusively predict changes in incidence based on temperature alone”, the thermal biology literature argues that **temperature change will promote transmission of certain diseases even as it limits transmission of others.**





# SHIFT ALSO REQUIRED FOR CONTROL

NEW CLIMATE AND DISEASE REALITIES —> NEW TOOLS NEEDED

The New York Times

Gene Drives Offer  
Against Disease

nature

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nature > news > article

NEWS | 18 April 2022

## Biotech firm announces results from first US trial of genetically modified mosquitoes

Oxitec reports that its insects behaved as planned – but a larger trial is needed to learn whether they can reduce wild mosquito populations.



A woman in Tanzania under a mosquito tent with a relative who was being treated for malaria. With gene drives, it may be possible to kill off a mosquito population or make the population resistant to malaria parasites. Uriel Sinai for The New York Times

The New York Times

## It Takes a Mosquito to Fight a Mosquito

In Australia, China and elsewhere, scientists are fighting disease-carrying mosquitoes by introducing another type, carrying just a harmless form of bacteria.

Jan. 8, 2019

## Tiny Bacterium Called *Wolbachia* Could Defeat Dengue

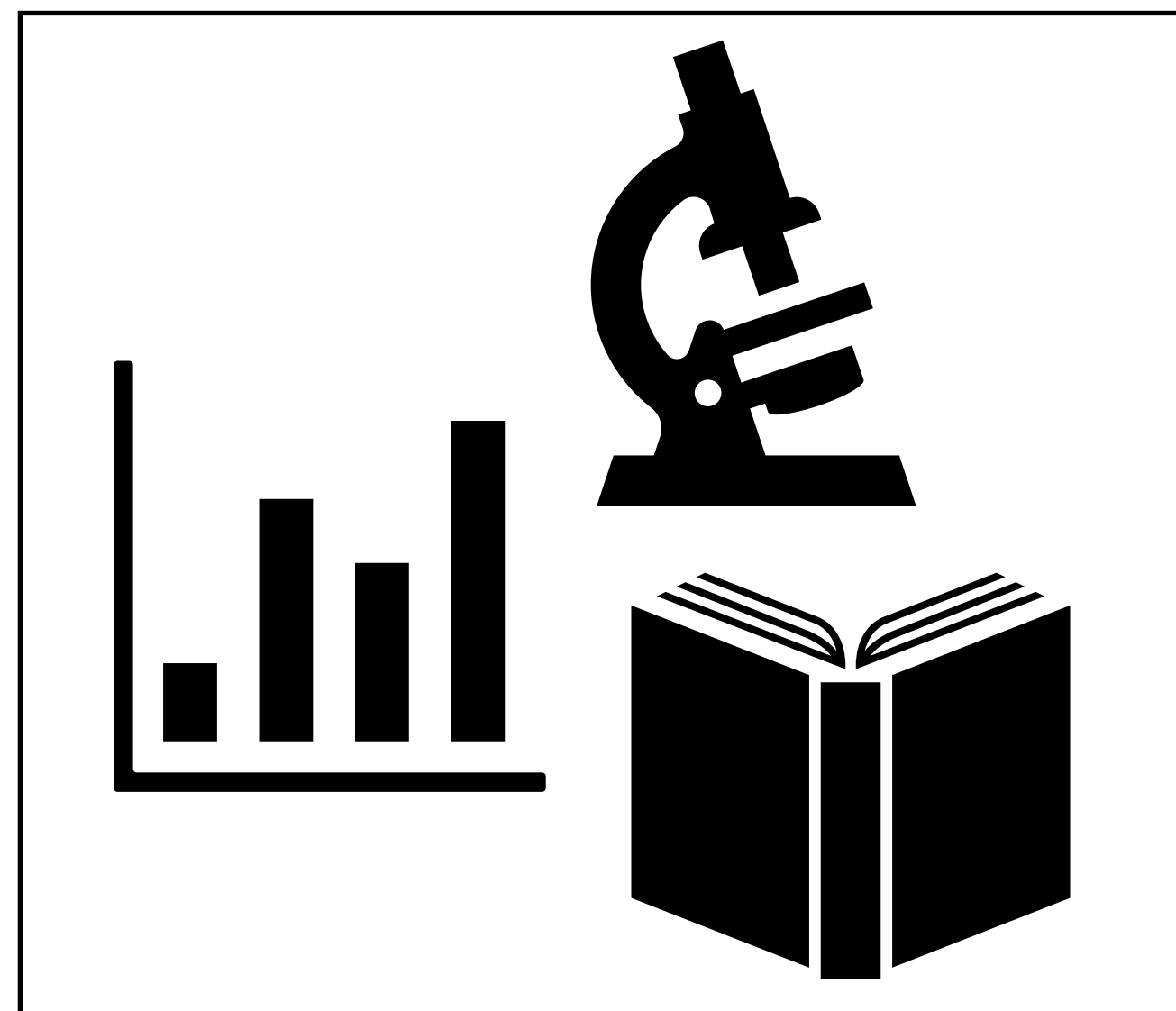
Scientists are immunizing mosquitoes against disease with the help of a common microbe



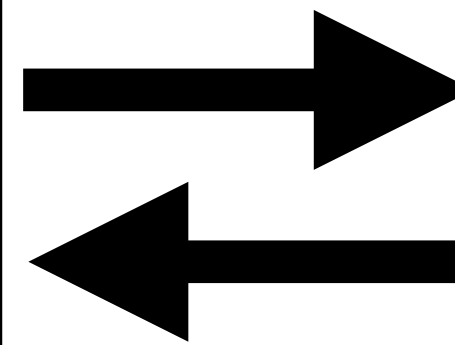
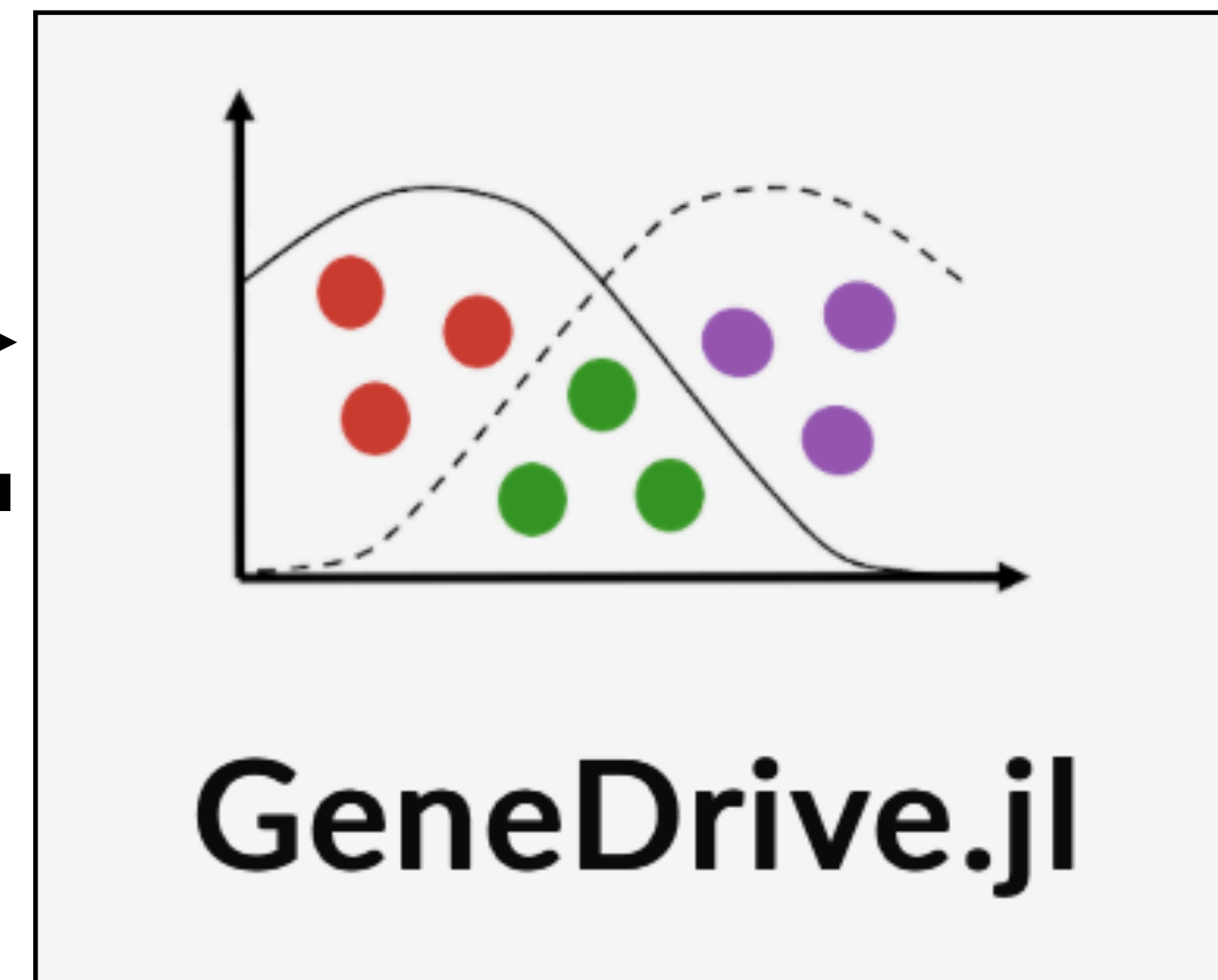
# SOFTWARE INFORMED BY SCIENCE

PHILOSOPHY OF DEVELOPMENT AND DESIGN

RESEARCH



SOFTWARE







# SOFTWARE

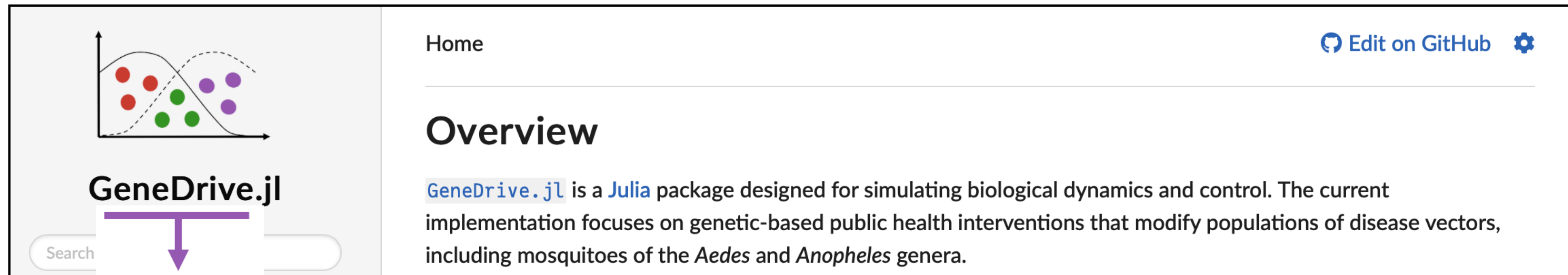



# GENEDRIVE.JL

## SOFTWARE INFORMED BY SCIENCE

Operational focus: genetic-based tools

- Characterize environmentally sensitive dynamics
- Define optimal policies for biocontrol



Home [Edit on GitHub](#) 

### Overview

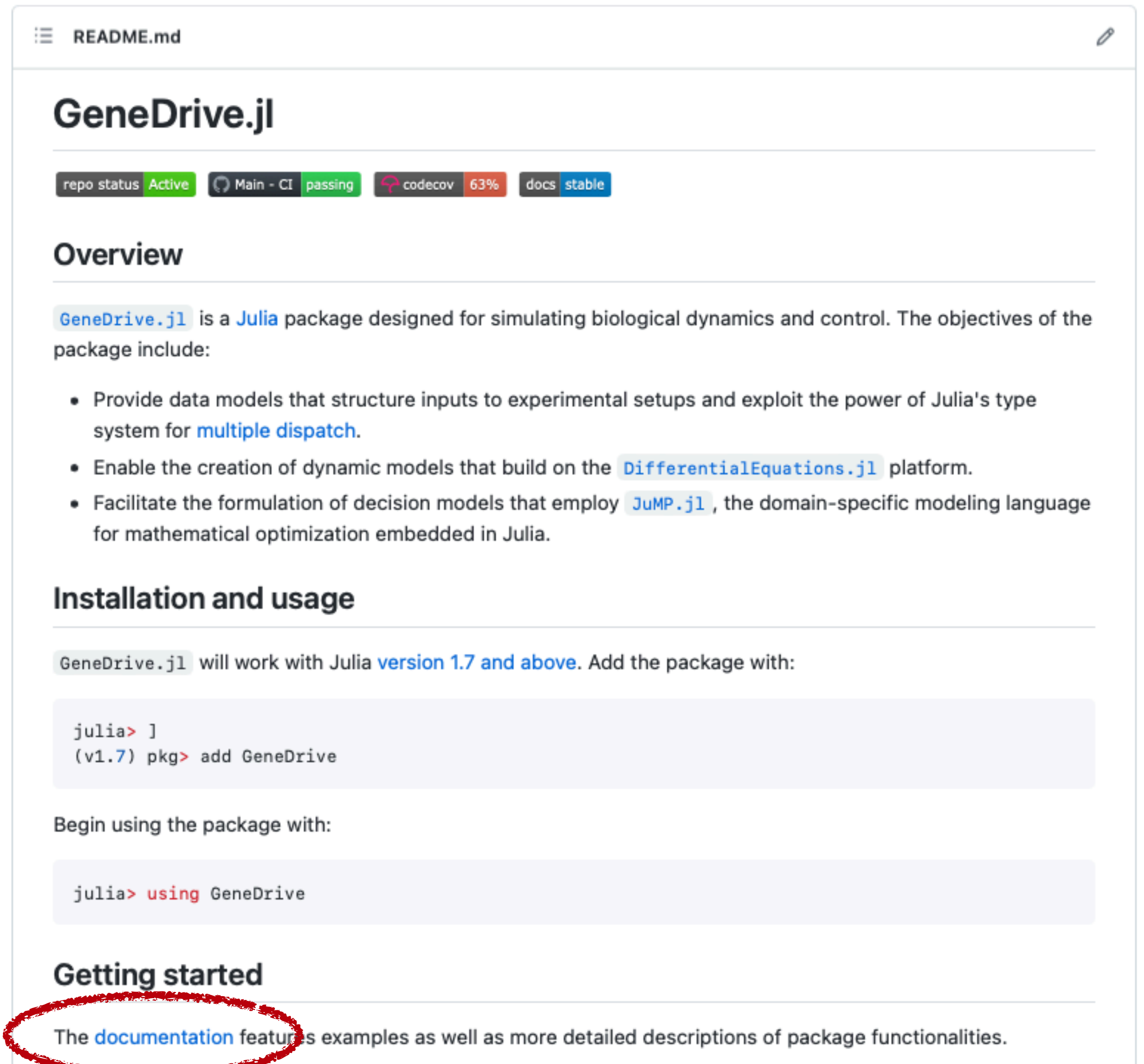
[GeneDrive.jl](#) is a [Julia](#) package designed for simulating biological dynamics and control. The current implementation focuses on genetic-based public health interventions that modify populations of disease vectors, including mosquitoes of the *Aedes* and *Anopheles* genera.



# GENEDRIVE.JL

## SOFTWARE INFORMED BY SCIENCE

- Open source
- Extensible and scalable
- Documentation including examples
- Collaborators welcome

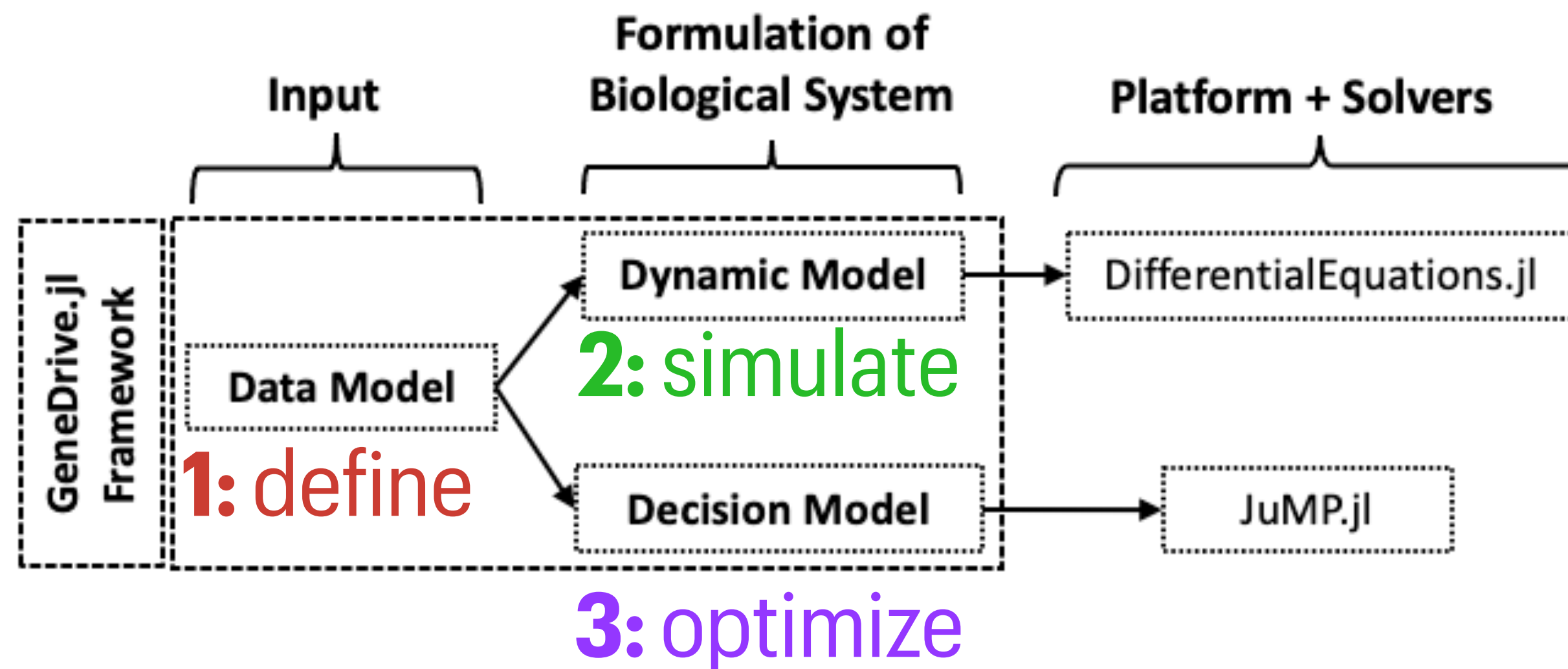


The screenshot shows the README.md file for the GeneDrive.jl package. At the top, it displays the package name "GeneDrive.jl" and a row of status badges: "repo status Active", "Main - CI passing", "codecov 63%", and "docs stable". Below this is an "Overview" section that describes the package as a Julia tool for biological dynamics simulation, listing its goals: providing data models, enabling dynamic models on the DifferentialEquations.jl platform, and facilitating decision models using JuMP.jl. The "Installation and usage" section provides instructions to install the package using Julia 1.7 or above and shows the terminal commands: `julia> ]`, `(v1.7) pkg> add GeneDrive`, and `julia> using GeneDrive`. The "Getting started" section begins with the sentence "The [documentation](#) features examples as well as more detailed descriptions of package functionalities.", where the word "documentation" is circled in red.



# GENEDRIVE.JL

## A THREE-PART FRAMEWORK



1. **Data model:** An abstraction used to define simulation inputs, store them, and dispatch methods.
2. **Dynamic model:** An ordinary differential equation formulation of the biological system.
3. **Decision model:** A discretized implementation of the biological system as a nonlinear mathematical program.





# RESEARCH AREA #1

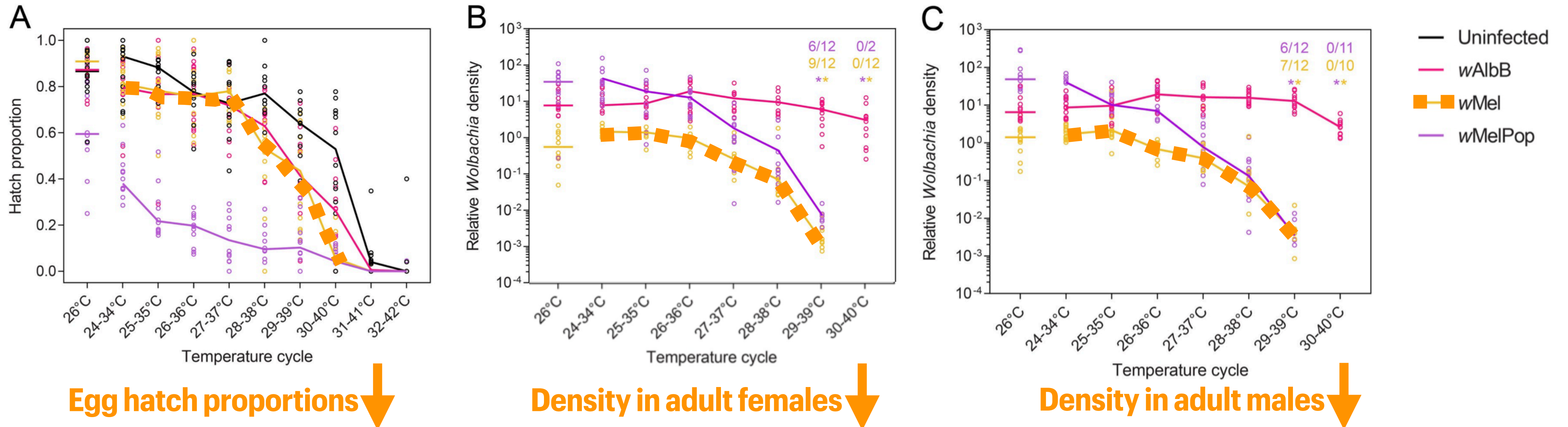


# TECHNOLOGICAL: WHAT TOOLS ARE (CLIMATE) APPROPRIATE?

## THERMAL BIOLOGY OF GENETIC-BASED INTERVENTION: WOLBACHIA (wMEL)

Most widely deployed bacterial strain

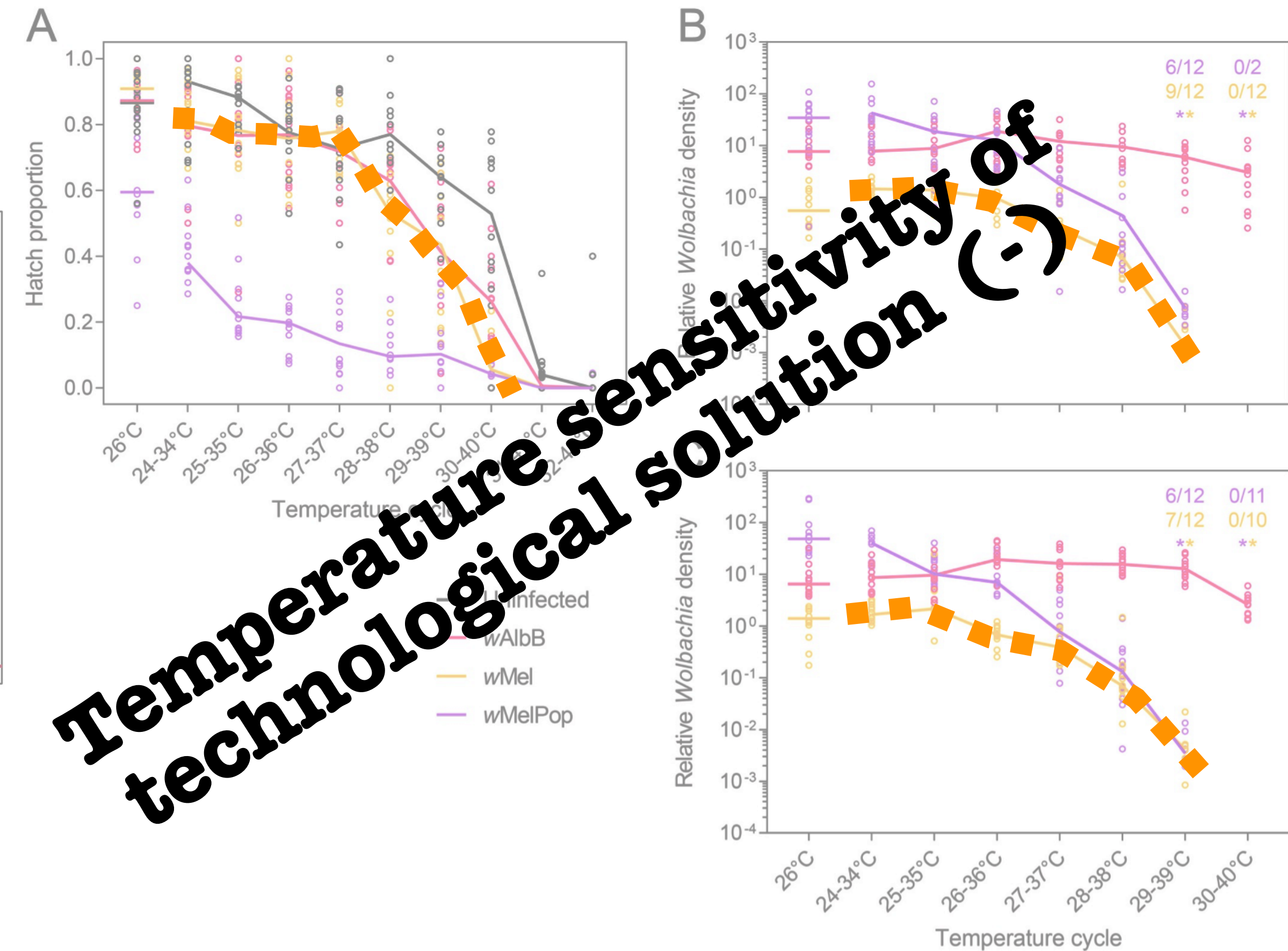
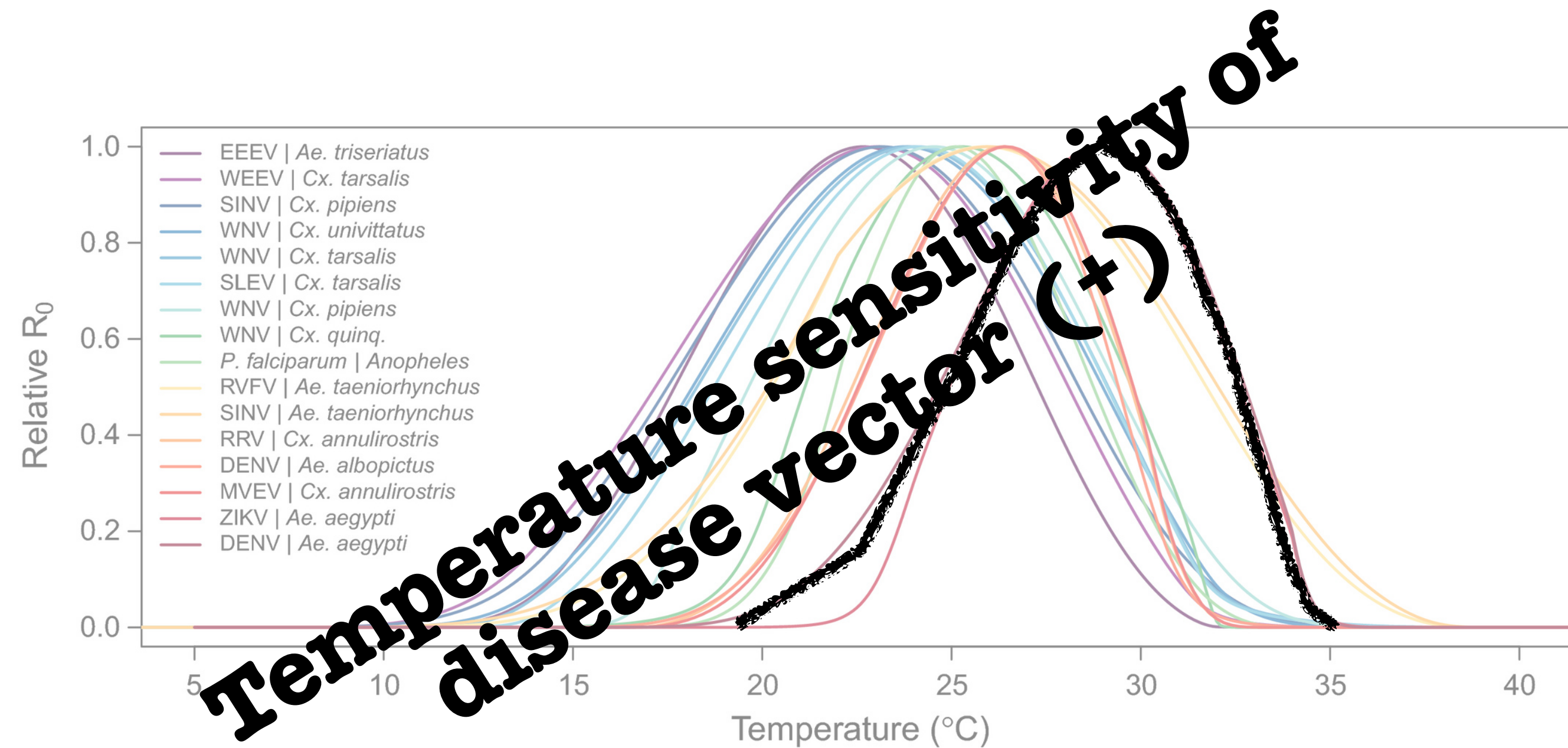
- *Wolbachia* infection: maternally inherited, and pathogen blocking.
- Key phenotype enabling use as a public health tool is weakened by cyclical heat stress.
- High temperatures can **cause the infection to fall out of the population entirely past 35°C**





# TECHNOLOGICAL: WHAT TOOLS ARE (CLIMATE) APPROPRIATE?

## THERMAL BIOLOGY OF GENETIC-BASED INTERVENTION: WOLBACHIA (wMEL)



Temperature sensitivity of technological solution (-)

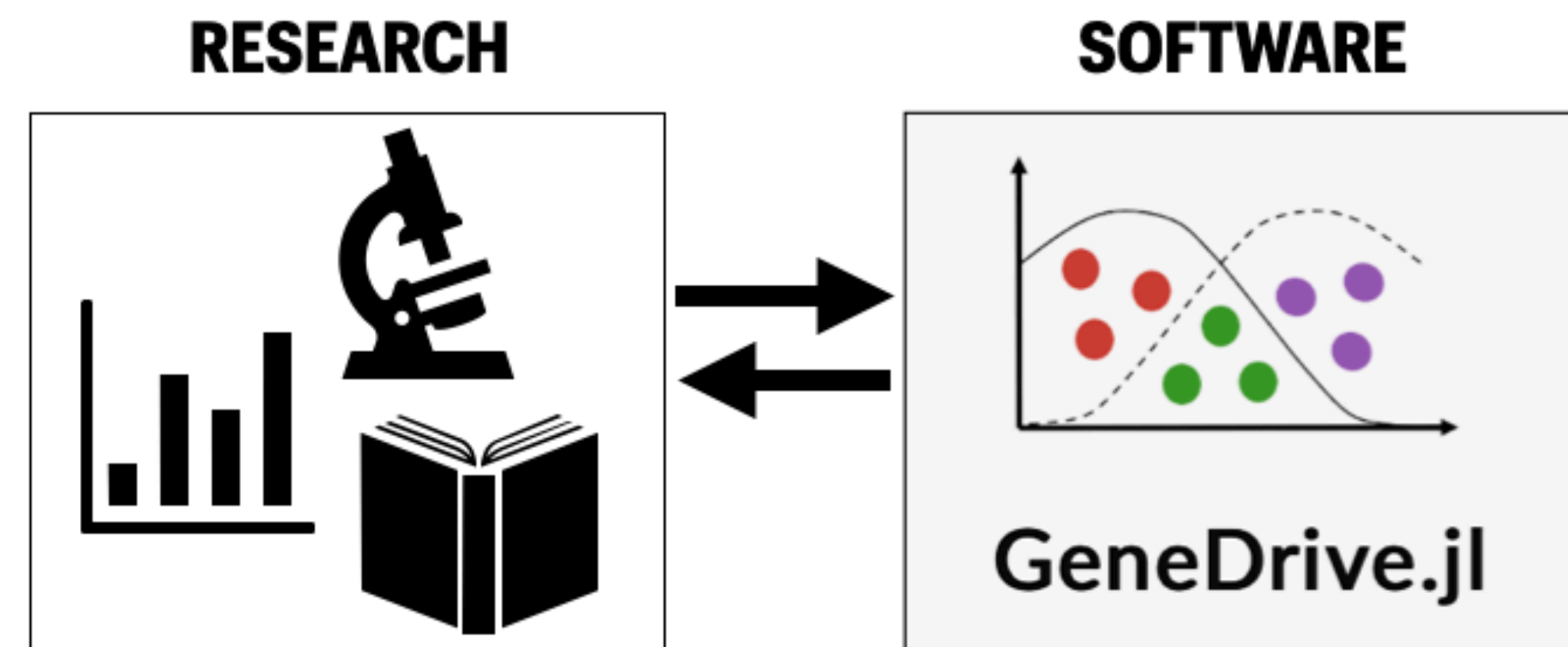


# EVALUATING WHETHER TOOLS ARE (CLIMATE) APPROPRIATE

## THE CASE FOR SOFTWARE

- Empirical research is essential: both laboratory and field studies furnish fundamental insights.
- To evaluate findings in the **necessary context (e.g., future climate realities)**, models are required.
- Models facilitate hypothesis testing:
  - ❖ **Explore uncertainties** in empirical data
  - ❖ **Extrapolate** to include new scenarios

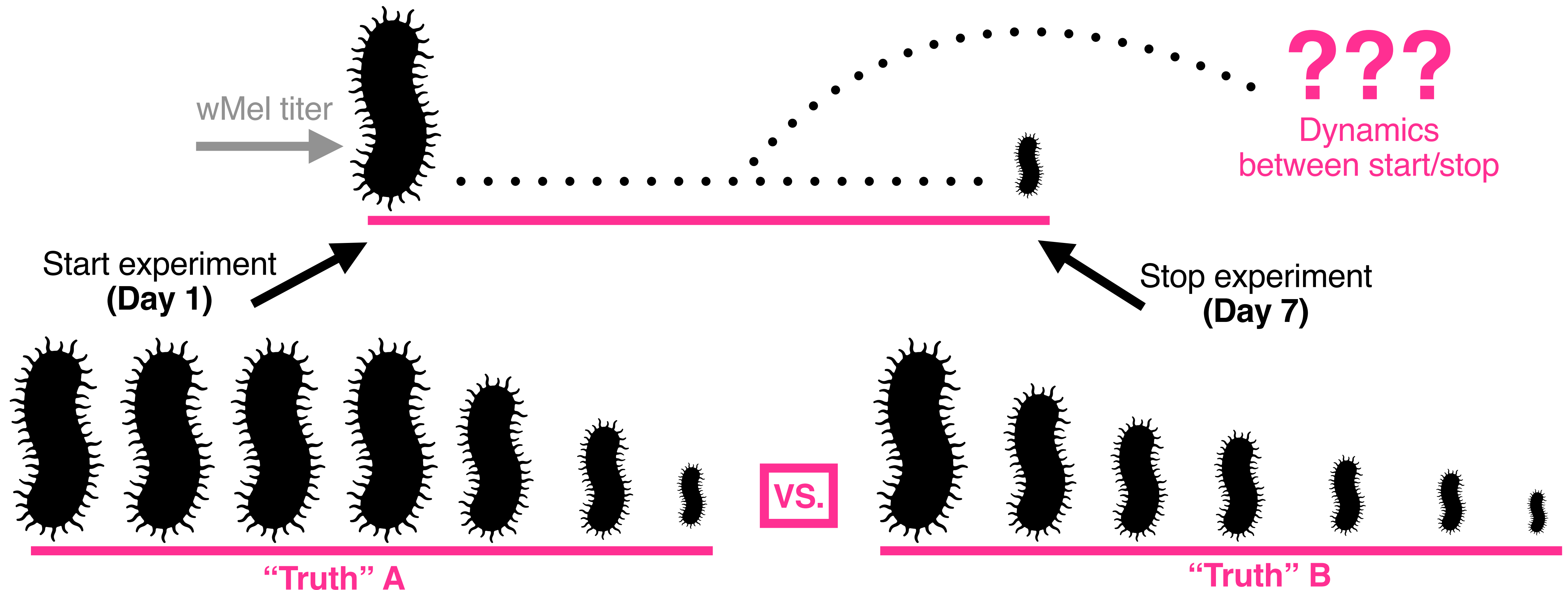
Formalize in software for “modular” modifications,  
scientific replicability





# REQUIREMENT: EXPLORE UNCERTAINTIES

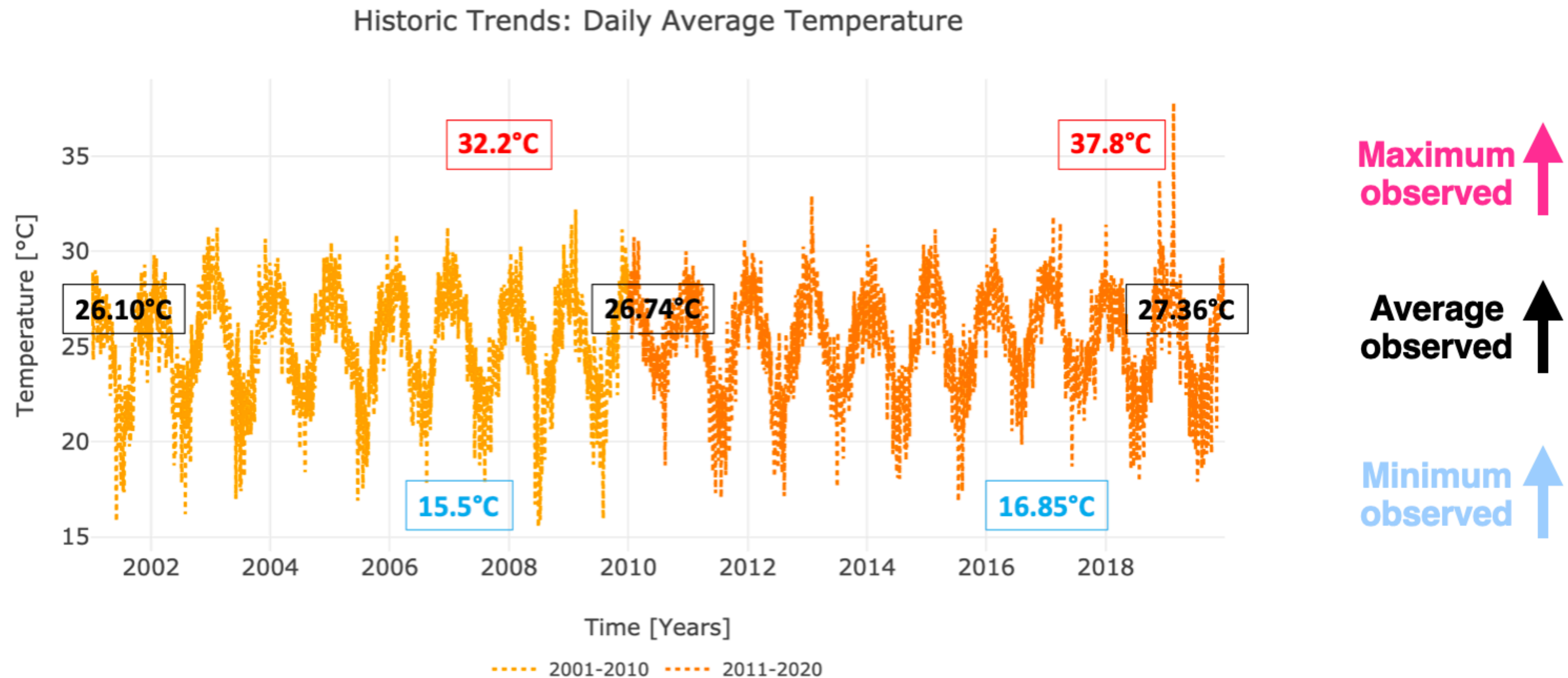
SOFTWARE TO ELUCIDATE THE TEMPORAL RESOLUTION OF TEMPERATURE EFFECTS





# REQUIREMENT: EXTRAPOLATE TO NEW SCENARIOS

SOFTWARE TO EXAMINE INTERVENTION EFFICACY UNDER FUTURE TEMPERATURES





# SOFTWARE DESIGNED TO MEET RESEARCH REQUIREMENTS

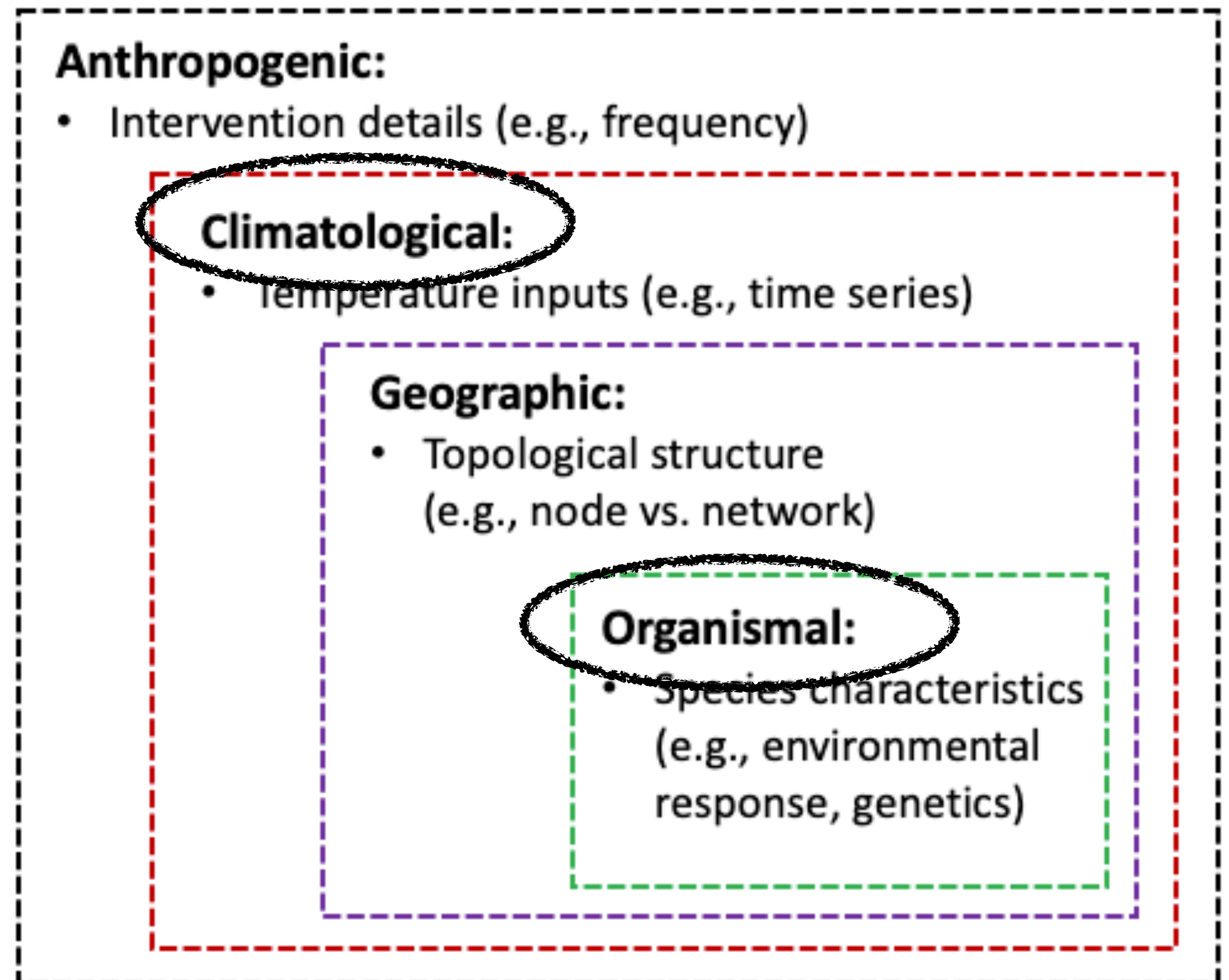
## GENEDRIVE.JL: THE DATA MODEL

### In principle:

- Straightforward to change or add new information as it becomes known
- Composability enables co-evolution with scientific discovery

### In practice (for this research):

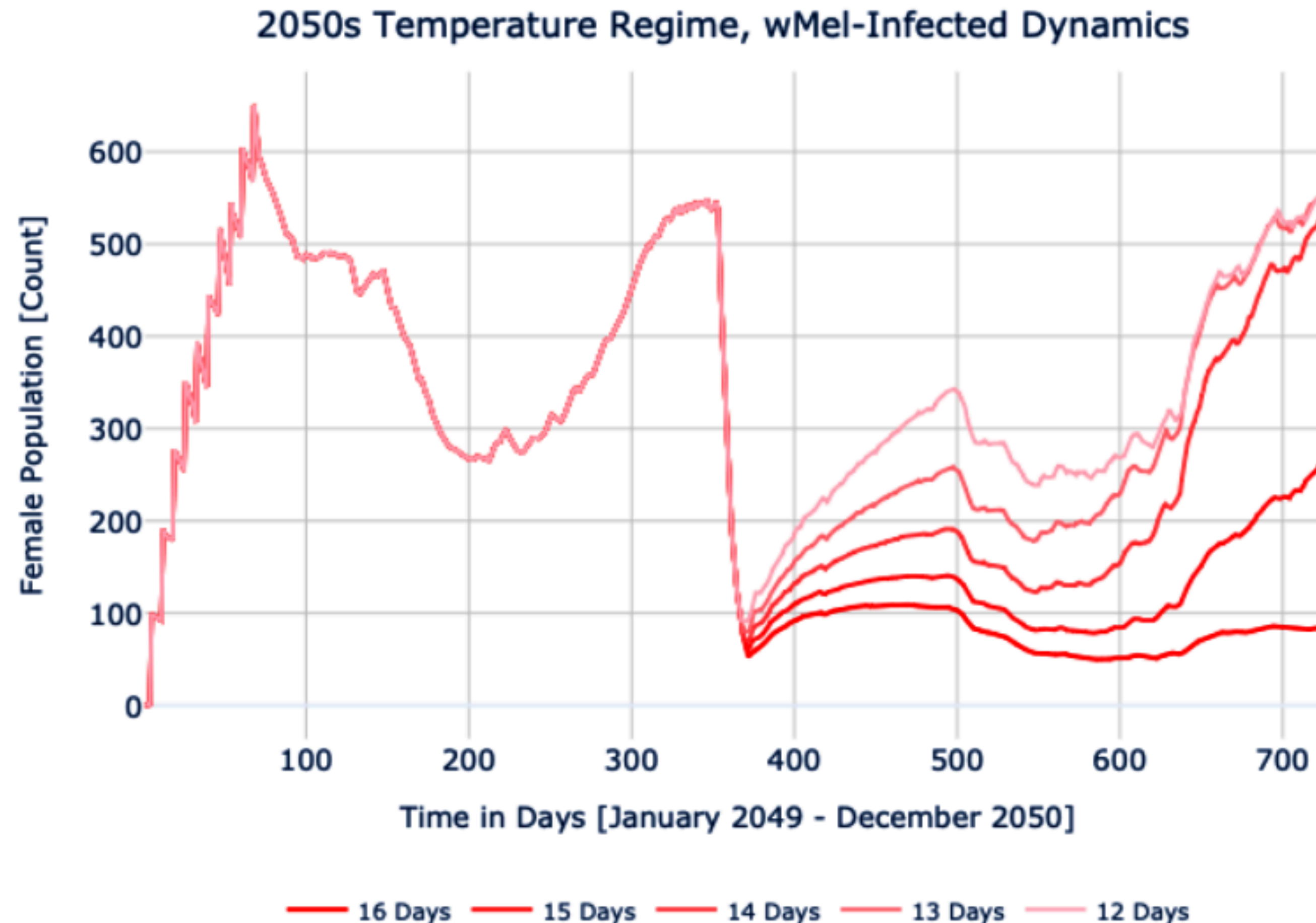
- Alternative functional forms: Wolbachia's thermal sensitivity, distinct from its *Ae. aegypti* host
- Alternative data inputs: Temperature time series for 2030s, 2050s, etc. (averages and heatwaves)





# FINDING: WOLBACHIA IS ROBUST TO FUTURE CLIMATES

## THERMAL BIOLOGY OF GENETIC-BASED INTERVENTION: WOLBACHIA (wMEL)







# RESEARCH AREA #2

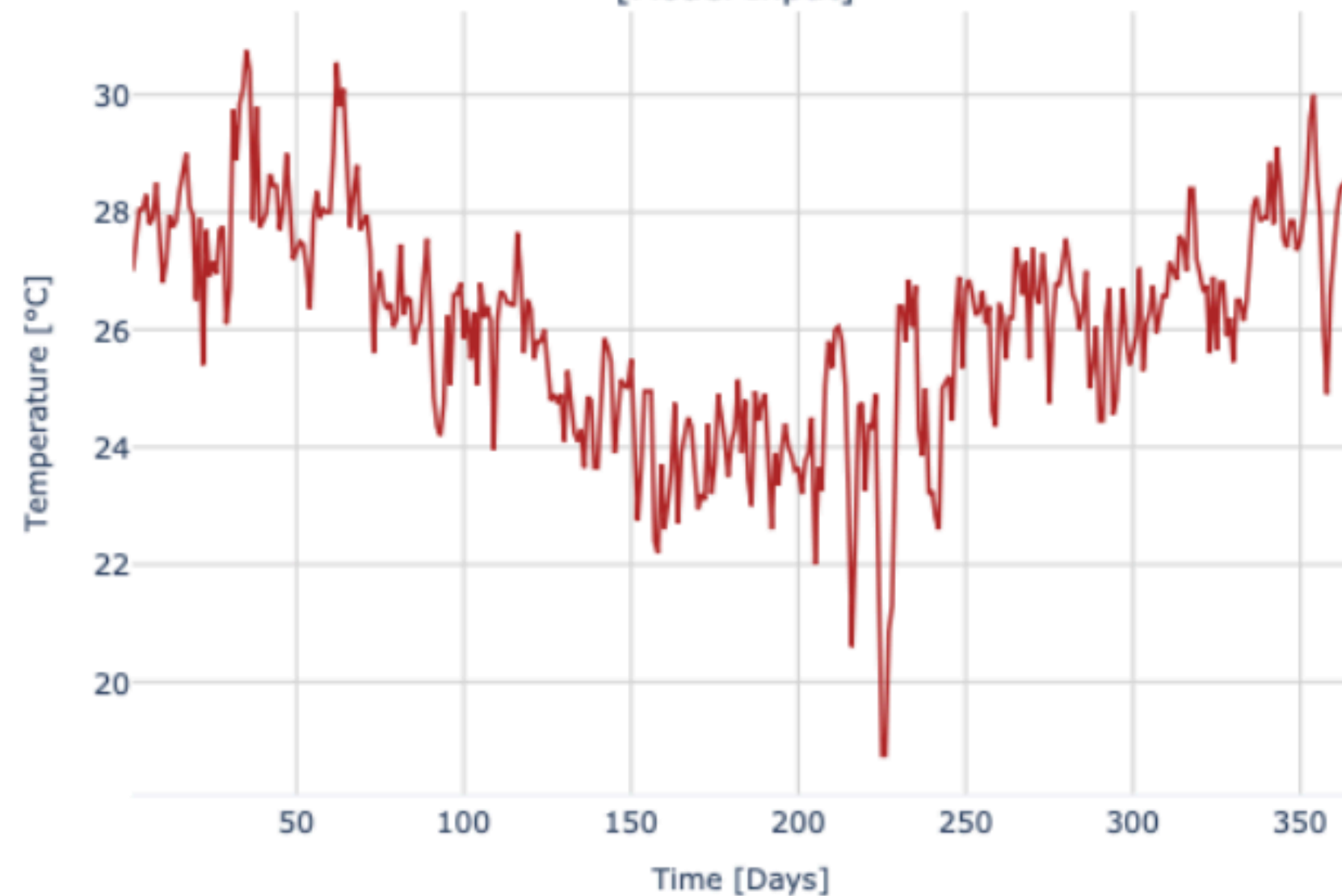


# STRATEGIC: WHAT IS THE “BEST” ALLOCATION OF TOOLS?

## DEFINING “BEST”: THE ECOLOGICAL CONSIDERATIONS

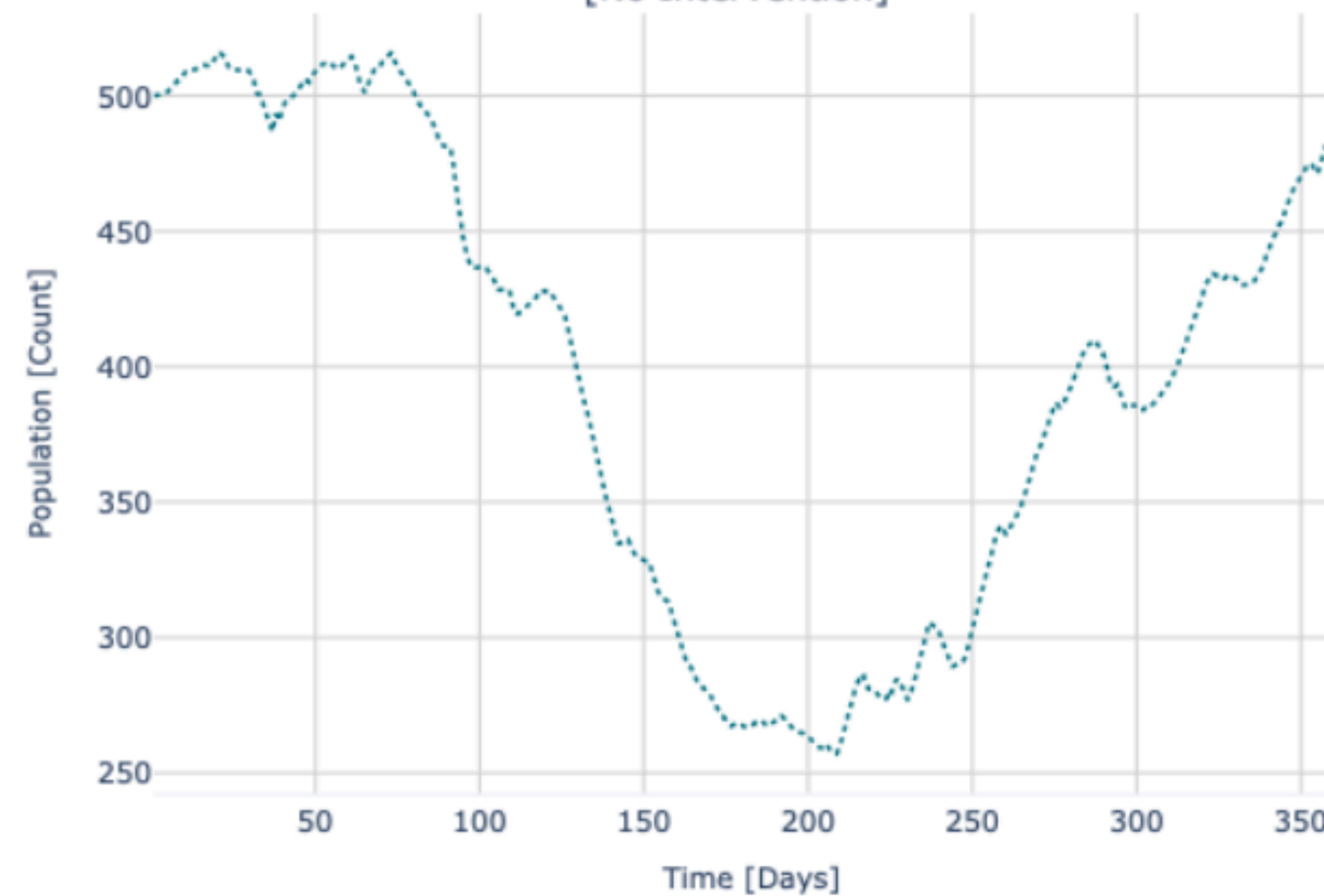
Location-specific factors are changing with shifting climates and land use practices (e.g., urbanization).

Timeseries of Daily Temperature  
[Model Input]



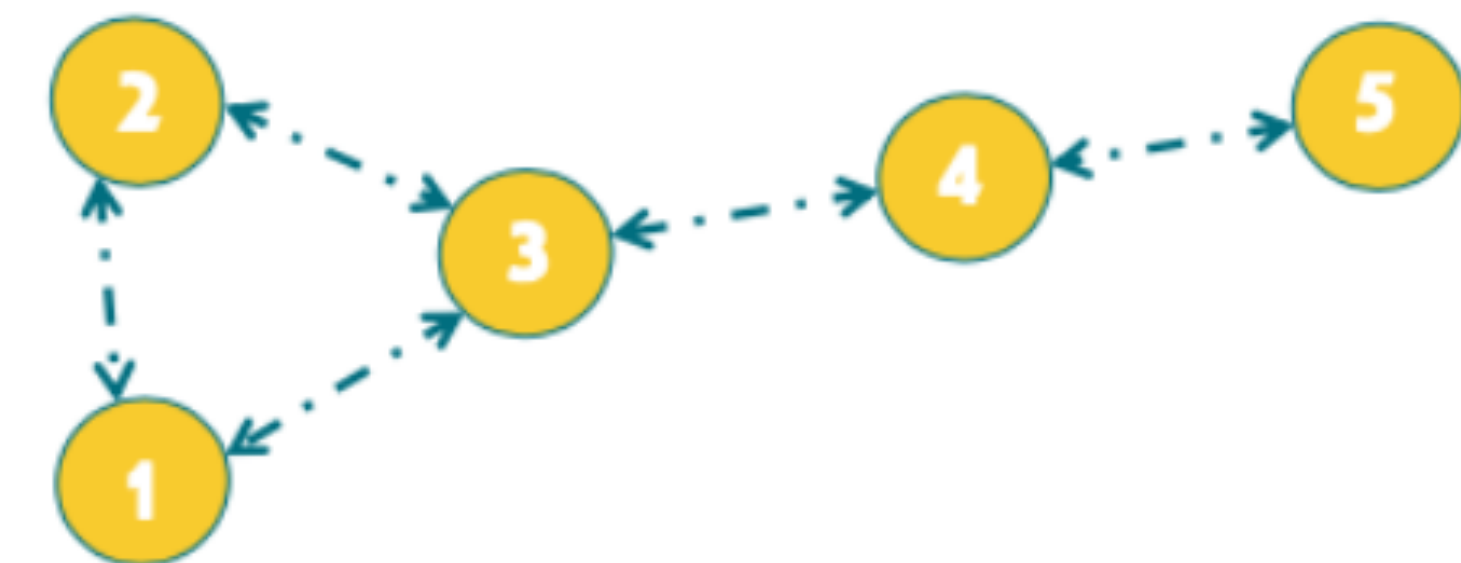
Seasonal, daily temperatures in the location of interest

Baseline: Wild Aedes Aegypti Female Dynamics  
[No Intervention]



Species lifecycle and adaptations in that location

Topology of the 5-Node Network



Geographic connectedness, spatial distribution



# STRATEGIC: WHAT IS THE “BEST” ALLOCATION OF TOOLS?

## DEFINING “BEST”: THE HUMAN HEALTH AND ECONOMIC CONSIDERATIONS

- The efficacy of a given public health intervention translates to lives saved and illness prevented.
- Efficiency gains also equate to cost reductions: deployment demands resources
  - ❖ Potentially fewer or more infrequent trips to the field (labor cost)
  - ❖ Smaller numbers of organisms released (material cost)
- Community comfort factor?

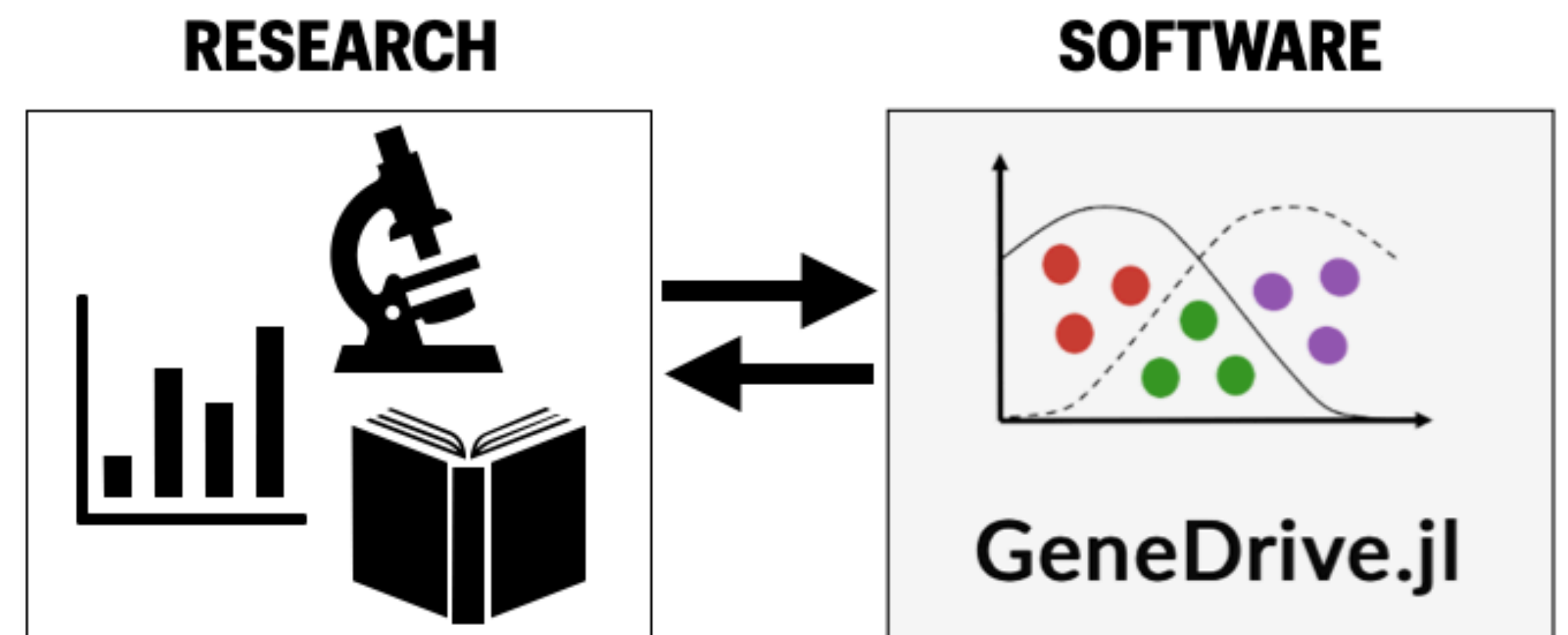


# EVALUATING WHAT STRATEGY IS “BEST”

## THE CASE FOR SOFTWARE

- **Again:** Empirical research is essential; both laboratory and field studies furnish fundamental insights.
- But models help evaluate findings in the **necessary context (e.g., alternative definitions of “best”)**.
- Models facilitate hypothesis testing:
  - ❖ **Explore differences** in goals (objectives)
  - ❖ **Extrapolate** to include new scenarios

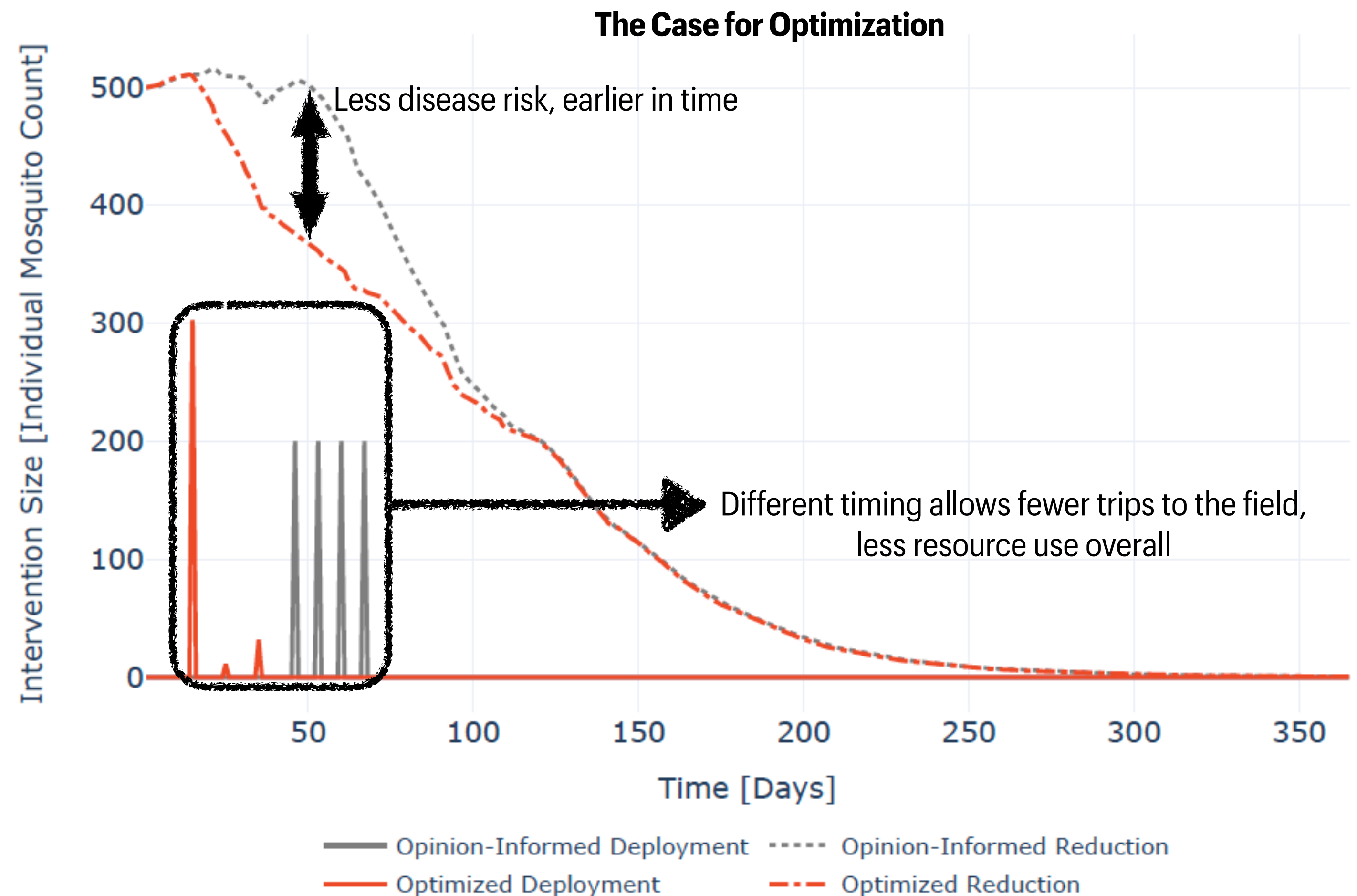
Coherence within a single framework



# REQUIREMENT: IMPROVE ON THE STATE OF THE ART

## SOFTWARE TO MAXIMIZE AND ITERATE OBJECTIVES, CONSIDERING RELEVANT DETAILS

- Presently, interventions designed based on the results of field work & expert opinion.
- Seeking:
  - ❖ A cheaper, more reproducible way to strategize
  - ❖ Ability to update when details change (as they do)
- Answer: optimization?





# SOFTWARE DESIGNED TO MEET RESEARCH REQUIREMENTS

## GENEDRIVE.JL: THE DECISION MODEL

**Objective function:** Various. Here, minimize wild vector (females,  $F$ ) using minimal releases (genetically modified,  $c_{\hat{g}t}$ )

**Equality constraints:** Lifecycle dynamics

**Parameterization:** Ecological and genetic details

**Inequality constraints:** Operational restrictions on deployments of  $c_{\hat{g}t}$

$$\min_{c_{\hat{g},t}} J(F_{n,g,t}, \alpha c_{\hat{g},t})$$

$$E_{g,t,i} = E_{g,t,i-1} - E_{g,t,i}(\mu_E + q_E i_E)$$

$$c_{\hat{g},t} \leq R_{n,t} D_{max}^{day} \quad \forall t \in \mathcal{T}_d,$$

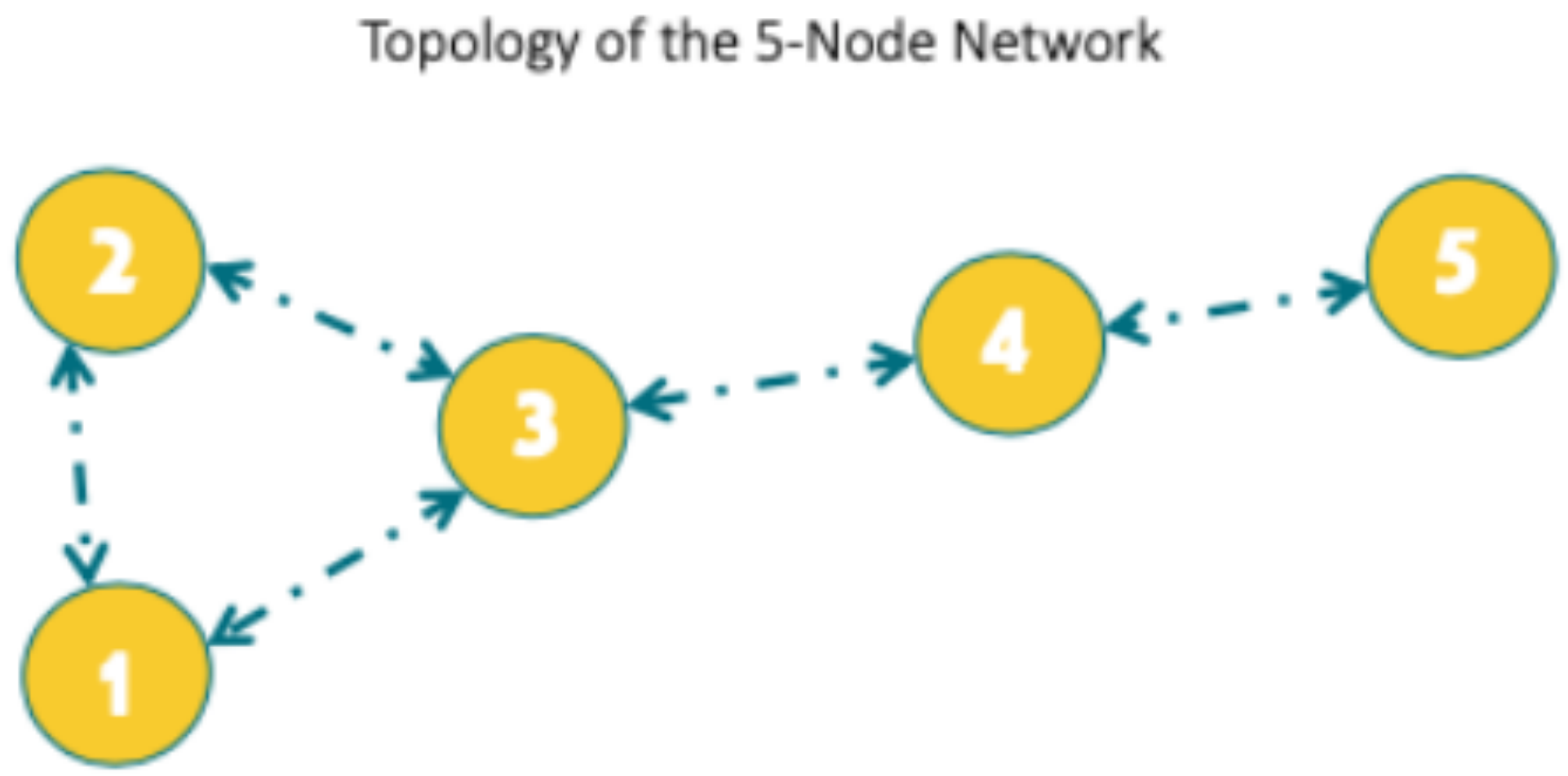
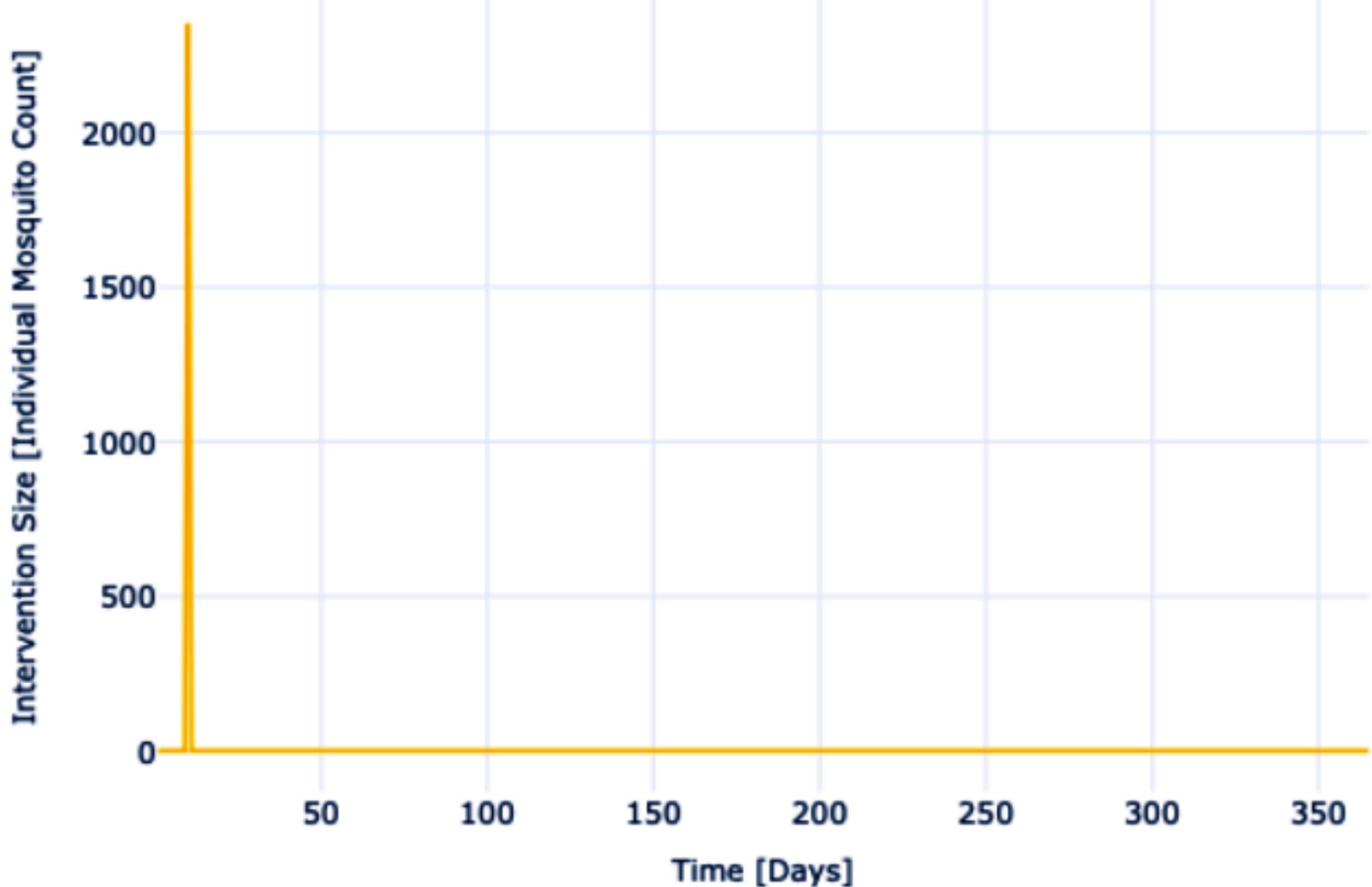
$$c_{\hat{g},t} \geq R_{n,t} D_{min}^{day} \quad \forall t \in \mathcal{T}_d,$$

$$\sum_{t \in \mathcal{T}_d} c_{\hat{g},t} \leq D_{max}^{trial}$$

# FINDING: CAPABLE OF EXPLORING DIFFERENT OBJECTIVES

## RISK REDUCTION ACROSS ENTIRE REGION (NETWORK) OF INTEREST

Deployment Schedule Over One Year:  
Wolbachia (Adult Males)



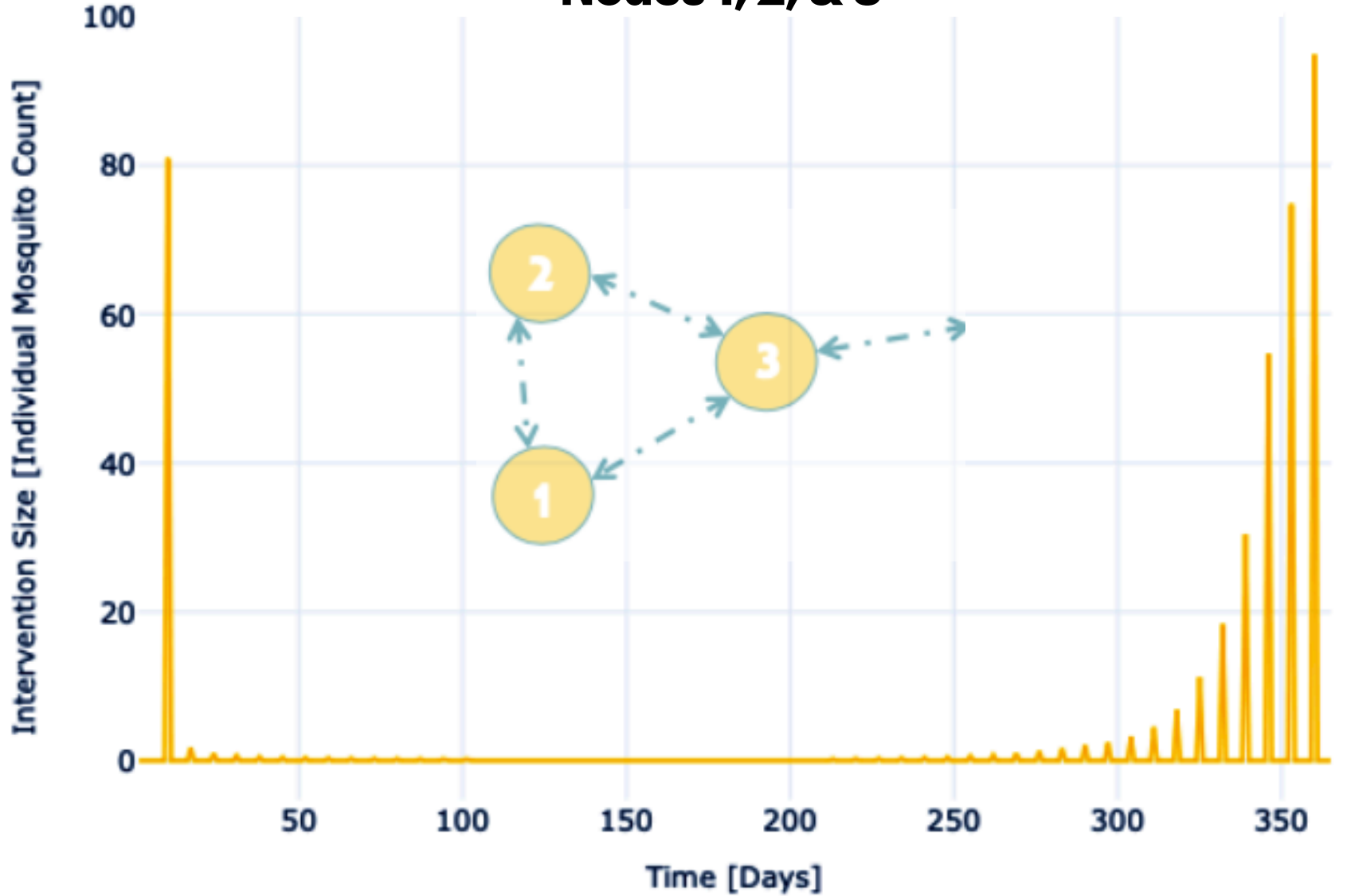


# FINDING: CAPABLE OF EXPLORING DIFFERENT OBJECTIVES

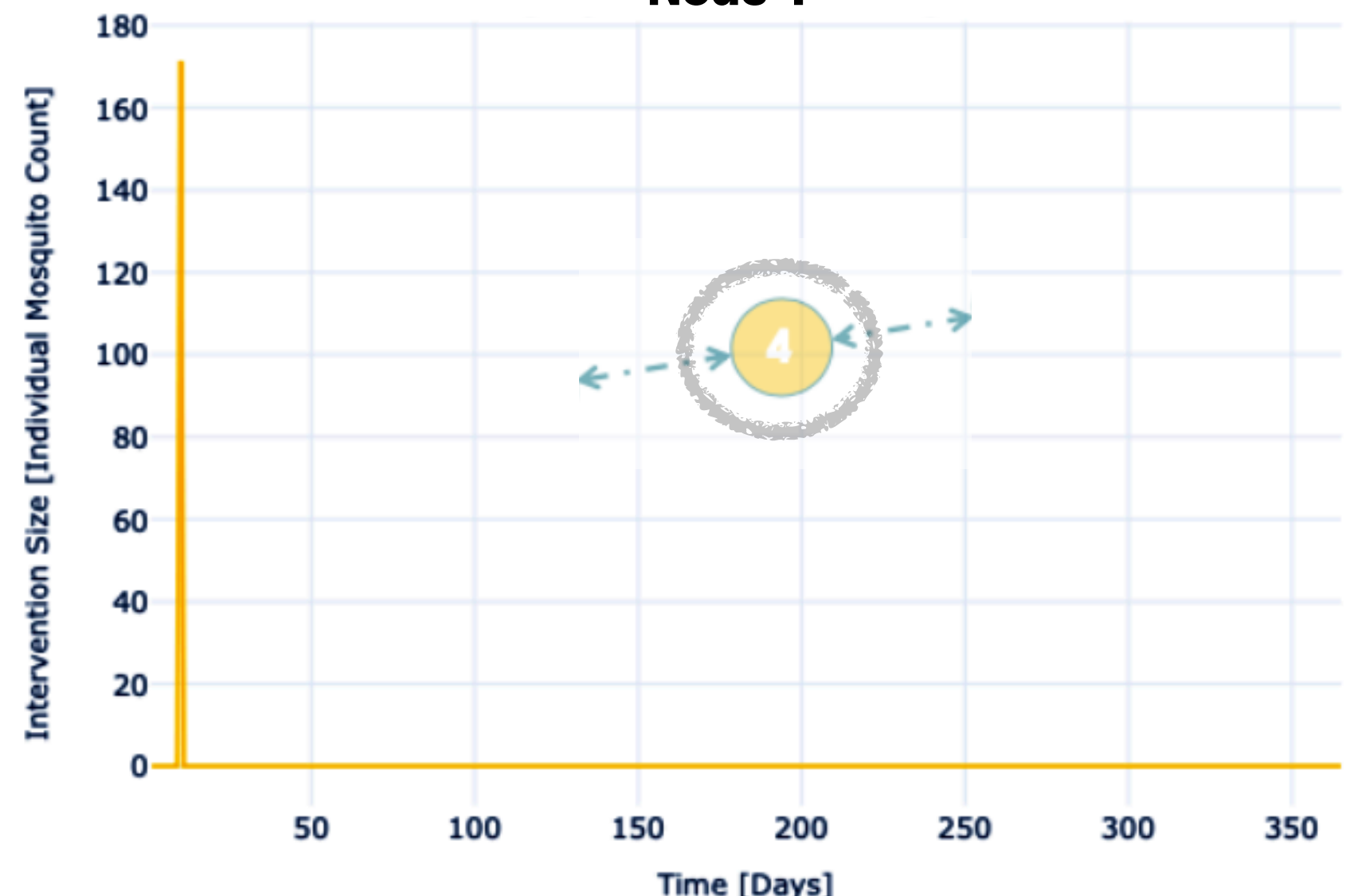
## RISK REDUCTION PRIORITIZING SINGLE LOCATION OF INTEREST IN A REGION (NETWORK)

Release policy in each area of the network is dependent on its connectivity to the node of interest.

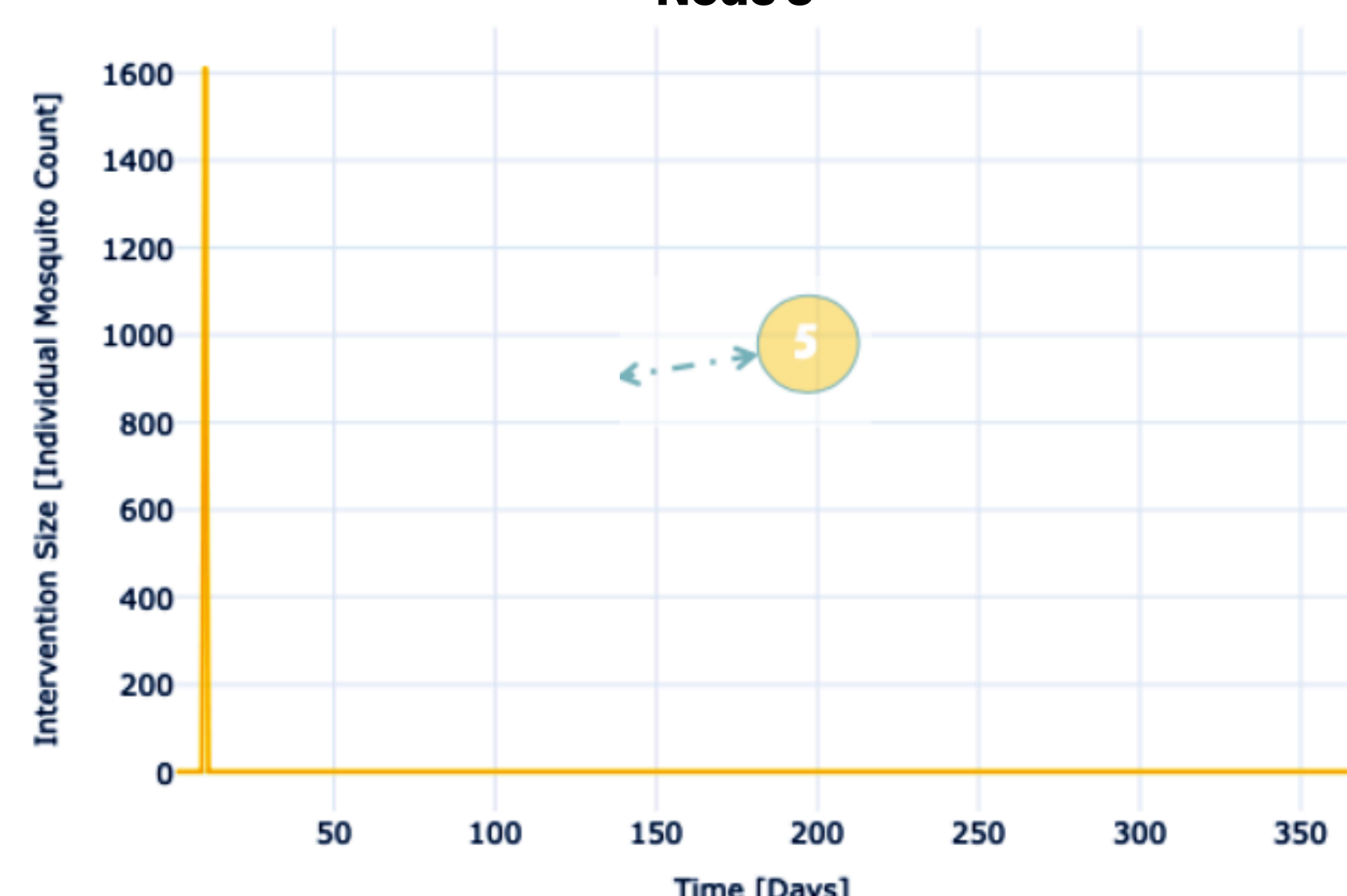
Deployment Schedule Over One Year:  
Nodes 1, 2, & 3



Deployment Schedule Over One Year:  
Node 4



Deployment Schedule Over One Year:  
Node 5

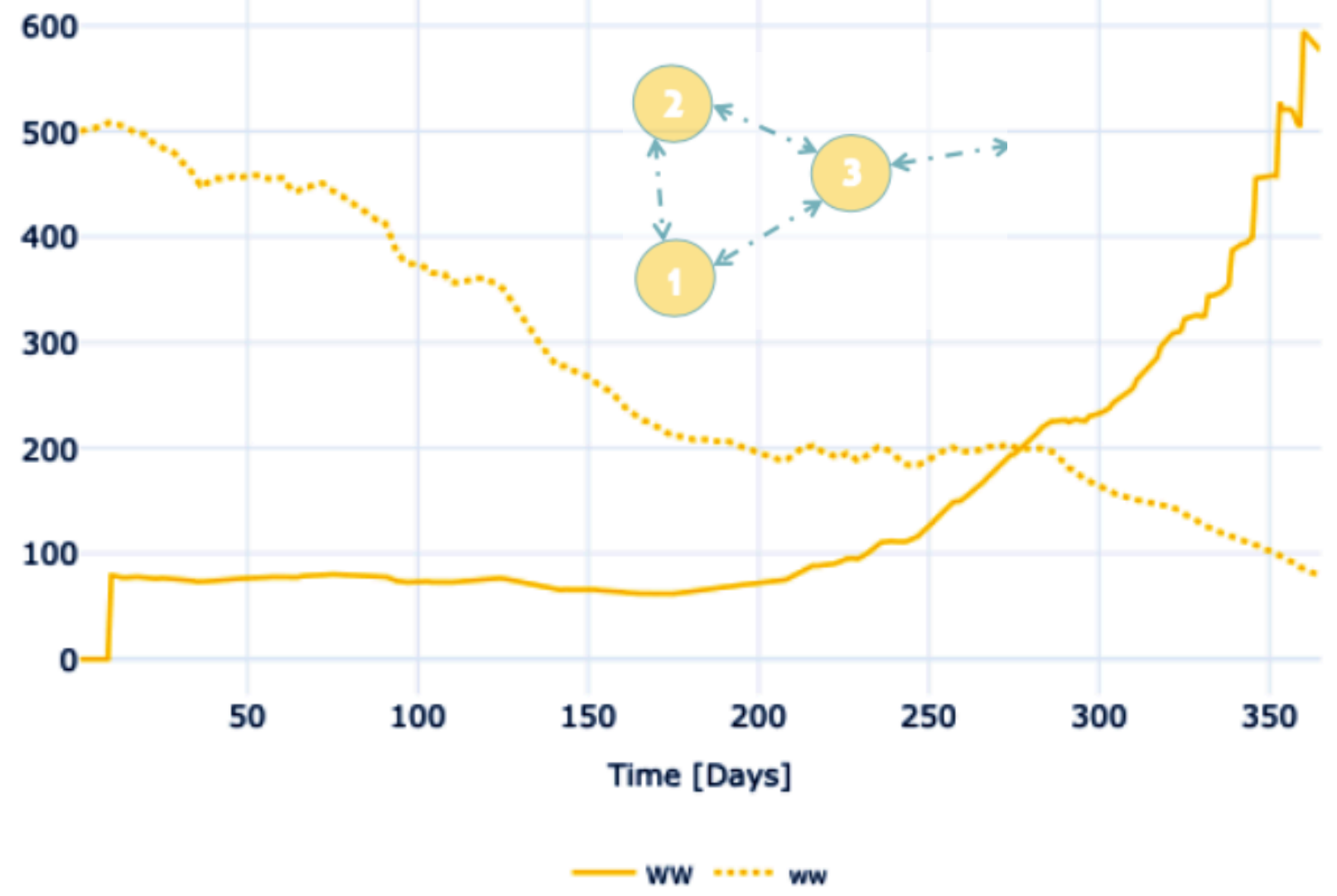


# FINDING: CAPABLE OF EXPLORING DIFFERENT OBJECTIVES

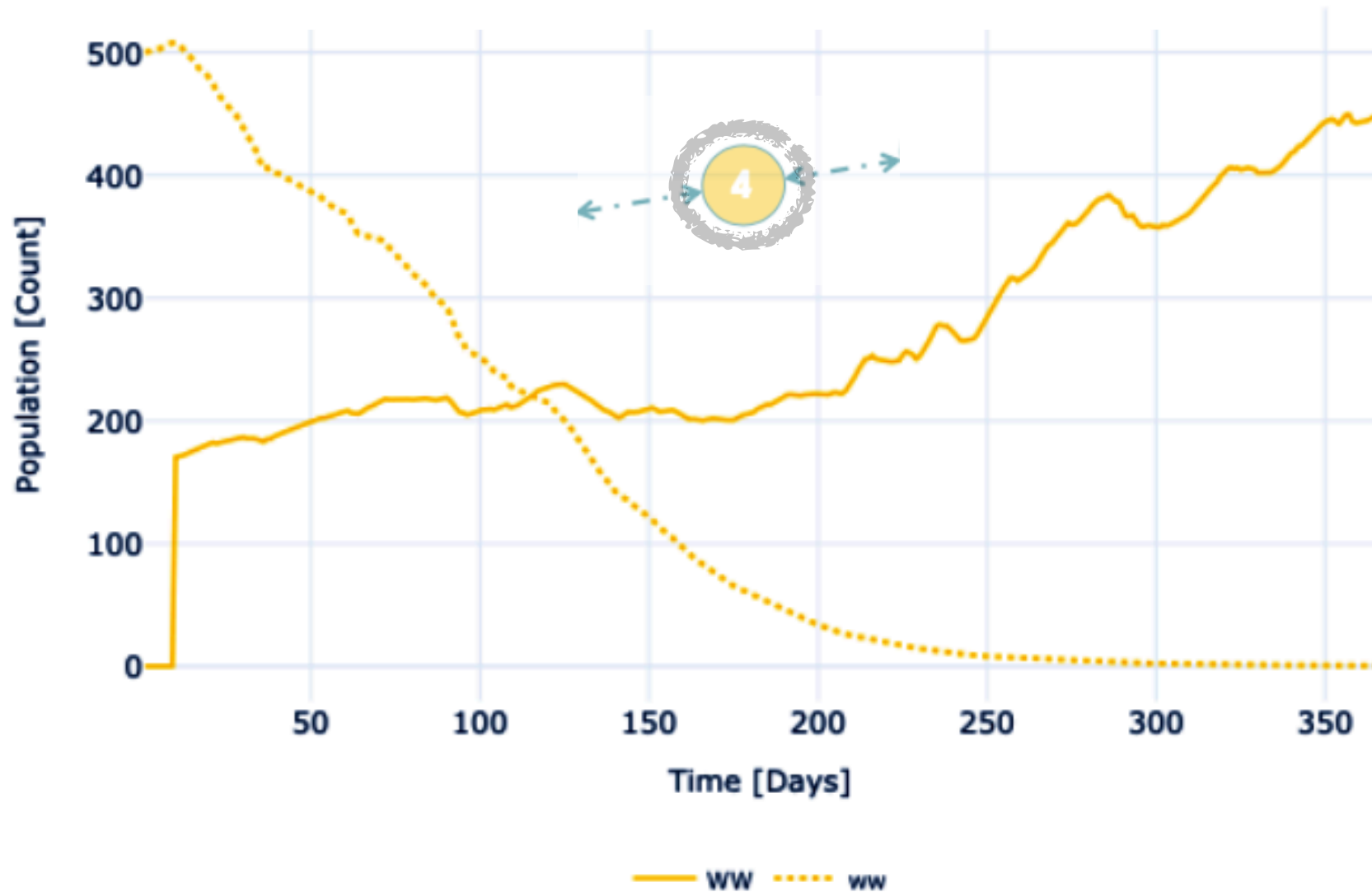
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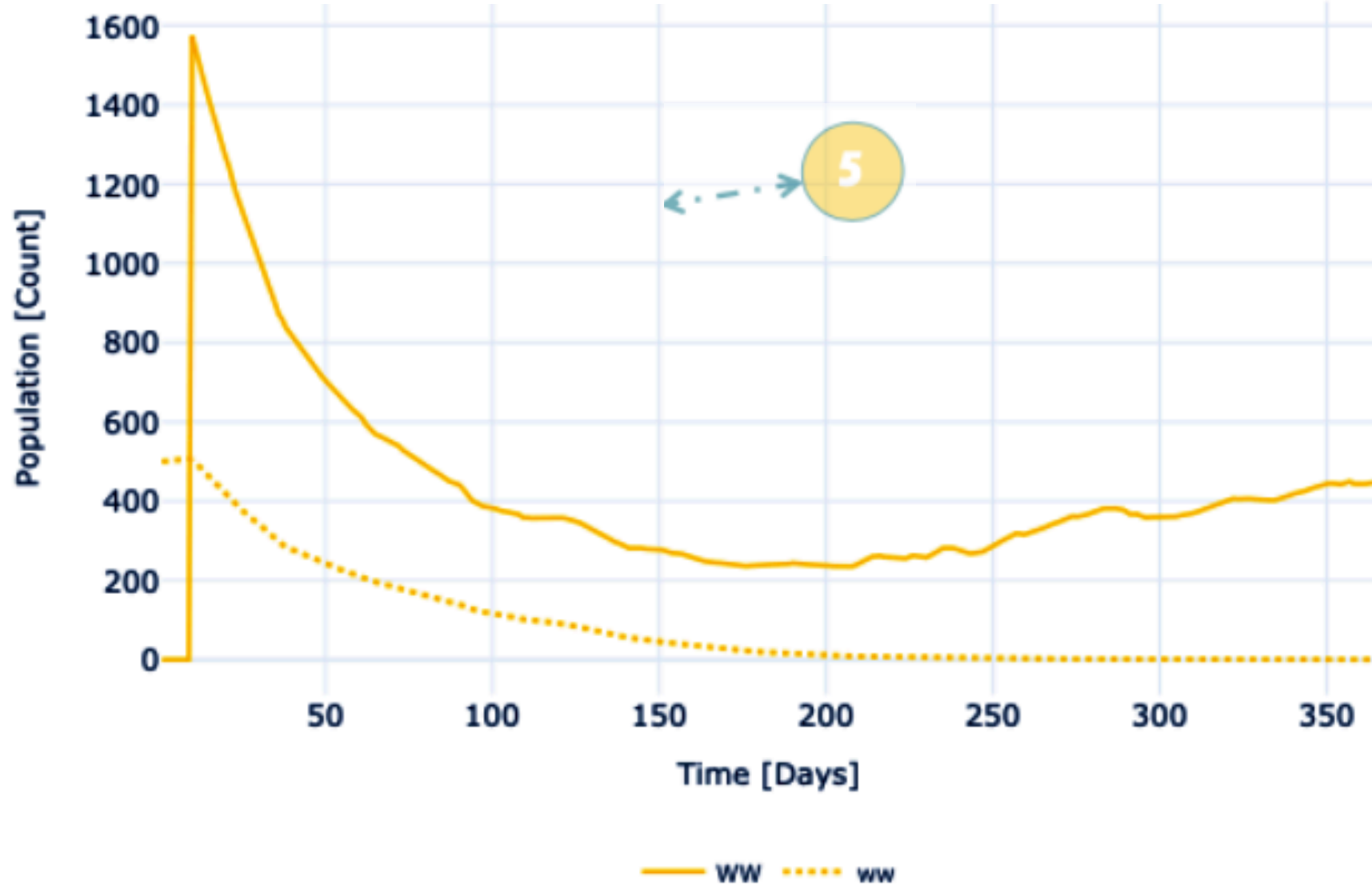
Corresponding Change in Disease Vector Over One Year: Nodes 1, 2, & 3



Corresponding Change in Disease Vector Over One Year: Node 4



Corresponding Change in Disease Vector Over One Year: Node 5

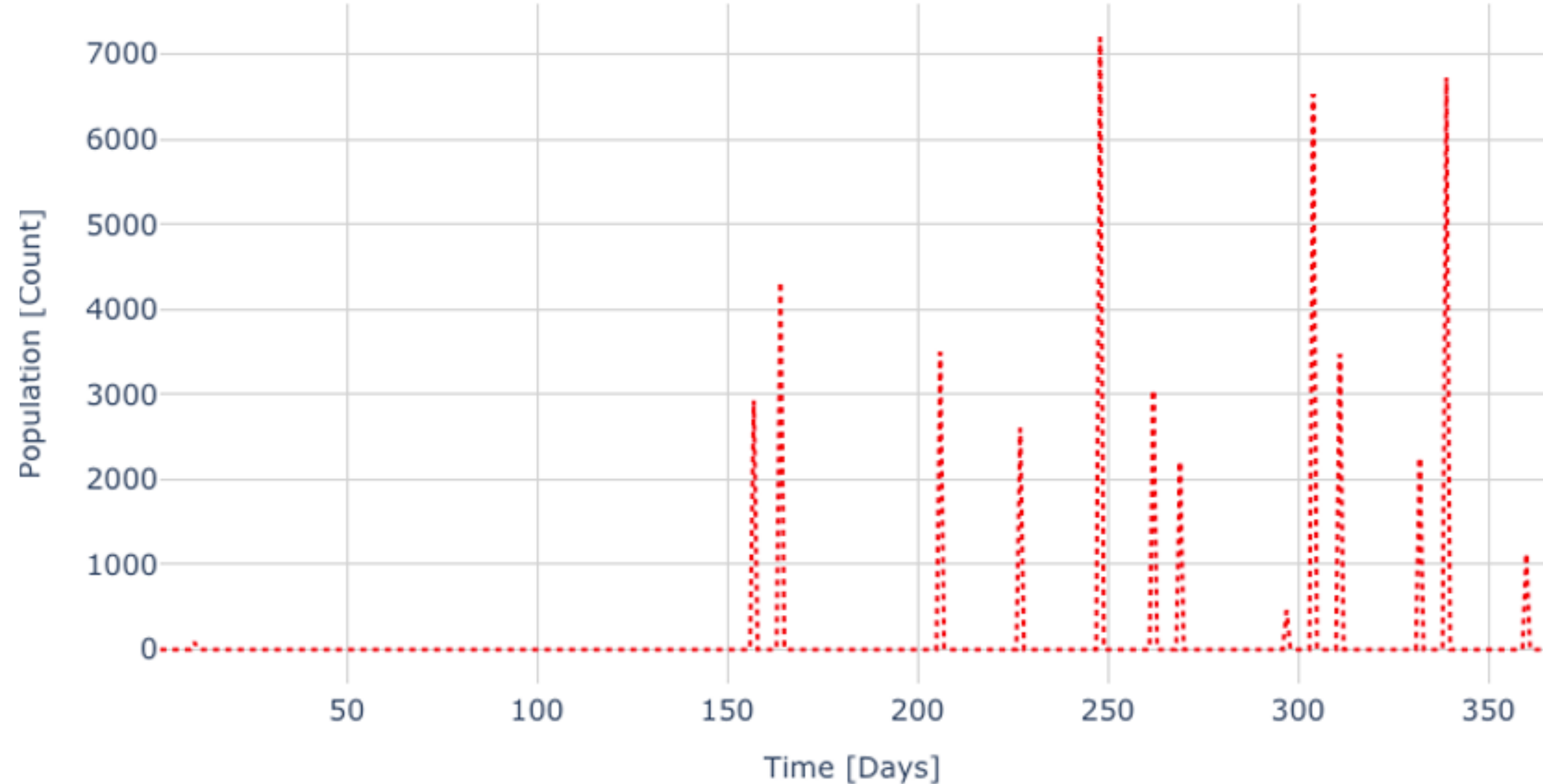




# FINDING: CAPABLE OF EXTRAPOLATING TO NEW SCENARIOS

ALTERED STRATEGY TO ACCOUNT FOR CHANGING CLIMATE

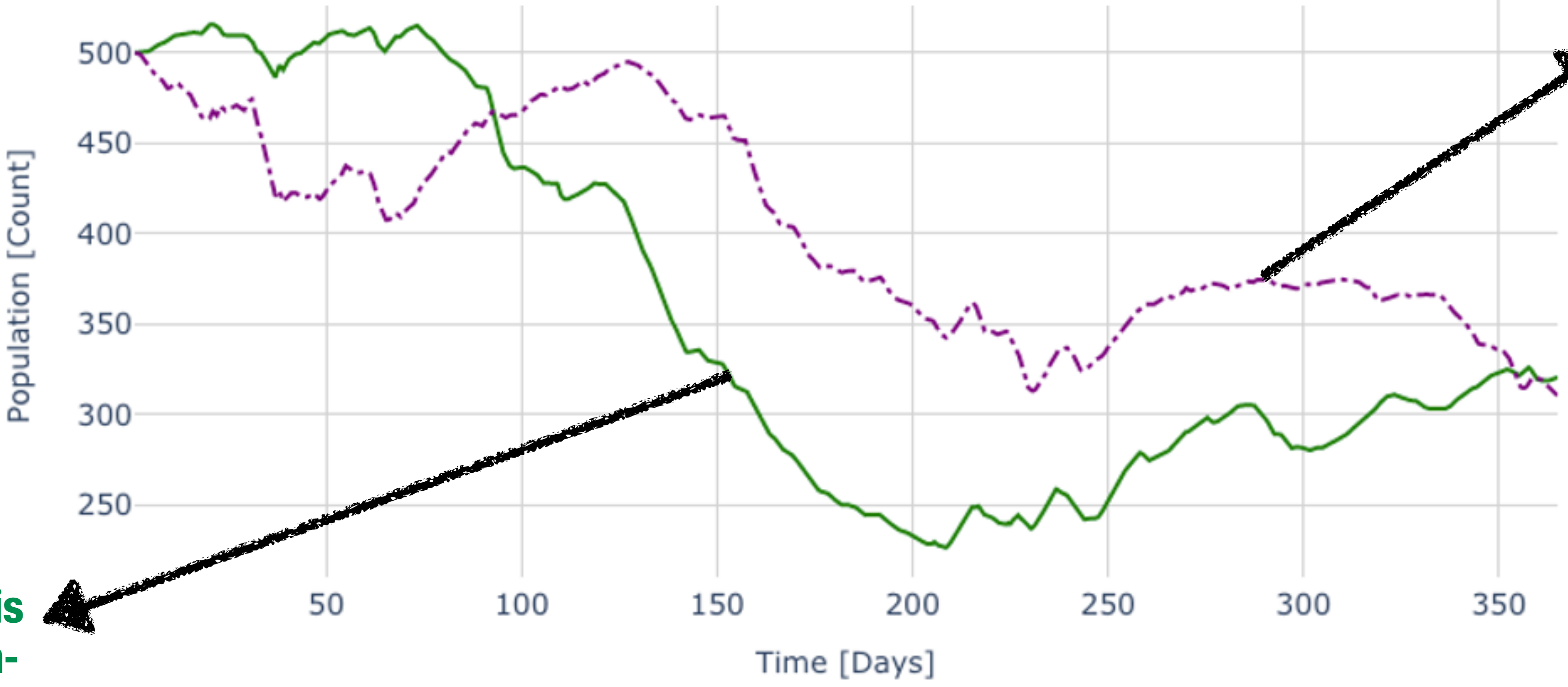
Schedule of Deployments Over One year (RIDL Males)



# FINDING: CAPABLE OF EXTRAPOLATING TO NEW SCENARIOS

## ALTERED STRATEGY TO ACCOUNT FOR CHANGING CLIMATE

Corresponding Change in Disease Vector Over One Year: The 2°C Difference



Result when strategy is optimized for location-specific temperature.

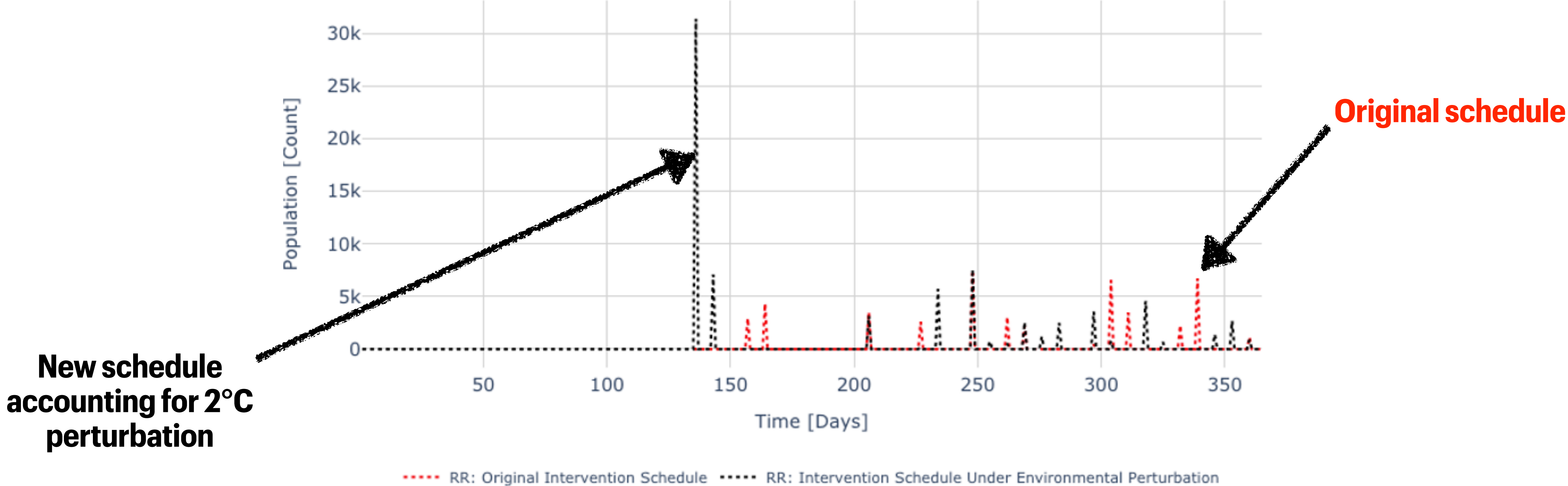
Result when annual average temperature increases 2°C, but same strategy is used.



# FINDING: CAPABLE OF EXTRAPOLATING TO NEW SCENARIOS

## ALTERED STRATEGY TO ACCOUNT FOR CHANGING CLIMATE

Schedule of Deployments Over One year (RIDL Males):  
The 2°C Difference





## **CURRENT FUNDING:**

- **MICROSOFT RESEARCH PHD FELLOWSHIP**

## **COLLABORATORS:**

- **LARA M. KUEPPERS, UC BERKELEY**
- **GORDANA RAŠIĆ, QUEENSLAND INSTITUTE OF MEDICAL RESEARCH (QIMR)**
- **JOHN M. MARSHALL, UC BERKELEY**
- **DAVID ANTHOFF, UC BERKELEY**
- **MICHAEL REDDY, MICROSOFT RESEARCH - HEALTH FUTURES**

# **THANK YOU**