

CIMMID: Climate integrated model of mosquitoborne infectious disease

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Managed by Triad National Security, LLC, for the U.S. Department of Energy's NNSA.



- 700 million cases of mosquito-born ID a year globally
- Highly sensitive to both climate change and landscape heterogeneity
- Fundamentally hard to model at largescale



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 - Isolate static landscape effects from secular trend in environmental data
- Locally calibrated climate model

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Los Alamos National Lab

DD-STE: Science, Technology & Engineering

ALD-SC: Scientific Computing

T: Theoretical Division

Lab Mission

<u>Team</u>

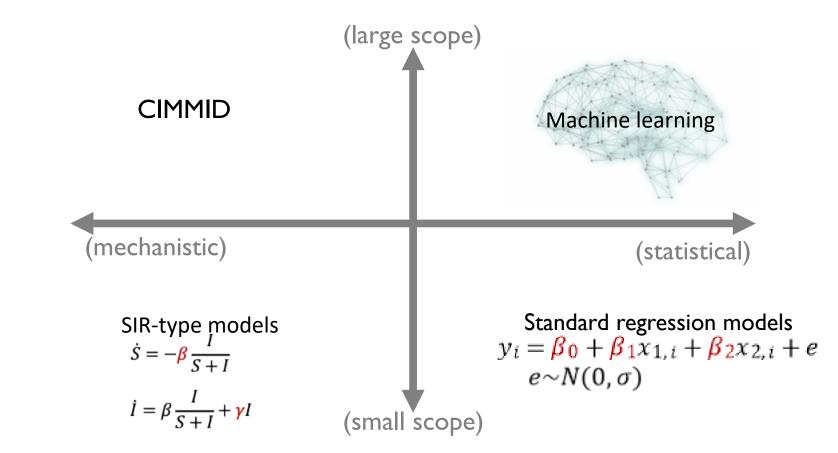
- ~20 staff
- 5-15 students
- Mostly domain experts
- PI: Dr. Carrie Manore

T6: Theoretical Biology and Biophysics

"We serve the nation by applying worldchanging science and technology to current and emerging national and global security challenges."

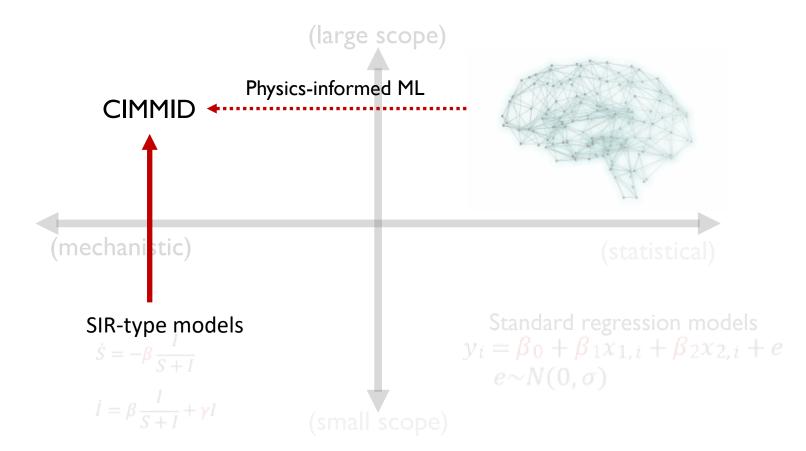


Modeling Philosophy





Modeling Philosophy





CIMMID Workflow

Los Alamos

- 1. Select pathogen, region (e.g. North America), and time frame.
- 2. Divide space into unique, contiguous units based on their potential for supporting relevant mosquitos in (1).
- Calibrate climate model to historical data for each unit in (2) and predict env. data to end of the study period.
- 4. Predict mosquito density dynamics overstudy time frame given (3).
- 5. Predict human risk given (4). Human risk model

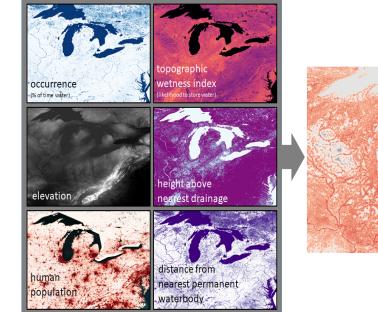


Hydrological-population units (hydropop units or HU)

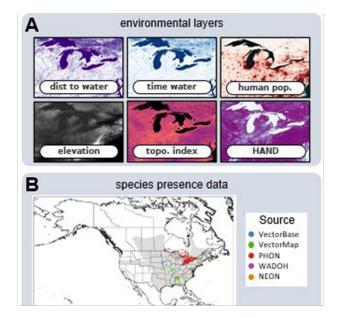
Earth System Model calibration

Process Based Model of mosquito development (PBM)

Mosquito abundance ~ landscape features + environmental variables (static) (dynamic)

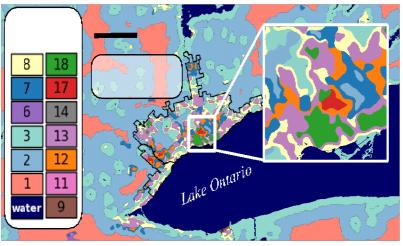


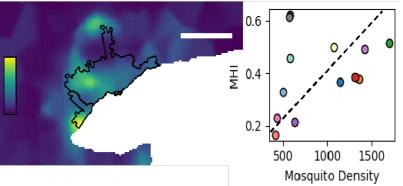


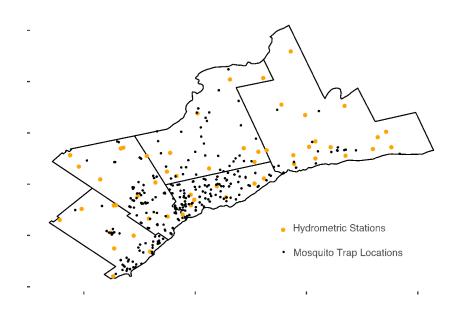




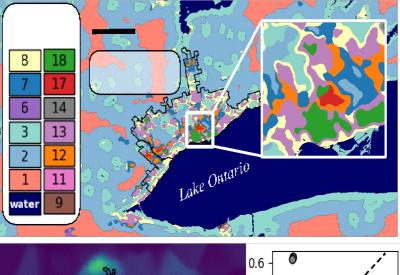
- Compute spatial Mosquito Habitat Index (MHI)
 - Satellite data: Static landscape features
 - Topology
 - Hydrological properties
 - Elevation
 - ..
 - Mosquito presence data
- Cluster values of MHI with human population density to define Hydropop Units (HUs)
 - HUs define long-term stable "homogeneity" regions
- Find contagious spatial units where MHI are all in the same cluster
 - Additional constraints (e.g. max/min HU area)

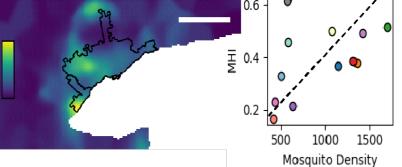




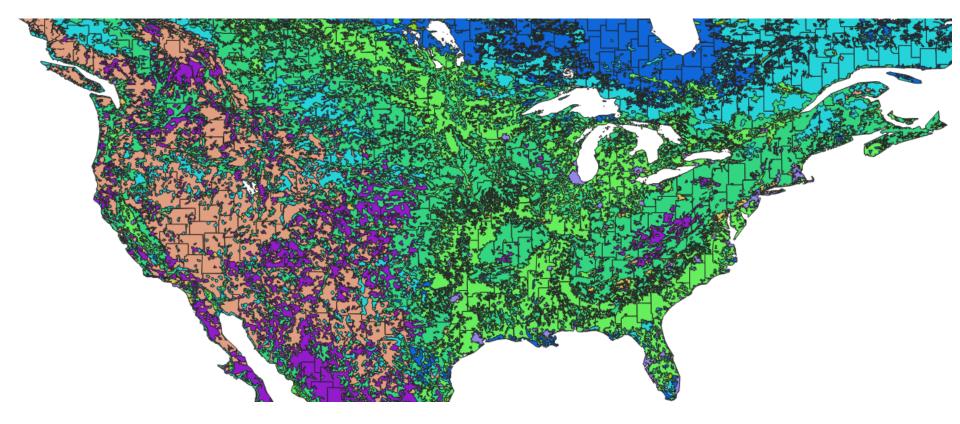


HUs facilitate simulation in regions with limited data







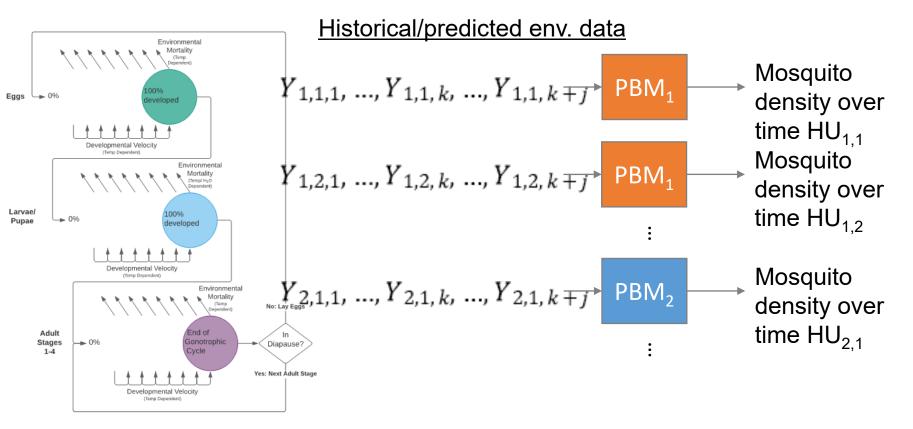




CIMMID core tech: Earth System Model (ESM) calibration

Historical env. dataPredicted env. data• Computational
bottleneck
$$HU_{1,1}$$
 $Y_{1,1,1}$, ..., $Y_{1,1,k}$ $ESM_{1,1}$ $Y_{1,1,k+1}$, ..., $Y_{1,1,K}$ • Limits spatial
resolution of HUs $HU_{1,2}$ $Y_{1,2,1}$, ..., $Y_{1,2,k}$ $ESM_{1,2}$ $Y_{2,k+1}$, ..., $Y_{2,K+j}$ • Emulator $HU_{2,1}$ $Y_{2,1,1}$, ..., $Y_{2,1,k}$ $ESM_{2,1}$ $Y_{2,1,k+1}$, ..., $Y_{1,1,K}$ • Exploit spatial
structure $HU_{2,1}$ $Y_{2,1,1}$, ..., $Y_{2,1,k}$ $ESM_{2,1}$ $Y_{2,1,k+1}$, ..., $Y_{1,1,K}$ • Exploit spatial
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structure $HU_{2,1}$ $Y_{2,1,1}$, ..., $Y_{2,1,k}$ $ESM_{2,1}$ $Y_{2,1,k+1}$, ..., $Y_{1,1,K}$ • ELM
homogeneity
regions

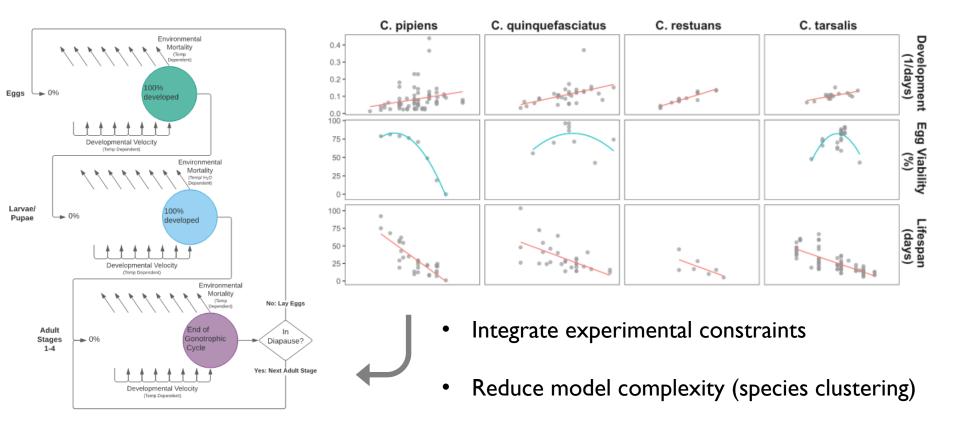
CIMMID core tech: Process-Based Model of Mos. Dev.





D P Shutt et al., A Process-based Model with Temperature, Water, and Lab-derived Data Improves Predictions of Daily Culex pipiens/restuans Mosquito Density, Journal of Medical Entomology, 2022

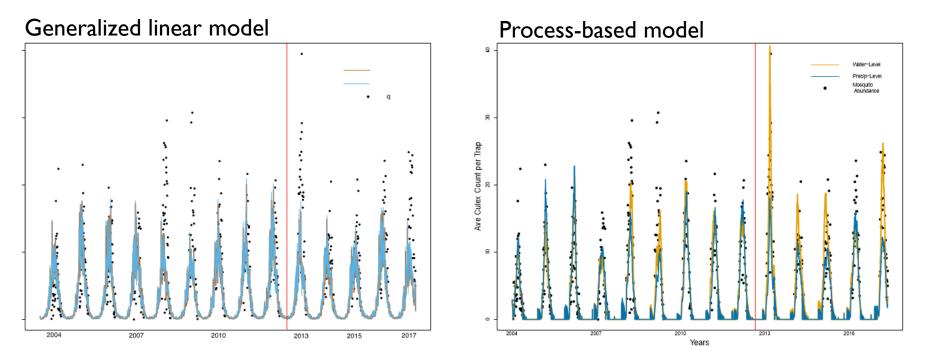
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CIMMID core tech: Process-Based Model of Mos. Dev.





 $\frac{dS_h}{dt} = -\lambda_h(t)S_h$ $\frac{dE_h}{dt} = \lambda_h(t)S_h - \langle \mathbf{A}E_h \rangle$ $\frac{dI_h}{dt}$ $\frac{dR_h}{dt} = \gamma_h I_h$ $\frac{dS_v}{dt}$ $= h_v(N_v)N_v - \lambda_v(t)S_v - \mu_v S_v$ $\frac{dE_v}{dt} = \lambda_v(t)S_v - \bigotimes E_v - \mu_v E_v$ $\frac{dI_v}{dt} = \langle \mathcal{A} E_v - \mu_v I_v.$



Human risk dynamics

 $= -\lambda_h(t)S_h$ $= \lambda_h(t)S_h - \langle \mathbf{X}\mathbf{E}_h \rangle$

 $= \gamma_h I_h$

=

$$h_v(N_v)N_v - \lambda_v(t)S_v - \mu_v S_v$$

$$= \lambda_v(t)S_v - \langle E_v - \mu_v E_v \rangle$$

 $= \quad \langle \mathcal{F}_v - \mu_v I_v.$

Mosquito risk dynamics



 dS_h

 $\frac{dt}{dE_h}$

 $\frac{dI_h}{dt}$

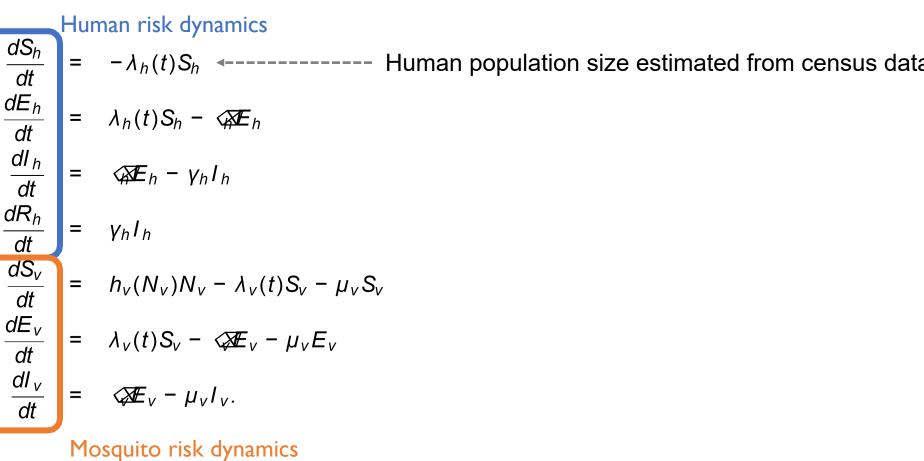
 dR_h

dt

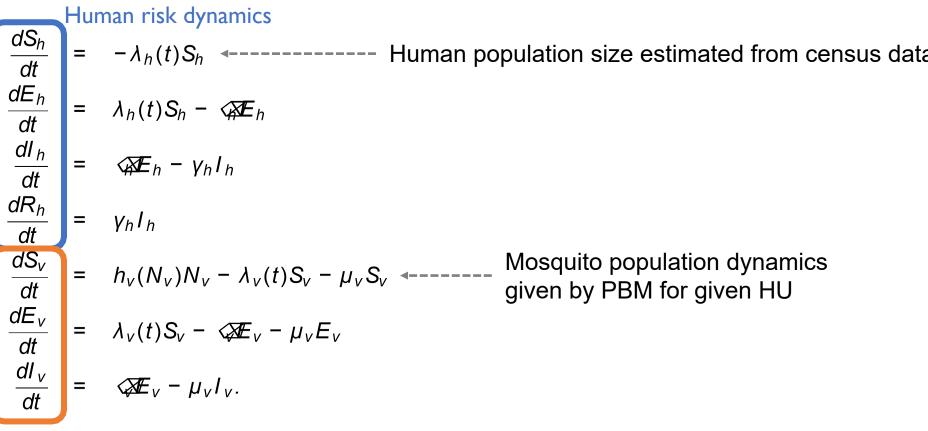
 dS_v

 $\frac{\frac{d}{dt}}{\frac{dE_v}{dt}}$

dt







Mosquito risk dynamics

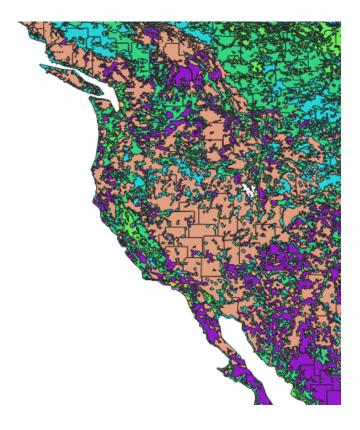
 $-\mu_v S_v$

Human risk dynamics dS_h = dt $\frac{dE_h}{dt}$ $\lambda_h(t)S_h - \langle \mathbf{x}\mathbf{E}_h \rangle$ = $\frac{dI_h}{dt}$ $\langle \mathbf{X} \mathbf{E}_h - \mathbf{y}_h \mathbf{I}_h$ = dR_h = Yh h dt $\frac{dS_{v}}{dt}$ $\frac{dE_{v}}{dt}$ $\frac{dU_{v}}{dt}$ $h_v(N_v)N_v - \lambda_v(t)S_v$ = $\lambda_v(t)S_v - \langle E_v - \mu_v E_v \rangle$ = $\langle \mathcal{F}_v - \mu_v I_v.$ = dt

- Force of infection terms
 - Overall biting rate
 - Generally known or constrained
 - Behavioral ecology of biting
 - Periodicity estimated from historical case count data
 - Coupling between HUs
 - Birds for West Nile Virus
 - Infected humans for Dengue

Mosquito risk dynamics

CIMMID: Bringing it all together



Using tons of different data we can

- Define local landscape heterogeneity
- Predict environmental variables under climate change
- Predict mosquito populations
- Predict human risk given human density and infected mosquito populations

Next steps

- Validating large-scale runs of CIMMID for WNV in US and Canada
- Integrating mitigation strategies into models to bring closer to "closed loop"



CIMMID core tech: Development status

ELM calibration Mosquito model Hydropop units Human risk model Count data Develop emulator Implement as partial Coupling for prototype diff equation WNV model Broader control ELM homogeneity Validation of Harden random parameters regions effects model for HUestimation for biting behavior Automatic cluster Validation on class specific estimation detection known watershed Validation of historical systems Validation with A. case count predictions aegypting Mature Developing In-progress







CIMMID Team 2019





National Institutes of Health





LABORATORY DIRECTED RESEARCH & DEVELOPMENT

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manore@lanl.gov



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