

# Teaching Modeling with Mosquito-Borne Disease Epidemics\*

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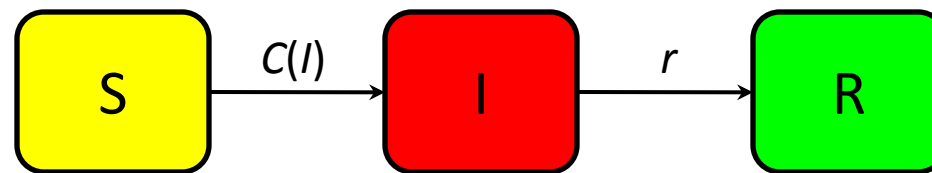
\*Part of the Symbiosis Project funded by the Howard Hughes Medical Institute

## Overall Goals

- To create a *context* which fosters the symbiosis of biology, statistics, and mathematics
  - Multiple entry points for many different students
  - Multiple activities possible and available
- To create a *context* that can be used with (and become familiar to) a wide range of students
  - Freshman use it superficially; Seniors more depth
  - An “ $F=ma$ ”-like alternative for the Life Sciences

# SIR Epidemics

- A population of constant size  $N$  is partitioned into  $S$  = Susceptibles,  $I$ =Infecteds,  $R$ =Recovered



- $C(I)$  = Contact rate of infection (as a function of  $I$ )
- $r$  = rate of recovery
- There are more sophisticated models
  - We stay simple for pedagogical purposes
  - Simple models are still highly applicable!

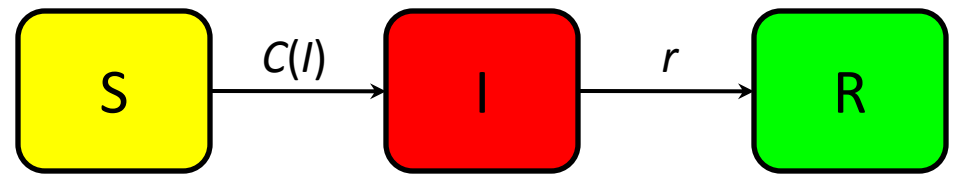
# SIR Epidemic Model

- SIR models are either Systems of DE's or Systems of Difference Equations

$$\frac{dS}{dt} = -C \langle \vec{S} \rangle$$

$$\frac{dI}{dt} = C \langle \vec{S} \rangle - rI$$

$$\frac{dR}{dt} = rI$$



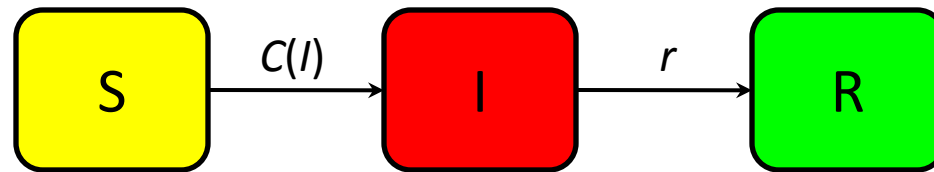
$$S_{t+1} = S_t \Delta t - C \langle \vec{S} \rangle_t$$

$$I_{t+1} = I_t \Delta t + C \langle \vec{S} \rangle_t - rI_t$$

$$R_{t+1} = R_t \Delta t + rI_t$$

# Types of SIR Epidemics

- Epidemic type often defined by how the disease is spread (Human-Human contact, Insect-borne, water borne, airborne, etc)



- Human-Human Contact:  $C(I) = b I$
- Mosquito-Human Contact:  $C(I) = c$

Assuming  
homogenous  
mixing of the  
populations

## Which Epidemic is it??

- There have been many instances where one epidemic type has been confused for another
  - Yellow fever is Mosquito-borne (Carlos Finlay, **1889**),
    - Proven to be Mosquito-borne in Cuba in **1900** ( Maj. Walter Reed). Experiment cost Jesse Lazear his life.
    - Practically no cases of Yellow fever in Cuba since **1901**
  - Yellow fever officially remains a human-human transmitted disease (via poor sanitation) until **1905**.
    - Last Yellow Fever epidemic, **New Orleans, 1905**
    - Extensive outbreak in spite of elaborate sanitation efforts



## Which Epidemic is it??

- This is still an Important Question – St. Louis Encephalitis, Dengue Fever, etc.
- And is a question we can explore
  - **Pedagogically:** Via Simulations (Netlogo) that generate data for each epidemic type (via models)
  - **Statistically:** Form Hypothesis, Analyze Data, Get  $p$ -value, Infer epidemic type
  - **Mathematically:** Why we should not be surprised that epidemic types are often confused?

# Pedagogically: The Simulations

- Simple S-I-R interactions among  $N$  agents
  - All Simulations: Recovery with a fixed rate  $r$  depending on how long an individual is ill (on average)
  - Human-Human: Infections with a given probability when a Susceptible meets an Infected
  - Mosquito-Human: Infections with a given probability when a Susceptible meets a Mosquito (assuming all Mosquitoes are infected)
- We simulate a “village” near a river (  $N = 500$  )

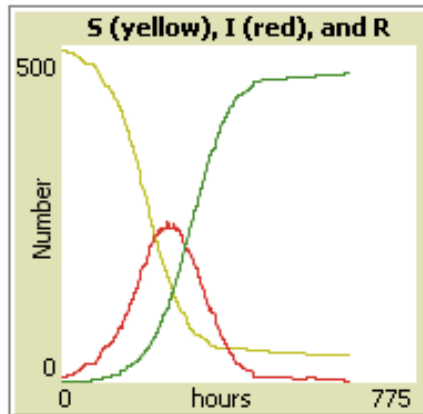


SetUp    Go    Step

TicksPerHour 8

ProbabilityOfInfection 0.28

A white susceptible (S) becomes a pink infected with "ProbabilityOfInfection" upon meeting an infected. Infecteds (I) turn red while symptomatic and then turn dark green once they recover (R).



# Pedagogically: The Simulations

- But there is more to the story
  - People work, eat, drink water during the day
  - People sleep at night ( violates homogeneity assumptions)
  - When they are sick enough, they want medical attention (hospital)
- Given a randomly-generated epidemic, can a student determine what type of epidemic it is?
  - Our “real world” is not so pristine as the models
  - But features of the models do “survive”

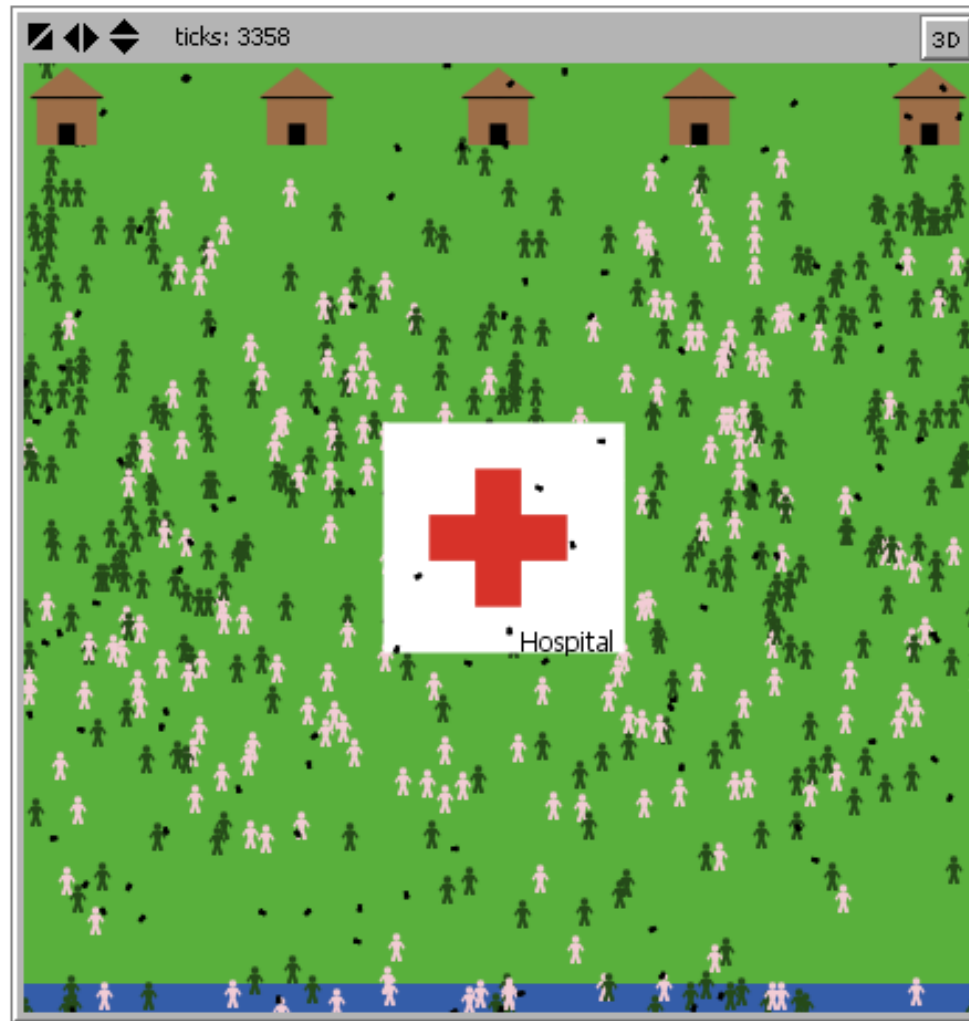
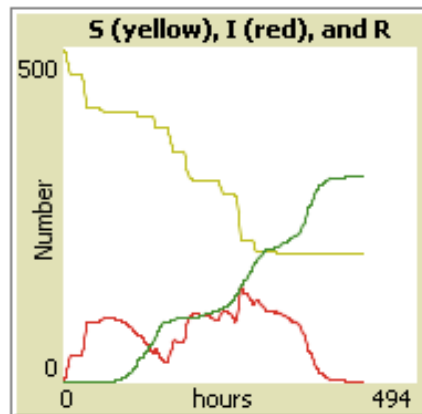
## Human-Human Epidemic

SetUp   Go   Step

TicksPerHour 8

ProbabilityOfInfection 0.28

A white susceptible (S) becomes a pink infected with "ProbabilityOfInfection" upon meeting an infected (I). Once symptomatic, infecteds turn red and head to the Hospital. Red infecteds turn dark green once they recover (R).



## Statistically: The Data

- Mosquito-Human epidemics are likewise set in “real world” circumstances
  - Model assumptions are violated
  - But same assumptions violated for all models!
- Although the models are developed mathematically, they are compared *statistically*
  - Especially given that assumptions are often violated
  - And that epidemics of different types tend to “look alike” (later)

## Mosquito-Human Epidemic

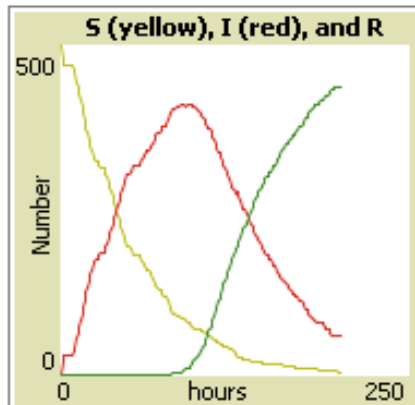
SetUp    Go    Step

TicksPerHour 8

ProbabilityOfInfection 0.28

On Reduce\_Mosquitos  
Off

Once bitten, a white susceptible (S) becomes a pink infected (I) with "ProbabilityOfInfection". Once symptomatic, infecteds turn red and head to the Hospital. Red infecteds turn dark green once they recover (R).



ticks: 1624    3D

Hospital

The simulation window shows a 3D view of a village with several houses and a central hospital marked with a red cross. The ground is green, and a blue river is at the bottom. Numerous human figures are scattered throughout the village. Some are white (susceptible), some are pink (infected), and some are red (symptomatic). A mosquito is visible near the top left. The top left corner shows navigation icons and the tick count (1624). The top right corner shows the 3D view toggle.

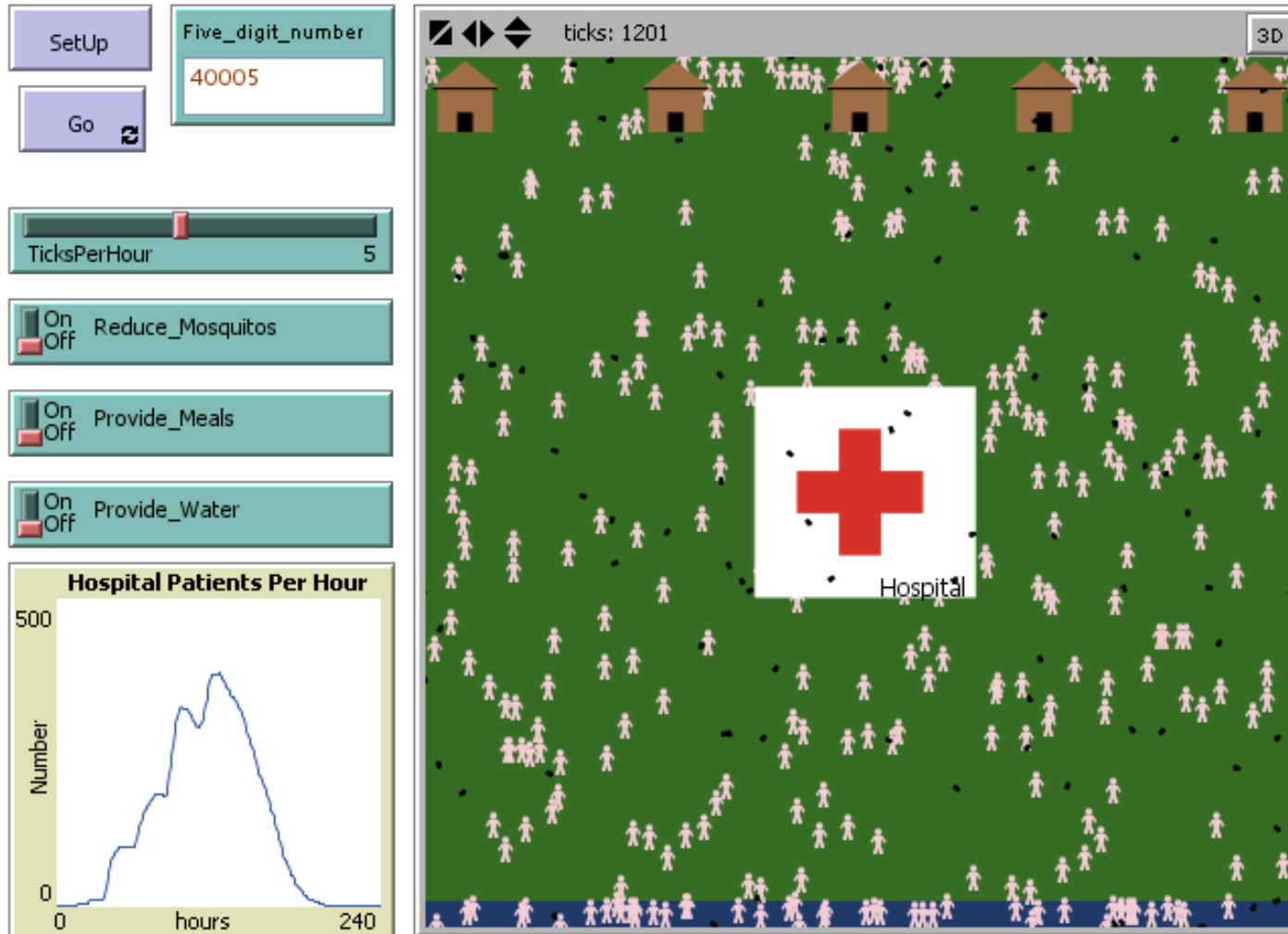


# Statistically: Analyzing The Data

- EpidemicVector simulation randomly simulates one of 4 epidemic types ( Human-Human, Mosquito-Human, Food Borne, Water Borne)
  - Based on a 5 digit number the student provides
  - They must determine which epidemic is generated
- Simulation mechanics:
  - People eat/drink during the day, sleep in huts at night
  - Once sufficiently ill, they go to the hospital
  - Only the patient count per hour is available for analysis



## Epidemic Simulations: 4 possible Methods of Transmission



# Statistically: Analyzing The Data

- Students can apply any of 3 interventions
  - Reduce the mosquito population
  - Provide untainted food to some of the population
  - Provide sterile water to some of the population
- Question: Which intervention reduces severity of the epidemic?
  - Simple approach: collect “max patient counts” in each simulation
  - Test for significance of difference between means of Experimental (with intervention) and Control
  - We have them do so with the randomization test

## The Randomization Test

Enter Data

Click 'Enter Data', click centers of disks, Click 'Enter Data' again to end entry.

Go Simulate

SimNum 500

Original Diff in Means  
25.8

Current Diff in Means  
7

Count	Trials
1	500
Proportion	0.0020

Proportion is the percentage of trial means whose magnitude exceeds that of the original.

Significant

InSignificant

Adata: 1 2 3 4 5 6 7

Bdata: 8 9 10 11 12 13 14

Load Data

ticks: 0 normal speed

Group	Mean	Values
Left	343	348, 358, 307, 341, 314
Right	336	371, 358, 348, 341, 344

Difference in Means Histogram

Number of means that occur in a given subinterval.-->

Subintervals of possible

powered by NetLogo

Result of Simulation with the data associated with the button **Significant**

## The Randomization Test

Enter Data

Click 'Enter Data', click centers of disks, Click 'Enter Data' again to end entry.

Go Simulate

SimNum 500

Original Diff in Means  
5.7

Current Diff in Means  
3.1

Count 237 Trials 1000

Proportion 0.237

Proportion is the percentage of trial means whose magnitude exceeds that of the original.

Significant

InSignificant

Adata  
1 2 3 4 5 6 7

Bdata  
8 9 10 11 12 13 14

Enter numbers separated by spaces into Adata and Bdata. Then press 'Load Data'.

Load Data

ticks: 0 normal speed

338	358	351	348
351	371	348	344
341	358	331	351
344	338	351	354
371	341	358	344

Mean = 351.1 Mean = 348

Difference in Means Histogram

Number of means that occur in a given subinterval.-->

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Result of Simulation with the data associated with the button **InSignificant**

# Mathematically: Linearization

- Mosquito-Human model is the *linearization* of the Human-Human model (in simple forms)

Human  
Human

$$\frac{dS}{dt} = -bIS$$

$$c = b(S_0 + S_0)$$

Mosquito  
Human

$$\frac{dS}{dt} = -cS$$

$$\frac{dI}{dt} = bIS - rI$$

$$\frac{dI}{dt} = cS - rI$$

$$\frac{dR}{dt} = rI$$

$$\frac{dR}{dt} = rI$$

# Mathematically: Linearization

- Linearization is ‘valid’ as long as  $R(t)$  is small
  - Thus, epidemic types often look “the same” for relatively long periods of time
  - Long term: Linearization (Mosquito) eventually infects everyone in the population (but may take “arbitrarily long” to do so )
- Deep Mathematical Question: On what interval is a linearization a ‘valid’ approximation?



Goal: Contexts that become “familiar” and can be used in many situations

- Freshman, General Majors: Simulation generates data for statistical testing
- Sophomore, Math Majors: Linear difference equation as both (a) programming assignment and (b) excuse to study sequences and series
- Junior/Senior: Systems of DE's, Linearization of DE's, statistical modeling, time series analysis

# Thank you!

Website: <http://math.etsu.edu/Symbiosis/epidemics>

- Data available from command line in Netlogo
- Netlogo interface with R can be used with Desktop Version of Netlogo (not enabled with applets)