



# Week 3: Stochasticity

Dr. Rachel Sippy  
University of Cambridge

# Week 3 Overview

- Monday, August 9:
  - Guest lecture & R session by Megan O'Driscoll
  - Stochastic models
  - Guided practice in R
- Tuesday, August 10:
  - Step-by-step model building
  - Building a COVID-19 model
  - Guided practice in R
- Wednesday, August 11:
  - Comparing models to data & evaluating models
  - Guided practice in R

# Post Questions in the Chat!

(or ask over microphone)

# Workshop Schedule

Time	Topics
2:00–2:10 pm	Greetings
2:10–3:00 pm	Guest lecture: COVID-19 with Megan O'Driscoll
3:00–3:10 pm	Break
3:10–4:00 pm	Stochasticity
4:00–4:10 pm	Break
4:00–5:00 pm	R Session

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Stochasticity

# Objectives

- Understand the difference between deterministic and stochastic models
- Understand options for adding stochasticity to a model

# Modeling Methods

Deterministic

Stochastic



# Modeling Methods

## Deterministic

- Describe what happens “on average” in a population
- Use average (single) value/transition rates in models

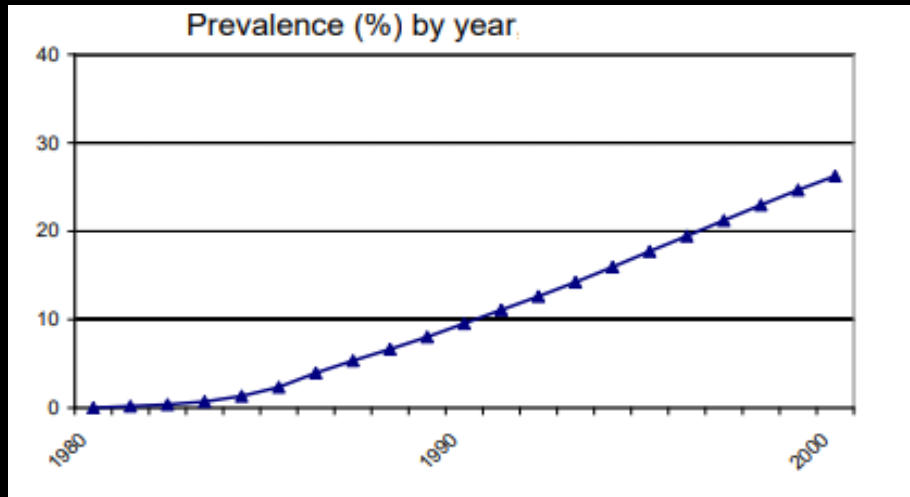
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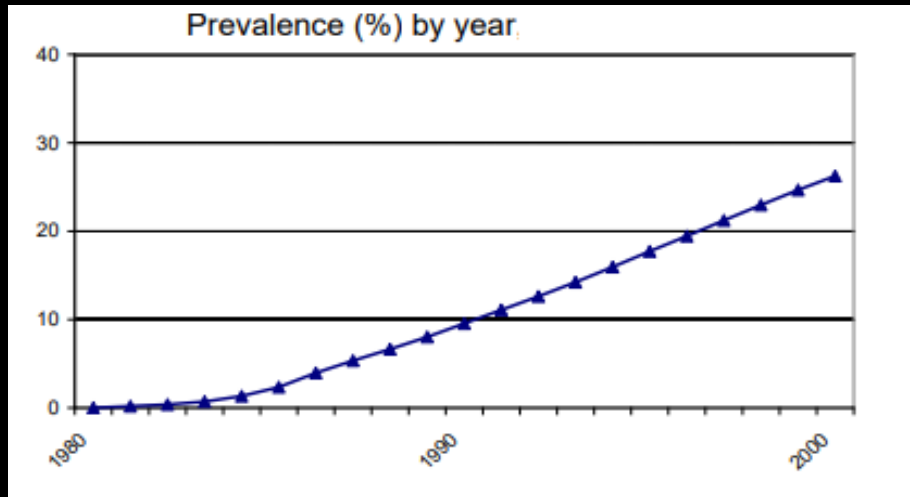
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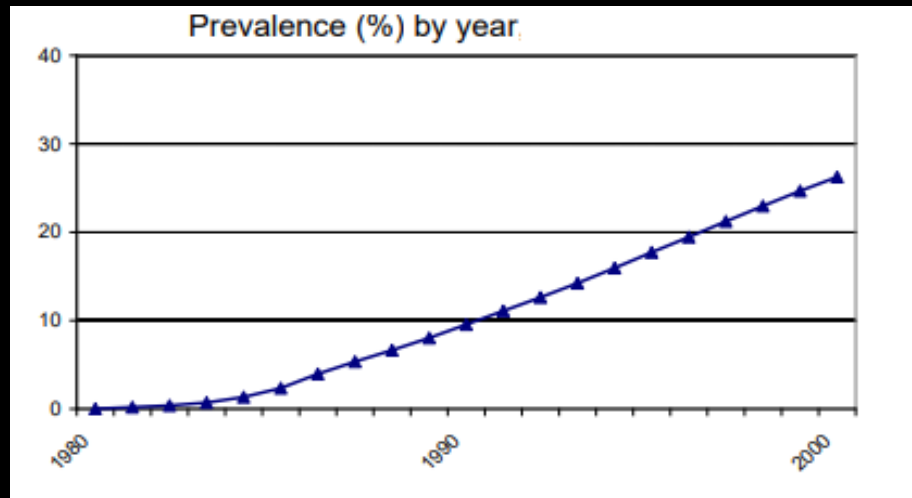
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- Use many (distribution) values/transition rates in models

# Modeling Methods

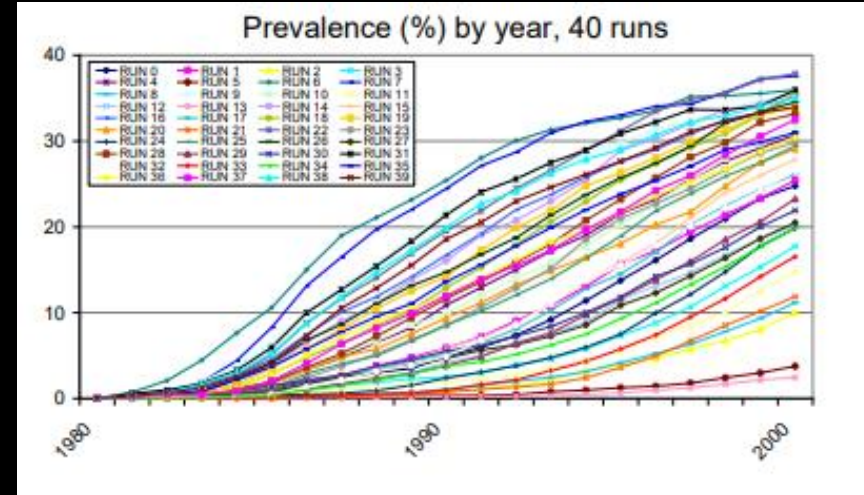
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## Stochastic

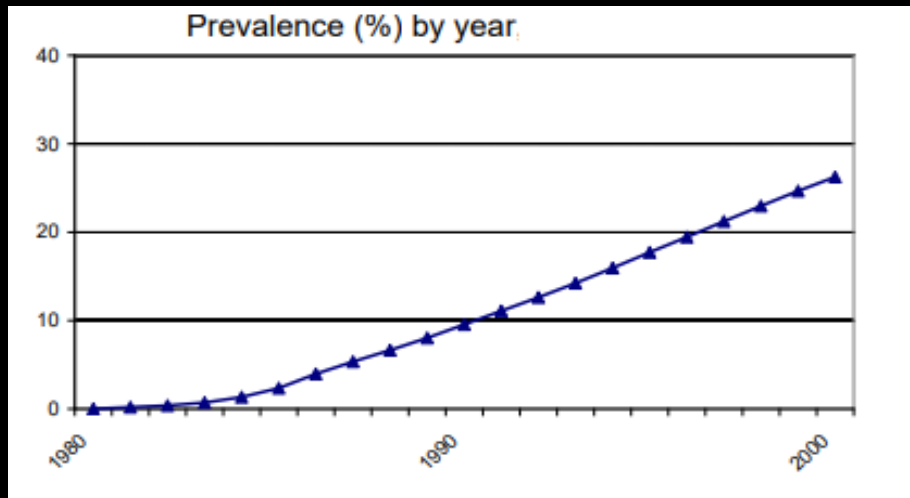
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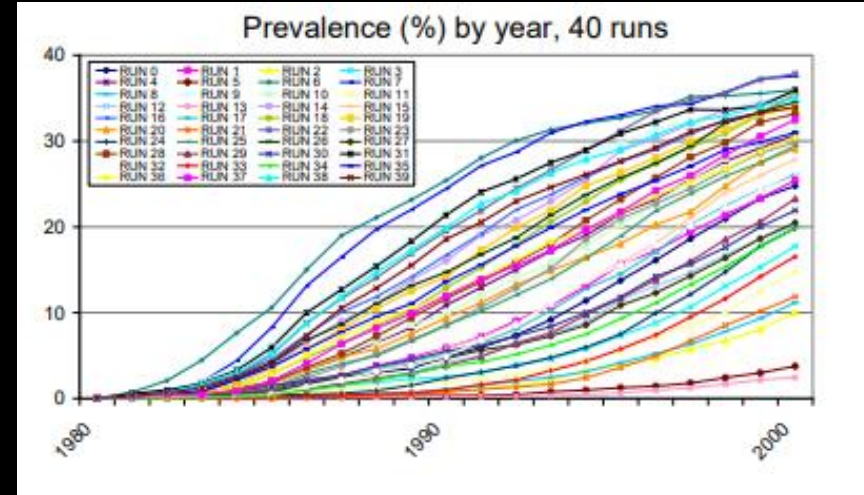
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## Stochastic - Random!

- Describes range of possible outcomes by incorporating chance
- Use many (distribution) values/transition rates in models



# Stochastic Modeling

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    - random error

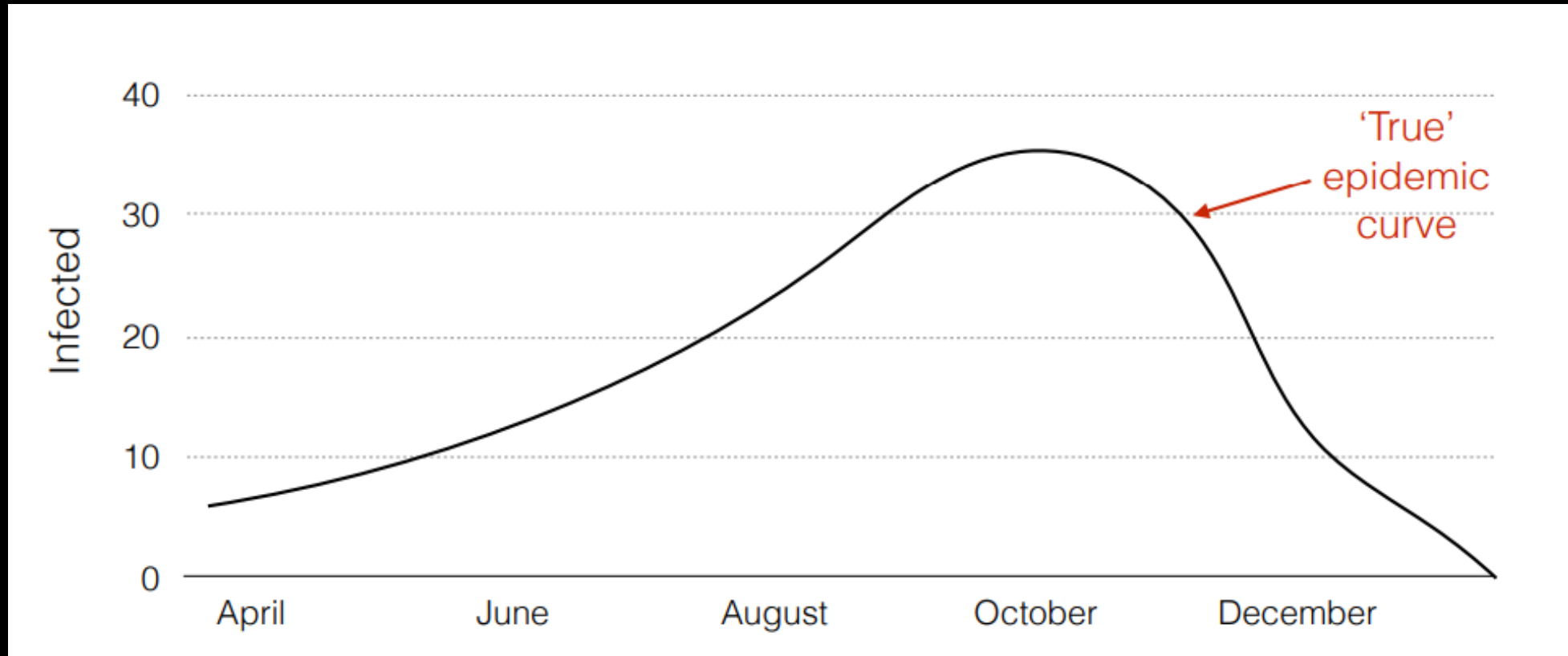
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    - infection event is a random event
    - will a susceptible contact an infected? will transmission occur?

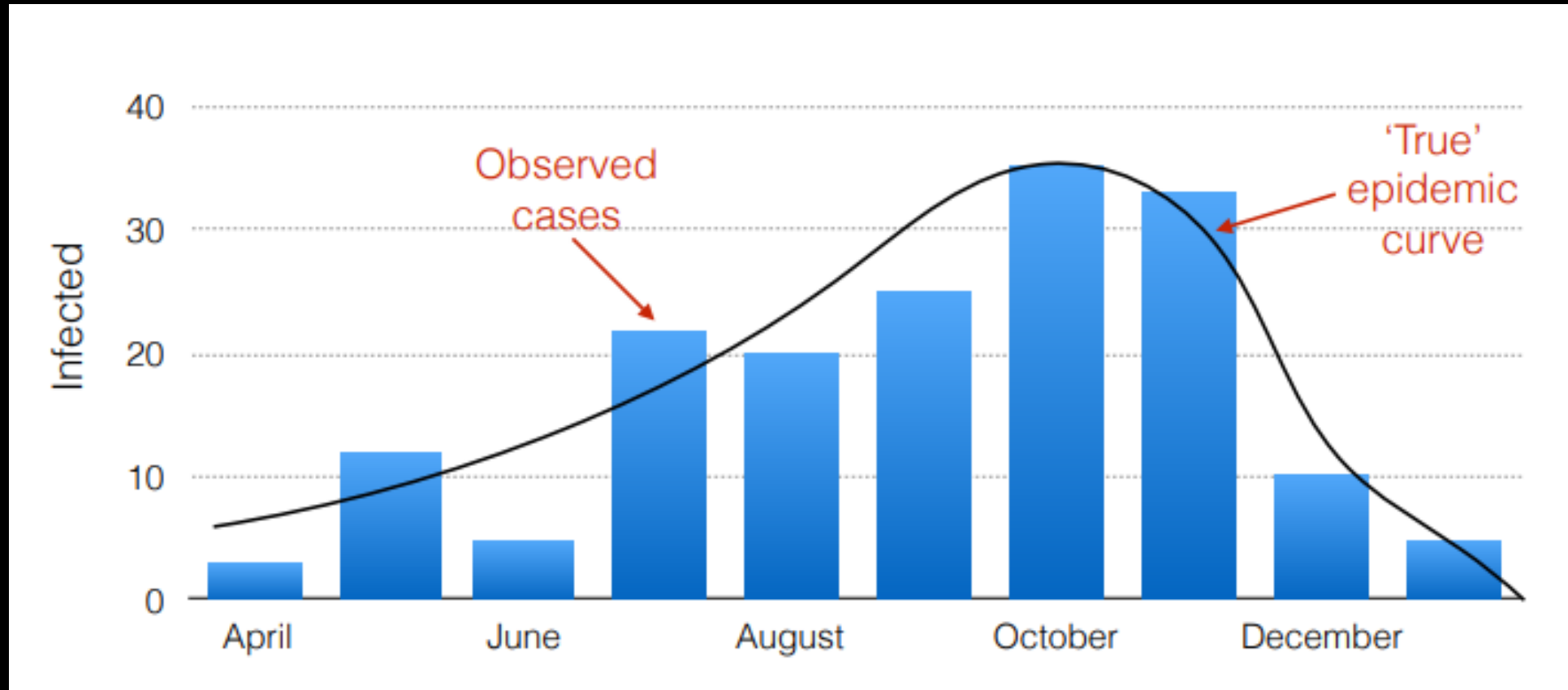
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    - will a susceptible contact an infected? will transmission occur?
    - can have profound impact on epidemic dynamics

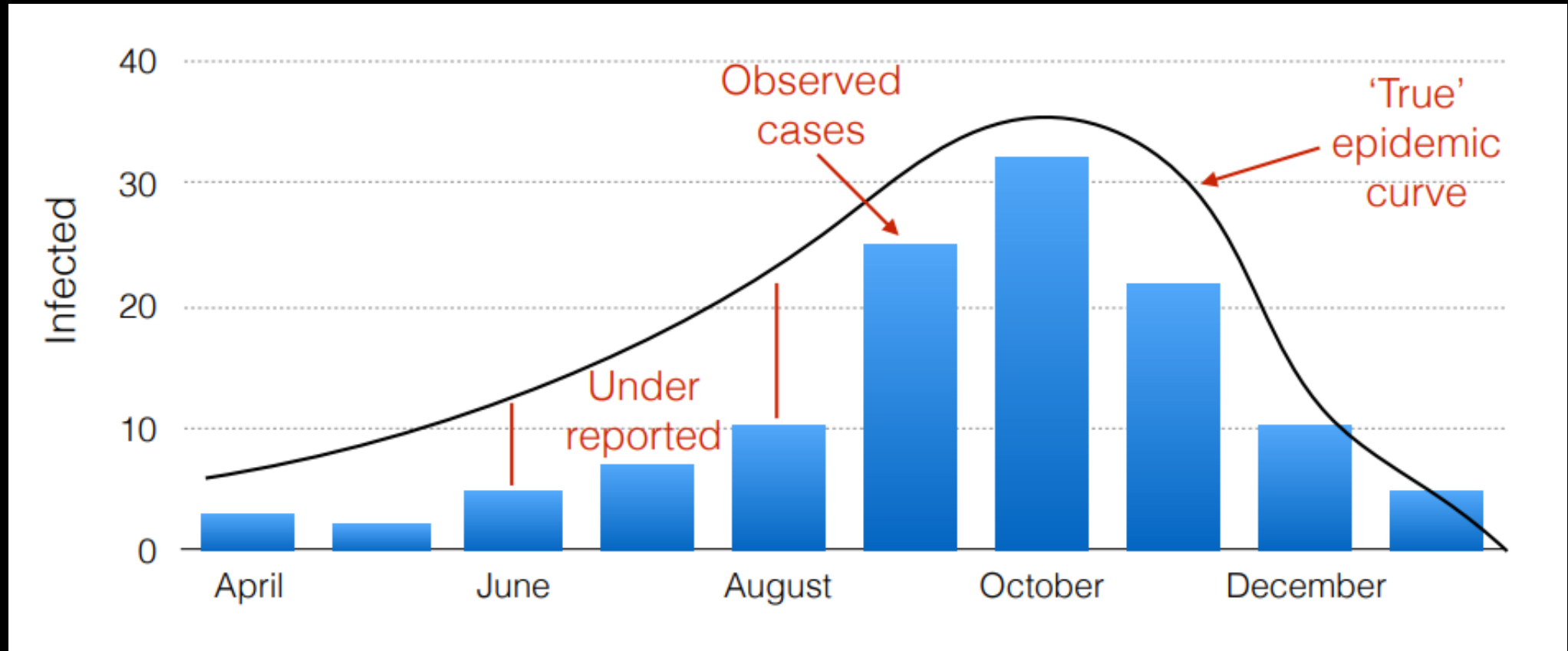
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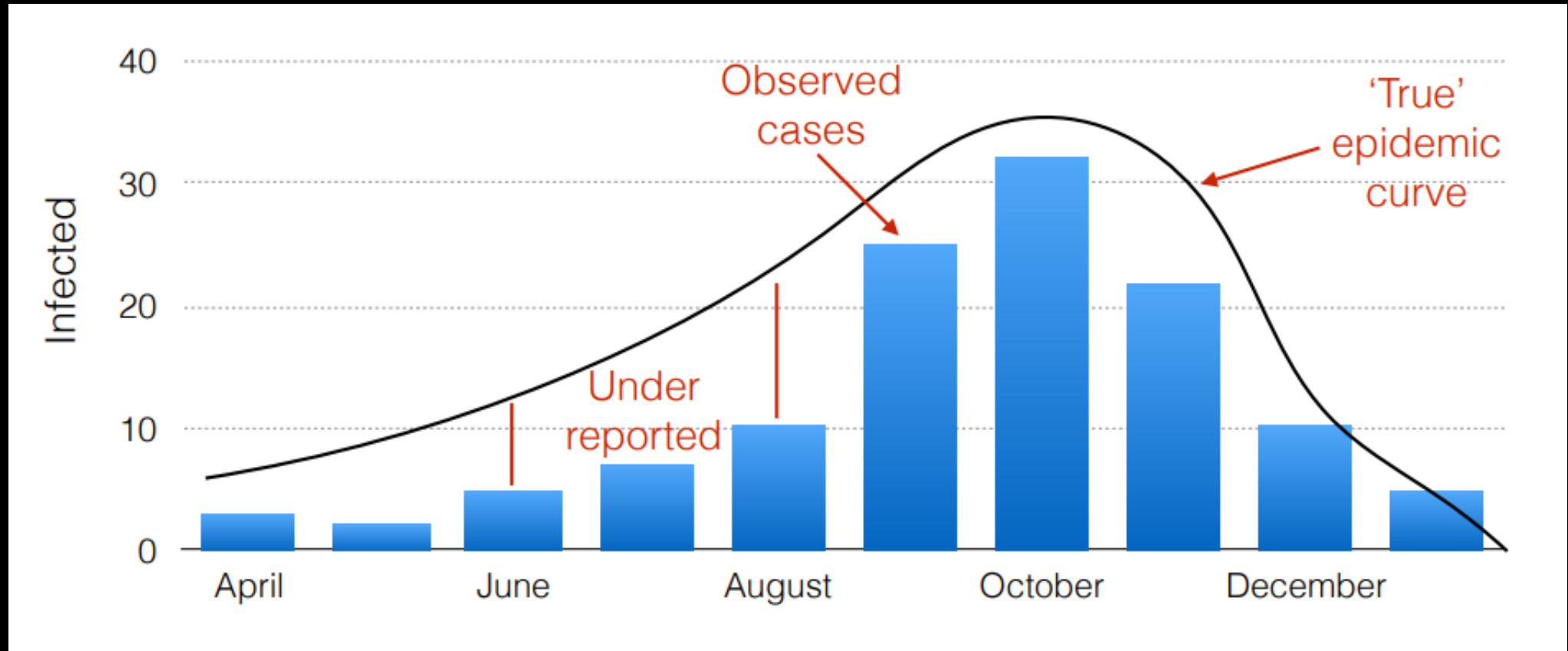
# Observational Stochasticity

$$I_t = \rho C_t$$

$I_t$ : number infected

$\rho$ : reporting rate

$C_t$ : case count

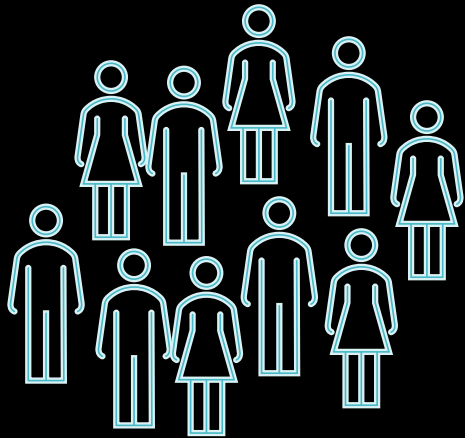


# Process Stochasticity

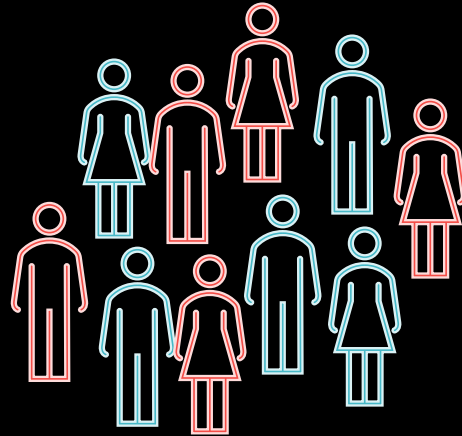
- Infection events are random events!
  - number of individuals you are in contact with
  - whether disease is transmitted during a contact
- These events can have a major impact on an epidemic
  - we want to build this uncertainty into our model

# Modeling Results

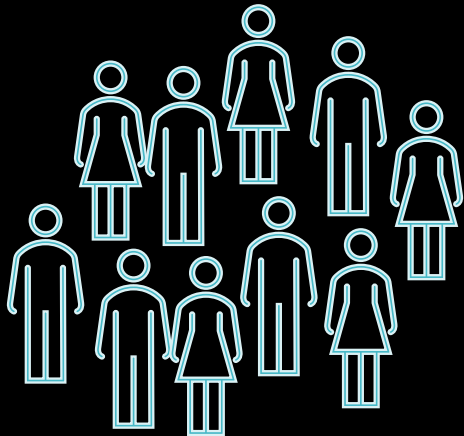
Deterministic



Probability of  
infection=50%



Stochastic

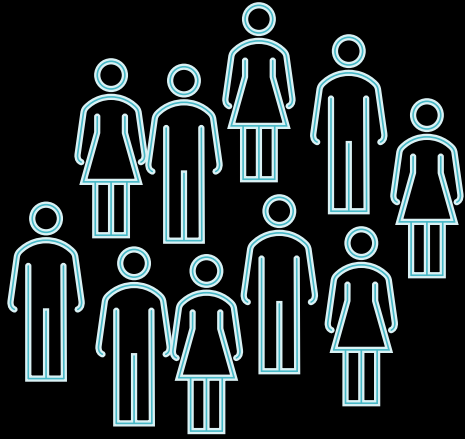


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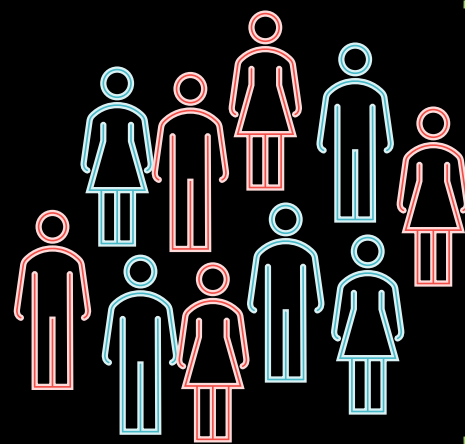
Starting population

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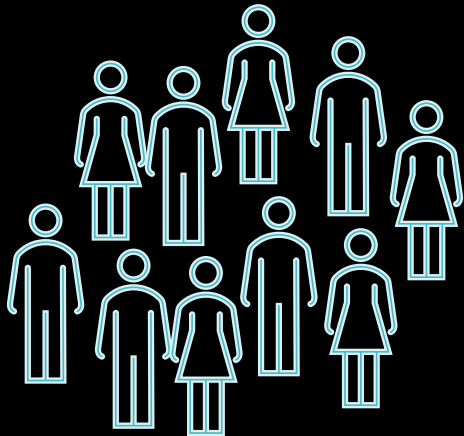


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This is what the  
outcome will be  
no matter how  
many times we  
run the model!

Stochastic

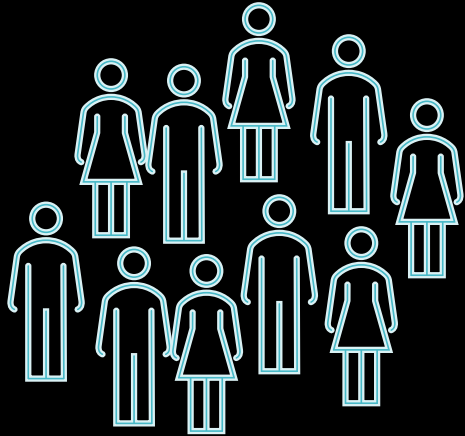


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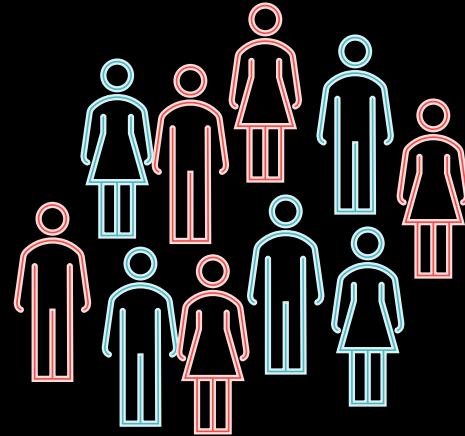
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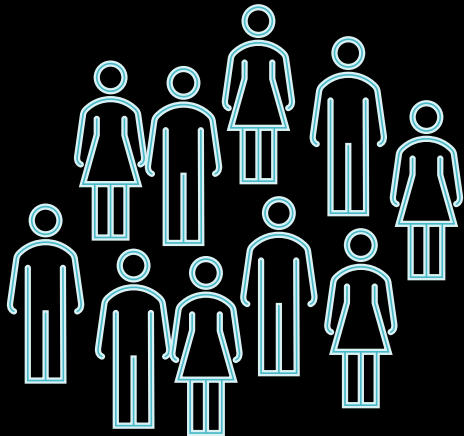
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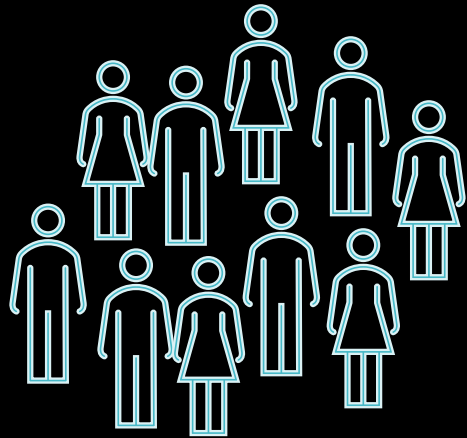
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Flip a coin for  
each person  
Probability of  
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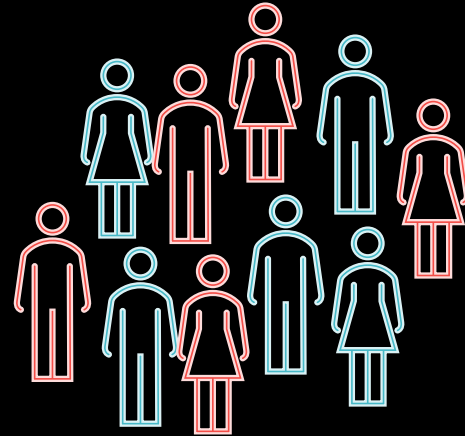


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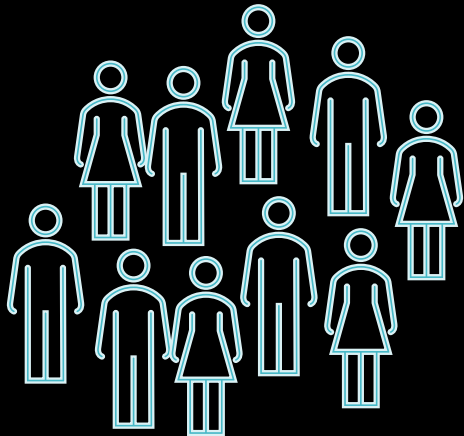
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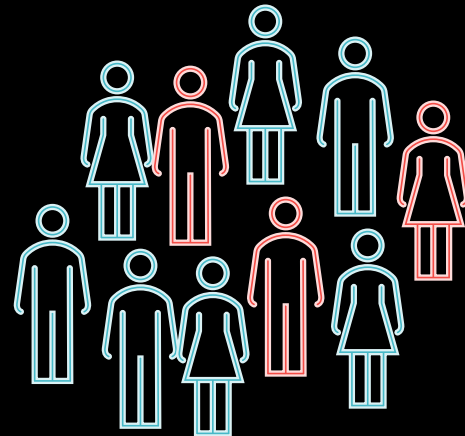


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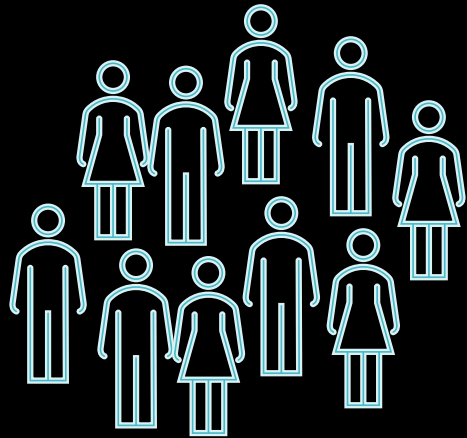
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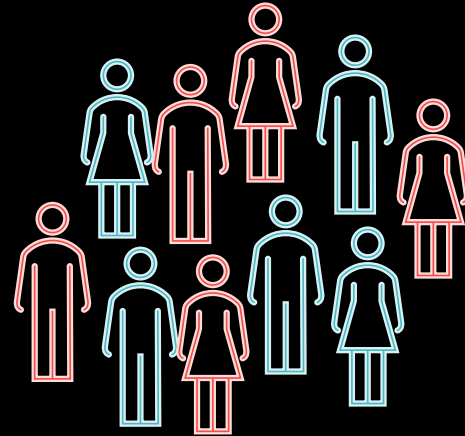


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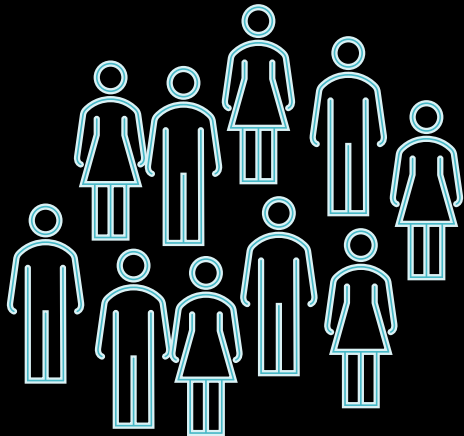
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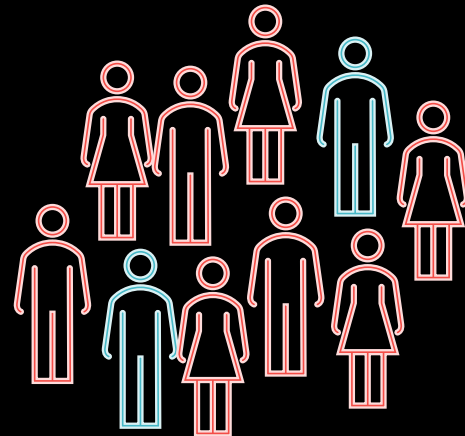


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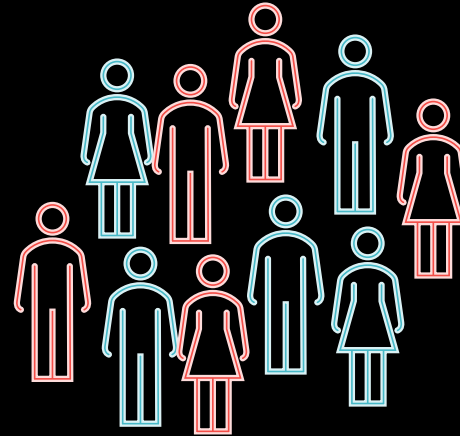


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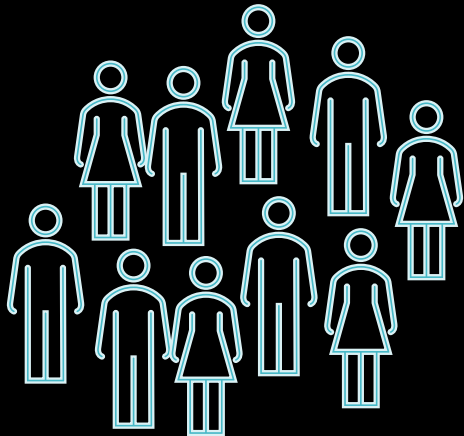
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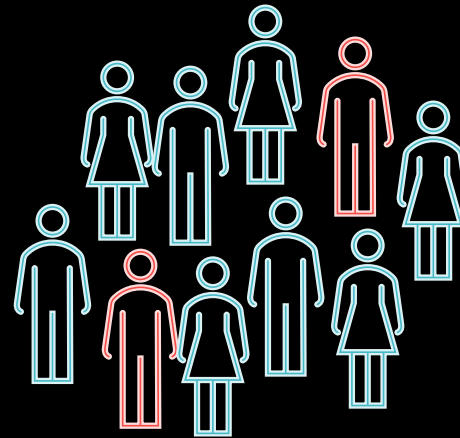
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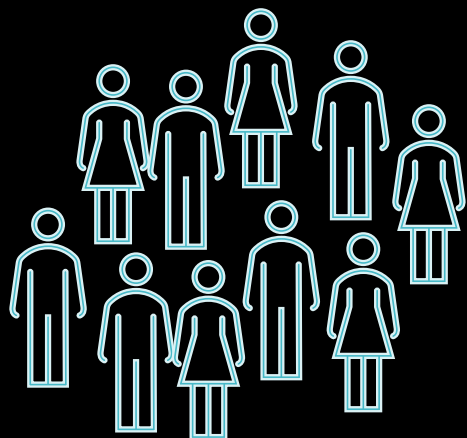


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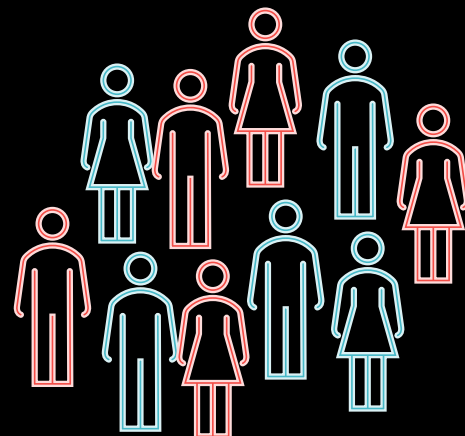


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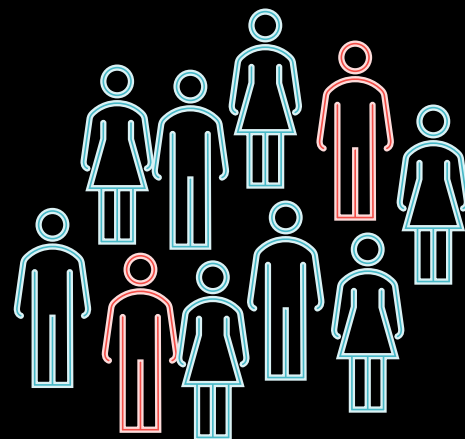
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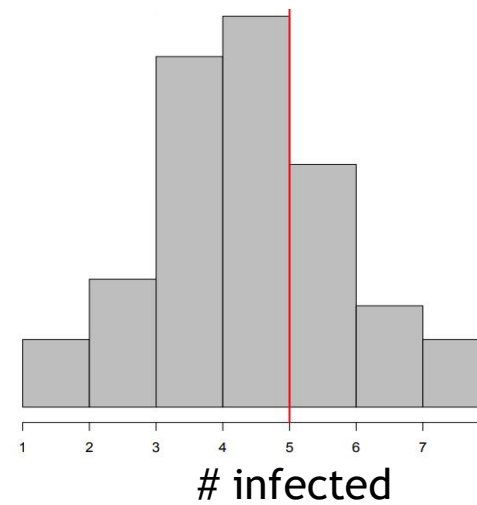


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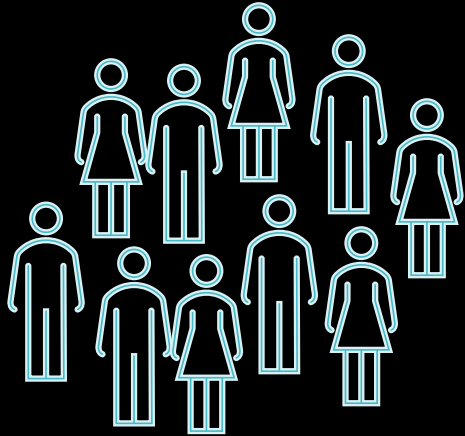
Starting population

`rbinom(200,10,0.5)`

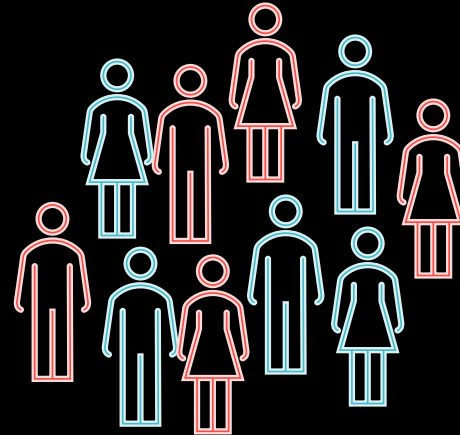


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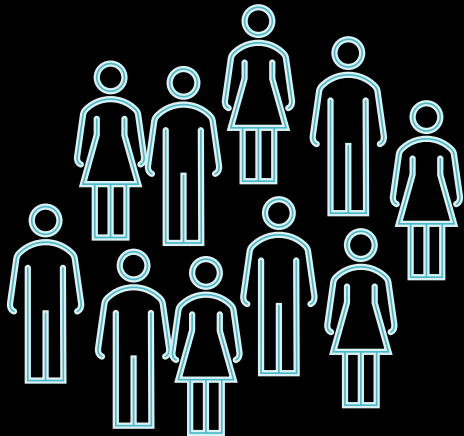


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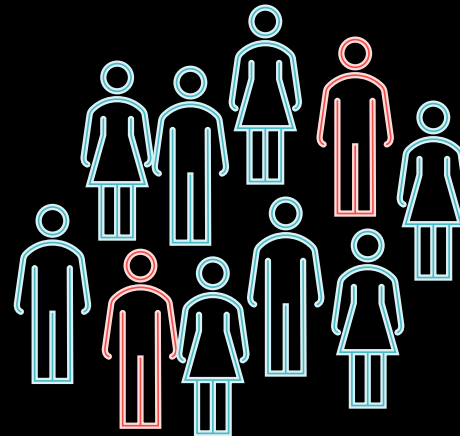
Average # infected will be 5

Stochastic

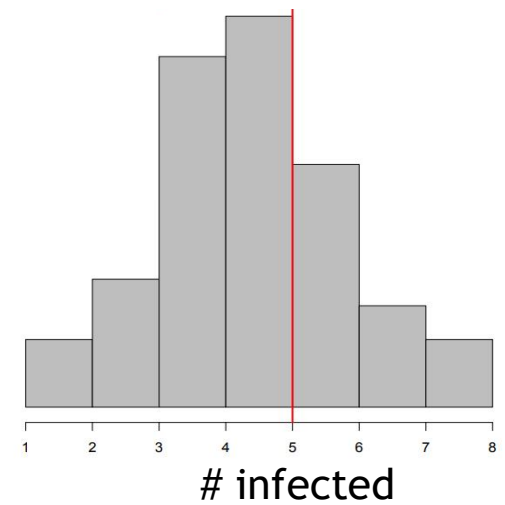


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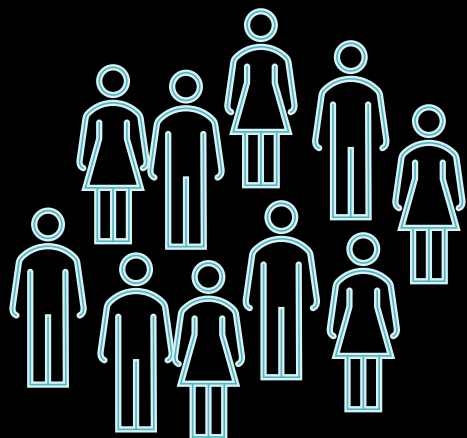


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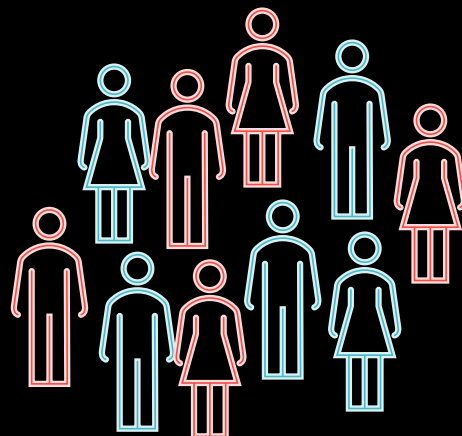


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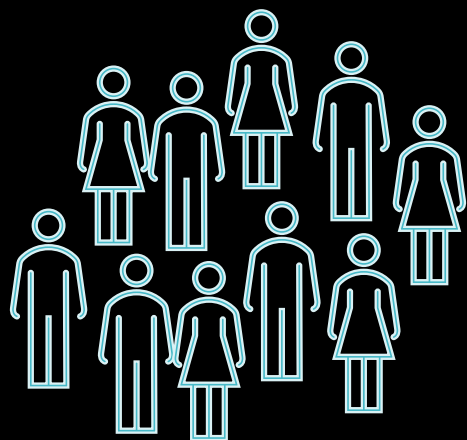
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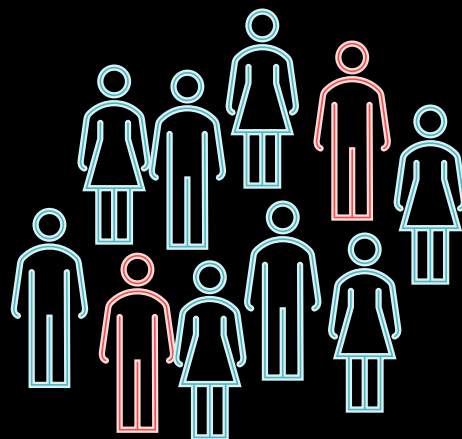
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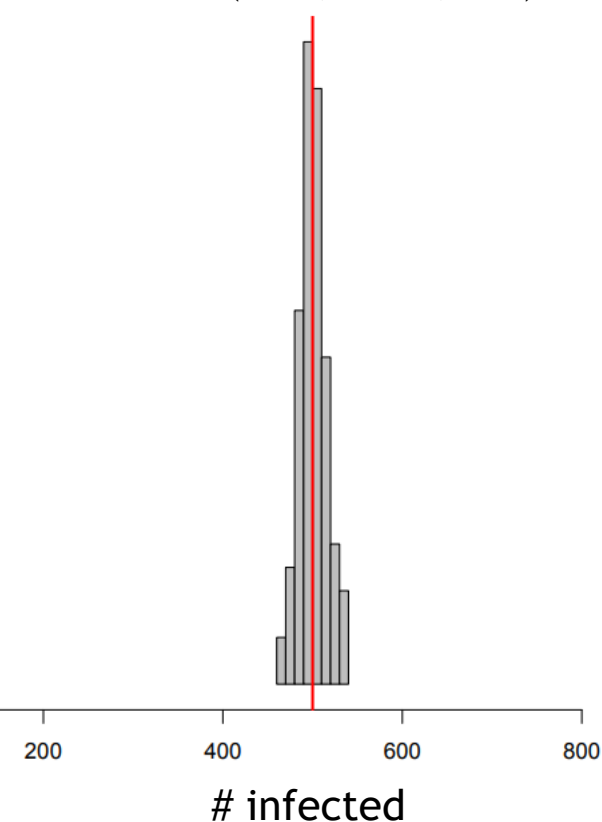


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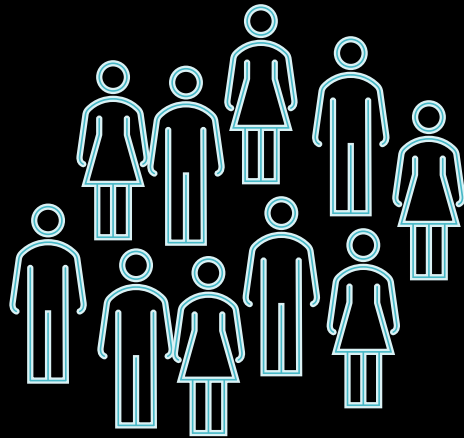
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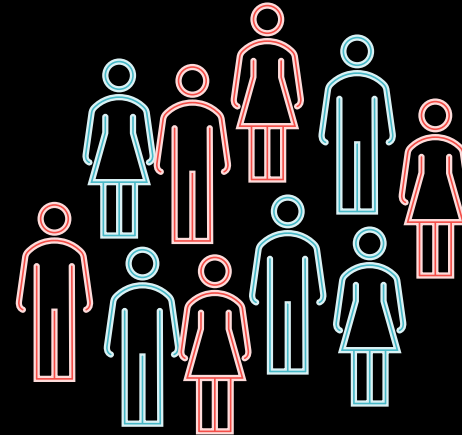


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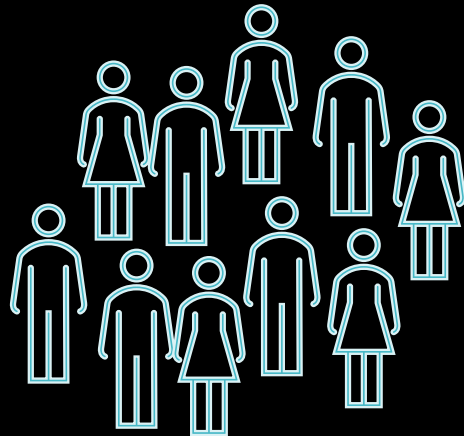
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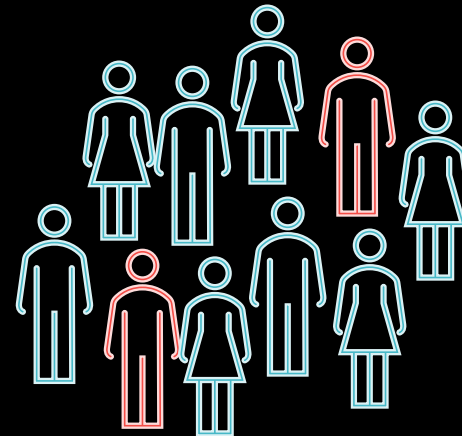
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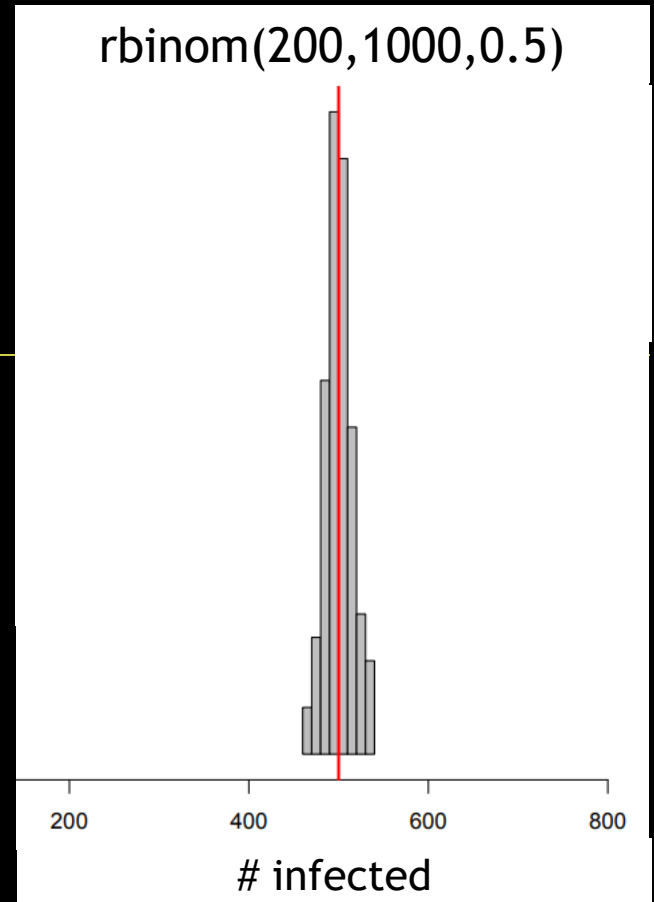


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Starting population

If we have many people, variance will be small - in this case deterministic predictions approximate reality



# Process Stochasticity in Models

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- Depends on modeling methods being used, multiple ways to include in models
- Model able to produce different results - even without changing our input values
  - adding stochasticity causes this
- Why do we want to add stochasticity?
  - adds variability to each simulation/modeling run
  - demonstrate extremes of disease dynamics
  - more realistic, more robust

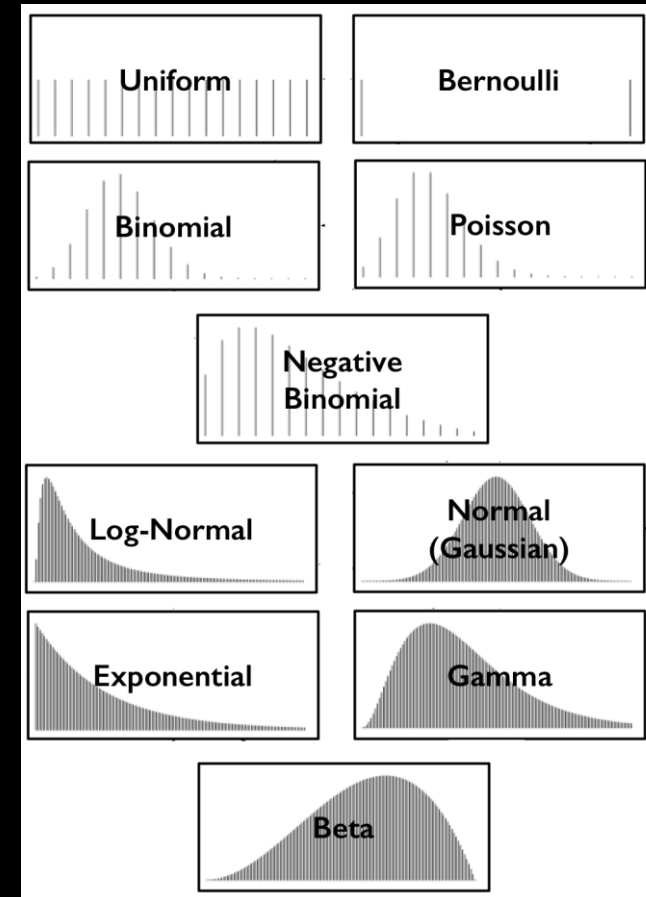
# Stochasticity Through Distributions

- Instead of using a single value for our model parameters, we can use a distribution of values



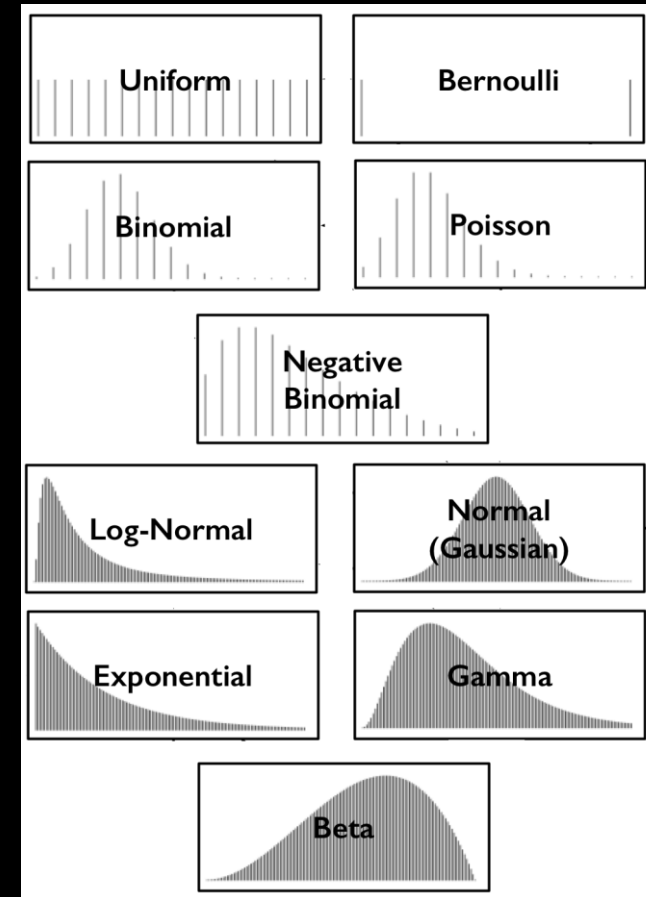
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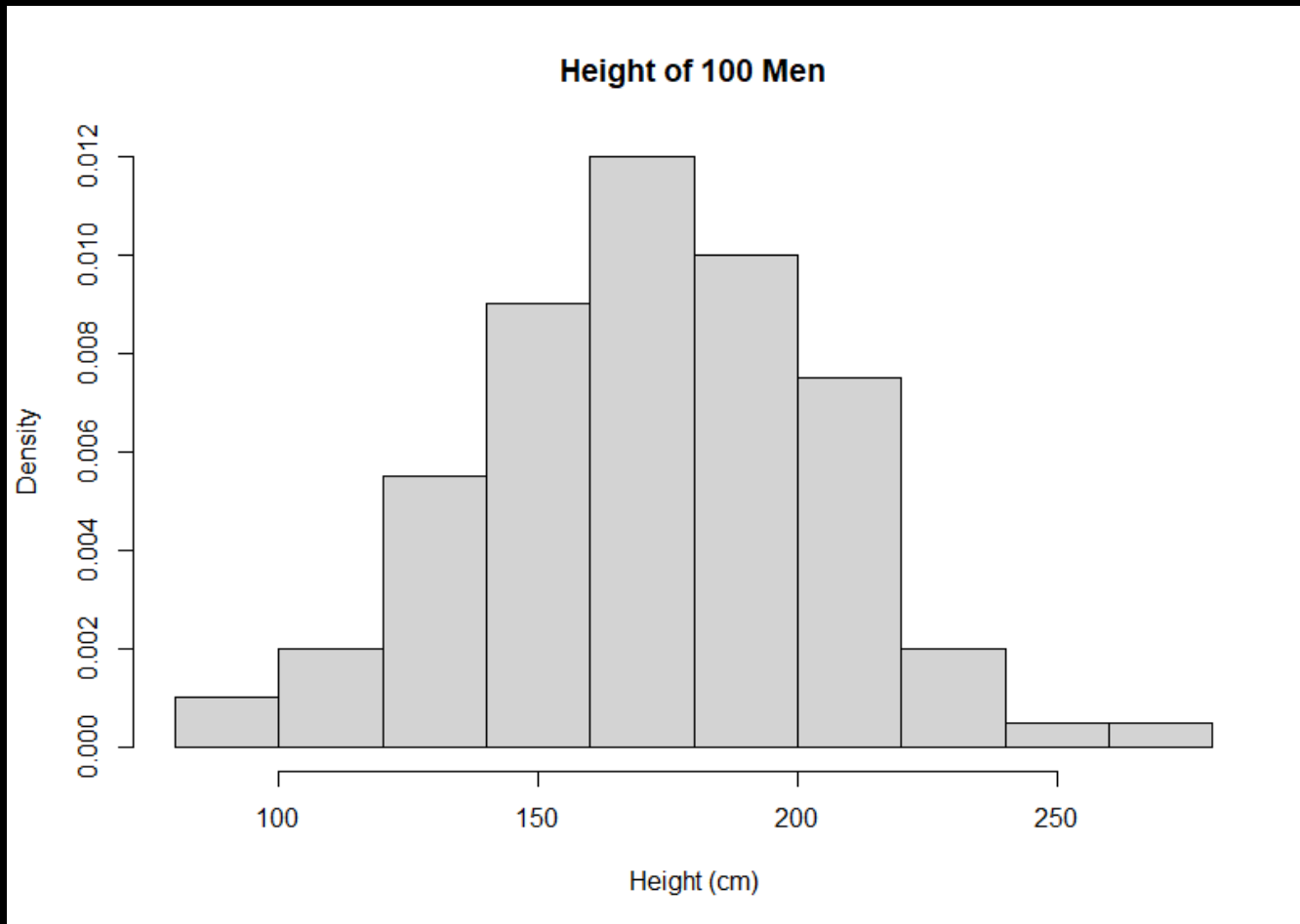
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# Stochasticity Through Distributions

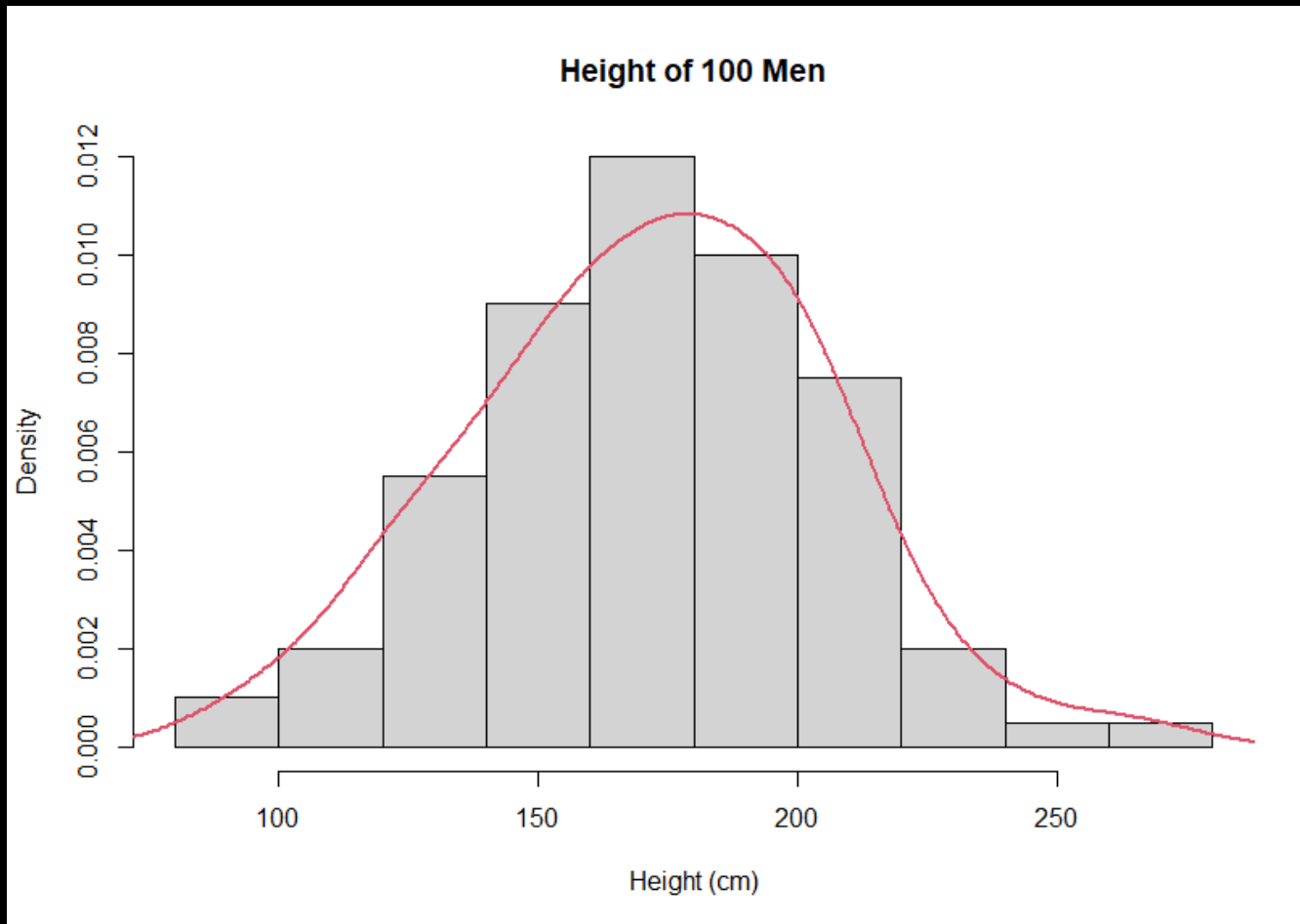
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  - represents probability that different outcomes will occur





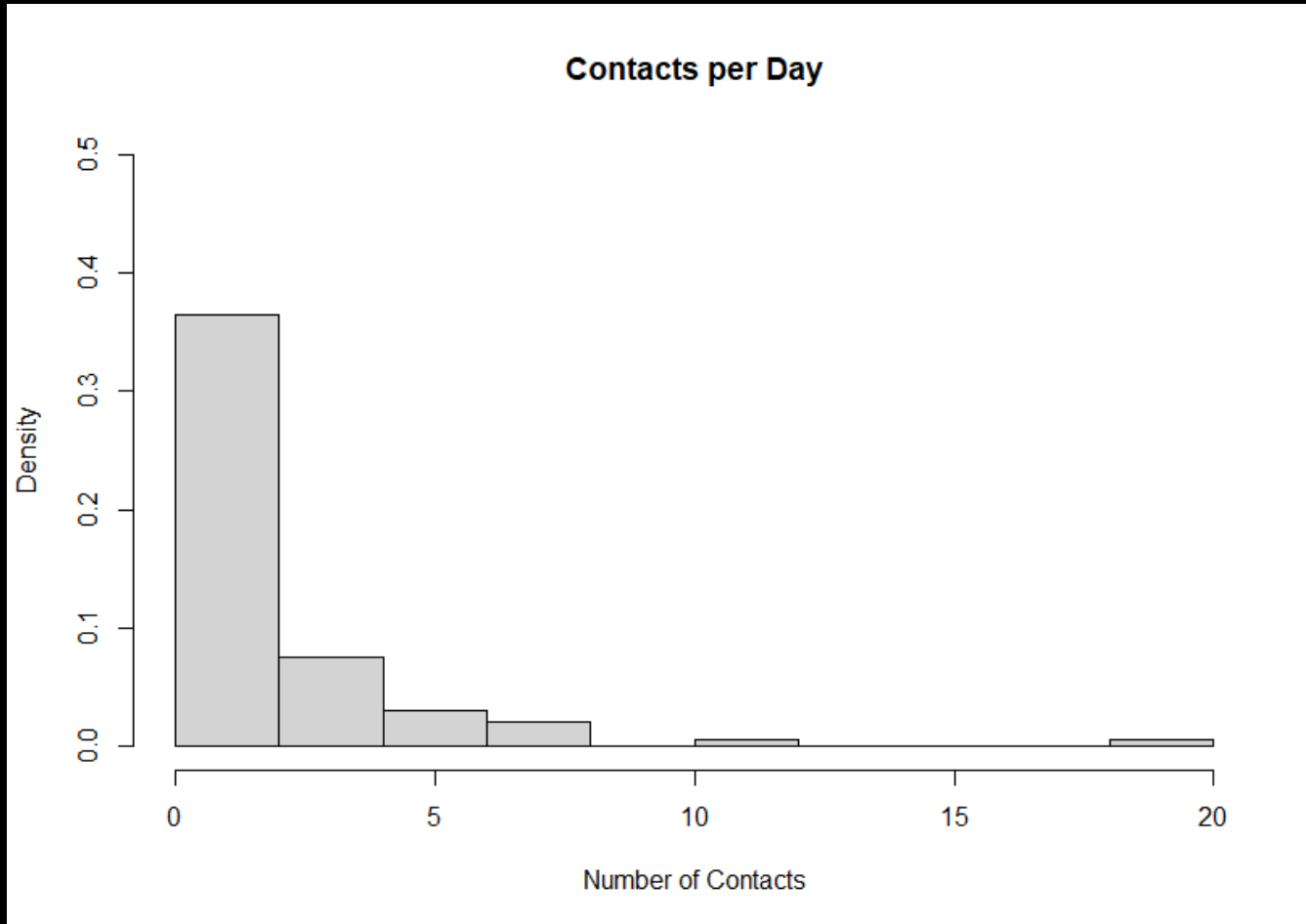
## Distributions Can Represent Real-life Phenomena

- Histogram of heights of 100 men



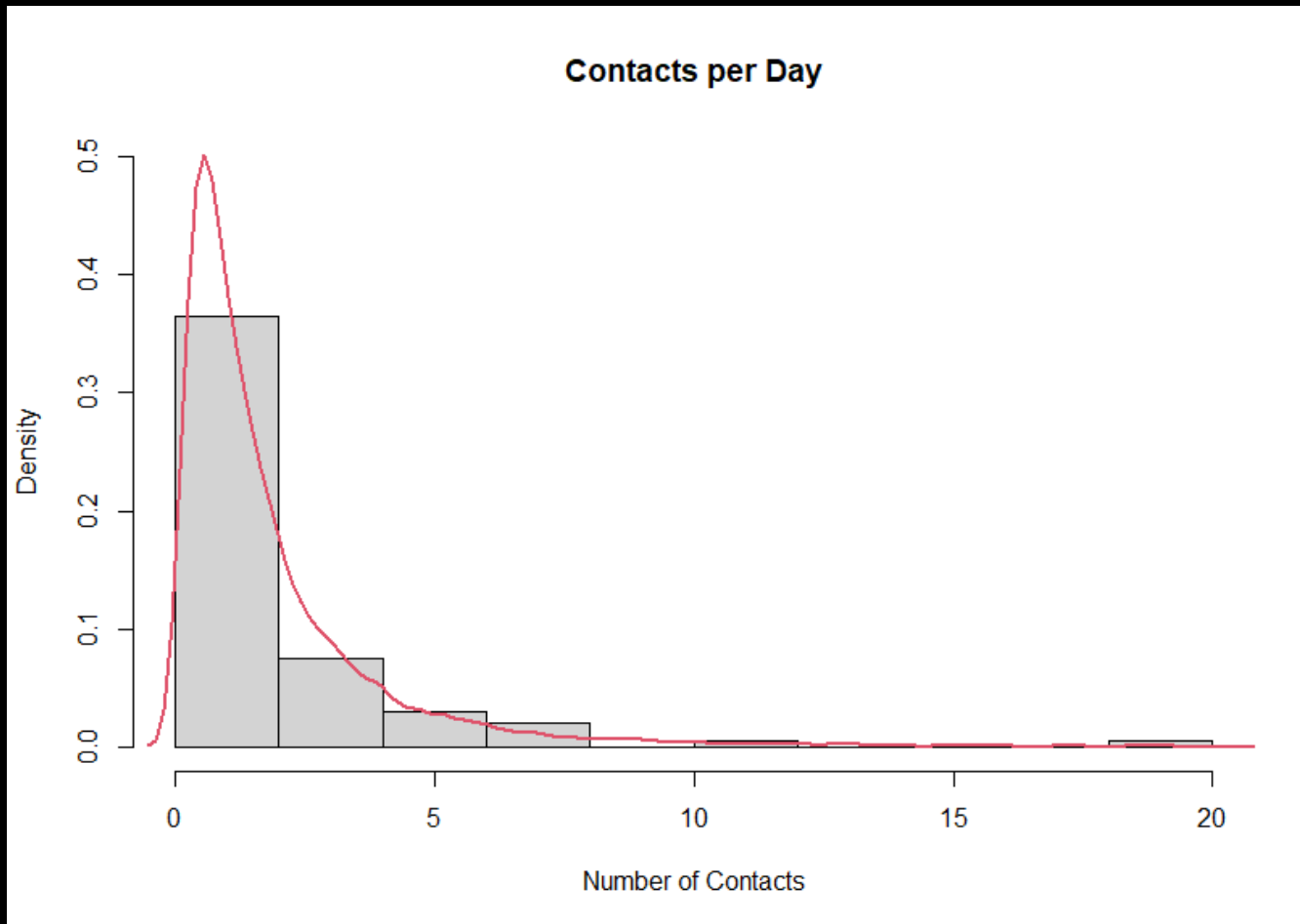
## Distributions Can Represent Real-life Phenomena

- Histogram of heights of 100 men
- Approximately normal!
- $Height \sim N(170, 30)$



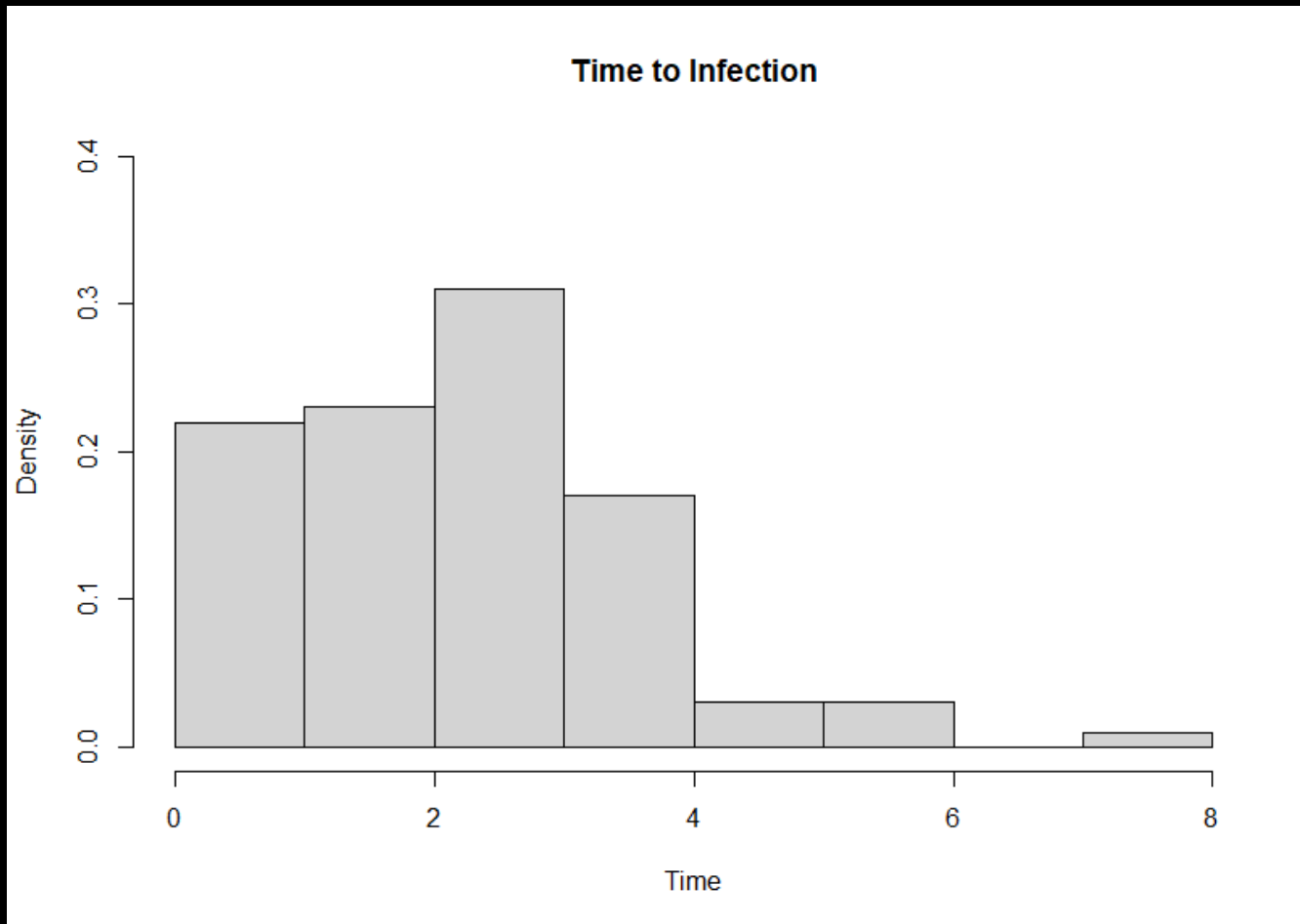
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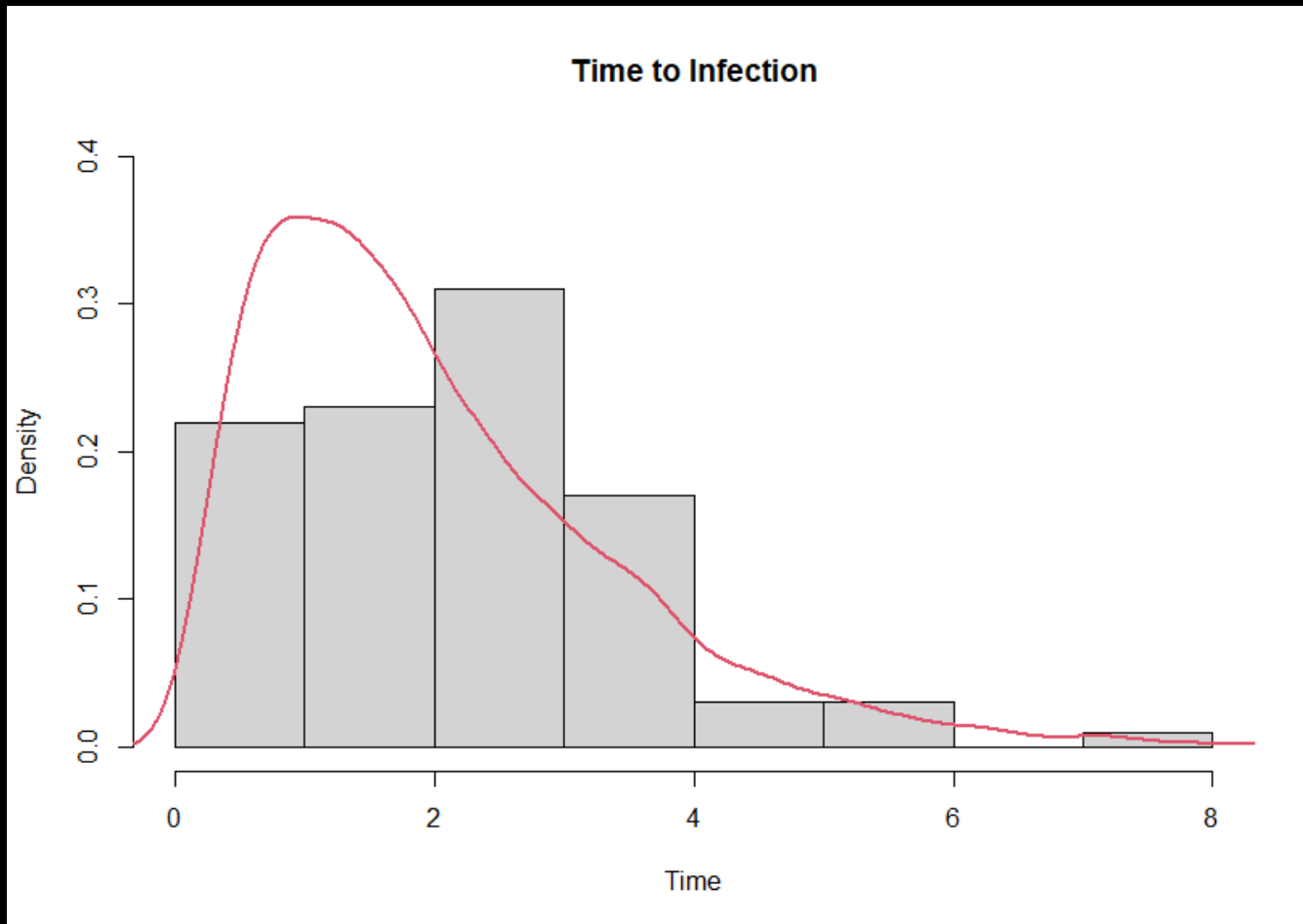
## Distributions Can Represent Real-life Phenomena

- Histogram of daily number of contacts
- Lognormal distribution!
- $C \sim \text{LogNormal}(0.2)$



## Distributions Can Represent Real-life Phenomena

- Histogram of time to infection



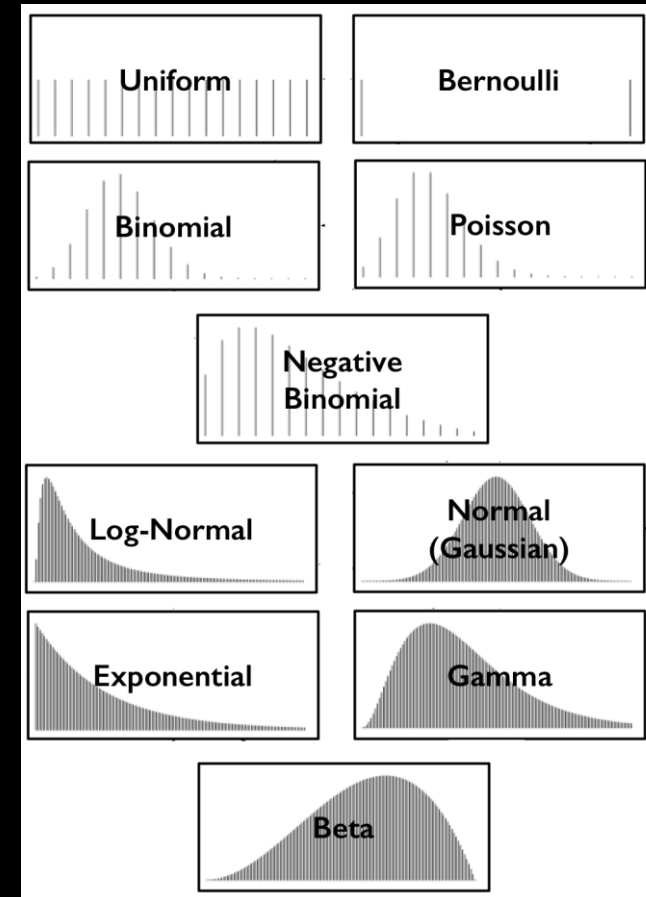
## Distributions Can Represent Real-life Phenomena

- Histogram of time to infection
- Gamma distribution!
- $tInf \sim \text{Gamma}(2, \text{rate} = 1)$



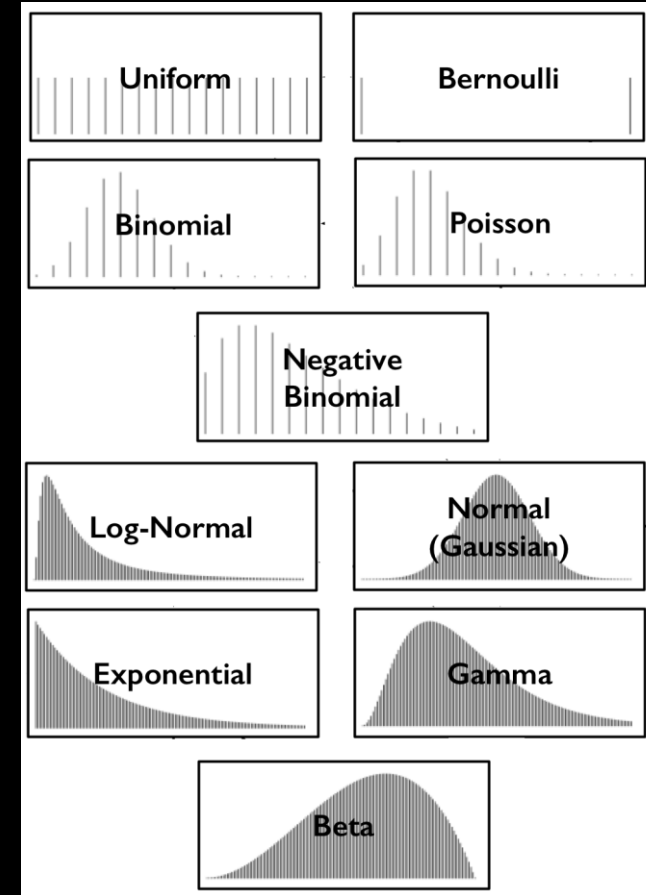
# Stochasticity Through Distributions

- Instead of using a single value for our model parameters, we can use a distribution of values
- What are distributions?
  - represents probability that different outcomes will occur
- We can tell the model to select one of the values from the distribution for each run/simulation
- We can include a distribution when making calculations/estimates



# Stochasticity Through Distributions

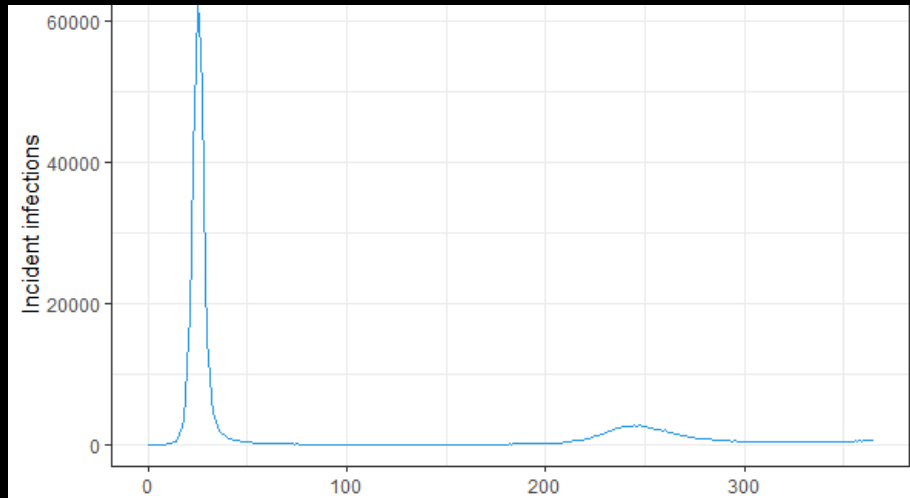
- Example: modeling force of infection ( $\lambda$ )
  - $\lambda(t) = \beta S(t)I(t)$
  - $I(t + 1) =$   
 $\sim \text{NegBin}(\lambda(t), S(t))$



# Comparison

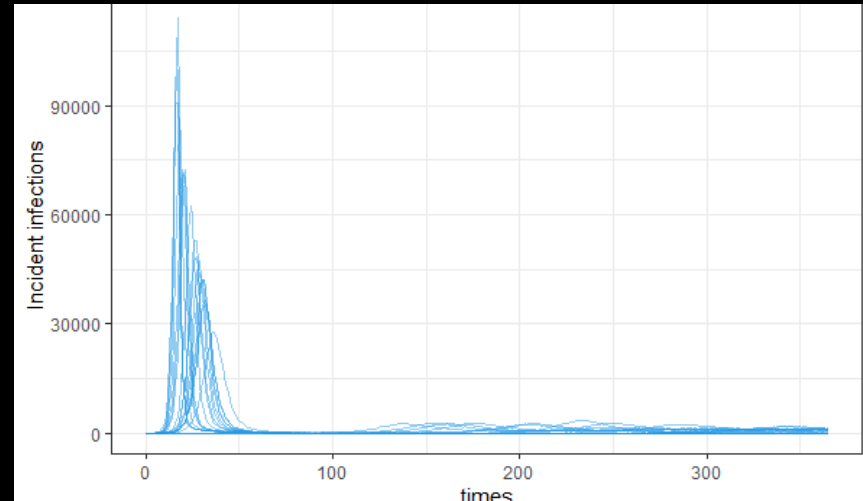
## Deterministic

- Use average (single) transition rates in models
  - $\beta = 1$



## Stochastic - Random!

- Use many (distribution) transition rates in models
  - $\beta \sim N(1, 0.25)$



# Modeling with Stochasticity

- Important part of infectious disease dynamics
- Can impact trajectory of disease
- Many ways to integrate stochasticity
  - directly add chance to your equations

# Questions?

10 minute break

# Workshop Schedule

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