

Model Fitting: The Basic Concept

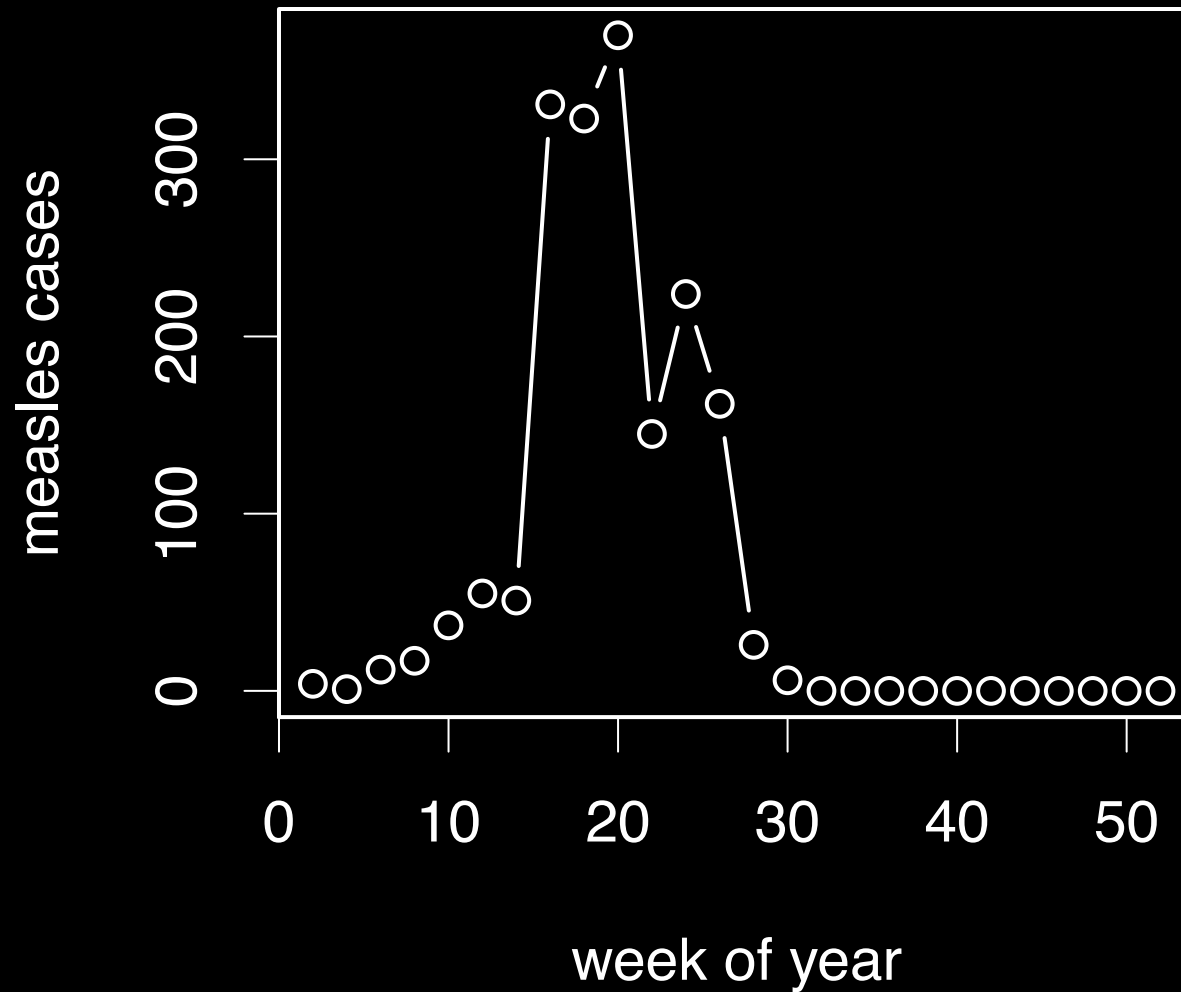


Cara Brook

E²M², Centre ValBio

Ranomafana National Park, Madagascar

What happened in Niamey?

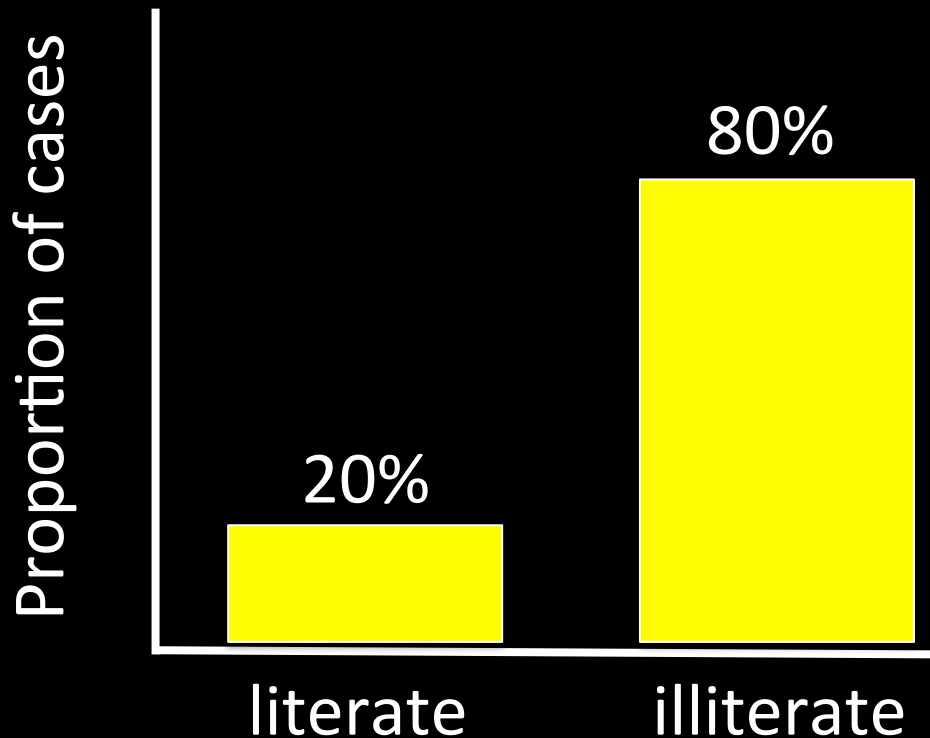


Traditional statistics is **data**-driven...

- We might ask questions about these data:
 - What proportion of cases occurred in males vs. females?
 - In winter vs. spring?
 - In Maradi vs. Dosso arrondissement?
- Goal: find **correlations** that **imply causality**

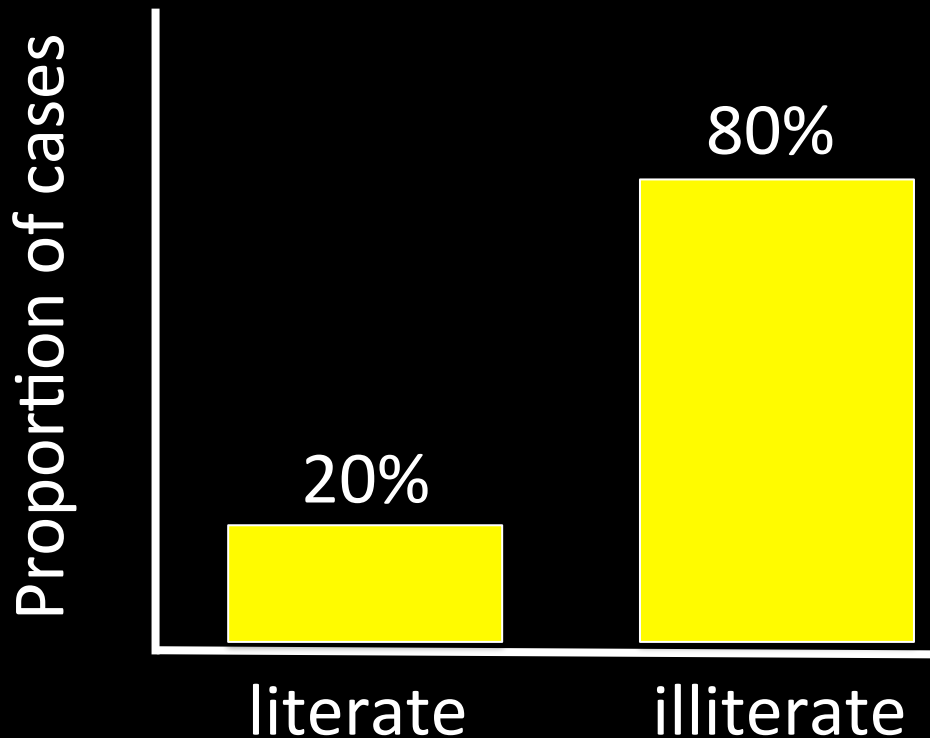
Traditional statistics is **data**-driven...

- Goal: find **correlations** that **imply causality**
- Imagine you discover:



Traditional statistics is **data**-driven...

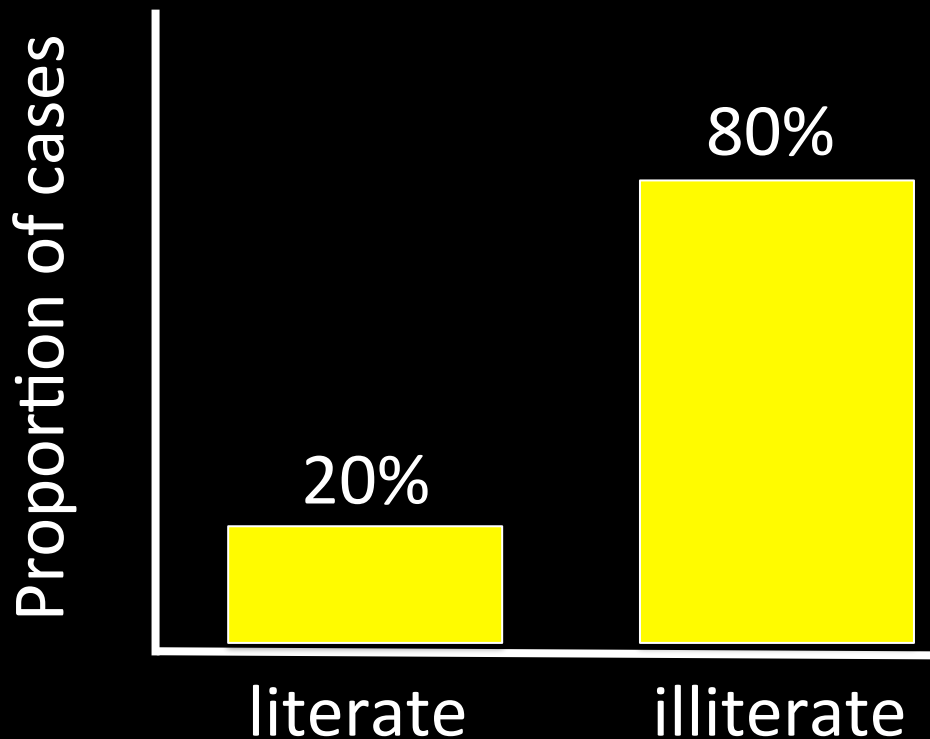
- Goal: find **correlations** that **imply causality**
- Imagine you discover:



*Does literacy
protect
against
measles?*

Traditional statistics is **data**-driven...

- Goal: find **correlations** that **imply causality**
- Imagine you discover:



**Must account for
confounding, bias,
and random error**

- *Literacy is
confounded with
age!*

Mechanistic modeling is **process**-driven...

- We want to understand **what** happened, **when** it happened, and **why** it happened

Mechanistic modeling is **process**-driven...

- We want to understand **what** happened, **when** it happened, and **why** it happened
- Allows us to scale up from **individual**-level processes to **population**-level patterns

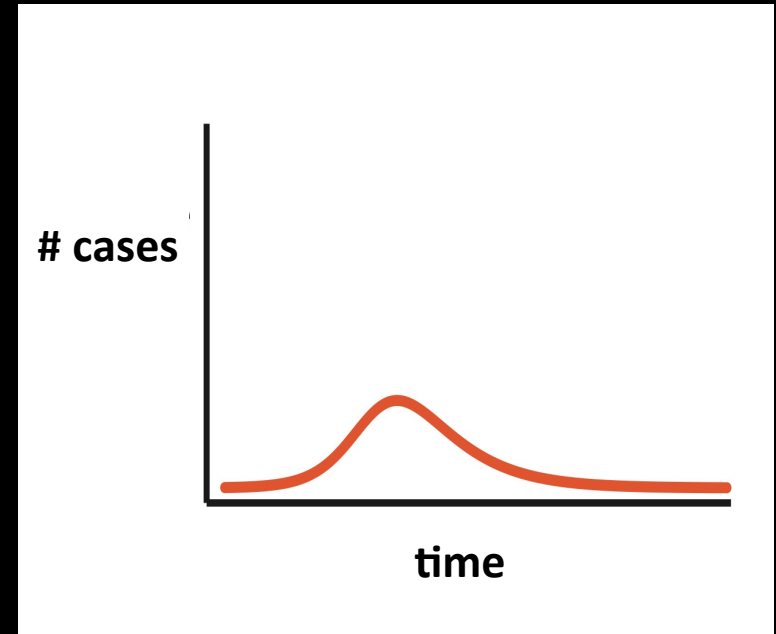
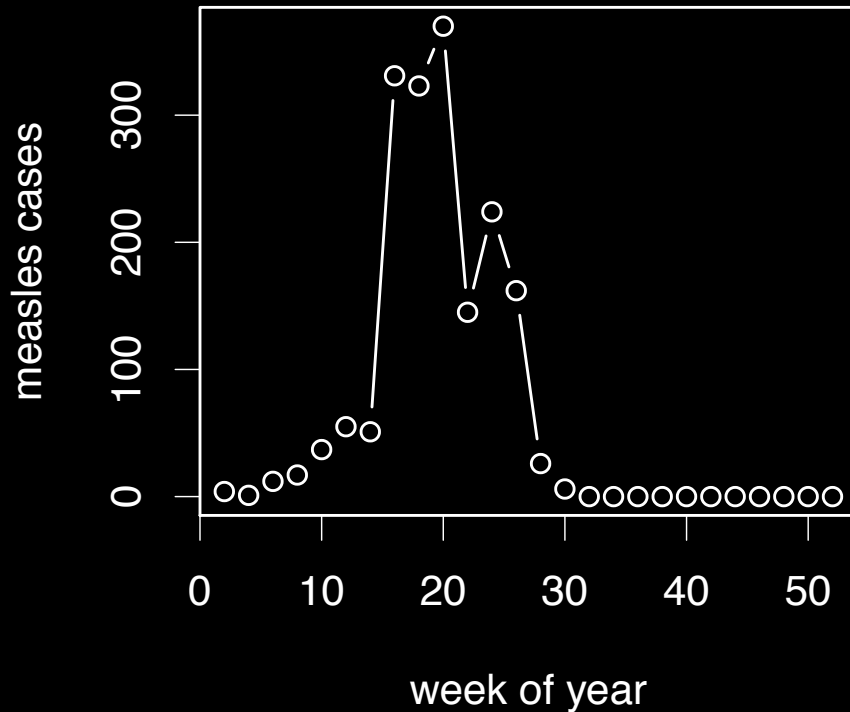
Mechanistic modeling is **process**-driven...

- We want to understand **what** happened, **when** it happened, and **why** it happened
- Allows us to scale up from **individual**-level processes to **population**-level patterns
- We start by building a **model** that uses explicit **processes** to recover the same outcomes (“**states**”) as our **data**

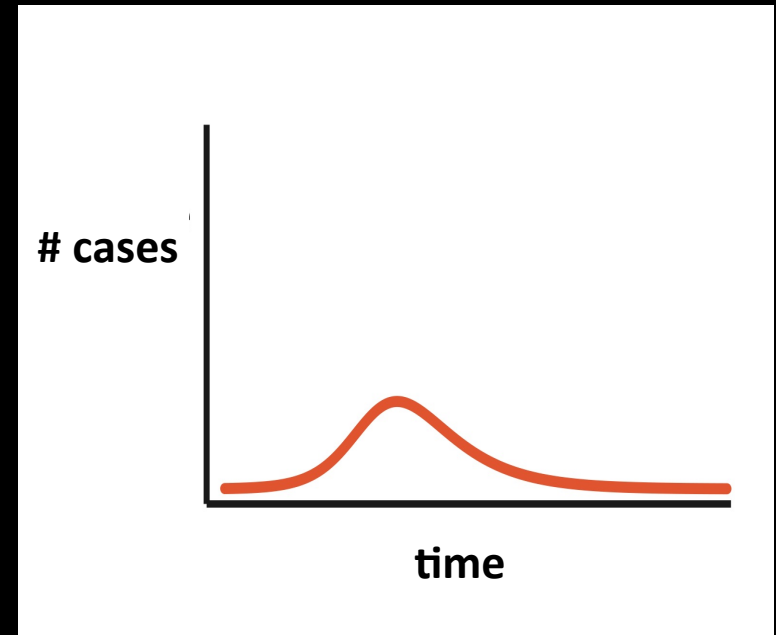
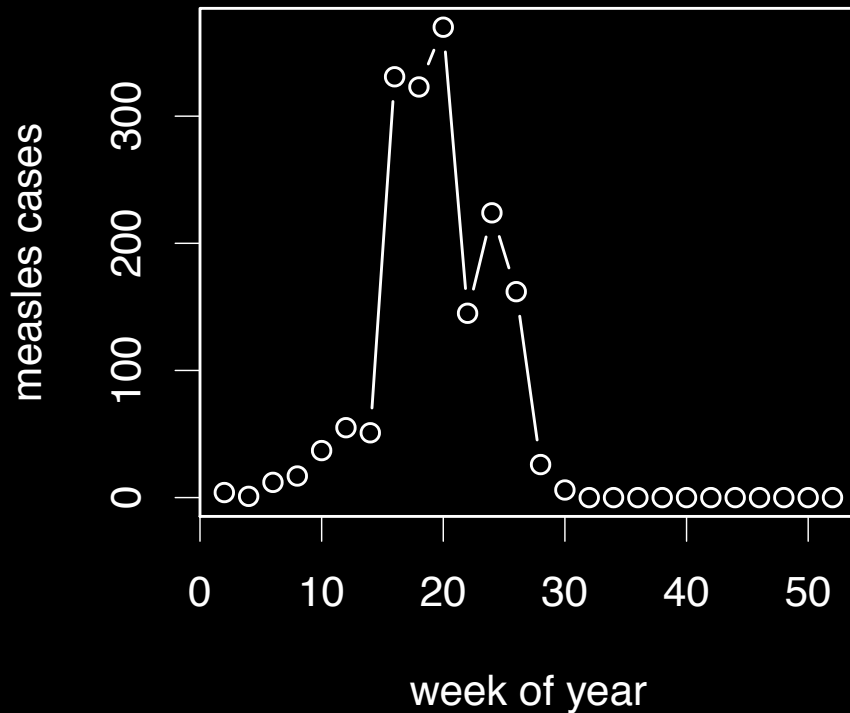
Mechanistic modeling is **process**-driven...

- We want to understand **what** happened, **when** it happened, and **why** it happened
- Allows us to scale up from **individual**-level processes to **population**-level patterns
- We start by building a **model** that uses explicit **processes** to recover the same outcomes (“**states**”) as our **data**
- *What state variables are captured in our data?*

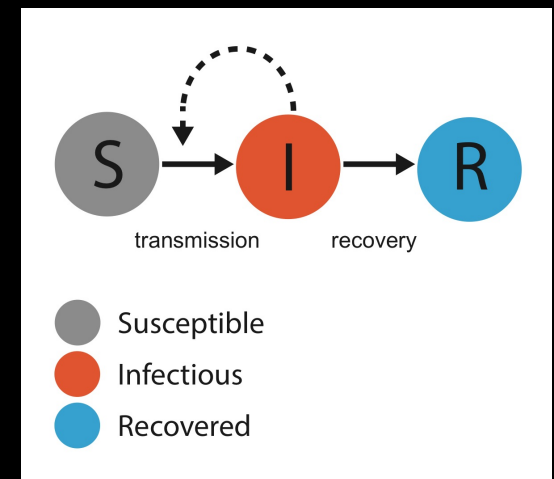
These data give us **infecteds** over time...



These data give us **infecteds** over time...



What *processes* contribute to the “infected” state in our system?



How to fit a model to data

1. Build a model that uses explicit processes to recover the same states as the data.
2. Use any statistical tool (i.e. maximum likelihood, least squares) to ask, assuming our model is true, how likely are we to recover the observed data?
3. Optimize the parameters behind the processes to make the model most likely to recover the data.
4. If need be, restructure your model to better match your data.

How to fit a model to data

1. Build a model that uses explicit processes to recover the same states as the data.

Discrete time models are simple:

$$S_{t+1} = S_t - \text{beta} * I_t * S_t / N$$

$$I_{t+1} = I_t + \text{beta} * I_t * S_t / N - \text{gamma} * I_t$$

where beta = transmission coefficient and gamma = recovery rate

How to fit a model to data

1. Build a model that uses explicit processes to recover the same states as the data.

If we set the timestep = $1/\text{gamma}$, we can reduce the system to:

$$S_{t+1} = S_t - \text{beta} * I_t * S_t / N_t$$

$$I_{t+1} = \text{beta} * I_t * S_t / N_t$$

This means we make the assumption that there are no overlapping infectious generations.

How to fit a model to data

1. Build a model that uses explicit processes to recover the same states as the data.

$$I_2 = I_1 + \text{beta} * I_1 * S_1 / N - \text{gamma} * I_1$$

$$I_2 = 1 + (.2)(1)(9/10) - (1)(1)$$

$I_1 = 1$ person

$S_1 = 9$ persons

$N = 10$

$\text{beta} = .2 \text{ hour}^{-1}$

$\text{gamma} = 1 \text{ hour}^{-1}$

$\Delta t = 1$ hour

How to fit a model to data

1. Build a model that uses explicit processes to recover the same states as the data.

$$I_2 = I_1 + \text{beta} * I_1 * S_1 / N - \text{gamma} * I_1$$

$$I_2 = 1 + (.2)(1)(9/10) - (1)(1)$$

$$I_2 = 1 + (.2)(1)(9/10) - 1$$

$$I_2 = .18 \text{ at 2 hours}$$

cancel!

$I_1 = 1$ person

$S_1 = 9$ persons

$N = 10$

$\text{beta} = .2 \text{ hour}^{-1}$

$\text{gamma} = 1 \text{ hour}^{-1}$

$\Delta t = 1 \text{ hour}$

How to fit a model to data

1. Build a model that uses explicit processes to recover the same states as the data.

$$I_2 = I_1 + \text{beta} * I_1 * S_1 / N - \text{gamma} * I_1$$

$$I_2 = 1 + (.2)(1)(9/10) - (1)(1)$$

$$I_2 = 1 + (.2)(1)(9/10) - 1$$

$$I_2 = .18 \text{ at 2 hours}$$

cancel!

$I_1 = 1$ person

$S_1 = 9$ persons

$N = 10$

$\text{beta} = .2 \text{ hour}^{-1}$

$\text{gamma} = 1 \text{ hour}^{-1}$

$\Delta t = 1 \text{ hour}$

$$I_2 = 1 + (.2)(2)(1)(9/10) - (1)(1)(2)$$

$I_1 = 1$ person

$S_1 = 9$ persons

$N = 10$

$\text{beta} = .2 \text{ hour}^{-1}$

$\text{gamma} = 1 \text{ hour}^{-1}$

$\Delta t = 2 \text{ hours}$

How to fit a **model** to **data**

1. Build a model that uses explicit processes to recover the same states as the data.

$$I_2 = I_1 + \text{beta} * I_1 * S_1 / N - \text{gamma} * I_1$$

$$I_2 = 1 + (.2)(1)(9/10) - (1)(1)$$

$$I_2 = \textcircled{1} + (.2)(1)(9/10) - \textcircled{1}$$

$$I_2 = .18 \text{ at 2 hours}$$

cancel!

$I_1 = 1$ person

$S_1 = 9$ persons

$N = 10$

$\text{beta} = .2 \text{ hour}^{-1}$

$\text{gamma} = 1 \text{ hour}^{-1}$

$\Delta t = 1 \text{ hour}$

$$I_2 = 1 + (.2)(2)(1)(9/10) - (1)(1)(2)$$

$$I_2 = \textcircled{1} + (.2)(2)(1)(9/10) - \textcircled{2}$$

$$I_2 = -.64 \text{ at 4 hours}$$

Don't cancel!

$I_1 = 1$ person

$S_1 = 9$ persons

$N = 10$

$\text{beta} = .2 \text{ hour}^{-1}$

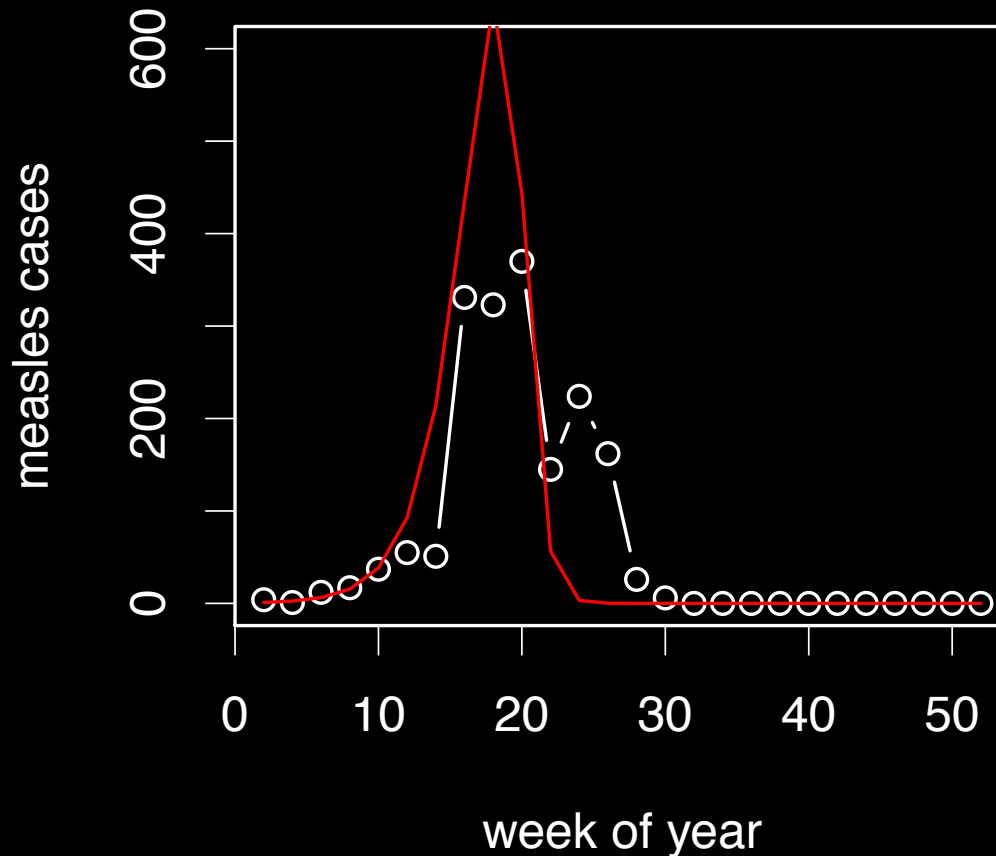
$\text{gamma} = 1 \text{ hour}^{-1}$

$\Delta t = 2 \text{ hours}$

How to fit a **model** to **data**

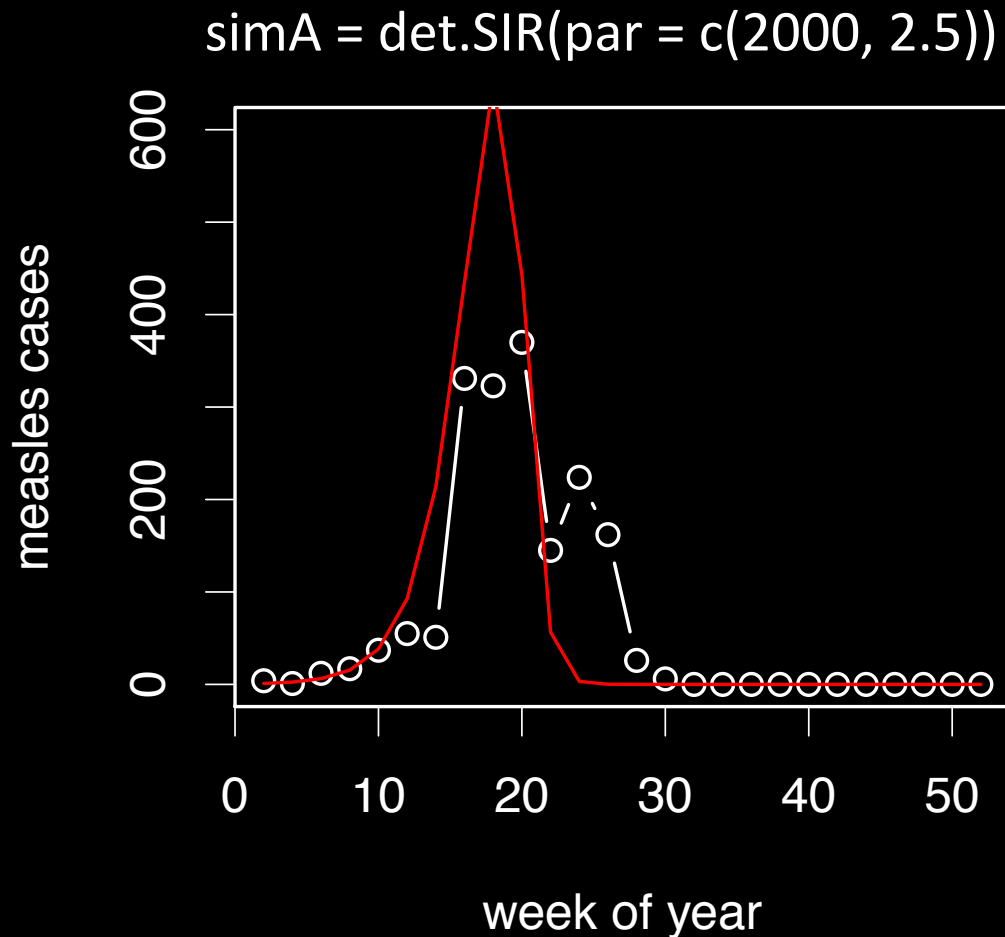
1. Build a model that uses explicit processes to recover the same states as the data.

`simA = det.SIR(par = c(2000, 2.5))`



How to fit a **model** to **data**

1. Build a model that uses explicit processes to recover the same states as the data.



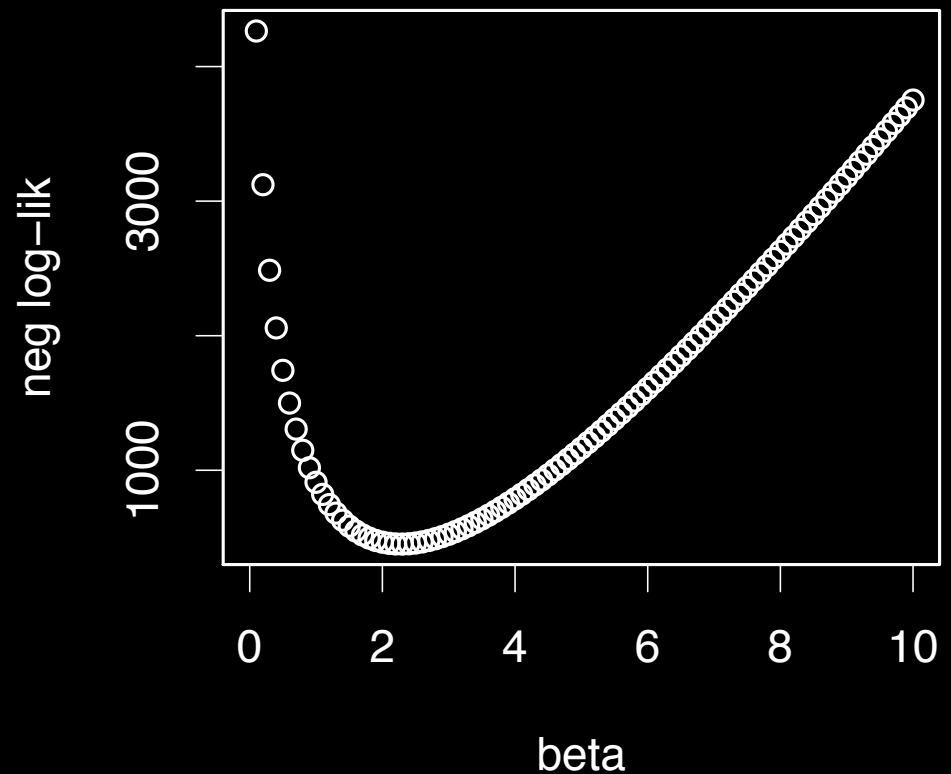
Model does a pretty good job, but but it overshoots our data by quite a bit.

What does this suggest about our guess for β ?

How to fit a model to data

2. Use any maximum likelihood to ask, assuming our model is true, how likely are we to recover the observed data?

`beta_guess[which.min(ll)] = 2.3`

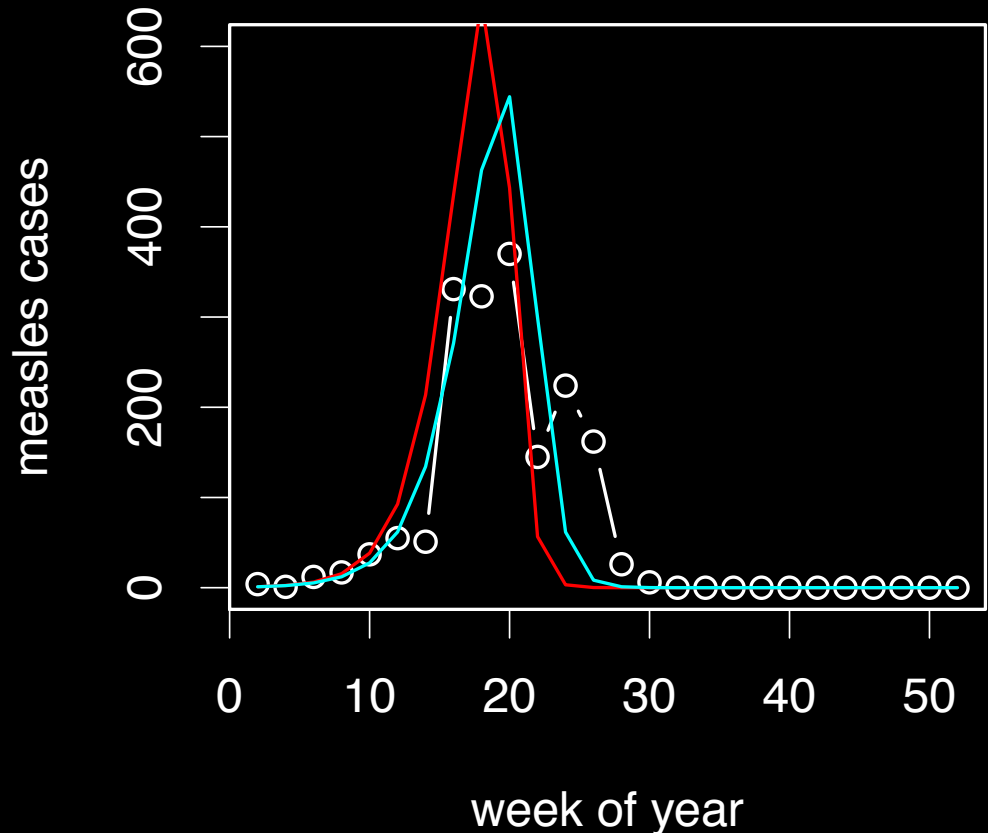


How to fit a **model** to **data**

2. Use any maximum likelihood to ask, assuming our model is true, how likely are we to recover the observed data?

```
simB = det.SIR(par = c(2000, 2.3))
```

New beta fits even better!



How to fit a **model** to **data**

3. Optimize the parameters behind the processes to make the model most likely to recover the data.

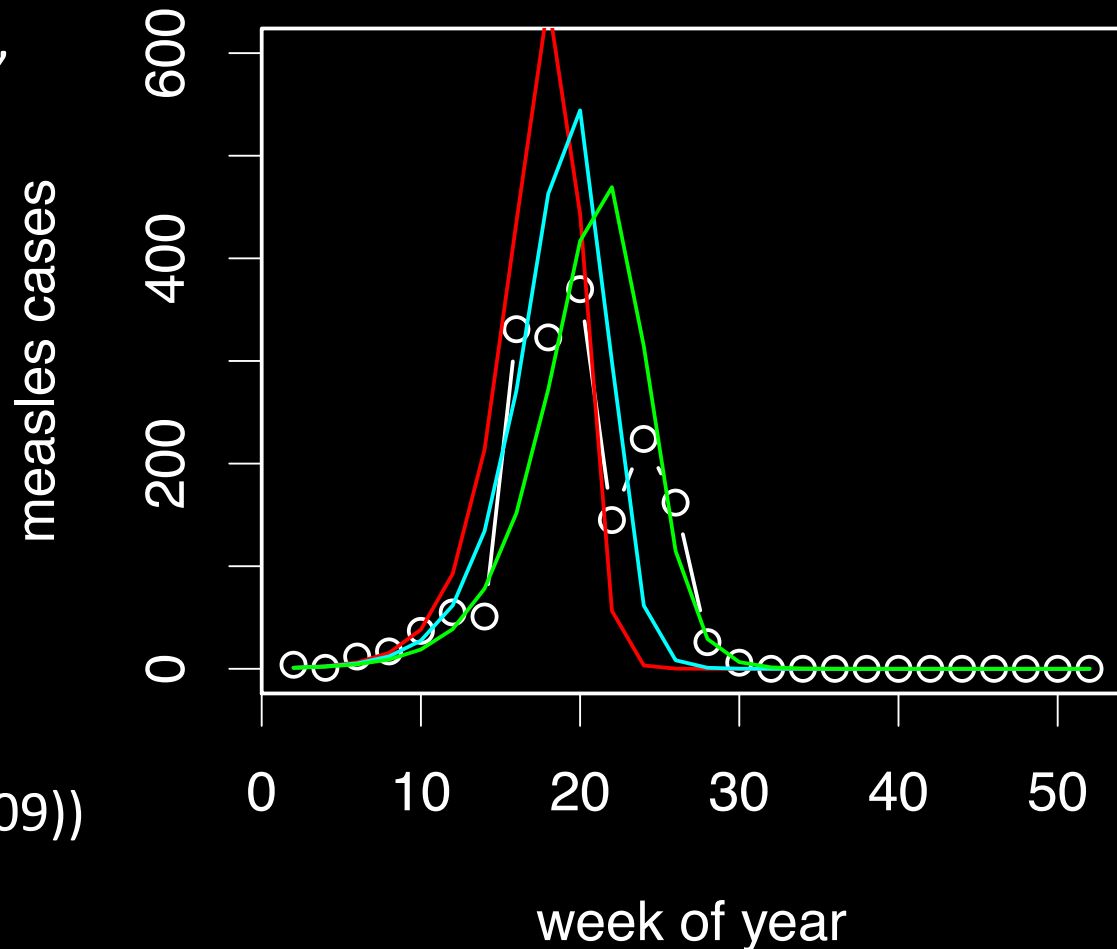
```
out <- optim(par = c(2000, 2.5),  
  likelihood,  
  method = "Nelder-Mead",  
  l = dat$cases)
```

```
out$par = c(2152, 2.09)
```

Looks great!

But why not perfect?

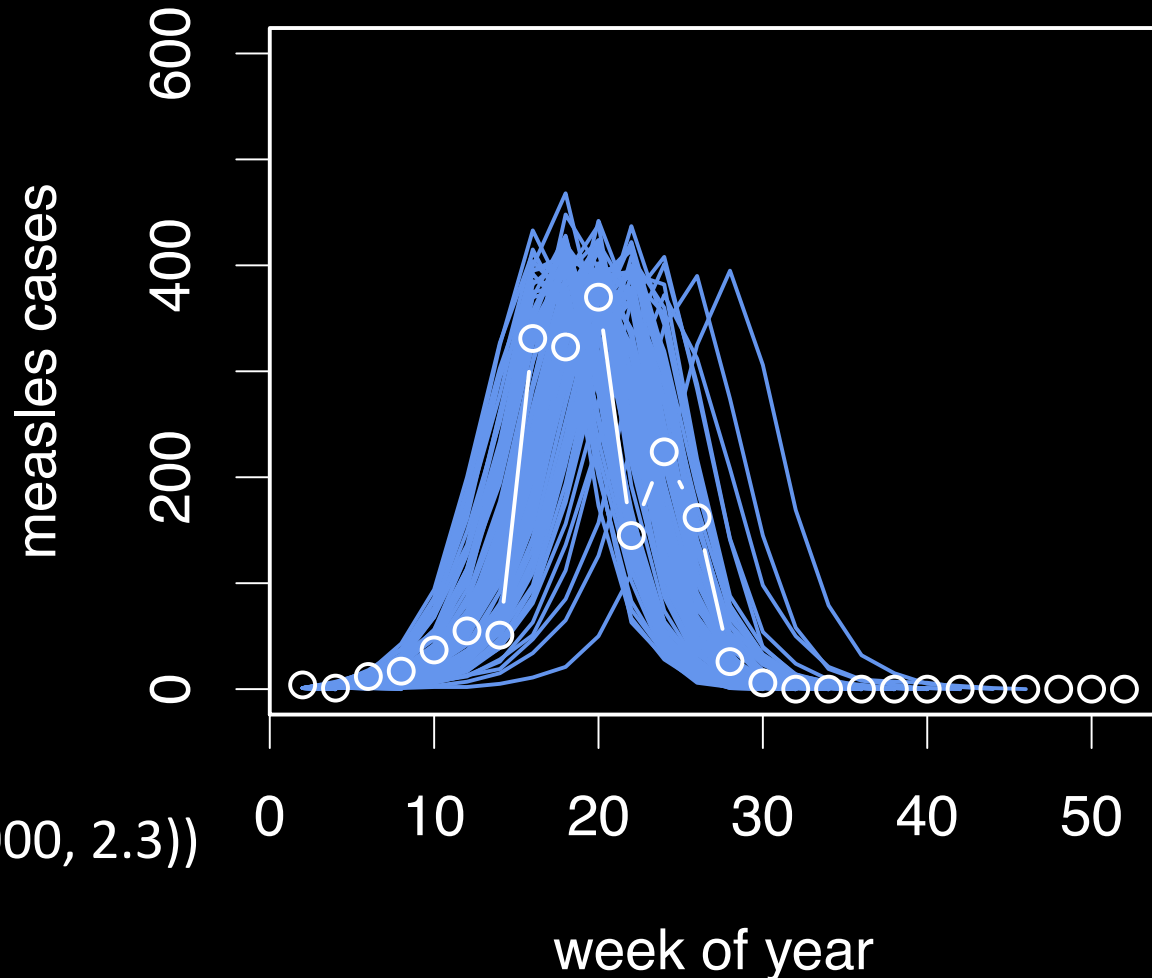
```
simC = det.SIR(par = c(2152, 2.09))
```



How to fit a model to data

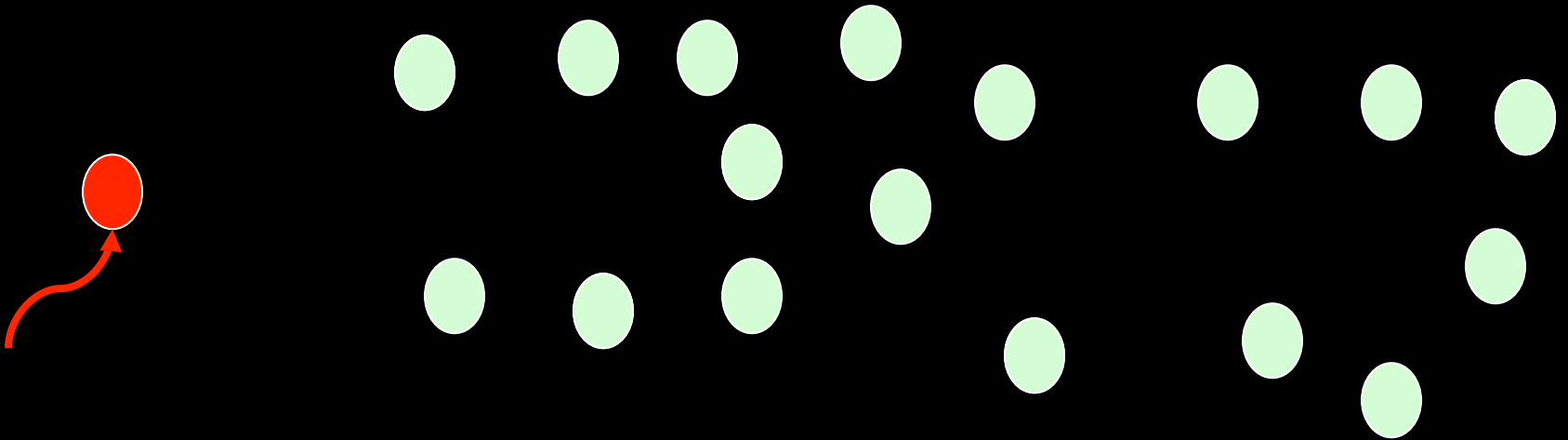
4. If need be, restructure your model to better match your data.

```
for (i in 1:100) {  
  stoch = stoch.SIR(par=c(2000, 2.3))  
}
```



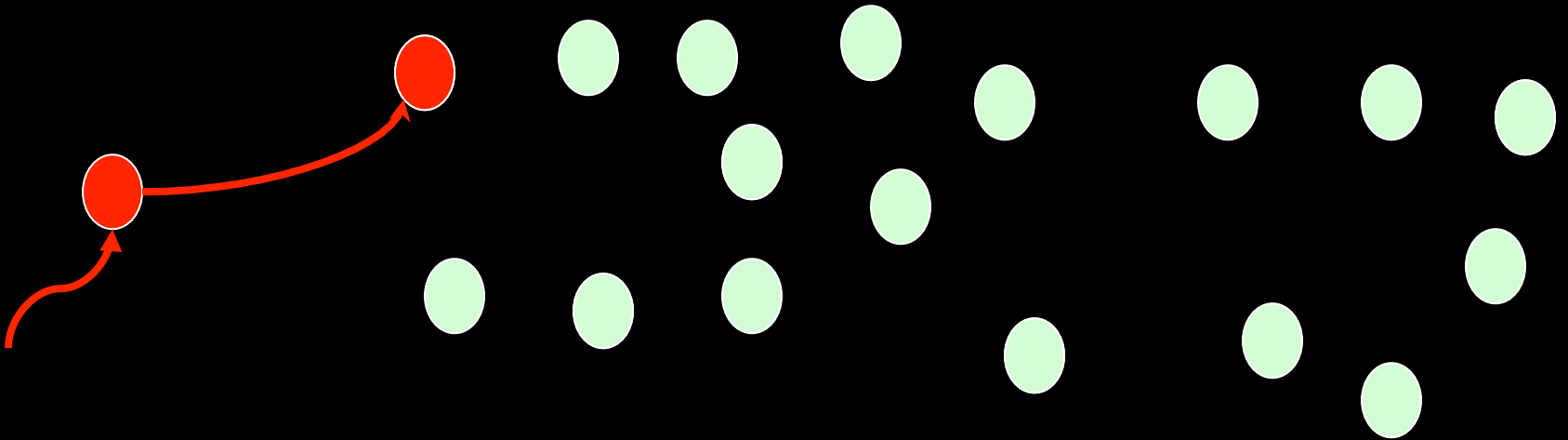
Insights from fitting dynamic models: R_0

- The basic reproduction number for a pathogen
- Defined as the number of new infections generated by one existing infection in a completely susceptible host population
- In these discrete time models, $R_0 = \beta$



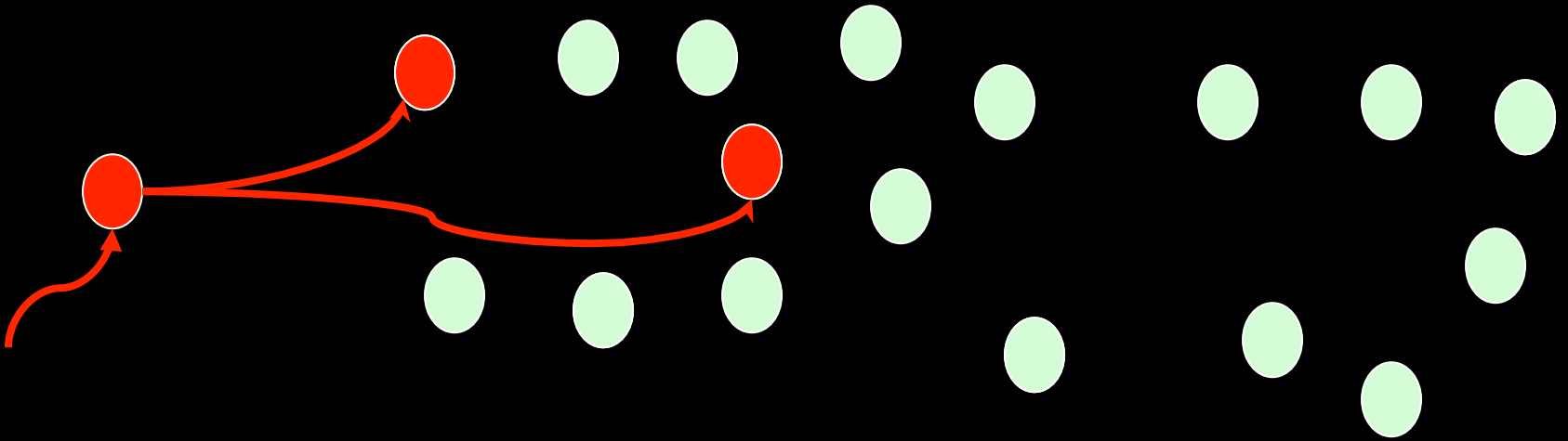
Insights from fitting dynamic models: R_0

- The basic reproduction number for a pathogen
- Defined as the number of new infections generated by one existing infection in a completely susceptible host population
- In these discrete time models, $R_0 = \beta$



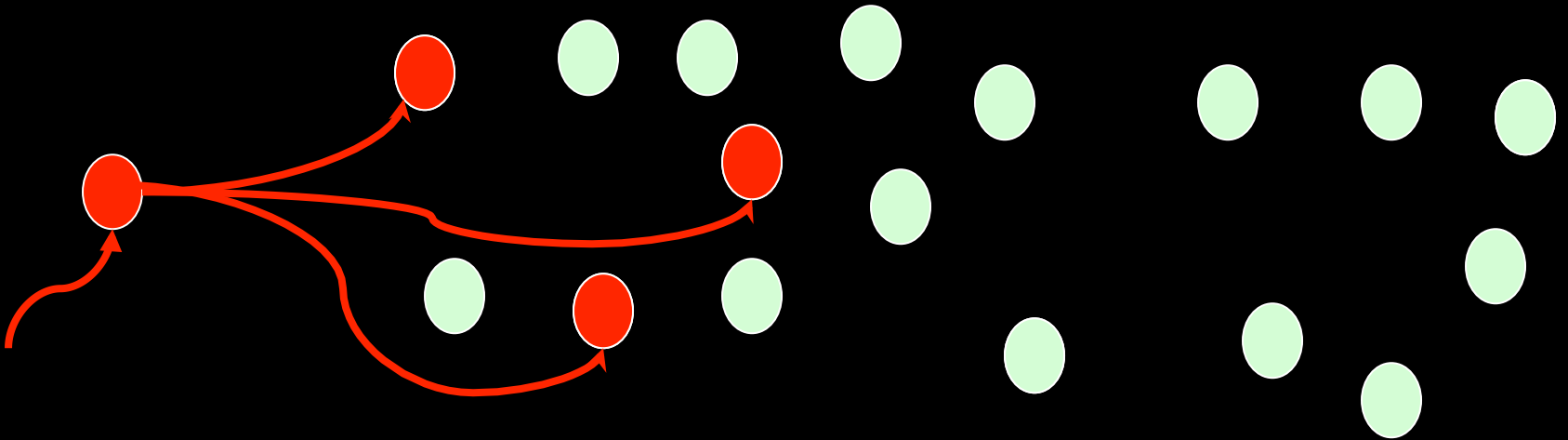
Insights from fitting dynamic models: R_0

- The basic reproduction number for a pathogen
- Defined as the number of new infections generated by one existing infection in a completely susceptible host population
- In these discrete time models, $R_0 = \beta$



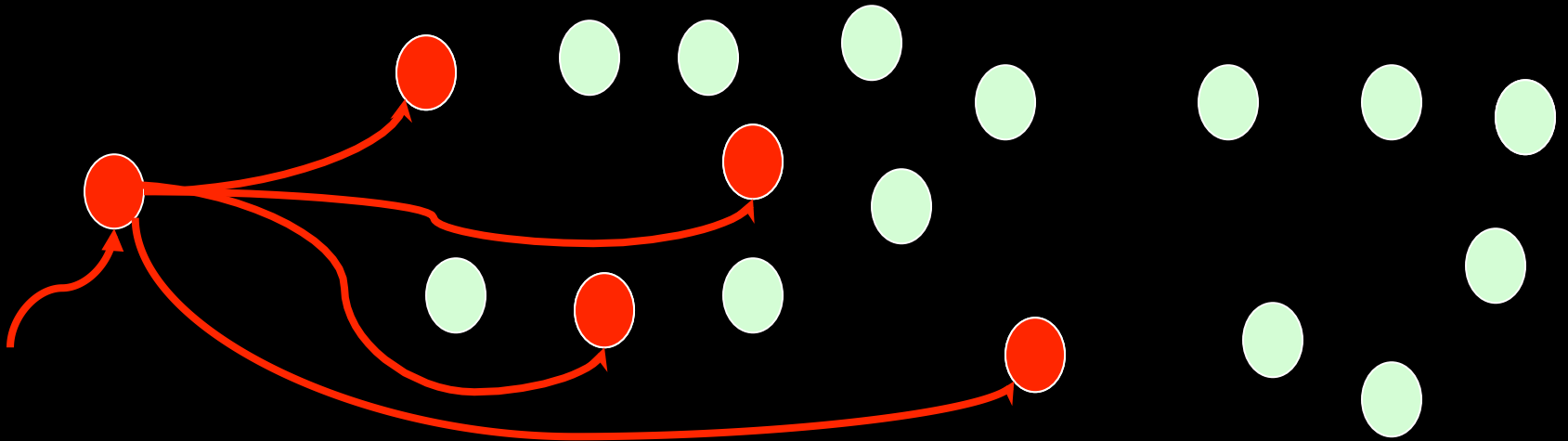
Insights from fitting dynamic models: R_0

- The basic reproduction number for a pathogen
- Defined as the number of new infections generated by one existing infection in a completely susceptible host population
- In these discrete time models, $R_0 = \beta$



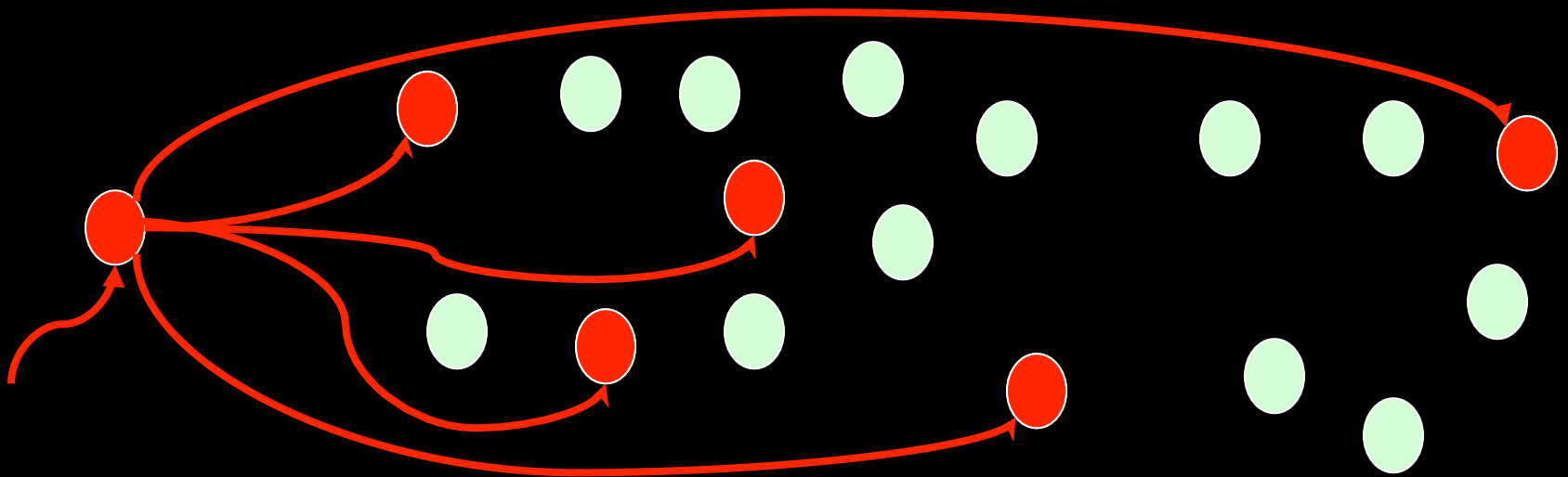
Insights from fitting dynamic models: R_0

- The basic reproduction number for a pathogen
- Defined as the number of new infections generated by one existing infection in a completely susceptible host population
- In these discrete time models, $R_0 = \beta$



Insights from fitting dynamic models: R_0

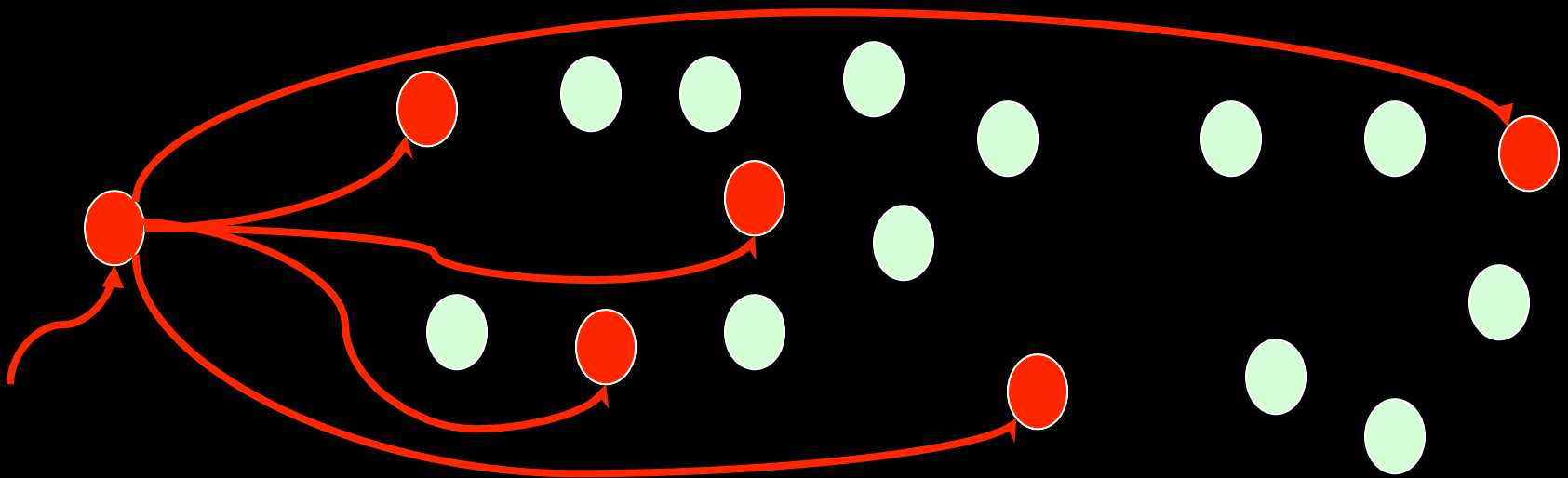
- The basic reproduction number for a pathogen
- Defined as the number of new infections generated by one existing infection in a completely susceptible host population
- In these discrete time models, $R_0 = \beta$



$$R_0 = 5$$

Insights from fitting dynamic models: R-effective

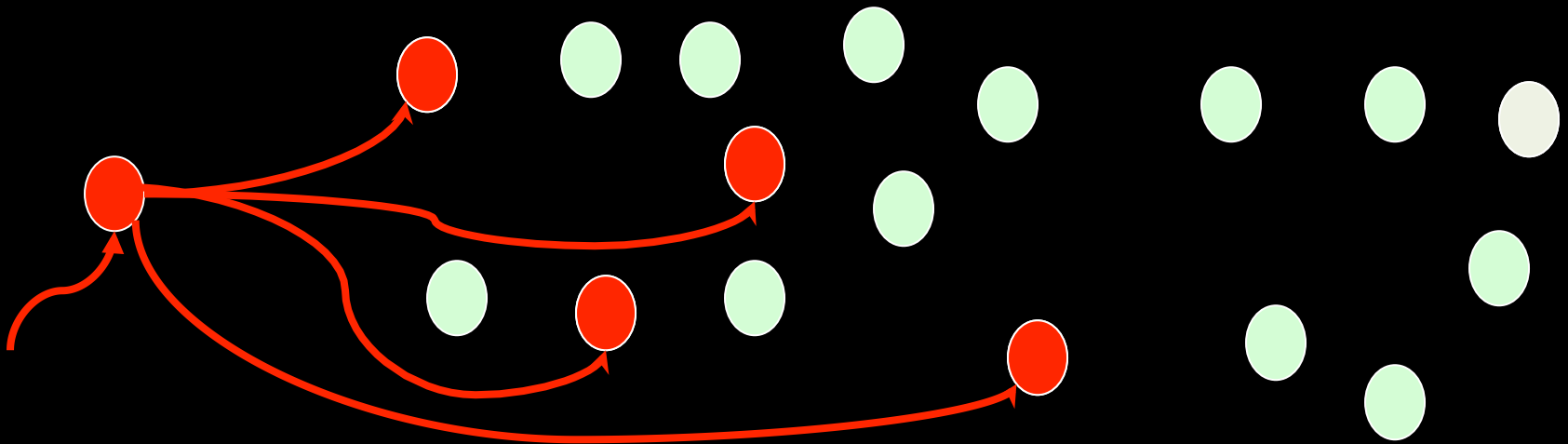
- The number of new infections generated by one existing infection in a **partially susceptible** host population
- $R_{\text{eff}} = R_0 * (S/N)$
- Changes over time!



$$R_{\text{eff}} = (5) * (12/17) = 3.53$$

Insights from fitting dynamic models: R-effective

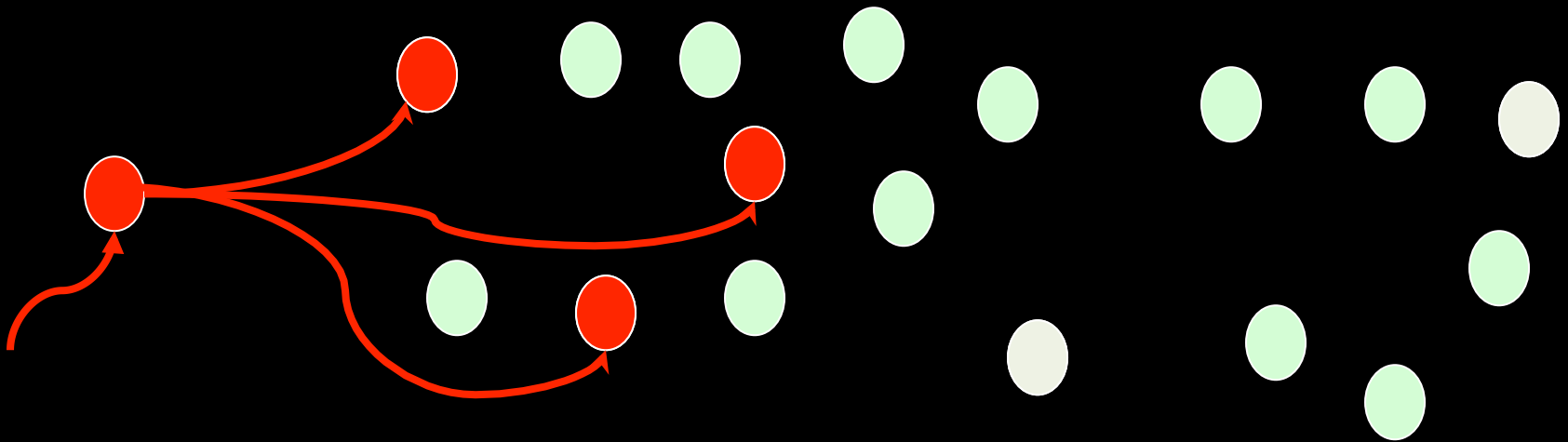
- The number of new infections generated by one existing infection in a **partially susceptible** host population
- $R_{eff} = R_0 * (S/N)$
- Changes over time!



$$R_{eff} = (5) * (13/17) = 3.82$$

Insights from fitting dynamic models: R-effective

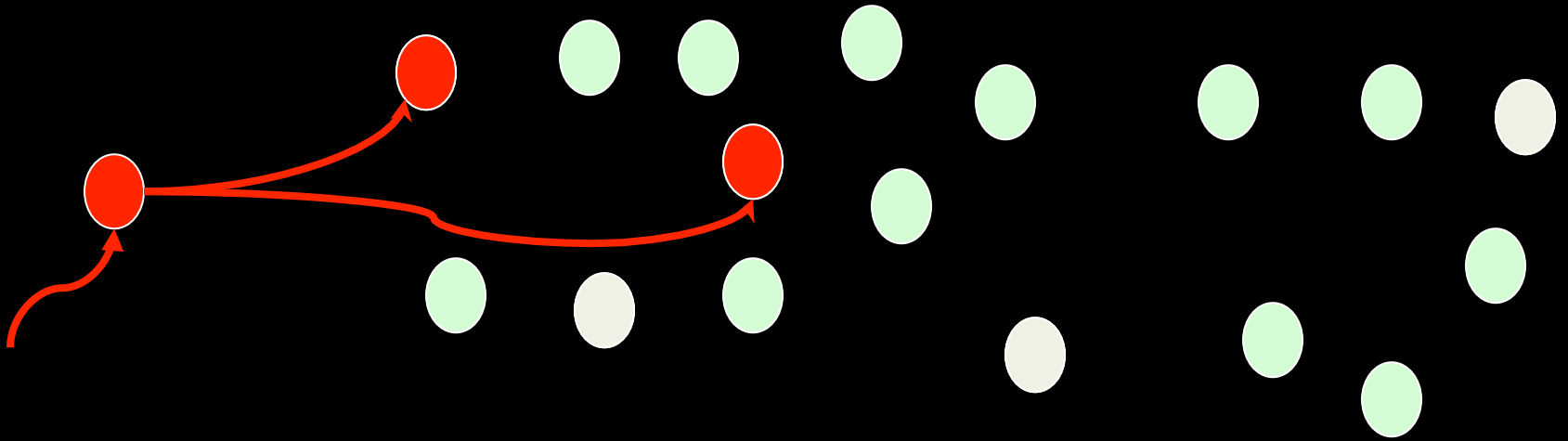
- The number of new infections generated by one existing infection in a **partially susceptible** host population
- $R_{eff} = R_0 * (S/N)$
- Changes over time!



$$R_{eff} = (5) * (14/17) = 4.12$$

Insights from fitting dynamic models: R-effective

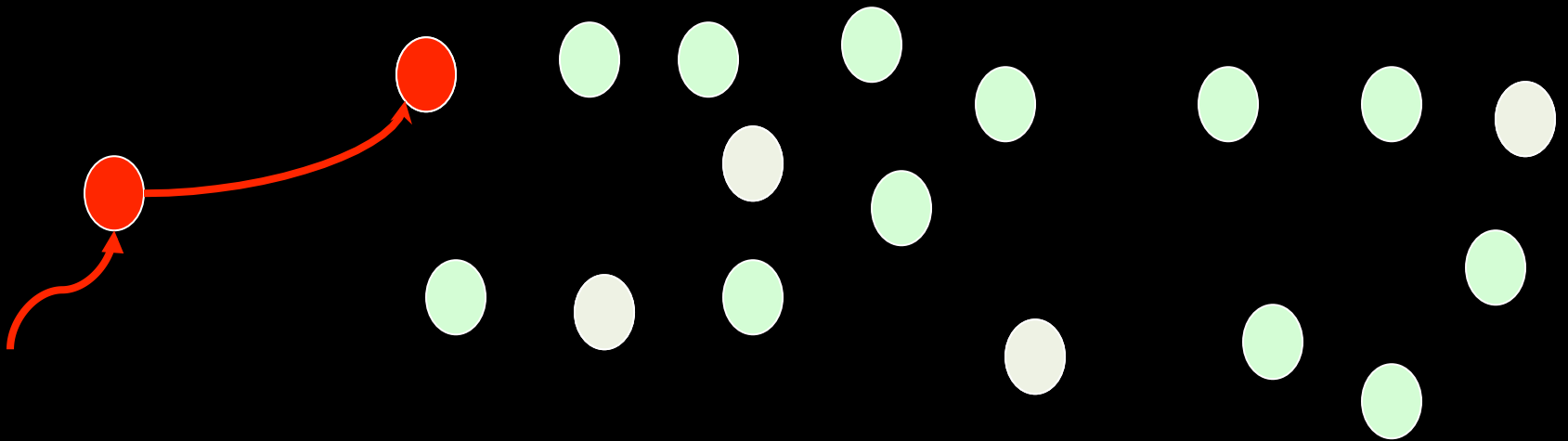
- The number of new infections generated by one existing infection in a **partially susceptible** host population
- $R_{eff} = R_0 * (S/N)$
- Changes over time!



$$R_{eff} = (5) * (15/17) = 4.41$$

Insights from fitting dynamic models: R-effective

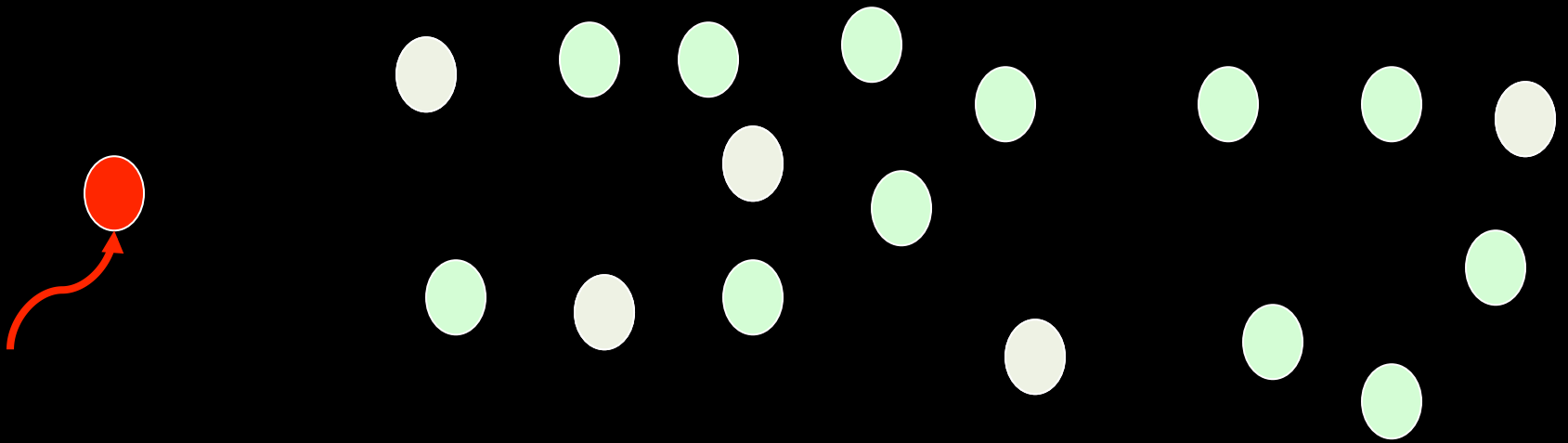
- The number of new infections generated by one existing infection in a **partially susceptible** host population
- $R_{eff} = R_0 * (S/N)$
- Changes over time!



$$R_{eff} = (5) * (16/17) = 4.71$$

Insights from fitting dynamic models: R-effective

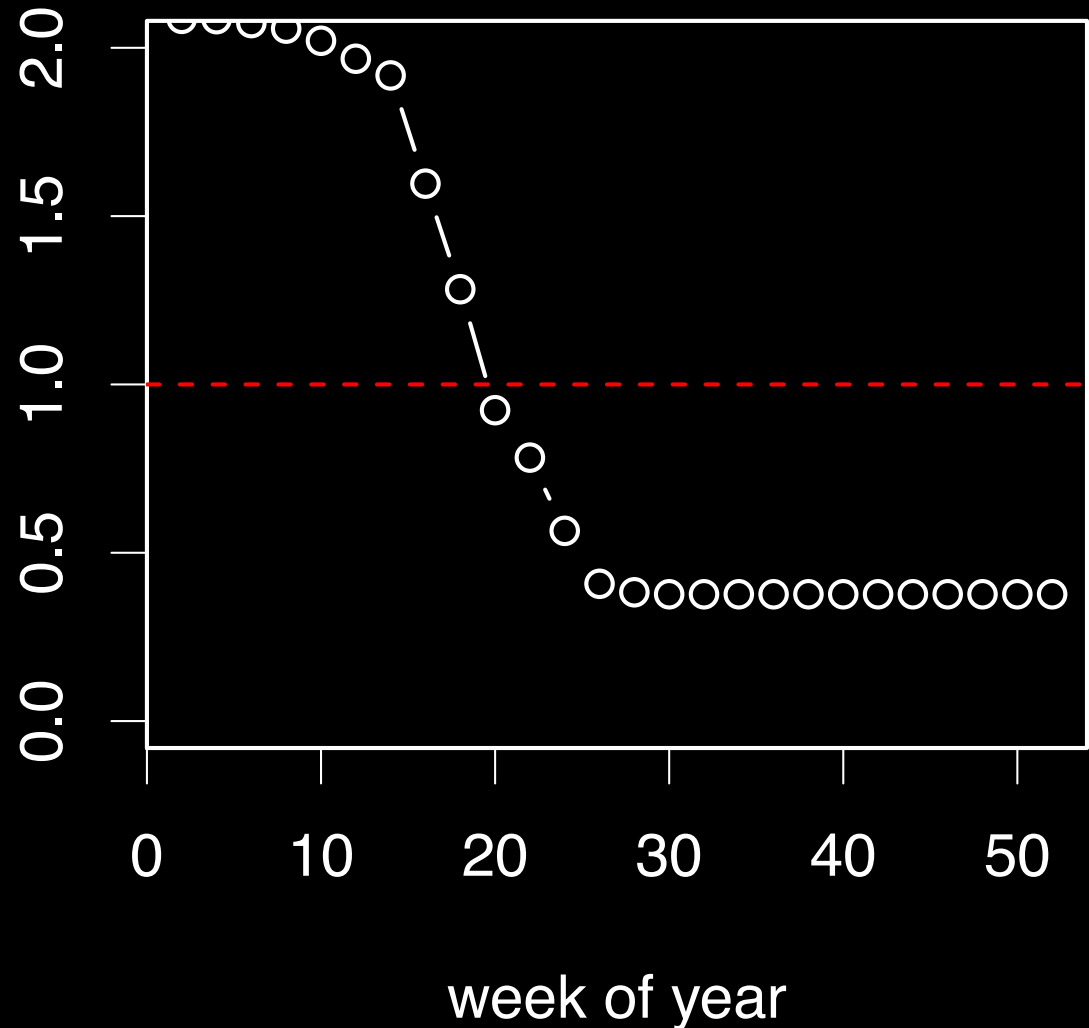
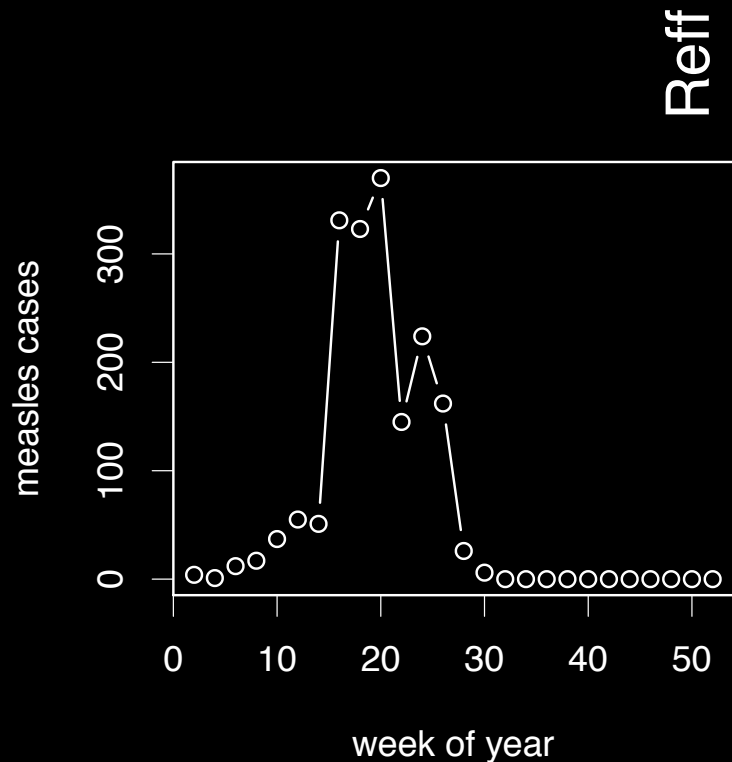
- The number of new infections generated by one existing infection in a **partially susceptible** host population
- $R_{eff} = R_0 * (S/N)$
- Changes over time!



$$R_{eff} = (5) * (17/17) = 5$$

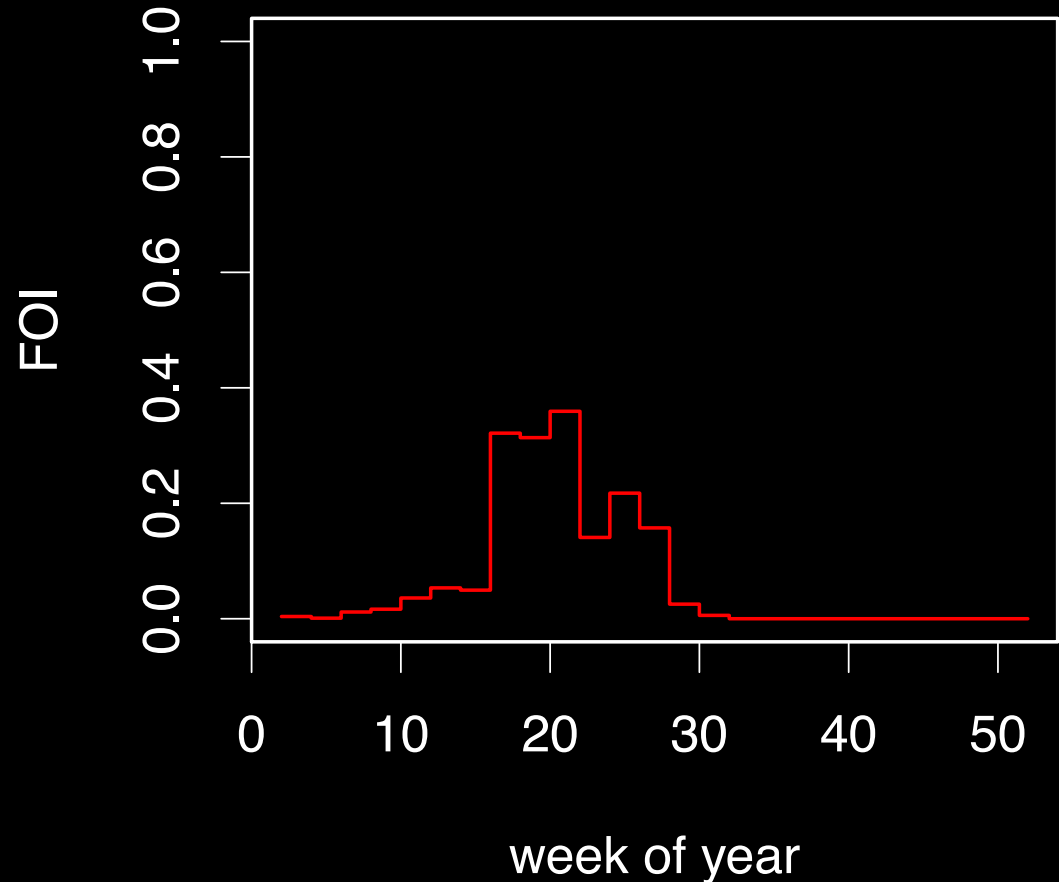
Insights from fitting dynamic models: R-effective

An epidemic spreads
when $R_{eff} > 1$
and declines when
 $R_{eff} < 1$



Insights from fitting dynamic models: Force of Infection

- Defined as the rate at which susceptibles become infected
- $FOI = R_0 * (I/N)$
- Changes over time, but constant across each timestep



Mechanistic modeling is **process**-driven...

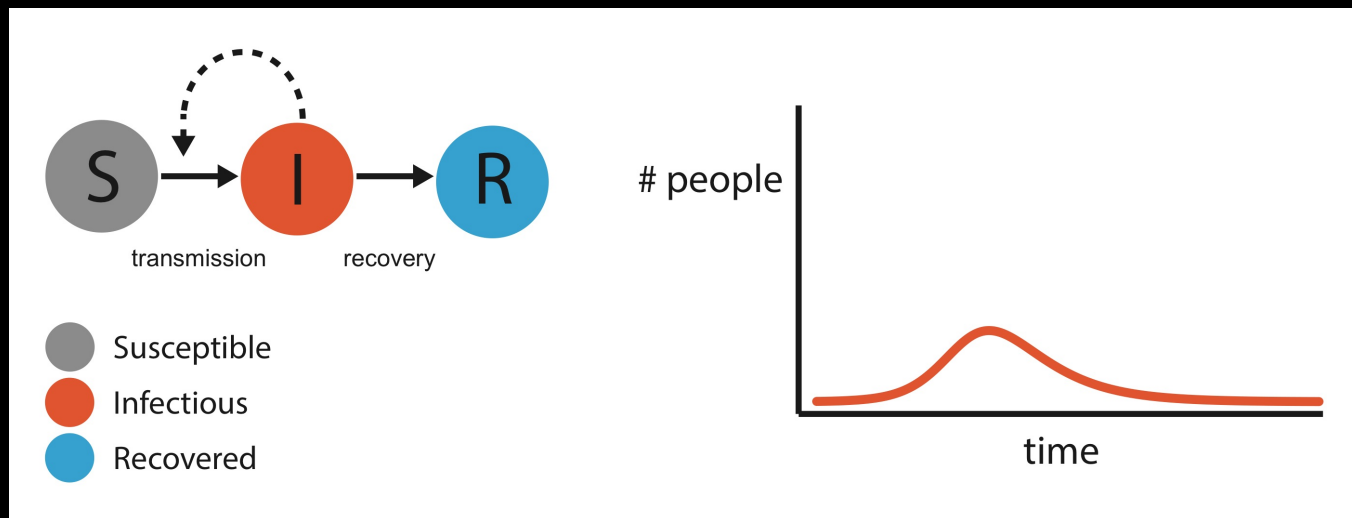
- We want to understand **what** happened, **when** it happened, and **why** it happened
- Allows us to scale up from **individual**-level processes to **population**-level patterns
- We start by building a **model** that uses explicit **processes** to recover the same outcomes (“**states**”) as our **data**

Mechanistic modeling is **process**-driven...

- Test “what if” scenarios not amenable to experimentation

Mechanistic modeling is **process**-driven...

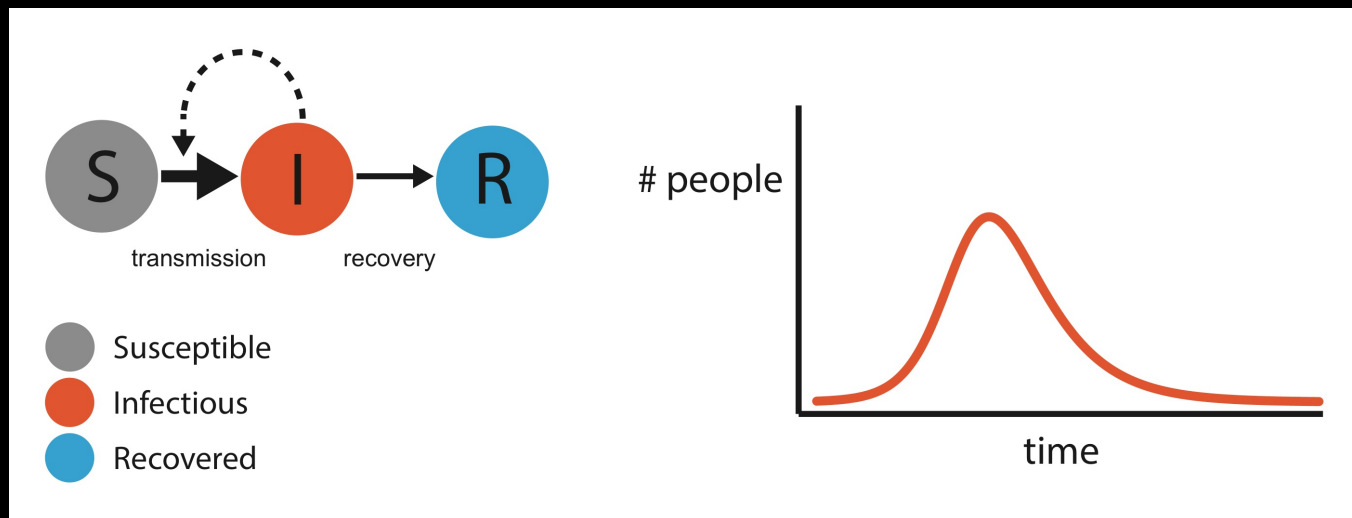
- Test “what if” scenarios not amenable to experimentation



Mechanistic modeling is **process**-driven...

- Test “what if” scenarios not amenable to experimentation

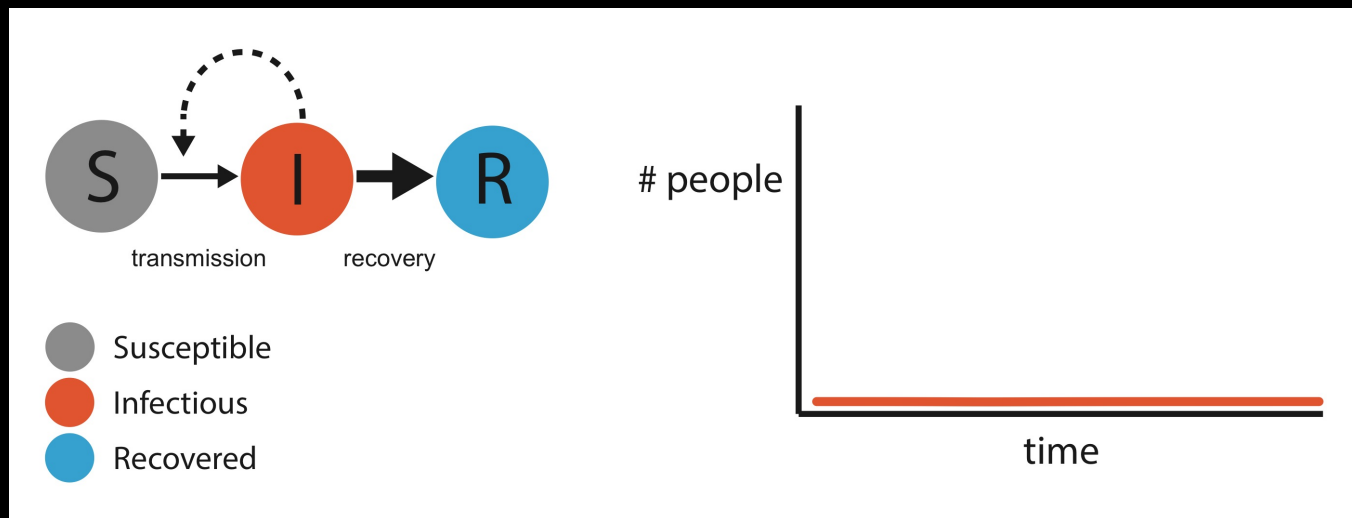
What if each person exposed 50% more people?



Mechanistic modeling is **process**-driven...

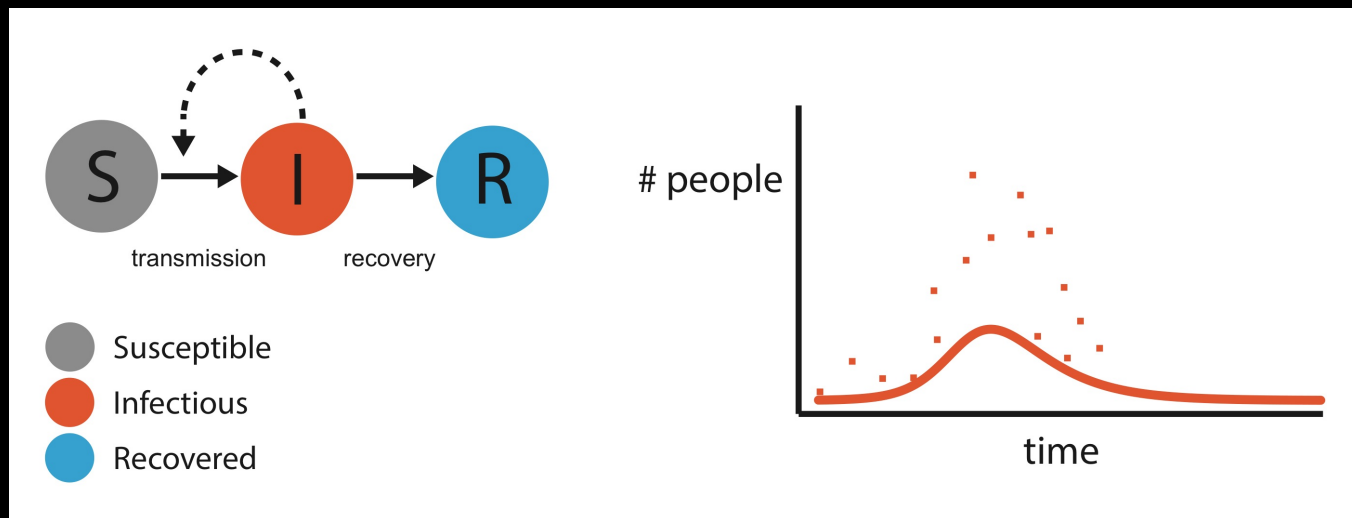
- Test “what if” scenarios not amenable to experimentation

What if we treated people and doubled the rate of recovery?



Mechanistic modeling is **process**-driven...

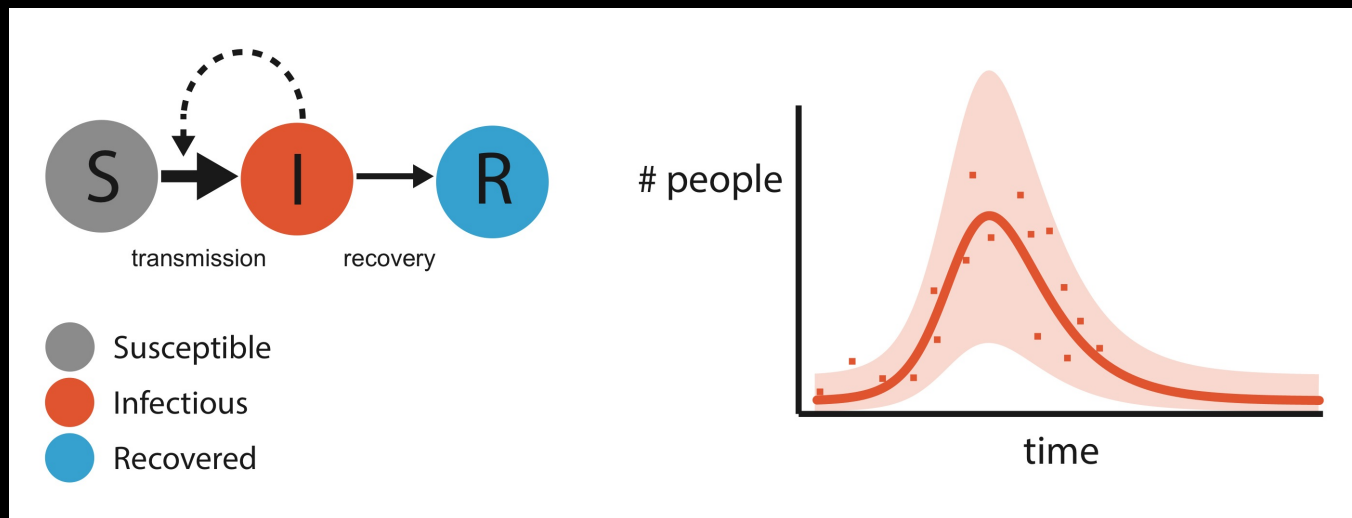
- Test “what if” scenarios not amenable to experimentation
- Estimate parameters that are difficult to measure by fitting models to available data



Mechanistic modeling is **process**-driven...

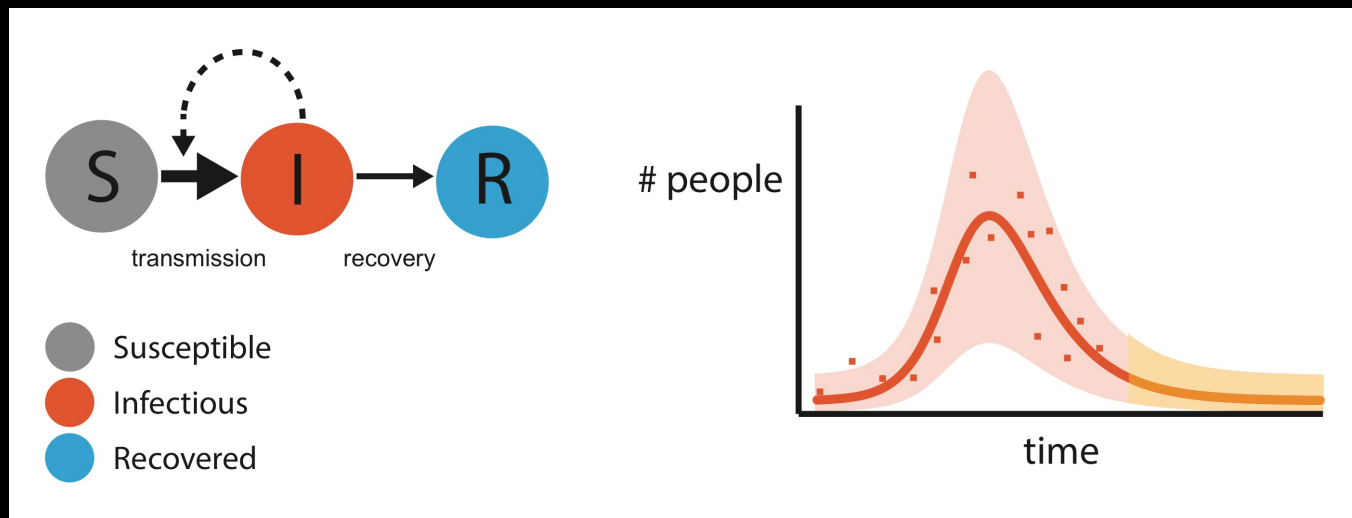
- Test “what if” scenarios not amenable to experimentation
- Estimate parameters that are difficult to measure by fitting models to available data

Estimate transmission rate or other model parameters
(with confidence intervals)



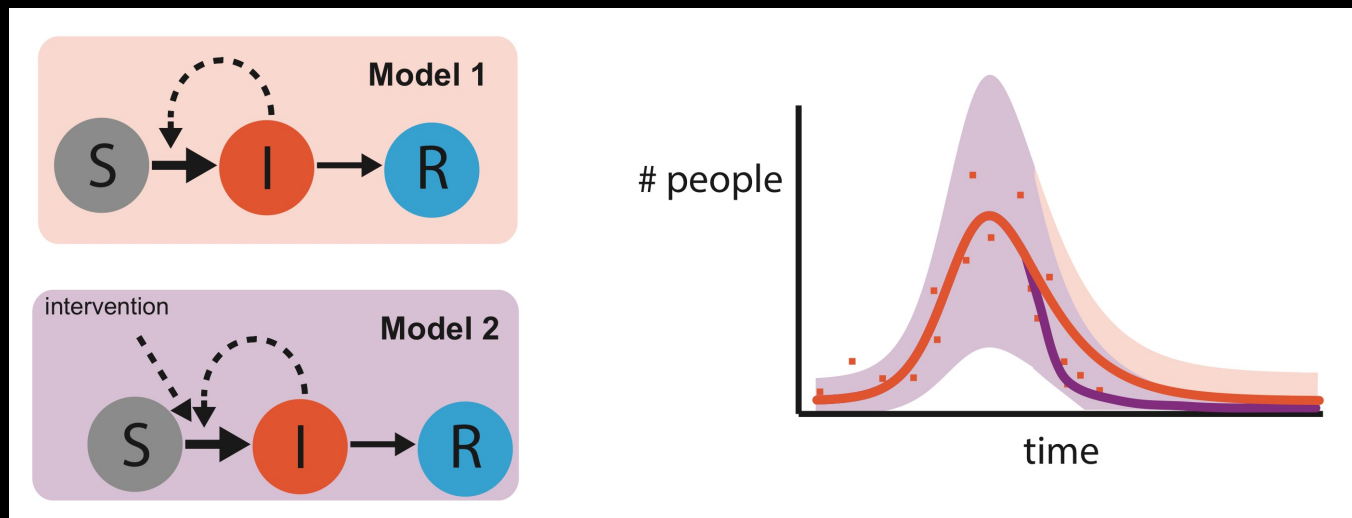
Mechanistic modeling is **process**-driven...

- Test “what if” scenarios not amenable to experimentation
- Estimate parameters that are difficult to measure by fitting models to available data
- Forecast forward in time



Mechanistic modeling is **process**-driven...

- Test “what if” scenarios not amenable to experimentation
- Estimate parameters that are difficult to measure by fitting models to available data
- Forecast forward in time
- Select between models of differing hypotheses



Mechanistic modeling is **process**-driven...

- **Estimate**: time series of state variables/ parameters of interest → **DATA**
- **Inference**: Build **model** to recapture **data**. Fit to optimize parameters and “infer” the process underlying the data
- Model assessment: Assess **plausibility** or **model comparison**
- End goal: **explain** observed patterns or **predict**