

### Model Me This: COVID-19 Scientific Predictions and Where We Go From Here

Tuesday, March 30

### Featuring: Tim Scheibe, PhD

Lab Fellow, Earth Scientist Environmental Dynamics & Simulations

### **DEMYSTIFYING COVID:**

A Special Edition Seminar Series



COMMUNITY SCIENCE & TECHNOLOGY SEMINAR SERIES @PNNL



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### **DEMYSTIFYING COVID:**

A Special Edition Seminar Series



EVERY TUESDAY IN MARCH 5:00-6:00 P.M.



MARCH02 Hindsight is 2020: The Science **Behind COVID-19** 

### **Presented by Steve Wiley**

What lessons have we learned over the last few months? What's left for us to uncover? And seriously what is the difference between a cold, a flu, and COVID symptoms?



MARCH09

What Do Bats Have to Do with It?

### **Presented by Amy Sims**

Bats, pangolins, and humans—oh my! This talk will explore the role wild animals play in the emergence of new diseases.



MARCH**16 Behind the Mask: The Science on** Stopping the Spread

### **Presented by Katrina Waters**

What measures keep our communities safe? And why do some strange, sometimes serious health effects linger even after COVID-19 has gone, including a loss of taste and smell or COVID toe? Join us to find out.



MARCH23

### **Presented by Kristin Omberg**

If you're confused about COVID-19 testing and vaccines, you're not alone. This talk will explore the science behind the 400+ diagnostic tests and 200+ vaccine candidates produced over the last year.



### **Presented by Tim Scheibe**

Using mathematical models, scientists across the globe are beginning to arrive at a more complete picture of how and why COVID-19 spread across geographical locations and human populations.

### Testing, Testing, 1, 2, 3 (And What's Up With The New Vaccine, Anyways?)

### Model Me This: COVID-19 Scientific Predictions and Where We Go from Here





# COMMUNITY REPRESENTATIVES





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# TODAY'S SPEAKER



# Tim Scheibe

**Lead Scientist** River Corridor Scientific Focus Area Project



### EVERY TUESDAY IN MARCH 5:00-6:00 P.M.



# When you hear the word "model", what do you think about?

Start the presentation to see live content. For screen share software, share the entire screen. Get help at pollev.com/app















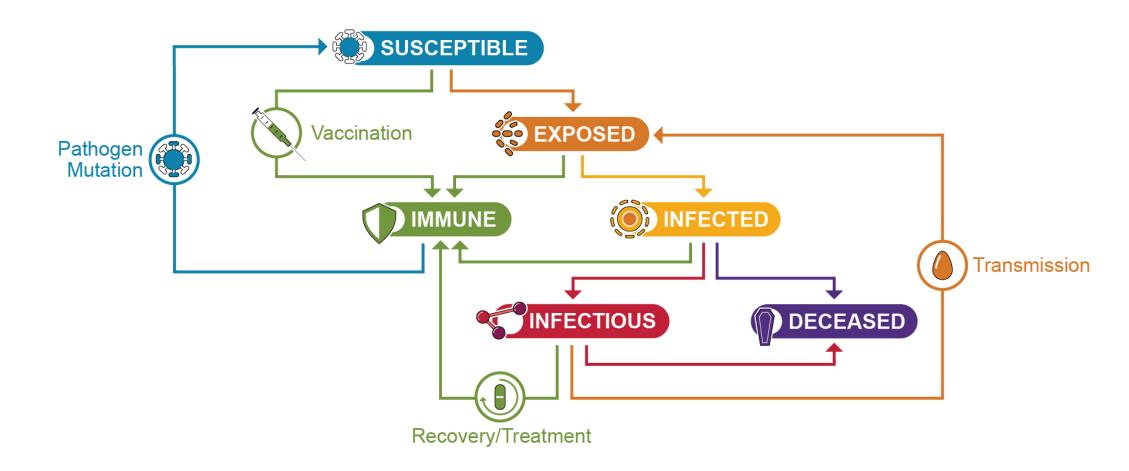






# **Today's discussion:** models and scientific prediction

### VIRAL INFECTION OVERVIEW





# Terminology

## **Pandemic**

• An outbreak of a disease that is prevalent over a whole country or world

### **Transmission**

Spreading a disease

## **Mutation**

• A genetic change in a disease that might or might not impact its transmission, symptoms or prior immunity

### Infectious/Contagious

• The state of being able to transmit a disease to another person

## Vaccine

• A non-harmful (hopefully) proxy of the disease agent that that can induce a protective immune response

## Model

• A representation of a disease or disease process that can recapitulate key aspects

### Reservoir

• Any person, animal, plant, soil or substance in which an infectious agent normally lives and multiplies



## What is a model?

"An approximate representation of a complex system that we use to learn about, encode knowledge of, and predict the behavior of that system"



### **Descriptive Model: "Tim Scheibe is...** ...a nerdy scientist who programs computers"

### "All models are wrong, some are useful."

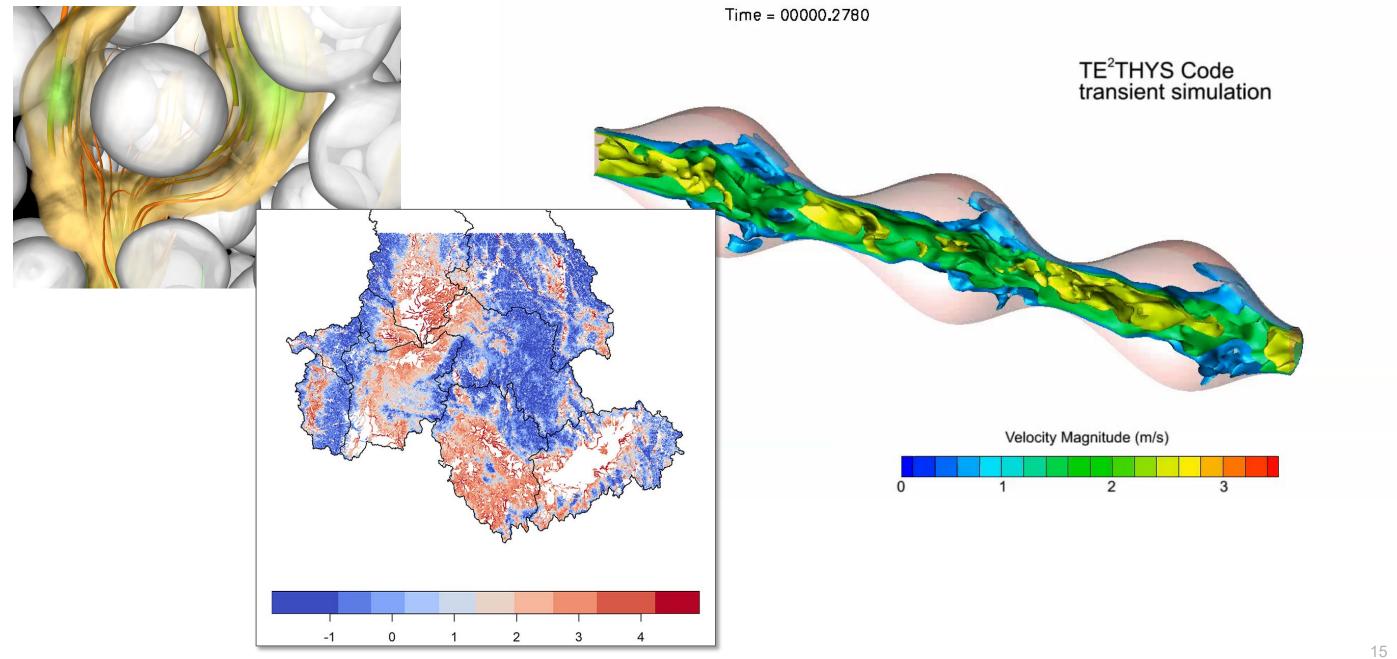
### **George Box** Statistician







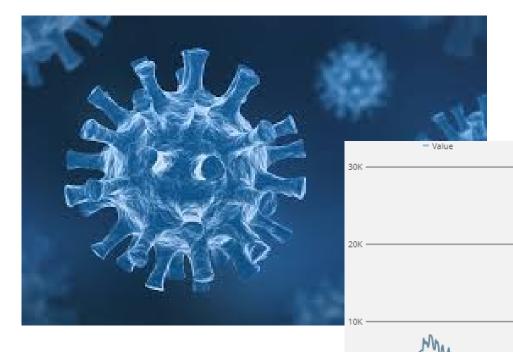
# Some of "my" models





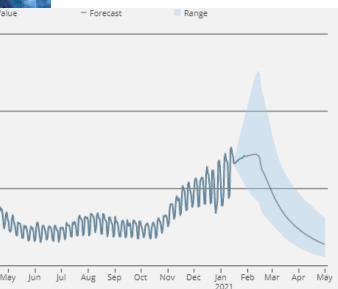
### Can we use mathematical models to learn about, encode knowledge of, and predict the spread of diseases like COVID-19?

• "An approximate representation of a complex system that we use to learn about, encode knowledge of, and predict the behavior of that system."



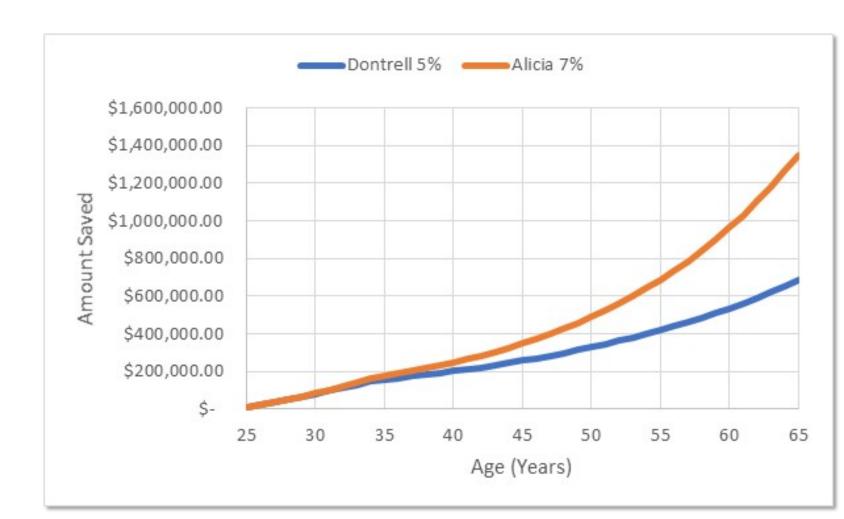
### "All models are wrong, some are useful."

### George Box Statistician



# A very simple mathematical model – $R_0$ ("r-zero" or "r-not")

Have you ever heard about "The Power of Compound Interest?"



Pacific

Northwest





# A very simple mathematical model – $R_0$

- $R_0$  is the basic reproduction number the average number of new people infected by one infected person.
  - Dontrell had COVID-19 and spread it to one other person in his home
  - Alicia had COVID-19 but quarantined early and did **not** spread it to anyone else
  - Olivia had COVID-19 and spread it to two other people before she became symptomatic
  - Brandon had COVID-19 without ever getting symptoms and unknowingly spread it to six other people
- The average number of new infections per person for this example is:

• 
$$R_0 = (1 + 0 + 2 + 6)/4 = 2.25$$

 $I_{t+1} = R_o I_t$ 

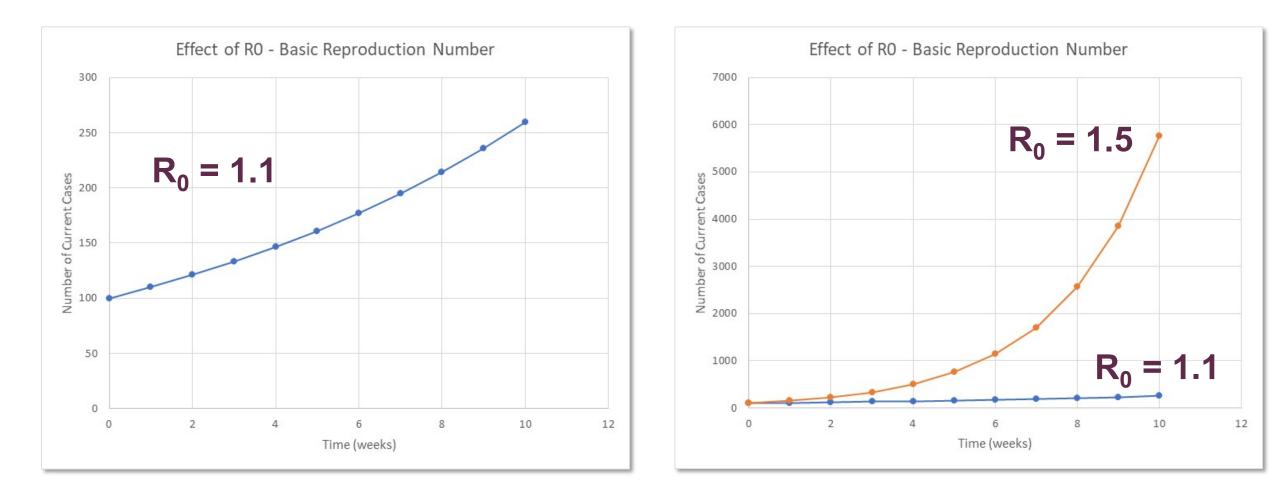




# A very simple mathematical model – R<sub>0</sub>

### • What can we learn from this model?

Like compound interest, a small change in R0 leads to a big difference in the number of infections over the same period.

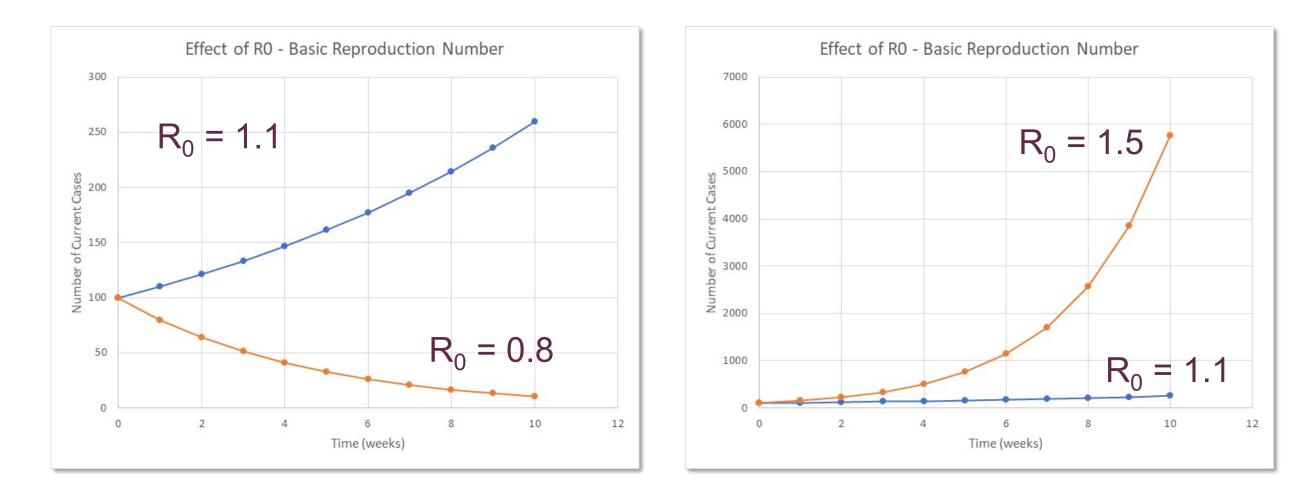






# A very simple mathematical model – R<sub>0</sub>

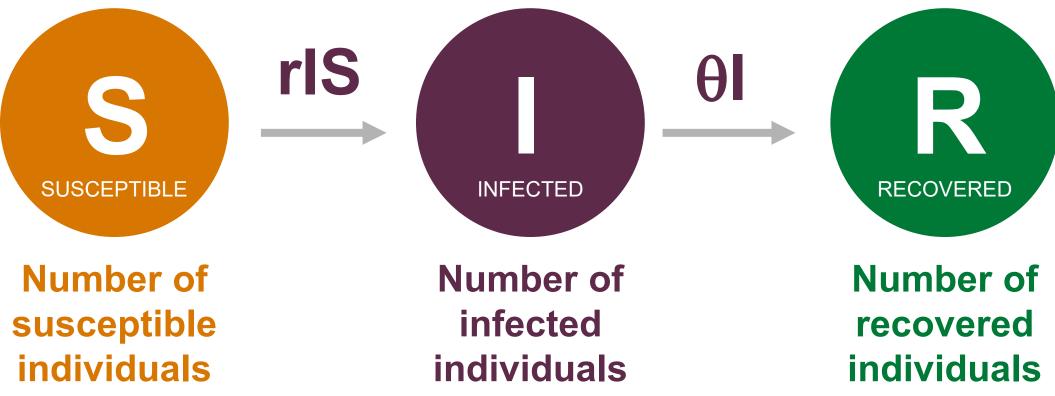
- What can we learn from this model?
  - If R<sub>0</sub> is less than 1.0, the pandemic will be controlled











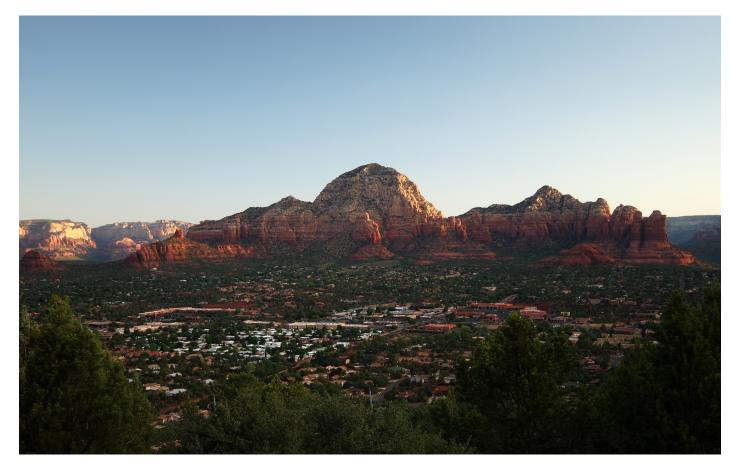
*r* = infection rate parameter  $\theta$  = recovery rate parameter



# A hypothetical example

### Imagine...

- A small city (population 10,000) remotely located, like Sedona, AZ
- One resident went to Phoenix and was exposed to COVID-19 early in the pandemic
- How could that one infection affect the city over one year's time?
- Assume no other COVID-19
   brought in from elsewhere



Sedona, Arizona (from Wikipedia)



# What can we learn from this model?

### **Model configuration:**

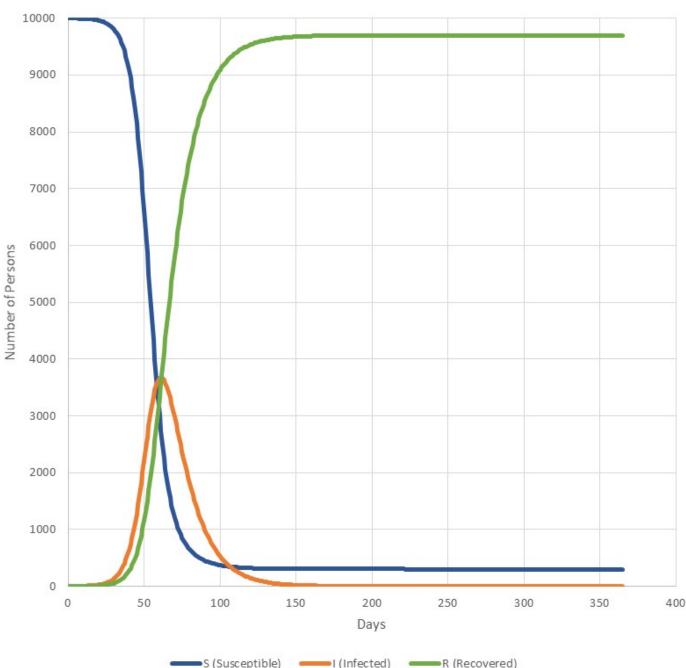
- Total population = 10,000
- Initial infections = 1 at time = 0 days
- Modeled time period = 1 year (365 days)

### **Parameters:**

- Infection rate r = 0.000025
- Recovery rate q = 0.07143

### **Insights:**

- Recovery lags behind infection
- Infection increases until the number of susceptible people decreases sufficiently, then drops
- Number of susceptible people doesn't go to zero • (not everyone gets infected) – herd immunity at ~97% infection



### COVID-19 Model

R (Recovered)



# What else can we learn from this model?

### **Change infection rate:**

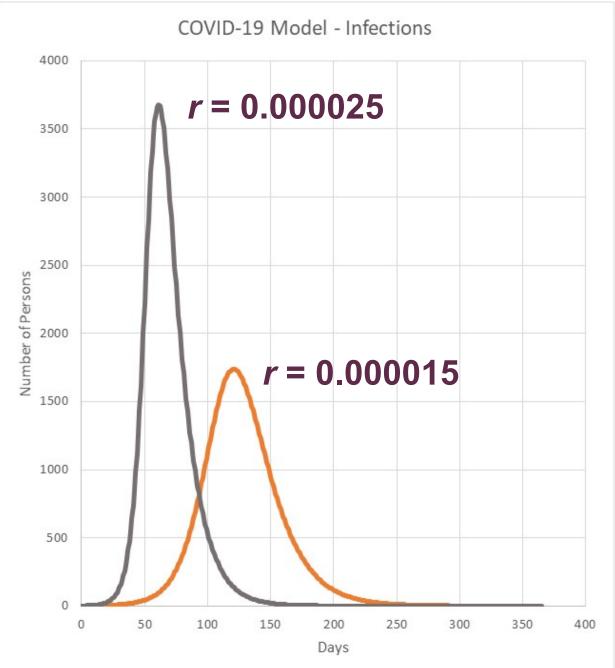
- Old infection rate r = 0.000025
- New infection rate r = 0.000015

### **Reflects less-contagious conditions:**

- Reduced interactions (social distancing)
- Reduced virus spread (mask wearing, personal hygiene)

### Insights:

- Flattened the curve peak is both lower and later
- Herd immunity is reached with less total infections (about 83%)





# What else can we learn from this model?

### **Change infection rate:**

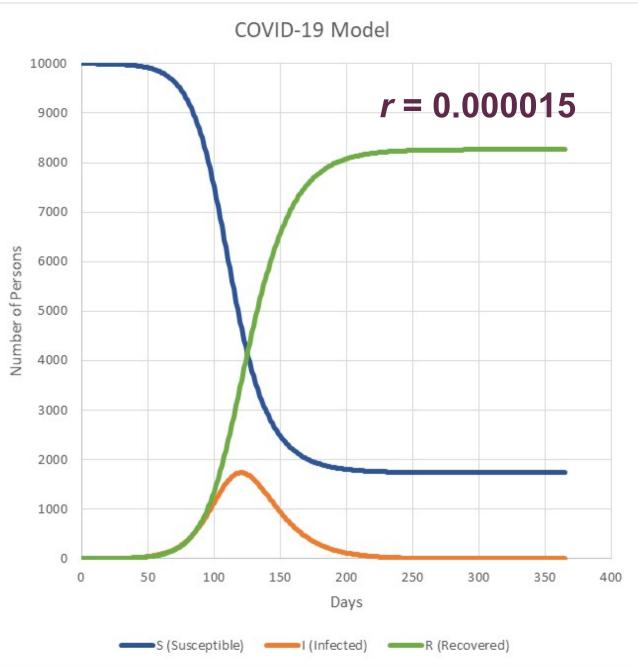
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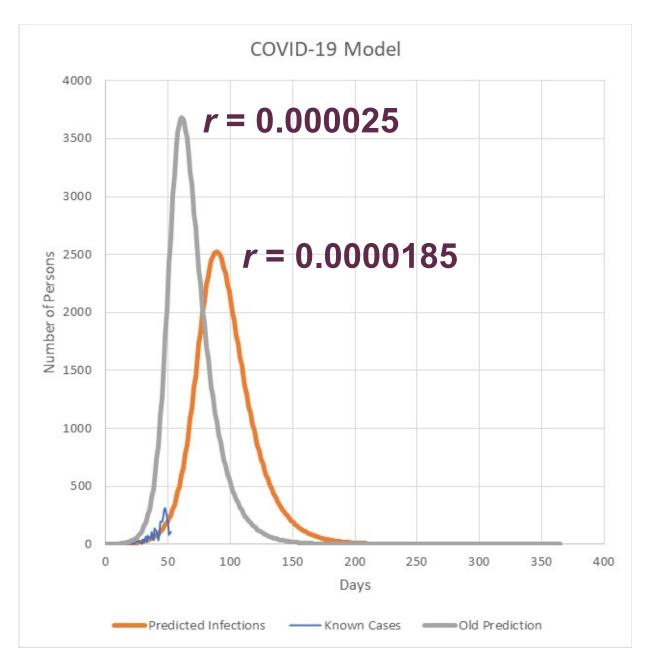
# **So...** what are the "correct" parameters?

### Initial estimates:

- Based on "prior knowledge" (from experience with other • diseases)
- If recovery takes 14 days on average, then q = 1/14 = 0.07143
- $R_{o}$  = the "basic reproduction number," or the expected • number of new infections from a single infection in a fully susceptible population. If this number is 3.5 in a population of 10,000, then r = 0.000025

### "Fitting" the model to early data:

- Once we start measuring the number of cases (testing), we can use those data to modify our model parameters so that the model better matches the observations
- Calibration or Parameter Estimation



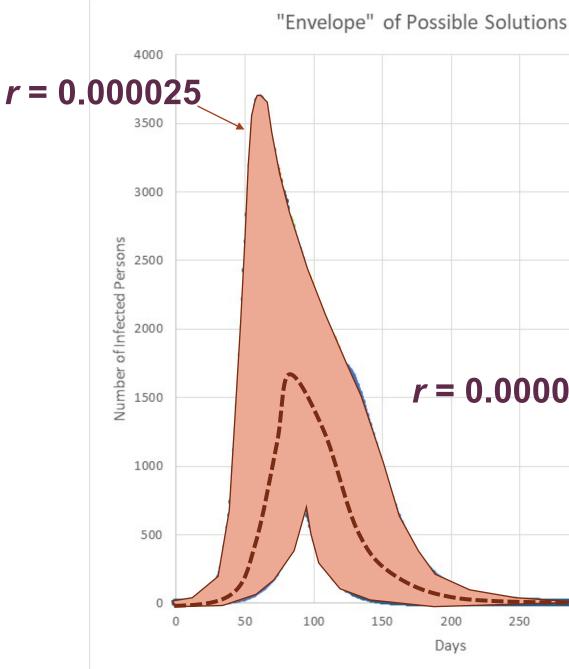




# So... what are the "correct" parameters?

### **Model uncertainty**

- Maybe we don't know the correct value, but we think we know likely minimum and maximum
- A range of unknown model parameters can be used to give an "envelope" of predictions



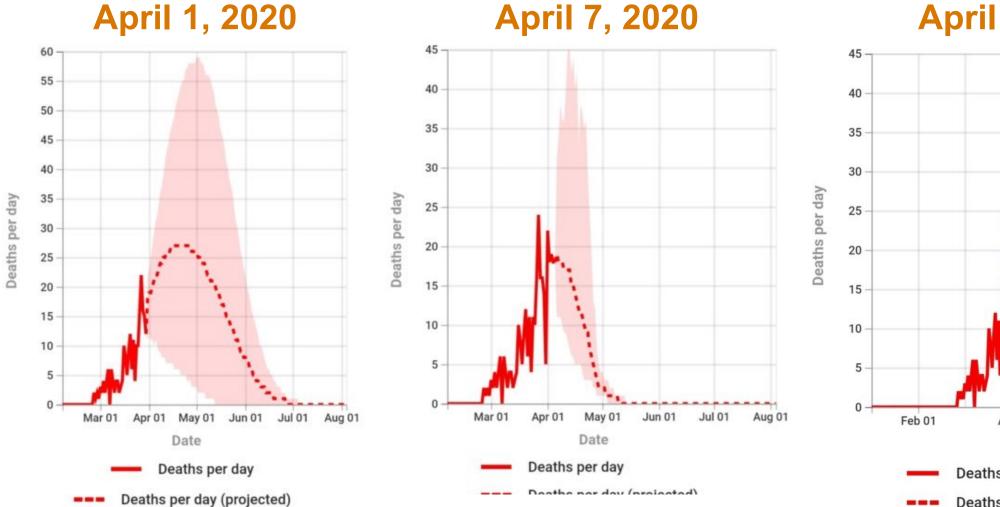


# *r* = 0.000015 250 300 350 400



## **Some real model predictions**







### Deaths per day (projected)



## **Some real model predictions**



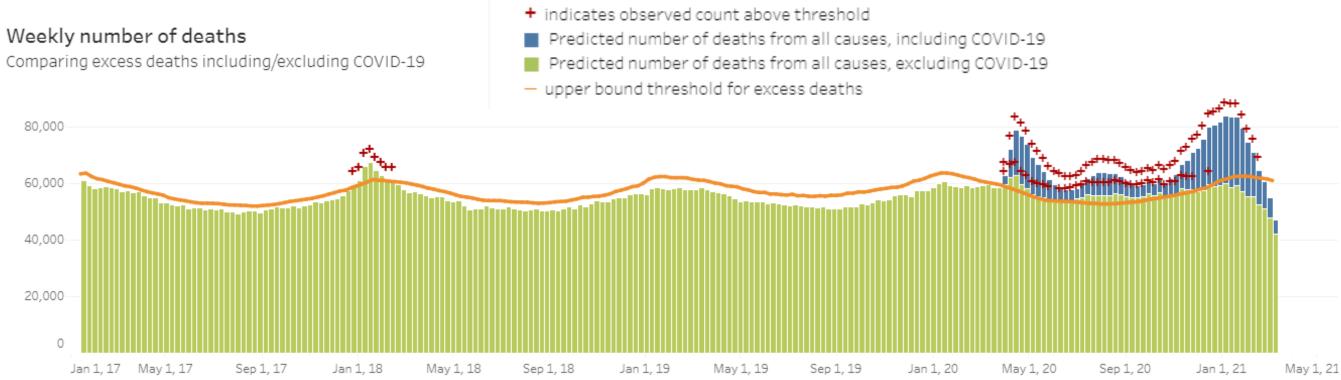
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# Statistical model – excess deaths

**Excess deaths:** The difference between the **observed** numbers of deaths in specific time periods and **expected** numbers of deaths in the same time periods (based on historical trends from 2013 to present)

Total excess deaths since 2/1/2020 across the United States: 529,900–647,117 (61,713–160,437 non-COVID excess deaths)



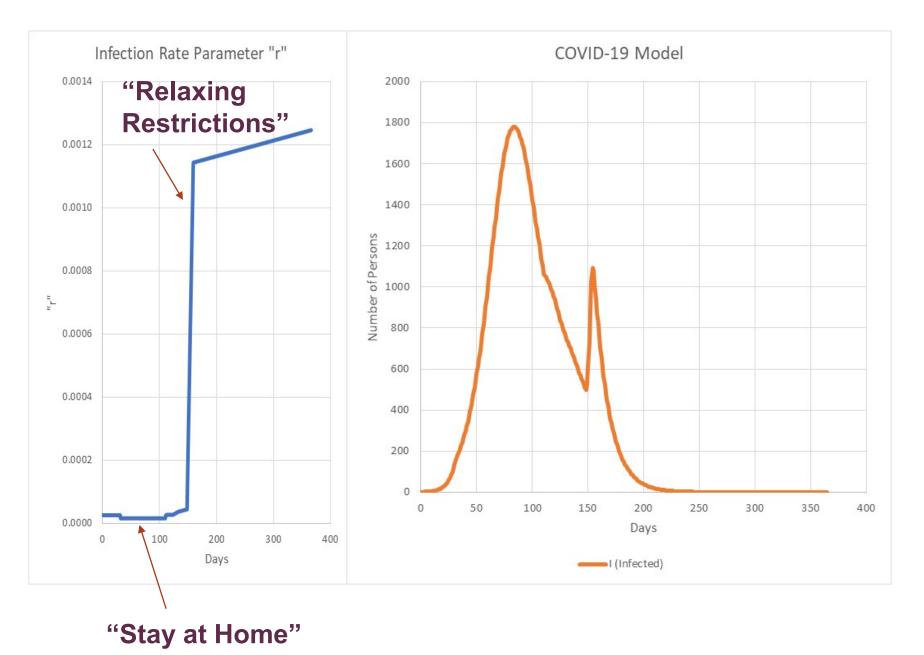
https://www.cdc.gov/nchs/nvss/vsrr/covid19/excess deaths.htm



# Let's explore our assumptions!

**Assumption:** The rate of infection (our parameter "r") is a single number

- It could vary over time because of public policies and behaviors, because of mutations in the virus, or because of environment (temperature, humidity)
- It could vary between subpopulations (large families, people in assisted living, health care workers, children in schools)

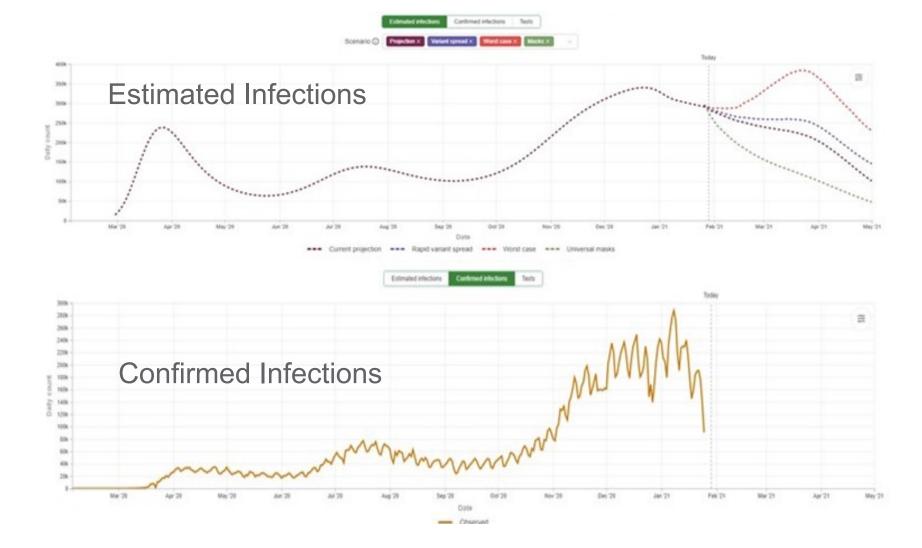




# Let's explore our assumptions!

# **Assumption:** Our testing data represent all infections

- People with mild symptoms may choose not to be tested
- Tests may not be available (especially early in the pandemic)
- Some people could be infected and never show symptoms (asymptomatic) – less likely to be tested

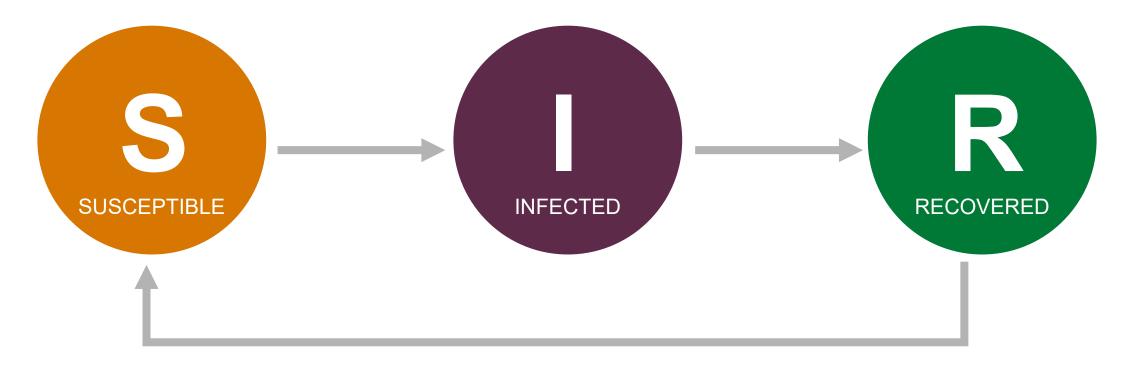




# Let's explore our assumptions!

### **Assumption: Recovered persons are immune**

- Immunity may be weak, allowing re-infection of a recovered person
- Immunity may only last for a period of time (Months? Years?)
- New mutations/strains may not be susceptible to antibodies





# **Can models predict** the future?

### That depends...

- How well are the underlying processes known? Are the assumptions valid? Are the important factors represented?
- How well are the parameters known? What supporting data/information do we have available? How uncertain are the data?
- Can the model reflect changes in future conditions?





### Hurricane Katrina



# **Can models predict the future?**

### A better question is: Can models help us learn and support policy decisions?

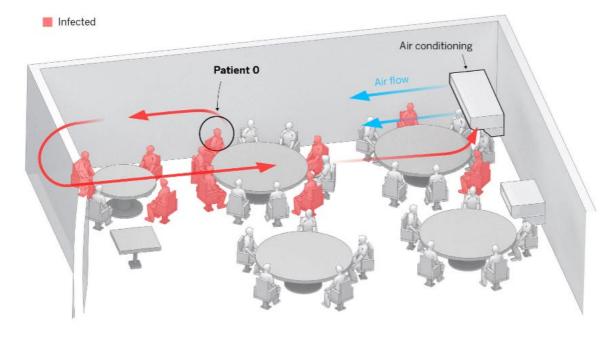
- Flattening the curve applying social distancing and masks to reduce "r"
  - Reduce total infections (and deaths)
  - Achieve herd immunity at lower level of infection
  - Reduce impacts on health care infrastructure
  - Limit infections and deaths while developing vaccines
- Vaccines: Reduce S without increasing I
- Asking the right questions of our models may reveal new data or information we need to collect (example: asymptomatic spread, possibility of reinfection, impact of new variants)



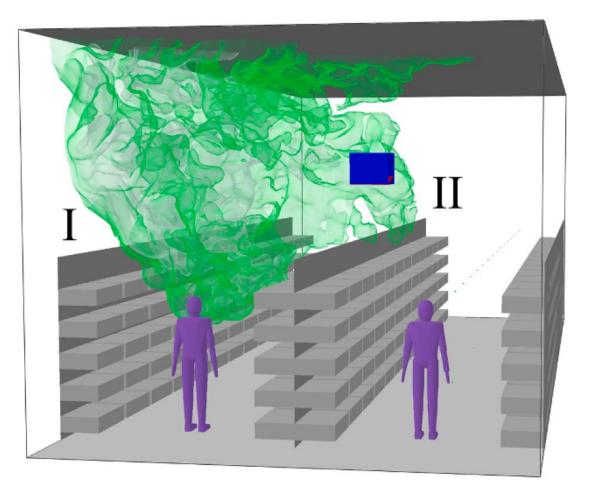


# Other models relevant to COVID-19

## Hydrodynamics and aerosol transport (computational fluid dynamics)



- Avoid internal air recirculation
- Use filters
- Increase external air flow
- Reduce background music
- Hold events outdoors



Vuorinen et al. (2020) Safety Science, doi:10.1016/j.ssci.2020.104866



# Summary: some modeling best practices

- 1. Understand your assumptions and how they affect the model outputs.
- 2. Test your model solutions and verify your code.
- 3. Realize that model predictions are inherently uncertain. Consider alternatives and interpret the results accordingly.
- 4. Understand that data may also have errors or assumptions.
- 5. Update the model and parameters as new information and data become available predictions *should* change over time.
- 6. Use the model to learn and guide additional research/data collection.
- 7. Work in teams to assure the best information and methods are used. Openness is essential!

### "All models are wrong, some are useful."

George Box Statistician



# **SUBMIT YOUR QUESTIONS VIA THE DISCUSSION CHAT**







**EVERY TUESDAY** IN MARCH 5:00-6:00 P.M.



# **THANK YOU FOR ATTENDING!**



# Malin Young

**Associate Laboratory Director Earth & Biological Sciences** 









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# Thank you



