

# Estimation and control of epidemics via Safe Blues

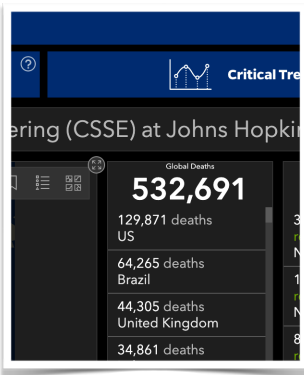
Yoni Nazarathy



Based on joint work with Raj Dandekar (MIT),  
Shane Henderson (Cornell), Marijn Jansen (UQ),  
Sarat Moka (UQ), Chris Rackauckas (MIT),  
Peter Taylor (Melbourne), and Aapeli Vuorinen (UQ, Melbourne).



**Presented as a Zoom Public Lecture at BrisScience, July 6, 2020**



**WARNING: COVID can break 30% of hearts**



$$\text{Deaths : } p = \frac{532,691}{11,367,743} \approx 4.5 \%$$

Number of people you love :  $n \approx 10$

Assumed proportion infected :  $F = 0.75$

Chance to lose a loved one :  $1 - (1 - F \times p)^n \approx 30 \%$

If  $p = 1 \%$  :  $1 - (1 - F \times p)^n \approx 7.3 \%$

If  $n = 20$  :  $1 - (1 - F \times p)^n \approx 50 \%$

# We are fighting back...



# But social distancing hurts...





So governments are doing their best\*  
to find the sweet spot...

\* With the exception of some governments across the Americas

# How?

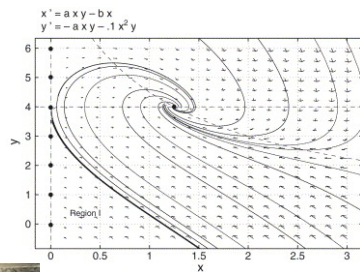
Clinical aspects...

Economic aid...

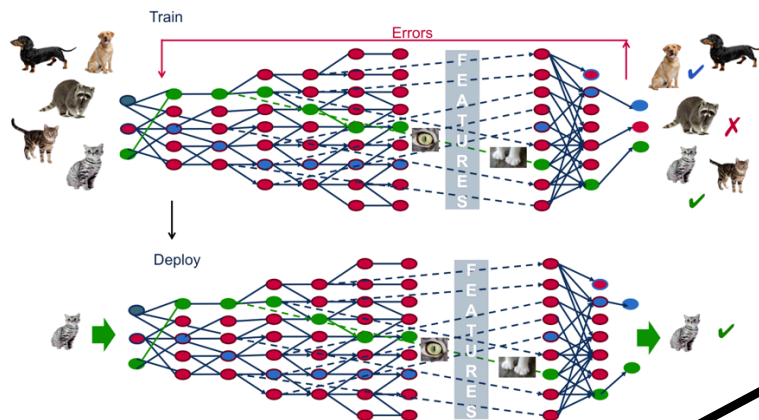
Modelling to  
inform policy...



1918-1920  
Spanish Flu



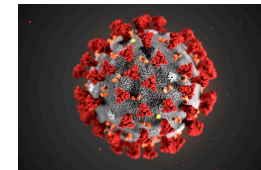
1928-1933  
SIR Model  
(Kermack–McKendrick theory)



1955 - 2015+  
Computational Statistics  
(ML/DS/AI)



2000+  
Smart Mobile Devices

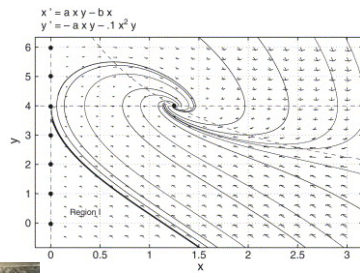


2019-2020+  
COVID-19 Pandemic

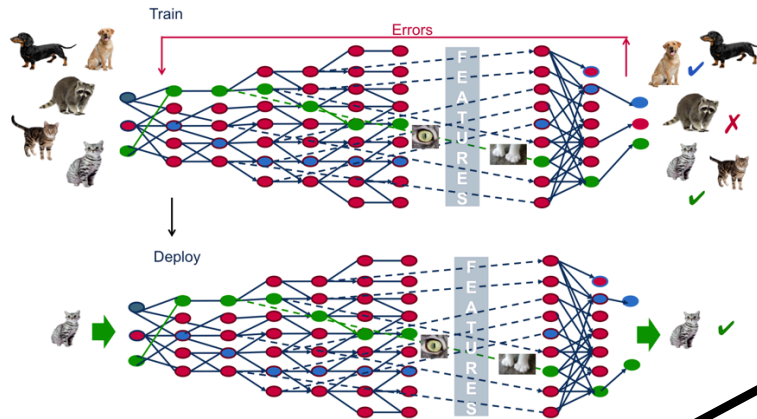
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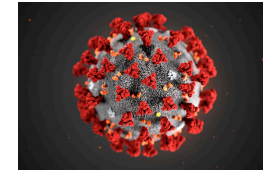
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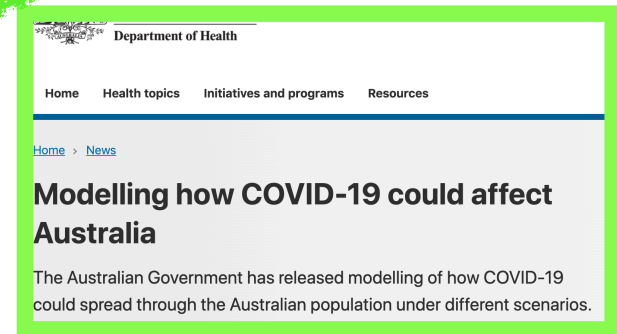
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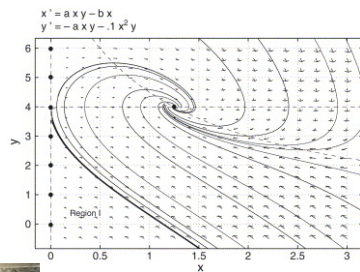


1918-1920

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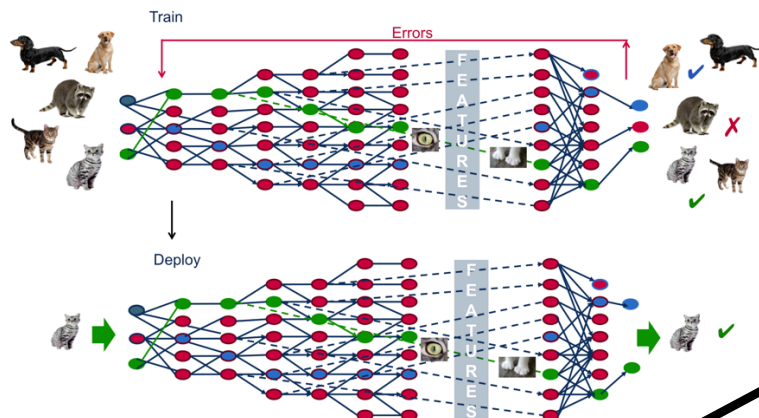
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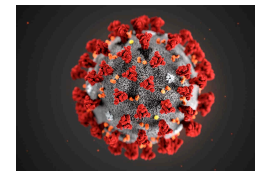
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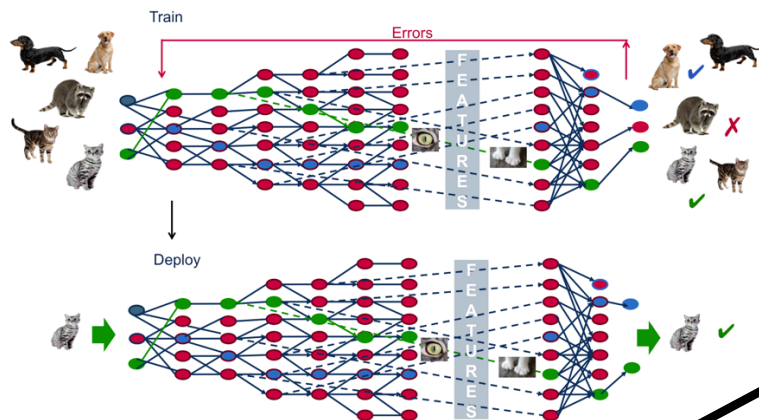
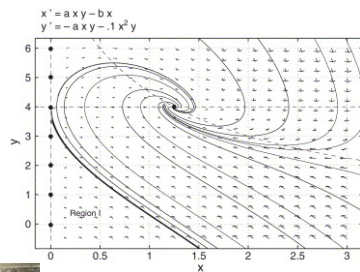


1918-1920

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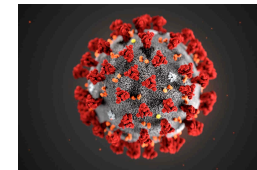
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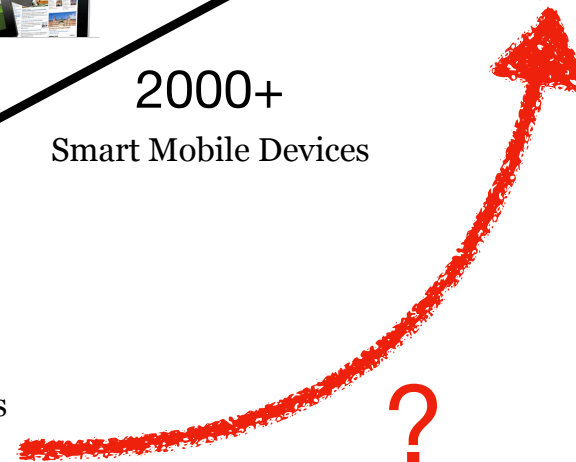
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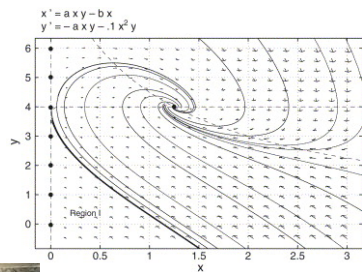


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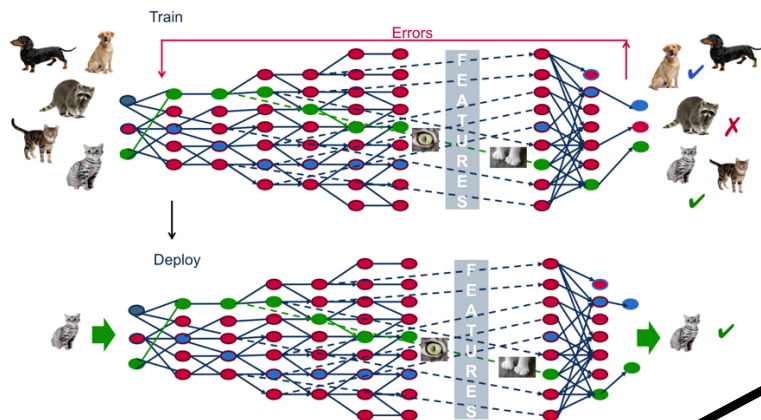




1918-1920  
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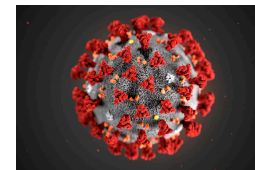
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2019-2020+  
COVID-19 Pandemic

 **Safe Blues**

Quantifying an epidemic with a single number,  
the reproductive number: **R**

**R** = The number of secondary infections induced by an infected individual

**R** < 1 : Exponential Decay

```
julia> 1000*0.7*0.7*0.7*0.7*0.7  
168.06999999999994
```

**R** > 1 : Exponential Growth

```
julia> 1000*1.4*1.4*1.4*1.4*1.4  
5378.239999999998
```

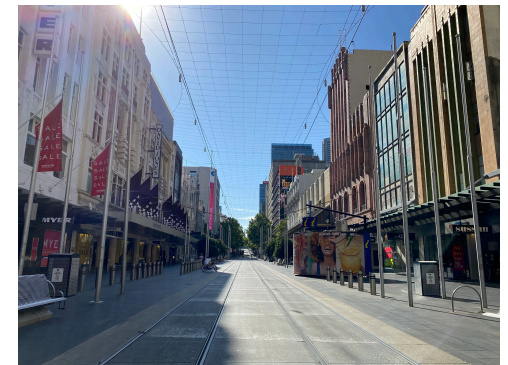
What is **R** for COVID?



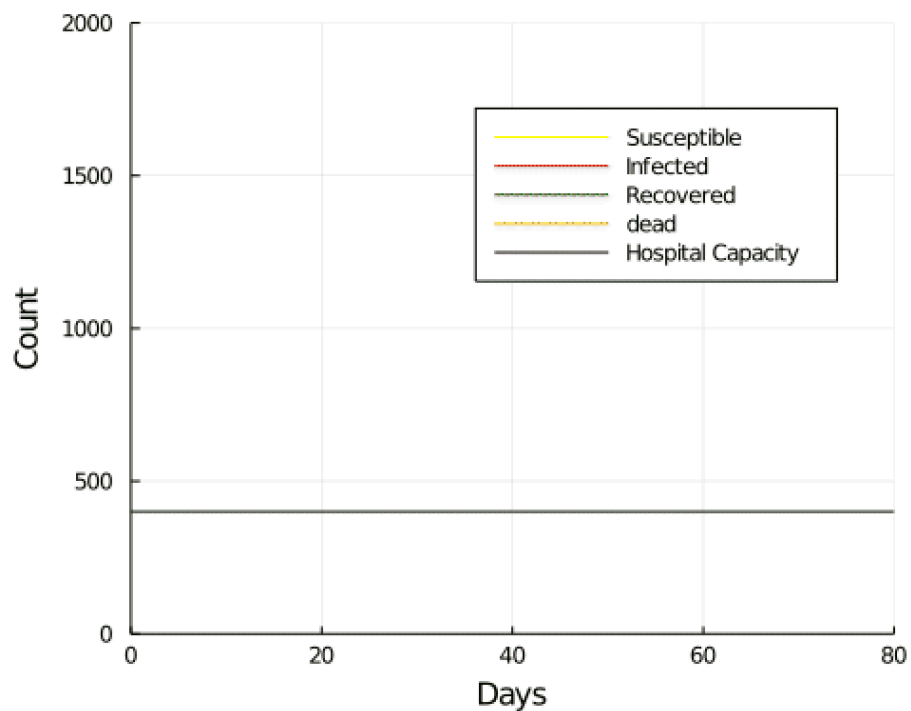
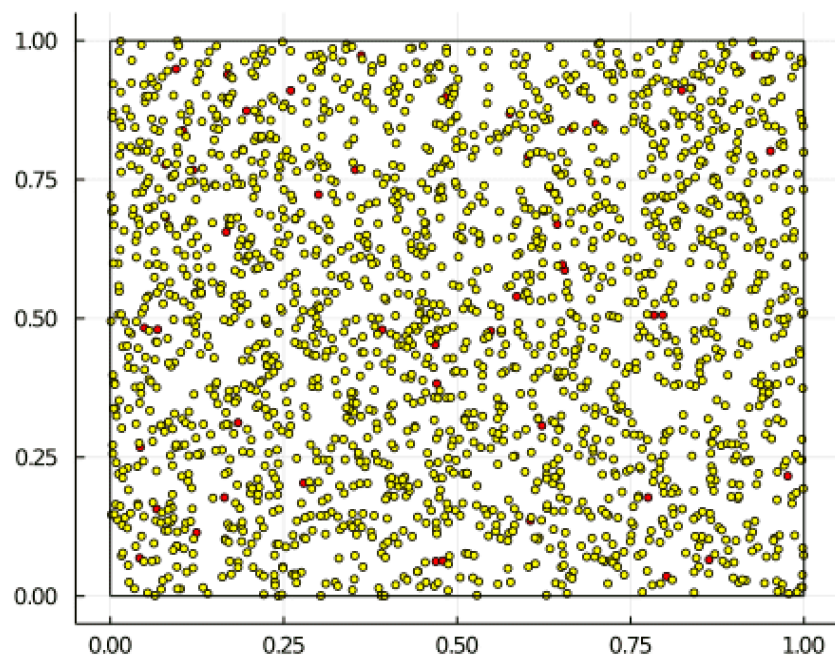
# What affects **R**?

biology  
environment  
conduct  
**contact**

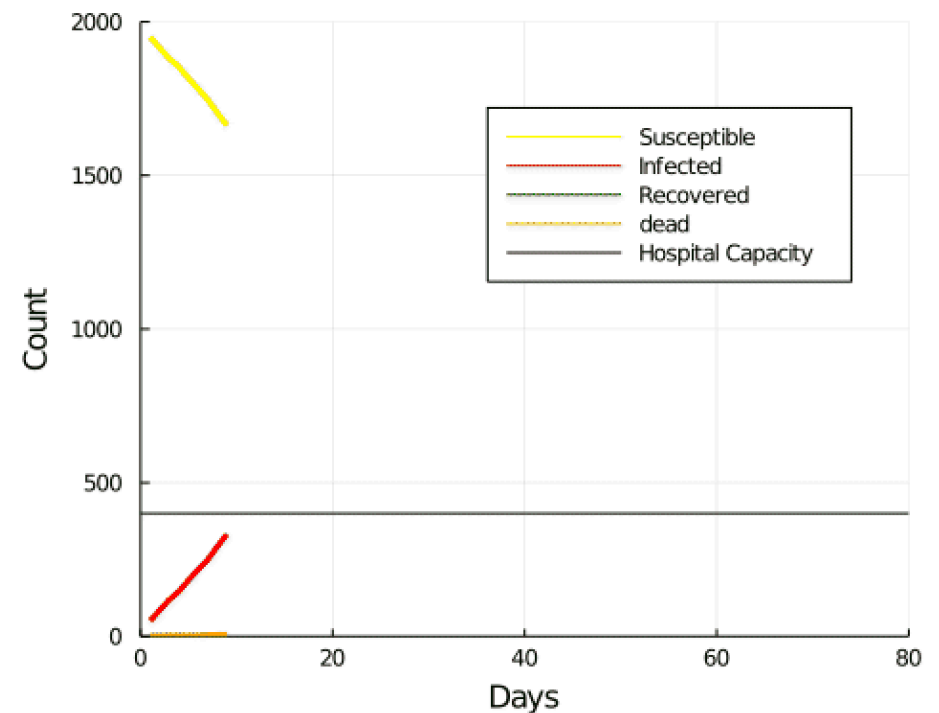
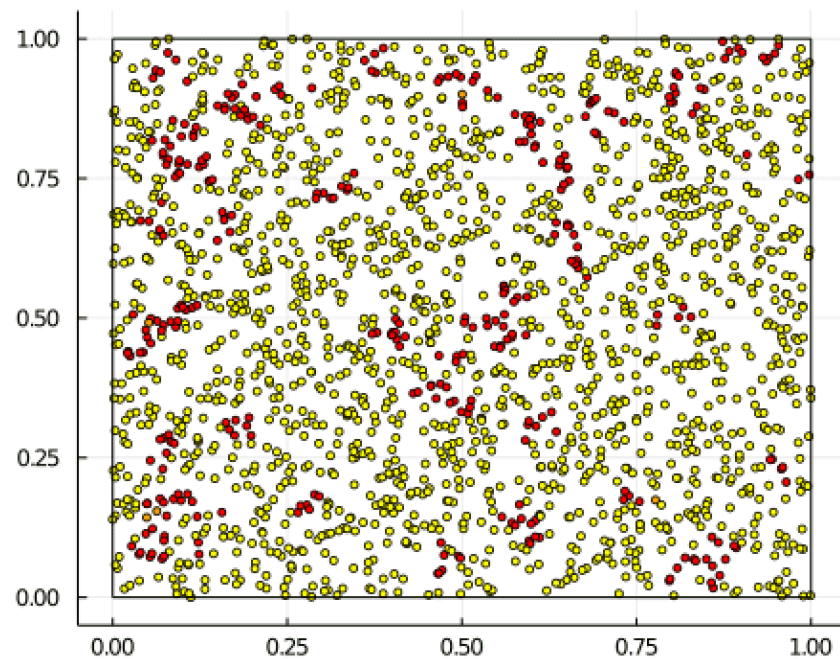
**Social  
distancing  
level**



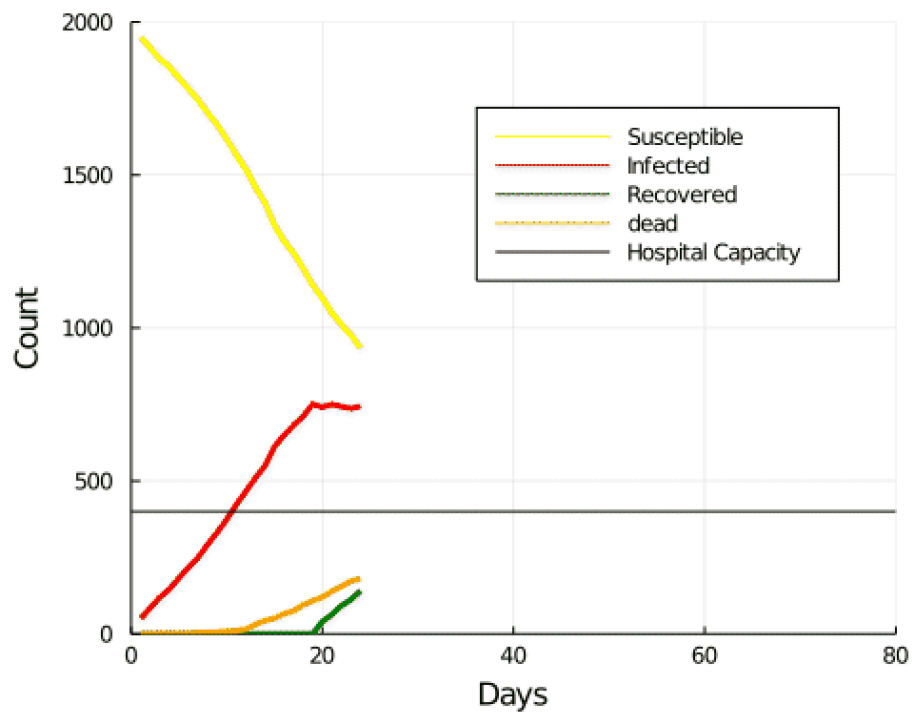
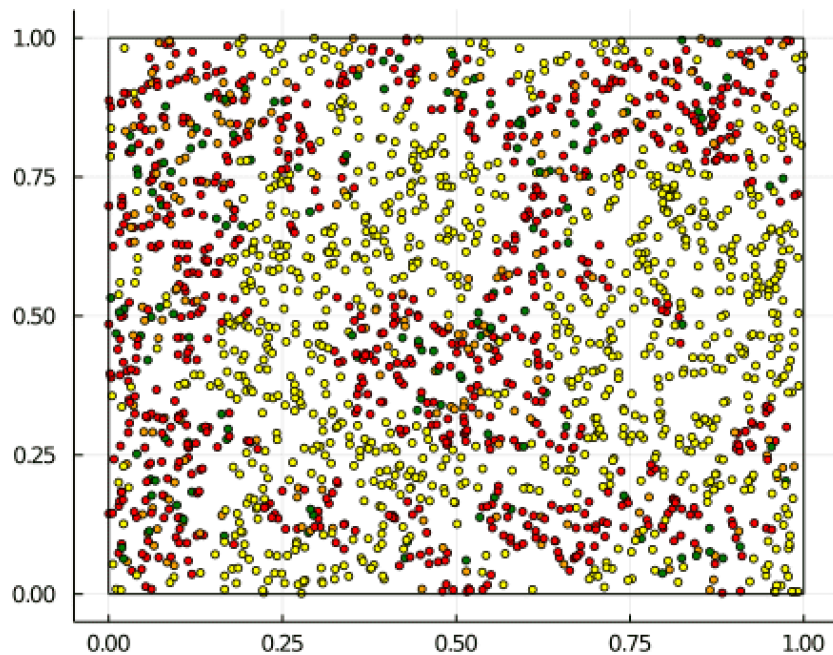
# Illustration: SIR Toy Model



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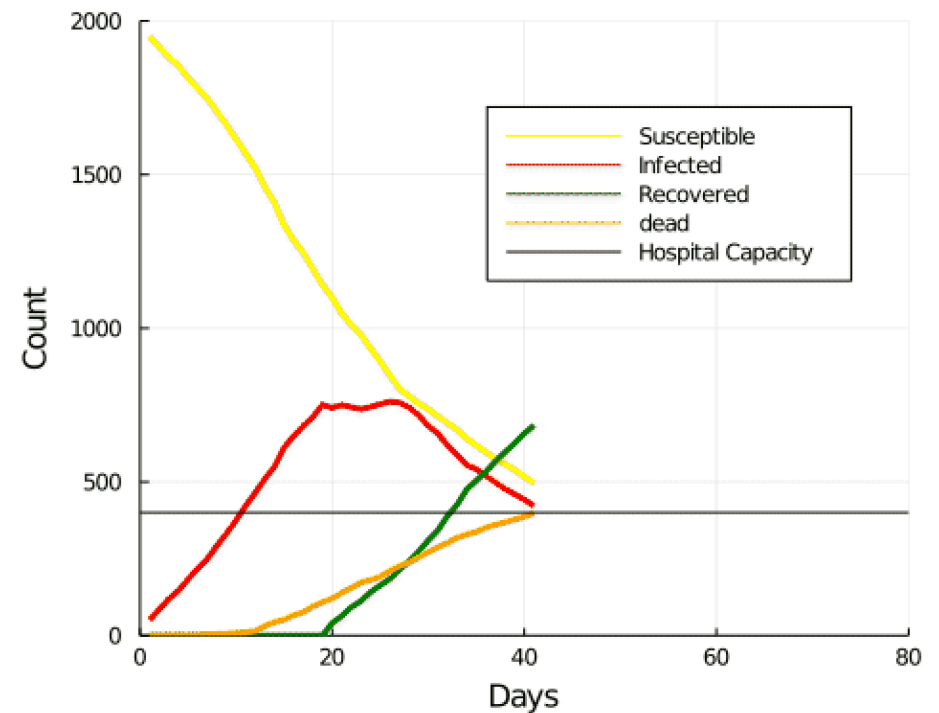
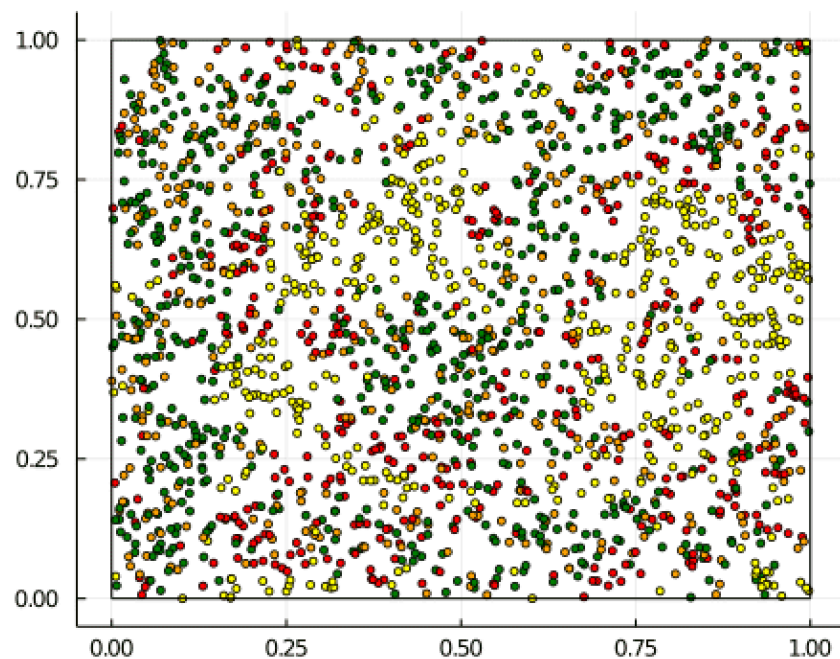


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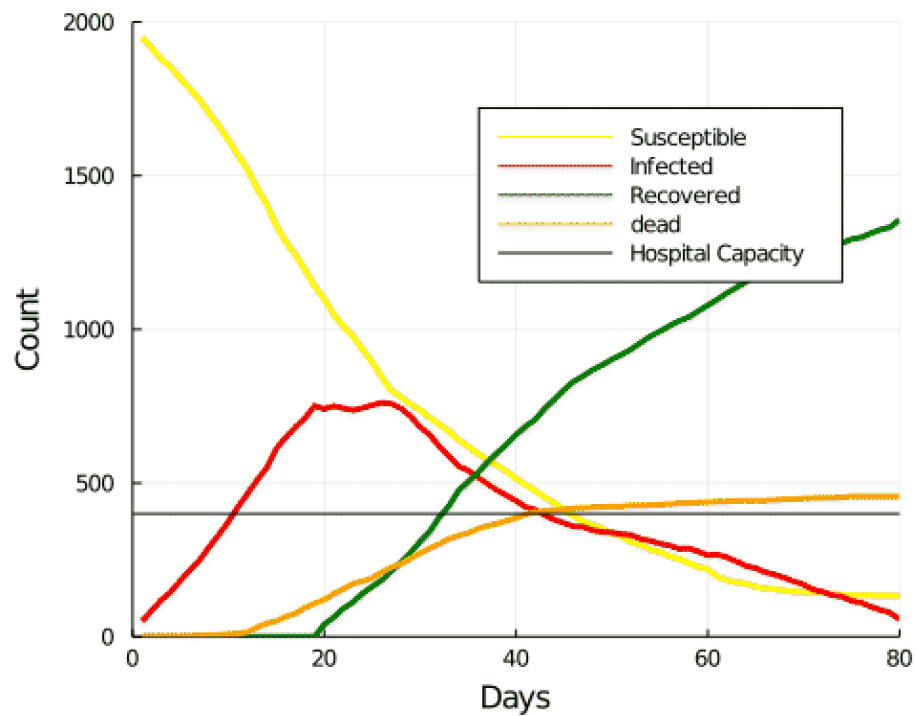
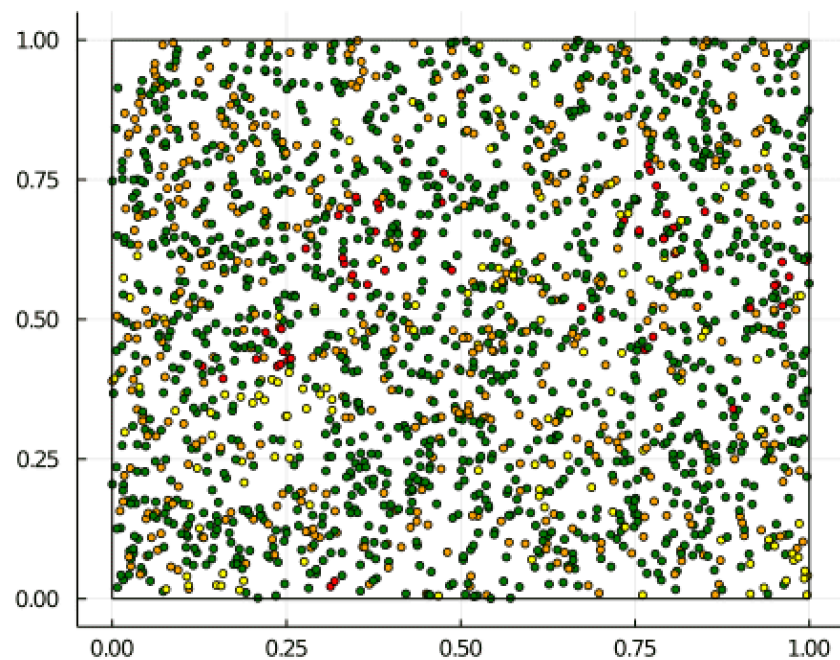




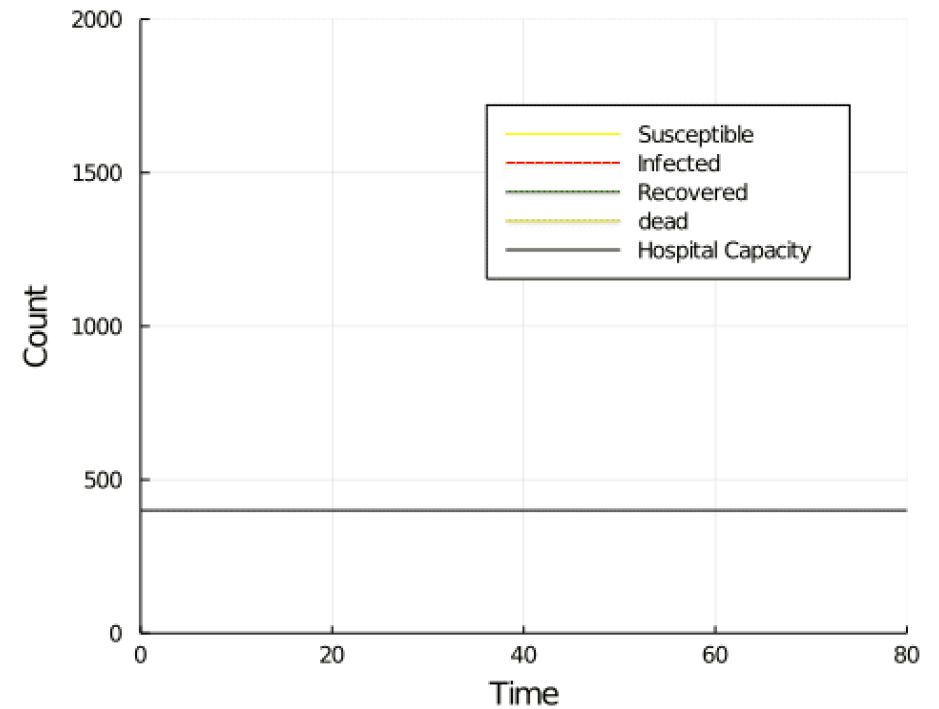
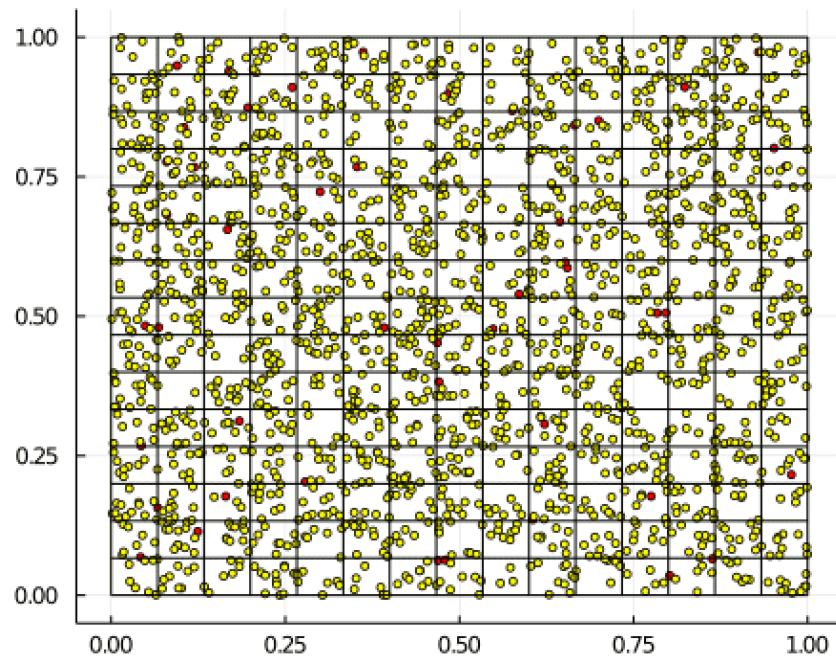
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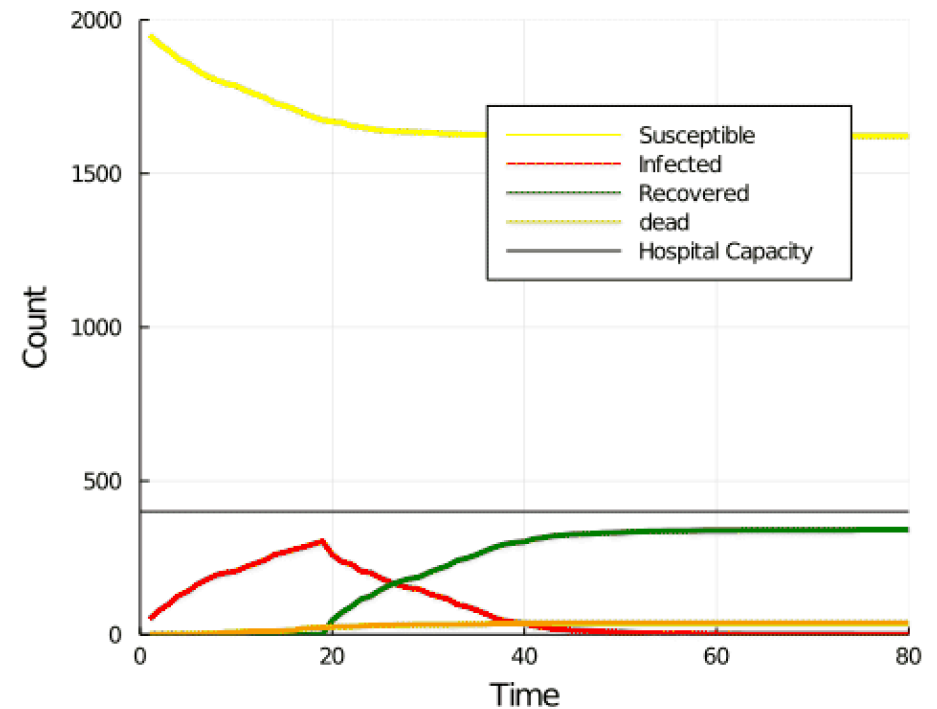
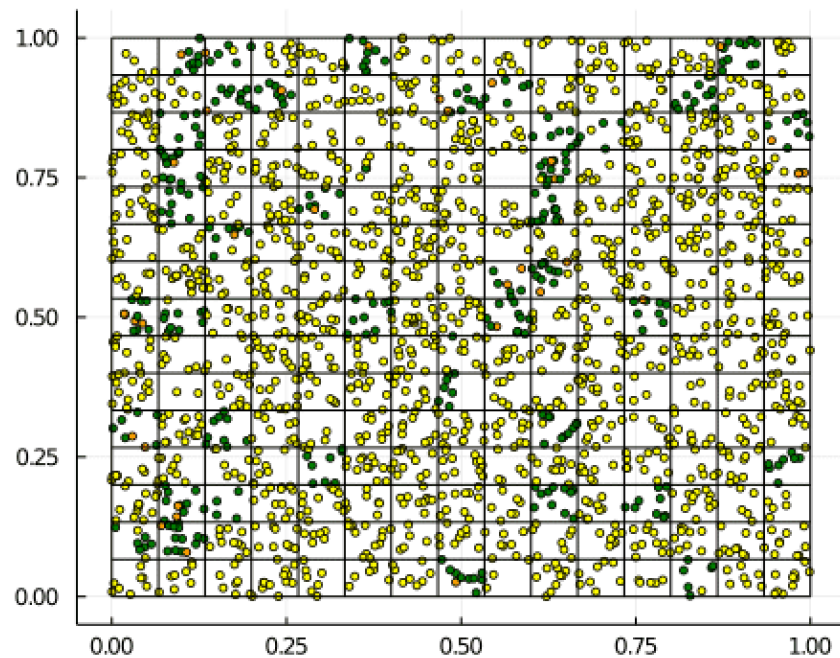
# Illustration: SIR Toy Model



# Illustration: SIR Toy Model with Social Distancing

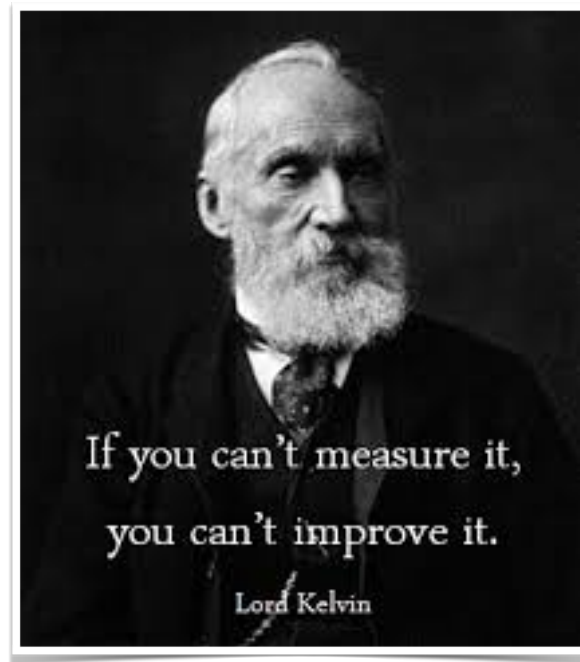


# Illustration: SIR Toy Model with Social Distancing





But how does social distancing work  
for non-toy models?



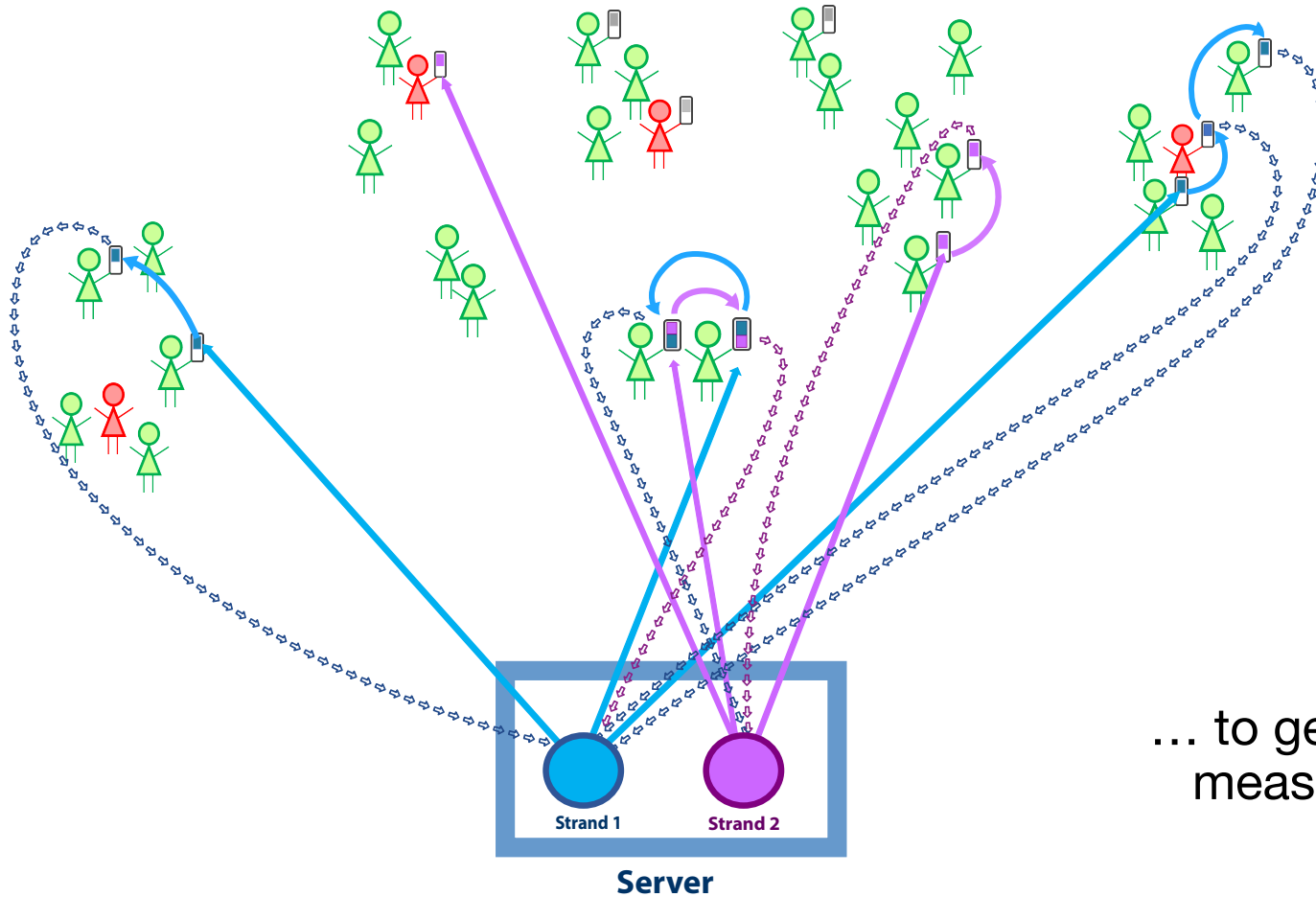
Big Problem: COVID measurements  
are noisy, partial, and **delayed!**



**Safe Blues: A Method for Estimation and Control in the Fight  
Against COVID-19**

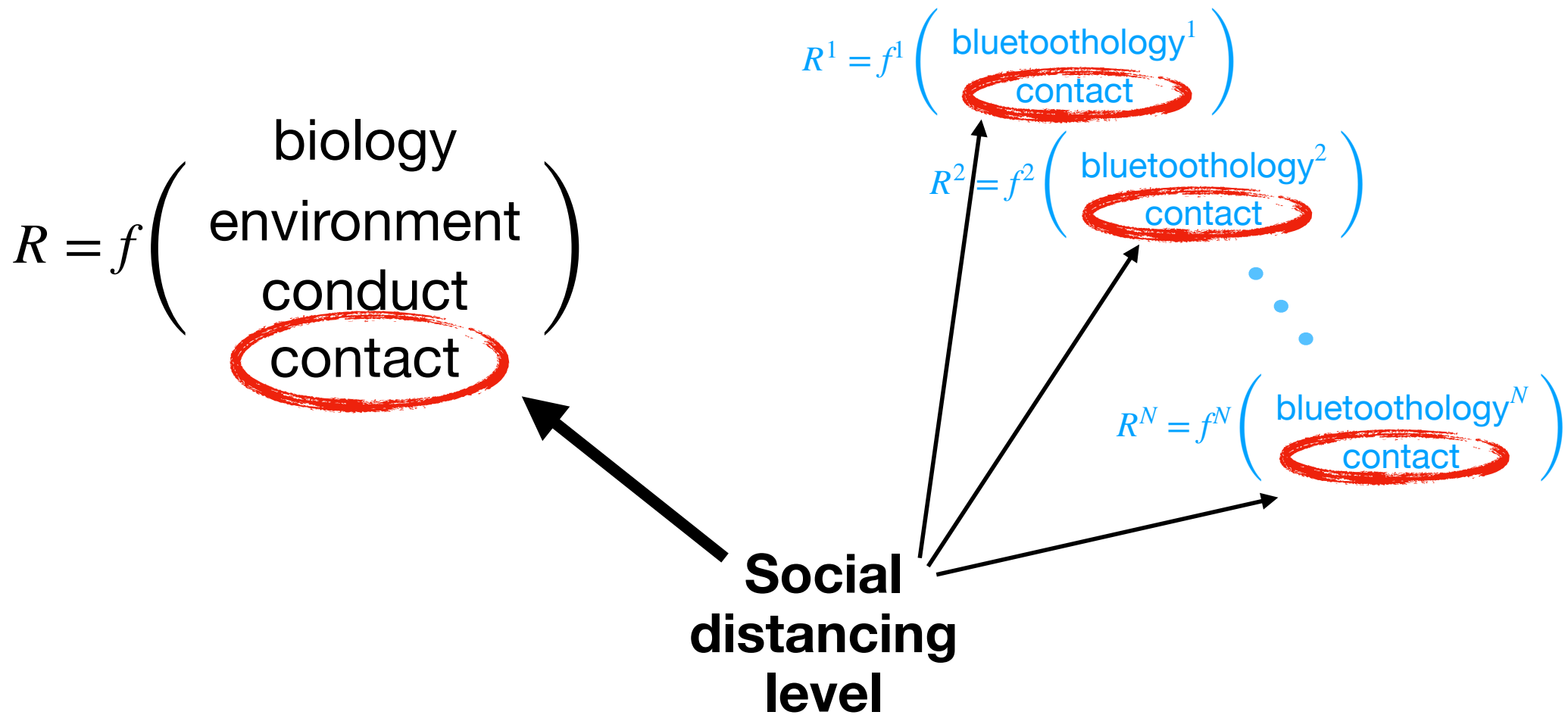
[safeblues.org](https://safeblues.org)

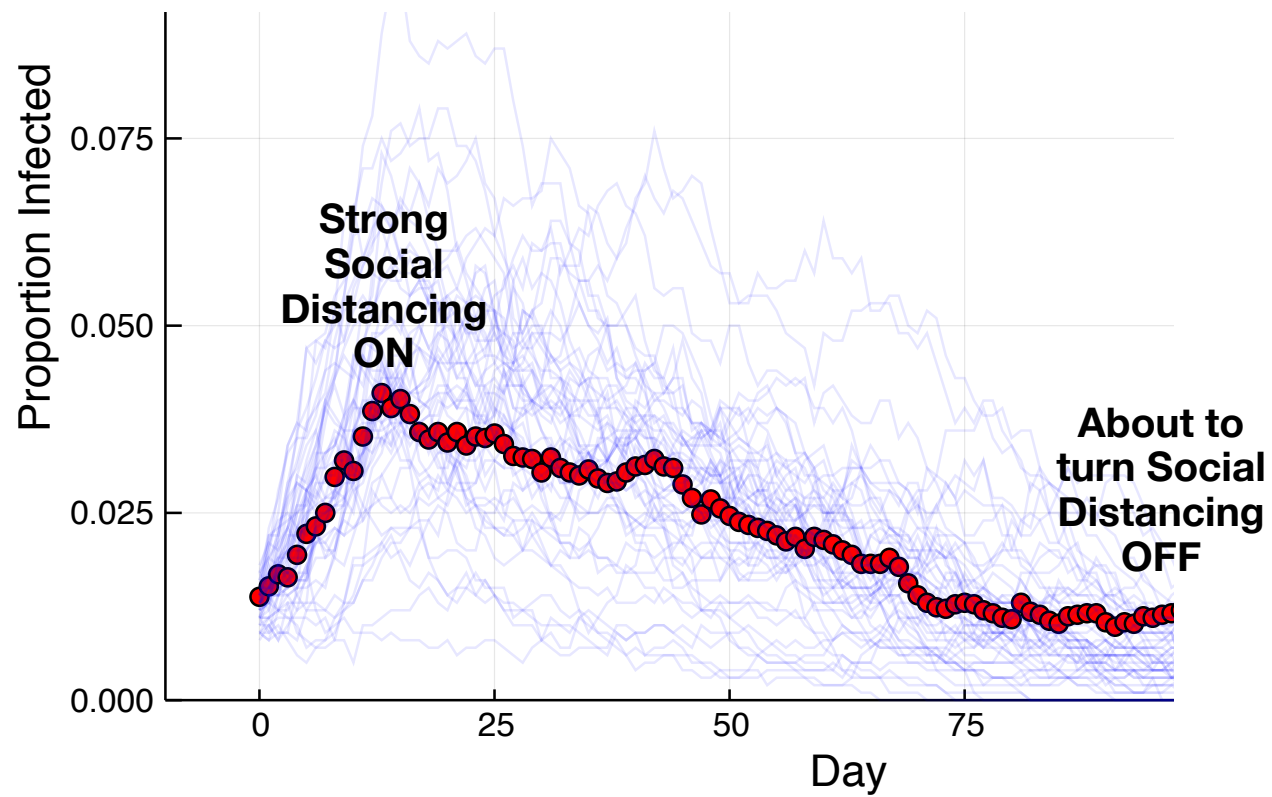
Safe Blues is about injecting many virtual safe viruses...



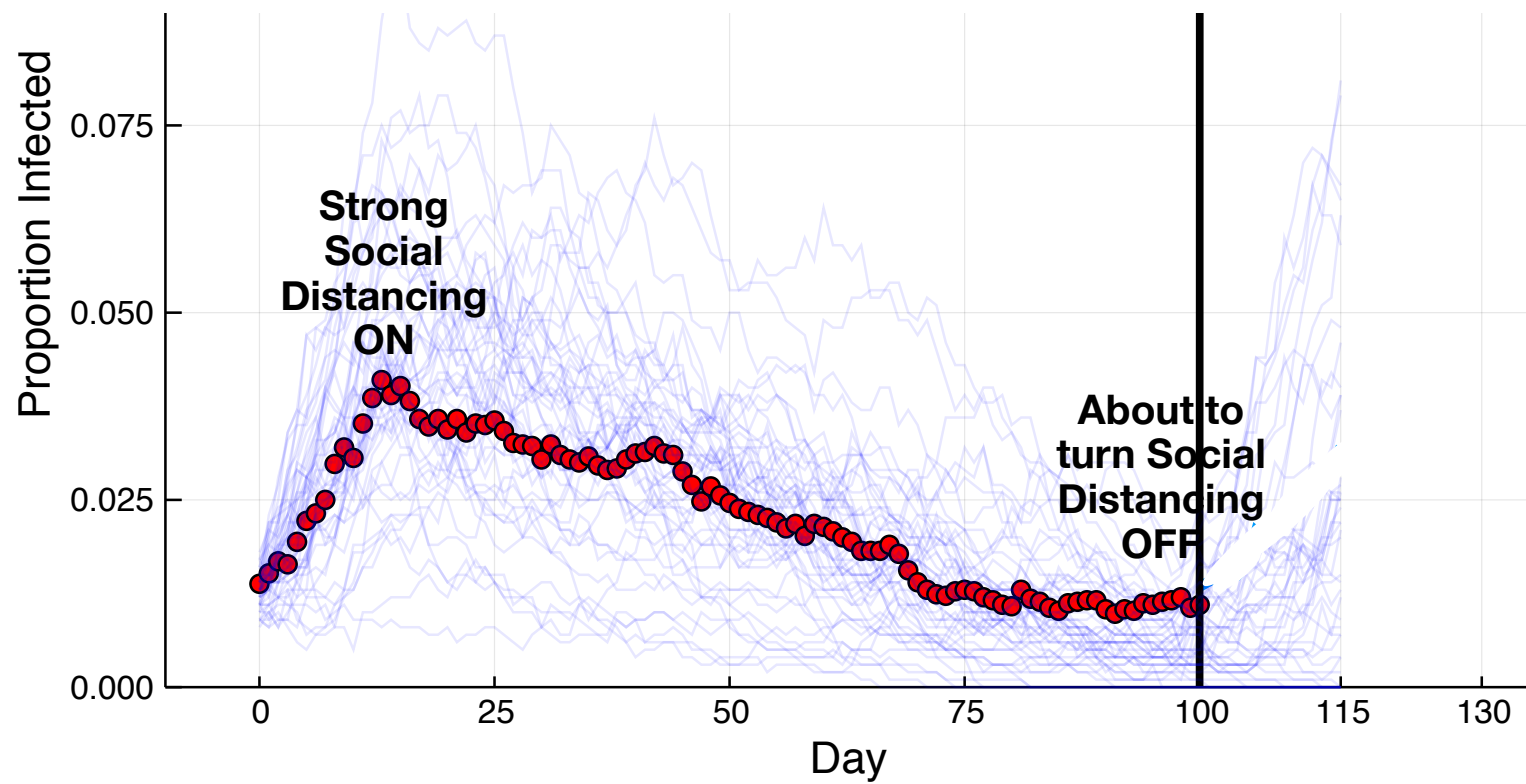
... to get **real time** measurements

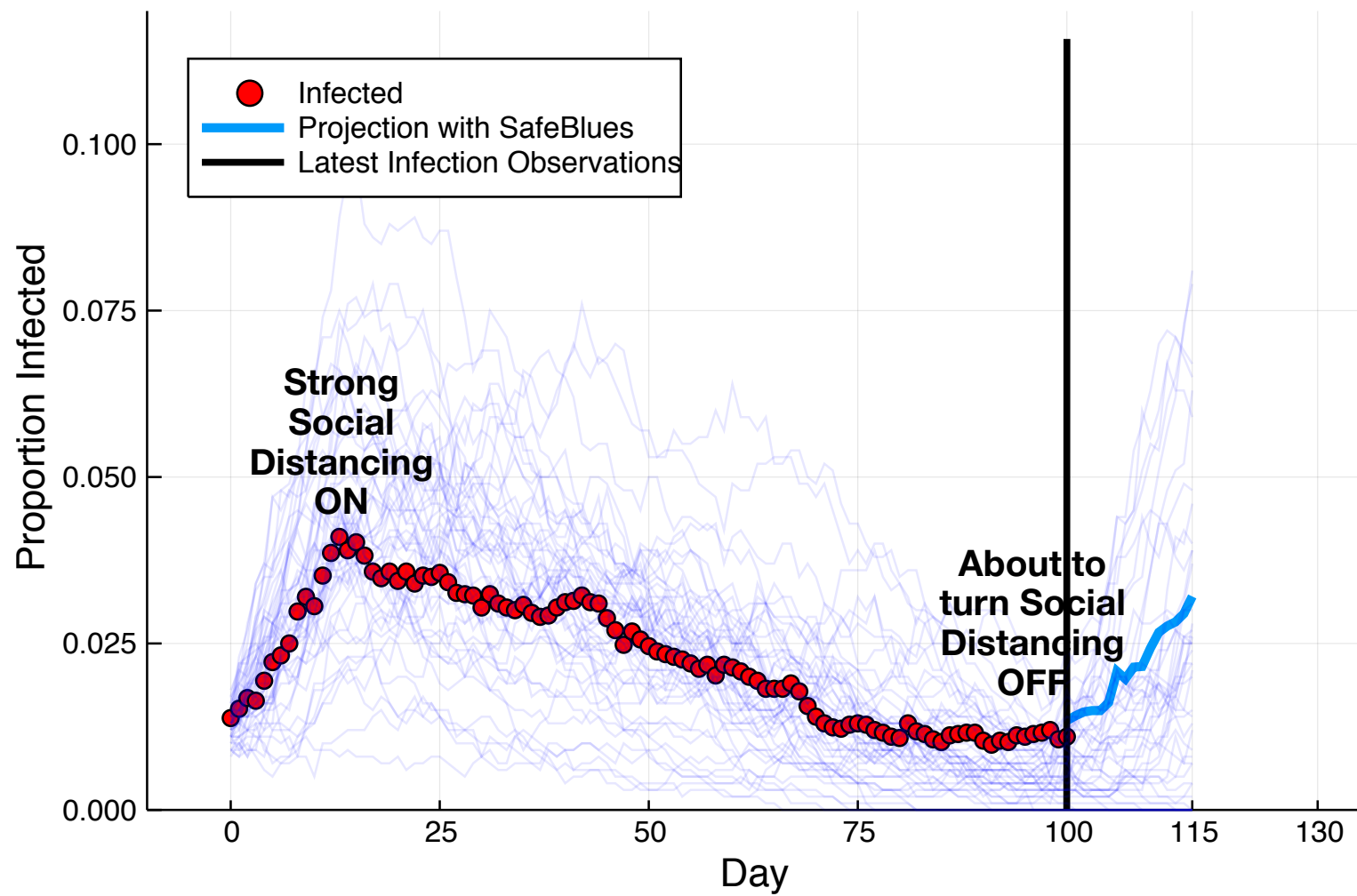
Safe Blues = virtual safe viruses

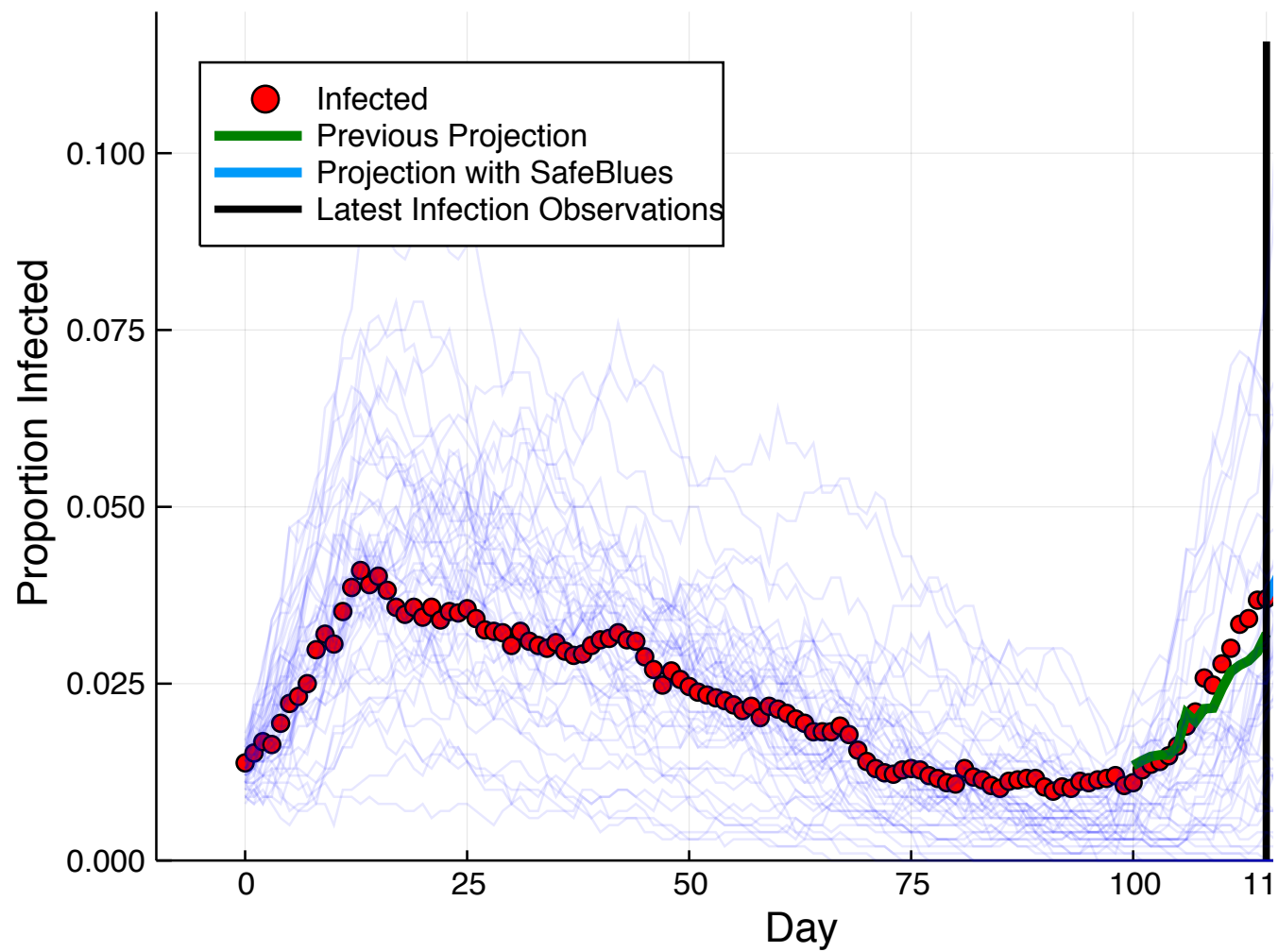


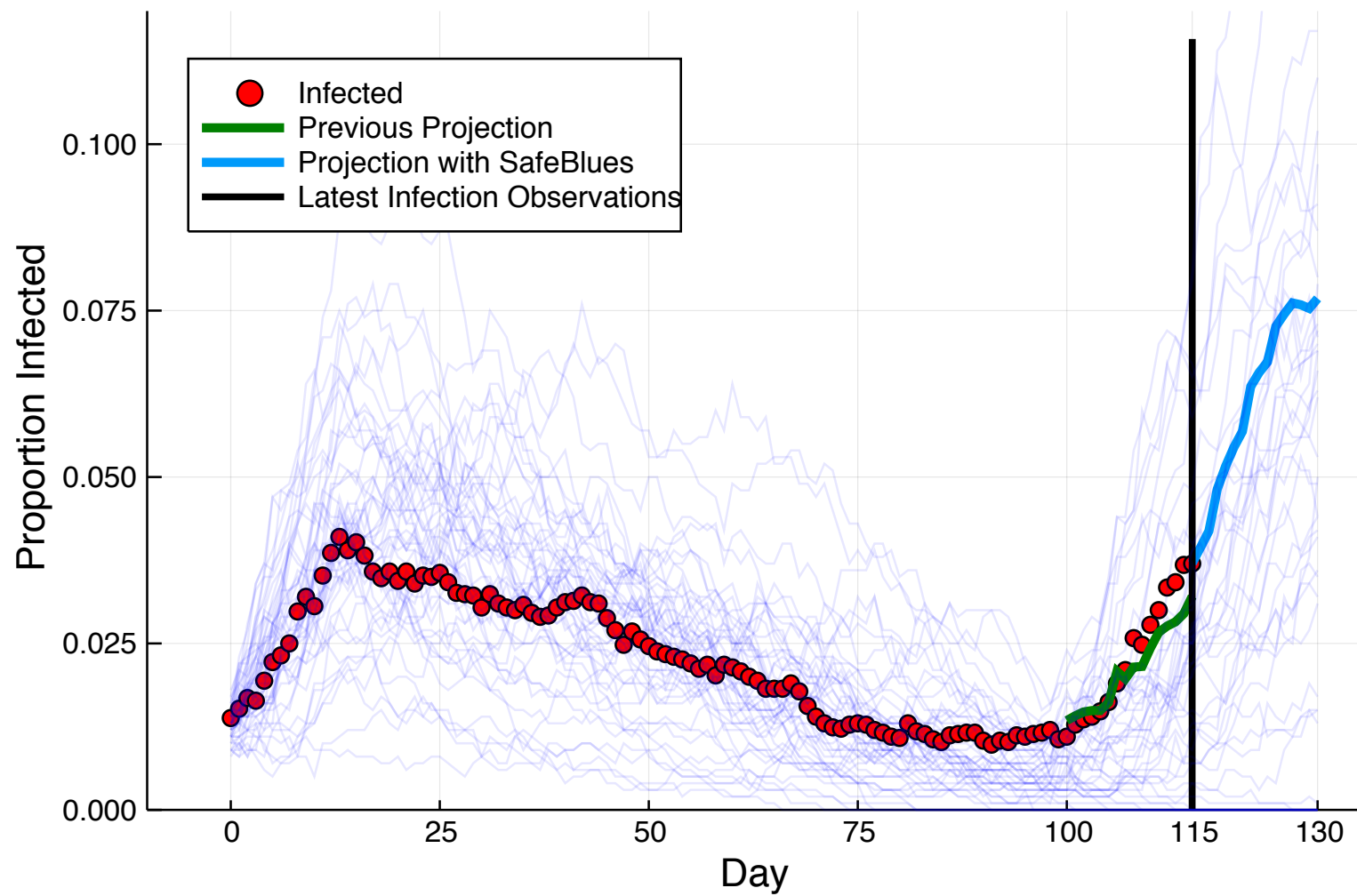




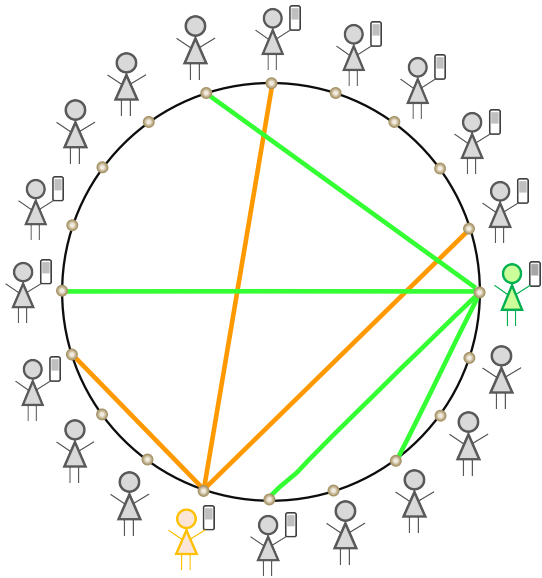




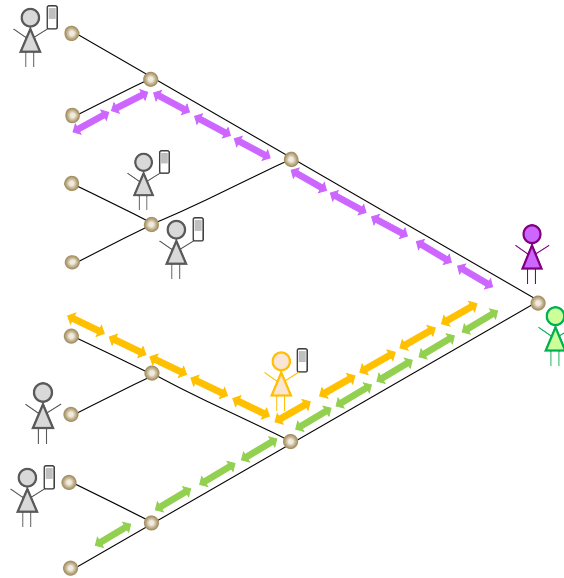




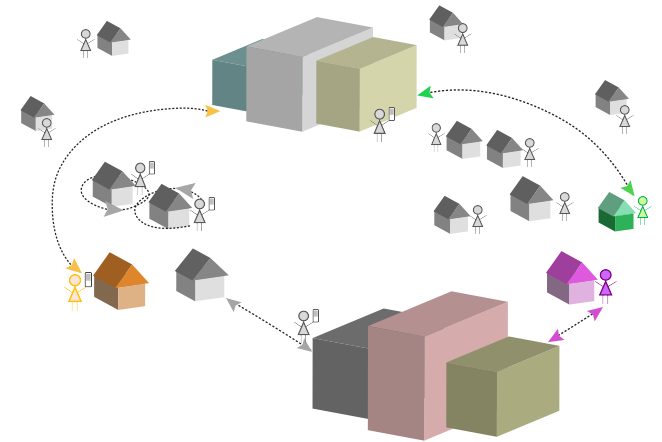
# How we tested Safe Blues...



## Model I

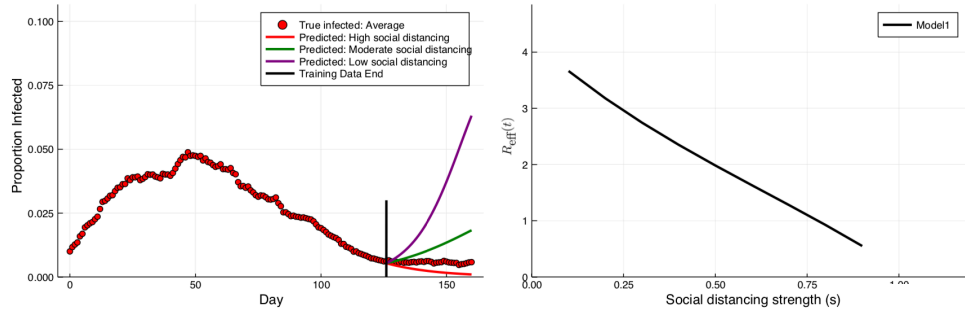


## Model II



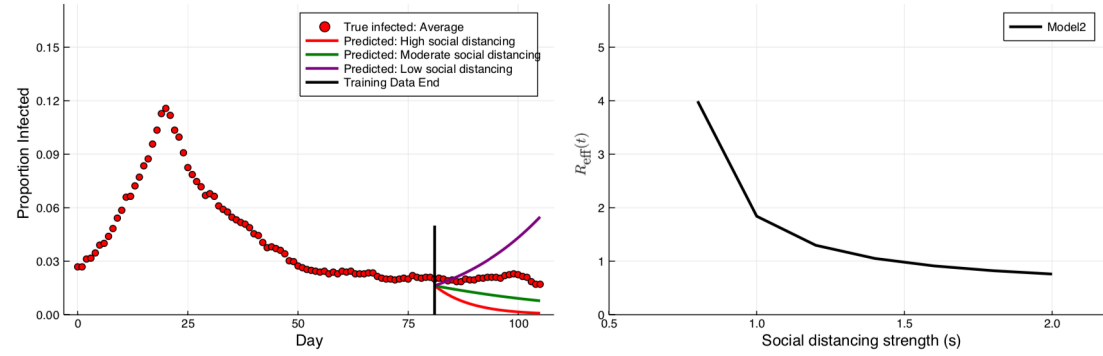
### Model III

# Understanding the effect of various social distancing regimes

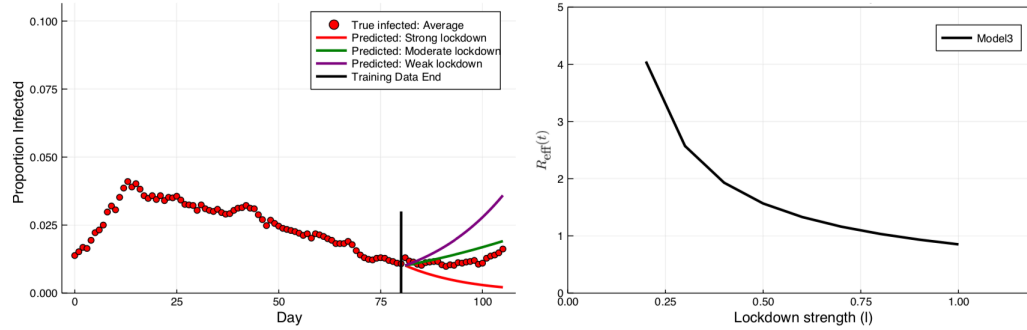


(a) Model I.

$$\begin{aligned} S' &= -C\beta(p)\delta SI, & \tilde{S}' &= -C\beta(p)\tilde{S}\tilde{I}, \\ I' &= C\beta(p)\delta SI - \gamma(p)\delta_\gamma I, & \tilde{I}' &= C\beta(p)\tilde{S}\tilde{I} - \gamma(p)\tilde{I}, \\ R' &= \gamma(p)\delta_\gamma I, & \tilde{R}' &= \gamma(p)\tilde{I}. \end{aligned}$$



(b) Model II.



(c) Model III.



# Safe Blues is not a contact tracing app

## ... but can be added to such an app

Currently working towards an Android App  
and an experiment based on “Trace Together”...



# Thank you

If you think Safe Blues is a good idea,  
then please let your government know about it.



[safeblues.org](https://safeblues.org)



@ynazarathy