

Dynamics of COVID-19: Near- and Long-Term Challenges

Joshua S. Weitz

School of Biological Sciences and School of Physics
Center for Microbial Dynamics and Infection
Quantitative Biosciences Program
Georgia Institute of Technology
Atlanta GA, USA

Technical References:

Shield Immunity: Weitz et al., medrxiv 10.1101/2020.04.01.20049767v1)

Asymptomatic transmission: Park, Cornforth, Dushoff & Weitz, *Epidemics*, in press (& medrxiv 2020.03.09.20033514v1)

R0 estimation: Park et al. (in review & available on medrxiv: 2020.01.30.20019877v3)

Generation intervals: Park et al., *Epidemics* 27:12 (2019)

Epidemics and behavior change: Eksin, Paarporn and Weitz *Epidemics* 27: 96-105 (2019)

Speed-strength and identifiability problems: Weitz & Dushoff, *Scientific Reports* 5: 8751 (2015)

Email: jsweitz@gatech.edu

Web: <http://ecothery.biology.gatech.edu>



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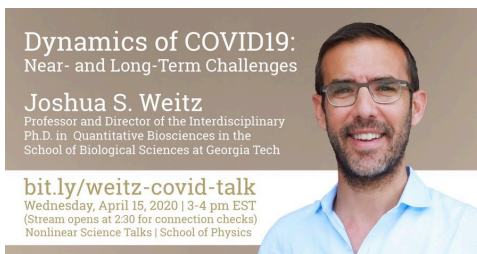
This work is the result of ongoing interdisciplinary collaborations:

Georgia Tech Team

Ashley Coenen
Dr. Stephen Beckett
Dr. David Demory
Marian Dominguez-Mirazo
Dr. Joey Leung
Guanlin Li
Andreea Magalie
Daniel Muratore
Rogelio Rodriguez-Gonzalez
Dr. Adriana Sanz
Shashwat Shivam
Conan Zhao

Collaborators

Clio Andris, GT
Daniel Cornforth, GT
Jonathan Dushoff, McMaster
Ceyhun Eksin, Texas A&M
Benjamin Lopman, Emory
Alicia Kraal, Emory
Kristen Nelson, Emory
Sang Woo Park, Princeton
Yorai Wardi, GT



As advertised



Current status

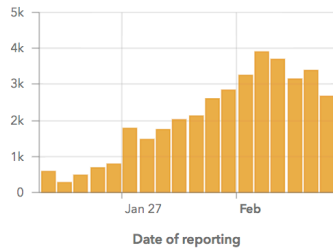
Novel coronavirus (2019-nCoV) situation as of 09 February 2020, 16:00 (CET)

37,499
confirmed cases

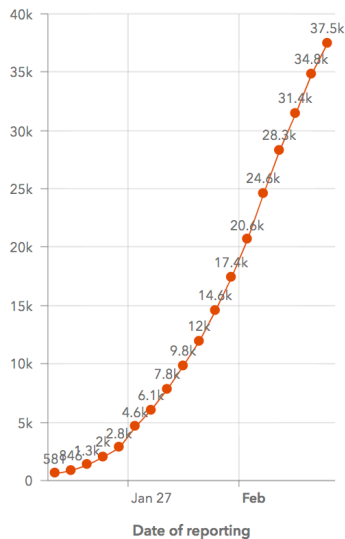
813
deaths

Cases from
25
countries

Cases by date of report

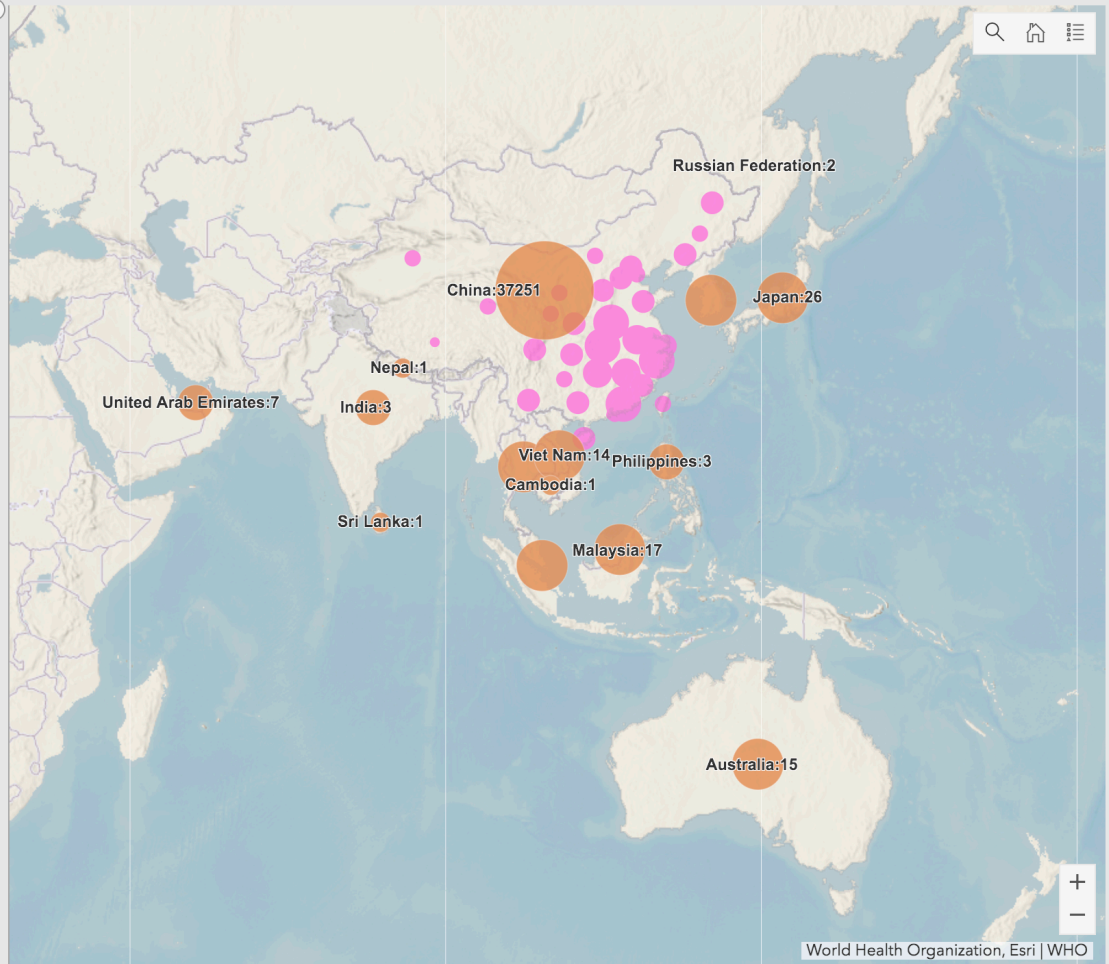


Cumulative confirmed cases



Countries with confirmed cases

<u>China</u> : 37251 Cases
<u>Singapore</u> : 43 Cases
<u>Thailand</u> : 32 Cases
<u>Republic of Korea</u> : 27 Cases
<u>Japan</u> : 26 Cases
<u>Malaysia</u> : 17 Cases
<u>Australia</u> : 15 Cases
<u>Germany</u> : 14 Cases
<u>Viet Nam</u> : 14 Cases
<u>United States of America</u> : 12 Cases
<u>France</u> : 11 Cases
<u>Canada</u> : 7 Cases
<u>United Arab Emirates</u> : 7 Cases
<u>United Kingdom</u> : 4 Cases
<u>India</u> : 3 Cases
<u>Italy</u> : 3 Cases
<u>Philippines</u> : 3 Cases
<u>Russian Federation</u> : 2 Cases
<u>Spain</u> : 2 Cases
<u>Cambodia</u> : 1 Cases
<u>Finland</u> : 1 Cases
<u>Nepal</u> : 1 Cases
<u>Sri Lanka</u> : 1 Cases
<u>Sweden</u> : 1 Cases
<u>Belgium</u> : 1 Cases

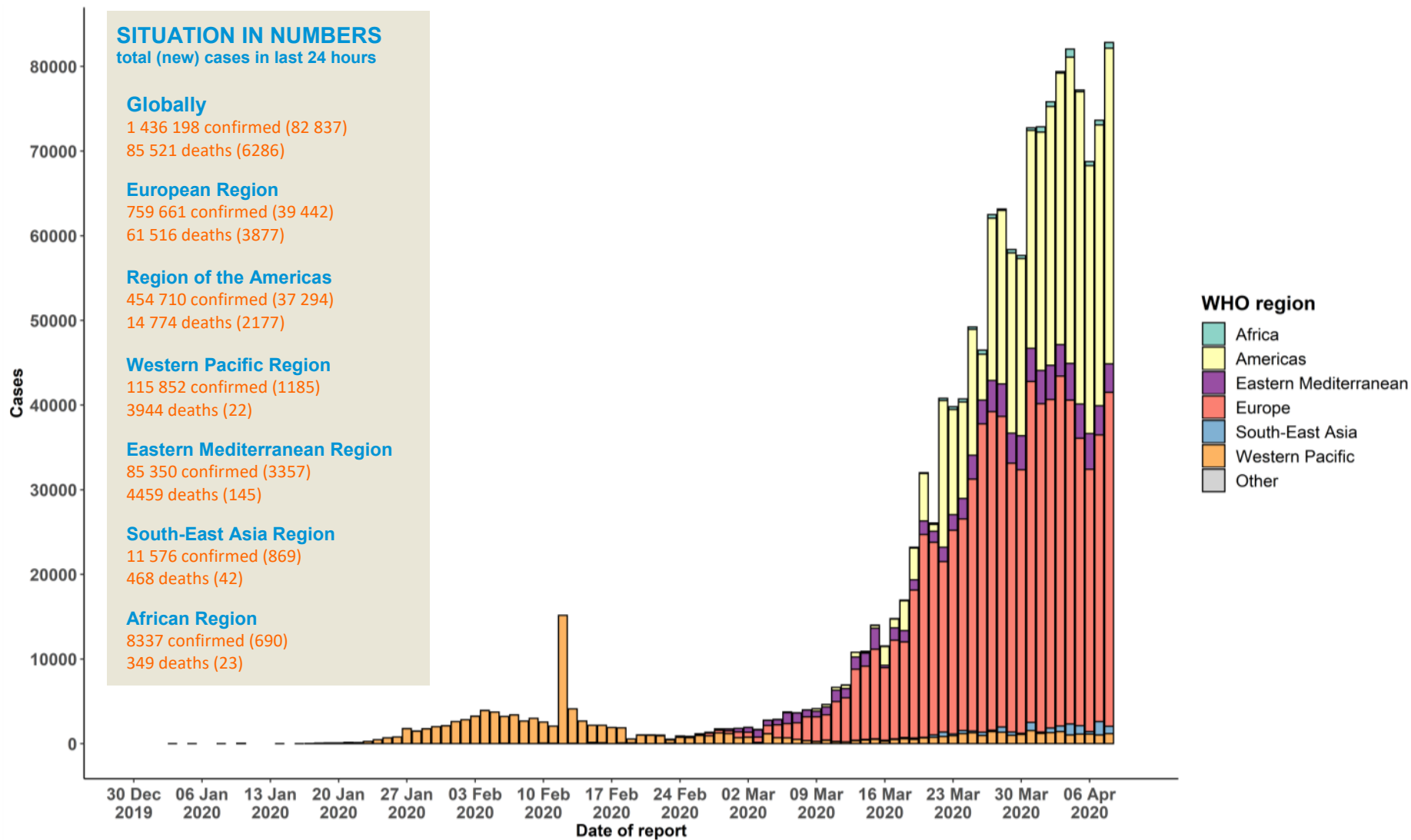


The boundaries and names shown and the designations used on this map do not imply the expression of any opinion whatsoever on the part of the World Health Organization concerning the legal status of any country, territory, city or area or of its authorities, or concerning the delimitation of its frontiers or boundaries. Dotted and dashed lines on maps represent approximate border lines for which there may not yet be full agreement.

Data source: [WHO](#), National Health Commission of the People's Republic of China
Map production: WHO Health Emergencies Programme
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World Health Organization Situation Report

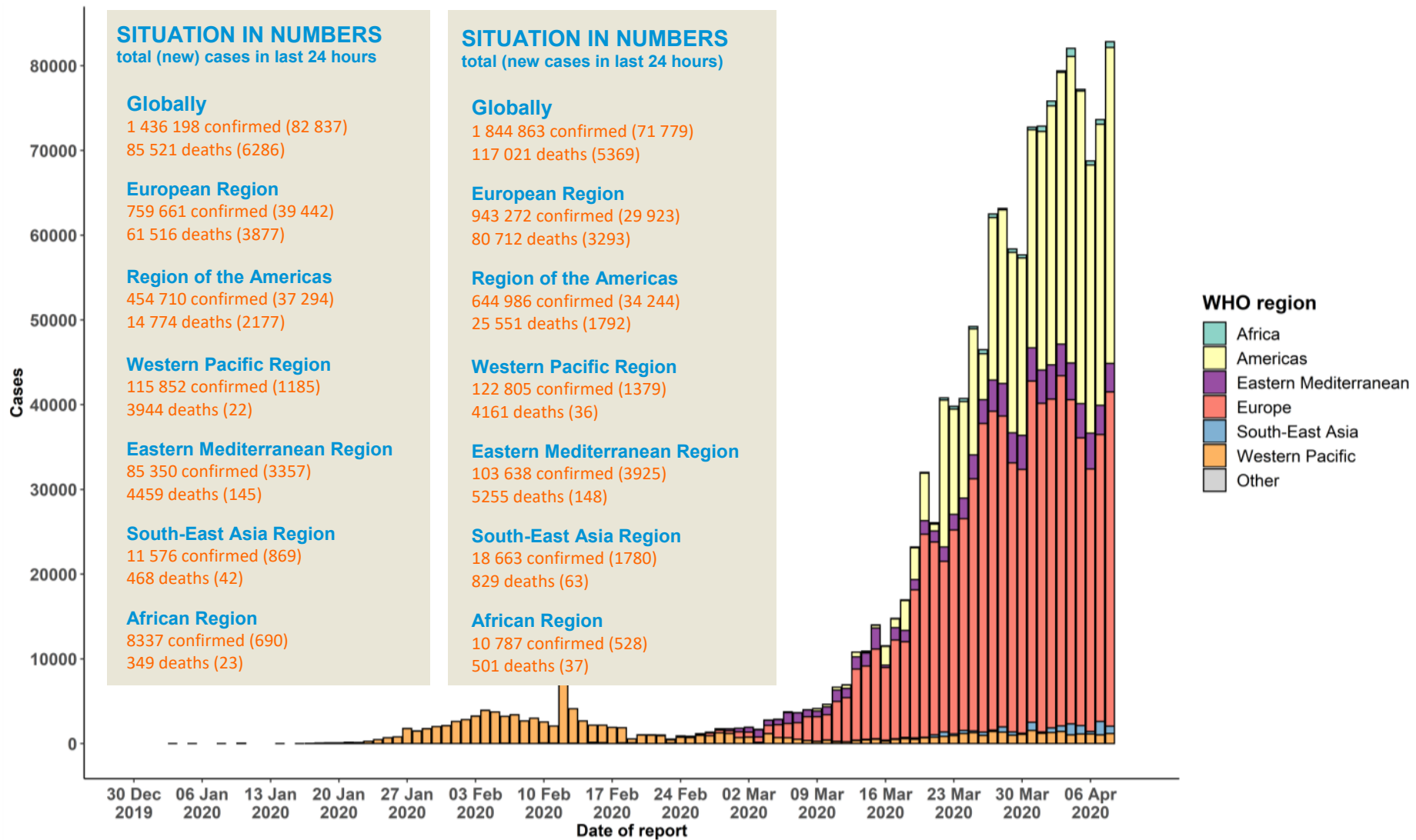
Source: WHO, Feb 9, 2020



World Health Organization Situation Report

Source: WHO, April 9, 2020 – 2 months later

April 14, 400K new cases and 30K new deaths in 5 days



World Health Organization Situation Report

Source: WHO, April 9, 2020 – 2 months later

Today's Talk

Part 1 – Dynamical Foundations of Epidemics

strength, speed, and size

Part 2 – Dynamics and Control

how we got to where we are now

Part 3 – Long-term strategies

how we might get out, in the absence of pharmaceutical interventions

Today's Talk

Part 1 – Dynamical Foundations of Epidemics

strength, speed, and size

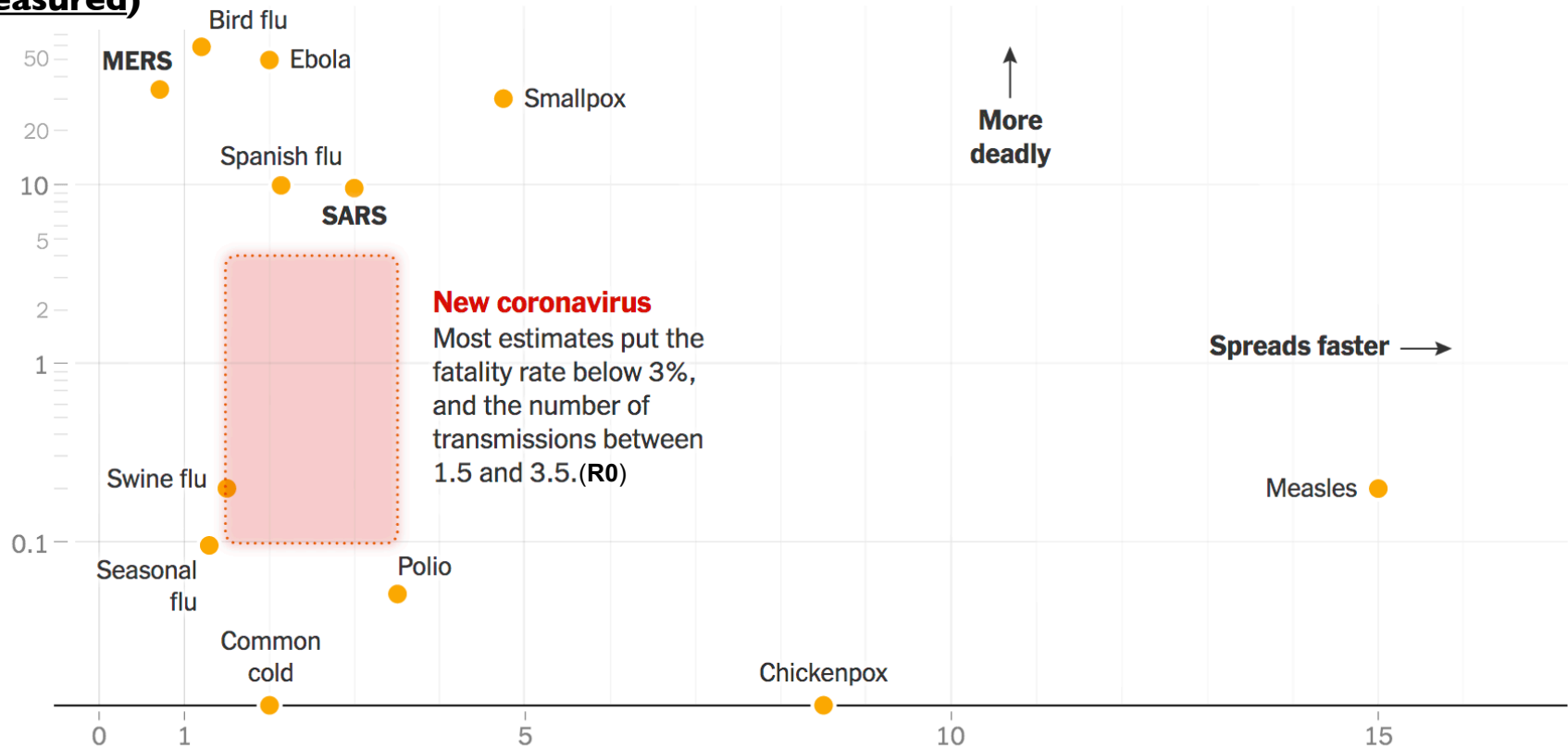
Part 2 – Dynamics and Control

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**Fatality rate
(directly
measured)**

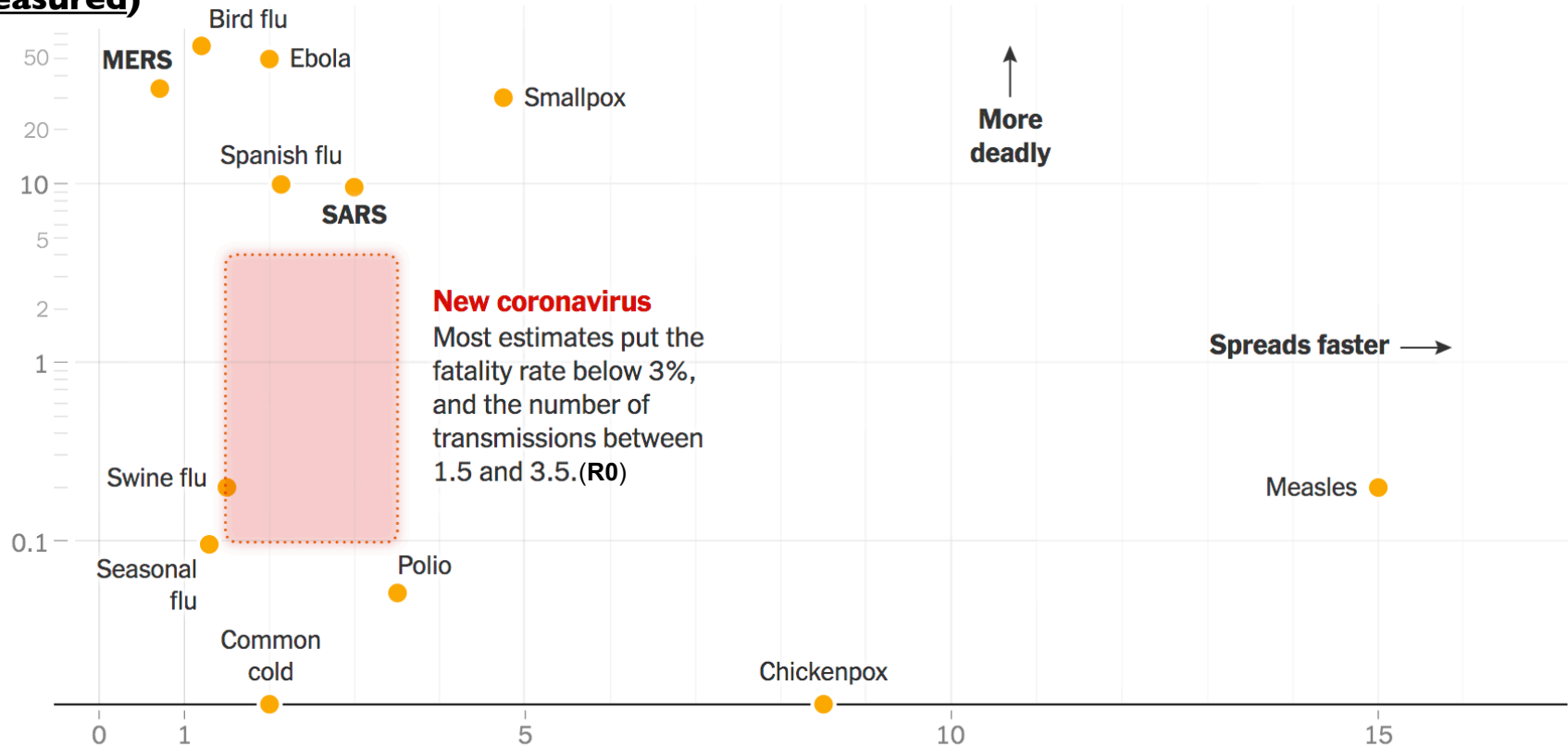


Basic reproductive number, " R_0 "
Equal to the average number of new infections per sick person
Indirectly measured

Early Question Motivating Work:

How certain should we be about estimates of the strength – R_0 – of a disease at the outset of an outbreak?

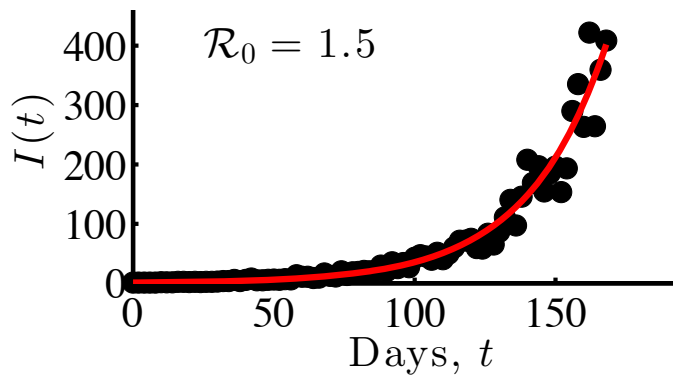
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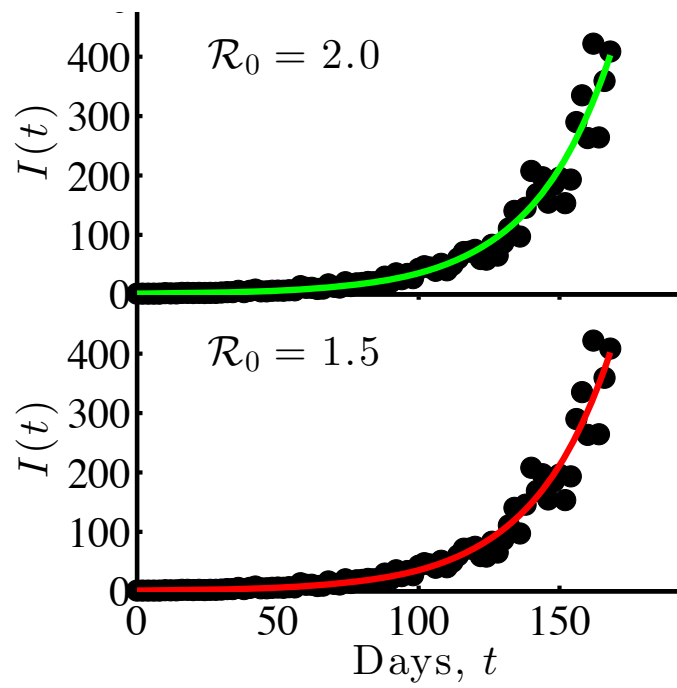


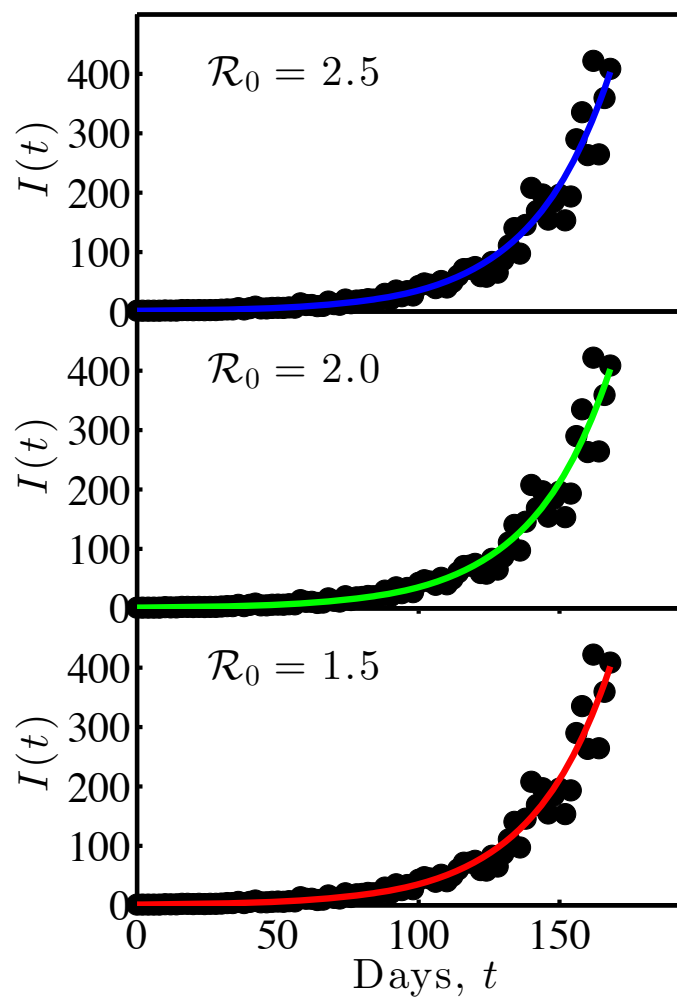
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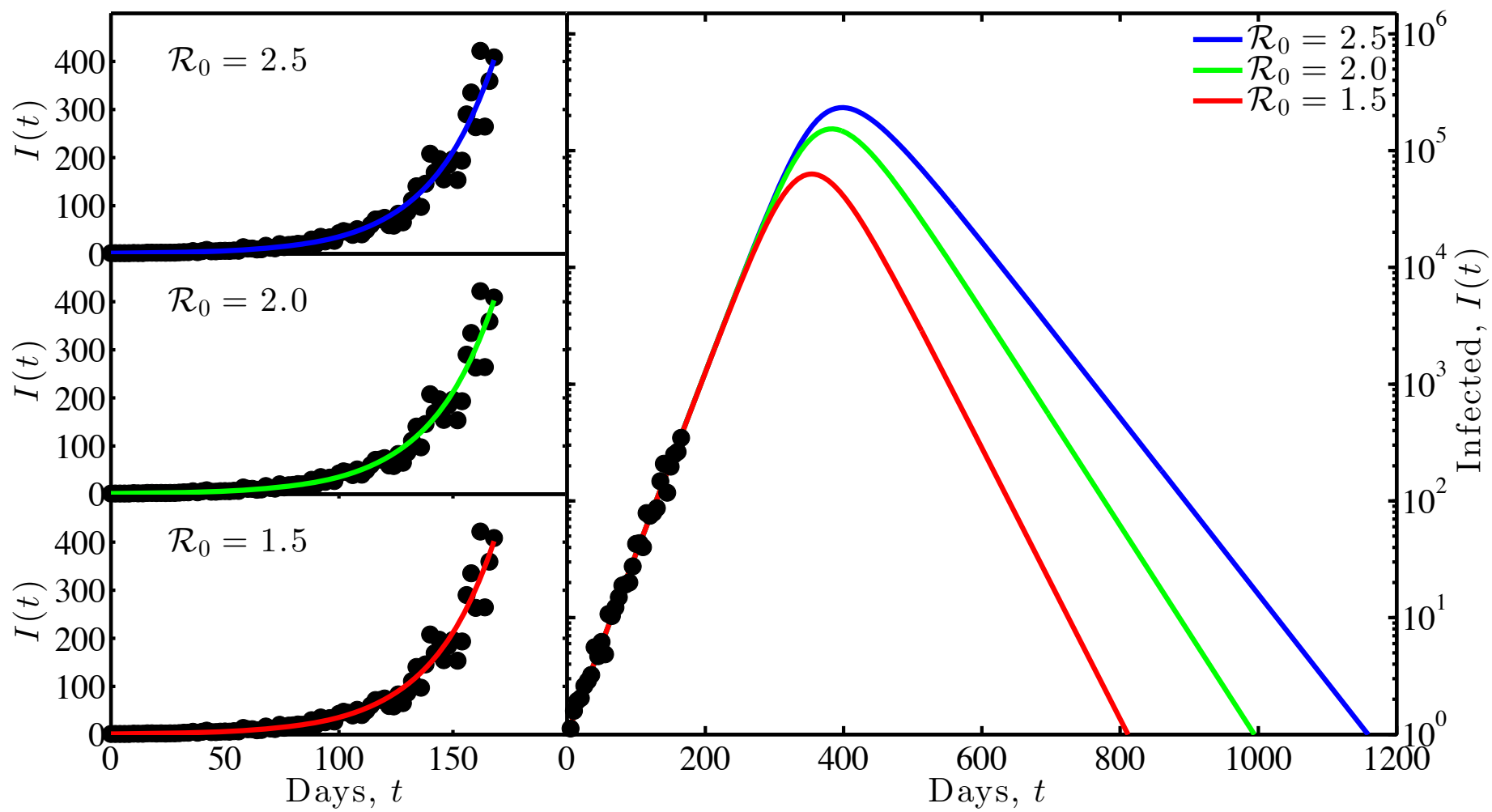
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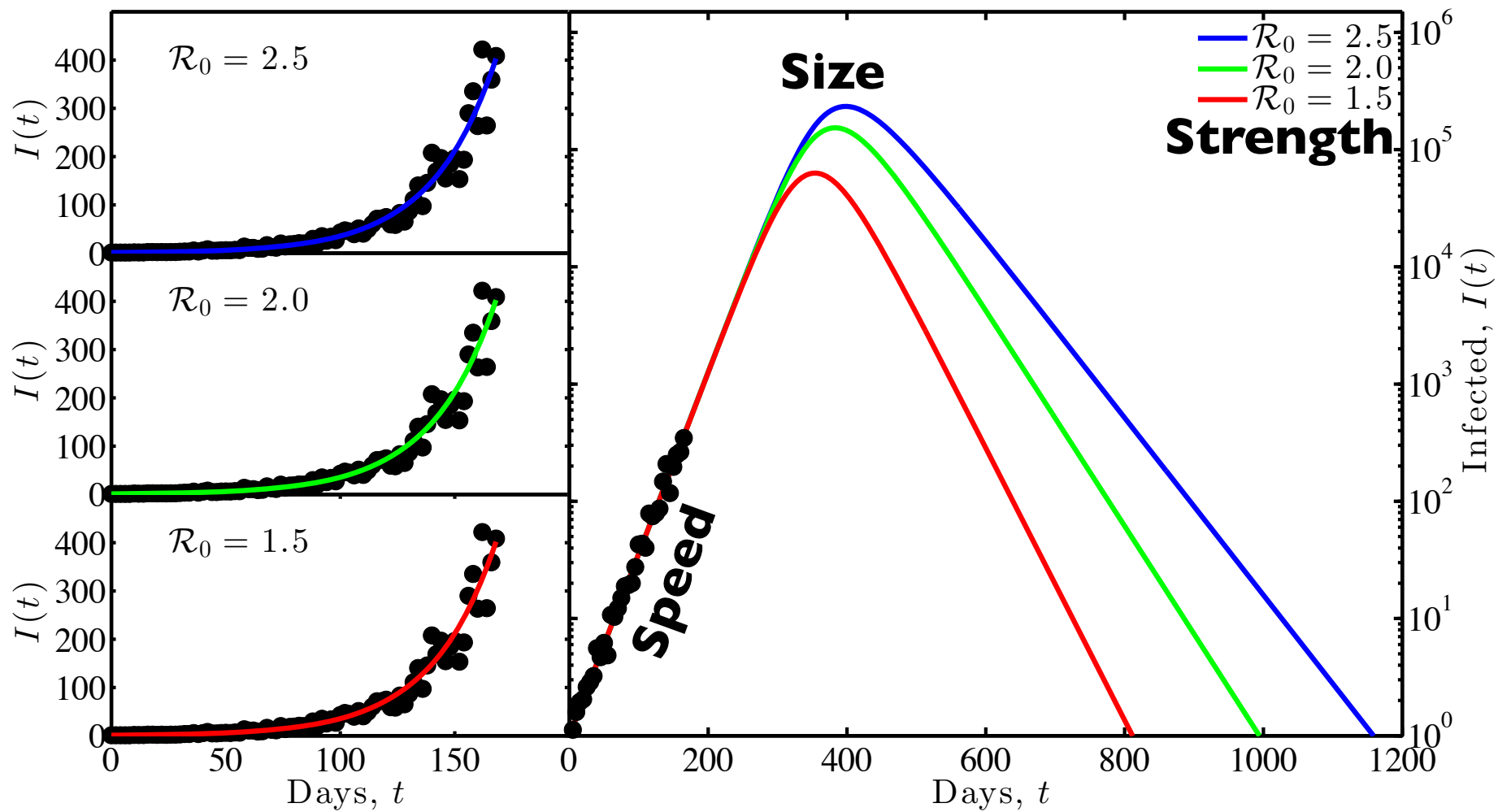
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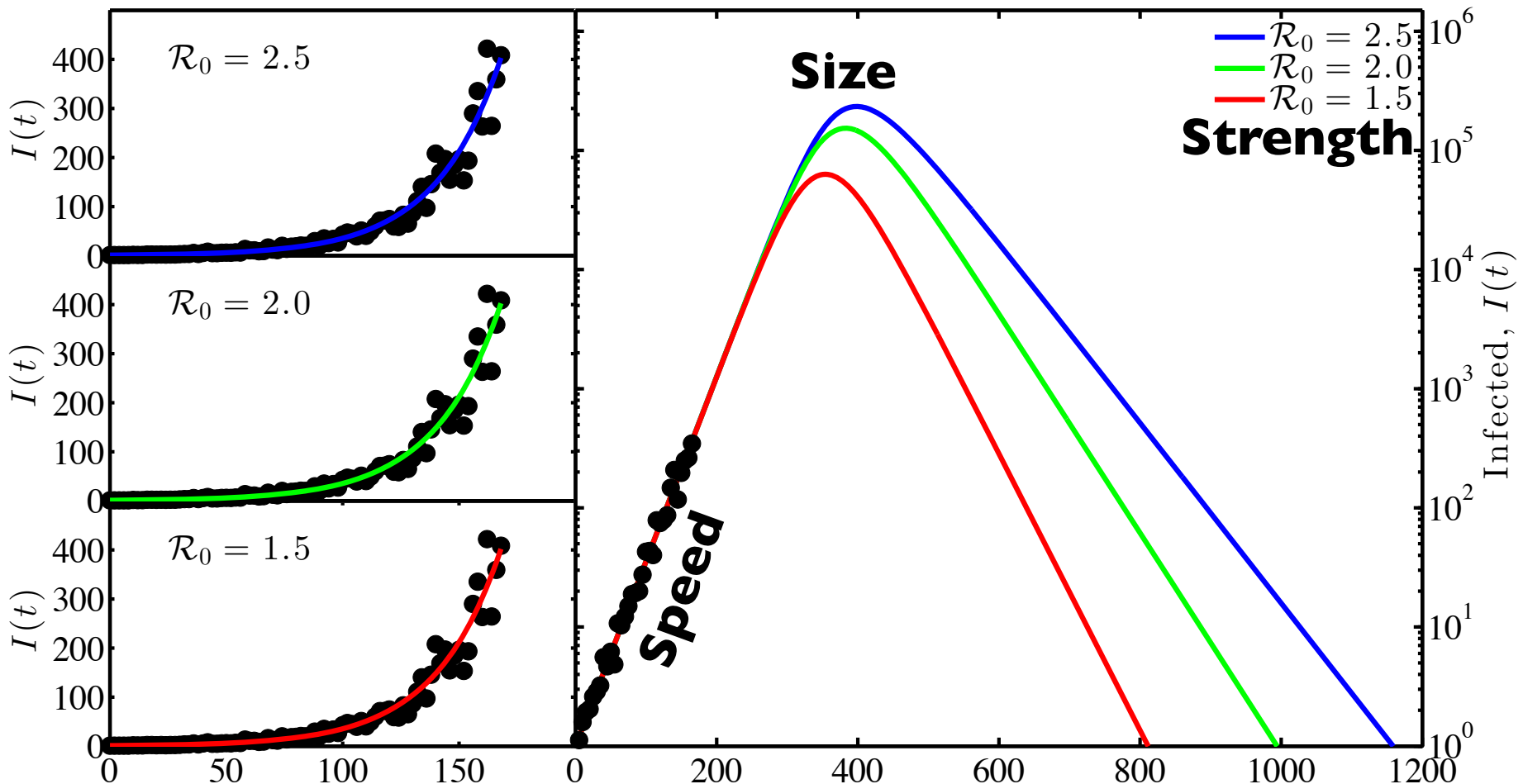






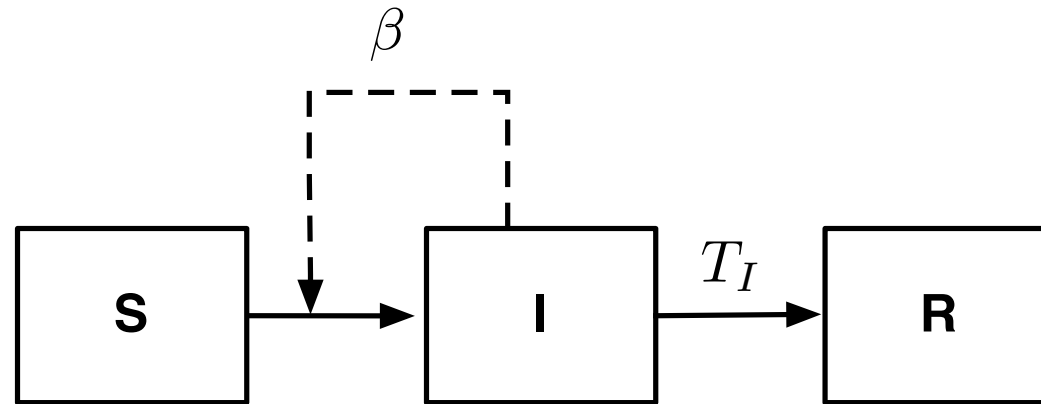






Tentative conclusion: Many values of R_0 can be compatible with the same observed rate of increase in cases – even if projected outbreak sizes are different.

SIR Model - Basics



Population “Classes”

S – The number of susceptible individuals

I – The number of infectious individuals

R – The number of “removed” individuals (through recovery or, possibly, death)

Mechanisms

Infection: Requiring contact between a **S** and a **I** individual at rate β .

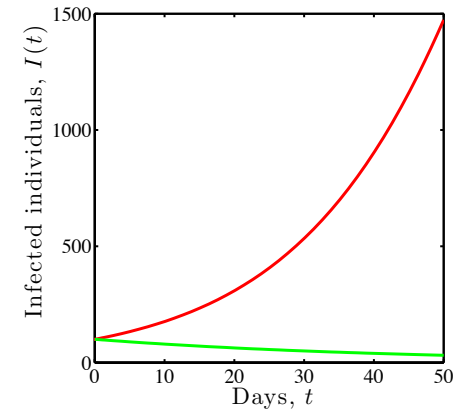
Recovery: After a period of infectiousness of average duration T_I .

SIR Model – Initial Dynamics

Depend on Basic Reproductive Number, R_0

The expected number of cases, initially changes like:

$$\dot{I} = \frac{I}{T_I} (\mathcal{R}_0 - 1)$$



SIR Model – Initial Dynamics

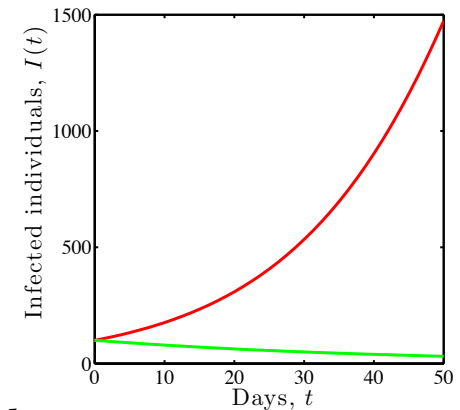
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where

$$\mathcal{R}_0 \equiv \underbrace{\text{infections per time}}_{\beta} \times \underbrace{\text{infectious period}}_{T_I}$$



SIR Model – Initial Dynamics

Depend on Basic Reproductive Number, R_0

The expected number of cases, initially changes like:

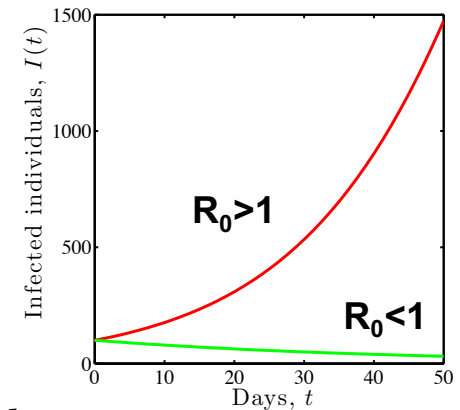
$$\dot{I} = \frac{I}{T_I} (\mathcal{R}_0 - 1)$$

where

$$\mathcal{R}_0 \equiv \underbrace{\text{infections per time}}_{\beta} \times \underbrace{\text{infectious period}}_{T_I}$$

such that

- Disease spreads whenever the average number of new cases exceeds unity, i.e: $\mathcal{R}_0 > 1$
- The increase is exponential



Estimating, R_0 , for 2019-nCoV

	Basic reproductive number $\mathcal{R}_0$
Study 1	1.5–3.5
Study 2	2.5 (1.5–3.5)*
Study 3	2.92 (95% CI: 2.28–3.67)
Study 4	3.8 (95% CI: 3.6–4.0)
Study 5	2.2 (90% CI: 1.4–3.8)
Study 6	5.47 (95% CI: 4.16–7.10) [‡]
Study 7	2.0–3.1

Bedford <i>et al.</i> [4]
Imai <i>et al.</i> [5]
Liu <i>et al.</i> [6]
Read <i>et al.</i> [8]
Riou and Althaus [10]
Zhao <i>et al.</i> [9]
Majumder and Mandl [7]

Many model choices:

Branching process

SEIR model (like SIR but with an asymptomatic class)

Exponential growth...

Estimating, R_0 , for 2019-nCoV

	Basic reproductive number $\mathcal{R}_0$	Mean generation interval $\bar{G}$ (days)	Generation-interval dispersion κ	
Study 1	1.5–3.5	10	1	Bedford <i>et al.</i> [4]
Study 2	2.5 (1.5–3.5)*	8.4	unspecified [†]	Imai <i>et al.</i> [5]
Study 3	2.92 (95% CI: 2.28–3.67)	8.4	0.2	Liu <i>et al.</i> [6]
Study 4	3.8 (95% CI: 3.6–4.0)	7.6	0.5	Read <i>et al.</i> [8]
Study 5	2.2 (90% CI: 1.4–3.8)	7–14	0.5	Riou and Althaus [10]
Study 6	5.47 (95% CI: 4.16–7.10) [‡]	7.6–8.4	0.2	Zhao <i>et al.</i> [9]
Study 7	2.0–3.1	6–10	0	Majumder and Mandl [7]

Many model choices & latent assumptions:

Branching process

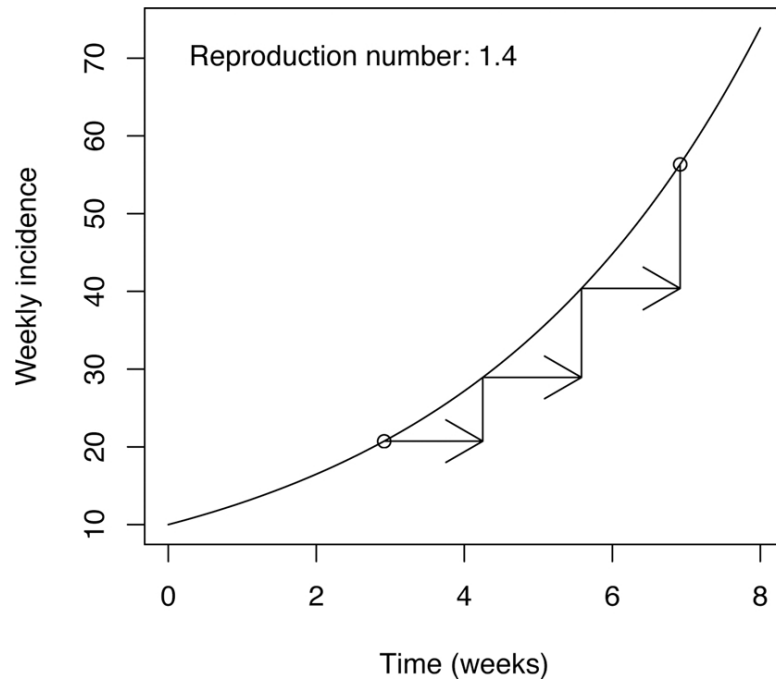
SEIR model (like SIR but with an asymptomatic class)

Exponential growth...

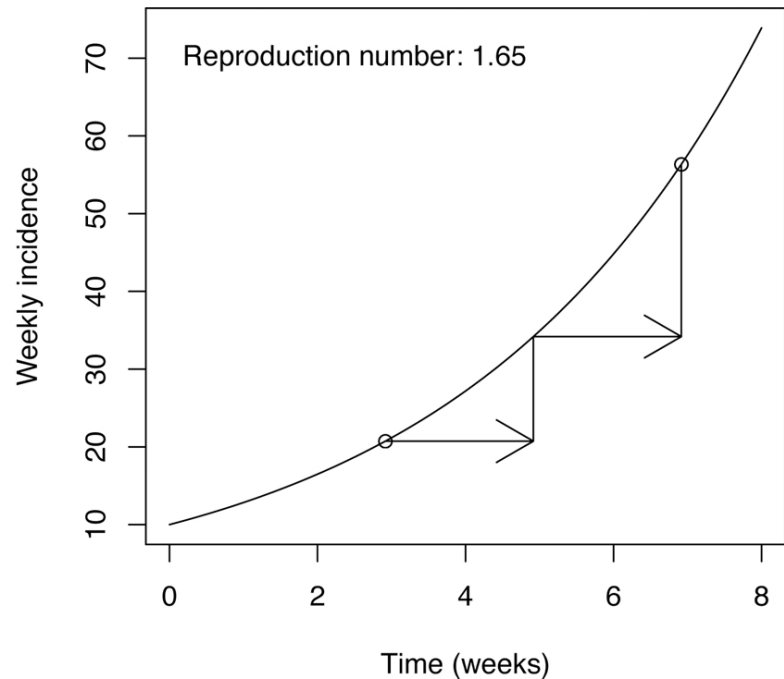
The brief answer is that speed and strength are linked.

The implicit link between speed and strength

Shorter generation intervals



Longer generation intervals



Epidemics 27 (2019) 12–18



Contents lists available at ScienceDirect

Epidemics

journal homepage: www.elsevier.com/locate/epidemics

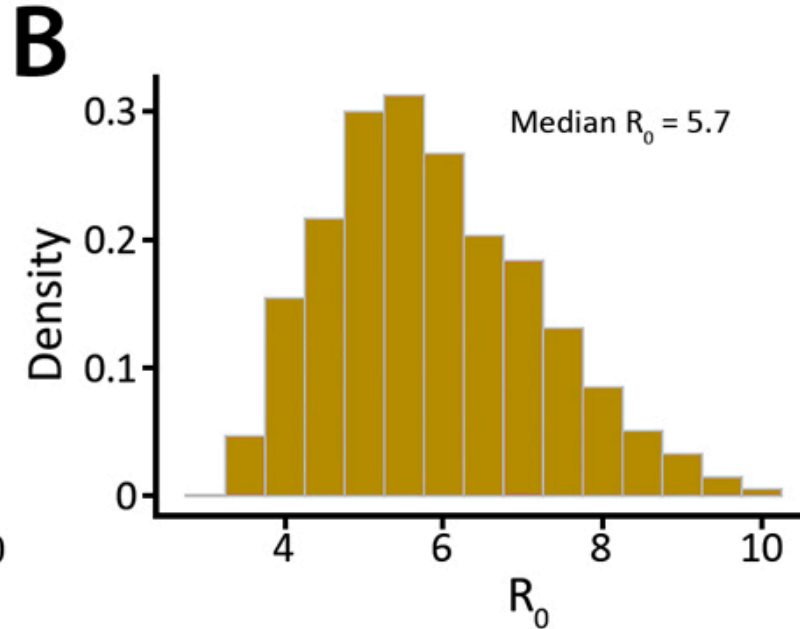
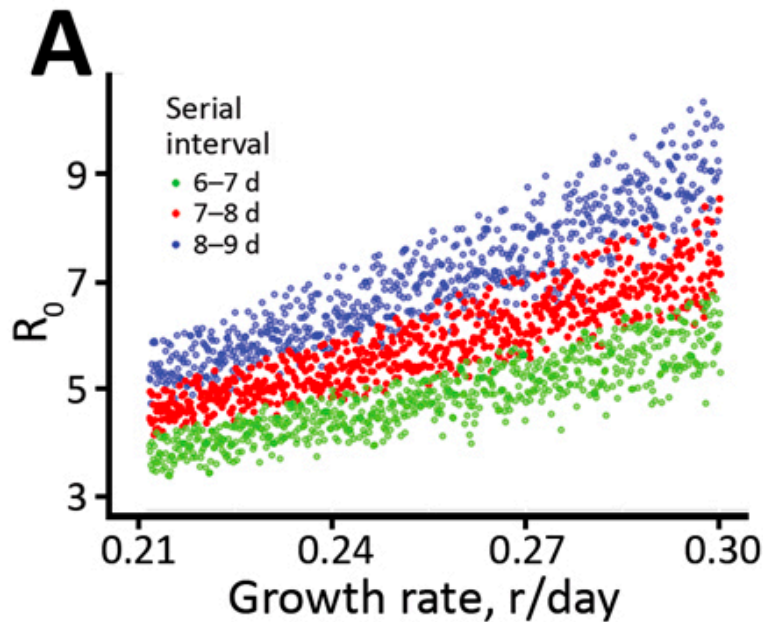


A practical generation-interval-based approach to inferring the strength of epidemics from their speed

Sang Woo Park^a, David Champredon^{b,c}, Joshua S. Weitz^{d,e}, Jonathan Dushoff^{a,*}



This link helps sort through putatively large R_0 claims (assumptions matter!)



EMERGING INFECTIOUS DISEASES®

ISSN: 1080-6059

EID Journal > Volume 26 > Early Release > Main Article > Figure 5



Disclaimer: Early release articles are not considered as final versions. Any changes will be reflected in the online version in the month the article is officially released.

Volume 26, Number 7—July 2020

Research

High Contagiousness and Rapid Spread of Severe Acute Respiratory Syndrome Coronavirus 2

Steven Sanche¹, Yen Ting Lin¹, Chonggang Xu, Ethan Romero-Severson, Nick Hengartner, and Ruian Ke²

Author affiliations: ¹Los Alamos National Laboratory, Los Alamos, New Mexico, USA

Forbes

EDITORS' PICK | 107,243 views | Apr 7, 2020, 05:34pm EDT

The COVID-19 Coronavirus Disease May Be Twice As Contagious As We Thought

Pooled estimates via a speed-strength relationship (technically using generation intervals)

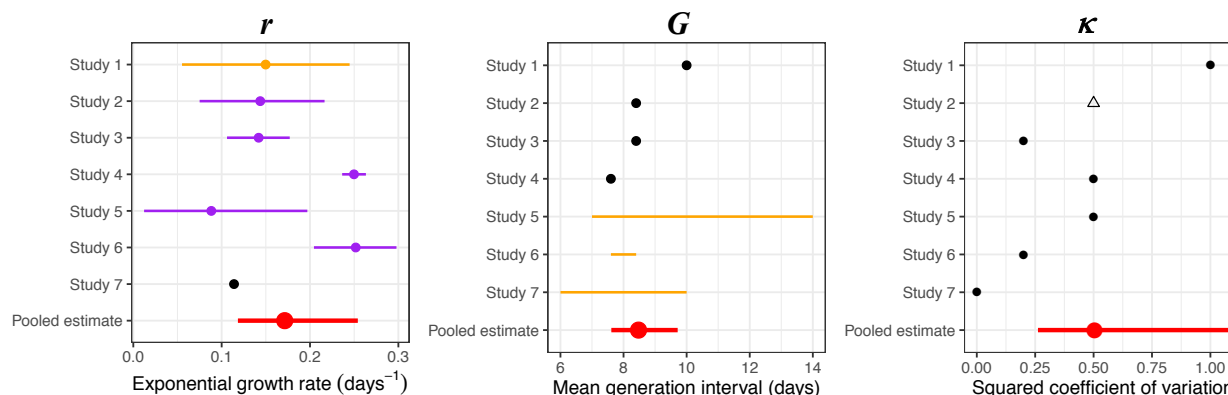
Sang Woo Park



Jonathan Dushoff

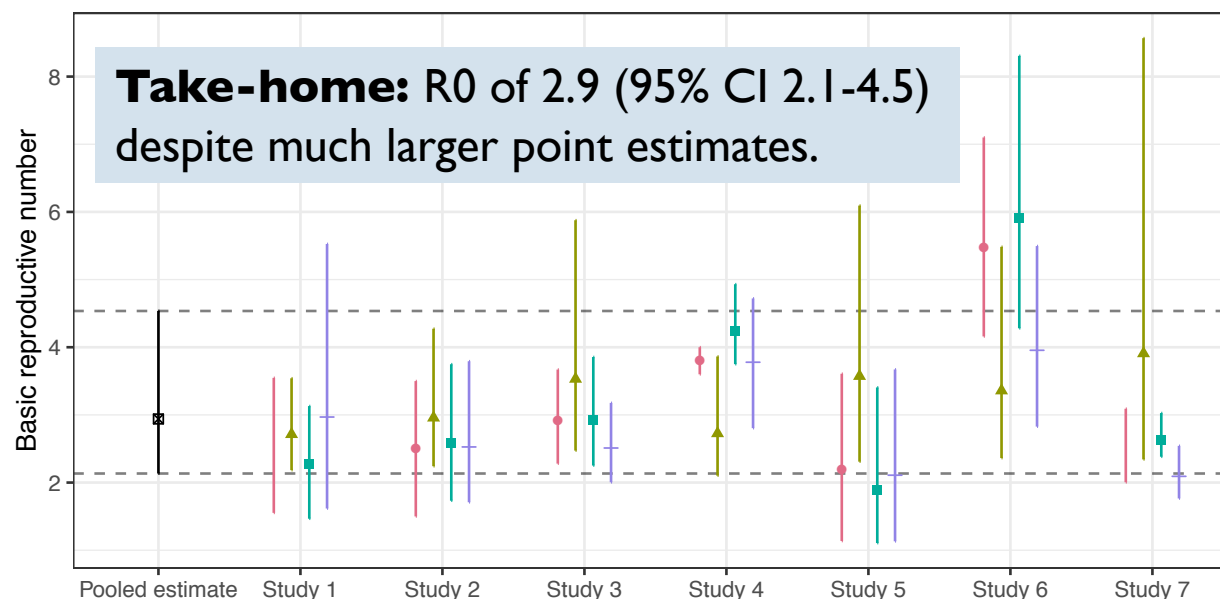


Step 1: estimate latent uncertainty in ‘parameters’.



Step 2: incorporate different types of uncertainty into R_0 estimates by study or as part of a ‘pooled’ estimate (using a Bayesian multi-level model)

$$\mathcal{R}_0 = (1 + \kappa T \bar{G})^{1/\kappa}$$



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how we might get out, in the absence of pharmaceutical interventions

Conditions for epidemic growth

$$\mathcal{R}_0 \equiv \overbrace{\beta}^{\text{infections per time}} \times \overbrace{T_I}^{\text{infectious period}}$$

Where infections per time, β , is a product of:

- Contacts by infectious individuals per unit time
- Probability of contact with a susceptible (S_0/N)
- Probability that the contact transmits the disease

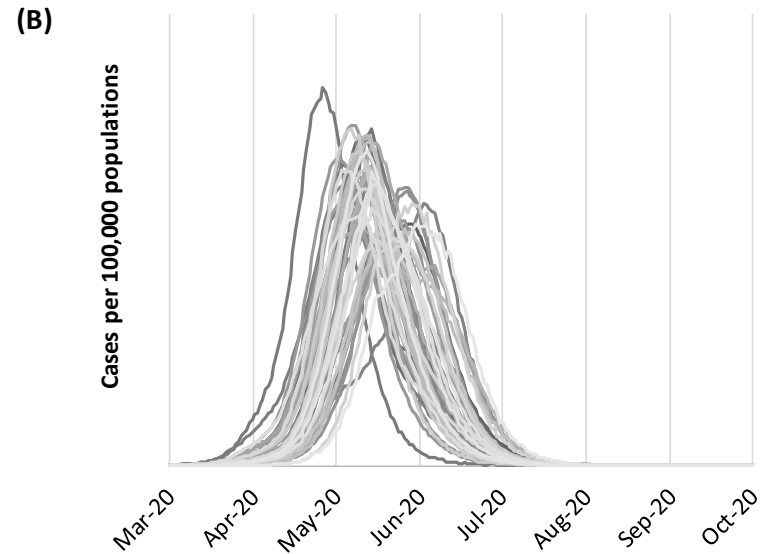
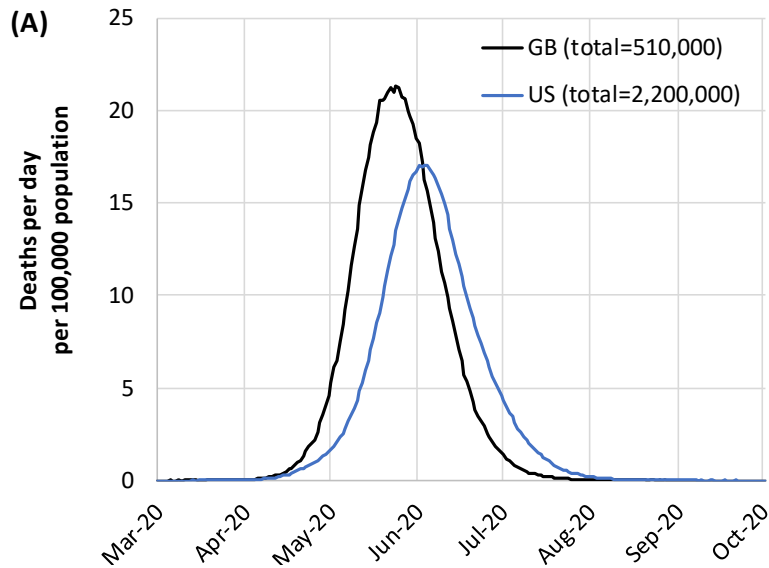
Conditions for epidemic growth also suggest opportunities for **control**

$$\mathcal{R}_0 \equiv \overbrace{\beta}^{\text{infections per time}} \times \overbrace{T_I}^{\text{infectious period}} \quad \textbf{Hospitalization \& treatment}$$

Where infections per time, β , is a product of:

- Contacts by infectious individuals per unit time **Contact tracing & targeted isolation**
- Probability of contact with a susceptible (S_0/N) **Quarantine, travel control, closures**
- Probability that the contact transmits the disease **Process engineering & PPE (masks)**

The Imperial Model Forecast – The Scale of the Problem



The unmitigated problem. This goes back to strength-size.

A strength of $R_0 \sim 2.4$ implies $\sim 80\%$ infected (in a 'mean-field' scenario).

Population of 330M $\times 80\% \times \sim 0.8\%$ IFR $\sim 2\text{M}+$ fatalities.



The reality of the Ebola outbreak is not reflected by model projections of high case numbers.

EPIDEMIOLOGY

Models overestimate Ebola cases

Rate of infection in Liberia seems to plateau, raising questions over the usefulness of models in an outbreak.

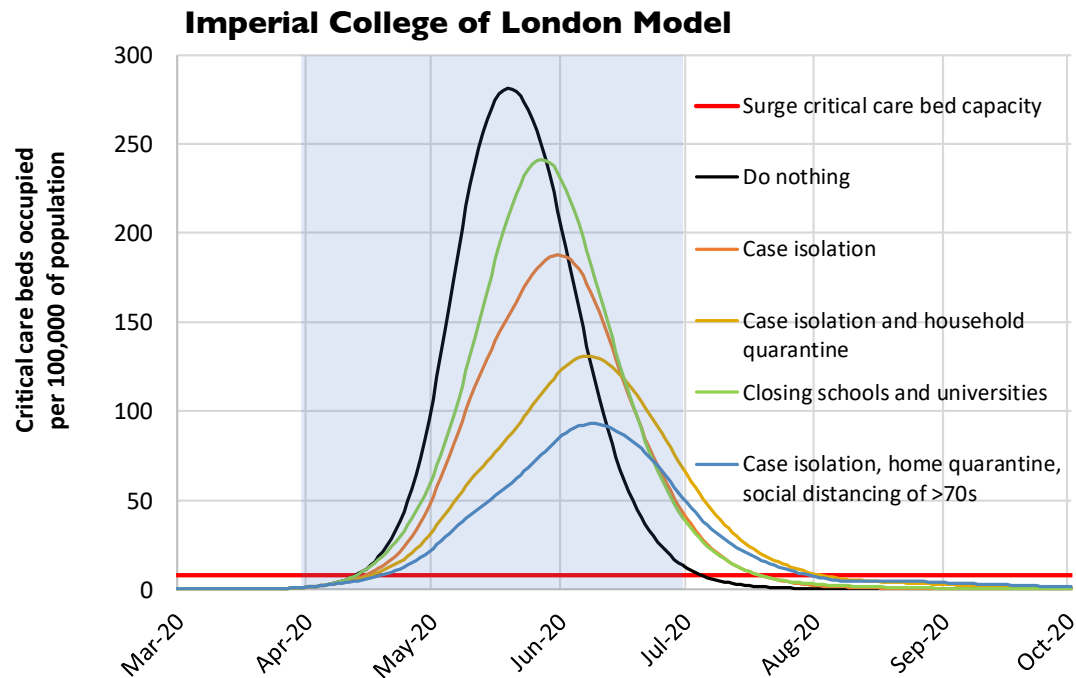


The reality of the Ebola outbreak is not reflected by model projections of high case numbers.

EPIDEMIOLOGY

Models overestimate **Coronavirus** cases

Rate of infection in Liberia seems to plateau, raising questions over the usefulness of models in an outbreak.



Model Purpose: to explain the need, scope, and potential outcome of **interventions**.

Different measures both reduce and 'flatten' the curve (i.e., reducing total burden)



Popular

Latest

TECHNOLOGY

Don't Believe the COVID-19 Models



That's not what they're for.

ZEYNEP TUFEKCI APRIL 2, 2020



Popular

Latest

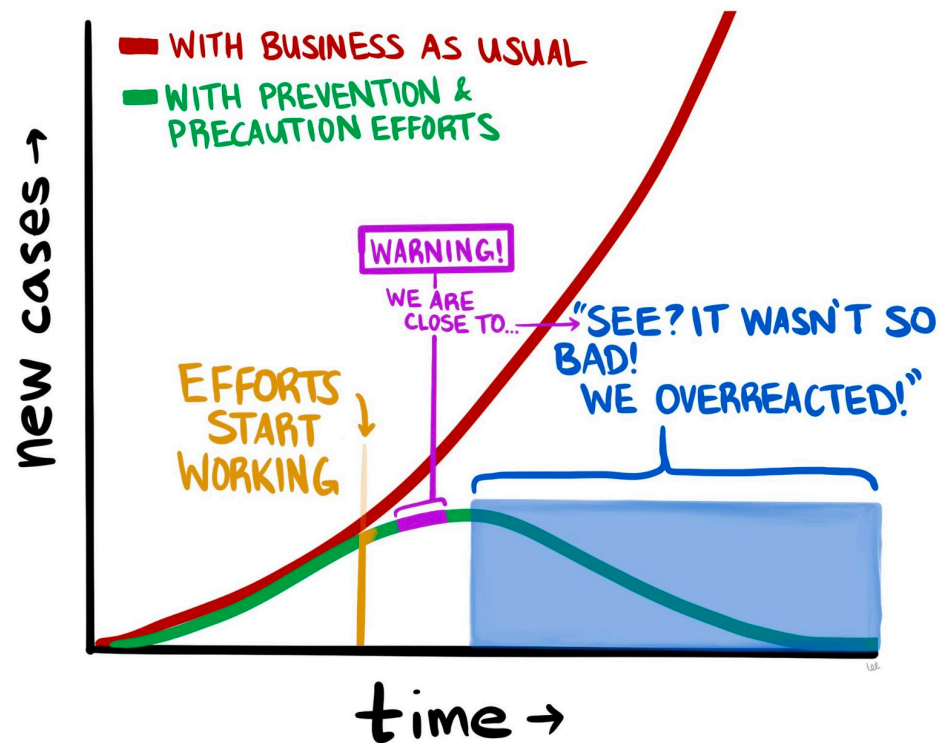
TECHNOLOGY

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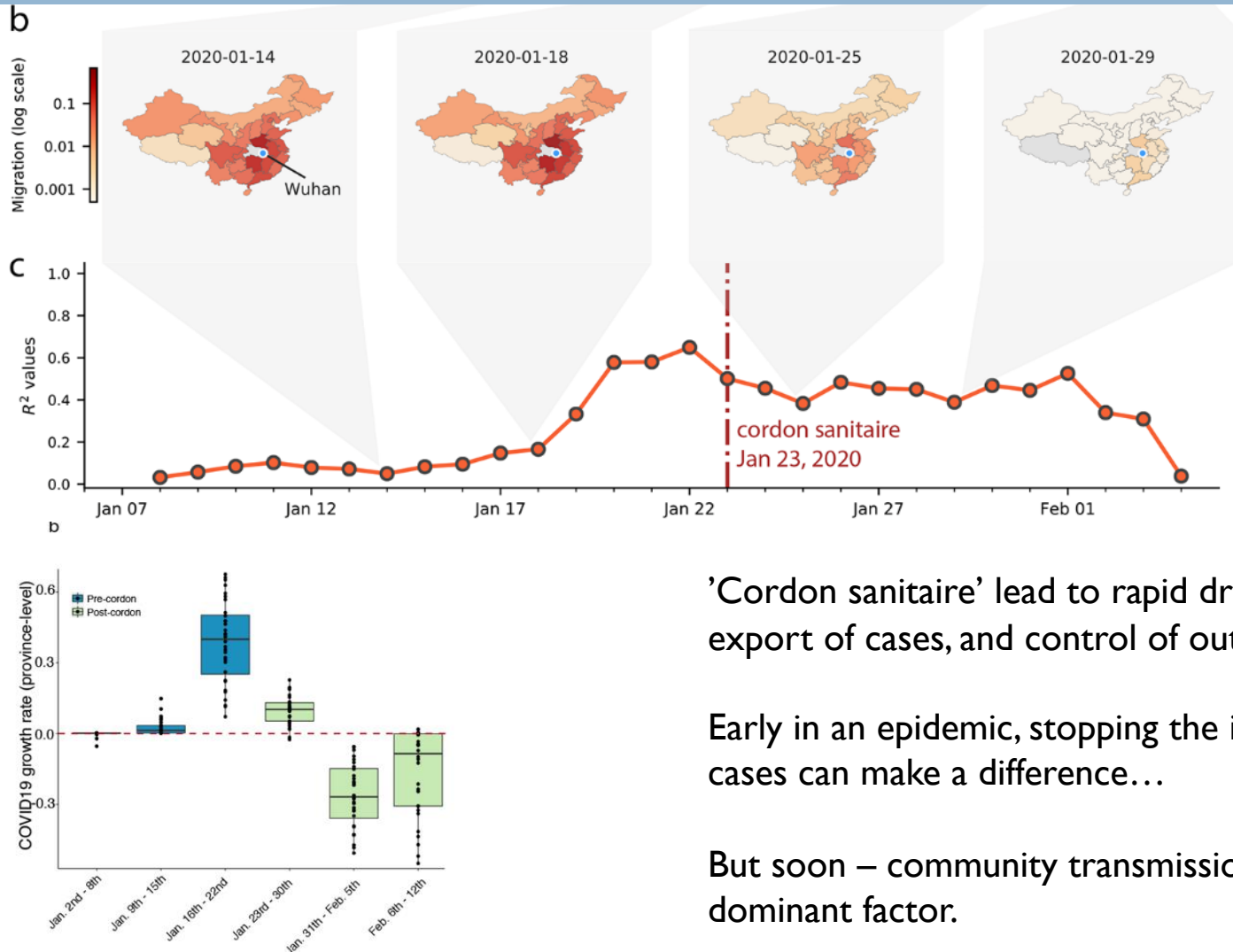
ZEYNEP TUFEKCI APRIL 2, 2020



What Evidence Is There on the Impact of Interventions?

Measures of Impact of Interventions

Scarpino and colleagues (Science 3/25/20)



'Cordon sanitaire' lead to rapid drop in export of cases, and control of outbreak.

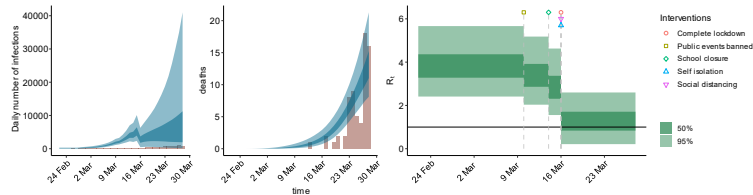
Early in an epidemic, stopping the import of cases can make a difference...

But soon – community transmission is the dominant factor.

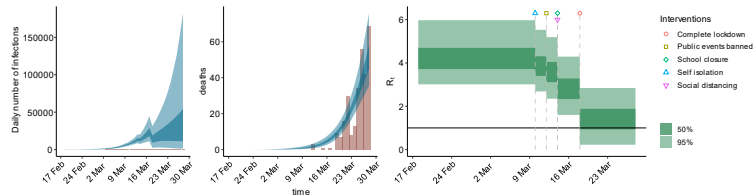
Measures of Impact of Interventions

Imperial College London Report, 3/30/20

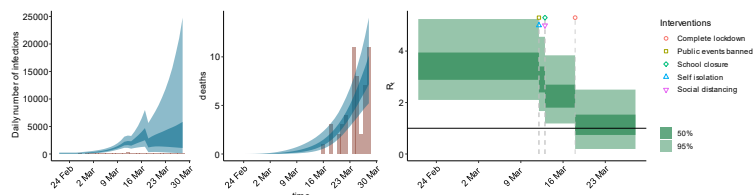
(A) Austria



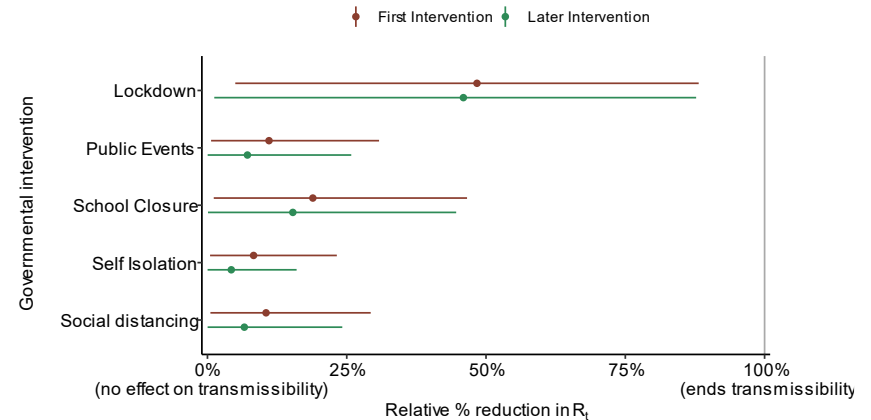
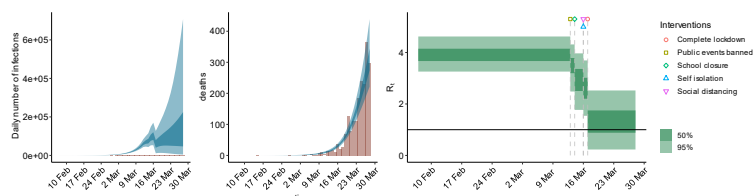
(B) Belgium



(C) Denmark



(D) France



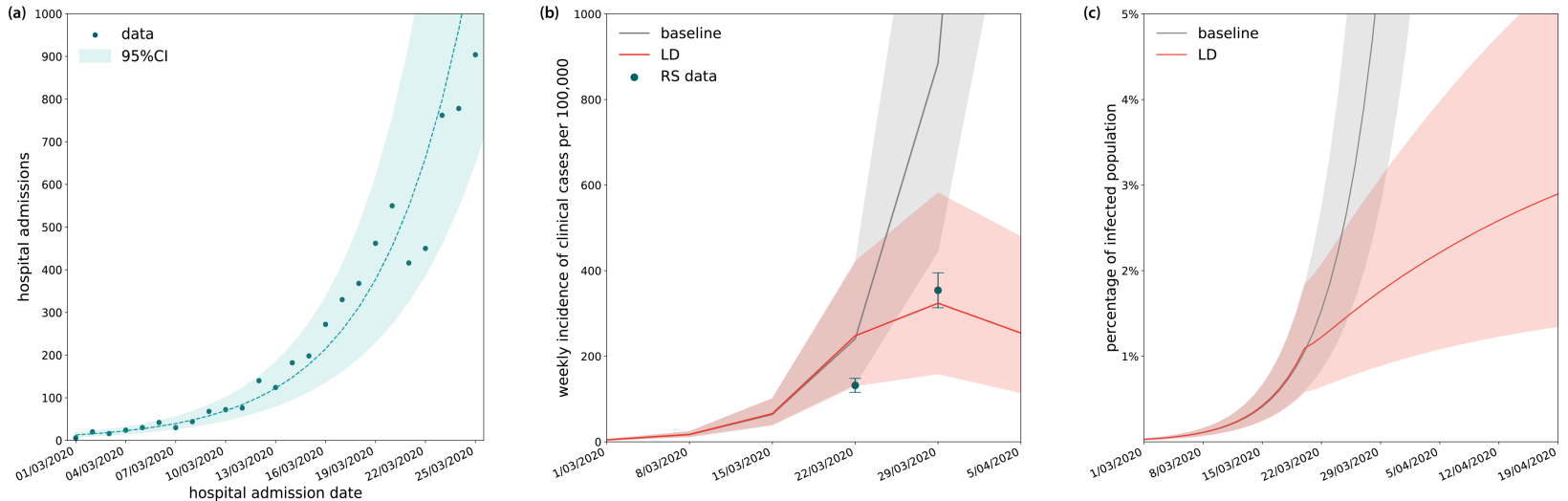
“[E]stimate that interventions across all 11 countries will have averted 59,000 deaths up to 31 March [95% credible interval 21,000-120,000].

Many more deaths will be averted through ensuring that interventions remain in place until transmission drops to low levels.”

Flaxman et al.

Measures of Impact of Interventions (4/12/20)

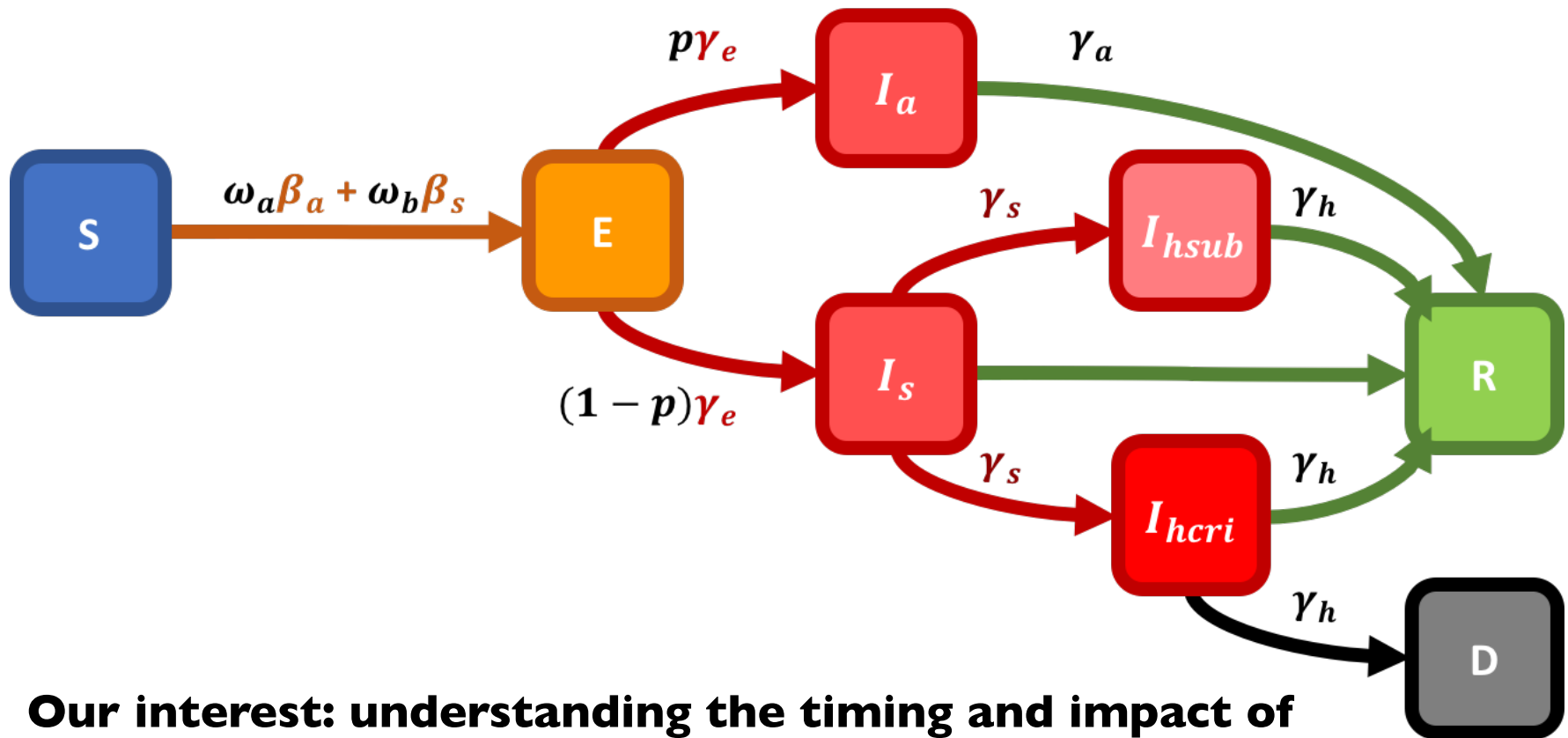
INSERM – Ile De France (Vittoria Colizza & team)



Take-away:

Lockdown interventions have rapidly shifted the curve from perilous exponential-like growth to severe (2-5% infected).

But there are still deep uncertainties which make COVID-19 hard to predict and control



Our interest: understanding the timing and impact of asymptomatic cases on epidemic potential...

Asymptomatic cases may be a significant driver of transmission (and bias estimates of R_0 and control)

Renewal equation formalism

$$\dot{S} = -i(t)$$
$$i(t) = \mathcal{R}_a S(t) \int_0^\infty i_a(t-\tau) g_a(\tau) d\tau + \mathcal{R}_s S(t) \int_0^\infty i_s(t-\tau) g_s(\tau) d\tau.$$

and assuming exponential growth
with an observed rate r yields:

$$\frac{1}{\mathcal{R}_0} = \int \exp(-r\tau) g(\tau) d\tau.$$

Speed r : measured

Generation interval : assumed (and
informed by clinical data)

Leads to an estimate of:

Strength R_0

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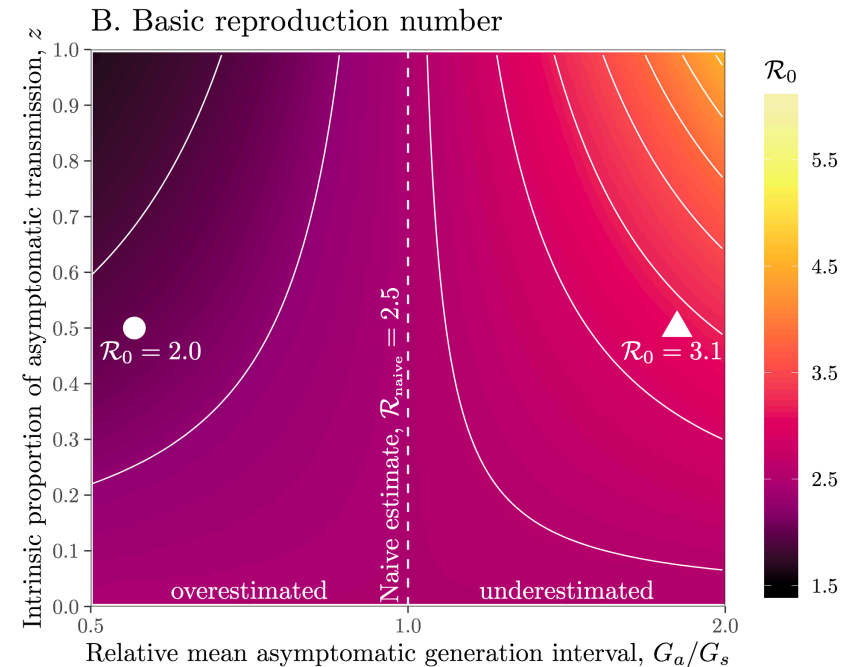
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Epidemics (in press) – w/SW Park, D. Cornforth & J. Dushoff

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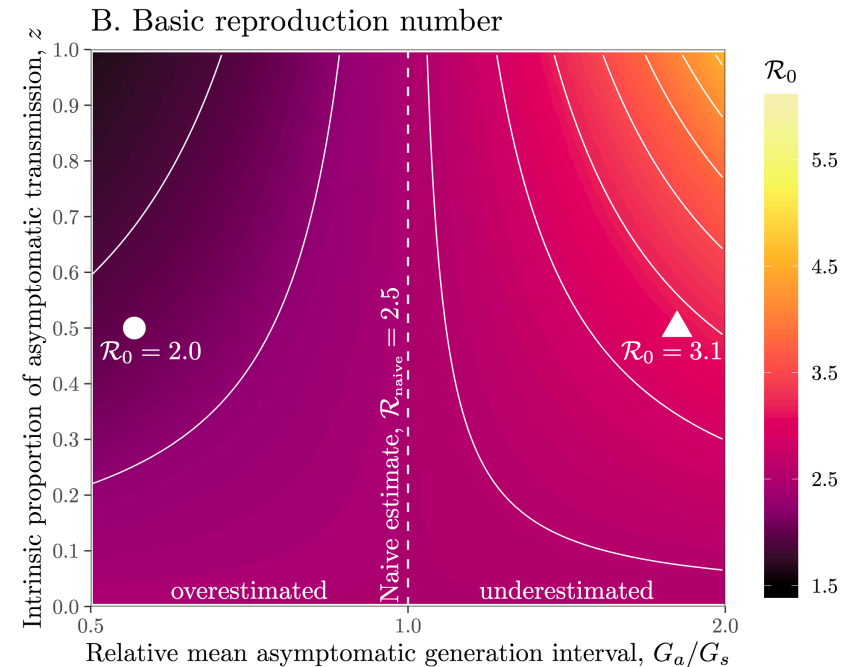
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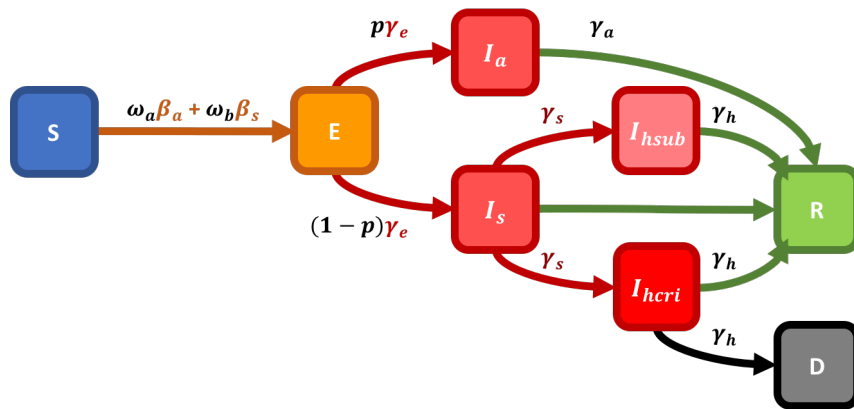
**Asymptomatic transmission
is a double-edged sword:**

- (i) Many more cases**
- (ii) Cases are not as bad**

Measures of Impact of Interventions

Georgia – Weitz Group Modeling

COVID-19 model
w/asymptomatic transmission

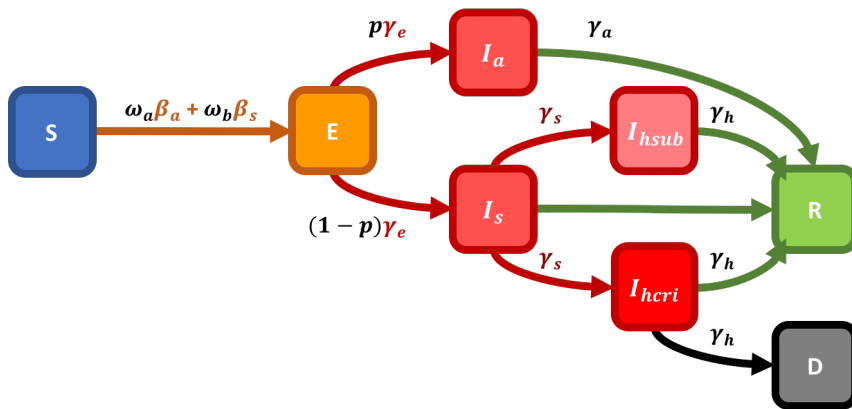


and GA age demographics
along with age-stratified risk
(via Imperial estimates).

Measures of Impact of Interventions

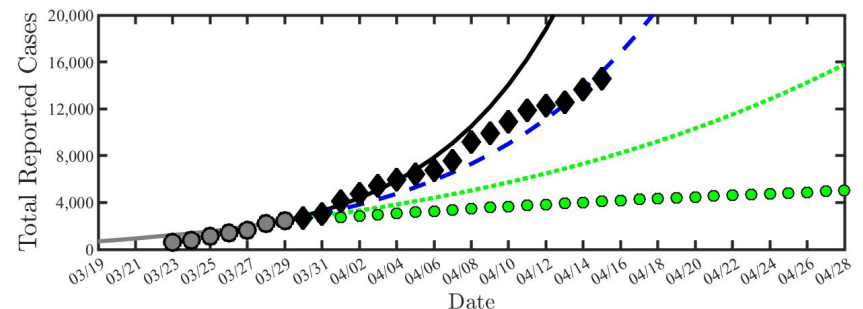
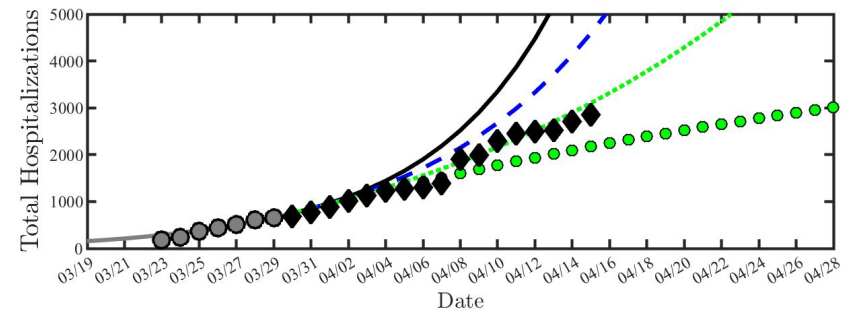
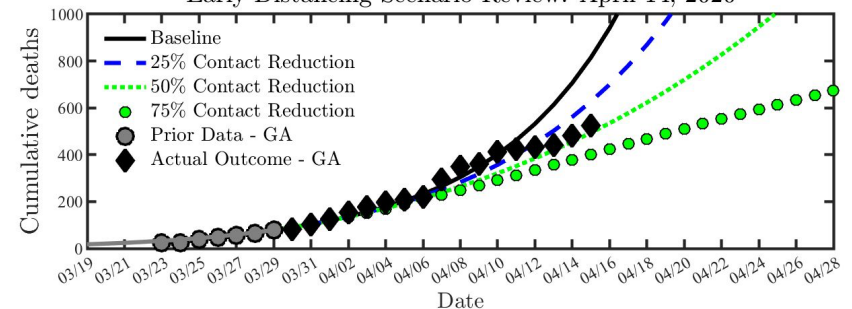
Georgia – Weitz Group Modeling

COVID-19 model
w/asymptomatic transmission



and GA age demographics
along with age-stratified risk
(via Imperial estimates).

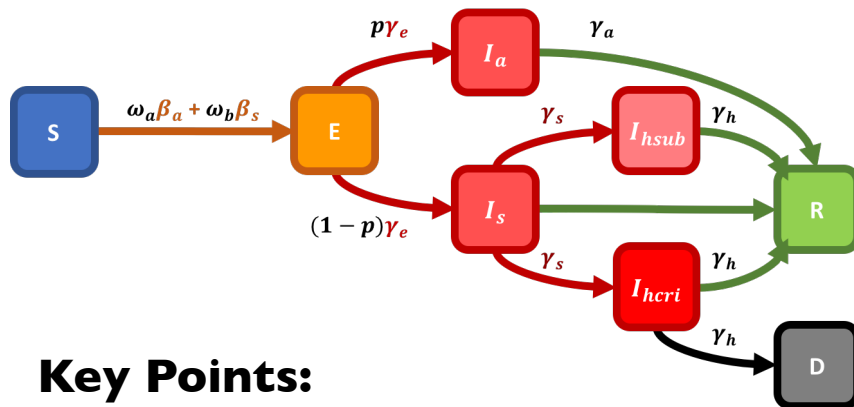
COVID-19 Near-Term Epidemic, Georgia Assessment, March 28, 2020
Early Distancing Scenario Review: April 14, 2020



Measures of Impact of Interventions

Georgia – Weitz Group Modeling

COVID-19 model w/asymptomatic transmission

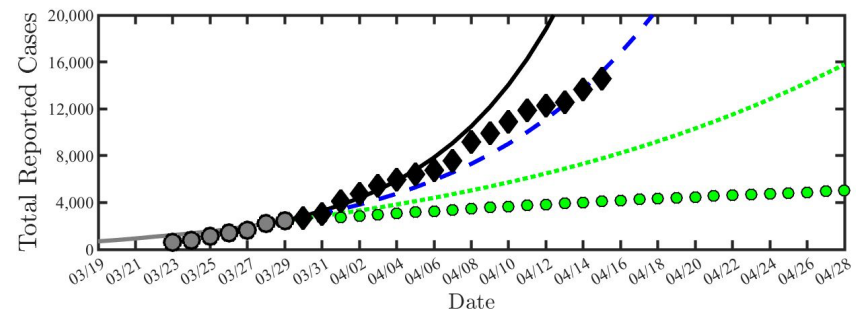
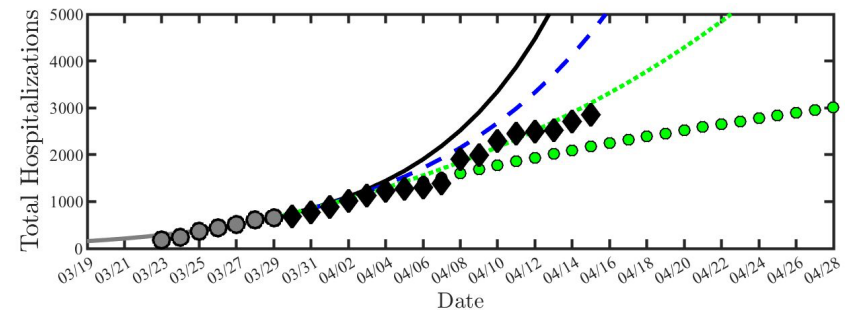
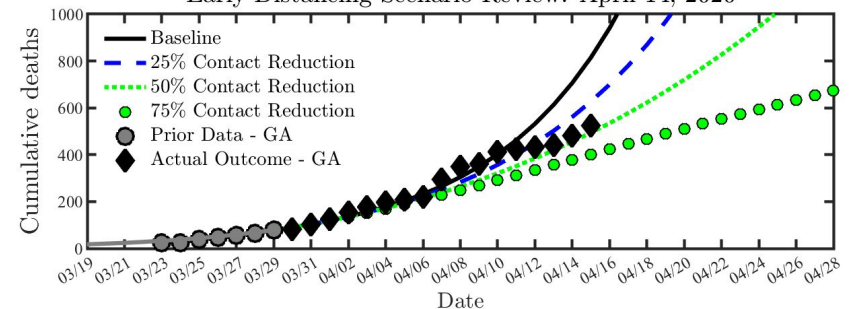


Key Points:

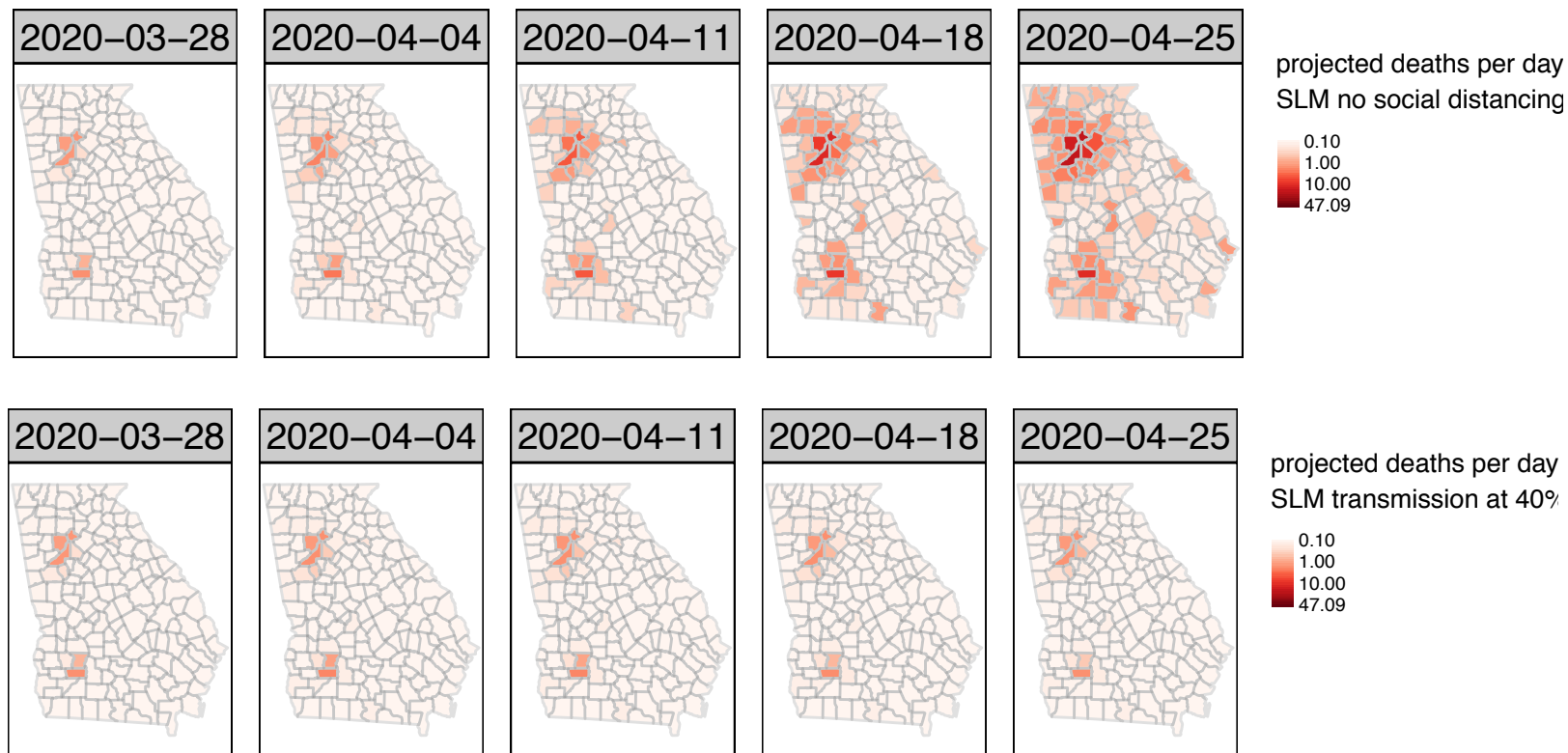
Reported deaths and hospitalizations show effect of large-scale orders (averting far worse outcomes)

Case ascertainment rate may be 1/10 to 1/30 suggesting prevalence ~1%-4% in Georgia vs. 'reported' prevalence of 0.1%.

COVID-19 Near-Term Epidemic, Georgia Assessment, March 28, 2020
Early Distancing Scenario Review: April 14, 2020



Ongoing Work – Moving from State to County-Level Metapopulation Models



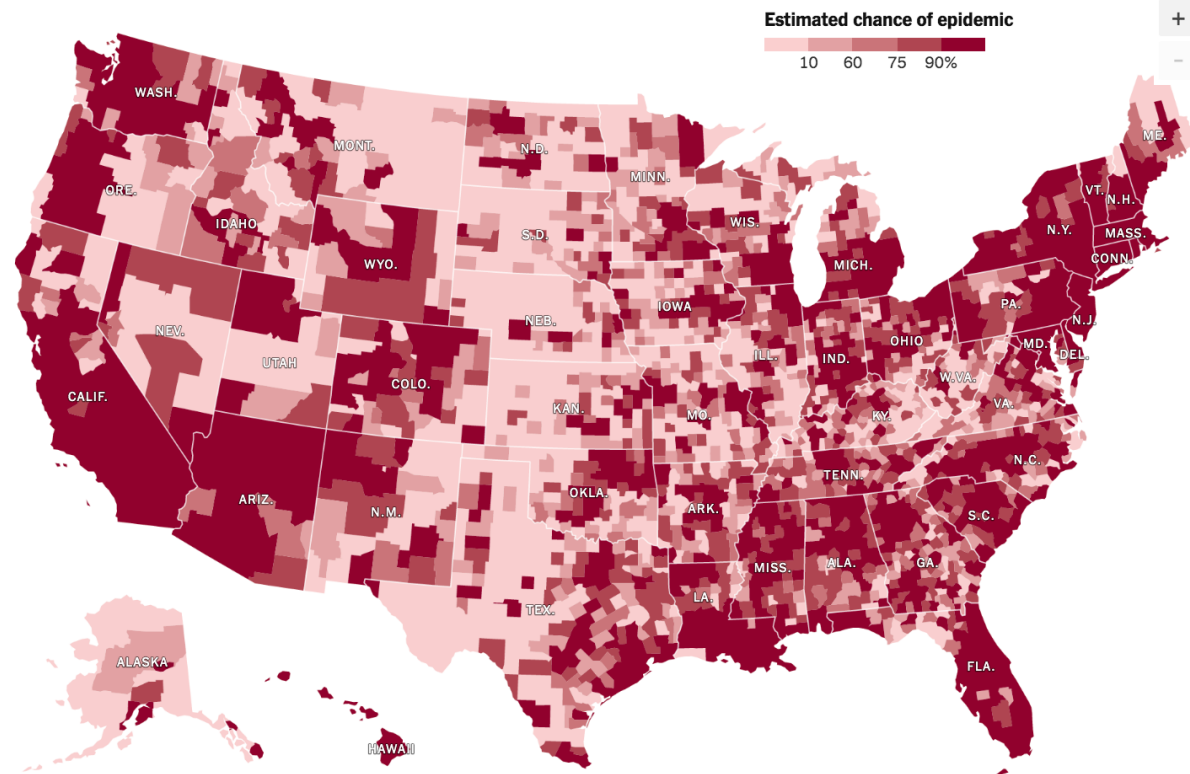
Dr. Stephen Beckett (GT, Bio Sci lead) w/Prof. Clio Andris (GT, IC and Design)
Report in progress – to be released at weitzgroup.github.io
See weitzgroup.github.io/cases.html & [deaths.html](https://weitzgroup.github.io/deaths.html) for datasets by county.

Does “My” Country Have an Epidemic?

Hint: Yes. (Lauren Meyers et al., UT-Austin)

Does My County Have an Epidemic? Estimates Show Hidden Transmission

By James Glanz, Matthew Bloch and Anjali Singhvi April 3, 2020



By The New York Times • Source: Emily Javan, Spencer Fox and Lauren Ancel Meyers, the University of Texas at Austin

Today's Talk

Part 1 – Dynamical Foundations of Epidemics

strength, speed, and size

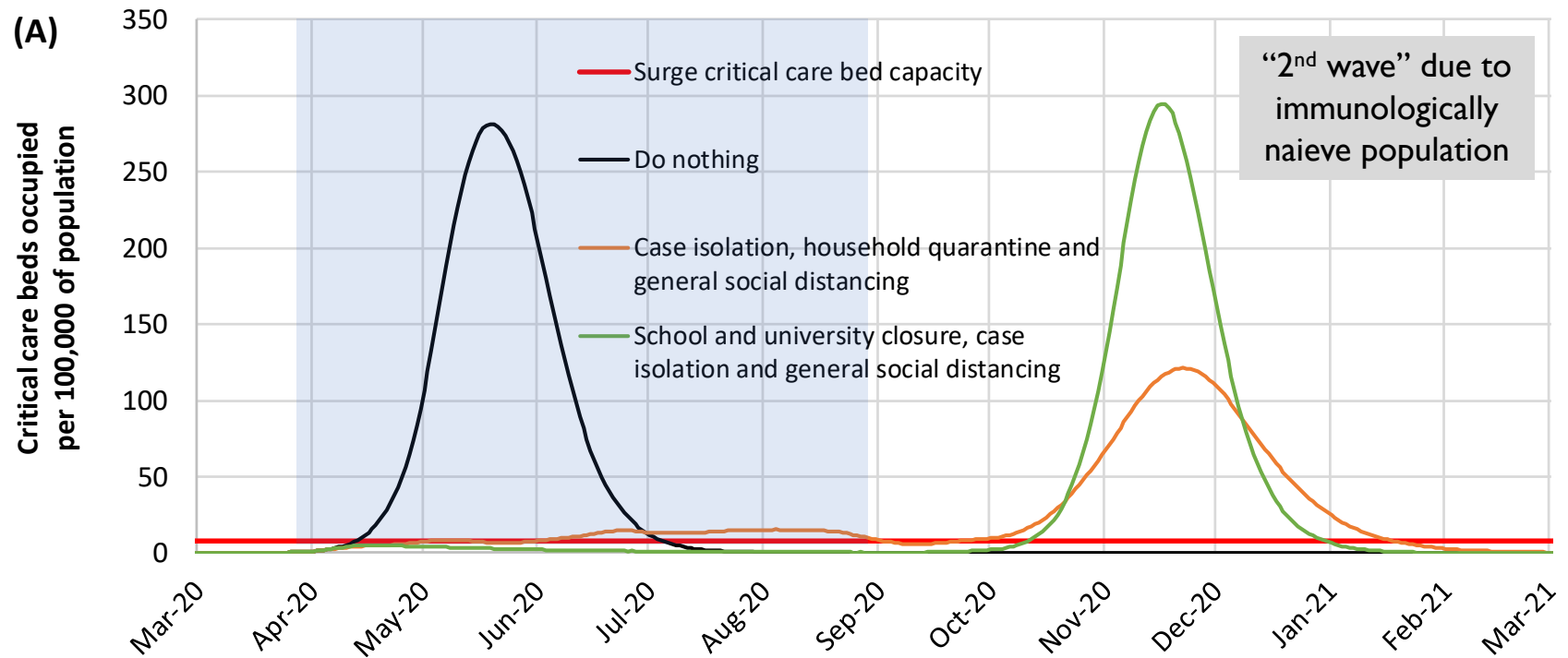
Part 2 – Dynamics and Control

how we got to where we are now

Part 3 – Long-term strategies

how we might get out, in the absence of pharmaceutical interventions

Imperial: Long-Term Prospects in the Absence of a Vaccine



Source: Imperial College London, 3/16 Report



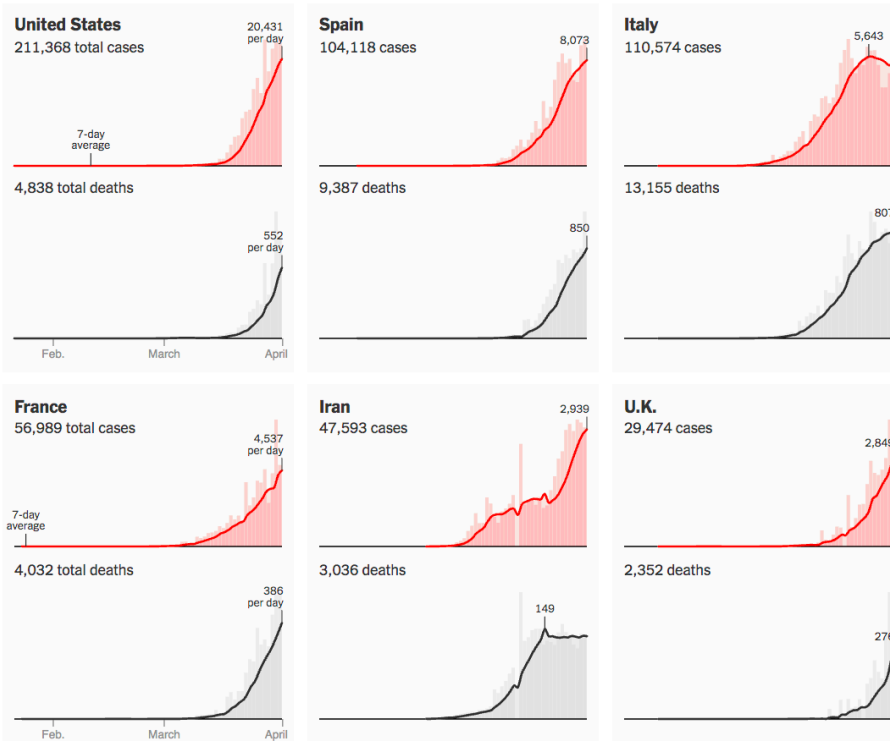
Current “Zero-Sum” Dichotomy: Infection Control or Economic Engagement



“Was the Dow to die for today, or what?!”

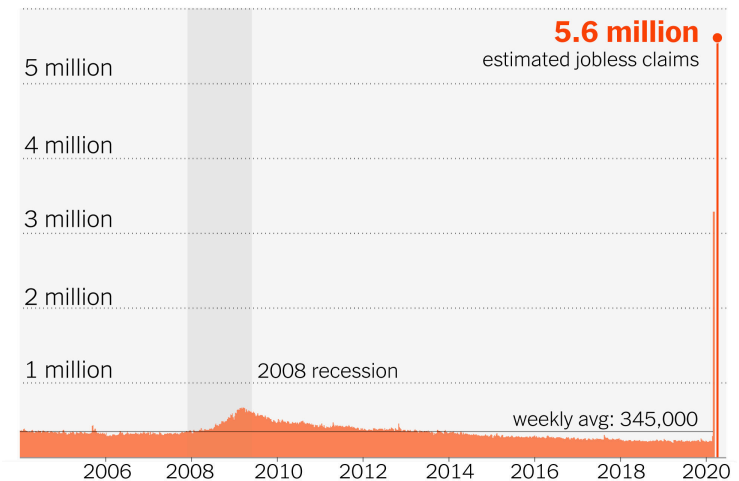
Source: Conde Nast

Are Countries Flattening the Curve for the Coronavirus?



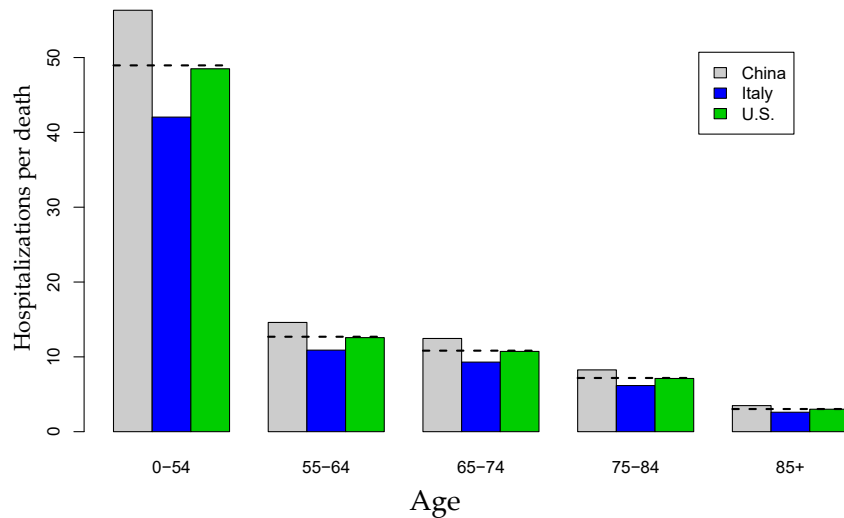
TheUpshot Jobless Claims Hit 3.3 Million in the Last Report. This Week’s Will Probably Be Worse.

By Quoc Trung Bui April 1, 2020



The Perils of 'Exit' Scenarios and Forecasting (IHME Model via U of Washington)

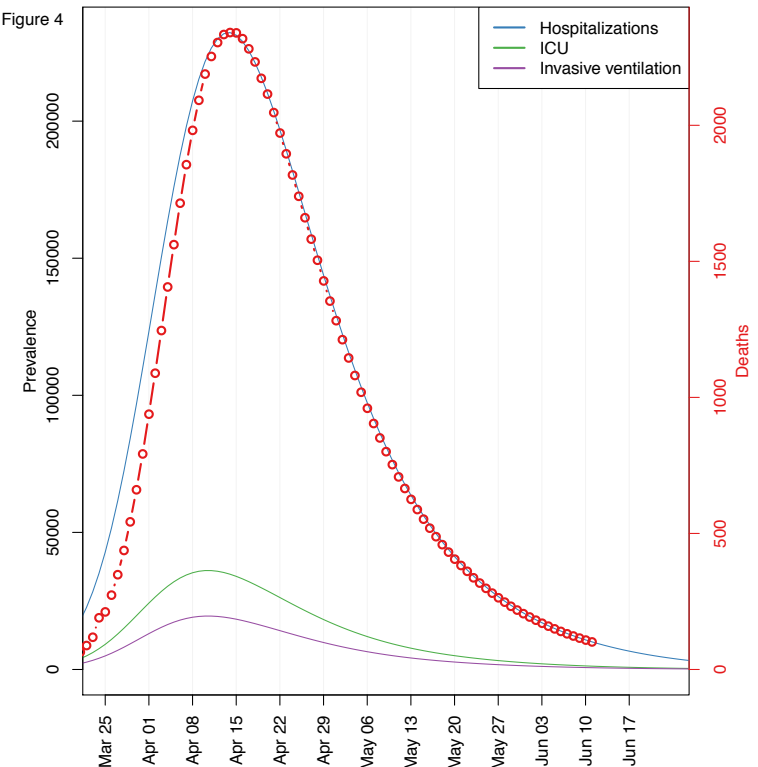
On March 28 – Predicted
April 15th as Peak
w/~2000 deaths per day



IHME Model Chris Murray



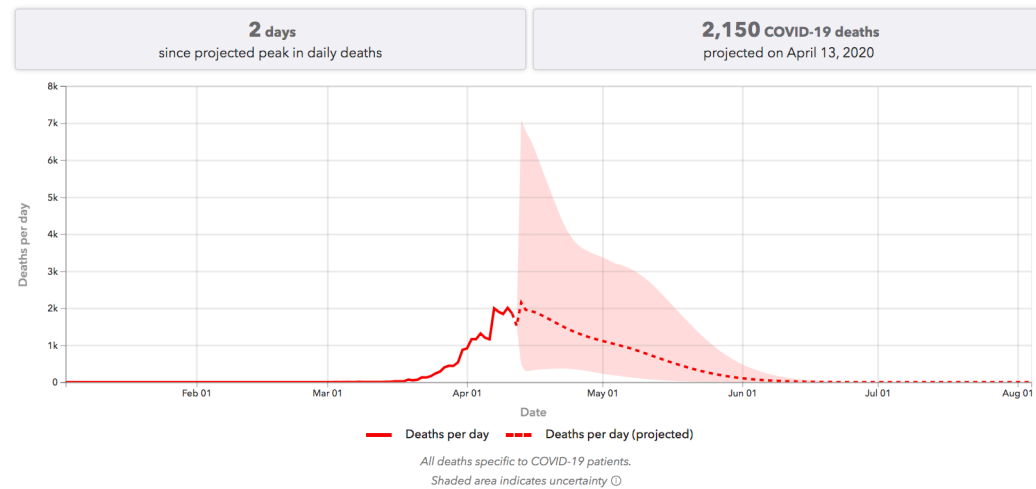
Figure 4



The Perils of 'Exit' Scenarios and Forecasting (IHME Model via U of Washington)

But now, the tail extends
and there is no uncertainty
after June 1 (!!)

IHME Model Chris Murray



The Perils of 'Exit' Scenarios and Forecasting (IHME Model via U of Washington)

But now, the tail extends and there is no uncertainty after June 1 (!!)

2. Functional Form for Covid-19

We considered several functional forms to model the death rate of the Covid-19 virus. Based both currently available data, the log rate starts slowly, increases quickly, and then flattens again as either social distancing or saturation goes into effect. This is the classic sigmoid shape. We first tried building the analysis using the sigmoidal function

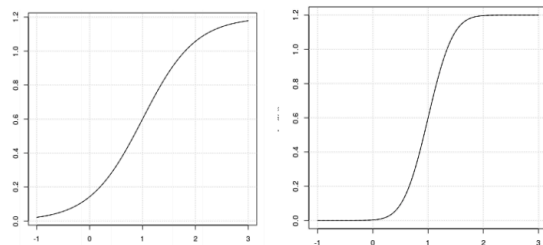


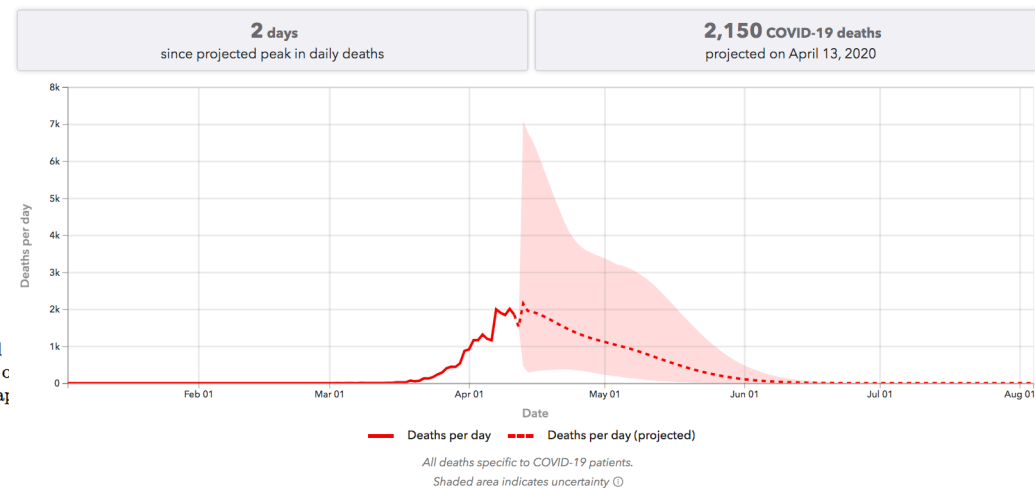
Figure 1. Exit function $\tilde{D}$ (left) and ERF function D (right). The ERF function fits the available Covid-19 data better than Exit.

$$\tilde{D}(t; \alpha, \beta, p) = \frac{p}{1 + \exp(-\alpha(t - \beta))}$$

where p controls the level, β the shift, and α the growth. We quickly discovered that the ERF error function provided a better fit to the data:

$$D(t; \alpha, \beta, p) = \frac{p}{2} (\Psi(\alpha(t - \beta))) = \frac{p}{2} \left(1 + \frac{2}{\sqrt{\pi}} \int_0^{\alpha(t - \beta)} \exp(-\tau^2) d\tau \right)$$

IHME Model Chris Murray



Buyer beware:

A curve fitting model based on an 'ERF' function should be used only for short-term predictions and its utility has already passed the expiration date.

Hint: the future is not that certain.

Thoughtful Exit Scenarios and Forecasting (INSERM/EPICX Lab)

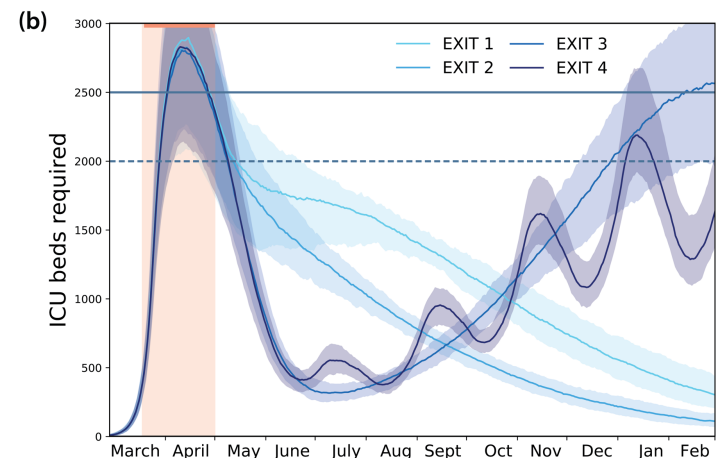
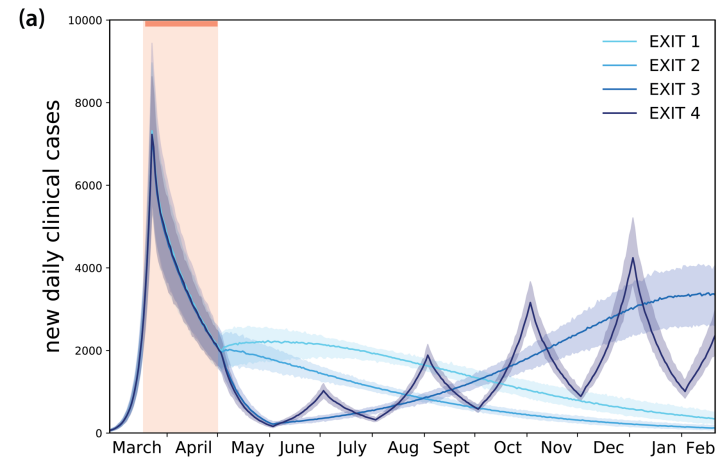
53

	March	April	May	June	July	Aug	Sept	Oct	Nov	Dec	Jan	Feb
LD(Apr)												
LD(May)												
LD(June)												
LD(Apr)+Strict												
LD(Apr)+Mod												
LD(Apr)+Mild												
LD(Apr)+SC,SI												
Exit 1												
Exit 2												
Exit 3												
Exit 4												
Exit 1 (1m after)												
Exit 2 (1m after)												
Exit 3 (1m after)												
Exit 4 (1m after)												

Figure 3. Scenarios (color code as in Table 1; CI refers to case isolation).

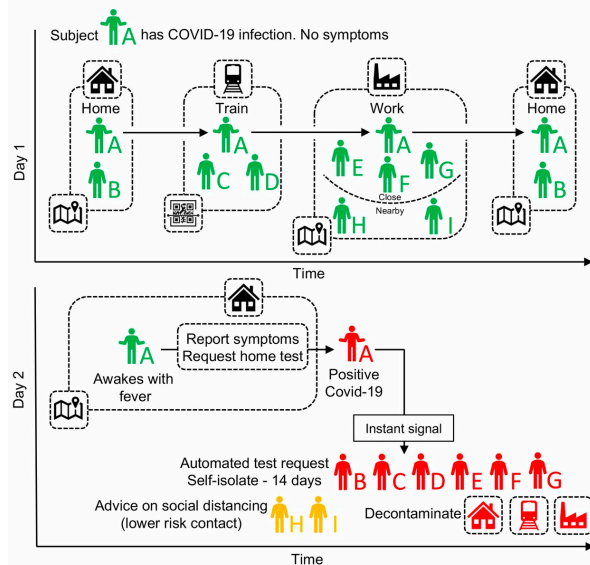
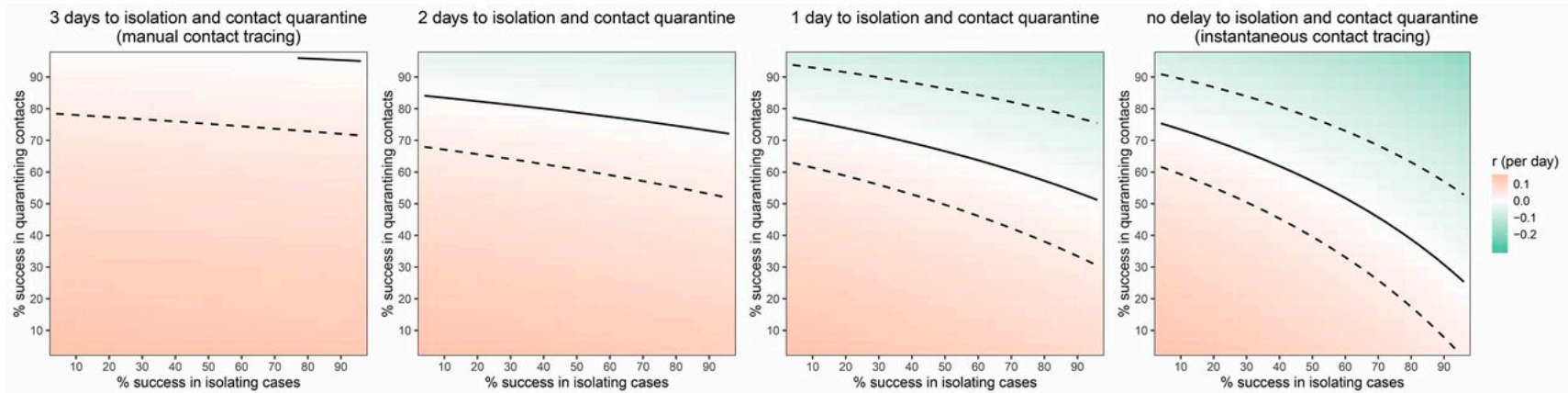
Take-away:

Sustained infection control is needed for ~1 year to prevent large-scale waves of new cases.



Testing and Tracing as a Means of Prevention not just Pathology

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Science

RESEARCH ARTICLES

Cite as: L. Ferretti *et al.*, *Science* 10.1126/science.abb6936 (2020).

Quantifying SARS-CoV-2 transmission suggests epidemic control with digital contact tracing

Luca Ferretti^{1*}, Chris Wymant^{1*}, Michelle Kendall¹, Lele Zhao¹, Anel Nurtay¹, Lucie Abeler-Dörner¹, Michael Parker², David Bonsall^{1,3†}, Christophe Fraser^{1,4†‡}

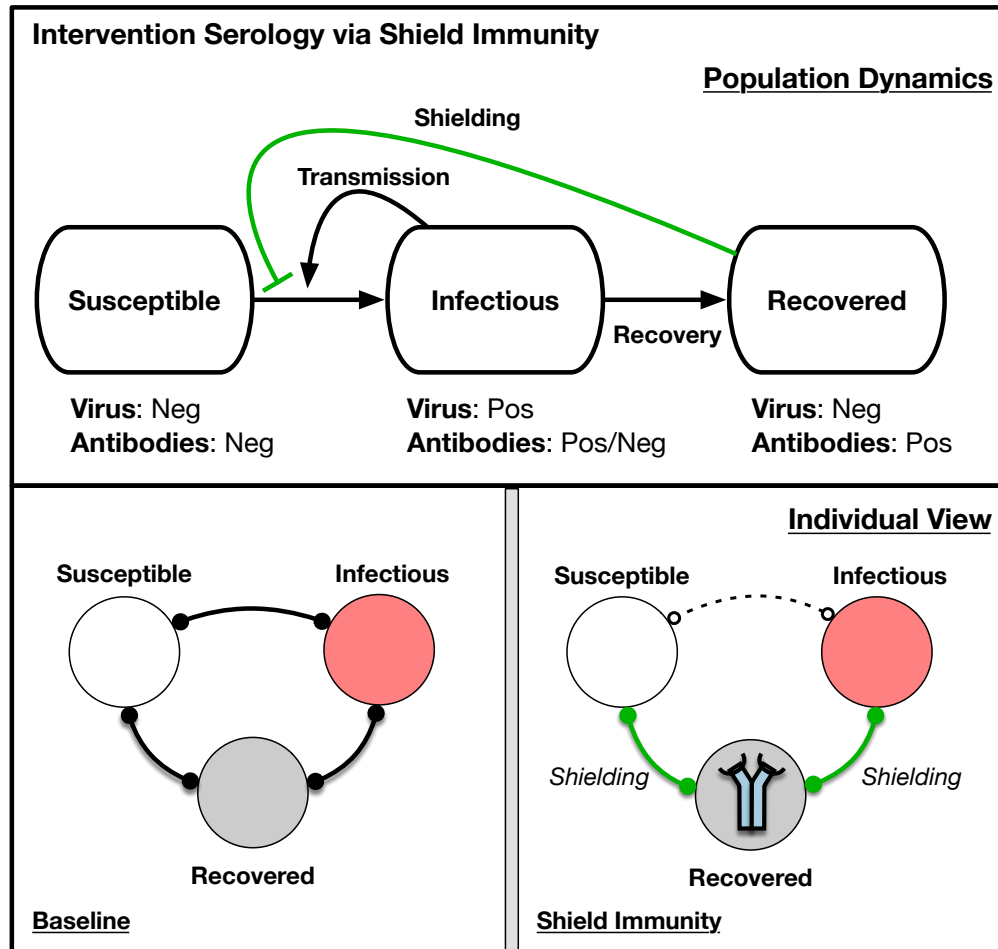
¹Big Data Institute, Li Ka Shing Centre for Health Information and Discovery, University of Oxford, Oxford, UK. ²Wellcome Centre for Ethics and the Humanities and Ethox Centre, University of Oxford, Oxford, UK. ³Oxford University NHS Trust, University of Oxford, Oxford, UK. ⁴Wellcome Centre for Human Genetics, University of Oxford, Oxford, UK.

**The Bulk of Interventions
Focus on Reducing New Transmission
by Closures and/or Tracing**

**but what about all those who have already
been sick and recovered...**

**could these infections (far from herd immunity)
help reduce collective risk?**

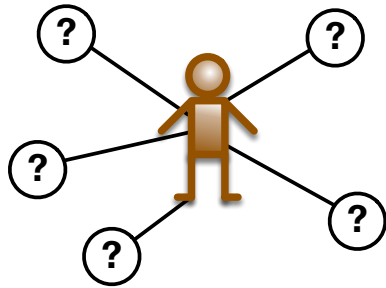
Our 'Shield Immunity' Proposal: Large-Scale Serological Testing to Reduce Transmission and Enable Economic Development



Weitz et al., medrxiv (in press, to be announced soon)

What is 'Shield Immunity'?

**Baseline: Unknown Infection Status
of Others in the Community**

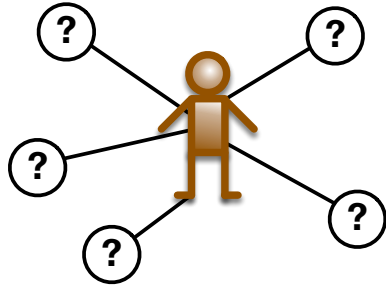


Disease status:

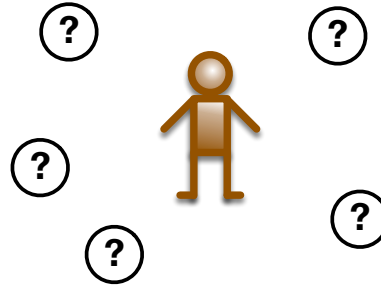
⓪ Unknown

What is 'Shield Immunity'?

**Baseline: Unknown Infection Status
of Others in the Community**



**Extreme Social Distancing:
Limited/No Interactions**

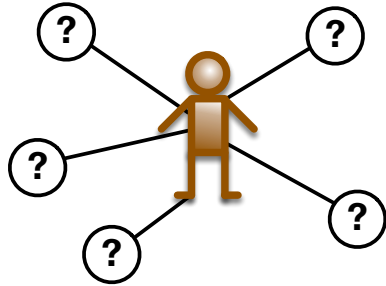


Disease status:

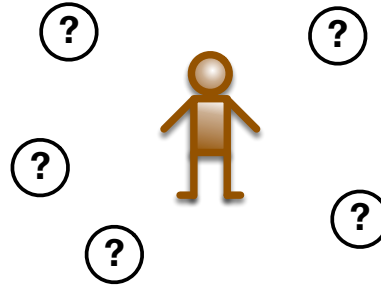
⓪ Unknown

What is 'Shield Immunity'?

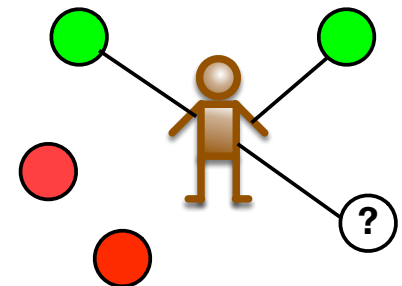
Baseline: Unknown Infection Status of Others in the Community



Extreme Social Distancing: Limited/No Interactions



Immune Shielding: Preferentially Interact with Recovered Individuals



Disease status:



Unknown

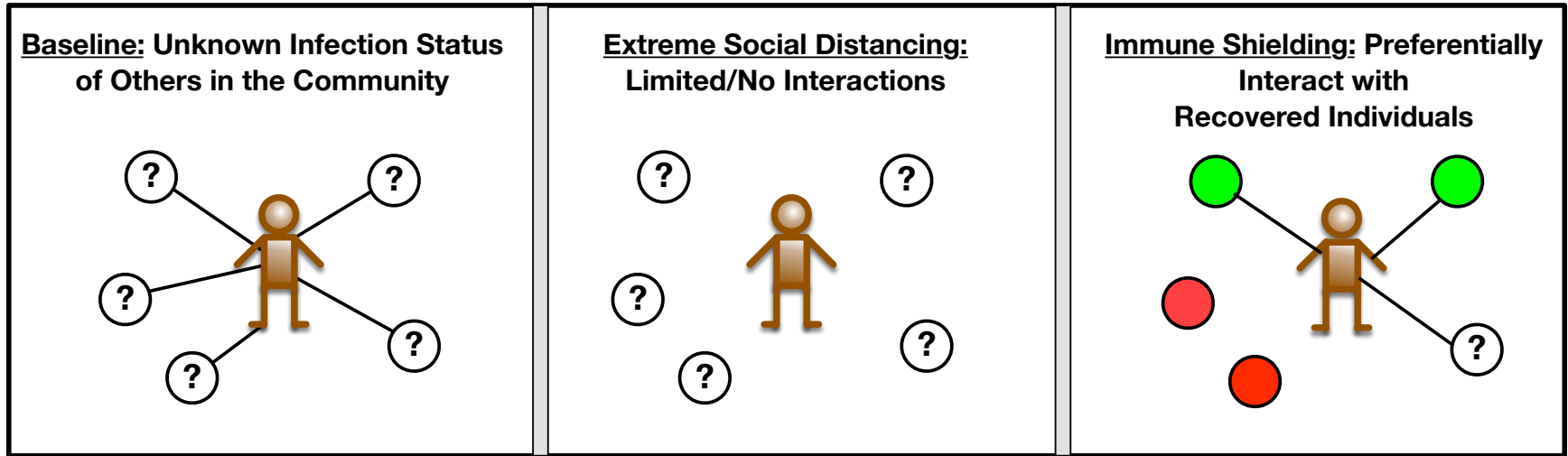


Infectious



Recovered

What is 'Shield Immunity'?



Disease status:



Unknown



Infectious



Recovered

Take-away: the scale and type of testing matters



Infectious

PCR provides a snapshot,
i.e., 'are you shedding virus now'?



Recovered

Serological testing for antibodies provides a history,
i.e., 'have you been infected recently or in the past'?
Recovered implies immune (duration still unknown).

Shield Immunity Dynamics

SIR model w/shielding

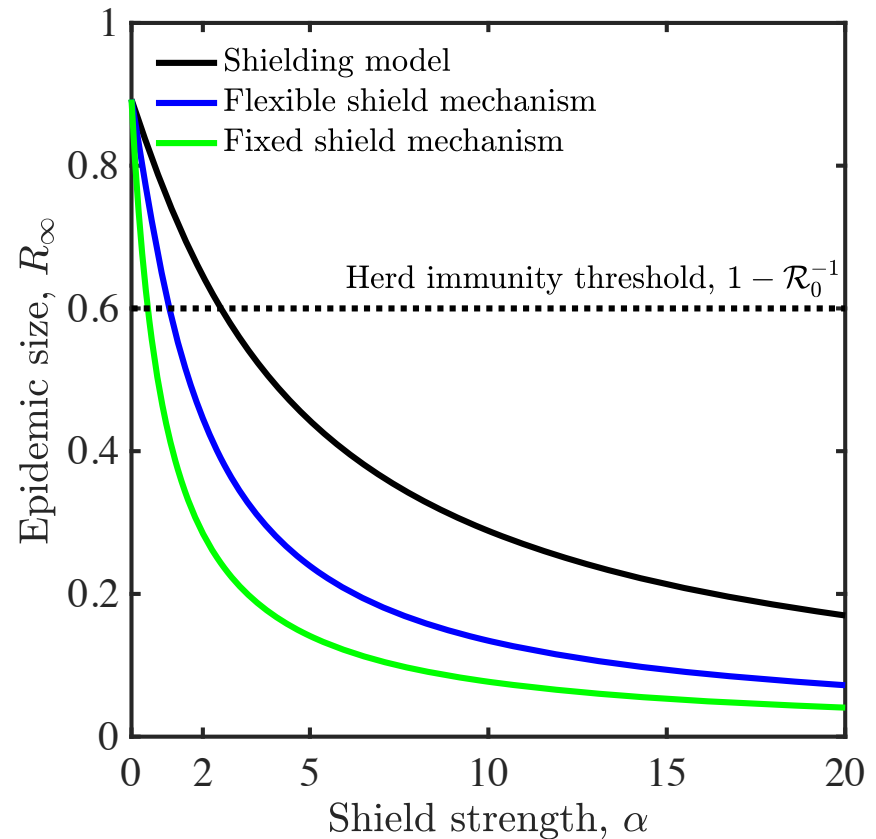
Recovered individuals have preferential interactions relative to other individuals.

This dilutes the susceptible fraction at levels well below the herd immunity threshold.

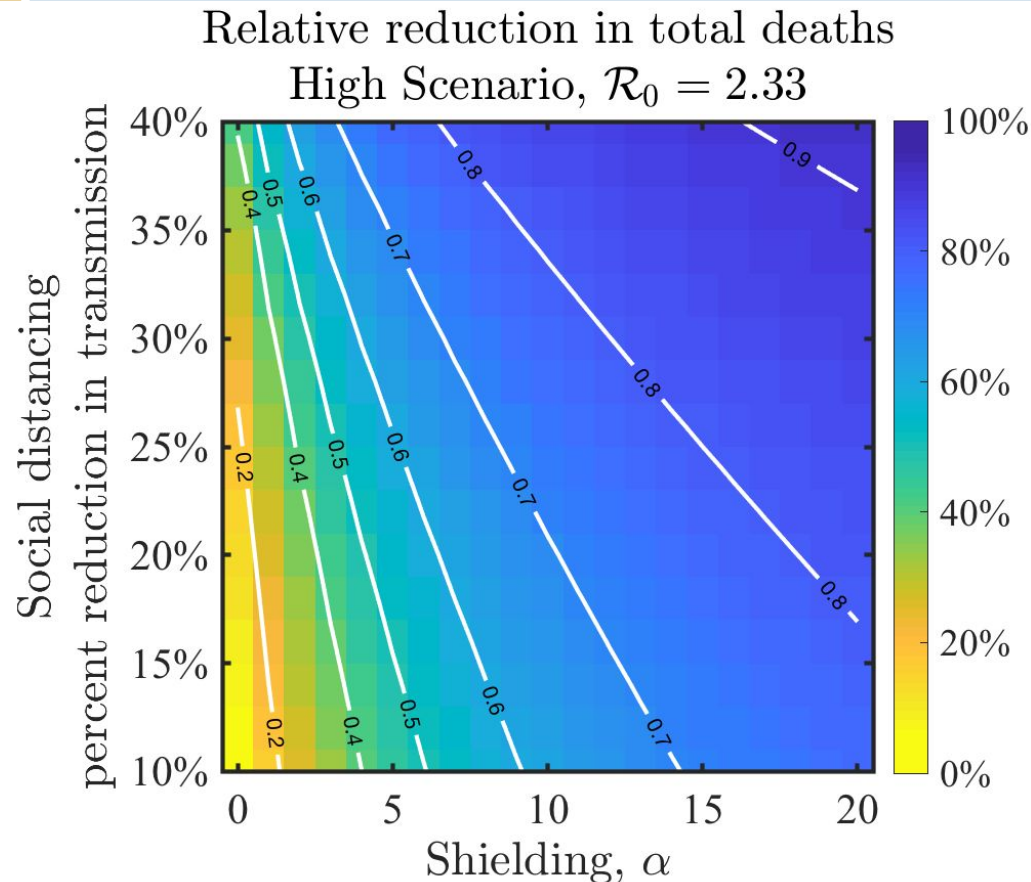
$$\dot{S} = -\beta \frac{SI}{1 + \alpha R}$$

$$\dot{I} = \beta \frac{SI}{1 + \alpha R} - \gamma I$$

$$\dot{R} = \gamma I$$



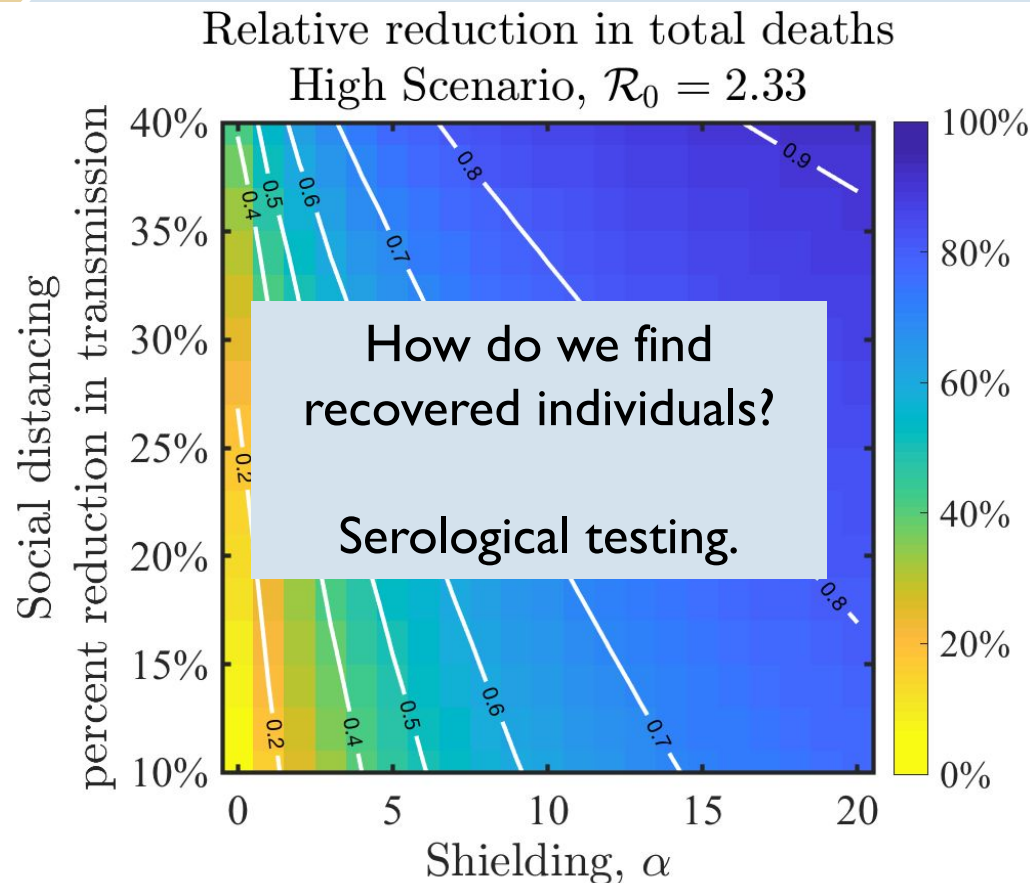
What could we do if we could identify recovered individuals?



Recovered individuals could elevate their interactions relative to other individuals – we call this ‘shield strength’.

Combining (less intense) social distancing and shielding could have significant population-wide benefits.

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Closing Thoughts

Part I – Dynamical Foundations of Epidemics

COVID-19 has epidemiological features of high R_0 , relatively fast spread, elevated fatality rates, and asymptomatic transmission that increase its epidemic potential.

Closing Thoughts

Part 1 – Dynamical Foundations of Epidemics

COVID-19 has epidemiological features of high R_0 , relatively fast spread, elevated fatality rates, and asymptomatic transmission that increase its epidemic potential.

Part 2 – Dynamics and Control

Interventions have made an enormous impact (averting tens/hundreds of thousands of fatalities). But, even if a peak has been averted, we remain immunologically naive.

Closing Thoughts

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Part 3 – Long-term strategies

Testing (PCR and serology) & contact tracing are needed at scale! Strategic and coordinated responses – we have a long road ahead, we'll be better off working collectively.

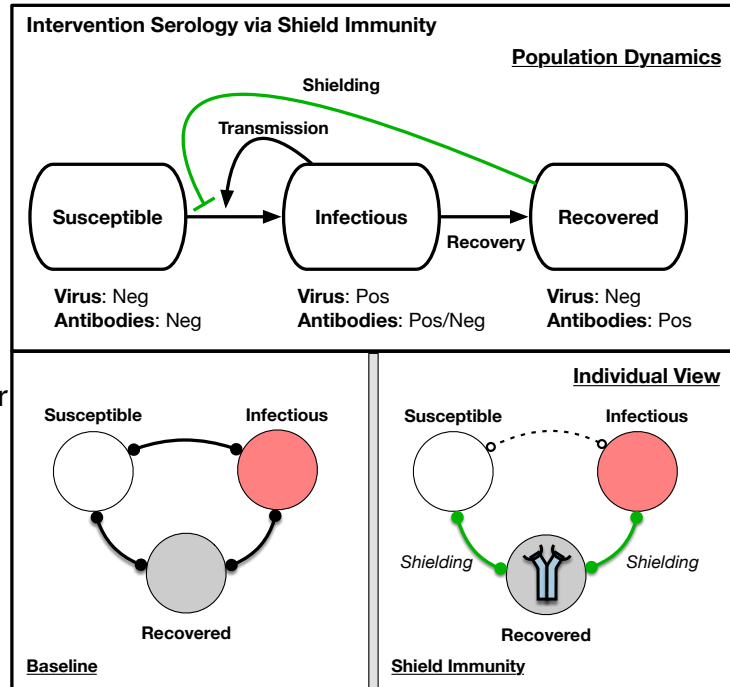


Georgia Tech Team

Ashley Coenen
Dr. Stephen Beckett
Dr. David Demory
Marian Dominguez-Mirazo
Dr. Joey Leung
Guanlin Li
Andreea Magalie
Daniel Muratore
Rogelio Rodriguez-Gonzalez
Dr. Adriana Sanz
Shashwat Shivam
Conan Zhao

Collaborators

Clio Andris, GT
Daniel Cornforth, GT
Jonathan Dushoff, McMaster
Ceyhun Eksin, Texas A&M
Benjamin Lopman, Emory
Alicia Kraal, Emory
Kristen Nelson, Emory
Sang Woo Park, Princeton
Yorai Wardi, GT



Code:

https://github.com/WeitzGroup/covid_shield_immunity

Preprint (now in press, to be announced soon)

<https://www.medrxiv.org/content/10.1101/2020.04.01.20049767v1>

Tweet thread:

<https://twitter.com/joshuasweitz/status/1245071163744645121?s=20>

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NSF, ARO, NIH, and the Simons Foundation