

A Complex Systems Theory of Disease

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Juan Genoves

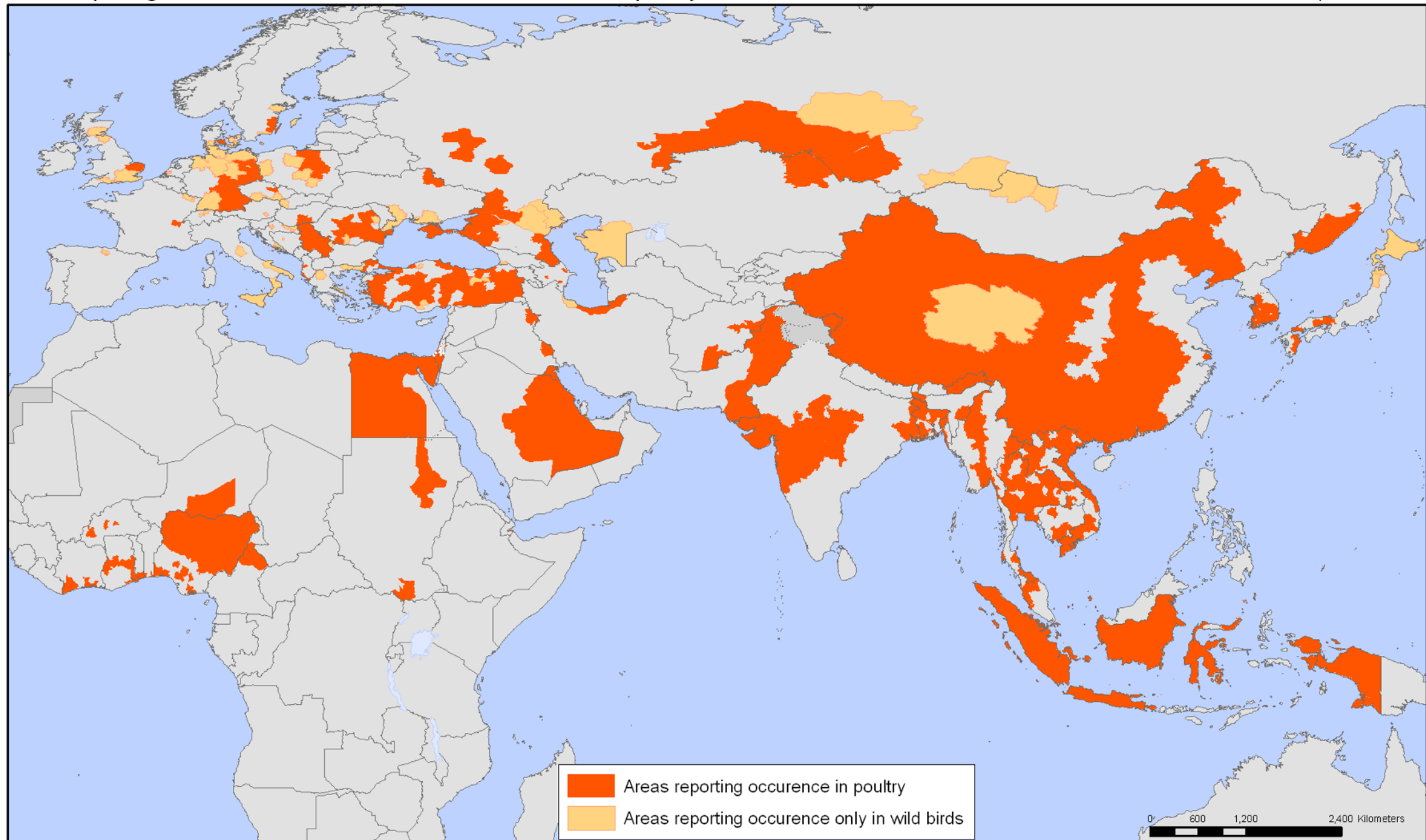
Where will the next pandemic emerge?



Where will the next **influenza** pandemic emerge?

Areas reporting confirmed occurrence of H5N1 avian influenza in poultry and wild birds since 2003

Status as of 14 November 2008
Latest available update



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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 lines on maps represent approximate border lines for which there may not yet be full agreement.

Data Source: World Organisation for Animal Health (OIE) and national governments

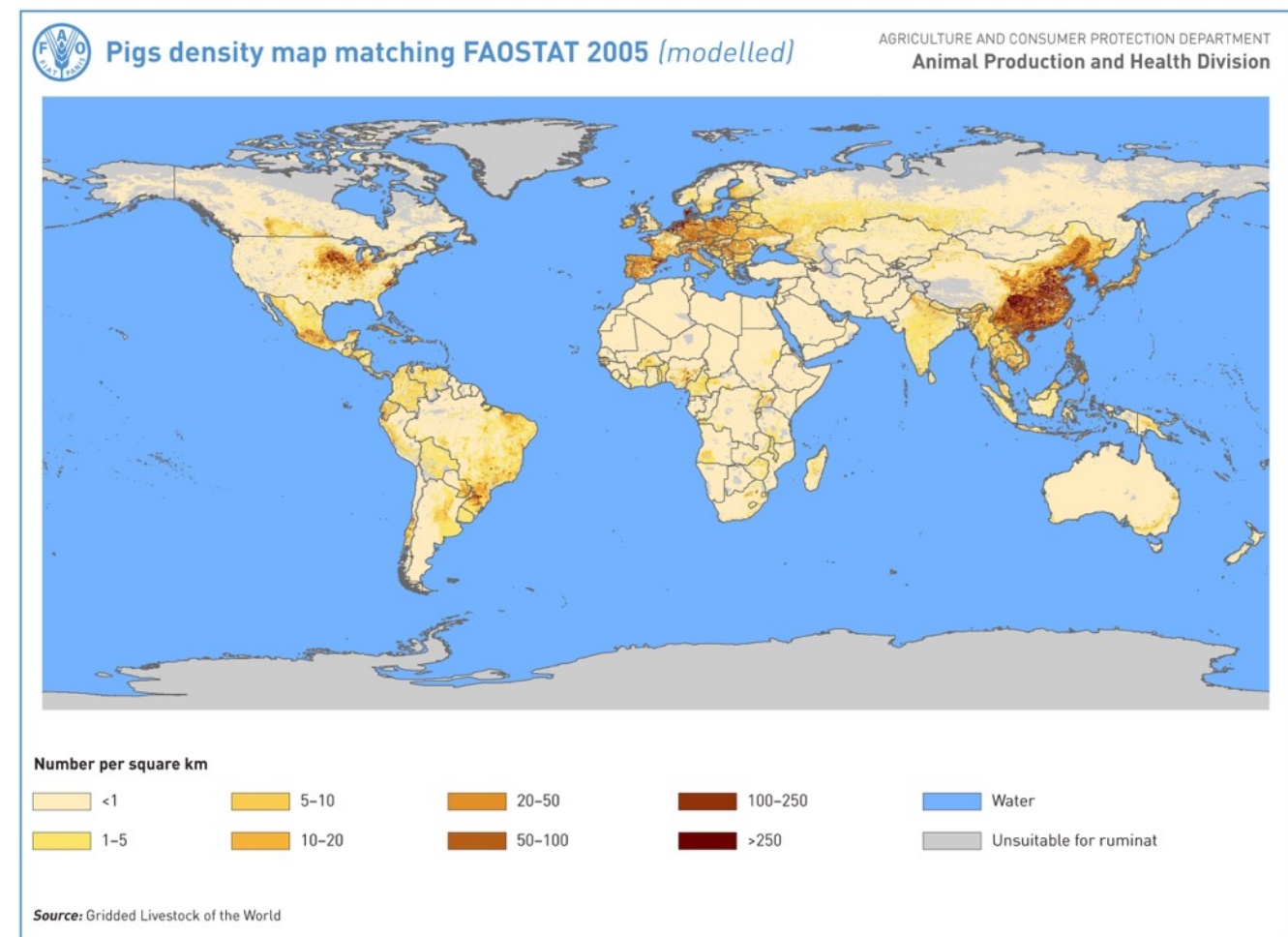
Map Production: Public Health Information and Geographic Information Systems (GIS), World Health Organization

What factors are important for pandemic flu?

Air travel



Density of pig farms

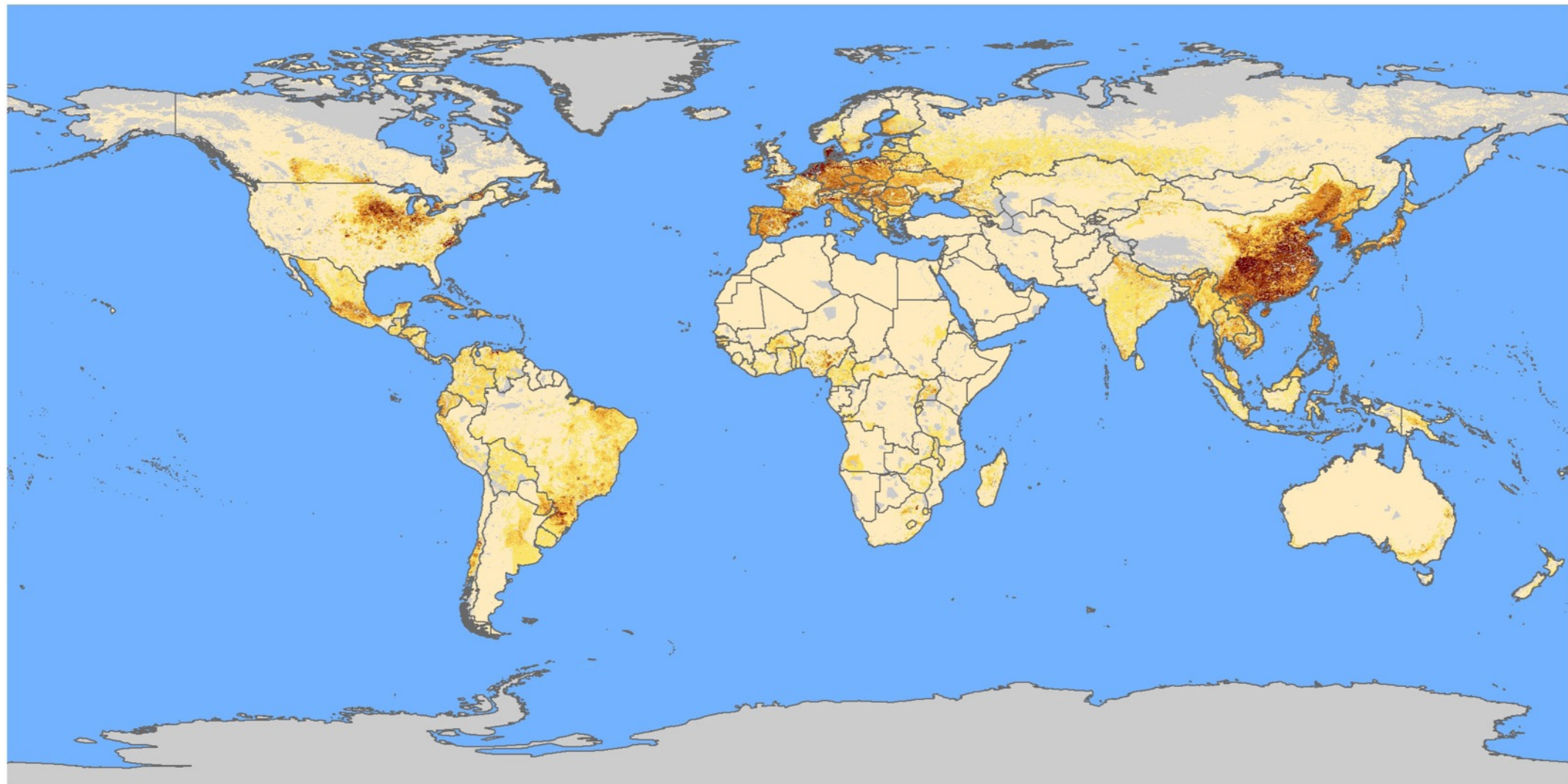


Worldwide pig density



Pigs density map matching FAOSTAT 2005 (*modelled*)

AGRICULTURE AND CONSUMER PROTECTION DEPARTMENT
Animal Production and Health Division



Number per square km

<1

5-10

20-50

100-250

Water

1-5

10-20

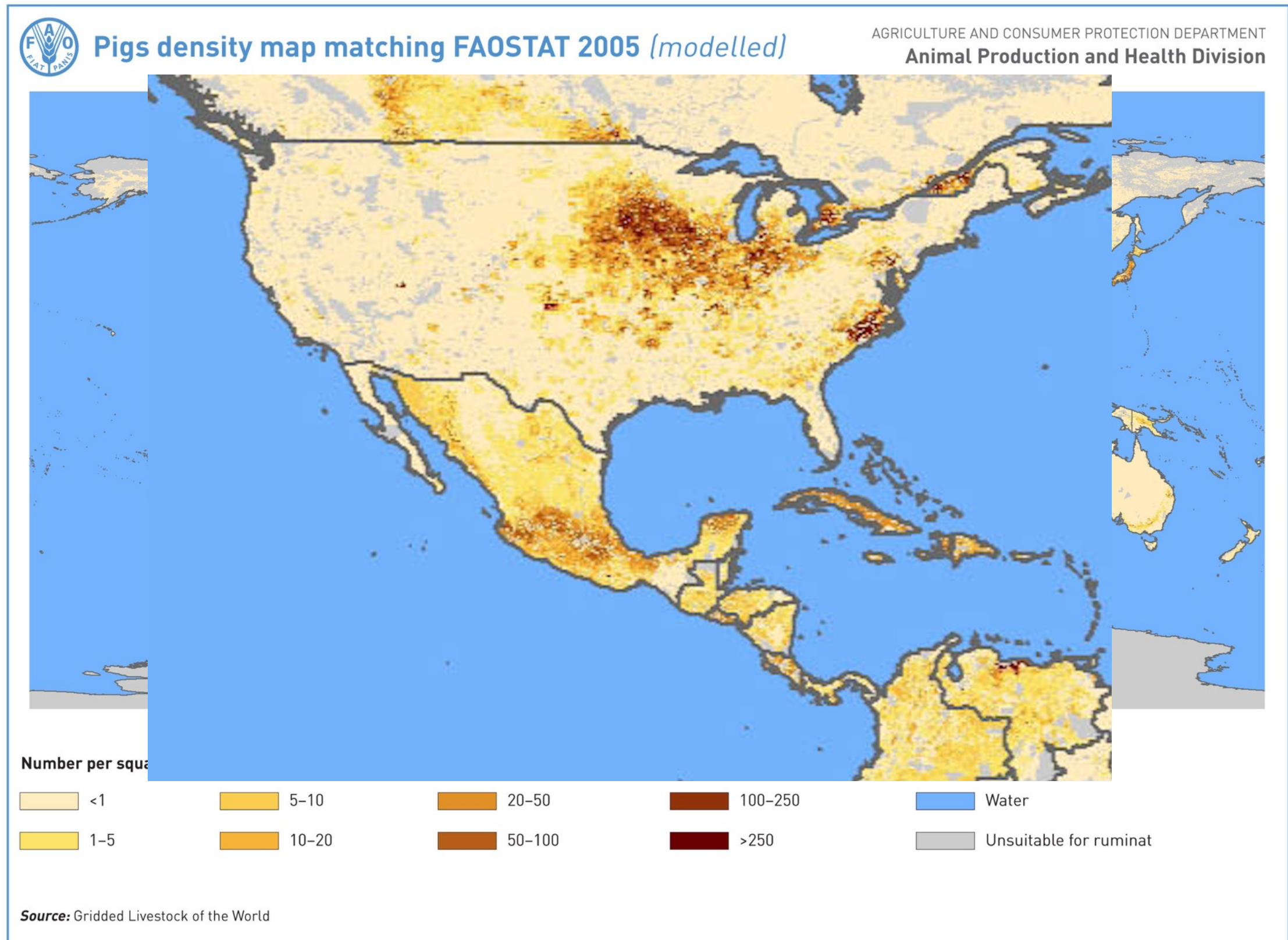
50-100

>250

Unsuitable for ruminat

Source: Gridded Livestock of the World

Worldwide pig density



and it happened again with Ebola



and it happened again with Ebola



TABLE I. — Age distribution of persons positive for either Lassa (LAS), Ebola (EBO) or Marburg (MAR) virus antibodies.

Age (years)	Nb tested	LAS-positive (prevalence %)	EBO-positive (prevalence %)	MAR-positive (prevalence %)
0-9	49	5 (10 %)	2 (4 %)	0
10-19	68	11 (16 %)	5 (7 %)	0
20-29	108	21 (19 %)	6 (6 %)	1
30-39	94	16 (17 %)	5 (5 %)	1
40-59	88	9 (10 %)	6 (7 %)	1
60 plus	26	5 (16 %)	2 (8 %)	2
Total	433	67 (16 %)	26 (6 %)	5 (1 %)



and it may happened again with Chagas



Brumpt et al. 1912
Salazar et al. 2015

and it may happened again with Chagas

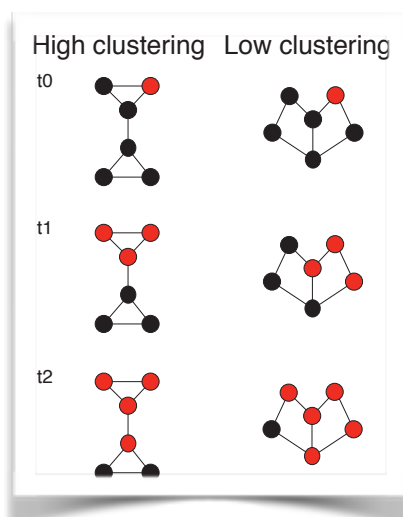
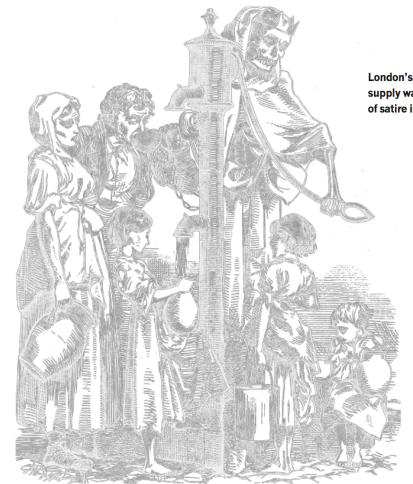


Brumpt et al. 1912
Salazar et al. 2015

Modeling infectious diseases

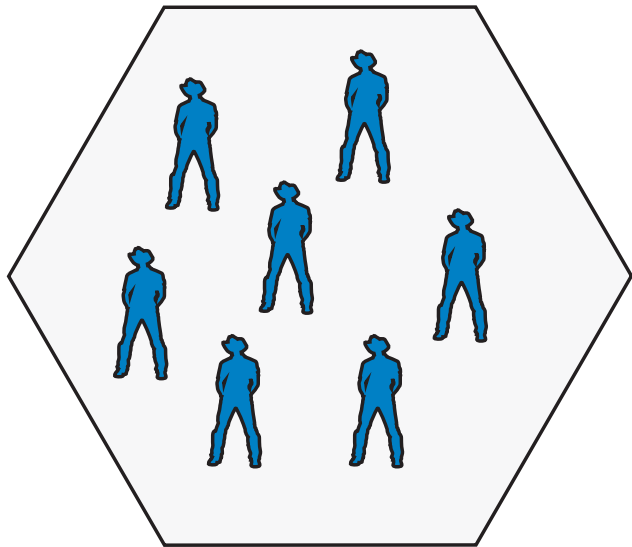


Inequality & disease



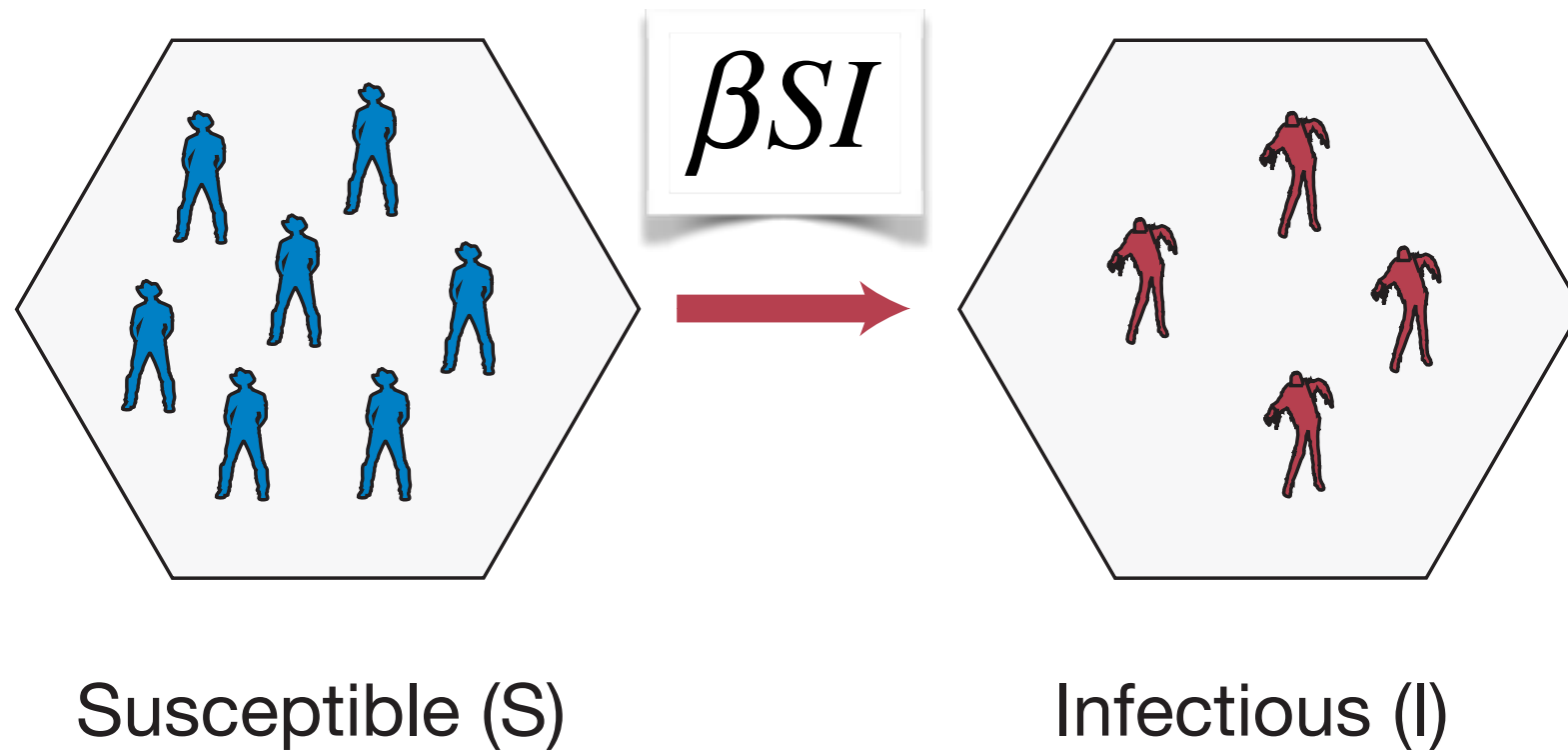
Social clustering & Ebola

Compartmental models

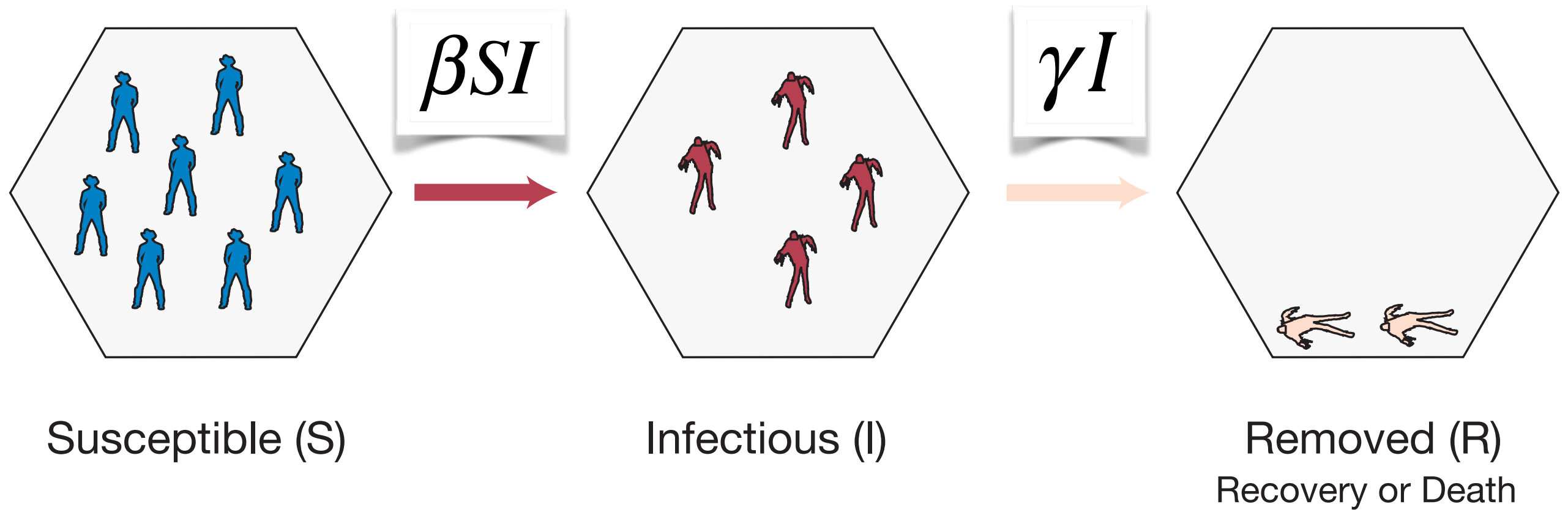


Susceptible (S)

Compartmental models - Mass Action Assumption



Compartmental models



Lions, Tigers, and Boxes ... oh my

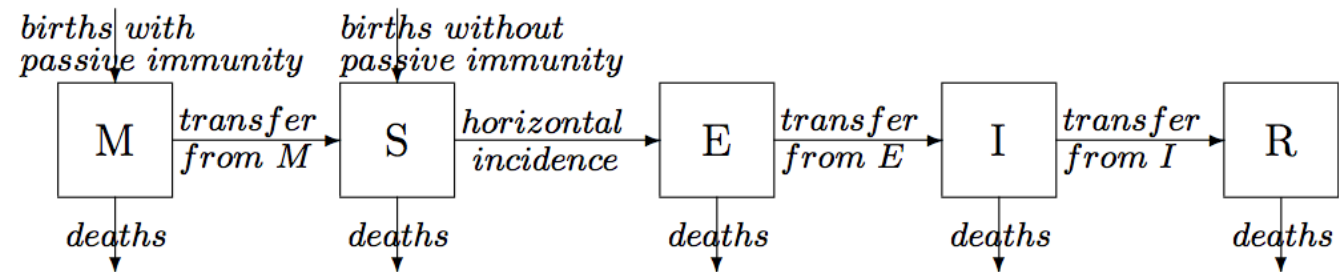
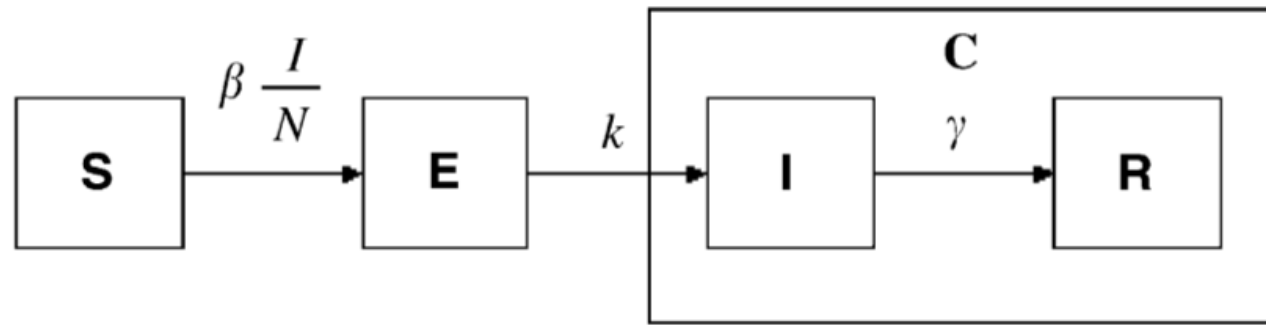


Fig. 1 The general transfer diagram for the MSEIR model with the passively immune class M , the susceptible class S , the exposed class E , the infective class I , and the recovered class R .

$$S'(t) = \mu \cdot (1 - wP - aP) - \beta[I_s(t) + I_a(t)]S(t) - \nu S(t) \quad (1)$$

$$I'_s(t) = \beta\sigma[I_s(t) + I_a(t)]S(t) - \gamma_s I_s(t) - \nu I_s(t) \quad (2)$$

$$I'_a(t) = \beta(1 - \sigma)[I_s(t) + I_a(t)]S(t) + \beta[I_s(t) + I_a(t)]V(t) - \gamma_a I_a(t) - \nu I_a(t) \quad (3)$$

$$V'(t) = \mu \cdot aP - \beta[I_s(t) + I_a(t)]V(t) - \nu V(t) \quad (4)$$

$$R'(t) = \mu \cdot wP + \gamma_s I_s(t) + \gamma_a I_a(t) - \nu R(t) \quad (5)$$

Lions, Tigers, and Boxes ... oh my

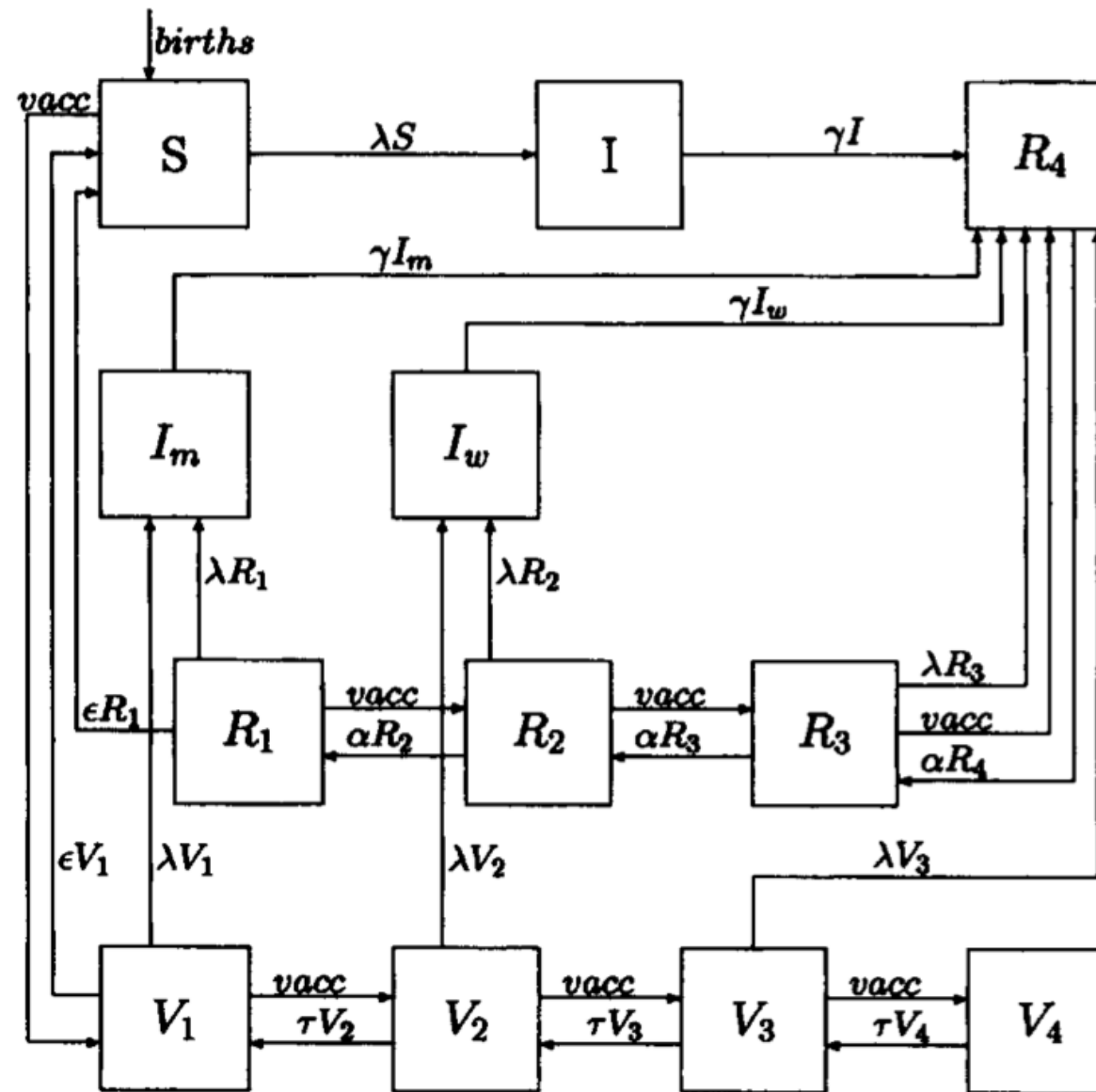
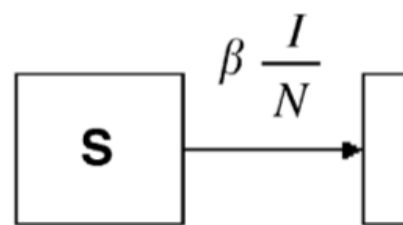
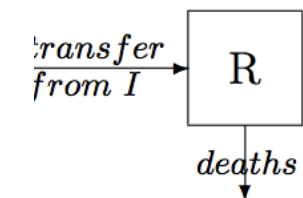


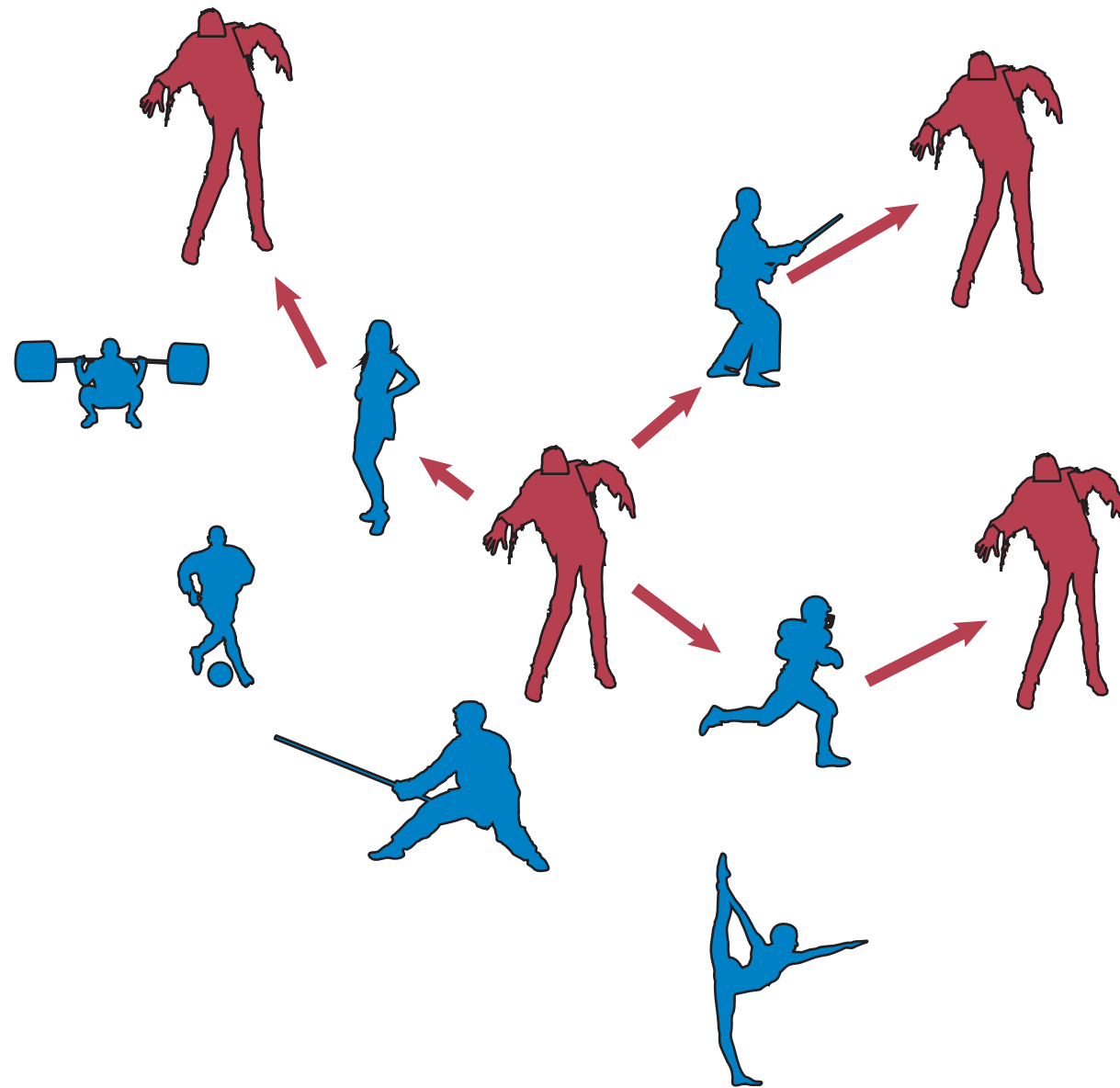
FIG. 3. Transfer diagram for the pertussis model with vaccination.



by immune class M , the
recovered class R .

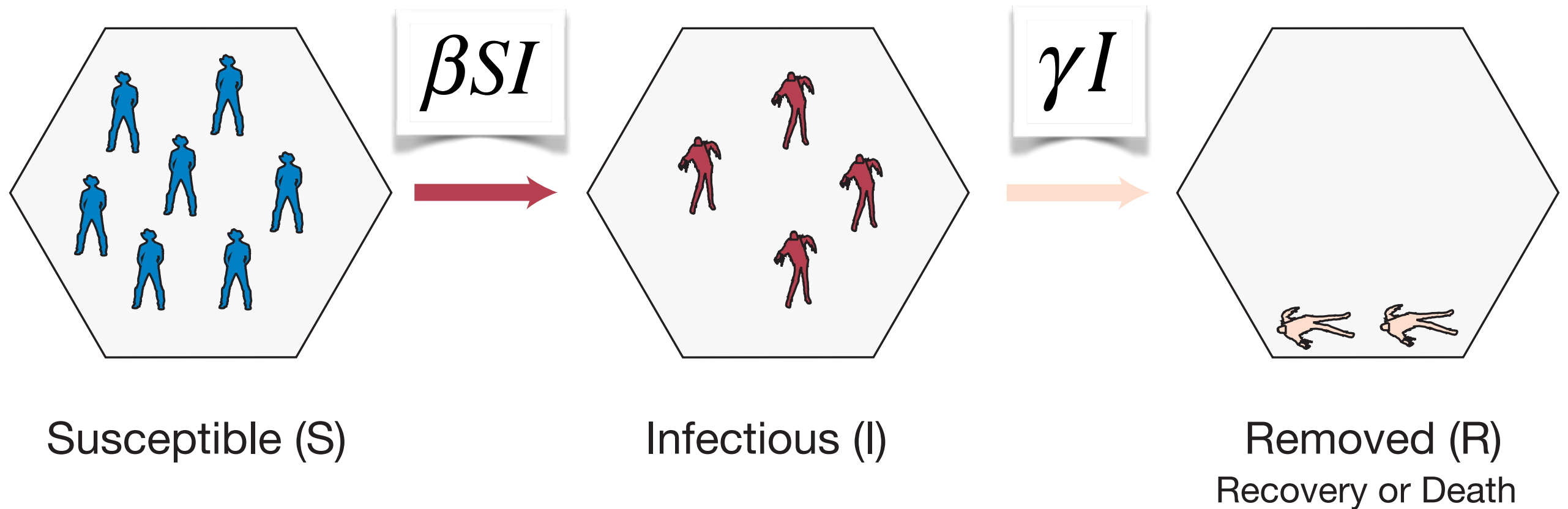
$$\begin{aligned}
 S'(t) &= \mu \cdot (1 - \beta \frac{I}{N}) \\
 I'_s(t) &= \beta \sigma [I_s(t) - I_a(t)] \\
 I'_a(t) &= \beta (1 - \sigma) [I_s(t) - I_a(t)] \\
 V'(t) &= \mu \cdot aP \\
 R'(t) &= \mu \cdot wP
 \end{aligned}$$

Reproduction Number



Expected number of secondary cases
in the beginning of an outbreak

Compartmental models



$$R_0 = \frac{\text{Infection rate}}{\text{Recovery rate}} = \frac{\beta S}{\gamma}$$

Reproduction Numbers

Disease	R_0
Measles & Whooping Cough	5 - 18
Chicken Pox	7 - 12
Polio	5 - 7
Smallpox	1.5 - 20+
Seasonal flu	1.1 - 1.5
Ebola	1.1 - 3

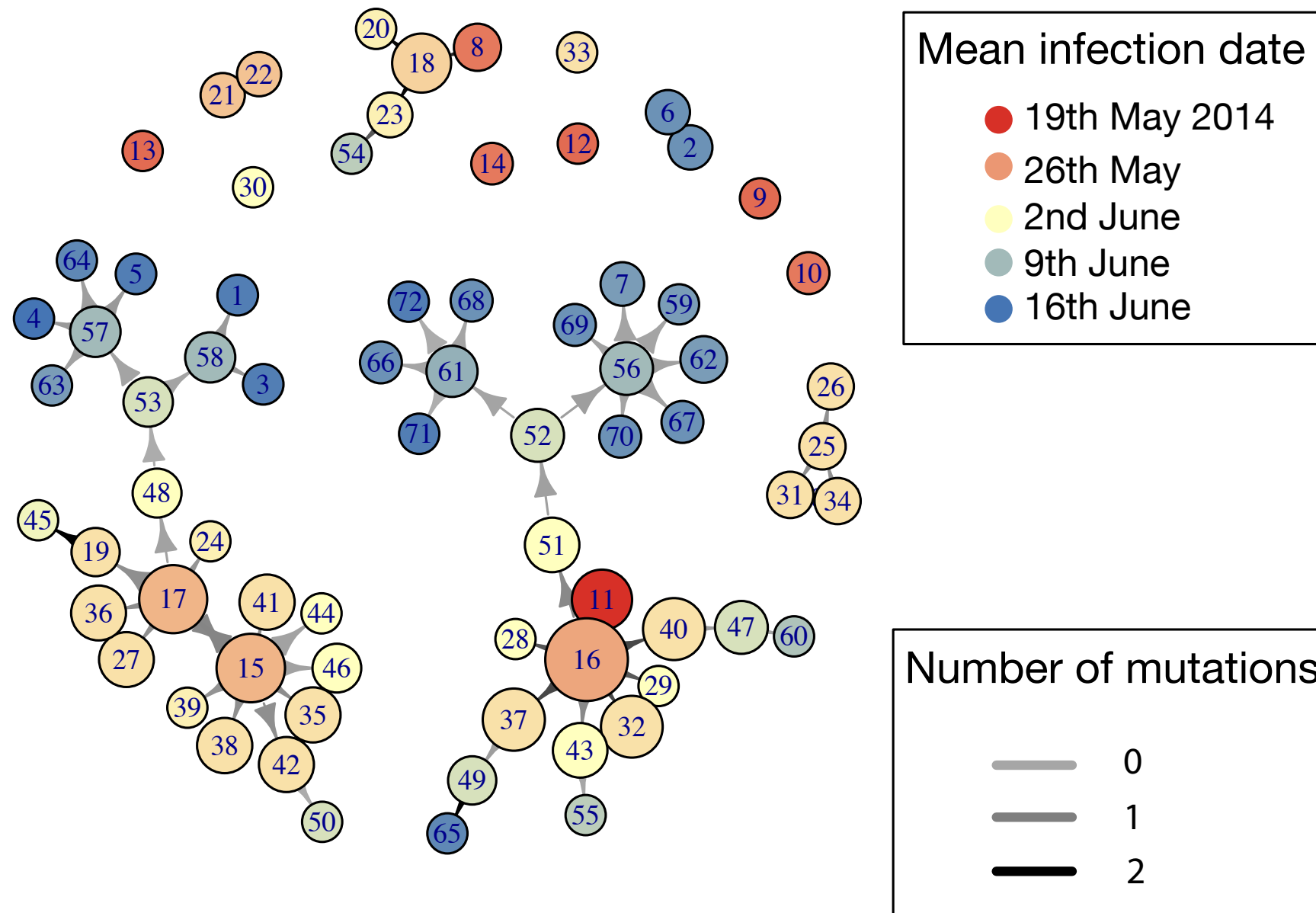
Problems with the Reproduction Number

Ebola virus genomic data



Modified from Gire et al. 2014

Reconstructed transmission network



Posterior number of secondary infections

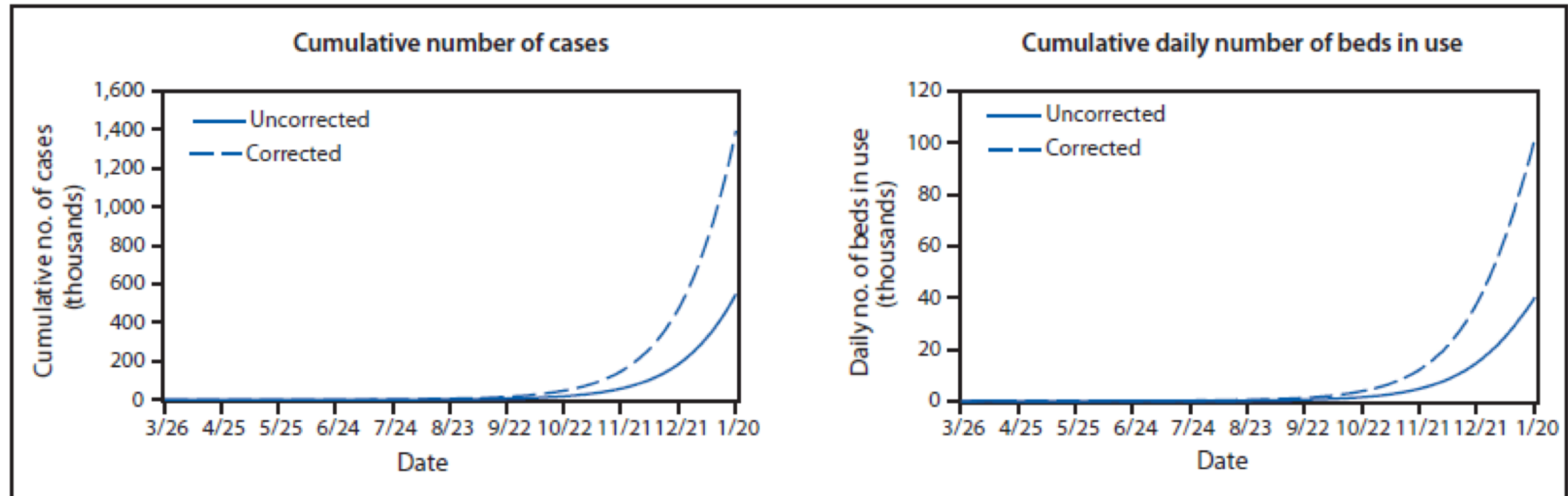


Problems with the Reproduction Number

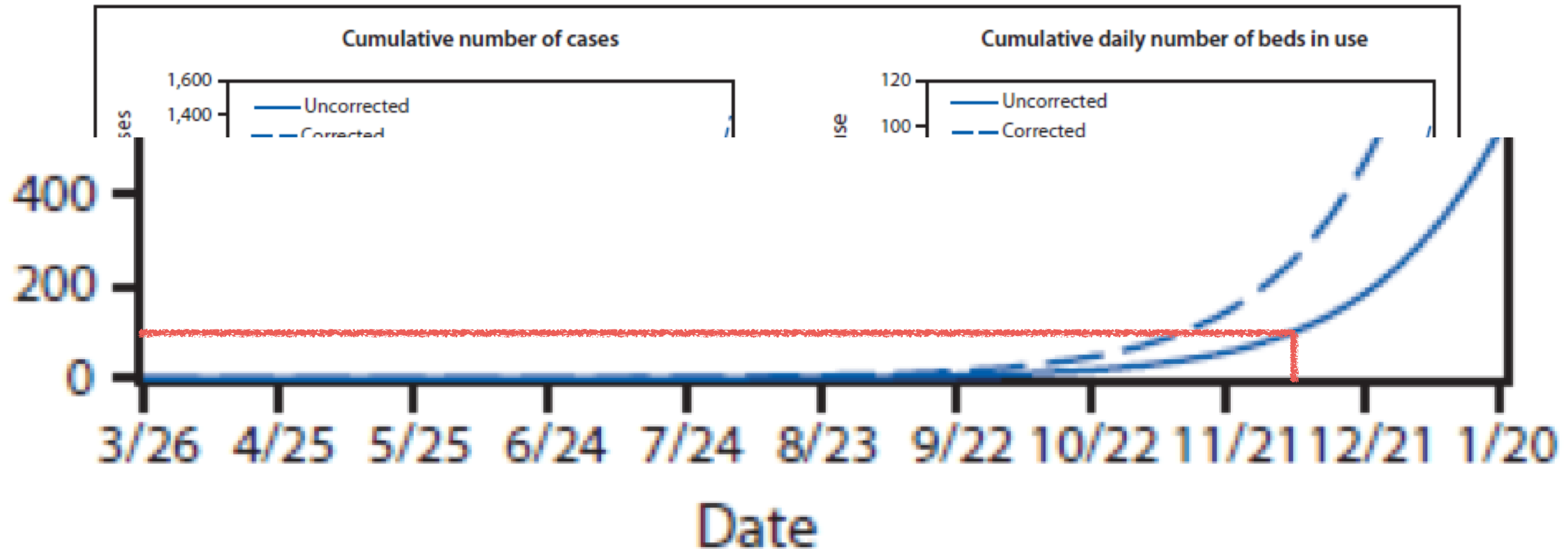
Contact Patterns Vary



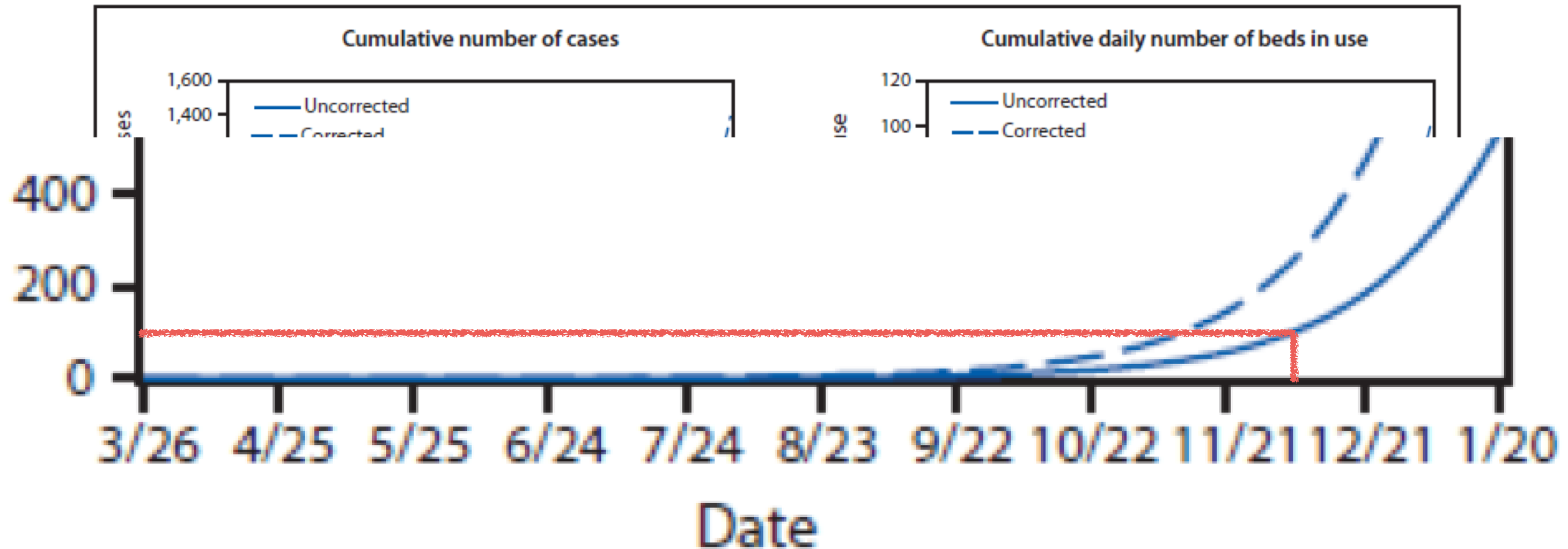
Ebola Predictions from mass-action models



Ebola Predictions from mass-action models

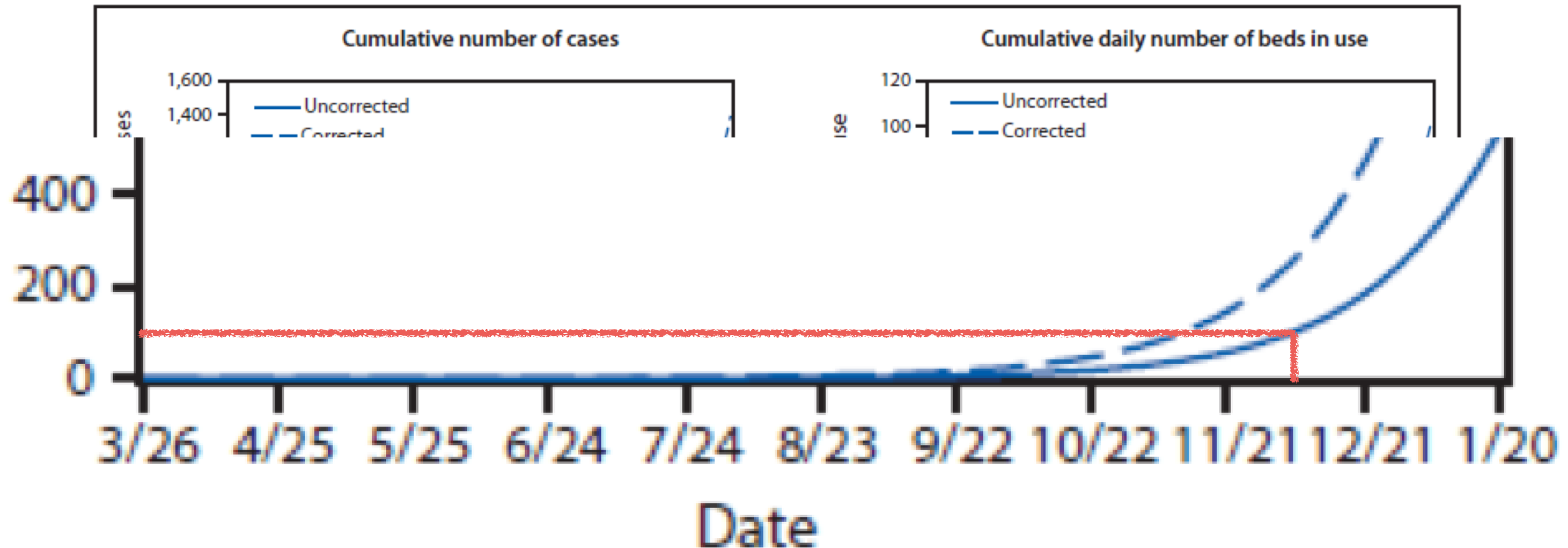


Ebola Predictions from mass-action models



Estimated - 100,000

Ebola Predictions from mass-action models



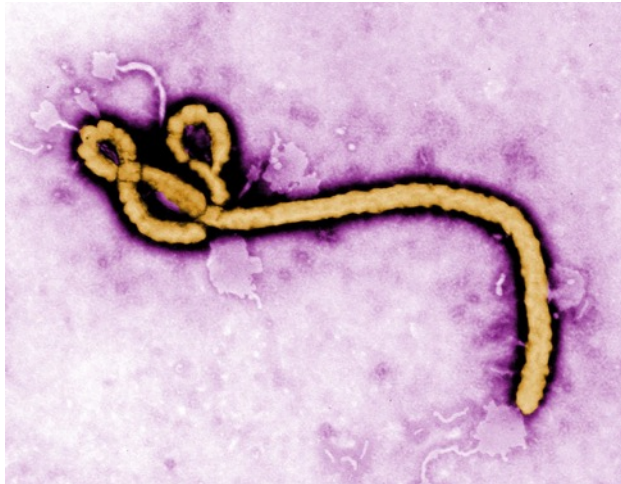
Estimated - 100,000

Observed - 10,000

Why do these problems exist?

Why do these problems exist?

1.



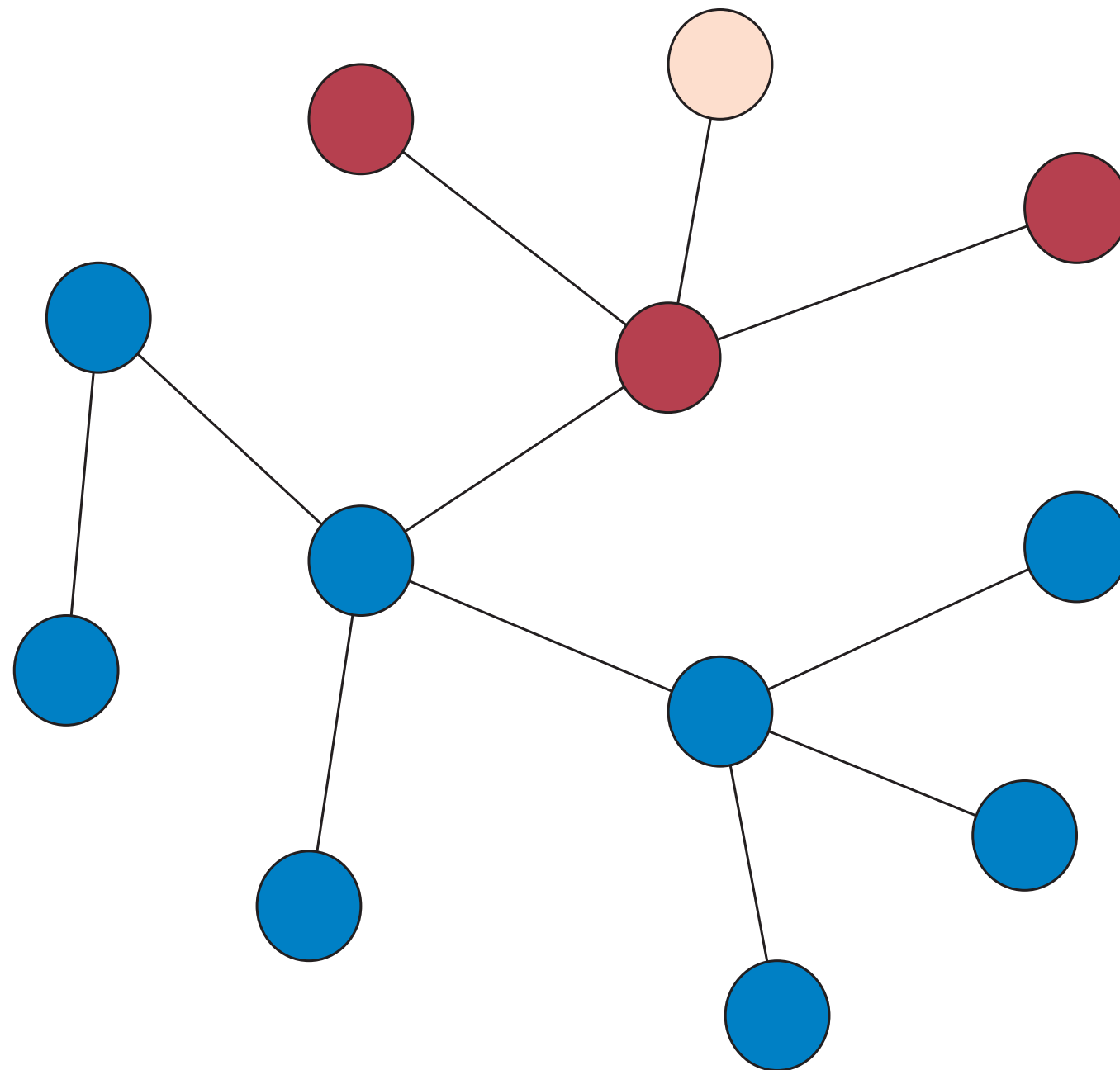
Intrinsic properties of the pathogen

2.



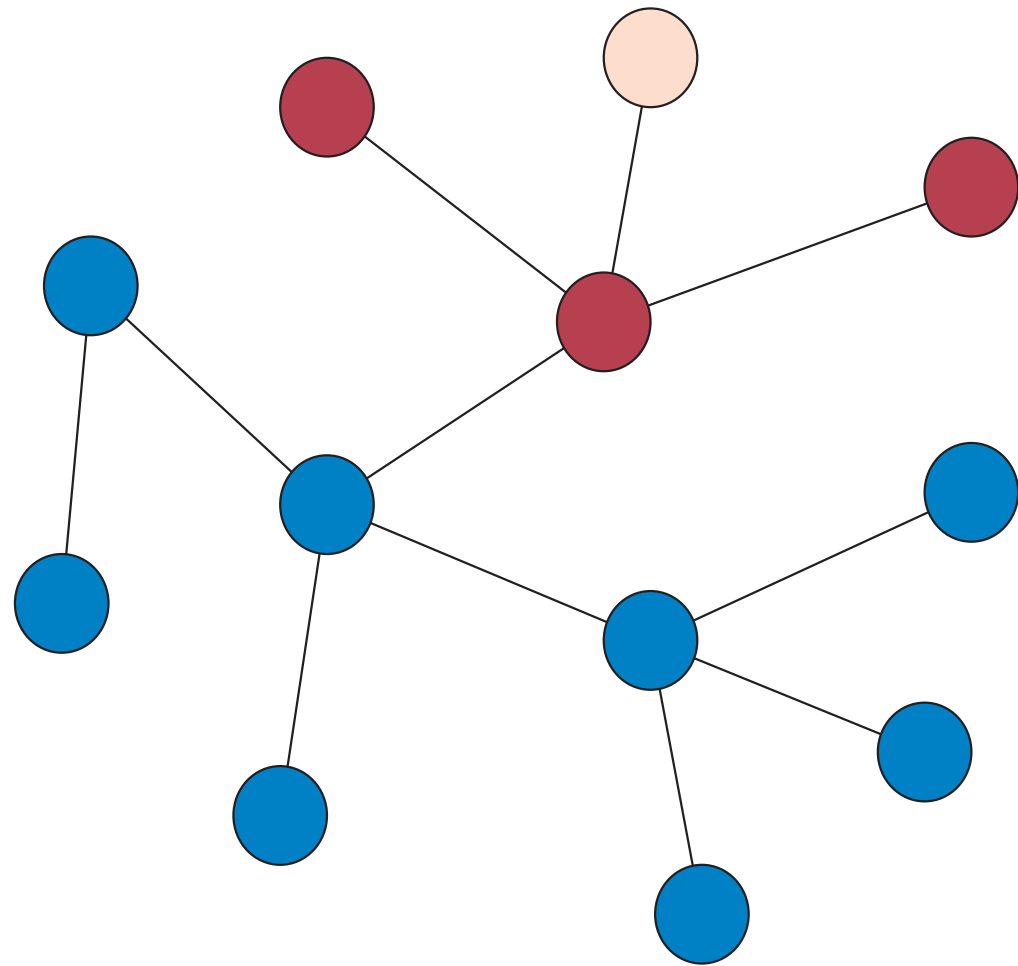
Contact patterns of the host

Network Epidemiology



Network Epidemiology

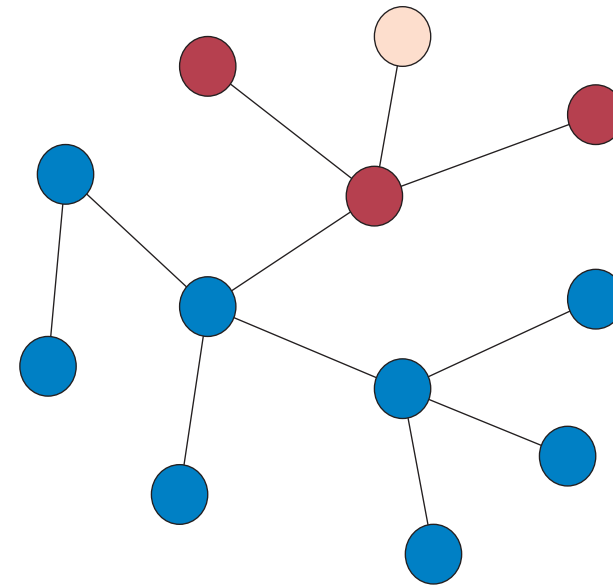
$$R_0 = T \left(\frac{\langle K^2 \rangle - \langle K \rangle}{\langle K \rangle} \right)$$



Network Epidemiology Vs.

Standard Calculation

$$R_0 = T \left(\frac{\langle K^2 \rangle - \langle K \rangle}{\langle K \rangle} \right)$$



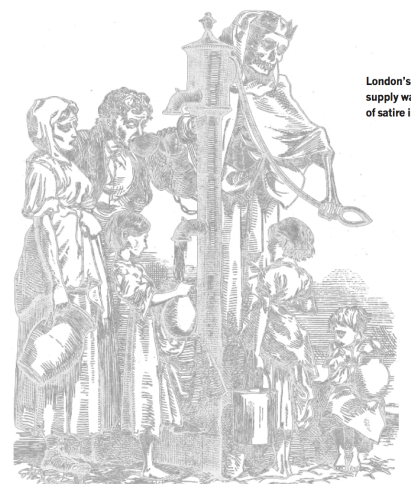
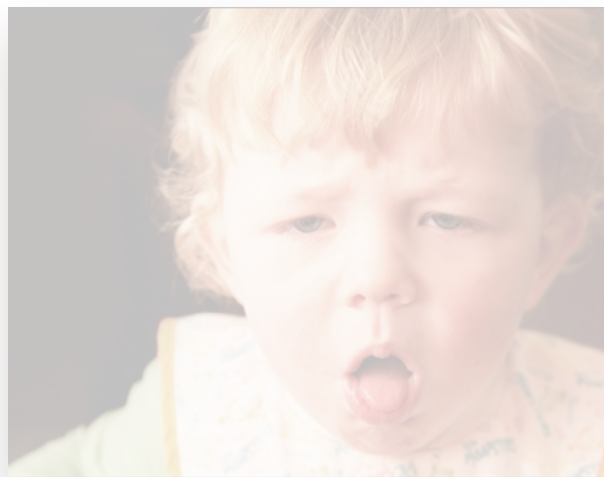
$$R_0 = \frac{\beta S}{\gamma}$$



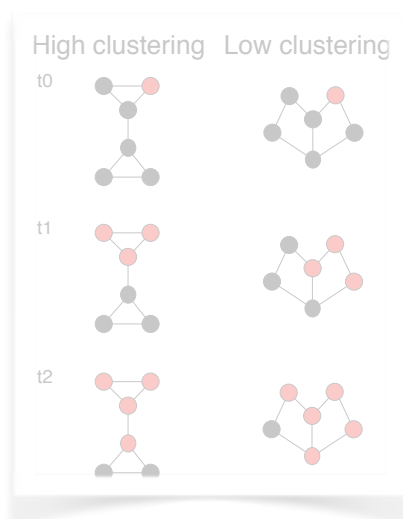
Model complexity



Modeling infectious diseases

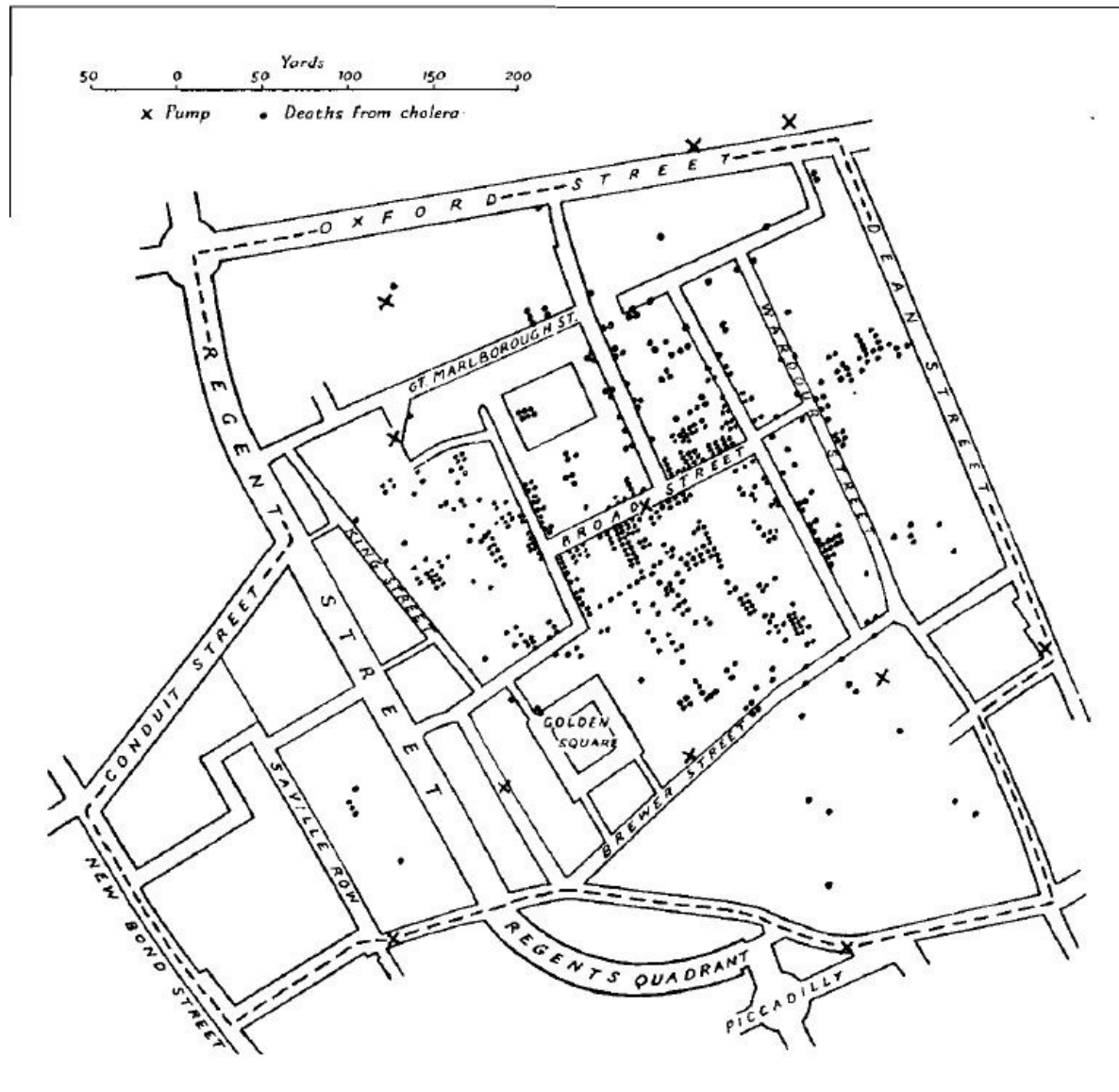


Inequality & disease



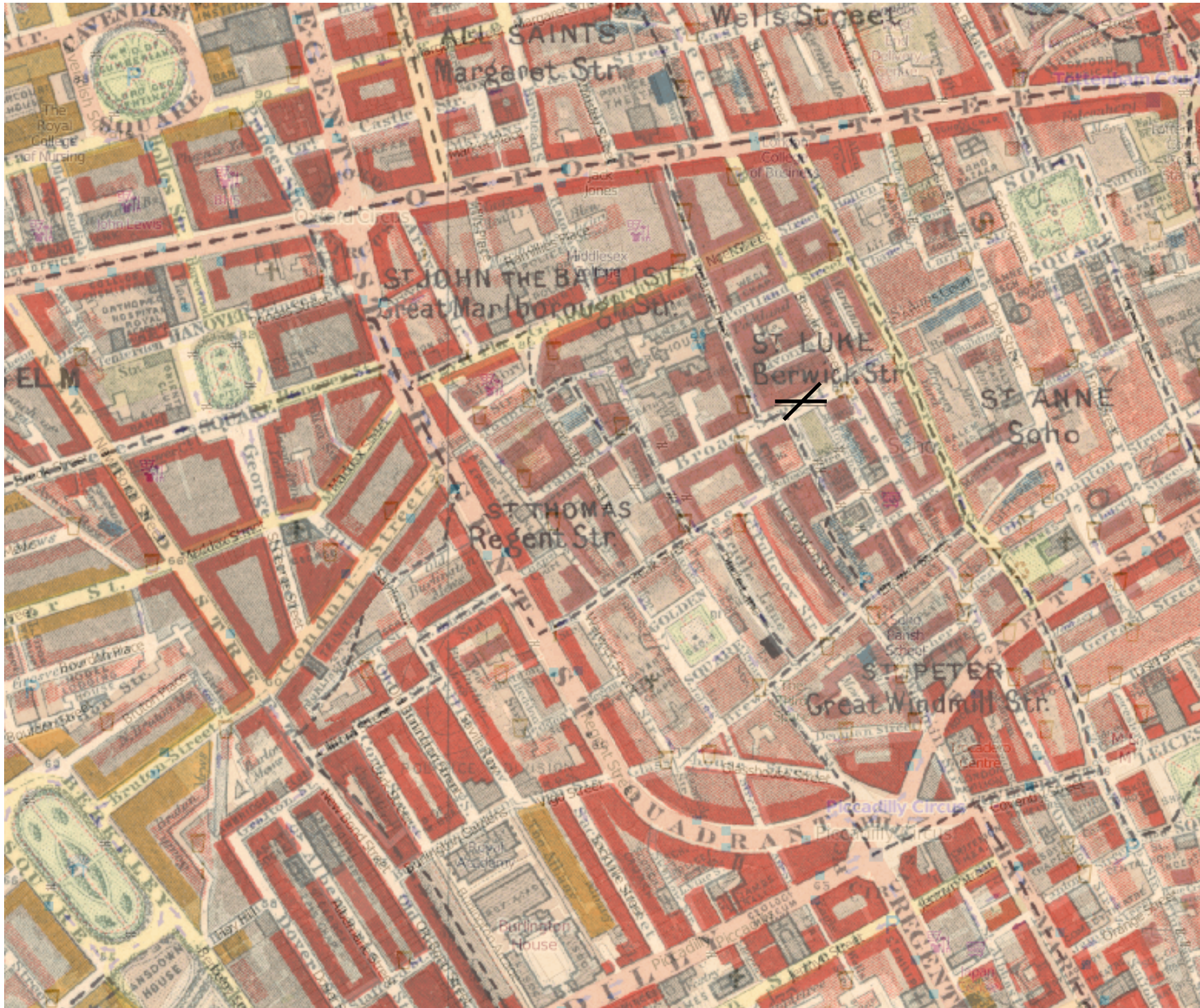
Social clustering & Ebola

The Cholera Outbreak of 1854



Snow 1855

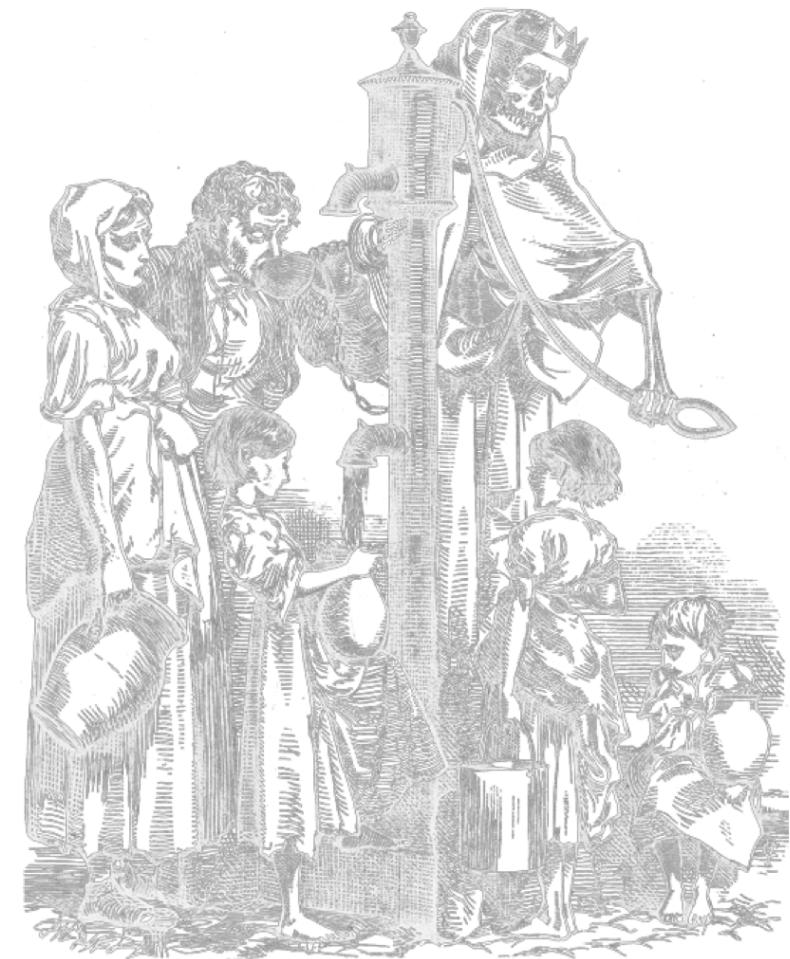
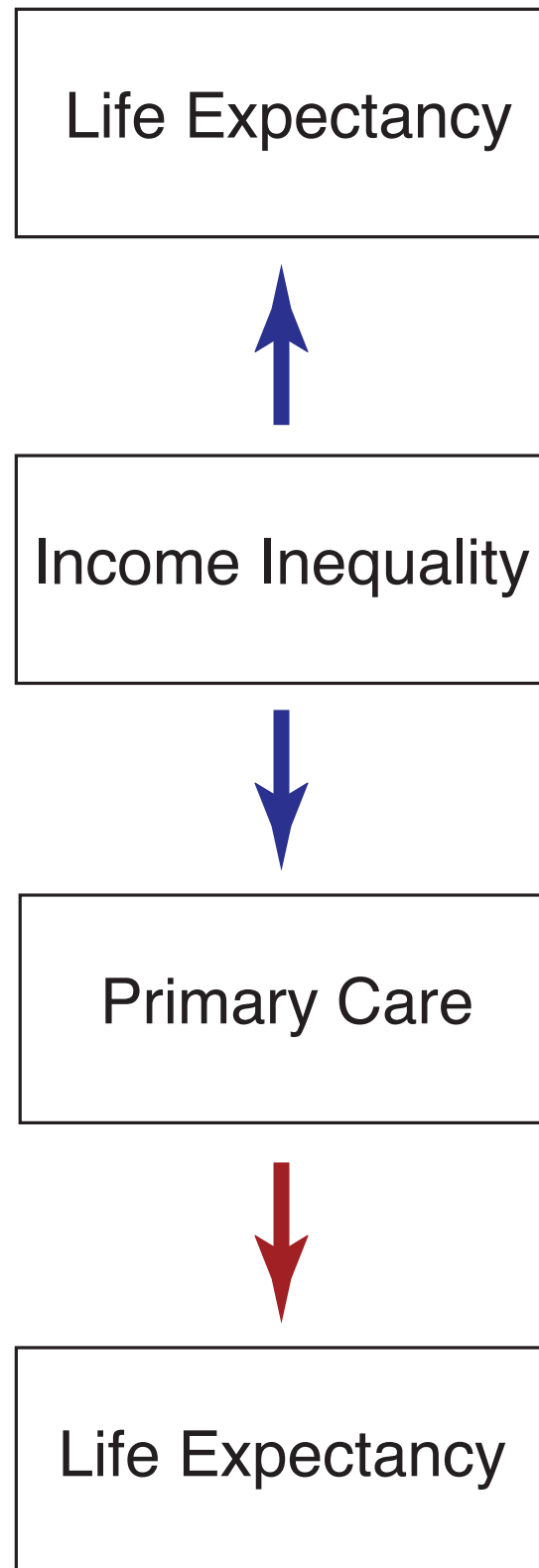
Poverty and the Cholera Outbreak of 1854



WEALTHY	
WELL-TO-DO	
COMFORTABLE	
POOR & COMFORTABLE (MIXED)	
POOR	
VERY POOR	
SEMI-CRIMINAL	

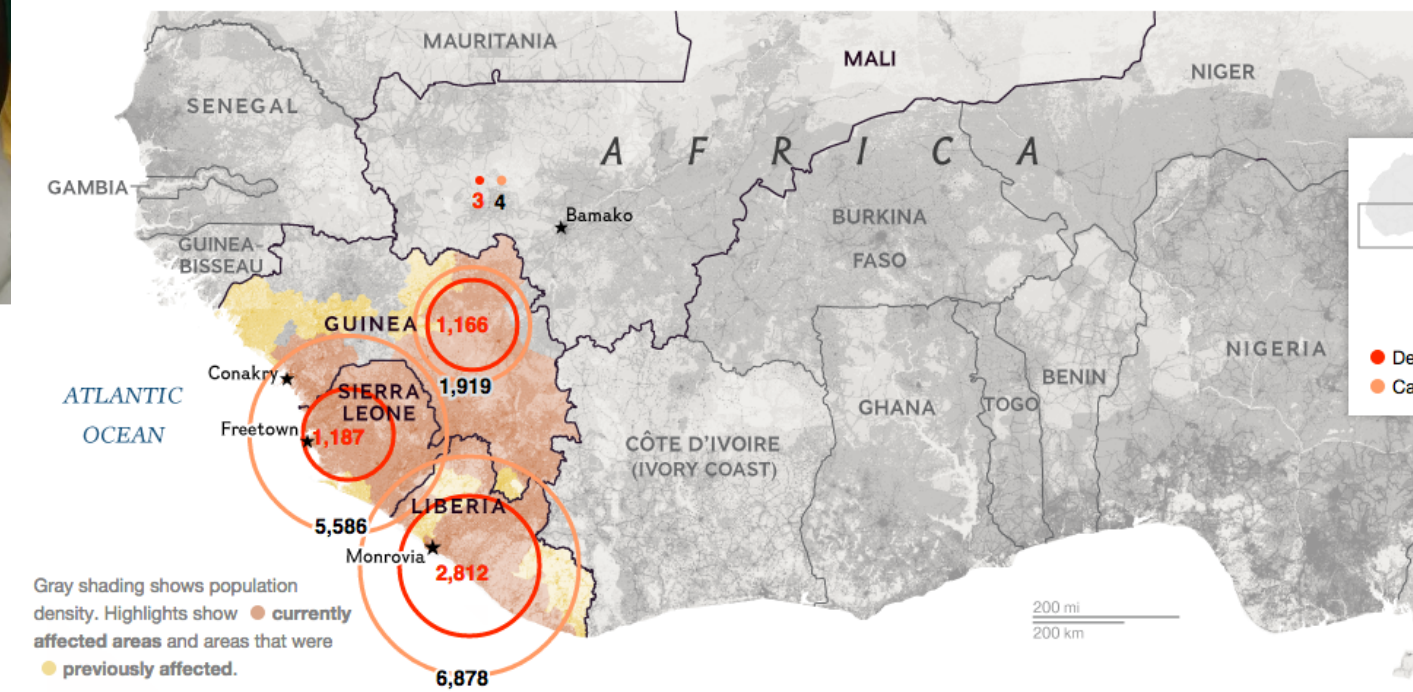
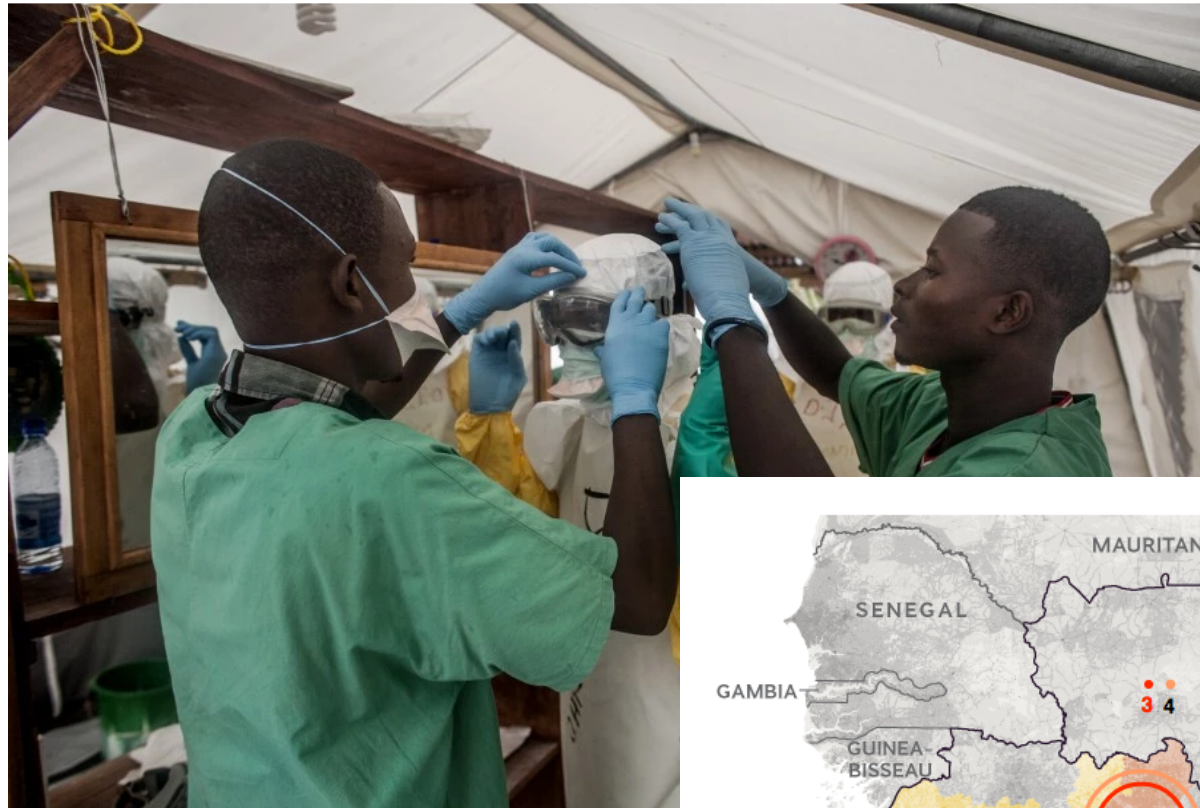
Booth 1898

Modern Inequality and health

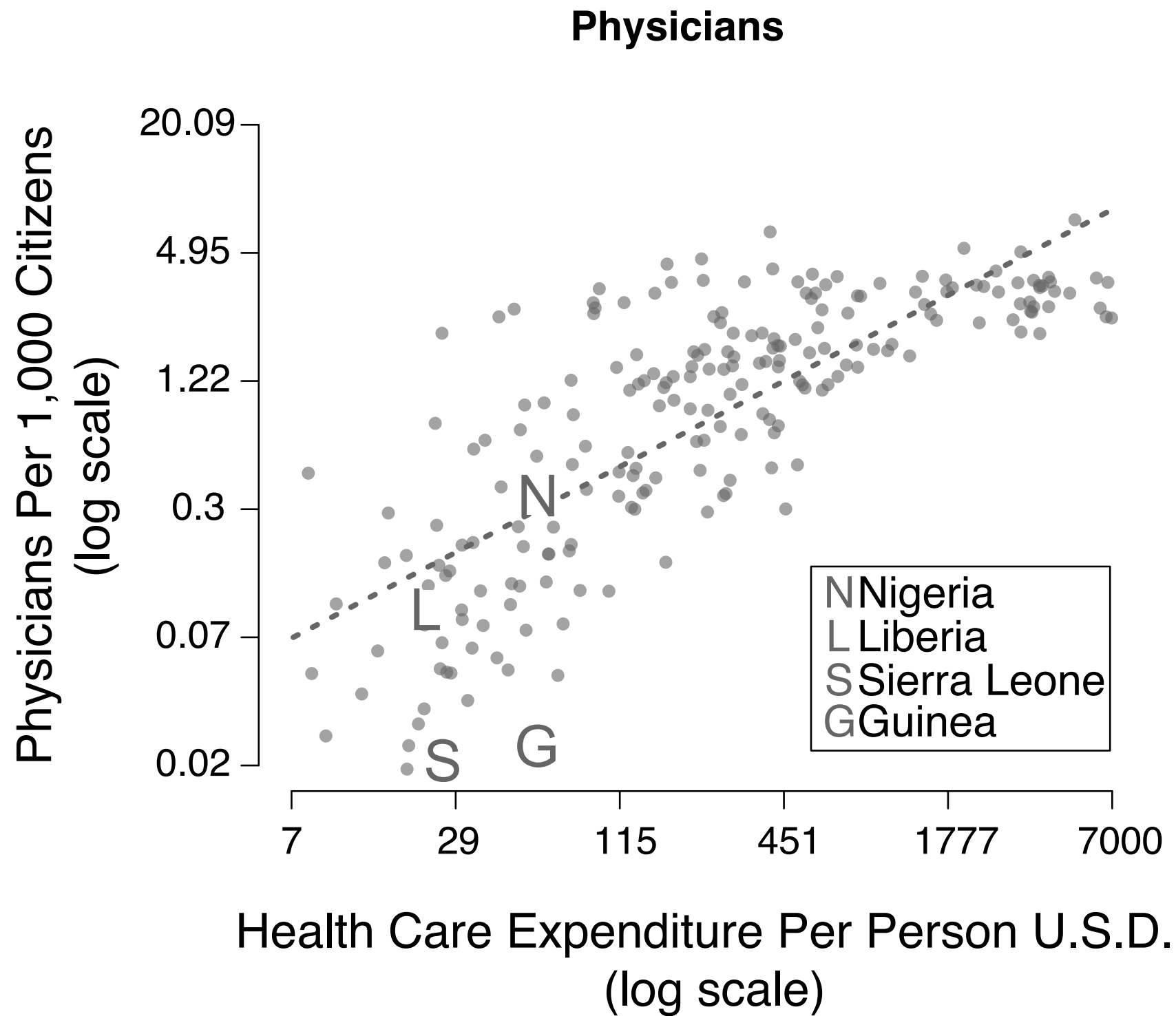


modified from Shi et al. 1999

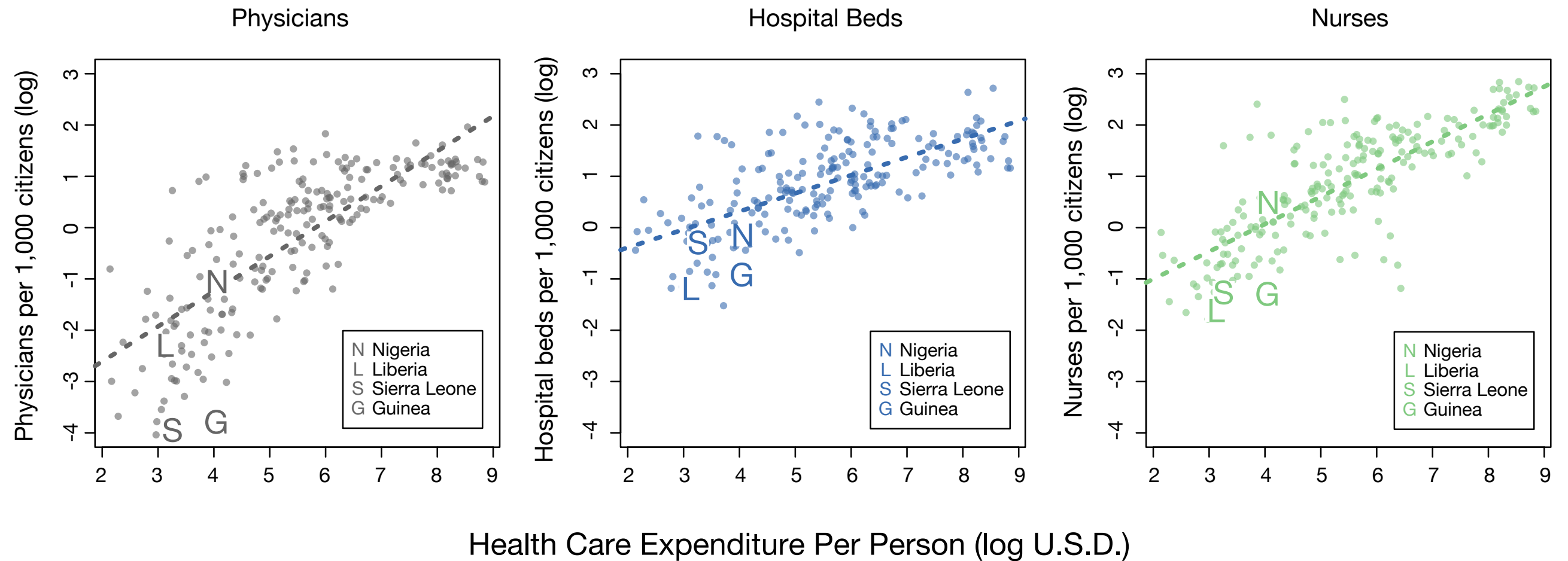
The current Ebola outbreak



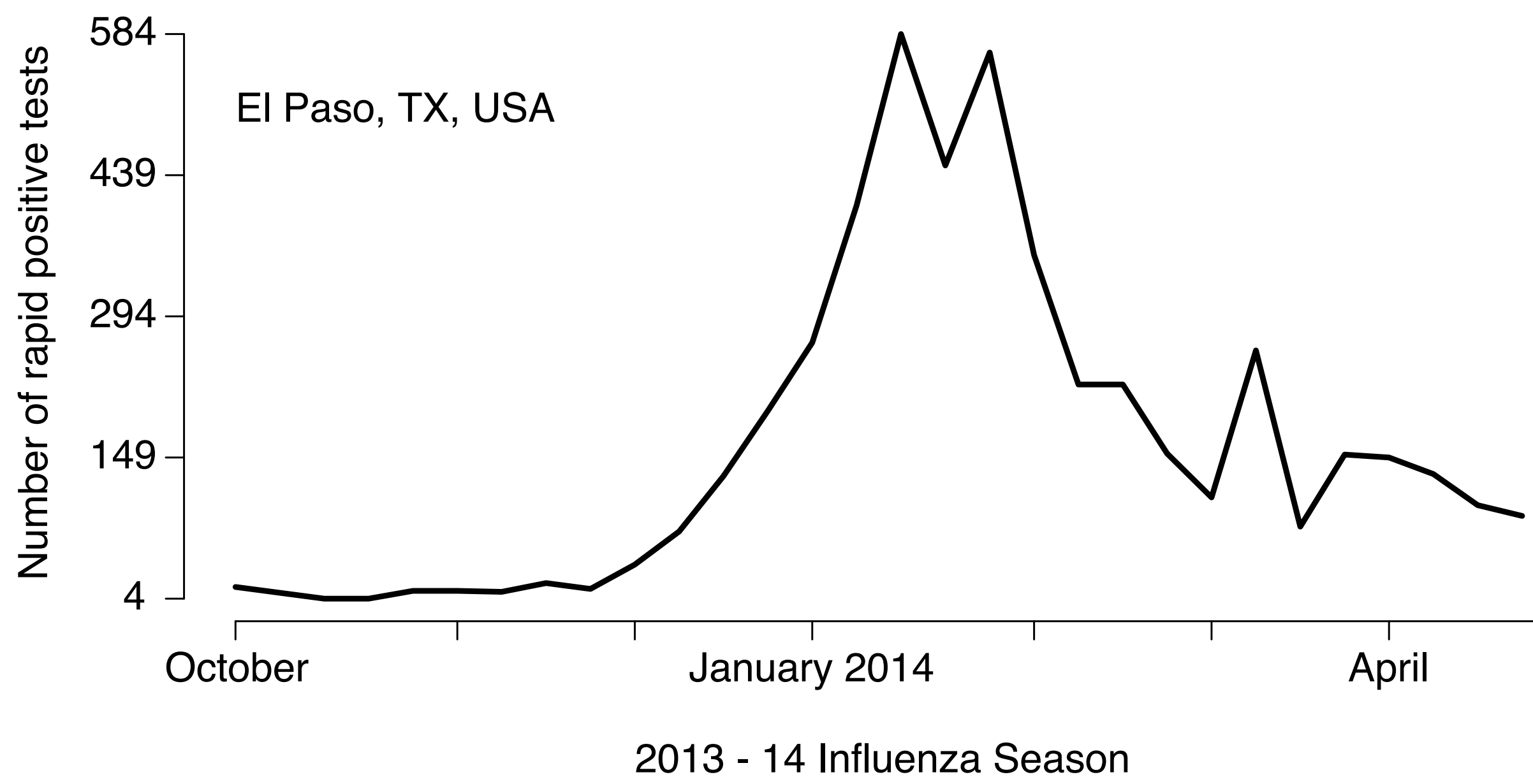
Poverty and the current Ebola virus outbreak



Poverty and the current Ebola outbreak



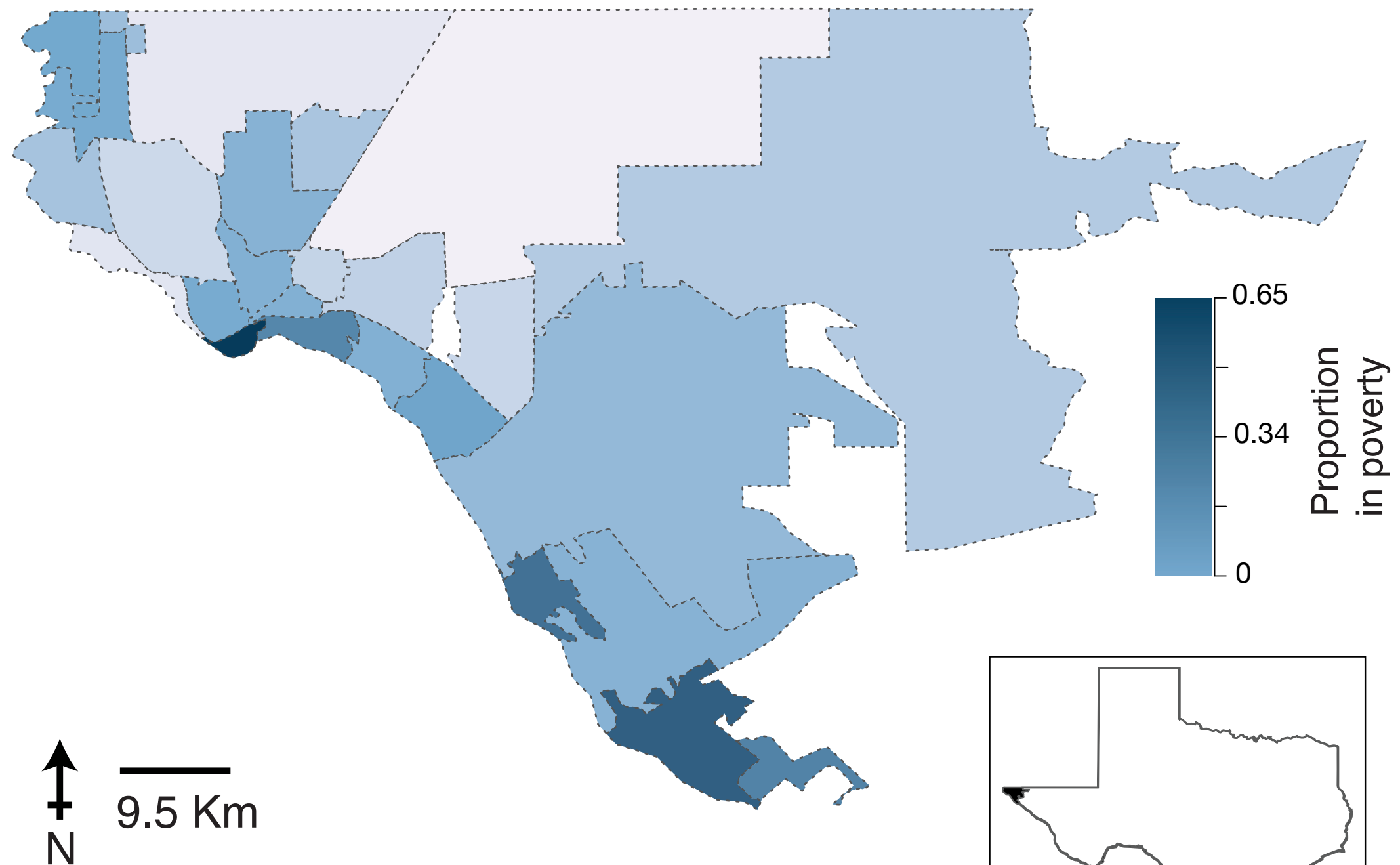
Influenza in El Paso, TX



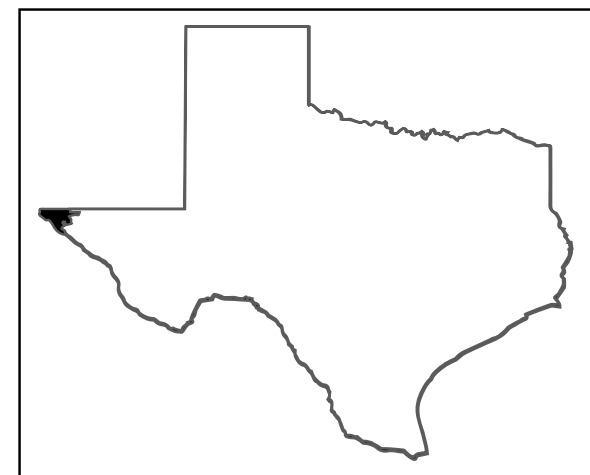
El Paso, TX 2014 - 15 Influenza Season



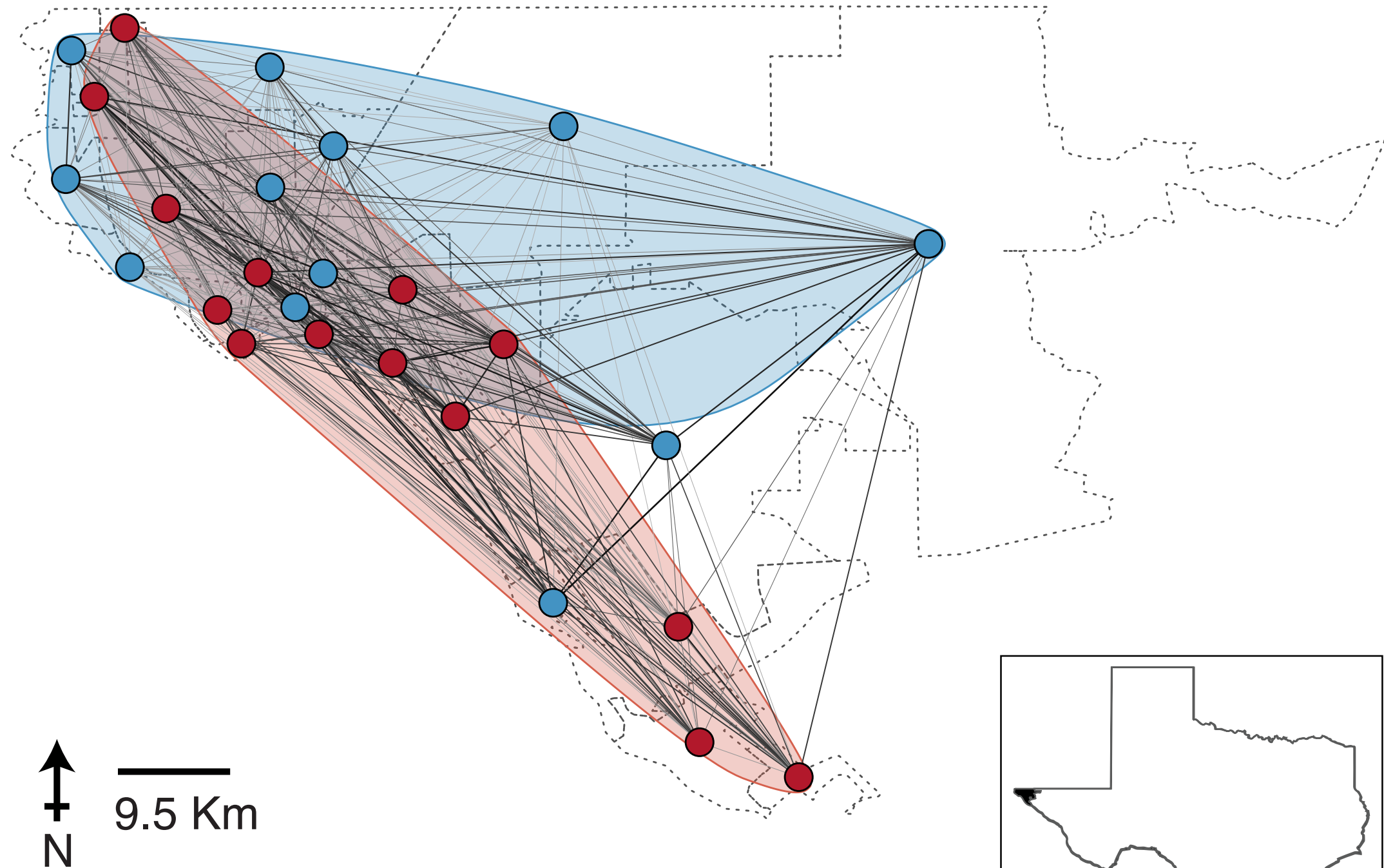
Poverty in El Paso



El Paso, TX, USA

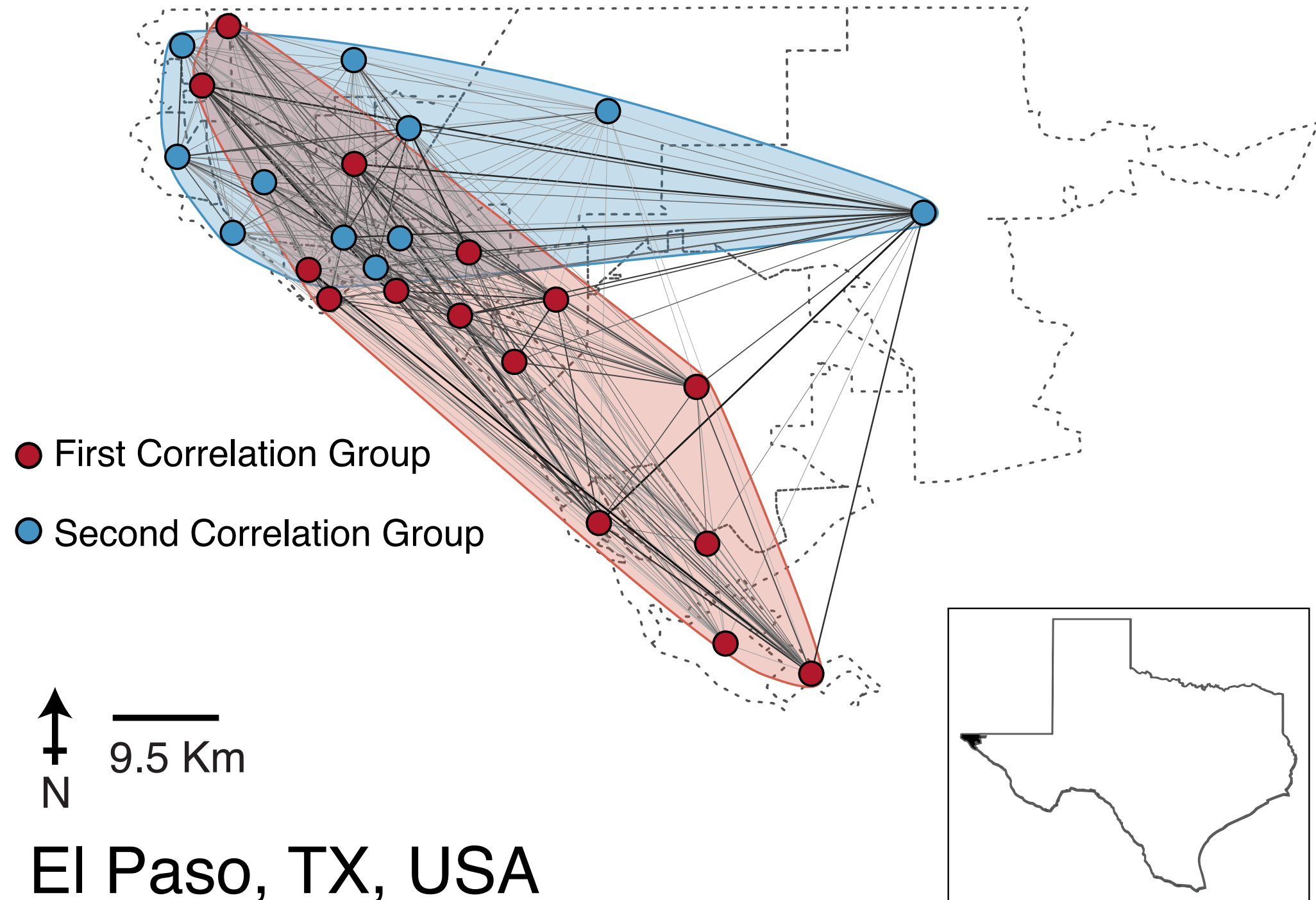


Poverty and flu in El Paso

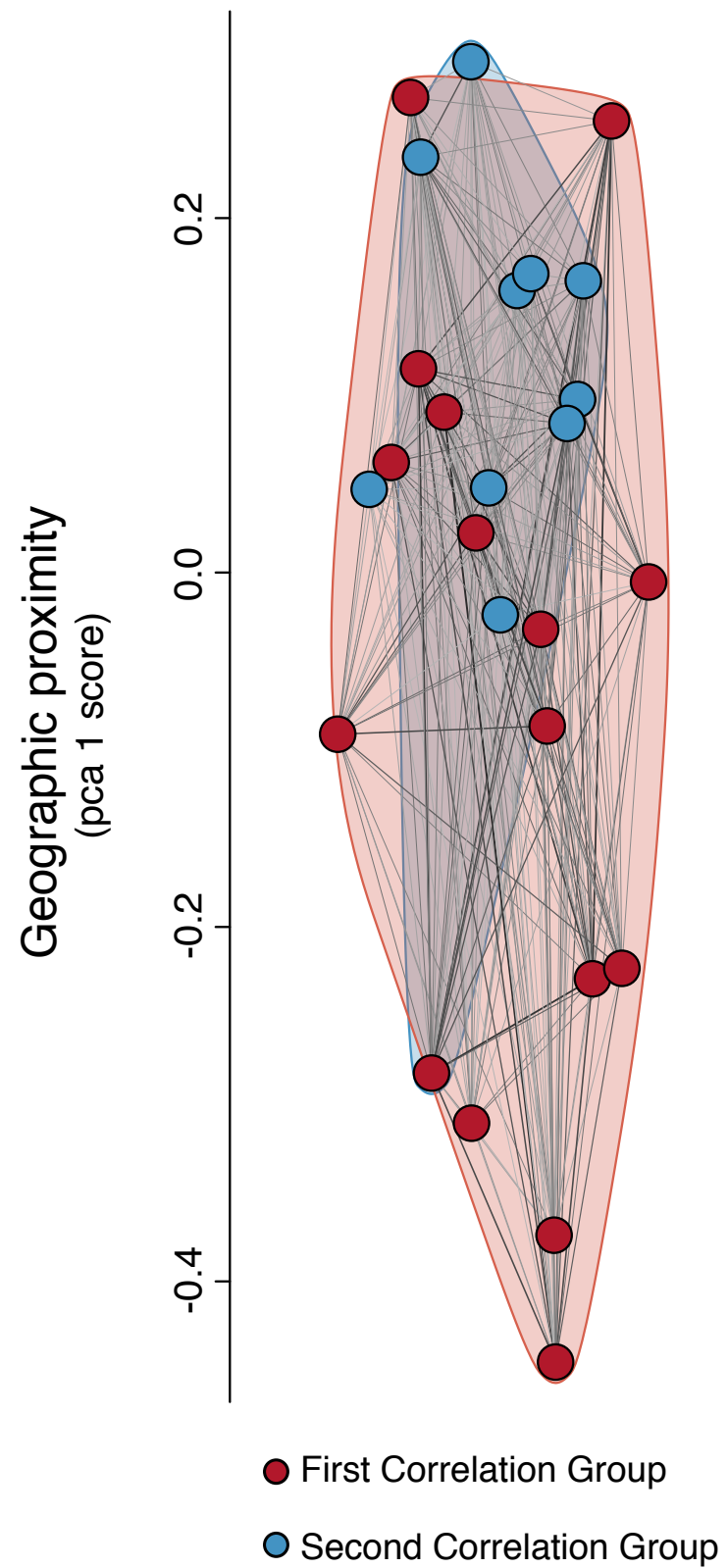


El Paso, TX, USA

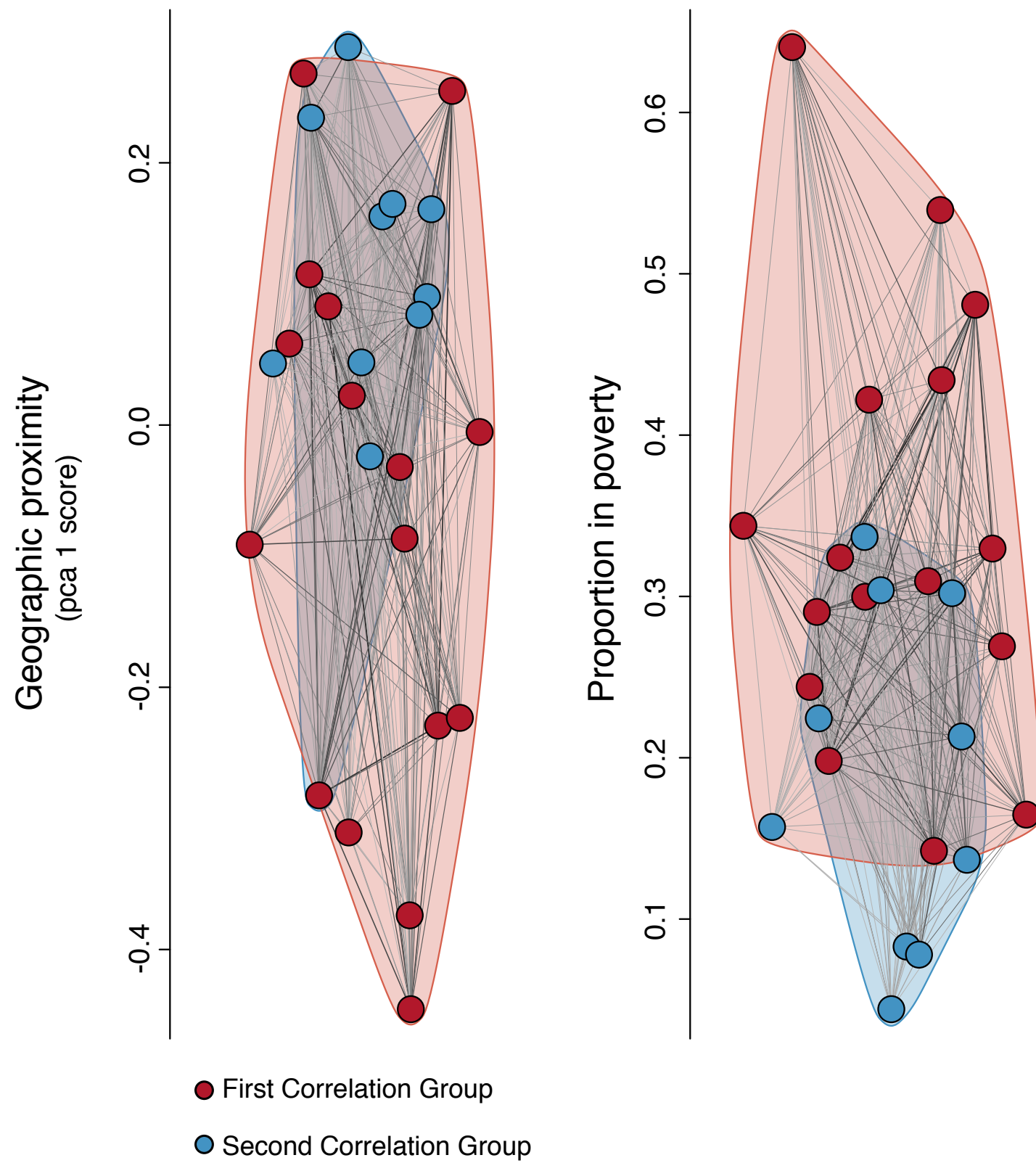
At least two strongly correlated groups



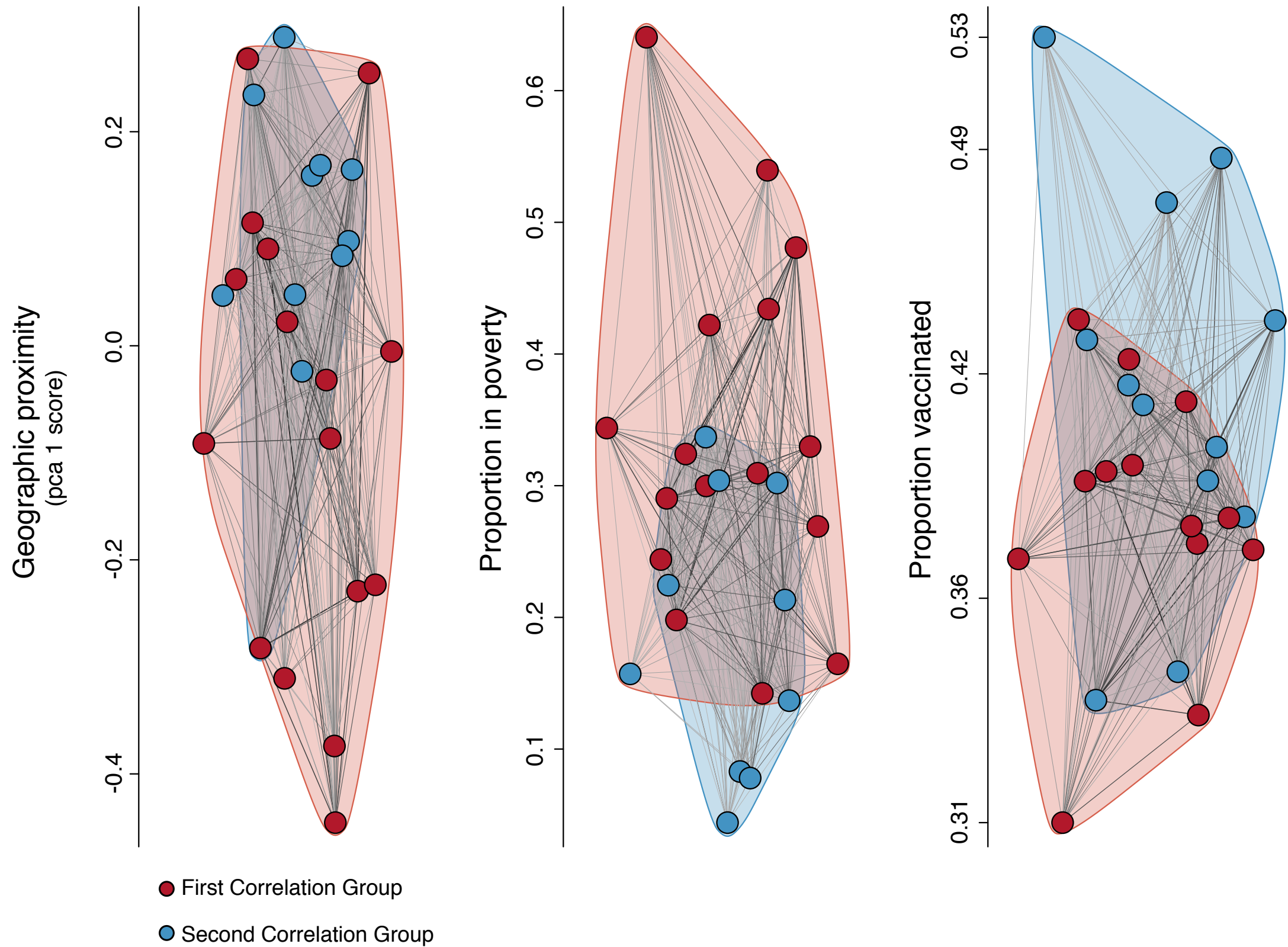
It's not geographic



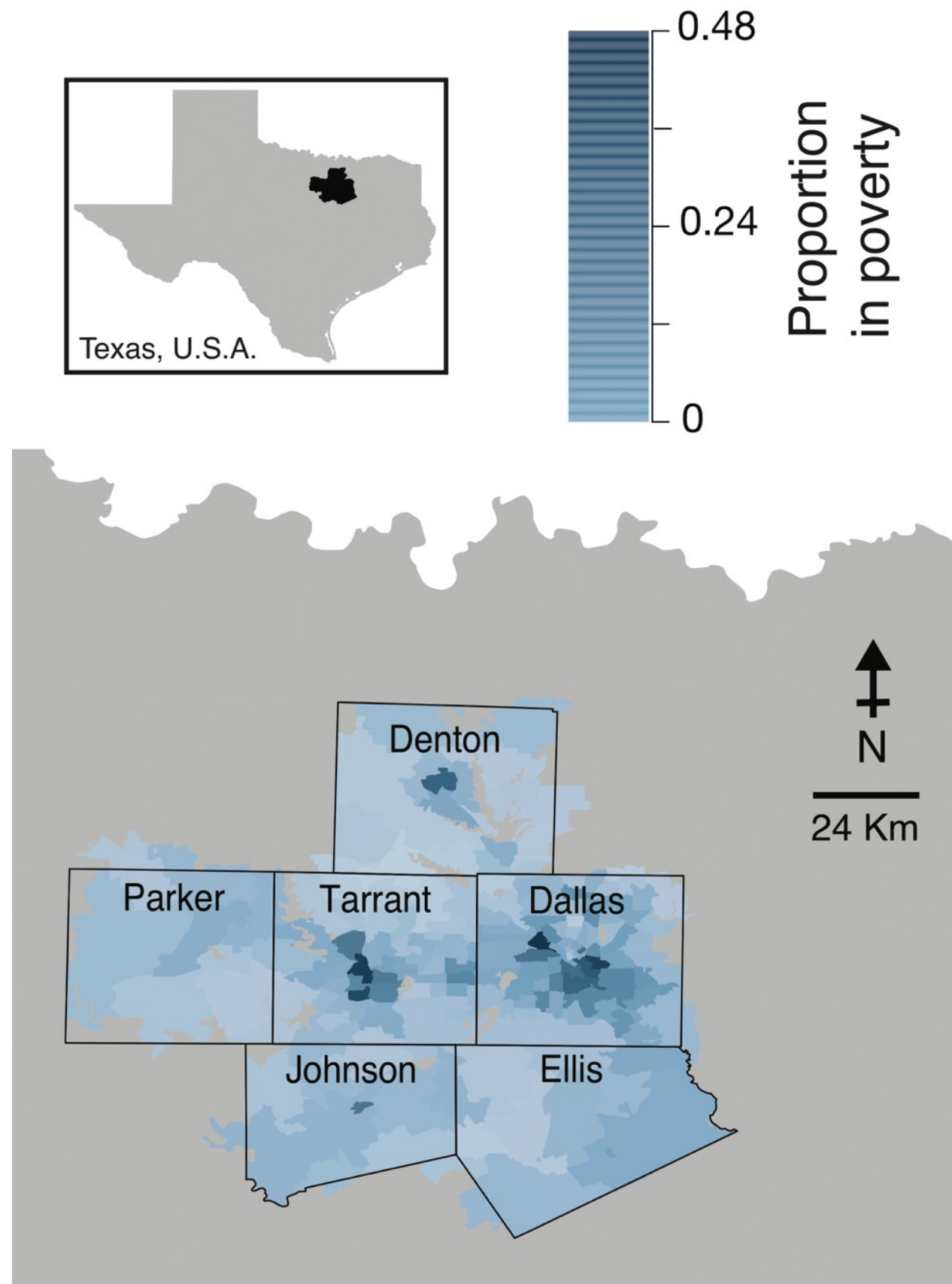
It's poverty and ...



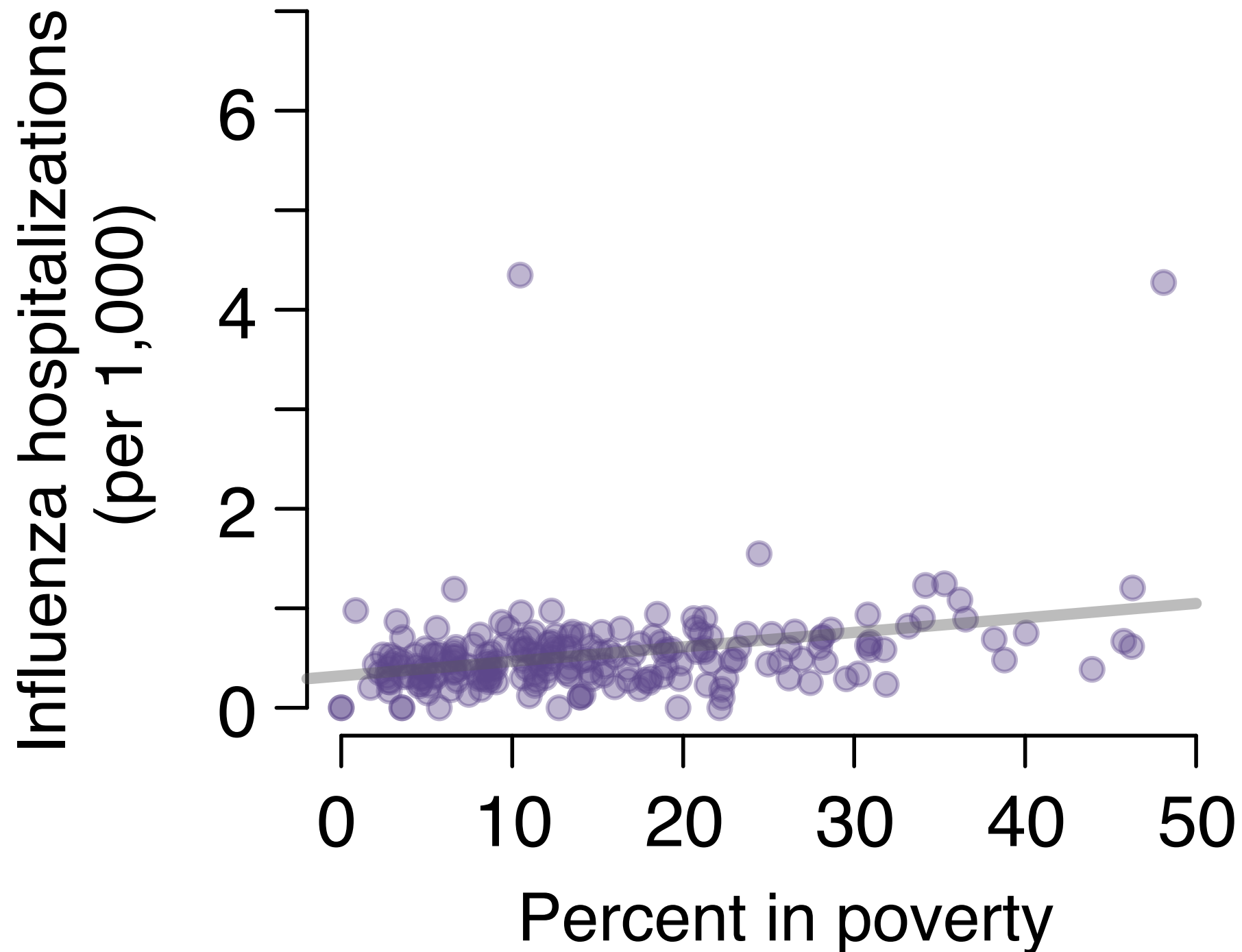
It's poverty and vaccination



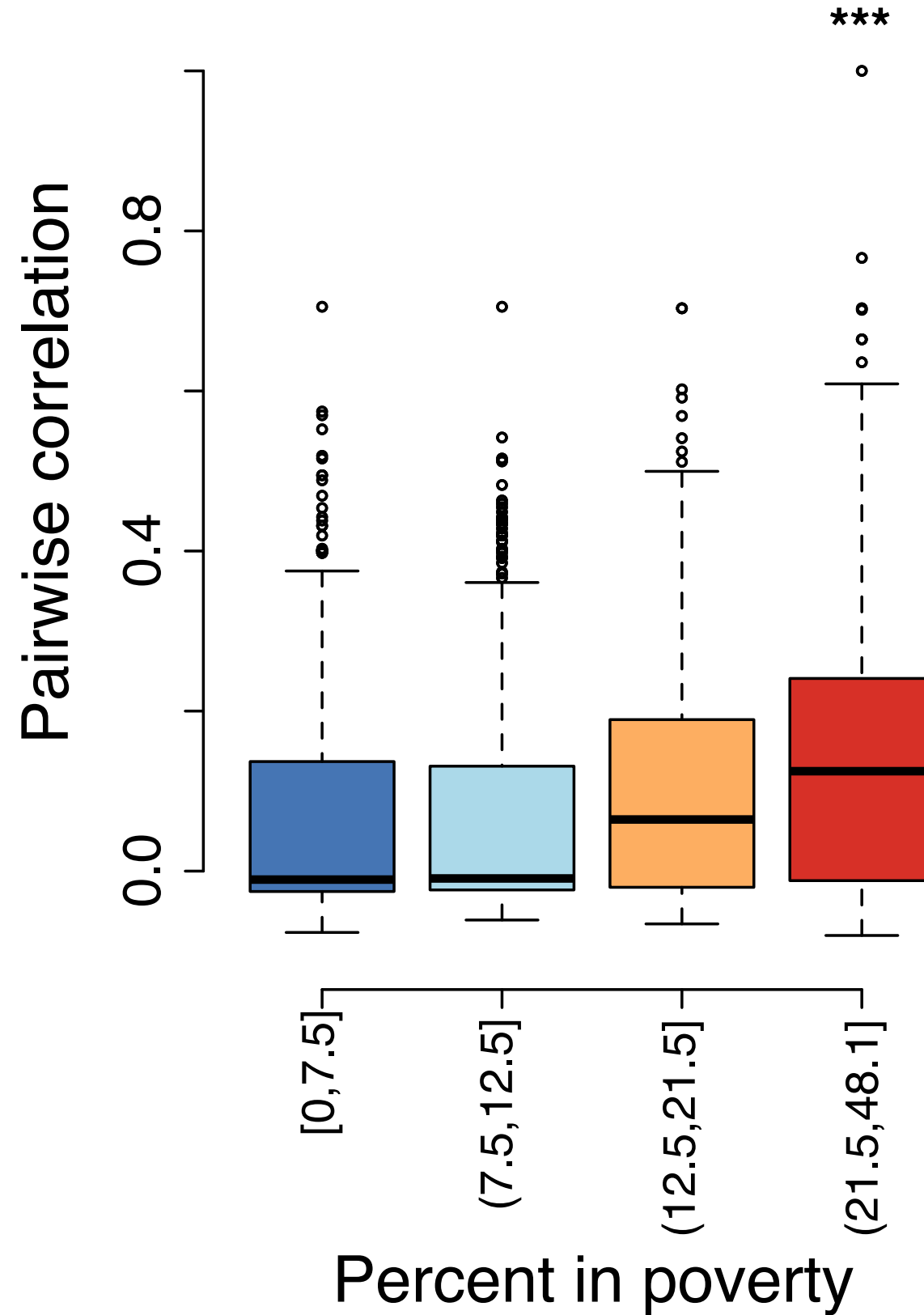
Poverty and influenza in Dallas, TX



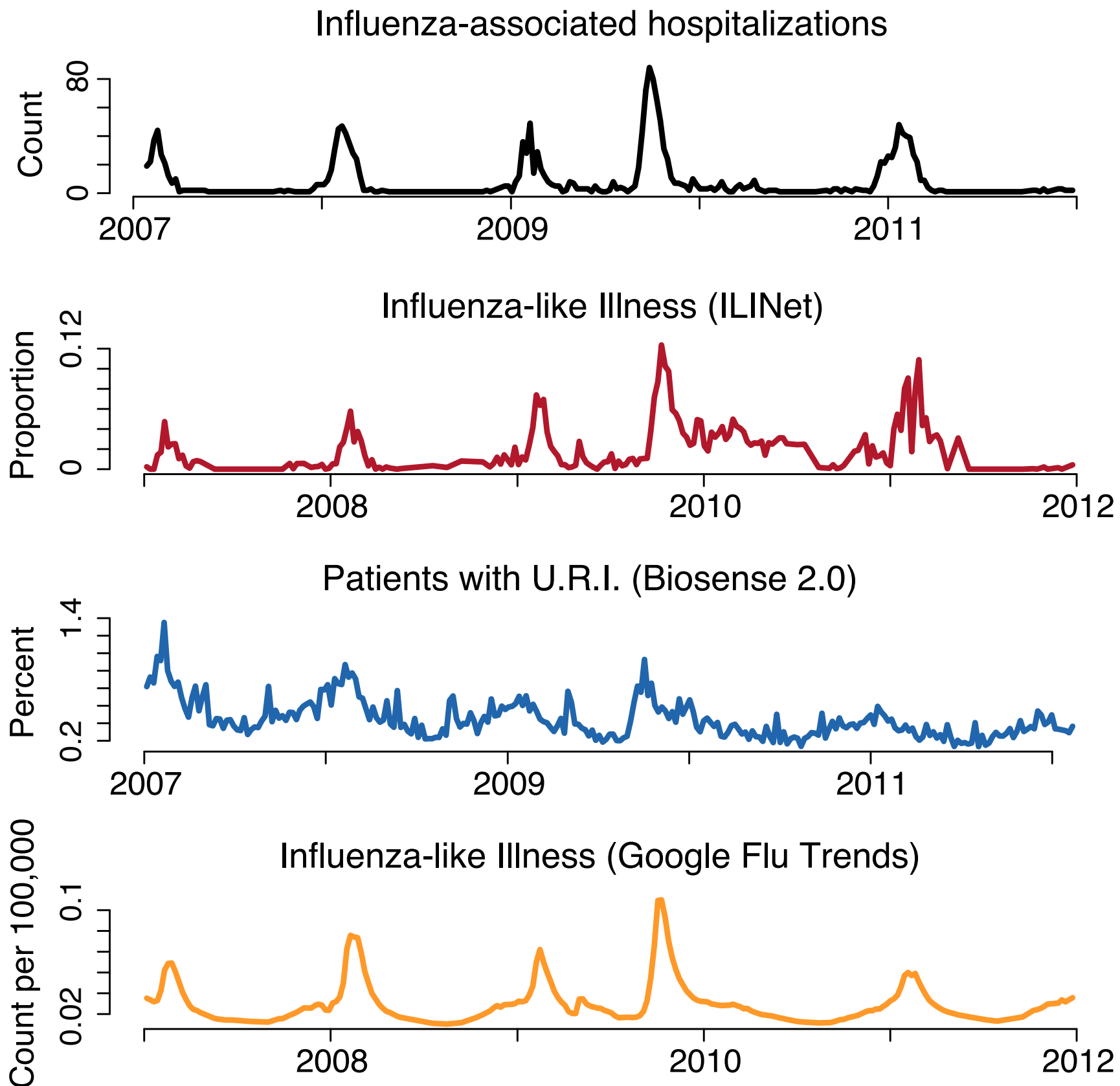
Higher hospitalization rates in poorer zip codes



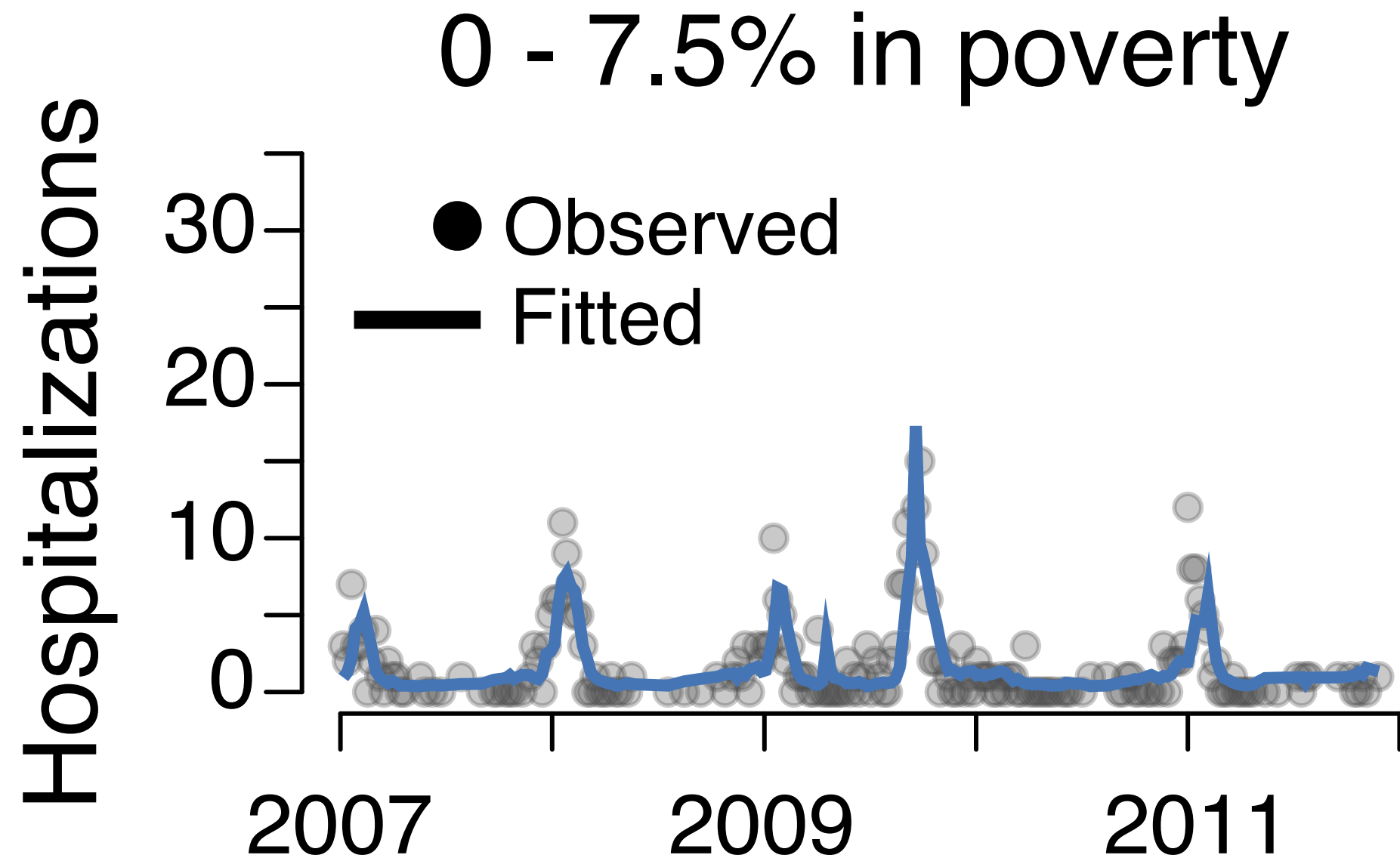
Poorer zip codes are in sync



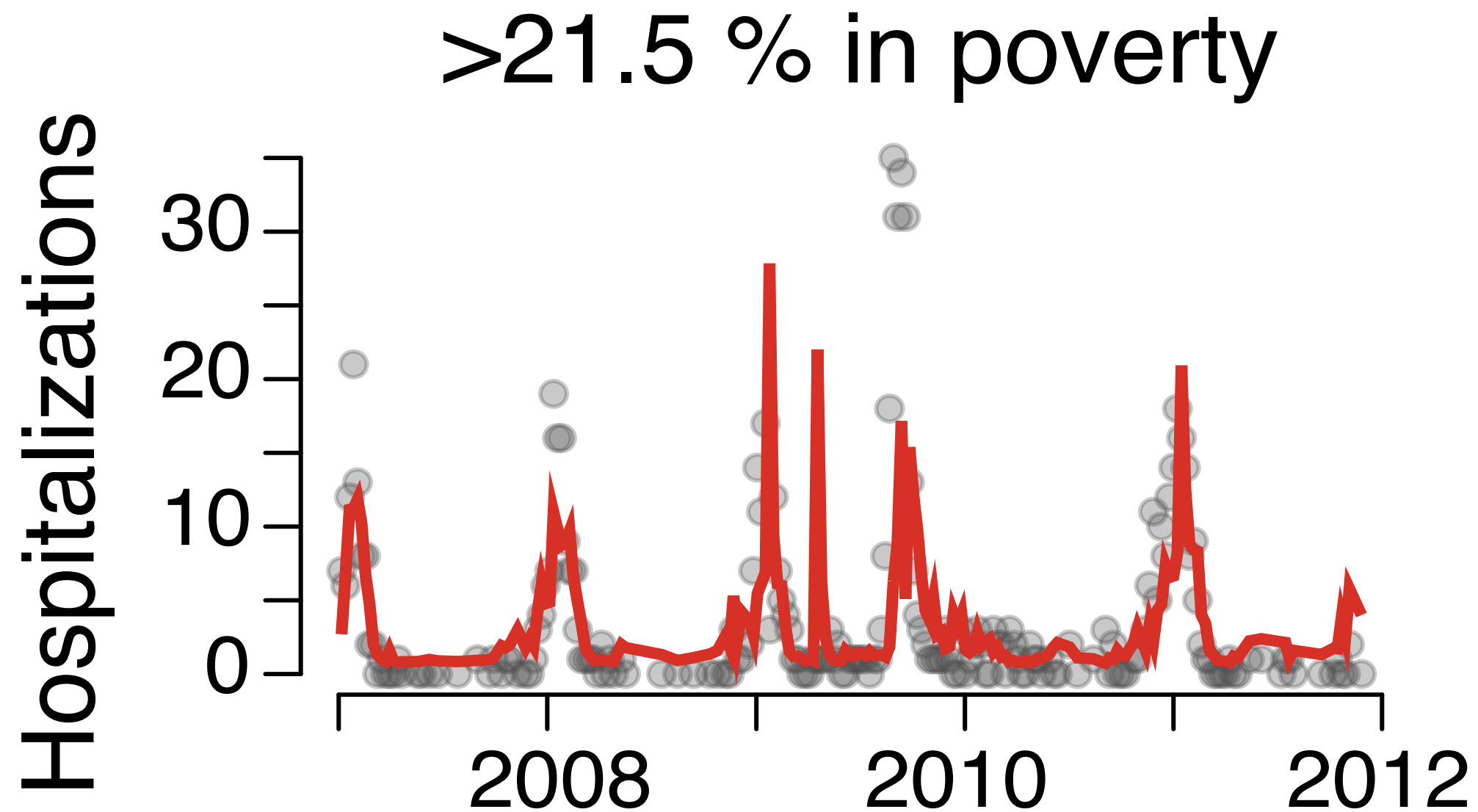
Predicting hospitalizations



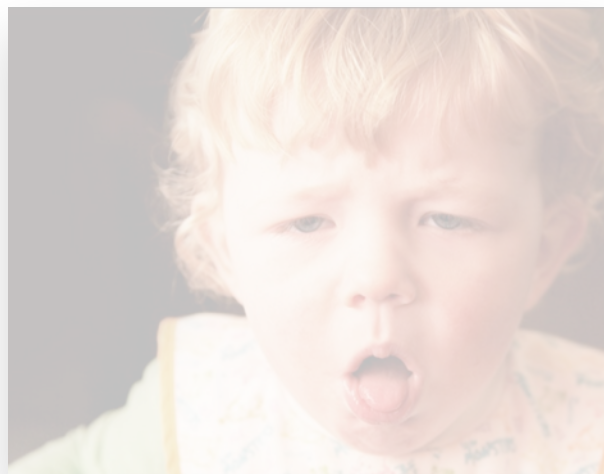
Predicting hospitalizations in the richest areas



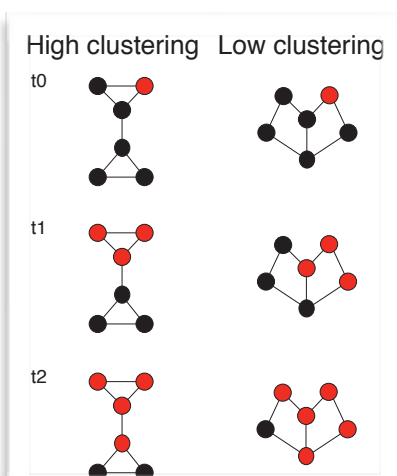
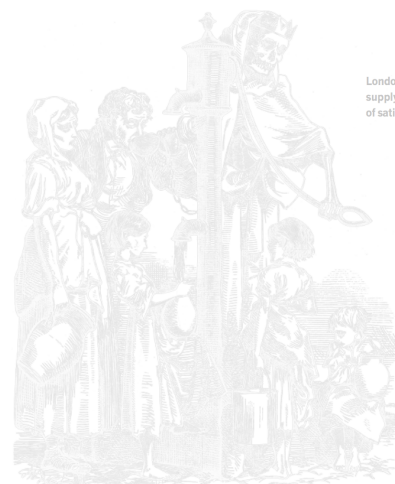
Predicting hospitalizations in the poorest areas



Modeling infectious diseases



Inequality & disease

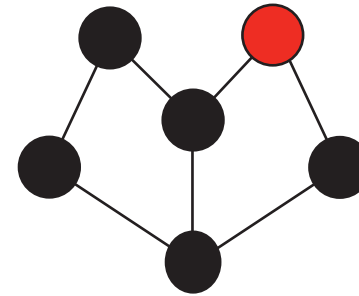
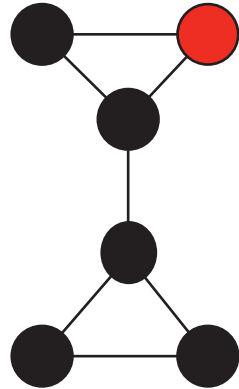


Social clustering & Ebola

Clustering and disease transmission

High clustering Low clustering

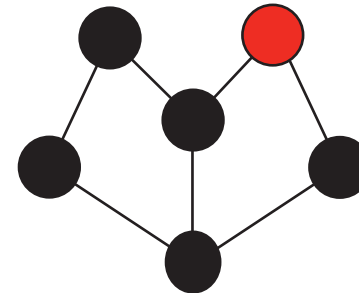
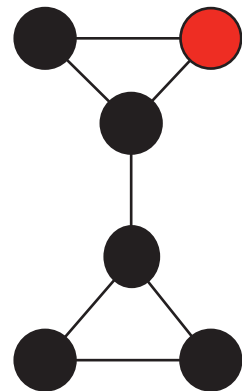
t_0



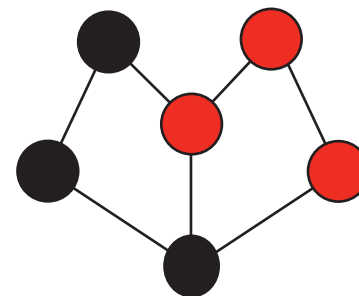
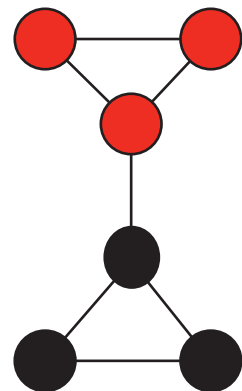
Clustering and disease transmission

High clustering Low clustering

t0



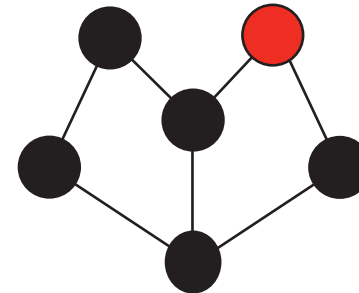
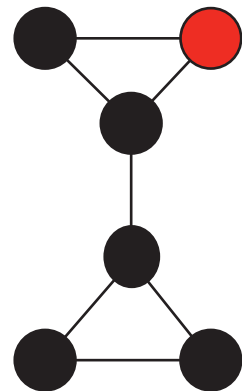
t1



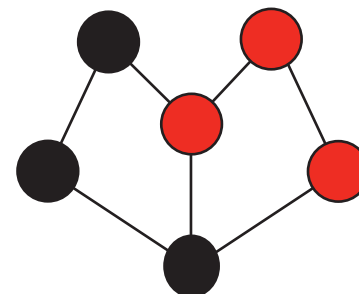
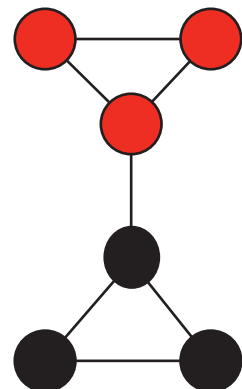
Clustering and disease transmission

High clustering Low clustering

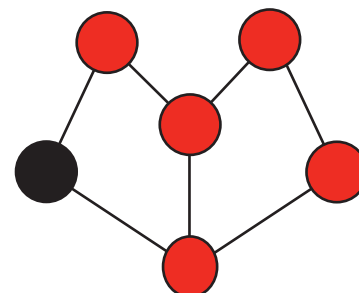
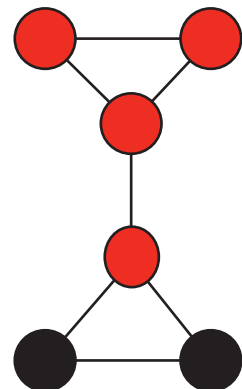
t0



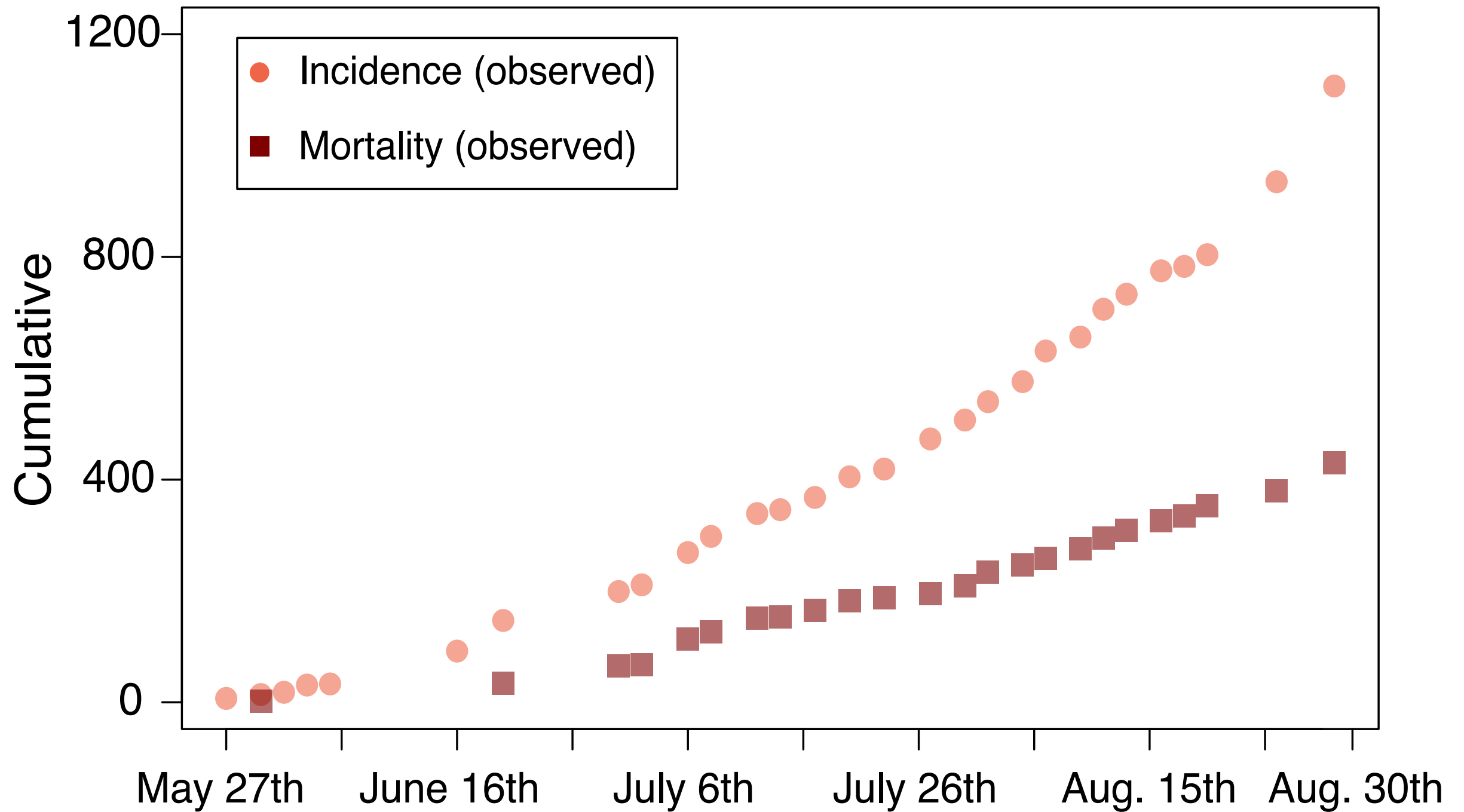
t1



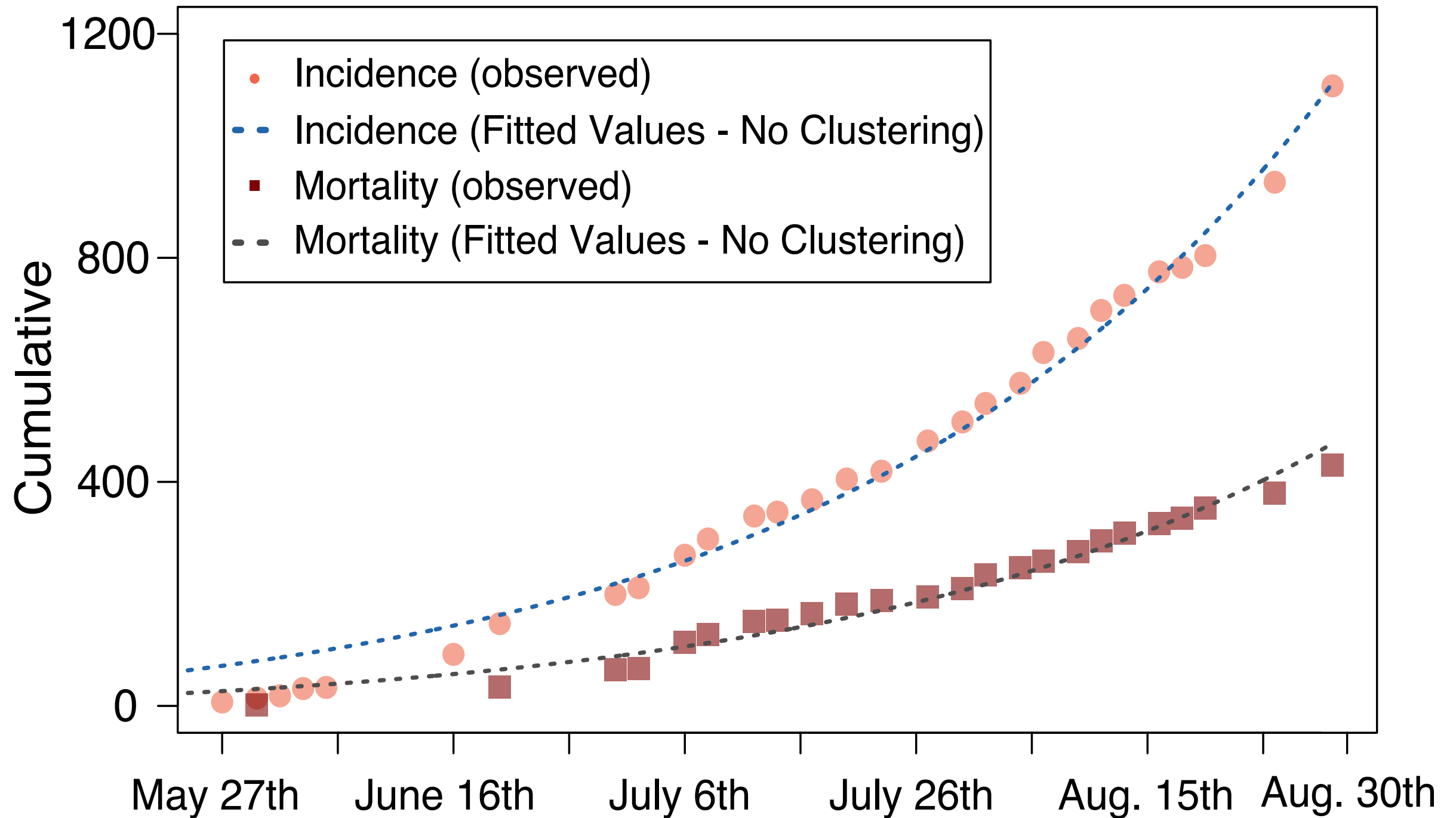
t2



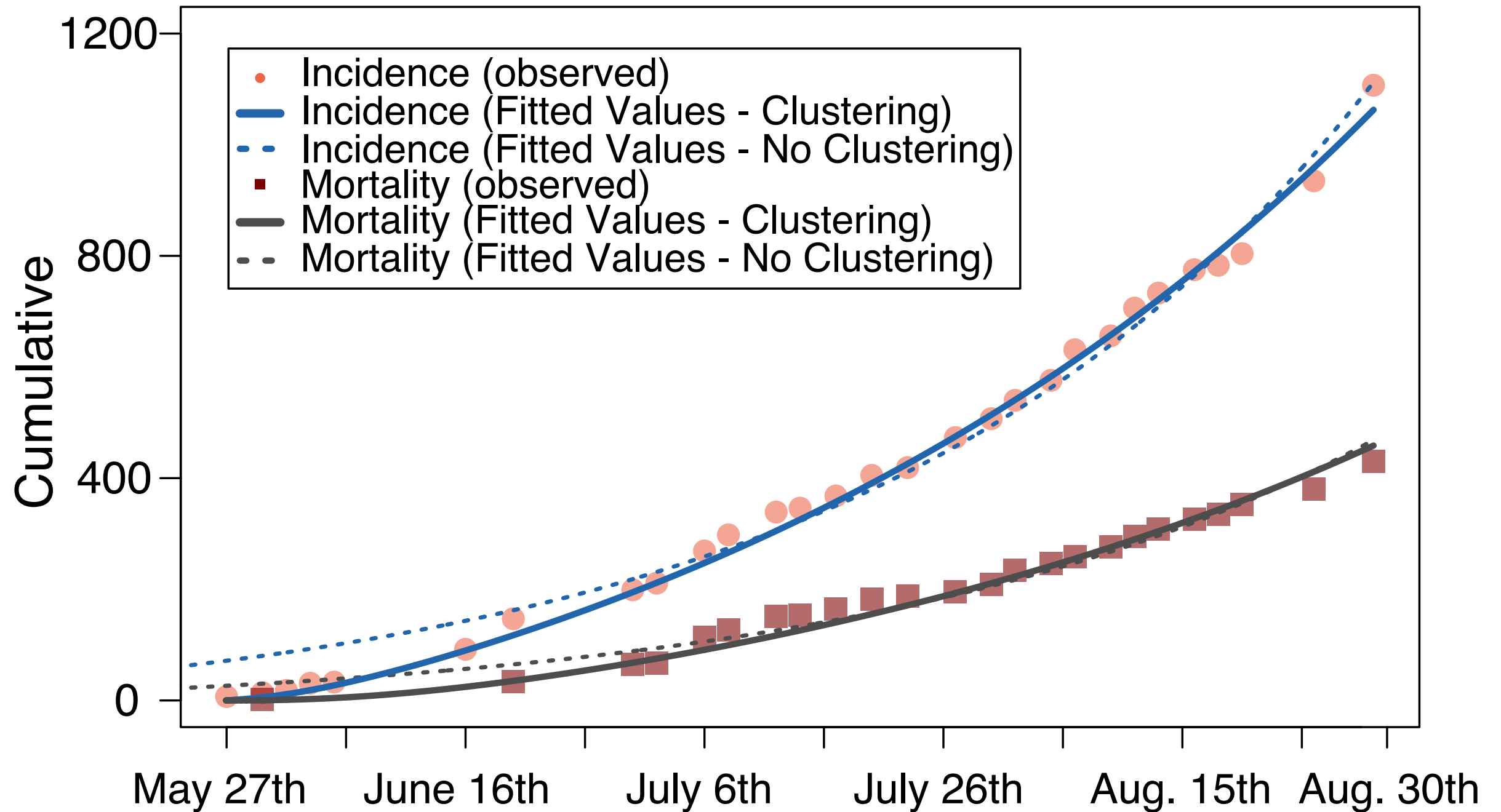
Outbreak in Sierra Leone



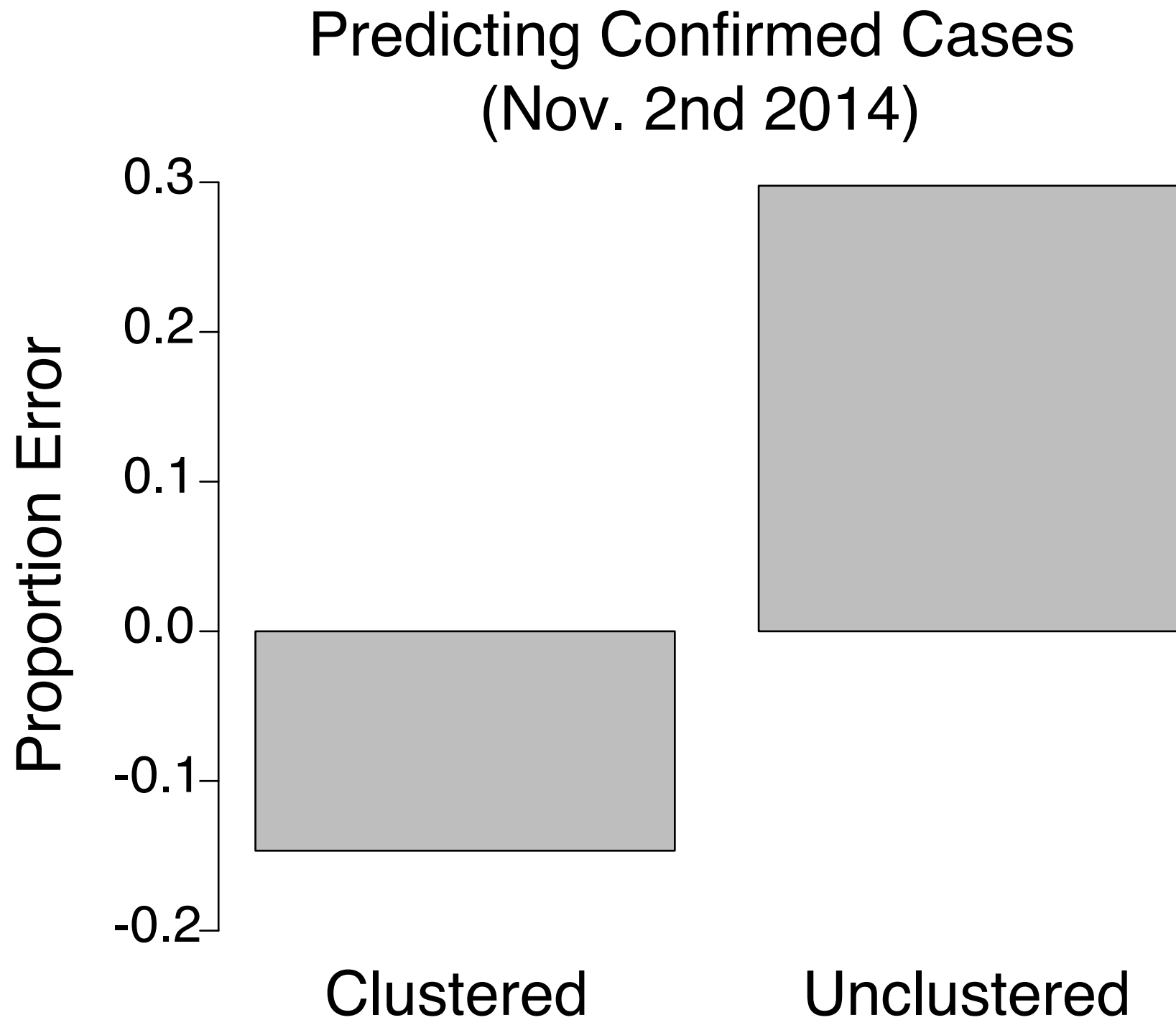
Evidence for clustered transmission



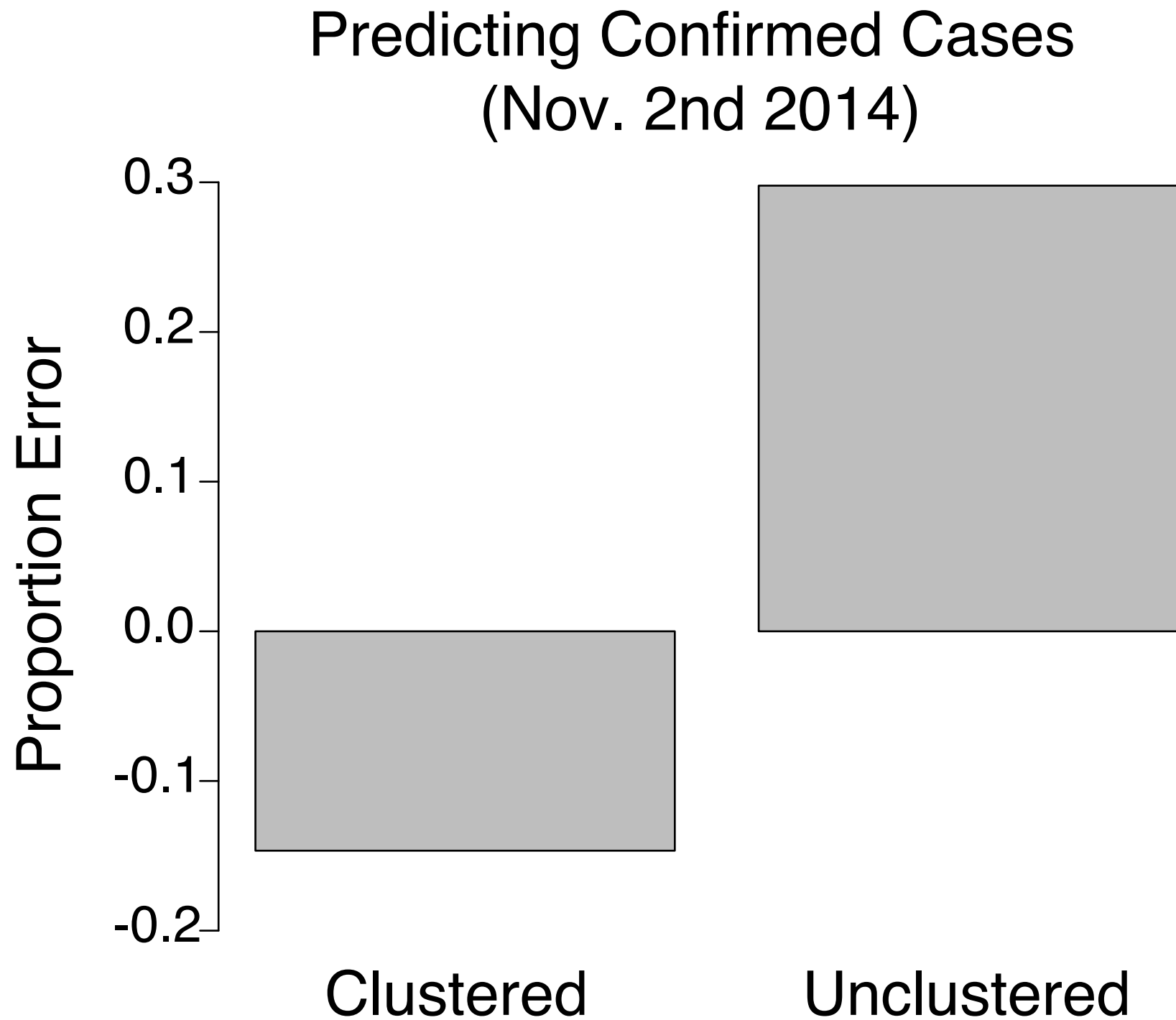
Evidence for clustered transmission



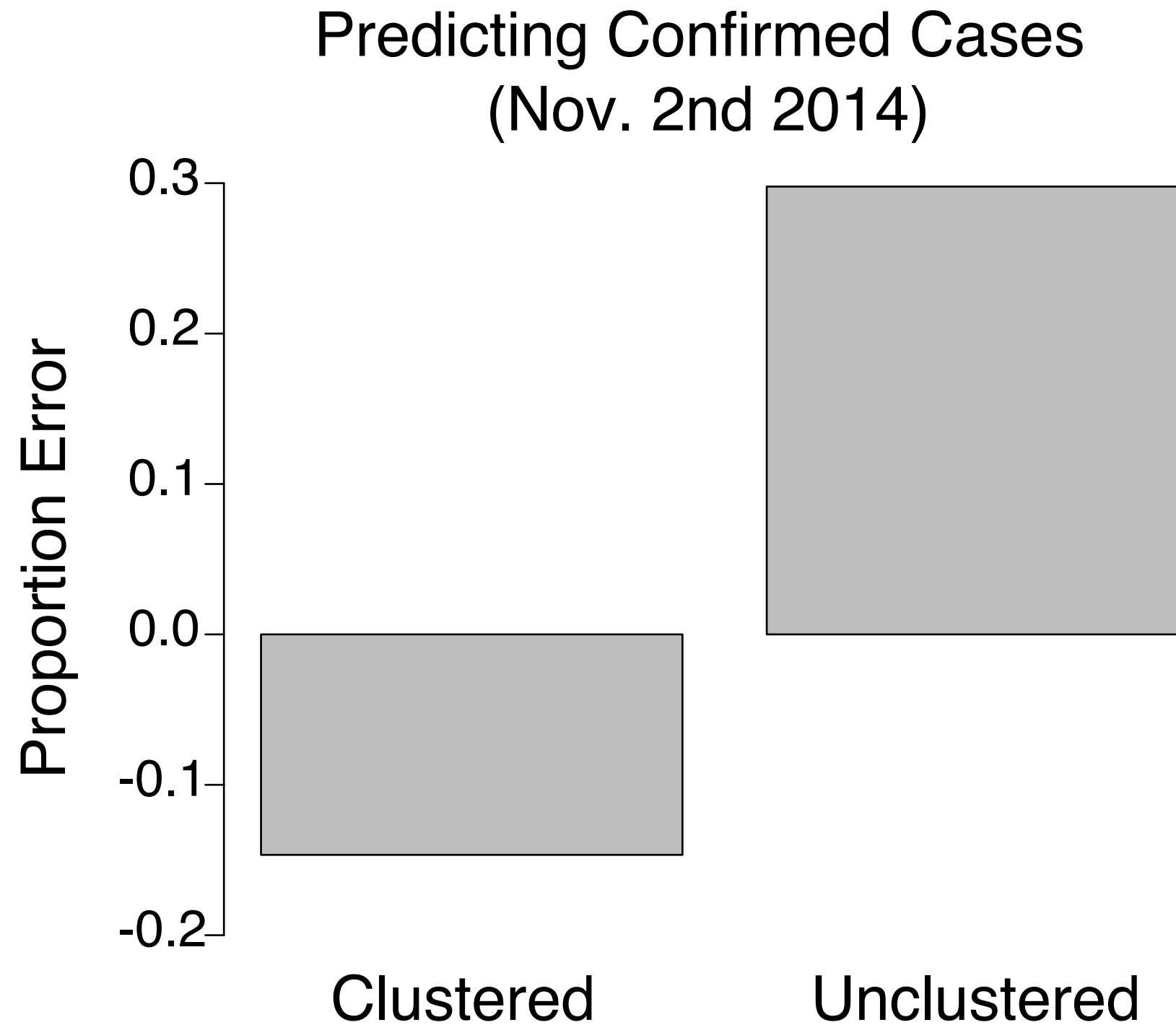
1. Dynamic importance of clustering



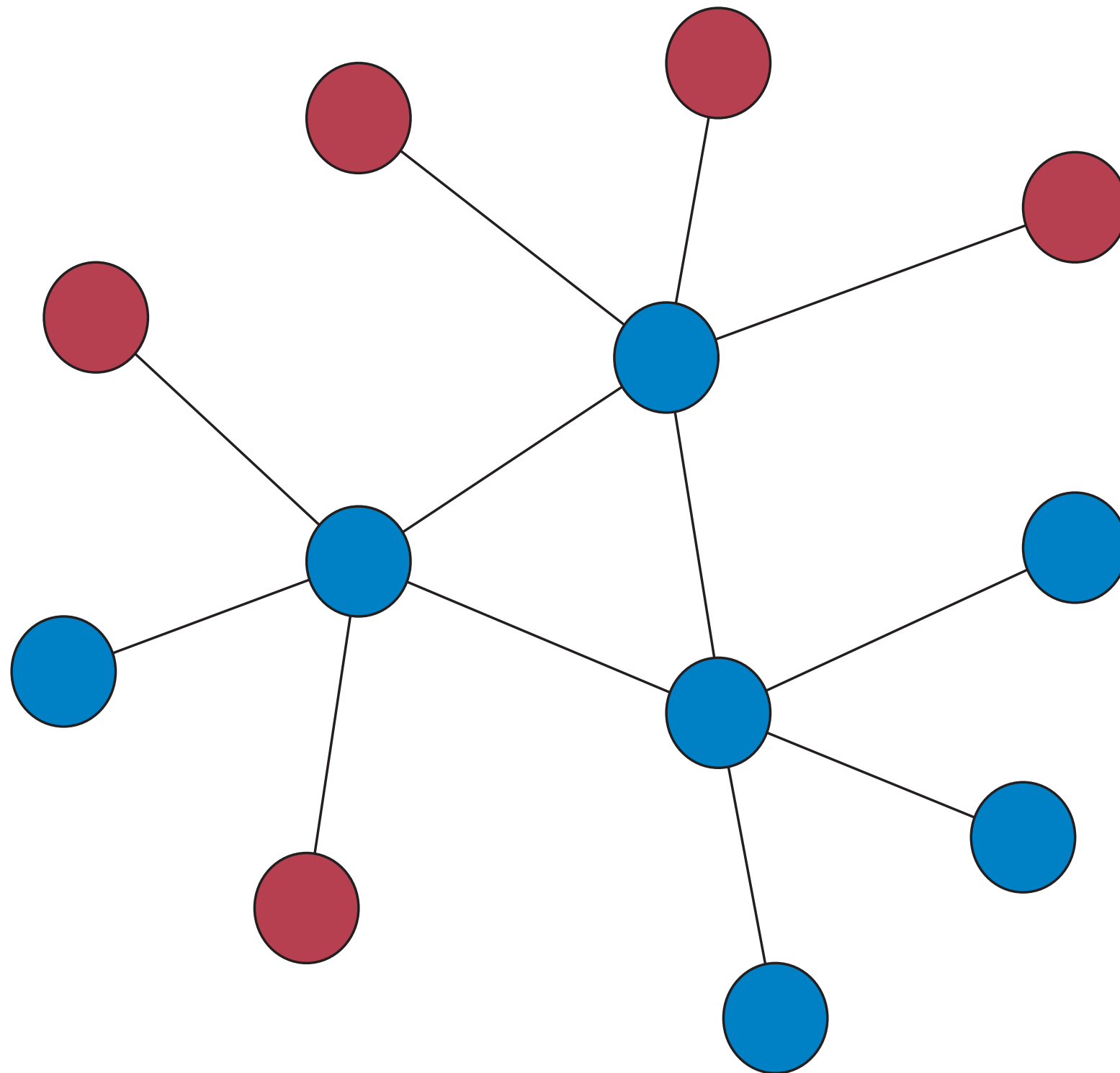
2. No evidence for interventions



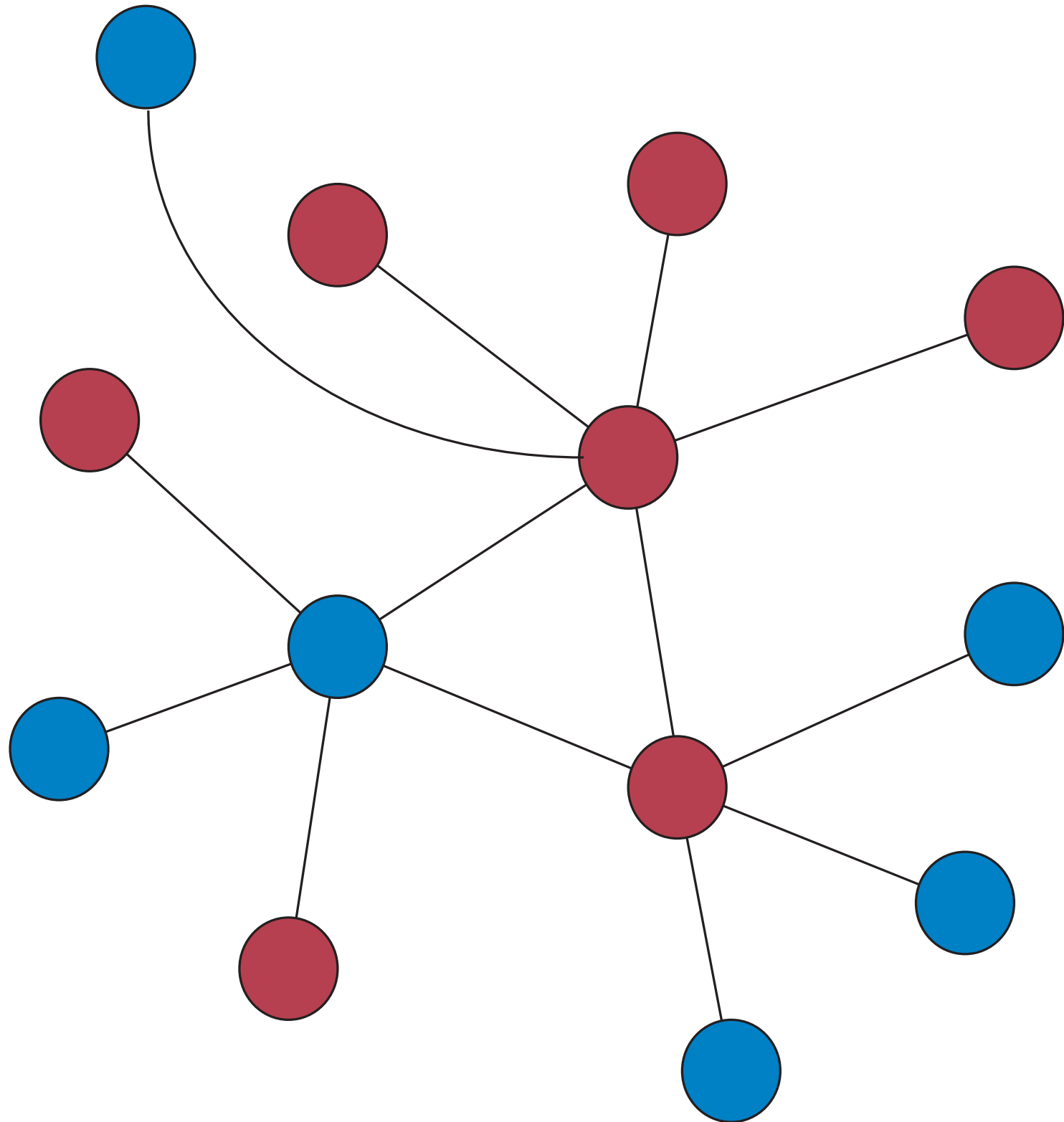
3. Perhaps we can forecast outbreaks



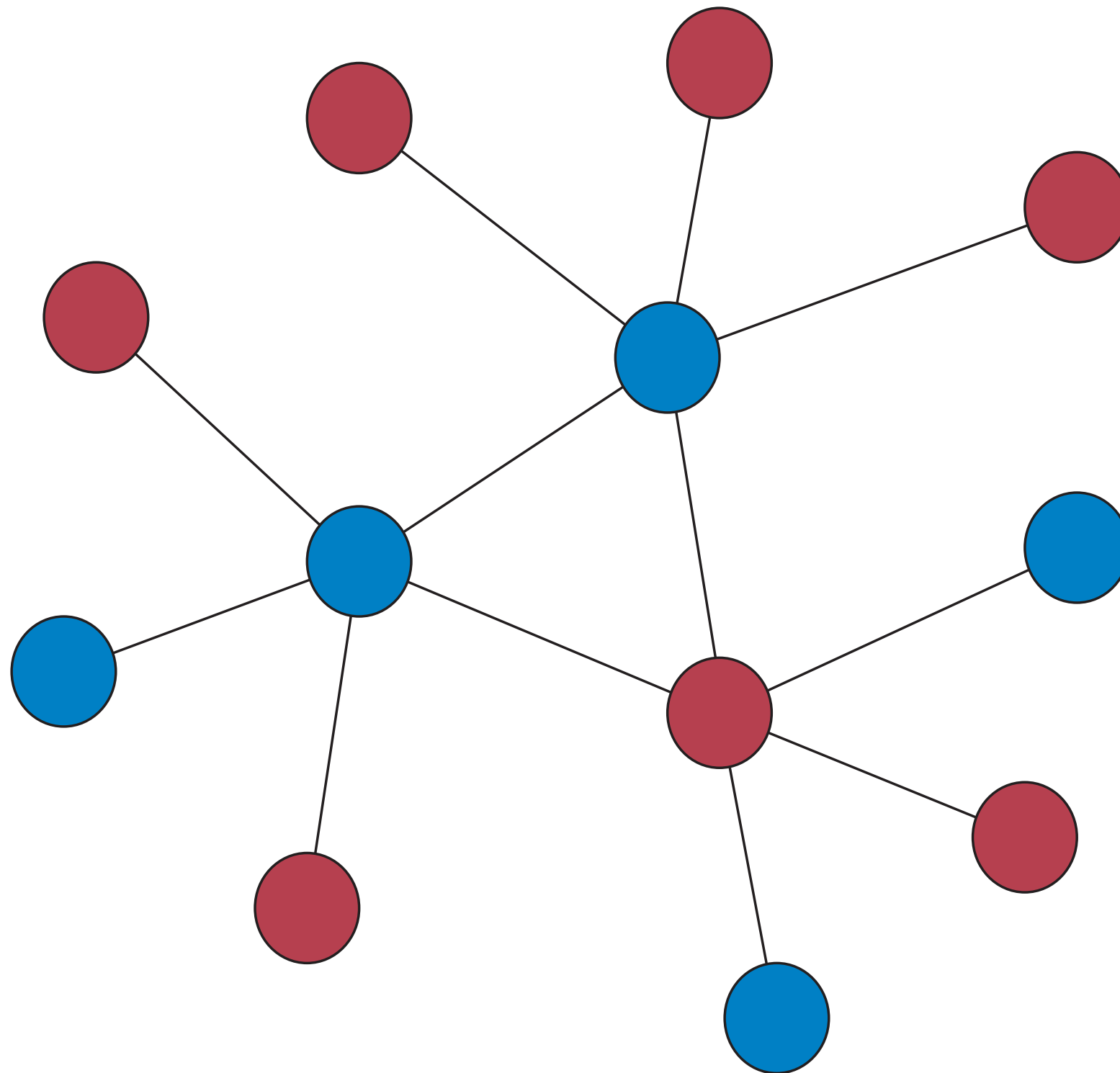
What happens when you replace sick workers?



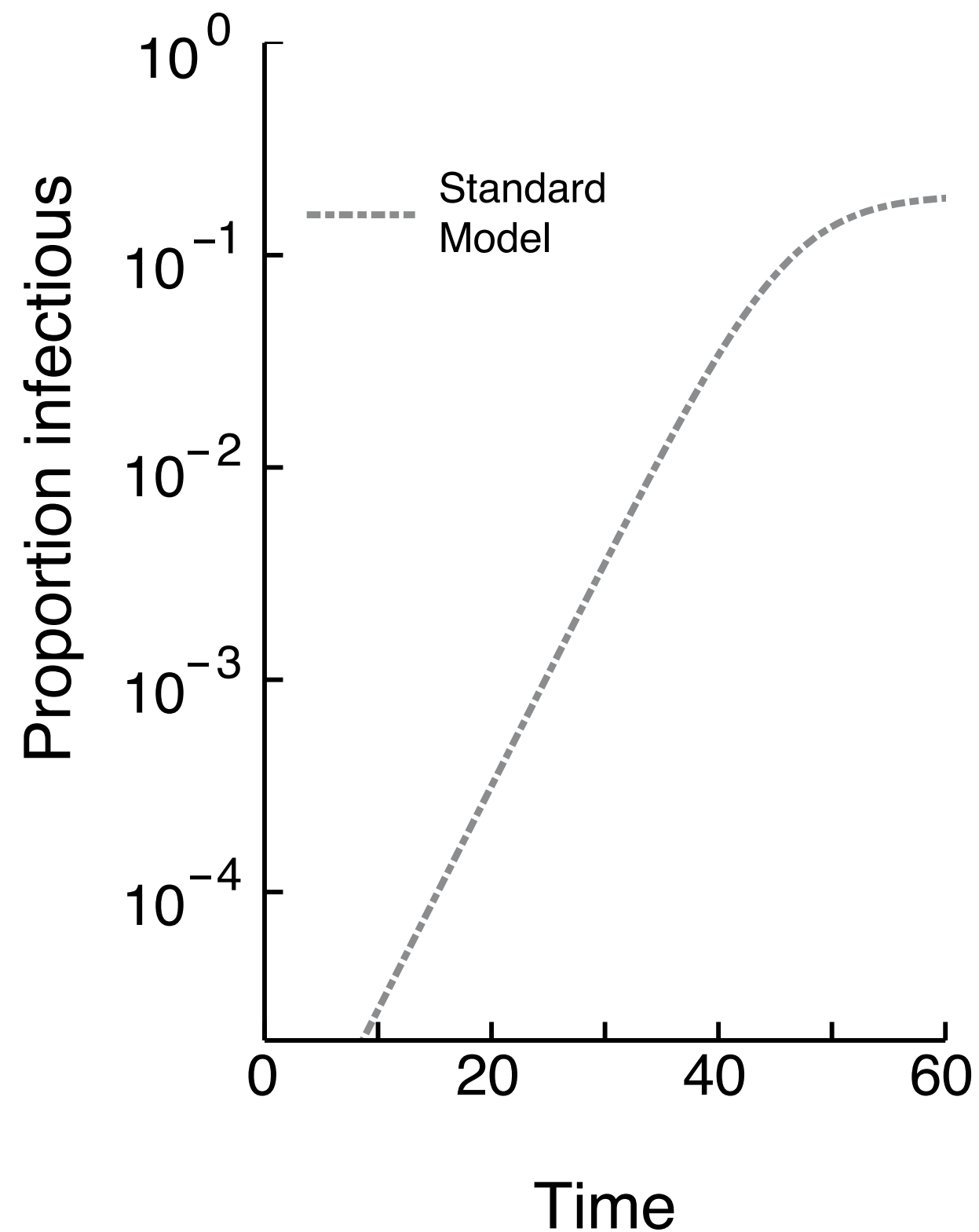
What happens when you replace sick workers?



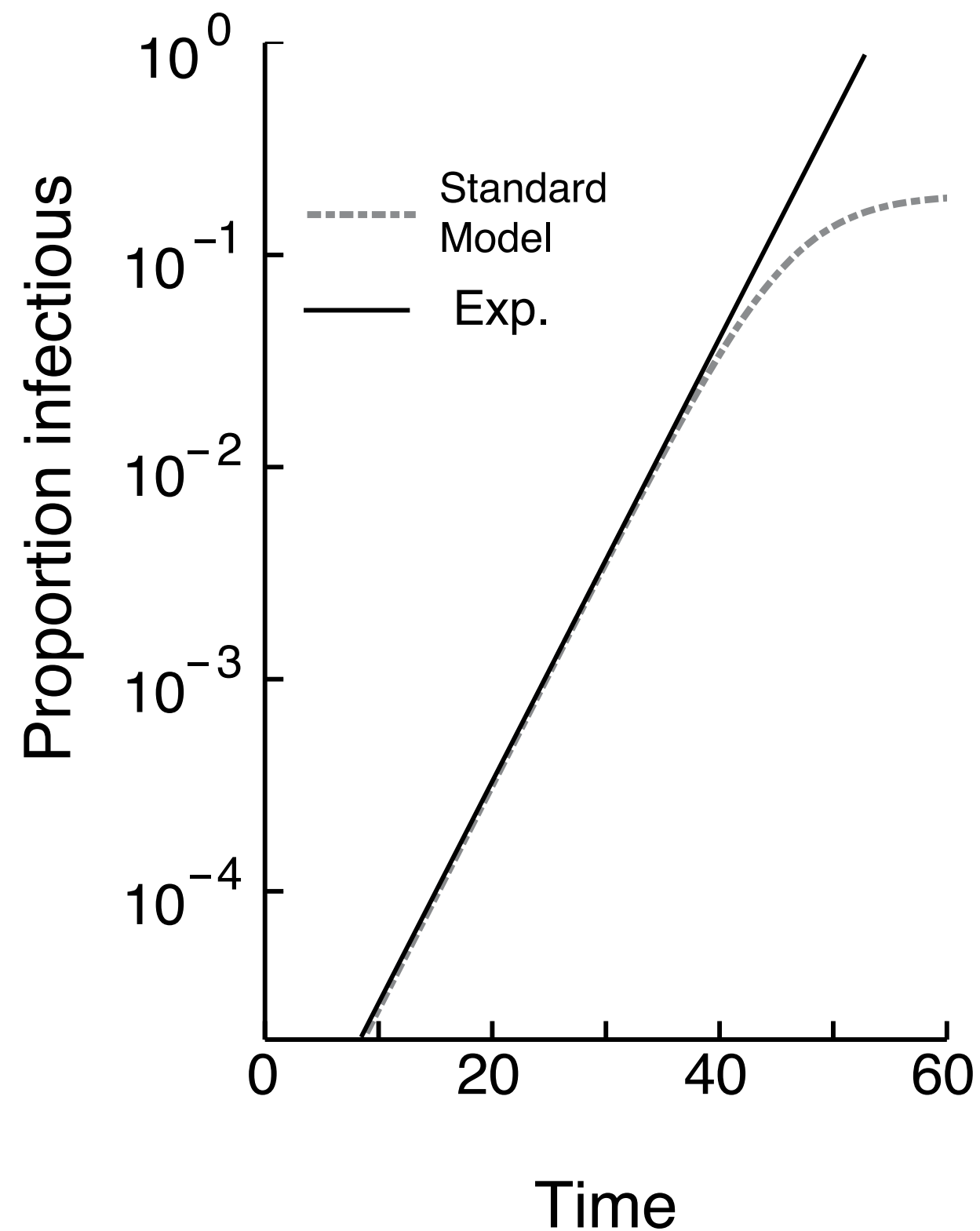
What happens when you replace sick workers?



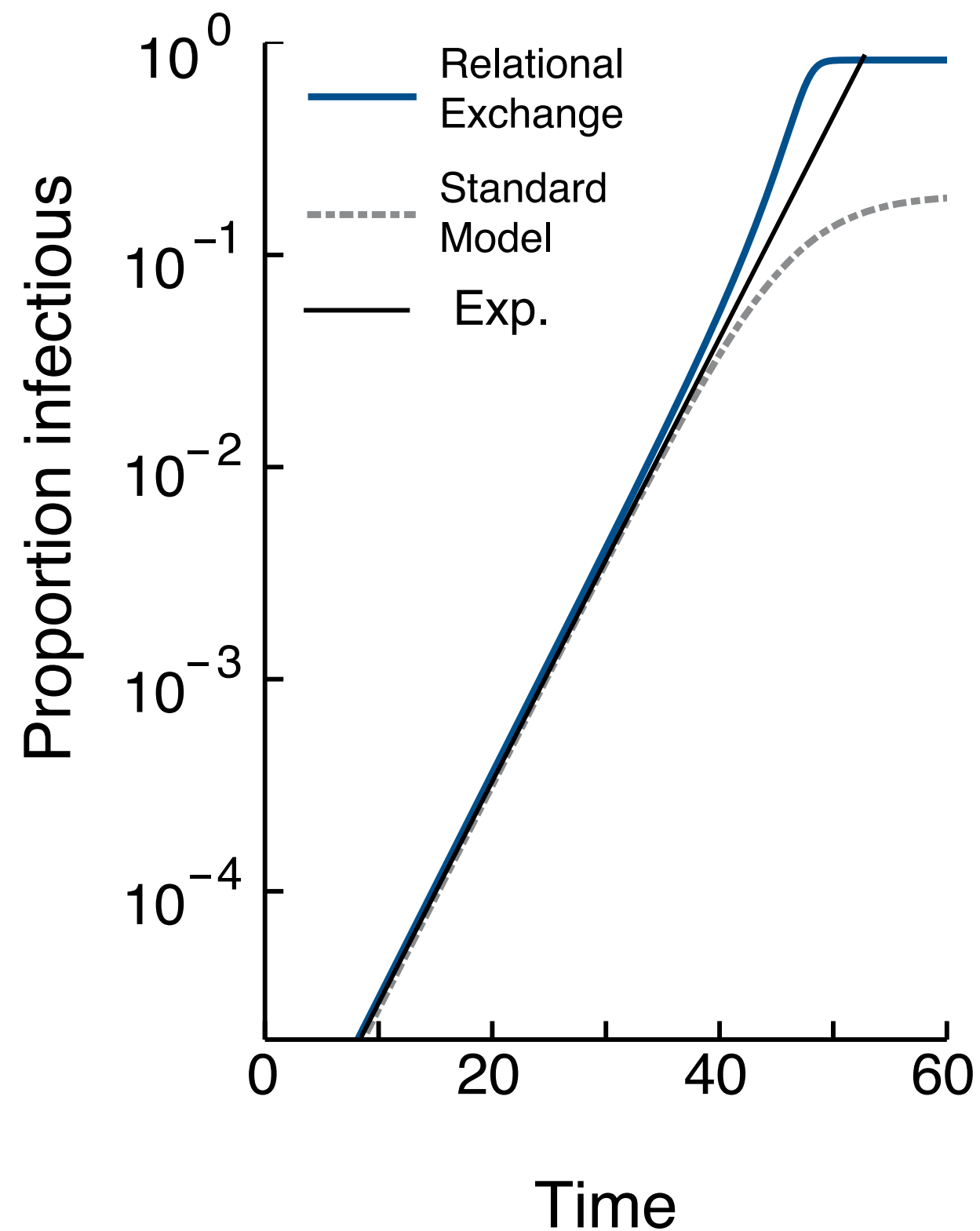
How does *Relational Exchange* work?



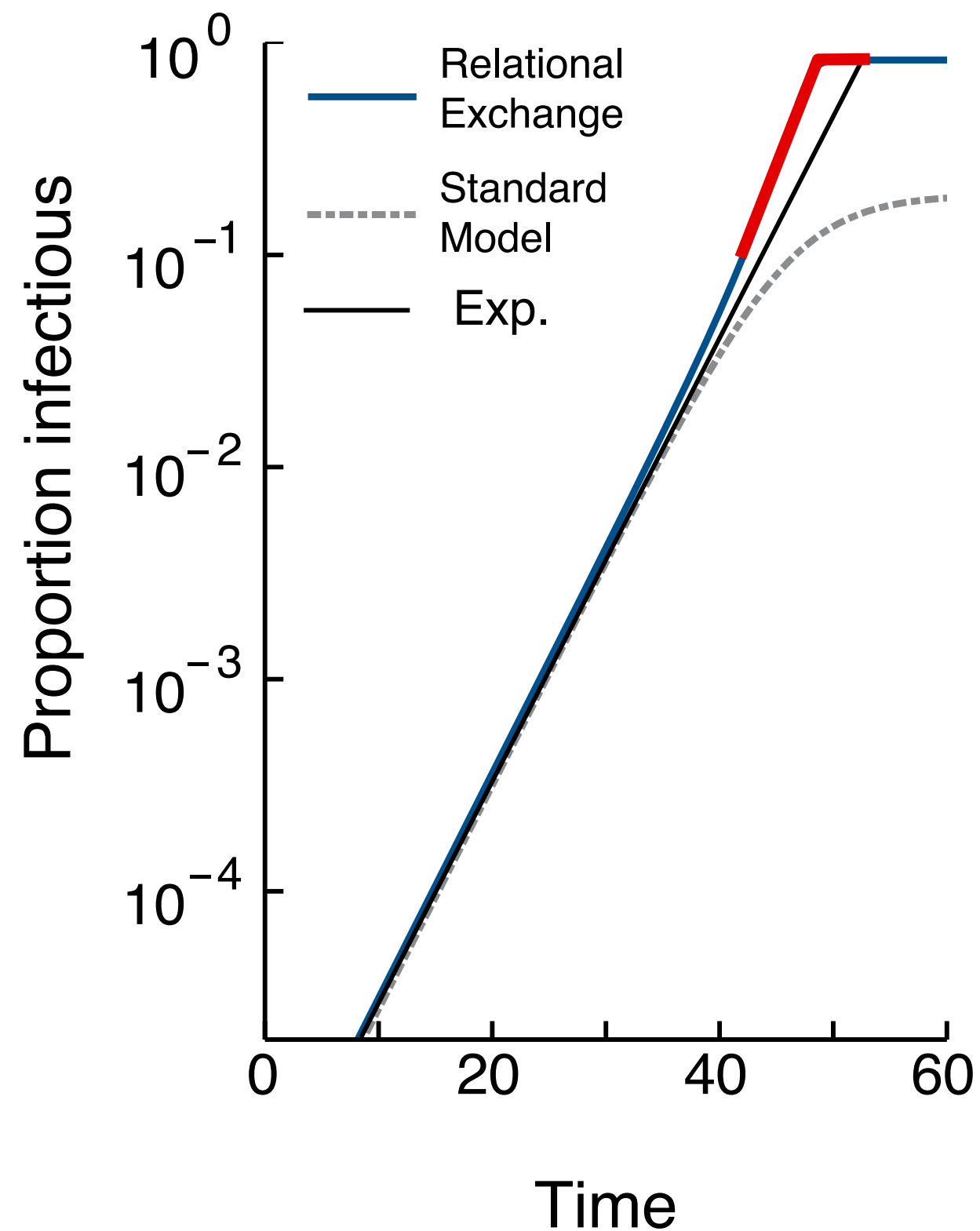
How does *Relational Exchange* work?



Accelerating exponential growth

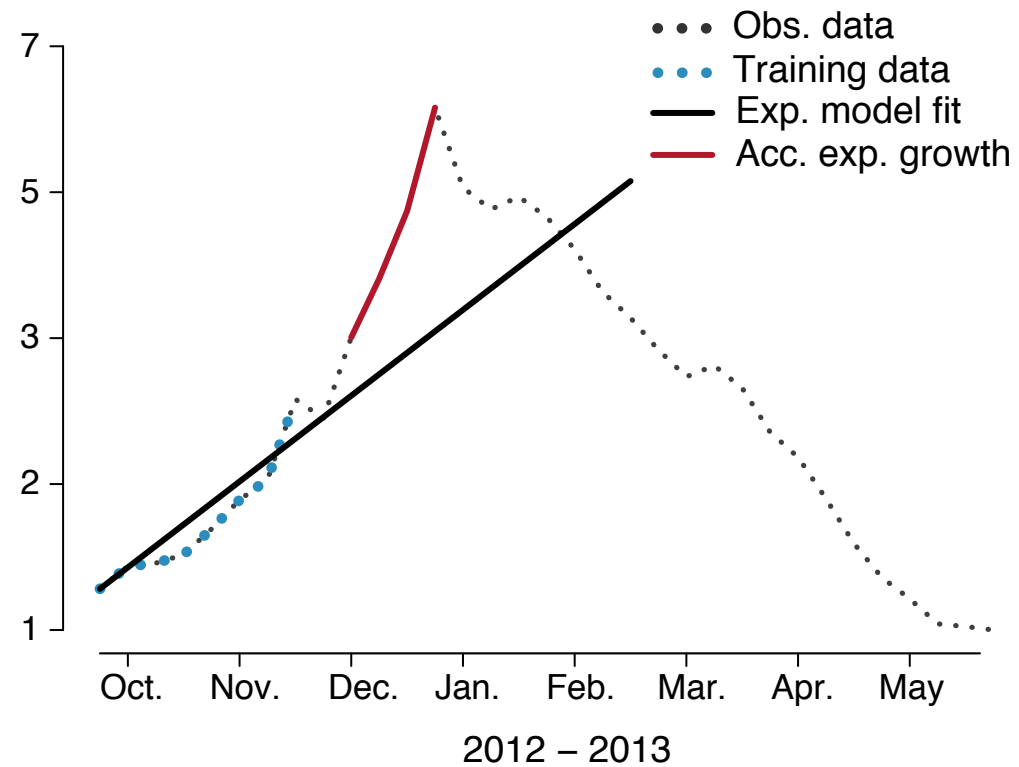


Accelerating exponential growth

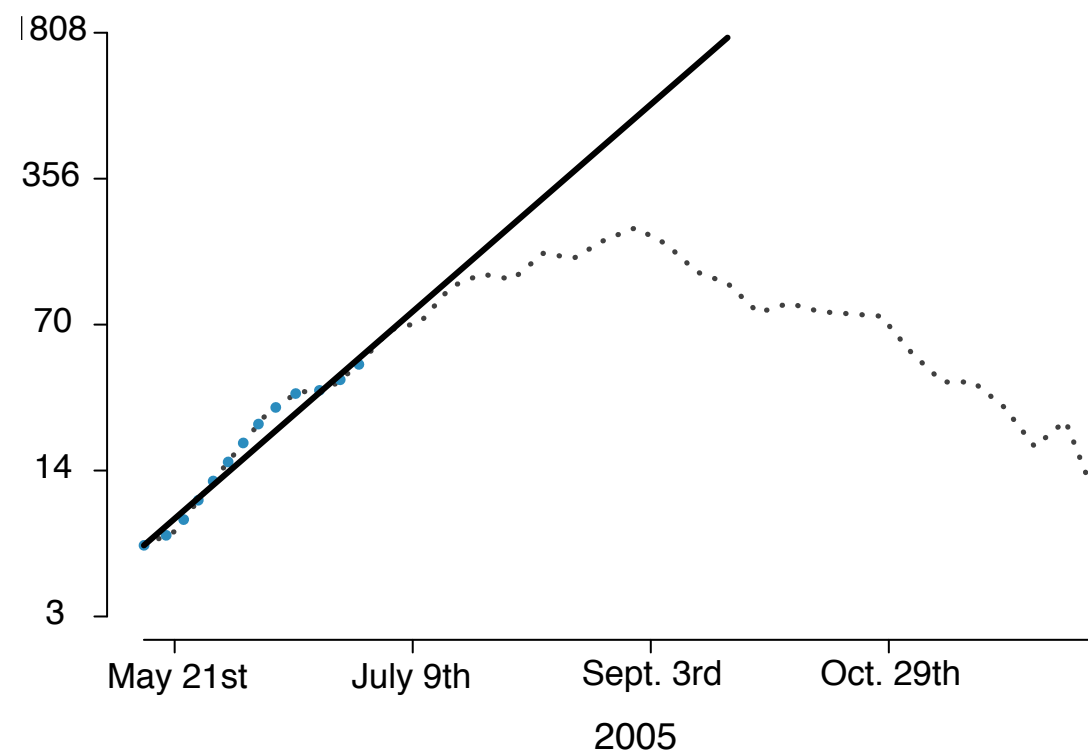


Empirical evidence for *Relational Exchange*

Empirical evidence for *Relational Exchange*

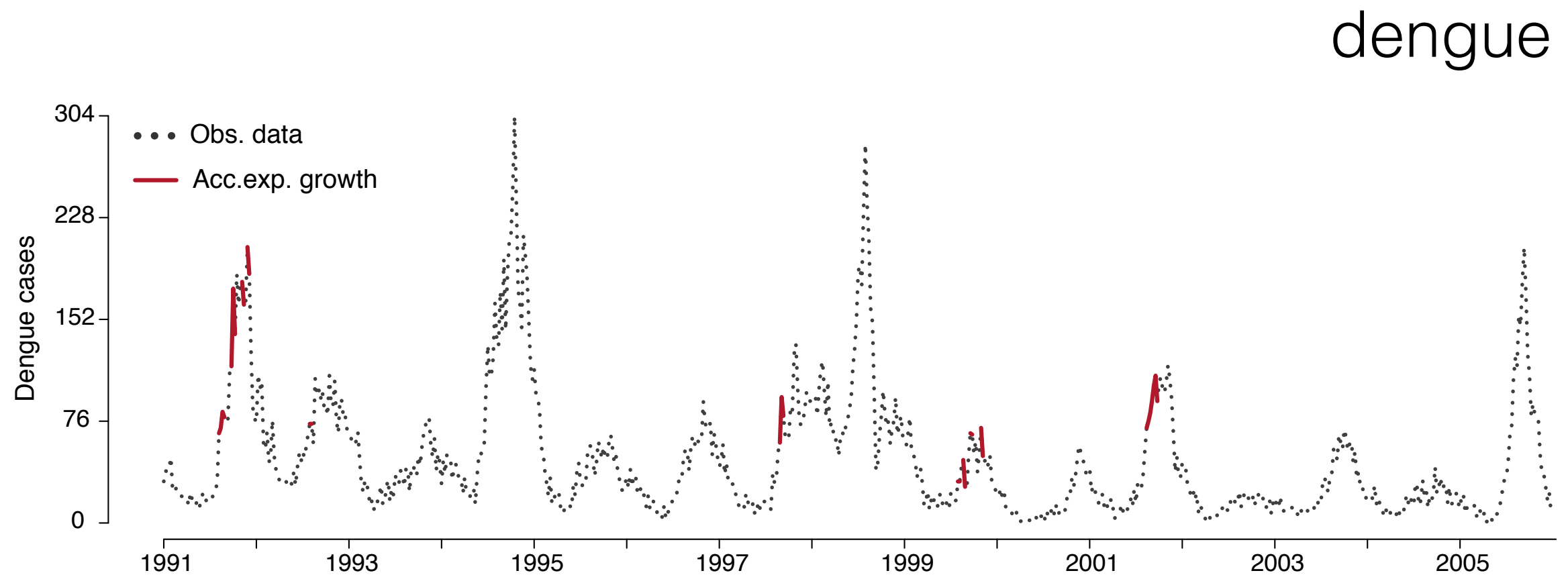
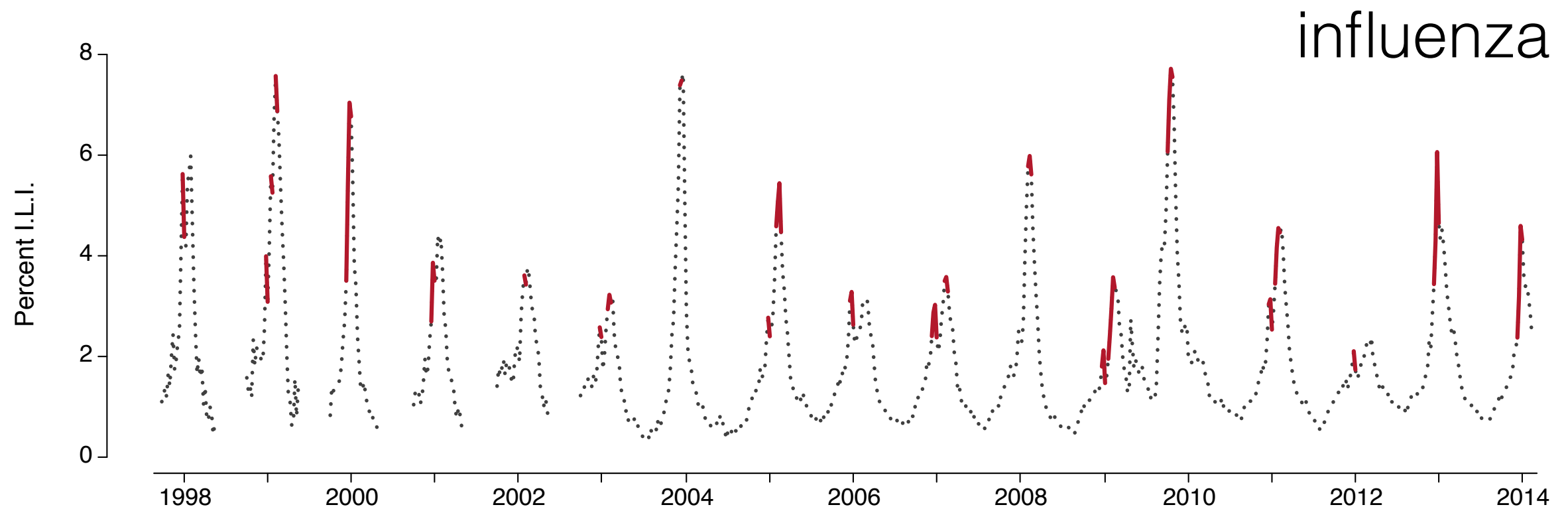


influenza



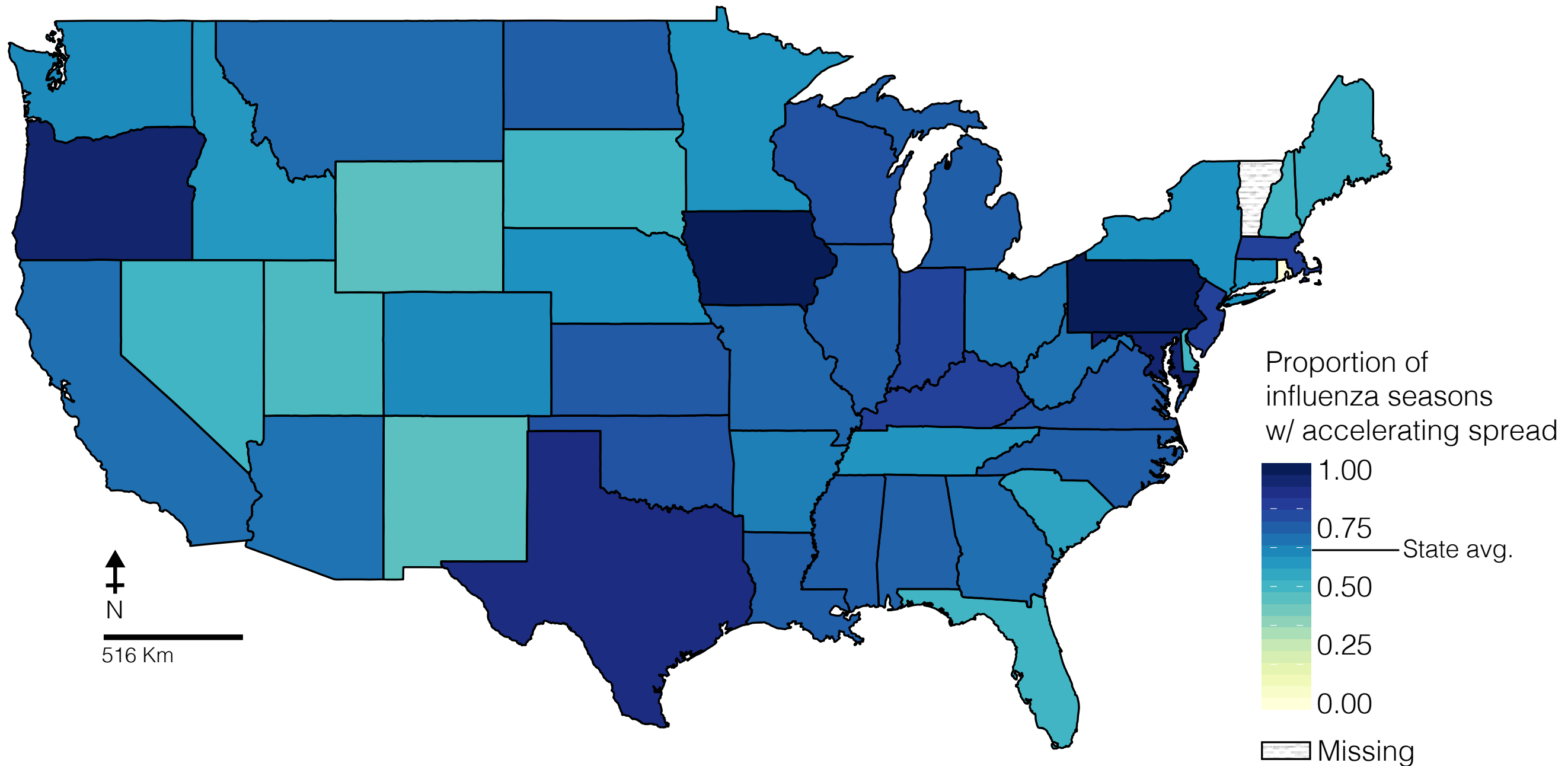
dengue

Empirical evidence for *Relational Exchange*



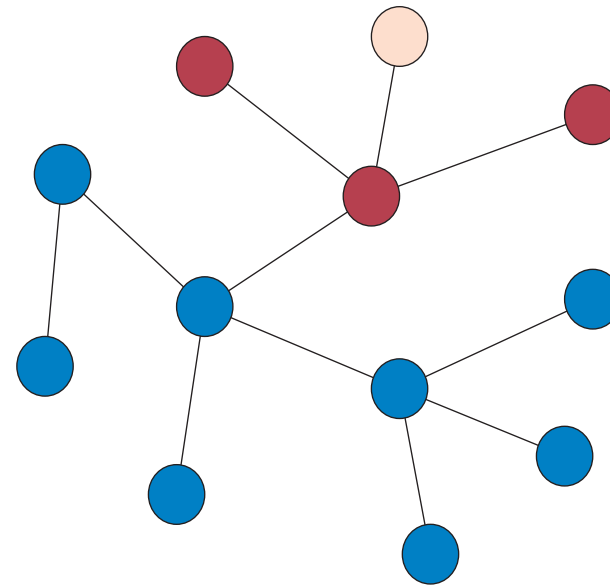
Empirical evidence for *Relational Exchange*

Influenza in the U.S.A. 1926 - 1951



Why do predictions fail?

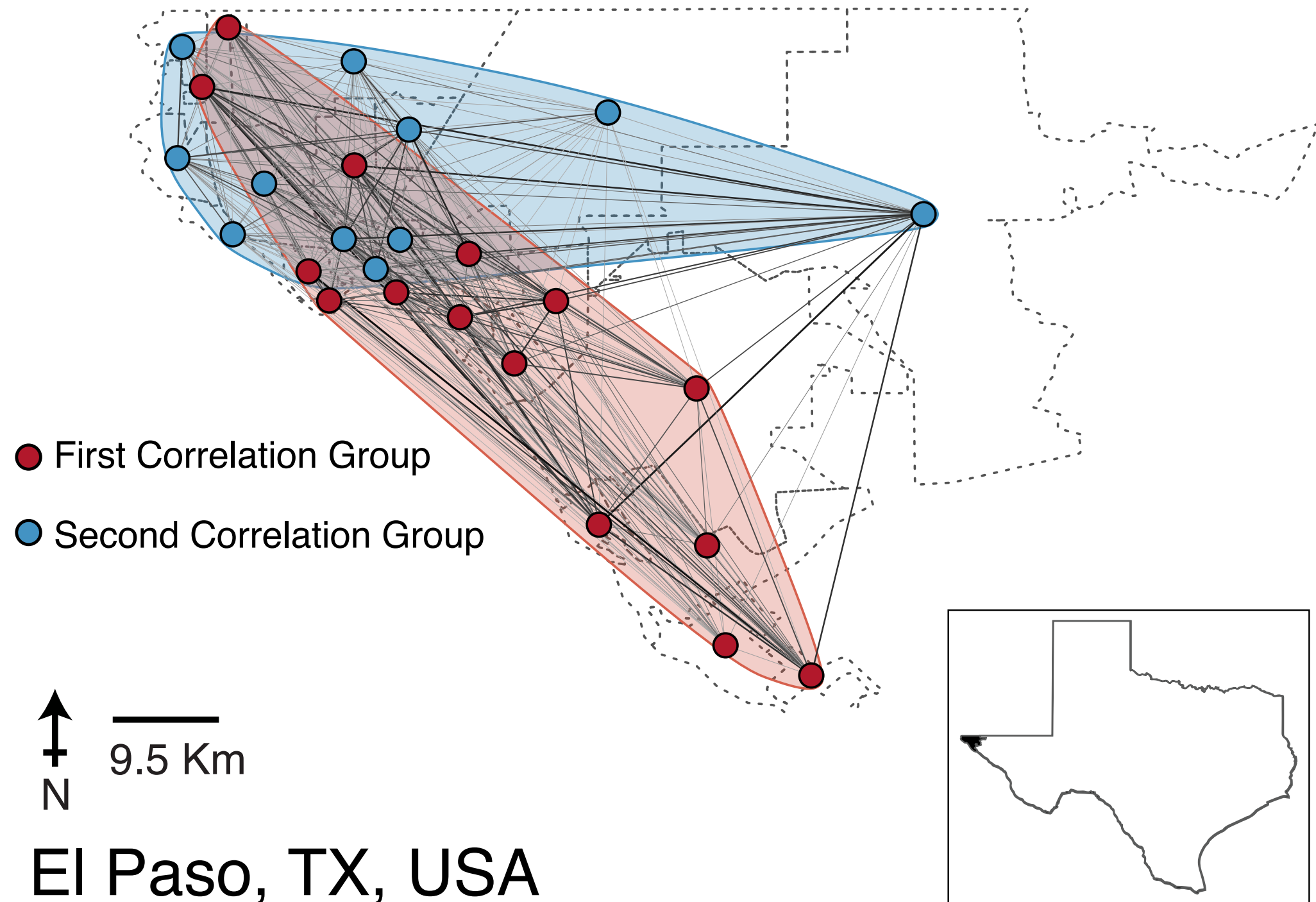
$$R_0 = T \left(\frac{\langle K^2 \rangle - \langle K \rangle}{\langle K \rangle} \right)$$



$$R_0 = \frac{\beta S}{\gamma}$$



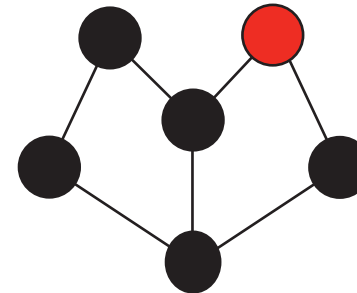
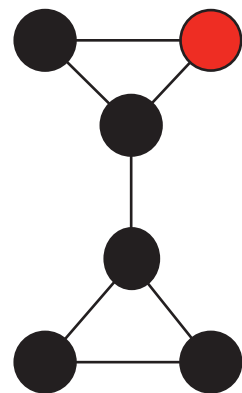
Why do predictions fail?



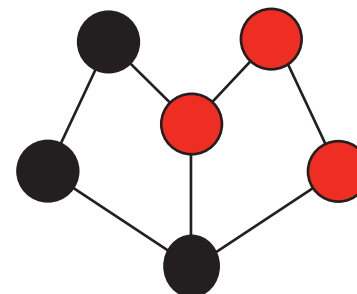
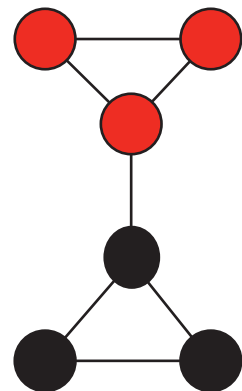
Why do predictions fail?

High clustering Low clustering

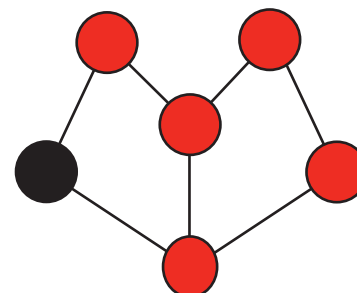
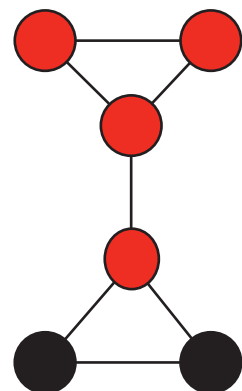
t0



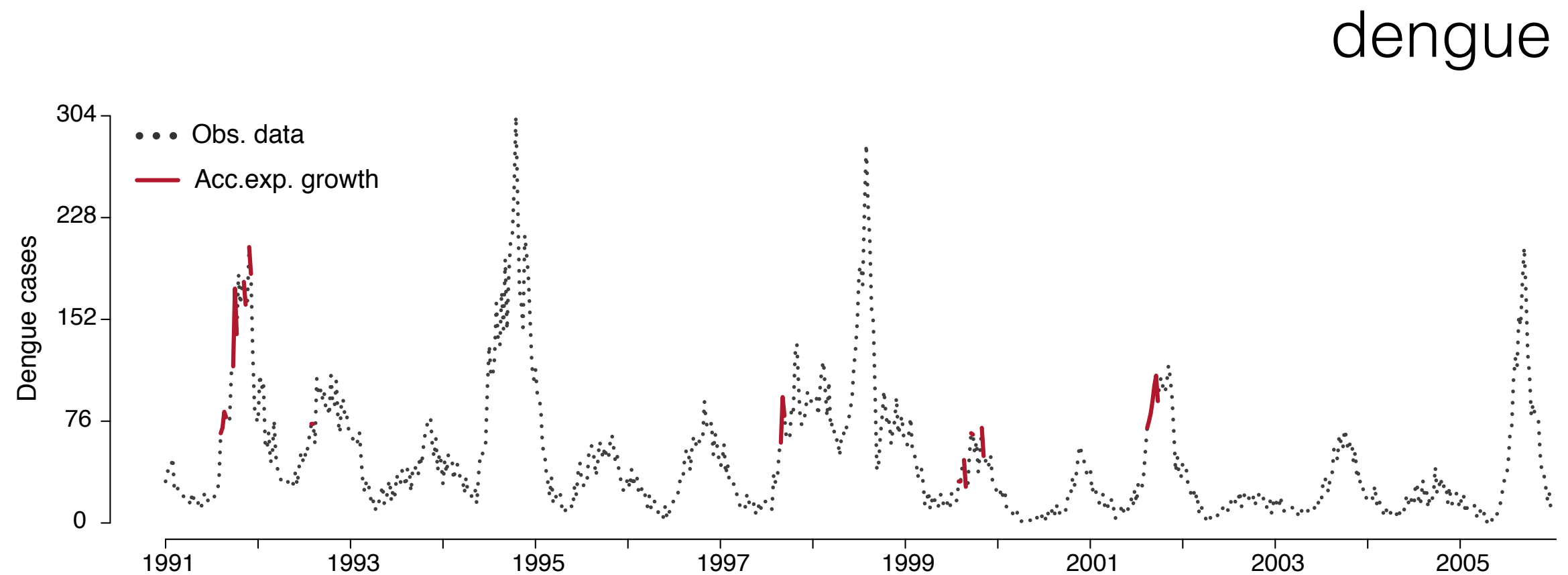
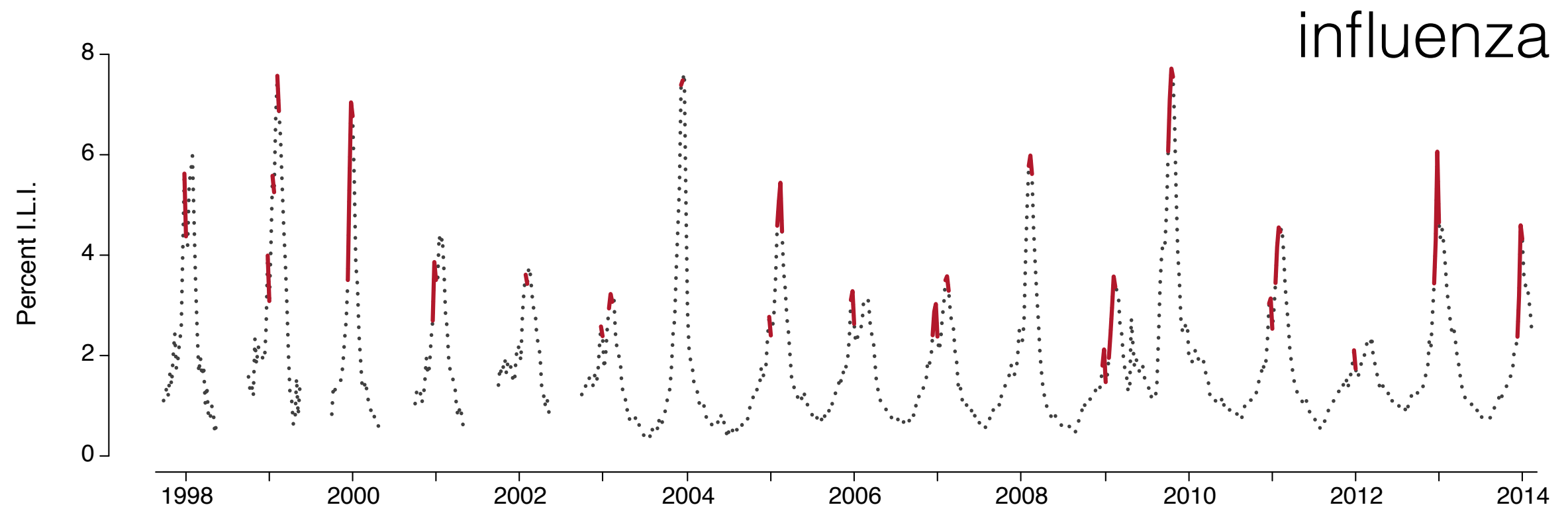
t1



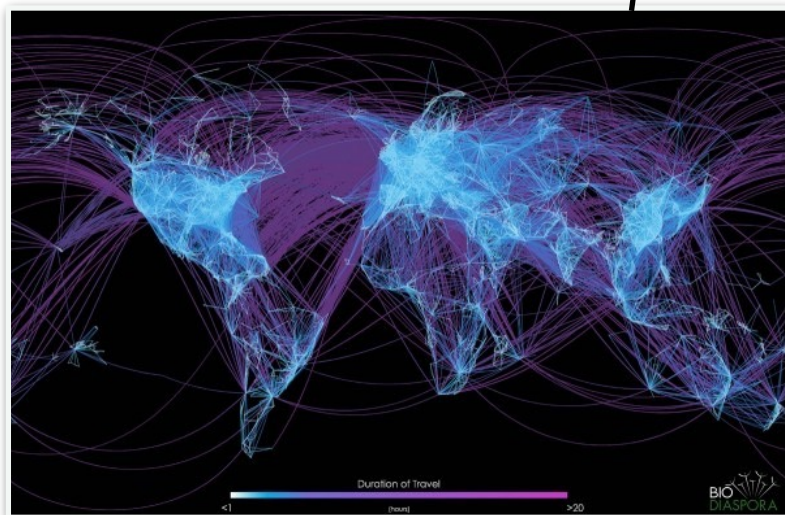
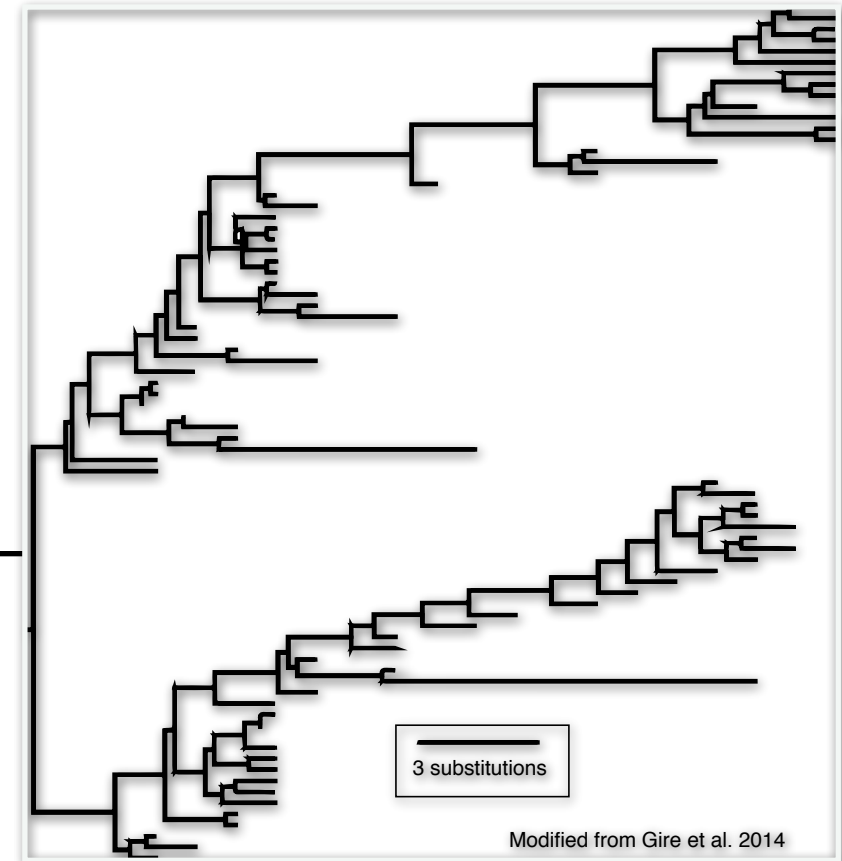
t2



Why do predictions fail?



Toward a complex systems theory of disease



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