

Biological Contagion

Principles of Complex Systems

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Biological
Contagion

**Sealie &
Lambie
Productions**



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Outline

Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

A confusion of contagions:

- ▶ Was Harry Potter some kind of virus?
- ▶ What about the Da Vinci Code?
- ▶ Did Sudoku spread like a disease?
- ▶ Language? The alphabet? [7]
- ▶ Religion?
- ▶ Democracy...?



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

A confusion of contagions:

- ▶ Was Harry Potter some kind of virus?
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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

A confusion of contagions:

- ▶ Was Harry Potter some kind of virus?
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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

A confusion of contagions:

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

A confusion of contagions:

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

A confusion of contagions:

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

A confusion of contagions:

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

Naturomorphisms

- ▶ “The feeling was contagious.”
- ▶ “The news spread like wildfire.”
- ▶ “Freedom is the most contagious virus known to man.”
—Hubert H. Humphrey, Johnson’s vice president
- ▶ “Nothing is so contagious as enthusiasm.”
—Samuel Taylor Coleridge



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

Optimism according to Ambrose Bierce: (田)

The doctrine that everything is beautiful, including what is ugly, everything good, especially the bad, and everything right that is wrong. ...



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

Optimism according to Ambrose Bierce: (田)

The doctrine that everything is beautiful, including what is ugly, everything good, especially the bad, and everything right that is wrong. ... **It is hereditary, but fortunately not contagious.**



Eric Hoffer, 1902–1983

There is a grandeur in the uniformity of the mass.

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



The spread of fanaticism

Hoffer's acclaimed work: "**The True Believer:**
Thoughts On The Nature Of Mass Movements" (1951) [8]

Quotes-aplenty:

- ▶ "We can be absolutely certain only about things we do not understand."
- ▶ "Mass movements can rise and spread without belief in a God, but never without belief in a devil."
- ▶ "Where freedom is real, equality is the passion of the masses. Where equality is real, freedom is the passion of a small minority."

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



The spread of fanaticism

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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The spread of fanaticism

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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The spread of fanaticism

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



CONFORMITY

WHEN PEOPLE ARE FREE TO DO AS THEY PLEASE,
THEY USUALLY IMITATE EACH OTHER.

www.despair.com

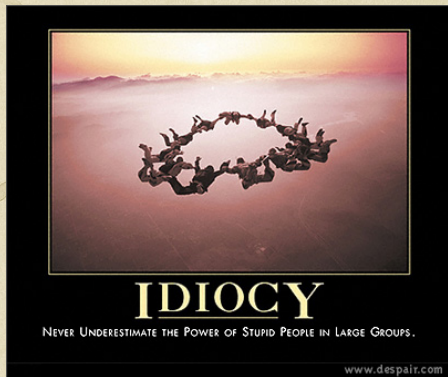
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“When people are free to do as they please, they usually imitate each other.”

—Eric Hoffer
“The Passionate State of Mind” [9]



The collective...



despair.com

“Never Underestimate
the Power of Stupid
People in Large
Groups.”

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Definitions

- ▶ (1) The spreading of a quality or quantity between individuals in a population.
- ▶ (2) A disease itself:
the plague, a blight, the dreaded lurgi, ...
- ▶ from Latin: *con* = 'together with' + *tangere* 'to touch.'
- ▶ Contagion has unpleasant overtones...
- ▶ Just **Spreading** might be a more neutral word
- ▶ But contagion is kind of exciting...

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

Definitions

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- ▶ But contagion is kind of exciting...



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

Definitions

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 - ▶ But contagion is kind of exciting...



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

Definitions

- ▶ (1) The spreading of a quality or quantity between individuals in a population.
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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

Definitions

- ▶ (1) The spreading of a quality or quantity between individuals in a population.
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Examples of non-disease spreading:

Interesting infections:

- ▶ [Spreading of buildings in the US... \(田\)](#)



- ▶ [Viral get-out-the-vote video. \(田\)](#)

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

Two main classes of contagion

1. Infectious diseases
2. Social contagion



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

Two main classes of contagion

1. **Infectious diseases**

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

Two main classes of contagion

1. **Infectious diseases:**
tuberculosis, HIV, ebola, SARS, influenza, ...
2. **Social contagion**



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

Two main classes of contagion

1. **Infectious diseases:**
tuberculosis, HIV, ebola, SARS, influenza, ...
2. **Social contagion:**
fashion, word usage, rumors, riots, religion, ...



Outline

Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References

Biological
Contagion

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



The standard **SIR model** ^[14]

- ▶ = basic model of disease contagion
- ▶ Three states:
 1. **S** = Susceptible
 2. **I** = Infective/Infectious
 3. **R** = Recovered or Removed or Resistant
- ▶ $S(t) + I(t) + R(t) = 1$
- ▶ Presumes random interactions (mass-action principle)
- ▶ Interactions are independent (no memory)
- ▶ Discrete and continuous time versions

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

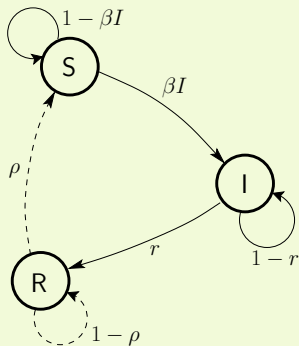
Model output

Conclusions

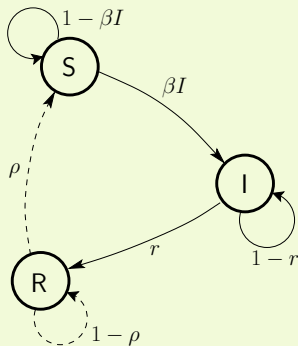
Predicting social
catastrophe

References

Discrete time automata example:



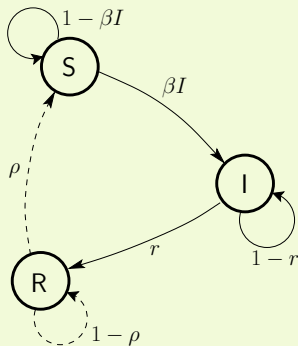
Discrete time automata example:



Transition Probabilities:



Discrete time automata example:

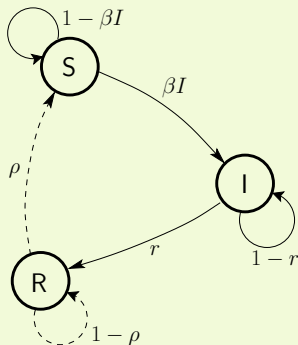


Transition Probabilities:

β for being infected given
contact with infected



Discrete time automata example:



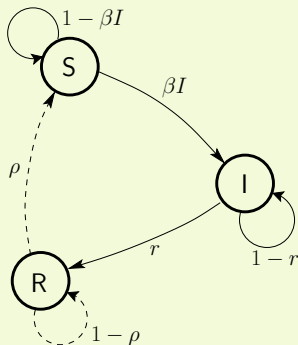
Transition Probabilities:

β for being infected given
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r for recovery



Discrete time automata example:



Transition Probabilities:

β for being infected given

contact with infected

r for recovery

ρ for loss of immunity



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

Original models attributed to

- ▶ 1920's: Reed and Frost
- ▶ 1920's/1930's: Kermack and McKendrick^[10, 12, 11]
- ▶ Coupled differential equations with a mass-action principle



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Independent Interaction models

Differential equations for continuous model

$$\frac{d}{dt}S = -\beta IS + \rho R$$

$$\frac{d}{dt}I = \beta IS - rI$$

$$\frac{d}{dt}R = rI - \rho R$$

β , r , and ρ are now **rates**.

Reproduction Number R_0 :

- ▶ R_0 = expected number of infected individuals resulting from a single initial infective
- ▶ Epidemic threshold: If $R_0 > 1$, 'epidemic' occurs.

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Reproduction Number R_0

Discrete version:

- ▶ Set up: One Infective in a randomly mixing population of Susceptibles
- ▶ At time $t = 0$, single infective random bumps into a Susceptible
- ▶ Probability of transmission = β
- ▶ At time $t = 1$, single Infective remains infected with probability $1 - r$
- ▶ At time $t = k$, single Infective remains infected with probability $(1 - r)^k$

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Reproduction Number R_0

Discrete version:

- ▶ Expected number infected by original Infective:

$$R_0 = \beta + (1 - r)\beta + (1 - r)^2\beta + (1 - r)^3\beta + \dots$$

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Reproduction Number R_0

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$$\begin{aligned}R_0 &= \beta + (1 - r)\beta + (1 - r)^2\beta + (1 - r)^3\beta + \dots \\ &= \beta \left(1 + (1 - r) + (1 - r)^2 + (1 - r)^3 + \dots \right)\end{aligned}$$

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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$$= \beta \left(1 + (1 - r) + (1 - r)^2 + (1 - r)^3 + \dots \right)$$

$$= \beta \frac{1}{1 - (1 - r)}$$

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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For S_0 initial infectives ($1 - S_0 = R_0$ immune):

$$R_0 = S_0\beta/r$$

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Independent Interaction models

For the continuous version

- ▶ Second equation:

$$\frac{d}{dt}I = \beta SI - rI$$

- ▶ Number of infectives grows initially if

$$\beta S(0) - r > 0$$

- ▶ Same story as for discrete model.

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

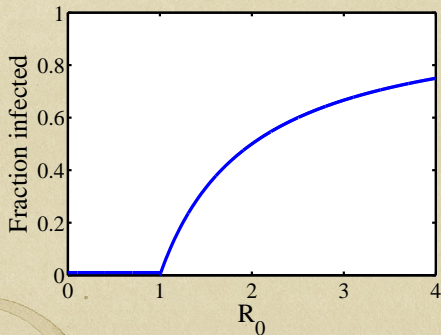
Predicting social
catastrophe

References



Independent Interaction models

Example of epidemic threshold:



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

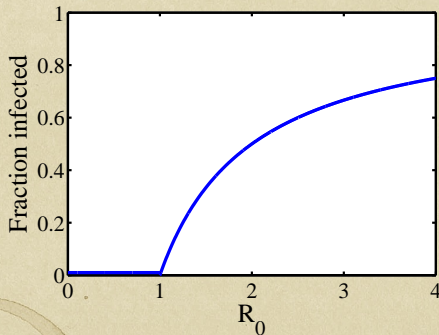
Predicting social
catastrophe

References



Independent Interaction models

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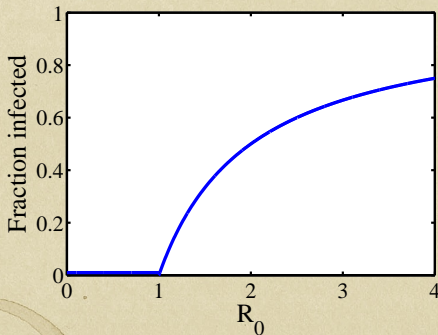


- ▶ Continuous phase transition.



Independent Interaction models

Example of epidemic threshold:



- ▶ Continuous phase transition.
- ▶ Fine idea from a simple model.



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

Many variants of the SIR model:

- ▶ **SIS**: susceptible-infective-susceptible
- ▶ **SIRS**: susceptible-infective-recovered-susceptible
- ▶ compartment models (age or gender partitions)
- ▶ more categories such as 'exposed' (**SEIRS**)
- ▶ recruitment (migration, birth)



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Outline

Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References

Biological
Contagion

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Disease spreading models

For novel diseases:

1. Can we predict the size of an epidemic?
2. How important is the reproduction number R_0 ?

R_0 approximately same for all of the following:

- ▶ 1918-19 "Spanish Flu" ~ 500,000 deaths in US
- ▶ 1957-58 "Asian Flu" ~ 70,000 deaths in US
- ▶ 1968-69 "Hong Kong Flu" ~ 34,000 deaths in US
- ▶ 2003 "SARS Epidemic" ~ 800 deaths world-wide

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Size distributions

Size distributions are important elsewhere:

- ▶ earthquakes (Gutenberg-Richter law)
- ▶ city sizes, forest fires, war fatalities
- ▶ wealth distributions
- ▶ 'popularity' (books, music, websites, ideas)
- ▶ Epidemics?

Really, what about epidemics?

- ▶ Simply hasn't attracted much attention.
- ▶ Data not as clean as for other phenomena.

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

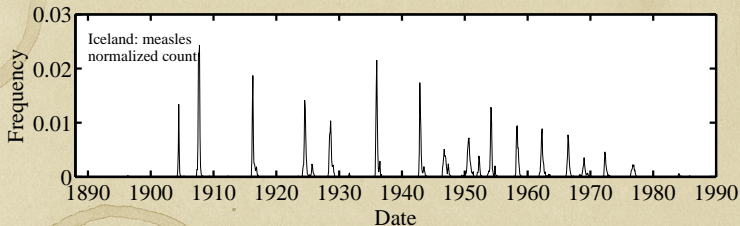
Predicting social
catastrophe

References



Feeling Ill in Iceland

Caseload recorded monthly for range of diseases in Iceland, 1888-1990



- ▶ Treat outbreaks separated in time as 'novel' diseases.

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

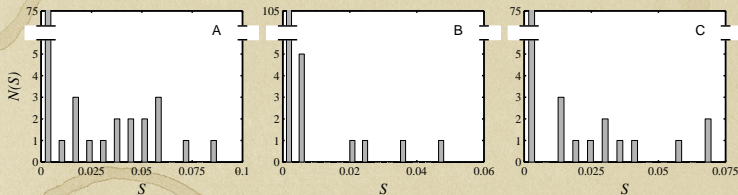
Predicting social
catastrophe

References



Really not so good at all in Iceland

Epidemic size distributions $N(S)$ for
Measles, Rubella, and Whooping Cough.

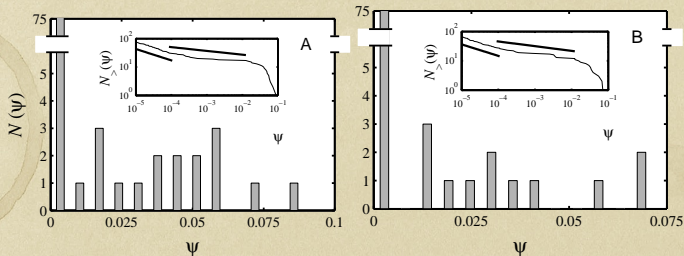


Spike near $S = 0$, relatively flat otherwise.



Measles & Pertussis

Biological
Contagion



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

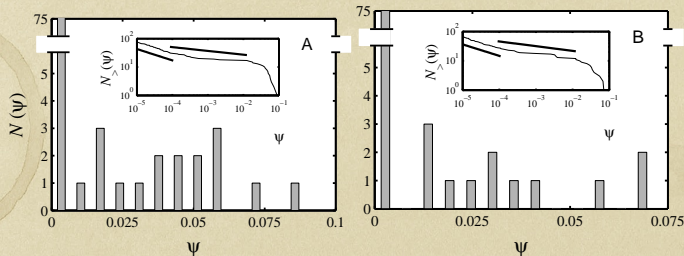
Conclusions

Predicting social
catastrophe

References



Measles & Pertussis



Insert plots:

Complementary cumulative frequency distributions:

$$N(\psi' > \psi) \propto \psi^{-\gamma+1}$$

Limited scaling with a possible break.

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Power law distributions

Measured values of γ :

- ▶ measles: 1.40 (low Ψ) and 1.13 (high Ψ)
- ▶ pertussis: 1.39 (low Ψ) and 1.16 (high Ψ)

- ▶ Expect $2 \leq \gamma < 3$ (finite mean, infinite variance)
- ▶ When $\gamma < 1$, can't normalize
- ▶ Distribution is quite flat.



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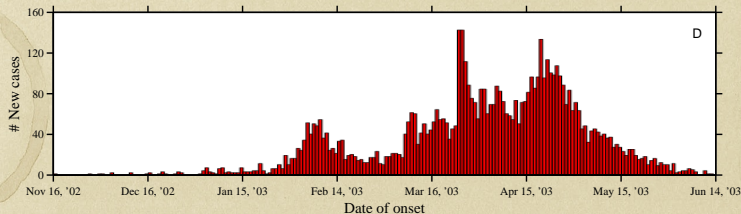
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Resurgence—example of SARS



- ▶ Epidemic slows...
- ▶ Epidemic discovers new 'pools' of susceptibles:
Resurgence.
- ▶ Importance of rare, stochastic events.

Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

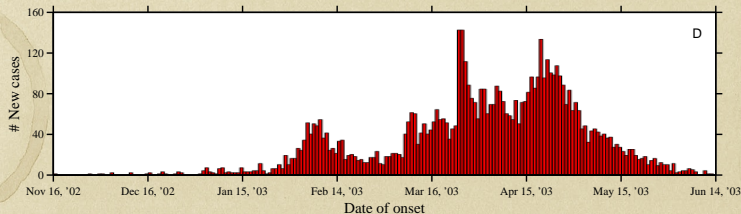
Conclusions

Predicting social catastrophe

References



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Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

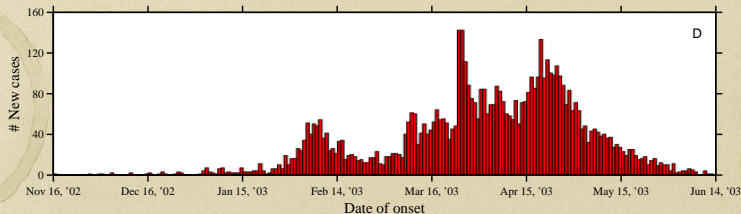
Conclusions

Predicting social catastrophe

References



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Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

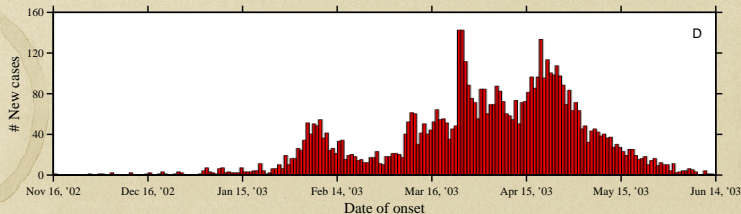
Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

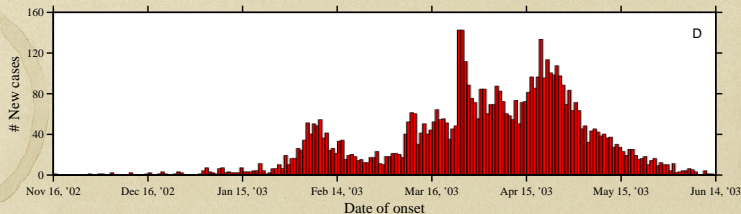
Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Outline

Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References

Biological
Contagion

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



The challenge

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

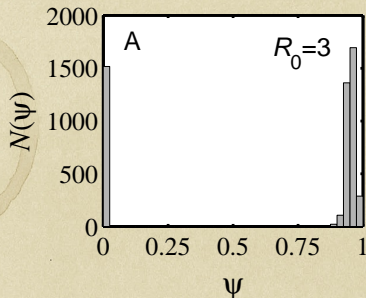
References

So... can a simple model produce

1. **broad epidemic distributions**
and
2. **resurgence ?**



Size distributions

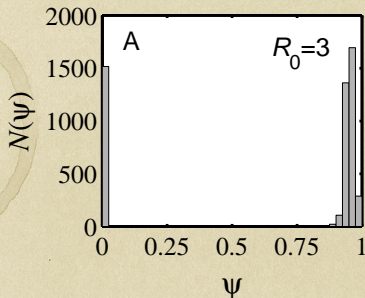


Simple models typically produce **bimodal** or **unimodal** size distributions.

- ▶ This includes network models: random, small-world, scale-free, ...
- ▶ Exceptions:
 1. Forest fire models
 2. Sophisticated metapopulation models



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

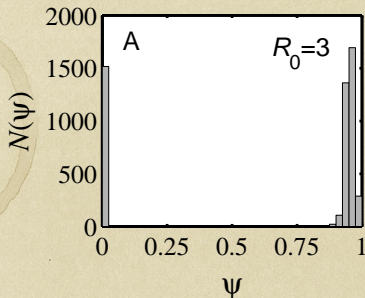
Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

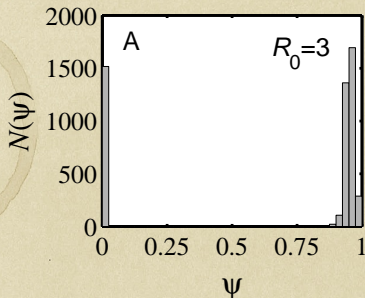
Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

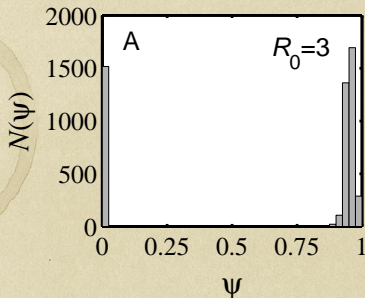
Conclusions

Predicting social catastrophe

References



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Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References



Burning through the population

Forest fire models:^[15]

- ▶ Rhodes & Anderson, 1996
- ▶ The physicist's approach:
"if it works for magnets, it'll work for people..."

A bit of a stretch:

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Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References



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Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References



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Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References



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Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References



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Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References



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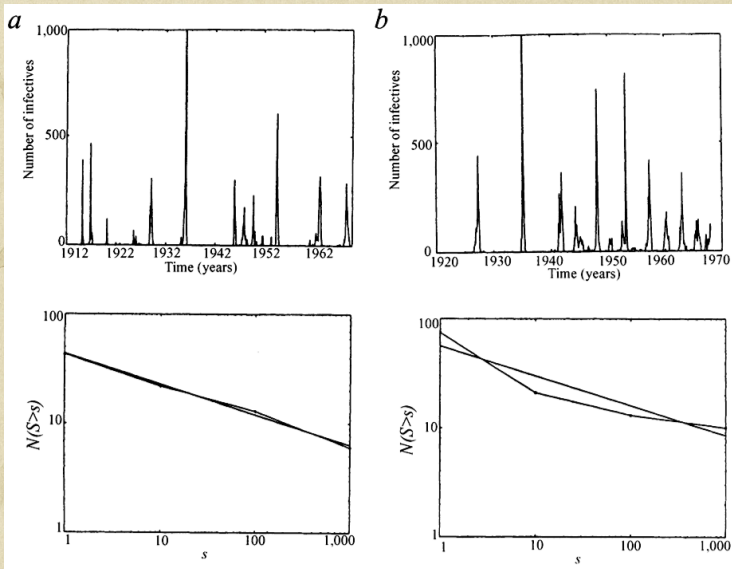
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Size distributions



From Rhodes and Anderson, 1996.

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Sophisticated metapopulation models:

- ▶ Multiscale models suggested earlier by others but not formalized (Bailey ^[1], Cliff and Haggett ^[4], Ferguson et al.)
- ▶ Community based mixing (two scales)—Longini. ^[13]
- ▶ Eubank et al.'s EpiSims/TRANSIMS—city simulations. ^[6]
- ▶ Spreading through countries—Airlines: Germann et al., Colizza et al. ^[5]

Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References

GLEAMviz
The Global Epidemic and Mobility Model

NEWS TEAM PUBLICATIONS PRESS

VISION CHALLENGES APPROACH MODEL SIMULATOR CASE STUDY

A MULTIDISCIPLINARY APPROACH TO EPIDEMIC ANALYSIS

$$D_i^c = \sum_j D_{ij}^{c-1} (k_j - 1) P_{ij}(k_j) (1 - R_0^{-1/k_j}) \left(1 - \sum_l \frac{D_{il}^c}{P_{il}} \right)$$

- ▶ GLEAM (田):
Global pandemic simulations by Vespignani et al.



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

- ▶ Vital work but perhaps hard to generalize from...
- ▶ ⇒ Create a simple model involving multiscale travel
- ▶ Very big question: **What is N ?**
- ▶ Should we model SARS in Hong Kong as spreading in a neighborhood, in Hong Kong, Asia, or the world?
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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Outline

Biological
Contagion

Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

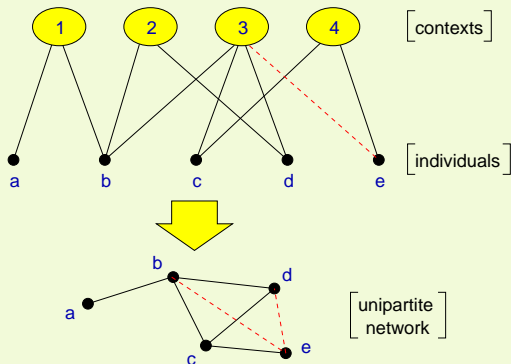
Predicting social
catastrophe

References



Improving simple models

Contexts and Identities—Bipartite networks



- ▶ boards of directors
- ▶ movies
- ▶ transportation modes (subway)

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

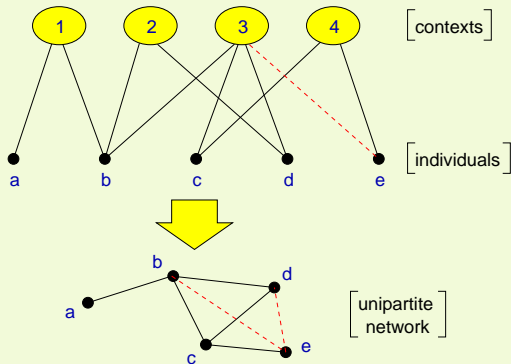
Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

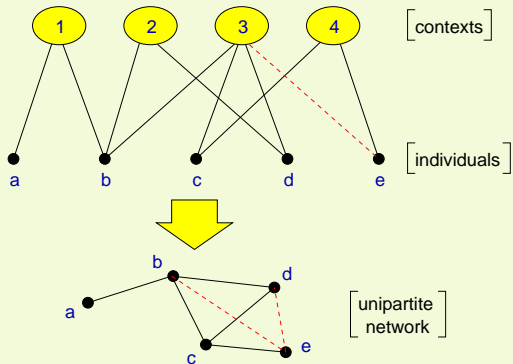
Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

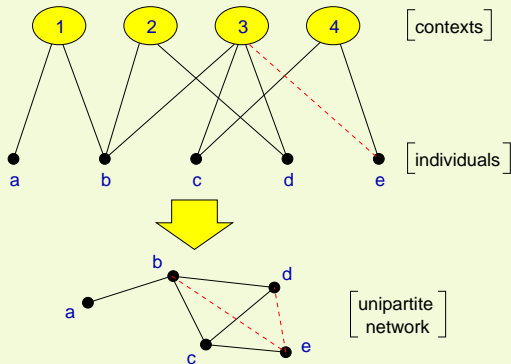
References

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Improving simple models

Idea for social networks: incorporate **identity**.

Identity is formed from attributes such as:

- ▶ Geographic location
- ▶ Type of employment
- ▶ Age
- ▶ Recreational activities

Groups are crucial...

- ▶ formed by people with at least one similar attribute
- ▶ Attributes \leftrightarrow Contexts \leftrightarrow Interactions \leftrightarrow Networks. [17]

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

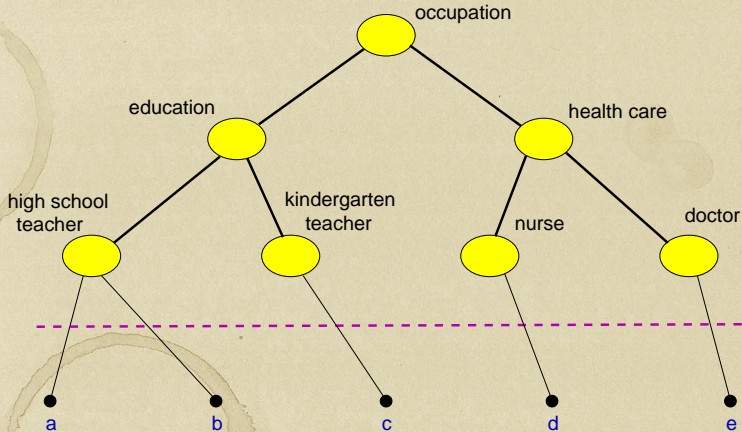
Conclusions

Predicting social
catastrophe

References



Infer interactions/network from identities



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

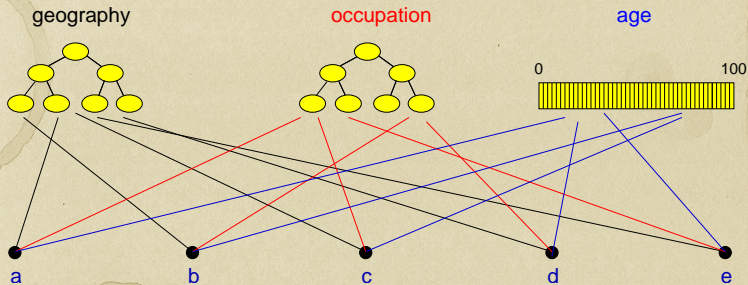
Predicting social
catastrophe

References



Distance makes sense in identity/context space.

Generalized context space



(Blau & Schwartz ^[2], Simmel ^[16], Breiger ^[3])



A toy agent-based model

Geography—allow people to move between contexts:

- ▶ Locally: standard SIR model with random mixing
- ▶ discrete time simulation
- ▶ β = infection probability
- ▶ γ = recovery probability
- ▶ P = probability of travel
- ▶ **Movement distance:** $\Pr(d) \propto \exp(-d/\xi)$
- ▶ ξ = typical travel distance



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- ▶ β = infection probability
- ▶ γ = recovery probability
- ▶ P = probability of travel
- ▶ **Movement distance:** $\Pr(d) \propto \exp(-d/\xi)$
- ▶ ξ = typical travel distance



A toy agent-based model

Geography—allow people to move between contexts:

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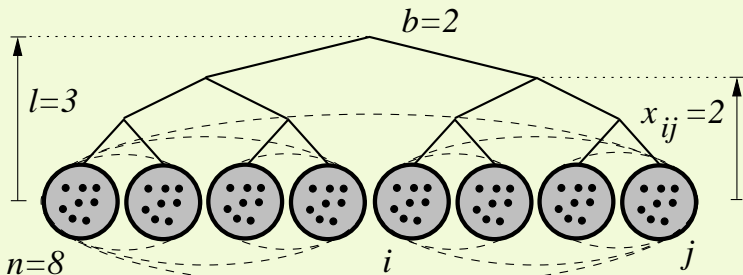
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A toy agent-based model

Schematic:



Outline

Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References

Biological
Contagion

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



- ▶ Define P_0 = Expected number of infected individuals **leaving** initially infected context.
- ▶ Need $P_0 > 1$ for disease to spread (independent of R_0).
- ▶ Limit epidemic size by restricting frequency of travel and/or range



Model output

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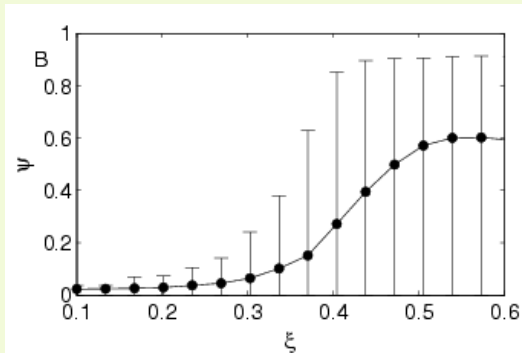


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Varying ξ :



- Transition in expected final size based on typical movement distance

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

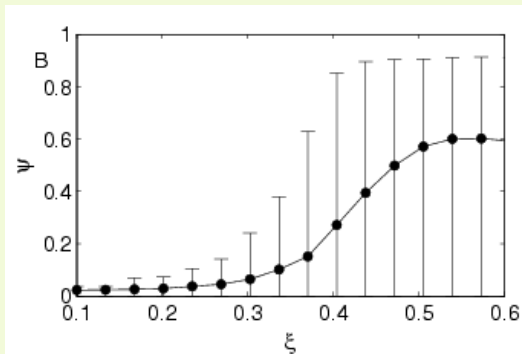
Conclusions

Predicting social
catastrophe

References



Varying ξ :



- Transition in expected final size based on typical movement distance (**sensible**)

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

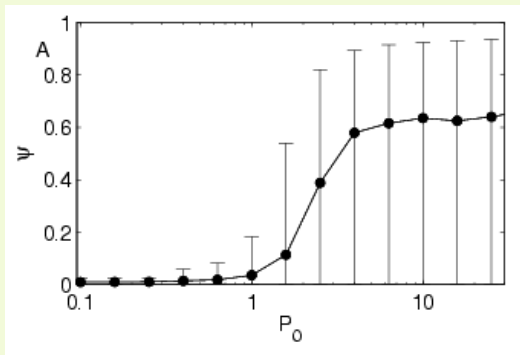
Conclusions

Predicting social
catastrophe

References



Varying P_0 :



- ▶ Transition in expected final size based on typical number of infectives leaving first group

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

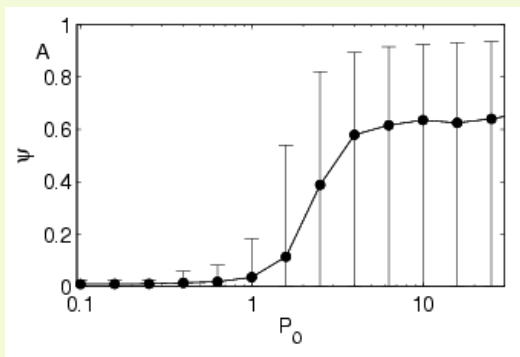
Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

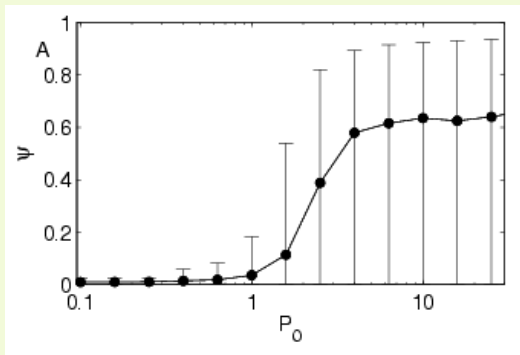
Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

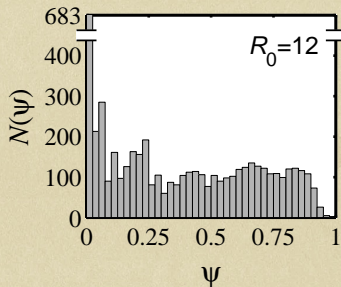
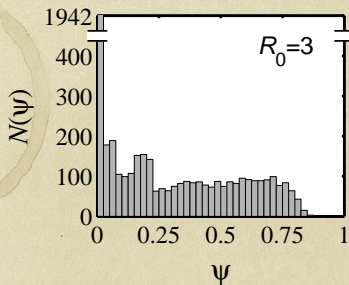
Conclusions

Predicting social
catastrophe

References



Example model output: size distributions



- ▶ Flat distributions are possible for certain ξ and P .
- ▶ Different R_0 's may produce similar distributions
- ▶ Same epidemic sizes may arise from different R_0 's

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

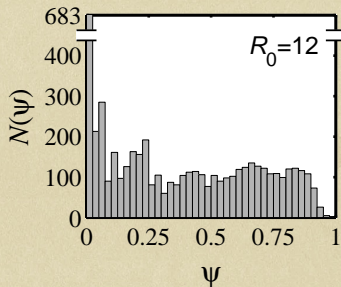
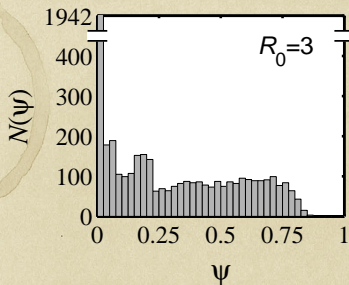
Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

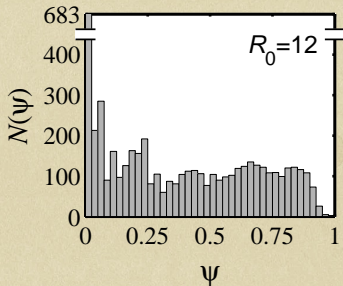
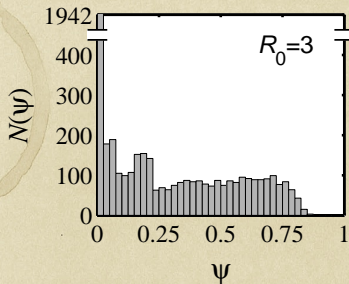
Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

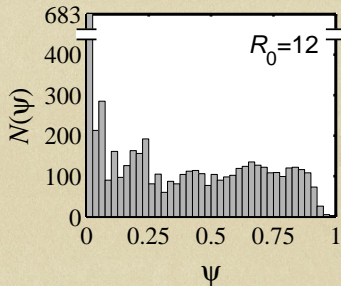
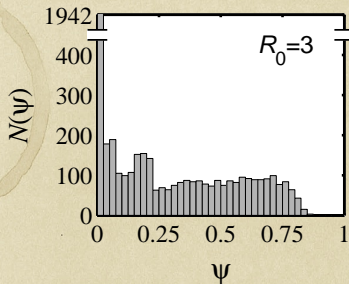
Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

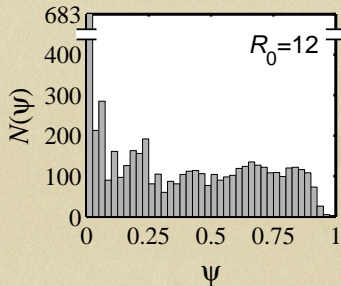
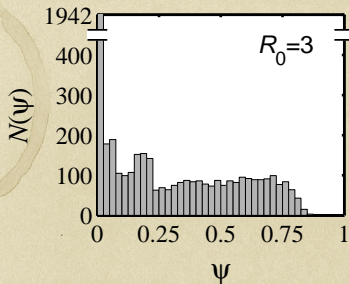
Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Model output—resurgence

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

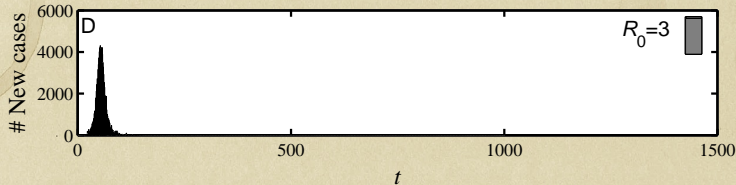
Model output

Conclusions

Predicting social
catastrophe

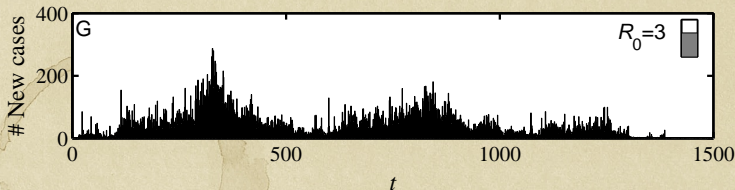
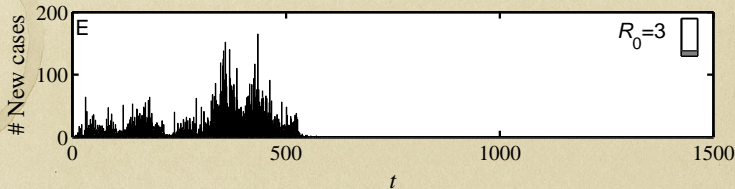
References

Standard model:



Model output—resurgence

Standard model with transport:



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



The upshot

Simple multiscale population structure

Biological
Contagion

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



The upshot

Simple multiscale population structure
+
stochasticity

Biological
Contagion

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



The upshot

Simple multiscale population structure
+
stochasticity

leads to

resurgence

+
broad epidemic size distributions

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Outline

Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References

Biological
Contagion

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Conclusions

- ▶ **For this model, epidemic size is highly unpredictable**
- ▶ Model is more complicated than SIR but still simple
- ▶ We haven't even included normal social responses such as travel bans and self-quarantine.
- ▶ The reproduction number R_0 is not terribly useful.
- ▶ R_0 , however measured, is not informative about
 1. how likely the observed epidemic size was,
 2. and how likely future epidemics will be.
- ▶ Problem: R_0 summarises one epidemic after the fact and enfold movement, the price of bananas, everything.

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References



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Conclusions

- ▶ Disease spread highly sensitive to population structure
- ▶ Rare events may matter enormously
- ▶ More support for controlling population movement



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- ▶ Disease spread highly sensitive to population structure
- ▶ Rare events may matter enormously (e.g., an infected individual taking an international flight)
- ▶ More support for controlling population movement (e.g., travel advisories, quarantine)



Conclusions

What to do:

- ▶ Need to separate movement from disease
- ▶ R_0 needs a friend or two.
- ▶ Need $R_0 > 1$ and $P_0 > 1$ and ξ sufficiently large for disease to have a chance of spreading

More wondering:

- ▶ Exactly how important are rare events in disease spreading?
- ▶ Again, what is N ?



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

Valiant attempts to use SIR and co. elsewhere:

- ▶ Adoption of ideas/beliefs (Goffman & Newell, 1964)
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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Outline

Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social catastrophe

References

Biological
Contagion

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Predicting social catastrophe isn't easy...

“Greenspan Concedes Error on Regulation”

- ▶ ... humbled Mr. Greenspan admitted that he had put too much faith in the self-correcting power of free markets ...
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- ▶ Rep. Henry A. Waxman: “Do you feel that your ideology pushed you to make decisions that you wish you had not made?”
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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



New York Times, October 23, 2008 (田)

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Predicting social catastrophe isn't easy...

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Economics, Schmeconomics

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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“I’ve been dealing with these big mathematical models of forecasting the economy ...

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I could forecast the economy better than any way I know.”



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Economics, Schmeconomics

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References

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Greenspan continues:

“The trouble is that we can’t figure that out. I’ve been in the forecasting business for 50 years. I’m no better than I ever was, and nobody else is. Forecasting 50 years ago was as good or as bad as it is today. And the reason is that human nature hasn’t changed. We can’t improve ourselves.”

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Economics, Schmeconomics

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Economics, Schmeconomics

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Economics, Schmeconomics

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Economics, Schmeconomics

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Economics, Schmeconomics

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Economics, Schmeconomics

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Economics, Schmeconomics

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Jon Stewart:

“You just bummed the @*!# out of me.”



wildbluffmedia.com

- ▶ From the Daily Show (田) (September 18, 2007)
- ▶ The full interview is here (田).

Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



Economics, Schmeconomics

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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



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Introduction

Simple disease
spreading models

Background

Prediction

More models

Toy metapopulation models

Model output

Conclusions

Predicting social
catastrophe

References



References V

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