Biological Contagion

Principles of Complex Systems Course 300, Fall, 2008

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A confusion of contagions:

- Is Harry Potter some kind of virus?
- What about the Da Vinci Code?
- Does Sudoku spread like a disease?
- ► Religion?
- Democracy...?

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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

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Social contagion

Optimism according to Ambrose Bierce: (III)

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

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Social contagion

Optimism according to Ambrose Bierce: (⊞)

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.

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Eric Hoffer, 1902–1983

There is a grandeur in the uniformity of the mass. When a fashion, a dance, a song, a slogan or a joke sweeps like wildfire from one end of the continent to the other, and a hundred million people roar with laughter, sway their bodies in unison, hum one song or break forth in anger and denunciation, there is the overpowering feeling that in this country we have come nearer the brotherhood of man than ever before.

ightharpoonup Hoffer (\boxplus) was an interesting fellow...

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Hoffer's acclaimed work:

"The True Believer:

Thoughts On The Nature Of Mass Movements" (1951) [3]

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."

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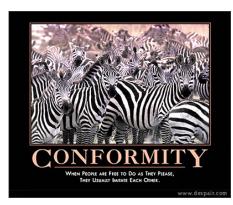
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"When people are free to do as they please, they usually imitate each other."

—Eric Hoffer "The Passionate State of Mind" [4]

despair.com

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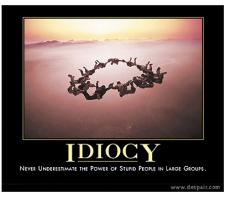


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"Never Underestimate the Power of Stupid People in Large

Groups."



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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...

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- ► Spreading of buildings in the US. (⊞)
- ightharpoonup Spreading of spreading (\boxplus).
- ▶ Viral get-out-the-vote video. (⊞)

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Two main classes of contagion

- Infectious diseases
- 2. Social contagion

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Two main classes of contagion

- 1. Infectious diseases
- 2. Social contagion

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Two main classes of contagion

- 1. Infectious diseases
- 2. Social contagion

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Two main classes of contagion

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

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Two main classes of contagion

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

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The standard SIR model^[8]

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

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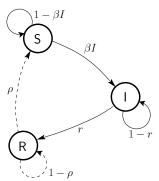




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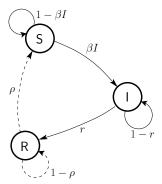
Discrete time automata example:







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Transition Probabilities:

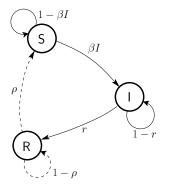
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Discrete time automata example:



Transition Probabilities:

β for being infected given contact with infected

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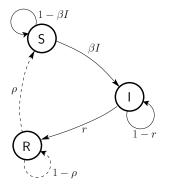
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Discrete time automata example:



Transition Probabilities:

 β for being infected given contact with infected r for recovery

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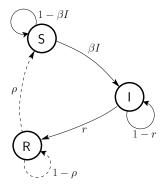
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Discrete time automata example:



Transition Probabilities:

 β for being infected given contact with infected r for recovery ρ for loss of immunity

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Original models attributed to

- ▶ 1920's: Reed and Frost
- ▶ 1920's/1930's: Kermack and McKendrick [5, 7, 6
- Coupled differential equations with a mass-action principle



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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 :

- ▶ R₀ = expected number of infected individuals resulting from a single initial infective
- \triangleright Epidemic threshold: If $R_0 > 1$, 'epidemic' occurs

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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 $(1 r)^k$

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Discrete version:

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Discrete version:

Expected number infected by original Infective:

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

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Expected number infected by original Infective:

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

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

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Expected number infected by original Infective:

$$R_0 = \beta + (1 - r)\beta + (1 - r)^2\beta + (1 - r)^3\beta + \dots$$
$$= \beta \left(1 + (1 - r) + (1 - r)^2 + (1 - r)^3 + \dots \right)$$
$$= \beta \frac{1}{1 - (1 - r)}$$

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Expected number infected by original Infective:

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

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

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Expected number infected by original Infective:

$$R_0 = \beta + (1 - r)\beta + (1 - r)^2\beta + (1 - r)^3\beta + \dots$$
$$= \beta \left(1 + (1 - r) + (1 - r)^2 + (1 - r)^3 + \dots \right)$$
$$= \beta \frac{1}{1 - (1 - r)} = \beta / r$$

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

$$R_0 = S_0 \beta / r$$

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For the continuous version

Second equation:

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

Number of infectives grows initially if

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

Same story as for discrete model.

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Number of infectives grows initially if

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

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$$\beta S(0) - r > 0 \Rightarrow \beta S(0) > r \Rightarrow \beta S(0)/r > 1$$

Same story as for discrete model.

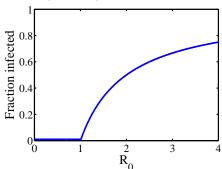
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Example of epidemic threshold:



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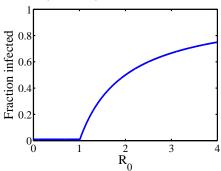
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Example of epidemic threshold:



Continuous phase transition.

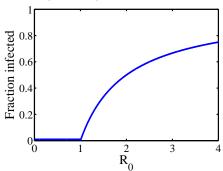
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Example of epidemic threshold:



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

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- 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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For novel diseases:

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

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For novel diseases:

- 1. Can we predict the size of an epidemic?
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References

For novel diseases:

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

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- ▶ 1918-19 "Spanish Flu" ~ 500,000 deaths in US

- ▶ 2003 "SARS Epidemic" ~ 800 deaths world-wide

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

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- ightharpoonup 1918-19 "Spanish Flu" \sim 500,000 deaths in US
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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?

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Size distributions are important elsewhere:

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Size distributions are important elsewhere:

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

Power laws distributions are common but not obligatory...



Size distributions

Really, what about epidemics?

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

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Really, what about epidemics?

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

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Really, what about epidemics?

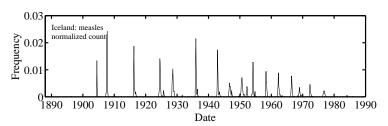
- Simply hasn't attracted much attention.
- Data not as clean as for other phenomena.

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Feeling III in Iceland

Caseload recorded monthly for range of diseases in Iceland, 1888-1890



Treat outbreaks separated in time as 'novel' diseases.

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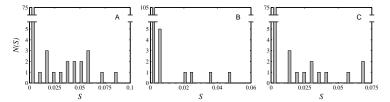
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Epidemic size distributions N(S) for Measles, Rubella, and Whooping Cough.



Spike near S = 0, relatively flat otherwise.

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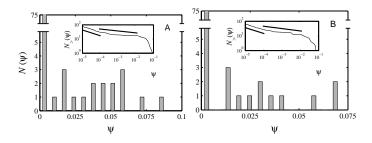
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Measles & Pertussis



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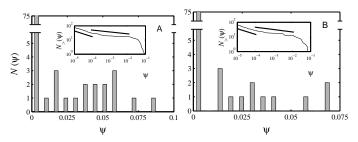
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Measles & Pertussis



Insert plots:

Complementary cumulative frequency distributions:

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

Limited scaling with a possible break.

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- ► measles: 1.40 (low Ψ) and 1.13 (high Ψ)
- ▶ pertussis: 1.39 (low Ψ) and 1.16 (high Ψ)
- ▶ Expect $2 \le \gamma < 3$ (finite mean, infinite variance)
- ▶ When γ < 1, can't normalize
- Distribution is quite flat.

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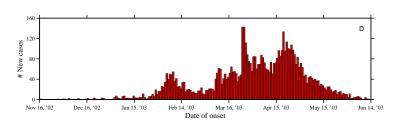
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- ► Epidemic slows...
- ► Epidemic discovers new 'pools' of susceptibles: Resurgence.
- ► Importance of rare, stochastic events.

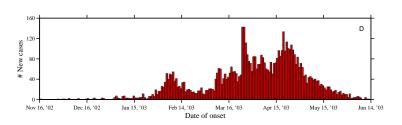
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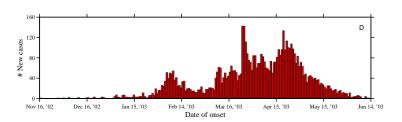
- ► Epidemic slows...
- ► Epidemic discovers new 'pools' of susceptibles: Resurgence.
- Importance of rare, stochastic events.

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- Epidemic slows... then an infective moves to a new context.
- ► Epidemic discovers new 'pools' of susceptibles: Resurgence.
- Importance of rare, stochastic events.

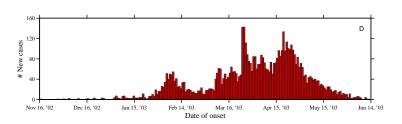
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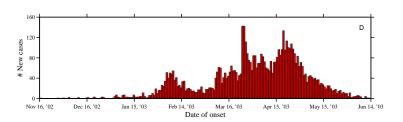
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- Epidemic slows... then an infective moves to a new context.
- ► Epidemic discovers new 'pools' of susceptibles: Resurgence.
- ► Importance of rare, stochastic events.

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The challenge

So... can a simple model produce

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

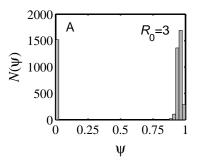
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Simple models typically produce bimodal or unimodal size distributions.

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

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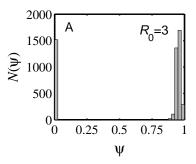
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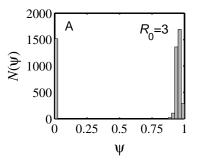
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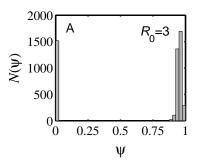
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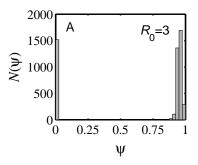
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Burning through the population

Forest fire models: [9]

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

A bit of a stretch:

- Epidemics ≡ forest fires spreading on 3-d and 5-d lattices
- Claim Iceland and Faroe Islands exhibit power law distributions for outbreaks.
- Original forest fire model not completely understood.

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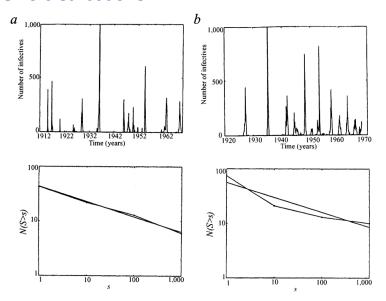
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From Rhodes and Anderson, 1996.

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- Eubank et al.'s EpiSims/TRANSIMS—city simulations.
- Spreading through countries—Airlines: Germann et al., Corlizza et al.
- ▶ Vital work but perhaps hard to generalize from...
- ► ⇒ Create a simple model involving multiscale travel
- Multiscale models suggested by others but not formalized (Bailey, Cliff and Haggett, Ferguson et al.)

Simple disease spreading models Background Prediction





- Community based mixing: Longini (two scales).
- ► Eubank et al.'s EpiSims/TRANSIMS—city simulations.
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- Very big question: What is N?
- Should we model SARS in Hong Kong as spreading
- ► For simple models, we need to know the final size beforehand...

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References

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- Should we model SARS in Hong Kong as spreading
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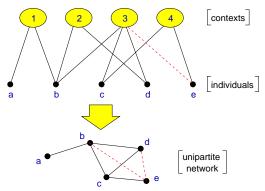
References

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Contexts and Identities—Bipartite networks



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

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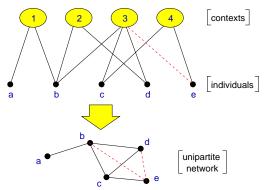
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Contexts and Identities—Bipartite networks



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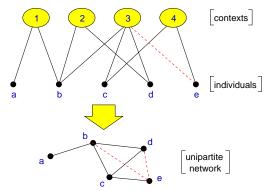
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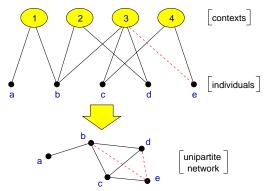
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....

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Contexts and Identities—Bipartite networks



- boards of directors
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Idea for social networks: incorporate identity.

Identity is formed from attributes such as:

- Geographic location
- Type of employmen
- Age
- Recreational activities

Groups are crucial..

- formed by people with at least one similar attribute
- ► Attributes ⇔ Contexts ⇔ Interactions ⇔ Networks. [11]

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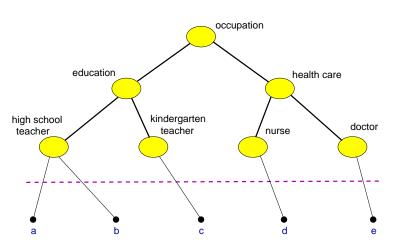
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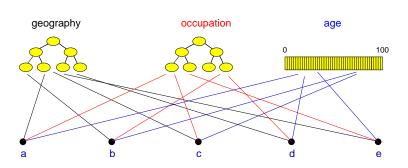


Distance makes sense in identity/context space.

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Generalized context space



(Blau & Schwartz [1], Simmel [10], Breiger [2])

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- Locally: standard SIR model with random mixing
- discrete time simulation
- $\triangleright \beta = \text{infection probability}$
- $ightharpoonup \gamma = recovery probability$
- ► *P* = probability of travel
- ▶ Movement distance: $Pr(d) \propto exp(-d/\xi)$
- \triangleright ξ = typical travel distance

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- ▶ Movement distance: $Pr(d) \propto exp(-d/\xi)$
- \triangleright ξ = typical travel distance

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- Locally: standard SIR model with random mixing
- discrete time simulation
- $ightharpoonup \beta$ = infection probability
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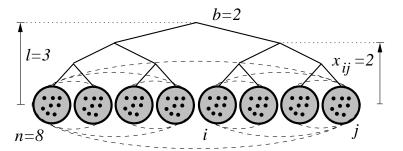
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A toy agent-based model

Schematic:



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- ▶ Define P₀ = 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

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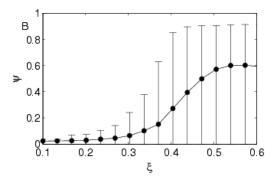


- ▶ Define P₀ = Expected number of infected individuals leaving initially infected context.
- Need $P_0 > 1$ for disease to spread (independent of R_0).
- Limit epidemic size by restricting frequency of travel and/or range

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Varying ξ :



 Transition in expected final size based on typical movement distance Introduction

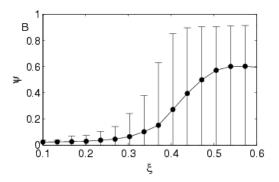
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Deference





Varying ξ :



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

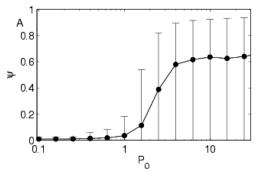
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Varying P_0 :



- Transition in expected final size based on typical number of infectives leaving first group
- ▶ Travel advisories: ξ has larger effect than P_0 .

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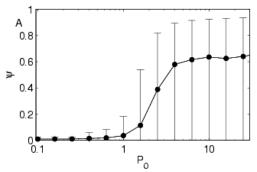
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Varying P_0 :



- Transition in expected final size based on typical number of infectives leaving first group (also sensible)
- ▶ Travel advisories: ξ has larger effect than P_0 .

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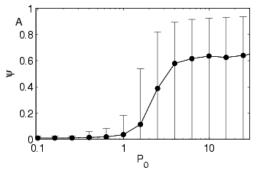
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Varying P_0 :



- Transition in expected final size based on typical number of infectives leaving first group (also sensible)
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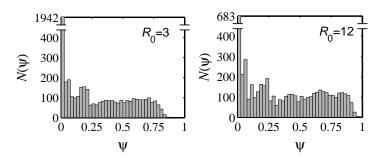
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Example model output: size distributions



- ► Flat distributions are possible for certain ξ and P.
- ▶ Different R₀'s may produce similar distributions
- ▶ Same epidemic sizes may arise from different R_0 's

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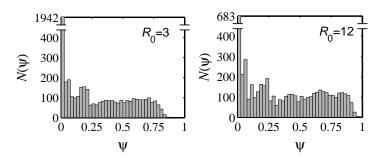
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Example model output: size distributions



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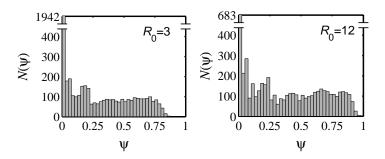
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Example model output: size distributions



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

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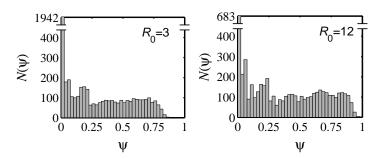
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Example model output: size distributions



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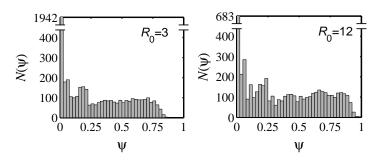
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Example model output: size distributions



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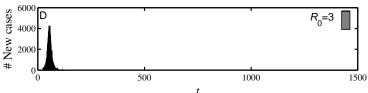
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Standard model:



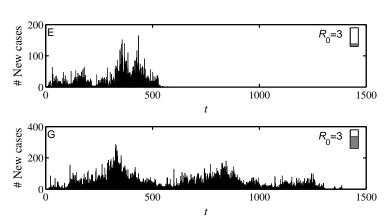
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Model output—resurgence

Standard model with transport:



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Multiscale population structure

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Multiscale population structure

+ stochasticity Introduction

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+

stochasticity

leads to

resurgence

broad epidemic size distributions

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For this model, epidemic size is highly unpredictable

- We haven't even included normal social responses
- ▶ The reproduction number R_0 is not very useful.
- \triangleright R_0 , however measured, is not informative about
- \triangleright Problem: R_0 summarises one epidemic after the fact

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- For this model, epidemic size is highly unpredictable
- Model is more complicated than SIR but still simple
- ▶ The reproduction number R_0 is not very useful.
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- 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 R₀ is not very useful.
- R₀, however measured, is not informative about
 1. how likely the observed epidemic size was.
 2. and how likely future epidemics will be.
- ▶ Problem: R₀ summarises one epidemic after the fact and enfolds movement, everything.

Simple disease spreading models Background





- Model is more complicated than SIR but still simple
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Simple disease spreading models Background

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- 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

More support for controlling population movement



- 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

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References





- Disease spread highly sensitive to population structure
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- More support for controlling population movement





- 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)

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- Need to separate movement from disease
- $ightharpoonup R_0$ needs a friend or two.
- Need R₀ > 1 and P₀ > 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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What to do:

- Need to separate movement from disease
- ► R₀ needs a friend or two.
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More wondering:

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What to do:

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What to do:

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- Again, what is N?

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- Adoption of ideas/beliefs (Goffman & Newell, 1964)
- Spread of rumors (Daley & Kendall, 1965)
- Diffusion of innovations (Bass, 1969)
- Spread of fanatical behavior (Castillo-Chávez & Song, 2003)

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