

Epidemics in Networks

Introduction

Joel C. Miller & Tom Hladish

20–22 July 2016

Introduction

Why model disease spread?

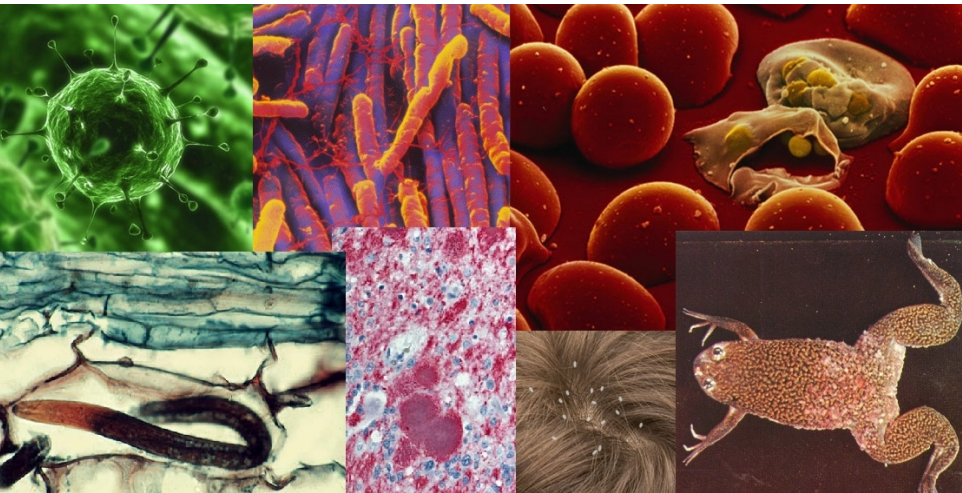
Disease spread

Networks

Key Epidemic Quantities

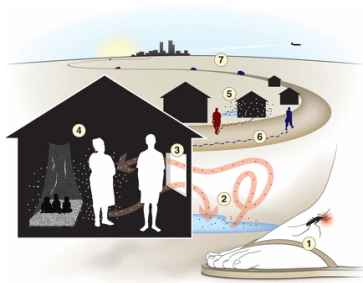
References

Infectious Diseases



Common thread:

enter a host → multiply in host → spread to another host



Recently eliminated diseases

- ▶ Smallpox

Recently eliminated diseases

- ▶ Smallpox
Eliminated by vaccination

Recently eliminated diseases

- ▶ Smallpox
Eliminated by vaccination
- ▶ Rinderpest (livestock)

Recently eliminated diseases

- ▶ Smallpox
Eliminated by vaccination
- ▶ Rinderpest (livestock)
Eliminated by vaccination

Recently eliminated diseases

- ▶ Smallpox
Eliminated by vaccination
- ▶ Rinderpest (livestock)
Eliminated by vaccination
- ▶ SARS

Recently eliminated diseases

- ▶ Smallpox
Eliminated by vaccination
- ▶ Rinderpest (livestock)
Eliminated by vaccination
- ▶ SARS
Eliminated by contact reduction

Recently eliminated diseases

- ▶ Smallpox
Eliminated by vaccination
- ▶ Rinderpest (livestock)
Eliminated by vaccination
- ▶ SARS
Eliminated by contact reduction
- ▶ Influenza A H1N1, pre-swine flu version

Recently eliminated diseases

- ▶ Smallpox
Eliminated by vaccination
- ▶ Rinderpest (livestock)
Eliminated by vaccination
- ▶ SARS
Eliminated by contact reduction
- ▶ Influenza A H1N1, pre-swine flu version
Outcompeted by new strain

Nearly eliminated diseases

- ▶ Polio

Nearly eliminated diseases

- ▶ Polio
- ▶ Guinea Worm

Recent emerging diseases

- ▶ HIV

Recent emerging diseases

- ▶ HIV
- ▶ SARS

Recent emerging diseases

- ▶ HIV
- ▶ SARS
- ▶ 2009 Influenza A H1N1

Recent emerging diseases

- ▶ HIV
- ▶ SARS
- ▶ 2009 Influenza A H1N1
- ▶ Ebola

Recent emerging diseases

- ▶ HIV
- ▶ SARS
- ▶ 2009 Influenza A H1N1
- ▶ Ebola
- ▶ MERS

Recent emerging diseases

- ▶ HIV
- ▶ SARS
- ▶ 2009 Influenza A H1N1
- ▶ Ebola
- ▶ MERS
- ▶ Zika

Introduction

Why model disease spread?

Disease spread

Networks

Key Epidemic Quantities

References

Why model disease spread?

We want to build mathematical models for infectious disease spread that:

- ▶ Predict future disease dynamics so that policy makers can prepare resources.
- ▶ Identify critical/efficient targets for intervention.
- ▶ Identify gaps in our knowledge.

Art is a lie that makes us realize truth

Pablo Picasso



Art is a lie that makes us realize truth

Pablo Picasso



It can scarcely be denied that the supreme goal of all theory is to make the irreducible basic elements as simple and as few as possible without having to surrender the adequate representation of a single datum of experience.

A. Einstein

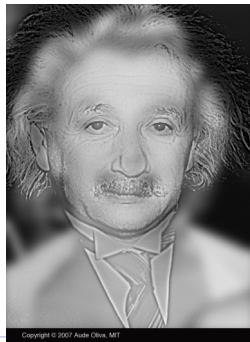
Art is a lie that makes us realize truth

Pablo Picasso



Everything should be made as simple as possible, but not simpler.

"A. Einstein"



How complex should a model be?

Modeling $\not\equiv$ mountain climbing

$$\dot{S} = -\beta kIS$$

$$\dot{I} = \beta kIS - \gamma I$$

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

$\not\equiv$



How complex should a model be?

Modeling \neq mountain climbing

$$\begin{aligned}\dot{S} &= -\beta kIS \\ \dot{I} &= \beta kIS - \gamma I \\ \dot{R} &= \gamma I\end{aligned}$$

\neq



- ▶ “Because it’s there” isn’t a good reason to include something in a model.
- ▶ Only include things that could affect decisions/improve policy.

How complex should a model be?

Modeling \neq mountain climbing

$$\dot{S} = -\beta kIS$$

$$\dot{I} = \beta kIS - \gamma I$$

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

\neq



- ▶ “Because it’s there” isn’t a good reason to include something in a model.
- ▶ Only include things that could affect decisions/improve policy.
- ▶ Sometimes intuition is good enough — it’s usually a simple mathematical model.

How complex should a model be?

Modeling \neq mountain climbing

$$\dot{S} = -\beta kIS$$

$$\dot{I} = \beta kIS - \gamma I$$

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

\neq



- ▶ “Because it’s there” isn’t a good reason to include something in a model.
- ▶ Only include things that could affect decisions/improve policy.
- ▶ Sometimes intuition is good enough — it’s usually a simple mathematical model.
- ▶ But when there are feedbacks or opposing effects, I don’t trust mine.

Introduction

Why model disease spread?

Disease spread

Networks

Key Epidemic Quantities

References

Disease spread

There are two major features that affect population-scale disease spread:

Disease spread

There are two major features that affect population-scale disease spread:

- ▶ Relation between mode of transmission and population structure.

Disease spread

There are two major features that affect population-scale disease spread:

- ▶ Relation between mode of transmission and population structure.
- ▶ How the immune system responds to exposure.

Mode of transmission

Potential spread mechanisms:

- ▶ Water & food contamination

Mode of transmission

Potential spread mechanisms:

- ▶ Water & food contamination

Cholera

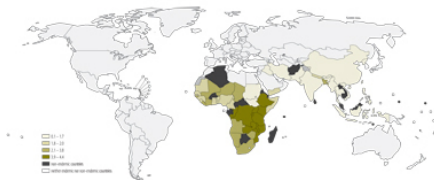


Mode of transmission

Potential spread mechanisms:

- ▶ Water & food contamination

Cholera



Mode of transmission

Potential spread mechanisms:

- ▶ Water & food contamination

Guinea Worm



Mode of transmission

Potential spread mechanisms:

- ▶ Water & food contamination

Guinea Worm



Mode of transmission

Potential spread mechanisms:

- ▶ Water & food contamination
- ▶ Vectors

Malaria, Chikungunya, Dengue, West Nile, Zika



Mode of transmission

Potential spread mechanisms:

- ▶ Water & food contamination
- ▶ Vectors

Chagas



Mode of transmission

Potential spread mechanisms:

- ▶ Water & food contamination
- ▶ Vectors

Lyme



Mode of transmission

Potential spread mechanisms:

- ▶ Water & food contamination
- ▶ Vectors

also Anaplasmosis, Babesiosis, Borrelia, Rocky Mountain Spotted Fever, Crimean-Congo Hemorrhagic Fever, ...



Mode of transmission

Potential spread mechanisms:

- ▶ Water & food contamination
- ▶ Vectors
- ▶ Direct contact

Influenza, SARS, MERS, Ebola, . . .

Mode of transmission

Potential spread mechanisms:

- ▶ Water & food contamination
- ▶ Vectors
- ▶ Direct contact

HIV, Gonorrhea, Chlamydia, **Zika**, ...

Immune response

The response of the immune system determines what effect an exposure has on an individual and whether that individual will transmit to others.

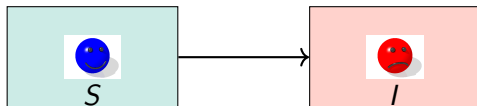
Some possible outcomes of infection:

Immune response

The response of the immune system determines what effect an exposure has on an individual and whether that individual will transmit to others.

Some possible outcomes of infection:

- Remains infected forever: SI



HIV, Tuberculosis, Hepatitis,

Immune response

The response of the immune system determines what effect an exposure has on an individual and whether that individual will transmit to others.

Some possible outcomes of infection:

- ▶ Remains infected forever: SI
- ▶ Gains permanent immunity: SIR



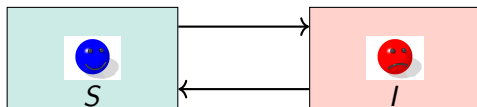
Measles, Mumps, Rubella, Pertussis, ...

Immune response

The response of the immune system determines what effect an exposure has on an individual and whether that individual will transmit to others.

Some possible outcomes of infection:

- ▶ Remains infected forever: SI
- ▶ Gains permanent immunity: SIR
- ▶ Recovers but can be reinfected: SIS



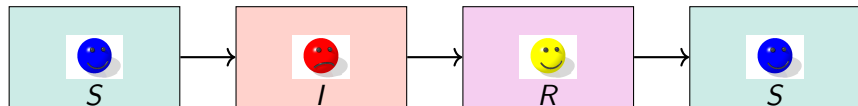
Many parasites (e.g., lice), Many bacteria, Many STDs, ...

Immune response

The response of the immune system determines what effect an exposure has on an individual and whether that individual will transmit to others.

Some possible outcomes of infection:

- ▶ Remains infected forever: SI
- ▶ Gains permanent immunity: SIR
- ▶ Recovers but can be reinfected: SIS
- ▶ Recovers with temporary immunity: SIRS



Dengue (sort of), Pertussis, Influenza,

Immune response

The response of the immune system determines what effect an exposure has on an individual and whether that individual will transmit to others.

Some possible outcomes of infection:

- ▶ Remains infected forever: SI
- ▶ Gains permanent immunity: SIR
- ▶ Recovers but can be reinfected: SIS
- ▶ Recovers with temporary immunity: SIRS

Introduction

Why model disease spread?

Disease spread

Networks

Key Epidemic Quantities

References

What is a contact network?

A network is a collection of individuals who are joined together based on interactions that may spread the disease in question. These connections (edges) may be:

What is a contact network?

A network is a collection of individuals who are joined together based on interactions that may spread the disease in question. These connections (edges) may be:

- ▶ Transient (sex workers or random encounter in crowded market)

What is a contact network?

A network is a collection of individuals who are joined together based on interactions that may spread the disease in question.

These connections (edges) may be:

- ▶ Transient (sex workers or random encounter in crowded market)
- ▶ Weighted (sharing an office versus brief daily conversation)

What is a contact network?

A network is a collection of individuals who are joined together based on interactions that may spread the disease in question.

These connections (edges) may be:

- ▶ Transient (sex workers or random encounter in crowded market)
- ▶ Weighted (sharing an office versus brief daily conversation)
- ▶ Clustered (*... Cause your friends are my friends and my friends are your friends, the more we get together the happier we'll be*)

What is a contact network?

A network is a collection of individuals who are joined together based on interactions that may spread the disease in question.

These connections (edges) may be:

- ▶ Transient (sex workers or random encounter in crowded market)
- ▶ Weighted (sharing an office versus brief daily conversation)
- ▶ Clustered (*... Cause your friends are my friends and my friends are your friends, the more we get together the happier we'll be*)
- ▶ Heterogeneously distributed

What is a contact network?

A network is a collection of individuals who are joined together based on interactions that may spread the disease in question.

These connections (edges) may be:

- ▶ Transient (sex workers or random encounter in crowded market)
- ▶ Weighted (sharing an office versus brief daily conversation)
- ▶ Clustered (*... Cause your friends are my friends and my friends are your friends, the more we get together the happier we'll be*)
- ▶ Heterogeneously distributed
- ▶ Directed.

What is a contact network?

A network is a collection of individuals who are joined together based on interactions that may spread the disease in question.

These connections (edges) may be:

- ▶ Transient (sex workers or random encounter in crowded market)
- ▶ Weighted (sharing an office versus brief daily conversation)
- ▶ Clustered (*... Cause your friends are my friends and my friends are your friends, the more we get together the happier we'll be*)
- ▶ Heterogeneously distributed
- ▶ Directed.
- ▶ ...

Network definition

- ▶ A network is a collection of nodes which are joined into pairs by edges.
- ▶ Two nodes that are joined together are called neighbors. The number of neighbors a given node has is its degree, k .
- ▶ There is no real difference between the definitions of “network” and “graph”.
- ▶ I will tend to use the terminology “partner” for neighbor and “partnership” for edge.

Network Properties

There are a number of things we can measure:

- ▶ **Degree distribution:** $P(k)$, the proportion of nodes with degree k .

High degree nodes tend to be infected early and in turn infect more nodes. So the early growth is more affected by the presence of high-degree nodes than by the average degree.

Network Properties

There are a number of things we can measure:

- ▶ Degree distribution: $P(k)$, the proportion of nodes with degree k .
- ▶ **Clustering**: frequency of short cycles [not common in sexual networks].

Clustering tends to slow the spread of a disease, but often does not significantly affect whether a disease occurs or how large it gets.

Network Properties

There are a number of things we can measure:

- ▶ Degree distribution: $P(k)$, the proportion of nodes with degree k .
- ▶ Clustering: frequency of short cycles [not common in sexual networks].
- ▶ **Dynamic networks**: Partnerships may change in time. Individuals may enter/leave the population.

Changing partnerships reduces the effect of local “susceptible depletion”

Network Properties

There are a number of things we can measure:

- ▶ Degree distribution: $P(k)$, the proportion of nodes with degree k .
- ▶ Clustering: frequency of short cycles [not common in sexual networks].
- ▶ Dynamic networks: Partnerships may change in time. Individuals may enter/leave the population.
- ▶ **Assortativity**: Individuals may actively select similar partners. In particular, partners with similar degree.

Assortative mixing by degree tends to make it easier for a disease to get established because the core of high-degree nodes provides a good place to spread. However, it often reduces the total size of the epidemic because the low degree nodes tend to connect only to low degree nodes.

Network Properties

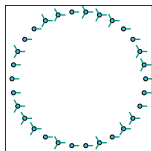
There are a number of things we can measure:

- ▶ Degree distribution: $P(k)$, the proportion of nodes with degree k .
- ▶ Clustering: frequency of short cycles [not common in sexual networks].
- ▶ Dynamic networks: Partnerships may change in time. Individuals may enter/leave the population.
- ▶ Assortativity: Individuals may actively select similar partners. In particular, partners with similar degree.
- ▶ **Edge weights**: some edges may have higher transmission probabilities than others.

Edge weights and many other effects are generally less significant (but what if weights inversely correlated with degree?)

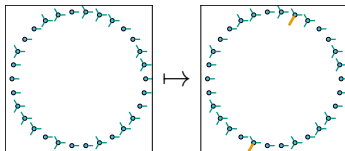
Configuration Model

The simplest model capturing a heterogeneous degree distribution:



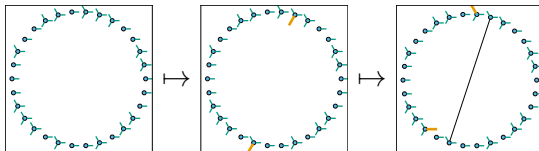
Configuration Model

The simplest model capturing a heterogeneous degree distribution:



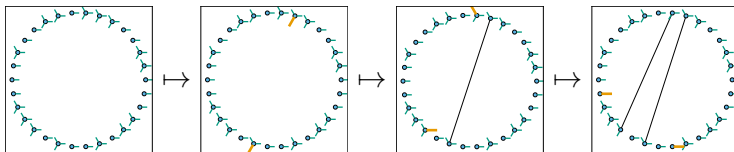
Configuration Model

The simplest model capturing a heterogeneous degree distribution:



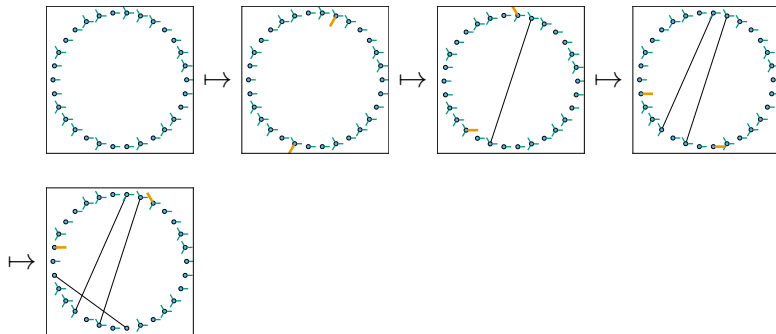
Configuration Model

The simplest model capturing a heterogeneous degree distribution:



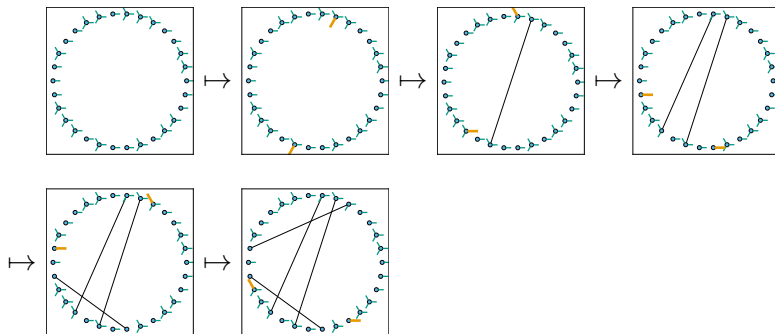
Configuration Model

The simplest model capturing a heterogeneous degree distribution:



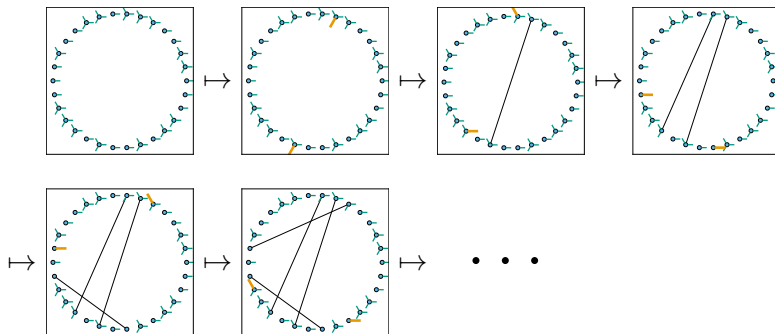
Configuration Model

The simplest model capturing a heterogeneous degree distribution:



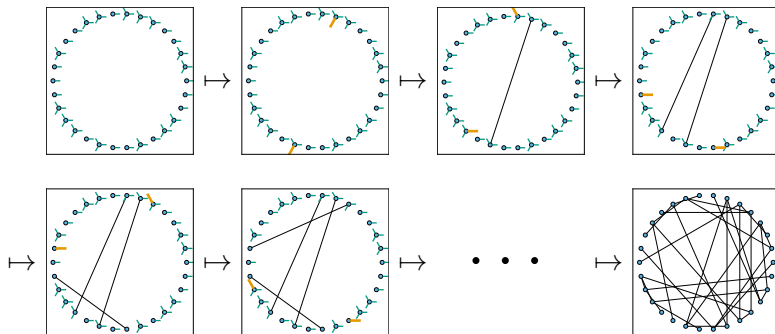
Configuration Model

The simplest model capturing a heterogeneous degree distribution:



Configuration Model

The simplest model capturing a heterogeneous degree distribution:

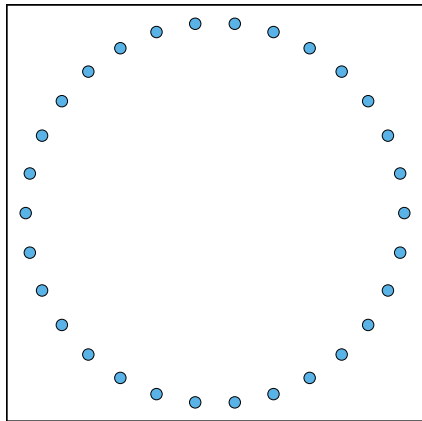


Size Bias

Do your friends have more friends than you do (on average)?

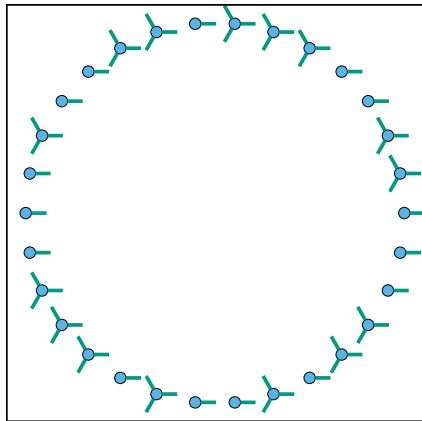
Size Bias

Do your friends have more friends than you do (on average)?



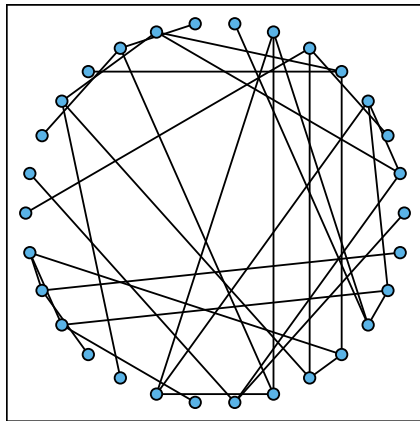
Size Bias

Do your friends have more friends than you do (on average)?



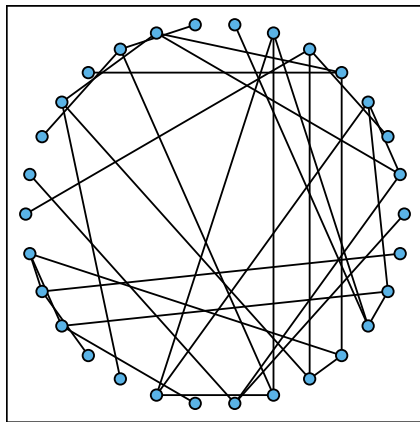
Size Bias

Do your friends have more friends than you do (on average)?



Size Bias

Do your friends have more friends than you do (on average)?



The probability a partner has degree k is $P_n(k) = kP(k) / \langle K \rangle$.

Size Bias

- ▶ A random individual has degree k with probability $P(k)$
- ▶ What about a random partner? What is the probability $P_n(k)$ a partner has degree k ?

Size Bias

- ▶ A random individual has degree k with probability $P(k)$
- ▶ What about a random partner? What is the probability $P_n(k)$ a partner has degree k ?
- ▶ Because of how partners are selected, a random partner is likely to have higher degree than a random individual $[1, 2]$.
- ▶ In fact $P_n(k) = kP(k)/\langle K \rangle$ where $\langle K \rangle$ is the average degree.

Size Bias

- ▶ A random individual has degree k with probability $P(k)$
- ▶ What about a random partner? What is the probability $P_n(k)$ a partner has degree k ?
- ▶ Because of how partners are selected, a random partner is likely to have higher degree than a random individual $[1, 2]$.
- ▶ In fact $P_n(k) = kP(k) / \langle K \rangle$ where $\langle K \rangle$ is the average degree.
- ▶ Note that the degrees of a random individual and a random neighbor of a random individual have different distributions, but a random neighbor and a random neighbor's random neighbor are both from $P_n(k)$.

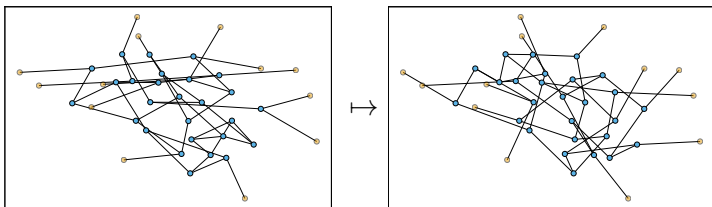
Size Bias

I cannot stress enough that if $P(k)$ is the probability a random individual has k partners, then

$$P_n(k) = kP(k) / \langle K \rangle$$

is the probability a random partner has k partners.

“Annealed” version



- ▶ The annealed network version assumes that at every moment the network looks like a Configuration model network.
- ▶ However, at every moment, an individual changes all of its partners.
- ▶ In practice this is appropriate if partnerships are so short or disease transmission so rare that an individual is unlikely to ever transmit to the same individual twice or transmit back to its infector.
- ▶ People who use the term “annealed network” call the static version a “quenched network”.

Social networks

- ▶ facebook
- ▶ linkedin
- ▶ twitter
- ▶ ...

These may be more appropriate for spread of ideas or opinions.

Contact networks

- ▶ The network of physical interactions.
- ▶ Often highly clustered.
- ▶ Appropriate for respiratory diseases.
- ▶ Often measured by giving people devices that measure physical proximity.

Sexual networks

- ▶ Appropriate for sexually transmitted diseases.
- ▶ Often low clustering.
- ▶ Often highly heterogeneous.
- ▶ Transient partnerships may play a large role.

Location–Location networks

- ▶ Cities connected by travel of people between them [spread of H1N1].
- ▶ Farms connected by movement of animals [foot and mouth].
- ▶ Habitats connected by bird migrations [West Nile].

Empirical networks

A number of attempts have been made to measure networks in “the wild”. Each case has its own peculiarities.

Empirical networks

A number of attempts have been made to measure networks in “the wild”. Each case has its own peculiarities.

- ▶ Polymod [3]: 7290 participants across 8 European countries recorded information about their contacts during a day.

Empirical networks

A number of attempts have been made to measure networks in “the wild”. Each case has its own peculiarities.

- ▶ Polymod [3]: 7290 participants across 8 European countries recorded information about their contacts during a day.
- ▶ Hospital interactions [4]: Employees, patients, and visitors at a pediatric hospital in Rome wore proximity detectors over a week-long period.

Empirical networks

A number of attempts have been made to measure networks in “the wild”. Each case has its own peculiarities.

- ▶ Polymod [3]: 7290 participants across 8 European countries recorded information about their contacts during a day.
- ▶ Hospital interactions [4]: Employees, patients, and visitors at a pediatric hospital in Rome wore proximity detectors over a week-long period.
- ▶ School interactions [5]: Students and employees at a high school wore proximity detectors.

Empirical networks

A number of attempts have been made to measure networks in “the wild”. Each case has its own peculiarities.

- ▶ Polymod [3]: 7290 participants across 8 European countries recorded information about their contacts during a day.
- ▶ Hospital interactions [4]: Employees, patients, and visitors at a pediatric hospital in Rome wore proximity detectors over a week-long period.
- ▶ School interactions [5]: Students and employees at a high school wore proximity detectors.
- ▶ Tasmanian Devils [6, 7]: Contacts between Tasmanian Devils were measured through collars with proximity detectors.

Empirical networks

A number of attempts have been made to measure networks in “the wild”. Each case has its own peculiarities.

- ▶ Polymod [3]: 7290 participants across 8 European countries recorded information about their contacts during a day.
- ▶ Hospital interactions [4]: Employees, patients, and visitors at a pediatric hospital in Rome wore proximity detectors over a week-long period.
- ▶ School interactions [5]: Students and employees at a high school wore proximity detectors.
- ▶ Tasmanian Devils [6, 7]: Contacts between Tasmanian Devils were measured through collars with proximity detectors.
- ▶ Lion interactions [8]: observations of within pride, between pride, and nomadic lion interactions.

Empirical networks

A number of attempts have been made to measure networks in “the wild”. Each case has its own peculiarities.

- ▶ Polymod [3]: 7290 participants across 8 European countries recorded information about their contacts during a day.
- ▶ Hospital interactions [4]: Employees, patients, and visitors at a pediatric hospital in Rome wore proximity detectors over a week-long period.
- ▶ School interactions [5]: Students and employees at a high school wore proximity detectors.
- ▶ Tasmanian Devils [6, 7]: Contacts between Tasmanian Devils were measured through collars with proximity detectors.
- ▶ Lion interactions [8]: observations of within pride, between pride, and nomadic lion interactions.
- ▶ Other wildlife [9].

Empirical networks

A number of attempts have been made to measure networks in “the wild”. Each case has its own peculiarities.

- ▶ Polymod [3]: 7290 participants across 8 European countries recorded information about their contacts during a day.
- ▶ Hospital interactions [4]: Employees, patients, and visitors at a pediatric hospital in Rome wore proximity detectors over a week-long period.
- ▶ School interactions [5]: Students and employees at a high school wore proximity detectors.
- ▶ Tasmanian Devils [6, 7]: Contacts between Tasmanian Devils were measured through collars with proximity detectors.
- ▶ Lion interactions [8]: observations of within pride, between pride, and nomadic lion interactions.
- ▶ Other wildlife [9].
- ▶ Romantic networks [10]

Sample location–location networks

- ▶ Livestock movement between farms [11] (and many ongoing studies).

Sample location–location networks

- ▶ Livestock movement between farms [11] (and many ongoing studies).
- ▶ Patient movement between hospitals: movement of patients in Orange County [12], movement of patients in The Netherlands [13].

Sample location–location networks

- ▶ Livestock movement between farms [11] (and many ongoing studies).
- ▶ Patient movement between hospitals: movement of patients in Orange County [12], movement of patients in The Netherlands [13].
- ▶ Individual movement between wards within a hospital [14] (and others that I recall seeing, but can't find).

Sample location–location networks

- ▶ Livestock movement between farms [11] (and many ongoing studies).
- ▶ Patient movement between hospitals: movement of patients in Orange County [12], movement of patients in The Netherlands [13].
- ▶ Individual movement between wards within a hospital [14] (and others that I recall seeing, but can't find).
- ▶ Travel through airline networks [15] (and many other papers by Colizza and Vespignani).

Sample location–location networks

- ▶ Livestock movement between farms [11] (and many ongoing studies).
- ▶ Patient movement between hospitals: movement of patients in Orange County [12], movement of patients in The Netherlands [13].
- ▶ Individual movement between wards within a hospital [14] (and others that I recall seeing, but can't find).
- ▶ Travel through airline networks [15] (and many other papers by Colizza and Vespignani).
- ▶ Seasonal population movements [16]: study of seasonal population movements for malaria control (phone data, census, satellite imagery).

Agent-based models

A number of groups have done large-scale simulations of populations

- ▶ Vancouver [17]: Simulations of individual contacts within the city of Vancouver (N)

Agent-based models

A number of groups have done large-scale simulations of populations

- ▶ Vancouver [17]: Simulations of individual contacts within the city of Vancouver (N)
- ▶ EpiSims [18]: Simulation of all individual movements through Portland, OR (1.6 million people). Later extended to a large number of other cities/regions (≈ 17 million).

Agent-based models

A number of groups have done large-scale simulations of populations

- ▶ Vancouver [17]: Simulations of individual contacts within the city of Vancouver (N)
- ▶ EpiSims [18]: Simulation of all individual movements through Portland, OR (1.6 million people). Later extended to a large number of other cities/regions (≈ 17 million).
- ▶ Epicast (based on “Scalable Parallel Short-range Molecular dynamics”: SPASM) [19]: Simulation of individual movement throughout the US (≈ 300 million).

Agent-based models

A number of groups have done large-scale simulations of populations

- ▶ Vancouver [17]: Simulations of individual contacts within the city of Vancouver (N)
- ▶ EpiSims [18]: Simulation of all individual movements through Portland, OR (1.6 million people). Later extended to a large number of other cities/regions (≈ 17 million).
- ▶ Epicast (based on “Scalable Parallel Short-range Molecular dynamics”: SPASM) [19]: Simulation of individual movement throughout the US (≈ 300 million).
- ▶ Thailand [20]: Simulated individual interactions in Thailand with the goal of identifying strategy to control pandemic influenza (500000 people).

Agent-based models

A number of groups have done large-scale simulations of populations

- ▶ Vancouver [17]: Simulations of individual contacts within the city of Vancouver (N)
- ▶ EpiSims [18]: Simulation of all individual movements through Portland, OR (1.6 million people). Later extended to a large number of other cities/regions (≈ 17 million).
- ▶ Epicast (based on “Scalable Parallel Short-range Molecular dynamics”: SPASM) [19]: Simulation of individual movement throughout the US (≈ 300 million).
- ▶ Thailand [20]: Simulated individual interactions in Thailand with the goal of identifying strategy to control pandemic influenza (500000 people).
- ▶ South Africa: Simulation by George Seage’s group at HSPH for HIV transmission (≈ 6 million?)

Introduction

Why model disease spread?

Disease spread

Networks

Key Epidemic Quantities

References

Lots of things to think about

For SIR, we are typically interested in

- ▶ \mathcal{P} , the probability of an epidemic.
- ▶ \mathcal{A} , the “attack rate”: the fraction infected (better named the attack ratio)
- ▶ \mathcal{R}_0 , the average number of infections caused by those infected early in the epidemic.
- ▶ $I(t)$, the time course of the epidemic.

For SIS, we are typically interested in

- ▶ \mathcal{P}
- ▶ $I(\infty)$, the equilibrium level of infection
- ▶ \mathcal{R}_0
- ▶ $I(t)$

Introduction

Why model disease spread?

Disease spread

Networks

Key Epidemic Quantities

References

References I

- [1] Scott L. Feld.
Why your friends have more friends than you do.
[American Journal of Sociology](#), 96(6):1464–1477, 1991.
- [2] N.A. Christakis and J.H. Fowler.
Social network sensors for early detection of contagious outbreaks.
[PLoS ONE](#), 5(9):e12948, 2010.
- [3] Joël Mossong, Niel Hens, Mark Jit, Philippe Beutels, Kari Auranen, Rafael Mikolajczyk, Marco Massari, Stefania Salmaso, Gianpaolo Scalia Tomba, Jacco Wallinga, Janneke Heijne, Malgorzata Sadkowska-Todys, Magdalena Rosinska, and W. John Edmunds.
Social contacts and mixing patterns relevant to the spread of infectious diseases.
[PLoS Medicine](#), 5(3):381–391, 2008.
- [4] Anna Machens, Francesco Gesualdo, Caterina Rizzo, Alberto E Tozzi, Alain Barrat, and Ciro Cattuto.
An infectious disease model on empirical networks of human contact: bridging the gap between dynamic network data and contact matrices.
[BMC infectious diseases](#), 13(1):185, 2013.
- [5] Marcel Salathé, Maria Kazandjiev, Jung Woo Lee, Philip Levis, Marcus W. Feldman, and James H. Jones.
A high-resolution human contact network for infectious disease transmission.
[Proceedings of the National Academy of Sciences](#), 107(51):22020–22025, 2010.
- [6] Rodrigo K Hamede, Jim Bashford, Hamish McCallum, and Menna Jones.
Contact networks in a wild tasmanian devil (*sarcophilus harrisii*) population: using social network analysis to reveal seasonal variability in social behaviour and its implications for transmission of devil facial tumour disease.
[Ecology Letters](#), 12(11):1147–1157, 2009.
- [7] Rodrigo Hamede, Jim Bashford, Menna Jones, and Hamish McCallum.
Simulating devil facial tumour disease outbreaks across empirically derived contact networks.
[Journal of Applied Ecology](#), 49(2):447–456, 2012.

References II

- [8] Meggan E Craft, Erik Volz, Craig Packer, and Lauren Ancel Meyers.
Disease transmission in territorial populations: the small-world network of serengeti lions.
[Journal of the Royal Society Interface](#), 8(59):776–786, 2011.
- [9] Meggan E Craft and Damien Caillaud.
Network models: an underutilized tool in wildlife epidemiology?
[Interdisciplinary perspectives on infectious diseases](#), 2011, 2011.
- [10] Peter S. Bearman, James Moody, and Katherine Stovel.
Chains of affection: The structure of adolescent romantic and sexual networks.
[The American Journal of Sociology](#), 110(1):44–91, 2004.
- [11] I.Z. Kiss, D.M. Green, and R.R. Kao.
The network of sheep movements within great britain: network properties and their implications for infectious disease spread.
[Journal of the Royal Society Interface](#), 3(10):669, 2006.
- [12] Susan S Huang, Taliser R Avery, Yeohan Song, Kristen R Elkins, Christopher C Nguyen, Sandra K Nutter, Alaka S Nafday, Curtis J Condon, Michael T Chang, David Chrest, et al.
Quantifying interhospital patient sharing as a mechanism for infectious disease spread.
[Infection control and hospital epidemiology: the official journal of the Society of Hospital Epidemiologists of America](#), 31(11):1160, 2010.
- [13] Tjibbe Donker, Jacco Wallinga, and Hajo Grundmann.
Patient referral patterns and the spread of hospital-acquired infections through national health care networks.
[PLoS Computational Biology](#), 6(3):e1000715, 2010.
- [14] A Sarah Walker, David W Eyre, David H Wyllie, Kate E Dingle, Rosalind M Harding, Lily O'Connor, David Griffiths, Ali Vaughan, John Finney, Mark H Wilcox, et al.
Characterisation of clostridium difficile hospital ward-based transmission using extensive epidemiological data and molecular typing.
[PLoS medicine](#), 9(2):e1001172, 2012.

References III

- [15] Duygu Balcan, Vittoria Colizza, Bruno Gonçalves, Hao Hu, José J Ramasco, and Alessandro Vespignani.
Multiscale mobility networks and the spatial spreading of infectious diseases.
[Proceedings of the National Academy of Sciences](#), 106(51):21484–21489, 2009.
- [16] Amy Wesolowski, Nathan Eagle, Andrew J Tatem, David L Smith, Abdisalan M Noor, Robert W Snow, and Caroline O Buckee.
Quantifying the impact of human mobility on malaria.
[Science](#), 338(6104):267–270, 2012.
- [17] Lauren Ancel Meyers, Babak Pourbohloul, Mark E. J. Newman, Danuta M. Skowronski, and Robert C. Brunham.
Network theory and SARS: predicting outbreak diversity.
[Journal of Theoretical Biology](#), 232(1):71–81, January 2005.
- [18] C. L. Barrett, S. G. Eubank, and J. P. Smith.
If smallpox strikes Portland. . . .
[Scientific American](#), 292(3):42–49, 2005.
- [19] Timothy C. Germann, Kai Kadau, Ira M. Longini Jr., and Catherine A. Macken.
Mitigation strategies for pandemic influenza in the United States.
[Proceedings of the National Academy of Sciences](#), 103(15):5935–5940, 2006.
- [20] Ira M Longini, Azhar Nizam, Shufu Xu, Kumnuan Ungchusak, Wanna Hanshaoworakul, Derek AT Cummings, and M Elizabeth Halloran.
Containing pandemic influenza at the source.
[Science](#), 309(5737):1083–1087, 2005.