

Physics of Life Data Epidemiology

Lect 15: temporal networks

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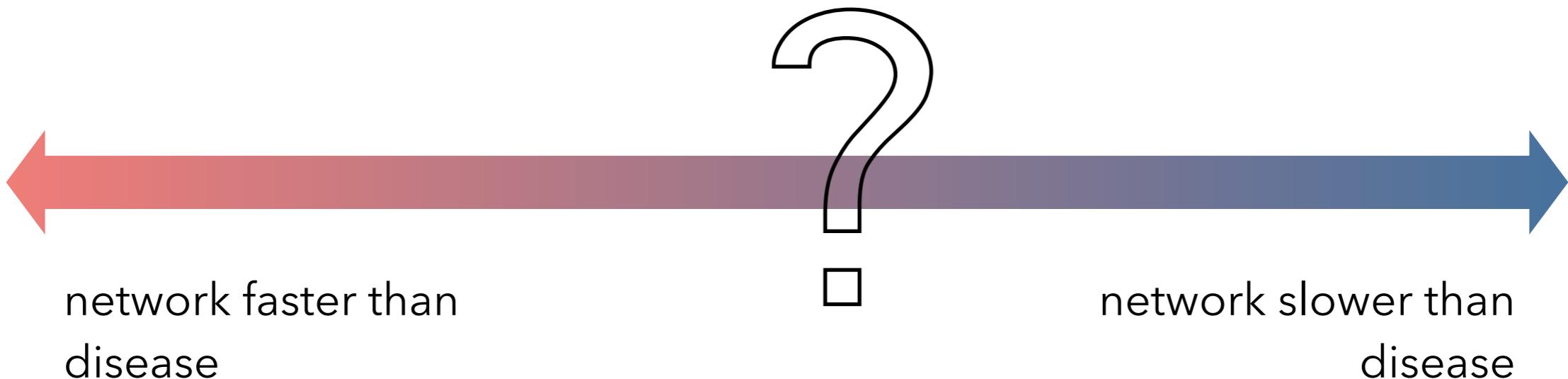
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importance of contact dynamics for an epidemic

**heterogeneous mean-field
approach**

**Individual-based
mean-field approach**

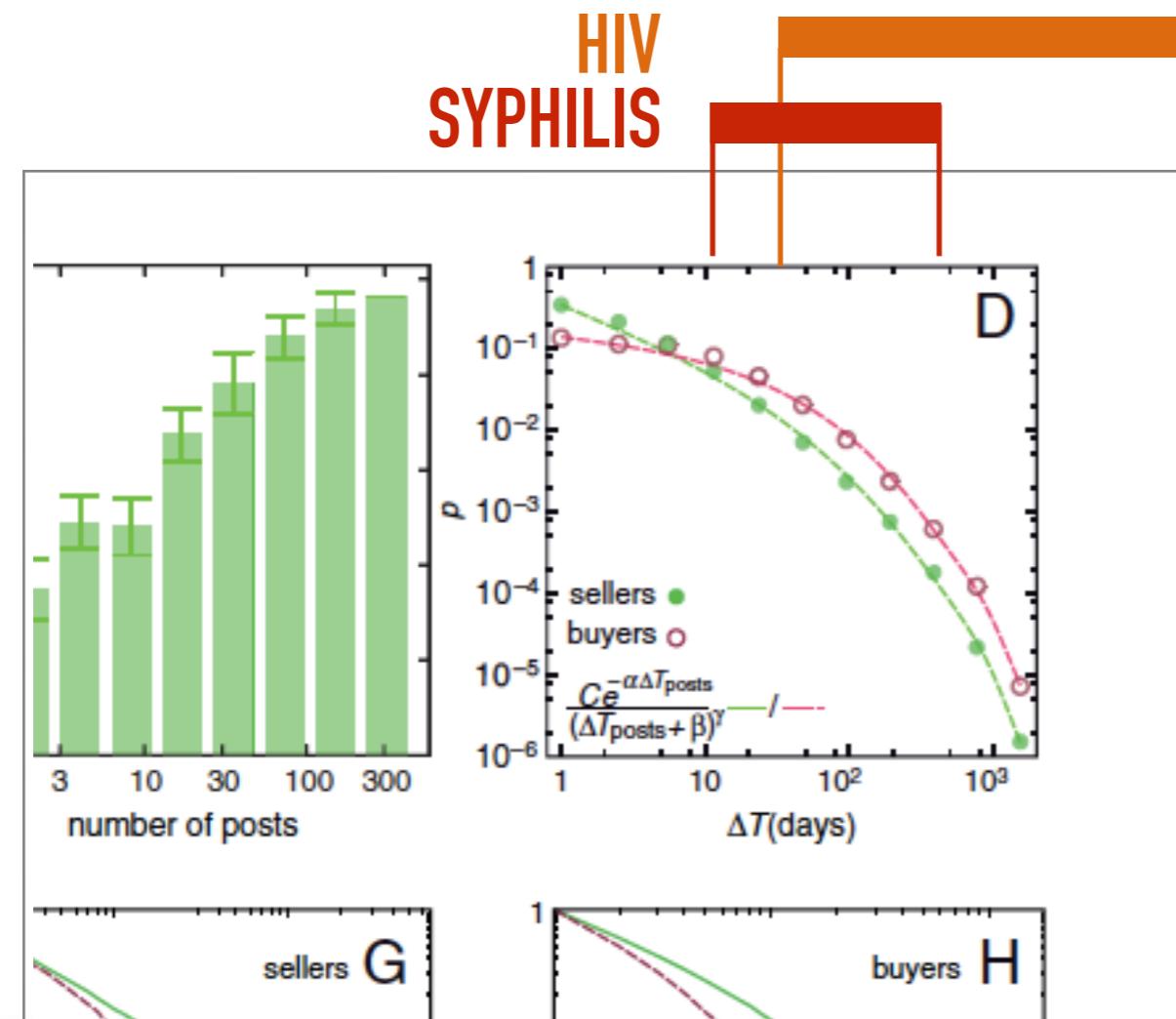


average infectious duration μ^{-1}

average inter contact time τ

importance of contact dynamics for an epidemic

time scale separation not applicable in many cases



internet mediated prostitution

[LEC. Rocha, et al, PNAS 2009]

approaches to temporal network epidemiology

Bottom-up: generative models

activity driven model, and its extensions

Top-down approaches: Randomised Reference Models

compare the epidemics on real data with the outcome in suitable null models

approaches to temporal network epidemiology

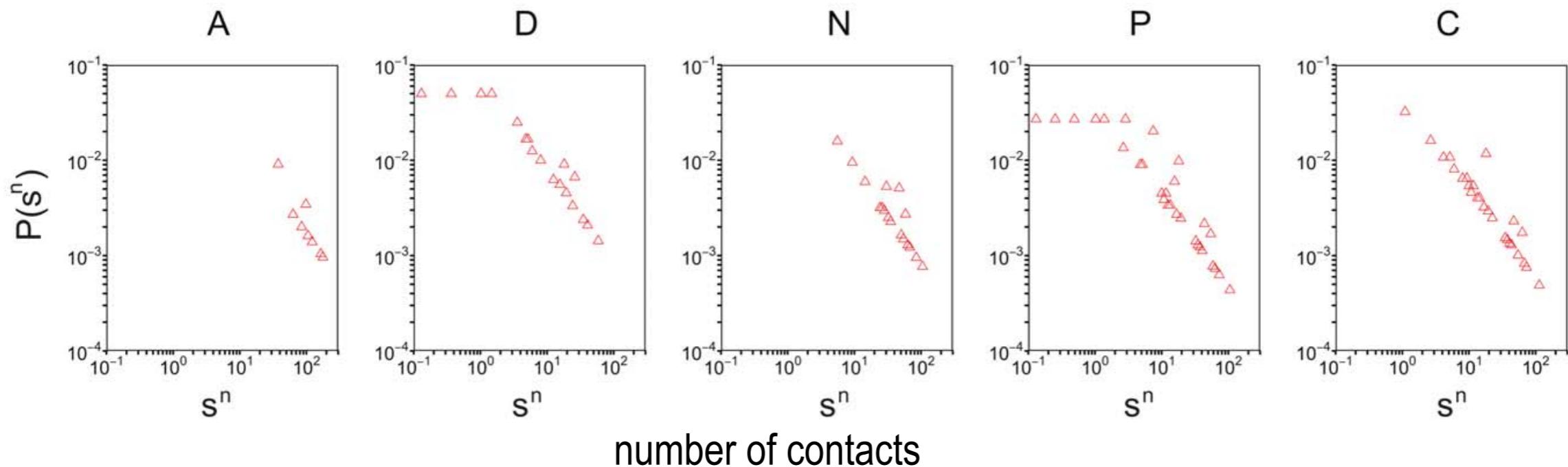
Bottom-up: generative models

activity driven model, and its extensions

Top-down approaches

what the degree of static networks hides

probability density function of number of contacts



[Isella et al PLOS ONE 2011]

Modelling heterogeneous activation frequency: **Activity driven model**

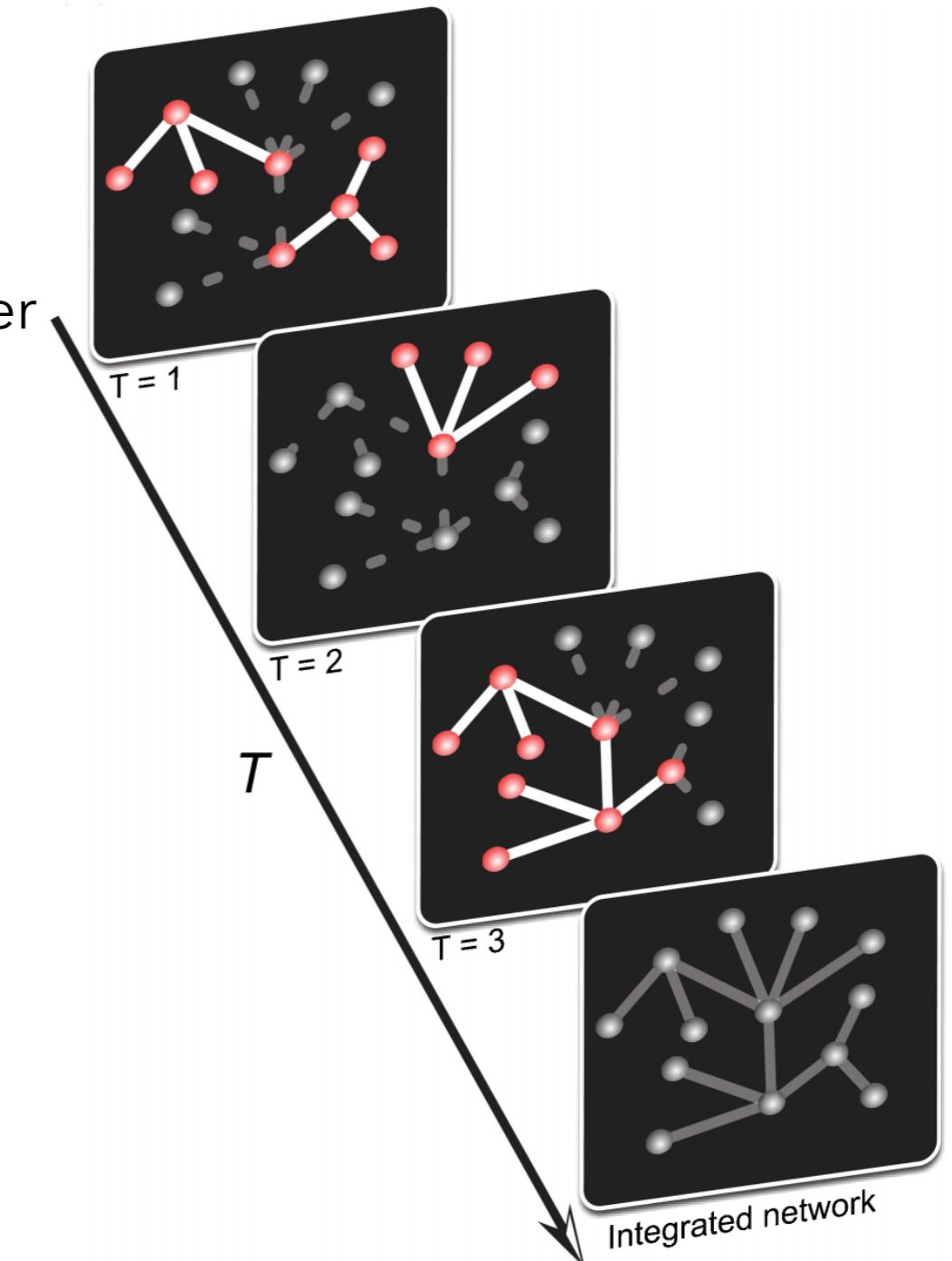
[Perra et al, Sci Rep 2012]

activity driven model

[Perra et al, Sci Rep 2012]

Model ingredients:

- discrete time (time step Δt)
- N : number of nodes
- x_i : activity potential, $\epsilon \leq x_i \leq 1$. This is e.g. number of activation of i during Δt normalised over the total number of activation.
- $F(x)$: distribution of activity potential
- $a_i = \eta x_i$: activation rate, with η rescaling factor chosen to tune the average number of active nodes per unit time in the system $\tilde{N} = \eta \langle x \rangle N$
- m : number of connections made at each activation

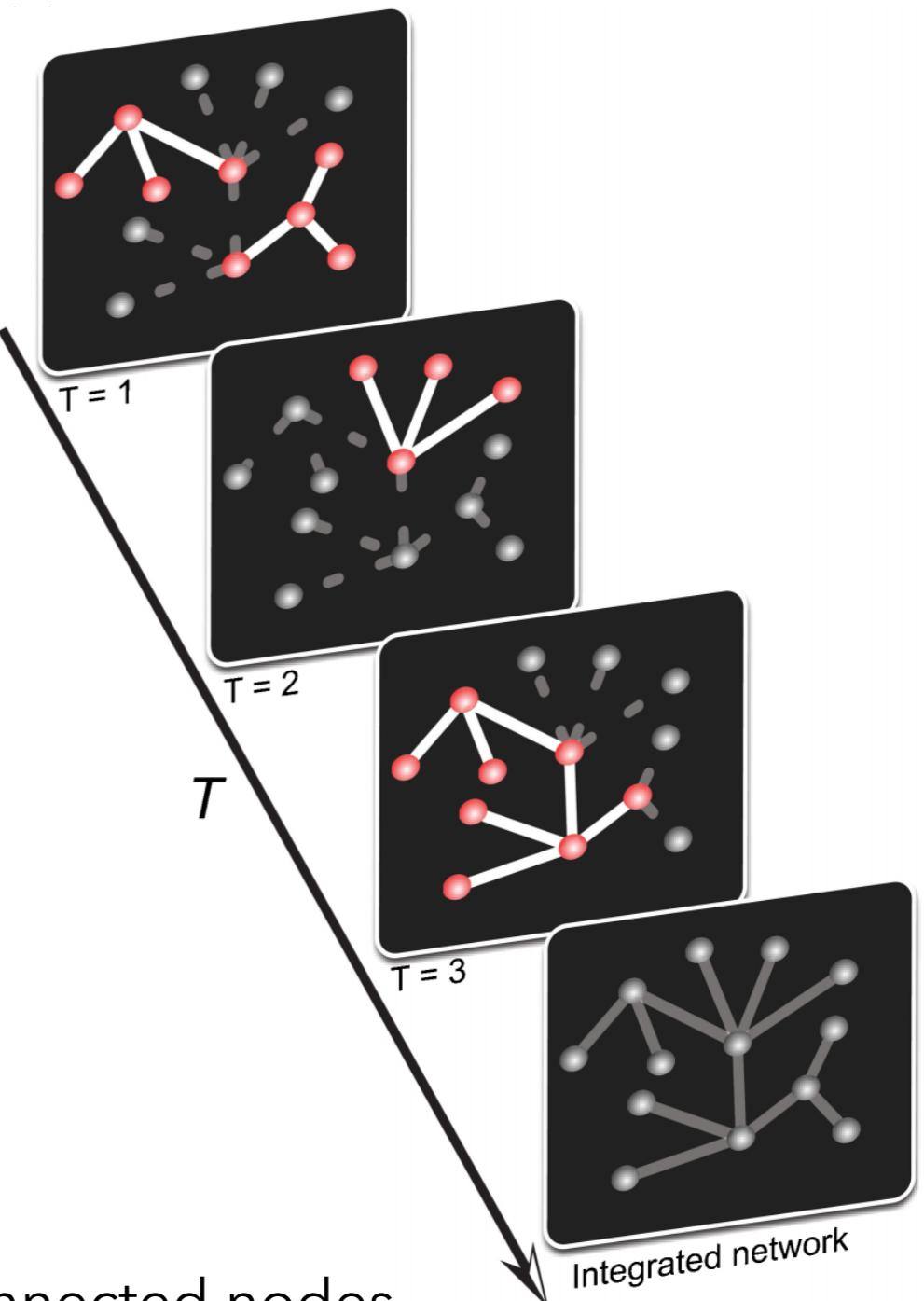


activity driven model

[Perra et al, Sci Rep 2012]

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Model algorithm:

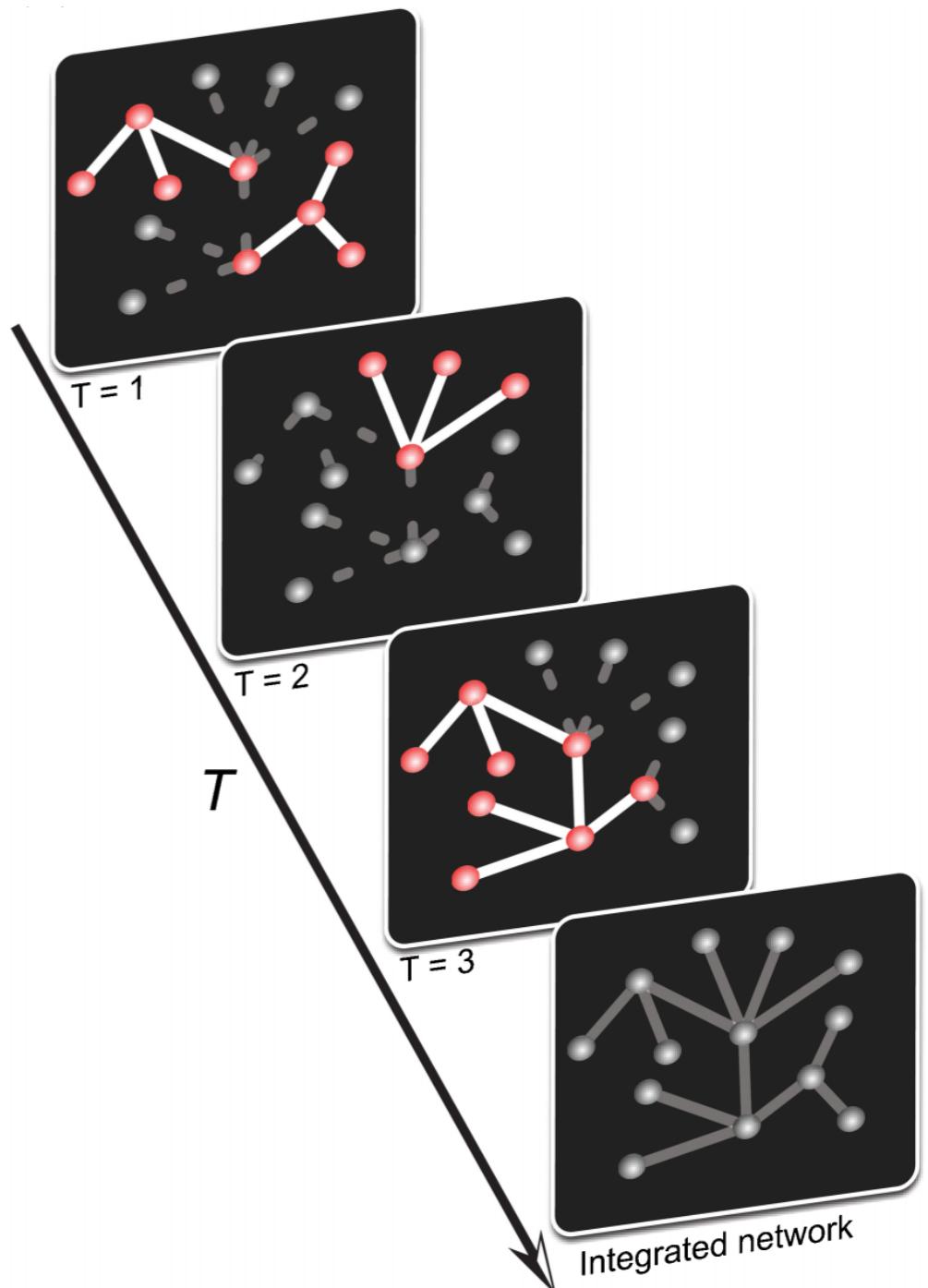
- each time step t : the network G_t starts with N disconnected nodes
- node i activates with prob $a_i \Delta t$ and makes m links with other randomly selected nodes. Non-active nodes can still receive connections
- At the next time step $t + \Delta t$ all the edges are deleted → All links last $\tau_i = \Delta t$

activity driven model

[Perra et al, Sci Rep 2012]

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At each time steps:

- number of edges $E_t = m \eta \langle x \rangle N$
- average degree $\langle k \rangle_t = \frac{2E_t}{N} = 2m \eta \langle x \rangle$
- The network is homogeneous!

activity driven model

[Perra et al, Sci Rep 2012]

Integrated network over a time window T

- degree of a node i in the aggregated network $k_T(i) = k_T^{\text{out}}(i) + k_T^{\text{in}}(i)$
- $k_T^{\text{out}}(i)$: i makes $Ta_i m$ links. How many different nodes does it connect to? (I don't count repeated links with the same node). Urn problem: # of different ball extracted from a urn with N balls after $Ta_i m$ extractions
 - prob each ball is extracted $p = 1 - \left[1 - \frac{1}{N}\right]^{Ta_i m}$
 - prob of extracting d balls is a Binomial $P(d) = \binom{N}{d} p^d (1-p)^{N-d}$
 - average # balls $k_T^{\text{out}} = Np = N \left[1 - e^{-Ta_i m/N}\right]$, if $N \rightarrow \infty$ and $\frac{T}{N} \rightarrow 0$

activity driven model

[Perra et al, Sci Rep 2012]

Integrated network over a time window T

- degree of a node i in the aggregated network $k_T(i) = k_T^{\text{out}}(i) + k_T^{\text{in}}(i)$
- $k_T^{\text{in}}(i)$: nodes that make connections with i among whose were not target by i (already counted in $k_T^{\text{out}}(i)$)
 - prob a node was not target by i : $\left[1 - \frac{1}{N}\right]^{Ta_i m} = e^{-Ta_i m/N}$
 - average number of links coming form these nodes $T\langle a \rangle m N$
 - They connect to i with probability $\frac{1}{N}$
 - $k_T^{\text{in}}(i) = T\langle a \rangle m e^{-Ta_i m/N}$

activity driven model

[Perra et al, Sci Rep 2012]

Integrated network over a time window T

- degree of a node i in the aggregated network $k_T(i) = k_T^{\text{out}}(i) + k_T^{\text{in}}(i)$

$$k_T(i) = N \left[1 - e^{-Ta_i m/N} \right] + mT \langle a \rangle e^{-Ta_i m/N} \simeq N \left[1 - e^{-Ta_i m/N} \right] = N \left[1 - e^{-T\eta x_i m/N} \right]$$

$$\text{if } N \rightarrow \infty \text{ and } \frac{T}{N} \rightarrow 0$$

we write the activity potential as a function of the degree $x(k) = -\frac{N}{\eta m T} \ln \left(1 - \frac{k}{N} \right)$

$$P_T(k) dk \sim F(x) dx \quad P_T(k) \sim F \left[x(k) \right] \frac{dx(k)}{dk} = \frac{1}{Tm\eta} \frac{1}{1 - \frac{k}{N}} F \left[-\frac{N}{\eta m T} \ln \left(1 - \frac{k}{N} \right) \right]$$

$$\frac{k}{N} \rightarrow 0, \text{ valid if } T \rightarrow 0$$

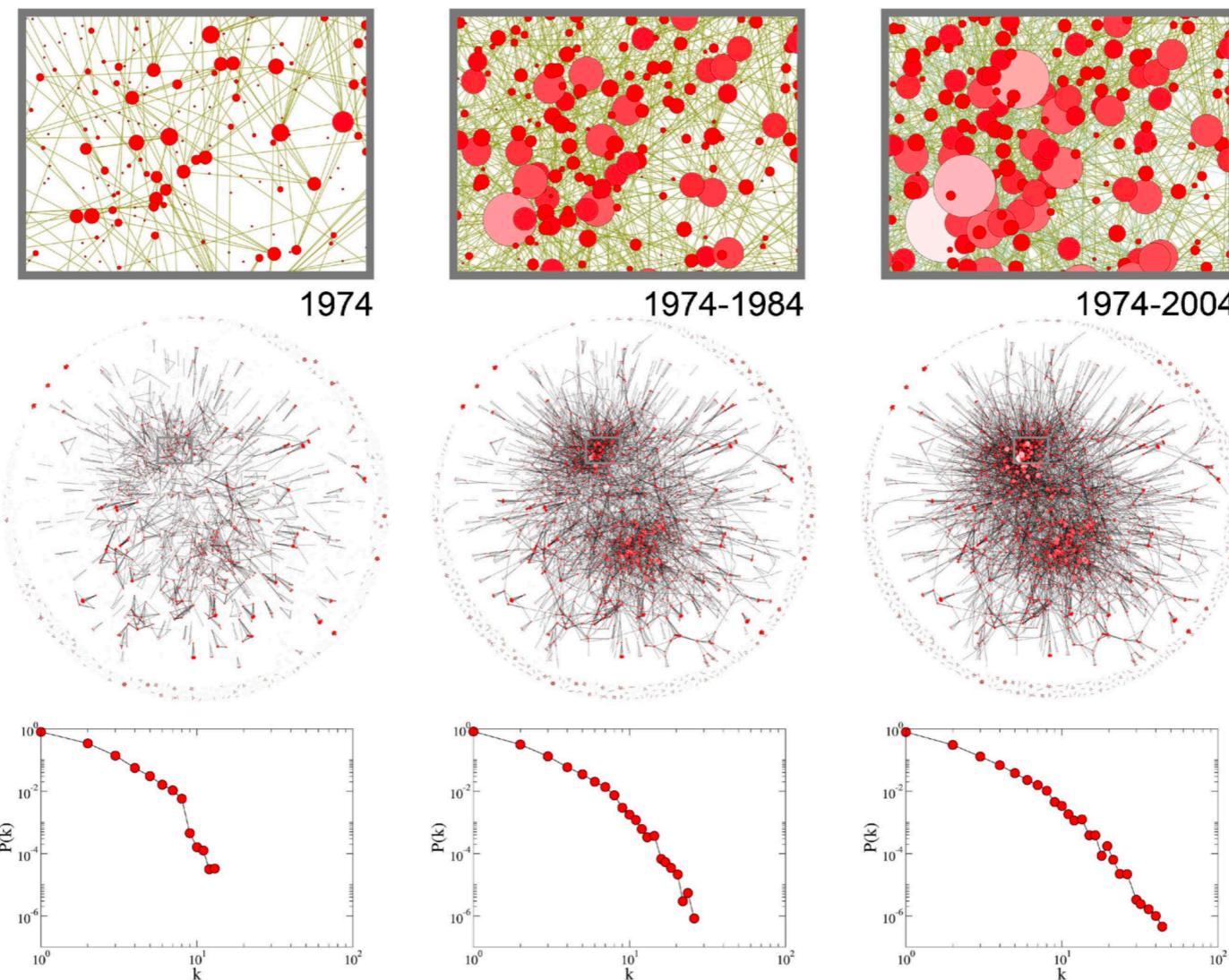
$$P_T(k) \sim \frac{1}{Tm\eta} F \left[\frac{k}{Tm\eta} \right]$$

activity driven model

[Perra et al, Sci Rep 2012]

$$P_T(k) \sim \frac{1}{Tm\eta} F \left[\frac{k}{Tm\eta} \right]$$

heterogenous topology in the aggregated network, over a window T , result from a heterogeneous activity potential



activity driven model

Effect of network dynamics on epidemic spreading

- activity block approximation
- SIR dynamics
- probability of transmission per contact λ
- for simplicity let's assume $m = 1$

$$I_a^{t+\Delta t} = -\mu \Delta t I_a^t + I_a^t + \lambda (N_a^t - I_a^t) a \Delta t \int da' \frac{I_{a'}^t}{N} + \lambda (N_a^t - I_a^t) \boxed{\int da' \frac{I_{a'}^t a' \Delta t}{N}}$$
$$\int da I_a^{t+\Delta t} = I^{t+\Delta t} = I^t - \mu \Delta t I^t + \lambda \langle a \rangle I^t \Delta t + \lambda \theta^t \Delta t$$
$$\theta^{t+\Delta t} = \theta^t - \mu \theta^t \Delta t + \lambda \langle a^2 \rangle I^t \Delta t + \lambda \langle a \rangle \theta^t \Delta t$$

$$\begin{aligned}\partial_t I &= -\mu I + \lambda \langle a \rangle I + \lambda \theta, \\ \partial_t \theta &= -\mu \theta + \lambda \langle a^2 \rangle I + \lambda \langle a \rangle \theta\end{aligned}$$

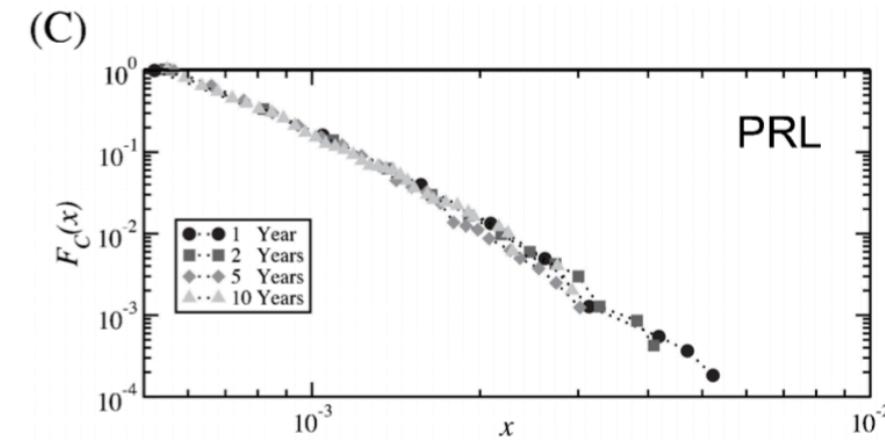
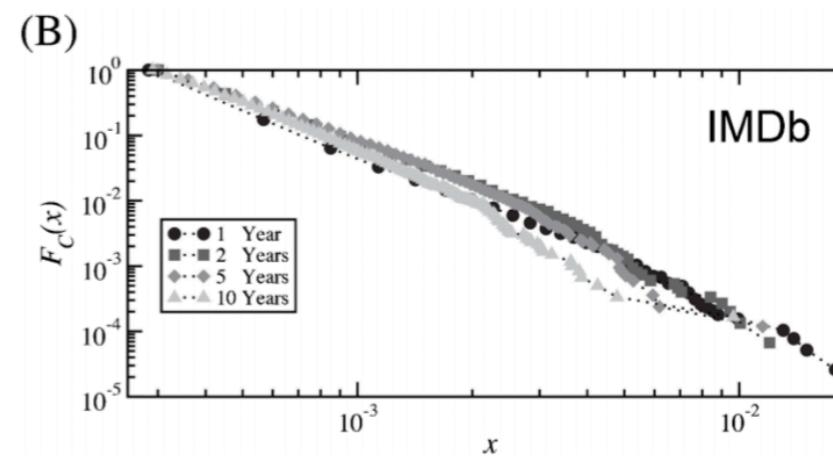
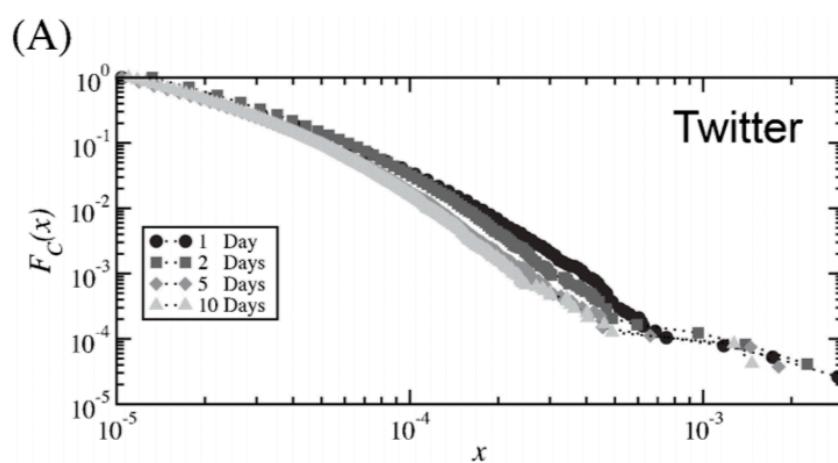
activity driven model

$$\begin{aligned}\partial_t I &= -\mu I + \lambda \langle a \rangle I + \lambda \theta, \\ \partial_t \theta &= -\mu \theta + \lambda \langle a^2 \rangle I + \lambda \langle a \rangle \theta\end{aligned}$$

$$J = \begin{pmatrix} -\mu + \lambda \langle a \rangle & \lambda \\ \lambda \langle a^2 \rangle & -\mu + \lambda \langle a \rangle \end{pmatrix} \quad \Lambda_{(1,2)} = \lambda \langle a \rangle - \mu \pm \lambda \sqrt{\langle a^2 \rangle}$$

$$\frac{\lambda}{\mu} > \frac{1}{\langle a \rangle + \sqrt{\langle a^2 \rangle}} + \mathcal{O}\left(\frac{1}{N}\right)$$

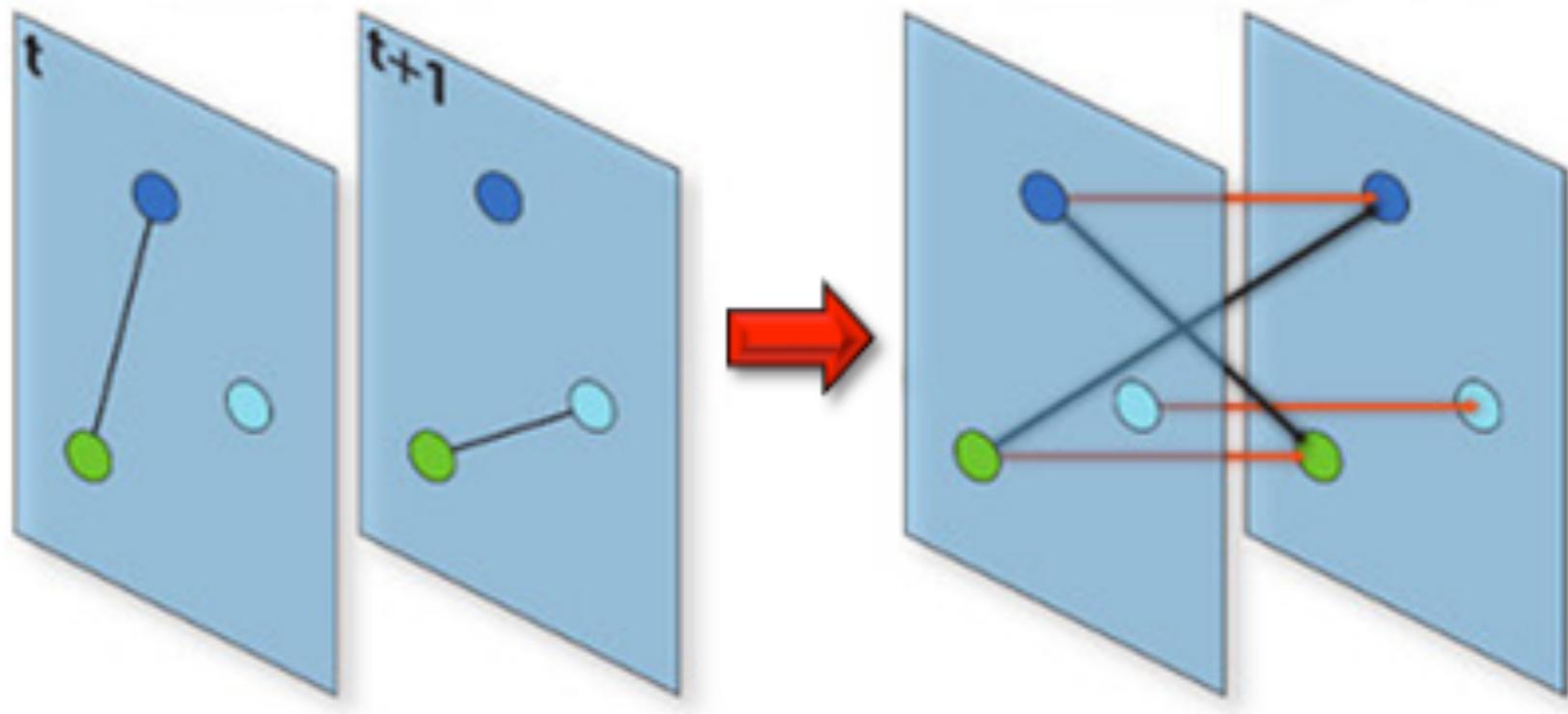
heterogeneities in the activation rate lower the epidemic threshold



activity driven model

- A model that captures a realistic property of human behaviour (face-to-face, sexual contacts, phone call, email, tweets)
- humans have heterogeneous activity rate
- the contact network at a certain instant of time is sparse, with homogeneous degree
- the aggregated network over a certain window is well connected with heterogeneous degree
- pattern of activation unfolds at the same time scale of the spreading process
- calculation possible in the activity-block approximation (same scheme as the degree block approximation)
- contact heterogeneities lower the epidemic threshold

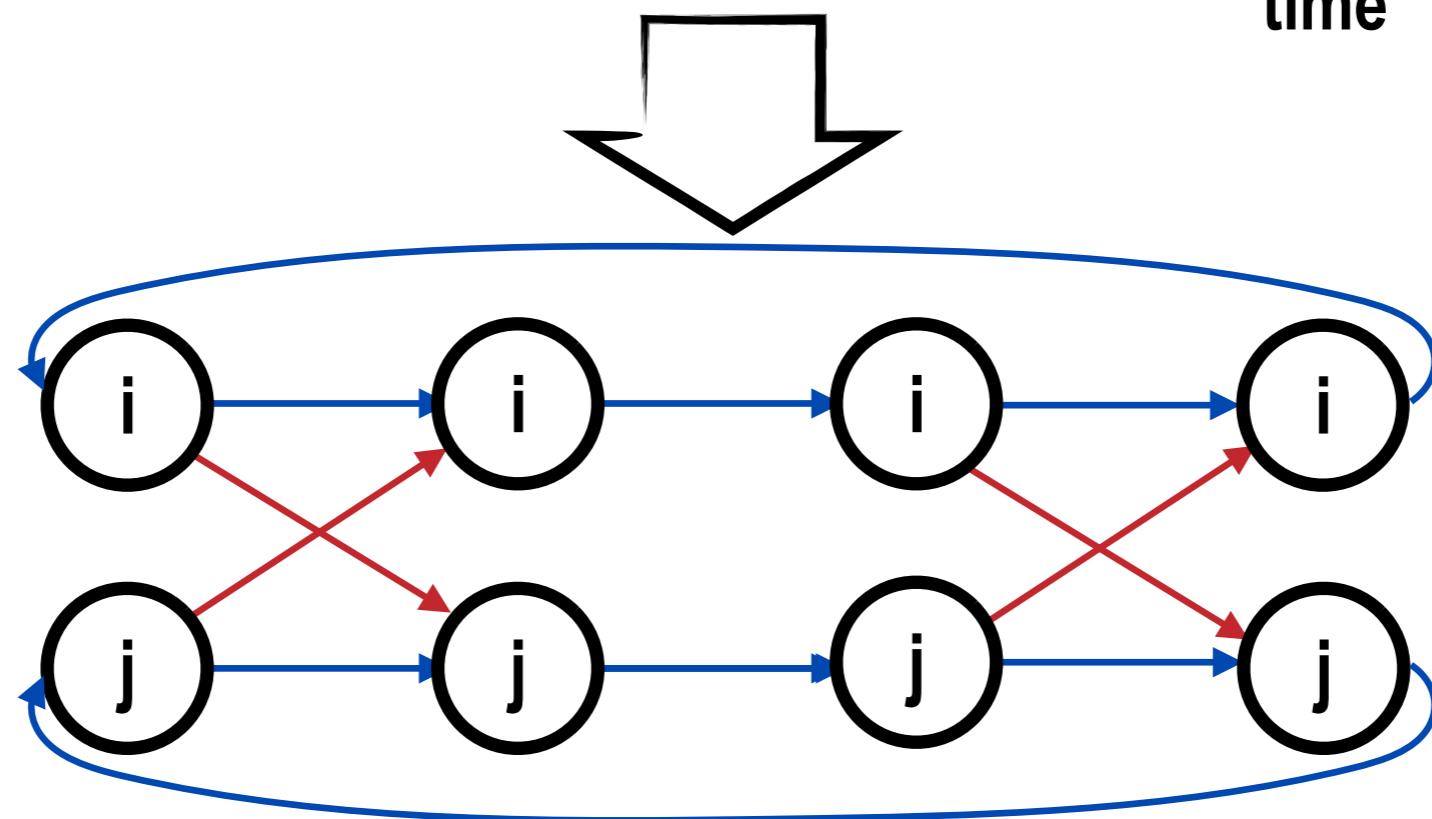
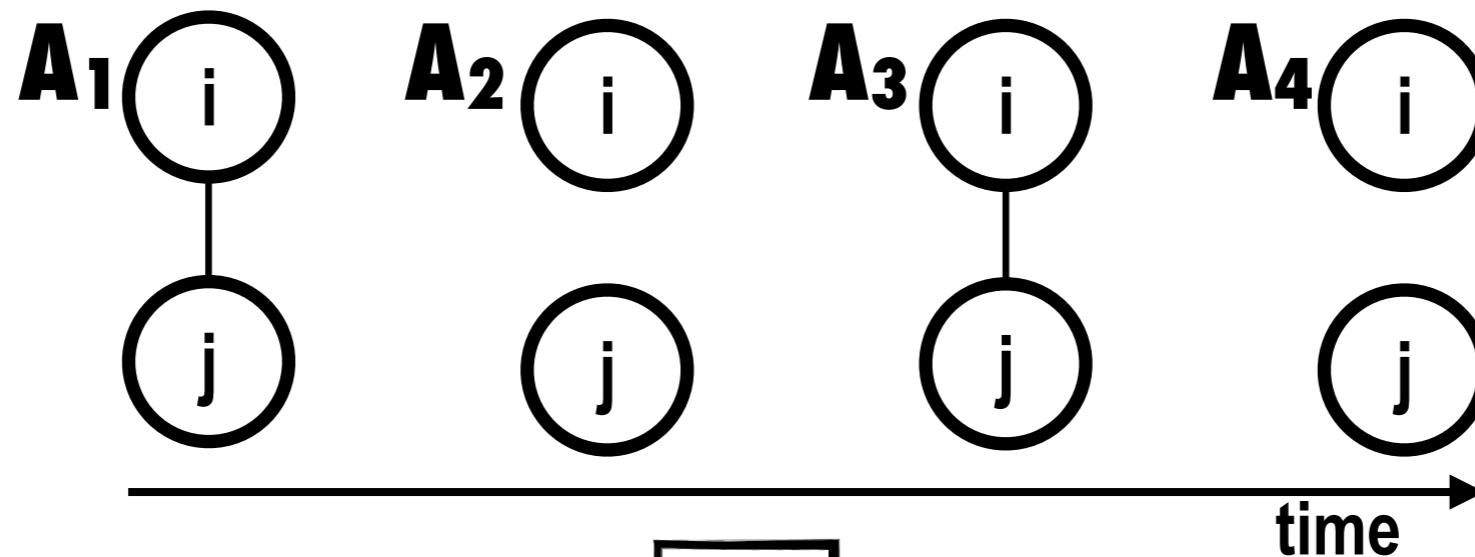
individual-based mean field approach



multilayer formalism + individual-based mean field approach = **Infectious propagator approach**

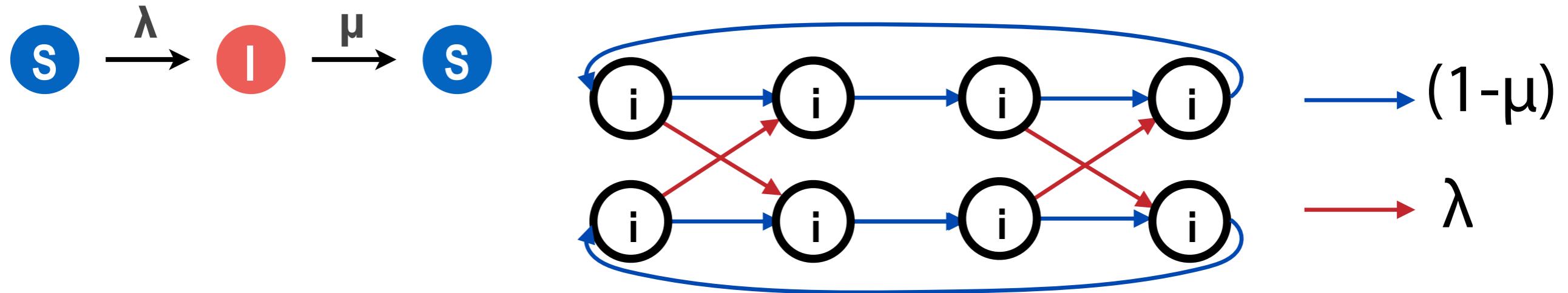
[multilayer: De Domenico et al PRX 2013, Valdano et al PRX 2015]

infectious propagator approach



[Valdano et al PRX 2015]

infectious propagator approach



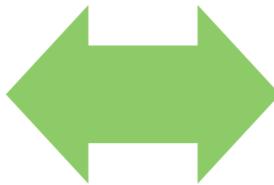
$NT \times NT$ supra-adjacency matrix

$$M = \begin{pmatrix} 0 & 1 - \mu + \lambda A_1 & 0 & \cdots & 0 \\ 0 & 0 & 1 - \mu + \lambda A_2 & \cdots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & 0 & \cdots & 1 - \mu + \lambda A_{T-1} \\ 1 - \mu + \lambda A_T & 0 & 0 & \cdots & 0 \end{pmatrix}$$

A green arrow points to the value '0' in the bottom-left corner of the matrix, which is highlighted with a green circle. Below the matrix, the text 'N x N blocks' is written.

infectious propagator approach

threshold on
temporal network



threshold on
STATIC network

epi threshold
 $\rho[\mathbf{M}(\lambda_{\text{thr}}, \mu)] = 1$

dim = NT
dim = N
[Powell, arXiv, 2011]

$$\rho[\mathbf{P}(\lambda_{\text{thr}}, \mu)] = 1$$

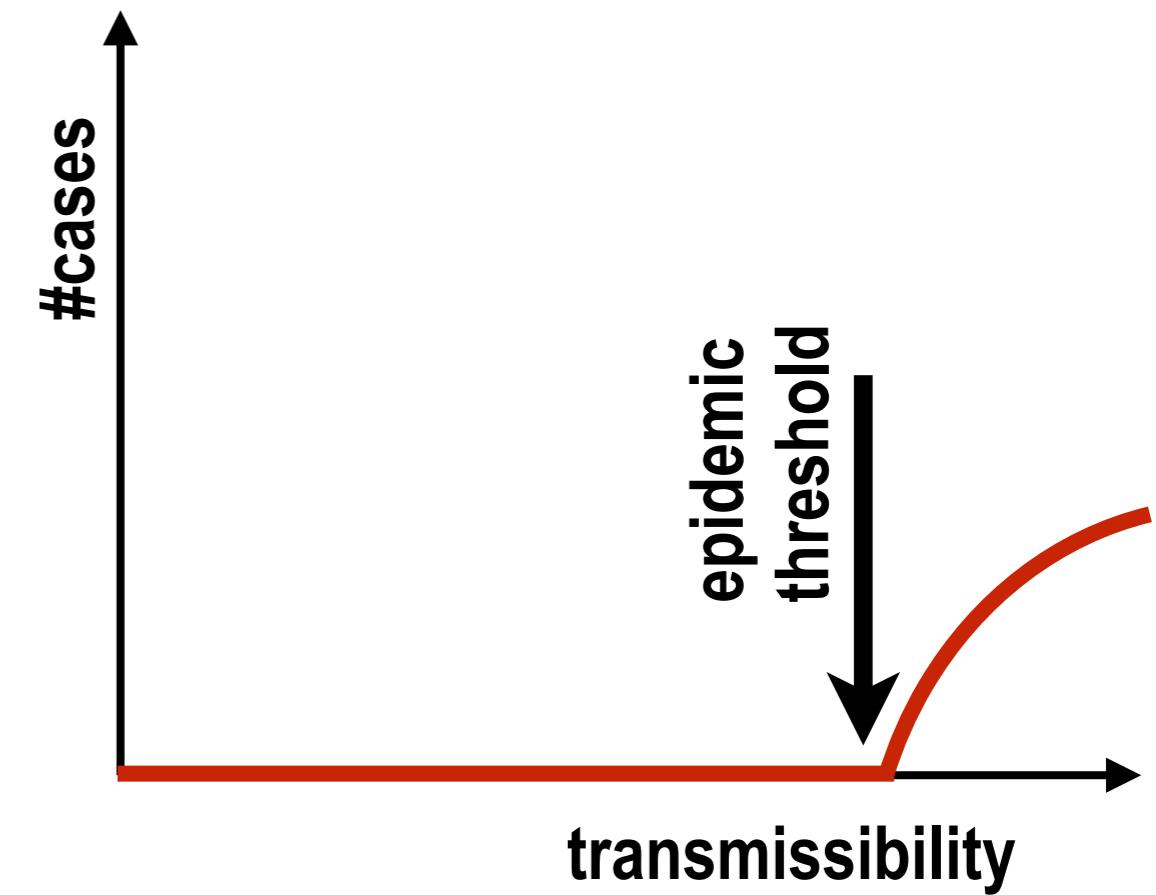
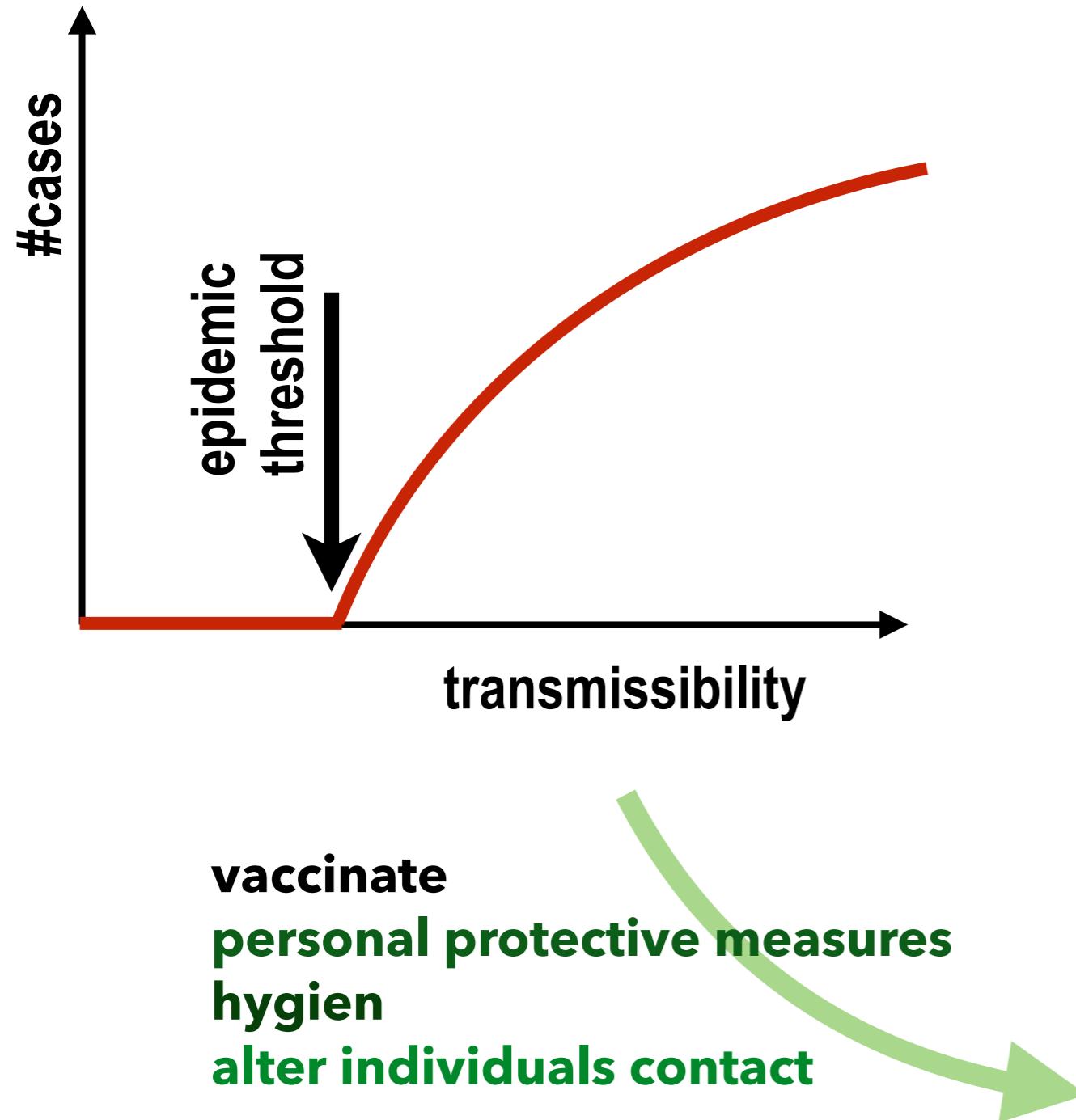
$$\mathbf{P} = (1 - \mu + \lambda \mathbf{A}_1) (1 - \mu + \lambda \mathbf{A}_2) \cdots (1 - \mu + \lambda \mathbf{A}_T)$$

[Lentz et al, PRL 2013]

infection
propagator

[Valdano et al PRX 2015]

reduce system's vulnerability

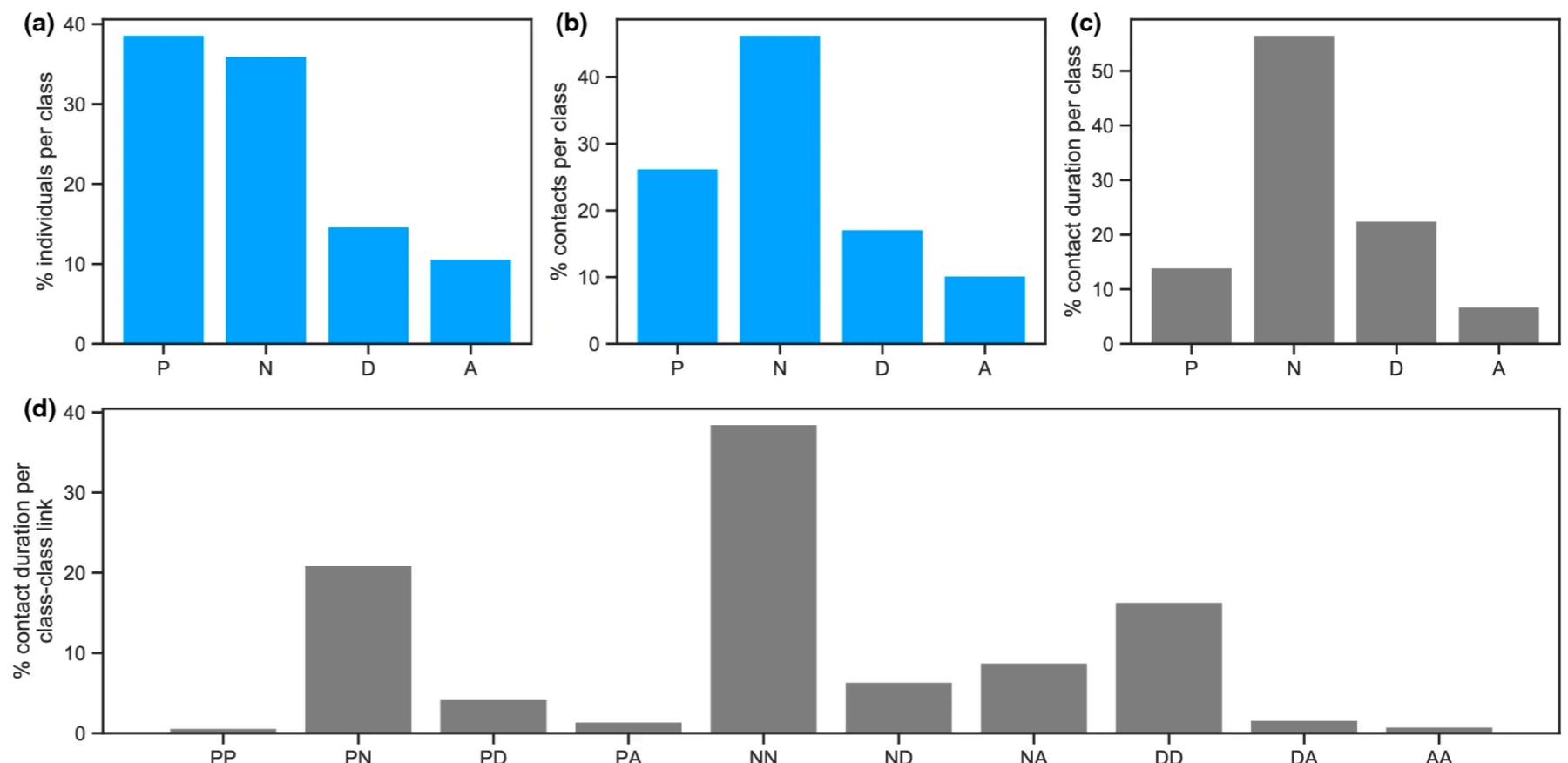


spread of nosocomial diseases

Sociopatterns.org

[Vanhems et al PLoS ONE 2013; Valdano et al Sci Rep 2021]

- geriatric ward
- 75 participants
- **27 nurses (N), 11 doctors (D), 8 administrative staff (A), 29 patients (P)**
- duration 4 days and 4 nights



spread of nosocomial diseases

