

# **Epidemic networks - modelling the relevant structure**

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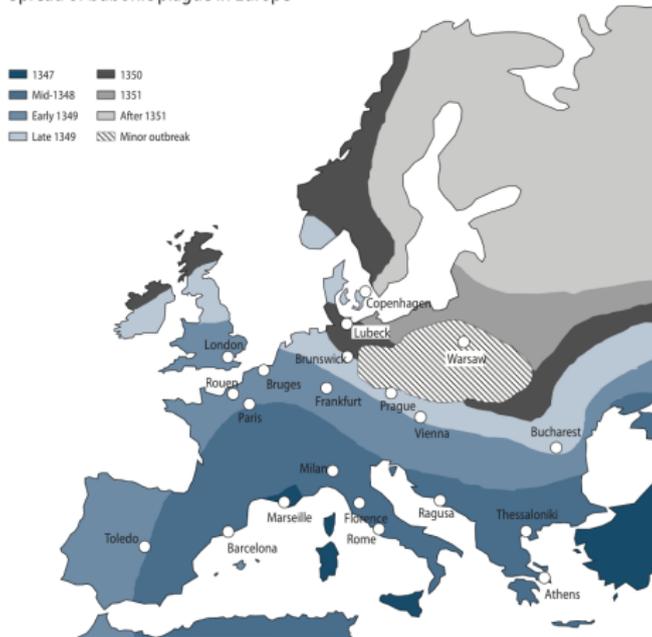
# Outline

- ▶ **Motivation**
- ▶ **Random graph models**
  - ▶ how do we generate 'realistic' networks?
- ▶ **Bi-partite graph models**
  - ▶ **Hypergraphs**

# Motivation

Historical epidemics were wave-like / spatial:

Spread of bubonic plague in Europe



(picture from WHO)

# Motivation

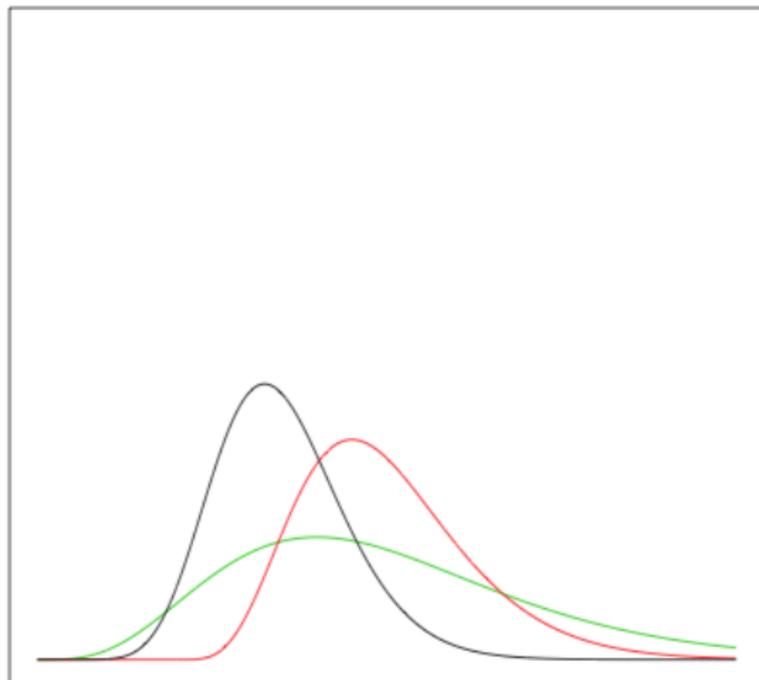
Modern epidemics are spatially heterogeneous but transmission rates are not a simple function of Euclidian distance – with all that implies for control.



# Approximations in Models

Approximation of infection process in SEIR models.

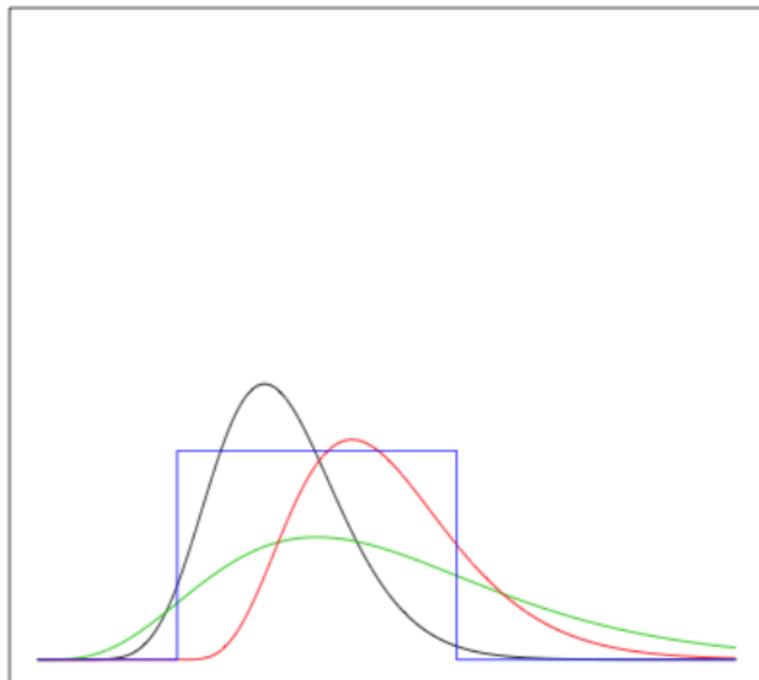
viral load, symptoms, infectivity



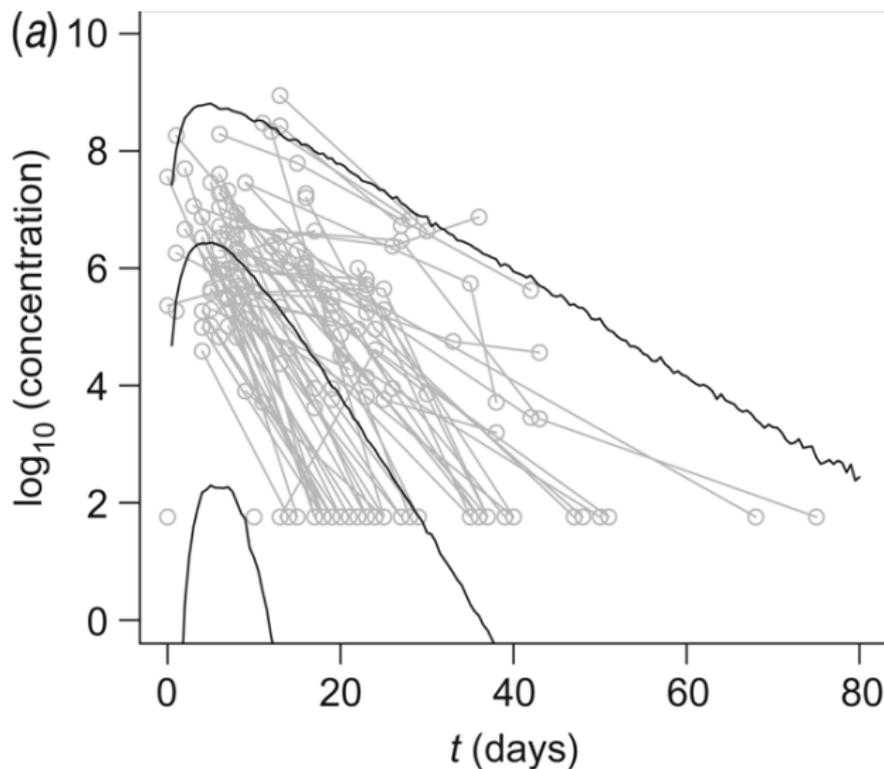
# Approximations in Models

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viral load, symptoms, infectivity



# Norovirus Shedding data



†

# Contact Heterogeneity

- ▶ consider individuals  $i, j, \dots \in \{1, \dots, N\}$ ;
- ▶  $\lambda_{ij}$  is the infection rate between  $i$  and  $j$ 
  - ▶ when  $i$  is infectious and  $j$  is susceptible.

# Contact Heterogeneity

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- ▶  $\lambda_{ij}$  is the infection rate between  $i$  and  $j$ 
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- ▶  $\lambda_{ij} = d(i, j)\rho(i|x_i)\psi(j|x_j)$

# Networks: Definitions

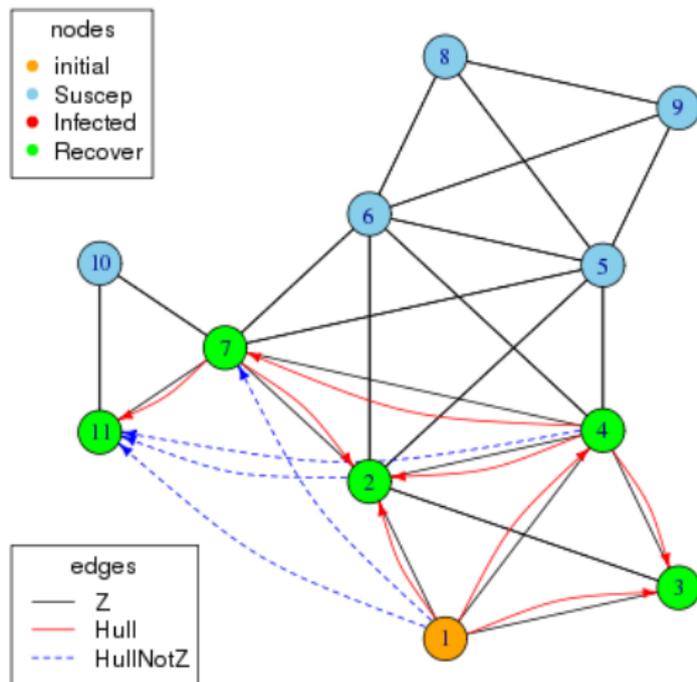
- ▶ consider individuals  $i, j, \dots \in \{1, \dots, N\}$ ; let  $G_{ij} = 1$  if  $i$  and  $j$  make contact capable of spreading disease.
- ▶  $\mathbf{G} = (G_{ij})$  is the *adjacency matrix* of a network / graph  $G \in \mathcal{G}$ , where  $\mathcal{G}$  is the set of all undirected graphs with  $N$  nodes.
- ▶  $k_i := \sum_j G_{ij}$  is the *degree* of node  $i$
- ▶ Individual  $i$  is in disease state  $X_i(t)$ , a discrete random variable; for e.g. network SIR dynamics we have:

$$\mathbb{P}[X_i(t + \delta) = I | X_i(t) = S] = \sum_j \tau G_{ij} \mathbb{I}_{\{X_j(t)=I\}} \delta + o(\delta) ; \quad (1)$$

$$\mathbb{P}[X_i(t + \delta) = R | X_i(t) = I] = \gamma \delta + o(\delta) .$$

# Inference on Network: from observed epidemic

Newman example 1



# Networks: Models

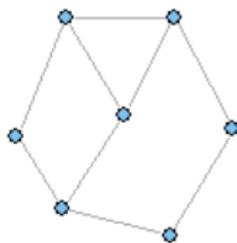
- ▶ Known or random
- ▶ Maths or simulation

The degree distribution  $P(k)$  the probability a randomly chosen node has degree  $k$  is the major factor affecting epidemics on graphs.

- ▶ Erdős Renyi
- ▶ Configuration model
- ▶ Small world

# Graphs with same degree distribution

572



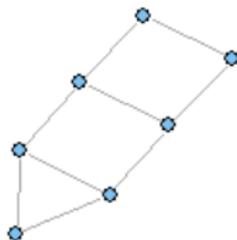
1 automorphisms

573



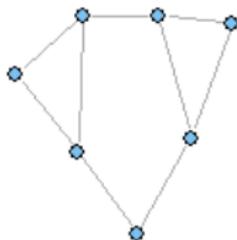
2 automorphisms

574



2 automorphisms

575



2 automorphisms

# Graph - Automorphisms

atlas number	572	573	574	575	576
automorphisms	1	2	2	2	4
number isomorphic	5040	2520	2520	2520	1260
clustering	0.6052	0.6225	0.5762	0.5905	0.5164
transitivity	.2	0	.2	.4	.6

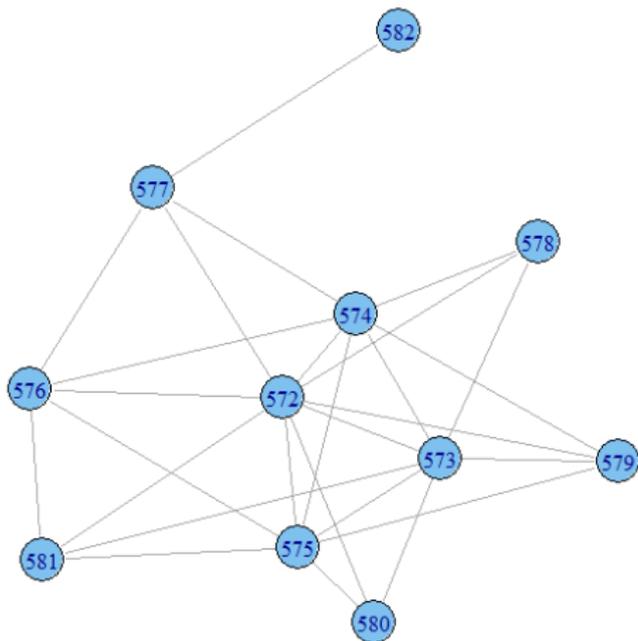
number	577	578	579	580	581	582
auto.	4	4	4	6	6	144
isomor.	1260	1260	1260	840	840	35
cluster	0.5701	0.6082	0.6082	0.5861	0.5861	0.2286
transit	.4	0	.2	.2	0	1.

all graphs with 7 nodes; 3 nodes of degree 2 and 4 nodes degree 3

# Rewiring algorithm

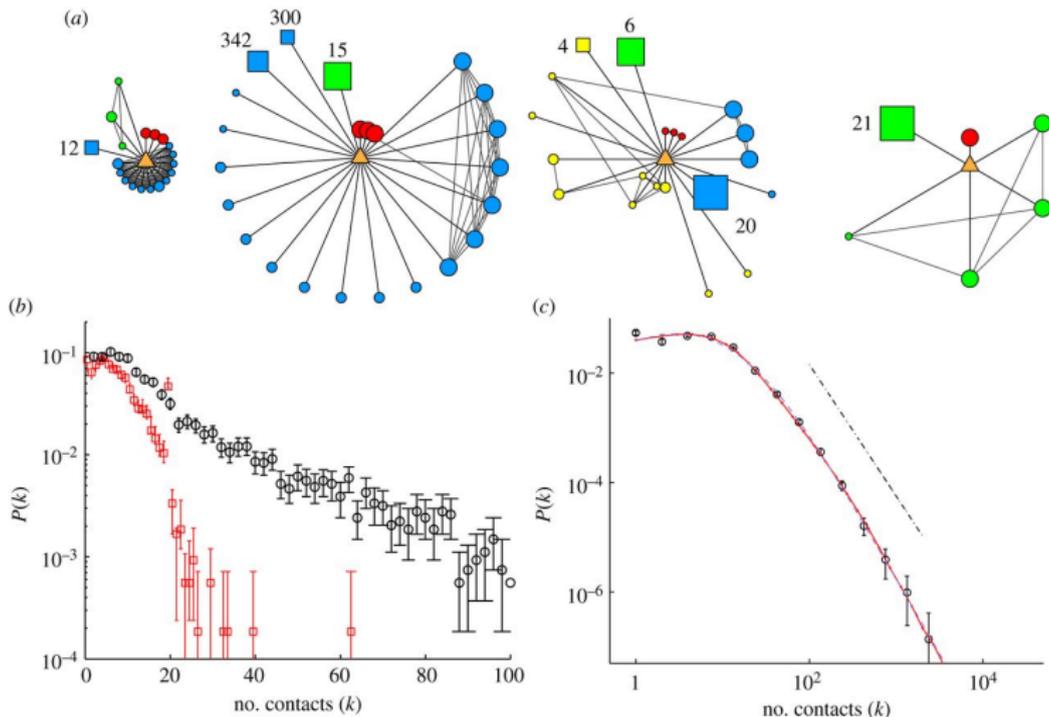
swap pairs of edges

rewires



# Networks: Heterogeneity

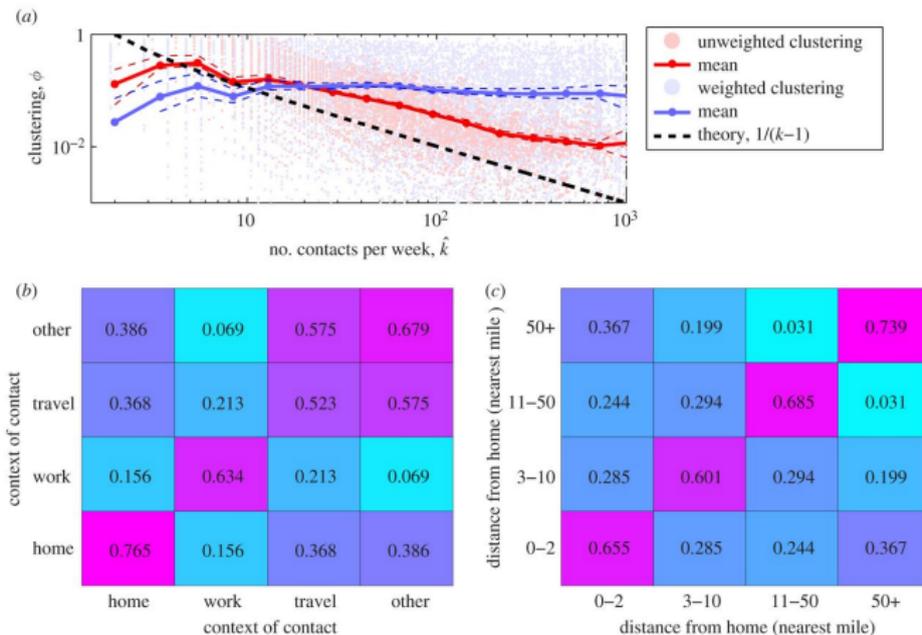
Measurement of direct contact patterns shows complex local structure and significant heterogeneity in degree<sup>†</sup>.



<sup>†</sup>L. Danon, T. House, J. M. Read and M. J. Keeling, "Social encounter networks: collective properties and disease transmission," *Journal of the Royal Society Interface*. **9**:76 (2012) 2826-2833.

# Networks: Clustering

Measurement also shows significant transitivity / clustering, generated more by context than space<sup>†</sup>.



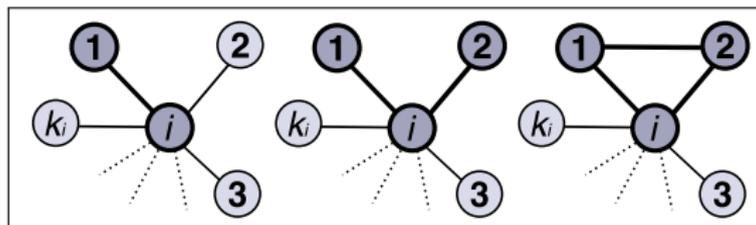
<sup>†</sup>L. Danon, **T. House**, J. M. Read and M. J. Keeling, "Social encounter networks: collective properties and disease transmission," *Journal of the Royal Society Interface*. **9:76** (2012) 2826-2833.

# Networks: Heterogeneity and Clustering

Suppose we want to generate full networks with given heterogeneity and clustering; a natural approach is to promote the  $G_{ij}$  to random variables. We want to specify:

$$\begin{aligned}M[G] &:= \sum_{i,j} G_{ij} = N\langle k \rangle \quad (\text{total links}), \\L[G] &:= \sum_{i,j,k \neq j} G_{ij}G_{ik} = N\langle k(k-1) \rangle \quad (\text{lines}), \\T[G] &:= \sum_{i,j,k} G_{ij}G_{jk}G_{ki} = \quad (\text{triangles}).\end{aligned} \tag{2}$$

Note, these are not independent:



# Networks: Measures

We will define a probability measure using a Hamiltonian framework so that the probability given to  $G \in \mathcal{G}$  is

$$\pi(G|\boldsymbol{\theta}) = \frac{e^{-H(G;\boldsymbol{\theta})}}{\mathcal{Z}_{\boldsymbol{\theta}}} , \text{ where } \mathcal{Z}_{\boldsymbol{\theta}} = \sum_{G \in \mathcal{G}} e^{-H(G;\boldsymbol{\theta})} . \quad (3)$$

We will also write  $\tilde{\pi}$  for the ensemble probability of a network taking a value of properties  $\mathbf{F}[G] = (F_a[G])$

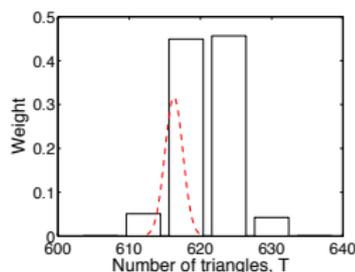
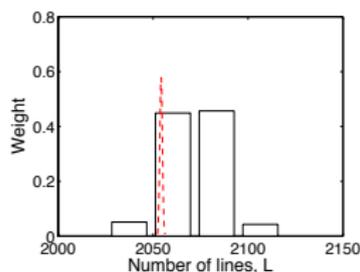
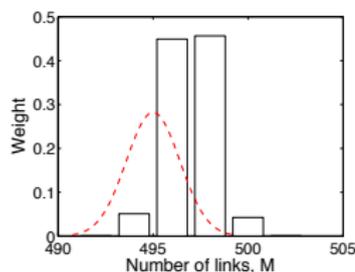
$$\tilde{\pi}(\mathbf{f}|\boldsymbol{\theta}) = \sum_{G \in \mathcal{G}} \pi(G|\boldsymbol{\theta}) \prod_a \mathbb{I}_{\{f_a = F_a[G]\}} . \quad (4)$$

It is easy to sample according to  $\pi$  in a Monte Carlo scheme, but hard to sample according to  $\tilde{\pi}$  for combinatorial reasons.

# Networks: A heterogeneous clustered measure

The following measure works if parameters are tuned appropriately:<sup>†</sup>

$$H(G; \theta, \beta) = \beta_m (M[G] - \theta_m N(N - 1))^2 + \beta_l (L[G] - M[G]((\theta_l - 1) + N^{-1}M[G]))^2 + \beta_t (T[G] - \theta_t L[G])^2 .$$



(Red dashed line – target; Black histograms – simulations)

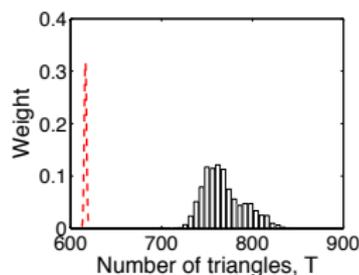
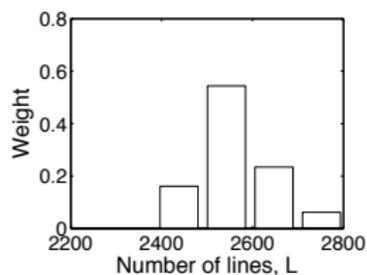
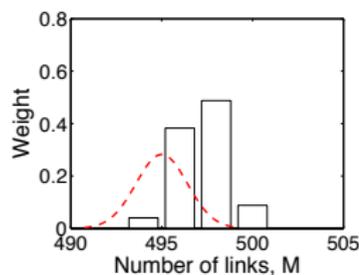
<sup>†</sup> T. House, "Heterogeneous clustered random graphs," Europhysics Letters **105** (2014) 68006.

# Networks: A heterogeneous clustered measure

Removing just one term causes failure:

$$H(G; \theta, \beta) = \beta_m (M[G] - \theta_m N(N - 1))^2$$

~~$$+ \beta_l (L[G] - M[G]((\theta_l - 1) + N^{-1}M[G]))^2 + \beta_t (T[G] - \theta_t L[G])^2 .$$~~



(Red dashed line – target; Black histograms – simulations)

# ERGM and Phase transitions in graphs

The theory explains a host of difficulties encountered by applied workers: many distinct models have essentially the same MLE, rendering the problems “practically” ill-posed. We give the first rigorous proofs of “degeneracy” observed in these models. showing that for many models, the extra sufficient statistics are useless: most realizations look like the results of a simple Erdős-Rényi model. We also find classes of models where the limiting graphs differ from Erdős-Rényi graphs. A limitation... it works only for dense graphs. †

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†Chatterjee and Diaconis, 2011, Annals of Statistics

# Motivation for bipartite models

- ▶ consider individuals  $i, j, \dots \in \{1, \dots, N\}$ ;
- ▶  $\lambda_{ij}$  is the infection rate between  $i$  and  $j$ 
  - ▶ when  $i$  is infectious and  $j$  is susceptible.
- ▶  $\lambda_{ij} = d(i, j)\rho(i|x_i)\psi(j|x_j)$
- ▶ Ebola know the village and household level matter
- ▶ 3 level schools, workplace, household

## Example bipartite graph

A bipartite graph  $G = (U, V, E)$  is two disjoint sets  $U$  and  $V$  comprising the nodes or vertices<sup>†</sup>, and a set of edges  $E$  where each edge is a pair of nodes  $(u, v)$ ,  $u \in U, v \in V$ . In general the two sets  $U$  and  $V$  can be of the same type so that all vertices in  $U$  and  $V$  are in some larger set of vertices, here  $U$  and  $V$  are distinct with  $U$  representing individuals who may become infected and  $V$  an abstract set of possible contacts, which may include physical premises such as schools, houses or work places and can include a temporal aspect.

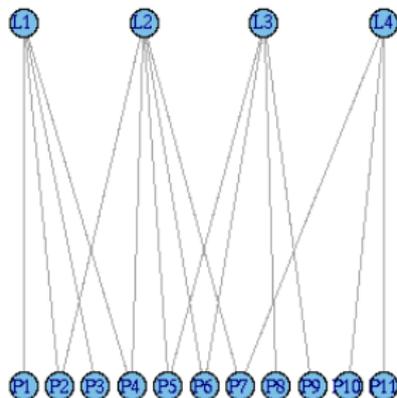
A convenient representation is the adjacency matrix

$\mathbf{A} = (a_{ij}, i \in U, j \in V)$  where  $a_{ij} = 1$  if and only if  $(i, j) \in E$  and  $a_{ij} = 0$  otherwise.

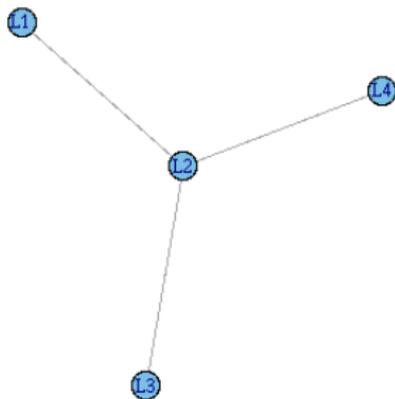
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<sup>†</sup>the terms are used interchangeably

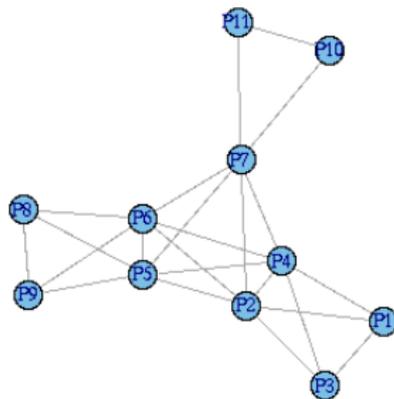
# Example bipartite graph



Top



Lower



# Bipartite representation of a household model

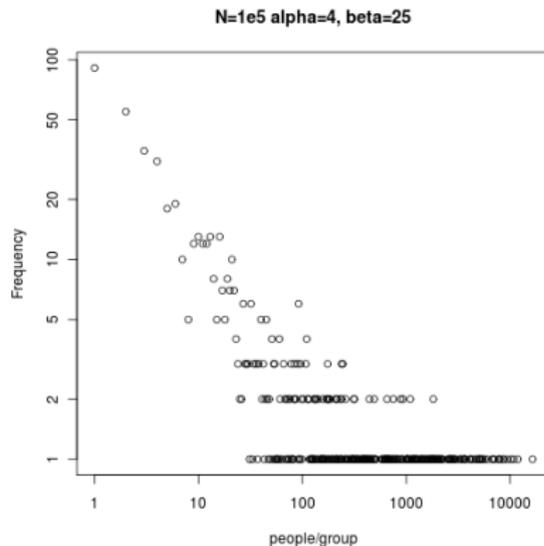
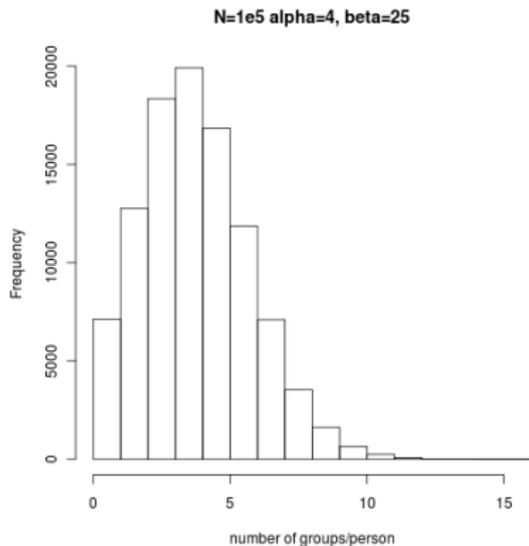
$\lambda_G$	$\lambda_H$	$\lambda_H$	$\lambda_H$	$\lambda_H$
1	1			
1	1			
1		1		
1		1		
1		1		
1			1	
1			1	
1			1	
1				1
1				1
1				1
1				1

other standard models can also be represented

# Random bipartite graphs

- ▶ Random Intersection Graphs
  - ▶ Theoretical results but not realistic
- ▶ Hypergraphs
  - ▶ Theoretical results
- ▶ Indian Buffet Process
  - ▶ Realistic but theory challenging

# Example degree distributions of IBP

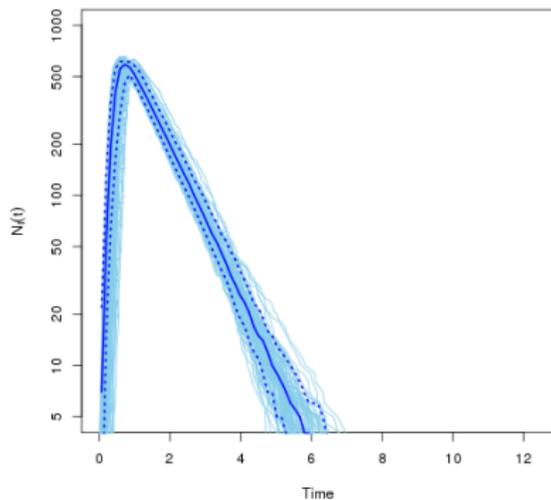


Distribution of marginal sums for an example IBP,  $N = 10^5$ ,  $\alpha = 4$ ,  $\beta = 25$ .

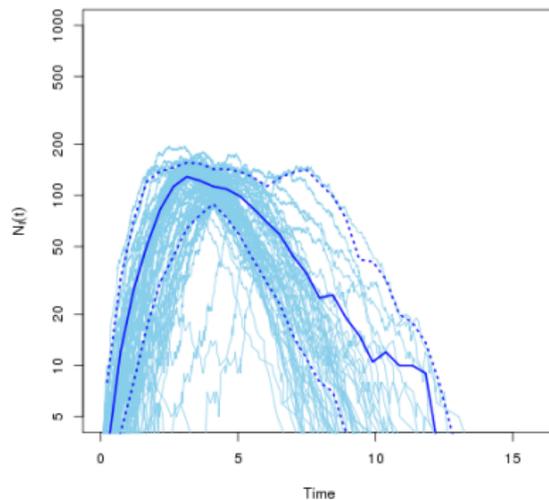
The left hand plot is a standard histogram, the right hand a count of counts.

# Example epidemics on Hypergraphs

HGeg4c,beta=1, gamma=1



HGeg4a,beta=1, gamma=1



# Conclusions

- ▶ Bipartite Graphs are a useful model
- ▶ Lots of interesting problems in distributions of graphs and hypergraphs

# Acknowledgements / Thanks

- ▶ WIDER (environment)
- ▶ EPSRC (my funding)
- ▶ The organisers, ECOFECT and your sponsors