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# Contact patterns of inpatients in a regional healthcare system

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KTH, CSC, Computational Biology

May 17, 2008, PhyDIS Workshop

<http://www.csc.kth.se/~pholme/>

- $R_0$ —the expectation value of number of others an infectious individual will infect in a susceptible population.
- The crudest approximation: There can be an epidemic in a population if  $R_0 > 1$ .

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# Epidemiology 101

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- The crudest approximation: There can be an epidemic in a population if  $R_0 > 1$ .

# $R_0$ of Smallpox

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| Outbreak               | $R_0$   |
|------------------------|---------|
| Boston, USA (1721)     | 4.3     |
| Burford, UK (1758)     | 3.4     |
| Paris, France (1766)   | 4–5     |
| Warrington, UK (1773)  | 4.0–5.3 |
| Chester, UK (1774)     | 5.8     |
| London, UK (1836–1870) | ~ 5     |
| Europe (1958–1973)     | 10–12   |
| Kosovo (1972)          | 10.8    |

# Stay out of hospitals!

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- Infectious diseases are a big threat to public health.
- Hospitals have a key-role in spreading of many kinds of disease.
- In epidemics of smallpox, TB, Ebola and SARS hospitals have played a crucial role.
- Other pathogens like MRSA, norovirus or *Mycoplasma pneumoniae* can be endemic within a health care system.

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# network epidemiology

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- This work: How does the relevant network look like for nosocomial infections.
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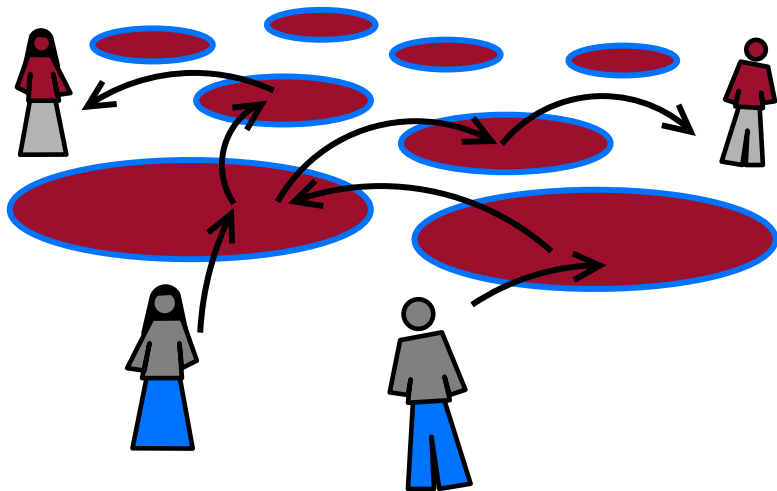
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# A health care system



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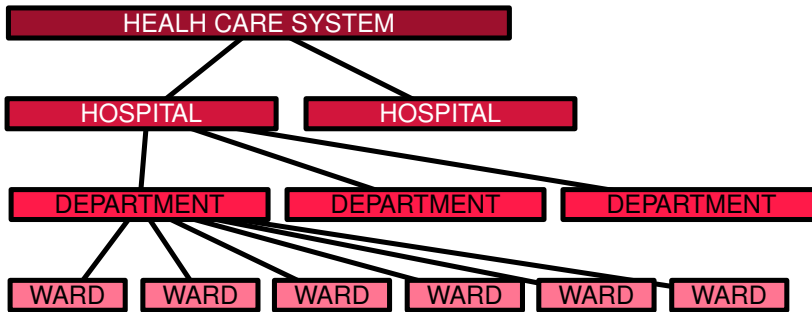
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- All hospitalizations of people in the Stockholm region of Sweden the years 2001 and 2002.
- 1.7 million inhabitants.
- 570,382 hospitalizations.
- 295,108 patients.
- 702 wards.



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- What we want: Construct simple graphs where edges represent “high enough” probability of disease transmission.
- Measure network structure → use theoretical results to say something about how contact patterns affect disease spread. “high enough” probability of disease transmission.
- Could be improved to valued graphs.



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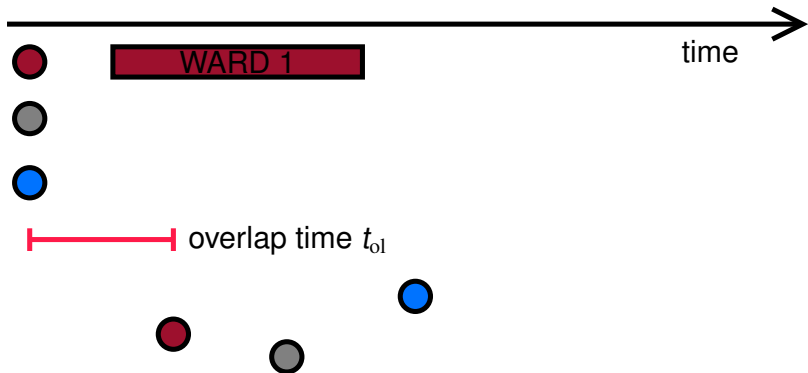
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# Constructing the networks

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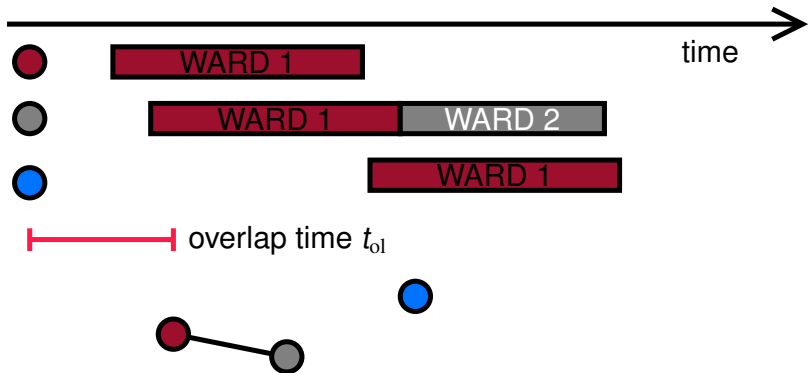
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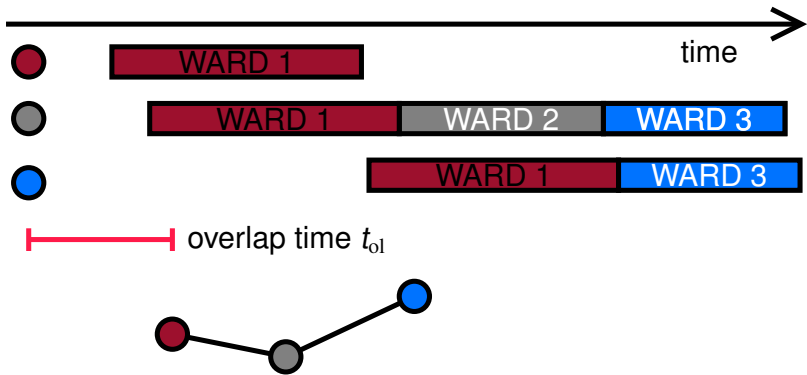
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# Constructing the networks



# Corrected reproduction number

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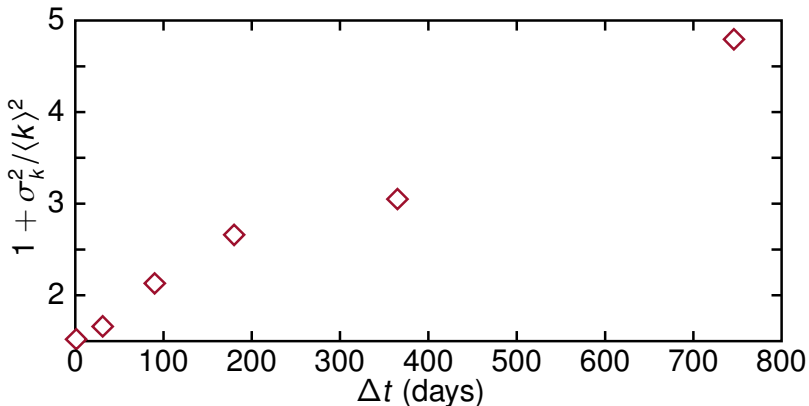
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Fluctuations in number of contacts  $\Rightarrow R_0$  needs to be corrected by a factor  $1 + \sigma_k^2 / \langle k \rangle^2$





# Clustering coefficient

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- Measuring the density of triangles.
- High clustering coefficient — slow growth.





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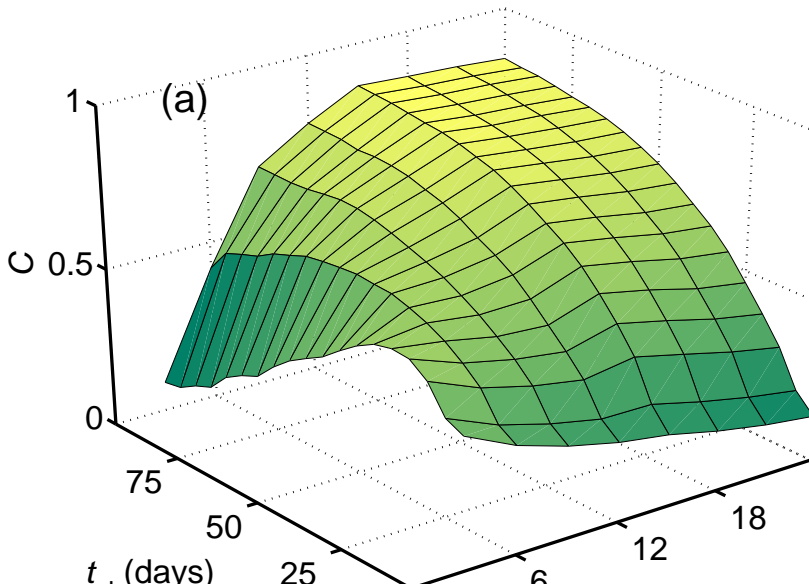
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- Measuring the correlations of degree at either side of an edge.
- Large assortativity — high probability of epidemic outbreaks, low outbreak sizes (if an outbreak occurs).
- Large disassortativity — low probability of epidemic outbreaks, large outbreak sizes.

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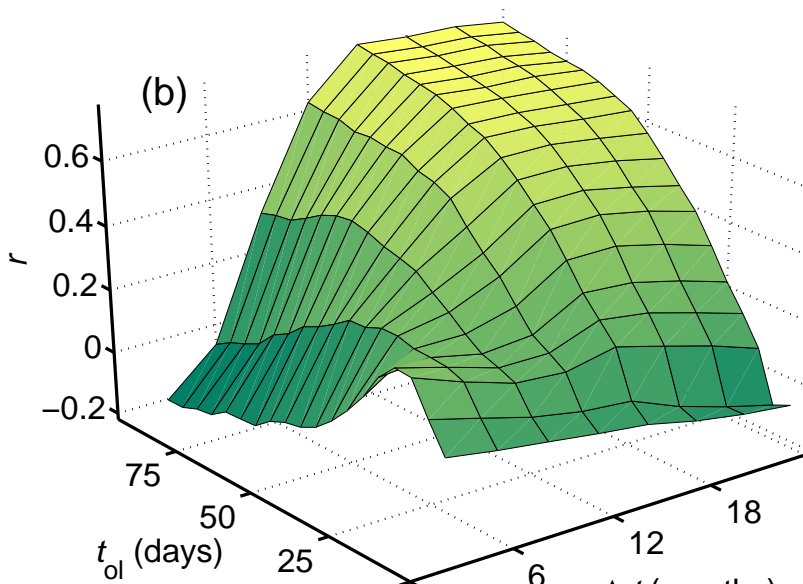
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# An agent-based model

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- A population of  $N$  individuals.
- A health care system with  $N_w$  wards of equal capacity.
- Each healthy (non-hospitalized) agent is, with probability  $p_1$ , hospitalized at a random ward.
- A hospitalized patient is assigned a duration  $t \in P_t$  of the hospital stay.
- After a  $t$  days the patient is, with probability  $p_2 > p_1$ , rehospitalized at a random ward.

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# Results for the model

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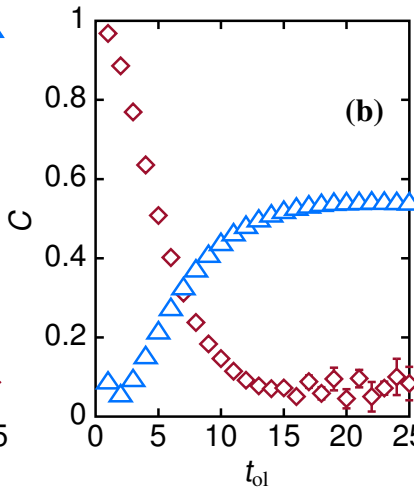
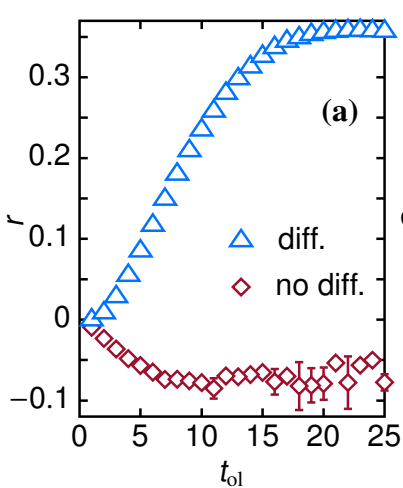
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# Summary & conclusions

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- Disease spreading within a health care system is important.
- A health care system for a population of 1.7 million → proximity networks of patients.
- Power-law distribution of hospitalization times & a skewed degree distribution.
- Both clustering and assortative mixing coefficients increase with both sampling and overlap times.
- The differentiation of hospitalization times per ward is necessary to explain this
- Future work include e.g. dynamic modeling of disease spreading.

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