

# Stochastic Modelling of the Spatial Spread of Influenza in Germany

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# Influenza

- Worldwide one of the **most common and severe** infectious diseases.
  - Major epidemics and pandemics of the 20th century: **Spanish Flu** (1918-20), **Asian Flu** (1957-58), **Hong Kong Flu** (1969)
  - Annual number of deaths caused by influenza in Germany is twice as high as those caused by road accidents, nevertheless **low vaccination rates**.
  - Steadily **new antigen mutants** of the influenza virus coming up.
  - In the opinion of experts the next pandemic is just a question of time (caused e.g. by avian flu).
- ⇒ Emergency plans are essential.

# Motivation

- Mathematical modelling of the spread of epidemics dates back to 1760 (Daniel Bernoulli) and has been well developed since then.
- Globalization has induced "new era" of epidemiology: People travel more frequently, faster and further than in former times.
- Hufnagel et al. (2004) have successfully simulated the world-wide spatial spread of the SARS epidemic in 2003 using a spatial SIR model based on global air traffic data.

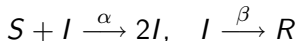
# Outline

- 1 Introduction
- 2 Model
- 3 Implementation and Initialization
- 4 Simulation
- 5 Conclusion

## Standard SIR Model

- Divide population of size  $N$  into **susceptible** ( $S$ ), **infected** ( $I$ ), and **removed** ( $R$ ) individuals.

- Transitions:



$\alpha$ : contact rate of an infectious individual sufficient to spread the disease,

$\beta$ : reciprocal average infectious period.

- Infection dynamics:

$$ds/dt = -\alpha sj, \quad dj/dt = \alpha sj - \beta j, \quad (1)$$

$$s = S/N, \quad j = I/N, \quad r = R/N = 1 - s - j.$$

## Standard SIR Model (cont.)

Crucial parameter: the **basic reproduction number**  $\rho = \alpha/\beta$ , *the average number of persons directly infected by an infectious case during its entire infectious period after entering a totally susceptible population.*

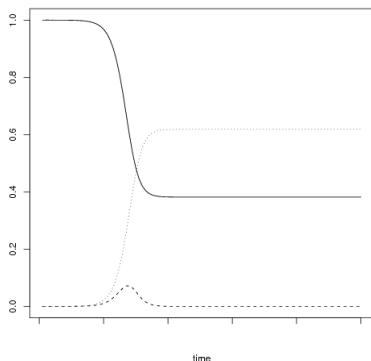
$\rho^{-1} > s(0)$ : no epidemic will occur

$\rho^{-1} > s(t)$ : epidemic decays,

i.e. major epidemic will occur when, in the early stages of an outbreak, each infective on average produces more than one further infective.

## Standard SIR Model (cont.)

Evolution of proportions of susceptible (solid), infected (dashed), and removed (dotted) individuals in the standard SIR model;  $\rho = 1.5$ .



# Stochastic SIR Model

Infection and recovery processes are of rather stochastic than deterministic character.

⇒ Write (1) in terms of Langevin equations:

$$\begin{aligned}\frac{ds}{dt} &= -\alpha sj + \frac{1}{\sqrt{N}} \sqrt{\alpha sj} \xi_1(t) \\ \frac{dj}{dt} &= \alpha sj - \beta j - \frac{1}{\sqrt{N}} \sqrt{\alpha sj} \xi_1(t) + \frac{1}{\sqrt{N}} \sqrt{\beta j} \xi_2(t),\end{aligned}$$

where  $\xi_1(t)$  and  $\xi_2(t)$  are independent Gaussian **white noise** forces, modelling fluctuations in disease transmission and recovery (Hufnagel et al., 2004).

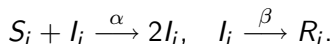


# Spatial SIR Model

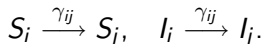
**Problem:** Assumption of homogeneous mixing is not given in our fully connected world anymore!

**Idea:** Introduce network of subregions  $i = 1, \dots, n$  of sizes  $N_i$ .

- **Local** infection dynamics within a subregion is given by stochastic SIR model as before:



- **Global** dispersal between nodes of network is rated in a connectivity matrix  $\gamma = (\gamma_{ij})_{ij}$ :



## Spatial SIR model (cont.)

The system of stochastic differential equations now changes to

$$\begin{aligned} \frac{ds_i}{dt} &= -\alpha s_i j_i - \sum_k \gamma_{ik} s_i + \sum_k \gamma_{ki} s_k + \frac{1}{\sqrt{N_i}} \sqrt{\alpha s_i j_i} \xi_1^{(i)}(t) \\ &\quad + \frac{1}{\sqrt{N_i}} \sqrt{\sum_k \gamma_{ik} s_i} \xi_4^{(i)}(t) - \frac{1}{\sqrt{N_i}} \sqrt{\sum_k \gamma_{ki} s_k} \xi_5^{(i)}(t) \\ \frac{dj_i}{dt} &= \alpha s_i j_i - \beta j_i - \sum_k \gamma_{ik} j_i + \sum_k \gamma_{ki} j_k - \frac{1}{\sqrt{N_i}} \sqrt{\alpha s_i j_i} \xi_1^{(i)}(t) + \frac{1}{\sqrt{N_i}} \sqrt{\beta j_i} \xi_2^{(i)}(t) \\ &\quad + \frac{1}{\sqrt{N_i}} \sqrt{\sum_k \gamma_{ik} j_i} \xi_4^{(i)}(t) - \frac{1}{\sqrt{N_i}} \sqrt{\sum_k \gamma_{ki} j_k} \xi_5^{(i)}(t) \\ \frac{dr_i}{dt} &= \beta j_i - \frac{1}{\sqrt{N_i}} \sqrt{\beta j_i} \xi_2^{(i)}(t). \end{aligned}$$

for  $i = 1, \dots, n$ , where  $\xi_1(\mathbf{t})$ ,  $\xi_2(\mathbf{t})$ ,  $\xi_4(\mathbf{t})$ , and  $\xi_5(\mathbf{t})$  denote independent vector-valued white noise forces standing for fluctuations in transmission, recovery, and outbound and inbound traffic, respectively.

# Keeping the System Closed

Area of  $n$  regions is assumed to be closed. i.e. we have to require

$$\sum_{i=1}^n \left( \frac{ds_i}{dt} + \frac{dj_i}{dt} + \frac{dr_i}{dt} \right) = 0 .$$

⇒ Introduce a weak form of dependence to the white noise forces such that the above equality holds almost surely.

## Numerical Scheme

Define functions  $a_j$  and  $b_{jk}$ ,  $1 \leq j \leq 3$ ,  $1 \leq k \leq 5$ , such that

$$ds_i(t) = a_1(t, s_i(t))dt + \sum_{k=1}^5 b_{1k}(t, s_i(t))dW_k^{(i)}(t)$$

$$dj_i(t) = a_2(t, j_i(t))dt + \sum_{k=1}^5 b_{2k}(t, j_i(t))dW_k^{(i)}(t)$$

$$dr_i(t) = a_3(t, r_i(t))dt + \sum_{k=1}^5 b_{3k}(t, r_i(t))dW_k^{(i)}(t).$$

## Numerical Scheme (cont.)

For example, for  $i = 1, \dots, n$ :

$$\begin{aligned}
 ds_i = & \underbrace{\left( -\alpha s_i j_i - \sum_k \gamma_{ik} s_i + \sum_k \gamma_{ki} s_k \right)}_{:=a_1(t, s_i(t))} dt + \underbrace{\frac{1}{\sqrt{N_i}} \sqrt{\alpha s_i j_i}}_{:=b_{11}(t, s_i(t))} \xi_1^{(i)}(t) dt \\
 & + \underbrace{\frac{1}{\sqrt{N_i}} \sqrt{\sum_k \gamma_{ik} s_i}}_{:=b_{14}(t, s_i(t))} \xi_4^{(i)}(t) dt - \underbrace{\frac{1}{\sqrt{N_i}} \sqrt{\sum_k \gamma_{ki} s_k}}_{:=b_{15}(t, s_i(t))} \xi_5^{(i)}(t) dt,
 \end{aligned}$$

$$b_{12}(t, s_i(t)) = b_{13}(t, s_i(t)) = 0, \quad \xi_k^{(i)}(t) dt = dW_k^{(i)}(t).$$

## Numerical Scheme (cont.)

Apply **Euler-Maruyama approximation scheme** to numerically solve the system of SDEs at discrete, equidistant instants  $0, \delta, 2\delta, \dots$  in the time domain:

$$s_i(m\delta) = s_i((m-1)\delta) + a_1((m-1)\delta, s_i((m-1)\delta))\delta + \sum_{k=1}^5 b_{1k}((m-1)\delta, s_i((m-1)\delta)) \Delta W_k^{(i)}(m)$$
$$j_i(m\delta) = j_i((m-1)\delta) + a_2((m-1)\delta, j_i((m-1)\delta))\delta + \sum_{k=1}^5 b_{2k}((m-1)\delta, j_i((m-1)\delta)) \Delta W_k^{(i)}(m)$$
$$r_i(m\delta) = r_i((m-1)\delta) + a_3((m-1)\delta, r_i((m-1)\delta))\delta + \sum_{k=1}^5 b_{3k}((m-1)\delta, r_i((m-1)\delta)) \Delta W_k^{(i)}(m)$$

for  $m \geq 1$ ,  $i = 1, \dots, n$  and  $\Delta W_k^{(i)}(m) := W_k^{(i)}(m\delta) - W_k^{(i)}((m-1)\delta)$ .

## Application: Influenza in Germany

- Consider the 438 districts of Germany.
- Data about incidences of influenza taken from Robert Koch Institute.
- Build up **connectivity matrix** to describe the strength between parts of Germany, considering
  - dispersal between **adjacent regions**, caused e.g. by commuters (info from Federal Statistical Office Germany),
  - domestic **train traffic** (ICE),
  - domestic **air traffic** ([www.oagflights.com](http://www.oagflights.com)),

where each of these components is provided with a weight regulating its influence.

## Parameter choice

- We assume that the **basic reproduction number**  $\rho = \alpha/\beta$  depends on population density:

$$\rho(d_i) = 1.0179 + 10^{-5} \cdot d_i,$$

where  $d_i$  is the population density of region  $i$ . (Disease is more likely to spread in areas with high population densities.)

- **Infectious period** of influenza: 4-5 days, hence we choose  $\beta = 2/9$ .
- We obtain the **contact rates**  $\alpha_i$  via  $\beta \cdot \rho(d_i)$ ,  $i = 1, \dots, n$ .



# Simulation

## ① Simulation 1:

- Starting values based on surveillance data from week 5/2005.
- Animation shows **proportion of infectives** and **time trend**.
- Surprisingly good agreement with actual course of the influenza epidemic 2005.

## ② Simulation 2:

- Artificial starting values in three districts.
- Simulation based on **three different connectivity matrices**:
  - ① local, train and air,
  - ② local and train,
  - ③ only local.

# Conclusion

- Spatial extension of classical SIR model.
- Implementation of certain sum-to-zero constraints and numerical approximation.
- Parameter choice based on external knowledge.
- Simulation of the spread of an influenza epidemic in Germany.

# Construction Sites

- Prevalence data of influenza is highly underreported.
  - Simulate underreporting.
  - Other data sources (Sentinella).
  - Consider different diseases.
- Model does not take into account important factors like
  - weather/temperature,
  - seasonal holiday travel,
  - measures undertaken to lower transmission rate.

## Future Work

Main purposes will be

- finding surveillance strategies in case of a sudden outbreak of an epidemic (isolation, vaccination, observation of migration),
- more formal statistical inference on model parameters based on available data from surveillance databases.

## References

- Dargatz, C., Georgescu, V. and Held, L. (2005) Stochastic Modelling of the Spatial Spread of Influenza in Germany. Technical Report, Munich University.
- Hufnagel, L., Brockmann, D. and Geisel, T. (2004). Forecast and Control of Epidemics in a Globalized World. *Proceedings of the National Academy of Sciences*, **101**, 15124-15129.