Computer Simulations

A practical approach to simulation

Semra Gündüç

gunduc@ankara.edu.tr

Ankara University Faculty of Engineering,

Department of Computer Engineering

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Introduction to Epidemics

Computational simulation of epicemics provide a valuable tool for the study the dynamics of epidemics.

- The spread of infectious deseases such as influenza, smallpox, HIV/AIDS are studied.
- It is assumed that for contamination direct contacts between the contaminated and the susceptible individuals are necessary.
- The validity of assumptions can be tested by using available epidemics data.



Introduction to Epidemics

Computational tools of studying epidemics varies.

- Epidemics can be modelled by using different techniques such as
 - ordinary diferential equations (mean-field),
 - partial differential equations (spatial distribution is taken into consideration),
 - Cellularautomatom (simple local rules),
 - stochastic dynamics (Agent based simulation, random walk, Monte-Carlo techniques)



Introduction to Epidemics

Computational tools of studying epidemics can be used to study many apparently different pnenomena.

- Modeling of social systems
 - naming, opinion spread, population dynamics, rumour propagation.
 - . . .
- Finance Modeling economics
- Population dynamics

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Introduction to Epidemics

Elements of the Epidemic Simultion:

- Susceptible
- Infected
- removed

The dynamics of the epidemics spread:

- Susceptible individuals become infected at rate proportional to the number of infected neighbours.
- Infected individuals become healty at a fixed rate δ
- Removed individuals become susceptible at a fixed rate α

An ordinary differential equation approach

- u : Fraction of infected
- v : Fraction of removed
- 1 u v : Fraction of susceptible

Assumption: Large, mixed, homogeneous population (Mean-Field Approximation)

$$\frac{du}{dt} = u \cdot (1 - u - v)) - \delta u$$
$$\frac{dv}{dt} = \delta u - \alpha v$$

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An ordinary differential equation approach

- If $\alpha = 0$, recovered individuals do not get contaminated again.
 - If δ is big, epidemics dies out
 - for $\delta < \delta_c$: Epidemic spreads linearly with time. where δ_c is a critical value of δ . Approach to steady state distribution.
- If $\alpha > 0$, problem is more complicated.

Referance : Dolgoarshinnykh, R. G. and Lalley, S. P. (2006), "Critical scaling for the SIS stochastic epidemic," J. Appl. Probab., 43, 892–898.

An ordinary differential equation approach

t_Final = 30; // Time to end calculation dt = 0.01; // Time steps t = 0; // Start clock while(t < t_Final) {</pre> u1 = u0 + (u0 * (1.0 - u0 - v0) - delta * u0)v1 = v0 + (delta * u0 - alpha * v0) * dt;u0 = u1;v0 = v1;t = t + dt;fprintf(out1," %f %f %f %f \n",t,1-u0-v0,u0,v0) }

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An ordinary differential equation approach

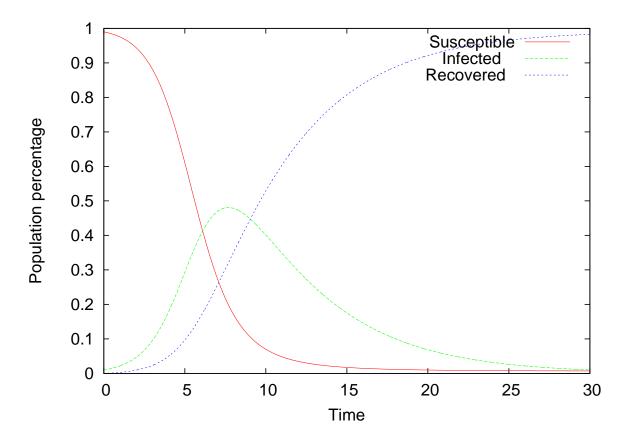


Figure 1: Epidemic spread $\alpha = 0.0$, $\delta = 0.2$.

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An ordinary differential equation approach

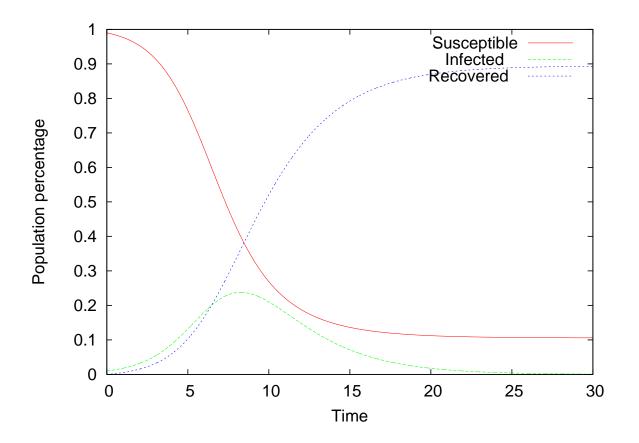


Figure 2: Epidemic spread $\alpha = 0.0$, $\delta = 0.4$.

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An ordinary differential equation approach

- Measle spread in Glasgow 1929
 - Measles took 24 weeks to spread over Glasgow. approximtely 150 meters per week
 - This is approximately linear spred rate of rakoon.
- Speread of muskrats over central Europa during a period of 27 years.

Spatial distribution is important. Mean-Field approach do not count

the spatial distribution of population.



Effects of spatial distribution of population

Simple spatial models of epidemics are known to exhibit critical thresholds in one dimension: Roughly, when the density of the initially infected set exceeds a certain level, the epidemic evolves in a markedly different fashion than its branching envelope for contact between the contaminated and susseptible person.

Effects of spatial distribution of population

Referances:

- 1. Lalley, S., "Spatial Epidemics: Critical Behavior in One Dimension," arXiv:math/0701698v2 (2007).
- 2. Aldous, D., "Brownian excursions, critical random graphs and the multiplicative coalescent," Ann. Probab., 25, 812–854 (1997).
- Martin-Löf, A., "The final size of a nearly critical epidemic, and the first passage time of a Wiener process to a parabolic barrier," J. Appl. Probab., 35, 671–682 (1998).

Partial differential equation approach

- Partial differential approach models similar phenomena. Only difference is that spatial distribution is taken into consideration.
- P(x,t) : Population density.

$$\frac{\partial P(x,t)}{\partial t} = \delta \Delta^2 P(x,t) + \alpha P(x,t)$$

The solution leads to linear spread reate. Partial differential equation solution approximates random walk.