

Local Outbreak Dynamics

A Basic Model for the Simulation of Local Infectious Disease Outbreaks

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Global Pandemic Influenca H1N1 Simulations





Simple FHS-Simulation for local H1N1 out brakes

Interest in an easy to use simulation instrument for analyzing local disease dynamics



Project Goals

Simulation Instrument

- Model of local outbrakes caused mainly by contact interactions
- Possibility to parameterize the setting (environment, type of virus, etc.)
- User-friendly interface with animation and visualization of simulation results

Scenario Analysis

Analysis of different control measures (i.e. hand wash, gloves, patient/HCW ratio, screening, isolation, etc.)

Awareness Training

- Tool and concept for the sensitizing of HCW



Basic Model Cases



Transmission of methicillinresistant Staphylococcus areus within an intensive care unit (KSSG)



Transmission of norovirus within a geriatric ward (SSH)



Basic Model Agents and Setting









Basic Model Modeling Paradigm

Systemdynamic (SD):



Rate R(t) = f(A, B, c, ...)

«Systemic Agents»:



Agentbased (ABM):





Basic Model Modelling Paradigm

Advantages

- Use of abstract systemic descriptions regarding the transmission mechanism (no need for detailed interaction datas)
- Possibility to have discrete events (i.e. change of shift, patient entry, etc.)
- Patients, HCW and visitors have a history
- Animation for better plausibilization through experts



Simulation Application

Application with a user-friendly interface is built with the open source modeling software GlobalSimulate[®]





Summary & Outlook

Goal

Creation of an easy to configure and to use instrument for simulating local disease dynamics

Status

Basic model building on a model paradigm in between SD and ABM

To dos

- Plausibilization and pointwise validation of the specific models
- Scenario analysis: effectiveness of different measures and strategies
- Further development of the user-interaction (GUI)
- Concept for awareness trainings with the simulation model



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