

Halle-Germany (Martin-Luther-University)



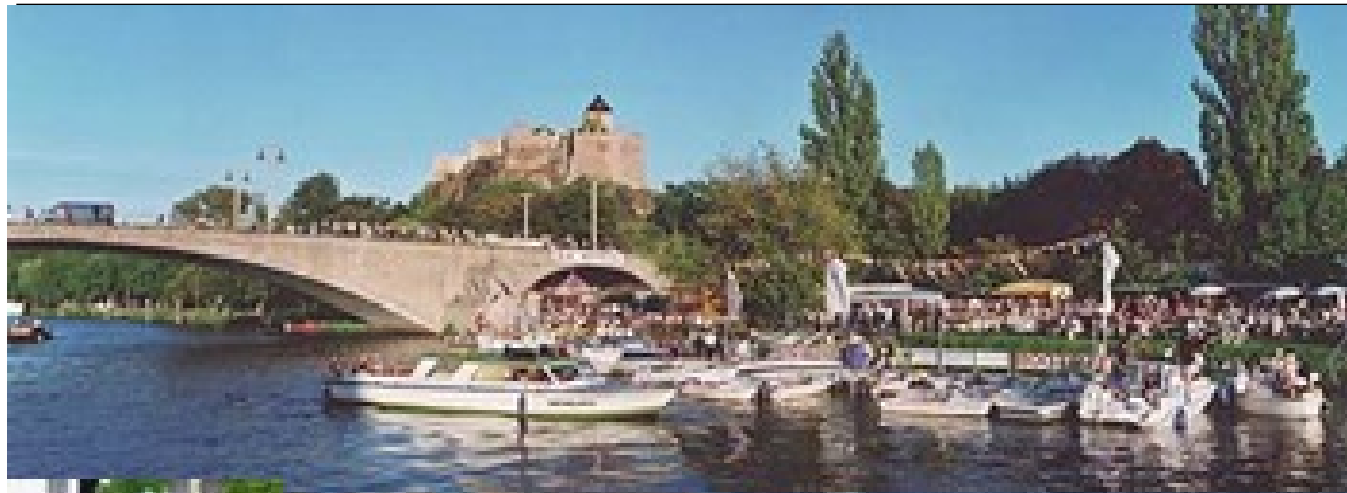
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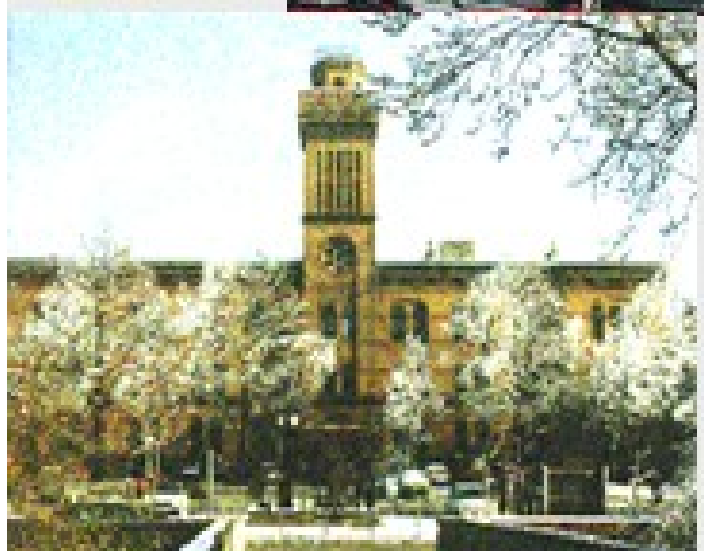
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JERUSALEM >> HALLE >> TEL AVIV

Halle
1200
years





EXACT SOLUTION OF A STOCHASTIC SIR-MODEL

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MODEL KEY WORDS

- **SIR**: Susceptible-Infected-Recovered Model
- **SIR**: Evolution of this three species
- **SIR**: Simple tractable model capturing relevant features
- **SIR** Non-linear model
- **SIR** Out of equilibrium
- **SIR** Stochastic description

Infection produces further infections



28/09/09

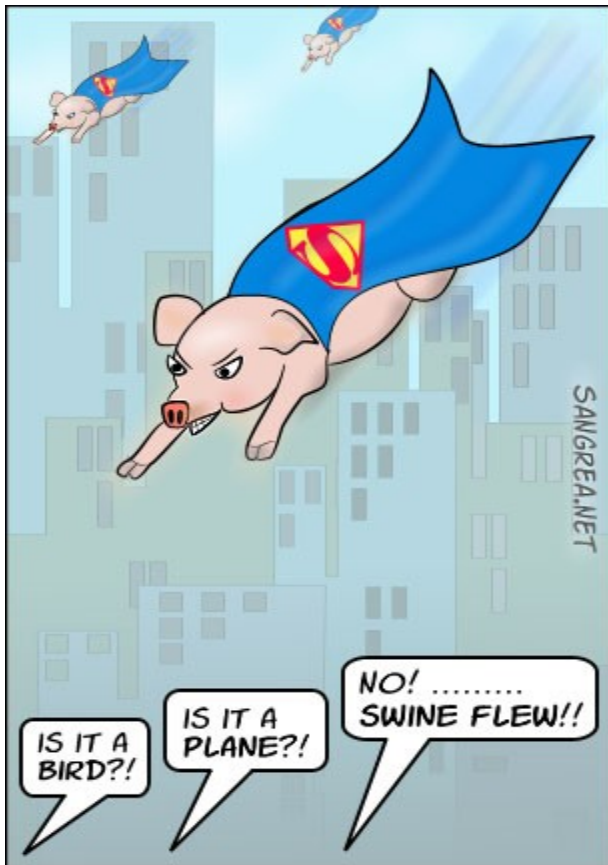
Newspaper Roanoke Time

REGION READY FOR FLU

Halle:

,Organized' Vaccination

Flew versus Flu



COMPLEX SYSTEMS

Comprise many
components

Interacting
non-linearly

System far from
equilibrium



Natural home of big
surprises

Emergent behavior

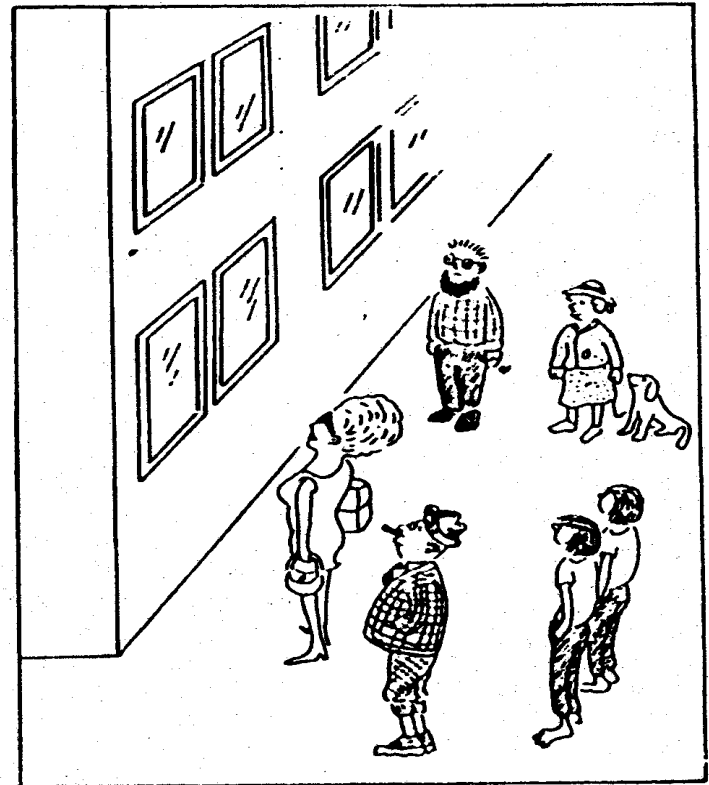
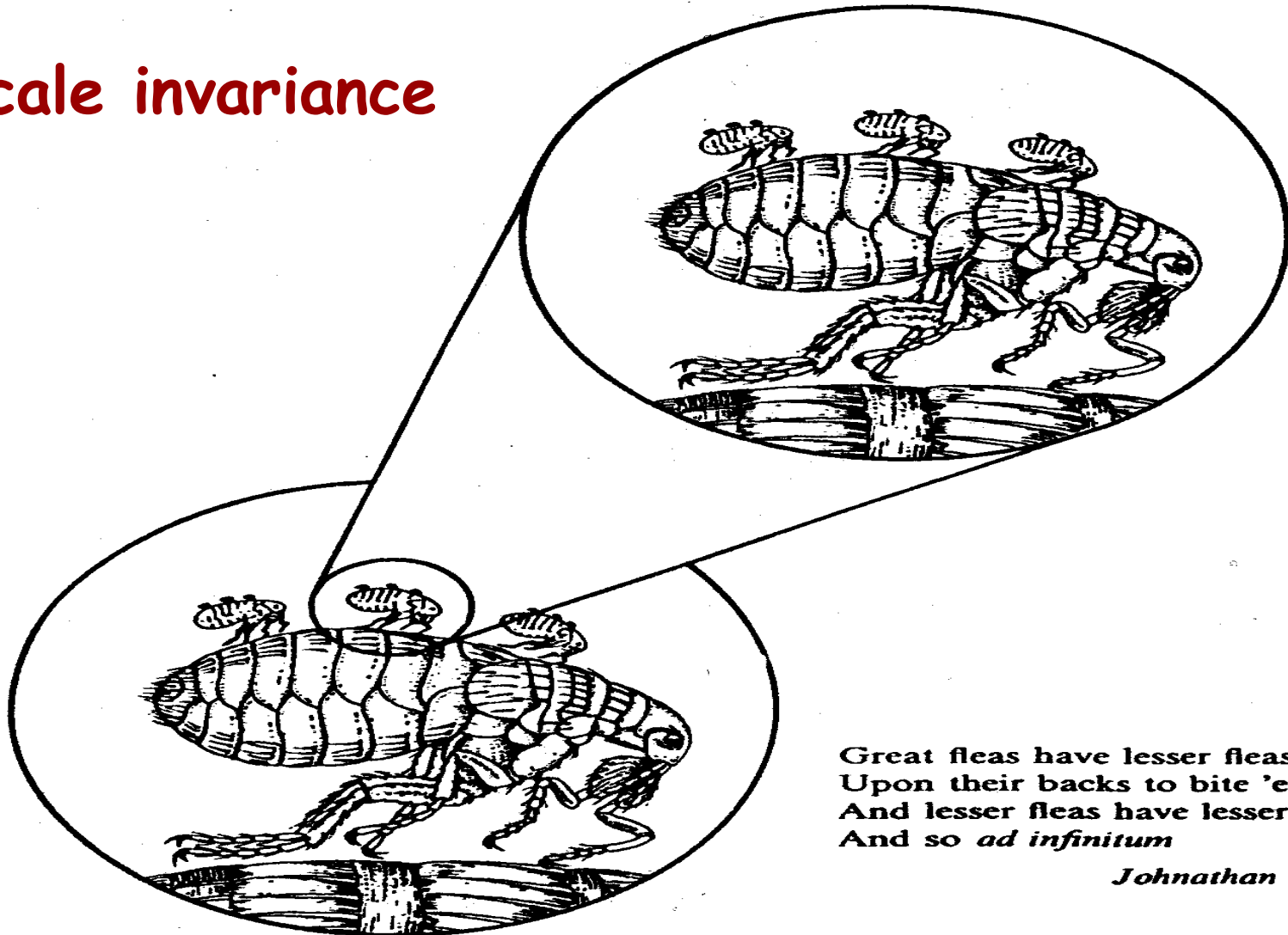


Abbildung 2.8: Ein Phasenübergang in der neugierigen Menge. (a) "Paramagnet";
(b) "Ferromagnet" Flash Mob

Scale invariance



Great fleas have lesser fleas
Upon their backs to bite 'em.
And lesser fleas have lesser still.
And so *ad infinitum*

Johnathan Swift

Fig. 16.3 Scale invariance.

Collective Behavior



Spreading of Diseases

DESCRIPTION

Many degrees of freedom

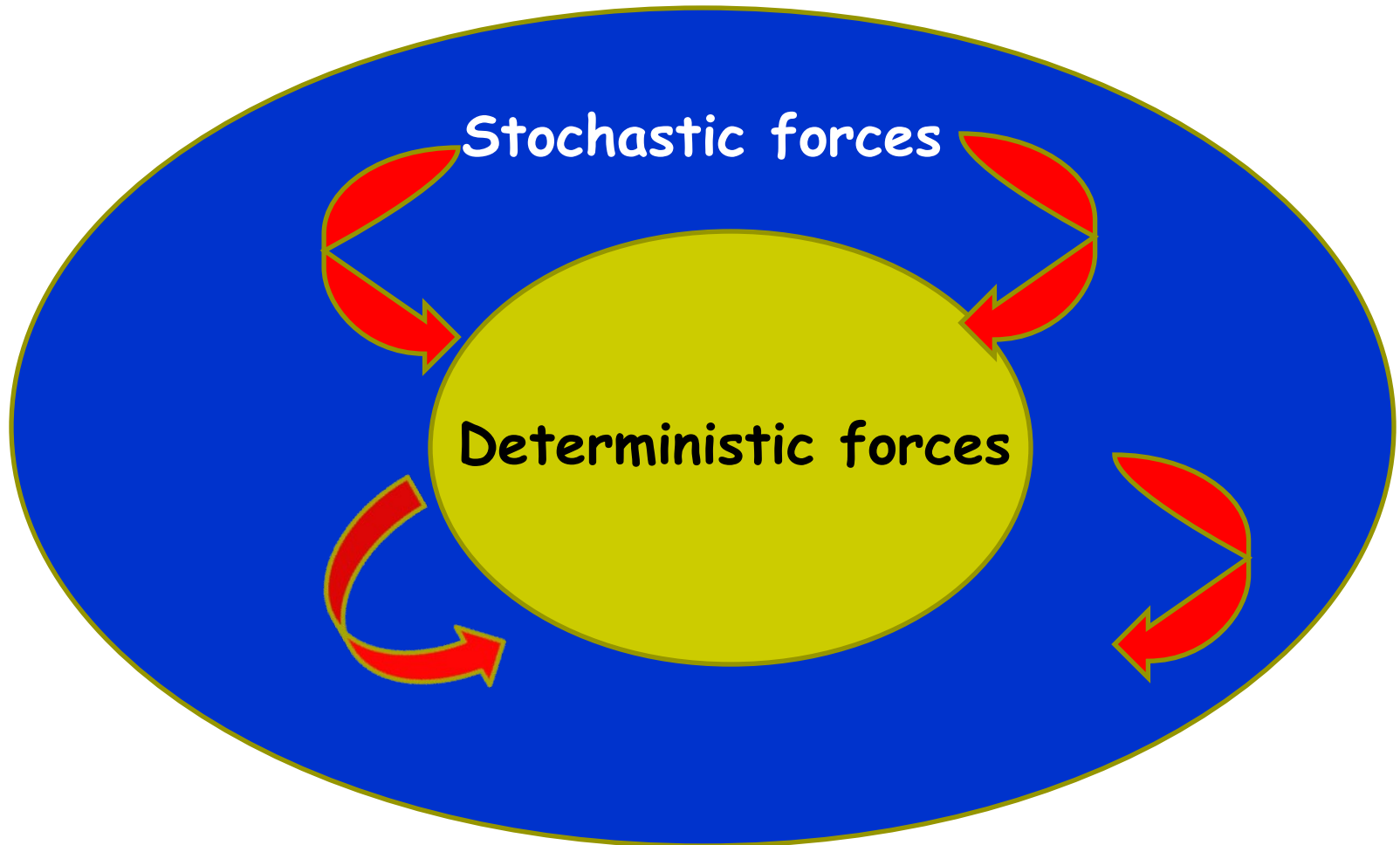
Not all of them specified by deterministic forces

Different time scales

Stochastic descriptions

- Langevin equation
- Fokker-Planck equation
- **Master equation**
- **Simulations (MC,MD)**

DESCRIPTION



MODEL

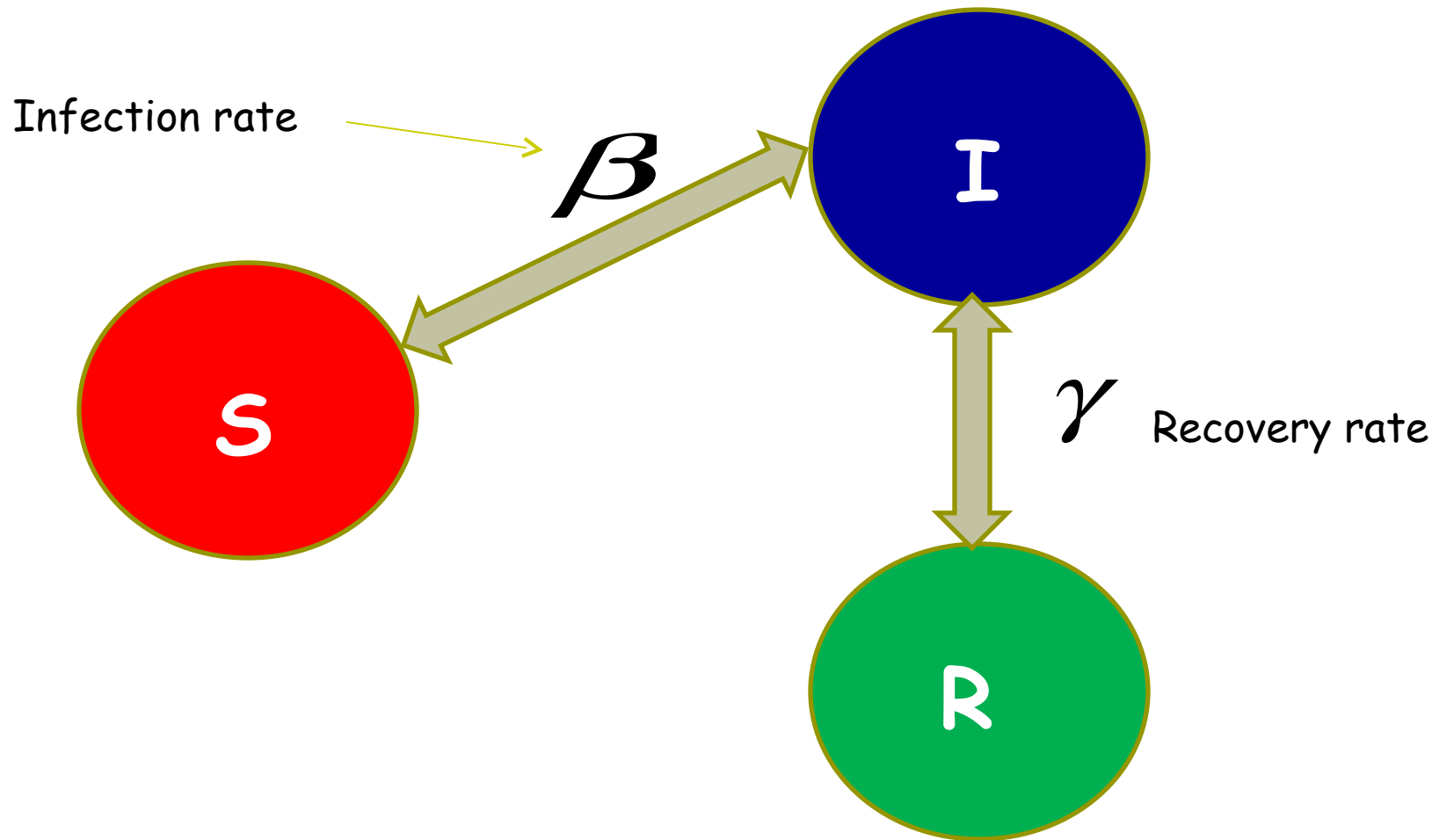
Three state model

S-susceptibles: healthy individuals can catch diseases

I-infectives: can transmit disease provided contacts

R-recovered: immunized or removed out

THREE STATE MODEL



MFA

deterministic equations

$S \rightarrow I \rightarrow R$

$$S(t) + I(t) + R(t) = N$$

$$\frac{\partial S(t)}{\partial t} = -\beta S(t) I(t)$$

$$\frac{\partial I(t)}{\partial t} = \beta S(t) I(t) - \gamma I(t)$$

$$\frac{\partial R(t)}{\partial t} = \gamma I(t)$$

STOCHASTIC MODEL

- ❖ Deterministic evolution neglect fluctuations
- ❖ Individuals are represented by nodes which are either of the three states S I R
- ❖ Contact defined by links between nodes, minimal connectivity
- ❖ Each node, state variable occupation number $n = 0, 1$:
- ❖ If node i is in the S state: $n_S(i, t) = 1$ otherwise $n = 0$
- ❖ Mapping master equation in a quantum formulation

Master-Equation

- PROCESS-DYNAMICS
- CHANGE OF A CERTAIN CONFIGURATION
- MARKOV PROCESS, NO MEMORY
- UNIFORM IN TIME
- CONTINUOUS IN TIME

$$\partial_t P(S_1, \dots, S_N, t) = \sum_i w(-S_i \rightarrow S_i) P(-S_i) - w(S_i \rightarrow -S_i) P(S_i)$$

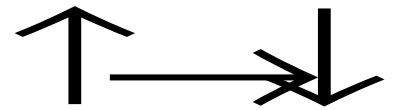
→ **GAIN**

LOST

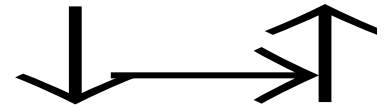
$$\partial_t P(S) = L P(S)$$

Simple Example

RATE : $\lambda(T, \dots)$



RATE : $\nu(T, \dots)$



Description

$p_{\uparrow}(t)$, $p_{\downarrow}(t)$ prob

Markov process

$$p_{\uparrow}(t + \Delta t) = p_{\downarrow}(t) \gamma \Delta t + p_{\uparrow}(t) (1 - \lambda \Delta t)$$

$$\frac{dp_{\uparrow}}{dt} = \gamma p_{\downarrow} - \lambda p_{\uparrow}$$

$$\frac{dp_{\downarrow}}{dt} = -\gamma p_{\downarrow} + \lambda p_{\uparrow}$$

General.

$$\frac{d}{dt} p_a = L_{ab} p_b$$

$$L_{ab} = \begin{pmatrix} -\lambda & \gamma \\ \lambda & -\gamma \end{pmatrix}$$

Quantum Language

OTHER LANGUAGE: CREATION AND ANNIHILATION OPERATOR

$$|0\rangle \equiv |\uparrow\rangle \quad |1\rangle \equiv |\downarrow\rangle$$

$$d^\dagger |0\rangle = |1\rangle$$

$$d |1\rangle = |0\rangle$$

$$n = d^\dagger d = 0, 1$$

MAPPING TO SPINS :

$$S = 1 - 2d^\dagger d = 1 - 2n$$

$$S |1\rangle = -1 |1\rangle \quad S |0\rangle = +1 |0\rangle$$

Mapping

$$\frac{\partial}{\partial t} p(\vec{n}, t) = L p(\vec{n}, t)$$

$$|F(t)\rangle = \sum_n p(\vec{n}, t) |\vec{n}\rangle \implies \frac{\partial}{\partial t} |F(t)\rangle = \hat{L} |F(t)\rangle$$

$$\langle \vec{n}' | \hat{L} | \vec{n} \rangle = L_{n'n}$$

$$d^+ |0\rangle = |1\rangle \quad \lambda\text{-process} \quad \uparrow \rightarrow \downarrow$$

$$d |1\rangle = |0\rangle \quad \gamma\text{-process} \quad \downarrow \rightarrow \uparrow$$

$$\hat{L}_f = \sum_i [\gamma(1 - d_i^+) d_i + \lambda(1 - d_i) d_i^+]$$

SIR Model

Master Eq. for the probability density



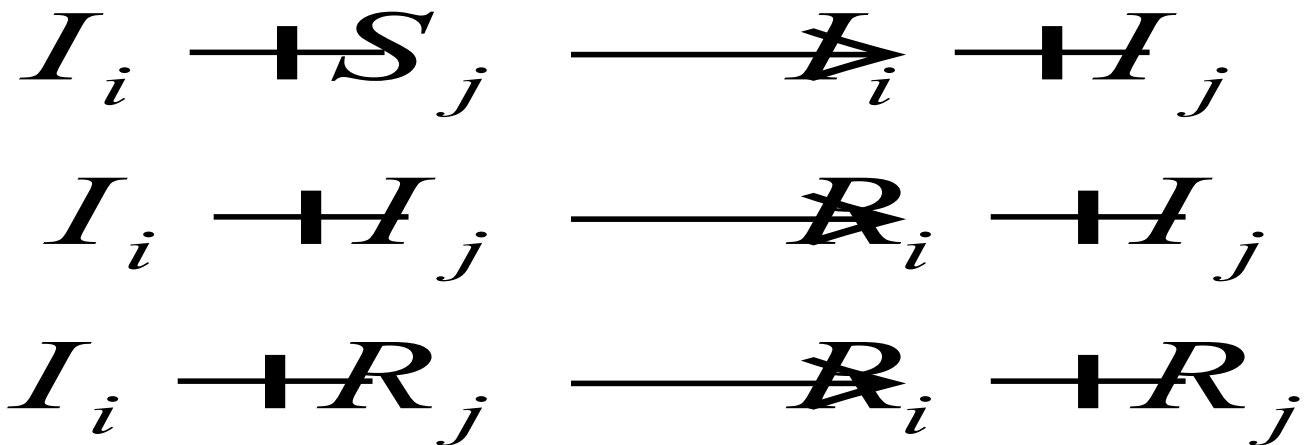
$$p(S, I, R, t) \equiv p(\vec{n}, t); \quad \frac{\partial p(\vec{n}, t)}{\partial t} = L p(\vec{n}, t)$$

$$n_i = \langle \cdot, \cdot \rangle \quad i = S, I, R \quad \left| F(t) \right\rangle = \sum_{\vec{n}} p(\vec{n}, t) \left| \vec{n} \right\rangle; \quad \frac{\partial}{\partial t} \left| F \right\rangle = - \hat{H} \left| F \right\rangle \quad \hat{H} = -\hat{L}$$

$$n_S(i, t) = \langle \cdot \rangle$$

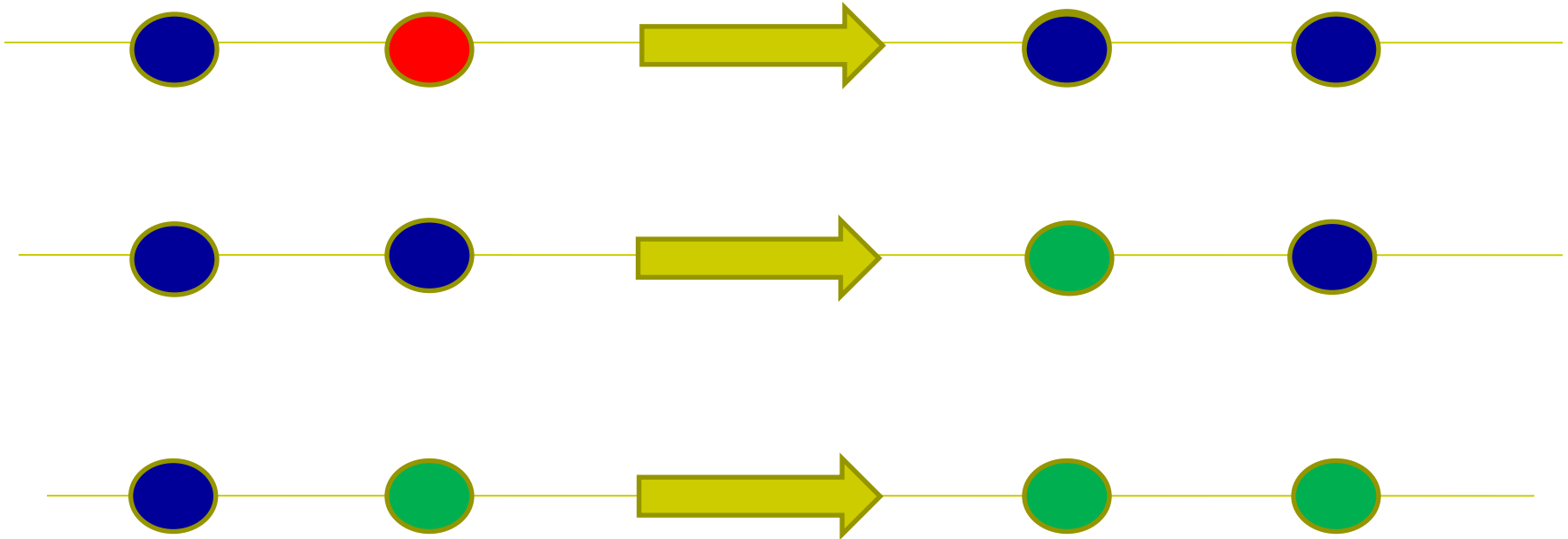
DYNAMICS

S → I → R



DYNAMICS

S → I → R



EVOLUTION OPERATOR

$$-\hat{H} = \beta \sum_i \left[b_{i+1}^+ a_{i+1} + b_{i-1}^+ a_{i-1} - A_{i+1} (1 - B_{i+1}) - A_{i-1} (1 - B_{i-1}) \right] B_i + \gamma \sum_i (b_i - B_i)$$

Creation of **I** and annihilation of **S**

provided lattice site i occupied with **I**

$$A_l = a_i^+ a_i \Rightarrow 0, 1$$

Particle number operator **S** state
 B Particle number of **I** state

Evolution Equation

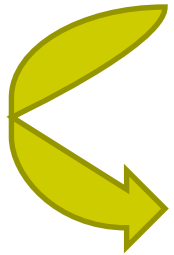
$$\frac{d}{dt} \langle A_r \rangle = -\beta \left[\langle A_r B_{r-1} \rangle + \langle A_r B_{r+1} \rangle \right]$$

$$\frac{d}{dt} \langle B_r \rangle = \beta \left[\langle A_r B_{r-1} \rangle + \langle A_r B_{r+1} \rangle \right] - \gamma \langle B_r \rangle$$

Hierarchy of Equations, MFA decoupling: $\langle AB \rangle = \langle A \rangle \langle B \rangle$

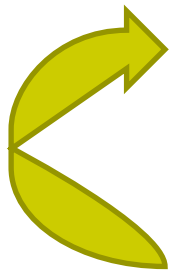
N-Point Cluster Functions

S surrounded by **I** at the edges



$$H_r(n) = \langle A_r A_{r+1} \dots A_{r+n-1} B_{r+n} \rangle$$

$$= \langle \text{red circle} \text{ red circle} \dots \text{red circle} \text{ blue circle} \rangle$$



$$G_r(n) = \langle \text{blue circle} \text{ red circle} \dots \text{red circle} \text{ blue circle} \rangle$$

$$G_r(n) = \langle B_{r-1} A_r \dots A_{r+n-1} B_{r+n} \rangle$$

$$A_i = a_i^+ a_i \rightarrow \text{red circle with } S \text{ inside} \quad B = b_i^+ b_i \rightarrow \text{blue circle with } I \text{ inside}$$


N-Point Cluster Functions

$$H_r(n) = \langle A_r A_{r+1} \dots A_{r+n-1} B_{r+n} \rangle$$

$$G_r(n) = \langle B_{r-1} A_r \dots A_{r+n-1} B_{r+n} \rangle$$

Coupled Equations (Use $A B = 0$)

$$\frac{d}{dt} H(n) = -[\gamma + \beta] H(n) + \beta [H(n+1) - G(n)]$$


$$\frac{d}{dt} G(n) = -2[\gamma + \beta] G(n) + 2\beta G(n+1)$$

N-Point Cluster Functions

$$\frac{d}{dt} H(n) = -[\gamma + \beta] H(n) + \beta [H(n+1) - G(n)]$$

$$\frac{d}{dt} G(n) = -\gamma [\gamma + \beta] G(n) + \gamma \beta G(n+1)$$



$$n_S(i, t) = \langle A_i(t) \rangle$$

$$n_I(i, t) = \langle B_i(t) \rangle$$

$$\frac{d}{dt} n_S(t) = -2\beta H(1, t)$$

$$\frac{d}{dt} n_I(t) = 2\beta H(1, t) - \gamma n_I(t)$$

N-Point Cluster Functions

$$H(n) = \langle [n \otimes A] [1 \otimes B] \rangle$$

$$G(n) = \langle [1 \otimes B] [n \otimes A] [1 \otimes B] \rangle$$

Increase by infection with β

$$\langle [n+1] \otimes A \ 1 \otimes B \rangle \rightarrow \langle [n] \otimes A \ 1 \otimes B \rangle$$

Recovery of infected individual $I \rightarrow R$ with rate γ

Infection of S by the adjacent I with β

$$\frac{d}{dt} H(n) = -[\gamma + \beta] H(n) + \beta [H(n+1) - G(n)]$$

Infection of S at the left border

N-Point Cluster Functions

$$H(n) = \langle [n \otimes A] [1 \otimes B] \rangle$$

$$G(n) = \langle [1 \otimes B] [n \otimes A] [1 \otimes B] \rangle$$

$$\frac{d}{dt} G(n) = -2[\gamma + \beta] G(n) + 2\beta G(n+1)$$

Coupled equations

$$\frac{d}{dt} H(n) = -[\gamma + \beta] H(n) + \beta [H(n+1) - G(n)]$$

$$\frac{d}{dt} G(n) = -2[\gamma + \beta] G(n) + 2\beta G(n+1)$$

- Competition between growth and reduction processes
- Coupling between H and G



Nontrivial steady state

Solution for arbitrary initial condition

Solution uncorrelated random initial distribution

$$n_S(t) = n_S(0) - 2\beta\tau n_S(0)n_I(0) \left\{ \left[\left(1 - \exp\left(-\frac{t}{\tau}\right) \right) \right] \right. \\ \left. \times [1 - \beta\tau n_I(0)] + \left[1 - \exp\left(-\frac{2t}{\tau}\right) \right] \frac{\beta\tau n_I(0)}{2} \right\}.$$

$$n_I(t) = n_I(0)[\exp(-\gamma t) + 2\beta\tau n_S(0)f(t)],$$

$$f(t) = \frac{\exp(-t/\tau) - \exp(-\gamma t)}{\gamma\tau - 1} [1 - \beta n_I(0)\tau] \\ + \frac{\beta n_I(0)\tau}{\gamma\tau - 2} [\exp(-2t/\tau) - \exp(-\gamma t)].$$

Solution

Stationary solution

$$\frac{n_S^*}{n_S(0)} = \left[\frac{\gamma + \beta[1 - n_S(0) - n_I(0)]}{\gamma + \beta[1 - n_S(0)]} \right]^2,$$

Relaxation time

$$\tau = \frac{1}{\gamma + \beta[1 - n_S(0)]}.$$

- Due to fluctuations, isolated regions of **susceptible** individuals evolve
- Finite stationary distribution of the **S** type even for large population size
- Relaxation time and stationary distribution depends on initial conditions
- Highly nonergodic, far from equilibrium situation
- $n_S(t)$ strictly monotonically decreasing (no **S** generated)
- $n_I(t)$ exhibit maximum, stationary state $n_I=0$

MFA

$$\frac{\partial S(t)}{\partial t} = -\beta S(t) I(t)$$

$$\frac{\partial I(t)}{\partial t} = \beta I(t) \left(S(t) - \frac{\gamma}{\beta} \right)$$

$$\frac{\partial R(t)}{\partial t} = \gamma I(t)$$

increase or decrease

Max. if $S(0) > \gamma/\beta$

$S(t') = \gamma/\beta$

$I(t') = I_{\max}$

t' different

$$\frac{dR}{dt} = -\frac{dU(R)}{dR}$$

$$U(R) = -\frac{S_0 \gamma^2}{\beta} \exp\left(-\frac{\beta}{\gamma} R\right) + \frac{\gamma R}{2} (R - 2N)$$

Overdamped motion in a potential

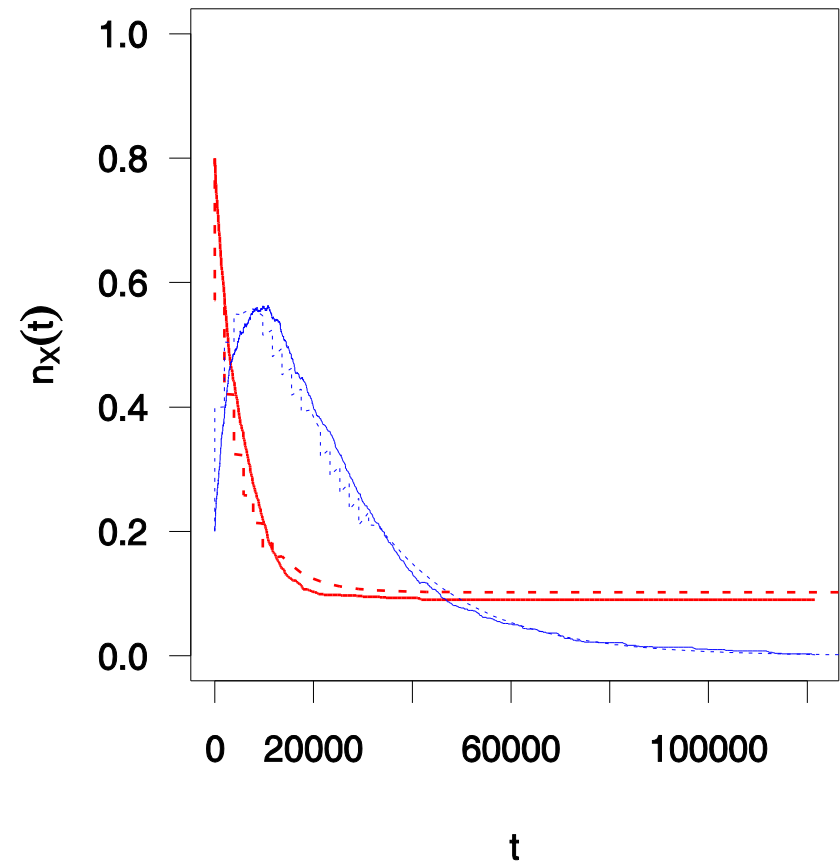
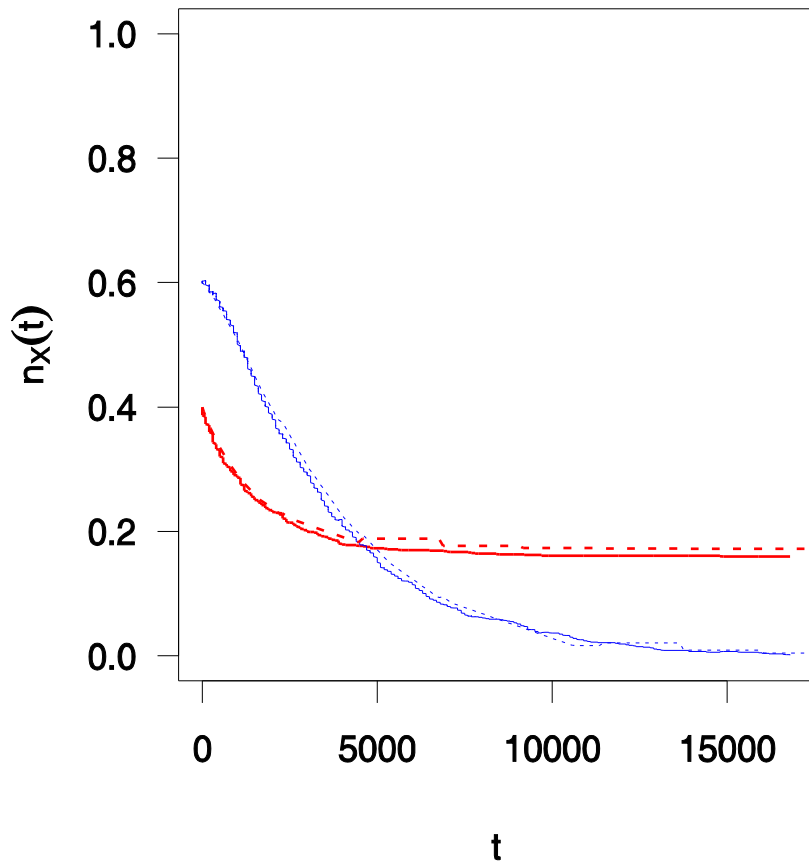
$$S(t \rightarrow \infty) \rightarrow S^* \approx S_0 \exp\left(-\frac{\beta}{\gamma} N\right)$$

MONTE CARLO SIMULATION

- Initially, each site is occupied independently and randomly by S with $n_s(0)$, I with $n_I(0) = 1 - n_s(0)$
- Update: Choose arbitrary site j
- If j occupied by an I, then I decays to R with $\frac{\gamma}{\beta + \gamma}$
- If not decay, then with prob $\frac{1}{2}$ adjacent site (left or right)
- Is this site occupied by S, then decay to I with $\frac{2\beta}{2\beta + \gamma}$

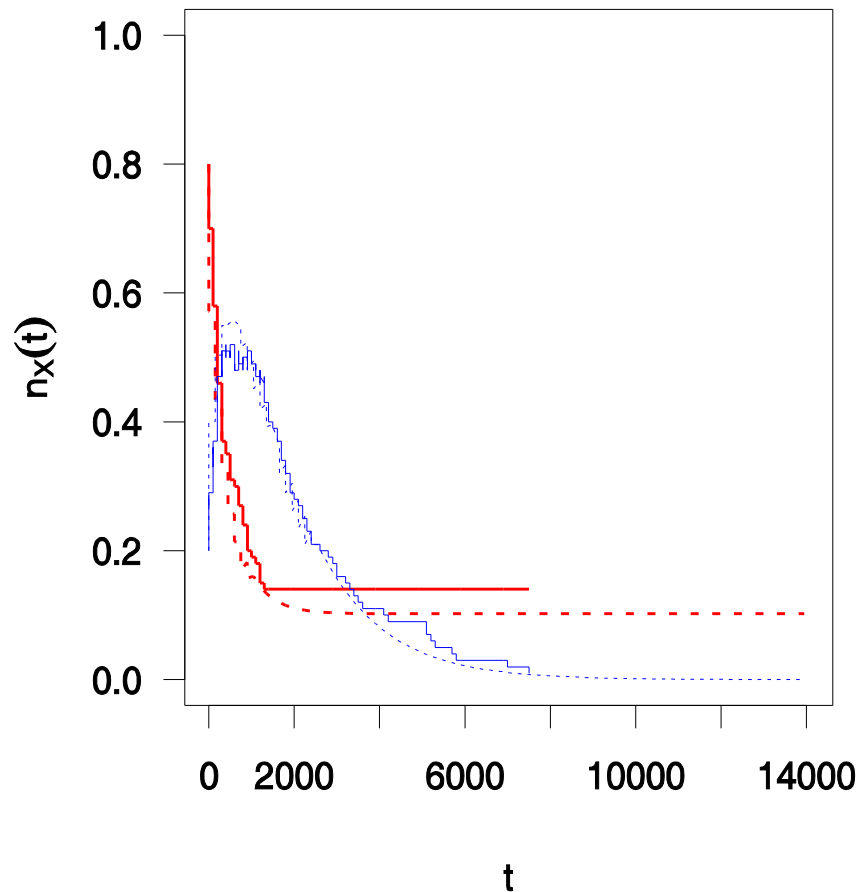
Numerical Results

Solid line: simulation; fixed $N = 10^3$

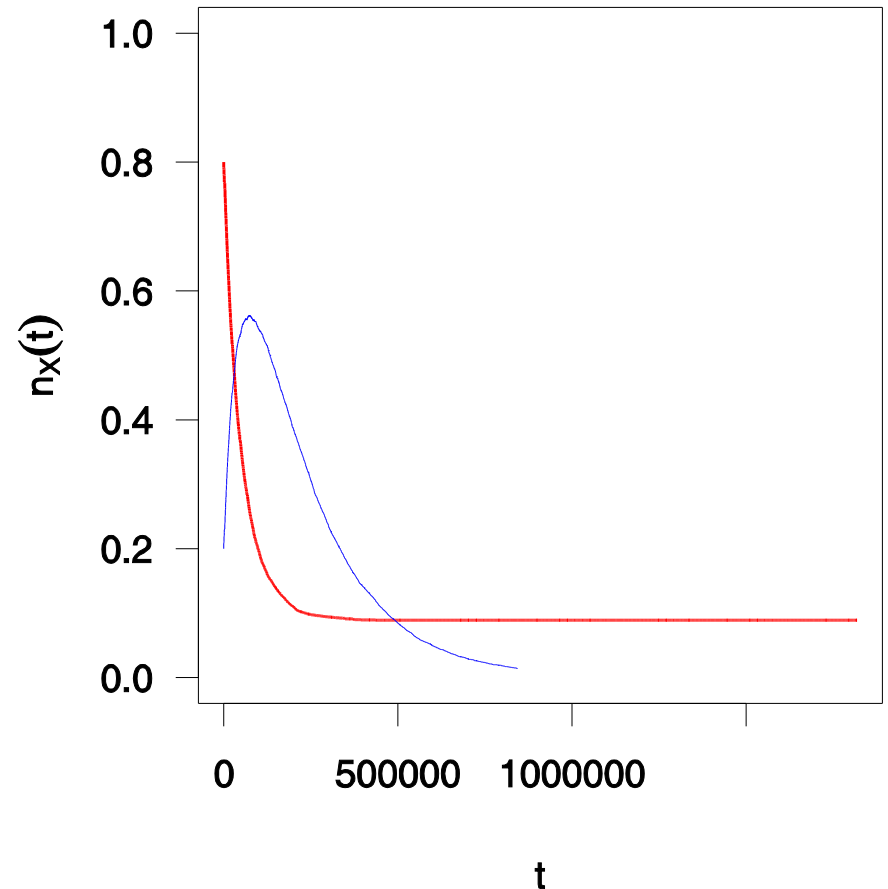


Numerical Results

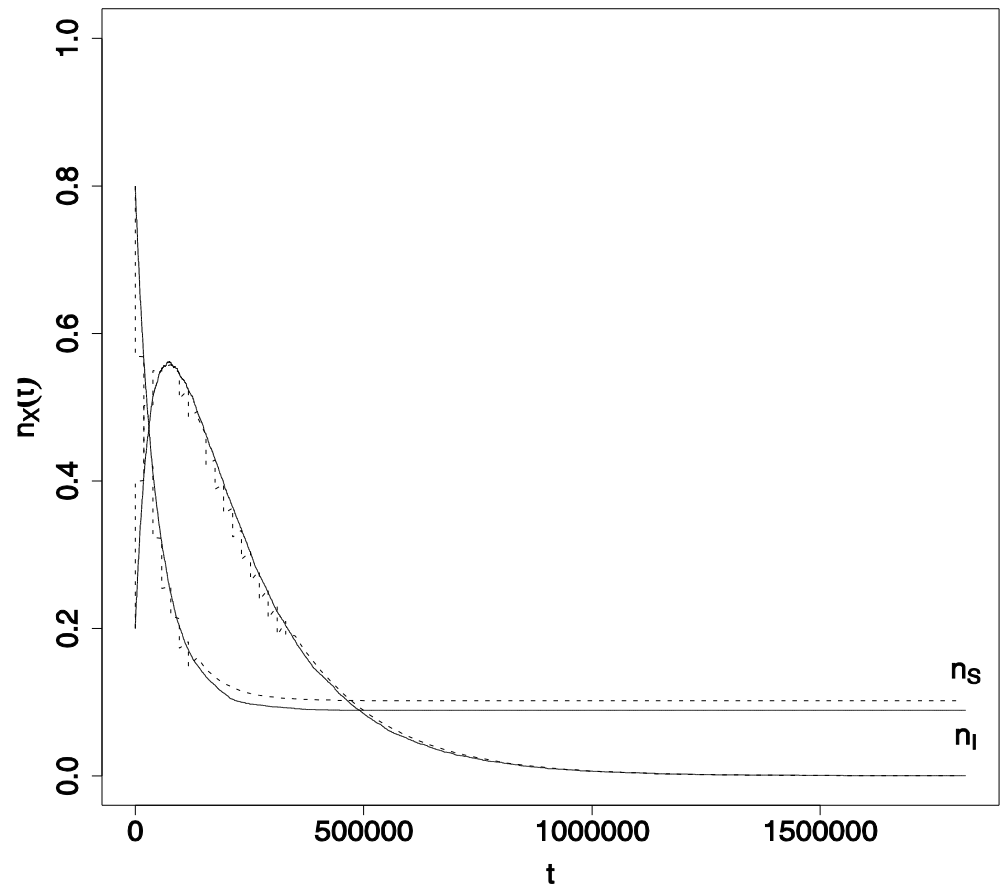
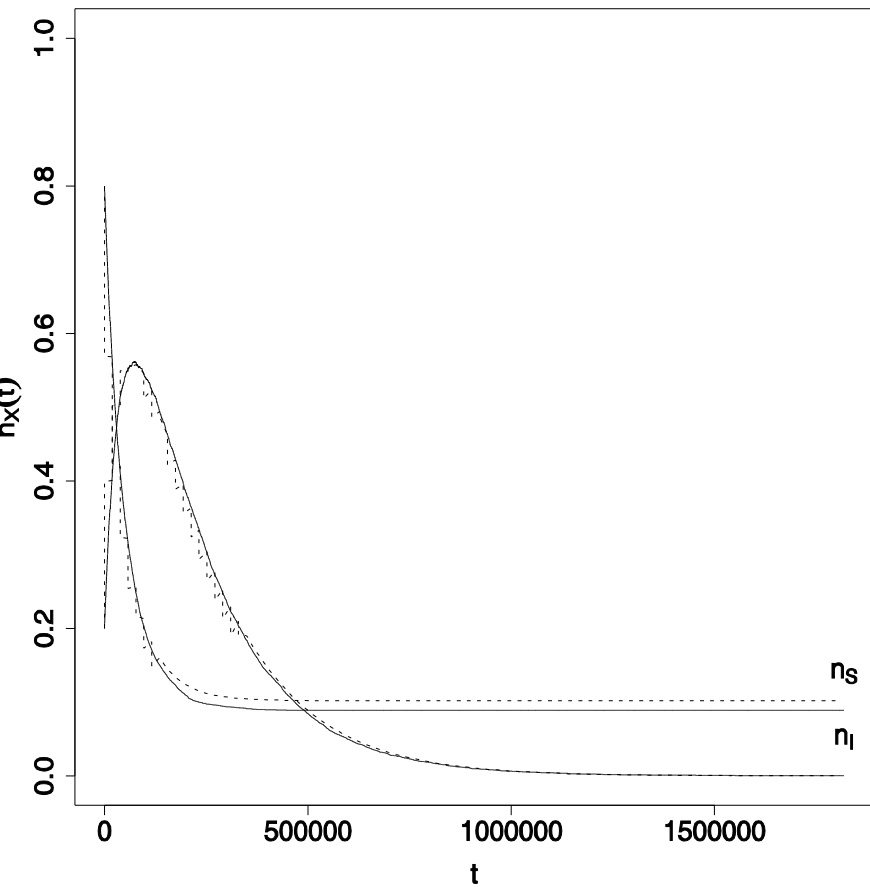
Different N: small N



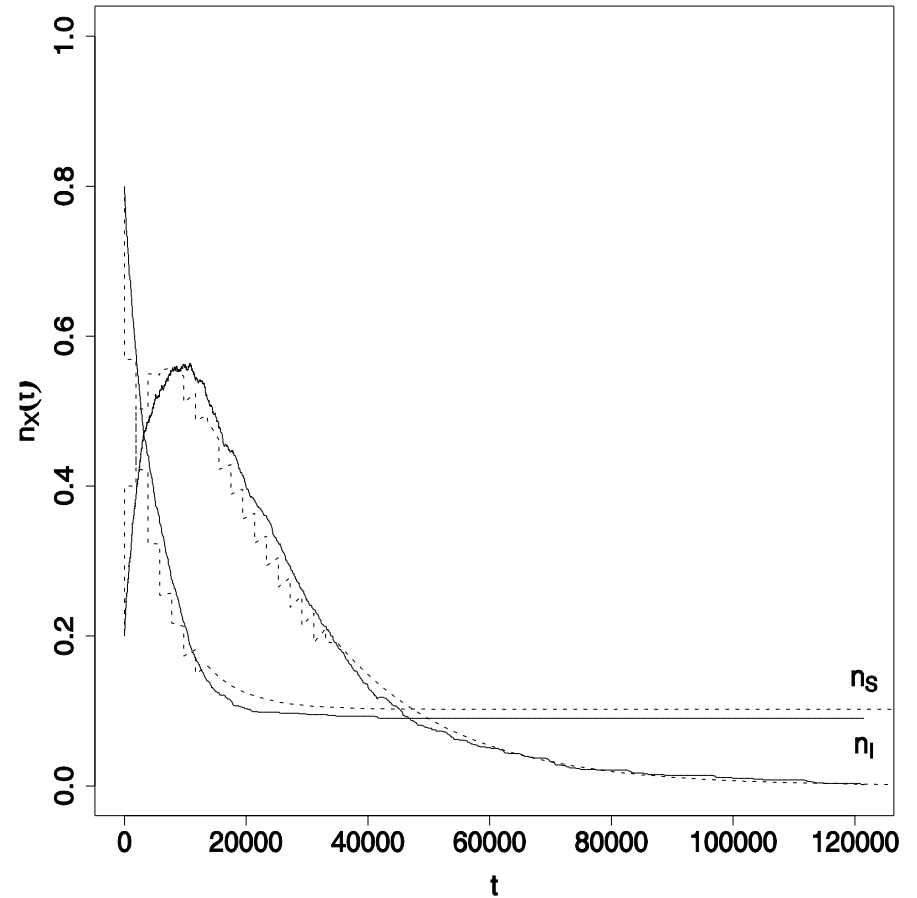
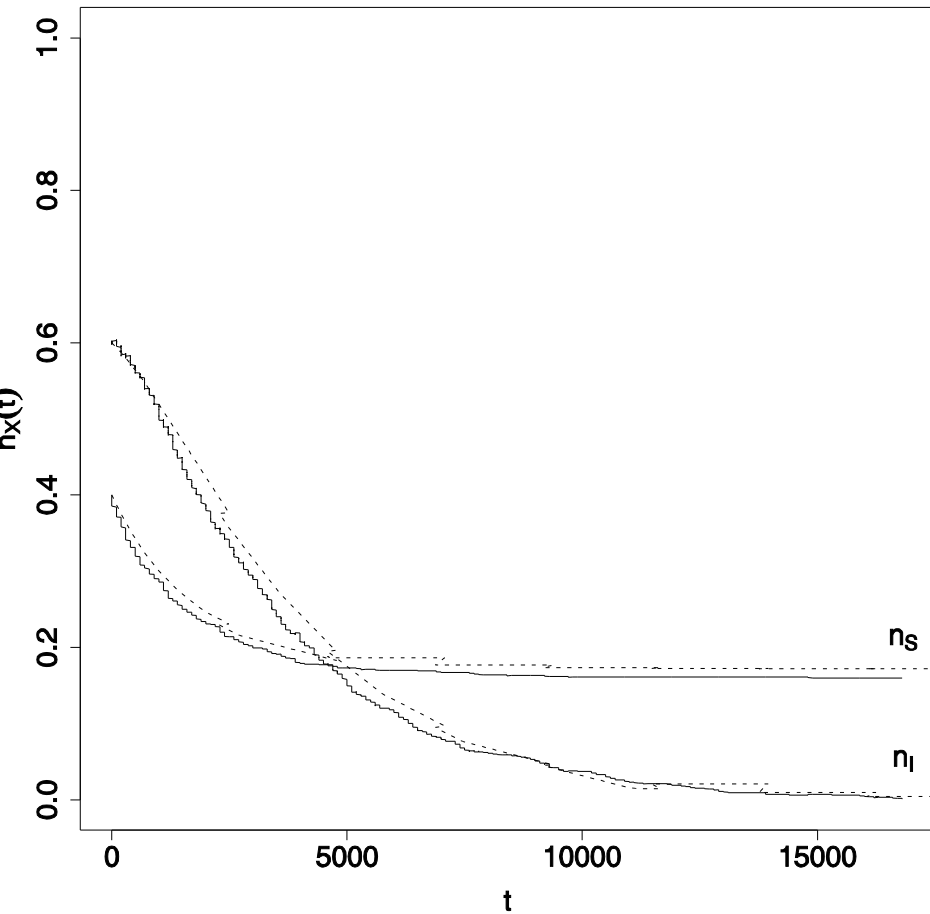
large N ($= 10^4$) fluctuations irrelevant



MONTE CARLO SIMULATION



MONTE CARLO SIMULATION

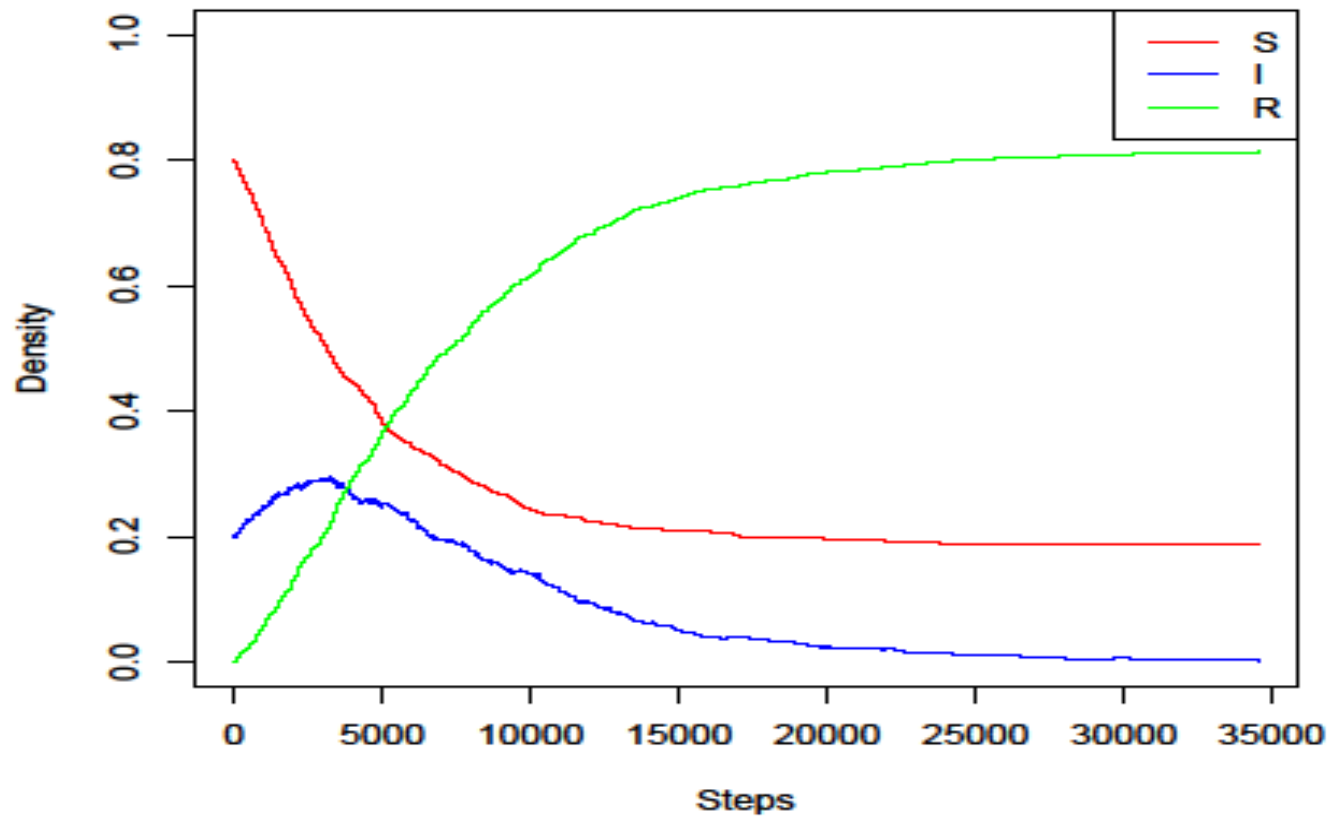




Different parameters (1-D)

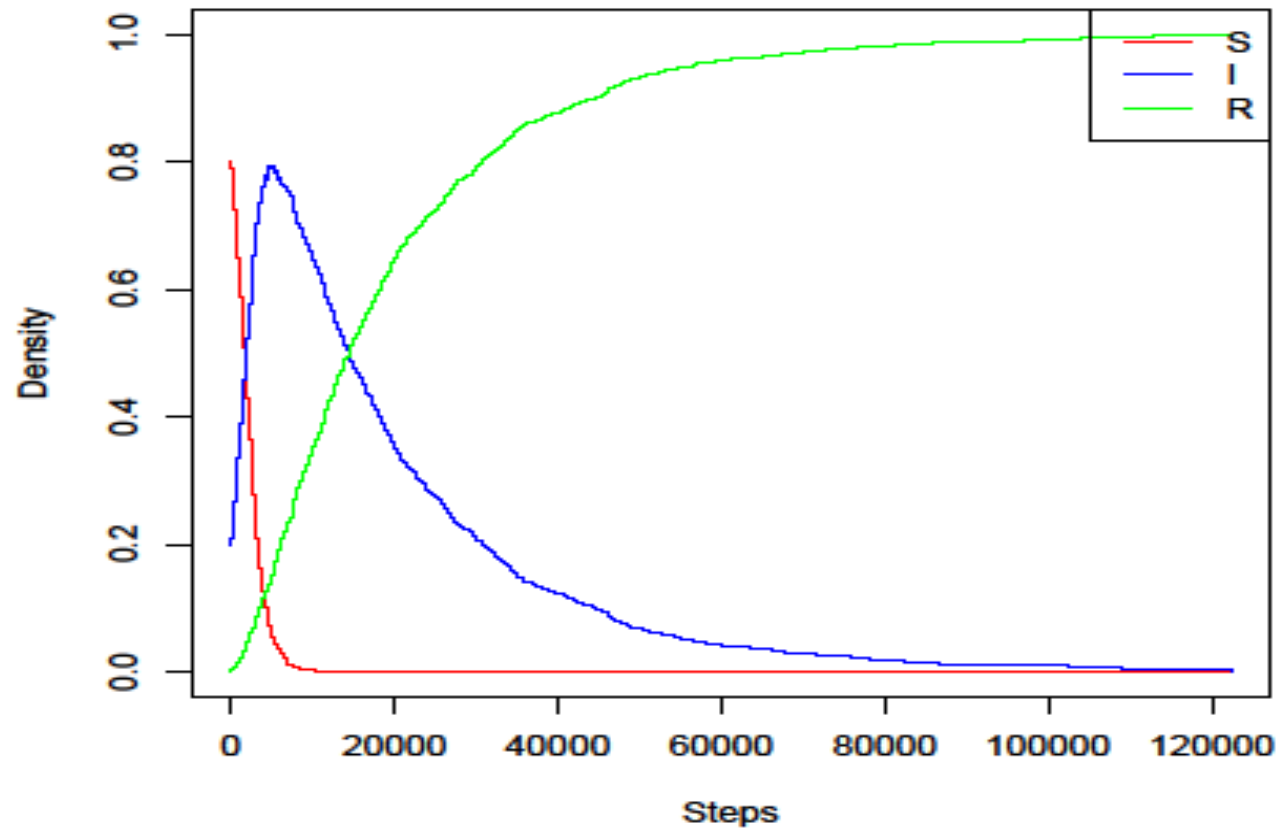
3-D Simulation

$N = 1000$, $\beta = 0.7$, $\gamma = 0.5$, $B_0 = 200$



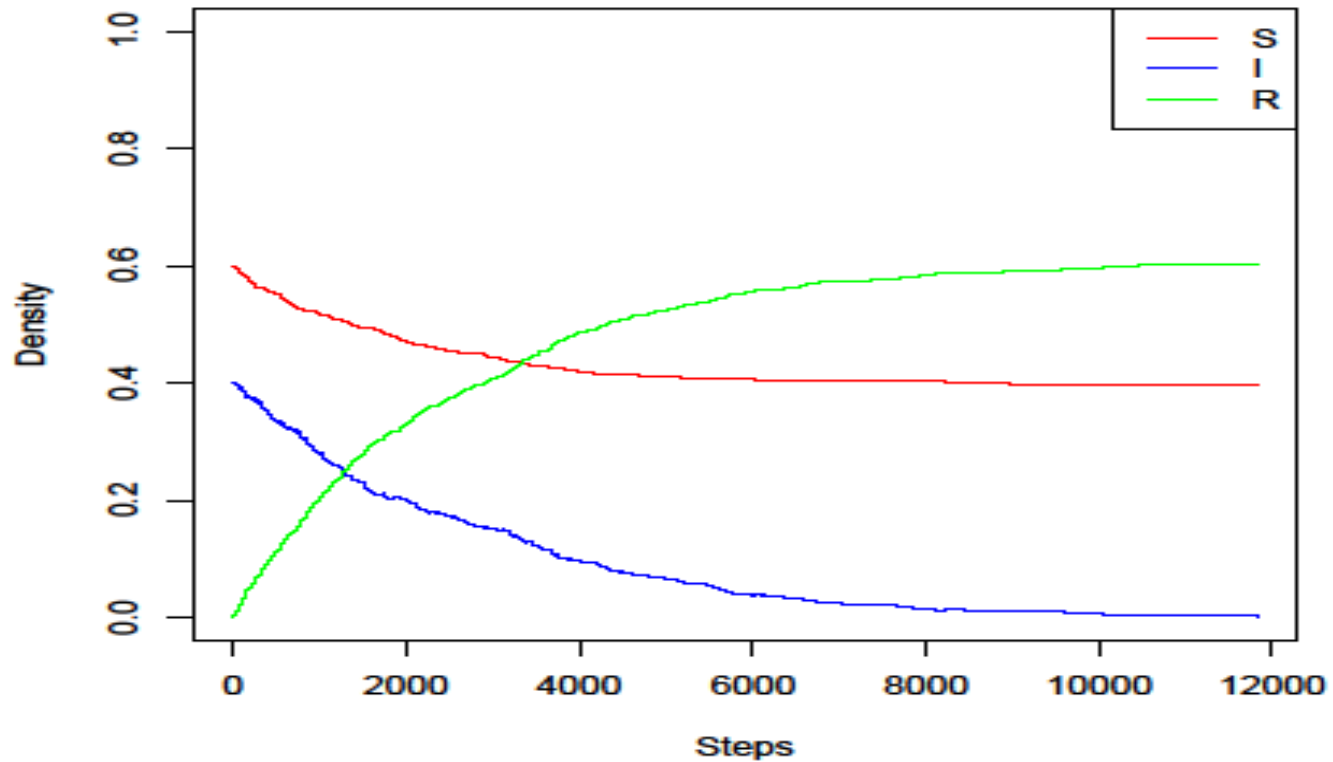
3-D Simulation

$N = 1000$, $\beta = 0.9$, $\gamma = 0.1$, $B_0 = 200$



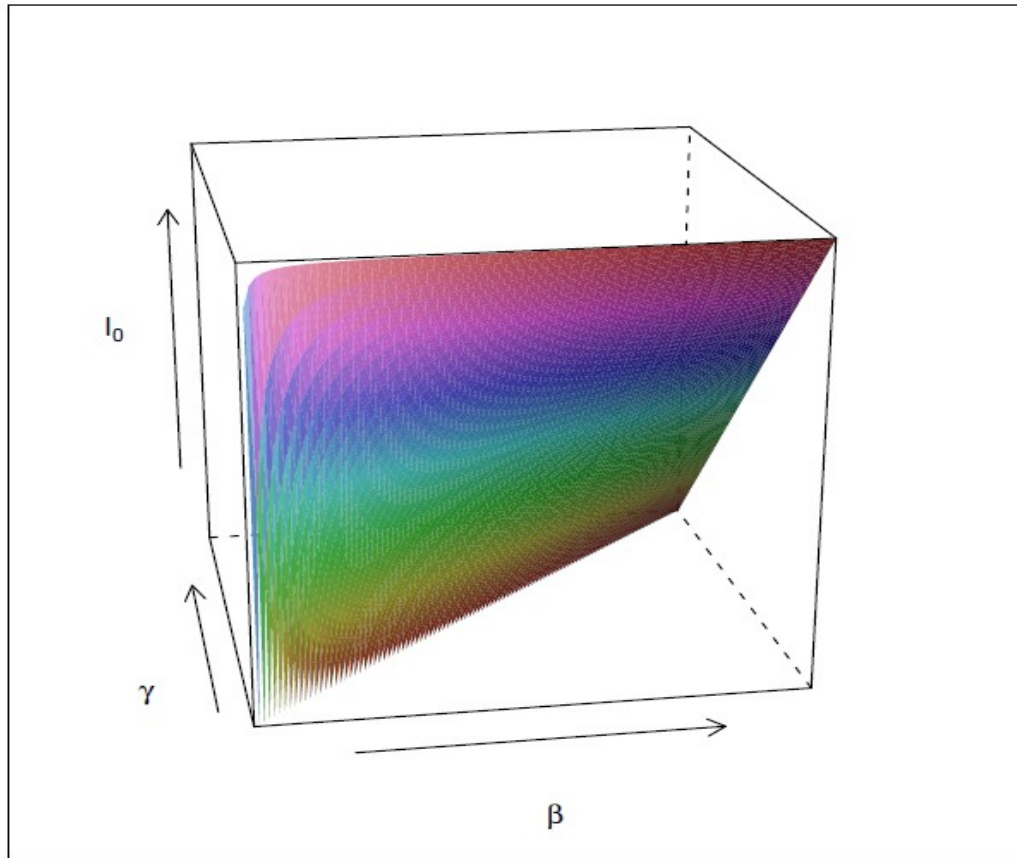
3-D Simulation

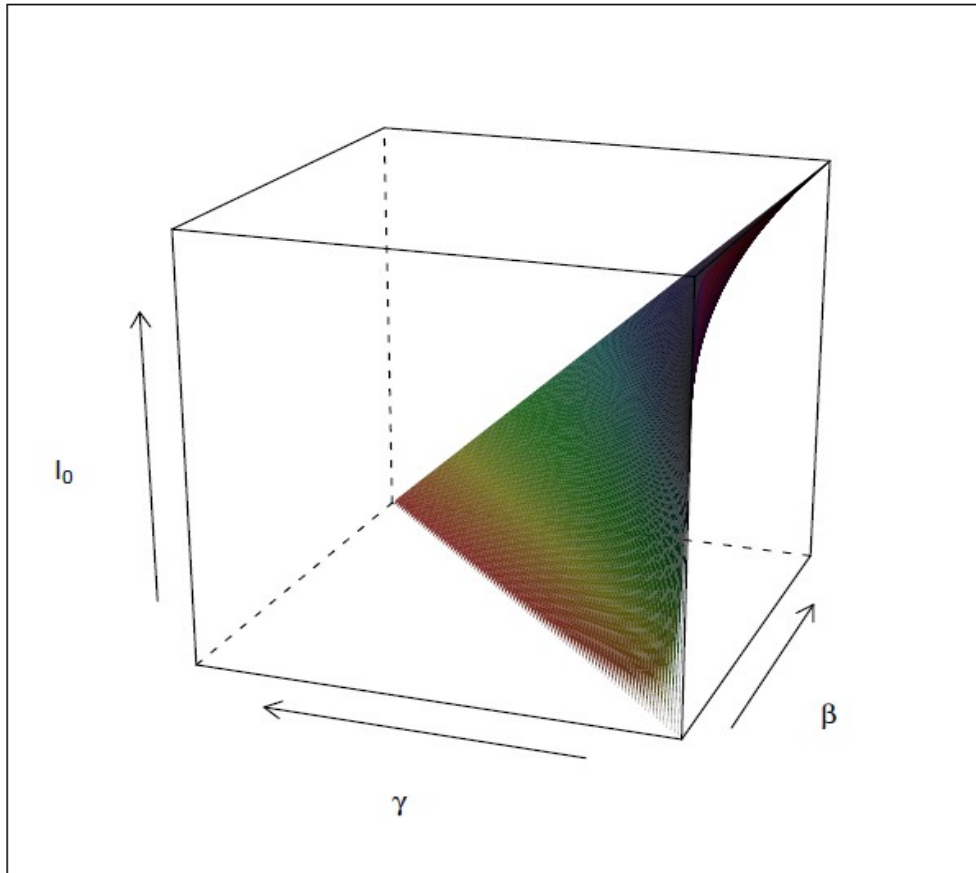
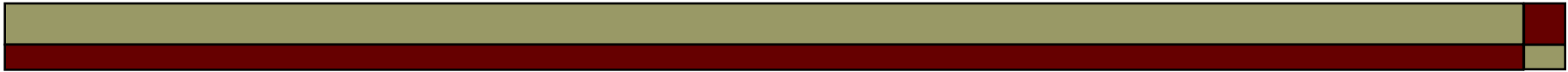
$N = 1000$, $\beta = 0.2$, $\gamma = 0.5$, $B_0 = 400$

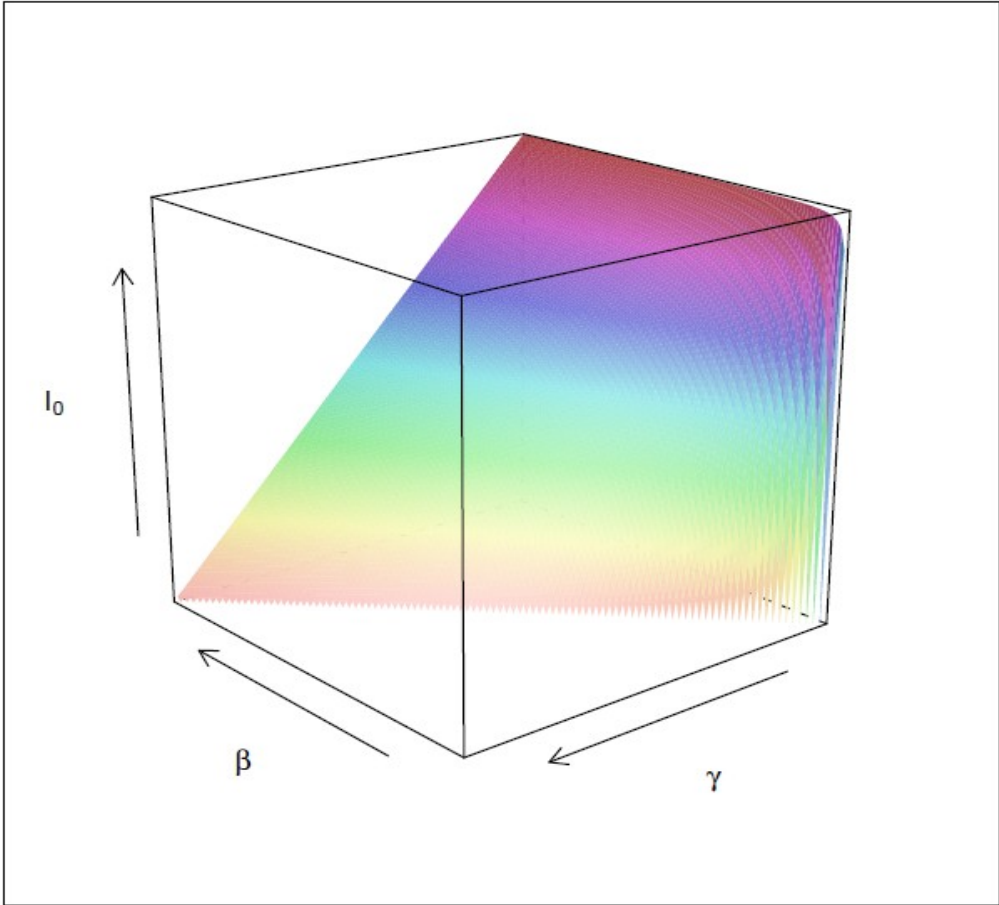
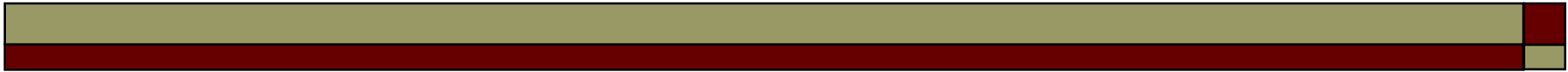


Phase diagram

Initial configuration
Infection rate B
Recovery rate γ







CONCLUSION

- SIMPLIFIED SIR MODEL & STOCHASTIC DYNAMICS
- INCOMPLETE CONTACT & FLUCTUATIONS
- MASTER EQUATION & QUANTUM FORMUL.
- PAULI-OPERATORS & COMMUTE/ANTICOM

EXACT SOLUTION

- ✓ DEPENDING ON PARAMETERS \rightarrow DIFFERENT BEHAVIOR
- ✓ INFECTIOUS \rightarrow MAXIMUM \leftrightarrow CONTINUOUS DECAY
- ✓ SIGNIFICANT DIFFERENCE TO MFA
- ✓ COMPARISON \leftrightarrow MONTE CARLO SIMULATION
- ✓ FLUCTUATIONS IRRELEVANT \leftrightarrow LARGE POPULATION

PROBLEMS

- Hopping of individuals (in particular S)
- Cluster size of I ($\langle BBBBBBB \rangle$)
- Empty sites
- Delay time for infection $S \rightarrow I$
- After waiting time $R \rightarrow S$ (S generated)
- Immunization
- Higher Dimensions (field theoretical approach)
- Critical Dimension
- Scale free networks (higher connectivity)

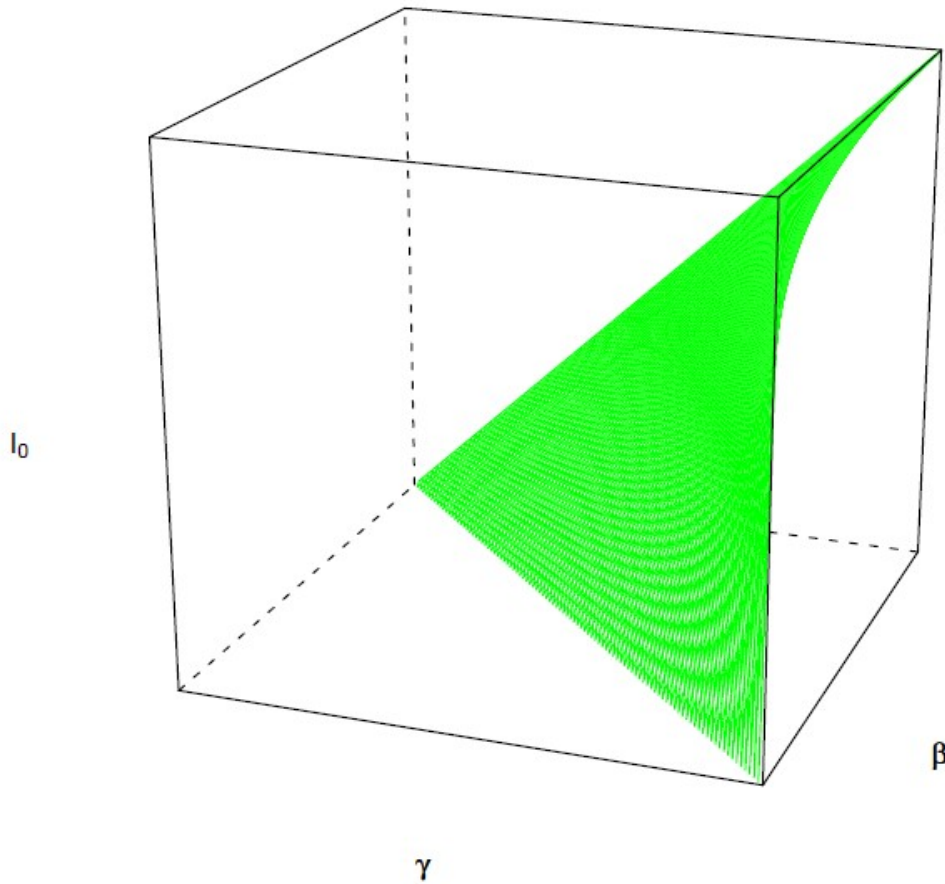


THE END

Key Words

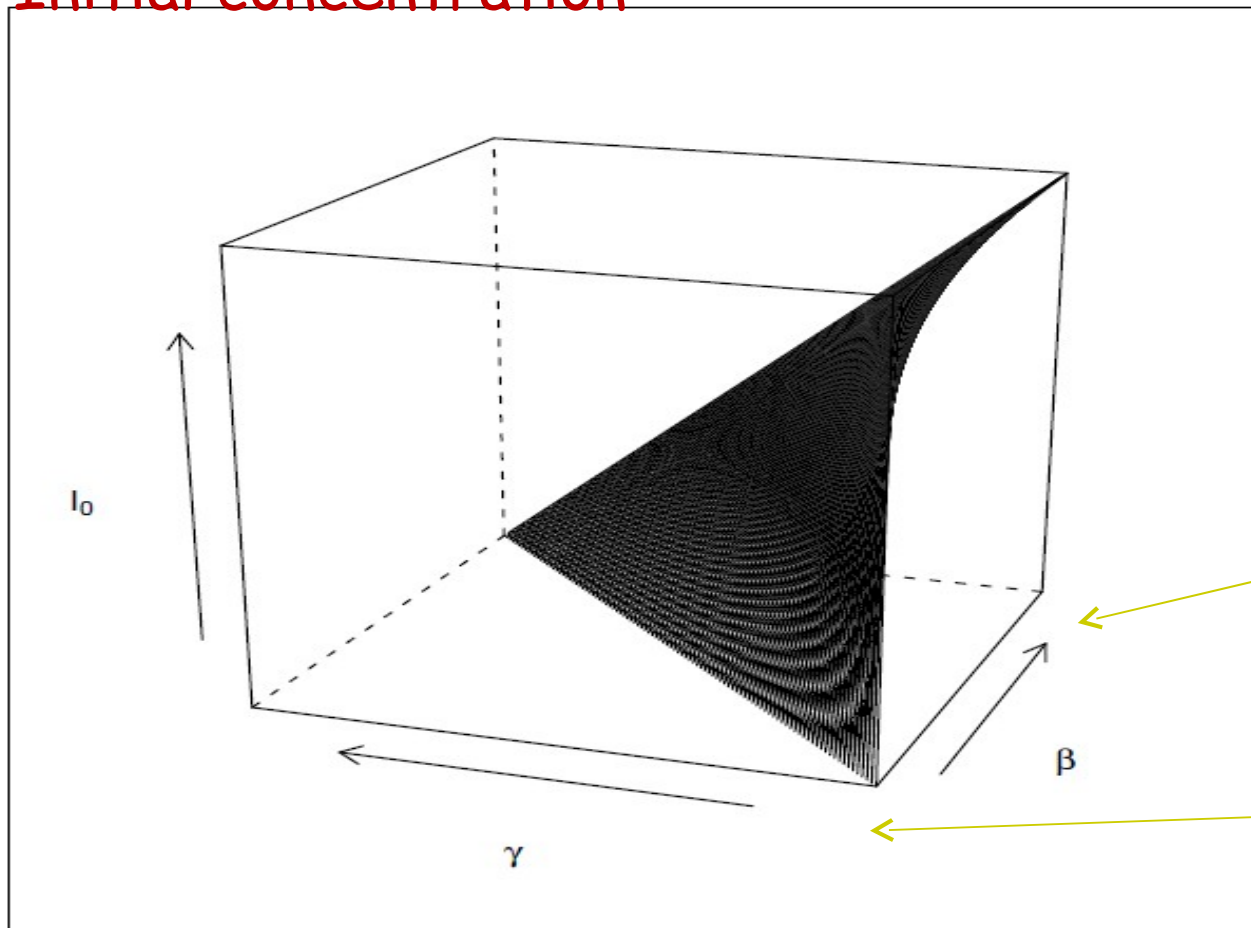
- Uncorrelated random initial conditions: analytical solution
- **I** population may increase initially before decaying to zero.
- Due to fluctuations, isolated regions of **susceptible** individuals evolve
- Finite stationary distribution of the **S** type even for large population size.
- Simulations, Mean-field, Master Equation

„Phase diagram“



Phase diagram

Initial concentration



Infection rate

Recovery rate