

Stochastic Epidemic Modelling

STOR-i Summer Research Project

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Introduction



Coronavirus Map. Image Credit: VK Studio/Shutterstock.com

Aims

- ▶ Create a simulation of the general stochastic SIR epidemic and its extensions.
- ▶ Work on epidemic inference using Bayesian statistics
- ▶ Design a model for a ward to include spatial structure

SIR Model

The population is divided into three categories:

- **Susceptible (S)** – Individuals who are not yet infected.
- **Infectious (I)** – Individuals who are currently infected with the pathogen and can transmit it to others.
- **Recovered (R)** – Individuals previously infected with the pathogen, but no longer transmit it, and are now immune.



SIR Model Equations

- ▶ **Infection Rate (β):** Determines how easily the disease spreads.

$$\mathbb{P}(\text{Infection}) = 1 - \exp(-\beta I(t-1))$$

- ▶ **Recovery Rate (γ):** Determines recovery time of infected individuals.

$$\mathbb{P}(\text{Recovery}) = 1 - \exp(-\gamma)$$

- ▶ **Basic Reproduction Number (R_0)**

$$R_0 = \frac{\beta}{\gamma}$$

SIR Plot

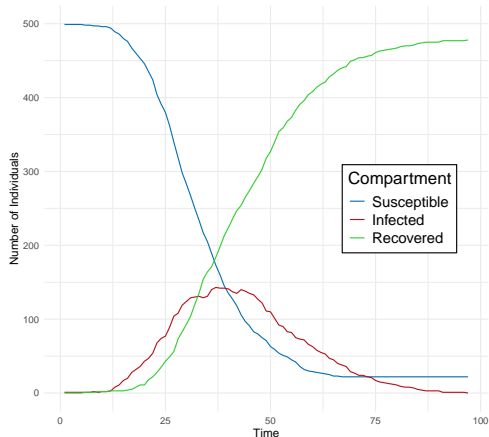
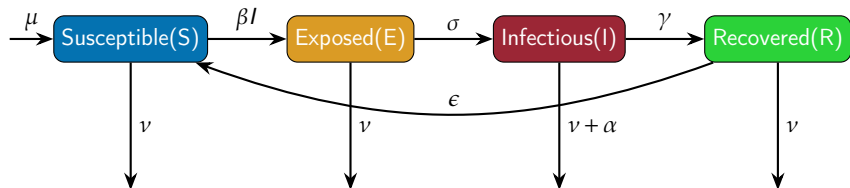


Figure: Stochastic SIR Model with Individual Tracking

Extensions of the SIR Model - SEIRS

The SEIRS model extends the SIR model and exhibits periodicity due to waning immunity over time. It also introduces a latency period where an individual carries the pathogen but is not infectious.



SEIRS Plot with a Varying Population

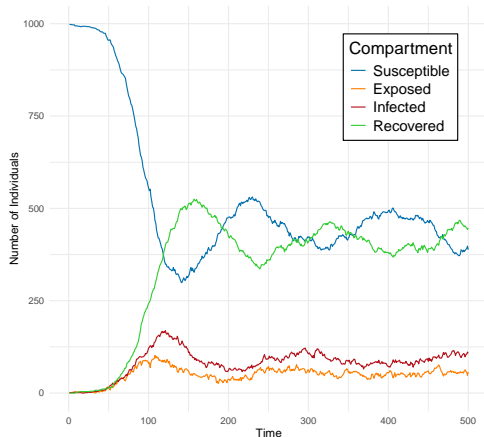


Figure: Stochastic SEIRS Model with Birth and Death rates

Inference with Random-Walk Metropolis MCMC

If given infection time data, Markov chain Monte Carlo (MCMC) methods can be applied to infer the rates of infection and recovery.

Inference with Random-Walk Metropolis MCMC

To estimate β and γ , the Random-Walk Metropolis algorithm is applied:

- ▶ **Proposal:** New β and γ values proposed from normal distributions centered at current values, with standard deviations λ_β and λ_γ .
- ▶ **Acceptance:** A proposed value is accepted based on:

$$\log(\alpha_\theta) = \log \Pi(\theta_{\text{proposed}} | X) - \log \Pi(\theta_{\text{current}} | X)$$

where θ is β or γ , and $\Pi(\theta | X)$ is the posterior distribution of θ .

- ▶ **Adaptive Tuning:** Every 100 iterations, λ_β and λ_γ are adjusted to maintain a 30% acceptance rate.

Trace Plots and Histograms

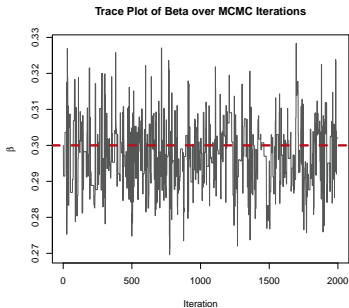


Figure: Beta Trace Plot

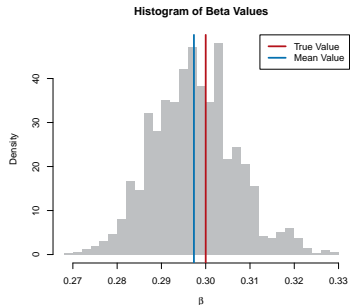


Figure: Beta Histogram

Trace Plots and Histograms

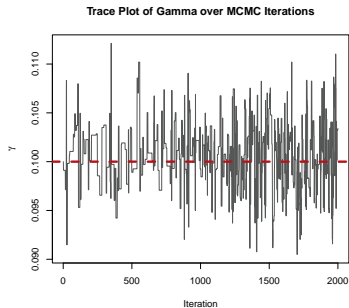


Figure: Gamma Trace Plot

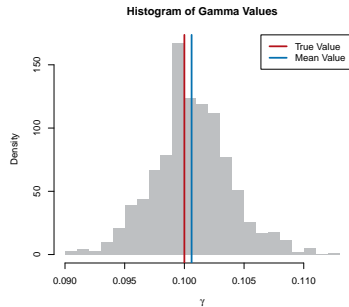


Figure: Gamma Histogram

Ward Layout

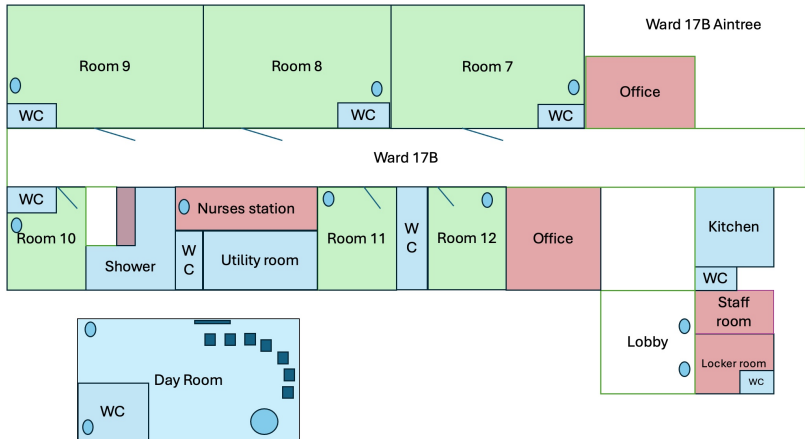


Figure: Layout of Aintree ward 17B

Calculation of Infectious Pressure with Distance

$$\lambda(t)_j = \beta_1 \underbrace{\sum_{i=1}^N I(t)_i f(D_{ij})}_{\text{Whole Ward}} + \beta_2 \underbrace{\sum_{i=1}^N r_{ij} I_i(t)}_{\text{Individual Room}}$$

- ▶ $\lambda(t)$ = infection rate (pressure)
- ▶ β = infection parameter
- ▶ $I(t)$ = vector of infection status
- ▶ $f(D)$ = distance function
- ▶ r = indicator if in the same room

Application to Aintree ward 17B

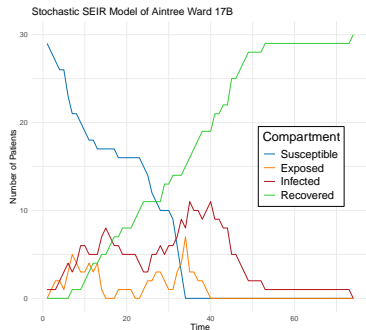


Figure: SEIR Model of Aintree ward 17B with random room allocation, assuming 30 patients.



Figure: Time of infection against Euclidean distance from room of the initially infected individual.

Model Summary

- ▶ Discrete Time
- ▶ Stochastic
- ▶ Markov Chain
- ▶ Agent-Based
- ▶ Spatial Structure

Conclusions

- ▶ **Stochastic epidemic models** provide insight into how infectious diseases spread in various settings.
- ▶ The **Random-Walk Metropolis MCMC algorithm** enables estimation of model parameters, such as infection and recovery rates, from observed data.
- ▶ The **application to Aintree Ward 17B** demonstrates the utility of these models in hospital settings, aiding in understanding and controlling outbreaks.

References

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- [3] Gelman, A., Gilks, W. R., and Roberts, G. O. (1997). *Weak convergence and optimal scaling of random walk Metropolis algorithms*
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