

## Stochastic Modelling and Random Processes

### Example sheet 5

#### 1. Contact process

[14]

Consider the CP  $(\eta_t : t \geq 0)$  on the complete graph  $\Lambda = \{1, \dots, L\}$  (i.e.  $q(i, j) = \lambda$  for all  $i \neq j$ ) with state space  $S = \{0, 1\}^L$  and transition rates

$$c(\eta, \eta^i) = \eta(i) + \lambda(1 - \eta(i)) \sum_{j \neq i} \eta(j),$$

and generator given by  $(\mathcal{L}f)(\eta) = \sum_{i \in \Lambda} c(\eta, \eta^i)(f(\eta^i) - f(\eta))$ .

Recall that  $\eta, \eta^i \in S$  are connected states such that the state of individual  $i$  is flipped:

$$\eta^i(k) = \begin{cases} 1 - \eta(k) & , k = i \\ \eta(k) & , k \neq i \end{cases}$$

- (a) Let  $N(\eta) := \sum_{i \in \Lambda} \eta(i) \in \{0, \dots, L\}$  be the number of infected individuals in configuration  $\eta$ . For any function  $f : \{0, \dots, L\} \rightarrow \mathbb{R}$  show that we can write for the composed function  $f \circ N : S \rightarrow \mathbb{R}$

$$(\mathcal{L}f \circ N)(\eta) = \lambda(L - N)N[f(N + 1) - f(N)] + N[f(N - 1) - f(N)]$$

for all  $\eta \in S$ , where we use the simplified notation  $N = N(\eta)$  on the right-hand side.

Hint: Use  $N(\eta^i) = N(\eta) \pm 1$  if  $\eta(i) = 0, 1$ , respectively, and  $(1 - \eta(i))\eta(i) = 0$ .

Convince yourself that this implies that  $(N_t : t \geq 0)$  with  $N_t := N(\eta_t)$  is a Markov chain on  $\{0, \dots, L\}$  and write down its generator  $\mathcal{L}f(n)$ .

- (b) Is the process  $(N_t : t \geq 0)$  irreducible, does it have absorbing states? Give all stationary distributions. Is the process ergodic?
- (c) Assume that  $\mathbb{E}(N_t^k) = \mathbb{E}(N_t)^k$  for all  $k \geq 1$ . This is called a **mean-field assumption**, meaning basically that we replace the random variable  $N_t$  by its expected value. Use this assumption and the usual evolution equation for functions of Markov chains to derive the **mean-field rate equation** for  $\rho(t) := \mathbb{E}(N_t)/L$ ,

$$\frac{d}{dt} \rho(t) = h(\rho(t)) := -\rho(t) + L\lambda(1 - \rho(t))\rho(t).$$

- (d) Analyze this equation by finding the stable and unstable stationary points via  $h(\rho^*) = 0$ , and give the limiting behaviour of  $\rho(t)$  as  $t \rightarrow \infty$  depending on the parameter  $\lambda > 0$ . What is the prediction for the stationary density  $\rho^*$  depending on  $\lambda$ ?

## 2. Simulation of Contact Processes

Consider again the contact process  $(\eta_t : t \geq 0)$ , but now with connections only between nearest neighbours, i.e.  $q(i, j) = q(j, i) = \lambda \delta_{j, i+1}$ , and periodic boundary conditions.

The critical infection rate  $\lambda_c$  can be defined such that the infection on the infinite lattice  $\Lambda = \mathbb{Z}$  started from the fully infected lattice dies out for  $\lambda < \lambda_c$ , and survives for  $\lambda > \lambda_c$ . It is known numerically up to several digits, depends on the dimension, and is around 1.65 in our case.

All simulations of the process should be done with initial condition  $\eta_0(i) = 1$  for all  $i \in \Lambda$ . You should use the Gillespie algorithm or the random sequential update from handout 4.

- (a) Simulate the process for  $L = 128, 256, 512, 1024$  and parameters  $\lambda = 1.62, \dots, 1.68$  with 0.01 increments (7 values) with at least 500 realizations each.

- i. For each  $L$ , plot the number of infected individuals  $N_t = \sum_{i \in \Lambda_L} \eta_t(i)$  averaged over realizations as a function of time up to time  $10 \times L$  for all values of  $\lambda$  as above in a single double-logarithmic plot. Use the curvature of the plots to estimate  $\lambda_c(L)$ .
- ii. Plot your estimates of  $\lambda_c(L)$  with error bars  $\pm 0.01$  against  $1/L$ . Extrapolate to  $1/L \rightarrow 0$  to get an estimate of  $\lambda_c = \lambda_c(\infty)$  with a reasonable error bar.

This approach is called **finite size scaling**, in order to correct for systematic **finite size effects** which influence the critical value.

- (b) Let  $T$  be the hitting time of the absorbing state  $\eta = \mathbf{0}$ , i.e. the lifetime of the infection. Measure the lifetime of the infection for  $\lambda = 1$  and  $\lambda = 1.8$  by running the process until extinction of the epidemic.

For  $\lambda = 1 < \lambda_c$  we expect  $T \propto C \log L$  + small fluctuations for some  $C > 0$ .

- i. Use large system sizes e.g.  $L = 128, 256, 512, 1024$  (or larger), confirm that  $\mathbb{E}(T)$  scales like  $\log L$  and determine  $C$  by averaging at least 200 realizations of  $T$  for each  $L$ .
  - ii. Then shift your data  $T_i$  for each  $L$  by  $T_i - \mathbb{E}(T)$  and plot the ‘empirical tail’ of the distribution of the shifted data, comparing to the **Gumbel distribution** (all in one plot with log-scale on the y-axis). Look up the Gumbel distribution on Wikipedia, with mean 0 only one parameter needs fitting. Discuss why this could be a good model for the noise here (check google for **extreme value statistics**)
- (c) For  $\lambda = 1.8 > \lambda_c$  we expect  $T \sim \text{Exp}(1/\mu)$  to be an exponential random variable with mean  $\mu = \mathbb{E}(T) \propto e^{CL}$  for some  $C > 0$ .
- i. Use \*small\* system sizes e.g.  $L = 8, 10, 12, 14$  (see how far you can go), and confirm that  $\mathbb{E}(T)$  scales like  $e^{CL}$ . Determine  $C$  by averaging at least 200 realizations of  $T$ .
  - ii. Then rescale your data  $T_i$  for each  $L$  by  $T_i/\mathbb{E}(T)$  and plot the ‘empirical tail’ of the distribution of the rescaled data, comparing to the theoretical tail  $e^{-t}$  (all in one plot with log-scale on the y-axis).

*Recall: The **empirical tail** of data  $T = (T_1, \dots, T_M)$  is the statistic  $\text{tail}_t(T) = \frac{1}{M} \sum_{i=1}^M \mathbb{1}_{T_i > t}$ . This decays from 1 to 0 as a (random) function of time  $t$ .*