

# SIR epidemics with stochastic infectious periods

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**SIR models** : spread of an epidemic among a closed and homogeneous population, according to the scheme :



- **S** : healthy individuals, but susceptible to be contaminated.
- **I** : infected individuals, who can contaminate the healthy ones (independently of each other).
- **R** : infected individuals whose infection period is over. They take no longer part to the infection process (removed).

# Stochastic infectious periods

Let  $X$  be a continuous time Markov process on  $\mathcal{E} \cup \{\star\}$  where

- $\mathcal{E}$  is a set of transient states,
- the state  $\star$  is absorbing.

When an individual  $i$  gets infected, a process  $X_i$  with the same evolution rules as  $X$  takes off to modulate his infectious period :

- The initial state of  $X_i$  is given by the cdf  $\alpha(\cdot)$  on  $\mathcal{E}$ .
- When  $X_i = x \in \mathcal{E}$ , his instantaneous contamination rate is  $S(t)\beta(x)$ .
- When  $X_i$  is absorbed, the infected individual is removed.

$S(0) = n$ ,  $R(0) = 0$  and  $I(0)$  may be random, as well as the initial state of the initial infection processes.

# Final epidemic outcome

The epidemic terminates at time  $T = \inf\{t \geq 0 \mid I(t) = 0\}$ .

Statistics of interest :

- the final number  $S(T)$  of susceptibles,

- the final severity  $\mathcal{A}(T) = \int_0^T \sum_{i=1}^{I(u)} a(X_i(u)) du = \sum_{i=1}^{R(T)} D_i$

where  $D_i$  is the total severity due to the  $i$ -th infected individual.

Assumptions :

- The processes  $\beta(X)$  and  $a(X)$  are càdlàg,
- $\tau_\star = \inf\{t \geq 0 \mid X(t) = \star\}$  is a.s. finite,
- The different versions  $X_i$  that activate throughout the epidemic are independent of each other.

# Martingales

Let  $\mathcal{F}$  be the filtration generated by the whole epidemic process.

## Theorem

For each  $k \in \{0, 1, 2, \dots, n\}$  and  $\theta \geq 0$ , the process

$$\left\{ \binom{S(t)}{k} \rho(k, \theta)^{S(t)} e^{-\theta A(t)} \prod_{i=1}^{I(t)} q(k, \theta, X_i(t)) \mid t \in \mathbb{R}^+ \right\}$$

is a martingale with respect to  $\mathcal{F}$  when

$$q(k, \theta, x) = \mathbb{E} \left[ e^{-\int_0^{\tau_\star} (\theta a(X(u)) + k\beta(X(u))) du} \mid X(0) = x \right],$$

$$\rho(k, \theta) = 1 - \int_{\mathcal{E}} d\alpha(y) + \int_{\mathcal{E}} q(k, \theta, y) d\alpha(y).$$

# Final epidemic outcome

Applying the optional stopping theorem, for all  $k \in \{0, \dots, n\}$  and  $\theta \geq 0$ ,

$$\mathbb{E} \left[ \binom{S(T)}{k} \rho(k, \theta)^{S(T)} e^{-\theta \mathcal{A}(T)} \right] = \binom{n}{k} \rho(k, \theta)^n \mathbb{E} \left[ \prod_{i=1}^{I(0)} q(k, \theta, X_i(0)) \right].$$

▷ A triangular system to determine the distribution of  $S(T)$  :

$$\sum_{s=k}^n \binom{s}{k} \rho(k, 0)^s \mathbb{P}(S(T) = s) = \binom{n}{k} \rho(k, 0)^n \mathbb{E} \left[ \prod_{i=1}^{I(0)} q(k, 0, X_i(0)) \right].$$

▷ The moments of the final severity. In particular,

$$\mathbb{E}[\mathcal{A}(T)] = \left( n - \mathbb{E}[S(T)] \right) \mathbb{E}[D \mid X(0) \sim \alpha] + \mathbb{E} \left[ \sum_{i=1}^{I(0)} \mathbb{E}[D \mid X_i(0)] \right].$$



# Example

## Classical SIR model :

- Infectious periods of duration  $L \sim F(\cdot)$ ,
- Contamination rate  $\beta S(t)$  per infected individual.
- Probability  $p$  of detection for any newly infected individual.

Here,

- $X$  is a Markov process with transient state space  $\mathcal{E} = \mathbb{R}^+$ ,  
 $X = s$  if the infectious period started  $s$  units of time ago ( $X(0) = 0$ ).
- $\beta(x) = \beta$  and  $a(x) = 1 \quad \forall x \in \mathbb{R}^+$ .

The coefficients  $q(k, \theta) \equiv q(k, \theta, 0)$  are given by

$$q(k, \theta) = \mathbb{E} \left[ e^{-(k\beta + \theta)D} \right] = \hat{F}(k\beta + \theta),$$

and  $\rho(k, \theta) = p + (1 - p)q(k, \theta)$ .

# Example

## Brownian infectious periods :

- $X$  is a Brownian motion with negative drift  $-d$  and variance  $\sigma^2$ ,
- $\beta(x) = \gamma x$  and  $a(x) = ax$  (for fixed  $\gamma, a > 0$ ),
- $X(0) > 0$  and absorption occurs at time  $\tau_\star = \inf\{t > 0 \mid X(t) = 0\}$ .

Here,

$$q(k, \theta, x) = e^{\frac{d}{\sigma^2}x} Ai\left(\frac{\frac{d^2}{\sigma^2} + 2(k\gamma + a\theta)x}{(2\sigma(k\gamma + a\theta))^{\frac{2}{3}}}\right) \left[ Ai\left(\frac{\frac{d^2}{\sigma^2}}{(2\sigma(k\gamma + a\theta))^{\frac{2}{3}}}\right) \right]^{-1}$$

where  $Ai$  is the standard Airy function :

$$Ai(y) = \frac{1}{\pi} \int_0^\infty \cos\left((t^3/3) + yt\right) dt.$$

# Example

Moreover,

$$\mathbb{E} [\tau_{\star} | X(0) = x] = \frac{x}{d} \quad \text{and} \quad \mathbb{E} [D | X(0) = x] = \frac{ax\sigma^2}{2d^2} + \frac{ax^2}{2d}.$$

Illustration :

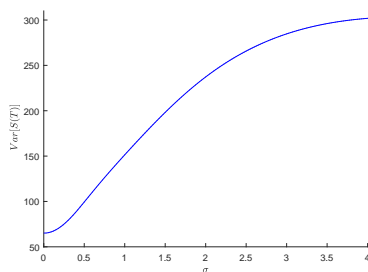
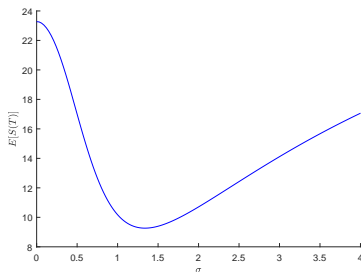


FIGURE –  $S(0) = 37$ ,  $I(0) = 3$ ,  $X(0) = 1$ ,  $p = 0$ ,  $d = 0.8$  and  $\beta = 0.05$ .

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# Markov-modulated fluid flows (MMFF)

An MMFF is a process  $(Y, \varphi) = \{(Y(t), \varphi(t)) \mid t \in \mathbb{R}^+\}$  in which

- $\varphi$  is a Markov process on a finite space  $\mathcal{S}$ , with generator  $Q$ .  
To each phase  $i \in \mathcal{S}$  one associates a rate  $c_i \neq 0$ .
- $Y$  takes its values in  $[0, B]$  and has the following dynamics :

$$\frac{d}{dt} Y(t) = \begin{cases} c_{\varphi(t)} & \text{if } 0 < Y(t) < B, \\ \max(0, c_{\varphi(t)}) & \text{if } Y(t) = 0, \\ \min(0, c_{\varphi(t)}) & \text{if } Y(t) = B. \end{cases}$$

Denoting by  $C$  the diagonal matrix of rates, we write  $Q$  and  $C$  according to the subdivision  $\mathcal{S}_+ = \{i \in \mathcal{S} \mid c_i > 0\}$  and  $\mathcal{S}_- = \{i \in \mathcal{S} \mid c_i < 0\}$  :

$$Q = \begin{bmatrix} Q_{++} & Q_{+-} \\ Q_{-+} & Q_{--} \end{bmatrix}, \quad C = \begin{bmatrix} C_+ & \\ & C_- \end{bmatrix}.$$

# SIR model with fluid infectious periods

We consider the class of SIR model where

- $X = (Y, \varphi)$  is a two-sided MMFF and  $\mathcal{E} = \{(y, i) \mid y \in [0, B], i \in \mathcal{S}\}$ .
- $[0, B]$  is partitioned into  $[v_0, v_1[, [v_1, v_2[, \dots, [v_{N-1}, v_N]$  and

$$\begin{aligned}\beta(x, i) &= \frac{\beta_{I,i}}{n} \quad \text{when } x \in [v_I, v_{I+1}[ , \\ a(x, i) &= a_{I,i} \quad \text{when } x \in [v_I, v_{I+1}[ .\end{aligned}$$

- The initial distribution vector is  $\alpha = [\alpha_0 \ \alpha_1 \ \dots \ \alpha_N]$  where

$$(\alpha_I)_i = \mathbb{P} (Y(0) = v_I, \varphi(0) = i) .$$

- $(Y, \varphi)$  is absorbed at time  $\tau_\star = \inf\{t > 0 \mid Y(t) = 0\}$ .

# Coefficients $q(k, \theta, x)$

let  $\delta_y = \inf\{t > 0 \mid Y(t) = y\}$ . The coefficients  $q(\cdot)$  we need here may be written as

$$q(k, \theta, (v_l, i)) = \mathbb{E} \left[ e^{-\int_0^{\delta_0} (\theta a(Y(t), \varphi(t)) + k \beta(Y(t), \varphi(t))) dt} \mid Y(0) = v_l, \varphi(0) = i \right].$$

To compute them, we start by determining

$$W_{ij}(l) = \mathbb{E} \left[ e^{-\int_0^{\delta_{v_l-1}} (\theta a(Y(t), \varphi(t)) + k \beta(Y(t), \varphi(t))) dt} \mathbb{1}_{\varphi(\delta_{v_l-1})=j} \mid Y(0) = v_l, \varphi(0) = i \right]$$

for  $0 \leq l \leq N$ ,  $i \in \mathcal{S}$  and  $j \in \mathcal{S}_-$ . Then, for  $1 \leq l \leq N$ ,

$$q(k, \theta, (v_l, i)) = [W(l)W_-(l-1)W_-(l-2) \cdots W_-(1)\mathbf{1}]_i.$$

# Mean final severity

The total severity due to a single infected individual is

$$\begin{aligned} D &= \int_0^{\tau_\star} a(Y(u), \varphi(u)) du \\ &= \sum_{r=0}^{N-1} \sum_{j \in \mathcal{S}} a_{r,j} \int_0^{\tau_\star} \mathbb{1}_{Y(u) \in [v_r, v_{r+1}], \varphi(u)=j} du. \end{aligned}$$

To compute  $\mathbb{E} [\mathcal{A}(T)]$ , we need to determine

$$\mathbb{E} [D \mid Y(0) = v_l, \varphi(0) = i] = \sum_{r=0}^{N-1} \sum_{j \in \mathcal{S}} a_{r,j} M_{ij}(l, r),$$

where

$$M_{ij}(l, r) = \mathbb{E} \left[ \int_0^{\tau_\star} \mathbb{1}_{Y(u) \in [v_r, v_{r+1}], \varphi(u)=j} du \mid Y(0) = v_l, \varphi(0) = i \right].$$



# Basic reproduction number

The matrices  $M(l, r)$  also provide us with one of the most frequently used estimators of the virulence of the epidemic :

$R_0$  = average number of infections made by a single infected individual facing infinitely many susceptibles.

When  $S = \infty$ , an infected individual at level  $x \in [v_l, v_{l+1}[$  in phase  $j$  makes contaminations at rate  $\beta_{l,j}$ . So,

$$\begin{aligned} R_0(\alpha) &= \mathbb{E} \left[ \int_0^{\tau_*} \sum_{r=0}^{N-1} \sum_{j \in \mathcal{S}} \beta_{r,j} \mathbb{1}_{Y(u) \in [v_r, v_{r+1}], \varphi(u)=j} du \right] \\ &= \sum_{l=0}^N \sum_{r=0}^{N-1} \sum_{i \in \mathcal{S}} \sum_{j \in \mathcal{S}} (\alpha_l)_i \beta_{r,j} M_{ij}(l, r). \end{aligned}$$

# Matrices $M(l, r)$

For  $x > 0$  define

$$\begin{aligned}
 \Gamma_{ij}(x) &= \mathbb{P}(\delta_0 < \delta_x, \varphi(\delta_0) = j \mid Y(0) = 0, \varphi(0) = i) & (i \in \mathcal{S}_+, j \in \mathcal{S}_-), \\
 \Gamma_{ij}^*(x) &= \mathbb{P}(\delta_x < \delta_0, \varphi(\delta_x) = j \mid Y(0) = x, \varphi(0) = i) & (i \in \mathcal{S}_-, j \in \mathcal{S}_+), \\
 \Lambda_{ij}(x) &= \mathbb{P}(\delta_x < \delta_0, \varphi(\delta_x) = j \mid Y(0) = 0, \varphi(0) = i) & (i, j \in \mathcal{S}_+), \\
 \Lambda_{ij}^*(x) &= \mathbb{P}(\delta_0 < \delta_x, \varphi(\delta_0) = j \mid Y(0) = x, \varphi(0) = i) & (i, j \in \mathcal{S}_-), \\
 A_{ij}(x) &= \mathbb{P}(\varphi(\delta_x) = j \mid Y(0) = x, \varphi(0) = i) & (i \in \mathcal{S}_+, j \in \mathcal{S}_-).
 \end{aligned}$$

$$M_+(l, r) = (I - \Gamma(v_r - v_l)\Gamma^*(v_l))^{-1} \Lambda(v_r - v_l)M_+(r, r) \quad (r > l),$$

$$M_-(l, r) = \Gamma^*(v_l)M_+(l, r)$$

$$M_-(l, r) = (I - \Gamma^*(v_l - v_{r+1})A(v_l))^{-1} \Lambda^*(v_l - v_{r+1})M_-(r+1, r) \quad (r < l-1).$$

$$M_+(l, r) = A(v_l)M_-(l, r)$$

## Proposition ( $r < N - 1$ )

$M_-(r, r) = \Gamma^*(v_r)M_+(r, r)$  and  $M_+(r + 1, r) = A(v_{r+1})M_-(r + 1, r)$ .

The matrices  $M_+(r, r)$  and  $M_-(r + 1, r)$  are given by

$$\begin{bmatrix} M_+(r, r) \\ M_-(r+1, r) \end{bmatrix} = \begin{bmatrix} L(w_r) \\ \tilde{L}(w_r) \end{bmatrix} + \begin{bmatrix} \Gamma(w_r)\Gamma^*(v_r) & \Lambda(w_r)A(v_{r+1}) \\ \Lambda^*(w_r)\Gamma^*(v_r) & \Gamma^*(w_r)A(v_{r+1}) \end{bmatrix} \begin{bmatrix} M_+(r, r) \\ M_-(r+1, r) \end{bmatrix}$$

where  $w_r = v_{r+1} - v_r$  and

- $L_{ij}(x)$  is the average time spent in phase  $j \in \mathcal{S}$  before reaching level zero or  $x$ , starting from  $(x, i \in \mathcal{S}_+)$ ,
- $\tilde{L}_{ij}(x)$  is the average time spent in phase  $j \in \mathcal{S}$  before reaching level zero or  $x$ , starting from  $(x, i \in \mathcal{S}_-)$ .

## Proposition ( $r = N - 1$ )

$$M_-(N-1, N-1) = \Gamma^*(v_{N-1})M_+(N-1, N-1),$$

$$M_+(N, N-1) = [(-Q_{++})^{-1} \ 0] + (-Q_{++})^{-1}Q_{+-}M_-(N, N-1).$$

The matrices  $M_+(N-1, N-1)$  and  $M_-(N, N-1)$  are given by

$$\begin{bmatrix} M_+(N-1, N-1) \\ M_-(N, N-1) \end{bmatrix} = \begin{bmatrix} L(w_{N-1}) \\ \tilde{L}(w_{N-1}) \end{bmatrix} + \begin{bmatrix} \Lambda(w_{N-1}) \\ \Gamma^*(w_{N-1}) \end{bmatrix} \begin{bmatrix} (-Q_{++})^{-1} & 0 \end{bmatrix}$$

$$+ \begin{bmatrix} \Gamma(w_{N-1})\Gamma^*(v_{N-1}) & \Lambda(w_{N-1})(-Q_{++})^{-1}Q_{+-} \\ \Lambda^*(w_{N-1})\Gamma^*(v_{N-1}) & \Gamma^*(w_{N-1})(-Q_{++})^{-1}Q_{+-} \end{bmatrix} \begin{bmatrix} M_+(N-1, N-1) \\ M_-(N, N-1) \end{bmatrix}.$$

## Example

The process  $(Y, \varphi)$  is an MMFF on the level space  $[0, 1]$  and ruled by the matrices

$$Q = \begin{bmatrix} -1 & 1 \\ 1 & -1 \end{bmatrix}, \quad C = \begin{bmatrix} 2 & \\ & -1 \end{bmatrix}.$$

The infectious periods start at level zero in the ascending phase, and

$$\beta_{l,i} = \begin{cases} \frac{\gamma}{n} \frac{l}{N} & \text{if } x \in [v_l, v_{l+1}[ \text{ and } i \in \mathcal{S}_+, \\ 0 & \text{else,} \end{cases}$$

$$a(x, i) = \begin{cases} \frac{\alpha l}{N} & \text{if } x \in [v_l, v_{l+1}[ \text{ and } i \in \mathcal{S}_-, \\ 0 & \text{else,} \end{cases}$$

# Example

$N$	10	20	50	100	200	500	1000	2000
$\mathbb{E}[S(T)]$	7.70	6.84	6.37	6.23	6.16	6.12	6.10	6.10
$\mathbb{V}ar[S(T)]$	128	121	116	114	114	113	113	113
$\mathbb{E}[D]$	0.64	0.67	0.69	0.70	0.70	0.70	0.70	0.70
$\mathbb{E}[\mathcal{A}(T)]$	20.6	22.2	23.2	23.5	23.7	23.8	23.8	23.8
$R_0$	2.70	2.85	2.94	2.97	2.99	2.99	3.00	3.00

TABLE –  $n = 37$ ,  $m = 3$ ,  $\gamma = 1.5$ ,  $\alpha = 1$ . Here,  $\mathbb{E}[\tau_\star] = 3.6$ .

# End

Thank you for your attention.