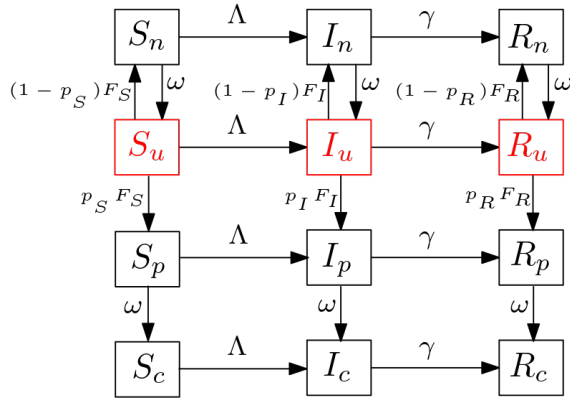


# Notes on a Simple Epidemic Model

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## 1 Method

### 1.1 model and parameters



The model is

$$dS_u/dt = -\Lambda S_u - F_S S_u + \omega S_n, \quad (1)$$

$$dS_n/dt = -\Lambda S_n + (1 - p_S) F_S S_u - \omega S_n, \quad (2)$$

$$dS_p/dt = -\Lambda S_p + p_S F_S S_u - \omega S_p, \quad (3)$$

$$dS_c/dt = -\Lambda S_c + \omega S_p, \quad (4)$$

$$dI_u/dt = \Lambda S_u - F_I I_u + \omega I_n - \gamma I_u, \quad (5)$$

$$dI_n/dt = \Lambda S_n + (1 - p_I) F_I I_u - \omega I_n - \gamma I_n, \quad (6)$$

$$dI_p/dt = \Lambda S_p + p_I F_I I_u - \omega I_p - \gamma I_p, \quad (7)$$

$$dI_c/dt = \Lambda S_c + \omega I_p - \gamma I_c, \quad (8)$$

$$dR_u/dt = \gamma I_u - F_R R_u + \omega R_n, \quad (9)$$

$$dR_n/dt = \gamma I_n + (1 - p_R) F_R R_u - \omega R_n, \quad (10)$$

$$dR_p/dt = \gamma I_p + p_R F_R R_u - \omega R_p, \quad (11)$$

$$dR_c/dt = \gamma I_c + \omega R_p, \quad (12)$$

$$dN/dt = \omega(S_n + I_n + R_n), \quad (13)$$

$$dP/dt = \omega(I_p + R_p), \quad (14)$$

Symbol	Description	Unit	Value
$N_0$	Total population size	people	$10^6$
$\omega$	Rate of onward flow from the awaiting to reported or untested compartments	1/day	-
$\gamma$	Recovery rate	1/day	1/3
$\rho$	Per capita testing intensity	1/day	0.01
$\eta_w$	Relative probability of transmission for isolated awaiting individuals	-	-
$\eta_c$	Relative probability of transmission for isolated confirmed individuals	-	-
$\Lambda$	Force of infection	1/day	-
$p_S$	Probability of false positive for susceptible	-	0
$p_I$	Probability of being infected and tested positive	-	1
$p_R$	Probability of being recovered and tested positive	-	0.5
$W_S, W_I, W_R$	Relative testing weight	-	Random testing: $W_S = W_I = W_R = 1$ , Non-random testing: $W_S = 0.3, W_I = W_R = 1$

Table 1: The underlying parameters of model, Eqs. (1) to (12).

## 2 Results

### 2.1 On the calculation of $\mathcal{R}_0$

- DFE is given by solving the following system <sup>1</sup>

$$\begin{aligned} S_u + S_n &= N_0 \\ F_S S_u - \omega S_n &= 0 \end{aligned}$$

- DFE:

$$\begin{aligned} S_u &= (1 - \frac{\rho}{\omega})N_0, \\ S_n &= \frac{\rho}{\omega}N_0, \\ I_j = R_j &= 0 \quad \text{for all } j. \end{aligned}$$

- At the DFE,  $F_I^* = \frac{\rho}{(1-\rho/\omega)}W_I/W_S$  also note that  $\partial F_I^*/\partial \rho > 0$ .

$$F = \beta/N_0 \begin{bmatrix} S_u & \eta_w S_u & \eta_w S_u & \eta_c S_u \\ S_n & \eta_w S_n & \eta_w S_n & \eta_c S_n \\ 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 \end{bmatrix}, \quad (15)$$

$$V = \begin{bmatrix} F_I + \gamma & -\omega & 0 & 0 \\ -(1-p_I)F_I & \omega + \gamma & 0 & 0 \\ -p_I F_I & 0 & \omega + \gamma & 0 \\ 0 & 0 & -\omega & \gamma \end{bmatrix}, \text{ thus} \quad (16)$$

$$V^{-1} = \begin{bmatrix} \frac{\omega + \gamma}{\omega \gamma + \gamma F_I + \gamma^2 + \omega F_I p_I} & \frac{\omega}{\omega \gamma + \gamma F_I + \gamma^2 + \omega F_I p_I} & 0 & 0 \\ \frac{(1-p_I)F_I}{\omega \gamma + \gamma F_I + \gamma^2 + \omega F_I p_I} & \frac{F_I + \gamma}{\omega \gamma + \gamma F_I + \gamma^2 + \omega F_I p_I} & 0 & 0 \\ \frac{p_I F_I}{\omega \gamma + \gamma F_I + \gamma^2 + \omega F_I p_I} & \frac{\omega p_I F_I}{(\omega \gamma + \gamma F_I + \gamma^2 + \omega F_I p_I)(\omega + \gamma)} & \frac{1}{\omega + \gamma} & 0 \\ \frac{\omega F_I p_I}{(\omega \gamma + \gamma F_I + \gamma^2 + \omega F_I p_I)\gamma} & \frac{\omega^2 F_I p_I}{(\omega \gamma + \gamma F_I + \gamma^2 + \omega F_I p_I)(\omega + \gamma)\gamma} & \frac{\omega}{(\omega + \gamma)\gamma} & \frac{1}{\gamma} \end{bmatrix}. \quad (17)$$

The particular form of  $F$  with two rows of zeros at the bottom, simplifies  $G$  as

$$G = \begin{bmatrix} G_{11} & G_{12} \\ 0 & 0 \end{bmatrix}, \text{ where } G_{11} = C \begin{bmatrix} A S_u & B S_u \\ A S_n & B S_n \end{bmatrix}. \quad (18)$$

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<sup>1</sup>Note when  $I_u = R_u = 0$ ,  $F_S = \frac{\rho N_0}{W_S S_u}$ .

The Disease-Free Equilibrium (DFE) for the SIR model, Eqs. (1) to (12), is given by solving the coupled system including  $S_u + S_n = N_0$  and  $F_S S_u - \omega S_n = 0$ . The DFE is

$$S_n^* = \frac{\rho}{\omega} N_0, \quad S_u^* = N_0 - S_n^*, \text{ and } I_j = R_j = 0 \text{ for all } j. \quad (19)$$

The basic reproduction number,  $\mathcal{R}_0$ , was calculated by using the next generation matrix method developed by Van den Driessche and Watmough (2002).  $\mathcal{R}_0$  is

$$\mathcal{R}_0 = (A \times S_u^* + B \times S_n^*) \times C, \quad (20)$$

where

$$\begin{aligned} A &= \gamma(\omega + \gamma) + (\gamma\eta_w + \omega\eta_c p_I) F_I, \\ B &= (\omega + (F_I + \gamma)\eta_w) \gamma + \frac{(\eta_w \gamma + \eta_c \omega) \omega p_I F_I}{\omega + \gamma}, \\ C &= \frac{\beta/\gamma}{N_0(\gamma(\omega + \gamma) + F_I(\gamma + \omega p_I))}. \end{aligned}$$

Note that the block matrix  $G_{12}$  does not influence  $\mathcal{R}_0$  defined as the spectral radius of  $G$ . All matters here are the eigenvalues of  $G_{11}$ , which are 0 and  $\mathcal{R}_0$  (20).

## 2.2 Sensitivity of $\mathcal{R}_0$ with respect to the underlying parameters

**note 1.** The tricky one is the  $\partial \mathcal{R}_0 / \partial \rho$ ; Following JD comment and some analysis, we can prove that analytically.

**Jonathan's comment:** "There should be a logical route to a clear proof about  $\rho$ . The total amount of time spent in  $I_x$  when starting from a given starting point should not depend on  $\rho$  (or anything but  $\gamma$ ). And this should be reflected in the column sums of  $V^{-1}$ . Since both of the  $\eta$ 's are  $< 1$ , all we should then need to prove is that the (nonzero) values on the first row of  $V^{-1}$  decrease with  $\rho$  to show that the elements of  $FV^{-1}$  decrease with  $\rho$ , which presumably shows that  $\mathcal{R}_0$  decreases as well. This needs to be filled ink but should work."

**note 2.** Notice that matrix  $F$  is a rank one matrix, thus  $FV^{-1}$  is of a rank one. This is why  $FV^{-1}$  has only one non-zero eigenvalue. Specifically,  $F$  is a rank one

means that it can be written as matrix product of a column vector and a row vector as follows

$$F = \beta/N_0 \begin{bmatrix} S_u \\ S_n \\ 0 \\ 0 \end{bmatrix} [ 1\eta_w\eta_w\eta_c ] .$$

**Proposition 1.** Let  $f$  and  $g$  be smooth real functions such that (i)  $f(x) \leq g(x)$  for all  $x \in (x_0, x_0 + \epsilon)$  and (ii)  $f(x_0) = g(x_0)$ , then the definition of the derivative yields

$$f'(x_0) = \lim_{h \rightarrow 0} \frac{f(x_0 + h) - f(x_0)}{h} = \lim_{h \rightarrow 0} \frac{f(x_0 + h) - g(x_0)}{h} \leq \lim_{h \rightarrow 0} \frac{g(x_0 + h) - g(x_0)}{h} = g'(x_0).$$

Based on the note 2, Also notice that  $FV^{-1}$  is of rank one matrix since it can be written as matrix product of a column vector and a row vector. Thus, the next generation matrix  $G = FV^{-1}$  has only one non-zero eigenvalue given by the trace  $G$ <sup>2</sup>.

Thus,

$$\text{trace}(G) = \text{trace}(G_{11}) \leq \frac{\beta}{\gamma} \frac{1}{N_0} (S_u^* + S_n^*) \quad \forall \rho \in (0, \epsilon). \quad (21)$$

It is easy to show that  $\text{trace}(G) = \frac{\beta}{\gamma} \frac{1}{N_0} (S_u^* + S_n^*) = 1/\gamma$ . Thus, following the Prop.1,  $\partial \mathcal{R}_0 / \partial \rho \leq 0$ .

## References

Van den Driessche, P. and Watmough, J. (2002). Reproduction numbers and sub-threshold endemic equilibria for compartmental models of disease transmission. *Mathematical biosciences*, 180(1-2):29–48.

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<sup>2</sup>see the argument in <https://www.physicsforums.com/threads/eigenvalues-of-a-rank-1-matrix.682216/>