

Notes on network and generation

Suppose we have one infected individual and one susceptible individual. Infected individual makes effective contact at a constant rate β and has a constant recovery rate of λ . Then, we can write the probability that the susceptible remains uninfected at time t as follows:

$$s(\tau) = \exp(-\beta \int_{t=0}^{\tau} \exp(-\lambda t) dt)$$

Solving the integral, we have

$$s(\tau) = \exp\left(-\frac{\beta}{\lambda}(1 - \exp(-\lambda\tau))\right)$$

Define $K(\tau) = F(\tau)\beta$ as the rate of secondary case, where $F(\tau) = \exp(-\lambda\tau)$ is the probability that an infectious individual has not recovered t time units after infection.

We define network generation interval as follows:

$$g(\tau) = \frac{K(\tau)s(\tau)}{\int_0^{\infty} K(\tau)s(\tau)d\tau}$$

which can be simplified into

$$g(\tau) = \frac{F(\tau)s(\tau)}{\int_0^{\infty} F(\tau)s(\tau)d\tau}$$

Let's try to solve the integral... First, let's do substitution:

$$\begin{aligned} dF &= -\lambda \exp(-\lambda\tau) d\tau \\ &= -\lambda F d\tau \end{aligned}$$

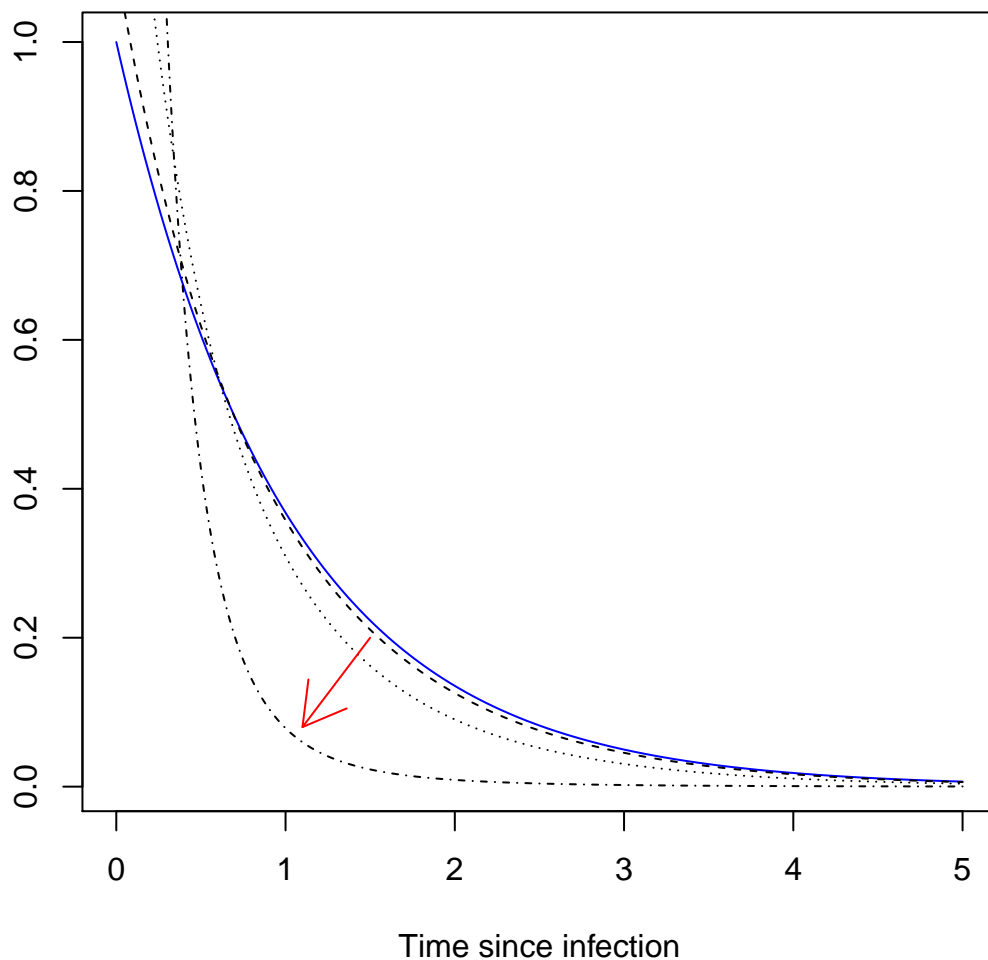
Substitute it:

$$\begin{aligned} &\int_0^{\infty} F(\tau)s(\tau)d\tau \\ &= -\frac{1}{\lambda} \int_1^0 \exp\left(-\frac{\beta}{\lambda}(1 - F)\right) dF \\ &= \frac{1}{\beta} \left[\exp\left(-\frac{\beta}{\lambda}(1 - F)\right) \right]_0^1 \\ &= \frac{1}{\beta} \left(1 - \exp\left(-\frac{\beta}{\lambda}\right) \right) \end{aligned}$$

Therefore, the generation interval is

$$g(\tau) = \beta \left(1 - \exp\left(-\frac{\beta}{\lambda}\right) \right)^{-1} \exp(-\lambda\tau) \exp\left(-\frac{\beta}{\lambda}(1 - \exp(-\lambda\tau))\right)$$

Comparison of intrinsic generation with network generation distribution for SIR model:



As we increase effective transmission rate, network generation follows the red arrow. Let's try to write a general equation.

In a homogeneous population, generation interval is defined as follows (Svensson 2015):

$$g(\tau) = \frac{P(X + Y > \tau) - P(X > \tau)}{E(Y)}$$

where X is the duration of the latent period and Y is the duration of the infection period. To check that it works, let's try *SEIR* model where both latent and infectious periods are exponentially distributed with rate γ and λ :

$$\begin{aligned}
P(X + Y > \tau) &= 1 - \int_0^\tau \int_0^s f_Y(s-t) f_X(t) dt ds \\
&= 1 - \int_0^\tau \int_0^s \lambda \exp(-\lambda(s-t)) \gamma \exp(-\gamma t) dt ds \\
&= 1 - \int_0^\tau \gamma \lambda \exp(-\lambda s) [\exp((\lambda - \gamma)t) / (\lambda - \gamma)]_0^s ds \\
&= 1 - \int_0^\tau \frac{\gamma \lambda}{\lambda - \gamma} \exp(-\lambda s) (\exp((\lambda - \gamma)s) - 1) ds \\
&= 1 - \frac{\gamma \lambda}{\lambda - \gamma} \int_0^\tau (\exp(-\gamma s) - \exp(-\lambda s)) ds \\
&= 1 - \frac{\gamma \lambda}{\lambda - \gamma} \left[\exp(-\lambda s) / \lambda - \exp(-\gamma s) / \gamma \right]_0^\tau \\
&= 1 - \frac{1}{\lambda - \gamma} \left[\gamma \exp(-\lambda s) - \lambda \exp(-\gamma s) \right]_0^\tau \\
&= 1 - \frac{1}{\lambda - \gamma} \left[\gamma \exp(-\lambda \tau) - \lambda \exp(-\gamma \tau) - \gamma + \lambda \right] \\
&= \frac{\gamma \exp(-\lambda \tau) - \lambda \exp(-\gamma \tau)}{\gamma - \lambda}
\end{aligned}$$

And we also have

$$\begin{aligned}
P(X > \tau) &= 1 - \int_0^\tau \gamma \exp(-\gamma t) dt \\
&= 1 + (\exp(-\gamma t))_0^\tau \\
&= \exp(-\gamma \tau)
\end{aligned}$$

Then,

$$\begin{aligned}
\frac{P(X + Y > \tau) - P(X > \tau)}{E(Y)} &= \lambda \left(\frac{\gamma \exp(-\lambda \tau) - \lambda \exp(-\gamma \tau)}{\gamma - \lambda} - \exp(-\gamma \tau) \right) \\
&= \lambda \gamma \left(\frac{\exp(-\lambda \tau) - \exp(-\gamma \tau)}{\gamma - \lambda} \right)
\end{aligned}$$

Got the same result as Svensson, which is a good sign. Now, how do we apply this to a network.. Going back to Trapman et al, we have the following survival function:

$$\begin{aligned}
s(\tau) &= \exp \left(-\beta \left(\int_0^\tau F(t) dt \right) \right) \\
&= \exp \left(-\beta \left(\int_0^\tau P(X + Y > t) - P(X > t) dt \right) \right)
\end{aligned}$$