

# Multi-type branching process model

Mary Bushman

July 21, 2018

Define a multitype branching process with  $X(n) = (X_w(n), X_m(n), X_c(n))$  where  $X_w(n)$ ,  $X_m(n)$ , and  $X_c(n)$  are the number of wildtype infections, mutant infections, and coinfections (wildtype + mutant) at time step  $n$ .

Let  $R_w$  and  $R_m$  represent the  $R_0$  values for wildtype and mutant infections, respectively. Let  $R_c$  be the  $R_0$  value for a coinfection (wildtype + mutant), with  $R_c = dR_w + (1-d)R_m$  where  $d$  is the “dominance” of the mutant genotype.

Let  $c$  be the probability that both wildtype and mutant are transmitted from a coinfection. In the event that only one strain is transmitted from a coinfection, let  $b$  be the probability that it is the mutant.

Let  $Y_{ij}$  be the number of progeny of type  $j$  from an individual of type  $i$ . Then let

$$\begin{aligned} Y_{ww} &\sim \text{Poisson}(R_w) \\ Y_{mm} &\sim \text{Poisson}(R_m) \\ Y_{cw} &\sim \text{Poisson}((1-b)(1-c)R_c) \\ Y_{cm} &\sim \text{Poisson}(b(1-c)R_c) \\ Y_{cc} &\sim \text{Poisson}(cR_c) \end{aligned}$$

All other  $Y_{ij}$  are assumed to be zero.

Define  $R_{ij} = E[Y_{ij}]$ .

Let  $\mu_1$  be the mutation probability (probability that a wildtype infection becomes a coinfection) and let  $\mu_2$  be the reversion probability (probability that a mutant infection becomes a coinfection).

Let  $p_w(k_w, k_m, k_c)$  be the probability that a wildtype infection gives rise to  $k_w$  wildtype infections,  $k_m$  mutant infections and  $k_c$  coinfections. To find this probability it is necessary to condition on whether a mutation occurs in the wildtype infection, turning it into a coinfection. Thus,

$$p_w(k_w, k_m, k_c) = (1 - \mu_1)Pr(k_w, k_m, k_c | \text{no mutation}) + \mu_1 Pr(k_w, k_m, k_c | \text{mutation})$$

.

For shorthand, define  $z(x) = \begin{cases} 1 & \text{if } x = 0 \\ 0 & \text{otherwise} \end{cases}$

Thus, we have

$$p_w(k_w, k_m, k_c) = z(k_m)z(k_c)(1 - \mu_1) \frac{R_{ww}^{k_w} e^{-R_{ww}}}{k_w!} + \mu_1 \frac{R_{cw}^{k_w} e^{-R_{cw}}}{k_w!} \frac{R_{cm}^{k_m} e^{-R_{cm}}}{k_m!} \frac{R_{cc}^{k_c} e^{-R_{cc}}}{k_c!}$$

Similarly,

$$p_m(k_w, k_m, k_c) = z(k_w)z(k_c)(1 - \mu_2) \frac{R_{mm}^{k_m} e^{-R_{mm}}}{k_m!} + \mu_2 \frac{R_{cw}^{k_w} e^{-R_{cw}}}{k_w!} \frac{R_{cm}^{k_m} e^{-R_{cm}}}{k_m!} \frac{R_{cc}^{k_c} e^{-R_{cc}}}{k_c!}$$

$$p_c(k_w, k_m, k_c) = \frac{R_{cw}^{k_w} e^{-R_{cw}}}{k_w!} \frac{R_{cm}^{k_m} e^{-R_{cm}}}{k_m!} \frac{R_{cc}^{k_c} e^{-R_{cc}}}{k_c!}$$

We then use these probability distributions to define the probability generating functions:

$$f_w(t_w, t_m, t_c) = \sum_{k_w=0}^{\infty} \sum_{k_m=0}^{\infty} \sum_{k_c=0}^{\infty} p_w(k_w, k_m, k_c) t_w^{k_w} t_m^{k_m} t_c^{k_c}$$

$$f_m(t_w, t_m, t_c) = \sum_{k_w=0}^{\infty} \sum_{k_m=0}^{\infty} \sum_{k_c=0}^{\infty} p_m(k_w, k_m, k_c) t_w^{k_w} t_m^{k_m} t_c^{k_c}$$

$$f_c(t_w, t_m, t_c) = \sum_{k_w=0}^{\infty} \sum_{k_m=0}^{\infty} \sum_{k_c=0}^{\infty} p_c(k_w, k_m, k_c) t_w^{k_w} t_m^{k_m} t_c^{k_c}$$

Each generating function maps  $[0, 1]^3 \rightarrow [0, 1]$ .

Then denote  $g(t_w, t_m, t_c) = (f_w(t_w, t_m, t_c), f_m(t_w, t_m, t_c), f_c(t_w, t_m, t_c))$ .

Note that  $g(t_w, t_m, t_c)$  maps  $[0, 1]^3 \rightarrow [0, 1]^3$ .

If we select some values  $t_w, t_m, t_c \in [0, 1]$  and call these  $t_w(0), t_m(0), t_c(0)$ , and define  $(t_w(n), t_m(n), t_c(n)) = g(t_w(n-1), t_m(n-1), t_c(n-1))$  then

$$\lim_{n \rightarrow \infty} (t_w(n), t_m(n), t_c(n)) = (q_w, q_m, q_c)$$

is a fixed point for  $g(t_w, t_m, t_c)$  on  $[0, 1]^3$ . Furthermore,  $q_i$  is the probability of eventual extinction for a multitype branching process beginning with a single infection of type  $i$ . For a multitype branching process beginning with  $(x_w, x_m, x_c)$  wildtype, mutant and coinfections, the ultimate extinction probability is

$$q_w^{x_w} q_m^{x_m} q_c^{x_c}$$

Approximate calculations of  $f_w(t_w, t_m, t_c)$ ,  $f_m(t_w, t_m, t_c)$ , and  $f_c(t_w, t_m, t_c)$  are easy to implement in R, as is the algorithm to converge to the fixed point of the function  $g(t_w, t_c, t_m)$ . Thus, for any set of parameters  $(R_w, R_m, d, c, b, \mu_1, \mu_2)$  we can easily calculate the ultimate extinction probability for starting conditions  $(x_w, x_m, x_c)$ . (Generally we will assume  $x_w = 1, x_m = x_c = 0$ , but the calculations are equally straightforward for other starting conditions.)