## Malaria Transmission Dynamics with Mosquito Treatment: Reproduction Number Analysis

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April 8, 2025

# 1 Closed-Form Expression for the Basic Reproduction Number, $R_0$

When the system is linearized about the disease-free equilibrium (DFE), we set

$$I_H = E_{1M} = E_{2M} = I_M = E_{1T} = E_{2T} = I_T = 0,$$

and the human population is entirely susceptible, i.e.  $S_H=1$ . For the mosquito population, we assume that the total population is normalized:

$$S_M + S_T = 1.$$

In the mosquito subsystem the DFE is obtained by solving the linear system:

$$\begin{pmatrix} -(t+g) & h \\ t & -(h+g) \end{pmatrix} \begin{pmatrix} S_M^* \\ S_T^* \end{pmatrix} = \begin{pmatrix} -g \\ 0 \end{pmatrix}.$$

A short calculation yields

$$S_M^* = \frac{h+g}{t+h+g}, \quad S_T^* = \frac{t}{t+h+g}.$$

Next, following the standard next-generation approach the infection cycle is "completed" by tracking the passage from an infectious human to an infected mosquito and back to a newly infected human.

#### **Untreated Branch**

An infectious human infects a mosquito (via a bite) at rate ac. In the untreated branch, the infection passes sequentially through two latent stages:

$$\begin{split} E_{1M} &= \frac{ac\,S_M^*}{t + s_{1M} + g}\,I_H, \\ E_{2M} &= \frac{s_{1M}}{s_{2M} + g}\,E_{1M}, \\ I_M &= \frac{s_{2M}}{g}\,E_{2M}. \end{split}$$

Thus the passage (or survival) probability for the untreated branch is

$$\pi_M = \frac{ac \, S_M^*}{t + s_{1M} + g} \cdot \frac{s_{1M}}{s_{2M} + g} \cdot \frac{s_{2M}}{g}.$$

#### **Treated Branch**

Similarly, the treated branch follows:

$$\begin{split} E_{1T} &= \frac{ac\,S_T^*}{s_{1T} + g}\,I_H, \\ E_{2T} &= \frac{s_{1T}}{s_{2T} + g}\,E_{1T}, \\ I_T &= \frac{s_{2T}}{g}\,E_{2T}, \end{split}$$

with passage probability

$$\pi_T = \frac{ac \, S_T^*}{s_{1T} + g} \cdot \frac{s_{1T}}{s_{2T} + g} \cdot \frac{s_{2T}}{g}.$$

### Additional Contribution: Untreated Mosquitoes Receiving Treatment During Latency

Experimental evidence (see slide 10) indicates that some mosquitoes in the untreated branch are treated during the latent period. Their contribution is given by:

$$\pi_{extra} = \frac{ac \, S_M^*}{t + s_{1M} + g} \cdot \frac{t}{t + s_{1M} + g} \cdot \frac{s_{1T}}{s_{1T} + g} \cdot \frac{s_{2T}}{s_{2T} + g}.$$

#### Overall Expression for $R_0$

An infectious human transmits to mosquitoes (at rate ac) and, in turn, an infectious mosquito transmits to humans at rate ab. With the mosquito-to-human ratio m and the human recovery rate r, the total number of secondary human infections produced by one infected human is

$$R_0 = \frac{m \, a \, b}{r} \Big[ \pi_M + \pi_T + \pi_{extra} \Big].$$

Substituting the expressions for  $\pi_M$ ,  $\pi_T$ , and  $\pi_{extra}$  and noting that ac appears in each term, we obtain

$$R_0 = \frac{m a^2 b c}{r g} \left[ \frac{S_M^* s_{1M} s_{2M}}{(t + s_{1M} + g)(s_{2M} + g)} + \frac{S_T^* s_{1T} s_{2T}}{(s_{1T} + g)(s_{2T} + g)} + \frac{S_M^* t s_{1T} s_{2T}}{(t + s_{1M} + g)(s_{1T} + g)(s_{2T} + g)} \right].$$

Finally, substituting

$$S_M^* = \frac{h+g}{t+h+g}, \quad S_T^* = \frac{t}{t+h+g},$$

the closed-form symbolic expression for  $R_0$  becomes

$$R_{0} = \frac{m a^{2} b c}{r g} \left[ \frac{(h+g) s_{1M} s_{2M}}{(t+h+g)(t+s_{1M}+g)(s_{2M}+g)} + \frac{t s_{1T} s_{2T}}{(t+h+g)(s_{1T}+g)(s_{2T}+g)} + \frac{(h+g) t s_{1T} s_{2T}}{(t+h+g)(t+s_{1M}+g)(s_{1T}+g)(s_{2T}+g)} \right].$$
(1)

This is the desired closed-form symbolic expression for  $R_0$  in terms of the model parameters, now correctly incorporating the extra contribution.

## 2 Closed-Form Endemic Equilibrium (EE)

At endemic equilibrium all time derivatives vanish. In addition to the normalizations

$$S_H + I_H = 1$$
,  $S_M + E_{1M} + E_{2M} + I_M + S_T + E_{1T} + E_{2T} + I_T = 1$ ,

we write the equilibrium conditions for each compartment.

#### 2.1 Human Compartments

The human dynamics are given by

$$0 = \frac{dS_H}{dt} = -m a b (I_M + I_T) S_H + r I_H,$$
  
$$0 = \frac{dI_H}{dt} = m a b (I_M + I_T) S_H - r I_H.$$

Since  $S_H = 1 - I_H$ , one immediately finds

$$\frac{I_H}{1 - I_H} = \frac{m a b \left(I_M + I_T\right)}{r},$$

so that

$$I_H = \frac{m \, a \, b \, (I_M + I_T)}{r + m \, a \, b \, (I_M + I_T)}, \quad S_H = 1 - I_H.$$

#### 2.2 Mosquito Compartments

For the untreated branch, the equilibrium equations are:

$$0 = \frac{dS_M}{dt} = g + h S_T - ac I_H S_M - t S_M - g S_M,$$

$$0 = \frac{dE_{1M}}{dt} = ac I_H S_M - (t + s_{1M} + g) E_{1M},$$

$$0 = \frac{dE_{2M}}{dt} = s_{1M} E_{1M} - (s_{2M} + g) E_{2M},$$

$$0 = \frac{dI_M}{dt} = s_{2M} E_{2M} - g I_M.$$

Thus, we obtain

$$E_{1M} = \frac{ac I_H S_M}{t + s_{1M} + g}, \quad E_{2M} = \frac{s_{1M} E_{1M}}{s_{2M} + g}, \quad I_M = \frac{s_{2M} E_{2M}}{g}.$$

For the treated branch, the equilibrium equations are:

$$0 = \frac{dS_T}{dt} = t S_M - ac I_H S_T - h S_T - g S_T,$$

$$0 = \frac{dE_{1T}}{dt} = ac I_H S_T + t E_{1M} - (s_{1T} + g) E_{1T},$$

$$0 = \frac{dE_{2T}}{dt} = s_{1T} E_{1T} - (s_{2T} + g) E_{2T},$$

$$0 = \frac{dI_T}{dt} = s_{2T} E_{2T} - g I_T.$$

From the first equation we solve for  $S_T$  in terms of  $S_M$ :

$$t S_M = (ac I_H + h + g) S_T \implies S_T = \frac{t S_M}{ac I_H + h + g}.$$

Similarly, we obtain

$$E_{1T} = \frac{ac I_H S_T + t E_{1M}}{s_{1T} + g}, \quad E_{2T} = \frac{s_{1T} E_{1T}}{s_{2T} + g}, \quad I_T = \frac{s_{2T} E_{2T}}{g}.$$

Finally, the equation for  $S_M$  is found from the  $S_M$  balance:

$$ac I_H S_M = g + h S_T - (t+g) S_M,$$

or equivalently,

$$S_M = \frac{g + h S_T}{ac I_H + t + g}.$$

Substituting

$$S_T = \frac{t \, S_M}{ac \, I_H + h + a},$$

yields a closed-form solution for  $S_M$ :

$$S_M \left( ac \, I_H + t + g \right) = g + \frac{ht \, S_M}{ac \, I_H + h + g}.$$

Solving for  $S_M$ , we obtain

$$S_{M} = \frac{g(ac I_{H} + h + g)}{(ac I_{H} + t + g)(ac I_{H} + h + g) - ht}.$$

Then,

$$S_T = \frac{t}{ac I_H + h + g} S_M.$$

#### 2.3 Summary of the Endemic Equilibrium

Collecting the results, the closed-form endemic equilibrium is given by:

$$\begin{split} S_{H} &= \frac{r}{r + m \, a \, b \, (I_{M} + I_{T})}, \\ I_{H} &= \frac{m \, a \, b \, (I_{M} + I_{T})}{r + m \, a \, b \, (I_{M} + I_{T})}, \\ S_{M} &= \frac{g (ac \, I_{H} + h + g)}{(ac \, I_{H} + t + g) (ac \, I_{H} + h + g) - ht}, \\ S_{T} &= \frac{t}{ac \, I_{H} + h + g} \, S_{M}, \\ E_{1M} &= \frac{ac \, I_{H} \, S_{M}}{t + s_{1M} + g}, \quad E_{2M} = \frac{s_{1M} \, E_{1M}}{s_{2M} + g}, \quad I_{M} = \frac{s_{2M} \, E_{2M}}{g}, \\ E_{1T} &= \frac{ac \, I_{H} \, S_{T} + t \, E_{1M}}{s_{1T} + g}, \quad E_{2T} = \frac{s_{1T} \, E_{1T}}{s_{2T} + g}, \quad I_{T} = \frac{s_{2T} \, E_{2T}}{g}. \end{split}$$

The consistency condition in the human equation,

$$\frac{I_H}{1 - I_H} = \frac{m a b \left(I_M + I_T\right)}{r},$$

must be satisfied. In principle the term  $I_M + I_T$  (denoted by  $\Lambda$ ) can be expressed explicitly in terms of  $I_H$  and parameters using the above relations, leading to a nonlinear (typically quadratic) equation in  $I_H$  that can be solved in closed form. Once  $I_H$  is determined, all other compartments are explicitly given by the formulas above.

## 3 Results

The following results are obtained for the symbolic expressions of the basic reproduction number,  $R_0$ , and the endemic equilibrium (EE) of the model. For comparison, results from the matrix method are also provided.

#### 3.1 Model Dynamics

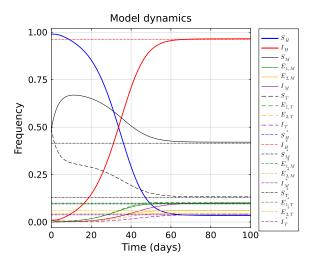


Figure 1: Convergence to endemic equilibrium with default parameters. The EE is obtained from the symbolic expressions.

#### 3.2 Basic Reproduction Number

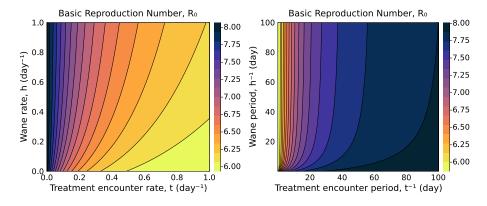


Figure 2:  $R_0$  vs treatment rates (t, h) and treatment periods (1/t, 1/h). Results are obtained from the matrix method.

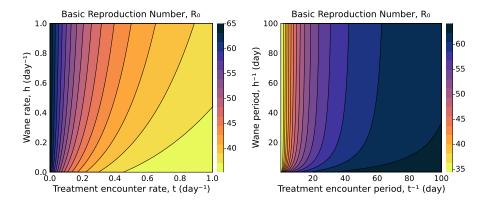


Figure 3:  $R_0$  vs treatment rates (t,h) and treatment periods (1/t,1/h). Results are obtained from the symbolic expressions.

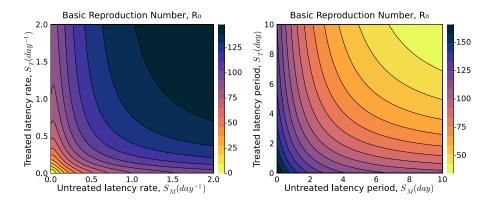


Figure 4:  $R_0$  vs latency rates  $(s_{1M}, s_{2M})$  and treatment rates  $(s_{1T}, s_{2T})$ . Results are obtained from the matrix method.

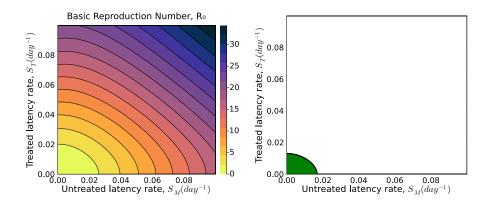


Figure 5:  $R_0$  vs latency rates  $(s_{1M}, s_{2M})$  and  $R_0 > 1$  cut. Results are obtained from the symbolic expressions.

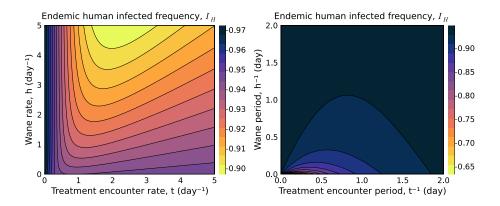


Figure 6: Endemic equilibrium of infections humans  $I_H^*$  vs treatment rates (t,h) and treatment periods (1/t,1/h). Results are obtained from the matrix method.

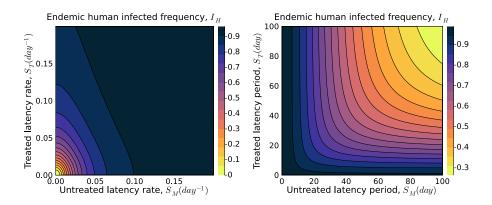


Figure 7: Endemic equilibrium of infections humans  $I_H^*$  vs latency rates  $(s_{1M},s_{2M})$  and treatment rates  $(s_{1T},s_{2T})$ . Results are obtained from the matrix method.